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Transcriptional Profiling of the Sperm Storage Organs of *Drosophila melanogaster*

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Transcriptional profiling of the sperm storage organs of *Drosophila melanogaster*

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Abstract
The occurrence of female sperm storage across taxa indicates the importance of this complex and dynamic process. Organs responsible for sperm storage (SSOs) and proteins expressed therein, are important in fundamental aspects of reproduction and could play a major role in evolutionary processes such as post-mating sexual selection. Given the essential role of SSOs, it is surprising that the process of sperm storage is so poorly understood. This study investigated the transcriptome of female *Drosophila melanogaster* SSOs (seminal receptacle and spermathecae). Spermathecae were enriched for proteases and metabolic enzymes while the seminal receptacle was enriched for genes involved in localization, signaling and ion transport. Differences in functional gene categories indicate that these organs play unique roles in sperm storage.

Keywords: sperm storage, microarray, spermathecae, seminal receptacle

Introduction

Female sperm storage is a reproductive strategy utilized, to some degree, in most internally fertilizing animals. Sperm storage allows for the female to acquire sperm before it is needed for fertilization. The duration of sperm storage is highly variable, ranging from a few hours (e.g. mouse), to decades (e.g. honey bee). The reproductive benefits of this process are significant as contact with a suitable male does not necessary correspond with egg availability. For females that multiply mate, sperm storage can offer an additional selection step via sperm competition and/or cryptic female sperm choice, as well as give the female a mechanism to compensate for infertile/genetically incompatible mates.

Organs dedicated to the process of sperm storage (SSOs) have evolved in diverse phyla. In insects, SSOs typically appear as sac-like structures (spermathecae) or long tubules (seminal receptacles). These organs are responsible for maintaining sperm viability, organizing the sperm in storage, and facilitating the proper release of sperm from storage (reviewed in Neubaum & Wolfner, 1999b; Bloch-Qazi et al., 2003). Judging from the rate of evolution of spermatheca and seminal receptacle proteins, they could be playing a major role in mediating evolutionary processes such as sperm competition, cryptic female sperm choice and co-evolution between male and female reproductive proteins (Prokupek et al., 2008, unpublished data). Inference from observations of stored sperm over time, and sequential structural changes in the female reproductive tract and SSOs after mating, indicates that the functions of SSOs are both dynamic and complicated (Bloch-Qazi & Wolfner, 2003; Adams & Wolfner, 2007). Unfortunately, little is known about how SSOs function even in a well-studied genetic model organism like *Drosophila melanogaster*.

D. melanogaster females multiply-mate and store sperm in two distinct SSOs (Lefevre & Jonsson, 1962; DeVries, 1964; Pitnick et al., 1999). Copulation in *D. melanogaster* results in the transfer of 4000–6000 sperm (Kaplan et al., 1962), approximately a quarter of the transferred sperm are stored, and 30–80% of stored sperm are used for fertilization (Kaplan et al., 1962; Fowler, 1973). *D. melanogaster* have two types of SSOs; the seminal receptacle (SR) and paired spermathecae (ST). The SR stores the majority of the sperm (65–80%) and is the first SSO to store and to release sperm (Gilbert, 1981; Neubaum & Wolfner, 1999b). The ST is the long-term SSO of *Drosophila*. Sperm accumulate slower in the ST and are utilized after the sperm in the SR are at least partially depleted. The process of sperm storage in *D. melanogaster* begins before copulation has ended (Lefevre & Jonsson, 1962; Fowler, 1973; Gilbert et al., 1981), and is complete within six hours of mating (Lefevre & Jonsson, 1962; Gilbert, 1981; Neubaum & Wolfner, 1999a; Tram & Wolfner, 1999; Bloch-Qazi & Wolfner, 2003). Within 90 minutes of mating, ovulation begins and

by hour three, post-mating, the female is laying fertilized eggs (Heifetz et al., 2000), indicating that the SR is actively storing and releasing sperm simultaneously. By 10 h post-mating a noticeable decline is seen in the sperm stores of the SR, and by 48 h the SR sperm stores are ~50% used, whereas the ST sperm stores are only ~15% depleted (Neubaum & Wolfner, 1999a).

It is unclear why *Drosophila* species require both the ST and the SR for optimal sperm storage. One suggestion is that the SR is more efficient at storing sperm, but the sperm are more susceptible to displacement by the sperm of other males (Civetta, 1999; Pitnick et al., 1999; Price et al., 1999). The ST may protect sperm from displacement, thus acting as a reservoir for the sperm from the first male to mate. The ST secretes proteins into the lumen of the ST, where sperm are stored, which could function to keep sperm viable for fertilization, or even quiesce the sperm for long-term storage (Anderson, 1945; Filosi & Perotti, 1975; Pitnick et al., 1999). ST secretions may play a role in capacitating sperm for fertilization; secretions leaving the ST would enter the reproductive tract in the area sperm must pass through in route to the egg (Pitnick et al., 1999). In this scenario, the presence of the ST can be explained by the hypothesis that ST originally functioned as secretory organs and evolved sperm storage function over time (Pitnick et al., 1999).

A study by Prokupek et al. (2008) identified a pool of genes enriched for expression in the ST. Genes encoding for serine proteases accounted for a large proportion (11/43) of the genes identified. In mammals, sperm is capacitated (activated) after encountering extracellular proteases, antioxidants and anti-bacterial proteins in the testis, (Cooper & Yueng, 2006) as well as mucins and glycoproteins in the female reproductive tract (Suarez & Pacey, 2006). The proteases identified in the ST may play a similar role in sperm capacitation as those found in the mammalian testis, giving credence to the theory that the ST originally evolved as secretory organs for sperm capacitation. These proteases could also be interacting with protease inhibitors passed to the female via the seminal fluid at the time of mating (Mueller et al., 2004).

Transcriptome gene expression analysis using microarrays provides a means to investigate gene expression and identify proteins in organs, such as SSOs, for which little functional information is available. This data can also be used to group genes by function, allowing for the identification of operational categories of proteins that play significant functional roles, essentially creating expression profiles for each organ. The expression profiles of SSOs are useful in the inference of specific sperm-related processes and the roles of these organs at different time points after mating.

A study was conducted by Allen & Spradling (2008) which included microarrays of ST from female *D. melanogaster*. This study investigated a gene that affects ST development as well as genes expressed in the ST. It was discovered that products secreted by ST are necessary for sperm maturation and function, paralleling the role of the mammalian epididymis and female reproductive tract. The gene Hr39 was found to function in the development of ST and parovaria, likely playing a role in the control of gene expression within ST of adult females. Hr39 in *Drosophila* is a putative hormone receptor closely related to the Sf1 nuclear hormone receptor of mammals. Mammalian Sf1 is active during embryogenesis for proper reproductive tract development (Allen & Spradling, 2008). The developmental similarities between Hr39 and Sf1 are one indicator of a close connection between Dipteran and mammalian reproductive biology. *D. melanogaster* is a model organism for studies of reproduction from which insight into human reproduction can be obtained.

Microarray studies investigating the genetic changes occurring in mated compared to virgin female *Drosophila* provide insight into endogenous changes following mating (Lawniczak & Begun, 2004; McGraw et al., 2004, 2008; Mack et al., 2006). In these studies over 1700 genes show a difference in expression in the female at 1–3 h post-mating, though the magnitude of most of these changes is small (less than twofold). These transcript-level changes were not confined to the reproductive tract, but occurred throughout the entire body. A microarray study of the lower reproductive tract was conducted by Mack et al. (2006). Mack et al (2006) observed a peak of expression in the lower reproductive tract occurring at 6 h post-mating. Specifically, the number of transcript-level changes was eight times higher at 6 h than at 3 h post-mating.

Although the previously listed studies provided valuable information in terms of expression profiles of the post-mated female, there is still limited information detailing post-mating changes within SSOs, and no information comparing SSOs. In this study, transcriptome analysis was conducted for each of the female SSOs of *D. melanogaster*, before and after mating to create detailed organ-specific transcription profiles. Additionally, this study asks the following questions:

- Do SSOs respond to mating?
- Are there organ-specific differences in the transcription profiles which indicate unique biological functions?

The gene expression data from this project provides insight into female sperm storage likely to be applicable to a wide range of species. The present study identified genes differentially expressed within and between each SSO of *D. melanogaster* at two time points post-mating, providing the most detailed information on sperm storage gene expression patterns available for any species.

Results and discussion

In this study gene expression changes were investigated in the SSOs of *D. melanogaster* at two time points

(3 and 6 h) after mating. A pre-mating (virgin) state was used as a baseline for comparison to gene expression patterns after mating. The 3 h post-mating time point represents the time needed for the complete entrance of sperm into storage, as well as the start of egg deposition and fertilization (Heifetz et al., 2000; Heifetz et al., 2001). The inclusion of the 6 h time point was based on a previous study showing a peak in change of gene expression in the lower reproductive tract 6 h after the completion of mating (Mack et al., 2006). Distinctively different patterns of gene expression were found for the two types of organs, and for pre- vs. post-mating time points; differentiation was greatest between organs compared to the effect of mating. The data on expression patterns produced for the two SSOs will be valuable for future functional research of sperm storage and studies on the role of these organs in evolution.

Average expression

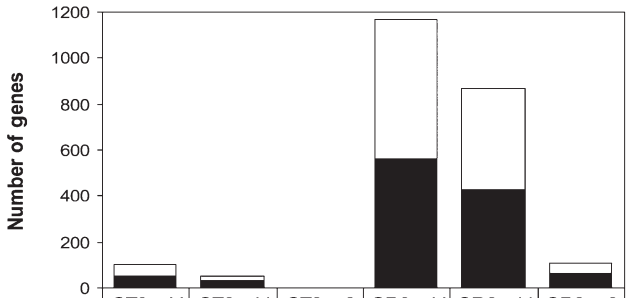
Genes included in analysis were those predicted to be 'present' by Gene Chip Operating Software (Affymetrix, Santa Clara, CA, USA). Genes were ranked by average expression, and highly expressed genes in each category were examined in more detail. Although genes expressed as lower levels are likely to play important roles, by looking at the genes with the highest expression we identified genes which are potentially organ-specific. Expressed genes were classified (Supporting Information Table S1) using Gene ontology (a controlled vocabulary for categories of genes), and identified in FlyAtlas which is a compilation of expression data from multiple tissues (Chintapalli et al., 2007).

The highest expressed gene at all time points in the SR is of unknown function (*CG18628*). Genes of unknown function are of interest in terms of their potential to play organ-specific roles. A BLAST search determined that this gene is highly similar to a serine protease with a secretion signal. *CG18628* was also identified using FlyAtlas as being up-regulated in the testis compared to the whole body (Supporting Information Table S1), suggesting that the function of this protein is important in both male and female reproductive tracts. It is possible that *CG18628* is a serine protease whose function is important for sperm and sperm storage. The high presence of *CG18628* in the virgin SR may indicate that this protease is creating and maintaining a specific environment within the SR.

Proteases accounted for approximately 10 percent of the top 100 expressed genes of the ST. At all three time points in the ST, the gene with the highest expression was a serine protease (*CG9897* in virgin and *CG17239* in ST3 and ST6). *CG17239* remains the highest expressed gene in ST at least 3 days after mating (Allen & Spradling, 2008), alluding to the importance of this particular protein for ST function. *CG17239* is found on the long arm of chromosome 2, in a cluster of 4 proteases (*CG17239*, *CG17012*, *CG17234*, and *CG17240*), all of which have secretion signals and are expressed in ST. This cluster of proteases has been previously described as being regulated by mating (Lawniczak & Begun, 2007). A study by Arbeitman et al. 2004 identified *CG17012* as being specifically expressed in the spermathecae and parovaria. In total, ten of the top 100 expressed genes in the ST were serine proteases; eight were described by FlyAtlas as present only in the ST (Supporting Information Table S1). ST proteins with secretion signals could directly interact with male seminal fluid proteins or sperm in the lumen of this organ. Additionally, these proteins may exit the ST, and function in reproductive processes (such as sperm capacitation) outside the ST. In general, serine proteases have been identified as important molecules in reproduction, both in the seminal fluid and in the female reproductive tract (Swanson et al., 2001; Lung et al., 2002; Lawniczak & Begun, 2004, 2007; Ravi-Ram & Wolfner, 2007). Serine proteases were highly over-represented among genes identified in a hybrid-selected ST cDNA library (Prokupek et al., 2008). More serine proteases were highly expressed in the ST but, at 3 h post-mating five of the 10 highly expressed serine proteases in the ST exhibit a similar pattern of expression in the SR. This pattern of gene expression suggests some similarity in function of the SR at 3 h post-mating and ST.

All three of the yolk protein (YP) genes in *D. melanogaster* (*YP1-YP3*) were highly expressed in the ST at all time points, but only *YP1* had high expression in the SR (Supporting Information Table S1). YP genes are expressed in the fat bodies and follicle cells. YPs are the most abundant molecules in yolk and the most abundant soluble proteins in the female body (Barnett et al., 1980; Brennan et al., 1982). The presence of YPs in the SSOs may indicate that these organs contribute to yolk production for eggs as suggested by Allen & Spradling (2008). Alternatively, YPs might play a novel role in the SSOs by protecting sperm from oxidative damage. In honeybees, the antioxidant activity of vitellogenins (similar to yolk proteins) promotes individual survival (Seehuus et al., 2006), suggesting YPs might similarly promote the survival of sperm. It is important to note that the ST is surrounded by closely associated fat body tissue; the presence of YPs in this, and previous, studies could be due to a failure to completely remove all fat cells from the exterior of the ST during dissection. However, the SR has no observable associated fat body and it does express *YP1*. Moreover, a conscious effort was made to remove ST fat body in the present study and yet the genes were expressed at a high level.

A high number of ion transport, muscle contraction, metal binding genes and immunity genes were highly expressed in the SR compared to ST, suggesting they might play a more prominent role in this organ. Perhaps ion transport genes act to generate and maintain a suitable environment in the SR before and after mating. Highly expressed muscle contraction genes may play a role in the



	ST3 – V	ST6 – V	ST3 – 6	SR3 – V	SR6 – V	SR3 – 6
□ down	51	17	0	609	437	46
■ up	50	31	0	559	428	63

Figure 1. Post-mating gene expression changes in the sperm storage organs. Bars indicate the number of differentially expressed genes (abs (FC) □ 2; FDR □ 0.05) at two time points post-mating. Black shading in each bar represents the proportion of genes up-regulated in mated females; white represents the proportion of genes down-regulated in mated females. ST = spermathecae, SR = seminal receptacle, V = virgin, 3 = 3 h post-mating, 6 = 6 h post-mating.

mechanism of sperm entrance into or release from the SR. On the other hand, lipid metabolism and oxoreductase gene categories were similarly represented in both SSOs suggesting some common functions for both these organs, possibly in maintaining optimal conditions for sperm function. Both the ST and SR showed a high number of genes involved in nucleic acid binding, most of which are ribosomal proteins, and are not thought to play organ-specific roles.

Differential gene expression

Differential gene expression was measured by pairwise comparisons of virgins to the two time points post-mating for ST and SR (Supporting Information Table S2). After mating, a higher number of genes were differentially expressed within the SR (1444) compared to the ST (104) (Figure 1). In comparison with our study, far fewer genes were differentially expressed (two-fold or greater) in the whole body of *D. melanogaster* (46 genes), or lower reproductive tract (38 genes) after mating, compared to virgins (Lawniczak & Begun, 2004; McGraw et al., 2004). A possible explanation for the paucity of genes differentially expressed in the whole-body after mating, or in the reproductive tract, is that opposing gene expression patterns in different organs or tissues negate each other (McGraw et al., 2004). The high number of differentially expressed genes in the SR compared to the ST can possibly be explained by the fact that the SR is both actively storing and releasing sperm at 3 h post-mating, and by 6 h is releasing sperm at a steady rate when the ST is presumably only storing sperm at both of these time points. The large number of differentially expressed genes after mating in the SR suggests that this organ has especially complicated and diverse functions. Other features of the ST and SR could account for the observed difference in gene expression of the two organs. For example, structurally the organs are quite different; the SR is more flexible, whereas the ST is chitinous.

RT-PCR

RT-PCR was performed for three genes (*CG9897*, *CG17239,* and *CG18628*), and data was compared with the microarray data in the form of differential expression (fold change). Patterns of gene expression (direction of expression changes) showed good correspondence between microarrays and RT-PCR (Supporting Information Table S3). Another indicator of the validity of our microarray data is that it matches well with comparable data (Allen & Spradling, 2008). As one example, the most abundant genes in the ST in the present study were also found to be highly expressed in Allen & Spradling (2008).

Over-represented gene categories

All genes were subjected to statistical analyses using Robust Microarray Analysis, an empirical Bayesian model, false discovery rate calculations, Gene Set Enrichment Analysis, and DAVID. No statistically significant over-represented gene categories were found up-regulated in the mated ST to virgin comparisons. At 3 h post-mating in ST, two categories of down-regulated genes were over-represented; lipid metabolism and electron transport (Supporting Information Table S4).

At 3 h post-mating in the SR to virgin comparisons, significantly up-regulated categories of genes over-represented in the SR included genes involved in defense, fertilization, female gamete generation, nervous system development (only 6 h vs. virgin) and regulation of translation. In the SR, down-regulated over-represented gene categories included electron transport, oxidative phosphorylation (only 3 h vs. virgin), and coenzyme metabolism (only 3 h vs. virgin).

Organ-specific changes in gene expression

To evaluate organ-specific patterns of gene expression the ST was compared to the SR and values of differential expression were calculated for each time point (Supporting Information Table S2). Comparing virgin SR and ST, a total of 1669 genes were differentially expressed; 641 genes up-regulated and 1028 down-regulated in STV compared to SRV. For 3 h post-mating SR and ST, a total of 1316 genes were differentially expressed; 500 genes up-regulated and 816 down-regulated in ST3 compared to SR3. At 6 h post-mating SR and ST, a total of 2353 genes were differentially expressed; 987 genes up-regulated and 1366 down-regulated in ST6 compared to SR6.

Analysis of the data described in the preceding paragraph revealed interesting patterns. Over-represented categories of genes up-regulated in STV compared to SRV include multiple categories of metabolism (i.e. lipid, monosaccharide) and electron transport. In SRV vs. STV

comparisons the following categories were up-regulated: development (organ, nervous system, embryonic), nerve impulse transmission and ion transport. Categories up-regulated in ST3 include biosynthesis of macromolecules and protein targeting, while categories up-regulated in SR3 include ion transport, protein kinase regulation, development (tissue, embryonic) and cell mobility. For ST6, metabolism (carboxylic acid, lipid, monosaccharide, coenzyme), biosynthesis, heme binding and peptidase activity were induced. SR6 shows an increase of nerve impulse transmission, development, chemosensory behavior, ion transport, protein kinase regulation, and cell differentiation (Supporting Information Table S4). *CG18628*, a gene of unknown function, whose expression is highest in the SR at all time points (discussed previously), is differentially expressed between the two organs. The expression of this gene is more than 10-fold higher in the SR compared to the ST at all time points. Additionally, *CG18628* is down-regulated in ST3 relative to virgin females. These expression patterns have been confirmed with RT-PCR (Supporting Information Table S3). Given the tissue specificity of this gene, as well as the interesting patterns of differential expression, *CG18628* could be playing a very interesting role in SR (and testicular) function and warrants further investigation.

The large number of genes differentially expressed between organs indicates organ-specific and temporal diversification of biological functions. The induction of metabolism-related genes in ST may be associated with the preparation of sperm for long-term-storage. In contrast, the SR shows overrepresentation of up-regulated nervous system impulse genes, cell motility, ion transport, protein kinases activity and chemosensory genes, all of which could play integral roles in establishing the internal milieu of the organ and the preparation of sperm for fertilization (including timing of release and sperm capacitation).

Gene Set Enrichment Analysis (GSEA)

Gene Set Enrichment Analysis (GSEA) was also used to classify over-represented categories of genes, characterized by Gene Ontology. Over-represented gene categories identified by GSEA were similar to those found by the DAVID analysis in the present study. For example, an up-regulation of serine-type endopeptidase activity was observed in STV compared to SR3, ST3 compared to SR3 and STV, SR3 compared to SR6, and in ST6 compared to ST3. Metallopeptidase activity, NADH dehydrogenase activity, transport, alcohol metabolism, neurotransmitter transport and cell adhesion were higher in the SR compared to ST at all three time points. No categories of genes were consistently up-regulated for all three time points in comparisons of ST and SR, but categories of endopeptidase activity, neprilysin activity and ER targeting were induced in STV and ST3 compared to the SR at corresponding time points. In comparisons of ST3 and SR6 vs. virgin comparisons, genes involved in cytochrom-c oxidase activity, metalloendopeptidase activity, tricarboxylic acid cycle, neuropeptide receptor activity and cellular metabolic processes were all down-regulated.

GSEA was also used to identify pathways which were significantly differentially expressed in comparisons of the SSO sample sets (Supporting Information Table S5). Carbohydrate metabolic pathways were down-regulated in ST3 and ST6 compared to virgin. A similar, but not as strong, trend was observed in comparisons of SR3 and SR6 to virgin. In ST6 compared to SR6, carbohydrate metabolism was highly up-regulated in the ST. Three pathways related to protein modification, and four pathways related to lipid metabolism were down-regulated in ST3 and ST6 compared to virgin, a trend not observed in SR comparisons. In ST6 relative to SR6, several amino acid, carbohydrate, and lipid metabolism pathways were found induced. The induction of metabolic processes observed in the ST could be due to preparation for the long term sperm storage. A number of pathways related to the metabolism and biodegradation of xenobiotics were down-regulated in ST3 and ST6 comparisons to virgin. A similar, but not as strong, trend was observed between virgin to SR3 and SR6 comparisons. Xenobiotic biodegradation and metabolism pathways include metabolism by cytochrome P450s. The down-regulation of *P450* genes, and possible implications, are discussed later in the 'cytochrome P450' section.

Differential expression data was also compared to data from Gershman et al (2007), and Guertin et al. (2006) which provides a list of insulin signaling target genes. Gershman identified 995 genes as potential transcriptional targets to dFOXO. Guertin identified 986 genes potentially regulated by dTOR. The different clusters of differentially expressed genes (see below), along with the differential expression data were compared to the insulin signaling gene lists to identify mating induced changes due to insulin signaling which is a key regulator of metabolism. There was no strong evidence for up or down-regulation of signaling in any comparison (data not shown). This suggests that within the SSOs, mating does not have a disproportionate effect on insulin signaling target genes.

Cluster analysis

Shared expression patterns of the 2054 differentially expressed genes were investigated using hierarchical cluster analysis (Figure 2). By clustering genes based on expression pattern similarity six sub-clusters of genes were created. Cluster 1 was the largest containing 1021 genes, clusters 2, 3, and 4, 5 and 6 contained 357, 471, 194, five and six genes respectively (Supporting Information Figure S1, a–f). Each cluster was analyzed using DAVID to detect overrepresentation of categories of genes.

Cluster 1 contained a significant overrepresentation of genes in the categories of metabolism (carboxylic, lipid,

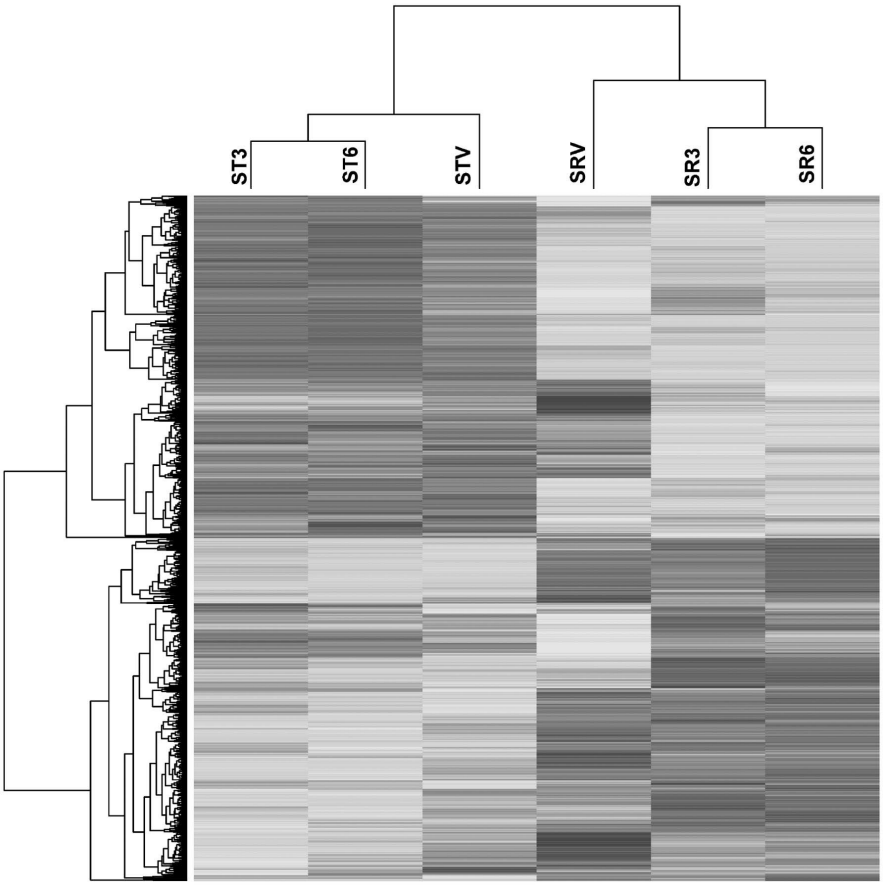


Figure 2. Clustering of the expression patterns of genes identified as significantly differentially expressed (abs (FC) □ 2, FDR □ 0.05). ST = spermathecae, SR = seminal receptacle, V = virgin, 3 = 3 h post-mating, 6 = 6 h post-mating. Shading on the heatmap corresponds to expression levels (dark = low, light = high expression).

monosaccharide, coenzyme and protein), electron transport, protein targeting to the ER, alcohol catabolism (the ALDH gene is associated with control of redox state) and signal peptidase activity. Gene categories in cluster 2 were primarily involved in developmental processes (including imaginal disc morphogenesis, organ and mesoderm development), muscle contraction and transport. The gene categories in cluster 3 were female gamete generation, chemosensory, kinase activity, cell fate determination and defense responses to other organisms. Tissue development was the only significantly over-represented category of cluster 4. Cluster 5 consisted of five genes, which included pepsin, an odorant receptor, a gene of unknown function, a gene involved in localization, and a Toll receptor that plays an important role in innate immunity. Cluster 6 contained four serine-protease genes, a gene of unknown function, and a gene involved in bacterial defense.

Defense genes

Genes involved in defense were up-regulated in the SR at both time points compared to virgin levels, a phenomenon not observed in the ST samples. Additionally, in the SR, four genes (*Drs*, *Def*, *TcpIV*, *Rel*) involved with defense/immunity are highly expressed (top 100 expressed genes) (Supporting Information Table S1). *Drs* maintains high levels in the ST at all times (pre and post-mating). Drosomyocin has been found previously to be constitutively expressed in both types of SSOs in *D. melanogaster* (Ferrandon et al., 1998), which suggests that *Drs* may play a role in immunity/defense that is not directly related to mating. Perhaps *Drs* expressed within the SSOs is released into the reproductive tract, participating in broad scale immunity of the reproductive tract.

Sperm capacitation

Sperm entering the female reproductive tract must undergo a series of biochemical transformations, called capacitation, before they are able to bind to the egg. In mammals, protein kinases are involved in the regulation of intracellular Ca^{2+} during sperm capacitation (Breitbart & Naor, 1999). One of the first of a series of steps in capacitation is an influx of calcium (as well as biocarbonate and hydrogen peroxide), which collectively activate adenylyl cyclase to produce cyclic AMP. In the current study, protein kinase regulator activity is up-regulated in SR3 and SR6 compared to virgin SR and compared to ST3 and SR6 respectively. Regulation of protein kinases may be a mechanism to control the rate of capacitation within the SR perhaps similar to the role of kinases in mammals.

The ST has previously been identified as important for sperm capacitation (Allen & Spradling, 2008). Females without ST mated and stored sperm successfully, but re-

mained sterile suggesting that ST secretions mix with sperm in the reproductive tract to make them fertilization competent regardless of their eventual storage site.

Cytochrome P450 genes

A total of 27 cytochrome *P450* genes were differentially regulated in ST and SR. When compared to virgin, one *P450* gene is up-regulated and nine are down-regulated in ST3, zero up and two down-regulated in ST6, six up and 16 down-regulated in SR3, and three up and 16 down-regulated in SR6. In both organs there is down-regulation of the *P450* genes in response to mating. GSEA analysis also finds a down-regulation of *P450* metabolic pathways in mated compared to virgin females. The down-regulation of *P450* genes associated with mating was observed in a study of whole female *Drosophila* (McGraw et al., 2004).

In insects, cytochrome *P450* genes code for proteins with diverse functions, including detoxification (Feyereisen, 1999; Wilson, 2001). Given that sperm are allogeneic antigens in the female reproductive tract, the viability of spermatozoa in a female reproductive tract over extended periods of time may be associated with immunologic recognition and immune tolerance. The down-regulation of detoxification genes such as *P450*s may be a primary example of such immune tolerance. A decrease in immune response associated with sperm storage has been previously documented in ants (Baer et al., 2006).

It would be counter-productive for the female to 'detoxify' certain molecules or metabolites received from males at the time of mating (see argument in *P450* genes) but, as seen by the up-regulation of specific defense genes, there is an increase in some forms of immunity following mating. Perhaps, the decrease in *P450*s is a mechanism of 'forced immune tolerance', induced by seminal proteins against the female's best interest. The down-regulation of *P450*s could be a male mechanism to circumvent female counter adaptations to the 'toxic' control of female function. Male accessory gland proteins, such as Acp62F have been found to be toxic to females, decreasing female lifespan. Various ways that males reduce the survival females with Acps, and the female counter-responses, could be an example of sexually antagonistic coevolution. Sexually antagonistic coevolution is currently a major topic in evolution and the female *P450* gene response to mating might fall into this context.

Hormone and signaling genes

Cytochrome *P450* genes are also involved in the biosynthesis of ecdysteroids and Juvenile Hormone (JH) (Feyereisen, 1999; Wilson, 2001). Mating is associated with an increase in JH synthesis (Moshitzky et al., 1996), the down-regulation of *P450*s could be one way females regulate JH levels after mating.

After mating both SSOs organs express endocrine system genes including JH epoxide hydrolases (*JHeh2* and *JHeh3*), genes coding for enzymes that inactivate juvenile hormone. *JHeh2* was expressed at higher levels in the SR at all time points relative to ST. Earlier studies found down-regulation of this gene at 6 h post-mating, compared to virgins, in the lower reproductive (Mack et al., 2006). Our study also found the levels as 6 h to be lower than virgin levels for both organs, but the difference was not statistically significant. *JHeh3* expression was higher at all time points post-mating in the ST relative to the SR. *JHeh3* was significantly up-regulated in ST3, and significantly down-regulated ST6 compared to virgin. The possible significance of high expression of these genes is that they might counter the effects of up-regulation of *JHamt* (a JH biosynthesis pathway gene) after mating in the present study. Expression of *JHamt* increased > four-fold at the 3 h and 6 h time points in both organs compared to virgin ($P < 0.1$). The reproductive and evolutionary implications of differential expression of *JHeh3* and *JHeh2* in response to mating are discussed in more detail in Prokupek et al. (2008).

Acp70A (sex peptide – SP) enters the SSOs bound to the sperm tail, and is released over time by proteolytic cleavage. SP acts by stimulating JH synthesis which in turn induces egg-laying and decreases a female's receptivity to remating; it also makes an important contribution to the cost of mating to females (Aigaki et al., 1991; Chapman et al., 2003; Liu & Kubli, 2003; Wigby & Chapman, 2005). A connection may exist between JH catabolism in the SSOs and JH synthesis induced by SP. Specifically, females may be acting to counter a detrimental physiological effect induced by males. The sex peptide receptor (CG16752 or SPR) (Yapici et al., 2008), required for the effect of SP, was expressed in both the ST and SR in all three time points. The level of this protein appeared to be stable within the ST at all time points, but fluctuated within the SR, being significantly down-regulated in SR3 compared to virgin. In SR6, SPR return to levels not significantly different from virgin levels.

Interactions with Acps

During the time of mating male *Drosophila* pass to females a number of proteins via their seminal fluid, some of which enter the SSOs. These proteins, in particular, male accessory gland proteins (Acps), are responsible for eliciting a suite of post-mating responses in the female. Female responses include increases in ovulation, ovipositioning, and mating refractoriness, as well as sperm storage and decreases in female lifespan (Wolfner, 1997; Wolfner, 2002). At least one Acp (Acp36DE) has been associated with proper sperm storage. Acp36DE associates with the sperm mass and localizes at the openings of the SSOs as well as within them (Bertram et al., 1996; Neubaum & Wolfner, 1999a). Female genes found to be differentially expressed in the present study might directly interact with Acps. For example, the cleavage of SP is accomplished by a trypsin member of the serine protease family (Peng

et al., 2005). Most Acps are no longer detected after 6 h post-mating, but SP continues to be cleaved from sperm for the entire time sperm are in storage (Peng et al., 2005). The identity of the specific serine protease involved in the release of SP is not known, but it is quite possibly female derived. Four of the seven Acps which enter the SSOs are serine protease inhibitors (serpins) (Mueller et al., 2004). Serpins bind to serine proteases, blocking their proteolytic function, and might interact with proteases in the ST. Proteases found within SSOs could be interacting with male proteins in cascades, similar to those seen in immune function and blood or sperm coagulation (Malm et al., 2000; Overduin & de Beer, 2000; Kim et al., 2008).

Discussion

Microarrays were used to investigate the transcriptome of *Drosophila melanogaster* SSOs in virgin females and at two time points post-mating. At each time point, high numbers of genes and gene sets were differentially expressed between the two SSOs, indicating that each organ plays a unique role in the process of sperm storage. ST were enriched for genes involved in proteolysis and metabolism. The products of these genes could be interacting with male proteins such as protease inhibitors or could be enzymes that proteins that synthesize lipids and carbohydrates for sperm maintenance. Other ST proteins, such as yolk proteins, could protect sperm perhaps by acting as an antioxidant. The SR exhibited a number of over-represented genes involved in localization, signaling, and ion transport. These genes could be working to maintain a homeostatic environment, as well as serving communication roles between the SR and oocytes to ensure the proper timing of fertilization. More genes were differentially expressed in the SR perhaps due to the dual role of sperm storage and sperm release played by this organ at the time points used in the present study. Further investigation of the roles of proteins in these two organs will allow for a more complete understanding of genes playing essential roles in the process of sperm storage, maintenance, and use.

Experimental procedures

Stock maintenance

The Canton-S (CS-C) stock of *Drosophila melanogaster* was used in the present study. Flies were reared in larval density-controlled vials, on a standard *Drosophila* diet in a temperature controlled environment (25 □C) with a standard 12/12 light/dark cycle. Adults were collected and held as virgins, and used for sample collection at 4–7 days old.

Mating conditions and organ sample collection

The ST sample included both spermathecae from each female and the attached spermathecal duct; the SR sample included the single seminal receptacle from the same females as the ST collections. For mated female collections, a single female was placed with a single male until mating was observed. Immediately after the mating had ended naturally the female and male were separated and post-mating time began.

Tissues were collected from virgin, 3 h post-mating, and 6 h post-mating females. Females were dissected on a cold plate in RNase later (Ambion, Austin, TX, USA). The ST and SR were removed and immediately placed into TRIzol (Invitrogen, Carlsbad, CA, USA) for RNA extraction. From each dissection event, conducted over a standardized period of time, we obtained approximately 160 ST and 80 SRs.

RNA extraction

The freshly dissected tissues from 80 females were ground in TRIzol and liquid nitrogen using mortar and pestle. An additional aliquot of TRIzol was used to rinse the mortar and pestle and the rinsate was added to the sample to ensure sample retention. RNA was extracted from freshly dissected ST and SR samples using RNeasy microprep kit (Qiagen, Hilden, Germany) and quantified using a NanoDrop (ND-1000) spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA). To obtain sufficient quantities of RNA from ST and SR, the RNA for each microarray sample was pooled from 10 independent dissection periods and extractions. Each sample used for one microarray chip (ST or SR) contained the RNA from the organs of ~800 females.

Microarray assays

ST and SR microarray cRNA sample preparation used a two-cycle target and labeling kit (Affymetrix) which is designed for starting RNA levels of □ 100 ng for each microarray chip. Even with pooling, there were not sufficient quantities of RNA to perform cDNA synthesis without amplification. For both the ST and SR 100ng of total RNA were used for amplification prior to cDNA synthesis. The Gene Chip *Drosophila* 2.0 array (Affymetrix), which includes probe sets to 18 880 *D. melanogaster* genes, was used to measure gene expression. Hybridization to the *Drosophila* Gene Chips was performed by the Microarray Core Facility at the University of Nebraska-Lincoln. Three biological replicates were run for each treatment (virgin ST and SR, 3 h post-mating ST and ST, 6 h post-mating ST and SR).

Transcriptional profiling

The quality assessment of microarrays included QA-plots and the calculation of Pearson correlation coefficients between replicates. One chip (seminal receptacle, 3 h time point) was removed because the inter-chip variation was too high. In one set of analyses, the Gene Chip Operating Software (Affymetrix) made calls for each as 'absent', 'marginal' or 'present' based on hybridization intensity. Independently of the Affymetrix software, Robust Multichip Averaging (RMA) procedure (Irizarry et al., 2003) was used for background correction and quantile normalization of the measured light intensities. The corrected light intensity values were used as statistical estimates for transcript levels.

To detect temporal patterns of female-expressed genes, pairwise comparisons were performed between virgins, 3 h, and 6 h post-mated females both in SR and ST. Patterns of gene expression changes between organs were detected by pairwise comparisons between the ST and SR at each time point. Differential expression of genes was determined using both fold change (FC), and false

discovery rate (FDR). FDR, being defined as the expected proportion of false positives among the declared significant results, is a multiple test correction method (Benjamini & Hochberg, 1995; Pawitan et al., 2005). The cut-offs used for differentially expressed genes were FC □ 2, and a FDR q-value □ 0.05.

In order to search for shared expression patters, we used complete linkage hierarchical cluster analysis. Log fold changes for the six treatment combinations compared to the average of the six treatment combinations were used when computing distances. Any gene which was determined to be differentially expressed in the pairwise comparisons was included in the analysis. Clustering was visualized by a heat map (Figure 2), created in R (R Development Core Team, 2008).

Gene set analysis

Unifying biological themes that are bases on the statistical significance of genes sets, as opposed to single genes, were identified by two different approaches. The Database for Annotation, Visualization and Integrated Discovery (DAVID) (http://david.abcc.ncifcrf.gov) was used to classify gene function for the genes present in all lists using the most relevant Gene ontology (GO) associated terms. DAVID was used to determine significant enrichment of known functional annotations within our differentially expressed gene lists. DAVID calculates the probability of observed representation of genes within a given category compared to a pre-defined genome list corresponding to the same category.

In contrast to DAVID, Gene Set Enrichment Analysis (GSEA) (Subramanian et al., 2005) analyzes the enrichment of all genes in a set like biochemical pathways and Gene ontology (Ashburner et al., 2000). Since a number of genes may show contradicting regulation, or several individual genes do not display significant differential regulation, GSEA provides an alternative approach to excluding genes that do not show differential expression. By ranking all genes using fold change, p-value and related metrics, the enrichment of a gene set in the up- down-regulated or unchanged domain is estimated. A normalized enrichment score and an FDR q-value are robust, unbiased statistical measures for the whole set. Significantly up-regulated gene sets are assigned high enrichment scores, while down-regulated sets receive extreme negative scores. Comparing these scores to those obtained from random permutations gives an estimate of statistical significance. Multiple test correction is performed by False Discovery Rate (FDR < 0.25) (Benjamini et al., 2001; Reiner et al., 2003).

Biochemical pathways were obtained from the Kyoto Encyclopedia of Genes and Genomes Pathways Database (KEGG, Okuda et al., 2008). Gene ontology sets for biological processes, molecular functions, and cellular localizations (Ashburner et al., 2000) were downloaded from FlyBase (Drysdale, 2008; Tweedie et al., 2008) (http://flybase.org/). Both databases were reformulated for GSEA using PERL programs.

RT-PCR

The reproducibility of microarray-determined transcript abundance was tested using quantitative real-time RT-PCR Taqman assays (Applied Biosystems, Foster City, CA, USA). Gene expression was examined for three genes (*CG9897*, *CG17239*, and *CG18628*). Mating and sample collection was highly similar to that used in the microarray sample collection. Thirty females were used for RNA extraction for each sample and RNA extraction was performed using the MagMax-96 for Microarrays Kit (Ambion). Primers and probes for each gene and for the housekeeping gene RP49 were ordered as pre-designed TaqMan Gene Expression Assays (ABI). cDNA construction used the High-capacity cDNA reverse transcription kit (ABI). For the assay, 12.5 ng of cDNA was mixed with TaqMan fast universal PCR master mix (2X) and the run was performed on an ABI 7500 FAST thermocycler, under the specified conditions for a Fast run type. The gene-expression levels obtained by RT-PCR were normalized to ribosomal protein 49 (rp49). Differential expression was calculated from the normalized expression values, and compared to the microarray differential expression data.

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Supporting Information (following pages)

Figure S1. Heat maps and boxplots of genes clustered by expression patterns.
(6 pages)

Table S1. Categories of genes highly expressed in the ST and SR
(1 page)

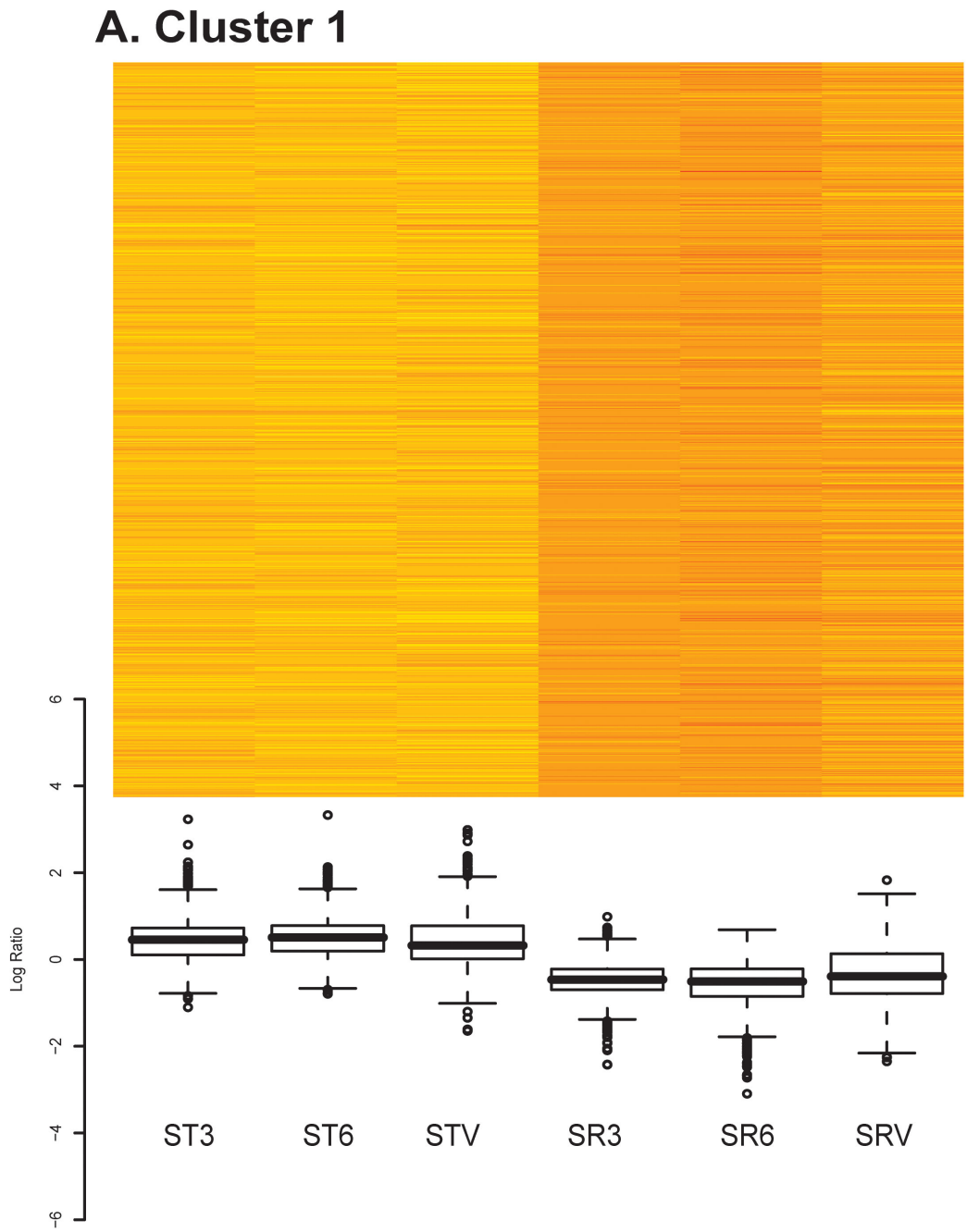
Table S2. Differential expression data from pairwise comparisons
(238 pages)

Table S3. Differential expression results for microarray compared to qRT-PCR
(1 page)

Table S4. Categories of differentially regulated genes identified by DAVID
(1 page)

Table S5. Signifcantly up-regulated pathways analyzed by Gene Set Enrichment Analysis (GSEA)
(2 pages)

Figure S1, A–F. Heat maps and boxplots of genes clustered by expression patterns.



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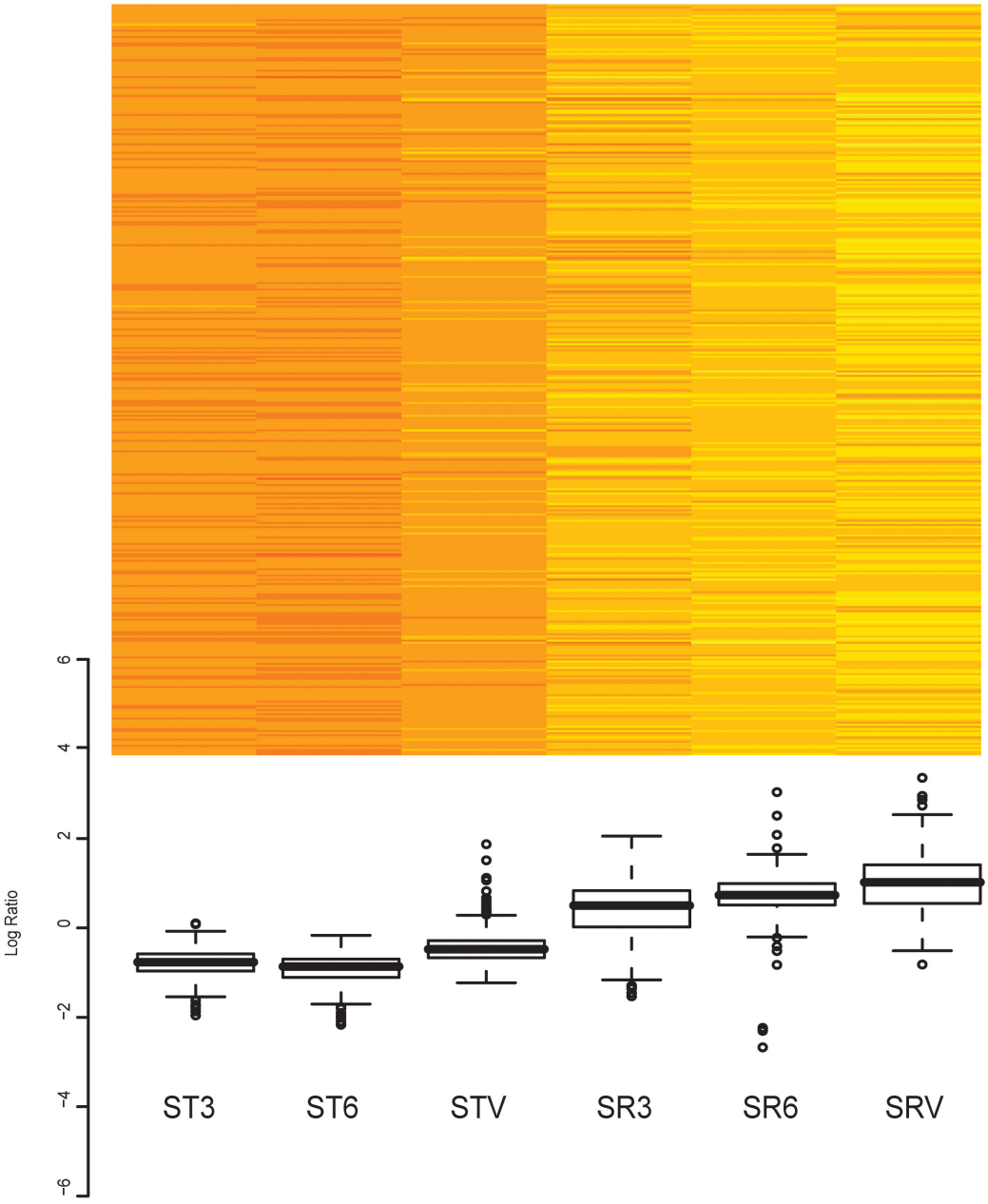

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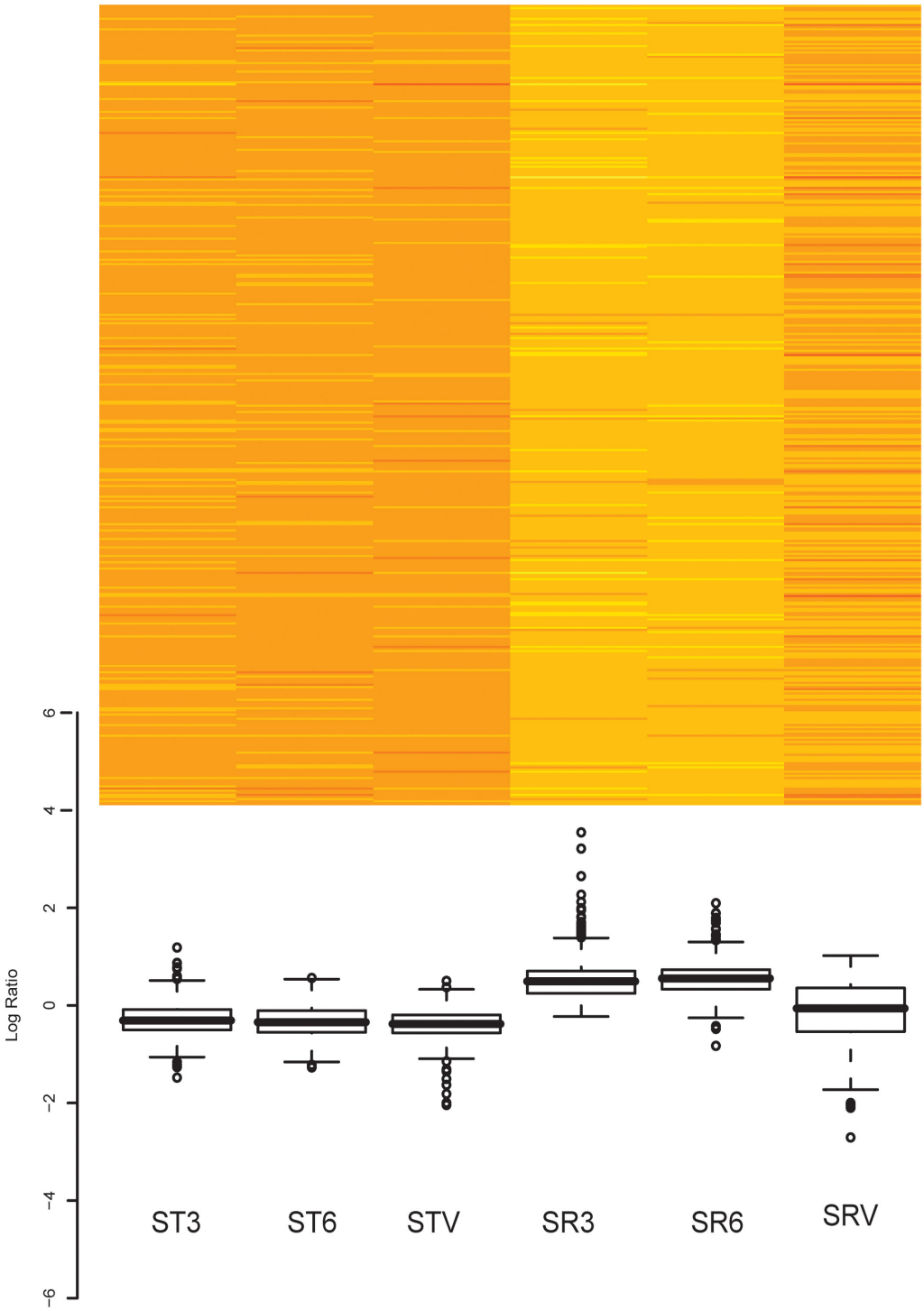

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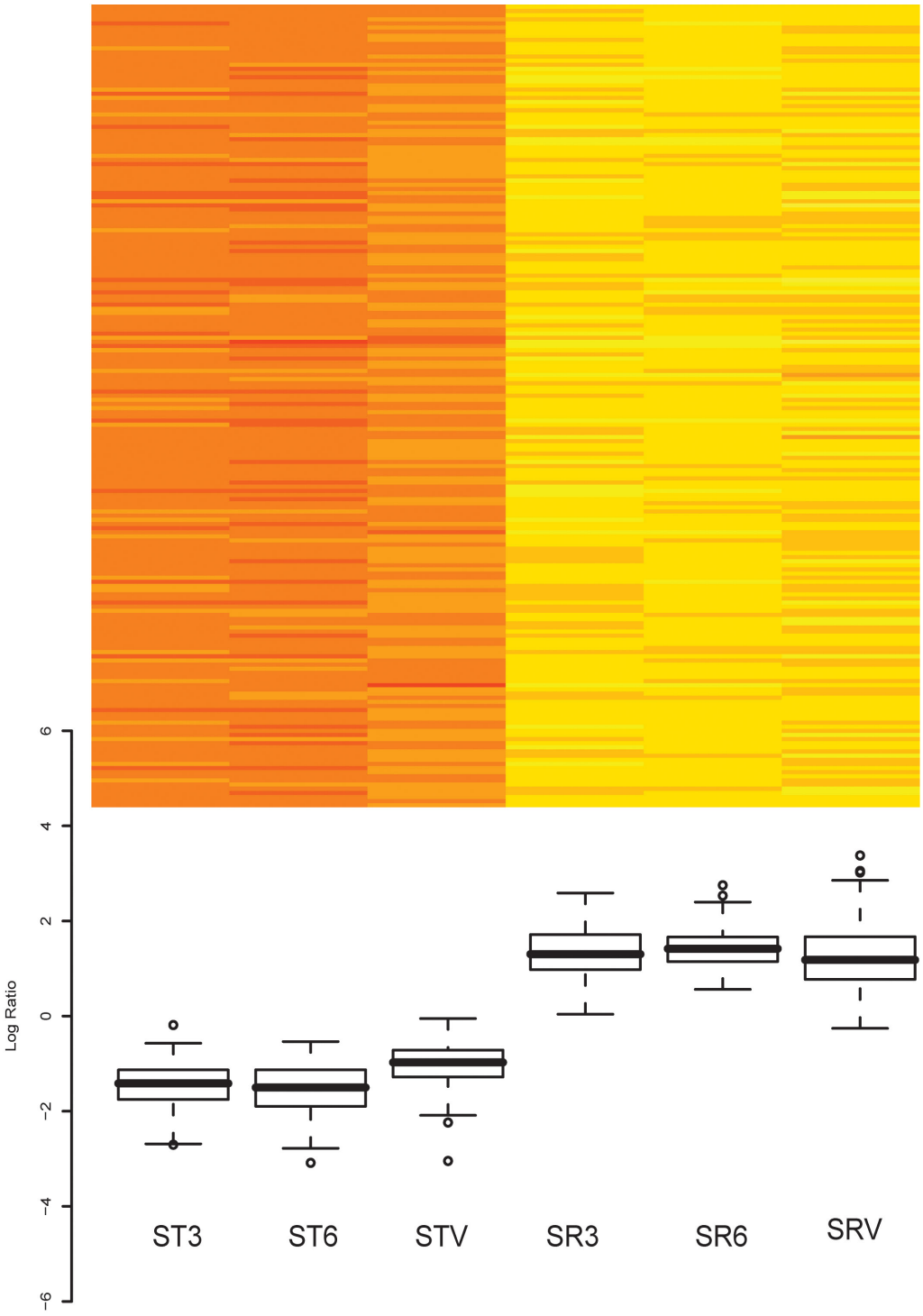

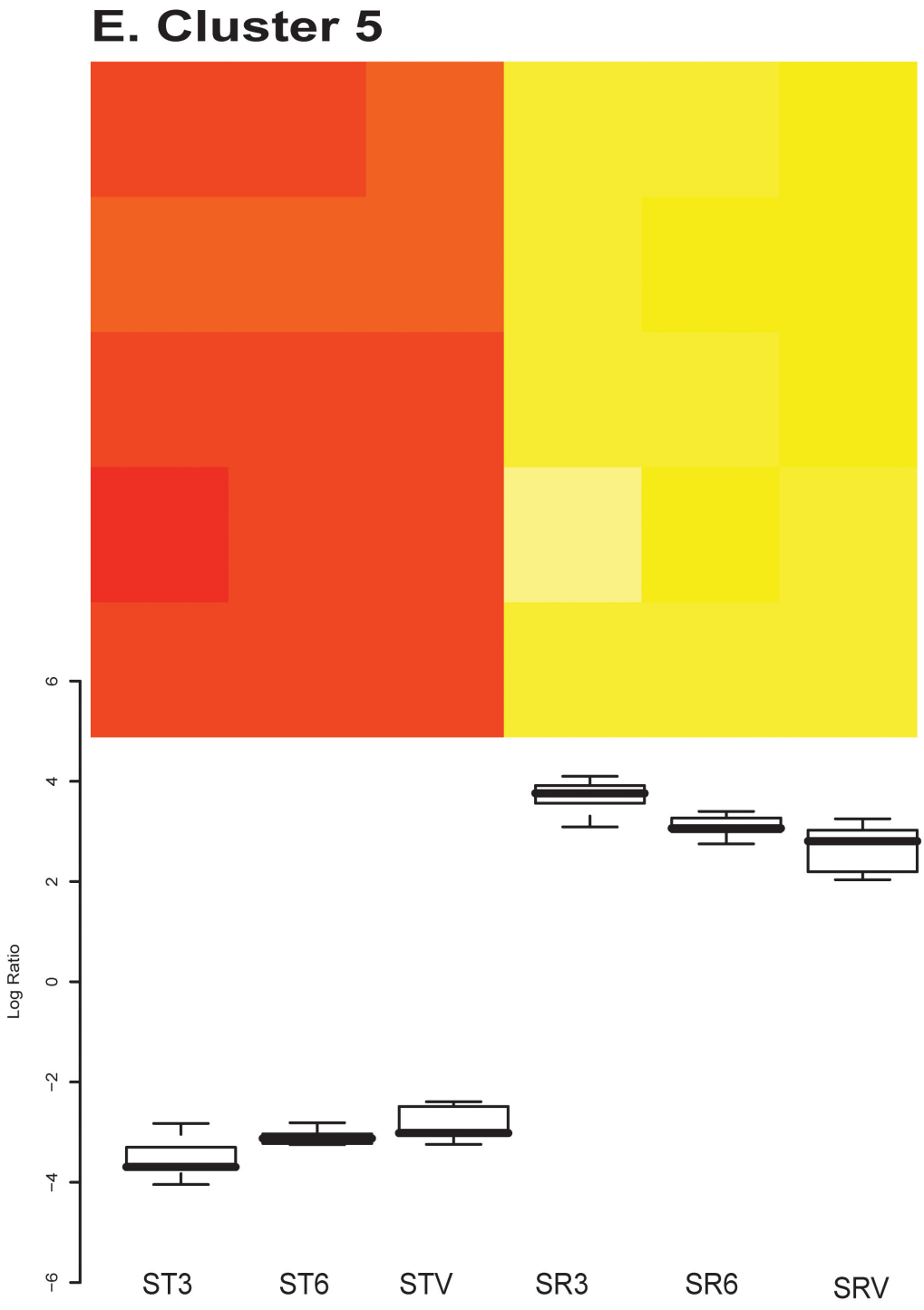
E. Cluster 5
Log Ratio
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0
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ST3
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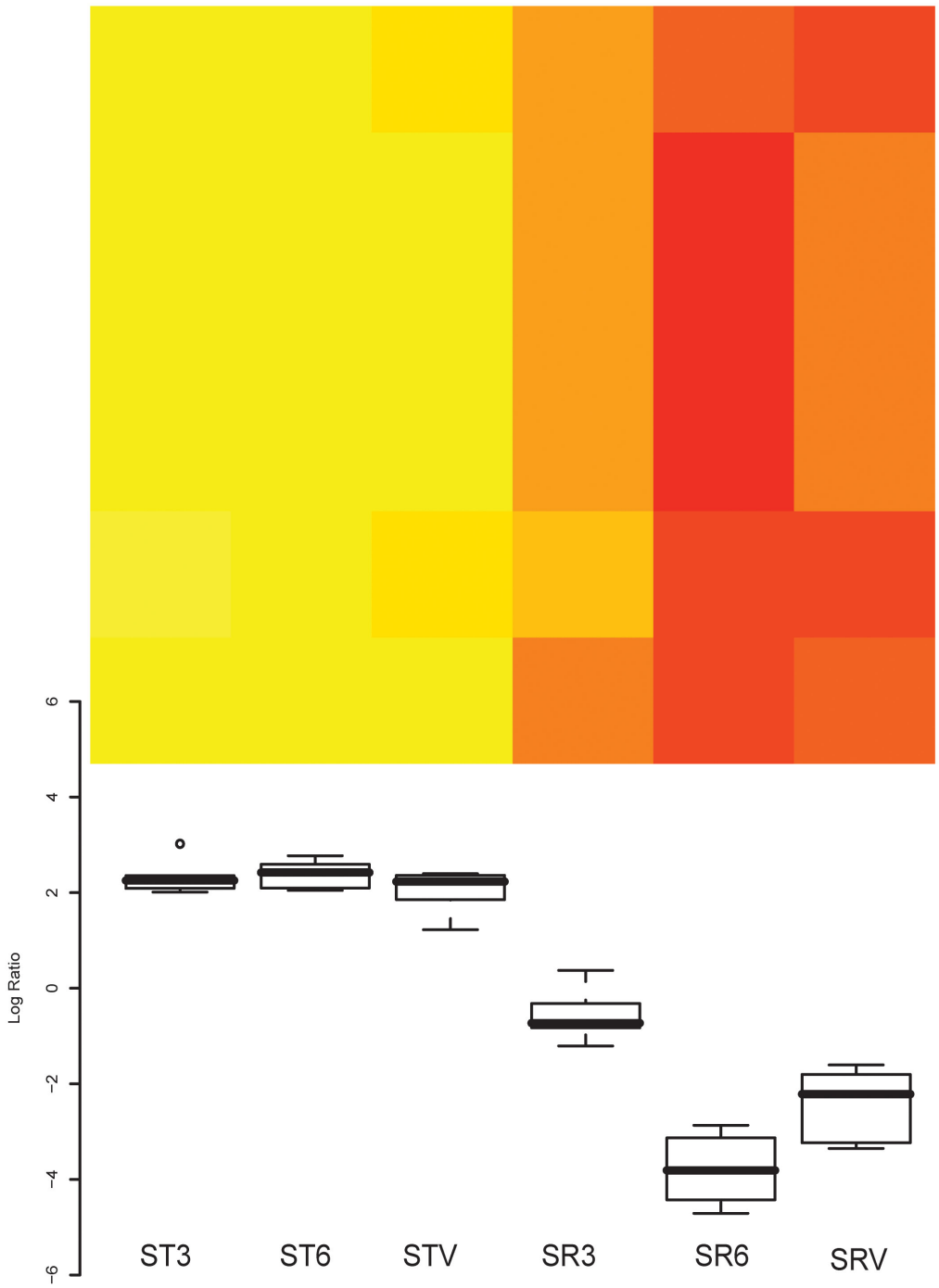
SR6
SRV

F. Cluster 6



Supplementary table 1: Categories of genes highly expressed in the ST and SR

		Genes highly expressed in ST			Genes highly expressed in			
		STV	ST3	ST6	SRV	SR3	SR6	Upregulated Compared to whole body(Flyatlas)
Serine Protease								
	CG1299	2501	3861	4042	334	60	144	Head, crop, H.gut (a,l),
	CG9897	1	14	11	1487	807	5681	ST(m,v)
	CG10469	89	134	93	98	398	214	H.gut, M.tubule (a,l), ST(m)
	CG13318	508	590	459	42	9	16	Head, ST (m,v)
	CG17012	4	15	23	413	36	157	ST (m,v)
	CG17234	73	2	3	737	2	1180	ST (m,v)
	CG17239	12	1	1	552	27	1858	ST (m,v)
	CG18125	137	5	6	1206	59	2754	ST (m,v)
	CG30371	26	10	16	1182	118	2563	Crop, M.tubule (a,l), SG (a,l), ST (m,v)
	CG31681	6	7	5	380	40	1576	ST (m,v)
	CG32277	27	35	34	2063	691	5569	ST (m,v)
	CG32834	20	26	12	2296	1199	6506	ST (m,v)
Female gamete generation/ lipid particle								
	Yp1	7	11	7	81	25	41	None
	Yp2	28	41	14	253	61	115	None
	Yp3	34	67	24	219	107	173	Head
Ion transport								
	blw	113	148	139	60	66	45	Head, Crop, M.gut(a,l), H.gut(a,l), M.tubule, TAG, SG(a,l), ST (m,v)
	CG10413	717	723	798	159	23	29	Crop, M.gut (a,l), H.gut(a,l), M.tubule(a,l), Ovary, Acc, TAG, SG, ST(m,v)
	CG4462	2523	2615	3470	193	16	38	Brain, Head, M.gut(a,l), H.gut(a,l), M.tubule, TAG, SG(a,l),
	CG5687	1516	2352	3002	45	768	160	Brain, Head, TAG
	CG8029	255	323	298	69	96	69	Brain, M.gut(a,l), H.gut(a,l), M.tubule(a,l), SG (a,l),
	Esp	974	1312	1344	78	90	39	H.gut(a,l), ST(m,v)
	l(2)01810	546	735	904	59	56	44	M.gut (a,l), H.gut (a,l), M.tubule, SG,
	Mpcp	81	112	106	54	121	102	Brain, Head, Crop, M.gut(a,l), H.gut(a,l), M.tubule(a,l), Acc, TAG, SG(a,l), ST(m,v)
	sesB	115	128	133	40	89	60	H.gut(a,l), M.tubule, TAG, SG, M.gut(l), ST(m,v)
	vha55	233	325	412	118	126	74	Brain, Head, M.gut, H.gut(a,l), M.tubule (a,l), Ovary, TAG, SG(a,l).
	vha68-2	170	216	216	22	33	19	Brain, Head, M.gut, H.gut(a,l), M.tubule (a,l), Ovary, TAG, SG(a,l).
	VhaPPA1-1	265	345	299	66	168	107	Brain, M.gut (a,l), H.gut (a,l), M. tubule (a,l), TAG, SG (a,l)
Transmembrane receptor								
	CG8444	396	506	418	86	361	327	Brain, Head, Crop, M.gut(a,l), H.gut(a,l), M.tubule (a,l), Fatbody, TAG, SG (a,l)
	dscam	57	68	62	303	2099	1856	Crop, M.gut (a,l), M. tubule, Acc, Fatbody, SG(a,l), ST(m)
Carbohydrate metabolism								
	CG7998	110	155	148	65	163	124	Brain, Crop, M.gut(a,l) , H.gut(a,l), M. Tubule(a), SG(a), ST(m,v)
	tps1	68	127	76	486	296	690	Brain, M. tubule, ST(m,v)
	treh	175	266	314	26	169	49	Brain, Head, M.gut, H.gut, Testis, Acc, TAG, SG(a,l),
Lipid metabolism/transport								
	CG3523	76	107	82	262	54	97	Head, Crop, M.gut (l), ST (v,m)
	RfaBp	97	104	51	343	95	104	Head, Fatbody, ST(m,v)
	Sap-r	85	84	73	85	65	55	Head, Crop, M.gut(a,l), H.gut(a,l), M.tubule(a,l), Acc, SG(a,l), ST(m,v)
Oxoreductase								
	Adh	22	42	26	39	34	33	Head, Crop, M.tubule, Fatbody, SG (a)
	Aldh	66	118	68	91	136	98	Head, Crop, M.tubule, SG(a), ST (m,v)
	ERp60	77	48	47	21	7	7	M.tubule(a,l), Acc, Fatbody, SG(a,l), M.gut(l), H.gut(l), ST(m,v)
	Gapdh2	141	301	186	104	128	88	Head, Crop, M.gut (l), H.gut (a,l), M.tubule, Ovary, TAG, SG (a), ST(m,v)
	Gpdh	202	327	226	218	94	137	Head, M.tubule(a), SG(a), ST(m,v)
Immunity/Defense								
	Def	1216	1100	655	18	17	15	ST (m,v)
	Drs	29	31	31	29	114	57	Fatbody, SG(a), ST(m,v)
	Rel	767	630	796	741	76	175	Head, Crop, M.gut(a,l), H.gut(a,l), M. tubule(a,l), Acc, Fatbody, SG(a), ST(m,v)
	TepIV	212	190	174	55	58	27	Head, Crop, M.gut, H.gut (a,l), M.tubule (l), Fatbody, ST(m,v)
Chemosensory								
	pst	140	133	112	178	85	89	Head, Crop, M.gut(a,l), H.gut(a,l), M.tubule, Acc, Fatbody, SG(a,l), ST(m,v)
	visgun	283	400	342	100	99	80	Brain, Head, Crop, H.gut(a,l), M.tubule(a,l), Acc, Fatbody, TAG, SG(a,l), ST(m,v)
Metal ion binding								
	Aph-4	448	1102	1625	97	1152	274	M.tubule(a,l), Acc
	FER1hch	78	126	104	64	44	48	Head, M.gut(a,l), H.gut(a,l), M.tubule(a,l), Fatbody, TAG, ST(m,v)
	FER2lch	119	165	166	73	67	56	Head, M.gut(a,l), H.gut, M.tubule(a,l), TAG, SG(a,l), ST(m,v)
	MtnA	102	154	137	7	153	76	M.gut, H.gut, M.tubule(a,l), SG,
	Vha13	263	352	402	94	143	101	M.gut(a,l), H.gut(a,l), M.tubule(a,l), TAG, SG(a,l)
	Vha16	139	139	160	17	20	12	M.gut(a,l), H.gut(a,l), M.tubule(a,l), SG(a,l),
	Vha26	136	187	178	32	41	18	Brain, M.gut(a,l), H.gut(a,l), M.tubule(a,l), TAG, SG(a,l),
Muscle contraction/development								
	Mp20	687	1186	1490	44	102	58	Crop, M.gut(a,l), H.gut(a,l),
	TM1	363	782	1205	20	238	100	Acc, SG(l)
	Tm2	559	1192	1596	50	141	64	H.gut(a,l),
	wupa	142	267	430	2	4	2	Crop, H.gut(a,l), Acc
	Mhc	245	459	582	38	72	35	Head, Crop, H.gut(a,l)
	Zeelin1	335	640	1012	62	64	40	Crop, H.gut(a,l)
Cell morphogenesis								
	mfas	1916	3014	3032	186	37	96	Brain, head, crop, H.gut(a,l), testis, Acc, TAG
	moe	406	241	355	115	83	59	Head, crop, M.gut(a,l), H.gut(a,l), M.tubule(a,l), Ovary, Acc, TAG, SG(a,l), ST(m,v)
Receptor binding								
	Idgf4	190	238	206	12	45	34	Head, H.gut(a,l), Fatbody, TAG, ST(m,v)
	Idgf1	582	432	362	356	78	262	Head, Crop, H.gut(a,l), Fatbody, ST(m,v)
Other								
Signal transduction	CG18067	25	3	2	16	50	31	Head, H.gut, Acc, Fatbody, TAG, SG, ST(m,v)
Unknown (Serine protease)	CG18628	997	2706	1414	1	1	1	Testis
sugar binding	CG3244	1096	1137	1056	170	71	92	Head, Crop, H.gut(a,l),ST(m)
Carbon-oxygen lyase	CG5379	355	83	458	6845	2495	11985	Brain, M.tubule, TAG, ST(m,v)
Calcium ion binding	CG6426	2	8	4	748	271	1130	Head, Crop, H.gut(a,l), ST(m,v)
Water transport	CG7777	166	202	153	77	179	86	Ovary, M.tubule (l), TAG, SG(a), H.gut(l), ST(m,v)
neurotransmitter	Chc	192	156	212	256	97	82	Brain, Head, Crop, M.gut(a,l), H.gut(a,l), M.tubule(a,l), Ovary, Acc, Fatbody, TAG, SG(a,l), ST(m,v)
Cysteine proteinase	Cp1	83	151	127	51	122	105	Head, M.gut, H.gut(l), Acc, M.tubule(l), ST(m,v)
protease inhibitor	cys	257	282	310	217	84	135	Head, Crop, H.gut(a,l), M.tubule, Acc, Fatbody, SG(a,l), ST(m,v)
Chaperone binding	Hsc70-4	74	46	80	19	21	28	Crop, H.gut(a,l), M.gut(a,l), M.tubule(a,l),Acc, SG(a,l), ST(m.v)
JH metabolism	jhamt	6320	2401	2095	1555	74	81	Brain, Head
Intracellular signaling cascade	regucalcin	614	1591	1316	88	120	119	Brain, Head, M.tubule(a,l), testis, Acc, Fatbody, SG(l),
metalloprotease	sda	1826	1111	1286	227	52	61	Brain, Head, M.gut(a,l), H.gut(a,l), M.tubule, Acc, TAG, SG(a,l)), ST(m,v)
Transmembrane Transporter	Sec61beta	138	79	70	318	227	232	M.gut(a,l), H.gut(l), M.tubule, Acc, Fatbody, SG(a,l), ST(m,v)
Serine protease inhibitor	Spn5	155	217	213	11	15	9	Head, H.gut(l), ST(m,v)
actin binding	twinstar	144	110	123	74	57	50	Crop, M.gut(a,l), H.gut(a,l), M.tubule(l), Sg(l)

Ranked gene expression of genes in the spermathecae and seminal receptacle. For example, CG1299 is the 2501th most expressed in the virgin spermathecae (STV); it is the 334th most expressed gene in virgin seminal receptacle (SRV). Shading indicates that the gene was among the top 100 expressed genes. Brain; Head(including brain); Crop; M.gut = Midgut; H.gut = Hindgut; M.tubule = Malpighian tubule; Ovary; Testis; Male Accessory Glands = Acc; Fatbody; ST = Spermatheca; Thoracico-abdominal Ganglia = TAG, Salivary Glands = SG; l = larval, a = adult; m = mated; v= virgin; SR = seminal receptacle. V= virgin; 3 = 3 hours post-mating; 6 = six hours post-mating.

Supplementary table 2 : Differential expression data from pairwise comparisons.

Fold change (FC) is given as log2 fold change along with the associated false discovery rate (FDR). Fold change is reported as relative to the first category in the listed comparison, for example in the comparison of STV - SRV, Gpdh is upregulated .0887 fold (log2) in STV compared to SRV. Data which is inconsistant between two probe sets for the same gene is flagged using red font. Inconsistant is defined as : at least two probe sets for the same gene being statistically significantly differentiated and showing opposite directions of gene expression. FC = fold change; FDR = False Discovery Rate; ST = spermatheca; SR = seminal receptacle; V = virgin; 3 = 3 hours post-mating; 6 = 6 hours post-mating.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	**Gene Title**	**Affy ID**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**
Gpdh	5' gene	1616608_a_at	0.0887	0.7110	-0.8703	0.0772	-0.2668	0.2227	0.2403	0.5515	0.4495	0.0379	0.2093	0.2663	-0.3177	0.7707	-0.4874	0.2790	-0.1697	0.7516
CG33057 /// mkg-p	monkey king prote	1622892_s_at	0.0757	0.8101	0.3595	0.1255	0.4645	0.0137	-0.0779	0.9200	-0.2333	0.3322	-0.1553	0.4948	-0.1978	0.8202	0.0640	0.9077	0.2618	0.4838
IM3 /// RpS10b	Immune induced r	1622893_at	1.7724	0.0725	0.2353	0.7853	1.1728	0.0376	0.9557	0.5008	0.9782	0.1908	0.0225	0.9808	0.0670	0.9898	-0.4238	0.7510	-0.4908	0.6969
CG15120	CG15120	1622894_at	1.8690	0.0157	1.1158	0.0606	1.5284	0.0000	0.5250	0.2171	-0.1176	0.6873	-0.6427	0.0125	0.1709	0.9450	-0.8038	0.2679	-0.9748	0.2131
CG32075	CG32075	1622895_at	0.4997	0.0401	0.1760	0.7080	0.4559	0.0806	-0.0230	0.9831	0.7043	0.0122	0.7274	0.0064	-0.3522	0.7823	0.3394	0.5563	0.6915	0.2248
pinta	CG13848	1622896_at	0.5257	0.1446	0.5303	0.2581	0.7831	0.0239	-0.2323	0.7477	-0.8919	0.0121	-0.6595	0.0272	-0.6667	0.7116	-1.0949	0.1397	-0.4282	0.5964
eve	even-skipped	1622897_at	0.2479	0.2356	0.2754	0.1328	0.3386	0.0333	0.0772	0.9130	-0.1654	0.4786	-0.2426	0.2249	0.0484	0.9571	-0.0598	0.8685	-0.1082	0.7071
mRpS10	mitochondrial ribo	1622898_a_at	-0.1057	0.6528	0.2280	0.4558	-0.1338	0.8319	-0.2381	0.6015	0.1472	0.5527	0.3853	0.0645	0.1331	0.9689	0.5102	0.5897	0.3771	0.7068
Rad9	Rad9	1622899_at	-0.0867	0.7145	-0.0150	0.8831	0.0662	0.7710	-0.0878	0.9196	0.0309	0.9342	0.1187	0.6672	-0.1302	0.8202	0.1019	0.7052	0.2321	0.3437
Gr59d	Gustatory recepto	1622900_at	0.0181	0.9485	0.0306	0.7627	0.0216	0.9082	-0.0047	0.9956	-0.1413	0.5286	-0.1366	0.4995	0.0536	0.9400	-0.1107	0.6417	-0.1643	0.4685
CG9993	CG9993	1622901_at	3.0824	0.2479	-2.1806	0.0289	0.0703	0.7050	2.4729	0.0056	2.2246	0.0008	-0.2483	0.5472	0.0943	0.9946	-3.1831	0.3168	-3.2774	0.3271
Pbprp5	Pheromone-bindir	1622902_at	0.1281	0.4950	-0.0439	0.6794	0.0356	0.8713	0.2585	0.3428	0.1957	0.2103	-0.0628	0.7045	-0.0720	0.9457	-0.0239	0.9640	0.0481	0.9125
Tango10	Transport and Gol	1622903_s_at	0.0907	0.7311	0.0136	0.9540	-0.3591	0.0675	-0.0485	0.9420	0.1288	0.5239	0.1774	0.3032	0.1665	0.8235	0.1145	0.7567	-0.0519	0.9031
mfr	misfire	1622904_at	-0.0927	0.6185	-0.1303	0.4953	0.1349	0.4827	0.0689	0.9406	-0.1581	0.5834	-0.2271	0.3517	-0.2536	0.6749	-0.1055	0.7115	0.1481	0.5808
---	---	1622905_at	0.0445	0.7848	0.1240	0.4821	0.0964	0.6165	0.0710	0.9254	0.0143	0.9679	-0.0568	0.8276	-0.0501	0.9422	-0.0664	0.8117	-0.0163	0.9582
---	---	1622906_at	0.1148	0.6498	0.4120	0.4317	0.8159	0.0051	-0.0351	0.9675	-0.0410	0.8943	-0.0058	0.9840	-0.4612	0.7547	0.3843	0.5591	0.8456	0.2009
CG30120 /// CG5189	CG30120 /// CG5	1622907_at	0.0006	0.9983	-0.0341	0.9008	0.0478	0.8734	0.1960	0.7225	-0.1427	0.5997	-0.3386	0.1259	0.0187	0.9914	-0.0571	0.9270	-0.0759	0.8884
CG10962	CG10962	1622908_a_at	1.3367	0.0060	1.8665	0.0027	2.6349	0.0001	0.9924	0.1119	0.2994	0.4383	-0.6930	0.0436	0.3411	0.7848	0.8856	0.1061	0.5444	0.3212
Pi3K21B	dPI 3-kinase	1622909_at	1.1505	0.0272	0.6698	0.1177	-0.0068	0.9858	-0.2128	0.8068	0.6267	0.0727	0.8395	0.0146	0.4780	0.7768	0.0817	0.9451	-0.3963	0.6140
---	---	1622910_at	-0.2166	0.2802	0.1625	0.4947	-0.0512	0.7739	-0.1744	0.5839	-0.2634	0.1038	-0.0890	0.5828	0.1581	0.8594	0.1735	0.6605	0.0154	0.9788
---	---	1622911_at	0.3333	0.0691	0.0462	0.6610	0.0686	0.6722	0.0117	0.9857	-0.0033	0.9883	-0.0151	0.9344	-0.1417	0.7337	-0.3178	0.0869	-0.1760	0.3360
Ucp4A	Ucp4A	1622912_at	-0.3510	0.5285	-0.0910	0.8363	0.3805	0.0397	0.1554	0.6584	-0.2404	0.1528	-0.3957	0.0169	-0.4463	0.8141	-0.0570	0.9648	0.3893	0.6443
PIP5K59B	PIP5K59B	1622913_a_at	0.0345	0.9013	0.0903	0.5797	0.2733	0.1003	0.0742	0.9319	-0.0939	0.7613	-0.1680	0.4979	0.0934	0.8653	0.0159	0.9649	-0.0775	0.7604
---	---	1622914_at	0.1115	0.4818	-0.0236	0.8285	0.0096	0.9660	-0.0842	0.8432	-0.0178	0.9369	0.0663	0.6884	-0.0404	0.9590	-0.1002	0.6949	-0.0598	0.8353
Tsp66A	Tetraspanin 66A	1622915_at	0.1515	0.3802	-0.0077	0.9519	0.1586	0.4139	0.0098	0.9893	0.1013	0.5822	0.0915	0.5898	0.0462	0.9589	0.1128	0.6904	0.0666	0.8352
CG11581	CG11581	1622916_at	0.0914	0.6314	0.2961	0.0409	0.1815	0.4684	-0.0828	0.9252	-0.1130	0.7121	-0.0302	0.9257	-0.0327	0.9774	-0.0482	0.9075	-0.0155	0.9703
CG32308	CG32308	1622917_a_at	0.1720	0.2867	0.0057	0.9635	0.2287	0.2753	-0.0068	0.9942	0.0351	0.8928	0.0419	0.8506	-0.0932	0.8903	0.1016	0.7305	0.1948	0.4592
---	---	1622918_at	0.0389	0.8220	0.1674	0.3450	-0.0968	0.5924	-0.0158	0.9840	-0.1387	0.4701	-0.1229	0.4834	0.0762	0.9174	0.0657	0.8500	-0.0104	0.9805
CG14640	CG14640	1622919_at	0.2897	0.1240	0.1621	0.2537	0.0337	0.8988	0.0848	0.8844	0.1989	0.3241	0.1142	0.5617	0.2005	0.8270	-0.0347	0.9552	-0.2352	0.5592
trp	Cosens-Manning r	1622920_at	0.1686	0.3630	-0.0866	0.6110	0.0722	0.7487	-0.0821	0.8732	0.1725	0.3426	0.2546	0.1099	-0.1250	0.8270	-0.0082	0.9865	0.1168	0.6502
CG32022	CG32022	1622921_at	0.0629	0.8638	0.3491	0.3911	0.2788	0.0707	0.0422	0.9263	0.0079	0.9702	-0.0342	0.8264	0.1076	0.9589	0.3067	0.6336	0.1991	0.7764
CG34139	CG34139	1622922_at	0.1871	0.5706	0.0504	0.5985	0.0825	0.6412	-0.0121	0.9917	-0.1297	0.6332	-0.1176	0.6378	0.0279	0.9862	0.0163	0.9812	-0.0116	0.9848
wrapper	wrapper	1622923_at	0.0493	0.8158	-0.1111	0.3437	0.0546	0.7441	0.1371	0.7885	0.0608	0.8146	-0.0763	0.7354	0.0130	0.9895	-0.0116	0.9739	-0.0246	0.9330
CG18540	CG18540	1622924_at	0.3416	0.2296	0.0219	0.8534	0.2659	0.2840	0.2591	0.5531	0.5444	0.0240	0.2853	0.1561	0.0610	0.9640	0.0920	0.8546	0.0310	0.9517
nej	CREB binding pro	1622925_at	0.8704	0.1205	0.3691	0.6464	-0.1742	0.6408	-0.2844	0.7424	1.1434	0.0093	1.4278	0.0021	0.3624	0.8909	0.4751	0.6656	0.1126	0.9356
CG5776	CG5776	1622926_at	0.9863	0.0420	0.1940	0.5930	0.2122	0.4375	-0.2827	0.5419	0.9024	0.0035	1.1851	0.0006	-0.4148	0.7726	0.0578	0.9531	0.4726	0.4563
CG8642	CG8642	1622927_at	-0.1386	0.3842	-0.1147	0.4922	-0.4072	0.0515	-0.1172	0.7791	-0.0478	0.8277	0.0695	0.7047	-0.0255	0.9742	-0.0500	0.8548	-0.0245	0.9295
CG7940	CG7940	1622928_at	-0.1757	0.3153	-0.0802	0.7603	-0.0455	0.8439	-0.0293	0.9727	0.0371	0.8989	0.0664	0.7794	-0.1600	0.8202	0.1714	0.5747	0.3315	0.2674
CG31222	CG31222	1622929_at	-0.0052	0.9788	-0.0058	0.9655	0.0280	0.8954	-0.2300	0.6046	-0.0141	0.9678	0.2159	0.2862	-0.1105	0.8379	-0.0302	0.9326	0.0804	0.7538
CG1957	CG1957	1622930_a_at	0.3958	0.1153	0.1783	0.2512	0.1162	0.5394	-0.3344	0.4704	0.0736	0.8112	0.4080	0.0700	-0.1748	0.7686	-0.0735	0.8244	0.1014	0.7240
CG5146	CG5146	1622931_at	0.2217	0.5534	0.4074	0.0577	0.4561	0.0324	-0.0077	0.9953	0.0471	0.8959	0.0548	0.8588	-0.0017	0.9995	0.3510	0.3259	0.3527	0.3491
sn	fascin	1622932_s_at	-0.2269	0.8250	-0.3598	0.5040	-0.9040	0.0261	-0.2897	0.8721	0.2876	0.6887	0.5773	0.3109	0.1285	0.9742	0.1875	0.9009	0.0590	0.9691
CG10344	CG10344	1622933_at	-1.4424	0.0018	0.4066	0.3987	0.9426	0.0297	-0.0299	0.9854	-1.3565	0.0036	-1.3267	0.0024	-0.4655	0.7653	0.5799	0.3738	1.0454	0.1433
su(f)	lethal3Des	1622934_at	0.4265	0.1196	0.3831	0.0750	0.5560	0.0177	0.0145	0.9879	-0.1881	0.4172	-0.2027	0.3228	-0.0157	0.9914	-0.0335	0.9476	-0.0178	0.9704
---	---	1622935_at	-0.1564	0.3094	0.0599	0.7558	0.3444	0.0341	0.0554	0.9314	-0.1382	0.4910	-0.1936	0.2612	-0.2010	0.7644	0.0086	0.9885	0.2097	0.4755
---	---	1622936_at	0.1635	0.4089	-0.0151	0.9503	-0.0323	0.8602	0.0002	0.9998	0.2282	0.1898	0.2280	0.1437	-0.0177	0.9862	0.0235	0.9505	0.0413	0.8982
CG13698	CG13698	1622937_at	2.4924	0.0011	1.8133	0.0052	2.1461	0.0000	-0.0290	0.9838	0.5943	0.0774	0.6233	0.0432	-0.3242	0.6955	0.0081	0.9924	0.3323	0.3546
CG17506	CG17506	1622938_at	0.0829	0.7632	0.0461	0.8458	0.0635	0.6883	0.0529	0.9033	0.1120	0.4451	0.0590	0.6905	-0.0475	0.9761	0.0493	0.9353	0.0968	0.8429
emb	Exportin	1622939_at	0.3315	0.1677	0.0408	0.9365	-0.3619	0.1444	-0.1692	0.7121	0.6367	0.0086	0.8059	0.0018	0.1987	0.8972	0.3260	0.5910	0.1273	0.8669
CG12698	CG12698	1622940_at	0.1259	0.5789	-0.0300	0.7559	0.2821	0.0604	0.1008	0.8578	0.0487	0.8531	-0.0521	0.8212	-0.1185	0.8202	-0.1108	0.6380	0.0077	0.9835
CG17765	CG17765	1622941_at	-0.0176	0.9535	-0.0180	0.9502	-0.3550	0.0844	-0.0044	0.9952	-0.0059	0.9793	-0.0015	0.9933	0.3423	0.7131	0.0155	0.9834	-0.3268	0.4085
CG31959	CG31959	1622942_at	0.3172	0.3337	0.0589	0.6286	0.0892	0.7437	-0.1357	0.8449	0.0017	0.9962	0.1374	0.5917	0.0053	0.9974	-0.0900	0.8619	-0.0953	0.8407
CG2371	CG2371	1622943_s_at	0.0217	0.9441	0.4112	0.1069	0.3816	0.0227	-0.0173	0.9863	0.0068	0.9849	0.0241	0.9363	-0.1864	0.7492	0.3309	0.1709	0.5173	0.0741
CG8735	CG8735	1622944_at	-0.1215	0.5467	0.1173	0.3883	-0.1697	0.4279	0.0800	0.8875	0.0016	0.9951	-0.0784	0.6962	0.2504	0.7726	0.3946	0.2731	0.1441	0.7371
Aats-lys	Lysyl-tRNA synthe	1622945_a_at	0.4847	0.0311	0.9101	0.0421	1.0541	0.0009	0.3037	0.2596	0.2099	0.1854	-0.0938	0.5490	0.2343	0.7810	0.6446	0.0878	0.4103	0.2757
CG6908	CG6908	1622946_at	1.8053	0.0086	1.7158	0.0119	1.1610	0.0585	0.2027	0.8822	0.4197	0.3801	0.2170	0.6532	0.6952	0.7576	0.2054	0.8799	-0.4898	0.6318
porin	Voltage-depender	1622947_s_at	-0.5595	0.0056	-0.2589	0.3010	-0.5664	0.0072	-0.1681	0.6471	-0.3123	0.0806	-0.1442	0.3817	0.1321	0.8378	0.0318	0.9404	-0.1003	0.7431
CG17721	CG17721	1622948_at	0.1375	0.4858	-0.2774	0.5352	0.1589	0.3686	0.1676	0.6663	0.3314	0.0760	0.1638	0.3337	-0.2366	0.8400	0.0182	0.9842	0.2548	0.6257
peb	hindsight	1622949_at	2.5202	0.0108	2.6959	0.0198	3.4111	0.0000	0.7374	0.3016	0.3720	0.3870	-0.3654	0.3423	0.1967	0.9457	0.5145	0.5776	0.3178	0.7520
CG10069	CG10069	1622950_a_at	0.9225	0.0081	0.7783	0.0332	0.6919	0.0016	-0.1824	0.6189	0.1051	0.5983	0.2875	0.0798	-0.0827	0.9400	0.0125	0.9846	0.0952	0.8217
---	---	1622951_at	-0.0734	0.6862	0.0018	0.9897	0.1582	0.4334	0.1397	0.7850	-0.0306	0.9175	-0.1703	0.3844	-0.1632	0.7953	-0.0170	0.9689	0.1462	0.6082

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1622952_at	0.6102	0.0157	0.2871	0.7089	-0.0976	0.8256	0.1607	0.7327	1.2134	0.0006	1.0527	0.0006	0.3033	0.9246	0.5330	0.6489	0.2297	0.8745
---	---	1622953_at	0.0590	0.7145	0.0909	0.4771	0.0638	0.6722	0.0822	0.8315	0.0951	0.5479	0.0129	0.9442	-0.0632	0.8999	0.0933	0.6414	0.1564	0.4049
---	---	1622954_at	0.1302	0.4122	0.3784	0.0520	-0.1096	0.5231	-0.2807	0.4568	-0.2886	0.1566	-0.0079	0.9755	0.1716	0.7387	0.0895	0.7205	-0.0821	0.7445
---	---	1622955_at	0.3096	0.1899	0.0320	0.7900	0.1529	0.6515	0.1374	0.7604	0.0631	0.7899	-0.0743	0.7183	0.0826	0.9405	0.0532	0.9202	-0.0294	0.9512
CG31542	CG31542	1622956_at	-0.0393	0.8478	0.1360	0.3735	0.1539	0.4367	-0.0119	0.9932	-0.1625	0.5887	-0.1506	0.5853	-0.0247	0.9816	0.0367	0.9222	0.0613	0.8414
Rep3	Rep3	1622957_s_at	0.0094	0.9728	-0.2408	0.1861	-0.1765	0.3995	0.2477	0.5038	0.4202	0.0388	0.1725	0.3371	-0.0222	0.9885	0.0384	0.9438	0.0606	0.8949
CG30288	CG30288	1622958_at	-0.0186	0.9438	-0.0087	0.9515	0.1615	0.2852	-0.0074	0.9922	-0.0620	0.7411	-0.0547	0.7528	-0.1805	0.7464	-0.1005	0.7053	0.0800	0.7729
CG12928	CG12928	1622959_at	0.7898	0.3911	-2.0723	0.0371	-0.8528	0.0799	0.8506	0.0325	2.0192	0.0001	1.1686	0.0005	-0.2965	0.9467	-0.8527	0.5422	-0.5561	0.7135
decay	caspase-3 protein	1622960_at	-0.2975	0.9030	-0.1594	0.1411	-1.0637	0.0616	-0.3264	0.9466	-0.8431	0.5709	-0.5167	0.7254	0.1936	0.9816	-0.9890	0.6250	-1.1826	0.5507
CG40162	CG40162	1622961_at	-0.0760	0.6217	0.0505	0.6073	0.0199	0.9061	0.0050	0.9943	-0.0057	0.9801	-0.0107	0.9551	0.0617	0.9238	0.2173	0.2924	0.1556	0.4791
CG17221	CG17221	1622962_a_at	-0.1911	0.3869	0.3880	0.0403	0.6467	0.0470	0.0903	0.9060	-0.4948	0.0500	-0.5850	0.0161	-0.0653	0.9246	0.1864	0.4180	0.2517	0.2917
---	---	1622963_at	0.1655	0.2861	0.1059	0.6249	0.0873	0.5997	0.0198	0.9761	0.1434	0.3950	0.1236	0.4219	0.0767	0.9309	-0.0252	0.9583	-0.1019	0.7596
CG34345	CG11053	1622964_at	0.2744	0.4469	-0.0981	0.3223	0.0233	0.9325	0.1338	0.8126	0.1054	0.6789	-0.0285	0.9170	-0.1563	0.8655	-0.3094	0.3941	-0.1532	0.7128
---	---	1622965_s_at	0.1091	0.5530	0.1603	0.3531	0.3640	0.0243	0.1108	0.8817	-0.1155	0.6926	-0.2263	0.3303	-0.1210	0.8145	-0.0563	0.8461	0.0648	0.8039
RpS28a	Ribosomal protein	1622966_at	0.3907	0.1467	0.3687	0.0556	0.1962	0.3821	-0.2621	0.4500	-0.0454	0.8528	0.2168	0.1974	-0.0695	0.9421	-0.1374	0.6762	-0.0679	0.8614
---	---	1622967_at	0.0976	0.4941	0.1351	0.2254	0.3511	0.0325	0.1848	0.5912	0.0760	0.7059	-0.1089	0.5201	0.0424	0.9451	0.0084	0.9808	-0.0340	0.8947
p115	p115	1622968_at	0.4738	0.1631	0.8410	0.0227	1.2958	0.0002	0.3146	0.5839	0.3859	0.1847	0.0713	0.8301	0.0331	0.9848	0.8365	0.0557	0.8033	0.0776
CG30484	CG30484	1622969_at	0.1650	0.3513	0.2786	0.2470	0.1977	0.2524	-0.0736	0.8932	-0.0539	0.8112	0.0196	0.9302	-0.0121	0.9923	-0.0472	0.9174	-0.0351	0.9323
form3	formin 3	1622970_at	-1.5127	0.0090	-3.5897	0.0141	-3.3553	0.0001	0.2373	0.8794	1.1961	0.0324	0.9589	0.0497	0.1380	0.9603	-0.7485	0.3259	-0.8865	0.2738
CG15087	CG15087	1622971_at	-0.7547	0.0363	-0.0825	0.8535	-0.0736	0.7776	-0.1205	0.8405	-0.1837	0.4275	-0.0632	0.8015	-0.0036	0.9990	0.3483	0.4317	0.3518	0.4412
---	---	1622972_at	0.5240	0.0163	0.6020	0.0141	0.5804	0.0065	-0.1329	0.7307	0.0086	0.9743	0.1415	0.3755	-0.1260	0.8049	-0.0535	0.8546	0.0725	0.7705
CG8031 /// DyakCG8031	CG8031	1622973_at	0.9167	0.0041	0.4530	0.1887	0.3952	0.0768	0.0347	0.9653	0.2035	0.3478	0.1688	0.3941	-0.0404	0.9848	-0.2815	0.5847	-0.2411	0.6443
CG4000	CG4000	1622974_at	0.9623	0.1731	0.8276	0.1298	0.9106	0.0070	0.0016	0.9988	-0.9372	0.0073	-0.9389	0.0042	-0.1187	0.9781	-0.9663	0.3358	-0.8476	0.4203
CG13477	CG13477	1622975_at	-0.0075	0.9710	0.1735	0.3285	0.2009	0.4110	0.0373	0.9620	-0.0252	0.9314	-0.0625	0.7882	-0.1075	0.9226	0.1259	0.7807	0.2334	0.5476
mod(mdg4)	Modifier67.2	1622976_at	-0.4396	0.2485	-0.3848	0.3946	-0.3670	0.2784	0.2700	0.7293	-0.1778	0.6508	-0.4478	0.1500	0.2771	0.8461	-0.0469	0.9620	-0.3240	0.6121
CG14635	CG14635	1622977_at	0.0927	0.6687	0.0475	0.8479	0.0468	0.7936	-0.0123	0.9895	0.0397	0.8926	0.0520	0.8340	-0.1262	0.8692	-0.1115	0.7601	0.0147	0.9760
CG6434	CG6434	1622978_at	-0.0847	0.6461	-0.0939	0.4901	-0.0149	0.9387	0.0084	0.9931	0.0236	0.9289	0.0152	0.9492	-0.0287	0.9762	0.0491	0.8830	0.0779	0.7754
CG7188	CG7188	1622979_a_at	-0.0662	0.7169	0.1516	0.4138	0.4227	0.0111	-0.0938	0.7927	0.0368	0.8442	0.1306	0.3236	-0.2993	0.6483	0.2949	0.2488	0.5942	0.0607
Gbp	GTP-binding-prote	1622980_at	0.3412	0.0919	0.0880	0.7239	0.4812	0.0215	0.1875	0.6144	0.3842	0.0452	0.1967	0.2396	-0.1172	0.8814	0.2189	0.4729	0.3360	0.2784
usp	Ultraspiracle	1622981_at	-0.1737	0.6934	0.5393	0.3392	0.2668	0.0783	-0.2761	0.3082	-0.5957	0.0035	-0.3196	0.0320	0.1746	0.9460	0.2077	0.8446	0.0330	0.9794
p38b	stress-activated p	1622982_at	-0.3559	0.1441	0.7142	0.0157	0.6978	0.0030	-0.0651	0.8942	-0.7848	0.0012	-0.7197	0.0010	-0.1122	0.8960	0.1770	0.6049	0.2893	0.3803
Nup154	tulipano	1622983_a_at	-0.3019	0.2265	-0.4622	0.2291	0.2589	0.3126	0.1000	0.8220	0.3213	0.0670	0.2213	0.1521	-0.5421	0.6927	0.1511	0.8431	0.6932	0.2338
Sod2	mitochodnrial SOI	1622984_at	0.0824	0.8042	0.0906	0.6248	0.1669	0.3566	0.0005	0.9994	-0.6735	0.0072	-0.6740	0.0042	-0.0123	0.9946	-0.6477	0.1248	-0.6354	0.1548
CG14325	CG14325	1622985_at	0.1092	0.7606	-0.0275	0.8269	0.0206	0.9053	0.1553	0.6854	0.0991	0.6142	-0.0561	0.7765	-0.0148	0.9916	-0.1686	0.6310	-0.1538	0.6621
---	---	1622986_at	-0.0106	0.9653	0.2396	0.3152	0.1841	0.4460	-0.0424	0.9639	-0.0911	0.7598	-0.0487	0.8670	0.1524	0.8202	0.1750	0.5411	0.0226	0.9533
CG15325	CG15325	1622987_at	0.0057	0.9853	0.1292	0.4229	0.0119	0.9498	-0.0273	0.9759	0.1460	0.5423	0.1732	0.4046	0.0634	0.9309	0.1654	0.4999	0.1020	0.7027
---	---	1622988_at	-0.0967	0.7110	0.2291	0.2762	0.1703	0.3702	-0.0952	0.8942	-0.0479	0.8806	0.0474	0.8645	0.1764	0.8461	0.2657	0.4895	0.0893	0.8549
CG5611	CG5611	1622989_at	-0.2988	0.6539	0.0465	0.9591	0.5692	0.0190	-0.0715	0.9436	-0.4321	0.1335	-0.3606	0.1615	-0.7519	0.7387	-0.3581	0.7518	0.3938	0.7152
CG5381	CG5381	1622990_at	-0.1874	0.3060	-0.4252	0.1944	-0.5729	0.1198	-0.0274	0.9803	0.2159	0.4409	0.2434	0.3237	0.2077	0.8875	0.0504	0.9551	-0.1573	0.8184
Ret	Ret oncogene	1622991_s_at	3.6183	0.0024	2.5542	0.0586	3.5908	0.0002	0.1464	0.8192	0.1333	0.6342	-0.0131	0.9674	-0.3431	0.9088	-0.5365	0.6411	-0.1934	0.8943
twe	twine	1622992_at	-0.6456	0.0684	-0.8289	0.0679	-1.1575	0.0098	-0.2360	0.7431	0.3319	0.2989	0.5678	0.0498	-0.2973	0.8510	0.0243	0.9846	0.3216	0.6476
---	---	1622993_at	0.1397	0.4934	0.1032	0.4728	0.0408	0.8155	-0.0072	0.9922	-0.0340	0.8626	-0.0267	0.8800	0.0275	0.9816	0.0180	0.9682	-0.0095	0.9838
---	---	1622994_at	-0.0660	0.7731	-0.2847	0.1648	-0.4155	0.0131	0.2350	0.6504	0.2810	0.2631	0.0460	0.8769	0.1188	0.8049	0.0292	0.9269	-0.0896	0.6867
mRpS17	mitochondrial ribo	1622995_at	0.1799	0.3864	0.0514	0.8929	0.2857	0.1910	0.2052	0.6490	0.1355	0.5641	-0.0697	0.7722	-0.0336	0.9862	0.1093	0.8550	0.1430	0.7815
RN-tre	tre oncogene-relat	1622996_at	0.1846	0.4173	-0.4081	0.2486	-0.6369	0.0209	-0.1320	0.8645	0.4952	0.0740	0.6272	0.0188	0.1088	0.9246	-0.0871	0.8754	-0.1959	0.6389
CG34367	CG13141	1622997_at	-0.1433	0.6257	0.0863	0.4728	0.0006	0.9982	-0.0826	0.9102	-0.0458	0.8834	0.0368	0.8953	0.0305	0.9841	0.2224	0.5374	0.1919	0.6054
pfk	piefke	1622998_a_at	-0.2422	0.1057	-0.0702	0.8240	-0.3505	0.0573	-0.0147	0.9857	0.2741	0.1433	0.2889	0.0862	0.1747	0.7770	0.2218	0.4032	0.0471	0.8984
1-Sep	septin	1622999_at	0.2868	0.2258	-0.0258	0.9010	-0.3949	0.0676	-0.0660	0.9232	0.5024	0.0232	0.5683	0.0084	0.1645	0.8450	0.0319	0.9541	-0.1325	0.7398
CG33093	CG33093	1623000_at	0.3313	0.3853	0.0564	0.6416	0.1158	0.4931	0.0247	0.9819	-0.1519	0.5912	-0.1766	0.4752	-0.0782	0.9589	-0.3277	0.4495	-0.2495	0.5886
CG31723	CG31723	1623001_at	0.0302	0.8656	-0.0901	0.5545	0.0393	0.8961	0.0598	0.9558	0.2788	0.3430	0.2190	0.4179	-0.2054	0.7230	0.1601	0.5404	0.3655	0.1680
CG31404	CG31404	1623002_at	0.3813	0.0431	0.1955	0.4335	0.2598	0.2742	0.2088	0.6800	0.1266	0.6305	-0.0821	0.7488	0.1515	0.8098	-0.1029	0.7364	-0.2543	0.3458
---	---	1623003_at	-0.0102	0.9710	-0.0922	0.5297	0.3423	0.0597	-0.0970	0.8908	-0.0179	0.9600	0.0791	0.7614	-0.1690	0.7230	0.1439	0.4976	0.3130	0.1552
CG18449	CG18449	1623004_a_at	0.1324	0.4290	0.0014	0.9919	-0.1210	0.5211	-0.0109	0.9901	0.1891	0.3433	0.2000	0.2571	0.0670	0.9095	0.0601	0.8291	-0.0069	0.9846
---	---	1623005_at	-0.0335	0.8367	0.0109	0.9173	0.0532	0.7822	0.0418	0.9584	0.0522	0.8484	0.0104	0.9688	0.0311	0.9588	0.0514	0.8127	0.0203	0.9313
SerT	cocaine-sensitive	1623006_at	-2.1317	0.0012	-0.0326	0.8651	-1.2688	0.0013	-1.0028	0.0316	-2.5862	0.0001	-1.5834	0.0003	0.2210	0.8012	-0.4367	0.2168	-0.6577	0.1080
DmirCG11940 /// jog	CG11940	1623007_a_at	0.2735	0.1183	0.1175	0.3725	-0.0747	0.7587	0.1204	0.8046	0.1994	0.3092	0.0790	0.6992	0.0978	0.8889	0.0157	0.9719	-0.0821	0.7964
CG5565 /// DsimCG5565	CG5565	1623008_at	0.0381	0.8811	-0.1310	0.3583	-0.1967	0.4606	0.0887	0.8645	0.2138	0.2481	0.1250	0.4772	-0.0587	0.9467	0.0026	0.9973	0.0612	0.8614
cyp33	cyclophilin-33	1623009_at	0.1523	0.3052	0.0012	0.9993	0.0868	0.6318	0.0344	0.9502	0.3823	0.0206	0.3478	0.0191	-0.1067	0.9300	0.0357	0.9557	0.1424	0.7527
Or30a	Odorant receptor	1623010_at	0.1109	0.5484	0.1954	0.4409	0.2247	0.2257	0.0000	0.9999	0.0881	0.7224	0.0881	0.6940	-0.0518	0.9515	0.0383	0.9237	0.0901	0.7608
CG34372	CG13552	1623011_at	0.1701	0.5012	-0.1093	0.3160	0.0204	0.9505	0.1463	0.7819	0.1819	0.4216	0.0356	0.8919	-0.2084	0.7953	-0.1176	0.7763	0.0908	0.8307
mRpL55	tom thumb	1623012_at	-0.6221	0.0071	0.2017	0.5846	-0.0752	0.7294	0.0912	0.8537	-0.2558	0.1504	-0.3471	0.0367	0.2905	0.7758	0.4647	0.2752	0.1743	0.7298
CG3907 /// DyakCG3907	CG3907	1623013_s_at	-0.4441	0.0631	-0.3347	0.0674	-0.3673	0.1703	0.0988	0.8802	-0.0771	0.7761	-0.1759	0.3995	0.1049	0.9309	0.0033	0.9979	-0.1016	0.8375
CG18537	CG18537	1623014_at	-0.3218	0.1601	-1.6532	0.0176	-1.8611	0.0013	-0.0908	0.9554	1.8257	0.0022	1.9166	0.0011	0.0358	0.9816	0.1192	0.7985	0.0834	0.8637
CG7671	CG7671	1623015_at	0.2474	0.1396	0.1567	0.7479	0.2662	0.2135	-0.0082	0.9937	0.3943	0.0616	0.4025	0.0370	-0.3214	0.7673	0.0991	0.8807	0.4205	0.3728
CG1299	CG1299	1623016_at	-3.0375	0.0020	-5.1150	0.0019	-4.8012	0.0000	0.5026	0.4631	1.0851	0.0105	0.5825	0.0804	0.1135	0.9683	-0.8336	0.2517	-0.9471	0.2247
CG7149	CG7149	1623017_at	0.3950	0.3539	0.3142	0.3697	0.4326	0.0909	-0.0179	0.9937	0.2476	0.6336	0.2655	0.5644	-0.1623	0.9061	0.2725	0.6044	0.4348	0.3880
CG12780	CG12780	1623018_at	-0.4376	0.8243	0.1290	0.4858	-0.2871	0.2396	-0.5246	0.9083	-1.4254	0.3356	-0.9008	0.5250	-0.0629	0.9914	-0.7602	0.5800	-0.6973	0.6170
Unc-89	PROJECTIN-like	1623019_a_at	-2.7544	0.0033	-3.1291	0.0048	-3.3917	0.0001	0.0220	0.9860	-0.2889	0.3429	-0.3109	0.2475	0.2621	0.9246	-0.5962	0.5374	-0.8583	0.3678
CG31937	CG31937	1623020_at	-0.0828	0.6982	0.8563	0.0535	1.5476	0.0003	0.1205	0.8385	-1.0379	0.0014	-1.1584	0.0005	-0.4760	0.6557	-0.2165	0.6427	0.2595	0.5699

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Prosbeta3	20s proteasome	1623021_at	0.0894	0.7977	0.0941	0.6625	0.3732	0.0420	0.1198	0.7135	0.0736	0.6616	-0.0462	0.7802	-0.1660	0.8999	-0.0039	0.9980	0.1621	0.7767
CG15820	CG15820	1623022_at	0.0983	0.7141	-0.2088	0.4735	-0.3708	0.2497	-0.1133	0.8815	0.4016	0.1185	0.5148	0.0331	-0.1480	0.9449	0.0052	0.9980	0.1532	0.8559
Slip1	Slip1	1623023_at	-0.1820	0.3997	-0.1988	0.6072	-0.5825	0.0214	-0.2419	0.5128	0.1574	0.4373	0.3993	0.0299	0.0182	0.9916	0.0642	0.9174	0.0460	0.9344
---	---	1623024_at	0.1734	0.6161	0.0307	0.7923	-0.0009	0.9965	-0.0440	0.9598	-0.1001	0.7050	-0.0561	0.8296	0.0241	0.9816	0.0252	0.9444	0.0012	0.9979
CG3457	CG3457	1623025_at	0.2342	0.3355	0.1228	0.5175	0.0230	0.9180	-0.1179	0.8721	-0.0381	0.9149	0.0798	0.7780	0.0382	0.9816	-0.0275	0.9648	-0.0657	0.8998
CG7573	CG7573	1623026_a_at	0.0316	0.8803	0.0887	0.4065	0.3746	0.1165	0.0125	0.9909	-0.2037	0.3971	-0.2162	0.3103	-0.1672	0.7893	0.0335	0.9377	0.2008	0.4659
CG6271 /// CG6277	CG6271 /// CG62	1623027_s_at	0.0738	0.8376	0.2925	0.2688	0.4013	0.0159	0.0685	0.9345	-0.2151	0.3907	-0.2836	0.1937	0.0528	0.9653	0.2371	0.4753	0.1843	0.5995
CG14872	CG14872	1623028_at	1.9992	0.0042	1.6122	0.0995	1.9864	0.0003	0.6377	0.3248	-0.0982	0.8394	-0.7359	0.0357	0.2450	0.9171	-0.5680	0.4976	-0.8130	0.3299
CG31535	CG31535	1623029_at	0.1829	0.3536	-0.0213	0.9061	0.4363	0.0928	0.1936	0.4962	0.1132	0.4729	-0.0804	0.5946	-0.2717	0.7307	-0.0365	0.9470	0.2351	0.5033
CG11726	CG11726	1623030_at	0.2015	0.3622	0.0891	0.4399	-0.1023	0.6138	-0.0273	0.9715	-0.0144	0.9604	0.0129	0.9582	0.0539	0.9514	-0.0310	0.9411	-0.0850	0.7849
RecQ4	RecQ4	1623031_a_at	0.1164	0.6017	-0.2234	0.6813	-0.5305	0.2636	-0.5705	0.1861	0.2356	0.3771	0.8061	0.0049	-0.1887	0.9412	-0.0507	0.9690	0.1380	0.8998
CG2336	CG2336	1623032_at	0.1659	0.3854	-0.0074	0.9511	0.0014	0.9946	0.0658	0.9154	0.0210	0.9401	-0.0448	0.8414	0.0219	0.9884	0.0586	0.9068	0.0367	0.9360
---	---	1623033_at	0.0340	0.8691	0.0966	0.5120	0.0661	0.7858	-0.0821	0.8512	-0.2673	0.0939	-0.1852	0.1930	-0.0286	0.9831	-0.0253	0.9617	0.0032	0.9953
CG7026	CG7026	1623034_at	-0.0786	0.5805	0.0647	0.5926	0.0738	0.7667	0.0522	0.9297	-0.0953	0.6199	-0.1475	0.3545	-0.2256	0.8013	-0.1102	0.8215	0.1153	0.7984
NaPi-T	Na[+]-dependent i	1623035_at	-0.2615	0.4997	-0.1437	0.5930	0.0155	0.9369	0.2110	0.8426	-0.2564	0.5459	-0.4674	0.1841	0.0900	0.9449	-0.2867	0.4753	-0.3767	0.3543
CG10249	CG10249	1623036_a_at	0.5236	0.3978	-0.0625	0.9189	0.0237	0.9628	0.3638	0.7121	0.6770	0.1293	0.3132	0.4603	0.4258	0.8553	0.2417	0.8546	-0.1840	0.8889
CG5594	CG5594	1623037_a_at	0.0676	0.7997	-0.8707	0.0386	-0.4858	0.0207	0.2491	0.6506	0.7432	0.0127	0.4942	0.0434	-0.1787	0.8608	-0.3076	0.4531	-0.1289	0.7933
CG31448	CG31448	1623038_at	0.2621	0.1648	-0.1394	0.5244	0.0164	0.9531	0.0327	0.9748	0.2951	0.2449	0.2625	0.2484	0.0128	0.9914	-0.0422	0.9157	-0.0550	0.8739
CG3409	CG3409	1623039_at	0.9076	0.0729	1.1770	0.0212	0.9090	0.0099	-0.1488	0.7556	0.3681	0.0762	0.5169	0.0127	0.2104	0.9138	0.6915	0.2964	0.4811	0.4942
CG4449	CG4449	1623040_at	-0.6565	0.0916	-0.4996	0.0144	-0.2867	0.0929	-0.3918	0.5133	-0.0119	0.9804	0.3800	0.1787	-0.5587	0.3166	0.0965	0.7943	0.6552	0.0625
CG13745	CG13745	1623041_at	-0.1424	0.6341	0.0984	0.4756	-0.0440	0.8372	-0.1951	0.7140	-0.1015	0.7198	0.0936	0.7172	0.0023	0.9987	0.1218	0.6666	0.1194	0.6683
---	---	1623042_at	0.1486	0.3380	0.2574	0.0802	0.5507	0.0391	0.1293	0.7647	-0.1138	0.5686	-0.2431	0.1396	-0.2478	0.7707	0.0735	0.8903	0.3213	0.3878
CG7852	CG7852	1623043_s_at	-0.1505	0.6080	-0.0365	0.9610	-0.4826	0.0494	-0.2265	0.6482	-0.0300	0.9293	0.1965	0.3765	0.1152	0.9616	0.0332	0.9782	-0.0820	0.9300
Cp15	s15 chorion gene	1623044_at	-0.1732	0.7840	0.1096	0.4442	-0.2236	0.2118	-0.2708	0.7271	-0.6657	0.0594	-0.3949	0.2029	0.0611	0.9677	-0.1474	0.7567	-0.2085	0.6282
Ip259	Intronic Protein 25	1623045_at	0.1452	0.4173	0.6056	0.0126	0.9506	0.0012	0.1730	0.5744	-0.4205	0.0159	-0.5935	0.0021	-0.1093	0.9031	0.0020	0.9989	0.1113	0.7728
CG32727	CG32727	1623046_at	0.2128	0.2025	-0.1054	0.6166	0.0590	0.7811	-0.0403	0.9494	0.0492	0.8248	0.0895	0.6183	-0.1428	0.7956	-0.2695	0.2288	-0.1267	0.6129
---	---	1623047_at	0.9820	0.0464	0.6680	0.2558	-0.2817	0.2369	-0.1305	0.8816	0.4895	0.1013	0.6200	0.0281	0.7433	0.6258	0.3839	0.5635	-0.3593	0.5984
Lcch3	GABA receptor be	1623048_a_at	-0.4202	0.0587	-0.4330	0.1386	-0.5733	0.2236	-0.3333	0.7833	0.2079	0.7215	0.5412	0.2271	-0.1212	0.8725	-0.0076	0.9911	0.1136	0.7417
CG9731	CG9731	1623049_at	0.5734	0.0648	-0.1408	0.8024	-0.0579	0.8189	0.2022	0.5423	0.5510	0.0072	0.3487	0.0314	0.0630	0.9816	-0.0554	0.9587	-0.1183	0.8932
HLH54F	HLH54F	1623050_at	-0.0018	0.9964	0.0268	0.8057	-0.2494	0.2891	-0.1090	0.8869	0.0299	0.9350	0.1389	0.5921	-0.0186	0.9862	0.0045	0.9935	0.0232	0.9449
CG18125	CG18125	1623051_at	2.2226	0.0161	0.7504	0.0062	4.7722	0.0000	3.9713	0.0091	2.6499	0.0042	-1.3214	0.0503	0.0678	0.9101	1.3641	0.0010	1.2963	0.0023
---	---	1623052_at	-0.0040	0.9883	0.0658	0.6009	0.0283	0.8819	-0.1278	0.8190	-0.2501	0.2548	-0.1223	0.5744	0.1174	0.8122	-0.0258	0.9382	-0.1432	0.5007
CG5174	CG5174	1623053_a_at	0.6344	0.0158	0.9636	0.0077	0.9428	0.0006	0.3105	0.3909	-0.1843	0.3771	-0.4948	0.0147	0.3959	0.6660	0.2403	0.5321	-0.1556	0.7087
---	---	1623054_at	0.0121	0.9411	-0.0211	0.9441	0.1519	0.4848	0.1203	0.8053	0.1097	0.6083	-0.0107	0.9657	-0.0370	0.9653	-0.0353	0.9204	0.0016	0.9972
CG9254	CG9254	1623055_at	0.1540	0.5825	-0.0613	0.6003	-0.0911	0.6434	-0.2207	0.4752	-0.0522	0.7980	0.1685	0.2599	-0.0740	0.9142	-0.1376	0.5897	-0.0636	0.8348
---	---	1623056_at	0.2957	0.1681	0.2676	0.2202	0.3594	0.0898	-0.1037	0.8830	-0.0698	0.8145	0.0339	0.9074	0.0367	0.9701	-0.0704	0.8338	-0.1072	0.7030
---	---	1623057_at	0.3407	0.0738	0.0000	1.0000	0.4255	0.0623	-0.0066	0.9934	0.0640	0.7210	0.0706	0.6578	-0.2006	0.7822	-0.1281	0.7215	0.0725	0.8602
CG32676	CG32676	1623058_at	-0.7528	0.0948	-0.1595	0.8664	-0.5219	0.0996	0.0454	0.9457	-0.4753	0.0201	-0.5206	0.0083	0.1674	0.9657	0.0722	0.9665	-0.0953	0.9486
---	---	1623059_at	0.0177	0.9317	0.0424	0.6454	0.0368	0.8633	-0.0210	0.9777	-0.1066	0.6117	-0.0856	0.6648	0.0262	0.9742	0.0310	0.9225	0.0048	0.9881
Cpr12A	CG15757	1623060_at	-0.0787	0.7401	0.0573	0.5764	0.1297	0.4113	0.0585	0.9307	0.0946	0.6729	0.0361	0.8773	0.1222	0.8202	0.1594	0.4807	0.0372	0.9031
CG10561	transcription unit 3	1623061_at	-0.6465	0.0051	-0.2773	0.1819	-0.2853	0.0983	-0.0365	0.9518	-0.2751	0.1010	-0.2386	0.1107	0.2044	0.6955	0.2047	0.3417	0.0003	0.9992
CG13535	CG13535	1623062_at	0.0940	0.7487	0.2014	0.1226	0.1360	0.4102	-0.0412	0.9384	-0.0914	0.5912	-0.0502	0.7712	-0.0539	0.9610	0.0762	0.8571	0.1301	0.7060
dpr20	dpr20	1623063_at	0.0972	0.6708	-0.1684	0.4183	-0.1587	0.4219	-0.1388	0.7138	-0.0807	0.6821	0.0581	0.7581	-0.1608	0.8215	-0.2970	0.3010	-0.1362	0.6731
---	---	1623064_at	0.2282	0.3047	-0.1384	0.4350	0.1681	0.4289	0.4054	0.3501	0.2702	0.2806	-0.1352	0.5870	0.0636	0.9342	-0.2185	0.3709	-0.2822	0.2731
CG13692	CG13692	1623065_at	0.3799	0.1059	-0.0050	0.9785	-0.0648	0.7465	0.2417	0.5330	0.2064	0.3124	-0.0354	0.8825	0.0220	0.9831	-0.1240	0.6414	-0.1461	0.5763
Irk3	Inwardly rectifying	1623066_at	-1.0299	0.1692	0.2158	0.1046	-0.2977	0.1369	-0.3604	0.8196	-1.9046	0.0084	-1.5443	0.0131	0.1276	0.9081	-0.4994	0.1894	-0.6270	0.1390
---	---	1623067_at	-0.0032	0.9883	0.0642	0.6776	-0.1919	0.2937	-0.0277	0.9753	-0.0631	0.8150	-0.0354	0.8921	0.0796	0.9030	0.0455	0.8990	-0.0340	0.9176
Cyp4e3	Cytochrome P450	1623068_at	0.7628	0.3844	0.0564	0.7653	0.2742	0.1287	-0.0334	0.9937	-0.9017	0.2670	-0.8683	0.2306	-0.0956	0.9628	-1.5514	0.0285	-1.4558	0.0441
CG17544	CG17544	1623069_s_at	-0.2005	0.5969	-0.5354	0.0281	0.1307	0.6064	-0.0864	0.8987	-0.2727	0.2185	-0.1864	0.3625	-0.8504	0.3738	-0.6574	0.1969	0.1930	0.7504
Shaw	Shaker cognate w	1623070_a_at	0.2037	0.3778	0.0751	0.7479	0.0952	0.6448	-0.0622	0.9502	-0.0800	0.8147	-0.0178	0.9593	0.0109	0.9952	-0.0721	0.9205	-0.0831	0.8959
---	---	1623071_at	-0.2208	0.1937	0.0690	0.5196	0.2698	0.1333	0.0803	0.8994	-0.1505	0.4884	-0.2307	0.2131	-0.0024	0.9977	0.0872	0.6713	0.0895	0.6567
CG7709	CG7709	1623072_at	-0.1085	0.5539	0.3027	0.2222	0.2332	0.2108	-0.1388	0.8676	-0.2976	0.3176	-0.1588	0.5898	0.0518	0.9460	0.2020	0.3737	0.1502	0.5378
ctp	dynein	1623073_s_at	-0.8393	0.0027	-0.4791	0.1597	-0.3337	0.0795	0.1206	0.7611	0.0361	0.8732	-0.0845	0.6271	0.0840	0.9409	0.4740	0.1649	0.3900	0.2830
---	---	1623074_at	0.3505	0.2593	-0.0118	0.9673	0.0274	0.8923	-0.0491	0.9266	0.1213	0.4680	0.1703	0.2343	-0.0977	0.9429	-0.1782	0.7068	-0.0805	0.8890
---	---	1623075_at	0.1056	0.4875	0.0287	0.8432	-0.0678	0.7471	0.1114	0.7508	0.0729	0.6747	-0.0385	0.8241	0.1078	0.8743	0.0412	0.9216	-0.0666	0.8461
CG32803	CG32803	1623076_at	0.0306	0.9375	-0.4112	0.1492	-0.4552	0.0247	-0.2357	0.6615	0.0164	0.9676	0.2521	0.2799	-0.1980	0.8236	-0.4136	0.2477	-0.2156	0.5856
---	---	1623077_at	0.1298	0.5795	0.0922	0.5306	0.1069	0.5573	-0.0303	0.9653	0.1123	0.5750	0.1425	0.4099	-0.0895	0.9078	0.1054	0.7439	0.1950	0.4931
---	---	1623078_at	0.1075	0.6212	0.0060	0.9651	0.1838	0.2393	0.1661	0.6086	0.0238	0.9168	-0.1423	0.3400	0.0018	0.9992	-0.1829	0.4943	-0.1847	0.4931
---	---	1623079_at	-0.1370	0.4444	-0.0122	0.9174	0.1360	0.3890	0.0517	0.9311	-0.0701	0.7335	-0.1218	0.4637	-0.0004	0.9998	0.0878	0.7183	0.0882	0.7134
CG18410	CG18410	1623080_at	-1.6038	0.0039	-1.0614	0.0637	-1.3755	0.0007	0.0552	0.9602	-0.2344	0.4366	-0.2896	0.2693	0.4191	0.7506	0.3153	0.6012	-0.1038	0.8934
Jhebp29	CG3776	1623081_at	0.1268	0.6090	0.6454	0.0461	0.9112	0.0015	0.0419	0.9598	-0.8362	0.0031	-0.8781	0.0015	-0.2180	0.8076	-0.2074	0.6021	0.0106	0.9856
---	---	1623082_s_at	0.2127	0.2627	-0.0856	0.4839	0.0492	0.7704	0.0183	0.9761	0.0673	0.6991	0.0490	0.7686	0.0107	0.9926	-0.2018	0.4452	-0.2125	0.4325
CG1698	CG1698	1623083_at	-2.3009	0.0010	-2.4352	0.0083	-2.7383	0.0006	0.0222	0.9869	-0.1527	0.6814	-0.1748	0.5922	0.2947	0.8884	-0.3510	0.6937	-0.6456	0.4258
sgg	Glycogen Synthas	1623084_at	-0.1913	0.5139	0.0522	0.7842	0.0691	0.7125	0.1546	0.6698	0.0384	0.8656	-0.1161	0.4789	0.3681	0.7694	0.4846	0.3522	0.1166	0.8699
CG12637	CG12637	1623085_at	0.2479	0.2236	0.0367	0.8266	0.1721	0.3060	0.1103	0.8409	0.1661	0.4348	0.0558	0.8087	-0.1970	0.7492	-0.1017	0.7424	0.0954	0.7527
Gnf1	germline transcrip	1623086_at	-0.5037	0.0366	0.0177	0.8803	-0.1269	0.4303	-0.1533	0.7857	-0.3590	0.1209	-0.2057	0.3332	-0.0462	0.9523	0.1892	0.4101	0.2354	0.3205
Gef64C	Guanine nucleotid	1623087_at	-4.0579	0.0011	-4.1597	0.0115	-4.1557	0.0001	0.7542	0.6065	0.8861	0.2278	0.1319	0.8800	0.6311	0.7726	0.8396	0.3684	0.2085	0.8693
CG10855	CG10855	1623088_at	-0.3019	0.0706	-0.1392	0.5411	0.1856	0.3949	0.0141	0.9857	-0.0424	0.8595	-0.0565	0.7812	-0.1370	0.8882	0.0479	0.9350	0.1849	0.6389
Mst84Db	Male-specific RNA	1623089_at	-0.0425	0.8506	0.0881	0.6453	-0.0364	0.8656	-0.1416	0.7511	-0.2172	0.2650	-0.0755	0.7136	-0.0294	0.9742	-0.0723	0.8006	-0.0429	0.8896

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG30390	CG30390	1623090_at	-0.1668	0.4682	-0.2397	0.3913	-0.5667	0.0058	-0.2053	0.5234	0.0930	0.6184	0.2983	0.0524	-0.0349	0.9809	-0.0546	0.9096	-0.0197	0.9666
CG12644	CG12644	1623091_at	0.1938	0.2453	0.0054	0.9629	0.2345	0.1826	0.0337	0.9688	0.0072	0.9826	-0.0265	0.9231	-0.0315	0.9653	-0.0601	0.8126	-0.0286	0.9161
Pcp	Gart intron gene	1623092_at	-3.8768	0.0054	-3.9699	0.0012	-4.4718	0.0000	-0.7364	0.3890	-1.3697	0.0120	-0.6333	0.1409	-0.3145	0.9342	-1.6154	0.1747	-1.3009	0.3011
CG13188	CG13188	1623093_at	0.0662	0.6703	0.1983	0.3740	0.0421	0.8484	-0.0787	0.9149	-0.0856	0.7564	-0.0069	0.9808	0.1402	0.7644	0.0681	0.7783	-0.0721	0.7502
Tsp42Ei	tetraspanin 42E	1623094_at	-1.8164	0.0046	-1.3169	0.0488	-1.6261	0.0001	-0.6681	0.0555	-0.4886	0.0230	0.1795	0.3271	-0.2503	0.8940	0.0536	0.9631	0.3040	0.6988
---	---	1623095_at	0.0731	0.7431	0.1883	0.1673	0.1826	0.3286	0.1232	0.8815	-0.1455	0.6453	-0.2687	0.2915	0.0053	0.9943	0.0389	0.8782	0.0336	0.8905
Tsp68C	Tetraspanin 68C	1623096_a_at	0.2498	0.1101	0.2649	0.1644	0.5447	0.0270	-0.1368	0.7138	-0.0998	0.5898	0.0370	0.8511	-0.0998	0.8449	0.0650	0.8117	0.1647	0.4457
CG31920	CG31920	1623097_at	0.0881	0.6685	0.0804	0.5727	0.0271	0.8956	0.0194	0.9805	0.0904	0.6827	0.0710	0.7330	-0.0341	0.9729	0.1700	0.5100	0.2041	0.4243
CG30323	CG30323	1623098_at	-0.1331	0.4091	-0.0368	0.7193	-0.0088	0.9645	-0.1237	0.6837	-0.0235	0.9040	0.1002	0.4524	-0.2507	0.6898	-0.1771	0.5075	0.0735	0.8202
---	---	1623099_at	0.2200	0.2135	0.0120	0.9101	0.1771	0.2965	0.1298	0.7492	0.1670	0.3533	0.0372	0.8566	0.0217	0.9788	-0.0634	0.7906	-0.0851	0.6867
Aac11	Aac11	1623100_at	-0.1663	0.3457	-0.0597	0.8934	-0.4255	0.0700	-0.3068	0.3534	0.1084	0.6006	0.4152	0.0224	0.0455	0.9816	0.2754	0.5336	0.2299	0.6166
Cbp80	cap binding protei	1623101_at	0.0364	0.8962	0.1527	0.5977	0.0559	0.8140	-0.1566	0.7333	0.0585	0.8183	0.2151	0.2418	-0.0924	0.9467	0.0761	0.9062	0.1685	0.7247
Gip	GIP-like	1623102_at	1.2801	0.0170	0.5402	0.2256	1.3175	0.0004	0.0491	0.9677	-0.4848	0.1269	-0.5339	0.0643	-0.4948	0.7498	-0.9675	0.1395	-0.4727	0.4900
CG14946	CG14946	1623103_at	-2.3602	0.0290	-3.2965	0.0016	-3.6257	0.0000	-0.2822	0.5848	0.3911	0.1359	0.6733	0.0117	-0.0148	0.9977	-0.8634	0.4495	-0.8486	0.4710
Smc5	Smc5	1623104_at	-0.2645	0.2177	-0.1298	0.3818	0.1366	0.4736	-0.1299	0.8090	-0.2610	0.2216	-0.1310	0.5300	-0.2786	0.6935	0.0118	0.9829	0.2904	0.3276
CG18672	CG18672	1623105_at	0.0198	0.9343	0.0431	0.6416	0.1396	0.5465	0.0537	0.9438	0.1538	0.5037	0.1001	0.6574	0.0383	0.9489	0.0982	0.6075	0.0599	0.7764
CtBP	C-terminal Bindin	1623106_s_at	0.1915	0.6418	-0.3659	0.0818	-0.7556	0.0278	0.2315	0.5405	0.7892	0.0026	0.5577	0.0073	0.6027	0.6935	0.2759	0.6961	-0.3268	0.6294
CG10459	CG10459	1623107_at	-0.1803	0.3984	-0.0636	0.7844	0.2259	0.4115	0.1389	0.8544	-0.0282	0.9438	-0.1672	0.5299	-0.2557	0.7152	0.0499	0.9115	0.3056	0.3079
geminin	geminin	1623108_at	0.1403	0.5431	-0.1295	0.5926	-0.0741	0.7004	0.1879	0.6701	0.1065	0.6514	-0.0814	0.7144	0.1264	0.8692	0.0101	0.9870	-0.1163	0.7413
CG4593	CG4593	1623109_at	0.6594	0.0124	0.8022	0.0426	0.9450	0.0043	0.3858	0.4304	0.3042	0.2595	-0.0816	0.7861	0.2163	0.8395	0.3418	0.4355	0.1255	0.8183
---	---	1623110_s_at	0.0115	0.9684	0.0000	1.0000	-0.0328	0.8437	-0.0215	0.9807	-0.0235	0.9399	-0.0020	0.9940	0.0164	0.9862	-0.0931	0.7065	-0.1095	0.6414
CG14362	CG14362	1623111_at	-0.0140	0.9413	-0.1791	0.3703	0.1100	0.5954	0.2207	0.5744	0.3638	0.0738	0.1431	0.4523	-0.0192	0.9845	0.0550	0.8558	0.0742	0.7754
CG4509	CG4509	1623112_at	-1.4836	0.0183	-3.0423	0.0022	-4.0787	0.0000	-0.5429	0.4258	1.5908	0.0021	2.1337	0.0004	0.3595	0.5765	-0.0633	0.8792	-0.4227	0.1701
CG13167	CG13167	1623113_at	0.1136	0.6736	-0.0215	0.8423	0.2988	0.1886	0.0422	0.9471	0.0887	0.6636	0.0465	0.8194	-0.1094	0.9305	0.0841	0.8857	0.1935	0.6588
---	---	1623114_at	0.2279	0.1707	-0.0503	0.6535	-0.1044	0.5161	0.1096	0.7608	0.2853	0.0688	0.1757	0.2045	-0.0213	0.9798	-0.0418	0.8778	-0.0204	0.9386
CG14071	CG14071	1623115_at	0.0418	0.8810	-0.0472	0.8433	-0.0018	0.9935	0.1133	0.8551	0.0798	0.7669	-0.0335	0.9007	-0.0554	0.9555	-0.0453	0.9198	0.0101	0.9838
---	---	1623116_at	-0.9511	0.0614	-1.8167	0.0095	-1.5491	0.0211	0.3770	0.7749	1.1477	0.0436	0.7707	0.1172	-0.1748	0.8012	-0.0964	0.7898	0.0785	0.8289
CG31781	CG31781	1623117_at	-1.9411	0.0008	-1.8337	0.0108	-3.1561	0.0000	-0.7698	0.0390	-0.5769	0.0127	0.1929	0.3021	0.3932	0.6749	-0.4217	0.2481	-0.8149	0.0702
CG4168	chaoptic-like	1623118_s_at	0.5933	0.0227	0.4016	0.0562	0.5116	0.0261	0.0446	0.9308	-0.2471	0.0999	-0.2917	0.0368	0.3293	0.6566	0.0942	0.8105	-0.2351	0.4523
DnaJ-60	DnaJ-like-60	1623119_at	-0.1479	0.6085	0.1125	0.2766	-0.2212	0.3691	0.0524	0.9518	-0.2279	0.3491	-0.2803	0.1893	0.1982	0.8222	-0.0460	0.9387	-0.2441	0.5271
---	---	1623120_at	0.1177	0.5580	0.1115	0.3823	0.0278	0.8792	-0.1091	0.7964	-0.1762	0.3100	-0.0671	0.7133	0.0732	0.9340	-0.1380	0.6512	-0.2112	0.4678
CG5521	CG5521	1623121_at	-0.5313	0.2778	0.3657	0.1216	0.3097	0.0711	-0.1094	0.9323	-0.6425	0.0936	-0.5331	0.1181	-0.0488	0.9695	0.2203	0.5199	0.2691	0.4273
CG17294	CG17294	1623122_at	-0.2968	0.1182	-0.3278	0.0879	-0.3533	0.0782	-0.3058	0.5067	-0.0461	0.8891	0.2597	0.2348	-0.2053	0.7220	-0.1022	0.7187	0.1031	0.7128
---	---	1623123_at	-0.0814	0.6791	0.0201	0.9278	0.0169	0.9350	-0.0202	0.9825	-0.0136	0.9677	0.0066	0.9802	-0.1448	0.8243	-0.0197	0.9648	0.1251	0.6743
CG7376	CG7376	1623124_at	-0.2394	0.5732	-0.4751	0.1276	-0.1279	0.5989	-0.1783	0.7167	0.3984	0.0770	0.5767	0.0113	-0.4735	0.7070	0.0081	0.9941	0.4816	0.3752
CG4459	CG4459	1623125_at	1.2738	0.3180	0.1385	0.4458	0.0814	0.5852	0.1041	0.8143	0.0392	0.8652	-0.0650	0.7278	0.2529	0.9711	-1.1557	0.5249	-1.4087	0.4335
CG13912	CG13912	1623126_at	2.1023	0.0011	1.6616	0.0216	2.3623	0.0002	0.5101	0.3752	0.2436	0.4747	-0.2665	0.3767	-0.2283	0.8968	-0.2531	0.7425	-0.0248	0.9828
mthl12	Mth-like 12	1623127_a_at	-0.0282	0.9059	0.0730	0.5448	0.1252	0.5602	-0.1204	0.7809	-0.0984	0.6217	0.0221	0.9203	-0.1828	0.7826	-0.0062	0.9925	0.1766	0.5615
CG13455	CG13455	1623128_at	0.0510	0.7442	0.0587	0.7071	-0.0540	0.7616	-0.0363	0.9578	0.0768	0.7157	0.1130	0.5219	0.1370	0.7994	-0.0204	0.9557	-0.1574	0.4980
---	---	1623129_at	-0.0409	0.8831	-0.0031	0.9810	0.1020	0.5949	0.0584	0.9308	-0.2259	0.2525	-0.2843	0.1056	0.0548	0.9457	0.0778	0.8028	0.0231	0.9463
CG30457	CG30457	1623130_at	0.2129	0.4762	0.2482	0.3126	0.1890	0.3650	0.0356	0.9761	-0.0782	0.8364	-0.1137	0.7171	-0.0284	0.9848	-0.0997	0.8250	-0.0713	0.8783
---	---	1623131_at	0.1563	0.4290	0.0728	0.6027	0.3537	0.1464	0.0476	0.9314	0.2482	0.1282	0.2005	0.1695	-0.1565	0.8133	0.0435	0.9185	0.2000	0.4815
---	---	1623132_at	0.0359	0.9156	-0.0205	0.8398	0.1007	0.5030	0.1807	0.6908	0.0553	0.8380	-0.1254	0.5421	0.0011	0.9998	0.1743	0.6601	0.1732	0.6577
CG10508	CG10508	1623133_a_at	-1.6364	0.0060	-1.6038	0.0677	-1.8229	0.0002	-0.2441	0.6926	-0.1354	0.6774	0.1088	0.7204	-0.0546	0.9852	-0.0399	0.9716	0.0147	0.9891
CG12506	CG12506	1623134_at	0.0696	0.6317	-0.0296	0.7709	0.0031	0.9916	-0.0511	0.9254	-0.0273	0.9045	0.0239	0.9041	0.0514	0.9277	-0.0366	0.8940	-0.0880	0.6607
---	---	1623135_at	0.3301	0.0831	0.0547	0.8308	0.0953	0.5368	0.1637	0.6822	0.2933	0.1141	0.1295	0.4628	0.0419	0.9514	0.0471	0.8710	0.0052	0.9874
---	---	1623136_at	0.2117	0.3603	-0.0581	0.5299	0.0819	0.7890	0.1131	0.8507	0.0407	0.8906	-0.0724	0.7623	0.0398	0.9816	-0.1659	0.6850	-0.2057	0.6018
dpr8	dpr8	1623137_s_at	-0.0058	0.9741	0.1325	0.4074	0.0979	0.5660	-0.1781	0.5854	-0.0334	0.8806	0.1446	0.3426	-0.0369	0.9705	0.1661	0.5259	0.2030	0.4342
pds5	CG17509	1623138_at	0.4203	0.3074	0.6341	0.1688	0.1836	0.3189	-0.0788	0.9329	0.0892	0.7927	0.1681	0.5343	0.4213	0.7324	0.3480	0.5258	-0.0733	0.9197
dor	deep orange	1623139_at	-0.2743	0.4595	0.0071	0.9873	0.3884	0.0355	0.1975	0.7469	0.1168	0.7051	-0.0806	0.7872	-0.1553	0.9342	0.3640	0.5636	0.5193	0.3963
CG17379	CG17379	1623140_at	0.3243	0.2315	0.0143	0.9517	0.2940	0.1064	0.1348	0.7293	0.1644	0.3536	0.0296	0.8862	-0.0571	0.9514	-0.1569	0.5994	-0.0998	0.7576
---	---	1623141_s_at	0.4833	0.0672	0.2158	0.3203	0.3458	0.0806	-0.0221	0.9828	0.1450	0.5724	0.1671	0.4577	-0.1198	0.7822	-0.0807	0.7029	0.0392	0.8779
---	---	1623142_at	0.2325	0.1290	0.0156	0.9113	0.0137	0.9602	0.1238	0.8137	0.1088	0.6388	-0.0150	0.9548	0.1327	0.8293	0.0494	0.8991	-0.0833	0.7863
---	---	1623143_s_at	0.0873	0.6043	-0.0775	0.5118	0.1094	0.6480	0.0785	0.8817	0.1251	0.5106	0.0466	0.8191	-0.0366	0.9742	-0.0035	0.9955	0.0330	0.9330
ci	Cubitus interruptu	1623144_at	-0.5795	0.2209	-1.8092	0.0097	-1.8992	0.0009	-0.4123	0.5680	0.1870	0.6549	0.5993	0.0727	-0.3295	0.8270	-1.0514	0.1021	-0.7218	0.2668
---	---	1623145_s_at	-0.0193	0.9440	-0.2609	0.3361	-0.6298	0.0097	-0.0505	0.9643	0.5732	0.0596	0.6236	0.0277	0.2091	0.7891	0.0154	0.9797	-0.1937	0.5830
CG10991	CG10991	1623146_at	0.1792	0.3946	0.0638	0.5281	0.1951	0.4977	0.0676	0.9339	0.0224	0.9505	-0.0451	0.8730	0.0018	0.9986	-0.0791	0.7250	-0.0810	0.7135
CG31855	CG31855	1623147_at	-0.1779	0.5604	0.1066	0.6303	0.3531	0.0674	0.0269	0.9803	-0.2000	0.4749	-0.2268	0.3566	0.0574	0.9760	0.0296	0.9689	-0.0277	0.9686
CG12990	CG12990	1623148_at	0.0179	0.9323	-0.0243	0.8239	-0.1520	0.3664	0.0526	0.9311	0.0880	0.6664	0.0353	0.8662	0.1089	0.8692	0.0712	0.8393	-0.0377	0.9171
CG33233	CG33233	1623149_at	0.1036	0.6174	-0.2108	0.3495	-0.1085	0.6692	0.1514	0.7142	0.3005	0.1102	0.1491	0.3956	0.0447	0.9717	0.0358	0.9450	-0.0089	0.9859
RpLP2	rpA1 gene.	1623150_at	0.2331	0.1547	0.6133	0.0269	0.7195	0.0038	0.0171	0.9803	-0.3353	0.0502	-0.3524	0.0266	0.0093	0.9933	0.0048	0.9935	-0.0045	0.9921
CG2010	CG2010	1623151_a_at	-2.2601	0.0067	-2.2271	0.0118	-2.8948	0.0001	-0.3468	0.4239	-0.5298	0.0344	-0.1829	0.4151	0.3189	0.8824	-0.7119	0.3773	-1.0308	0.2251
beat-IIIa	beat-IIIa	1623152_at	0.1856	0.3301	-0.0700	0.7633	0.1413	0.4840	0.2328	0.5988	0.1843	0.4246	-0.0485	0.8516	0.1115	0.8297	-0.0391	0.9064	-0.1507	0.5040
---	---	1623153_s_at	-0.0961	0.7703	0.1765	0.3593	0.1004	0.5759	0.0803	0.9380	-0.1540	0.6519	-0.2343	0.4074	-0.0965	0.8917	0.2106	0.4280	0.3071	0.2655
CG1271	CG1271	1623154_a_at	0.8152	0.0345	-0.0292	0.8453	0.1627	0.4806	-0.1139	0.8794	0.1531	0.5818	0.2670	0.2455	-0.2296	0.8192	-0.6849	0.1033	-0.4553	0.2830
CG33284	CG33284	1623155_at	0.2479	0.1928	0.0948	0.5017	-0.0711	0.8004	0.0061	0.9956	0.2819	0.2804	0.2758	0.2356	0.0804	0.8609	0.0342	0.9039	-0.0462	0.8459
---	---	1623156_at	0.2983	0.2580	-0.0548	0.7560	0.0893	0.6411	0.1373	0.7711	-0.0048	0.9879	-0.1421	0.4498	0.0520	0.9635	-0.1199	0.7421	-0.1719	0.6052
---	---	1623157_at	-0.1617	0.4438	0.1242	0.4099	0.1794	0.2960	0.0241	0.9688	-0.1419	0.3870	-0.1659	0.2482	0.0434	0.9515	0.1637	0.4450	0.1203	0.6017
rho-4	rhomboid-4	1623158_s_at	1.0170	0.0237	1.4318	0.0070	0.5111	0.1242	0.4667	0.2352	0.7619	0.0064	0.2952	0.1554	1.3938	0.2892	1.1713	0.1012	-0.2225	0.7898

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1623159_at	0.0288	0.8693	-0.2357	0.1584	-0.3136	0.0973	-0.1135	0.8512	0.4108	0.0656	0.5243	0.0159	-0.2614	0.7070	-0.0893	0.8131	0.1721	0.5790
Mmp1	Matrix metalloprot	1623160_at	-0.3661	0.6566	-0.5140	0.3879	-0.6346	0.1346	0.3648	0.6354	1.1291	0.0095	0.7644	0.0311	0.6572	0.8270	0.8716	0.4983	0.2144	0.9020
CG15777	CG15777	1623161_at	-0.0321	0.8606	0.1831	0.3624	0.0496	0.8986	-0.1687	0.8544	-0.3256	0.3331	-0.1569	0.6455	-0.0336	0.9816	-0.1642	0.6512	-0.1306	0.7333
---	---	1623162_at	0.0562	0.7951	0.2155	0.3340	-0.1654	0.3802	-0.3959	0.1951	-0.1402	0.4674	0.2557	0.1204	0.1684	0.8191	0.1910	0.5422	0.0225	0.9588
CG33543	CG33543	1623163_at	0.1444	0.5021	-0.1197	0.3975	-0.0518	0.8671	0.0772	0.9295	0.3943	0.1214	0.3171	0.1644	0.0056	0.9964	0.0016	0.9989	-0.0040	0.9937
Eip75B	Ecdysone-induce	1623164_a_at	0.0862	0.7263	0.3511	0.4000	0.2873	0.2136	-0.0188	0.9857	-0.3573	0.1356	-0.3385	0.1138	0.1273	0.9405	-0.0107	0.9931	-0.1380	0.8368
thetaTry	thetaTrypsin	1623165_at	0.4012	0.3354	-0.1341	0.3133	0.1632	0.3333	0.1629	0.6922	0.1330	0.5085	-0.0299	0.8959	-0.1719	0.8940	-0.2962	0.5591	-0.1243	0.8406
CG6071	CG6071	1623166_at	-0.1826	0.5870	-0.0874	0.6227	0.1434	0.4958	0.1374	0.7539	0.1292	0.5212	-0.0082	0.9723	-0.1500	0.9066	0.1211	0.8455	0.2712	0.5786
---	---	1623167_at	0.0814	0.7928	0.2849	0.3510	0.2302	0.3774	0.0130	0.9937	-0.1075	0.7998	-0.1205	0.7481	0.0106	0.9950	-0.0022	0.9989	-0.0128	0.9841
TyrRII	CG16766	1623168_at	0.2211	0.2762	0.0031	0.9858	-0.0247	0.9015	-0.0339	0.9481	0.0275	0.8893	0.0615	0.6887	0.1196	0.9149	-0.0794	0.8893	-0.1990	0.6377
Rim	Rim	1623169_at	0.1449	0.4771	0.0334	0.7740	-0.0743	0.7066	0.0475	0.9463	0.1393	0.5078	0.0917	0.6562	0.0547	0.9390	0.0674	0.8132	0.0127	0.9701
---	---	1623170_at	-0.1821	0.5120	0.0000	1.0000	-0.3188	0.1838	-0.1532	0.8350	-0.0418	0.9147	0.1114	0.7048	-0.0519	0.9495	-0.0821	0.7883	-0.0302	0.9295
lilli	lilliputian	1623171_s_at	0.4048	0.2613	0.0929	0.6469	-0.4765	0.0806	-0.4238	0.4095	0.4502	0.1133	0.8741	0.0051	0.0096	0.9964	0.2239	0.7043	0.2143	0.7154
---	---	1623172_at	0.0554	0.7680	0.0468	0.6790	0.0986	0.6287	-0.0529	0.9252	-0.0630	0.7542	-0.0102	0.9617	0.0020	0.9984	0.0245	0.9372	0.0225	0.9342
CG10013 /// DvirCG10013	CG10013	1623173_at	0.1088	0.6901	0.0603	0.6587	-0.2689	0.2450	-0.1837	0.7138	0.0645	0.8210	0.2482	0.2250	0.1313	0.8202	0.1261	0.6264	-0.0052	0.9895
---	---	1623174_at	0.0539	0.8132	0.1020	0.6298	0.2360	0.3534	0.0861	0.8899	0.1365	0.5423	0.0504	0.8325	-0.0007	0.9998	0.1164	0.7490	0.1171	0.7422
Grx-1	Grx-1	1623175_at	0.0540	0.7633	0.0826	0.7087	0.1709	0.3639	-0.1231	0.8403	0.0342	0.9148	0.1573	0.4675	0.1173	0.8650	0.2034	0.4655	0.0861	0.7955
CG16812	CG16812	1623176_at	-0.1655	0.4356	0.1958	0.2482	0.5614	0.0128	0.3122	0.3837	0.0219	0.9402	-0.2903	0.1083	-0.0252	0.9764	0.3375	0.1054	0.3627	0.1116
CG33228	CG33228	1623177_at	0.1731	0.5257	0.0816	0.6902	0.2445	0.2758	-0.3431	0.2840	-0.0298	0.9092	0.3133	0.0661	-0.3762	0.7215	-0.0481	0.9470	0.3281	0.4764
CG14190	CG14190	1623178_at	0.1088	0.6347	0.1062	0.4748	-0.2144	0.1792	-0.0507	0.9413	-0.0734	0.7548	-0.0227	0.9252	0.1257	0.7714	-0.1057	0.5897	-0.2314	0.2259
CG15415	CG15415	1623179_at	-1.0100	0.0034	-0.2081	0.1746	-1.1059	0.0026	-0.4283	0.2763	-0.2411	0.3033	0.1871	0.3844	-0.0208	0.9860	-0.0526	0.8929	-0.0318	0.9319
Hr46	Complementation	1623180_a_at	-0.7691	0.0113	-0.3012	0.3052	-0.8464	0.0023	-0.1519	0.8064	0.1299	0.6378	0.2818	0.2002	0.2350	0.7485	0.3106	0.3075	0.0756	0.8517
---	---	1623181_at	0.3175	0.0780	0.1215	0.3693	0.2409	0.1902	0.1264	0.7929	0.1079	0.6200	-0.0184	0.9397	0.0120	0.9887	-0.0498	0.8389	-0.0618	0.7744
mRpL51	mitochondrial ribo	1623182_at	0.0632	0.8785	0.5026	0.1360	0.2491	0.2134	-0.1255	0.8403	-0.3488	0.1321	-0.2233	0.2886	0.1943	0.8846	0.1270	0.8551	-0.0673	0.9231
---	---	1623183_at	0.0161	0.9256	0.1445	0.4149	0.2318	0.1844	-0.0791	0.9215	-0.0712	0.8119	0.0080	0.9791	-0.0647	0.9305	-0.0992	0.7165	-0.0346	0.9169
---	---	1623184_at	0.1061	0.5243	0.0102	0.9220	-0.0117	0.9490	0.0844	0.8115	0.0085	0.9690	-0.0759	0.5882	0.0195	0.9816	-0.0527	0.8374	-0.0723	0.7437
jar	Myosin VI	1623185_s_at	0.3872	0.0591	-0.0041	0.9860	0.8254	0.0008	0.2380	0.4214	-0.0484	0.8112	-0.2864	0.0548	-0.4662	0.3166	-0.3818	0.1260	0.0844	0.7730
---	---	1623186_at	0.2850	0.2286	0.1727	0.3962	0.0012	0.9952	0.0128	0.9872	0.0245	0.9268	0.0117	0.9609	0.1331	0.8387	0.0784	0.8247	-0.0547	0.8813
---	---	1623187_at	-0.1038	0.5188	-0.0647	0.6744	0.2162	0.5441	0.1970	0.6465	0.0486	0.8583	-0.1484	0.4498	-0.0980	0.9400	-0.1525	0.7456	-0.0545	0.9211
CG17208	CG17208	1623188_at	0.1517	0.6595	0.2863	0.1334	0.2587	0.1013	-0.0552	0.9376	-0.2781	0.1749	-0.2230	0.2269	0.1094	0.9405	-0.0097	0.9925	-0.1191	0.8350
slik	polo kinase kinas	1623189_at	-0.5898	0.0950	0.1076	0.6553	0.2139	0.2497	-0.0369	0.9659	-0.3629	0.1110	-0.3260	0.1094	-0.0896	0.9589	0.3756	0.4446	0.4652	0.3569
CG9168	CG9168	1623190_at	-0.0085	0.9704	-0.0036	0.9802	0.1570	0.3987	-0.0387	0.9558	-0.1965	0.3014	-0.1578	0.3630	0.0132	0.9875	0.0113	0.9719	-0.0019	0.9953
Rab9	Rab9	1623191_at	-0.2561	0.4621	-0.1949	0.7642	0.0617	0.7851	-0.6125	0.2438	-1.5761	0.0010	-0.9636	0.0049	-0.8140	0.5461	-1.4808	0.0405	-0.6667	0.3017
---	---	1623192_at	0.1731	0.2321	-0.0508	0.6773	0.3467	0.1437	0.0403	0.9592	0.0280	0.9228	-0.0123	0.9621	-0.3459	0.6584	-0.0907	0.8306	0.2552	0.4367
CG33123	CG33123	1623193_at	1.1026	0.0022	1.0065	0.0854	0.7356	0.0029	0.1508	0.7658	0.9816	0.0015	0.8308	0.0019	0.3250	0.7768	0.8524	0.0953	0.5274	0.2997
CG7637	CG7637	1623194_at	0.7762	0.0836	0.4375	0.4019	0.5251	0.0947	0.4715	0.5357	0.9127	0.0292	0.4411	0.2144	0.3297	0.8097	0.5682	0.3040	0.2385	0.7097
CG17944	CG17944	1623195_at	0.0310	0.8885	0.0578	0.5375	-0.0476	0.7899	-0.0367	0.9518	-0.0604	0.7665	-0.0237	0.9084	0.0754	0.8999	-0.0568	0.8501	-0.1322	0.5724
---	---	1623196_s_at	-0.0323	0.8787	-0.0960	0.7341	0.0876	0.5774	-0.0091	0.9940	-0.0240	0.9485	-0.0149	0.9619	-0.2698	0.6955	-0.1481	0.6394	0.1217	0.7115
zormin	D-Titin	1623197_a_at	-2.5508	0.0017	-1.7900	0.0111	-2.6437	0.0001	-0.4312	0.2591	-0.9031	0.0027	-0.4719	0.0274	0.3318	0.8379	-0.2985	0.6894	-0.6303	0.3583
---	---	1623198_at	0.1369	0.5639	-0.0934	0.6403	0.1403	0.4841	-0.0051	0.9956	0.1083	0.5890	0.1134	0.5274	-0.0713	0.9457	-0.0635	0.8915	0.0077	0.9875
CG32352	CG32352	1623199_a_at	-0.7792	0.0087	-1.1642	0.0774	-1.2612	0.0003	0.0676	0.8920	0.2854	0.0878	0.2178	0.1427	0.0978	0.9599	-0.0865	0.9198	-0.1843	0.7774
fng	fringe	1623200_at	-1.5014	0.0006	-2.6130	0.0025	-2.6156	0.0001	0.1910	0.8559	0.9886	0.0173	0.7976	0.0267	0.1510	0.8714	-0.0537	0.9272	-0.2047	0.6067
CG5783	CG5783	1623201_at	-1.1055	0.0116	-1.4885	0.0125	-1.6403	0.0000	-0.1272	0.7863	-0.2435	0.2073	-0.1163	0.5393	-0.0075	0.9964	-0.5929	0.1577	-0.5853	0.1990
CG14113	CG14113	1623202_at	0.1478	0.5939	-0.1381	0.4016	-0.0210	0.9331	0.0402	0.9558	0.0231	0.9331	-0.0170	0.9451	-0.0656	0.9447	-0.1687	0.5776	-0.1031	0.7541
CG7456	CG7456	1623203_at	0.0085	0.9748	0.1643	0.4606	0.0935	0.6421	0.0775	0.9059	0.2378	0.2629	0.1603	0.4179	0.1024	0.9101	0.3730	0.2356	0.2705	0.4114
CG15728	CG15728	1623204_at	-0.0301	0.8902	0.0721	0.5773	-0.0458	0.8418	0.0323	0.9566	-0.0715	0.6979	-0.1038	0.4996	0.0263	0.9831	0.1527	0.6296	0.1264	0.6995
---	---	1623205_at	-0.0488	0.8353	0.0534	0.6138	0.1051	0.5714	-0.0764	0.8967	-0.2020	0.3025	-0.1256	0.5007	0.0040	0.9969	0.0195	0.9627	0.0155	0.9674
eya	clift	1623206_a_at	-0.2345	0.1775	-0.0167	0.9457	-0.1538	0.4015	-0.2379	0.5610	-0.4326	0.0451	-0.1947	0.3071	-0.0676	0.9340	-0.1206	0.6762	-0.0530	0.8812
CG3548	CG3548	1623207_at	0.0432	0.8979	-0.6647	0.2048	-0.9609	0.0044	-0.2406	0.5913	0.6694	0.0101	0.9100	0.0016	-0.0132	0.9962	-0.1936	0.7978	-0.1805	0.8044
Sox100B	Sox100B	1623208_at	0.0137	0.9554	0.0047	0.9695	0.0712	0.7613	0.1075	0.8605	0.1562	0.4989	0.0486	0.8483	-0.0821	0.9316	-0.0453	0.9265	0.0368	0.9330
CG8833	CG8833	1623209_at	-0.2313	0.5694	0.0320	0.8460	-0.0791	0.7069	-0.0765	0.8908	0.4954	0.0163	0.5719	0.0052	0.1245	0.9250	0.5642	0.1900	0.4397	0.3335
dimm	Dimmed	1623210_at	0.1174	0.5843	-0.0488	0.6206	0.1658	0.3205	0.1229	0.8140	-0.0104	0.9743	-0.1333	0.5002	-0.0057	0.9964	-0.1164	0.7338	-0.1107	0.7437
CG17234	CG17234	1623211_at	2.1806	0.0044	0.3957	0.0209	3.9526	0.0000	3.3724	0.0049	2.6804	0.0011	-0.6921	0.1588	0.0254	0.9746	0.9833	0.0048	0.9579	0.0050
tamo	tamo	1623212_s_at	-0.4313	0.4124	-1.3390	0.0740	-1.2321	0.0005	0.4125	0.2090	1.2587	0.0004	0.8462	0.0012	0.2797	0.9011	0.3109	0.7494	0.0311	0.9829
---	---	1623213_at	0.3682	0.0390	-0.0835	0.4973	0.1378	0.4195	0.3253	0.5515	0.4133	0.1423	0.0879	0.7799	0.0692	0.8909	-0.0403	0.8830	-0.1095	0.5892
CG9669	CG9669	1623214_at	0.6272	0.0161	0.6770	0.0232	0.8426	0.0031	0.1378	0.7307	0.3248	0.0725	0.1870	0.2449	0.0627	0.9558	0.5111	0.1242	0.4484	0.1987
---	---	1623215_at	0.1299	0.3972	0.2077	0.3611	0.0602	0.7834	-0.0485	0.9445	-0.1482	0.4776	-0.0997	0.6238	0.0364	0.9589	-0.0122	0.9709	-0.0486	0.8517
CG9629	CG9629	1623216_s_at	0.7161	0.0585	-0.1676	0.8266	0.7530	0.0079	0.0105	0.9955	-0.1403	0.7495	-0.1509	0.6988	-0.8257	0.6557	-1.0482	0.1498	-0.2225	0.8088
CG8743	CG8743	1623217_at	0.1206	0.6187	-0.1123	0.3605	0.2433	0.2118	0.1070	0.8221	-0.0054	0.9845	-0.1124	0.5351	-0.0343	0.9761	-0.0572	0.8857	-0.0229	0.9525
Alh	Alhambra	1623218_s_at	0.1724	0.5692	0.5209	0.4025	-0.1350	0.6199	-0.2775	0.6533	0.0522	0.8980	0.3297	0.2157	0.3034	0.8379	0.5505	0.3585	0.2471	0.7223
---	---	1623219_at	0.2550	0.1569	-0.0686	0.6196	-0.0233	0.9053	-0.1181	0.8449	-0.0560	0.8461	0.0622	0.8039	-0.0141	0.9862	-0.1052	0.6032	-0.0911	0.6577
---	---	1623220_at	0.2358	0.2482	0.0857	0.4996	0.0945	0.5526	0.0697	0.9343	0.2689	0.2809	0.1992	0.3868	0.0088	0.9916	0.0211	0.9473	0.0123	0.9665
---	---	1623221_at	0.0488	0.7705	0.1010	0.6594	0.2215	0.2373	-0.0373	0.9639	-0.0266	0.9309	0.0107	0.9689	0.1221	0.8655	0.1820	0.5446	0.0599	0.8779
His3.3B	anon-fast-evolvin	1623222_s_at	-0.2234	0.5802	0.6930	0.2275	0.2780	0.3846	0.1535	0.6650	-0.4749	0.0126	-0.6284	0.0022	0.5157	0.8033	0.3643	0.7090	-0.1514	0.8984
---	---	1623223_at	0.2556	0.2773	-0.0566	0.6335	-0.0117	0.9625	0.2107	0.5690	0.2229	0.2411	0.0122	0.9594	-0.1039	0.9174	-0.1532	0.6949	-0.0493	0.9181
Eig71Eb	Gene II	1623224_at	0.0888	0.6516	0.0447	0.7854	-0.0939	0.6415	-0.0445	0.9358	0.0539	0.7851	0.0984	0.5355	0.1546	0.8076	0.1386	0.6271	-0.0159	0.9686
CG7611	CG7611	1623225_s_at	0.1298	0.5459	0.0530	0.8880	0.0644	0.6799	-0.1503	0.7121	-0.2355	0.2023	-0.0853	0.6571	-0.2548	0.7768	-0.2391	0.5548	0.0157	0.9809
CG10154	CG10154	1623226_at	-0.5862	0.8159	1.5534	0.0293	0.6635	0.5119	-0.4584	0.9036	-2.0545	0.0944	-1.5961	0.1444	0.4017	0.9717	-0.0502	0.9935	-0.4519	0.9067
Hil	D-hillarin	1623227_s_at	-0.9713	0.0185	-1.8882	0.0123	-1.9635	0.0002	0.1949	0.8671	0.8402	0.0465	0.6453	0.0806	0.2353	0.8215	-0.1624	0.7506	-0.3977	0.3705

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1623228_at	0.0005	0.9985	0.0698	0.5128	0.1618	0.2889	0.1283	0.7409	-0.0071	0.9790	-0.1353	0.3941	0.0442	0.9646	0.0321	0.9407	-0.0122	0.9773
CG7881	CG7881	1623229_at	-0.4885	0.7524	-1.0173	0.0167	-0.4328	0.1818	0.4318	0.8405	-0.0911	0.9363	-0.5229	0.4970	-0.2536	0.9545	-0.8715	0.5179	-0.6179	0.6609
---	---	1623230_s_at	0.1577	0.4892	0.2869	0.0447	0.2948	0.1449	-0.1041	0.8293	-0.0390	0.8736	0.0650	0.7452	0.2479	0.7266	0.2213	0.4791	-0.0266	0.9510
dyn-p25	dynactin, p25 subu	1623231_at	-0.6619	0.0230	-0.2615	0.0752	-0.2173	0.1433	0.0448	0.9254	-0.1239	0.4019	-0.1687	0.1886	0.0286	0.9816	0.1946	0.4683	0.1659	0.5523
---	---	1623232_at	-0.1124	0.5089	-0.0826	0.5856	-0.1412	0.4407	-0.0149	0.9857	0.0129	0.9643	0.0279	0.9026	0.0252	0.9835	-0.0272	0.9521	-0.0524	0.8924
Rab26	Rab26	1623233_at	0.1085	0.5671	0.1286	0.4950	0.1469	0.3856	0.0326	0.9641	-0.0636	0.7861	-0.0962	0.6241	0.0071	0.9950	-0.0025	0.9973	-0.0096	0.9835
Calx	Na/Ca-exchange p	1623234_s_at	-2.6795	0.0008	-1.8026	0.0205	-2.6931	0.0001	-0.2357	0.4940	-0.3631	0.0522	-0.1274	0.4630	0.6322	0.7187	0.4937	0.5120	-0.1386	0.8900
sec10	sec10	1623235_at	0.1891	0.3805	-0.1066	0.7479	-0.3469	0.0892	-0.0615	0.9190	0.7577	0.0024	0.8192	0.0010	0.0727	0.9487	0.3794	0.2442	0.3067	0.3767
---	---	1623236_at	0.0788	0.7338	-0.0598	0.6882	-0.0081	0.9729	0.0391	0.9507	0.0365	0.8758	-0.0026	0.9906	-0.0376	0.9689	-0.1493	0.5776	-0.1117	0.6900
CG14030	CG14030	1623237_at	0.1192	0.6438	0.1692	0.3279	-0.3088	0.2133	-0.0771	0.8967	0.3229	0.0990	0.3999	0.0299	0.5644	0.5228	0.4748	0.2430	-0.0896	0.8751
CG5618	CG5618	1623238_at	-0.4451	0.1827	0.8708	0.0639	1.1492	0.0024	0.0359	0.9774	-0.6936	0.0327	-0.7295	0.0167	-0.3194	0.8202	0.6888	0.2169	1.0082	0.1158
prod	proliferation disru	1623239_at	-0.2560	0.3056	0.7527	0.0325	0.7760	0.0007	0.0264	0.9744	-0.7113	0.0043	-0.7377	0.0022	-0.0747	0.9441	0.2636	0.4094	0.3384	0.3048
---	---	1623240_at	0.0275	0.9416	0.0742	0.5387	0.0168	0.9218	-0.0316	0.9767	-0.1016	0.7478	-0.0700	0.8163	0.0242	0.9831	0.0314	0.9425	0.0071	0.9868
CG30104	CG30104	1623241_s_at	-0.2350	0.5027	0.6231	0.2816	0.2014	0.6664	-0.6151	0.5008	-1.9794	0.0024	-1.3643	0.0076	-0.1866	0.9168	-1.1158	0.0831	-0.9292	0.1575
---	---	1623242_at	0.0499	0.8077	0.0359	0.8608	-0.0687	0.7145	-0.1158	0.7764	0.0967	0.6054	0.2125	0.1624	0.0814	0.8909	0.0725	0.7985	-0.0089	0.9811
CG5656	CG5656	1623243_at	0.0013	0.9962	-0.0065	0.9599	0.0897	0.6653	-0.0965	0.9110	-0.0997	0.7613	-0.0032	0.9925	-0.1417	0.8558	-0.0909	0.8304	0.0508	0.9086
---	---	1623244_at	-0.2855	0.3929	-0.5355	0.1625	-1.9787	0.0002	-0.0844	0.9247	0.6759	0.0199	0.7603	0.0073	1.2253	0.2553	0.3288	0.5591	-0.8965	0.1271
---	---	1623245_at	0.0516	0.7793	0.0101	0.9379	0.2542	0.1937	-0.0500	0.9442	-0.1158	0.6064	-0.0658	0.7711	-0.0126	0.9898	0.0856	0.7269	0.0981	0.6719
CG6475	CG6475	1623246_at	0.0856	0.6094	-0.0131	0.9054	0.1951	0.3495	-0.2142	0.5837	-0.1543	0.4562	0.0599	0.7878	-0.1563	0.8122	-0.1750	0.5367	-0.0187	0.9626
CG10420	CG10420	1623247_at	0.9216	0.1113	1.9164	0.0063	1.5398	0.0012	0.4236	0.3328	1.1738	0.0014	0.7502	0.0059	0.5773	0.7726	2.0570	0.0391	1.4798	0.1184
---	---	1623248_s_at	-0.1048	0.6469	0.0210	0.9468	-0.1047	0.6795	-0.1803	0.6656	-0.1282	0.5495	0.0521	0.8172	-0.0017	0.9997	0.0007	0.9998	0.0024	0.9979
---	---	1623249_s_at	-0.1030	0.4680	-0.0129	0.9146	-0.0540	0.7677	0.0014	0.9985	-0.0911	0.5858	-0.0925	0.5385	0.0772	0.8680	-0.0479	0.8481	-0.1251	0.5145
CG18591	CG18591	1623250_at	0.1374	0.4169	0.1333	0.6226	0.3433	0.1018	0.0480	0.9298	-0.0396	0.8478	-0.0876	0.5816	-0.0482	0.9690	-0.0173	0.9758	0.0308	0.9463
CG14450	CG14450	1623251_at	-0.4604	0.0200	-0.5048	0.0843	-0.5702	0.0034	-0.2000	0.4786	0.0459	0.8053	0.2459	0.0716	0.0677	0.9015	0.2042	0.2960	0.1365	0.5142
---	---	1623252_a_at	-0.3156	0.5994	-0.1366	0.7964	-1.1104	0.0112	-0.2497	0.8631	0.1771	0.7757	0.4268	0.3691	0.3215	0.8903	0.1088	0.9391	-0.2127	0.8509
---	---	1623253_at	0.1901	0.3207	0.1306	0.4884	0.0012	0.9959	-0.1057	0.8605	0.0916	0.7136	0.1974	0.3147	0.1641	0.8400	0.2087	0.5527	0.0446	0.9231
CG31531	CG31531	1623254_s_at	-1.1413	0.0058	-0.7802	0.1068	-1.7063	0.0000	-0.2897	0.4335	-0.1473	0.4886	0.1423	0.4589	0.3664	0.7204	0.1192	0.8341	-0.2472	0.5858
---	---	1623255_at	0.2312	0.2992	0.1420	0.3969	0.2330	0.1824	0.0050	0.9956	-0.0211	0.9338	-0.0261	0.9042	-0.0294	0.9848	-0.0417	0.9407	-0.0122	0.9837
GstE1	glutathione-S-tran	1623256_at	0.5719	0.0691	0.3706	0.0948	0.4011	0.0401	0.2912	0.4116	0.2115	0.2852	-0.0797	0.7002	0.4003	0.6557	0.0937	0.8504	-0.3067	0.4078
CG17665	CG17665	1623257_a_at	-0.4296	0.0303	-0.5300	0.1447	-0.6318	0.0039	-0.0801	0.8794	-0.0395	0.8687	0.0407	0.8455	0.0356	0.9816	-0.1036	0.8326	-0.1392	0.7432
CG12493	CG12493	1623258_at	-0.0467	0.9583	0.0155	0.9203	-0.2620	0.1254	-0.4386	0.7413	-0.4062	0.5120	0.0325	0.9651	-0.0258	0.9911	-0.2041	0.6998	-0.1783	0.7416
---	---	1623259_at	0.2331	0.1543	-0.0089	0.9674	0.2448	0.1297	0.0444	0.9436	-0.0052	0.9844	-0.0496	0.8044	-0.1290	0.8378	-0.1749	0.5140	-0.0459	0.8977
---	---	1623260_at	0.0455	0.8307	-0.1552	0.4834	0.0146	0.9353	0.1756	0.6189	0.2913	0.0978	0.1157	0.4922	-0.0353	0.9779	0.0155	0.9769	0.0508	0.9032
RabX4	RabX4	1623261_at	-0.0222	0.9049	-0.0894	0.4844	-0.3900	0.1600	-0.0924	0.8871	-0.2242	0.3122	-0.1318	0.5393	0.1204	0.9326	-0.4053	0.3669	-0.5257	0.2666
CG13883	CG13883	1623262_at	0.3590	0.0452	0.1210	0.5184	0.2705	0.1015	-0.1766	0.7121	-0.0380	0.8983	0.1386	0.5090	0.0211	0.9875	-0.0943	0.8123	-0.1153	0.7440
---	---	1623263_at	0.0756	0.6710	0.0500	0.8523	-0.0714	0.7859	-0.1388	0.8067	0.0806	0.7655	0.2194	0.2811	-0.0916	0.8846	-0.0536	0.8754	0.0379	0.9086
CG3493	CG3493	1623264_at	0.0806	0.7857	0.6851	0.0305	0.8662	0.0013	0.2850	0.4420	0.0943	0.6827	-0.1907	0.2982	0.1952	0.8062	0.6993	0.0568	0.5042	0.1550
CG10513	CG10513	1623265_at	-3.4567	0.0129	-0.6914	0.6600	-2.4747	0.0033	-1.6898	0.1119	-3.0786	0.0011	-1.3888	0.0223	0.2617	0.9742	-0.0979	0.9784	-0.3596	0.8962
CG14180	CG14180	1623266_at	-0.1074	0.6431	0.0340	0.7562	0.0642	0.6737	-0.0028	0.9978	-0.0343	0.9252	-0.0314	0.9218	0.0497	0.9411	0.1277	0.5591	0.0780	0.7453
CG5237	CG5237	1623267_at	-0.2339	0.3604	-0.0432	0.7695	0.0197	0.9171	0.0048	0.9956	-0.0840	0.7266	-0.0888	0.6800	-0.1423	0.8037	-0.0817	0.7810	0.0607	0.8418
CG33785 /// CG33786	CG33786 /// CG3	1623268_a_at	0.0397	0.8923	-0.2943	0.3074	-0.3145	0.0757	0.2381	0.7121	0.8598	0.0103	0.6217	0.0257	0.1776	0.8076	0.5246	0.0943	0.3470	0.2697
---	---	1623269_at	-0.0685	0.7633	0.0202	0.8580	-0.0416	0.8315	0.1442	0.7814	0.1168	0.6253	-0.0274	0.9173	0.0564	0.9515	0.1056	0.7490	0.0493	0.8983
---	---	1623270_at	0.3413	0.0894	0.0639	0.7156	0.2148	0.3256	0.0395	0.9655	0.2246	0.3643	0.1852	0.4140	-0.0589	0.9296	-0.0895	0.7162	-0.0306	0.9181
janA	janus A	1623271_a_at	-0.2048	0.2090	0.1291	0.5600	0.3530	0.0772	0.0215	0.9744	-0.2406	0.1392	-0.2621	0.0745	-0.1190	0.8756	0.0937	0.8061	0.2127	0.4892
Acp53C14b	Acp53C14b	1623272_at	0.5873	0.1903	0.9178	0.1580	0.9333	0.0807	-0.4262	0.7556	-0.9362	0.1110	-0.5100	0.3452	-0.1781	0.9514	-0.6100	0.4952	-0.4319	0.6445
CG5432 /// dei	CG5432 /// delilah	1623273_at	0.0619	0.6913	-0.0224	0.8458	-0.0003	0.9984	-0.1009	0.8791	-0.0340	0.9143	0.0669	0.7912	-0.1913	0.7142	-0.1366	0.5488	0.0547	0.8444
bol	boule	1623274_s_at	-0.5102	0.0406	-0.2636	0.3337	-0.8805	0.0017	-0.2310	0.5751	0.0737	0.7718	0.3046	0.1059	-0.0494	0.9405	-0.0446	0.8785	0.0048	0.9880
Or35a	Odorant receptor	1623275_at	0.1557	0.3227	-0.0614	0.7929	-0.0770	0.6036	-0.0117	0.9931	0.1013	0.7486	0.1130	0.6861	-0.0005	0.9998	-0.0405	0.8884	-0.0400	0.8831
CG31663	CG31663	1623276_at	-0.4732	0.2002	-0.2415	0.2701	-0.5745	0.0278	-0.2252	0.6869	-0.5241	0.0487	-0.2990	0.1968	0.0451	0.9547	-0.3788	0.1095	-0.4238	0.1035
CG17100	CG17100	1623277_at	-0.1267	0.8167	-0.0536	0.7783	-1.1664	0.0019	-0.6031	0.1152	0.2330	0.3157	0.8361	0.0024	0.4638	0.7644	0.3389	0.6281	-0.1249	0.8887
---	---	1623278_at	-0.1593	0.2925	0.1780	0.3672	0.2645	0.1784	-0.0803	0.8578	-0.3537	0.0355	-0.2734	0.0620	0.1257	0.8480	0.2726	0.3064	0.1468	0.6189
---	---	1623279_at	0.2978	0.1368	0.0766	0.7713	0.3076	0.1181	0.0791	0.8868	0.1625	0.3996	0.0834	0.6693	-0.0599	0.9581	0.0583	0.9053	0.1182	0.7484
---	---	1623280_at	-0.0711	0.6920	-0.1319	0.3430	0.0848	0.6223	0.1875	0.6493	0.1024	0.6474	-0.0851	0.6832	-0.1763	0.7307	-0.1005	0.6819	0.0758	0.7674
His2B:CG17949 /// His2B:C	His2B:CG33868 //	1623281_s_at	-0.2069	0.6916	-2.0012	0.0360	-1.4509	0.0596	0.1416	0.9371	2.1749	0.0022	2.0333	0.0017	-0.3287	0.9195	0.1952	0.9083	0.5239	0.6621
CG31171	CG31171	1623282_at	0.1407	0.4771	0.1208	0.4347	0.1538	0.4004	-0.0515	0.9603	-0.0472	0.8994	0.0043	0.9901	0.1218	0.8486	0.1104	0.7162	-0.0114	0.9788
---	---	1623283_at	0.0388	0.8834	0.1416	0.5325	0.1516	0.3541	0.0562	0.9300	-0.0447	0.8544	-0.1009	0.5912	0.1345	0.8270	0.0981	0.7424	-0.0364	0.9174
CG10251	CG10251	1623284_at	-0.1467	0.5064	-0.0161	0.9022	0.0220	0.9046	0.0356	0.9436	-0.0612	0.7119	-0.0968	0.4784	0.0356	0.9717	0.0326	0.9353	-0.0030	0.9946
CG1637	CG1637	1623285_at	0.4452	0.1727	-0.5647	0.0286	-0.0341	0.9228	-0.0041	0.9956	0.2127	0.3392	0.2168	0.2725	-0.5179	0.6897	-0.7830	0.1374	-0.2651	0.6398
CG32299	CG32299	1623286_at	0.0708	0.6588	-0.0498	0.8217	0.0394	0.8194	-0.0753	0.8815	0.0809	0.6802	0.1562	0.3197	-0.1121	0.8541	-0.0803	0.8018	0.0317	0.9272
CG13921	CG13921	1623287_at	0.1411	0.7536	-0.2299	0.5285	-0.4864	0.1855	-0.3585	0.6578	0.0467	0.9320	0.4052	0.2457	-0.1854	0.9340	-0.2865	0.7304	-0.1011	0.9187
Pros26	Proteasome subu	1623288_at	0.0458	0.8540	0.3691	0.0985	0.6838	0.0027	0.2274	0.5233	-0.2462	0.1850	-0.4736	0.0122	-0.0500	0.9710	0.1021	0.8218	0.1521	0.6924
Obp83ef	Pseudogene 83e	1623289_at	-0.5859	0.2352	0.3121	0.7300	0.9094	0.0372	-0.1874	0.8327	-0.7529	0.0328	-0.5655	0.0641	-0.5588	0.8331	0.1509	0.9323	0.7097	0.5437
Osi4	Osiris	1623290_at	-0.1164	0.4634	-0.1525	0.3735	0.1384	0.3821	0.0996	0.8496	0.1828	0.3479	0.0831	0.6776	-0.1670	0.7810	-0.0737	0.8304	0.0933	0.7544
CG31673	CG31673	1623291_at	-0.3110	0.1193	0.2584	0.2222	0.3142	0.1036	-0.3027	0.5019	-0.9637	0.0026	-0.6609	0.0082	-0.3057	0.6332	-0.2302	0.3756	0.0755	0.8190
CG33128	CG33128	1623292_at	0.2537	0.4024	0.0250	0.8062	-0.2882	0.3343	-0.2516	0.5617	-0.0285	0.9273	0.2232	0.2657	0.0876	0.9246	-0.0960	0.8123	-0.1836	0.5808
CG4042	CG4042	1623293_at	-0.4119	0.2479	0.2933	0.1902	0.7911	0.0118	-0.1657	0.6122	-0.8390	0.0009	-0.6733	0.0014	-0.3957	0.7541	0.0996	0.9036	0.4953	0.3695
CG17669	CG17669	1623294_at	-0.1157	0.6784	0.1484	0.2215	0.1255	0.4205	-0.0968	0.8940	-0.1982	0.4276	-0.1014	0.6893	0.0006	0.9998	0.1337	0.6600	0.1331	0.6571
CG8564	CG8564	1623295_at	-0.0806	0.7888	0.1236	0.3952	0.2630	0.1655	-0.0228	0.9846	-0.2389	0.3847	-0.2161	0.3853	-0.0413	0.9487	0.0237	0.9401	0.0650	0.7745
CG32625	CG32625	1623296_at	0.6828	0.1459	-0.0012	0.9999	-0.4798	0.0427	-0.2186	0.6544	0.1747	0.4778	0.3933	0.0649	0.2812	0.8454	-0.2424	0.7200	-0.5236	0.3859

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18371	CG18371	1623297_at	-0.1127	0.5537	-0.1974	0.1896	-0.0418	0.7990	-0.1774	0.5633	-0.1371	0.3967	0.0403	0.8216	-0.1595	0.8122	-0.0977	0.7741	0.0618	0.8674
CG6067	CG6067	1623298_at	2.4980	0.0079	1.5314	0.0134	2.5633	0.0007	0.5405	0.7003	-0.1290	0.8810	-0.6695	0.2475	-0.5378	0.7215	-1.1984	0.0716	-0.6606	0.3047
Mmp2	Matrix metalloprot	1623299_at	-0.8668	0.0023	-0.3911	0.3636	-0.3504	0.0677	0.3035	0.6919	-0.2854	0.4371	-0.5889	0.0663	0.0093	0.9940	0.0426	0.9263	0.0333	0.9344
CG18586 /// CG5568	CG18586 /// CG5	1623300_at	0.0171	0.9361	-0.0353	0.8877	-0.0379	0.9218	-0.2046	0.5639	0.2290	0.2072	0.4335	0.0159	-0.3099	0.8049	-0.0487	0.9535	0.2612	0.6414
---	---	1623301_at	0.1089	0.5277	0.2133	0.1279	0.1956	0.2132	-0.0703	0.8815	-0.0584	0.7613	0.0119	0.9535	0.1114	0.8486	0.1127	0.6762	0.0013	0.9979
CG8067	CG8067	1623302_at	0.2901	0.3044	-0.2210	0.3755	-0.3779	0.0415	-0.1762	0.6147	0.3074	0.0811	0.4836	0.0083	-0.1180	0.9087	-0.3039	0.3986	-0.1859	0.6350
---	---	1623303_at	0.0631	0.7441	0.0778	0.4102	0.3006	0.0520	-0.1122	0.7979	0.0299	0.9029	0.1421	0.3892	-0.0228	0.9829	0.0025	0.9967	0.0253	0.9408
CG9107	CG9107	1623304_at	-0.1145	0.5235	0.0800	0.5415	-0.0341	0.8711	-0.1743	0.7735	-0.3732	0.1405	-0.1989	0.4004	-0.0978	0.8842	-0.1019	0.7299	-0.0041	0.9924
CG5091	CG5091	1623305_at	-0.0203	0.9528	-0.2022	0.3742	-0.3290	0.0883	-0.3660	0.3625	0.0099	0.9780	0.3758	0.0690	-0.1263	0.8690	-0.0713	0.8678	0.0551	0.8947
l(1)G0004	lethal (1) G0004	1623306_at	0.0668	0.8901	0.2734	0.2380	0.8133	0.0019	0.1361	0.8901	-0.5573	0.0918	-0.6934	0.0268	-0.4032	0.6984	-0.3582	0.4097	0.0450	0.9421
---	---	1623307_at	-0.0294	0.8750	0.0993	0.6229	0.2543	0.2385	0.1091	0.8815	-0.0700	0.8225	-0.1792	0.4463	0.1661	0.7685	0.1883	0.4302	0.0222	0.9471
CG33784	CG33784	1623308_at	0.2754	0.2733	-0.1526	0.5114	0.1992	0.2867	0.1517	0.8411	0.1784	0.5573	0.0268	0.9393	-0.2465	0.7230	-0.3314	0.2518	-0.0850	0.8209
NepYr	Neuropeptide Y re	1623309_at	0.9816	0.4538	-0.7700	0.0233	-0.0939	0.5515	0.7054	0.0457	0.8359	0.0022	0.1305	0.4973	0.1255	0.9862	-0.9059	0.6156	-1.0314	0.5637
---	---	1623310_at	-0.0040	0.9898	0.3267	0.0946	0.3088	0.0938	-0.0766	0.8794	-0.3300	0.0602	-0.2535	0.1006	-0.0504	0.9535	0.0012	0.9991	0.0516	0.8834
CG34127	CG34127	1623311_at	-0.0778	0.7964	0.2258	0.2033	0.0104	0.9729	-0.2002	0.7929	-0.1930	0.5647	0.0072	0.9854	0.0158	0.9924	0.1737	0.6879	0.1578	0.7179
CG11836	CG11836	1623312_s_at	-0.8168	0.0356	-1.4553	0.0169	-1.2620	0.0036	-0.0074	0.9952	0.8433	0.0051	0.8506	0.0029	-0.2010	0.9333	0.0948	0.9401	0.2958	0.7391
CG6114	SNF1-related	1623313_at	-0.0987	0.5347	0.0262	0.8627	0.0468	0.7865	-0.0634	0.8863	-0.0805	0.6300	-0.0172	0.9255	0.0061	0.9952	0.0820	0.7922	0.0759	0.8011
toy	twin of eyeless	1623314_at	-0.0733	0.6655	-0.2164	0.2947	0.0923	0.7375	0.0277	0.9728	0.0965	0.6750	0.0688	0.7553	-0.3322	0.6832	-0.1394	0.7162	0.1928	0.5913
---	---	1623315_at	0.0997	0.6176	-0.0078	0.9515	-0.1520	0.4839	0.1282	0.7749	0.1506	0.4384	0.0224	0.9227	0.1821	0.8215	0.0040	0.9959	-0.1781	0.6225
CG14785	CG14785	1623316_at	-0.2126	0.6509	0.0570	0.8599	-0.1483	0.4982	-0.2450	0.5242	-0.3965	0.0539	-0.1515	0.4222	0.0312	0.9898	-0.2256	0.7040	-0.2568	0.6498
CG15477	CG15477	1623317_at	0.2402	0.1321	0.1699	0.1326	0.3894	0.0273	-0.1685	0.5619	-0.1943	0.1897	-0.0259	0.8834	-0.1468	0.7893	-0.1267	0.6119	0.0201	0.9512
Met75Ca /// Met75Cb	Met75Ca /// Met7	1623318_s_at	0.1971	0.3562	0.1399	0.4406	-0.0749	0.7260	-0.0104	0.9883	0.0579	0.7757	0.0683	0.6998	0.1766	0.7726	0.0755	0.8280	-0.1011	0.7391
---	---	1623319_at	0.3374	0.0907	0.2335	0.1652	0.2480	0.1062	0.0399	0.9345	-0.0079	0.9715	-0.0478	0.7592	-0.0892	0.8963	-0.0979	0.7439	-0.0087	0.9840
CG3711	CG3711	1623320_at	-0.0434	0.9191	-0.3596	0.1468	-0.3848	0.0610	-0.1299	0.8007	0.3913	0.0627	0.5212	0.0126	0.0571	0.9816	0.1542	0.8248	0.0970	0.8940
CG40343	CG40343	1623321_at	-0.0214	0.9187	-0.1025	0.4427	0.1607	0.4086	0.1134	0.8115	0.0563	0.8089	-0.0570	0.7868	-0.0711	0.9238	-0.0228	0.9551	0.0483	0.8880
CG9067	CG9067	1623322_at	0.2870	0.1208	0.2422	0.2977	0.0516	0.7608	-0.0934	0.8132	0.1167	0.4739	0.2101	0.1299	0.1583	0.8016	0.0683	0.8493	-0.0900	0.7710
Gr10b	Gustatory recepto	1623323_at	-0.0552	0.7245	0.0180	0.8803	0.1920	0.3077	0.0353	0.9479	-0.0670	0.7018	-0.1023	0.4795	0.0145	0.9893	0.0208	0.9543	0.0063	0.9859
lin	lines	1623324_at	0.0676	0.8770	0.3465	0.0791	0.2763	0.1060	0.0078	0.9937	0.0800	0.7228	0.0723	0.7268	0.1753	0.8450	0.3718	0.3000	0.1965	0.6201
CG6847 /// DmirCG6847	CG6847	1623325_at	-0.8691	0.0833	-1.2131	0.1326	-0.8860	0.0258	0.5457	0.2101	0.1973	0.4712	-0.3483	0.1351	0.2397	0.9243	-0.2654	0.8059	-0.5051	0.5720
mge	maggie	1623326_a_at	0.3025	0.0626	-0.1444	0.6723	-0.3082	0.0623	0.0141	0.9842	0.4949	0.0093	0.4809	0.0064	0.2285	0.7893	0.1752	0.6619	-0.0533	0.9161
---	---	1623327_at	0.8823	0.0032	0.3582	0.7072	1.2228	0.0015	0.1696	0.6580	-0.1142	0.5658	-0.2838	0.0856	-0.5628	0.7772	-0.6678	0.4391	-0.1050	0.9313
CG7252	CG7252	1623328_at	0.0838	0.7490	0.1602	0.4409	0.3706	0.0246	-0.0976	0.7906	-0.1602	0.2896	-0.0626	0.6913	-0.1018	0.9101	0.0511	0.9188	0.1529	0.6617
CG14500	CG14500	1623329_at	0.4517	0.2455	-0.1150	0.5285	0.3816	0.1399	0.2238	0.5545	0.0757	0.7434	-0.1480	0.4190	-0.2164	0.7358	-0.3524	0.1838	-0.1360	0.6435
CG5568	CG5568	1623330_at	0.3057	0.0737	-0.2841	0.1078	0.2171	0.4982	0.4082	0.3032	0.4939	0.0404	0.0856	0.7317	-0.2300	0.6898	-0.1375	0.5808	0.0925	0.7307
CG32199	CG32199	1623331_at	-0.1312	0.3499	0.0031	0.9884	-0.1225	0.4082	-0.0477	0.9313	0.0065	0.9795	0.0542	0.7617	-0.0198	0.9893	0.0756	0.8592	0.0954	0.7972
Obp99d	Odorant-binding p	1623332_at	-0.0429	0.8747	0.4367	0.0745	0.1437	0.4042	-0.2424	0.5543	-0.4281	0.0486	-0.1857	0.3383	0.2090	0.7802	0.2037	0.5468	-0.0053	0.9924
l(1)G0007	lethal (1) G0007	1623333_a_at	-0.1097	0.8633	0.3461	0.6540	0.6600	0.1224	0.1748	0.8274	0.0240	0.9582	-0.1509	0.6322	-0.1420	0.9737	0.6130	0.5855	0.7550	0.4916
CG12913	CG12913	1623334_at	0.0563	0.7951	0.1285	0.4206	-0.0296	0.8786	-0.1187	0.8473	-0.1640	0.4886	-0.0453	0.8632	0.1422	0.8438	0.0653	0.8773	-0.0769	0.8375
CG5044	CG5044	1623335_a_at	0.3410	0.4725	0.2910	0.0985	0.7537	0.0089	-0.1916	0.6010	-0.9892	0.0008	-0.7976	0.0011	-0.6033	0.7049	-0.8700	0.1761	-0.2667	0.7238
CG13719	CG13719	1623336_at	0.1301	0.6002	-0.0810	0.6975	-0.1633	0.4397	0.0809	0.8817	0.0027	0.9916	-0.0782	0.6880	0.1439	0.8993	-0.0508	0.9404	-0.1947	0.6708
---	---	1623337_at	-0.0281	0.9250	-0.0229	0.9153	0.0281	0.8730	-0.0498	0.9463	-0.0237	0.9364	0.0261	0.9187	-0.0569	0.9530	-0.0155	0.9751	0.0414	0.9171
---	---	1623338_at	0.3249	0.2699	0.3355	0.2864	0.0782	0.6408	0.0074	0.9937	-0.1199	0.5601	-0.1273	0.4842	0.0624	0.9657	-0.0627	0.9175	-0.1251	0.7893
CG7498	CG7498	1623339_at	0.7293	0.0139	0.2657	0.0920	0.2158	0.1633	-0.3045	0.4455	0.4552	0.0423	0.7597	0.0028	-0.1100	0.8494	0.0418	0.9095	0.1518	0.5511
CG15333	CG15333	1623340_at	-0.0657	0.7121	-0.2312	0.1417	-0.3272	0.0940	0.1174	0.7415	0.1472	0.3570	0.0299	0.8716	0.1623	0.7500	0.0587	0.8408	-0.1036	0.6607
Osi17	Osiris	1623341_at	0.1316	0.4511	0.0834	0.5427	0.1576	0.3435	0.0147	0.9922	-0.1611	0.6445	-0.1757	0.5686	-0.0509	0.9538	0.0167	0.9689	0.0675	0.8375
CG8369	CG8369	1623342_at	-0.4348	0.0747	-1.3045	0.0301	-1.1112	0.0007	0.2068	0.5735	0.2815	0.1336	0.0747	0.7076	0.1555	0.9095	-0.5001	0.2924	-0.6556	0.2031
CG4049	R-gene	1623343_at	0.0024	0.9944	0.2398	0.2839	0.2162	0.3846	-0.0640	0.9436	-0.2657	0.3099	-0.2017	0.4036	0.1652	0.8480	0.1446	0.7274	-0.0205	0.9703
CG12856	CG12856	1623344_at	-0.1244	0.5122	0.1180	0.4150	0.1127	0.4773	-0.1814	0.6354	-0.2428	0.1945	-0.0614	0.7668	-0.0402	0.9679	-0.0221	0.9599	0.0181	0.9630
CG33474	CG33474	1623345_at	0.5769	0.1022	-0.1194	0.4979	0.3954	0.1015	0.1097	0.8498	-0.0111	0.9733	-0.1208	0.5631	-0.4284	0.6955	-0.6624	0.1401	-0.2340	0.6328
CG2861	CG2861	1623346_at	0.1477	0.3674	-0.2750	0.0599	0.0196	0.9049	0.1435	0.6280	0.2470	0.0911	0.1036	0.4520	-0.1542	0.7726	-0.2774	0.2145	-0.1233	0.6196
CG11674	CG11674	1623347_at	0.9187	0.3543	-1.5872	0.2012	-1.0715	0.0721	0.0452	0.9558	2.3181	0.0001	2.2729	0.0000	-0.6245	0.9137	-0.1761	0.9583	0.4484	0.8668
CG15058	CG15058	1623348_at	-0.1623	0.4831	0.1542	0.3389	-0.2849	0.1096	0.0145	0.9882	-0.1781	0.4523	-0.1926	0.3575	0.2959	0.5126	0.1006	0.6762	-0.1953	0.3803
---	---	1623349_x_at	-0.1175	0.8389	-0.8847	0.0521	-1.4559	0.0026	-0.0261	0.9803	0.9097	0.0047	0.9357	0.0025	0.5380	0.7927	0.1565	0.9062	-0.3815	0.6896
Lrr47	Leucine-rich repea	1623350_at	-0.0774	0.6778	0.1038	0.4689	0.0141	0.9607	-0.1608	0.6513	0.1483	0.4015	0.3091	0.0494	-0.0219	0.9901	0.3628	0.3287	0.3848	0.3248
CG3857	CG3857	1623351_at	-0.0919	0.6496	0.4505	0.1213	0.5051	0.0085	-0.2899	0.3793	-0.4530	0.0233	-0.1631	0.3399	-0.2242	0.6749	0.1849	0.3941	0.4091	0.0994
CG7016	CG7016	1623352_at	1.0932	0.0452	0.0029	1.0000	0.9926	0.0013	0.5991	0.1643	0.5241	0.0499	-0.0750	0.7981	-0.2343	0.9487	-0.4728	0.7003	-0.2385	0.8694
abs	abstract	1623353_at	-0.2468	0.1361	-0.3441	0.1136	-0.0308	0.8961	0.1453	0.8078	0.0428	0.8951	-0.1025	0.6807	-0.2290	0.5924	-0.0208	0.9460	0.2082	0.2855
---	---	1623354_at	0.0450	0.8519	0.1447	0.5406	-0.0042	0.9881	-0.0520	0.9345	0.0131	0.9632	0.0650	0.7469	0.0801	0.9390	0.1642	0.6385	0.0841	0.8372
v	vermilion	1623355_at	2.2322	0.0056	1.2395	0.1415	2.9839	0.0000	1.0056	0.2316	0.4307	0.4020	-0.5749	0.1975	-0.7983	0.6557	-0.6599	0.3567	0.1384	0.8905
CG11999	CG11999	1623356_at	0.5900	0.0081	0.7819	0.0095	1.1631	0.0007	0.1870	0.5953	0.4621	0.0179	0.2751	0.0855	-0.2700	0.7230	0.8160	0.0383	1.0859	0.0320
CG8616	CG8616	1623357_at	0.1162	0.5969	-0.0941	0.5906	-0.2797	0.1747	-0.2449	0.3793	-0.0015	0.9947	0.2434	0.0844	-0.1499	0.8102	-0.2108	0.4124	-0.0609	0.8575
CG30381	CG30381	1623358_at	0.0421	0.8250	0.4705	0.0790	0.2561	0.0929	-0.4037	0.2397	-0.6964	0.0051	-0.2927	0.1063	0.0008	0.9998	0.1037	0.8094	0.1030	0.7981
stwl	alpha transcript	1623359_at	-0.4384	0.1803	0.2401	0.6936	-0.7926	0.0036	-0.5094	0.2969	-0.2958	0.3072	0.2136	0.4271	0.5103	0.6557	0.4484	0.3259	-0.0619	0.9234
CG32534	CG32534	1623360_at	0.1781	0.3396	0.1476	0.5534	0.1527	0.5248	0.1400	0.7949	0.2388	0.2797	0.0987	0.6648	0.0449	0.9457	0.0142	0.9652	-0.0307	0.9112
---	---	1623361_at	0.1967	0.2931	0.2940	0.3247	0.1354	0.5989	0.0053	0.9943	0.0271	0.9013	0.0218	0.9099	0.1943	0.8033	0.1611	0.6483	-0.0331	0.9427
---	---	1623362_at	0.2075	0.3600	0.1267	0.6269	0.4417	0.1278	-0.1935	0.6864	-0.0518	0.8586	0.1416	0.5098	-0.1706	0.7230	-0.1097	0.6256	0.0610	0.8138
CG6432	CG6432	1623363_at	0.2615	0.8335	-1.4951	0.3801	1.3496	0.0904	2.6186	0.0978	1.0754	0.2507	-1.5432	0.0675	-0.5259	0.9365	-0.7201	0.7788	-0.1941	0.9471
CG4250	CG4250	1623364_at	-1.9885	0.0231	-2.8493	0.0121	-3.0079	0.0002	-0.1308	0.8251	-0.4648	0.0480	-0.3340	0.1028	0.1665	0.9689	-1.4064	0.1925	-1.5730	0.1823
Bgb	Big brother	1623365_at	-0.6317	0.0117	-0.5440	0.0638	-0.5242	0.0707	0.2736	0.5255	-0.0082	0.9809	-0.2817	0.1617	0.1390	0.8940	-0.0034	0.9977	-0.1424	0.7514

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
mRpL19	mitochondrial ribo	1623366_at	0.1353	0.6753	0.0632	0.8088	0.1468	0.5859	0.1011	0.8111	0.4358	0.0176	0.3348	0.0330	-0.0125	0.9952	0.3445	0.5232	0.3570	0.5092
---	---	1623367_at	0.3067	0.1235	-0.0432	0.7905	0.0480	0.7965	0.0455	0.9488	0.2120	0.2878	0.1665	0.3604	-0.1197	0.8521	-0.1165	0.6969	0.0032	0.9946
---	---	1623368_at	0.1176	0.6838	-0.0249	0.8209	0.1837	0.2976	0.2055	0.6642	0.1422	0.5602	-0.0633	0.8039	-0.0398	0.9717	0.0131	0.9796	0.0529	0.8915
CG31796	CG31796	1623369_at	-0.1184	0.6335	-0.0180	0.9101	-0.0001	0.9998	-0.0379	0.9596	0.1156	0.5878	0.1535	0.4004	0.0577	0.9624	0.1277	0.7506	0.0700	0.8797
wek	weckle	1623370_at	0.4581	0.0651	0.4752	0.0159	0.7513	0.0024	0.0200	0.9860	0.0329	0.9293	0.0129	0.9689	-0.0685	0.9506	0.3541	0.2632	0.4226	0.2140
CG11126	CG11126	1623371_at	-0.6463	0.0803	0.1607	0.4748	-0.5558	0.1022	-0.7516	0.3628	-1.0233	0.0365	-0.2716	0.5592	-0.1903	0.8202	-0.1898	0.6093	0.0005	0.9992
CG5873	CG5873	1623372_at	-3.3908	0.0263	-0.8138	0.3965	-3.7581	0.0002	-3.1831	0.0122	-4.3352	0.0003	-1.1521	0.0475	0.0722	0.9928	-1.8056	0.3022	-1.8778	0.3081
CG14806	CG14806	1623373_at	-0.1369	0.4104	0.1368	0.5250	-0.0978	0.7802	0.0104	0.9911	-0.2001	0.3297	-0.2105	0.2467	0.1530	0.8814	0.0207	0.9760	-0.1324	0.7749
Fim	fimbrin	1623374_a_at	1.1111	0.0180	1.3985	0.0108	1.2632	0.0005	0.1356	0.8028	0.3317	0.1256	0.1961	0.3228	0.1276	0.9459	0.5741	0.3000	0.4465	0.4457
CG17081 /// DmirCG17081	CG17081	1623375_at	-1.4086	0.0103	-0.9973	0.1174	-0.8687	0.0057	-0.4680	0.4612	-0.6302	0.0701	-0.1622	0.6474	-0.4997	0.6660	-0.2082	0.6926	0.2915	0.5578
Nmda1	N-methyl-D-aspar	1623376_s_at	0.0165	0.9380	-0.0390	0.8370	-0.4443	0.0282	-0.0038	0.9956	0.2992	0.0735	0.3030	0.0463	0.2915	0.6749	0.2332	0.4126	-0.0583	0.8813
CG14841	CG14841	1623377_at	-0.0242	0.9256	0.1027	0.4368	0.3090	0.0722	0.0316	0.9518	-0.1025	0.5025	-0.1341	0.3085	-0.0854	0.8940	0.0738	0.8114	0.1592	0.5230
Oli	Olig family	1623378_at	-0.0683	0.8748	-0.0131	0.9540	-0.0508	0.7926	-0.0501	0.9672	-0.1402	0.7034	-0.0901	0.8019	-0.0322	0.9848	-0.1046	0.8397	-0.0724	0.8918
---	---	1623379_at	-0.0262	0.9533	0.2446	0.5422	0.4144	0.0384	0.0134	0.9884	-0.3048	0.1610	-0.3182	0.1026	-0.0128	0.9952	-0.0766	0.9259	-0.0638	0.9296
---	---	1623380_at	-0.1209	0.5651	-0.2839	0.2226	-0.6214	0.0050	-0.4210	0.1601	0.3839	0.0409	0.8050	0.0010	0.1351	0.9168	0.3328	0.4598	0.1977	0.6887
---	---	1623381_at	0.1326	0.4141	0.1571	0.3155	0.1262	0.5123	-0.1759	0.5664	-0.1039	0.5366	0.0720	0.6585	-0.0046	0.9974	-0.0054	0.9941	-0.0008	0.9992
---	---	1623382_at	0.1513	0.3825	-0.2506	0.0738	0.1448	0.3572	0.2508	0.4141	0.3041	0.0766	0.0533	0.7781	-0.1558	0.7697	-0.1519	0.5163	0.0039	0.9924
Dhc64C	dynein	1623383_at	0.4138	0.1671	0.2586	0.3250	0.3239	0.3052	0.2414	0.7295	0.7182	0.0292	0.4769	0.0881	0.2427	0.8270	0.6217	0.1622	0.3790	0.4149
ytr	yantar	1623384_at	-0.0236	0.9547	0.3406	0.1175	0.0127	0.9441	-0.0854	0.8815	-0.0998	0.6493	-0.0144	0.9537	0.2343	0.8202	0.2481	0.5827	0.0138	0.9847
---	---	1623385_at	0.0669	0.7368	-0.0740	0.5564	0.1763	0.2664	0.0222	0.9726	0.0150	0.9505	-0.0072	0.9711	-0.2693	0.7095	-0.0690	0.8730	0.2003	0.5284
CG14438 /// DmirCG14438	CG14438	1623386_s_at	-0.1529	0.6920	0.1228	0.6755	-0.2383	0.4975	-0.1845	0.8837	0.2361	0.6198	0.4206	0.2786	0.1744	0.9092	0.4892	0.3580	0.3148	0.5868
Bap55	BRM-associated ¡	1623387_at	0.4501	0.0161	0.0530	0.9095	0.0518	0.7630	-0.0108	0.9863	0.5225	0.0051	0.5332	0.0027	-0.0059	0.9974	0.2221	0.6184	0.2281	0.6114
CG15861	CG15861	1623388_at	0.0782	0.7277	0.1290	0.4895	0.6151	0.0568	0.2466	0.6144	-0.0162	0.9659	-0.2628	0.2315	-0.1635	0.8890	0.1326	0.8174	0.2961	0.5227
Ddr	Discoidin domain	1623389_at	-0.0363	0.8955	-0.0676	0.7851	0.3557	0.0476	0.1279	0.8671	0.1682	0.5638	0.0402	0.9011	-0.1639	0.8215	-0.0392	0.9353	0.1248	0.7138
CG11768	CG11768	1623390_at	-0.0919	0.6554	-0.5279	0.0481	-0.2740	0.1871	-0.0158	0.9883	0.2205	0.3877	0.2364	0.2951	0.0026	0.9976	0.0929	0.6524	0.0903	0.6606
sisA	sisterless A	1623391_at	0.1281	0.5047	0.1737	0.1251	0.1652	0.2875	-0.0914	0.8272	-0.1668	0.3054	-0.0754	0.6507	-0.0675	0.9405	-0.0626	0.8754	0.0049	0.9923
---	---	1623392_at	-0.1909	0.3674	0.1199	0.5251	0.1014	0.5328	-0.1192	0.8671	-0.2227	0.3870	-0.1035	0.6952	-0.0109	0.9938	0.0693	0.8749	0.0802	0.8371
CG3277	CG3277	1623393_a_at	0.2253	0.1223	0.1998	0.3445	0.1764	0.5485	-0.2004	0.7121	-0.4117	0.0957	-0.2113	0.3512	0.0783	0.8861	-0.0751	0.7599	-0.1534	0.4697
CG3536	CG3536	1623394_at	-0.0685	0.7214	0.0088	0.9540	0.2847	0.1417	0.0247	0.9774	-0.1020	0.6772	-0.1268	0.5482	-0.1934	0.7475	0.0765	0.8172	0.2699	0.3008
---	---	1623395_at	-0.0091	0.9748	-0.1545	0.3495	-0.3882	0.0520	0.0607	0.9228	0.3407	0.0741	0.2800	0.0972	0.1523	0.8283	0.2077	0.4895	0.0555	0.8913
CG13875	CG13875	1623396_at	-2.5315	0.0012	-1.9085	0.0194	-2.7254	0.0000	-0.2310	0.6324	-0.4347	0.0687	-0.2037	0.3474	0.3544	0.7956	0.2670	0.6762	-0.0874	0.9121
CG40442	CG40442	1623397_at	0.2647	0.3101	-0.0472	0.8147	0.2681	0.2223	0.2624	0.5067	0.4536	0.0369	0.1911	0.3159	-0.1948	0.7997	-0.0560	0.9085	0.1388	0.6974
CG4830	CG4830	1623398_at	0.1546	0.6027	0.1368	0.3683	0.4929	0.0507	0.1354	0.8544	-0.1425	0.6310	-0.2779	0.2450	-0.0977	0.9457	-0.0636	0.9265	0.0341	0.9564
---	---	1623399_at	0.1879	0.3452	0.1088	0.5751	0.2392	0.1375	0.0346	0.9640	-0.0778	0.7470	-0.1124	0.5794	-0.1572	0.7768	-0.2291	0.3269	-0.0719	0.8073
---	---	1623400_at	0.2772	0.3043	0.0567	0.7771	0.0247	0.8982	0.1807	0.6972	0.0187	0.9531	-0.1620	0.4179	0.1146	0.9076	-0.1965	0.6007	-0.3111	0.3865
CG8092	CG8092	1623401_a_at	0.8450	0.1175	1.3374	0.0223	0.5586	0.0262	-0.5277	0.1627	0.1695	0.4795	0.6971	0.0053	0.3719	0.8235	0.6079	0.3716	0.2360	0.7724
Arc-p34	Arc-p34	1623402_at	0.1211	0.5803	-0.4744	0.1579	-0.6387	0.0054	0.1114	0.8543	0.8849	0.0025	0.7735	0.0027	0.3670	0.6762	0.3976	0.2603	0.0306	0.9525
Cklalpha	Casein Kinase 1	1623403_s_at	0.6790	0.1170	0.4790	0.0159	-0.5709	0.2587	-0.2650	0.5453	0.6177	0.0144	0.8827	0.0018	0.7936	0.7293	0.4221	0.7043	-0.3715	0.7437
CG16890	CG16890	1623404_at	0.0549	0.7717	0.1598	0.4567	0.2692	0.1180	0.0564	0.9502	0.0039	0.9913	-0.0525	0.8571	-0.1282	0.8427	0.0967	0.7625	0.2249	0.4059
pav	Pavarotti KLP	1623405_at	1.4456	0.0035	1.0845	0.0226	0.9636	0.0008	-0.0375	0.9819	0.9892	0.0199	1.0267	0.0104	0.1827	0.7633	0.8243	0.0204	0.6416	0.0469
---	---	1623406_at	-0.1704	0.4164	0.0913	0.7263	0.1699	0.3653	-0.0926	0.8578	-0.2343	0.2074	-0.1417	0.4166	0.0126	0.9929	0.2234	0.5214	0.2108	0.5525
CG10298	CG10298	1623407_at	0.0219	0.9079	0.0656	0.7577	-0.0345	0.8308	-0.1605	0.6869	-0.1568	0.4113	0.0037	0.9874	0.1046	0.8439	0.0692	0.8045	-0.0354	0.9065
CG2947 /// CG32789 /// Ds	anon-fast-evolving	1623408_s_at	0.4795	0.1331	0.6235	0.0675	0.2448	0.1580	-0.2076	0.6010	0.2061	0.3073	0.4137	0.0284	0.1351	0.8889	0.3185	0.3733	0.1834	0.6395
---	---	1623409_at	0.0271	0.8650	0.1883	0.3131	0.1201	0.5141	-0.1828	0.5680	-0.2212	0.1748	-0.0384	0.8390	-0.0970	0.8806	-0.0591	0.8642	0.0379	0.9110
Pxd	Peroxidase	1623410_at	-0.1551	0.4840	-0.8358	0.0151	-0.9367	0.0579	0.1928	0.9196	0.9491	0.1039	0.7563	0.1468	0.0500	0.9521	-0.1062	0.7068	-0.1562	0.5549
lola	longitudinals abse	1623411_at	-0.1063	0.7758	0.4765	0.2975	-0.2791	0.1604	-0.0975	0.9236	0.0142	0.9763	0.1117	0.7348	0.6721	0.5665	0.6904	0.1786	0.0184	0.9838
CG17150 /// DmirCG17150	CG17150	1623412_a_at	0.0962	0.5071	-0.0240	0.8883	0.0477	0.8524	-0.0263	0.9774	-0.0246	0.9404	0.0016	0.9956	0.0033	0.9976	-0.0455	0.9027	-0.0488	0.8844
---	---	1623413_at	0.1957	0.2887	0.0522	0.5772	0.1769	0.4047	0.0712	0.9074	0.1280	0.5462	0.0568	0.7979	-0.0198	0.9831	-0.0844	0.7346	-0.0646	0.8044
---	---	1623414_at	0.0328	0.8810	0.0471	0.6886	0.4695	0.0499	0.0163	0.9833	-0.1019	0.6009	-0.1182	0.4870	-0.1982	0.7893	0.0028	0.9977	0.2010	0.5476
dl	dorsal	1623415_at	-1.7963	0.0038	-2.6449	0.0270	-3.3746	0.0000	-0.2784	0.4141	0.9268	0.0011	1.2052	0.0002	0.5091	0.7893	0.1267	0.9204	-0.3824	0.6657
CG33265	CG33265	1623416_at	0.0611	0.8243	-0.0888	0.4951	0.0888	0.6558	0.2233	0.5744	0.2965	0.1440	0.0733	0.7384	0.0473	0.9589	0.1157	0.6939	0.0684	0.8364
---	---	1623417_at	0.0428	0.7982	0.0388	0.8647	0.1689	0.2829	-0.0767	0.9036	-0.0776	0.7511	-0.0009	0.9975	-0.1972	0.7070	-0.0981	0.6876	0.0991	0.6800
Cad96Ca	Cad96Ca	1623418_at	-2.3269	0.0006	-1.9816	0.0626	-2.8198	0.0000	-0.3575	0.5070	-0.3397	0.2347	0.0178	0.9599	0.4512	0.7707	-0.0283	0.9820	-0.4795	0.4822
CG31358	CG31358	1623419_s_at	0.1977	0.3571	0.1070	0.4268	0.0918	0.5626	-0.0927	0.8074	-0.0656	0.7075	0.0271	0.8801	0.0542	0.9599	-0.0719	0.8680	-0.1262	0.7154
---	---	1623420_at	0.0738	0.6187	-0.0828	0.4953	0.0189	0.9294	0.1959	0.6958	0.1925	0.4201	-0.0033	0.9907	-0.0012	0.9994	0.0275	0.9229	0.0287	0.9089
CG6347	CG6347	1623421_at	-0.2827	0.1778	-0.0496	0.8100	-0.1189	0.7206	0.0861	0.9242	0.1117	0.7234	0.0257	0.9393	-0.1898	0.7746	0.1250	0.7002	0.3148	0.2869
CG31908	CG31908	1623422_a_at	-0.2024	0.2692	0.3685	0.1708	0.3346	0.0380	0.0050	0.9946	-0.1607	0.3030	-0.1656	0.2313	-0.0526	0.9589	0.3263	0.2442	0.3788	0.2110
---	---	1623423_at	0.1872	0.2789	-0.0029	0.9816	0.1404	0.4821	-0.0392	0.9491	0.0549	0.7940	0.0941	0.5828	-0.0855	0.8692	0.0673	0.7985	0.1529	0.4776
babo	Activin A receptor	1623424_a_at	0.8098	0.0116	0.4124	0.0755	0.1069	0.6287	-0.3998	0.4529	0.5235	0.0729	0.9233	0.0042	-0.1193	0.8692	0.1206	0.7123	0.2398	0.4114
Buffy	48A-E Bcl-2-like	1623425_at	0.2634	0.2248	-0.3478	0.1917	-0.4142	0.0501	-0.3062	0.5039	-0.3843	0.1130	-0.0781	0.7714	-0.2314	0.7953	-0.9733	0.0330	-0.7419	0.0836
CG31633	CG31633	1623426_at	-0.6659	0.0081	-0.2762	0.1722	-0.3892	0.0629	-0.0255	0.9774	0.0483	0.8678	0.0738	0.7583	0.1252	0.8400	0.4518	0.0887	0.3267	0.2189
CG7179	CG7179	1623427_at	0.6742	0.1898	0.0331	0.8437	0.0263	0.9308	-0.0068	0.9937	0.1045	0.5997	0.1114	0.5306	0.1253	0.9549	-0.5381	0.4021	-0.6634	0.3171
CG14933 /// DyakCG14933	CG14933	1623428_at	2.0018	0.0028	0.7750	0.3258	1.3633	0.0111	0.5444	0.6034	0.0163	0.9832	-0.5281	0.2669	-0.0138	0.9970	-1.0810	0.2168	-1.0672	0.2553
CG11073	CG11073	1623429_at	-1.9956	0.0022	-1.7863	0.0607	-1.5383	0.0001	0.1801	0.8366	-0.1242	0.7510	-0.3043	0.3089	-0.1073	0.9598	-0.0530	0.9583	0.0543	0.9491
Phm	Peptidyl gycine alp	1623430_at	-0.1271	0.4968	-0.8718	0.0171	-0.6573	0.0263	0.2260	0.7045	0.5654	0.0449	0.3394	0.1648	0.1876	0.7683	0.0214	0.9620	-0.1662	0.5615
CG13423	CG13423	1623431_at	0.0140	0.9470	0.7450	0.2579	0.3501	0.0704	-0.3034	0.3837	-0.2577	0.1881	0.0457	0.8407	0.0608	0.9775	0.3492	0.5084	0.2885	0.5995
plexB	Plexin-B	1623432_s_at	-0.3220	0.3382	0.3554	0.5141	0.1500	0.4838	-0.4062	0.3265	-0.4941	0.0452	-0.0879	0.7329	-0.1024	0.9677	0.3021	0.6835	0.4046	0.5657
CG30369	CG30369	1623433_at	0.0961	0.7141	0.0130	0.9273	-0.0027	0.9923	0.0474	0.9558	0.1106	0.6749	0.0632	0.8078	-0.1776	0.8439	-0.1400	0.7489	0.0375	0.9431
CG3702	CG3702	1623434_at	0.1995	0.2789	-0.1381	0.7722	0.1281	0.5474	0.0588	0.9004	0.8134	0.0008	0.7546	0.0006	-0.1772	0.8882	0.4785	0.3006	0.6558	0.1911

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1623435_at	-0.0886	0.7107	0.0413	0.8234	-0.0675	0.7907	-0.1878	0.5735	-0.1688	0.3303	0.0190	0.9276	-0.2460	0.7893	-0.1283	0.7943	0.1177	0.8056
bun	short-sighted	1623436_a_at	0.0823	0.7848	0.3395	0.4389	0.1936	0.2764	-0.1648	0.6872	-0.5278	0.0132	-0.3631	0.0393	-0.0769	0.9665	-0.2976	0.5604	-0.2207	0.6816
CG9948	CG9948	1623437_at	0.2377	0.4150	-0.3689	0.1666	-0.7239	0.0048	-0.0382	0.9744	0.4811	0.0951	0.5193	0.0488	0.2581	0.7550	-0.0985	0.8340	-0.3567	0.3205
CG9505	CG9505	1623438_at	0.4237	0.4338	0.5392	0.4035	1.0305	0.0139	-0.1631	0.8874	-0.2171	0.6123	-0.0540	0.9086	-0.5367	0.8202	-0.1076	0.9451	0.4290	0.6877
CG17744	CG17744	1623439_at	0.1480	0.3524	0.0817	0.6611	-0.1094	0.5180	-0.2021	0.4734	-0.0195	0.9270	0.1827	0.1771	0.0180	0.9894	-0.0898	0.8061	-0.1079	0.7434
CG31848	CG31848	1623440_at	0.1913	0.4409	-0.0091	0.9551	0.1851	0.3871	0.0431	0.9500	0.0401	0.8736	-0.0030	0.9901	-0.2282	0.7464	-0.1296	0.7000	0.0986	0.7787
ph-p	Polyhomeotic	1623441_at	-0.3200	0.2475	-0.1594	0.6840	-0.2573	0.2913	0.0976	0.9116	0.0608	0.8696	-0.0368	0.9137	0.1993	0.8869	0.2639	0.6425	0.0646	0.9297
CG14088	CG14088	1623442_at	0.0401	0.8226	0.0792	0.4737	0.4207	0.0304	0.1067	0.8280	-0.0045	0.9879	-0.1112	0.5482	-0.0075	0.9946	0.0924	0.7719	0.0999	0.7410
---	---	1623443_at	-0.0061	0.9757	0.0986	0.4402	0.3442	0.0923	-0.0703	0.8830	-0.2256	0.1634	-0.1553	0.2921	-0.0476	0.9514	0.0350	0.9242	0.0826	0.7608
rn	roughened eye	1623444_at	-0.1847	0.2679	0.1782	0.4047	-0.0793	0.7631	0.0047	0.9956	-0.1863	0.4848	-0.1910	0.4229	0.1005	0.8882	0.0648	0.8626	-0.0357	0.9231
CG14095	CG14095	1623445_at	0.1193	0.7145	0.1857	0.3714	0.5044	0.0247	0.1423	0.8087	0.1172	0.6577	-0.0251	0.9303	-0.1478	0.8378	0.1338	0.6857	0.2817	0.3552
---	---	1623446_a_at	-0.5383	0.0800	0.0304	0.8425	-0.1196	0.5175	0.0702	0.9346	-0.4762	0.0636	-0.5463	0.0240	0.0677	0.9517	0.0415	0.9402	-0.0262	0.9568
CG12201	CG12201	1623447_a_at	0.0633	0.7235	0.3483	0.0412	0.0164	0.9330	-0.1601	0.6615	-0.1210	0.5137	0.0391	0.8472	0.1628	0.7506	0.1741	0.4246	0.0113	0.9737
CG5196	CG5196	1623448_at	0.0023	0.9915	0.4664	0.0341	0.8779	0.0036	0.0853	0.8794	-0.5258	0.0140	-0.6111	0.0043	-0.2480	0.7215	0.1132	0.7424	0.3612	0.2331
CG7840	CG7840	1623449_at	0.8248	0.0245	0.4276	0.1238	0.7508	0.0031	0.0610	0.9300	0.6689	0.0063	0.6079	0.0058	-0.2142	0.8628	0.2736	0.6082	0.4878	0.3424
Gr59e	Gustatory recepto	1623450_at	0.0869	0.6908	0.2356	0.1254	0.3566	0.1093	0.0457	0.9647	-0.1875	0.5157	-0.2332	0.3509	0.0160	0.9894	0.1838	0.4729	0.1678	0.5257
Ank2	Ank2	1623451_at	0.2207	0.1327	0.2115	0.3102	0.0053	0.9826	-0.2585	0.5597	-0.0104	0.9764	0.2481	0.2240	-0.0110	0.9914	-0.0321	0.9275	-0.0211	0.9462
CG13068 /// DyakCG13068	CG13068	1623452_at	0.4343	0.1433	0.2740	0.1335	0.0193	0.9576	-0.3490	0.5363	0.1304	0.7014	0.4794	0.0714	-0.0873	0.9108	-0.0281	0.9501	0.0591	0.8743
CG10447	CG10447	1623453_at	0.2403	0.2648	0.0277	0.8360	-0.1634	0.6393	-0.0123	0.9883	0.1823	0.3701	0.1946	0.2794	0.1523	0.8763	-0.0298	0.9622	-0.1822	0.6602
---	---	1623454_at	-0.0272	0.9014	-0.1010	0.4204	-0.2166	0.1876	-0.0686	0.8837	0.1124	0.5058	0.1810	0.2052	0.0415	0.9589	0.0680	0.8220	0.0265	0.9347
Imp	KH-domain protei	1623455_s_at	-0.2943	0.4455	0.5869	0.1074	0.3989	0.2175	-0.2486	0.5735	-0.5203	0.0292	-0.2717	0.1785	-0.1114	0.9689	0.3821	0.6394	0.4936	0.5334
CG12477	CG12477	1623456_at	0.1744	0.4463	0.1506	0.2158	0.1001	0.5134	-0.0596	0.9413	-0.0470	0.8792	0.0127	0.9658	0.0657	0.9411	0.0337	0.9402	-0.0320	0.9342
---	---	1623457_at	0.2030	0.2065	0.3556	0.1677	0.3775	0.0283	-0.0466	0.9311	-0.1862	0.2419	-0.1396	0.3371	-0.0789	0.8940	0.0500	0.8731	0.1289	0.5811
CG14899 /// DbuzCG14899	CG14899	1623458_at	-0.2777	0.3028	0.5126	0.0600	1.2117	0.0009	-0.0279	0.9757	-0.5110	0.0300	-0.4831	0.0240	-0.6047	0.5434	0.4051	0.3739	1.0098	0.0679
Spn4	neuroserpin	1623459_at	0.9457	0.0097	0.3582	0.0314	0.8050	0.0190	0.4355	0.2863	0.8989	0.0035	0.4634	0.0374	0.0190	0.9914	0.3230	0.3878	0.3039	0.4356
Aats-leu	Leucyl-tRNA synth	1623460_at	0.1423	0.5559	0.5858	0.1481	0.7692	0.0046	-0.0743	0.9117	-0.2666	0.2091	-0.1922	0.3228	-0.1333	0.9226	0.3547	0.4344	0.4880	0.2964
CG40239	CG40239	1623461_at	-0.1542	0.4396	-0.1023	0.6095	0.1042	0.5701	0.1888	0.7070	0.2175	0.3525	0.0287	0.9189	0.0126	0.9914	0.0803	0.8153	0.0676	0.8412
cato	cousin of atonal	1623462_at	0.0989	0.5623	0.2254	0.2521	0.1458	0.4256	-0.1024	0.8233	-0.1060	0.5831	-0.0036	0.9873	0.1945	0.7464	0.0804	0.8040	-0.1141	0.6830
tsr	cofilin	1623463_at	-0.3927	0.0227	-0.1832	0.1100	-0.2953	0.0762	-0.0393	0.9377	0.0249	0.9026	0.0641	0.6769	0.0976	0.8744	0.3026	0.1975	0.2051	0.4067
CG13217	CG13217	1623464_at	0.1361	0.4923	0.0089	0.9658	0.1705	0.2880	0.1137	0.8350	-0.0650	0.7977	-0.1787	0.3457	0.0275	0.9742	0.0135	0.9694	-0.0140	0.9650
CG6138	CG6138	1623465_at	0.0317	0.8606	0.0360	0.7699	0.0986	0.5076	-0.0353	0.9647	-0.0890	0.7157	-0.0538	0.8222	0.0120	0.9913	-0.0223	0.9521	-0.0343	0.9152
bab1	bric-a-brac	1623466_at	-1.3142	0.0080	-1.3448	0.0142	-1.9595	0.0002	-0.2326	0.6327	0.1625	0.5216	0.3951	0.0670	-0.1992	0.8270	-0.1534	0.7245	0.0458	0.9313
CG8087	CG8087	1623467_at	0.1358	0.5328	0.0785	0.5781	0.1384	0.5764	-0.0347	0.9683	-0.0783	0.7753	-0.0436	0.8704	-0.0542	0.9689	-0.0188	0.9768	0.0354	0.9445
CG2213	CG2213	1623468_at	-0.4148	0.3512	-0.6266	0.1009	-0.7414	0.0251	-0.1238	0.8913	0.2415	0.4446	0.3653	0.1783	-0.2092	0.8650	-0.3202	0.5283	-0.1110	0.8636
CG18331	CG18331	1623469_s_at	-0.0293	0.8836	0.0243	0.9166	0.1167	0.5599	0.1075	0.8028	0.0140	0.9562	-0.0935	0.5874	-0.0704	0.9460	0.0653	0.8857	0.1357	0.6990
Las	Lipoic acid syntha	1623470_at	-0.1756	0.4898	0.0656	0.5451	0.3023	0.0511	0.0677	0.8856	-0.4790	0.0093	-0.5467	0.0031	-0.2032	0.7979	-0.1958	0.5767	0.0074	0.9889
---	---	1623471_s_at	-0.2718	0.6973	0.6277	0.1937	-0.7387	0.1129	-0.3893	0.4568	-0.3063	0.2837	0.0830	0.7943	1.0295	0.7062	0.6743	0.5749	-0.3553	0.7956
wus	wurst	1623472_at	-0.7523	0.1529	0.2412	0.3309	-0.0961	0.7802	-0.3859	0.6854	-0.7865	0.0773	-0.4006	0.3205	-0.2764	0.8609	0.1351	0.8830	0.4115	0.5399
shark	Tyrosine kinase 7	1623473_at	-0.3216	0.2593	-0.1620	0.4656	-0.2891	0.0671	-0.0590	0.9236	-0.1392	0.4731	-0.0802	0.6798	0.1392	0.8888	0.1319	0.7731	-0.0073	0.9913
CG15479	CG15479	1623474_at	-0.2531	0.4916	0.0610	0.7141	-0.1452	0.4760	-0.2321	0.6175	-0.2711	0.2434	-0.0389	0.8882	-0.1356	0.8870	-0.0879	0.8594	0.0477	0.9231
Rm62	Lighten up	1623475_s_at	-0.4996	0.0993	0.2472	0.4734	0.0560	0.7861	-0.0332	0.9672	-0.6596	0.0080	-0.6264	0.0060	0.2422	0.8228	0.1308	0.8248	-0.1114	0.8472
---	---	1623476_at	0.1587	0.2911	0.0400	0.7088	0.2626	0.1638	-0.0164	0.9777	0.1173	0.4560	0.1336	0.3336	0.0199	0.9848	0.1071	0.6834	0.0872	0.7484
---	---	1623477_at	0.1176	0.3891	-0.0071	0.9506	-0.0174	0.9351	-0.0154	0.9860	0.2567	0.2029	0.2722	0.1307	0.0258	0.9775	0.0658	0.8153	0.0400	0.8932
CG16752 /// DsimCG16752	CG16752	1623478_at	-0.4356	0.2273	2.6193	0.0074	0.1959	0.6231	-2.0201	0.0174	-2.8473	0.0004	-0.8273	0.0426	0.3246	0.8023	0.2191	0.7257	-0.1055	0.8874
Rbf	Retinoblastoma-fa	1623479_at	0.1712	0.6034	0.5788	0.0925	0.2413	0.1402	0.0221	0.9817	0.2160	0.3520	0.1940	0.3544	0.3021	0.8001	0.7006	0.1497	0.3985	0.4356
CG5047	CG5047	1623480_at	0.1539	0.4699	0.4299	0.2790	0.2966	0.1594	0.0956	0.8908	-0.2538	0.2797	-0.3494	0.0940	-0.0436	0.9816	-0.1833	0.7058	-0.1397	0.7839
CG9066 /// DyakCG9066	CG9066	1623481_at	-0.0722	0.7532	-0.6050	0.0944	-0.5587	0.0076	-0.2138	0.5539	0.1422	0.4654	0.3560	0.0394	-0.1932	0.8331	-0.2807	0.4696	-0.0875	0.8618
CG9072	CG9072	1623482_at	0.2263	0.2167	0.0371	0.7525	-0.0005	0.9982	0.0283	0.9689	-0.0262	0.9215	-0.0545	0.7976	0.1879	0.8192	0.0552	0.9157	-0.1327	0.7307
CG17283	CG17283	1623483_at	0.0808	0.6211	0.0440	0.7543	0.2002	0.3844	0.0433	0.9423	-0.0238	0.9211	-0.0670	0.7108	-0.0184	0.9893	0.0394	0.9321	0.0579	0.8837
vg	vestigial	1623484_at	5.7693	0.0024	3.5803	0.0018	6.0361	0.0000	3.0343	0.0417	2.2476	0.0150	-0.7867	0.2978	0.6499	0.7423	0.0579	0.9663	-0.5920	0.4922
Xpac	Xeroderma pigme	1623485_at	0.1091	0.5934	0.3676	0.1470	0.3797	0.0333	-0.1167	0.8247	-0.0020	0.9945	0.1147	0.5712	-0.1489	0.8471	0.1751	0.6093	0.3240	0.3227
CG7900	CG7900	1623486_at	1.5986	0.0011	2.0579	0.0181	2.5304	0.0000	0.0018	0.9988	0.3310	0.1982	0.3292	0.1528	-0.4330	0.6749	0.7648	0.0825	1.1978	0.0385
dmt	dalmatian	1623487_at	-0.1265	0.6756	0.2789	0.0775	0.0482	0.8492	-0.2972	0.6391	-0.4345	0.1595	-0.1373	0.6704	-0.0784	0.8956	0.0349	0.9196	0.1133	0.6375
CG40164	CG40164	1623488_at	0.0039	0.9860	0.0366	0.7193	0.1755	0.2817	0.0423	0.9465	0.0121	0.9648	-0.0303	0.8864	-0.0509	0.9460	0.0776	0.7842	0.1285	0.6005
---	---	1623489_at	-0.0245	0.9374	-0.2965	0.3655	-0.6204	0.0106	-0.0351	0.9777	0.3329	0.2844	0.3680	0.1814	0.1044	0.8395	-0.0150	0.9652	-0.1195	0.5995
Ten-m	tenascin major	1623490_at	-1.4685	0.0156	-2.3275	0.0205	-2.1535	0.0001	-0.4985	0.2011	0.0211	0.9518	0.5197	0.0216	-0.7021	0.7363	-1.0034	0.2424	-0.3013	0.7738
CG15155	CG15155	1623491_at	-1.1289	0.2871	-0.8542	0.0393	-0.6122	0.0549	0.7242	0.7663	0.2955	0.8195	-0.4288	0.6943	0.2845	0.7893	0.2733	0.5636	-0.0111	0.9874
CG17010	CG17010	1623492_a_at	0.0030	0.9886	-0.0632	0.7717	-0.0619	0.7344	0.0118	0.9883	-0.0813	0.7096	-0.0931	0.6290	-0.0370	0.9816	-0.0450	0.9353	-0.0080	0.9885
CG12717	CG12717	1623493_at	-0.4493	0.3158	1.0546	0.0523	0.6921	0.0380	-0.4289	0.6166	-1.1024	0.0182	-0.6735	0.0798	0.0349	0.9862	0.4618	0.3018	0.4269	0.3671
ImpE3	Ecdysone-inducib	1623494_at	-0.6225	0.0157	-0.8413	0.0447	-0.5676	0.0098	0.2408	0.6010	0.0956	0.7234	-0.1451	0.5192	-0.1692	0.8122	-0.2268	0.4418	-0.0576	0.8874
l-2	inhibitor-2	1623495_at	0.4718	0.0229	-0.0331	0.7386	-0.0683	0.6454	-0.1748	0.5539	-0.0851	0.6132	0.0897	0.5497	-0.1035	0.8767	-0.4296	0.1052	-0.3261	0.2245
kz	kurz	1623496_at	0.6411	0.0963	0.1060	0.8079	0.5626	0.1422	0.3788	0.6130	0.9873	0.0168	0.6085	0.0724	0.0008	0.9999	0.4136	0.5252	0.4129	0.5332
HLHm3	split locus enhanc	1623497_at	-0.1769	0.9082	0.3669	0.3107	-0.4399	0.0986	-0.4184	0.8609	-0.7236	0.4119	-0.3053	0.7403	0.2358	0.9503	-0.3929	0.7769	-0.6287	0.6043
CG5857	CG5857	1623498_at	0.3937	0.0544	0.0843	0.8461	0.0256	0.9135	0.0038	0.9956	0.7031	0.0063	0.6993	0.0039	-0.0890	0.9635	0.1963	0.7551	0.2854	0.6166
CG12341	CG12341	1623499_at	0.4639	0.0236	0.2937	0.4266	0.1611	0.3769	0.0463	0.9445	0.6342	0.0058	0.5879	0.0049	0.2536	0.7644	0.3444	0.3223	0.0909	0.8418
CG13622	CG13622	1623500_at	0.3069	0.2358	0.2800	0.2734	0.3230	0.1254	0.0747	0.8875	0.0100	0.9711	-0.0647	0.7357	0.0689	0.9558	-0.0059	0.9941	-0.0748	0.8788
eIF4E-7	eIF4E-7	1623501_at	-0.0465	0.7883	0.0682	0.5443	-0.0568	0.7704	-0.1052	0.7425	-0.1507	0.2888	-0.0455	0.7700	0.0817	0.8888	0.0569	0.8500	-0.0248	0.9360
CG31910	CG31910	1623502_at	0.1824	0.4748	0.1450	0.3952	0.0183	0.9393	-0.2325	0.4950	-0.2532	0.1595	-0.0207	0.9253	-0.0296	0.9769	-0.1101	0.6839	-0.0806	0.7780
sda	slamdance	1623503_at	-2.5268	0.0043	-2.7108	0.0150	-2.8602	0.0000	0.1046	0.9116	0.7669	0.0184	0.6623	0.0213	0.2036	0.9296	0.7882	0.2740	0.5846	0.4465

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1623504_at	0.0650	0.7880	0.1062	0.5385	0.2327	0.2169	0.0025	0.9978	-0.1655	0.4881	-0.1680	0.4324	0.0272	0.9816	0.0466	0.9086	0.0194	0.9600
CR11700 /// Ubi-p5E	CR11700 /// Ubiqu	1623505_s_at	-0.1780	0.3269	0.1038	0.5372	-0.0216	0.9253	0.0186	0.9821	-0.1360	0.5124	-0.1546	0.3975	0.1199	0.8940	0.1259	0.7506	0.0060	0.9921
Pp4-19C	Protein phosphata	1623506_s_at	-0.0921	0.6945	-0.0374	0.7681	0.0489	0.7981	-0.1025	0.8087	-0.0723	0.7103	0.0301	0.8802	-0.1373	0.8971	0.0741	0.9018	0.2114	0.6191
CG33136	CG33136	1623507_at	0.0281	0.8695	-0.0510	0.7345	0.1284	0.4457	0.0790	0.8500	0.0860	0.6071	0.0070	0.9709	-0.0856	0.8689	-0.1001	0.6536	-0.0145	0.9614
CG30466	CG30466	1623508_at	-0.1994	0.5366	-0.7147	0.0245	-0.6226	0.0207	0.1978	0.7764	0.7274	0.0208	0.5296	0.0474	-0.1905	0.8479	0.0451	0.9462	0.2356	0.5958
CG15252	CG15252	1623509_at	-0.0234	0.9172	-0.0902	0.7495	0.0004	0.9989	0.0673	0.9518	0.1694	0.6244	0.1021	0.7653	0.1249	0.8906	0.1878	0.6073	0.0630	0.8932
CG7970	CG7970	1623510_at	0.9861	0.0112	0.7570	0.0185	0.8129	0.0008	0.0174	0.9838	0.0176	0.9518	0.0002	0.9995	-0.1321	0.8400	-0.2568	0.3297	-0.1247	0.6736
---	---	1623511_at	0.1860	0.2763	0.0207	0.8387	0.3105	0.0688	0.0774	0.8663	-0.0269	0.9044	-0.1043	0.5040	0.1090	0.8235	0.1691	0.3997	0.0601	0.8110
CG13291	CG13291	1623512_at	-0.0569	0.7420	-0.0294	0.8778	-0.2017	0.2100	-0.1219	0.7161	-0.0843	0.6182	0.0376	0.8293	-0.0631	0.9449	-0.2267	0.4089	-0.1635	0.5789
Upf1	Upf1	1623513_at	0.4866	0.6399	1.0701	0.2145	1.3503	0.0004	0.4865	0.4337	0.2846	0.4196	-0.2019	0.5469	0.2699	0.9555	0.8380	0.5727	0.5681	0.7228
jdp	jdp	1623514_a_at	-0.2308	0.3035	-0.2111	0.4655	-0.1816	0.3446	0.1179	0.8987	-0.0307	0.9456	-0.1485	0.6318	0.0239	0.9848	-0.0797	0.8343	-0.1035	0.7520
---	---	1623515_at	0.0870	0.5398	0.0016	0.9907	0.1079	0.5399	0.1141	0.7492	0.2275	0.1414	0.1133	0.4377	-0.0342	0.9506	-0.0435	0.8450	-0.0092	0.9709
Ent1	Equilibrative nucle	1623516_s_at	1.2602	0.0036	0.6734	0.0538	0.7955	0.0125	-0.2744	0.6247	-0.1102	0.7364	0.1642	0.5470	-0.3546	0.7230	-0.7327	0.0972	-0.3781	0.3914
CG4962	CG4962	1623517_at	0.2326	0.2450	0.0595	0.6243	0.5938	0.0035	0.2111	0.6463	0.1312	0.5885	-0.0799	0.7393	-0.2353	0.7149	-0.0229	0.9590	0.2124	0.4429
---	---	1623518_s_at	0.2008	0.2430	-0.2321	0.6425	-0.0102	0.9712	-0.1060	0.8137	-0.4882	0.0137	-0.3822	0.0240	-0.4556	0.7370	-0.9942	0.0945	-0.5386	0.3638
esc	extra sex combs	1623519_at	0.5989	0.3001	-0.0678	0.9096	-0.2896	0.4658	-0.6152	0.2117	0.6705	0.0311	1.2857	0.0011	-0.1902	0.9452	-0.1129	0.9328	0.0773	0.9474
CG10336	CG10336	1623520_a_at	0.3045	0.0620	-0.0531	0.6117	-0.0009	0.9982	-0.2830	0.5174	0.0581	0.8460	0.3411	0.0976	-0.1442	0.9132	-0.1758	0.7475	-0.0316	0.9630
CG11909	CG11909	1623521_at	3.4405	0.0028	2.0099	0.0804	3.5704	0.0001	1.4770	0.1848	-0.0151	0.9884	-1.4922	0.0215	-0.0816	0.9862	-1.6217	0.1364	-1.5400	0.1823
CG11668	CG11668	1623522_at	-0.1441	0.6167	0.0327	0.7668	0.2662	0.3417	0.1112	0.8830	-0.2227	0.4002	-0.3340	0.1458	-0.1666	0.7707	-0.0739	0.8153	0.0927	0.7417
CG5810	CG5810	1623523_at	-1.1349	0.0277	0.1598	0.7596	-1.9514	0.0012	-1.8873	0.0316	-2.0857	0.0019	-0.1984	0.6953	0.2254	0.9229	-0.9083	0.2217	-1.1337	0.1651
CG7031	CG7031	1623524_at	0.2214	0.3747	0.2183	0.4399	0.3413	0.0423	0.1938	0.7321	0.1016	0.7312	-0.0922	0.7340	0.0348	0.9717	0.0810	0.7922	0.0462	0.8915
Pcl	Polycomblike	1623525_at	-0.1776	0.3297	0.0561	0.8919	-0.1655	0.4346	-0.2951	0.4795	0.1189	0.6339	0.4140	0.0443	-0.0147	0.9916	0.3179	0.3158	0.3326	0.3175
CG32511	CG32511	1623526_at	0.2193	0.1408	-0.1115	0.5600	-0.0964	0.6316	0.0631	0.9387	0.1555	0.5480	0.0924	0.7183	0.0852	0.9076	-0.0516	0.8939	-0.1368	0.6263
CG40122	CG40122	1623527_at	0.3769	0.1476	0.2820	0.1233	0.3656	0.0517	0.1561	0.8128	0.0422	0.9070	-0.1139	0.6752	0.0649	0.9467	0.0032	0.9961	-0.0617	0.8774
CG31784	CG31784	1623528_a_at	0.0903	0.7560	-0.0563	0.7384	-0.1213	0.6423	0.2431	0.5675	0.3804	0.0815	0.1373	0.5140	0.0740	0.9590	0.0332	0.9617	-0.0408	0.9435
Or19a /// Or19b	Odorant receptor	1623529_s_at	0.1242	0.7186	0.0715	0.5091	0.3684	0.0629	-0.0560	0.9343	-0.1593	0.4418	-0.1033	0.6083	0.0833	0.9238	0.0374	0.9353	-0.0459	0.9086
---	---	1623530_at	0.2771	0.1036	0.1903	0.3627	0.1604	0.3036	0.0474	0.9514	-0.0743	0.7733	-0.1216	0.5665	0.1372	0.7324	-0.0280	0.9167	-0.1652	0.3518
Mst89B	Mst89B	1623531_at	-0.0199	0.9144	0.2479	0.1727	0.1914	0.3384	-0.1278	0.7753	-0.1619	0.4014	-0.0340	0.8784	-0.2150	0.7628	0.0832	0.8341	0.2981	0.3255
---	---	1623532_at	0.3391	0.6930	0.0146	0.9010	-0.4142	0.1265	-0.1019	0.9482	-0.1233	0.8229	-0.0215	0.9689	0.2618	0.9545	-0.5125	0.7461	-0.7743	0.5906
---	---	1623533_at	0.0124	0.9621	0.0258	0.7982	-0.3451	0.0380	0.0707	0.8946	0.0241	0.9226	-0.0466	0.8168	0.2403	0.6749	0.0408	0.9115	-0.1995	0.4114
---	---	1623534_at	0.3532	0.0535	0.1048	0.4315	0.4654	0.1009	-0.0523	0.9252	0.0989	0.5863	0.1512	0.3214	-0.3052	0.6927	-0.0649	0.8879	0.2403	0.4642
CG7164	CG7164	1623535_at	-0.0273	0.8802	-0.0044	0.9791	-0.0603	0.6892	-0.0275	0.9649	-0.0201	0.9317	0.0075	0.9711	0.0113	0.9923	-0.0501	0.9017	-0.0614	0.8611
CG11347	CG11347	1623536_s_at	-0.6233	0.4005	0.4780	0.6419	0.2457	0.5303	0.0397	0.9647	-0.6359	0.0155	-0.6755	0.0072	0.3377	0.9405	0.5161	0.7554	0.1783	0.9261
---	---	1623537_at	-0.2445	0.5251	-0.5020	0.3365	-0.5411	0.0366	-0.7293	0.1073	-0.5351	0.0511	0.1942	0.4418	-0.6922	0.6496	-0.8102	0.1742	-0.1180	0.8889
---	---	1623538_at	0.0923	0.7275	0.1522	0.4008	0.1221	0.4235	0.0793	0.8732	-0.0326	0.8892	-0.1119	0.5002	0.1687	0.8292	0.2011	0.5577	0.0324	0.9434
CG30039	CG30039	1623539_at	0.0578	0.6866	0.0157	0.9062	0.1336	0.4527	-0.0302	0.9643	-0.0215	0.9321	0.0087	0.9689	-0.0812	0.8940	-0.0038	0.9941	0.0774	0.7712
CG31704	CG31704	1623540_at	1.1021	0.0083	1.0163	0.1096	1.1940	0.0011	0.3013	0.6544	-0.0096	0.9839	-0.3109	0.2898	0.2195	0.8869	-0.1145	0.8938	-0.3340	0.5874
---	---	1623541_at	0.1119	0.6909	-0.1524	0.3687	0.0927	0.6687	0.1404	0.8578	0.2307	0.4301	0.0903	0.7723	-0.0306	0.9780	-0.1125	0.7072	-0.0819	0.7983
CG11373	CG11373	1623542_at	0.1502	0.4427	0.0416	0.7333	-0.0205	0.9262	-0.0932	0.8425	-0.1458	0.4173	-0.0526	0.7869	0.0262	0.9764	-0.1207	0.5923	-0.1469	0.5050
CG5807	CG5807	1623543_at	0.3860	0.1674	0.5020	0.0404	-0.1374	0.5492	-0.1597	0.8395	0.0187	0.9676	0.1783	0.5353	0.5486	0.3162	0.2422	0.3810	-0.3064	0.2918
comm3	comm3	1623544_at	-0.7273	0.5864	-0.5315	0.1301	-1.5509	0.0032	-0.3939	0.6166	-1.2967	0.0061	-0.9028	0.0183	0.5293	0.9031	-1.3209	0.3992	-1.8502	0.2585
mus209	proliferating cell n	1623545_at	0.7857	0.2273	-0.2685	0.8437	0.1188	0.8717	0.0018	0.9988	0.9504	0.0155	0.9486	0.0096	-0.3460	0.9514	0.0137	0.9980	0.3597	0.8737
Taf8	prodos	1623546_at	0.3358	0.0426	0.3120	0.1630	0.8028	0.0047	0.2172	0.4529	0.0470	0.8084	-0.1701	0.2260	-0.2202	0.6557	0.0540	0.8435	0.2742	0.1990
side	downstream of ey	1623547_at	-0.4329	0.0590	0.0965	0.7020	-0.2484	0.1985	-0.0760	0.9171	-0.3690	0.1024	-0.2930	0.1459	-0.0154	0.9914	-0.0464	0.9230	-0.0310	0.9431
CG18269	CG18269	1623548_at	0.1029	0.6947	0.0529	0.6815	-0.0752	0.6942	-0.1177	0.8543	0.0807	0.7716	0.1985	0.3496	0.0528	0.9717	-0.0206	0.9739	-0.0734	0.8861
CG1973	CG1973	1623549_at	0.3157	0.1595	0.5778	0.2488	0.4442	0.0656	-0.1787	0.7093	0.0873	0.7334	0.2660	0.1723	-0.0685	0.9742	0.3902	0.4576	0.4587	0.3898
Phlpp	CG10493	1623550_at	0.2111	0.2615	0.0962	0.4026	0.1628	0.4245	-0.0451	0.9218	-0.0300	0.8701	0.0151	0.9287	-0.0378	0.9657	-0.1217	0.6314	-0.0840	0.7568
CG18081 /// DyakCG18081	CG18081	1623551_at	0.1366	0.4050	0.1772	0.2358	-0.2210	0.1828	-0.0642	0.9273	0.0561	0.8345	0.1203	0.5625	0.4195	0.2652	0.1949	0.2924	-0.2246	0.2552
CG2893	CG2893	1623552_s_at	0.0142	0.9801	-0.4712	0.1912	-0.1034	0.8429	-0.1357	0.9009	0.3516	0.3254	0.4873	0.1205	-0.4683	0.8000	0.0528	0.9665	0.5211	0.5173
Miro	Mitochondrial Rho	1623553_a_at	0.0344	0.8298	-0.0696	0.6545	-0.1894	0.2538	-0.2305	0.4717	-0.2081	0.2269	0.0224	0.9151	-0.1349	0.7990	-0.3241	0.1392	-0.1892	0.4003
CG3708	CG3708	1623554_at	0.2191	0.3512	0.1801	0.2218	0.3231	0.0872	-0.0089	0.9941	-0.2144	0.3914	-0.2056	0.3600	-0.0311	0.9624	-0.0498	0.8393	-0.0187	0.9421
CG10131	CG10131	1623555_at	-2.2415	0.0293	-4.1123	0.0049	-4.0922	0.0000	-0.4130	0.3357	0.1297	0.6409	0.5428	0.0225	-0.4035	0.9101	-1.5504	0.2111	-1.1469	0.3809
CG32669	CG32669	1623556_at	0.0645	0.8927	0.2105	0.2880	0.2768	0.3606	-0.2149	0.7906	-0.5528	0.0945	-0.3379	0.2554	-0.1486	0.9457	-0.2269	0.7787	-0.0783	0.9328
CG31679	CG31679	1623557_at	0.2914	0.2007	0.0468	0.6675	0.3375	0.0955	0.1884	0.6560	0.0556	0.8295	-0.1328	0.4969	0.1025	0.8513	0.0429	0.9003	-0.0595	0.8354
Cpr49Ae	CG8505	1623558_at	-2.8441	0.0154	-3.8049	0.0314	-4.2394	0.0000	0.0472	0.9486	-0.6174	0.0091	-0.6646	0.0039	0.5543	0.8960	-1.5685	0.3071	-2.1228	0.2028
---	---	1623559_s_at	0.2596	0.1938	-0.8424	0.0412	-0.8098	0.0365	0.0123	0.9943	1.2694	0.0051	1.2570	0.0032	0.0118	0.9939	-0.0195	0.9739	-0.0313	0.9493
CG18048	CG18048	1623560_at	-0.5016	0.0403	-0.5446	0.1491	-0.2705	0.1215	-0.0138	0.9863	-0.2534	0.1817	-0.2396	0.1591	0.0317	0.9841	-0.0810	0.8749	-0.1127	0.7944
---	---	1623561_at	0.1117	0.6153	0.1475	0.5559	0.4316	0.0204	0.0804	0.8897	-0.0290	0.9149	-0.1094	0.5691	0.1182	0.9066	0.1332	0.7567	0.0150	0.9799
CG14431	CG14431	1623562_at	0.1579	0.5122	0.1006	0.4072	-0.0087	0.9726	-0.1472	0.8705	-0.1513	0.6741	-0.0041	0.9916	0.0478	0.9546	-0.0735	0.8187	-0.1213	0.6476
---	---	1623563_at	0.3629	0.1559	0.1203	0.5201	0.2070	0.3100	0.2018	0.6776	0.2412	0.2934	0.0395	0.8836	-0.1694	0.8302	-0.1265	0.7453	0.0429	0.9249
Syx8	syntaxin	1623564_at	-0.4224	0.0260	-0.4418	0.0985	-0.0969	0.6846	0.1073	0.7929	-0.2430	0.1437	-0.3503	0.0269	-0.1911	0.8461	-0.2531	0.5531	-0.0619	0.9121
CG17278	CG17278	1623565_at	-0.6547	0.1842	-1.1113	0.1171	-0.9931	0.0046	0.2609	0.6483	1.4655	0.0008	1.2046	0.0011	0.0812	0.9831	0.9265	0.2830	0.8454	0.3565
CG13957	CG13957	1623566_at	0.1374	0.4572	0.0926	0.8209	-0.1838	0.4073	0.0576	0.9220	0.3635	0.0463	0.3059	0.0581	0.2543	0.8012	0.2820	0.5176	0.0277	0.9646
---	---	1623567_at	0.0477	0.8033	0.0256	0.7992	-0.0393	0.8411	0.0711	0.9037	-0.0787	0.7234	-0.1497	0.3990	0.0333	0.9611	-0.0421	0.8767	-0.0754	0.7287
---	---	1623568_at	0.0199	0.9376	-0.1491	0.3887	-0.1655	0.3248	-0.0598	0.9023	-0.1028	0.5459	-0.0430	0.8091	0.0011	0.9994	-0.0665	0.8028	-0.0676	0.7846
CG14907	CG14907	1623569_at	-0.7192	0.0080	0.2428	0.0944	0.6717	0.2079	0.2143	0.7000	-0.3223	0.2058	-0.5366	0.0269	-0.2133	0.9320	0.5332	0.5244	0.7465	0.3678
CG7924	CG7924	1623570_at	-0.3730	0.2396	-0.2849	0.1001	-0.4420	0.0579	0.2790	0.5681	0.4968	0.0526	0.2178	0.3444	0.0237	0.9848	0.0126	0.9811	-0.0111	0.9819
CG9328	CG9328	1623571_a_at	-2.7184	0.0004	-1.4389	0.0327	-3.6717	0.0000	-1.1885	0.0044	-1.1165	0.0005	0.0720	0.7204	0.9061	0.2884	0.2355	0.6242	-0.6706	0.1573
xmas-2	xmas-2	1623572_at	0.2114	0.2625	0.1934	0.3940	0.8045	0.0040	0.0386	0.9540	-0.0498	0.8315	-0.0884	0.6390	-0.3313	0.6898	0.0498	0.9264	0.3811	0.2778

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11266	CG11266	1623573_s_at	-0.1669	0.6065	0.3601	0.5976	0.4119	0.1929	-0.1866	0.7696	-0.6824	0.0197	-0.4958	0.0456	-0.1832	0.9454	-0.1196	0.9258	0.0636	0.9563
CG11694	CG11694	1623574_at	0.2201	0.2309	0.0606	0.6096	0.3279	0.1309	0.0378	0.9465	-0.1617	0.3126	-0.1995	0.1586	0.0698	0.9514	-0.0061	0.9935	-0.0759	0.8654
---	---	1623575_at	-0.0751	0.7678	-1.8089	0.2415	-2.2090	0.0057	0.0265	0.9941	1.9196	0.0180	1.8931	0.0119	0.0969	0.8953	-0.0266	0.9519	-0.1235	0.6848
CG1850	CG1850	1623576_at	0.2197	0.2908	0.2163	0.4143	0.2649	0.3327	0.0338	0.9520	-0.0533	0.7784	-0.0872	0.5764	0.0061	0.9964	-0.1536	0.6437	-0.1597	0.6291
---	---	1623577_at	-0.0904	0.5687	0.0509	0.6255	0.1328	0.4273	-0.0490	0.9314	-0.2024	0.2335	-0.1533	0.3228	-0.0866	0.8439	-0.0212	0.9415	0.0654	0.7526
---	---	1623578_at	0.0372	0.8449	-0.0048	0.9673	-0.2122	0.2552	0.0194	0.9798	-0.0066	0.9809	-0.0260	0.9083	0.0291	0.9816	-0.0212	0.9650	-0.0503	0.9025
CG33773	CG33773	1623579_at	-0.1948	0.4478	-0.0587	0.7577	-0.0169	0.9408	0.0960	0.8671	-0.1851	0.3719	-0.2812	0.1201	0.0802	0.9246	-0.0441	0.9202	-0.1243	0.6917
alphaTub84B	alpha-Tubulin	1623580_at	-0.2352	0.3202	0.3243	0.1110	-0.0461	0.7723	-0.0638	0.9136	-0.1590	0.4034	-0.0952	0.6111	0.1770	0.8192	0.3786	0.2171	0.2016	0.5482
---	---	1623581_at	0.0691	0.7230	0.1219	0.4986	0.1586	0.3987	0.1556	0.7351	0.0096	0.9759	-0.1460	0.4493	0.1407	0.8331	0.1287	0.6769	-0.0120	0.9785
CG6364 /// DyakCG6364	CG6364	1623582_at	0.5884	0.0514	1.0031	0.0518	0.7970	0.0008	0.0460	0.9263	0.0538	0.7640	0.0078	0.9674	0.2127	0.8609	0.4746	0.3181	0.2619	0.6186
PNUTS	PNUTS	1623583_at	-0.5051	0.2276	-0.0680	0.9512	0.3764	0.0941	0.0663	0.9218	-0.0920	0.6945	-0.1584	0.4045	-0.2794	0.9246	0.3841	0.7399	0.6635	0.5194
DyakCG10473 /// hkl	CG10473 /// hook	1623584_at	-0.1094	0.7380	0.1428	0.5277	0.0636	0.7789	0.0212	0.9777	0.0384	0.8810	0.0172	0.9428	0.0408	0.9829	0.2572	0.5765	0.2165	0.6437
CG34383	CG14365	1623585_at	0.0821	0.6314	0.0702	0.5841	-0.0200	0.9207	0.0150	0.9857	0.0222	0.9340	0.0072	0.9761	0.0905	0.8882	0.0391	0.9170	-0.0514	0.8755
pyx	pyrexia	1623586_a_at	-1.0739	0.0961	-3.0246	0.0027	-2.1722	0.0003	0.3738	0.6893	1.7382	0.0026	1.3644	0.0045	-0.1998	0.9149	-0.3336	0.6387	-0.1338	0.8810
CG31730	CG31730	1623587_at	0.0613	0.7178	0.1888	0.3795	0.1618	0.4282	0.0436	0.9436	0.0531	0.8044	0.0094	0.9661	-0.0958	0.8906	0.0167	0.9692	0.1125	0.7017
EfTuM	Elongation factor `	1623588_at	-0.6114	0.0051	0.4415	0.1822	0.5697	0.0330	-0.0183	0.9777	-1.1367	0.0003	-1.1184	0.0002	-0.1107	0.9467	-0.0264	0.9760	0.0843	0.9028
CG5326	CG5326	1623589_a_at	0.2138	0.2479	0.2547	0.2362	0.3004	0.2705	0.0987	0.8640	-0.0358	0.8980	-0.1346	0.4918	-0.0067	0.9964	0.0800	0.8867	0.0867	0.8667
Smc5	Smc5	1623590_s_at	-0.2810	0.2716	-0.3653	0.1075	-0.5476	0.0051	-0.4163	0.1649	-0.1402	0.4592	0.2760	0.0897	-0.2778	0.6955	-0.1330	0.6823	0.1448	0.6441
CG17883	CG17883	1623591_a_at	0.0827	0.7141	0.0363	0.9337	-0.1365	0.3626	-0.4001	0.1976	-0.0278	0.9162	0.3723	0.0338	-0.1609	0.8764	-0.0264	0.9684	0.1345	0.7774
CG15894 /// DmirCG15894	CG15894	1623592_at	-0.0224	0.9157	0.3028	0.1695	0.2772	0.2187	0.1847	0.6015	-0.0088	0.9746	-0.1935	0.2258	0.1186	0.8235	0.1956	0.3669	0.0770	0.7638
---	---	1623593_at	-0.0668	0.7108	0.0914	0.5016	0.0510	0.7705	-0.1574	0.7409	-0.2582	0.2163	-0.1008	0.6387	0.0144	0.9893	0.0722	0.8018	0.0578	0.8408
blow	blown fuse	1623594_at	-2.8148	0.0029	-4.0494	0.0011	-3.2861	0.0001	0.4514	0.6450	0.5182	0.2782	0.0667	0.9078	0.1485	0.8374	-0.3474	0.2254	-0.4958	0.1271
CG5439	CG5439	1623595_at	-0.5994	0.0347	-0.6804	0.0096	-1.1040	0.0012	-0.1650	0.7028	-0.1069	0.6288	0.0581	0.7945	0.3975	0.6557	0.0181	0.9781	-0.3794	0.3057
Cpr72Eb	CG12255	1623596_at	0.0939	0.6207	0.0750	0.6388	0.0768	0.7475	-0.0533	0.9533	-0.0780	0.7998	-0.0246	0.9367	0.1720	0.7644	0.0319	0.9341	-0.1401	0.5838
CG17684	CG40072	1623597_s_at	0.1325	0.4597	0.0489	0.8237	-0.1236	0.4177	0.1061	0.7950	0.2803	0.0931	0.1741	0.2446	0.0460	0.9683	0.0498	0.9162	0.0038	0.9945
---	---	1623598_at	0.0882	0.6964	0.0286	0.9034	-0.0477	0.7627	-0.1578	0.8132	-0.0804	0.8037	0.0774	0.7932	0.1222	0.8756	0.0557	0.9054	-0.0665	0.8707
boca	boca	1623599_at	0.0196	0.9051	0.1256	0.3822	0.1842	0.2613	0.0988	0.7599	0.5225	0.0034	0.4238	0.0052	0.0929	0.8841	0.7160	0.0257	0.6231	0.0441
---	---	1623600_at	0.2200	0.4276	0.2320	0.2182	0.3374	0.0921	-0.0770	0.9053	-0.2538	0.2253	-0.1768	0.3591	-0.0089	0.9946	-0.0196	0.9687	-0.0107	0.9837
Amyrel	amyrel	1623601_at	0.2561	0.2984	0.0465	0.7973	0.2246	0.4082	0.0736	0.9011	0.1901	0.3286	0.1165	0.5341	-0.0221	0.9922	0.1650	0.8031	0.1871	0.7531
Ptp69D	Protein tyrosine pl	1623602_a_at	-0.2977	0.3633	-0.1623	0.5912	-0.5391	0.0110	-0.2551	0.4979	0.2097	0.2973	0.4648	0.0172	0.0621	0.9741	0.2111	0.6873	0.1490	0.7915
CG4793	CG4793	1623603_at	-0.1267	0.4053	-0.0807	0.6020	0.1156	0.4663	0.0343	0.9441	0.0153	0.9397	-0.0190	0.9103	-0.0333	0.9635	0.0977	0.6490	0.1310	0.5278
CG7742	CG7742	1623604_at	0.0433	0.8033	-0.0452	0.7868	-0.3313	0.1296	-0.3549	0.2703	-0.0631	0.7846	0.2918	0.0864	0.0562	0.9514	-0.0332	0.9402	-0.0894	0.7815
cbt	TGF-beta-inducibl	1623605_a_at	0.4521	0.0391	0.0778	0.8845	-0.1568	0.7277	0.0520	0.9473	0.5227	0.0265	0.4706	0.0260	0.2810	0.8692	0.1699	0.8550	-0.1111	0.9057
RpLP1	Ribosomal-proteir	1623606_at	0.4150	0.0329	1.0091	0.0120	0.7811	0.0031	0.0228	0.9749	-0.1891	0.2906	-0.2119	0.1804	0.1109	0.8298	-0.0666	0.8123	-0.1775	0.4213
Mgat1	N-acetylglucosam	1623607_at	0.8978	0.0027	1.0768	0.0278	0.8847	0.0006	-0.2843	0.3544	0.2503	0.1522	0.5346	0.0053	-0.0857	0.9467	0.5308	0.1626	0.6166	0.1432
---	---	1623608_at	0.0176	0.9483	0.0788	0.6453	0.1957	0.2516	-0.0988	0.7973	-0.0386	0.8487	0.0602	0.7168	-0.1884	0.7697	0.0932	0.7825	0.2816	0.3138
CG15172	CG15172	1623609_at	0.0080	0.9692	0.0494	0.8384	-0.0978	0.5552	-0.1480	0.8544	-0.0296	0.9452	0.1184	0.6990	0.1150	0.8122	0.1104	0.6059	-0.0046	0.9884
CG34357	CG14652	1623610_at	0.0829	0.5986	-0.0244	0.8080	-0.0928	0.5843	0.0307	0.9539	0.1139	0.4529	0.0832	0.5646	-0.0518	0.9460	-0.1589	0.5068	-0.1071	0.6724
---	---	1623611_at	-0.1300	0.4338	-0.1611	0.5396	-0.0987	0.7420	0.2400	0.7205	0.2175	0.4948	-0.0226	0.9534	0.2507	0.7633	0.1198	0.7802	-0.1309	0.7443
CG7722 /// DsimCG7722	CG7722	1623612_at	-0.7513	0.1432	2.2692	0.0657	-0.1842	0.6600	-2.5610	0.0416	-4.4453	0.0004	-1.8843	0.0093	0.0288	0.9943	-1.0324	0.2593	-1.0612	0.2798
CG13762	CG13762	1623613_at	0.1838	0.3890	0.2325	0.3039	0.2542	0.1861	-0.1502	0.8281	-0.0226	0.9544	0.1276	0.6402	0.0063	0.9942	0.0542	0.8380	0.0479	0.8509
---	---	1623614_at	0.2460	0.3496	0.0745	0.5224	0.3440	0.1481	0.0276	0.9666	-0.0702	0.7298	-0.0979	0.5695	-0.0118	0.9924	-0.0894	0.8117	-0.0776	0.8307
CG8675	CG8675	1623615_at	0.3556	0.1829	-0.1487	0.7090	0.4263	0.0903	0.2871	0.5179	0.2890	0.2160	0.0018	0.9952	-0.1967	0.8270	-0.2863	0.4462	-0.0896	0.8542
---	---	1623616_at	0.2426	0.3066	-0.0501	0.6408	0.0247	0.8886	-0.0396	0.9518	0.1126	0.5692	0.1522	0.3675	-0.0157	0.9900	-0.2732	0.3010	-0.2575	0.3569
---	---	1623617_at	0.3287	0.0981	0.0016	0.9901	-0.0969	0.7270	0.0001	0.9999	0.2268	0.2717	0.2267	0.2177	0.0374	0.9781	-0.1222	0.7504	-0.1596	0.6476
CG34356	CG12524	1623618_at	0.1561	0.3671	0.0868	0.5479	0.1029	0.5765	0.0663	0.8943	-0.0626	0.7523	-0.1289	0.4066	0.0923	0.9016	-0.1000	0.7611	-0.1923	0.4978
CG15684	CG15684	1623619_at	-0.1094	0.6266	0.1117	0.5015	0.1738	0.2974	-0.0093	0.9910	-0.0812	0.6956	-0.0719	0.7065	0.0091	0.9939	0.1283	0.6643	0.1192	0.6881
rho-6	rhomboid-6	1623620_a_at	0.0577	0.7961	0.0000	1.0000	0.3630	0.0653	0.1389	0.7556	0.0924	0.6730	-0.0464	0.8325	-0.1184	0.8672	0.0614	0.8789	0.1798	0.5448
---	---	1623621_at	-0.0115	0.9638	-0.0589	0.5751	0.1256	0.4296	0.4559	0.1332	0.3797	0.0441	-0.0762	0.6918	0.0887	0.9076	0.0408	0.9237	-0.0479	0.8982
CG12862	CG12862	1623622_a_at	0.0737	0.7303	0.2355	0.2074	0.2662	0.1943	-0.0324	0.9677	-0.1235	0.5908	-0.0911	0.6798	0.0119	0.9914	0.1275	0.6382	0.1156	0.6724
---	---	1623623_at	0.3968	0.1587	0.2464	0.4102	0.0122	0.9622	0.1399	0.8676	0.0749	0.8449	-0.0649	0.8498	0.2011	0.8049	-0.0621	0.9054	-0.2632	0.4548
CG14869	CG14869	1623624_at	-0.8771	0.0146	-0.8565	0.0354	-1.0875	0.0073	-0.5822	0.0584	-0.7047	0.0027	-0.1225	0.4652	-0.3702	0.7779	-0.5788	0.2972	-0.2086	0.7504
CG14542	CG14542	1623625_at	-0.4074	0.0508	0.0402	0.7853	0.3072	0.2871	0.3239	0.3348	-0.1383	0.4976	-0.4622	0.0157	0.0301	0.9898	0.3701	0.4833	0.3401	0.5325
CG11412	CG11412	1623626_a_at	-0.3513	0.1493	-0.0361	0.7089	-0.2843	0.2067	0.1202	0.8155	0.0353	0.8985	-0.0849	0.6895	0.2155	0.7726	0.2378	0.4696	0.0223	0.9626
---	---	1623627_at	-0.0420	0.8129	-0.0253	0.8797	-0.0946	0.6854	0.0481	0.9558	0.1351	0.5970	0.0870	0.7279	0.0185	0.9869	0.0713	0.8372	0.0528	0.8809
Gr36b	Gustatory recepto	1623628_at	0.2513	0.1338	0.0145	0.9432	0.1515	0.3266	-0.0343	0.9603	-0.0432	0.8548	-0.0089	0.9689	-0.1477	0.7686	-0.1597	0.4562	-0.0120	0.9709
alpha4GT2	alpha4GT2	1623629_at	-0.1135	0.5398	0.1505	0.3485	-0.1098	0.5590	-0.0203	0.9768	-0.0616	0.7650	-0.0413	0.8338	0.0618	0.8982	0.0964	0.6156	0.0346	0.8885
---	---	1623630_at	0.2028	0.2457	0.0554	0.6044	0.0608	0.6928	-0.2483	0.4599	0.0082	0.9772	0.2565	0.1141	0.0282	0.9774	0.0196	0.9591	-0.0086	0.9835
AP-47	AP-47	1623631_at	-0.2638	0.3041	0.4919	0.1990	0.5615	0.0063	0.2537	0.5079	-0.2264	0.2671	-0.4801	0.0163	0.2895	0.7644	0.6674	0.1089	0.3780	0.3690
CG6327	CG6327	1623632_s_at	0.1362	0.5423	0.7180	0.0321	0.2192	0.2877	-0.2861	0.4927	-0.5319	0.0246	-0.2458	0.2158	0.1460	0.8874	-0.0059	0.9941	-0.1519	0.7307
CG2034	CG2034	1623633_at	0.0708	0.7981	0.9828	0.0186	1.0193	0.0004	0.2014	0.7556	-0.4545	0.1028	-0.6559	0.0167	0.2358	0.7689	0.6163	0.0825	0.3805	0.2798
CG6527	CG6527	1623634_at	0.2424	0.2800	-0.0320	0.8960	-0.1564	0.4370	0.0766	0.8667	0.1180	0.4841	0.0414	0.8211	0.0629	0.9611	-0.0461	0.9364	-0.1089	0.8050
TotM	Turandot M	1623635_at	2.4991	0.0651	-0.4618	0.7808	1.6108	0.1130	1.7482	0.4611	1.9552	0.1269	0.2070	0.8930	-0.2519	0.9774	-1.0525	0.6417	-0.8006	0.7398
Obp19a	Odorant-binding p	1623636_at	0.1270	0.6020	0.1890	0.1573	0.0748	0.7892	-0.1571	0.7872	-0.2484	0.3044	-0.0913	0.7204	0.0477	0.9679	-0.0775	0.8590	-0.1252	0.7271
Lcp65Ab1 /// Lcp65Ab2	Larval cuticle prot	1623637_s_at	-0.0859	0.6908	0.0536	0.7997	0.0786	0.6228	0.1114	0.8383	-0.0202	0.9469	-0.1316	0.5040	0.0826	0.8890	0.0437	0.8960	-0.0389	0.8993
CG3894	dmNeural-like	1623638_a_at	-0.0373	0.9264	0.4893	0.0707	0.2807	0.3227	-0.0655	0.9441	-0.5495	0.0459	-0.4840	0.0484	0.2193	0.8049	0.1999	0.6156	-0.0194	0.9732
---	---	1623639_at	0.1102	0.6181	-0.2427	0.2019	-0.1354	0.5413	-0.0640	0.9504	0.1871	0.5392	0.2511	0.3340	-0.1370	0.8049	-0.0469	0.8903	0.0901	0.7362
CG32169	CG32169	1623640_at	-0.0525	0.8335	-0.0979	0.3913	-0.2513	0.1534	-0.0030	0.9961	0.1893	0.3526	0.1924	0.2879	0.0723	0.9375	0.0164	0.9735	-0.0559	0.8884
CG32856	CG32856	1623641_at	-0.1983	0.3850	0.1368	0.7861	0.1539	0.4905	-0.1147	0.8764	-0.2347	0.3658	-0.1201	0.6461	0.0751	0.9526	0.1075	0.8304	0.0324	0.9514

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG16723	CG16723	1623642_at	0.2357	0.2442	0.0250	0.8051	0.2597	0.2939	0.0751	0.9300	0.0805	0.7932	0.0054	0.9863	-0.1340	0.8122	-0.0220	0.9531	0.1121	0.6580
ldgf3	Imaginal disc grov	1623643_s_at	-0.6302	0.0098	-0.8627	0.0562	-0.7679	0.0023	-0.0537	0.8987	0.1543	0.2653	0.2080	0.0915	-0.1960	0.8465	-0.0565	0.9328	0.1395	0.7777
oho23B	Ribosomal protein	1623644_s_at	0.2491	0.1501	0.6858	0.0137	0.7827	0.0024	0.0631	0.9138	-0.5997	0.0065	-0.6628	0.0025	-0.0604	0.9330	-0.1550	0.5157	-0.0945	0.7154
RhoGAP5A	RhoGAP5A	1623645_a_at	-0.4308	0.4338	-0.6080	0.4347	-0.8687	0.0395	0.0627	0.9314	0.3373	0.1169	0.2746	0.1538	0.4833	0.8465	0.1254	0.9404	-0.3579	0.7683
CG15400	CG15400	1623646_at	-0.0325	0.8620	-0.1937	0.4996	-0.0373	0.8535	-0.0677	0.9247	-0.1154	0.6283	-0.0477	0.8476	-0.0924	0.9238	-0.1710	0.6191	-0.0786	0.8488
CG8243	CG8243	1623647_at	-0.2034	0.6329	-0.2583	0.6431	-0.2390	0.4072	0.0041	0.9956	0.2154	0.2073	0.2113	0.1668	-0.0552	0.9872	0.2718	0.7787	0.3270	0.7134
Psa	Puromycin sensiti	1623648_s_at	0.4662	0.2292	0.5265	0.0940	0.6074	0.0406	0.2677	0.4908	0.6436	0.0080	0.3759	0.0477	0.2681	0.8202	0.6403	0.1772	0.3722	0.4627
---	---	1623649_at	0.0517	0.8562	0.0678	0.4857	0.0153	0.9306	-0.0430	0.9435	0.0976	0.6042	0.1406	0.3761	-0.0239	0.9848	0.1357	0.6598	0.1596	0.5969
PNGase	PNGase	1623650_a_at	0.4841	0.0110	0.1616	0.5590	0.0858	0.6252	-0.1319	0.7608	0.1747	0.3586	0.3066	0.0688	-0.0840	0.9137	-0.0963	0.7699	-0.0124	0.9769
---	---	1623651_at	0.0253	0.9119	-0.0121	0.9189	-0.0885	0.5980	0.0010	0.9988	-0.0653	0.7138	-0.0663	0.6802	0.1302	0.8270	-0.0293	0.9419	-0.1594	0.5448
CG14740	CG14740	1623652_at	0.3150	0.2184	0.1334	0.5279	0.1613	0.4715	0.1660	0.7507	0.1366	0.5785	-0.0293	0.9156	0.0431	0.9742	-0.1045	0.8038	-0.1476	0.6844
VhaM9.7-2	VhaM9.7-2	1623653_at	-0.8532	0.0428	-0.9694	0.0102	-1.4200	0.0000	-0.1595	0.6678	-0.1922	0.2804	-0.0328	0.8762	0.2107	0.8554	-0.3591	0.4503	-0.5698	0.2474
---	---	1623654_at	0.0479	0.7917	-0.0507	0.7875	0.0580	0.7539	-0.0646	0.9218	-0.2032	0.3113	-0.1386	0.4615	-0.0563	0.9666	-0.0783	0.8785	-0.0221	0.9673
Yp2	yolk protein	1623655_at	1.2916	0.0202	0.3625	0.3707	1.3335	0.0022	0.4013	0.6869	0.8230	0.0774	0.4217	0.3174	-0.3721	0.7644	-0.1055	0.8924	0.2665	0.6395
CG1109	CG1109	1623656_s_at	0.2158	0.6369	0.2740	0.1433	0.4155	0.0674	0.1325	0.8844	0.2021	0.5456	0.0695	0.8469	0.0457	0.9851	0.4545	0.4095	0.4088	0.4777
CG14505	CG14505	1623657_at	0.1158	0.5180	0.0293	0.8606	0.0561	0.7389	0.0124	0.9873	0.0566	0.8066	0.0442	0.8378	0.0194	0.9862	-0.1527	0.5630	-0.1721	0.5117
CG13306	CG13306	1623658_at	-1.8582	0.0063	-1.6632	0.0136	-1.6780	0.0001	-0.0300	0.9803	-0.5420	0.0710	-0.5119	0.0580	0.1026	0.9653	-0.3567	0.5971	-0.4593	0.4835
Obp56d	Odorant-binding p	1623659_at	-2.1828	0.1991	-0.6182	0.6046	-2.8609	0.0065	-1.5692	0.2591	-2.9129	0.0045	-1.3437	0.0702	0.6734	0.9238	-1.3943	0.5727	-2.0677	0.3878
CG4892	CG4892	1623660_at	-0.0756	0.7916	0.0782	0.7383	0.1937	0.4821	-0.2622	0.7604	-0.3515	0.3495	-0.0893	0.8323	-0.2714	0.7506	-0.2015	0.6075	0.0699	0.8900
---	---	1623661_at	0.0914	0.6214	0.0720	0.6145	0.1889	0.2331	0.1834	0.6217	0.0799	0.7077	-0.1035	0.5730	0.0487	0.9405	0.0278	0.9314	-0.0210	0.9411
CG11044	CG11044	1623662_at	-2.5715	0.0041	-2.4734	0.0083	-2.7917	0.0000	-0.2703	0.5917	-1.1160	0.0018	-0.8457	0.0036	0.0218	0.9950	-0.9546	0.2168	-0.9763	0.2363
CG18063	CG18063	1623663_a_at	-0.1275	0.4002	-0.0894	0.3775	-0.0819	0.7393	0.1253	0.8532	0.1240	0.6531	-0.0012	0.9970	-0.0417	0.9677	0.0325	0.9404	0.0742	0.8256
CG10585 /// DyakCG10585	CG10585	1623664_at	-0.8386	0.0127	-0.4866	0.0371	-0.2244	0.2053	-0.1178	0.7678	-0.3959	0.0273	-0.2780	0.0694	-0.2913	0.7663	-0.0040	0.9973	0.2873	0.5058
---	---	1623665_at	0.0859	0.5583	-0.0316	0.7743	0.0885	0.6099	0.1638	0.6086	0.0915	0.6039	-0.0724	0.6642	0.0644	0.9521	-0.0181	0.9738	-0.0825	0.8412
CG10516	CG10516	1623666_at	2.2263	0.0006	1.1366	0.0473	2.0921	0.0004	0.4468	0.1807	0.6572	0.0066	0.2104	0.2467	-0.4406	0.8024	-0.4001	0.6108	0.0405	0.9716
---	---	1623667_at	0.0886	0.5437	0.2017	0.3503	0.3410	0.1703	0.1127	0.7833	-0.1545	0.3711	-0.2672	0.0788	-0.0986	0.9092	-0.0119	0.9841	0.0867	0.8209
sha	kojak	1623668_at	0.2689	0.1789	0.1279	0.4120	0.0722	0.7109	-0.1135	0.8487	0.0058	0.9859	0.1193	0.5827	-0.0230	0.9862	-0.0279	0.9531	-0.0049	0.9924
CG16836	CG16836	1623669_at	3.0298	0.0009	1.5922	0.1721	2.4546	0.0003	0.7862	0.3692	0.3339	0.5279	-0.4522	0.3170	-0.0795	0.9862	-1.0576	0.2924	-0.9781	0.3569
CG14562 /// DmirCG14562	CG14562	1623670_at	-0.6705	0.0813	-0.5147	0.1040	-0.4409	0.0239	0.0826	0.9179	0.2419	0.3333	0.1593	0.5010	0.1097	0.9340	0.4808	0.2389	0.3712	0.3895
---	---	1623671_at	0.0710	0.7030	0.1463	0.1910	0.2452	0.2523	-0.2304	0.4455	-0.1037	0.5592	0.1267	0.4099	-0.1992	0.7997	-0.0224	0.9666	0.1768	0.6166
---	---	1623672_at	0.1233	0.6318	0.0003	1.0000	0.0661	0.7273	0.0332	0.9688	-0.0302	0.9220	-0.0635	0.7981	-0.0607	0.9548	-0.0407	0.9353	0.0201	0.9659
CG10435	CG10435	1623673_at	-0.2894	0.1966	0.1975	0.3207	0.2007	0.3119	0.1847	0.7764	-0.3203	0.2396	-0.5050	0.0438	0.1711	0.7588	0.3061	0.1769	0.1350	0.5882
---	---	1623674_at	-0.0317	0.8664	-0.0070	0.9508	0.2046	0.2990	0.0204	0.9745	-0.0719	0.6888	-0.0923	0.5485	-0.0802	0.9246	0.0349	0.9402	0.1152	0.7224
Obp99b	turn on sex-specif	1623675_at	0.9179	0.3107	0.9911	0.5279	0.1933	0.7873	0.4462	0.8800	-0.2983	0.8114	-0.7444	0.4339	1.1672	0.7726	-0.2366	0.9321	-1.4038	0.4287
Doa	Mosaic suppresso	1623676_at	0.0645	0.6945	-0.0423	0.7334	-0.1312	0.5225	-0.0403	0.9649	0.2911	0.2301	0.3314	0.1254	-0.0367	0.9717	0.1463	0.6009	0.1830	0.5003
---	---	1623677_at	0.1177	0.6049	0.2105	0.2006	0.0176	0.9150	-0.1032	0.8431	-0.0990	0.6487	0.0042	0.9862	0.0287	0.9829	-0.0471	0.9210	-0.0758	0.8462
mRpS30	mitochondrial ribo	1623678_at	-0.1242	0.5539	-0.3614	0.4045	-0.5733	0.0027	-0.1488	0.6338	0.2785	0.0723	0.4273	0.0079	0.0372	0.9872	0.0710	0.9334	0.0338	0.9659
CG4789	CG4789	1623679_at	0.2663	0.2674	0.0378	0.9511	0.1423	0.5967	0.0381	0.9722	0.2757	0.3185	0.2375	0.3427	0.0189	0.9916	0.0092	0.9925	-0.0097	0.9886
CG32793	CG32793	1623680_at	0.0504	0.7790	0.0569	0.6971	-0.1247	0.6334	-0.0351	0.9674	0.0600	0.8356	0.0951	0.6880	-0.0153	0.9901	0.0475	0.9082	0.0628	0.8577
CG10163	CG10163	1623681_at	-1.9697	0.0075	-2.1559	0.0032	-2.4807	0.0004	-0.0717	0.9693	0.1544	0.7932	0.2261	0.6482	-0.0732	0.9168	-0.1192	0.6512	-0.0459	0.8905
Cam	calmodulin	1623682_a_at	-0.2989	0.4148	0.3093	0.1257	-0.0752	0.7635	-0.5405	0.1194	-0.4188	0.0507	0.1217	0.5554	0.1436	0.9352	0.4137	0.4695	0.2701	0.6580
mRpL33	mitochondrial ribo	1623683_at	0.0522	0.8309	0.6493	0.0755	0.5246	0.0291	-0.0106	0.9866	-0.7205	0.0014	-0.7099	0.0009	0.1223	0.9453	-0.1017	0.9009	-0.2239	0.7134
CG33960	CG33960	1623684_at	0.0514	0.8759	0.2160	0.1553	0.2534	0.4490	-0.0082	0.9949	-0.1185	0.6930	-0.1103	0.6873	-0.0196	0.9914	0.0859	0.8853	0.1055	0.8413
CG4064	CG4064	1623685_at	0.2713	0.2418	0.1307	0.2193	-0.0776	0.7677	0.0318	0.9688	0.0630	0.8102	0.0312	0.9029	0.1123	0.8609	-0.1029	0.7319	-0.2152	0.4114
CG7148	CG7148	1623686_at	-0.3602	0.1742	0.8540	0.0219	0.9619	0.0006	0.1225	0.7647	-0.9679	0.0007	-1.0904	0.0003	-0.0577	0.9677	0.2936	0.4379	0.3513	0.3671
CG15199	CG15199	1623687_at	1.5272	0.0012	1.1230	0.0948	1.8557	0.0049	0.0635	0.8889	-0.0593	0.7417	-0.1228	0.3849	-0.8795	0.7202	-0.6281	0.5552	0.2514	0.8474
CG3680	CG3680	1623688_at	0.0882	0.6849	0.3869	0.2691	0.4762	0.0091	-0.0011	0.9991	-0.2052	0.3798	-0.2041	0.3282	0.0825	0.9365	0.3522	0.2660	0.2697	0.4201
CG14642	CG14642	1623689_a_at	5.2792	0.0025	2.4883	0.0014	5.6936	0.0000	2.5707	0.1119	2.1435	0.0332	-0.4272	0.6697	-0.6650	0.5667	-0.5584	0.2705	0.1066	0.8813
CG17190	CG17190	1623690_at	-0.1912	0.6853	-0.3152	0.3083	-0.2204	0.3015	-0.1545	0.8908	-0.5821	0.1213	-0.4276	0.2045	-0.2059	0.8767	-0.6974	0.1647	-0.4915	0.3552
CG3964	CG3964	1623691_at	0.2054	0.2824	0.0795	0.7216	0.3161	0.0411	0.1893	0.5453	0.2862	0.0794	0.0969	0.5416	-0.0776	0.9030	0.0591	0.8527	0.1367	0.5801
rho-4	rhomboid-4	1623692_s_at	0.9806	0.0368	1.5089	0.0149	1.1611	0.0119	0.6865	0.2109	0.8021	0.0232	0.1156	0.7460	0.9662	0.5126	1.2810	0.0795	0.3147	0.6853
drongo	drongo	1623693_a_at	-0.1507	0.4837	-1.3243	0.0770	-1.0678	0.0039	-0.0544	0.9436	0.1594	0.4874	0.2138	0.2796	-0.1312	0.9547	-0.8020	0.2198	-0.6709	0.3334
Gas8	Growth arrest spe	1623694_at	0.1123	0.5268	-0.0670	0.5883	0.0627	0.7105	0.0032	0.9956	0.1423	0.4561	0.1391	0.4179	-0.0073	0.9928	-0.0202	0.9469	-0.0130	0.9622
CG11255	CG11255	1623695_a_at	0.4157	0.1179	0.5473	0.1792	0.5101	0.0236	-0.1347	0.8074	-0.2960	0.1759	-0.1614	0.4349	-0.0366	0.9862	-0.2034	0.7124	-0.1667	0.7697
CG15634	CG15634	1623696_at	0.0793	0.6409	-0.4564	0.0874	-0.0590	0.7986	0.4837	0.1526	0.4980	0.0230	0.0143	0.9565	-0.0697	0.9398	-0.1137	0.7273	-0.0440	0.9096
CG33465	CG33465	1623697_s_at	-1.5954	0.0100	-1.3569	0.0149	-1.7558	0.0001	-0.2365	0.6338	-0.9795	0.0026	-0.7430	0.0053	-0.0133	0.9952	-0.8854	0.0881	-0.8721	0.1170
Iap2	IAP-like protein	1623698_s_at	-0.6390	0.0281	-0.8399	0.1574	-0.9395	0.0018	0.1163	0.7994	0.1481	0.4394	0.0318	0.8845	0.0480	0.9859	-0.1252	0.8862	-0.1732	0.8182
sm	smooth	1623699_a_at	-0.6756	0.3289	-0.7544	0.3766	-1.2344	0.0022	0.2618	0.8085	0.7884	0.0699	0.5266	0.1702	0.5157	0.8692	0.7175	0.5880	0.2019	0.9067
CG32797	CG32797	1623700_at	0.1964	0.4068	-0.1722	0.3098	-0.2078	0.1904	0.0031	0.9956	0.1582	0.3290	0.1551	0.2827	-0.0807	0.9137	-0.1492	0.5889	-0.0685	0.8354
CG30120	CG30120	1623701_at	0.0615	0.8261	-0.0917	0.7887	-0.3411	0.1125	-0.2226	0.7380	-0.1691	0.5992	0.0535	0.8779	-0.0288	0.9823	-0.1865	0.5529	-0.1577	0.6263
CG3226	CG3226	1623702_at	-0.0617	0.8831	-0.0159	0.9660	0.0035	0.9910	0.2149	0.7441	0.3604	0.2112	0.1455	0.6213	0.1591	0.8875	0.3710	0.3706	0.2119	0.6400
---	---	1623703_at	-0.1470	0.4139	0.1659	0.3044	0.1619	0.3389	-0.0408	0.9622	-0.2457	0.2867	-0.2049	0.3268	0.0980	0.8875	0.0921	0.7715	-0.0059	0.9886
CG17841	CG17841	1623704_at	-0.3138	0.2648	-0.2333	0.5075	-0.2236	0.3963	-0.0307	0.9603	-0.3034	0.0677	-0.2727	0.0664	-0.0150	0.9952	-0.3045	0.6313	-0.2895	0.6474
---	---	1623705_at	-0.1307	0.4614	-0.0745	0.7829	-0.0970	0.7144	-0.0423	0.9465	-0.0347	0.8851	0.0077	0.9724	0.0286	0.9860	0.0148	0.9829	-0.0137	0.9829
CG9498	CG9498	1623706_at	-0.6935	0.3255	-0.3926	0.0606	-0.0690	0.7901	0.2278	0.9086	0.4807	0.4675	0.2530	0.7055	-0.1481	0.9238	0.7911	0.1268	0.9392	0.1028
CG32192	CG32192	1623707_a_at	0.1685	0.3533	-0.0660	0.5591	-0.1629	0.3407	-0.1159	0.7950	0.0801	0.7029	0.1960	0.2291	0.0958	0.8561	-0.0143	0.9682	-0.1101	0.6367
alph	alphabet	1623708_at	-0.4923	0.1781	-0.7696	0.1573	-1.0843	0.0031	-0.0176	0.9900	0.4753	0.1286	0.4929	0.0804	0.4394	0.7633	0.3958	0.5324	-0.0437	0.9613
CG14854	CG14854	1623709_at	-1.8402	0.0186	0.1187	0.6105	-1.7365	0.0133	-2.2602	0.0855	-2.3468	0.0085	-0.0866	0.9262	0.0131	0.9946	-0.1518	0.7883	-0.1649	0.7514
CG5847	CG5847	1623710_at	-1.4116	0.0265	-0.1236	0.5450	-0.4924	0.1100	-0.2044	0.8714	-1.4640	0.0059	-1.2595	0.0070	0.0390	0.9884	-0.4118	0.4939	-0.4508	0.4569

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Nach	pickpocket 4	1623711_at	0.4007	0.0640	0.1140	0.4955	0.1077	0.6146	-0.1835	0.6338	0.0466	0.8489	0.2301	0.1723	0.0538	0.9454	-0.0662	0.8353	-0.1200	0.6389
CG10132	CG10132	1623712_at	-1.1736	0.0064	-1.3025	0.0060	-1.6478	0.0001	-0.2761	0.3454	-0.0365	0.8702	0.2395	0.1110	-0.0794	0.9405	-0.2620	0.4230	-0.1826	0.6048
CG18581	CG18581	1623713_at	0.1128	0.4762	0.0109	0.9197	0.0212	0.9187	-0.0534	0.9255	-0.0014	0.9956	0.0521	0.7868	-0.0457	0.9506	-0.1092	0.6471	-0.0634	0.8170
CG8620	CG8620	1623714_at	-0.0737	0.8000	0.1913	0.8447	-0.2710	0.2804	0.1592	0.8234	0.7006	0.0197	0.5414	0.0358	0.6761	0.7633	0.9918	0.2753	0.3157	0.7763
CG4567	CG4567	1623715_at	-0.2819	0.2508	-0.6585	0.1467	-0.0698	0.8290	0.1924	0.5919	0.6596	0.0040	0.4672	0.0115	-0.3237	0.7839	0.2541	0.6458	0.5779	0.2731
Arp66B	Actin-related prote	1623716_at	-0.0296	0.8892	0.2027	0.3624	0.0782	0.7496	-0.1082	0.7733	0.1844	0.2427	0.2926	0.0438	0.0833	0.9422	0.5040	0.1488	0.4208	0.2543
CG32086	CG32086	1623717_at	0.1118	0.6473	-0.1757	0.1597	-0.0985	0.5109	-0.0456	0.9465	0.1435	0.4739	0.1891	0.2751	-0.1208	0.8680	-0.2063	0.4828	-0.0855	0.8097
upd2	unpaired 2	1623718_at	0.2899	0.1480	0.0871	0.4195	0.0502	0.7868	-0.1226	0.8070	0.1196	0.5811	0.2423	0.1738	0.0571	0.9324	0.0076	0.9863	-0.0495	0.8618
---	---	1623719_at	0.1176	0.5301	0.1457	0.3498	-0.0331	0.8661	-0.1360	0.7691	0.0406	0.8765	0.1765	0.3246	0.0335	0.9779	0.0304	0.9490	-0.0031	0.9953
CG30070	CG30070	1623720_at	-0.1134	0.5844	-0.3198	0.1598	-0.1511	0.4496	0.1083	0.8544	0.1357	0.5543	0.0274	0.9161	-0.3660	0.7142	-0.2197	0.6253	0.1463	0.7625
---	---	1623721_at	0.2430	0.1777	0.1399	0.4328	-0.1748	0.4958	-0.1648	0.7380	-0.0158	0.9616	0.1491	0.4720	0.1672	0.8480	-0.1018	0.8343	-0.2689	0.4746
nonA-l	no-on transient A	1623722_at	0.0885	0.7654	0.2517	0.2724	0.1797	0.3274	-0.0495	0.9507	0.0805	0.7614	0.1300	0.5499	0.2509	0.8049	0.4240	0.3012	0.1731	0.7156
---	---	1623723_at	0.1960	0.3683	-0.1243	0.3755	-0.2140	0.1990	0.1098	0.7451	0.3076	0.0440	0.1978	0.1339	-0.0840	0.8956	-0.1470	0.5565	-0.0631	0.8350
CG15027 /// DyakCG15027	CG15027	1623724_at	0.2334	0.4492	0.3505	0.4070	0.1947	0.3248	0.1892	0.8046	0.4412	0.1467	0.2519	0.3719	0.3587	0.7220	0.4097	0.3275	0.0509	0.9330
CG7914	CG7914	1623725_at	0.0207	0.9302	0.3872	0.0287	0.6068	0.0046	0.0209	0.9774	-0.3902	0.0358	-0.4111	0.0182	-0.2861	0.6557	-0.0730	0.8349	0.2132	0.4250
CG11755	CG11755	1623726_at	-0.1335	0.7336	0.0545	0.6185	0.1680	0.4106	-0.0805	0.9295	-0.2104	0.4471	-0.1299	0.6333	-0.0094	0.9964	0.2492	0.6406	0.2586	0.6257
Cyp313a2	Cyp313a2	1623727_at	0.0274	0.9072	0.0961	0.4653	-0.0808	0.7856	-0.1828	0.7890	-0.1706	0.5725	0.0122	0.9721	0.0484	0.9589	0.0840	0.8079	0.0356	0.9231
CG13689	CG13689	1623728_at	0.3820	0.1818	-0.1529	0.5376	-0.1662	0.5580	0.0008	0.9996	0.2103	0.6559	0.2095	0.6230	-0.2873	0.6955	-0.3526	0.2263	-0.0652	0.8723
CG4136	CG4136	1623729_at	0.0529	0.8623	-0.0081	0.9700	0.1336	0.4406	-0.0054	0.9955	-0.0812	0.7101	-0.0758	0.7039	-0.0895	0.9365	0.0035	0.9972	0.0931	0.8368
---	---	1623730_at	0.2537	0.2143	0.1398	0.4823	0.0561	0.7378	0.0236	0.9777	0.1677	0.4352	0.1441	0.4629	0.0864	0.8692	-0.0265	0.9377	-0.1129	0.6186
trk	trunk	1623731_at	0.0864	0.7938	-0.1137	0.5142	-0.0273	0.9477	-0.0268	0.9847	0.3085	0.3400	0.3353	0.2396	-0.2410	0.7826	0.0621	0.9157	0.3030	0.4356
CG31410	CG31410	1623732_at	0.4909	0.3751	0.0536	0.6273	-0.0162	0.9729	-0.1324	0.8796	-0.1233	0.7240	0.0091	0.9804	-0.2410	0.9174	-0.6175	0.4410	-0.3765	0.6666
Nmd3	transcription unit I	1623733_at	0.3700	0.1122	0.0959	0.8266	0.5600	0.0048	0.1019	0.8500	0.1254	0.5535	0.0235	0.9226	-0.0396	0.9816	-0.0095	0.9917	0.0301	0.9574
m6	E(spl) region trans	1623734_at	-0.2062	0.1969	0.0375	0.8147	0.0599	0.7445	-0.0370	0.9376	-0.0903	0.5436	-0.0534	0.7169	-0.0050	0.9964	0.1145	0.7200	0.1196	0.7011
E5	E5	1623735_at	-0.4649	0.0813	0.0992	0.5985	-0.4048	0.1047	-0.3178	0.4532	-0.4633	0.0491	-0.1455	0.5131	0.2323	0.7712	0.2078	0.5622	-0.0245	0.9612
trh	trachealess	1623736_at	-0.1873	0.8182	-0.1498	0.7402	-0.5439	0.0898	-0.2941	0.8507	-0.0207	0.9809	0.2734	0.6391	0.0524	0.9870	0.0227	0.9884	-0.0297	0.9809
Rab14	Rab-protein 14	1623737_a_at	-0.7287	0.0175	-0.2121	0.4737	-0.0224	0.9118	0.0633	0.9136	-0.6515	0.0044	-0.7148	0.0018	-0.0385	0.9816	-0.1086	0.8297	-0.0701	0.8932
---	---	1623738_at	0.2194	0.2707	0.0708	0.6226	0.3327	0.0409	-0.0507	0.9523	-0.2257	0.3459	-0.1750	0.4281	-0.0451	0.9611	-0.0205	0.9620	0.0246	0.9463
Karybeta3	Karybeta3	1623739_at	-0.0330	0.8836	0.1439	0.2411	0.1224	0.4875	0.0740	0.8640	-0.2667	0.0835	-0.3407	0.0213	0.0883	0.8800	-0.1014	0.6847	-0.1897	0.4051
CG32154	CG32154	1623740_at	-2.3186	0.0012	-0.1175	0.7905	-1.1864	0.0270	-0.5824	0.6338	-2.4051	0.0026	-1.8227	0.0053	0.3156	0.8424	-0.2985	0.6794	-0.6141	0.3576
CG13981 /// scu	CG13981 /// scully	1623741_at	0.5440	0.1665	0.2164	0.4010	0.5531	0.0046	0.1671	0.7675	-0.3370	0.1591	-0.5042	0.0270	-0.1836	0.8956	-0.6831	0.1769	-0.4994	0.3526
CG4813	CG4813	1623742_at	0.3518	0.3005	0.1734	0.3802	0.0350	0.8452	0.2832	0.2596	0.5571	0.0035	0.2739	0.0451	0.2619	0.8222	0.2685	0.6032	0.0066	0.9939
CG3191	CG3191	1623743_at	-0.2489	0.4978	0.3107	0.5470	0.7828	0.0050	0.0830	0.9136	-1.2404	0.0009	-1.3234	0.0004	-0.5158	0.7215	-0.6171	0.2880	-0.1013	0.9025
CG30411	CG30411	1623744_at	3.4325	0.0038	2.8987	0.0011	4.4450	0.0000	0.0655	0.9854	-0.7264	0.3792	-0.7918	0.2781	-1.5648	0.1902	-1.3782	0.0490	0.1866	0.8080
CG31528	CG31528	1623745_at	0.1175	0.6080	-0.0293	0.8520	0.2299	0.1450	0.1082	0.8384	0.0368	0.8910	-0.0713	0.7385	-0.0113	0.9943	-0.0751	0.8884	-0.0637	0.9004
PICK1	PICK1	1623746_a_at	0.3994	0.0306	0.1402	0.4449	0.8430	0.0014	-0.1017	0.8190	-0.0972	0.6125	0.0045	0.9840	-0.5486	0.2553	-0.1035	0.7056	0.4450	0.1013
CG5895	CG5895	1623747_at	-1.7552	0.0072	-1.5924	0.0077	-2.2743	0.0004	-0.7103	0.2975	-0.8626	0.0378	-0.1524	0.7208	-0.1315	0.9380	-0.7465	0.1484	-0.6151	0.2593
---	---	1623748_at	0.2321	0.3305	0.0212	0.8750	0.0985	0.6959	0.0349	0.9622	0.1095	0.6117	0.0746	0.7204	-0.0108	0.9914	-0.1299	0.5854	-0.1192	0.6224
---	---	1623749_at	0.1479	0.6055	0.1798	0.2486	0.0714	0.7065	-0.1347	0.8244	-0.1774	0.4710	-0.0427	0.8783	0.0881	0.9333	-0.0631	0.9019	-0.1513	0.6853
CG32246	CG32246	1623750_at	0.0963	0.6849	0.0221	0.8888	-0.0077	0.9837	0.0061	0.9956	0.0475	0.8791	0.0414	0.8814	0.0744	0.9428	-0.0466	0.9266	-0.1210	0.7425
---	---	1623751_at	0.0173	0.9192	-0.0437	0.6559	0.0250	0.8909	0.0647	0.9186	0.1961	0.3203	0.1314	0.4793	-0.0692	0.9342	0.0661	0.8592	0.1353	0.6394
CG5959	CG5959	1623752_at	0.1991	0.3055	0.0441	0.8606	0.0423	0.8832	0.0910	0.8942	0.2755	0.2257	0.1844	0.3819	0.1160	0.8558	0.0262	0.9505	-0.0898	0.7697
TepIII	Thiolester contain	1623753_at	-0.6661	0.0704	-1.3993	0.1103	-1.0205	0.0003	0.5590	0.2631	0.7615	0.0182	0.2025	0.4709	0.1740	0.9301	0.0662	0.9496	-0.1077	0.9031
Qm	Ribosome RL10	1623754_s_at	0.0521	0.7145	0.1682	0.1351	0.1841	0.2367	0.0041	0.9956	-0.0665	0.7195	-0.0705	0.6699	0.0230	0.9737	0.0415	0.8631	0.0185	0.9390
Ten-a	Tenascin accesso	1623755_at	-2.2261	0.0136	-0.1204	0.8396	-1.9397	0.0002	-1.4389	0.0433	-1.6875	0.0021	-0.2486	0.5239	0.3486	0.9137	0.1708	0.9225	-0.1778	0.9089
CG17239 /// DyakCG17239	CG17239	1623756_at	2.2918	0.0013	0.7426	0.0079	4.9153	0.0000	3.5900	0.0008	1.9385	0.0009	-1.6516	0.0010	0.0567	0.9238	0.4982	0.0365	0.4416	0.0613
Cry	drosocrystallin	1623757_at	-0.0628	0.7953	-0.3702	0.1419	-0.1824	0.3743	0.3252	0.3863	0.3362	0.1126	0.0110	0.9675	-0.0879	0.8609	-0.1555	0.4429	-0.0675	0.7774
CG13123	CG13123	1623758_at	0.5335	0.0211	0.2005	0.6208	0.2703	0.2089	-0.0193	0.9860	0.3447	0.1759	0.3640	0.1107	0.0254	0.9884	0.1285	0.7876	0.1031	0.8305
CG13428	CG13428	1623759_at	0.3484	0.5816	0.0472	0.7583	0.1999	0.3699	-0.0985	0.7929	-0.0680	0.6986	0.0305	0.8637	-0.2176	0.9294	-0.4949	0.5503	-0.2773	0.7631
Ugt86De	Ugt86De	1623760_at	0.0257	0.8880	-0.0732	0.4365	0.0908	0.5461	0.2042	0.5008	0.1442	0.3814	-0.0600	0.7273	-0.0770	0.8875	-0.1094	0.6170	-0.0323	0.9089
CG15347 /// DyakCG15347	CG15347	1623761_at	1.3192	0.0088	0.6848	0.0347	0.3688	0.4062	0.7396	0.3969	1.7874	0.0039	1.0478	0.0250	1.1146	0.2553	1.2725	0.0343	0.1579	0.7909
key	kenny	1623762_at	0.2692	0.2699	-0.3844	0.0250	-0.4259	0.1483	0.1797	0.8028	0.7601	0.0165	0.5805	0.0317	0.2639	0.8097	0.1355	0.8190	-0.1284	0.8218
CG14274	CG14274	1623763_at	0.2775	0.2754	0.2177	0.3153	0.1290	0.7009	-0.0052	0.9956	0.0759	0.7676	0.0811	0.7220	0.1664	0.8412	0.1252	0.7594	-0.0412	0.9316
Or85d	Odorant receptor	1623764_at	-0.0377	0.8479	-0.0731	0.5040	0.0452	0.8119	-0.0229	0.9767	-0.1206	0.5627	-0.0976	0.6183	-0.1310	0.8202	-0.0179	0.9634	0.1131	0.6607
CG31606	CG31606	1623765_at	0.1478	0.3869	0.0021	0.9884	-0.0338	0.8469	0.0269	0.9704	0.1786	0.3419	0.1518	0.3740	0.0834	0.8768	0.0434	0.8864	-0.0401	0.8896
CG14054	CG14054	1623766_at	-0.0774	0.7304	-0.2623	0.1722	0.0528	0.7997	0.4244	0.1703	0.4105	0.0358	-0.0139	0.9535	0.0081	0.9943	0.0703	0.8484	0.0622	0.8608
CG17839	CG17839	1623767_at	-2.2779	0.0084	-1.5860	0.1266	-1.7161	0.0001	-0.2452	0.8738	-1.0061	0.0640	-0.7609	0.1112	-0.0949	0.9816	-0.3422	0.7456	-0.2474	0.8256
CG31840	CG31840	1623768_at	0.0754	0.6426	-0.0040	0.9829	-0.0371	0.8157	0.0197	0.9777	0.0846	0.6743	0.0649	0.7318	0.1764	0.8033	0.0757	0.8509	-0.1006	0.7723
CG7322	CG7322	1623769_at	2.6634	0.0008	1.5400	0.1422	2.2433	0.0002	0.9754	0.1125	0.4560	0.2134	-0.5194	0.1127	0.1944	0.9514	-0.6639	0.4907	-0.8583	0.3743
CG1673 /// DanaCG1673 ///	CG1673	1623770_at	1.7900	0.0900	1.7958	0.0126	3.3498	0.0019	1.0670	0.2501	0.5233	0.3517	-0.5436	0.2756	-0.1905	0.9742	0.5906	0.7251	0.7811	0.6189
CHOp24	CHOp24	1623771_at	0.6541	0.0120	0.5838	0.0561	0.5992	0.0040	0.0525	0.9145	0.3698	0.0239	0.3174	0.0282	-0.0266	0.9875	0.3370	0.3697	0.3636	0.3552
mthl10	Mth-like 10	1623772_a_at	0.0100	0.9633	-0.0623	0.6072	-0.1362	0.4082	-0.0095	0.9904	-0.0940	0.6325	-0.0844	0.6405	-0.0559	0.9305	-0.1132	0.6090	-0.0572	0.8254
CG32082	CG32082	1623773_at	0.0086	0.9792	0.0214	0.9189	0.0087	0.9623	-0.1720	0.6763	-0.1815	0.3583	-0.0095	0.9689	0.0708	0.9368	0.0322	0.9440	-0.0386	0.9211
---	---	1623774_at	0.2281	0.2235	0.1534	0.2088	0.1156	0.4509	0.0038	0.9956	0.2068	0.2170	0.2030	0.1756	-0.0425	0.9589	0.0055	0.9924	0.0480	0.8813
CG31089	CG31089	1623775_at	0.1000	0.6576	-0.0655	0.6147	-0.1568	0.4028	0.0371	0.9592	0.0628	0.7847	0.0258	0.9103	-0.1110	0.8917	-0.0795	0.8494	0.0316	0.9416
dsx	intersex-62c	1623776_s_at	-0.3949	0.3598	0.7742	0.0126	0.0560	0.7869	-0.4550	0.1032	-0.4360	0.0164	0.0190	0.9252	0.2552	0.8479	0.7815	0.1489	0.5263	0.3552
CG1358	CG1358	1623777_s_at	-2.7369	0.0009	-6.0069	0.0021	-5.8319	0.0000	0.4555	0.6701	1.5220	0.0094	1.0665	0.0268	0.3072	0.8973	-1.4082	0.1174	-1.7154	0.0879
CG9162	CG9162	1623778_at	-0.0130	0.9563	-0.6029	0.0728	-0.0165	0.9715	0.0407	0.9803	0.7720	0.0604	0.7314	0.0484	-0.6040	0.3056	0.3308	0.2643	0.9348	0.0333
CG34420	CG5917	1623779_at	0.1013	0.5506	0.0504	0.7973	0.1766	0.2651	0.0588	0.9009	0.1130	0.4790	0.0543	0.7413	-0.1603	0.7893	-0.0902	0.7731	0.0701	0.8279

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1623780_at	-0.0365	0.8501	0.0073	0.9568	0.2607	0.1517	0.0278	0.9666	-0.0321	0.8920	-0.0600	0.7528	-0.1303	0.7726	0.0378	0.8981	0.1680	0.4003
---	---	1623781_at	-0.0319	0.8695	-0.0704	0.5348	0.0793	0.6375	0.0698	0.8791	0.0801	0.6484	0.0103	0.9587	-0.1148	0.8395	-0.0055	0.9924	0.1092	0.6677
SCAP	SCAP	1623782_at	0.0985	0.6886	-0.0503	0.7362	0.1156	0.4637	0.0184	0.9764	0.1452	0.3544	0.1268	0.3728	-0.1504	0.8222	-0.0186	0.9682	0.1319	0.6634
CG8489	CG8489	1623783_at	0.0414	0.8636	-0.1442	0.2397	-0.3064	0.0530	-0.0084	0.9909	0.1536	0.3405	0.1620	0.2558	0.0312	0.9812	-0.0604	0.8821	-0.0916	0.7849
CG5732	CG5732	1623784_at	-0.0128	0.9669	-0.1038	0.3764	0.1263	0.4386	0.2337	0.5969	0.0779	0.7718	-0.1557	0.4615	-0.0357	0.9611	-0.1068	0.6312	-0.0711	0.7677
pbl	PEBBLE	1623785_a_at	-0.9195	0.3800	-2.5004	0.0114	-2.9865	0.0001	-0.5812	0.0978	0.3976	0.0591	0.9788	0.0008	-0.0675	0.9914	-1.3220	0.3575	-1.2545	0.4011
---	---	1623786_at	0.0784	0.7484	-0.1207	0.4241	0.0978	0.6448	0.0425	0.9610	0.1392	0.5829	0.0967	0.6937	-0.1053	0.8525	-0.1057	0.6857	-0.0005	0.9992
CG7144	CG7144	1623787_at	0.2752	0.5476	1.2768	0.0239	1.1678	0.0012	0.1349	0.8628	0.0419	0.9142	-0.0930	0.7596	0.0882	0.9677	1.0846	0.0721	0.9965	0.1166
CG7461 /// DyakCG7461	CG7461	1623788_at	0.4474	0.0149	0.1418	0.2494	0.3291	0.0387	-0.0209	0.9761	0.0053	0.9837	0.0262	0.9003	-0.2520	0.5519	-0.2246	0.2369	0.0275	0.9194
CG31776	CG31776	1623789_at	0.2385	0.2430	0.0742	0.5807	0.1073	0.7180	-0.0912	0.9110	0.0010	0.9979	0.0922	0.7399	0.1865	0.8114	-0.0955	0.8215	-0.2820	0.3921
CG31901	CG31901	1623790_at	0.7694	0.7745	-0.4372	0.2402	-4.4920	0.0001	-3.1449	0.1119	-0.3057	0.8407	2.8392	0.0151	0.5238	0.9589	-1.6231	0.5989	-2.1469	0.4740
Rab9D /// Rab9Db /// Rab9	CG32673 /// CG3	1623791_s_at	-0.1459	0.3630	-0.3503	0.0493	-0.5818	0.1002	0.0220	0.9860	0.4411	0.1225	0.4191	0.1017	-0.0704	0.9238	0.1059	0.7012	0.1763	0.4854
Dhod	Dihydroorotate de	1623792_a_at	0.1706	0.3616	0.2556	0.3305	0.2398	0.1304	0.0175	0.9777	0.1840	0.2327	0.1665	0.2281	0.0349	0.9816	0.3526	0.3092	0.3177	0.3832
krz	kurtz	1623793_at	0.0374	0.9125	0.8556	0.2719	0.2293	0.1776	-0.2774	0.6531	-0.1812	0.5726	0.0962	0.7694	0.3716	0.8122	0.4661	0.4807	0.0946	0.9162
CG7928	CG7928	1623794_at	-0.0635	0.7832	-0.1239	0.3749	-0.1705	0.2427	-0.0161	0.9840	0.2557	0.1721	0.2718	0.1055	0.0670	0.9238	0.1719	0.4631	0.1050	0.6817
Cyp313a5	Cyp313a5	1623795_at	-0.0856	0.5570	0.0059	0.9752	-0.0189	0.9252	-0.0650	0.8975	-0.0398	0.8531	0.0252	0.8995	-0.0190	0.9862	0.0845	0.7886	0.1035	0.7155
CG12027	CG12027	1623796_at	0.0434	0.8527	0.1158	0.4637	0.2409	0.1733	0.0665	0.9106	-0.1047	0.6121	-0.1712	0.3156	0.0551	0.9503	0.0700	0.8445	0.0149	0.9706
CG5969	CG5969	1623797_at	-0.2221	0.4976	0.0273	0.9579	-0.5480	0.0057	-0.3572	0.2177	-0.2293	0.1853	0.1280	0.4341	0.0977	0.9623	-0.0081	0.9947	-0.1058	0.8932
CG5669 /// DereCG5669 ///	CG5669	1623798_at	0.1432	0.6506	0.0953	0.7516	0.0766	0.8123	-0.1208	0.9162	0.2009	0.6104	0.3217	0.3245	0.0031	0.9994	0.2339	0.6483	0.2309	0.6491
CG7557	CG7557	1623799_a_at	-0.2983	0.0990	-0.0084	0.9387	0.1734	0.2222	0.0663	0.9117	-0.1387	0.4859	-0.2049	0.2279	0.0123	0.9901	0.1376	0.5374	0.1253	0.5819
CG18682	CG18682	1623800_at	0.0122	0.9684	0.0474	0.6498	0.0225	0.9344	0.0313	0.9603	0.0913	0.6175	0.0600	0.7357	-0.1336	0.8841	-0.0404	0.9425	0.0932	0.8310
---	---	1623801_at	0.2672	0.1965	0.0765	0.8477	-0.2827	0.2920	-0.1941	0.5353	0.0243	0.9154	0.2184	0.1328	0.2643	0.8875	0.2200	0.8070	-0.0442	0.9661
CG1625	CG1625	1623802_at	0.4249	0.2608	0.3769	0.4514	0.0759	0.6891	-0.3970	0.5681	-0.2476	0.5123	0.1495	0.6902	-0.2004	0.8680	-0.3081	0.5342	-0.1077	0.8641
---	---	1623803_at	-0.0681	0.6676	-0.0617	0.6389	-0.1028	0.6255	0.0825	0.8640	0.0544	0.7961	-0.0281	0.8908	0.0486	0.9683	0.0406	0.9379	-0.0080	0.9874
hep	hemipterous	1623804_a_at	1.0616	0.0921	0.4593	0.5767	-0.4364	0.4477	-0.0186	0.9901	1.5542	0.0014	1.5728	0.0008	0.8671	0.7770	0.8931	0.5160	0.0260	0.9910
CG4875	CG4875	1623805_at	0.6354	0.0110	0.0026	0.9956	-0.0766	0.8024	0.1017	0.8219	0.4578	0.0176	0.3561	0.0313	0.1988	0.8608	-0.1447	0.8039	-0.3436	0.4653
Ada2b	Transcriptional ad	1623806_at	0.0244	0.9225	0.4293	0.1085	0.3888	0.1101	-0.0293	0.9672	-0.2592	0.1617	-0.2299	0.1660	0.0946	0.9404	0.1685	0.7012	0.0739	0.8903
milt	Milton	1623807_a_at	-1.0442	0.0306	-0.4127	0.4646	-0.3195	0.1336	0.0596	0.9512	-0.3518	0.1884	-0.4114	0.0875	-0.0885	0.9727	0.2738	0.7092	0.3623	0.6038
---	---	1623808_s_at	0.1624	0.4955	-0.0957	0.5054	0.1943	0.3205	0.1626	0.7664	0.0899	0.7451	-0.0727	0.7794	0.0643	0.9555	-0.0131	0.9841	-0.0775	0.8608
CG1354	CG1354	1623809_s_at	0.6280	0.0049	1.1516	0.0077	1.4171	0.0001	0.0030	0.9956	-0.4264	0.0162	-0.4293	0.0096	-0.1048	0.8400	0.1795	0.3972	0.2843	0.2059
CG17265	CG17265	1623810_at	-1.4368	0.0020	-0.9377	0.0521	-1.0582	0.0009	-0.3120	0.5280	-0.8826	0.0052	-0.5706	0.0217	-0.2984	0.6927	-0.5373	0.0933	-0.2389	0.4568
mRpL18	mitochondrial ribo	1623811_at	-0.1043	0.7121	1.0858	0.0314	1.2968	0.0014	-0.3490	0.2968	-0.8780	0.0017	-0.5290	0.0092	-0.4607	0.7423	0.3652	0.5539	0.8259	0.1875
CG10943	CG10943	1623812_at	0.0772	0.8923	-0.1663	0.2552	-0.1525	0.4592	0.0074	0.9956	0.0246	0.9630	0.0172	0.9688	-0.0469	0.9862	-0.2402	0.7341	-0.1933	0.7907
CG33307	CG33307	1623813_at	2.6427	0.0004	1.0310	0.1856	2.0128	0.0003	0.3471	0.6533	0.9483	0.0191	0.6012	0.0731	-0.5242	0.7611	-0.6874	0.3275	-0.1631	0.8640
CG13829	CG13829	1623814_at	0.0681	0.7560	0.0437	0.7992	-0.2186	0.3736	-0.0211	0.9835	0.1470	0.5602	0.1681	0.4460	0.1951	0.7611	0.1955	0.4740	0.0004	0.9992
CG8100	CG8100	1623815_at	0.3137	0.0969	0.0916	0.4600	0.2132	0.1783	0.0257	0.9658	0.0037	0.9879	-0.0220	0.9086	0.1140	0.8461	0.0104	0.9821	-0.1036	0.7045
CG1943 /// DyakCG1943	CG1943	1623816_s_at	-0.3266	0.0688	-0.1092	0.5510	-0.4403	0.0737	0.1585	0.6013	0.0855	0.6121	-0.0730	0.6419	0.5041	0.6483	0.2105	0.6676	-0.2936	0.5299
CG7916	CG7916	1623817_at	-0.3096	0.4404	-0.0799	0.5039	-0.3050	0.1306	-0.2142	0.8550	-0.3207	0.4693	-0.1065	0.8253	0.0913	0.9592	-0.1011	0.8952	-0.1924	0.7455
---	---	1623818_at	-0.1376	0.4147	-0.1986	0.3416	0.1300	0.4742	0.0728	0.9165	-0.0380	0.8979	-0.1108	0.6122	-0.2411	0.6575	-0.1156	0.6291	0.1255	0.5988
RanGap	Segregation disto	1623819_at	0.1250	0.6569	-0.0649	0.7848	0.1840	0.3437	-0.1881	0.5363	0.4008	0.0200	0.5889	0.0022	-0.3994	0.7233	0.2253	0.6793	0.6248	0.2189
CG12252	CG12252	1623820_at	0.2839	0.1825	-0.1744	0.4151	-0.2277	0.1910	-0.0875	0.8676	0.6413	0.0050	0.7289	0.0017	-0.1275	0.8795	0.1868	0.5855	0.3143	0.3444
CG12011	CG12011	1623821_at	-0.2111	0.8521	0.0219	0.9381	-0.0461	0.8599	-0.3442	0.9011	-0.6695	0.4754	-0.3254	0.7361	-0.1514	0.8815	-0.3214	0.4037	-0.1700	0.6942
---	---	1623822_at	-0.0650	0.6819	-0.0962	0.5149	-0.0287	0.8947	-0.0574	0.9353	0.0492	0.8566	0.1065	0.6138	-0.0131	0.9912	0.0133	0.9739	0.0264	0.9371
CG11227	CG11227	1623823_a_at	0.1418	0.4548	-0.0311	0.8126	-0.0866	0.6362	0.0876	0.8512	0.1997	0.2404	0.1121	0.4910	0.0104	0.9914	-0.0513	0.8584	-0.0616	0.8102
CG17242	CG17242	1623824_at	0.1543	0.6199	0.0348	0.7590	0.1197	0.5483	0.1239	0.8640	0.0780	0.8045	-0.0460	0.8800	0.0686	0.9361	-0.0047	0.9938	-0.0733	0.8299
CG17018	CG17018	1623825_s_at	0.2212	0.8601	-2.6343	0.1953	-2.8377	0.0185	-0.1938	0.8096	3.3473	0.0001	3.5412	0.0000	-0.1185	0.9914	0.2395	0.9538	0.3580	0.9197
CHIP	CHIP	1623826_at	-0.4969	0.0722	-0.2220	0.3584	-0.1752	0.2370	-0.0889	0.9089	-0.1863	0.4721	-0.0974	0.7109	-0.2358	0.6955	-0.0452	0.9064	0.1906	0.4620
fra	frazzled	1623827_a_at	0.0061	0.9862	-0.5334	0.2832	-0.7265	0.0026	-0.3339	0.3102	0.2603	0.1749	0.5942	0.0050	-0.0503	0.9848	-0.1469	0.8556	-0.0966	0.9057
---	---	1623828_at	-0.0636	0.7145	-0.0687	0.5878	0.0457	0.8082	0.0364	0.9610	-0.0132	0.9659	-0.0496	0.8276	-0.0105	0.9935	0.0326	0.9451	0.0430	0.9155
CG31414	CG31414	1623829_at	0.1666	0.3224	0.0553	0.5894	0.2612	0.1333	0.0727	0.8823	0.0115	0.9643	-0.0612	0.7349	0.0787	0.9264	0.0450	0.9178	-0.0337	0.9323
CG13350 /// DereCG13350	CG13350	1623830_at	0.1421	0.6779	-0.2742	0.4543	-0.4053	0.0744	-0.2211	0.6591	0.5371	0.0330	0.7582	0.0045	-0.3125	0.8378	0.2672	0.7061	0.5797	0.3680
---	---	1623831_x_at	-0.0904	0.8228	-2.5213	0.0138	-1.8936	0.0027	0.2808	0.5832	1.6802	0.0004	1.3994	0.0005	-0.5881	0.7233	-0.4499	0.5514	0.1382	0.8891
---	---	1623832_a_at	0.1543	0.4440	0.0950	0.4883	0.1819	0.3143	-0.0010	0.9988	0.0728	0.6932	0.0739	0.6570	0.0142	0.9895	-0.0613	0.8406	-0.0755	0.7774
CG7309	CG7309	1623833_at	0.2284	0.3346	0.0330	0.8184	0.2366	0.1452	0.0619	0.9149	0.0801	0.7015	0.0183	0.9353	-0.0531	0.9515	-0.0154	0.9723	0.0377	0.9173
CG18641	CG18641	1623834_at	0.0504	0.9280	0.1815	0.4690	0.4096	0.0237	0.2866	0.6932	-0.1574	0.6808	-0.4440	0.1373	-0.0686	0.9588	-0.0765	0.8863	-0.0080	0.9897
CG6592	CG6592	1623835_at	0.1524	0.4084	0.0520	0.6777	0.4743	0.0167	0.0181	0.9778	0.0627	0.7535	0.0446	0.8134	-0.0564	0.9400	0.1473	0.5399	0.2037	0.3859
RpL13	r-protein L13	1623836_at	0.0543	0.7168	0.3849	0.0247	0.5998	0.0192	0.1768	0.7375	-0.2983	0.1991	-0.4751	0.0303	0.0040	0.9952	-0.0248	0.9269	-0.0289	0.9025
Pez	Pez	1623837_at	0.2045	0.6486	-0.0700	0.8667	-0.3799	0.1432	0.0525	0.9401	0.6061	0.0093	0.5537	0.0084	0.4380	0.7683	0.2895	0.6803	-0.1484	0.8570
CG4629	CG4629	1623838_s_at	0.6459	0.1351	-0.0005	1.0000	1.6347	0.0093	0.7705	0.0316	0.7993	0.0025	0.0288	0.9034	-0.7292	0.8202	0.0450	0.9868	0.7742	0.5846
CG6415	CG6415	1623839_at	2.7701	0.0030	1.3242	0.1392	3.6555	0.0001	1.2878	0.1316	0.5995	0.2424	-0.6884	0.1309	-1.1063	0.5754	-0.8584	0.3208	0.2479	0.8237
CG18607	CG18607	1623840_at	0.3369	0.4671	0.2552	0.2908	0.2198	0.1908	0.0292	0.9774	-0.2384	0.3560	-0.2676	0.2387	0.1600	0.9361	-0.3041	0.6605	-0.4641	0.4776
CG15169	CG15169	1623841_at	0.0179	0.9351	0.0314	0.8670	-0.1962	0.3944	0.0404	0.9558	0.0584	0.8094	0.0180	0.9410	0.2992	0.7152	0.0828	0.8625	-0.2165	0.5499
PhKgamma	phosphorylase kin	1623842_a_at	0.7843	0.0460	-0.1268	0.6307	0.0585	0.8581	0.0720	0.9182	0.0897	0.7202	0.0177	0.9482	-0.1281	0.9342	-0.8370	0.1012	-0.7089	0.1801
CG5903	CG5903	1623843_at	-0.6680	0.0084	-0.3405	0.0953	-0.5039	0.0204	-0.0493	0.9515	-0.4980	0.0332	-0.4487	0.0323	0.0930	0.8956	-0.1981	0.4565	-0.2911	0.2855
elfless	elfless	1623844_at	0.1390	0.4853	0.0431	0.6832	0.1939	0.1921	0.0044	0.9956	-0.0476	0.8857	-0.0520	0.8560	-0.0005	0.9998	-0.0794	0.8059	-0.0790	0.7947
---	---	1623845_s_at	0.0884	0.5764	-0.1160	0.5963	-0.0988	0.6685	-0.1785	0.6844	0.0861	0.7160	0.2646	0.1457	-0.1885	0.7423	-0.1413	0.5825	0.0472	0.8877
disco-r	disco-related	1623846_at	-3.7786	0.0004	-3.9426	0.0022	-4.5849	0.0001	-0.0132	0.9952	0.7428	0.0987	0.7560	0.0636	-0.1206	0.9514	-0.2452	0.7162	-0.1246	0.8761
CG11906	Gene 6	1623847_at	0.1966	0.2678	0.3122	0.1177	0.3406	0.0656	0.0792	0.9042	0.1888	0.3898	0.1096	0.6132	0.1279	0.8042	0.3449	0.1130	0.2170	0.3221
---	---	1623848_a_at	0.1226	0.7953	-0.0430	0.9158	0.1724	0.6412	0.1717	0.6086	0.3031	0.0743	0.1314	0.4018	-0.3126	0.8717	-0.0527	0.9664	0.2599	0.7744

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG6726	CG6726	1623849_s_at	-0.5312	0.4339	0.3835	0.1538	0.1872	0.3281	-0.5043	0.5255	-1.2121	0.0106	-0.7078	0.0610	-0.0568	0.9831	-0.1120	0.9075	-0.0551	0.9503
---	---	1623850_at	0.0774	0.7265	-0.0251	0.8194	-0.0075	0.9797	0.0063	0.9949	0.1367	0.5210	0.1303	0.4985	0.0657	0.9503	0.0491	0.9198	-0.0166	0.9727
CG9400	CG9400	1623851_at	-1.4455	0.0552	0.0124	0.9864	-2.1524	0.0168	-1.8675	0.1022	-3.5839	0.0008	-1.7164	0.0119	0.6361	0.8145	-2.0280	0.0825	-2.6640	0.0559
CG10407	CG10407	1623852_at	-0.2953	0.1884	-0.5195	0.1755	-0.6924	0.0033	-0.0974	0.8300	-0.2783	0.1083	-0.1809	0.2461	-0.0318	0.9816	-0.4331	0.1504	-0.4013	0.2130
betaggt-I	type I Geranylgera	1623853_at	-0.5137	0.0462	0.5726	0.1221	1.2990	0.0003	0.2861	0.4751	-0.5648	0.0169	-0.8509	0.0016	-0.2623	0.7997	0.6075	0.1495	0.8698	0.0812
---	---	1623854_at	0.2393	0.1477	0.0184	0.8801	0.1166	0.5920	0.1578	0.7388	0.4965	0.0255	0.3388	0.0719	-0.0550	0.9233	-0.0580	0.8123	-0.0030	0.9924
CG11660	CG11660	1623855_s_at	0.2461	0.3483	-0.5319	0.3744	-0.0840	0.7074	0.1030	0.9185	1.0146	0.0062	0.9116	0.0060	-0.1049	0.9636	0.2536	0.7251	0.3585	0.5929
CG30395	CG30395	1623856_at	0.2622	0.1451	0.0979	0.4542	0.1468	0.3326	0.0258	0.9727	0.1799	0.3453	0.1541	0.3734	-0.0539	0.9267	-0.0867	0.6883	-0.0328	0.9025
Ranbp16	Ranbp16	1623857_at	0.0826	0.6287	-0.4202	0.1336	-0.4398	0.0184	-0.0651	0.9039	0.4486	0.0188	0.5137	0.0062	-0.0101	0.9928	-0.0162	0.9698	-0.0061	0.9880
NP15.6	NP15.6	1623858_at	-0.0354	0.9132	0.3392	0.2338	0.4037	0.0335	-0.0651	0.9311	-0.8493	0.0031	-0.7842	0.0026	0.0849	0.9619	-0.3318	0.5192	-0.4167	0.4095
mRpL49	mitochondrial ribo	1623859_at	-0.3826	0.1901	-0.2569	0.5425	-0.2585	0.2775	-0.0349	0.9648	-0.1315	0.5601	-0.0967	0.6551	0.0405	0.9862	-0.0040	0.9985	-0.0445	0.9533
CG15478	CG15478	1623860_at	0.2523	0.2016	0.0625	0.5802	-0.0507	0.8127	-0.0695	0.9018	0.0379	0.8762	0.1074	0.5504	-0.0762	0.9144	-0.1228	0.6508	-0.0466	0.8918
Eip55E	Ecdysone-induce	1623861_at	0.7892	0.0126	1.4512	0.0129	1.4011	0.0006	0.2012	0.6917	-0.0292	0.9296	-0.2304	0.2777	0.4242	0.6955	0.5903	0.1718	0.1662	0.7458
ninaA	neither inactivatio	1623862_at	0.1120	0.5383	-0.0704	0.5583	-0.0314	0.8647	0.0485	0.9422	0.0053	0.9849	-0.0432	0.8444	0.0361	0.9657	-0.1872	0.4046	-0.2233	0.3369
Tis11	Tis11 homolog	1623863_a_at	0.1592	0.7017	0.0888	0.6364	-0.1243	0.7171	-0.0332	0.9603	-0.1290	0.4888	-0.0957	0.5905	0.1526	0.9460	-0.1156	0.9125	-0.2682	0.7332
CG6321	CG6321	1623864_at	-0.3894	0.2315	-0.1444	0.5657	-0.5880	0.0021	-0.5278	0.1875	-0.3105	0.1974	0.2173	0.3250	-0.1306	0.8924	-0.0419	0.9425	0.0887	0.8484
---	---	1623865_at	0.0564	0.7167	0.2662	0.0993	0.2726	0.0818	-0.0376	0.9438	-0.2401	0.1145	-0.2025	0.1361	0.0211	0.9775	-0.0281	0.9193	-0.0492	0.8256
Cyp4ac2	Cyp4ac2	1623866_at	2.2786	0.0946	0.7935	0.5425	1.8758	0.0033	0.7608	0.0925	0.6802	0.0185	-0.0806	0.7860	-0.1564	0.9877	-0.8425	0.7572	-0.6862	0.8059
CG14168	CG14168	1623867_at	0.0026	0.9902	0.1333	0.2942	0.1241	0.6132	0.0928	0.8738	-0.0344	0.9026	-0.1272	0.5197	0.0885	0.8689	0.0729	0.7798	-0.0156	0.9596
CG30275	CG30275	1623868_a_at	-0.0498	0.8310	-0.0989	0.4045	-0.0812	0.6905	0.0154	0.9803	0.1024	0.5276	0.0870	0.5629	0.0192	0.9848	-0.0450	0.8967	-0.0643	0.8253
CtBP	C-terminal Bindin	1623869_at	-0.2003	0.5707	0.3416	0.3130	0.0913	0.8239	0.0247	0.9857	-0.2665	0.4142	-0.2912	0.3116	0.2495	0.8882	0.3333	0.6450	0.0838	0.9278
---	---	1623870_at	0.2443	0.1666	0.0939	0.4239	0.1529	0.3340	0.0798	0.8942	0.0273	0.9223	-0.0525	0.8165	0.0637	0.9095	-0.0297	0.9248	-0.0934	0.6670
CG18563	CG18563	1623871_at	0.8187	0.0855	-0.5269	0.4974	-0.0597	0.9348	0.7011	0.1119	1.2319	0.0013	0.5308	0.0314	0.2456	0.9457	-0.0458	0.9827	-0.2914	0.8325
CG13673	CG13673	1623872_at	0.2449	0.2124	-0.0728	0.7007	-0.0719	0.7431	-0.1374	0.7931	0.0111	0.9734	0.1485	0.4641	-0.0148	0.9911	-0.1834	0.5071	-0.1687	0.5512
Stim	Drosophila Strom	1623873_at	-0.4561	0.0228	-0.4370	0.1672	-0.4298	0.0310	-0.1695	0.5842	-0.0833	0.6374	0.0862	0.5847	-0.2616	0.7707	-0.1651	0.7034	0.0965	0.8444
CG14215	CG14215	1623874_at	0.3256	0.5576	-0.1682	0.8243	0.2547	0.4024	0.2889	0.4861	0.7119	0.0069	0.4230	0.0393	-0.2427	0.9400	0.0824	0.9599	0.3251	0.7815
CG3434 /// DwilCG3434	CG3434	1623875_at	0.4273	0.0338	0.3809	0.2784	0.4149	0.0344	-0.0286	0.9576	0.2308	0.1054	0.2594	0.0477	-0.1109	0.9317	0.1161	0.8359	0.2270	0.6152
---	---	1623876_at	0.1972	0.3330	0.0989	0.6696	0.3333	0.0453	0.1238	0.6998	0.0629	0.7121	-0.0609	0.6940	-0.1448	0.7707	0.0248	0.9425	0.1696	0.4370
elB	elbow B	1623877_a_at	-0.0056	0.9952	1.0539	0.2198	1.2387	0.0016	0.2903	0.6480	-0.5344	0.0836	-0.8247	0.0094	0.1823	0.9612	0.6194	0.5800	0.4370	0.7137
CG6123	CG6123	1623878_at	-0.0782	0.7170	-0.0572	0.7774	0.0159	0.9425	0.1122	0.8057	0.0696	0.7469	-0.0426	0.8387	-0.0245	0.9831	0.0551	0.8856	0.0795	0.8058
CG15764	CG15764	1623879_at	0.1301	0.4526	0.0555	0.7903	0.3660	0.0633	0.1706	0.5660	-0.0485	0.7951	-0.2191	0.1067	0.0189	0.9898	0.0308	0.9506	0.0118	0.9823
CG34422	CG7274	1623880_at	-0.2611	0.2201	-0.0196	0.8453	0.0943	0.6769	0.0977	0.9117	-0.1253	0.6977	-0.2230	0.3918	-0.1299	0.8692	-0.0340	0.9463	0.0960	0.8001
Sirt4	Sirt4	1623881_s_at	-0.3710	0.0894	-0.1017	0.7770	0.0791	0.6198	-0.1116	0.8189	-0.4810	0.0198	-0.3694	0.0369	-0.2261	0.7997	-0.2013	0.6122	0.0248	0.9640
CG14195	CG14195	1623882_at	0.0702	0.8695	0.0619	0.5709	0.1364	0.5393	0.0175	0.9838	-0.1620	0.4401	-0.1795	0.3317	-0.0383	0.9787	-0.0991	0.8208	-0.0608	0.8949
CG18661 /// DyakCG18661	CG18661	1623883_at	-1.2297	0.0207	-1.7971	0.0159	-1.5889	0.0006	0.3427	0.6394	0.3439	0.3449	0.0013	0.9981	0.1147	0.9093	-0.4301	0.2145	-0.5448	0.1538
CG16775	CG16775	1623884_at	0.2600	0.4899	-0.0219	0.8313	0.2416	0.3379	0.1645	0.8688	-0.0418	0.9325	-0.2063	0.5496	-0.1434	0.8438	-0.2592	0.3752	-0.1158	0.7353
alpha-Est1	fragment A	1623885_at	0.5450	0.1789	0.3312	0.7273	1.2233	0.0001	0.1527	0.8776	-0.4349	0.1994	-0.5876	0.0576	-0.7358	0.7220	-0.6133	0.4976	0.1226	0.9196
---	---	1623886_at	0.1045	0.6042	-0.0403	0.7090	0.0836	0.6242	0.1177	0.8350	0.1562	0.4856	0.0384	0.8795	-0.0056	0.9964	0.0783	0.8353	0.0839	0.8085
CG12367	CG12367	1623887_at	0.4284	0.2589	0.9581	0.0435	0.8452	0.0126	-0.4898	0.3941	-1.1365	0.0046	-0.6467	0.0327	-0.3423	0.7358	-0.5449	0.1940	-0.2025	0.6668
CG11847	CG11847	1623888_at	0.3075	0.1358	0.1796	0.6889	0.0026	0.9905	-0.0190	0.9774	0.7952	0.0012	0.8142	0.0006	0.1141	0.9400	0.7048	0.1297	0.5907	0.2189
CG1317	CG1317	1623889_at	-0.4351	0.1037	-0.6571	0.0638	-0.5981	0.0095	-0.0415	0.9466	0.0852	0.6693	0.1267	0.4427	-0.1087	0.9365	0.0661	0.9221	0.1748	0.7221
---	---	1623890_at	0.1461	0.4025	0.1351	0.2916	0.0915	0.5857	0.0049	0.9956	0.0046	0.9892	-0.0003	0.9994	-0.0199	0.9816	0.0098	0.9784	0.0296	0.9161
CSN7	Drosophila COP9	1623891_at	0.0769	0.6281	0.4105	0.0915	0.3814	0.0495	-0.0809	0.8470	-0.2029	0.1847	-0.1219	0.3937	0.1082	0.8906	0.1389	0.6714	0.0307	0.9412
Bsg25A	blastoderm-specif	1623892_at	0.1251	0.4787	0.3273	0.1961	0.3396	0.0747	-0.0425	0.9558	-0.1591	0.4618	-0.1167	0.5707	0.0226	0.9816	0.2057	0.3653	0.1831	0.4407
CG13784	CG13784	1623893_at	0.2993	0.4934	0.2195	0.5275	-0.2310	0.2226	-0.2490	0.4715	0.2055	0.2719	0.4545	0.0143	0.1651	0.9317	0.0517	0.9600	-0.1134	0.8947
CG8086	CG8086	1623894_a_at	0.2279	0.6355	0.3263	0.6725	0.7771	0.0018	0.2566	0.5570	-0.1711	0.4673	-0.4277	0.0402	-0.1989	0.9405	0.0823	0.9505	0.2812	0.7717
CG31912	CG31912	1623895_at	0.0899	0.7160	0.3822	0.3409	-0.0022	0.9922	0.0192	0.9860	0.0541	0.8736	0.0349	0.9105	0.3081	0.6749	0.3543	0.2239	0.0462	0.9118
CG34404	CG18496	1623896_a_at	-1.0946	0.0041	-0.3788	0.1037	-0.7441	0.0010	-0.0823	0.8732	-0.1597	0.3837	-0.0774	0.6797	0.3143	0.6749	0.3815	0.2005	0.0673	0.8699
---	---	1623897_at	0.0284	0.9283	-0.1749	0.3129	-0.0373	0.9190	0.2350	0.7007	0.4808	0.0877	0.2458	0.3391	-0.0681	0.9352	-0.0114	0.9821	0.0567	0.8751
---	---	1623898_at	0.1955	0.2122	-0.0054	0.9713	-0.0021	0.9924	-0.0727	0.8760	-0.0036	0.9884	0.0691	0.6814	-0.0009	0.9996	-0.0940	0.6844	-0.0931	0.6848
Takr99D	Tachykinin-like re	1623899_at	0.0345	0.8933	-0.2066	0.1579	-0.0631	0.7775	0.1563	0.7305	0.0306	0.9142	-0.1257	0.5243	-0.2021	0.6955	-0.1898	0.3668	0.0123	0.9701
CG14935	CG14935	1623900_a_at	1.7727	0.0030	1.1234	0.1160	2.5258	0.0003	0.5754	0.5636	-0.2197	0.7103	-0.7951	0.0834	-0.8227	0.6272	-0.7791	0.2507	0.0436	0.9670
sar1	sar1	1623901_s_at	0.3768	0.0299	1.3190	0.0087	1.4690	0.0002	0.0759	0.9116	-0.4773	0.0347	-0.5532	0.0118	-0.0643	0.9495	0.3725	0.2026	0.4368	0.1727
tsu	tsunagi	1623902_at	-0.2326	0.3792	-0.0157	0.9481	0.1904	0.4268	0.0026	0.9962	-0.2656	0.1529	-0.2682	0.1072	-0.1131	0.9411	0.1692	0.7692	0.2823	0.5757
CG18473	CG18473	1623903_at	1.0219	0.2773	-0.0585	0.9163	0.1143	0.4337	0.1519	0.9470	-0.4663	0.4902	-0.6182	0.2898	-0.0001	0.9999	-1.4263	0.1827	-1.4262	0.2136
Rpt3R	Rpt3R	1623904_a_at	0.0083	0.9697	-0.0801	0.6581	0.4163	0.0218	0.1472	0.6628	0.1848	0.2545	0.0376	0.8405	-0.1272	0.8541	0.1701	0.5684	0.2973	0.3061
---	---	1623905_s_at	0.0768	0.7820	0.2048	0.2516	0.1346	0.6011	-0.1249	0.8192	-0.2065	0.3433	-0.0816	0.7205	-0.0416	0.9742	-0.0530	0.9158	-0.0114	0.9837
Ripalpha	RPA-interacting p	1623906_at	-0.2445	0.3951	-0.2674	0.3736	-0.7288	0.0081	-0.6503	0.1582	-0.2708	0.3394	0.3795	0.1263	-0.2206	0.8400	-0.3547	0.4302	-0.1341	0.8095
armi	armitage	1623907_at	0.2589	0.5347	-0.5677	0.0755	-0.3193	0.3682	-0.2328	0.7596	0.5330	0.1026	0.7657	0.0169	-0.4700	0.7230	-0.2507	0.7000	0.2194	0.7416
CG32111	CG32111	1623908_at	0.3126	0.2429	-0.1379	0.5114	0.2961	0.3143	0.0133	0.9937	0.1195	0.7580	0.1062	0.7659	-0.1126	0.9096	-0.2319	0.5276	-0.1193	0.7753
inv	invected	1623909_s_at	-1.4632	0.0011	-0.1730	0.3377	-0.7017	0.0029	-0.7163	0.0492	-1.5505	0.0002	-0.8342	0.0015	-0.1441	0.8270	-0.2349	0.3850	-0.0908	0.7796
alphaTub85E	alpha-Tubulin	1623910_at	-0.7281	0.0822	0.4277	0.0497	0.0962	0.6486	-0.2503	0.7138	-1.0883	0.0047	-0.8380	0.0092	0.1028	0.8795	0.0607	0.8684	-0.0420	0.9067
---	---	1623911_at	0.0774	0.6988	0.0990	0.6147	0.0702	0.8277	0.1229	0.8281	0.0426	0.8840	-0.0803	0.7305	0.0031	0.9984	-0.0805	0.8353	-0.0835	0.8175
---	---	1623912_at	0.2065	0.4148	0.0566	0.7114	0.1486	0.5430	0.0284	0.9745	0.1489	0.5241	0.1205	0.5824	-0.1746	0.8097	-0.0290	0.9521	0.1456	0.6549
---	---	1623913_at	0.0231	0.9118	-0.0173	0.8979	-0.0861	0.5996	-0.0306	0.9659	0.0204	0.9406	0.0510	0.8112	0.0877	0.8814	0.1735	0.4379	0.0858	0.7401
RpL14	Minute(3)66D	1623914_at	0.4379	0.0797	1.4431	0.0060	1.4271	0.0002	0.1297	0.8446	-0.8087	0.0060	-0.9384	0.0019	0.2450	0.7464	0.0651	0.8893	-0.1799	0.6030
CG14511	CG14511	1623915_at	0.0001	0.9994	0.5428	0.0990	0.8306	0.0095	-0.0361	0.9745	-0.3166	0.2503	-0.2804	0.2563	-0.3604	0.7095	0.1939	0.6633	0.5544	0.1959
CG14694	CG14694	1623916_at	-1.5575	0.0791	-0.0736	0.7801	-0.8187	0.0606	-0.9777	0.2309	-1.6910	0.0048	-0.7133	0.0996	-0.1508	0.9657	-0.4082	0.7007	-0.2574	0.8274
Sh	Shaker	1623917_a_at	-0.1267	0.4452	-0.7330	0.2040	-0.8824	0.0142	-0.0078	0.9956	0.9416	0.0109	0.9494	0.0063	0.0727	0.9618	0.1361	0.8008	0.0634	0.9133

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Sras	severas	1623918_at	0.3744	0.1095	0.0549	0.7861	0.2398	0.1454	0.2509	0.4037	0.9757	0.0006	0.7249	0.0011	0.0303	0.9816	0.5693	0.0699	0.5390	0.1013
kuk	kugelkern	1623919_a_at	-1.0166	0.0833	-2.0201	0.0321	-1.9987	0.0021	-0.3300	0.4455	-0.0076	0.9826	0.3224	0.1237	-0.3843	0.8869	-1.0572	0.2807	-0.6729	0.5266
CG11775	CG11775	1623920_at	0.2269	0.2361	0.2967	0.2799	0.1604	0.2935	-0.1025	0.8671	-0.2129	0.3317	-0.1105	0.6137	0.1934	0.7644	-0.0437	0.9178	-0.2371	0.3953
rg	Drosophila A kina	1623921_at	0.0257	0.9219	-0.1136	0.4362	-0.1715	0.2393	-0.1284	0.8642	0.0360	0.9228	0.1645	0.5266	0.0187	0.9829	-0.0263	0.9341	-0.0451	0.8636
ect	Interferon-like pro	1623922_at	0.3225	0.2826	-0.1119	0.5749	0.0007	0.9977	0.0947	0.8385	0.1761	0.3177	0.0814	0.6515	0.0160	0.9916	-0.2409	0.5158	-0.2569	0.4875
gl	glass	1623923_a_at	0.0749	0.6545	-0.0831	0.4621	0.0801	0.5878	0.1733	0.6908	0.1791	0.3863	0.0058	0.9816	-0.1290	0.7550	-0.0201	0.9436	0.1089	0.5585
CG12325	CG12325	1623924_at	0.2187	0.5243	-0.1712	0.7562	-0.1141	0.7327	0.4076	0.5451	1.2468	0.0043	0.8393	0.0152	0.1875	0.9246	0.8064	0.2116	0.6190	0.3665
CG4477	CG4477	1623925_at	0.0929	0.6193	0.0174	0.8767	0.1946	0.1876	0.0468	0.9204	0.1282	0.3795	0.0814	0.5646	-0.0559	0.9238	-0.0087	0.9808	0.0472	0.8509
CG11604	CG11604	1623926_at	0.6376	0.0691	-0.4074	0.4267	0.0035	0.9946	0.1656	0.8599	1.2757	0.0033	1.1101	0.0037	-0.1557	0.9467	0.1916	0.8400	0.3474	0.6476
CG31925	CG31925	1623927_at	-0.1594	0.4848	0.1523	0.3736	0.0482	0.7644	-0.0263	0.9704	-0.2184	0.2271	-0.1921	0.2363	0.0375	0.9589	0.1234	0.5622	0.0859	0.7060
CG5910	CG5910	1623928_at	-0.4293	0.4676	-1.5221	0.0161	-0.7553	0.0538	-0.3008	0.5502	-0.1573	0.5769	0.1435	0.5804	-1.0307	0.6092	-1.2036	0.1517	-0.1729	0.8844
---	---	1623929_at	0.2725	0.2600	0.1175	0.7384	0.1922	0.3947	0.2738	0.6457	0.2491	0.4016	-0.0248	0.9461	0.0897	0.8801	-0.0202	0.9569	-0.1099	0.6553
---	---	1623930_s_at	0.3140	0.2329	0.1055	0.5030	0.3141	0.2343	0.0732	0.9314	0.0991	0.7402	0.0259	0.9344	-0.0426	0.9754	-0.1492	0.6955	-0.1066	0.7938
CanA-14F	protein phosphata	1623931_s_at	-1.1425	0.0361	0.5474	0.2114	0.6800	0.0052	-0.0657	0.9633	-1.0217	0.0149	-0.9560	0.0124	-0.0115	0.9964	0.7827	0.2217	0.7943	0.2497
Ent3	Equilibrative nucle	1623932_at	-0.3604	0.2204	0.0651	0.8071	-0.4298	0.1764	0.1964	0.7128	0.0315	0.9260	-0.1649	0.4755	0.1873	0.8686	-0.0002	1.0000	-0.1875	0.7096
Sh	Shaker	1623933_at	-0.7643	0.1451	-1.2041	0.0683	-1.9284	0.0002	-0.2151	0.8815	0.2548	0.6447	0.4698	0.2916	-0.0070	0.9984	-0.2555	0.7707	-0.2485	0.7677
blp	black pearl	1623934_at	0.1694	0.4980	-0.0342	0.8796	-0.2521	0.0971	-0.2018	0.6916	0.0797	0.7798	0.2815	0.1801	-0.0149	0.9914	-0.1230	0.7456	-0.1081	0.7764
sxe2	sex-specific enzyr	1623935_at	0.2206	0.3072	0.0464	0.7793	0.2057	0.2982	0.1529	0.6937	0.1043	0.5939	-0.0485	0.8103	-0.0204	0.9887	0.0600	0.8980	0.0804	0.8385
cl	clot	1623936_at	-0.1220	0.5568	0.4515	0.0726	0.8012	0.0015	0.0952	0.8220	-0.4784	0.0109	-0.5736	0.0029	-0.0289	0.9829	0.1152	0.7516	0.1441	0.6668
---	---	1623937_at	0.1441	0.4328	-0.1322	0.2362	0.0660	0.7604	0.0949	0.8535	0.2288	0.2168	0.1339	0.4440	-0.0632	0.9340	-0.0412	0.9121	0.0220	0.9486
---	---	1623938_at	-0.0101	0.9560	0.0378	0.7249	0.3366	0.0672	0.2067	0.5093	0.0392	0.8557	-0.1675	0.2620	-0.0965	0.8706	0.1043	0.6949	0.2009	0.4037
---	---	1623939_at	0.0947	0.6453	-0.0456	0.6642	0.0499	0.8430	0.0824	0.8671	0.3522	0.0480	0.2698	0.0835	-0.2426	0.7464	-0.1991	0.5455	0.0435	0.9198
CG9855	CG9855	1623940_at	-0.0258	0.9091	0.6828	0.1175	0.9071	0.0008	0.1246	0.7845	-0.2891	0.1221	-0.4137	0.0220	-0.2362	0.8141	0.4166	0.3026	0.6527	0.1438
---	---	1623941_at	0.0402	0.8807	-0.1184	0.4440	0.0130	0.9573	0.1715	0.8432	0.1294	0.7307	-0.0421	0.9143	-0.1450	0.7997	-0.2594	0.2584	-0.1144	0.6577
CG6974	CG6974	1623942_at	0.1086	0.5085	0.0012	0.9985	-0.1225	0.4475	0.1470	0.6872	0.2107	0.2131	0.0637	0.7249	-0.0920	0.9031	-0.0735	0.8438	0.0185	0.9637
CG17556	CG17556	1623943_at	0.0505	0.8294	0.3935	0.2610	0.0299	0.8832	-0.2075	0.6622	0.0898	0.7380	0.2972	0.1428	0.1291	0.8903	0.4182	0.2189	0.2891	0.4203
CG12605	CG12605	1623944_at	-0.0050	0.9789	-0.0194	0.8601	0.0313	0.8634	-0.0616	0.9009	-0.0535	0.7826	0.0080	0.9685	-0.1057	0.8626	-0.1397	0.5926	-0.0341	0.9197
CG3419	CG3419	1623945_at	0.2774	0.1583	0.3174	0.2799	0.5730	0.0048	-0.1125	0.7834	-0.1139	0.5272	-0.0014	0.9949	-0.3642	0.6868	-0.1573	0.7060	0.2069	0.6024
CG32970	CG32970	1623946_at	0.1920	0.3024	-0.0672	0.5719	-0.1288	0.5102	0.1458	0.6615	0.3341	0.0426	0.1883	0.1861	0.0480	0.9623	0.0346	0.9402	-0.0134	0.9764
Gr58c	Gustatory recepto	1623947_at	0.1132	0.5228	-0.0164	0.8839	0.2235	0.1351	0.2663	0.4529	0.3220	0.0946	0.0558	0.7967	-0.1203	0.8308	0.0749	0.8045	0.1952	0.4148
CG40174	CG40174	1623948_at	0.0302	0.8987	0.0563	0.6836	0.2478	0.1810	0.2615	0.5255	0.0368	0.9015	-0.2247	0.2463	-0.0799	0.9174	0.0322	0.9404	0.1121	0.7087
CG32816	CG32816	1623949_s_at	0.2858	0.3394	0.9166	0.0229	0.0804	0.8091	-0.4554	0.4690	-1.0378	0.0085	-0.5824	0.0585	0.6079	0.6092	-0.2657	0.6310	-0.8737	0.1192
Ama	amalgam	1623950_s_at	-0.4452	0.7769	-0.4700	0.1759	-0.2322	0.2447	0.3014	0.9481	-0.6737	0.6439	-0.9751	0.4225	-0.1349	0.9589	-0.5887	0.4255	-0.4538	0.5634
dpr2	dpr2	1623951_at	0.0106	0.9548	0.0992	0.4016	0.0572	0.8058	-0.0533	0.9413	-0.0964	0.6846	-0.0431	0.8589	-0.0335	0.9767	0.0792	0.8255	0.1127	0.7154
Su(z)12	Su(z)12	1623952_a_at	-0.2749	0.4229	-0.0725	0.8763	-0.1536	0.3322	-0.0990	0.8111	0.1216	0.4790	0.2206	0.1311	-0.0441	0.9831	0.3359	0.4881	0.3799	0.4353
CG11592	CG11592	1623953_at	0.2576	0.4612	0.4147	0.3500	0.2602	0.6803	-0.2109	0.7401	-0.1777	0.5557	0.0332	0.9232	-0.0791	0.9831	0.1696	0.8962	0.2487	0.8178
CoRest	CG33525	1623954_s_at	0.2667	0.6978	-0.4077	0.5655	-1.4673	0.0164	-1.1799	0.0500	0.5462	0.1134	1.7261	0.0006	-0.1487	0.9813	-0.0470	0.9863	0.1017	0.9585
CG5087	CG5087	1623955_at	-0.0059	0.9853	0.6324	0.0308	0.2540	0.1293	-0.1616	0.7187	-0.4892	0.0242	-0.3276	0.0735	0.1164	0.9056	0.2898	0.4056	0.1734	0.6469
pen	penguin	1623956_at	0.1669	0.5180	-0.1080	0.4321	0.7476	0.0078	0.4775	0.2511	0.5485	0.0342	0.0710	0.7989	-0.4722	0.5228	0.1289	0.7595	0.6011	0.1192
GstS1	glutathione-S-tran	1623957_s_at	2.2165	0.0005	1.3876	0.0040	1.8541	0.0001	0.3186	0.5455	0.4024	0.1392	0.0838	0.7832	-0.3054	0.6749	-0.5164	0.0912	-0.2111	0.4912
---	---	1623958_at	-0.0139	0.9332	-0.1533	0.3186	-0.1006	0.6045	0.0727	0.8723	0.1520	0.3427	0.0792	0.6207	-0.0707	0.9260	-0.0911	0.7667	-0.0204	0.9548
CG1532 /// DyakCG1532	CG1532	1623959_at	-1.0063	0.0027	-1.8038	0.0061	-1.4013	0.0016	0.2716	0.4511	0.6911	0.0045	0.4196	0.0247	0.0147	0.9946	-0.0273	0.9748	-0.0419	0.9525
---	---	1623960_s_at	1.4866	0.0012	1.1036	0.0654	1.6356	0.0003	0.1461	0.7958	0.3155	0.1672	0.1695	0.4324	-0.3187	0.8222	-0.0386	0.9684	0.2801	0.6621
Arpc3B	Arpc3B	1623961_at	1.9981	0.0015	1.0648	0.0069	1.2561	0.0005	0.3518	0.5808	1.0238	0.0077	0.6720	0.0286	0.1691	0.8513	0.2005	0.6156	0.0314	0.9523
---	---	1623962_x_at	0.0943	0.7425	0.1819	0.3227	0.0275	0.8781	-0.0645	0.9149	-0.0385	0.8789	0.0260	0.9099	-0.1605	0.8141	-0.0270	0.9526	0.1335	0.6623
CG32115	CG32115	1623963_at	1.8493	0.0421	0.2790	0.6553	1.5050	0.0059	0.3195	0.7647	0.3874	0.4087	0.0679	0.9011	-0.8928	0.6955	-1.0279	0.2686	-0.1352	0.9193
Or83c	Odorant receptor	1623964_at	0.2120	0.1385	-0.3622	0.1399	-0.2056	0.3764	0.0655	0.8937	0.3520	0.0373	0.2865	0.0535	-0.1436	0.8806	-0.1543	0.7090	-0.0106	0.9852
---	---	1623965_at	0.0302	0.9116	-0.0912	0.6071	0.2679	0.3070	0.2412	0.6673	0.1197	0.6976	-0.1215	0.6615	-0.2795	0.7230	0.0586	0.9105	0.3381	0.3307
Nrt	disabled	1623966_s_at	-0.5911	0.2505	-1.1600	0.1300	-0.5890	0.0694	0.4616	0.7098	0.5603	0.3277	0.0988	0.8823	-0.0726	0.9652	-0.1376	0.8117	-0.0650	0.9161
Shc	SHC-adaptor prot	1623967_at	-0.7478	0.0026	-0.1598	0.5065	0.0354	0.9261	0.2042	0.6471	-0.2583	0.2310	-0.4624	0.0248	0.0326	0.9877	0.2908	0.5591	0.2582	0.6140
CG10631	CG10631	1623968_at	0.1024	0.8212	0.1061	0.6811	-0.1651	0.5895	-0.0184	0.9884	0.1460	0.6679	0.1644	0.5821	0.3428	0.8270	0.1389	0.8812	-0.2040	0.7934
CG31200	CG31200	1623969_at	0.6268	0.5050	0.1974	0.3236	-0.0633	0.7417	-0.2647	0.7949	-0.6532	0.1145	-0.3886	0.3022	-0.1187	0.9816	-0.9247	0.4103	-0.8060	0.4924
CG13154	CG13154	1623970_at	-0.1164	0.6186	0.0143	0.8886	0.0185	0.9389	-0.0027	0.9978	-0.2208	0.3882	-0.2181	0.3403	-0.0218	0.9826	0.0693	0.8168	0.0910	0.7291
CG9150	CG9150	1623971_at	0.2161	0.3854	0.0823	0.6289	0.1481	0.3944	-0.0332	0.9712	-0.1192	0.6496	-0.0861	0.7305	0.0343	0.9689	-0.0073	0.9877	-0.0416	0.8950
CG4538 /// DereCG4538 ///	CG4538	1623972_a_at	0.4363	0.0798	0.3187	0.1796	0.2949	0.1899	-0.3169	0.3855	-0.3013	0.1434	0.0157	0.9523	-0.2458	0.8472	-0.4258	0.4203	-0.1800	0.7741
---	---	1623973_at	-0.0178	0.9214	0.0813	0.7077	0.1333	0.4528	0.0105	0.9931	0.0432	0.8940	0.0327	0.9099	0.0064	0.9939	0.0305	0.9170	0.0241	0.9266
CG15549	CG15549	1623974_at	0.1401	0.5948	-0.1666	0.4475	-0.1523	0.4982	0.2192	0.5156	0.1011	0.6047	-0.1180	0.4883	0.0269	0.9822	-0.1736	0.5529	-0.2004	0.4882
az2	Zinc finger protein	1623975_at	-0.2734	0.3101	-0.1659	0.1423	-0.2928	0.2062	0.1236	0.8501	0.2315	0.3393	0.1079	0.6621	0.2749	0.7726	0.3586	0.3737	0.0838	0.8811
CG30081	CG30081	1623976_at	0.0661	0.7591	0.1756	0.3659	0.1381	0.5094	-0.1512	0.7018	-0.1704	0.3561	-0.0192	0.9312	0.0040	0.9970	-0.0004	0.9998	-0.0044	0.9924
nvy	nervy	1623977_at	-1.1690	0.0272	-2.0503	0.0092	-2.3380	0.0000	0.4106	0.3996	0.8986	0.0062	0.4880	0.0504	0.0002	0.9999	-0.2156	0.6216	-0.2158	0.6222
CG10862	CG10862	1623978_at	0.0851	0.5877	-0.0454	0.6560	0.0764	0.6889	0.1062	0.8194	0.1042	0.6015	-0.0020	0.9928	-0.1281	0.8202	-0.1525	0.5266	-0.0244	0.9402
CG15482	CG15482	1623979_at	0.1210	0.5917	0.2405	0.2084	0.1304	0.4431	-0.2744	0.3863	-0.1996	0.2680	0.0748	0.6914	0.1028	0.8903	0.1201	0.7050	0.0173	0.9673
CG15905	CG15905	1623980_at	0.2487	0.0910	0.1622	0.4406	0.2203	0.3686	0.0799	0.9328	0.0373	0.9241	-0.0426	0.8995	-0.0291	0.9659	-0.0217	0.9411	0.0074	0.9809
CG7017	CG7017	1623981_at	-0.0686	0.8027	0.0198	0.8546	0.1105	0.5994	0.0773	0.9046	0.0190	0.9517	-0.0583	0.8046	-0.0691	0.9238	0.1417	0.5794	0.2108	0.3921
---	---	1623982_s_at	-0.0587	0.7409	0.0816	0.6506	-0.2219	0.1610	0.0145	0.9808	0.0590	0.7313	0.0445	0.7862	0.1896	0.6749	0.0732	0.7287	-0.1163	0.5460
CG9514	CG9514	1623983_at	-0.2755	0.1983	-0.2094	0.1849	-0.2099	0.3152	0.0401	0.9524	0.0872	0.6814	0.0471	0.8241	-0.0837	0.9296	0.0738	0.8631	0.1575	0.6328
MICAL	MICAL	1623984_s_at	-0.4065	0.2708	0.8938	0.2111	0.6704	0.0200	-0.0816	0.9471	-0.9857	0.0120	-0.9041	0.0108	0.1825	0.9333	0.3731	0.6186	0.1906	0.8287
l(3)j2D3	lethal (3) j2D3	1623985_at	-0.4891	0.0302	-0.0197	0.9515	-0.1975	0.1815	-0.0625	0.9314	-0.0874	0.7306	-0.0249	0.9252	-0.0863	0.9409	0.2382	0.5246	0.3245	0.3807
CG15046	CG15046	1623986_at	-1.3949	0.0038	-1.1328	0.0588	-0.6643	0.0286	0.2380	0.6692	-0.2955	0.2666	-0.5335	0.0316	-0.1727	0.9095	0.0336	0.9702	0.2063	0.7416

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG12679	CG12679	1623987_at	-0.2058	0.3138	-0.0524	0.8708	-0.1008	0.7039	0.0940	0.9185	-0.2042	0.4901	-0.2982	0.2386	-0.0080	0.9963	-0.1408	0.7439	-0.1328	0.7527
CG14691	CG14691	1623988_at	-0.1451	0.4689	0.2153	0.1363	0.3220	0.0876	0.0389	0.9626	0.0071	0.9826	-0.0319	0.9042	-0.1125	0.8446	0.1901	0.4172	0.3026	0.2149
CG5734	CG5734	1623989_at	-0.4613	0.1432	-0.1023	0.8489	-0.2318	0.2531	-0.1602	0.7432	-0.4226	0.0543	-0.2625	0.1716	0.0851	0.9717	0.0541	0.9561	-0.0311	0.9738
CG33220	CG33220	1623990_at	0.1591	0.5085	0.0898	0.4844	0.1234	0.4736	0.1279	0.8022	0.1885	0.3673	0.0606	0.7913	-0.0200	0.9871	-0.0303	0.9477	-0.0102	0.9835
coro	coronin	1623991_s_at	-0.9983	0.0023	-0.6440	0.0485	-1.3913	0.0001	-0.2465	0.4590	0.0210	0.9341	0.2674	0.0968	0.3761	0.6272	0.3215	0.3064	-0.0546	0.9026
CG7229	Gene 4	1623992_at	-0.1022	0.4971	-0.2950	0.0370	-0.0235	0.9118	-0.0153	0.9830	-0.0758	0.6874	-0.0606	0.7312	0.0163	0.9909	-0.1032	0.7737	-0.1194	0.7197
CG4291	CG4291	1623993_at	-0.0502	0.7695	0.0905	0.6403	0.1614	0.3334	-0.0107	0.9865	-0.1426	0.3605	-0.1319	0.3468	-0.0597	0.9092	-0.0567	0.8132	0.0030	0.9924
CG30416	CG30416	1623994_at	-0.1214	0.4725	-0.0971	0.3446	-0.0484	0.8302	0.2335	0.4861	0.1484	0.4233	-0.0851	0.6442	0.0512	0.9340	0.0196	0.9506	-0.0316	0.9080
RpL35A	Ribosomal protein	1623995_at	0.4355	0.0352	1.0949	0.0091	1.5123	0.0008	0.2620	0.6110	-0.5238	0.0478	-0.7858	0.0053	0.0163	0.9831	-0.0047	0.9910	-0.0210	0.9340
CG5001	CG5001	1623996_at	-1.5277	0.0010	-1.1241	0.0149	-1.0930	0.0006	0.2172	0.6755	-0.0474	0.8852	-0.2646	0.2258	0.2344	0.7707	0.4962	0.1418	0.2618	0.4620
RpS30	Ribosomal protein	1623997_s_at	0.4360	0.0228	0.8248	0.0223	0.6208	0.0225	0.0832	0.8908	-0.2095	0.3093	-0.2927	0.1078	0.1611	0.8076	0.0195	0.9655	-0.1416	0.6329
CG10565	CG10565	1623998_at	-0.1178	0.6775	0.5808	0.0375	1.1406	0.0009	0.3801	0.3744	-0.4360	0.0743	-0.8161	0.0033	-0.1739	0.8386	0.2766	0.4302	0.4505	0.2162
Dys	Dystrophin-like pr	1623999_at	-0.1616	0.5539	0.0999	0.6245	0.0636	0.8127	0.0623	0.9562	0.0123	0.9794	-0.0500	0.8915	0.0154	0.9884	0.0450	0.8963	0.0296	0.9259
CG13443	CG13443	1624000_at	0.1522	0.4898	0.0279	0.9099	0.2179	0.2985	0.1735	0.6457	0.0809	0.7014	-0.0926	0.6178	-0.1827	0.8283	-0.1219	0.7787	0.0608	0.9009
sbb	scribble	1624001_at	0.1386	0.6113	0.0656	0.7502	-0.0028	0.9956	0.0645	0.9598	0.1859	0.6163	0.1214	0.7369	0.0822	0.9467	-0.0076	0.9931	-0.0897	0.8524
CG31291	CG31291	1624002_a_at	0.1713	0.4857	0.1884	0.5006	-0.1212	0.5483	-0.0001	0.9999	0.0012	0.9980	0.0013	0.9978	0.2767	0.6749	-0.0665	0.8593	-0.3432	0.2210
CG10639	CG10639	1624003_at	0.1872	0.3631	0.5152	0.0755	0.7525	0.0155	-0.1280	0.8578	-0.8517	0.0060	-0.7237	0.0075	-0.3885	0.6706	-0.5442	0.1382	-0.1557	0.7072
rols	Antisocial	1624004_s_at	0.7592	0.2735	0.6043	0.3697	0.6890	0.0112	0.5691	0.2789	0.8256	0.0159	0.2565	0.3707	0.5042	0.8427	0.7098	0.5100	0.2056	0.8862
---	---	1624005_at	-0.0057	0.9748	-0.0119	0.9330	0.2542	0.1042	0.2313	0.5061	-0.0444	0.8528	-0.2757	0.0941	0.0271	0.9816	-0.0275	0.9492	-0.0547	0.8786
MED25	Mediator complex	1624006_at	0.1628	0.6896	-0.5096	0.0925	-0.3556	0.1224	0.1542	0.7239	1.1344	0.0006	0.9802	0.0006	0.0268	0.9916	0.4796	0.4262	0.4528	0.4683
CG6304	CG6304	1624007_at	0.0129	0.9649	-0.0248	0.8217	0.1389	0.4799	0.3264	0.1749	0.2441	0.0946	-0.0824	0.5690	0.2525	0.7215	0.1090	0.7606	-0.1435	0.6577
eIF2B-epsilon	eIF2B-epsilon	1624008_at	0.1817	0.3862	0.2571	0.1239	0.3308	0.0469	-0.0016	0.9978	0.0739	0.6682	0.0756	0.6246	0.0911	0.9092	0.2962	0.2787	0.2051	0.4835
CG34360	CG12802	1624009_at	0.4277	0.2421	0.3954	0.4102	-0.4191	0.0841	-0.6455	0.1634	0.2797	0.3262	0.9252	0.0037	-0.0246	0.9914	0.2079	0.7341	0.2325	0.6886
CG9062	CG9062	1624010_a_at	-0.2558	0.5763	0.7820	0.0637	0.8134	0.0040	0.1143	0.9154	-0.2025	0.5793	-0.3168	0.2992	0.0453	0.9841	0.7179	0.1498	0.6726	0.2094
Cpr76Bc	CG9295	1624011_at	0.1202	0.6617	0.1661	0.4070	-0.1272	0.4245	-0.1770	0.7805	-0.1737	0.5389	0.0032	0.9920	0.1352	0.8270	-0.1293	0.6414	-0.2646	0.3109
mRpL10	mitochondrial ribo	1624012_at	0.1251	0.7150	0.2983	0.1302	0.0284	0.8745	0.0141	0.9838	-0.2600	0.0980	-0.2741	0.0551	0.1786	0.8629	0.0103	0.9916	-0.1683	0.7222
CG11574	CG11574	1624013_at	-0.1798	0.3669	0.1776	0.3780	-0.0736	0.7561	-0.2863	0.5357	-0.2134	0.3875	0.0728	0.7869	0.0970	0.9095	0.1751	0.5880	0.0781	0.8411
Use1	Use1	1624014_at	0.1076	0.5496	-0.0283	0.9206	-0.1203	0.4272	0.0541	0.9314	0.2399	0.1959	0.1858	0.2687	-0.0068	0.9964	0.3313	0.3759	0.3382	0.3858
CG13732	CG13732	1624015_at	0.1108	0.4229	-0.0225	0.9188	0.1296	0.5374	0.0759	0.8817	0.2939	0.0925	0.2180	0.1608	0.1051	0.8940	0.0594	0.8910	-0.0456	0.9109
CG14238	CG14238	1624016_at	-0.1202	0.5969	0.0644	0.6435	0.1645	0.3328	0.0567	0.9228	-0.3060	0.0849	-0.3627	0.0295	-0.0857	0.9309	-0.0938	0.8218	-0.0081	0.9874
Ahcy89E	AdoHcyase-like	1624017_at	1.1151	0.6507	-3.4083	0.0026	-2.5467	0.0001	0.6651	0.3553	1.1639	0.0121	0.4988	0.1744	-0.3051	0.9816	-3.1802	0.2588	-2.8751	0.3375
---	---	1624018_at	0.0000	1.0000	-0.0115	0.9591	0.0344	0.8877	-0.0157	0.9883	-0.0747	0.8112	-0.0589	0.8399	-0.1623	0.7770	-0.0893	0.7671	0.0730	0.8129
RpL36	Viability	1624019_s_at	0.0557	0.7310	0.6005	0.0367	0.6372	0.0062	0.1119	0.7647	-0.3215	0.0494	-0.4335	0.0089	0.0119	0.9914	0.0280	0.9429	0.0161	0.9640
mod(mdg4)	Modifier67.2	1624020_at	-0.7456	0.0292	-0.5473	0.0875	-0.4891	0.0094	-0.0881	0.9311	-0.2252	0.4728	-0.1370	0.6590	-0.2108	0.7826	0.0174	0.9761	0.2282	0.5059
dlg1	Discs large	1624021_a_at	-0.3608	0.3312	0.4908	0.1067	-0.9078	0.0732	-0.6855	0.0840	-0.2225	0.3427	0.4630	0.0321	0.7452	0.7220	0.6695	0.4513	-0.0756	0.9517
Atac1	CG9200	1624022_at	0.2828	0.1207	-0.0243	0.8659	-0.0128	0.9539	0.0139	0.9865	0.3851	0.0591	0.3711	0.0444	-0.0003	0.9999	0.0969	0.7903	0.0972	0.7770
Pp1-87B	Protein phosphata	1624023_at	-0.1830	0.2109	0.2304	0.3895	-0.0449	0.7713	0.0089	0.9894	-0.0801	0.6445	-0.0891	0.5597	0.1581	0.8076	0.2058	0.4475	0.0476	0.8982
---	---	1624024_s_at	0.1687	0.3791	0.2142	0.2942	0.1616	0.3348	-0.0453	0.9426	-0.0130	0.9616	0.0323	0.8791	0.0577	0.9514	-0.0332	0.9411	-0.0910	0.7848
CG14674	CG14674	1624025_at	-0.0089	0.9812	0.1135	0.4728	0.4299	0.0519	0.1341	0.7670	-0.1762	0.3701	-0.3104	0.0735	-0.0829	0.9246	-0.0119	0.9829	0.0710	0.8509
---	---	1624026_at	0.0764	0.7169	0.0428	0.8156	0.1894	0.3320	0.0126	0.9882	-0.0358	0.8943	-0.0484	0.8298	-0.0484	0.9623	-0.0396	0.9305	0.0088	0.9849
CG4780	CG4780	1624027_at	1.0819	0.0043	1.3642	0.0297	1.3327	0.0002	0.1649	0.5735	0.5597	0.0036	0.3948	0.0103	0.1179	0.9491	0.8507	0.1251	0.7328	0.2056
CG4415	CG4415	1624028_at	-0.1398	0.4276	0.1056	0.5334	0.0715	0.8084	0.0375	0.9649	-0.1543	0.5251	-0.1917	0.3607	0.0817	0.9259	0.2116	0.4779	0.1299	0.6877
CG3608	CG3608	1624029_at	0.7450	0.0097	1.4114	0.0083	1.7158	0.0003	0.1819	0.6890	0.1825	0.3993	0.0006	0.9982	-0.1180	0.9246	0.7555	0.0810	0.8735	0.0729
rnh1	ribonuclease H1	1624030_at	-0.2465	0.4538	0.0049	0.9669	0.4007	0.0616	0.1824	0.6120	0.0858	0.6758	-0.0966	0.5917	-0.3017	0.7464	0.3719	0.3285	0.6736	0.1190
CG40139	CG40139	1624031_at	-0.2646	0.0899	-0.6662	0.0148	-0.5086	0.0583	0.1223	0.8449	0.6221	0.0150	0.4998	0.0236	-0.1287	0.8062	0.0320	0.9266	0.1606	0.4777
CG3887	CG3887	1624032_at	0.0440	0.8977	-0.3068	0.2114	-0.4836	0.0160	0.0661	0.8792	0.3372	0.0306	0.2711	0.0467	0.2175	0.8380	-0.0100	0.9925	-0.2275	0.6311
Appl	beta-amyloid-prot	1624033_at	-1.5539	0.0031	-0.3661	0.0345	-0.9309	0.0017	-0.4685	0.4475	-1.0606	0.0075	-0.5921	0.0531	0.0562	0.9516	-0.1594	0.5912	-0.2156	0.4568
Gr28a	Gustatory recepto	1624034_at	0.1837	0.3473	-0.0788	0.6554	-0.2231	0.3192	-0.0409	0.9603	-0.0203	0.9509	0.0206	0.9397	-0.1029	0.8837	-0.2211	0.4037	-0.1181	0.6892
LCBP1	LDLa domain con	1624035_a_at	0.2458	0.3613	0.4148	0.1648	0.5481	0.0096	0.0080	0.9943	-0.3540	0.1332	-0.3621	0.0872	-0.0133	0.9933	-0.1009	0.8353	-0.0876	0.8519
CG9307	CG9307	1624036_at	0.0360	0.8961	0.0947	0.6399	0.1498	0.5031	-0.0019	0.9978	0.0440	0.8495	0.0459	0.8214	-0.0415	0.9679	-0.0191	0.9661	0.0224	0.9539
Taf4	TBP-associated fa	1624037_s_at	0.2717	0.4471	-0.0177	0.9273	-0.4799	0.0133	-0.2167	0.5621	0.4341	0.0310	0.6508	0.0032	0.1942	0.8786	0.2012	0.7200	0.0070	0.9935
CG17024	CG17024	1624038_at	0.0083	0.9758	-0.2496	0.3736	-0.0218	0.9134	0.1745	0.7711	0.1524	0.5788	-0.0221	0.9448	0.0491	0.9521	-0.0906	0.7572	-0.1396	0.5969
CG10147	CG10147	1624039_at	0.0517	0.8908	0.4799	0.1700	0.4262	0.0804	-0.0644	0.9601	-0.3115	0.3667	-0.2470	0.4365	-0.0893	0.9390	0.0299	0.9597	0.1192	0.7780
CG12732	CG12732	1624040_at	0.1301	0.6896	-0.0180	0.8887	0.0761	0.7438	-0.1022	0.8234	-0.2574	0.1463	-0.1552	0.3411	-0.0812	0.9657	-0.2573	0.6416	-0.1761	0.7674
cul-4	cul-4	1624041_at	0.4965	0.0440	0.3658	0.3233	0.3797	0.0231	-0.0982	0.7857	0.2764	0.0672	0.3747	0.0127	-0.0328	0.9829	0.1673	0.6622	0.2002	0.5914
Gprk2	G protein-coupled	1624042_at	-0.2006	0.6604	1.0078	0.0944	-0.3379	0.2718	0.1317	0.8251	0.4848	0.0422	0.3532	0.0876	1.4580	0.3166	1.6395	0.0564	0.1815	0.8577
---	---	1624043_at	0.5448	0.0838	0.0638	0.6896	0.1436	0.5003	0.2549	0.3433	1.0298	0.0004	0.7749	0.0006	0.2626	0.7899	0.6068	0.1418	0.3442	0.4235
mRpL52	mitochondrial ribo	1624044_at	0.0027	0.9906	0.4252	0.1085	0.2732	0.2145	0.0597	0.9410	0.1547	0.5293	0.0951	0.6949	0.2010	0.8270	0.4802	0.1957	0.2792	0.4824
---	---	1624045_at	0.0519	0.8272	0.1716	0.5246	0.4209	0.0708	0.0502	0.9375	-0.1548	0.4228	-0.2051	0.2199	-0.0986	0.9142	-0.0414	0.9358	0.0573	0.8972
---	---	1624046_at	0.1657	0.2848	-0.0057	0.9615	-0.2852	0.1883	-0.1630	0.7929	0.1209	0.6757	0.2839	0.2105	0.0454	0.9309	-0.0232	0.9325	-0.0686	0.7235
CG31078	CG31078	1624047_at	0.0075	0.9799	-0.2043	0.1088	-0.0259	0.8790	0.1034	0.8578	0.1588	0.4611	0.0554	0.8123	-0.0425	0.9589	0.0193	0.9599	0.0618	0.8325
CG15185	CG15185	1624048_at	0.3276	0.1219	0.1102	0.3996	0.0278	0.8905	-0.2320	0.4690	-0.0408	0.8560	0.1912	0.2162	-0.0776	0.8953	-0.0926	0.7105	-0.0149	0.9633
CG13802 /// DmsR-2	CG13802 /// Drom	1624049_at	-0.1754	0.3883	0.1191	0.5432	0.0703	0.6672	-0.2212	0.5311	-0.4824	0.0171	-0.2612	0.1122	-0.0227	0.9821	-0.0669	0.8307	-0.0442	0.8918
CG6842	CG6842	1624050_at	-0.1481	0.3686	-0.4384	0.0384	-0.5396	0.0143	0.0463	0.9322	0.2911	0.0747	0.2448	0.0908	0.0759	0.8999	-0.1208	0.6111	-0.1967	0.3862
CG13850	CG13850	1624051_at	-0.2502	0.4276	0.2639	0.2552	0.1160	0.5705	-0.1010	0.8159	-0.3959	0.0279	-0.2949	0.0570	-0.0799	0.9607	0.0108	0.9923	0.0907	0.8844
CG6294	CG6294	1624052_at	-0.1106	0.7764	0.9141	0.2489	0.2868	0.3010	0.0775	0.9380	-0.4549	0.1186	-0.5324	0.0474	0.5498	0.7810	0.4154	0.6586	-0.1344	0.9095
CG33967	CG33967	1624053_at	-0.2528	0.5809	0.8666	0.0262	0.7297	0.0023	-0.0780	0.9375	-0.7882	0.0142	-0.7102	0.0138	0.0672	0.9677	0.2845	0.5289	0.2173	0.6428
Adgf-A	Adenosine deamir	1624054_at	-1.3510	0.0009	0.6154	0.3167	-1.5436	0.0192	-1.3711	0.0770	-2.1701	0.0012	-0.7990	0.0511	0.9049	0.6483	0.0180	0.9925	-0.8869	0.2834
---	---	1624055_at	-0.1598	0.4768	0.1624	0.5505	0.0773	0.7936	0.0349	0.9745	-0.2747	0.3111	-0.3096	0.1963	0.2949	0.6557	0.0482	0.9064	-0.2467	0.3695

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG12455	CG12455	1624056_a_at	-1.0721	0.0021	-0.7640	0.1047	-1.2103	0.0005	-0.7422	0.1119	-0.5918	0.0402	0.1504	0.5898	-0.0202	0.9893	-0.0712	0.8727	-0.0510	0.9057
CG16713	CG16713	1624057_at	2.5280	0.0040	0.3032	0.7291	1.6248	0.0008	0.8555	0.3030	0.7830	0.1076	-0.0725	0.9015	-0.6421	0.7945	-1.5534	0.1339	-0.9113	0.3880
fs(1)N	Nasrat	1624058_at	0.2965	0.6800	-1.1210	0.1382	-0.3857	0.3611	0.4783	0.2973	1.5521	0.0006	1.0738	0.0016	-0.6395	0.8395	0.0438	0.9868	0.6833	0.6254
CG32223	CG32223	1624059_at	0.6933	0.2102	0.5644	0.2543	-0.5839	0.0483	-0.5183	0.5799	0.4208	0.3910	0.9391	0.0352	0.7365	0.6706	0.5579	0.4245	-0.1786	0.8418
bab2	bric-a-brac	1624060_at	-0.6182	0.1724	-2.3453	0.0415	-2.7019	0.0006	0.1490	0.9223	1.3608	0.0100	1.2118	0.0102	0.4981	0.8332	-0.3320	0.7883	-0.8302	0.4093
Vm32E	Vitelline membran	1624061_at	0.1906	0.6946	-0.1291	0.3133	-0.1031	0.5087	-0.0766	0.9116	-0.0333	0.9128	0.0433	0.8647	-0.1830	0.8478	-0.3483	0.3712	-0.1653	0.7127
CG14962	CG14962	1624062_at	0.8046	0.0032	0.0230	0.9585	-0.3052	0.2460	-0.0158	0.9860	0.6345	0.0100	0.6503	0.0054	0.3277	0.7893	-0.0575	0.9460	-0.3852	0.4755
CG15699	CG15699	1624063_at	0.2636	0.3507	0.3930	0.1118	0.5767	0.0447	0.2036	0.7415	-0.1609	0.5863	-0.3646	0.1320	0.0869	0.9278	0.1662	0.6218	0.0793	0.8434
CG9915 /// DyakCG9915	CG9915	1624064_at	-0.0042	0.9827	-0.4780	0.0386	-0.5035	0.0236	-0.0847	0.8534	0.2423	0.1422	0.3270	0.0344	-0.1303	0.8540	-0.2420	0.4002	-0.1117	0.7391
sqh	myosin II	1624065_at	-0.1321	0.5982	0.3796	0.1472	0.3480	0.1354	0.1982	0.6382	0.0348	0.9026	-0.1633	0.3950	0.2401	0.7893	0.6821	0.0860	0.4420	0.2622
---	---	1624066_at	0.0702	0.7013	0.0956	0.4097	0.2736	0.1317	0.2139	0.5376	0.1640	0.3761	-0.0499	0.8063	0.0151	0.9862	0.0892	0.6930	0.0741	0.7492
CG6704	CG6704	1624067_at	0.4735	0.4662	0.4700	0.5196	0.4205	0.1661	-0.0399	0.9777	-0.1981	0.6246	-0.1583	0.6776	-0.0624	0.9913	-0.0878	0.9634	-0.0254	0.9892
---	---	1624068_at	0.1410	0.3594	0.0551	0.6316	0.1989	0.2475	0.0252	0.9734	0.0683	0.7607	0.0430	0.8419	0.0100	0.9914	0.1168	0.5825	0.1067	0.6214
CG7296	CG7296	1624069_at	0.4634	0.4914	-0.0993	0.5773	-0.4467	0.0597	-0.4059	0.2697	-0.0338	0.9108	0.3721	0.0592	-0.0991	0.9764	-0.4918	0.5618	-0.3927	0.6528
RpS9	Ribosomal protein	1624070_at	1.5921	0.0014	0.9656	0.0097	1.4497	0.0003	0.3186	0.3840	0.4644	0.0322	0.1458	0.4569	-0.3258	0.6955	-0.3663	0.2825	-0.0405	0.9344
CG14550	CG14550	1624071_at	0.1236	0.6967	0.1117	0.6406	-0.0467	0.8108	0.0177	0.9777	0.1665	0.3071	0.1488	0.3107	0.0544	0.9762	0.2171	0.6497	0.1627	0.7491
---	---	1624072_at	-0.0238	0.9325	0.1659	0.4567	0.1971	0.2300	0.1271	0.8202	0.0386	0.8973	-0.0884	0.7017	0.0051	0.9966	0.1563	0.6339	0.1513	0.6428
CG15800	CG15800	1624073_at	-0.0078	0.9696	0.1774	0.2696	-0.0401	0.8924	-0.2119	0.5280	-0.2372	0.1743	-0.0253	0.9047	0.1263	0.8973	0.0395	0.9470	-0.0868	0.8549
Obp56a	Odorant-binding p	1624074_at	0.2908	0.6685	0.0304	0.7623	-0.4374	0.0443	-0.4251	0.5249	-0.4335	0.2145	-0.0084	0.9850	0.0997	0.9748	-0.7279	0.3409	-0.8275	0.3027
CG31495	CG31495	1624075_at	0.0489	0.8295	0.1265	0.5873	0.1273	0.4706	-0.0855	0.8796	0.0203	0.9427	0.1058	0.5836	-0.1459	0.8795	-0.0315	0.9590	0.1144	0.7980
CG5849	CG5849	1624076_at	1.8488	0.0031	1.0596	0.0200	1.9243	0.0006	0.2177	0.8531	0.2071	0.6685	-0.0105	0.9849	-0.5940	0.6724	-0.5385	0.3285	0.0556	0.9435
CG13390	CG13390	1624077_at	-0.5990	0.0368	-0.0862	0.5363	0.7733	0.0047	0.3084	0.4586	-0.6407	0.0118	-0.9490	0.0013	-0.5667	0.5089	-0.2033	0.6443	0.3633	0.3813
CG5463	CG5463	1624078_at	-0.0517	0.8693	-0.2369	0.6913	0.1012	0.5494	-0.2931	0.5570	-0.0768	0.8094	0.2163	0.3611	-0.6750	0.3987	-0.2698	0.5539	0.4052	0.3668
Rab21	Rab21	1624079_a_at	-0.2909	0.1321	-0.3281	0.0873	-0.1830	0.4765	0.0176	0.9863	0.0101	0.9790	-0.0075	0.9802	-0.1686	0.8454	-0.0692	0.8963	0.0994	0.8237
Trn	transportin	1624080_s_at	0.5081	0.3611	-0.0876	0.8862	-0.5251	0.1620	-0.1010	0.9506	1.5590	0.0049	1.6601	0.0022	0.2490	0.9309	0.8640	0.3421	0.6149	0.5319
CG14359	CG14359	1624081_at	0.1507	0.4403	0.1173	0.4184	-0.0250	0.8872	-0.0979	0.8059	-0.1860	0.2419	-0.0880	0.5776	0.0853	0.8885	0.0188	0.9605	-0.0664	0.8166
Or33c	Olfactory receptor	1624082_at	0.0185	0.9261	0.0170	0.8687	0.3284	0.0751	0.1322	0.7121	0.0496	0.8044	-0.0826	0.6176	-0.2396	0.7220	-0.1001	0.7760	0.1396	0.6519
M(2)21AB	Methionine adeno	1624083_s_at	0.5368	0.0422	0.4284	0.0911	0.2328	0.2484	-0.2119	0.6120	0.0792	0.7499	0.2911	0.1186	-0.0116	0.9950	0.0000	1.0000	0.0117	0.9861
---	---	1624084_s_at	0.0527	0.7187	-0.0276	0.7863	0.0536	0.7543	0.0910	0.7981	0.0726	0.6556	-0.0185	0.9164	-0.0914	0.8439	-0.0405	0.8830	0.0510	0.8321
CG9876	CG9876	1624085_at	0.1534	0.5809	0.2729	0.3312	0.1373	0.5565	0.1093	0.8526	0.0497	0.8585	-0.0595	0.8040	0.1406	0.8122	0.0323	0.9341	-0.1083	0.6883
Vha36	Vacuolar H-ATPa	1624086_at	-1.0499	0.0276	-0.2598	0.1639	-0.4639	0.0235	-0.2244	0.5376	-1.1747	0.0004	-0.9503	0.0006	-0.0211	0.9923	-0.3206	0.5469	-0.2995	0.5811
---	---	1624087_at	-0.0496	0.8596	0.1300	0.5983	0.3139	0.0814	0.1413	0.8447	-0.1284	0.6715	-0.2697	0.2597	0.0309	0.9836	0.1776	0.6380	0.1466	0.7071
CG3108	CG3108	1624088_at	0.2376	0.2627	0.0946	0.4759	0.1733	0.4684	0.0844	0.9160	-0.0544	0.8696	-0.1388	0.5769	0.0896	0.8740	0.0814	0.7591	-0.0081	0.9835
---	---	1624089_at	-0.0791	0.6780	0.1893	0.3636	0.1119	0.5198	-0.0630	0.9314	-0.2253	0.3069	-0.1623	0.4284	0.0225	0.9841	0.1785	0.4954	0.1560	0.5637
CG12726	CG12726	1624090_at	0.8309	0.4685	0.1106	0.5039	0.3546	0.0776	-0.0375	0.9504	-0.0508	0.8044	-0.0132	0.9502	-0.0950	0.9862	-0.6563	0.6156	-0.5612	0.6744
---	---	1624091_at	-0.0908	0.6491	0.0580	0.6004	0.2258	0.2098	0.0197	0.9777	-0.2208	0.2023	-0.2405	0.1205	-0.0216	0.9869	-0.0103	0.9865	0.0113	0.9829
Jon99Ci	Jonah 99C Alpha	1624092_at	0.2726	0.3655	-0.0283	0.7679	-0.0581	0.8121	0.1010	0.8337	0.1887	0.3062	0.0877	0.6397	0.0839	0.9421	-0.0411	0.9433	-0.1250	0.7640
---	---	1624093_at	0.1877	0.2766	-0.1661	0.6811	-0.2545	0.2210	-0.2861	0.5944	0.5013	0.0688	0.7873	0.0067	-0.0095	0.9964	0.0458	0.9505	0.0554	0.9318
---	---	1624094_at	-0.0797	0.6553	0.1480	0.1961	0.0902	0.7260	-0.0733	0.9353	-0.1444	0.6270	-0.0711	0.8135	0.0934	0.8454	0.0798	0.7248	-0.0137	0.9622
---	---	1624095_at	0.2075	0.2237	0.3026	0.1026	0.1278	0.4814	0.0946	0.8005	0.0013	0.9951	-0.0933	0.5250	0.1391	0.7726	0.0113	0.9747	-0.1278	0.5612
---	---	1624096_at	0.0619	0.7497	0.0775	0.6174	0.0974	0.5783	0.0852	0.8507	0.1138	0.5138	0.0286	0.8834	0.0360	0.9816	0.1712	0.6512	0.1352	0.7353
wge	winged eye	1624097_s_at	-0.1585	0.8141	-0.8765	0.2730	-1.0332	0.0087	0.0297	0.9808	1.0216	0.0054	0.9918	0.0037	0.1719	0.9659	0.3631	0.7891	0.1912	0.8982
CG18675 /// tipE	CG18675 /// temp	1624098_s_at	-0.0316	0.8594	-0.1111	0.5396	-0.1774	0.3719	0.0153	0.9834	0.1756	0.3026	0.1602	0.2945	-0.0448	0.9495	0.1164	0.6053	0.1612	0.4597
CG1701	CG1701	1624099_at	0.1470	0.3436	-0.0300	0.8400	0.0167	0.9384	0.0025	0.9962	0.1724	0.3268	0.1700	0.2779	-0.2190	0.7644	-0.1434	0.6765	0.0755	0.8501
CG1529	CG1529	1624100_at	0.0187	0.9513	-0.2575	0.3248	-0.0540	0.8701	0.0122	0.9941	0.3150	0.3616	0.3027	0.3271	-0.0994	0.9342	0.0955	0.8584	0.1949	0.6392
Cyp6a23	Cyp6a23	1624101_at	1.1008	0.0252	-0.1997	0.4086	0.7428	0.0132	-0.0543	0.9507	-0.5590	0.0293	-0.5047	0.0282	-1.0207	0.3587	-1.8488	0.0222	-0.8282	0.1897
---	---	1624102_at	0.1723	0.4386	-0.0421	0.8399	-0.0963	0.5857	0.0035	0.9965	0.2415	0.3661	0.2380	0.3181	-0.0384	0.9657	-0.0384	0.9178	0.0000	1.0000
CG7246	CG7246	1624103_at	0.4394	0.2013	-0.0238	0.9480	0.6798	0.0216	0.4936	0.1680	0.7592	0.0048	0.2656	0.1714	-0.3540	0.7251	0.1246	0.8290	0.4786	0.2865
---	---	1624104_at	0.1043	0.5903	0.1271	0.3045	0.1597	0.4980	-0.0353	0.9598	-0.0956	0.6420	-0.0603	0.7641	0.1408	0.8508	-0.0719	0.8690	-0.2127	0.5083
CG8248	CG8248	1624105_at	-0.1247	0.3834	0.0124	0.9700	0.3930	0.1052	0.0375	0.9583	0.0360	0.8880	-0.0015	0.9950	-0.2395	0.8202	0.0551	0.9368	0.2946	0.5153
CG3790	CG3790	1624106_at	0.0864	0.6359	-0.0792	0.7322	0.2032	0.3484	0.2066	0.7130	0.2821	0.2717	0.0755	0.7916	-0.0251	0.9734	-0.0548	0.8216	-0.0297	0.9080
---	---	1624107_at	0.2798	0.2802	0.0609	0.5641	0.3524	0.0299	0.1660	0.7293	0.1881	0.3933	0.0222	0.9340	0.1739	0.7726	0.1844	0.4903	0.0105	0.9809
CG34114	CG34114	1624108_at	0.1446	0.2958	0.0649	0.5926	0.3539	0.0714	-0.0149	0.9802	-0.2022	0.1575	-0.1872	0.1436	-0.1190	0.8141	-0.0079	0.9842	0.1111	0.6216
CG31522	CG31522	1624109_s_at	-1.1651	0.0768	-1.4329	0.0374	-2.2568	0.0002	-0.3559	0.3619	-0.3253	0.1419	0.0306	0.9095	0.4108	0.8510	-0.6259	0.5018	-1.0367	0.2677
cv-c	crossveinless-c	1624110_at	0.3119	0.2178	0.1753	0.3372	0.3966	0.0282	0.0425	0.9521	-0.1313	0.5319	-0.1738	0.3340	-0.1105	0.9088	-0.0240	0.9655	0.0864	0.8430
mun	munin	1624111_at	0.1279	0.4199	0.1545	0.3955	0.1947	0.1838	0.0995	0.8902	0.0639	0.8365	-0.0356	0.9027	0.0404	0.9710	-0.1236	0.6949	-0.1640	0.5819
---	---	1624112_at	0.2588	0.1896	0.2073	0.2734	0.2332	0.2419	0.0830	0.8837	-0.0685	0.7678	-0.1516	0.4002	-0.0575	0.9404	-0.1130	0.6634	-0.0555	0.8572
CG7845	CG7845	1624113_at	0.5857	0.0885	0.2485	0.6223	0.3143	0.1221	0.2726	0.6796	0.7491	0.0233	0.4765	0.0847	0.1153	0.9449	0.3267	0.5324	0.2114	0.7090
Klp59C	Klp59C	1624114_at	0.1782	0.3122	-0.1189	0.5305	-0.0511	0.8526	-0.0534	0.9538	0.0516	0.8781	0.1050	0.6918	0.1157	0.8882	-0.0320	0.9496	-0.1477	0.6604
---	---	1624115_s_at	-0.0432	0.7884	0.3994	0.0488	0.3580	0.0515	-0.0684	0.9017	-0.0907	0.6535	-0.0223	0.9193	0.0983	0.8609	0.3903	0.0953	0.2921	0.2149
Mst36Fa	CG31801	1624116_at	0.1281	0.5947	-0.0202	0.8724	-0.4568	0.3929	-0.5332	0.7121	0.0789	0.9318	0.6122	0.3034	-0.0269	0.9860	-0.0175	0.9772	0.0094	0.9863
CG18397	CG18397	1624117_at	-0.2054	0.6572	-0.1028	0.7868	-0.3964	0.2298	0.0023	0.9988	0.5860	0.0949	0.5837	0.0659	0.4503	0.7726	0.7395	0.2489	0.2892	0.6966
Rab-RP3	rab-related proteir	1624118_at	-0.0669	0.7408	0.2058	0.1676	0.4334	0.0533	0.2117	0.5836	-0.4179	0.0399	-0.6295	0.0041	0.0273	0.9831	-0.0155	0.9769	-0.0428	0.9186
CG32486	CG32486	1624119_at	-0.4357	0.0490	0.6193	0.3313	-0.5458	0.0358	-0.2674	0.5350	-0.6686	0.0097	-0.4011	0.0507	0.9448	0.4207	0.3826	0.5591	-0.5622	0.3803
---	---	1624120_at	0.1407	0.3508	-0.0179	0.8884	-0.1079	0.5847	-0.0753	0.8578	-0.0778	0.6464	-0.0025	0.9893	-0.0932	0.8455	-0.1987	0.3000	-0.1054	0.6187
knk	knickkopf	1624121_at	-0.9967	0.0312	-0.9612	0.0889	-1.3925	0.0016	-0.3276	0.5492	-0.6692	0.0259	-0.3415	0.1760	-0.0390	0.9913	-0.5574	0.4503	-0.5183	0.4978
---	---	1624122_a_at	-1.6243	0.0022	-1.0854	0.0795	-2.3091	0.0000	-0.2935	0.3222	-0.4320	0.0205	-0.1385	0.3869	0.9283	0.4128	0.0604	0.9506	-0.8679	0.1791
CG13196	CG13196	1624123_at	0.2943	0.3858	-0.3895	0.3341	0.1018	0.5590	0.4374	0.5330	0.7500	0.0461	0.3127	0.3523	-0.1510	0.8222	-0.2300	0.4073	-0.0790	0.8214
GNBP1	osiris	1624124_at	-0.7570	0.0334	-0.9783	0.0170	-0.7886	0.0007	-0.1792	0.5012	-0.3349	0.0258	-0.1558	0.2185	-0.3561	0.6955	-0.5495	0.1440	-0.1934	0.6398

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
ft	fat	1624125_at	-0.5717	0.7109	-0.5783	0.0896	-0.3568	0.1358	-0.0061	0.9988	-0.4823	0.7303	-0.4762	0.7064	-0.2036	0.9113	-0.4367	0.5158	-0.2331	0.7554
CG2321	CG2321	1624126_at	-0.0720	0.7700	0.1381	0.3938	-0.0332	0.8581	-0.0025	0.9962	0.0467	0.8326	0.0492	0.8007	-0.0246	0.9823	0.0052	0.9931	0.0299	0.9342
---	---	1624127_at	0.0748	0.7840	0.0053	0.9869	-0.2065	0.2222	-0.0039	0.9956	-0.0101	0.9705	-0.0062	0.9779	0.1490	0.8609	0.0367	0.9477	-0.1123	0.7824
CG31769	CG31769	1624128_at	2.7102	0.0028	0.5969	0.5339	1.9845	0.0048	1.1268	0.2909	1.1176	0.0770	-0.0092	0.9906	-0.3874	0.8850	-1.1799	0.2278	-0.7925	0.4475
CG7708	CG7708	1624129_s_at	-0.7024	0.1410	-0.2093	0.0986	-0.5223	0.0783	-0.2661	0.6893	-0.3051	0.3298	-0.0391	0.9175	-0.1445	0.8480	-0.1768	0.5972	-0.0323	0.9416
Obp56h	Odorant-binding p	1624130_at	0.1022	0.5655	0.2006	0.3705	-0.0491	0.7807	-0.0802	0.8500	0.0141	0.9516	0.0943	0.5316	0.1224	0.8507	0.0470	0.9086	-0.0754	0.8230
---	---	1624131_at	0.0687	0.7748	0.2697	0.1272	0.3025	0.1586	0.1786	0.6107	0.0092	0.9733	-0.1694	0.2871	0.1193	0.9015	0.1803	0.6382	0.0609	0.9011
CG4447	CG4447	1624132_at	0.1762	0.4261	0.2396	0.1825	0.1212	0.4119	-0.1356	0.7658	0.0064	0.9822	0.1420	0.4365	-0.0662	0.8981	0.0062	0.9873	0.0724	0.7439
---	---	1624133_at	0.1431	0.3876	0.4462	0.0419	0.3021	0.0910	-0.2311	0.5019	-0.0294	0.9077	0.2017	0.2177	0.1480	0.7707	0.1946	0.3597	0.0466	0.8729
NC2alpha	NC2alpha	1624134_a_at	0.5600	0.0139	0.3752	0.0804	0.4478	0.0472	0.1462	0.7604	0.1663	0.4368	0.0201	0.9375	-0.0157	0.9869	0.0173	0.9620	0.0331	0.9121
HBS1	HBS1	1624135_at	-0.1826	0.5467	0.2433	0.3636	0.5484	0.0129	0.4044	0.3008	-0.0407	0.8966	-0.4451	0.0370	0.0846	0.9495	0.3761	0.3327	0.2914	0.4801
---	---	1624136_at	-0.1078	0.6635	0.0166	0.9154	0.0674	0.7220	-0.0463	0.9375	-0.0693	0.7298	-0.0230	0.9116	-0.1017	0.8968	-0.0232	0.9620	0.0785	0.8307
CG11911	CG11911	1624137_at	-0.0615	0.9523	-0.7612	0.1655	-0.6356	0.1994	0.1809	0.9311	-0.0151	0.9875	-0.1961	0.7725	-0.0207	0.9964	-0.4344	0.7439	-0.4138	0.7513
CG5800	CG5800	1624138_at	0.1445	0.6917	0.0057	0.9771	0.7209	0.0027	0.4398	0.2506	0.4674	0.0460	0.0276	0.9214	-0.2800	0.8122	0.5923	0.2101	0.8723	0.1092
PhKgamma	phosphorylase kin	1624139_at	0.0431	0.8517	-0.2871	0.1219	0.0525	0.8097	-0.1251	0.6673	-0.4808	0.0049	-0.3557	0.0115	-0.5982	0.3712	-0.8556	0.0391	-0.2574	0.4865
---	---	1624140_at	0.1884	0.2452	0.0287	0.7823	0.1459	0.3303	0.0643	0.9186	0.0731	0.7511	0.0088	0.9711	0.0237	0.9778	-0.0742	0.7604	-0.0978	0.6567
CG4901	CG4901	1624141_at	-0.1247	0.4410	-0.3772	0.3171	-0.2424	0.2180	-0.0395	0.9635	0.2473	0.2817	0.2868	0.1591	-0.0976	0.9296	0.0232	0.9685	0.1208	0.7744
l(2)s5379	lethal (2) s5379	1624142_a_at	0.4910	0.0364	1.0204	0.0287	0.6691	0.0019	-0.0796	0.8544	-0.1668	0.2906	-0.0871	0.5760	0.2071	0.7822	0.3564	0.2518	0.1493	0.6731
CG12071	CG12071	1624143_a_at	-0.0512	0.9288	-0.1806	0.5021	-0.3340	0.1633	-0.1564	0.7493	0.1180	0.6164	0.2744	0.1484	0.0085	0.9974	-0.1645	0.8356	-0.1730	0.8149
CG13566	CG13566	1624144_at	0.1114	0.5805	0.1534	0.3828	0.5906	0.0105	0.1765	0.6545	-0.0790	0.7214	-0.2554	0.1307	-0.0567	0.9589	0.1507	0.6660	0.2074	0.5341
isopeptidase-T-3	isopeptidase T-3	1624145_a_at	0.1560	0.3006	-0.1984	0.3552	-0.4521	0.0247	-0.1227	0.7023	0.4400	0.0096	0.5627	0.0019	0.0486	0.9677	0.0481	0.9225	-0.0006	0.9992
CG5174	CG5174	1624146_at	0.0230	0.9234	0.2984	0.1609	0.3224	0.0991	-0.2077	0.5149	-0.2453	0.1438	-0.0376	0.8473	-0.1954	0.8222	0.1073	0.8202	0.3027	0.4110
---	---	1624147_at	-0.1484	0.6301	0.1369	0.3568	-0.2607	0.1742	-0.1493	0.8578	-0.2384	0.4451	-0.0891	0.7911	0.0568	0.9646	0.0224	0.9685	-0.0343	0.9433
---	---	1624148_at	0.1824	0.3960	0.0652	0.6104	0.0985	0.5615	-0.0733	0.9186	0.0214	0.9485	0.0947	0.6820	-0.1209	0.8379	-0.1463	0.5672	-0.0254	0.9408
CG4669 /// DsimCG4669	CG4669	1624149_at	0.0154	0.9528	-0.0854	0.5075	0.0387	0.8321	0.1049	0.8164	0.0317	0.8959	-0.0732	0.6943	-0.0644	0.9142	-0.1218	0.5823	-0.0573	0.8261
FKBP59	FK506-binding pro	1624150_at	0.0486	0.7754	0.2564	0.3020	0.0560	0.7284	0.0193	0.9777	0.0896	0.6427	0.0703	0.6980	0.2318	0.7230	0.3644	0.1798	0.1326	0.6630
Ccn	Ccn	1624151_a_at	-1.8133	0.0041	-0.1726	0.3837	-0.8960	0.0077	-0.6509	0.1658	-1.5744	0.0007	-0.9235	0.0040	-0.0208	0.9841	-0.1143	0.6599	-0.0935	0.7296
CG11851	CG11851	1624152_at	1.0377	0.0014	0.9440	0.0918	1.0065	0.0007	-0.0603	0.9248	0.1547	0.4366	0.2150	0.2099	-0.1321	0.9105	-0.0237	0.9735	0.1084	0.8387
Tina-1	Troponin C-akin-1	1624153_at	0.3203	0.2509	1.4167	0.0440	1.3212	0.0007	0.1497	0.7777	-0.8359	0.0033	-0.9856	0.0010	0.4009	0.7733	0.3578	0.5753	-0.0431	0.9613
ial	Aurora B	1624154_at	0.2462	0.3930	-0.0218	0.9640	-0.0456	0.8067	-0.1991	0.8202	-0.0012	0.9980	0.1979	0.5593	0.0495	0.9514	-0.0397	0.9157	-0.0892	0.7492
---	---	1624155_at	0.1369	0.5568	0.0832	0.6025	0.1910	0.2089	0.0859	0.8721	0.0070	0.9804	-0.0789	0.6853	0.1484	0.8099	0.0612	0.8644	-0.0872	0.7734
Ugt86Da	Ugt86Da	1624156_at	-0.1182	0.6099	0.0816	0.7042	-0.6905	0.0161	-0.2544	0.4596	-0.5645	0.0090	-0.3101	0.0661	0.5778	0.5638	-0.3335	0.4684	-0.9113	0.0820
CG8520	CG8520	1624157_at	-0.8945	0.0034	-0.3608	0.0456	-0.6038	0.0028	-0.1679	0.5724	-0.2492	0.1009	-0.0813	0.5928	0.1279	0.8608	0.1907	0.5302	0.0627	0.8737
CG14481	CG14481	1624158_at	-0.4405	0.2024	-0.2041	0.4364	-1.1371	0.0030	-0.6282	0.3097	-0.6691	0.0681	-0.0409	0.9264	0.0350	0.9868	-0.5767	0.2163	-0.6117	0.2201
Cyp9h1	Cyp9h1	1624159_at	-0.6879	0.3640	-1.8030	0.0725	-1.9504	0.0002	-0.2590	0.7135	0.2320	0.4910	0.4910	0.0877	-0.1505	0.9709	-0.9518	0.3469	-0.8013	0.4558
CG11329	CG11329	1624160_at	0.3317	0.4062	-0.0675	0.6725	0.3070	0.3111	0.0168	0.9904	0.5796	0.0635	0.5628	0.0466	-0.3800	0.6749	0.1870	0.6437	0.5670	0.1538
---	---	1624161_at	0.1689	0.4396	-0.0450	0.6680	-0.1547	0.2972	-0.0078	0.9937	0.1528	0.4336	0.1606	0.3525	0.0075	0.9939	-0.1866	0.4006	-0.1942	0.3953
---	---	1624162_at	0.3788	0.0397	-0.0031	0.9807	0.3113	0.1055	-0.0233	0.9745	0.1226	0.5175	0.1459	0.3773	-0.0894	0.8960	0.0034	0.9950	0.0928	0.7522
---	---	1624163_at	0.1361	0.4316	0.0224	0.8535	0.0921	0.7106	0.0837	0.8289	0.0889	0.5818	0.0052	0.9776	0.0594	0.9514	0.0368	0.9364	-0.0226	0.9565
GlcAT-P	GlcAT-P	1624164_s_at	1.2912	0.0180	1.2724	0.0307	0.8799	0.0096	-0.4514	0.5008	-0.0336	0.9498	0.4177	0.1871	-0.2089	0.8956	0.0158	0.9906	0.2247	0.7434
Psn	Dpresenilin	1624165_a_at	0.1449	0.4402	0.0972	0.6822	0.4214	0.0176	0.2040	0.4356	0.3278	0.0299	0.1238	0.3475	0.0323	0.9816	0.3142	0.3236	0.2819	0.3953
---	---	1624166_x_at	0.0487	0.7344	-0.0106	0.9389	0.1788	0.3995	-0.0044	0.9956	-0.0850	0.6687	-0.0806	0.6555	-0.1073	0.8744	-0.0708	0.8445	0.0365	0.9211
---	---	1624167_at	0.0929	0.6664	-0.1714	0.4042	-0.1212	0.4192	0.2121	0.4324	0.2540	0.0875	0.0419	0.8013	-0.0950	0.9128	-0.1926	0.5453	-0.0976	0.7889
CG14411 /// DereCG14411	CG14411	1624168_at	0.1550	0.8588	0.4080	0.4860	-0.1036	0.7004	0.0835	0.8932	0.1508	0.4876	0.0673	0.7684	0.6896	0.8049	0.5509	0.6686	-0.1387	0.9329
Ilp7	Drosophila insulin	1624169_at	-0.5392	0.0163	0.0055	0.9772	-0.4836	0.0247	-0.1996	0.6837	-0.3796	0.0960	-0.1800	0.3959	-0.0458	0.9587	-0.1265	0.6319	-0.0807	0.7799
CG15368	CG15368	1624170_at	-0.1470	0.4869	0.7237	0.0351	0.1055	0.5927	-0.1307	0.7948	-0.9029	0.0017	-0.7722	0.0020	0.3665	0.5754	-0.0410	0.9291	-0.4075	0.1880
CG13488	CG13488	1624171_at	-0.1073	0.5968	0.0334	0.8722	-0.2454	0.1567	0.0999	0.8623	0.2968	0.1466	0.1969	0.2898	0.0855	0.8940	0.2028	0.3887	0.1172	0.6476
CG33966	CG33966	1624172_at	0.2123	0.6041	-0.0924	0.3957	0.0666	0.6928	-0.0538	0.9598	-0.0908	0.7909	-0.0370	0.9137	-0.1875	0.8846	-0.3241	0.5246	-0.1366	0.8245
CG7264	CG7264	1624173_at	-0.0532	0.8298	-0.1947	0.1300	-0.0659	0.7737	0.1485	0.8074	0.1914	0.4480	0.0429	0.8814	-0.0947	0.8882	-0.0793	0.8062	0.0154	0.9678
---	---	1624174_at	-0.0437	0.8461	-0.0262	0.7923	0.0494	0.8061	0.0689	0.8738	0.1790	0.2346	0.1102	0.4365	-0.1153	0.8909	0.0713	0.8750	0.1866	0.5801
caup	iroquois	1624175_at	-0.0927	0.8161	-0.0181	0.9189	0.0035	0.9894	0.0699	0.9558	-0.0916	0.8320	-0.1615	0.6428	-0.0028	0.9978	-0.0008	0.9994	0.0020	0.9962
CG10590	CG10590	1624176_at	0.3819	0.3024	1.0986	0.0523	0.9577	0.0011	-0.1102	0.8831	-0.1354	0.6356	-0.0252	0.9365	0.0955	0.9666	0.7419	0.2077	0.6464	0.2993
Cpr67Fa1	CG7941	1624177_at	0.3029	0.2256	0.0832	0.4737	0.1640	0.4311	0.0074	0.9943	0.0174	0.9576	0.0100	0.9711	-0.0202	0.9853	-0.1185	0.6704	-0.0983	0.7342
CG3777 /// DsimCG3777	CG3777	1624178_s_at	0.0680	0.7799	0.0206	0.8886	0.2858	0.2810	0.2899	0.4517	0.2252	0.2851	-0.0647	0.7811	0.0899	0.8597	0.0176	0.9587	-0.0723	0.7607
CG31680	CG31680	1624179_at	-0.2752	0.2493	0.1667	0.5885	0.0364	0.9036	-0.0404	0.9673	-0.2134	0.4266	-0.1730	0.4848	-0.0468	0.9716	0.1151	0.7718	0.1619	0.6435
Psf3	Psf3	1624180_at	-0.1975	0.6681	-0.4876	0.2370	-0.4537	0.0972	-0.1035	0.9491	0.0290	0.9674	0.1325	0.7943	-0.0844	0.9610	-0.1733	0.7656	-0.0889	0.8926
CG8062 /// DmirCG8062	CG8062	1624181_at	-0.0833	0.7140	0.4611	0.2223	0.9963	0.0012	0.0135	0.9922	-0.1726	0.5776	-0.1861	0.4985	-0.6055	0.3273	0.4101	0.1948	1.0157	0.0327
---	---	1624182_at	0.1451	0.3723	-0.0327	0.7923	0.2478	0.2394	0.1501	0.7582	0.1901	0.3746	0.0400	0.8714	0.0500	0.9451	0.0812	0.7572	0.0312	0.9182
Fas1	fasciclin I	1624183_a_at	-0.7531	0.0491	-0.9217	0.0308	-0.8362	0.0039	-0.0735	0.9223	-0.3543	0.1177	-0.2808	0.1664	0.0195	0.9928	-0.2770	0.6093	-0.2965	0.5819
---	---	1624184_at	0.2679	0.1051	0.3537	0.1522	0.7415	0.0088	0.1221	0.8194	0.0039	0.9899	-0.1182	0.5692	-0.2487	0.7726	0.0975	0.8460	0.3463	0.3622
CG1041	CG1041	1624185_at	0.7957	0.0055	0.9852	0.0507	1.0478	0.0004	0.1748	0.6160	0.2317	0.1811	0.0569	0.7666	0.0406	0.9800	0.4702	0.1798	0.4296	0.2527
CG16717	CG16717	1624186_at	-1.0882	0.0012	0.2508	0.0925	0.1758	0.4405	-0.2022	0.6577	-0.7836	0.0043	-0.5814	0.0098	-0.0905	0.9092	0.5903	0.0571	0.6808	0.0542
fs(1)Yb	no ovaries	1624187_at	0.2261	0.2507	-0.2385	0.4848	-0.0157	0.9459	0.1866	0.6441	0.3520	0.0752	0.1654	0.3594	0.0017	0.9997	-0.1640	0.7550	-0.1657	0.7444
CG31392	CG31392	1624188_at	-0.3825	0.0883	0.6881	0.0887	0.8759	0.0004	-0.0291	0.9777	-0.8658	0.0069	-0.8366	0.0048	-0.1040	0.8810	0.2706	0.2942	0.3746	0.1823
CG12236	CG12236	1624189_at	0.2780	0.5124	0.1252	0.5678	-0.4053	0.0563	-0.3629	0.4861	0.3050	0.2747	0.6680	0.0153	0.1121	0.9459	0.0814	0.9162	-0.0307	0.9675
CG31637	CG31637	1624190_at	-0.6221	0.1364	-0.3318	0.4672	-1.2188	0.0006	-0.3072	0.7023	0.0052	0.9922	0.3123	0.3579	0.5232	0.7305	0.2819	0.7012	-0.2413	0.7484
cmet	CENP-meta	1624191_at	0.4506	0.2133	1.3716	0.0291	-0.0425	0.8585	-0.4917	0.2994	0.4894	0.0805	0.9811	0.0027	0.8625	0.3517	1.5242	0.0222	0.6617	0.2051
r-cup	ryder cup	1624192_at	0.1131	0.6807	-0.0328	0.8037	0.1163	0.5037	0.0271	0.9777	0.1476	0.5724	0.1205	0.6248	-0.1498	0.7953	-0.0746	0.8122	0.0752	0.7969
CG3983 /// DyakCG3983	CG3983	1624193_s_at	0.2759	0.2214	-0.0282	0.9642	0.0237	0.9419	-0.0076	0.9937	0.4739	0.0276	0.4815	0.0163	-0.0647	0.9816	0.1756	0.8268	0.2403	0.7307

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Taf12L	ryan express	1624194_a_at	0.2915	0.1918	0.2347	0.3071	0.3319	0.1966	0.0486	0.9496	-0.1159	0.6253	-0.1644	0.4098	0.0913	0.9296	-0.0570	0.9115	-0.1483	0.6881
sug	sugarbabe	1624195_at	2.3517	0.0108	0.7620	0.2302	1.4684	0.0455	0.0077	0.9973	0.9294	0.1489	0.9217	0.1100	-0.7455	0.7768	-0.5739	0.6425	0.1715	0.9125
---	---	1624196_at	0.0452	0.8922	-0.0167	0.8996	-0.1521	0.3081	-0.1351	0.8265	0.0443	0.8906	0.1795	0.4201	0.0760	0.9250	0.0653	0.8628	-0.0106	0.9816
CG6652	CG6652	1624197_a_at	0.2716	0.2011	0.0848	0.5363	0.3349	0.0400	-0.0680	0.8908	0.0990	0.5842	0.1670	0.2661	-0.1887	0.8202	0.0023	0.9988	0.1909	0.6060
CG12582	CG12582	1624198_at	0.6830	0.0368	0.2233	0.5452	0.2389	0.2771	0.0550	0.9040	0.1748	0.2398	0.1198	0.3847	0.0433	0.9862	-0.1555	0.8351	-0.1988	0.7571
NPFR76F	Neuropeptide F-lil	1624199_at	1.3987	0.0696	2.2669	0.0074	1.4758	0.0002	-0.2667	0.8498	0.1784	0.7727	0.4451	0.3444	0.4529	0.7644	1.0659	0.1030	0.6131	0.3503
CG8679	CG8679	1624200_at	1.9915	0.0027	1.9330	0.0114	1.9085	0.0001	0.3286	0.3836	0.7091	0.0059	0.3805	0.0505	0.2952	0.8157	0.6128	0.2254	0.3176	0.5672
CG15498	CG15498	1624201_at	0.2901	0.1561	0.0922	0.5696	0.0682	0.7539	0.1112	0.7556	0.1763	0.2528	0.0651	0.6865	0.0275	0.9816	-0.0253	0.9533	-0.0528	0.8873
CG2127	CG2127	1624202_a_at	0.0139	0.9349	-0.0244	0.9308	-0.0319	0.8998	0.0733	0.9452	0.1937	0.5543	0.1205	0.7087	0.0689	0.9238	0.1086	0.6854	0.0397	0.9054
Gli	gliotactin	1624203_s_at	0.2807	0.6588	-0.1266	0.7192	-0.1287	0.5004	0.2769	0.6354	0.8364	0.0106	0.5595	0.0361	0.1996	0.9340	0.3022	0.7364	0.1026	0.9231
CG4483	CG4483	1624204_at	0.3275	0.2752	-0.0216	0.8386	0.1655	0.5085	0.0948	0.9185	0.0692	0.8514	-0.0256	0.9425	-0.1008	0.9237	-0.0723	0.8850	0.0285	0.9533
Tom40	Translocase of ou	1624205_at	-0.2095	0.2778	0.1994	0.4005	0.2514	0.1708	0.1634	0.6015	-0.1038	0.5406	-0.2672	0.0622	0.1264	0.8861	0.3512	0.2729	0.2249	0.5163
CG11300	CG11300	1624206_at	0.0567	0.7632	0.1176	0.6480	0.1045	0.4840	0.0753	0.8822	0.0396	0.8624	-0.0357	0.8599	0.0553	0.9296	-0.0353	0.9085	-0.0906	0.6848
btsz	bitesize	1624207_at	-0.3861	0.4572	0.5572	0.4162	0.5670	0.0404	-0.0479	0.9666	-0.5836	0.0564	-0.5357	0.0507	0.0416	0.9914	0.3353	0.7490	0.2937	0.7792
Arf51F	ADP ribosylation f	1624208_s_at	-0.4303	0.0626	0.4623	0.0695	-0.1944	0.2520	-0.2850	0.3521	-0.5232	0.0096	-0.2381	0.1274	0.2273	0.7324	0.3365	0.2217	0.1091	0.7410
---	---	1624209_at	0.4524	0.0806	0.3348	0.0588	0.4273	0.0321	0.1635	0.6998	-0.0438	0.8646	-0.2073	0.2343	0.0160	0.9898	-0.0332	0.9402	-0.0491	0.8949
CG32699	CG32699	1624210_at	0.0228	0.9505	0.8592	0.2350	-0.1359	0.6462	0.2222	0.7349	1.3519	0.0014	1.1298	0.0018	1.3211	0.3166	2.2182	0.0215	0.8972	0.2137
CG9005	CG9005	1624211_at	0.0682	0.8561	-0.0090	0.9882	0.6697	0.0348	0.0581	0.9496	-0.0318	0.9295	-0.0899	0.7450	-0.5111	0.7042	0.0135	0.9921	0.5246	0.3627
CG6784	CG6784	1624212_at	-0.0061	0.9804	-0.1294	0.3655	0.0185	0.9299	0.0405	0.9351	-0.0382	0.8405	-0.0787	0.5925	-0.1621	0.8033	-0.0763	0.8329	0.0858	0.7925
CG15535	CG15535	1624213_at	-0.3457	0.0573	-0.6081	0.0621	-0.6823	0.0018	-0.1609	0.6940	0.0517	0.8294	0.2125	0.2085	0.0619	0.9238	-0.0371	0.9104	-0.0989	0.6731
CG9437	CG9437	1624214_at	-0.0997	0.7518	0.3556	0.2314	0.8435	0.0123	0.3413	0.5841	-0.1327	0.7194	-0.4740	0.0947	-0.0745	0.9589	0.3659	0.3571	0.4403	0.2923
DIP1	klett	1624215_s_at	-0.4081	0.1142	-0.0769	0.7888	-0.4598	0.0107	-0.3241	0.3793	-0.1045	0.6572	0.2197	0.2401	-0.0355	0.9826	0.2066	0.6041	0.2421	0.5391
Ccp84Ae	cuticle cluster 4	1624216_at	0.1554	0.5240	0.1358	0.3549	0.1733	0.3850	-0.0341	0.9745	-0.0897	0.7769	-0.0556	0.8553	-0.0618	0.9503	-0.1081	0.7595	-0.0463	0.9095
---	---	1624217_at	0.2526	0.0888	0.1084	0.5221	0.1328	0.4483	0.0666	0.9136	0.0007	0.9981	-0.0659	0.7549	-0.0260	0.9726	-0.1177	0.5542	-0.0917	0.6569
---	---	1624218_at	0.2304	0.4199	0.0190	0.8946	0.0861	0.6574	0.1854	0.6622	0.3096	0.1267	0.1242	0.5316	0.1562	0.8270	0.0091	0.9884	-0.1470	0.6440
fz2	D-frizzled2	1624219_s_at	-0.2419	0.3751	0.4676	0.3005	-0.1664	0.4719	-0.1670	0.6506	-0.5711	0.0070	-0.4041	0.0193	0.3337	0.8076	0.0212	0.9844	-0.3125	0.6121
---	---	1624220_s_at	0.0013	0.9950	-0.2944	0.1874	-0.0859	0.6529	0.0545	0.9451	0.3236	0.1498	0.2691	0.1812	-0.2115	0.8013	-0.0607	0.9095	0.1508	0.7000
CG32987	CG32987	1624221_at	0.1300	0.5137	0.0386	0.8480	0.1418	0.5180	0.1548	0.7857	0.1250	0.6333	-0.0298	0.9174	-0.1567	0.8049	0.0095	0.9854	0.1662	0.5541
CG15814 /// DsimCG15814	CG15814	1624222_s_at	0.0509	0.8503	1.1700	0.0328	1.1605	0.0011	-0.1360	0.7409	-0.6808	0.0034	-0.5448	0.0056	-0.1337	0.9312	0.4685	0.3397	0.6022	0.2499
---	---	1624223_at	0.3191	0.4147	0.0807	0.7750	-0.0873	0.6710	-0.0936	0.9314	0.0444	0.9220	0.1380	0.6829	0.2355	0.7187	-0.0817	0.8166	-0.3173	0.2574
---	---	1624224_at	0.1985	0.2574	-0.3295	0.1349	-0.3068	0.0521	0.1615	0.6578	0.4351	0.0213	0.2736	0.0821	-0.0802	0.9543	-0.2952	0.4810	-0.2150	0.6271
danr	Distal antenna rel	1624225_at	0.0472	0.9157	-0.0921	0.5590	0.0650	0.7194	0.0922	0.9011	0.0448	0.8906	-0.0474	0.8657	-0.0053	0.9977	-0.0750	0.9049	-0.0697	0.9023
trus	toys are us	1624226_at	-0.2117	0.3323	0.7800	0.0426	1.3280	0.0002	0.2641	0.4517	-0.5144	0.0149	-0.7786	0.0014	-0.4305	0.5259	0.3579	0.2519	0.7884	0.0540
CG31066	CG31066	1624227_at	0.3261	0.1247	0.1333	0.4402	-0.0849	0.5999	-0.0434	0.9543	0.0913	0.7022	0.1347	0.4989	0.1812	0.8016	0.0960	0.8033	-0.0851	0.8209
mRpL54	mitochondrial ribo	1624228_at	0.0738	0.8192	0.0983	0.7729	0.0723	0.8363	-0.1625	0.8336	0.0763	0.8398	0.2388	0.3785	-0.1220	0.9467	0.0963	0.9110	0.2183	0.7342
CG15705	CG15705	1624229_at	0.0180	0.9297	0.0195	0.8546	0.0444	0.7887	-0.0977	0.8097	0.0185	0.9354	0.1162	0.4444	-0.0312	0.9618	0.0087	0.9787	0.0400	0.8689
Ubi-p63E	ubiquitin	1624230_s_at	-0.2889	0.1526	-0.2464	0.5283	-0.4680	0.0103	0.1164	0.8096	0.2034	0.2938	0.0870	0.6620	0.3479	0.7215	0.3232	0.4308	-0.0246	0.9688
Arf79F	ADP-ribosylation f	1624231_s_at	0.2744	0.1163	0.4771	0.0383	0.3864	0.0409	-0.0373	0.9438	0.1848	0.2212	0.2221	0.1005	0.1618	0.8114	0.4411	0.1199	0.2793	0.3307
CG13891	CG13891	1624232_at	0.3236	0.1469	0.1199	0.6040	0.2901	0.3248	0.2104	0.6869	0.1498	0.5680	-0.0607	0.8263	0.0971	0.9238	0.0238	0.9653	-0.0733	0.8699
Gr8a	Gustatory recepto	1624233_at	0.0421	0.8649	0.0583	0.5748	0.0599	0.7546	0.1006	0.8794	0.0726	0.7940	-0.0280	0.9195	0.0247	0.9816	-0.0833	0.7577	-0.1080	0.6601
CG7735	CG7735	1624234_at	0.0203	0.9081	0.2187	0.3761	0.1624	0.2723	-0.1421	0.7850	-0.1589	0.4789	-0.0168	0.9506	-0.0006	0.9998	0.2152	0.3623	0.2158	0.3809
CG4285	CG4285	1624235_at	0.1483	0.6853	-0.3215	0.1578	-0.1428	0.6005	0.2508	0.7187	0.4452	0.1567	0.1944	0.5270	-0.1469	0.8480	-0.2388	0.4588	-0.0919	0.8178
CG30002	CG30002	1624236_at	2.9231	0.0010	1.0656	0.1792	1.9483	0.0007	0.6246	0.4141	1.0224	0.0242	0.3978	0.3021	-0.5176	0.7726	-0.9969	0.1827	-0.4792	0.5581
CG15099 /// DmirCG15099	CG15099	1624237_at	-0.5554	0.0537	0.2407	0.5929	0.6151	0.0229	0.3591	0.4932	-0.1567	0.6099	-0.5157	0.0444	-0.0156	0.9939	0.6751	0.1345	0.6907	0.1512
---	---	1624238_at	0.1257	0.5039	0.0277	0.8867	0.0944	0.6935	-0.0083	0.9940	0.1050	0.6867	0.1133	0.6246	0.0306	0.9739	0.0910	0.7340	0.0604	0.8354
CG8335	CG8335	1624239_at	-0.0418	0.8339	-0.0860	0.3943	0.1215	0.6485	0.1082	0.8738	0.0017	0.9960	-0.1065	0.6662	-0.0319	0.9741	-0.0532	0.8778	-0.0213	0.9496
Uch	ubiquitin carboxyl	1624240_at	-0.1902	0.3735	0.3396	0.1510	0.5174	0.0052	-0.0867	0.8512	-0.6353	0.0034	-0.5486	0.0039	-0.0779	0.9342	0.1099	0.7660	0.1878	0.5612
NaCP60E	put. sodium chanr	1624241_at	0.0000	1.0000	-0.1510	0.6862	-0.5636	0.0621	-0.4620	0.2438	-0.1556	0.5356	0.3064	0.1432	0.0075	0.9976	-0.1986	0.7543	-0.2061	0.7371
CG32301	CG32301	1624242_at	0.1738	0.4821	-0.2220	0.3955	-0.0373	0.8242	0.2909	0.5220	0.2879	0.2266	-0.0030	0.9919	0.0867	0.9164	-0.0493	0.9105	-0.1360	0.6607
CG9265	CG9265	1624243_at	-0.8405	0.2908	-0.5439	0.3485	-0.9433	0.0194	-0.6859	0.2401	-1.6218	0.0013	-0.9359	0.0089	-0.1532	0.9764	-1.3868	0.2470	-1.2336	0.3311
S6k	dS6 kinase	1624244_at	0.7249	0.1969	0.2769	0.2028	-0.2332	0.4707	-0.1007	0.9228	0.5653	0.0741	0.6660	0.0257	0.3099	0.8609	0.2322	0.7978	-0.0777	0.9382
CG15239	CG15239	1624245_at	0.0851	0.6115	-0.0203	0.9204	0.0828	0.6480	0.0236	0.9745	0.1692	0.3583	0.1455	0.3847	-0.1030	0.8956	0.0515	0.9085	0.1544	0.6239
CG11334	CG11334	1624246_s_at	0.2814	0.3578	0.9171	0.0415	1.3536	0.0002	0.3196	0.2502	-0.0906	0.6217	-0.4102	0.0134	-0.0602	0.9775	0.5110	0.3010	0.5711	0.2778
CG5543	CG5543	1624247_at	0.2046	0.2643	0.2426	0.4349	0.4068	0.0758	0.2114	0.6341	0.2846	0.1901	0.0732	0.7595	0.0592	0.9589	0.2734	0.4029	0.2141	0.5381
Aef1	Adh enhancer fac	1624248_s_at	0.0360	0.9435	0.3225	0.4555	0.4022	0.0562	0.1144	0.8862	0.5013	0.0672	0.3869	0.1090	0.1855	0.9142	0.9125	0.1300	0.7269	0.2426
---	---	1624249_at	0.2145	0.4326	-0.1494	0.4062	-0.0293	0.8948	0.0096	0.9929	-0.0568	0.8366	-0.0665	0.7809	0.1095	0.8940	-0.1664	0.6137	-0.2759	0.3809
CG32432	CG32432	1624250_at	-0.0682	0.7442	0.0845	0.7238	0.3108	0.2304	-0.0370	0.9664	-0.0519	0.8645	-0.0149	0.9592	-0.1100	0.9298	0.0352	0.9587	0.1452	0.7558
CG14235 /// CG18809 /// D	CG18809 /// Cytoc	1624251_a_at	-0.0690	0.7449	0.1867	0.4809	0.0262	0.9211	0.0272	0.9838	-0.4205	0.1661	-0.4478	0.1005	0.0969	0.9238	-0.2954	0.3733	-0.3923	0.2614
CG12592 /// sle	CG12592 /// slend	1624252_s_at	0.5160	0.1223	0.3395	0.3747	-0.1246	0.3892	-0.0430	0.9376	0.8134	0.0010	0.8564	0.0005	0.4063	0.7387	0.7867	0.1299	0.3804	0.4776
CG9125	CG9125	1624253_at	0.3645	0.2435	0.3246	0.1035	0.7102	0.0149	-0.0965	0.9037	-0.1412	0.6244	-0.0447	0.8845	-0.2773	0.6898	-0.1100	0.7424	0.1673	0.5808
CG32070	CG32070	1624254_at	0.1614	0.3232	0.1252	0.4457	0.2281	0.1334	0.1221	0.8096	0.1773	0.3909	0.0552	0.8075	-0.0709	0.8940	-0.0259	0.9353	0.0450	0.8646
---	---	1624255_at	0.1304	0.3757	0.0284	0.8465	-0.0933	0.5994	-0.2081	0.4511	0.0306	0.8786	0.2387	0.0768	-0.0405	0.9555	-0.0604	0.8290	-0.0200	0.9464
CG16857	CG16857	1624256_at	-1.4389	0.0102	0.2198	0.7175	-0.4225	0.2397	-0.8381	0.2172	-2.1492	0.0008	-1.3111	0.0041	0.0127	0.9964	-0.3168	0.6718	-0.3294	0.6512
CG18259	CG18259	1624257_at	0.0833	0.6767	-0.1694	0.2837	-0.3657	0.0280	-0.1774	0.5437	0.2951	0.0561	0.4724	0.0047	0.0804	0.9132	-0.0248	0.9525	-0.1052	0.7194
---	---	1624258_at	0.2014	0.2212	-0.0194	0.8763	0.2467	0.1337	0.0910	0.8196	0.1743	0.2670	0.0833	0.5945	-0.1751	0.7644	-0.0723	0.8231	0.1028	0.7130
CG32040	CG32040	1624259_at	-1.1216	0.0060	-2.9592	0.0029	-1.6584	0.0122	1.1412	0.3359	1.5136	0.0342	0.3723	0.5869	-0.2214	0.8331	-0.3405	0.4354	-0.1191	0.8287
CG11977	CG11977	1624260_at	0.1686	0.3318	0.0916	0.6041	0.2305	0.1703	0.1939	0.5311	0.2792	0.0854	0.0853	0.5996	0.1104	0.8446	0.1521	0.5258	0.0417	0.8949
CG14118	CG14118	1624261_at	0.1793	0.2748	0.3790	0.0494	0.1031	0.7242	-0.0887	0.8738	-0.1586	0.4325	-0.0699	0.7391	0.1046	0.8810	0.1161	0.7002	0.0115	0.9788
phyl	phyllopod	1624262_at	-0.2242	0.2709	-0.2475	0.4342	-0.2759	0.1223	-0.1949	0.7304	-0.1322	0.6417	0.0628	0.8280	0.0189	0.9870	-0.2344	0.3701	-0.2533	0.3552

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG5933	CG5933	1624263_at	0.4333	0.0412	0.0783	0.7935	0.1950	0.4209	-0.2524	0.5238	0.3105	0.1331	0.5629	0.0088	-0.3015	0.7485	-0.0154	0.9841	0.2860	0.4912
CG13408	CG13408	1624264_at	0.0902	0.6349	0.1313	0.4455	0.1166	0.4241	0.0639	0.8908	-0.0522	0.7831	-0.1161	0.4284	-0.0114	0.9928	0.0242	0.9590	0.0356	0.9282
l(1)G0196	lethal (1) G0196	1624265_at	-0.0664	0.7665	0.1831	0.5445	-0.1474	0.5976	-0.3550	0.3722	-0.5146	0.0295	-0.1596	0.4516	-0.1068	0.9483	-0.2777	0.5988	-0.1709	0.7675
Art6	Arginine methyltra	1624266_at	0.1229	0.4615	0.1926	0.2983	0.4464	0.0600	0.1257	0.7690	-0.0900	0.6613	-0.2158	0.1826	-0.1225	0.8283	0.0994	0.7112	0.2219	0.3624
CG7182	CG7182	1624267_at	-0.2581	0.2493	0.5134	0.0985	0.9010	0.0165	-0.0010	0.9988	-0.5135	0.0145	-0.5125	0.0089	-0.4209	0.7215	0.2743	0.6041	0.6952	0.1865
---	---	1624268_at	-0.1718	0.4119	0.0429	0.6754	0.2326	0.2002	-0.0599	0.9266	-0.1821	0.3604	-0.1223	0.5181	-0.2031	0.7707	-0.0303	0.9499	0.1728	0.5862
gkt	glaikit	1624269_at	0.2801	0.0935	-0.1302	0.5054	0.0492	0.7840	0.1457	0.7838	0.5546	0.0196	0.4089	0.0431	-0.2353	0.7644	0.0550	0.9140	0.2903	0.3878
---	---	1624270_at	0.1761	0.4827	-0.0648	0.6469	-0.0516	0.7430	-0.0302	0.9672	-0.0197	0.9449	0.0105	0.9660	-0.0904	0.9201	-0.2215	0.4757	-0.1311	0.7011
CG11808	CG11808	1624271_at	-0.1525	0.3967	-0.3176	0.0710	0.0255	0.8939	-0.0317	0.9684	0.1512	0.4891	0.1829	0.3380	-0.2960	0.7187	0.0958	0.8336	0.3917	0.2668
CHKov1	CHKov1	1624272_at	-0.6051	0.0443	0.3922	0.2790	0.4192	0.0579	0.0905	0.9066	-0.3651	0.1397	-0.4556	0.0457	0.0255	0.9885	0.7053	0.0795	0.6798	0.1121
CG8034 /// DmirCG8034	CG8034	1624273_at	-0.5679	0.2124	-0.5335	0.4801	-0.2709	0.3817	-0.1285	0.9048	-0.6950	0.0514	-0.5665	0.0718	-0.4483	0.8374	-0.8454	0.3340	-0.3971	0.6886
CG31953	CG31953	1624274_at	0.2382	0.2339	0.1218	0.4645	0.0532	0.7535	-0.1830	0.5918	-0.0347	0.8809	0.1483	0.3509	0.0508	0.9587	0.0943	0.7779	0.0435	0.9079
sktl	Skittles	1624275_at	0.0881	0.8114	0.4244	0.1074	0.4503	0.0113	-0.0873	0.8663	-0.3019	0.0989	-0.2145	0.1888	-0.0704	0.9545	0.0487	0.9328	0.1191	0.7807
---	---	1624276_at	-0.0042	0.9892	-0.0574	0.6362	0.1291	0.4585	0.1587	0.6908	0.2643	0.1511	0.1056	0.5638	-0.0002	0.9999	0.0059	0.9875	0.0060	0.9846
CG8273	CG8273	1624277_at	0.1221	0.6699	0.4005	0.0410	0.2557	0.1438	-0.0768	0.8899	-0.0474	0.8437	0.0294	0.8953	0.0292	0.9816	0.3018	0.2818	0.2726	0.3601
mRpL2	mitochondrial ribo	1624278_at	-0.2247	0.2692	0.3713	0.0859	0.2873	0.1790	0.1327	0.8134	-0.4142	0.0651	-0.5469	0.0136	0.1575	0.8655	0.1635	0.6957	0.0059	0.9924
TotZ	Turandot Z	1624279_at	-0.1230	0.5766	0.0646	0.7068	0.1180	0.5736	-0.1111	0.8500	-0.1329	0.5649	-0.0219	0.9344	-0.0952	0.9095	-0.0261	0.9585	0.0691	0.8604
CG2263	CG2263	1624280_at	-0.0198	0.9212	1.3271	0.0054	1.3484	0.0002	0.0806	0.8895	-0.7575	0.0029	-0.8381	0.0011	0.0455	0.9589	0.7039	0.0276	0.6584	0.0438
---	---	1624281_at	0.1406	0.5460	-0.0518	0.6311	-0.0149	0.9625	0.1082	0.7930	0.1322	0.4502	0.0240	0.9060	0.0721	0.9421	0.0274	0.9567	-0.0447	0.9168
---	---	1624282_s_at	-0.1050	0.6710	-0.0792	0.4617	0.1245	0.4808	0.0571	0.9436	0.0227	0.9465	-0.0344	0.9003	-0.0677	0.9238	0.0355	0.9222	0.1032	0.6881
Cpr47Eb	CG13224	1624283_at	0.7245	0.0345	0.1639	0.3691	0.2367	0.5818	0.1505	0.8708	0.1874	0.5959	0.0369	0.9258	0.1817	0.8861	-0.0786	0.9156	-0.2603	0.6130
DyakCG10473 /// hkl	CG10473 /// hook	1624284_a_at	-0.1809	0.6390	0.5496	0.0785	0.4806	0.0152	-0.1201	0.8649	-0.2890	0.2491	-0.1689	0.4786	-0.1174	0.9137	0.3508	0.3409	0.4681	0.2312
CG3088	CG3088	1624285_at	0.4222	0.3206	0.0915	0.5175	-0.0628	0.8142	-0.0162	0.9852	0.2476	0.2039	0.2638	0.1296	-0.0771	0.9764	-0.0452	0.9650	0.0319	0.9743
janB	janus B	1624286_at	0.0227	0.9524	-0.3420	0.1116	-0.2511	0.1375	0.1969	0.5854	0.2981	0.1040	0.1012	0.5809	-0.1054	0.9238	-0.1459	0.7339	-0.0405	0.9371
CG17217	CG17217	1624287_at	0.0555	0.7404	-0.0059	0.9591	0.2114	0.2542	0.1230	0.7121	0.0594	0.7399	-0.0637	0.6893	-0.0480	0.9474	0.0468	0.8832	0.0948	0.7020
CG10654	CG10654	1624288_at	-0.0857	0.7110	-0.1094	0.4867	0.1088	0.5719	0.2025	0.6265	-0.1530	0.4764	-0.3554	0.0571	0.1155	0.8521	-0.0824	0.8008	-0.1978	0.4557
CG10133	CG10133	1624289_at	0.4318	0.2250	0.5910	0.0348	0.8432	0.0064	-0.0427	0.9639	-0.2716	0.2784	-0.2289	0.3123	-0.2022	0.8672	-0.1573	0.7943	0.0449	0.9469
CG4752	CG4752	1624290_at	-1.2303	0.0025	0.6505	0.1047	0.4623	0.0408	-0.5867	0.2037	-2.1557	0.0002	-1.5690	0.0004	-0.0479	0.9792	-0.0444	0.9485	0.0035	0.9965
CG6685	CG6685	1624291_at	-0.2256	0.2983	0.2074	0.2861	0.3024	0.1594	-0.4266	0.2495	-0.3912	0.0767	0.0354	0.8928	-0.3256	0.6955	0.1409	0.7199	0.4665	0.1983
Acp26Ab	Accessory gland-s	1624292_a_at	0.0026	0.9889	-0.0424	0.7385	0.0695	0.7652	-0.1192	0.8081	-0.1577	0.4366	-0.0384	0.8662	0.0548	0.9331	-0.0483	0.8712	-0.1031	0.6473
CG40497	CG40497	1624293_at	0.0529	0.8619	0.1862	0.3372	0.2193	0.1935	-0.2159	0.5799	-0.0040	0.9896	0.2119	0.2356	-0.2485	0.7215	-0.0210	0.9650	0.2275	0.4503
cngl	CNG channel-like	1624294_at	0.1796	0.2516	0.0568	0.5954	0.0752	0.6436	-0.0036	0.9956	-0.0485	0.8534	-0.0449	0.8483	-0.0185	0.9816	-0.0713	0.7323	-0.0527	0.8123
dpr4	dpr4	1624295_at	0.2682	0.0992	0.0825	0.5495	-0.1019	0.5914	0.0249	0.9745	0.1213	0.5589	0.0963	0.6230	0.1287	0.7721	-0.0517	0.8390	-0.1804	0.3552
CG14508	CG14508	1624296_at	0.6581	0.0767	0.4405	0.0824	0.4817	0.1790	0.0936	0.9029	0.2583	0.3031	0.1648	0.4881	-0.0489	0.9816	0.2107	0.7048	0.2596	0.6238
drl	linotte	1624297_at	1.9221	0.0040	3.1607	0.0287	2.9551	0.0009	0.0363	0.9872	-1.1584	0.0387	-1.1947	0.0217	0.1776	0.9611	0.0136	0.9951	-0.1640	0.9080
CG33257	CG33257	1624298_at	0.1738	0.4552	0.0025	0.9877	0.0731	0.7834	-0.0787	0.9445	0.0558	0.8997	0.1345	0.6914	-0.0777	0.9284	-0.0818	0.8244	-0.0041	0.9933
---	---	1624299_at	0.0396	0.8367	0.1547	0.2929	0.1332	0.4426	-0.0032	0.9960	-0.1619	0.4225	-0.1587	0.3798	-0.0485	0.9514	-0.0937	0.7340	-0.0453	0.8893
nmo	nemo	1624300_s_at	-0.6531	0.0476	-0.1676	0.3113	-0.4196	0.0613	-0.1641	0.7146	-0.2797	0.1696	-0.1156	0.5690	0.1857	0.8049	0.0009	0.9996	-0.1848	0.5795
Bre1	lethal (3) 01640	1624301_at	0.2411	0.4472	0.2428	0.3486	0.3899	0.0594	0.1473	0.8738	0.3339	0.3076	0.1866	0.5602	0.0629	0.9657	0.4839	0.2023	0.4209	0.2964
CG7903	CG7903	1624302_at	-0.7524	0.0068	-0.0892	0.6058	-0.0727	0.6373	-0.0027	0.9962	-0.3725	0.0591	-0.3697	0.0396	-0.1565	0.8558	0.0859	0.8614	0.2424	0.5092
---	---	1624303_at	0.2773	0.1942	0.2142	0.2710	0.2971	0.3096	-0.1152	0.8723	-0.0663	0.8358	0.0489	0.8667	0.0569	0.9409	-0.0154	0.9688	-0.0723	0.8036
---	---	1624304_s_at	-0.4790	0.0665	-0.6386	0.2579	-0.9441	0.0005	-0.3314	0.3847	-0.1313	0.5731	0.2001	0.3066	-0.0445	0.9849	-0.2470	0.6820	-0.2025	0.7451
Kr-h2	Kruppel homolog	1624305_at	-0.0723	0.7800	0.1065	0.6951	0.0608	0.7920	-0.2009	0.4749	0.4110	0.0144	0.6120	0.0015	-0.1707	0.8815	0.5543	0.1927	0.7251	0.1300
CG17195	CG17195	1624306_at	0.0859	0.6682	0.1486	0.4537	0.1385	0.5540	0.2409	0.6166	0.0839	0.7717	-0.1570	0.4965	0.0369	0.9781	-0.0211	0.9682	-0.0580	0.8949
CG10801	CG10801	1624307_at	0.2527	0.1946	0.1353	0.3435	0.1551	0.5945	-0.0881	0.9138	0.0478	0.8906	0.1360	0.5978	0.0445	0.9589	0.0714	0.8267	0.0269	0.9382
Sgs3	group IV	1624308_at	0.2126	0.3120	-0.0600	0.8523	0.3561	0.0605	0.1947	0.5680	0.1590	0.3737	-0.0357	0.8612	-0.1939	0.8076	-0.1553	0.6725	0.0387	0.9342
---	---	1624309_s_at	0.0956	0.5998	0.0711	0.5260	0.0362	0.8345	-0.0534	0.9117	0.0129	0.9550	0.0663	0.6794	0.1048	0.8740	-0.0479	0.9037	-0.1527	0.5808
CG4753	CG4753	1624310_s_at	1.0263	0.2251	1.2038	0.0845	2.4895	0.0003	0.6576	0.7009	-0.2337	0.8099	-0.8913	0.2041	-0.5769	0.7464	-0.1528	0.8897	0.4242	0.6016
Ccp84Ac	cuticle cluster 6	1624311_at	0.4053	0.1958	0.3385	0.1915	0.2304	0.4000	-0.0061	0.9956	0.0168	0.9743	0.0229	0.9572	0.2815	0.7673	0.0103	0.9911	-0.2712	0.5196
CG3326	fidgetin_DROME	1624312_at	0.0774	0.6399	-0.2952	0.1698	-0.4116	0.2258	-0.0781	0.9154	0.3906	0.0910	0.4687	0.0305	-0.0888	0.9514	-0.0958	0.8779	-0.0070	0.9924
---	---	1624313_at	0.4218	0.0365	-0.0621	0.6844	0.0498	0.7737	-0.0453	0.9538	0.1377	0.5492	0.1830	0.3509	-0.1226	0.8331	-0.3031	0.1967	-0.1805	0.4726
CG40397	CG40397	1624314_at	0.0613	0.6769	-0.0690	0.6190	-0.0381	0.8154	0.0526	0.9343	0.2434	0.1917	0.1908	0.2582	-0.1046	0.8417	0.0580	0.8407	0.1626	0.4648
---	---	1624315_at	0.0577	0.7870	0.1010	0.3331	0.1028	0.7425	-0.0031	0.9956	0.0850	0.6739	0.0881	0.6250	0.0115	0.9943	0.1061	0.8343	0.0946	0.8459
CG9737	CG9737	1624316_at	0.0316	0.9071	0.0130	0.9187	-0.0072	0.9729	0.0129	0.9883	-0.0351	0.9011	-0.0480	0.8393	0.0392	0.9526	-0.0323	0.9157	-0.0715	0.7527
---	---	1624317_at	0.0394	0.7916	-0.0166	0.8796	-0.0127	0.9550	-0.0635	0.8877	-0.0489	0.7924	0.0145	0.9386	-0.1211	0.8326	-0.1445	0.5630	-0.0234	0.9435
128up	upstream of RpIII1	1624318_at	0.4839	0.0582	0.4002	0.3473	0.1214	0.4617	0.0333	0.9603	0.5836	0.0062	0.5503	0.0049	0.2731	0.8049	0.5219	0.2430	0.2488	0.6179
CG12455	CG12455	1624319_at	0.0324	0.8663	0.0499	0.7156	0.0024	0.9945	0.0021	0.9984	0.1393	0.5689	0.1372	0.5345	0.0499	0.9449	0.0648	0.8190	0.0149	0.9629
mei-9	meiotic 9	1624320_at	-0.1157	0.6169	0.1293	0.3784	0.3679	0.0261	0.0241	0.9639	0.0308	0.8688	0.0067	0.9698	-0.1795	0.8157	0.1239	0.7400	0.3034	0.3545
Rac2	Rac2	1624321_at	0.5303	0.2849	-0.5810	0.1655	-1.7191	0.0073	-0.2732	0.8485	1.2073	0.0287	1.4805	0.0073	0.9350	0.6557	0.1346	0.9178	-0.8004	0.3559
Rh50	Rh50	1624322_at	-0.7231	0.4129	-0.5210	0.2856	-1.1353	0.0005	-0.7246	0.1614	-1.5306	0.0011	-0.8060	0.0115	-0.1049	0.9848	-1.3123	0.2707	-1.2074	0.3406
---	---	1624323_at	0.0856	0.5650	0.2489	0.3641	0.4583	0.0344	0.0323	0.9672	0.0255	0.9295	-0.0068	0.9788	-0.0796	0.9365	0.1957	0.5539	0.2753	0.3921
mRpS5	mitochondrial ribo	1624324_a_at	-0.4116	0.1400	-0.0698	0.7863	-0.1762	0.3360	-0.3442	0.3035	-0.5826	0.0099	-0.2384	0.1743	-0.2514	0.7772	-0.3543	0.3458	-0.1029	0.8324
pip	pipe	1624325_at	0.1672	0.3316	0.1743	0.1328	0.1672	0.3579	-0.1068	0.8726	-0.0836	0.7637	0.0233	0.9354	0.1466	0.7755	0.0663	0.8185	-0.0803	0.7537
Rep4	Caspase-activate	1624326_at	0.5507	0.0521	0.3714	0.2074	0.2811	0.1633	-0.0179	0.9860	0.2823	0.2485	0.3001	0.1679	-0.0447	0.9725	0.0237	0.9648	0.0684	0.8774
PpN58A	Protein phosphata	1624327_at	0.0880	0.7058	-0.1012	0.4155	0.0209	0.9263	0.0253	0.9688	0.1365	0.4305	0.1112	0.4866	-0.1105	0.8681	-0.0574	0.8792	0.0531	0.8831
1-Sep	septin	1624328_a_at	-0.0325	0.9426	0.2369	0.3492	0.4474	0.0345	-0.1959	0.7596	-0.1369	0.6597	0.0590	0.8534	-0.1419	0.9362	0.2330	0.7158	0.3748	0.5238
Klp59D	Klp59D	1624329_at	0.0816	0.7054	0.1935	0.1187	0.3280	0.0655	-0.0341	0.9491	-0.0797	0.6317	-0.0456	0.7841	0.0179	0.9875	0.1269	0.6512	0.1090	0.7071
CG5694	CG5694	1624330_s_at	0.4585	0.0325	-0.3482	0.1791	-0.6998	0.0162	-0.0879	0.8190	0.8420	0.0007	0.9300	0.0003	0.2372	0.8073	0.0350	0.9587	-0.2022	0.6423
l(2)k14505	lethal (2) k14505	1624331_at	0.1123	0.5664	0.4613	0.2831	0.2545	0.2703	0.0543	0.9463	-0.2847	0.2090	-0.3390	0.0952	0.2667	0.8202	0.2728	0.5929	0.0061	0.9945

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18217 /// CG4098	CG4098 /// CG18	1624332_s_at	-0.4928	0.0602	-0.5788	0.0238	-0.0852	0.6045	0.0003	0.9997	-0.2450	0.2683	-0.2453	0.2131	-0.3809	0.6903	-0.0419	0.9478	0.3389	0.3981
CG9903	CG9903	1624333_at	-0.1557	0.6408	-0.0269	0.8372	0.0745	0.6442	0.0103	0.9874	-0.0338	0.8703	-0.0441	0.8018	-0.0089	0.9952	0.0536	0.9273	0.0626	0.9026
---	---	1624334_at	0.0506	0.8601	0.0410	0.6840	0.0092	0.9619	0.1136	0.8284	0.0338	0.9034	-0.0799	0.7084	0.0790	0.8841	0.0274	0.9341	-0.0516	0.8444
CG11188	CG11188	1624335_at	0.5195	0.0798	0.2537	0.4458	0.3753	0.0296	0.3738	0.3318	0.7720	0.0049	0.3981	0.0503	0.2133	0.8270	0.4070	0.3000	0.1937	0.6577
---	---	1624336_at	0.1257	0.4676	0.1475	0.4271	0.0197	0.9216	-0.0374	0.9482	0.0324	0.8808	0.0699	0.6798	0.1408	0.8141	0.1123	0.6844	-0.0286	0.9344
RpS14b	Ribosomal protein	1624337_at	0.2154	0.4369	0.1438	0.6240	-0.0508	0.7738	0.0953	0.8103	0.1386	0.3918	0.0433	0.8073	0.1167	0.8982	0.0511	0.9231	-0.0656	0.8880
CG17301	CG17301	1624338_at	0.0622	0.7529	0.0307	0.7803	0.5068	0.0833	0.1508	0.7542	-0.0328	0.9096	-0.1835	0.3344	-0.2147	0.7633	0.0831	0.8353	0.2978	0.3299
BBS1	BBS1	1624339_at	0.0230	0.9264	0.0445	0.7848	0.0440	0.8409	0.0787	0.8698	-0.0065	0.9801	-0.0852	0.6178	-0.0612	0.9555	0.0712	0.8764	0.1323	0.7135
Gs1l	GS1-like protein	1624340_at	0.0402	0.8307	0.3931	0.1274	0.4392	0.0903	-0.0673	0.9300	-0.8639	0.0031	-0.7965	0.0027	-0.2359	0.8157	-0.5056	0.2119	-0.2698	0.5411
Osi10	Osiris	1624341_a_at	0.1199	0.4276	-0.1090	0.5487	-0.1537	0.3546	-0.0475	0.9228	0.0904	0.5731	0.1380	0.3064	0.1396	0.7810	0.1289	0.5750	-0.0107	0.9756
CG10911 /// DyakCG10911	CG10911	1624342_at	-0.3807	0.9297	-1.1866	0.0873	-3.8855	0.0001	-2.2489	0.5757	-1.5457	0.4735	0.7032	0.7550	0.5019	0.9734	-2.4632	0.5228	-2.9651	0.4370
Opbp	Optix-binding prot	1624343_at	0.1066	0.4379	0.2380	0.1205	0.1053	0.6021	0.1790	0.5869	-0.0246	0.9167	-0.2036	0.1741	0.1528	0.7826	0.1579	0.5252	0.0051	0.9901
CG17922	CG17922	1624344_at	-2.8490	0.0008	-0.2780	0.2390	-1.1478	0.0153	-1.0328	0.2506	-2.9002	0.0008	-1.8674	0.0028	-0.0609	0.9816	-0.3009	0.6264	-0.2400	0.7099
Nop56	Nop56	1624345_a_at	0.9201	0.0831	0.8466	0.1100	1.3083	0.0001	0.7057	0.2935	0.9212	0.0273	0.2155	0.5846	0.4338	0.7142	0.8563	0.0933	0.4225	0.4016
---	---	1624346_at	-0.1160	0.5968	0.0025	0.9886	0.0559	0.8606	-0.1152	0.8882	-0.0953	0.7742	0.0199	0.9549	-0.1781	0.7893	0.0467	0.9157	0.2248	0.4428
CG8398	CG8398	1624347_s_at	-0.1007	0.5706	0.0952	0.5618	-0.2992	0.2309	-0.2371	0.5150	-0.0339	0.8974	0.2032	0.2390	0.2187	0.7768	0.1125	0.7873	-0.1061	0.7915
Usp36	Ubiquitin specific	1624348_a_at	0.0801	0.8495	0.6013	0.0833	0.7093	0.0096	0.2488	0.5335	0.0174	0.9552	-0.2314	0.2153	0.0373	0.9884	0.5041	0.3510	0.4668	0.4076
---	---	1624349_at	-0.0193	0.9405	-0.2367	0.1463	-0.2548	0.0901	0.1715	0.7507	0.4610	0.0537	0.2895	0.1656	-0.0416	0.9514	-0.0873	0.7056	-0.0457	0.8657
---	---	1624350_at	0.0723	0.6978	0.0349	0.8165	0.2782	0.1946	0.0494	0.9314	0.0174	0.9447	-0.0320	0.8733	-0.0707	0.9238	-0.0313	0.9363	0.0394	0.9068
---	---	1624351_at	0.1162	0.5146	0.0967	0.6691	0.0914	0.7193	0.0363	0.9755	0.1388	0.6717	0.1025	0.7406	-0.1373	0.8668	0.0163	0.9781	0.1536	0.6638
dpr13	dpr13	1624352_at	0.1340	0.4869	0.0029	0.9962	0.2046	0.4178	0.0671	0.9012	0.0379	0.8704	-0.0291	0.8895	-0.1674	0.8600	-0.0010	0.9996	0.1663	0.6962
CG32654	CG32654	1624353_at	0.4535	0.1952	0.8871	0.1554	1.0489	0.0020	0.1187	0.8899	0.0998	0.7729	-0.0189	0.9587	-0.0792	0.9745	0.5188	0.3986	0.5980	0.3491
not	nonstop	1624354_at	0.4151	0.1992	0.0020	0.9905	-0.1729	0.4090	0.0491	0.9314	0.9142	0.0007	0.8652	0.0006	0.2356	0.8187	0.5068	0.2126	0.2712	0.5403
bl	bancal	1624355_at	-0.4312	0.1024	0.2563	0.3300	0.3252	0.1344	-0.0289	0.9777	-0.6623	0.0188	-0.6334	0.0142	-0.0365	0.9816	0.1315	0.7761	0.1680	0.6875
---	---	1624356_at	0.0919	0.6673	0.0821	0.4088	0.0064	0.9765	0.0603	0.9247	0.0111	0.9702	-0.0492	0.8195	0.0099	0.9898	0.0020	0.9957	-0.0079	0.9764
cona	corona	1624357_at	0.0803	0.5867	0.0236	0.8284	0.1184	0.5560	-0.1988	0.5680	-0.2112	0.2376	-0.0123	0.9571	-0.1168	0.8379	-0.2445	0.2825	-0.1277	0.6141
CG12531	CG12531	1624358_at	0.0154	0.9283	0.2080	0.3072	0.0485	0.7919	-0.1053	0.8217	-0.1707	0.3583	-0.0654	0.7385	0.0982	0.8486	-0.0122	0.9734	-0.1104	0.6328
CG40221	CG40221	1624359_at	0.1023	0.5744	0.0694	0.6025	0.0360	0.8295	-0.0177	0.9767	0.0459	0.8027	0.0636	0.6851	-0.1040	0.8692	-0.0165	0.9682	0.0876	0.7641
croc	FD1 transcription	1624360_at	0.0169	0.9626	-0.0235	0.8520	0.1278	0.4624	0.2089	0.7507	0.3110	0.2784	0.1021	0.7397	0.0678	0.9409	0.1638	0.5847	0.0960	0.7734
l(2)01424	lethal (2) 01424	1624361_s_at	-0.0434	0.9036	-0.1717	0.4108	-0.3712	0.0298	-0.0714	0.9149	0.4196	0.0498	0.4909	0.0167	0.0960	0.9246	0.3335	0.3111	0.2374	0.4978
Nplp4	neuropeptide-like	1624362_at	0.7465	0.1650	0.5251	0.3142	-0.1754	0.6315	-0.4188	0.3596	-0.2378	0.3719	0.1811	0.4638	0.2556	0.9267	-0.5084	0.6032	-0.7640	0.4114
CG15201	CG15201	1624363_at	-0.0085	0.9929	0.9881	0.0470	0.3515	0.4186	-0.4360	0.6015	-1.8258	0.0019	-1.3897	0.0037	0.1746	0.9636	-0.9469	0.3526	-1.1216	0.2960
CG4849	CG4849	1624364_at	0.0526	0.7856	-0.1458	0.6545	0.1273	0.4087	0.1263	0.7289	0.1554	0.3487	0.0290	0.8806	-0.1999	0.8141	0.0545	0.9208	0.2545	0.4851
CG10915	CG10915	1624365_at	-0.7066	0.2235	-0.3335	0.7492	-0.0947	0.5402	-0.0870	0.9200	-0.0750	0.8202	0.0120	0.9711	-0.3284	0.9142	0.2841	0.8444	0.6124	0.5917
CG17098	CG17098	1624366_at	-0.0113	0.9732	0.0792	0.5246	0.2126	0.3723	0.0236	0.9774	-0.0750	0.7574	-0.0986	0.6354	0.0430	0.9640	0.0993	0.7437	0.0563	0.8699
---	---	1624367_at	0.1402	0.4572	0.0515	0.7150	0.3163	0.1196	0.0010	0.9988	0.1402	0.4447	0.1392	0.3971	-0.1168	0.9066	-0.0400	0.9451	0.0768	0.8729
---	---	1624368_at	0.0354	0.8844	0.0278	0.8135	0.0974	0.5739	0.0721	0.8877	0.1030	0.5818	0.0309	0.8797	0.0259	0.9816	0.1410	0.6131	0.1151	0.6892
tomboy40	dtomboy40	1624369_at	0.1673	0.3961	0.2774	0.0843	0.1610	0.5560	-0.1569	0.7857	0.0288	0.9315	0.1857	0.4031	0.2223	0.6955	0.3506	0.1382	0.1283	0.6166
CG34372	CG13552	1624370_at	0.0993	0.6193	0.1454	0.4984	0.1291	0.4684	-0.0521	0.9467	-0.1458	0.5376	-0.0937	0.6857	-0.1809	0.7822	-0.0622	0.8781	0.1187	0.7091
CG7356	CG7356	1624371_a_at	-2.1636	0.0028	-0.9147	0.0938	-2.6497	0.0002	-0.9396	0.1321	-1.2927	0.0050	-0.3530	0.2989	0.7485	0.6660	0.0328	0.9796	-0.7157	0.3171
CG10908	Dm Derlin-1	1624372_at	0.4470	0.0217	0.7476	0.0231	0.8545	0.0014	-0.0585	0.9039	0.0450	0.8190	0.1034	0.4945	-0.2238	0.6960	0.5652	0.0415	0.7890	0.0320
Crg-1	Circadianly Regul	1624373_at	0.2505	0.0892	-0.1515	0.1995	0.0262	0.9059	0.2204	0.4464	0.5806	0.0038	0.3602	0.0189	-0.1139	0.8823	-0.0815	0.8353	0.0323	0.9371
CG12481	CG12481	1624374_at	0.0643	0.7895	0.0027	0.9886	0.2595	0.3161	0.1443	0.8025	-0.0390	0.9026	-0.1833	0.3933	0.1092	0.9011	-0.0081	0.9914	-0.1173	0.7514
insv	insensitive	1624375_at	-0.9519	0.0160	-2.2013	0.0414	-2.4851	0.0019	-0.1111	0.8967	1.5038	0.0008	1.6149	0.0004	0.1797	0.9514	0.0887	0.9501	-0.0909	0.9411
---	---	1624376_at	-0.0704	0.6638	-0.0621	0.5723	-0.0117	0.9652	0.0422	0.9387	0.1135	0.5042	0.0713	0.6704	-0.0439	0.9589	0.0788	0.7925	0.1227	0.6328
CG32594	E protein	1624377_s_at	-0.8288	0.0397	-0.5900	0.1717	-1.8019	0.0005	0.0769	0.9432	0.8914	0.0105	0.8145	0.0095	1.1418	0.3712	0.9956	0.1454	-0.1462	0.8776
east	enhanced adult se	1624378_at	0.3673	0.4509	-0.0699	0.9216	-0.1916	0.6631	0.0197	0.9838	0.7767	0.0048	0.7570	0.0032	0.2773	0.9238	0.3962	0.7198	0.1189	0.9295
---	---	1624379_s_at	-0.0285	0.8979	-0.4679	0.1261	-0.6086	0.0441	-0.0994	0.9436	0.8021	0.0493	0.9015	0.0196	-0.1275	0.8439	-0.0828	0.8094	0.0447	0.9031
CG1359	CG1359	1624380_at	-0.2796	0.4069	-0.1198	0.4140	-0.2899	0.1015	0.1103	0.7690	0.1123	0.5045	0.0020	0.9920	0.1931	0.8395	0.3034	0.4380	0.1103	0.8221
CG17658	CG17658	1624381_at	0.1179	0.4750	-0.1972	0.4121	-0.2612	0.1862	-0.2826	0.2870	0.4008	0.0184	0.6834	0.0011	-0.2134	0.8446	0.0782	0.9085	0.2916	0.5361
---	---	1624382_at	0.1213	0.4498	-0.0841	0.5247	-0.1263	0.4343	0.2596	0.3793	0.2908	0.0828	0.0313	0.8741	0.0126	0.9893	-0.0883	0.6881	-0.1009	0.6328
---	---	1624383_at	0.0191	0.9281	-0.0032	0.9845	0.0058	0.9767	0.0190	0.9805	0.0005	0.9986	-0.0185	0.9376	-0.0057	0.9952	-0.0065	0.9897	-0.0008	0.9987
CG13749	CG13749	1624384_at	0.5364	0.2252	-0.0697	0.7142	0.4755	0.0785	-0.0924	0.9125	-0.0158	0.9696	0.0765	0.7981	-0.4045	0.7062	-0.2264	0.6408	0.1781	0.7264
CG17735	CG17735	1624385_at	-0.1999	0.3924	0.1787	0.3972	0.0920	0.6699	-0.2017	0.6791	-0.1596	0.5074	0.0421	0.8762	-0.0803	0.9590	0.4287	0.3287	0.5090	0.2757
CSN1b	COP9 complex hc	1624386_at	-0.0968	0.5968	0.8116	0.0240	1.1026	0.0007	-0.0703	0.8942	-0.7595	0.0019	-0.6893	0.0017	-0.2842	0.7036	0.2696	0.3761	0.5538	0.1110
---	---	1624387_at	-0.3426	0.0855	-0.7541	0.0596	-1.1590	0.0001	-0.0680	0.9185	0.3332	0.1027	0.4012	0.0352	0.2433	0.7215	-0.1343	0.6723	-0.3776	0.2090
---	---	1624388_at	0.1520	0.4886	0.2095	0.3937	0.4426	0.0124	0.0890	0.8942	-0.1631	0.4856	-0.2521	0.2062	0.0289	0.9816	-0.0936	0.7846	-0.1225	0.6877
CG7526	CG7526	1624389_at	-2.4075	0.0086	-3.5073	0.0314	-3.7863	0.0000	0.0147	0.9874	0.3285	0.1391	0.3138	0.1142	0.2982	0.9416	-0.7539	0.5679	-1.0521	0.4098
thoc7	thoc7	1624390_a_at	0.3879	0.0401	0.2463	0.3716	0.0927	0.5677	-0.0256	0.9649	0.3904	0.0203	0.4160	0.0095	0.1681	0.8446	0.2489	0.4893	0.0809	0.8613
CG32389	CG32389	1624391_at	-0.0327	0.8805	0.0780	0.5173	-0.0486	0.7844	0.0490	0.9413	-0.0898	0.6800	-0.1388	0.4397	0.1232	0.8692	0.1493	0.6425	0.0261	0.9503
Mvl	malvolio	1624392_s_at	-0.2806	0.3494	1.0111	0.0334	-0.2255	0.6954	-0.0111	0.9934	0.0238	0.9517	0.0349	0.9129	1.1843	0.5134	1.3198	0.1320	0.1355	0.9104
w	white	1624393_at	-0.7507	0.0434	-2.1121	0.0047	-1.7920	0.0197	0.8005	0.5731	1.0316	0.1529	0.2311	0.7729	0.1900	0.8714	-0.3618	0.4368	-0.5518	0.2532
CG13332	CG13332	1624394_at	0.0473	0.8295	0.2592	0.1243	0.1065	0.5824	-0.0339	0.9745	-0.0918	0.7698	-0.0579	0.8483	0.1740	0.7947	0.0217	0.9627	-0.1523	0.6149
CG31102	CG31102	1624395_at	0.9963	0.0186	2.0796	0.0135	1.9635	0.0002	-0.2052	0.8436	-1.1117	0.0110	-0.9065	0.0165	-0.1683	0.9114	-0.1908	0.7703	-0.0225	0.9797
CG7927	CG7927	1624396_at	-0.1824	0.2560	0.3295	0.1942	0.1837	0.2168	-0.0168	0.9777	-0.0811	0.6301	-0.0644	0.6850	0.0023	0.9992	0.3312	0.2852	0.3289	0.3141
---	---	1624397_at	0.1420	0.5300	-0.0489	0.6514	0.1971	0.2285	0.1414	0.6993	0.1701	0.3171	0.0286	0.8862	0.0226	0.9831	-0.0987	0.7256	-0.1212	0.6435
CG8630	CG8630	1624398_at	-2.2819	0.0017	-1.8168	0.0302	-1.4304	0.0002	-0.0101	0.9883	0.0387	0.8531	0.0487	0.7869	-0.2755	0.8714	0.4192	0.5526	0.6946	0.3141
---	---	1624399_at	0.1328	0.6585	-0.1493	0.4973	0.3006	0.0464	0.1736	0.7351	0.3222	0.1564	0.1486	0.4985	-0.1502	0.8692	-0.0697	0.8993	0.0805	0.8681
cdc2rk	cdc2-related-kina	1624400_a_at	0.1154	0.5413	0.4589	0.1380	0.5491	0.0099	-0.0154	0.9857	0.0708	0.7580	0.0861	0.6657	-0.0772	0.9555	0.3453	0.3832	0.4224	0.3061

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1624401_at	0.1024	0.6241	0.1595	0.5185	0.2192	0.1729	-0.0614	0.8841	-0.0233	0.9069	0.0381	0.8135	-0.0650	0.9462	0.0717	0.8597	0.1367	0.6674
CG17906	CG17906	1624402_at	0.3305	0.2416	-0.0380	0.7009	0.3422	0.0383	0.2196	0.5068	0.2893	0.0991	0.0697	0.7071	-0.0904	0.9310	-0.1574	0.6768	-0.0670	0.8864
CG13071	CG13071	1624403_at	0.0514	0.8097	0.0711	0.6535	0.1522	0.3225	0.0313	0.9558	-0.0597	0.7396	-0.0910	0.5409	-0.1119	0.8692	-0.1689	0.5501	-0.0569	0.8763
Sk1	Sphingosine kinas	1624404_a_at	0.3600	0.2475	0.8005	0.0121	1.0289	0.0009	-0.0953	0.9066	-0.5322	0.0471	-0.4369	0.0647	-0.3274	0.7215	-0.0510	0.9341	0.2764	0.4851
CG7059	CG7059	1624405_a_at	1.1659	0.0020	2.9109	0.0094	2.0874	0.0036	0.4296	0.4420	-0.4391	0.1478	-0.8686	0.0071	1.2001	0.4377	1.1664	0.1410	-0.0337	0.9798
CG5168	CG5168	1624406_at	0.2258	0.1970	0.0097	0.9702	0.0500	0.8139	0.0576	0.9117	0.2519	0.1253	0.1943	0.1868	0.0201	0.9898	0.0357	0.9476	0.0156	0.9767
lin19	dCullin1	1624407_s_at	0.3070	0.0540	0.0243	0.8763	-0.2224	0.2367	-0.0671	0.8817	0.2955	0.0591	0.3627	0.0165	0.1979	0.7547	0.0058	0.9925	-0.1921	0.4892
NetA	NETRIN	1624408_at	-0.1572	0.6267	0.0241	0.8399	-0.0708	0.6932	0.0721	0.9466	-0.1230	0.7303	-0.1952	0.5069	-0.0273	0.9764	-0.1273	0.5882	-0.1000	0.6835
CG11315	CG11315	1624409_a_at	2.3096	0.0014	1.2835	0.0875	2.9039	0.0001	1.1705	0.1140	0.3070	0.5139	-0.8635	0.0366	-0.6130	0.6955	-0.7642	0.2286	-0.1512	0.8610
CG15658 /// DmirCG15658	CG15658	1624410_at	1.0122	0.0022	-0.2170	0.2452	-0.2059	0.2550	-0.0810	0.8862	0.5476	0.0116	0.6286	0.0038	-0.1767	0.7633	-0.7539	0.0222	-0.5772	0.0559
CG18437	CG18437	1624411_at	-0.0649	0.6916	-0.0063	0.9657	-0.0487	0.8793	0.0588	0.9393	-0.1712	0.4667	-0.2300	0.2562	0.3392	0.6903	0.2014	0.5889	-0.1378	0.7325
ap	Xasta	1624412_at	2.2409	0.0111	1.4817	0.0504	2.4434	0.0009	0.1623	0.9314	0.0345	0.9686	-0.1278	0.8434	-0.6384	0.7822	-0.4631	0.6762	0.1753	0.8984
CG14527	CG14527	1624413_at	0.3665	0.1468	-0.2272	0.6553	-0.0480	0.8429	-0.2882	0.6615	-0.0183	0.9702	0.2699	0.3512	-0.4715	0.6898	-0.6619	0.1599	-0.1904	0.7323
CG10041	CG10041	1624414_at	0.3328	0.1005	0.2319	0.4867	0.2056	0.2499	0.1495	0.7130	0.2348	0.2025	0.0853	0.6558	0.0828	0.9400	0.0366	0.9469	-0.0461	0.9211
CG8121	CG8121	1624415_s_at	-0.5865	0.1256	-0.3754	0.3387	-0.5581	0.0427	-0.1565	0.7021	-0.0124	0.9659	0.1441	0.4115	0.0168	0.9950	0.2486	0.7248	0.2318	0.7434
---	---	1624416_at	-0.0207	0.9230	-0.0769	0.7672	0.4352	0.0178	0.1514	0.7906	0.1070	0.6881	-0.0444	0.8714	-0.0920	0.9088	0.1542	0.6117	0.2462	0.3939
CG10186	CG10186	1624417_a_at	0.3115	0.2314	0.1132	0.4455	0.2857	0.2081	-0.0707	0.8844	-0.1465	0.3891	-0.0757	0.6593	-0.2582	0.7464	-0.2206	0.5259	0.0376	0.9361
bhr	bhringi	1624418_s_at	0.2070	0.4201	0.0394	0.8796	-0.1269	0.4780	0.0567	0.9179	0.4351	0.0183	0.3784	0.0205	0.1943	0.8395	0.2948	0.4585	0.1005	0.8418
Cct5	T-complex Chape	1624419_a_at	-0.2365	0.4639	-0.2555	0.1163	-0.1685	0.6115	0.1488	0.8541	0.6300	0.0375	0.4812	0.0680	0.0559	0.9816	0.6728	0.1948	0.6169	0.2655
CG17118	CG17118	1624420_at	0.0968	0.5334	0.0519	0.7725	-0.0209	0.9321	-0.1013	0.8196	-0.0517	0.8104	0.0497	0.7994	-0.0562	0.9409	-0.0116	0.9783	0.0446	0.8905
---	---	1624421_at	0.1334	0.5920	0.1502	0.3963	0.3512	0.0450	0.0089	0.9922	-0.0516	0.8336	-0.0605	0.7764	-0.0205	0.9822	-0.0078	0.9849	0.0127	0.9690
---	---	1624422_at	0.1073	0.4957	-0.1030	0.3904	-0.0465	0.8244	0.2106	0.7028	0.2647	0.3019	0.0542	0.8545	0.0721	0.8973	0.0727	0.7760	0.0006	0.9990
CG5945 /// DyakCG5945	CG5945	1624423_at	-0.2440	0.3105	-0.7136	0.0695	-0.9058	0.0026	-0.0878	0.9080	0.1466	0.5794	0.2345	0.2879	0.1481	0.9142	-0.2739	0.5907	-0.4219	0.3898
CG13587	CG13587	1624424_at	0.0473	0.8802	0.0131	0.9013	0.2033	0.3473	0.0293	0.9744	-0.2439	0.2702	-0.2732	0.1643	0.0154	0.9914	-0.0469	0.9231	-0.0622	0.8837
CG7384	CG7384	1624425_at	2.2534	0.0020	0.6756	0.2859	2.0565	0.0006	0.7316	0.1160	1.2964	0.0014	0.5649	0.0315	-0.6520	0.7506	-0.3445	0.7381	0.3076	0.7638
---	---	1624426_at	0.2538	0.3680	0.0174	0.8638	0.3227	0.2453	-0.0240	0.9695	-0.1541	0.3491	-0.1301	0.3861	-0.1958	0.8400	0.0387	0.9519	0.2345	0.5828
CG10646	CG10646	1624427_at	0.3428	0.1410	0.3611	0.2761	0.1037	0.6035	-0.0402	0.9477	0.4372	0.0196	0.4774	0.0082	0.2147	0.8235	0.2802	0.4954	0.0655	0.9057
Eig71Ei	Eig71Ei	1624428_at	0.1390	0.4170	0.0075	0.9547	-0.0289	0.9160	0.1290	0.7735	0.1949	0.3044	0.0659	0.7456	0.0452	0.9742	0.0821	0.8673	0.0369	0.9396
CG3967	CG3967	1624429_at	-0.1381	0.5478	0.0227	0.9683	-0.0659	0.7187	0.2989	0.3752	0.3818	0.0505	0.0829	0.6776	0.4414	0.7116	0.5396	0.2619	0.0982	0.8864
Or49a	Odorant receptor	1624430_at	0.1252	0.4387	0.0325	0.7454	0.0685	0.7811	-0.0297	0.9669	-0.0830	0.7010	-0.0533	0.8003	0.1135	0.8541	-0.0445	0.9085	-0.1580	0.5579
CG15908	CG15908	1624431_at	-0.4140	0.0999	-0.4031	0.2861	-0.2661	0.3703	0.2192	0.6115	0.3842	0.0776	0.1649	0.4149	0.0096	0.9952	0.2307	0.5635	0.2212	0.5862
Spz3	Spz3	1624432_at	-1.0560	0.0147	-1.9893	0.0177	-1.5209	0.0010	0.4797	0.5854	0.9592	0.0387	0.4795	0.2319	0.1911	0.8882	-0.1183	0.8690	-0.3094	0.5670
CG10589 /// DyakCG10589	CG10589	1624433_at	0.0140	0.9551	0.0384	0.7946	-0.0133	0.9470	0.1556	0.6856	0.0835	0.6827	-0.0721	0.7020	0.1313	0.7826	0.0435	0.8838	-0.0878	0.7040
tld	tolloid	1624434_at	0.7754	0.3209	0.1029	0.6935	0.3933	0.0812	0.3390	0.1973	0.1665	0.2997	-0.1725	0.2258	-0.0121	0.9979	-0.3955	0.7477	-0.3834	0.7495
Tsp42Ej	sunglasses	1624435_at	-2.2443	0.0009	-1.2389	0.0818	-2.3553	0.0004	-0.7026	0.4337	-1.0338	0.0423	-0.3311	0.4811	0.5386	0.7215	0.2542	0.7257	-0.2844	0.6816
---	---	1624436_at	0.0400	0.8192	-0.0767	0.7154	-0.0249	0.8938	0.2626	0.4331	0.2840	0.1203	0.0215	0.9240	0.2351	0.6749	0.0669	0.8244	-0.1682	0.4795
CG30025 /// CG30031 /// dy	CG30031 /// CG3(	1624437_s_at	0.1812	0.5074	0.0470	0.7556	-0.0565	0.8438	0.0663	0.9346	0.1543	0.5485	0.0880	0.7313	0.0589	0.9589	-0.0188	0.9723	-0.0777	0.8514
CG5017	CG5017	1624438_at	0.1556	0.3791	0.0793	0.7409	0.0906	0.6403	-0.2231	0.5008	0.0013	0.9960	0.2244	0.1520	-0.0636	0.9636	-0.0597	0.9193	0.0040	0.9953
CG4169	ubiquinol-cytochrc	1624439_at	-0.9111	0.0233	-0.1431	0.9034	-0.0044	0.9916	0.0376	0.9538	-1.1941	0.0004	-1.2317	0.0002	-0.0697	0.9898	-0.2193	0.8993	-0.1496	0.9245
---	---	1624440_at	0.2386	0.2463	-0.4328	0.4755	-0.9822	0.0045	-0.5197	0.3692	0.7794	0.0255	1.2991	0.0016	0.0383	0.9908	0.1225	0.9079	0.0842	0.9311
Eip93F	Eip93F	1624441_at	0.8715	0.3866	1.2713	0.0080	0.3001	0.6701	-0.2327	0.8498	0.3289	0.4832	0.5616	0.1596	0.5841	0.8760	0.5696	0.7400	-0.0145	0.9954
CG12586	CG12586	1624442_at	0.1507	0.3630	-0.0786	0.4514	-0.0341	0.8614	0.0406	0.9580	0.2443	0.2293	0.2038	0.2668	0.1334	0.7677	0.0840	0.6969	-0.0495	0.8393
CG6294 /// CG6299	CG6299 /// CG62	1624443_s_at	-0.7558	0.2384	0.7643	0.3704	0.8313	0.0422	0.1032	0.9218	-0.9583	0.0092	-1.0615	0.0034	0.1796	0.9717	0.5308	0.7151	0.3512	0.8256
Mst84Da	Male-specific RNA	1624444_at	0.1905	0.2502	0.0449	0.7929	0.1510	0.4403	0.1592	0.6394	0.0424	0.8437	-0.1168	0.4581	-0.0363	0.9672	-0.0101	0.9815	0.0262	0.9354
---	---	1624445_at	0.1576	0.3181	0.0560	0.5774	0.1654	0.3834	0.0990	0.7929	-0.0176	0.9356	-0.1166	0.4177	-0.0416	0.9589	0.0105	0.9801	0.0522	0.8640
---	---	1624446_at	0.0032	0.9895	0.1259	0.3353	0.1135	0.5714	-0.0467	0.9404	-0.0677	0.7515	-0.0210	0.9240	-0.0879	0.8956	-0.0029	0.9961	0.0851	0.7728
dco	double-time	1624447_s_at	0.3562	0.1559	-0.0442	0.8209	-0.0732	0.7521	-0.0963	0.8738	0.5774	0.0140	0.6737	0.0042	-0.0422	0.9721	0.1643	0.6156	0.2064	0.5176
CG6356	CG6356	1624448_at	-1.2502	0.0619	0.3968	0.1829	-0.1015	0.6095	-0.4340	0.6327	-1.9461	0.0018	-1.5121	0.0032	0.2455	0.8541	-0.2163	0.7361	-0.4617	0.4090
---	---	1624449_at	-0.0759	0.8023	0.2938	0.3869	0.2576	0.1286	-0.1247	0.7278	-0.5531	0.0050	-0.4284	0.0096	0.2446	0.7204	0.1914	0.5141	-0.0532	0.8918
Orct	Organic cation tra	1624450_at	1.9574	0.0014	0.5580	0.0248	1.2293	0.0003	0.5253	0.0772	1.2288	0.0003	0.7035	0.0014	-0.0354	0.9852	-0.0808	0.9023	-0.0454	0.9405
CG2879	CG2879	1624451_at	0.1314	0.5097	-0.0377	0.8459	0.2074	0.2197	0.0799	0.8809	0.0549	0.8064	-0.0250	0.9095	-0.0490	0.9421	-0.0438	0.8849	0.0052	0.9874
ple	tyrosine hydroxyla	1624452_a_at	-0.0745	0.6786	-0.5575	0.0110	-0.1164	0.5676	0.4879	0.1054	0.6009	0.0056	0.1129	0.5132	-0.0906	0.9095	0.0772	0.8394	0.1677	0.5789
---	---	1624453_at	0.1543	0.4954	0.0796	0.6736	-0.0477	0.8142	-0.0771	0.9200	-0.0303	0.9266	0.0469	0.8637	0.0963	0.8455	0.0194	0.9525	-0.0769	0.7431
CG10361	CG10361	1624454_at	1.3927	0.0009	0.5720	0.2750	1.6495	0.0012	0.2329	0.6195	-0.1360	0.5928	-0.3689	0.0785	-0.8181	0.5989	-0.9358	0.1579	-0.1178	0.9008
---	---	1624455_at	0.2585	0.2210	0.1068	0.3642	0.1315	0.4578	0.1242	0.7327	0.0369	0.8627	-0.0873	0.5902	0.0437	0.9657	0.0288	0.9486	-0.0149	0.9719
CG17154	CG17154	1624456_at	0.0224	0.9302	-0.0148	0.9453	-0.1217	0.4163	0.0124	0.9884	0.0298	0.9167	0.0174	0.9461	0.1159	0.8446	0.0747	0.8123	-0.0412	0.9025
CG5906	CG5906	1624457_at	0.0139	0.9581	0.0293	0.7719	-0.0014	0.9950	0.0513	0.9260	0.0452	0.8309	-0.0061	0.9766	0.0962	0.8653	0.0571	0.8546	-0.0391	0.9007
CG6083	CG6083	1624458_at	-0.6685	0.1310	-0.0133	0.9044	-0.3599	0.0297	-0.2030	0.8403	-0.9205	0.0226	-0.7175	0.0393	0.0051	0.9967	-0.2549	0.4094	-0.2601	0.4114
Sox21a	Sox21a	1624459_at	0.1692	0.3497	0.4324	0.0321	0.3454	0.1206	-0.0739	0.8830	0.0840	0.6655	0.1579	0.3101	0.1259	0.8427	0.1347	0.6326	0.0088	0.9842
CG31624 /// CG31988 /// D	CG31988 /// CG3	1624460_s_at	-0.0599	0.8538	0.0167	0.9178	0.1991	0.2021	0.2572	0.6081	-0.0502	0.8811	-0.3074	0.1706	-0.0119	0.9898	-0.0313	0.9175	-0.0194	0.9433
---	---	1624461_at	0.0090	0.9756	0.0666	0.8236	0.0578	0.8232	-0.0690	0.8844	-0.1062	0.5408	-0.0373	0.8418	0.0021	0.9992	0.0766	0.8550	0.0745	0.8501
CG32331	CG32331	1624462_at	0.1305	0.4868	-0.0822	0.6453	-0.0943	0.6739	-0.1133	0.8327	0.1336	0.5418	0.2469	0.1747	-0.1357	0.8504	-0.0336	0.9449	0.1021	0.7703
pr-set7	pr-set7	1624463_s_at	-0.1477	0.7245	-0.0225	0.9536	-0.1727	0.3274	0.1525	0.8218	0.3243	0.2181	0.1718	0.4986	0.1669	0.8930	0.2307	0.6460	0.0638	0.9199
---	---	1624464_s_at	-0.9792	0.0504	-1.2243	0.0248	-1.4483	0.0003	-0.0412	0.9665	-0.2875	0.2649	-0.2463	0.2895	0.1822	0.9168	-0.5012	0.4031	-0.6834	0.2757
Ugt	UDP-glucose:glyc	1624465_at	0.4704	0.1675	0.4560	0.3228	0.5492	0.0075	-0.0112	0.9903	0.7339	0.0043	0.7451	0.0024	-0.0523	0.9852	0.8150	0.1925	0.8673	0.2016
CG15117	CG15117	1624466_a_at	1.2619	0.0112	0.5362	0.1772	1.5004	0.0001	0.5303	0.1138	0.6095	0.0089	0.0792	0.7092	-0.4919	0.7106	-0.1432	0.8500	0.3487	0.5511
CG8046	CG8046	1624467_a_at	-2.0246	0.0079	-3.3232	0.0034	-3.7355	0.0000	0.0555	0.9712	1.0393	0.0157	0.9839	0.0124	-0.1638	0.9445	-0.6451	0.3549	-0.4812	0.5176
CG10026	CG10026	1624468_a_at	-0.2432	0.6539	0.3272	0.6888	-0.0679	0.8226	0.2351	0.6558	0.0538	0.8727	-0.1813	0.4484	0.7170	0.7485	0.6606	0.5020	-0.0563	0.9695
DmsR-1	Dromyosuppressi	1624469_at	-0.1897	0.2670	0.0583	0.7071	-0.1540	0.4617	-0.2221	0.5046	-0.1190	0.5276	0.1031	0.5532	0.1747	0.8097	0.1414	0.6721	-0.0333	0.9371

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31670	CG31670	1624470_at	-0.1059	0.5880	0.0077	0.9645	0.1382	0.4458	0.1382	0.7647	0.0614	0.7997	-0.0768	0.7136	-0.0998	0.8825	0.0195	0.9634	0.1193	0.6728
Act42A	42A actin.	1624471_s_at	-0.2981	0.2674	-0.3895	0.0629	-0.3550	0.1815	0.2057	0.7647	0.9334	0.0073	0.7277	0.0134	0.1716	0.8086	0.8328	0.0276	0.6612	0.0607
CG10486	CG10486	1624472_at	0.1802	0.6386	-0.1802	0.3798	-0.1188	0.6792	0.2552	0.7501	0.4253	0.2205	0.1701	0.6323	0.0933	0.9296	0.0200	0.9723	-0.0733	0.8729
DNAprim	DNA primase	1624473_at	0.1020	0.7917	0.0706	0.7090	0.3509	0.2376	-0.2542	0.5067	-0.0339	0.9037	0.2203	0.2252	-0.3538	0.8202	0.2565	0.7319	0.6103	0.3569
CG4287	CG4287	1624474_at	-0.0712	0.7604	-0.4472	0.0615	-0.4205	0.0356	-0.1374	0.8025	0.0135	0.9686	0.1510	0.4704	-0.1557	0.8472	-0.2975	0.3669	-0.1418	0.7067
---	---	1624475_at	0.4891	0.0383	0.6174	0.1463	0.8763	0.0039	0.2846	0.4728	0.5278	0.0216	0.2433	0.2013	0.0312	0.9862	0.5421	0.1838	0.5109	0.2398
Tom	Barbu	1624476_at	-0.3243	0.8005	-0.2691	0.1071	-0.3094	0.1043	-0.0342	0.9943	-0.6410	0.5530	-0.6068	0.5353	-0.0733	0.9774	-0.4610	0.4489	-0.3878	0.5460
---	---	1624477_at	0.4898	0.0876	0.1009	0.6285	0.2966	0.1590	0.0304	0.9573	0.0673	0.7006	0.0369	0.8308	-0.1535	0.8692	-0.1457	0.7400	0.0078	0.9902
Roc1a	Roc1a	1624478_at	-0.1349	0.4552	-0.0616	0.8230	-0.0281	0.9161	0.0832	0.8967	-0.0129	0.9687	-0.0961	0.6605	0.0338	0.9831	0.0965	0.8478	0.0628	0.9023
CG6800	CDK7-like	1624479_at	0.1925	0.2755	0.0023	0.9905	0.2639	0.2213	0.1462	0.8195	0.1775	0.5026	0.0312	0.9193	-0.0658	0.9514	-0.1257	0.7378	-0.0599	0.8918
mRpL43	mitochondrial ribo	1624480_at	-0.0051	0.9888	0.6041	0.0429	0.6751	0.0045	-0.0053	0.9955	-0.3451	0.0682	-0.3398	0.0476	-0.1001	0.9457	0.2695	0.5618	0.3696	0.4114
CG31534	CG31534	1624481_a_at	1.0152	0.0144	1.2307	0.0218	0.7603	0.0011	0.0290	0.9672	0.2715	0.1410	0.2424	0.1417	0.4315	0.6832	0.5156	0.2212	0.0841	0.8885
smid	smallminded	1624482_at	-0.0199	0.9160	0.3683	0.2676	0.3492	0.0375	0.0034	0.9956	-0.1923	0.2591	-0.1957	0.1965	-0.1082	0.9168	0.1312	0.7604	0.2394	0.5257
CG13472 /// DmirCG13472	CG13472	1624483_at	-0.1132	0.7181	0.0775	0.7927	0.5578	0.0368	0.3235	0.4752	0.1143	0.6807	-0.2093	0.3494	-0.0253	0.9914	0.5277	0.3066	0.5529	0.3082
BEAF-32	boundary element	1624484_a_at	0.4844	0.0107	0.6264	0.0885	0.3833	0.0350	-0.1472	0.6166	0.0922	0.5603	0.2393	0.0707	0.0799	0.9374	0.0781	0.8607	-0.0018	0.9979
CG6867	CG6867	1624485_at	1.0609	0.5870	-0.4146	0.1116	-1.1118	0.0054	-0.4583	0.5724	-0.8128	0.0547	-0.3545	0.3517	-0.1226	0.9914	-2.7742	0.2829	-2.6515	0.3308
CG12835	CG12835	1624486_at	0.2327	0.2283	0.2518	0.2003	0.3686	0.0671	0.1101	0.8189	0.0428	0.8624	-0.0673	0.7396	0.0915	0.9186	0.0996	0.7978	0.0081	0.9868
CG11475	CG11475	1624487_at	-0.3870	0.0819	-0.3636	0.0385	-0.0679	0.7445	0.2394	0.3921	-0.1004	0.5543	-0.3398	0.0241	-0.0663	0.9589	-0.2852	0.4380	-0.2189	0.5767
CG17734	CG17734	1624488_a_at	0.3205	0.0769	0.3690	0.0291	0.1105	0.4829	-0.0504	0.9182	0.2680	0.0770	0.3184	0.0261	0.0955	0.8814	0.3524	0.1418	0.2569	0.3042
CG11596	CG11596	1624489_a_at	0.2683	0.1604	0.2271	0.1312	0.4772	0.0110	0.0734	0.8770	0.0980	0.5795	0.0246	0.9001	0.0323	0.9781	0.1405	0.6441	0.1083	0.7375
hb	hunchback	1624490_s_at	-1.4424	0.0470	-1.3169	0.1416	-2.1788	0.0013	-0.6541	0.0433	-0.5263	0.0110	0.1277	0.4592	0.0093	0.9989	-0.5254	0.6382	-0.5347	0.6309
pdm2	miti-mere	1624491_a_at	-0.4514	0.3188	-0.0530	0.6453	-0.4088	0.1057	-0.0103	0.9956	-0.4730	0.2670	-0.4627	0.2233	0.0654	0.9611	-0.1824	0.6587	-0.2479	0.5320
CG9597	CG9597	1624492_at	0.1622	0.5751	0.4956	0.0445	0.3622	0.0521	-0.3511	0.3276	-0.1774	0.4059	0.1736	0.3644	-0.1456	0.8814	0.2817	0.4503	0.4273	0.2674
---	---	1624493_at	0.1287	0.4856	0.2842	0.3083	0.0058	0.9820	-0.1910	0.7327	-0.0150	0.9693	0.1760	0.4554	0.0566	0.9242	0.1341	0.5140	0.0775	0.7341
---	---	1624494_at	0.1782	0.3091	0.0570	0.5884	0.3046	0.2460	0.0566	0.9314	0.0908	0.6834	0.0342	0.8822	0.0164	0.9862	0.0015	0.9985	-0.0149	0.9621
---	---	1624495_at	0.0505	0.8493	-0.0562	0.5977	-0.0439	0.8244	0.0226	0.9728	0.0441	0.8346	0.0214	0.9151	-0.0410	0.9656	-0.1257	0.6497	-0.0847	0.7777
xl6	xl6	1624496_at	-0.2012	0.5749	-0.1122	0.8217	-0.3089	0.1336	-0.1241	0.8611	0.4296	0.0922	0.5537	0.0232	0.0757	0.9762	0.4812	0.4300	0.4055	0.5274
---	---	1624497_at	0.5452	0.2001	-0.0976	0.6092	-0.4184	0.2505	-0.1569	0.9022	-0.3566	0.4024	-0.1997	0.6384	-0.0896	0.9246	-1.1187	0.0158	-1.0291	0.0320
Cpr49Aa	CG30045	1624498_at	0.1157	0.5609	0.0399	0.8266	0.2545	0.2026	-0.0123	0.9889	-0.0191	0.9505	-0.0068	0.9788	-0.0106	0.9913	0.0457	0.8749	0.0563	0.8247
CG9911	CG9911	1624499_s_at	0.5640	0.0145	0.3598	0.1648	0.3773	0.0333	-0.0489	0.9314	0.6808	0.0025	0.7296	0.0011	0.0362	0.9816	0.5716	0.0968	0.5354	0.1393
---	---	1624500_at	0.0344	0.8885	-0.0140	0.9393	-0.1152	0.6751	0.0590	0.9436	0.0751	0.7961	0.0161	0.9573	0.1604	0.8815	0.2001	0.6586	0.0397	0.9451
CG12496	CG12496	1624501_at	0.2453	0.0996	0.0369	0.7345	-0.0024	0.9921	-0.0304	0.9641	-0.0367	0.8778	-0.0063	0.9776	-0.0093	0.9914	-0.0063	0.9870	0.0029	0.9929
CG1104	CG1104	1624502_a_at	0.4160	0.1180	0.8807	0.0745	0.8466	0.0012	0.1065	0.8222	0.0952	0.6465	-0.0113	0.9611	0.0749	0.9653	0.5466	0.2217	0.4717	0.3204
CG4786	CG4786	1624503_at	-1.8055	0.0006	-1.7580	0.0160	-2.5706	0.0004	0.0013	0.9988	0.6208	0.0169	0.6195	0.0105	0.7801	0.6824	0.7493	0.3269	-0.0308	0.9816
Mtch	Mitochondrial carr	1624504_s_at	0.2450	0.2025	0.6413	0.0370	0.6126	0.0038	-0.0148	0.9838	-0.2509	0.1293	-0.2360	0.1107	-0.1585	0.8235	-0.0156	0.9760	0.1428	0.6565
CG6113	CG6113	1624505_at	2.5816	0.0024	1.2698	0.0260	2.0759	0.0010	0.9940	0.2753	0.7134	0.1817	-0.2806	0.6042	0.2522	0.8444	-0.6346	0.2145	-0.8868	0.1261
Jon25Biii	Jonah 25B	1624506_at	0.2217	0.8067	-0.1370	0.4333	-0.1458	0.5785	0.0554	0.9470	0.0297	0.9289	-0.0258	0.9287	-0.0435	0.9928	-0.5890	0.6191	-0.5455	0.6468
CG31268	CG31268	1624507_at	0.1216	0.5444	0.1317	0.4154	0.1116	0.5305	0.0351	0.9603	0.0320	0.9022	-0.0031	0.9898	0.0492	0.9545	0.0668	0.8460	0.0176	0.9622
CG13526	CG13526	1624508_at	-0.0698	0.7021	-0.0170	0.8668	-0.1078	0.5445	0.0335	0.9518	0.0773	0.6568	0.0439	0.8002	-0.0405	0.9552	0.0063	0.9886	0.0468	0.8669
LvpL	hdl cuticle gene cl	1624509_at	0.0827	0.8413	-0.0342	0.8727	0.0243	0.9062	0.1823	0.7298	0.1202	0.6513	-0.0621	0.8165	0.0199	0.9914	-0.0601	0.9289	-0.0799	0.8918
sprt	sprite	1624510_at	-0.8737	0.0113	0.6003	0.4268	-0.1240	0.7034	-0.2934	0.8057	-1.2395	0.0173	-0.9461	0.0332	0.4888	0.7644	0.0945	0.9322	-0.3942	0.5965
---	---	1624511_s_at	0.1546	0.5568	-0.0159	0.9064	0.0316	0.8791	0.0402	0.9488	0.0775	0.6999	0.0373	0.8540	0.0138	0.9914	-0.0312	0.9476	-0.0450	0.9109
CG34386	CG14496	1624512_at	0.1847	0.4306	0.1700	0.5283	0.5181	0.0685	-0.0219	0.9857	-0.0655	0.8586	-0.0436	0.8975	-0.0664	0.9309	0.0668	0.8432	0.1333	0.6195
CG33228 /// CG4806	CG4806 /// CG33	1624513_at	0.1456	0.6703	-0.1597	0.7803	-0.0962	0.7005	0.0750	0.9141	0.8552	0.0029	0.7801	0.0026	-0.0224	0.9939	0.5481	0.3903	0.5705	0.3878
Vha16-3	Vha16-3	1624514_at	0.0481	0.8503	-0.0435	0.6652	-0.0184	0.9422	-0.1500	0.7688	-0.0742	0.7768	0.0758	0.7460	0.0060	0.9964	-0.0638	0.8897	-0.0697	0.8676
CG6461 /// DmirCG6461	CG6461	1624515_at	-1.4944	0.0006	-1.7623	0.0756	-1.9544	0.0008	-0.2456	0.4126	-0.1481	0.3836	0.0975	0.5501	-0.0906	0.9816	-0.4433	0.6458	-0.3527	0.7284
---	---	1624516_at	0.3500	0.2013	-0.0590	0.7659	0.1489	0.4946	0.1366	0.8891	0.1292	0.7388	-0.0075	0.9856	-0.0292	0.9848	-0.2874	0.3992	-0.2582	0.4683
Ect3	beta-galactosidas	1624517_at	2.9542	0.0072	2.4844	0.0452	5.4050	0.0000	1.0015	0.1544	0.6357	0.1336	-0.3658	0.3487	-2.0346	0.3771	0.0880	0.9663	2.1225	0.1267
CG13369	CG13369	1624518_at	-0.2581	0.2944	0.4000	0.5178	0.7003	0.0057	0.3403	0.2051	-0.4824	0.0095	-0.8228	0.0006	-0.0125	0.9964	0.1064	0.9113	0.1190	0.8896
Gbeta76C	guanine nucleotid	1624519_at	0.3455	0.0556	0.0532	0.7042	0.0735	0.7451	-0.1011	0.8921	0.1834	0.4832	0.2846	0.2019	0.0384	0.9588	0.0238	0.9432	-0.0146	0.9618
ftz-f1	Ftz interacting pro	1624520_a_at	0.7407	0.2061	0.2949	0.7004	-0.4498	0.0248	-0.5340	0.3294	0.5410	0.0884	1.0750	0.0032	0.0143	0.9970	0.1960	0.8860	0.1817	0.8890
CG17896	CG17896	1624521_a_at	0.1889	0.3005	0.2056	0.1194	0.5166	0.0203	-0.2618	0.4590	-0.6406	0.0058	-0.3788	0.0341	-0.5363	0.2892	-0.4975	0.0780	0.0388	0.9169
CG5302	CG5302	1624522_at	0.1522	0.6054	-0.0441	0.6935	0.1174	0.4350	0.1503	0.7647	-0.0049	0.9881	-0.1552	0.4421	0.1518	0.8049	0.0018	0.9988	-0.1500	0.5830
CG32683	CG32683	1624523_at	0.0770	0.6258	-0.1479	0.4099	0.0748	0.7093	0.0630	0.8908	0.1312	0.4089	0.0682	0.6711	-0.1327	0.7890	-0.1069	0.6391	0.0257	0.9304
---	---	1624524_at	0.0337	0.9042	0.0517	0.5762	0.0869	0.6140	-0.0521	0.9351	-0.0176	0.9507	0.0345	0.8776	0.0965	0.8326	0.0625	0.7932	-0.0340	0.8959
CG12998	CG12998	1624525_at	0.2736	0.7582	0.8728	0.4253	0.2656	0.5918	-0.0238	0.9840	-0.9545	0.0051	-0.9307	0.0034	0.4688	0.9022	-0.1996	0.9277	-0.6683	0.6578
---	---	1624526_s_at	0.1950	0.3915	-0.0773	0.5244	0.0601	0.8429	-0.0923	0.8337	0.0031	0.9899	0.0954	0.5629	-0.1636	0.8320	-0.0812	0.8546	0.0824	0.8408
Hey	Hairy/E(spl)-relate	1624527_at	-0.0517	0.8473	-0.0850	0.4907	-0.1022	0.7336	0.1536	0.7630	0.1376	0.5590	-0.0160	0.9542	0.0327	0.9816	-0.1242	0.7489	-0.1569	0.6602
Gyc-89Db	Guanylyl cyclase	1624528_at	0.1554	0.3607	0.0162	0.8845	0.0947	0.5999	0.0305	0.9620	-0.0339	0.8809	-0.0644	0.7208	-0.0446	0.9589	-0.0538	0.8764	-0.0093	0.9821
RpS17	ribosomal protein	1624529_at	0.3015	0.1894	1.2555	0.0066	1.3180	0.0006	0.1794	0.7826	-0.3596	0.1786	-0.5390	0.0323	0.1553	0.8222	0.1186	0.7198	-0.0367	0.9265
RhoBTB	RhoBTB	1624530_at	0.4253	0.0522	0.5060	0.0367	0.2759	0.1021	-0.4271	0.2435	-0.5071	0.0269	-0.0800	0.7298	-0.1907	0.7673	-0.3816	0.1508	-0.1909	0.4998
---	---	1624531_s_at	-0.1493	0.5934	-0.8983	0.0674	-0.5865	0.0180	0.4021	0.4337	0.9526	0.0055	0.5504	0.0360	0.0149	0.9939	-0.0462	0.9462	-0.0611	0.9168
CG7051 /// DmirCG7051	CG7051	1624532_a_at	0.1651	0.2624	0.1581	0.3054	0.0997	0.5252	0.0113	0.9894	0.1149	0.5920	0.1035	0.6004	-0.0340	0.9717	-0.0037	0.9941	0.0303	0.9282
trx	trithorax	1624533_s_at	0.3393	0.1545	-0.5455	0.1468	-0.7527	0.0173	0.1398	0.8663	0.8476	0.0110	0.7078	0.0149	0.3285	0.7628	-0.0617	0.9325	-0.3902	0.3986
ng1	nested	1624534_at	0.0613	0.6784	0.0007	0.9997	0.0076	0.9733	0.0115	0.9866	0.0836	0.6610	0.0721	0.6828	-0.0170	0.9835	-0.0491	0.8500	-0.0321	0.9026
CG33229	CG33229	1624535_at	-0.0814	0.7799	0.0484	0.7954	-0.3624	0.0671	-0.1745	0.7530	-0.1047	0.7053	0.0697	0.7959	0.2091	0.7726	-0.0211	0.9672	-0.2302	0.4765
CG30130	CG30130	1624536_at	0.1118	0.5885	-0.0994	0.5371	0.1585	0.3817	0.0906	0.8738	0.0095	0.9755	-0.0811	0.6987	-0.1337	0.8292	-0.0343	0.9341	0.0994	0.7409
CG9866	CG9866	1624537_s_at	-0.1439	0.6636	-0.2617	0.5737	-0.1946	0.1791	0.0406	0.9538	0.2268	0.2391	0.1862	0.2846	0.0554	0.9814	0.1096	0.8792	0.0542	0.9387
---	---	1624538_at	-0.0368	0.8102	0.1248	0.5251	0.3233	0.0868	-0.0110	0.9904	-0.0089	0.9776	0.0021	0.9931	-0.0782	0.9101	0.1569	0.5415	0.2351	0.3545

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG5871	CG5871	1624539_at	0.5084	0.0272	0.2074	0.1673	0.4050	0.0800	0.1507	0.8101	0.7908	0.0073	0.6401	0.0115	0.0173	0.9901	0.5441	0.0864	0.5268	0.1184
CG14642	CG14642	1624540_at	2.3354	0.0020	1.2233	0.0189	2.1377	0.0002	0.5215	0.1372	0.4595	0.0370	-0.0620	0.7944	-0.3819	0.7644	-0.3692	0.5067	0.0127	0.9880
RhoGAP19D	RhoGAP19D	1624541_at	0.6448	0.0920	0.0127	0.9864	0.3545	0.1812	-0.1669	0.7451	0.3025	0.1768	0.4694	0.0276	-0.3718	0.8270	-0.1503	0.8830	0.2215	0.7954
CG31757	CG31757	1624542_at	-1.1528	0.0116	-1.0646	0.0628	-1.6383	0.0033	-0.0687	0.9308	0.3571	0.1211	0.4258	0.0452	0.0964	0.9457	0.0064	0.9942	-0.0900	0.8761
---	---	1624543_s_at	-0.5241	0.3206	-0.5168	0.2768	-1.2109	0.0050	-0.4151	0.5926	0.4423	0.2626	0.8574	0.0230	0.2982	0.8589	0.5839	0.3802	0.2857	0.7075
---	---	1624544_at	-0.3637	0.0979	-0.9185	0.1580	-0.6549	0.0332	-0.0910	0.9182	0.7063	0.0177	0.7973	0.0062	0.0017	0.9998	0.2559	0.7299	0.2543	0.7271
CG32504	CG32504	1624545_at	0.0884	0.6202	-0.0909	0.4257	-0.2081	0.2224	-0.0865	0.8512	0.0646	0.7451	0.1511	0.3241	0.0511	0.9405	-0.0412	0.8962	-0.0923	0.6939
---	---	1624546_at	-0.0478	0.7846	-0.2041	0.2395	0.0559	0.7652	0.1551	0.7351	0.2215	0.2801	0.0664	0.7675	-0.0368	0.9701	0.0430	0.9111	0.0799	0.7918
B52	Serine/arginine ric	1624547_s_at	0.2460	0.3199	1.8139	0.0048	1.4417	0.0004	0.0141	0.9894	-0.4424	0.0735	-0.4565	0.0432	0.4574	0.5352	1.1233	0.0206	0.6659	0.0905
ppk19	pickpocket 19	1624548_at	0.1579	0.3642	-0.0060	0.9616	0.0840	0.6948	0.0462	0.9562	0.0343	0.9142	-0.0119	0.9674	0.1679	0.7387	0.0333	0.9204	-0.1346	0.5515
CG3523	Fatty acid synthet	1624549_at	1.0660	0.0059	-0.0447	0.8741	0.5186	0.0247	0.3474	0.2085	0.9068	0.0007	0.5594	0.0033	-0.1403	0.9135	-0.1389	0.8123	0.0014	0.9989
Trf	TBP-related factor	1624550_at	0.1910	0.2778	0.2497	0.4867	0.7250	0.0185	0.1977	0.6822	-0.2052	0.3705	-0.4029	0.0497	-0.1829	0.8541	0.0485	0.9411	0.2314	0.5995
CG31861	CG31861	1624551_at	0.3657	0.1450	0.2845	0.0618	0.4261	0.0692	0.1244	0.8815	0.0758	0.8352	-0.0486	0.8856	0.1796	0.7644	0.0316	0.9390	-0.1480	0.5853
CG3740	CG3740	1624552_at	-0.0034	0.9868	-0.0385	0.8147	0.0973	0.5063	0.0312	0.9653	0.0957	0.6573	0.0645	0.7571	-0.0812	0.8882	0.1032	0.6660	0.1844	0.4037
CG14098	CG14098	1624553_a_at	-0.1493	0.4148	-0.4280	0.0523	-0.0740	0.7013	0.1736	0.5680	0.3607	0.0284	0.1871	0.1787	-0.1210	0.8728	0.0107	0.9854	0.1318	0.6890
CG3703	CG3703	1624554_at	0.0934	0.5379	-0.3238	0.1348	-0.4770	0.0143	0.1615	0.8140	0.4989	0.0684	0.3374	0.1624	0.1010	0.8882	-0.1195	0.6952	-0.2205	0.4273
Ets97D	tiny	1624555_at	0.0488	0.8548	0.2138	0.5352	0.2226	0.1953	-0.0591	0.9329	0.0029	0.9920	0.0620	0.7894	-0.0304	0.9848	0.1825	0.6483	0.2129	0.5872
bol	boule	1624556_s_at	-1.5612	0.0008	-1.1006	0.0291	-1.4735	0.0008	-0.1140	0.8512	-0.1701	0.4606	-0.0561	0.8234	0.0794	0.9742	0.0727	0.9407	-0.0066	0.9952
Fen1	Flap endonucleas	1624557_at	0.4828	0.2314	-0.1356	0.6038	0.0238	0.9496	-0.0275	0.9803	0.9389	0.0046	0.9664	0.0024	-0.3316	0.8473	0.0838	0.9425	0.4154	0.5870
dro3	drosomycin-E	1624558_at	0.4093	0.0313	0.1782	0.3776	0.1773	0.2375	-0.0738	0.8917	-0.0356	0.8842	0.0382	0.8571	-0.0063	0.9942	-0.1061	0.6166	-0.0998	0.6394
Mlf	Myelodysplasia/m	1624559_a_at	-0.1331	0.5340	0.2336	0.2381	-0.0702	0.8128	0.0988	0.8678	-0.1320	0.5602	-0.2307	0.2198	0.3656	0.6955	0.3434	0.3724	-0.0222	0.9704
ergic53	ergic-53	1624560_at	0.7514	0.0107	0.6589	0.0386	0.6207	0.0040	-0.0277	0.9659	0.4764	0.0131	0.5041	0.0061	0.0394	0.9734	0.3663	0.1948	0.3269	0.2784
---	---	1624561_at	0.2410	0.1558	0.1528	0.2882	0.1245	0.5138	-0.0714	0.8792	-0.0408	0.8442	0.0306	0.8713	0.0049	0.9966	-0.0787	0.8444	-0.0836	0.8209
Mst33A	Mst33A	1624562_s_at	0.1374	0.3825	-0.0131	0.8988	-0.1635	0.4542	-0.3851	0.3294	0.0102	0.9773	0.3953	0.0571	0.0008	0.9998	-0.0443	0.9157	-0.0451	0.9033
CG5225	CG5225	1624563_at	0.2260	0.1695	-0.1454	0.1613	0.0657	0.7513	-0.0214	0.9838	0.0846	0.7780	0.1061	0.6832	-0.1055	0.8898	-0.2361	0.4038	-0.1305	0.6770
RpS14a	Ribosomal protein	1624564_s_at	0.1338	0.5511	0.3844	0.1326	0.2420	0.1306	-0.0247	0.9749	-0.0687	0.7658	-0.0440	0.8427	0.0244	0.9816	0.0345	0.9239	0.0101	0.9780
CG16711	CG16711	1624565_a_at	0.2656	0.7416	-0.3364	0.6676	-0.5784	0.1767	-0.0120	0.9937	1.2786	0.0025	1.2906	0.0014	0.3293	0.9296	0.5744	0.6669	0.2451	0.8818
---	---	1624566_at	0.2122	0.2885	0.0114	0.9165	0.0407	0.7978	0.0920	0.8058	0.2197	0.1391	0.1276	0.3512	-0.0560	0.9457	-0.0185	0.9640	0.0374	0.9123
Rab35	Rab35	1624567_at	-0.8456	0.0287	-0.5171	0.1584	-1.1869	0.0007	-0.4281	0.3425	-0.3100	0.2316	0.1180	0.6610	0.1003	0.9390	-0.2240	0.6027	-0.3243	0.4327
CG7985	CG7985	1624568_at	-1.8119	0.0049	-2.2543	0.0070	-1.8732	0.0001	0.3588	0.6701	0.6260	0.1162	0.2672	0.4839	-0.0240	0.9906	0.2667	0.5527	0.2907	0.5171
CG31087	CG31087	1624569_at	0.1415	0.7632	0.0133	0.8996	-0.4823	0.0723	-0.6808	0.1462	-0.2504	0.3882	0.4304	0.0893	0.0338	0.9885	-0.1870	0.7584	-0.2208	0.6974
TfIIA-L	Transcription fact	1624570_a_at	-0.4262	0.0232	0.4972	0.0420	0.5451	0.0089	-0.1758	0.6041	-0.7569	0.0017	-0.5811	0.0032	-0.1407	0.8236	0.0995	0.7490	0.2403	0.3752
CG32500 /// CG32857 /// C	CG32500 /// CG3	1624571_s_at	-0.4705	0.0815	0.0432	0.8522	0.4731	0.0110	0.1378	0.7596	-0.8593	0.0017	-0.9971	0.0006	-0.3226	0.7070	-0.3371	0.3340	-0.0145	0.9809
MED21	Mediator complex	1624572_a_at	-0.1716	0.4988	-0.5331	0.0953	-1.0749	0.0003	-0.1674	0.6242	0.2607	0.1220	0.4281	0.0123	0.2228	0.8111	-0.1054	0.8386	-0.3282	0.4037
CG4998 /// DmirCG4998	CG4998	1624573_at	0.1373	0.4222	-4.3513	0.0010	-2.6705	0.0084	1.4769	0.3921	4.3496	0.0017	2.8727	0.0060	-0.2342	0.8133	-0.0427	0.9492	0.1915	0.6677
CG14656	CG14656	1624574_at	0.1879	0.2277	0.1473	0.4170	0.0196	0.9438	0.0148	0.9838	0.0185	0.9378	0.0037	0.9858	0.1530	0.8497	0.0479	0.9277	-0.1051	0.7951
fru	fru-satori	1624575_a_at	0.1301	0.3603	0.1051	0.4621	0.0235	0.9370	0.1128	0.7596	0.0676	0.7121	-0.0452	0.7991	0.1007	0.8694	0.0310	0.9377	-0.0697	0.8202
CG12994	CG12994	1624576_at	0.2586	0.1737	0.4368	0.1485	0.2153	0.1871	-0.2316	0.5680	-0.4298	0.0444	-0.1982	0.2919	0.1235	0.8486	-0.1826	0.5129	-0.3061	0.2731
Ace	acetylcholinestera	1624577_at	0.0515	0.9413	0.0722	0.5608	-0.2280	0.1459	-0.2896	0.4590	-0.4989	0.0264	-0.2093	0.2743	0.1845	0.9444	-0.4585	0.5880	-0.6429	0.4322
spheroide	CG9675	1624578_at	-0.4033	0.8987	-0.1056	0.7775	-0.1523	0.4426	0.2636	0.9711	-1.5176	0.4273	-1.7812	0.2866	0.1209	0.9913	-1.1998	0.6314	-1.3207	0.5964
Cpr62Ba	CG13934	1624579_at	0.8316	0.0529	1.4043	0.0261	1.9646	0.0016	-0.0079	0.9922	-0.6105	0.0038	-0.6026	0.0024	-0.3663	0.8270	-0.1703	0.8619	0.1961	0.8239
CG9362	CG9362	1624580_at	-0.0612	0.8090	0.1400	0.5054	-0.2621	0.1601	-0.3120	0.3625	-0.3310	0.0907	-0.0190	0.9376	0.1847	0.8270	-0.0830	0.8654	-0.2676	0.4573
dpr7	dpr7	1624581_at	-1.3437	0.0069	-0.7061	0.0498	-0.8425	0.0019	-0.5049	0.2823	-0.6523	0.0261	-0.1474	0.5944	-0.2744	0.7215	-0.1580	0.6576	0.1164	0.7560
SIFR	SIFamide recepto	1624582_a_at	-0.0289	0.8993	0.1063	0.3799	0.0337	0.8639	-0.2924	0.4707	-0.3047	0.1610	-0.0123	0.9645	-0.0382	0.9657	-0.0609	0.8537	-0.0227	0.9463
CG6115 /// DyakCG6115	CG6115	1624583_at	-0.1665	0.3108	0.1137	0.6389	-0.0828	0.7207	0.1084	0.8640	-0.7334	0.0065	-0.8418	0.0021	0.2303	0.7769	-0.1147	0.7970	-0.3449	0.3406
CG32643	CG32643	1624584_at	0.0274	0.9035	0.0957	0.4731	0.0664	0.8070	-0.0409	0.9647	-0.0855	0.7712	-0.0446	0.8776	0.0053	0.9970	-0.0571	0.9161	-0.0624	0.8969
---	---	1624585_at	0.2291	0.1760	0.1562	0.4628	0.1665	0.4697	0.1261	0.8267	0.0573	0.8418	-0.0688	0.7809	-0.0981	0.8775	-0.0446	0.9064	0.0535	0.8717
Prat2	Phosphoribosylan	1624586_a_at	2.8252	0.0024	1.5996	0.0148	2.9052	0.0000	0.9775	0.3386	0.0927	0.9098	-0.8848	0.0931	-0.3222	0.7070	-1.0252	0.0276	-0.7029	0.0859
CG13323	CG13323	1624587_at	-0.0298	0.8909	0.2659	0.2521	0.0504	0.7610	0.1224	0.8403	-0.0040	0.9901	-0.1264	0.5755	0.3068	0.6695	0.2774	0.3259	-0.0294	0.9431
---	---	1624588_at	0.2074	0.2585	0.2874	0.4290	-0.0306	0.9222	-0.3260	0.4239	-0.1852	0.4272	0.1408	0.5199	-0.0406	0.9848	0.1056	0.8785	0.1462	0.8039
cenG1A	centaurin gamma	1624589_a_at	0.0951	0.9150	-0.0742	0.9211	-0.6122	0.0132	0.1161	0.7929	0.2481	0.1663	0.1320	0.4359	0.7474	0.7726	0.2335	0.8857	-0.5139	0.6769
CG17134	CG17134	1624590_at	0.3995	0.0451	1.6295	0.0074	2.4733	0.0010	0.5410	0.0978	0.6100	0.0075	0.0690	0.7397	-0.2402	0.9101	1.4168	0.0761	1.6570	0.0679
CG9222	CG9222	1624591_at	0.0926	0.7051	0.0833	0.5319	0.0069	0.9843	0.0306	0.9744	0.0111	0.9756	-0.0196	0.9471	0.0894	0.9238	0.0891	0.8252	-0.0003	0.9996
CG5188	CG5188	1624592_at	0.2895	0.2035	0.5133	0.0723	0.6385	0.0057	-0.1484	0.8028	-0.1093	0.6886	0.0391	0.8909	-0.1681	0.7633	0.1075	0.6789	0.2756	0.2552
CG14332	CG14332	1624593_at	0.1147	0.6697	0.0563	0.6804	0.1016	0.5130	0.1655	0.6954	0.0984	0.6549	-0.0671	0.7523	-0.0489	0.9469	0.0471	0.8850	0.0960	0.7035
Sk2	Sphingosine kinas	1624594_at	0.1636	0.6964	0.2746	0.6975	0.1706	0.4576	-0.0166	0.9819	0.0547	0.7965	0.0713	0.6944	0.1326	0.9677	0.2684	0.8084	0.1358	0.9089
Rbp2	anon-fast-evolving	1624595_a_at	-0.3278	0.4950	0.9621	0.1940	0.9813	0.0061	-0.2755	0.7138	-0.4993	0.1436	-0.2238	0.4965	-0.2850	0.8940	0.9181	0.2346	1.2031	0.1562
CG31463	CG31463	1624596_at	0.0670	0.6618	-0.0450	0.6681	-0.0495	0.8129	-0.0148	0.9838	0.0886	0.6455	0.1034	0.5365	0.0705	0.8940	0.0274	0.9309	-0.0431	0.8699
---	---	1624597_at	-0.0276	0.8827	-0.0285	0.8314	-0.0679	0.6938	0.0533	0.9436	0.0833	0.7436	0.0299	0.9086	0.0508	0.9646	0.0772	0.8551	0.0264	0.9514
---	---	1624598_at	0.5055	0.0668	0.0131	0.9122	0.1867	0.3384	-0.0089	0.9937	-0.1390	0.5480	-0.1301	0.5356	0.0585	0.9530	-0.1372	0.6762	-0.1956	0.5298
CG12656	CG12656	1624599_at	0.2497	0.5040	0.0474	0.9597	0.6008	0.0914	-0.0569	0.9314	-1.1683	0.0005	-1.1115	0.0004	-0.5192	0.8461	-1.0882	0.3137	-0.5690	0.6331
futsch	Futsch	1624600_at	-0.0782	0.6187	0.1989	0.1142	0.2275	0.2199	-0.1094	0.8126	-0.2074	0.2570	-0.0979	0.5913	0.0070	0.9928	0.1047	0.5716	0.0977	0.6052
---	---	1624601_at	0.1188	0.5463	0.1371	0.3741	0.0888	0.6959	-0.0021	0.9973	0.0070	0.9790	0.0092	0.9668	0.1177	0.8692	0.0018	0.9989	-0.1159	0.7197
CG13235	CG13235	1624602_at	-0.0254	0.9192	-0.1786	0.2708	-0.2979	0.0713	0.1084	0.7840	0.3480	0.0393	0.2396	0.1001	-0.0869	0.9095	-0.0952	0.7755	-0.0082	0.9849
CG10754 /// DlitCG10754	CG10754	1624603_at	0.3771	0.0811	0.2299	0.3972	0.1262	0.4448	-0.0882	0.8190	0.0092	0.9690	0.0973	0.5031	0.1262	0.8510	-0.0342	0.9401	-0.1604	0.5877
CG5629	CG5629	1624604_a_at	-0.4604	0.1718	0.1016	0.4233	-0.3987	0.0920	-0.3783	0.3452	0.2753	0.2329	0.6536	0.0073	-0.0880	0.9514	0.6390	0.1376	0.7270	0.1251
---	---	1624605_at	0.2848	0.1688	0.0477	0.7696	-0.0987	0.7201	0.1210	0.8578	0.2572	0.2924	0.1362	0.5707	0.1576	0.8629	-0.0416	0.9445	-0.1992	0.6150
CG13062	CG13062	1624606_at	0.2312	0.2853	0.0617	0.5890	0.0815	0.7042	-0.0009	0.9994	-0.0508	0.8789	-0.0499	0.8637	0.0648	0.9653	-0.1424	0.7692	-0.2073	0.6306
---	---	1624607_at	-0.2413	0.3420	0.5313	0.0613	0.1858	0.5273	-0.1149	0.8123	-0.7297	0.0032	-0.6148	0.0041	0.1920	0.8960	0.1229	0.8754	-0.0692	0.9273

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
cpo	couch potato	1624608_s_at	-0.3463	0.6390	-0.8181	0.1510	-1.3254	0.0149	0.4923	0.5044	1.3828	0.0045	0.8905	0.0191	0.9312	0.7116	0.8449	0.4203	-0.0864	0.9535
CG31279	CG31279	1624609_at	0.0409	0.8893	0.0117	0.9241	0.2450	0.1209	0.1843	0.7543	0.0542	0.8723	-0.1301	0.6130	-0.0166	0.9875	0.0671	0.8352	0.0837	0.7652
---	---	1624610_at	0.1073	0.6063	0.0696	0.5561	-0.1022	0.5333	-0.0896	0.8747	-0.1215	0.5694	-0.0320	0.8925	0.0978	0.8270	0.0497	0.8438	-0.0481	0.8387
CG2852 /// DyakCG2852	CG2852	1624611_a_at	0.2422	0.1410	0.2573	0.0705	0.7460	0.0008	0.2126	0.4511	0.2609	0.0895	0.0483	0.7765	-0.2925	0.5905	0.3262	0.1634	0.6187	0.0447
Gr59c	Gustatory recepto	1624612_at	0.0890	0.6226	0.0469	0.6494	0.1670	0.3038	0.0379	0.9513	-0.0902	0.6378	-0.1281	0.4244	-0.0034	0.9980	-0.1069	0.7692	-0.1035	0.7674
CG5986	CG5986	1624613_at	0.0107	0.9650	0.4962	0.1573	0.3421	0.1303	-0.0721	0.9289	-0.0023	0.9947	0.0698	0.7932	0.1005	0.9357	0.4410	0.2481	0.3404	0.3977
CG14894	CG14894	1624614_at	-0.1950	0.2325	0.5152	0.0142	0.6047	0.0106	0.1276	0.7121	0.0025	0.9909	-0.1251	0.3923	0.0030	0.9978	0.7015	0.0285	0.6984	0.0410
---	---	1624615_at	0.3631	0.1116	0.0753	0.6494	0.3840	0.0433	0.0544	0.9375	0.2097	0.3044	0.1553	0.4111	-0.0695	0.8972	-0.0698	0.7770	-0.0003	0.9992
Trh	Tryptophan hydro	1624616_at	0.2818	0.0787	0.0154	0.8992	0.3835	0.0929	0.2230	0.5223	0.0350	0.8873	-0.1880	0.2525	-0.0034	0.9984	0.0024	0.9985	0.0058	0.9921
CG1665	CG1665	1624617_at	1.2999	0.0039	1.1303	0.0281	1.9668	0.0000	0.3855	0.3353	0.1301	0.6081	-0.2554	0.2164	-0.4586	0.6960	-0.0200	0.9834	0.4387	0.3884
chrw	chrowded	1624618_at	-0.8223	0.0169	-1.5487	0.0036	-1.1063	0.0046	0.4532	0.4199	0.6221	0.0507	0.1689	0.5874	0.0189	0.9862	0.0011	0.9991	-0.0177	0.9613
CG18477 /// CG31780	CG31780 /// CG1	1624619_s_at	0.7546	0.0390	0.1039	0.8112	0.1670	0.7282	0.0074	0.9956	0.0294	0.9505	0.0220	0.9567	-0.1308	0.9589	-0.3768	0.6218	-0.2461	0.7665
Klp61F	kinesin-like protei	1624620_at	0.5642	0.2683	-0.5333	0.3092	-0.4243	0.0456	-0.2611	0.5735	0.7182	0.0089	0.9793	0.0014	-0.1332	0.9461	-0.1991	0.7903	-0.0658	0.9371
---	---	1624621_at	0.0105	0.9654	0.0532	0.7003	0.0106	0.9614	-0.0891	0.8190	-0.0942	0.5671	-0.0051	0.9787	0.0235	0.9836	0.0259	0.9509	0.0024	0.9960
ACXE	ACXE	1624622_at	0.1342	0.6299	0.2558	0.1871	-0.1303	0.5572	-0.2540	0.5552	-0.1314	0.5869	0.1226	0.5804	0.1573	0.8823	0.0076	0.9935	-0.1498	0.7492
CG32972	CG32972	1624623_at	0.0677	0.6640	0.1021	0.5948	0.0977	0.5320	0.0916	0.7902	0.0607	0.7082	-0.0309	0.8495	0.0670	0.9380	0.1269	0.6686	0.0598	0.8668
mud	mushroom body d	1624624_at	0.1538	0.4958	-0.2229	0.1688	-0.0623	0.7163	0.1052	0.8707	0.2655	0.2408	0.1603	0.4527	-0.0896	0.9095	-0.2548	0.3534	-0.1653	0.5808
CG17150 /// DmirCG17150	CG17150	1624625_a_at	0.0383	0.7975	0.0085	0.9378	0.0714	0.7538	-0.0182	0.9790	-0.0120	0.9634	0.0062	0.9770	0.0458	0.9405	0.0206	0.9463	-0.0252	0.9231
---	---	1624626_at	-0.2770	0.3000	0.0089	0.9471	-0.0052	0.9843	-0.0944	0.8897	-0.1967	0.4012	-0.1023	0.6655	-0.1053	0.8943	0.0106	0.9854	0.1159	0.7341
twin	twin	1624627_s_at	0.3882	0.1283	-0.0496	0.7525	-0.2911	0.3491	0.0187	0.9777	0.7470	0.0017	0.7282	0.0011	0.2527	0.8395	0.2604	0.6394	0.0077	0.9931
CG10396	CG10396	1624628_at	-0.1538	0.5106	-0.0582	0.6455	0.1161	0.5368	0.0237	0.9745	-0.0329	0.8943	-0.0567	0.7800	-0.0825	0.8534	-0.0244	0.9341	0.0581	0.7931
CG32119	CG32119	1624629_at	0.0163	0.9551	-0.1460	0.3311	0.2845	0.1869	-0.0269	0.9819	0.1078	0.7482	0.1347	0.6435	-0.3111	0.6483	-0.0480	0.9085	0.2631	0.3513
CG30103	CG30103	1624630_at	0.0720	0.7145	-0.0363	0.7876	0.0990	0.5521	0.0458	0.9311	-0.0612	0.7396	-0.1070	0.4706	-0.0391	0.9643	-0.0494	0.8850	-0.0104	0.9778
---	---	1624631_x_at	-0.5054	0.0438	-1.0436	0.1243	-1.4498	0.0006	-0.2714	0.7894	0.7014	0.0924	0.9728	0.0170	0.0065	0.9978	0.1391	0.8537	0.1326	0.8514
Klp3A	Kinesin-Like-Prote	1624632_at	-0.2398	0.6770	-1.6258	0.0793	-1.5249	0.0034	-0.2278	0.7135	1.0837	0.0031	1.3114	0.0008	-0.3113	0.9174	-0.1856	0.9075	0.1257	0.9316
---	---	1624633_at	0.2288	0.4079	0.1456	0.3975	0.1140	0.6824	-0.2168	0.7293	0.0428	0.9127	0.2596	0.3072	-0.1977	0.8023	0.0203	0.9704	0.2179	0.5299
CG40050	CG40050	1624634_at	0.1067	0.4847	-0.0454	0.7304	0.2855	0.0595	0.2707	0.3443	0.1711	0.3041	-0.0996	0.5361	-0.0651	0.8987	-0.0592	0.8081	0.0059	0.9849
CG14210	CG14210	1624635_at	0.1733	0.5876	0.0235	0.9612	0.1712	0.3049	0.2940	0.7447	0.5630	0.1536	0.2690	0.4762	0.1270	0.9036	0.4131	0.2610	0.2861	0.4683
ACXD	ACXD	1624636_at	-0.5071	0.2252	2.0444	0.0116	1.6169	0.0031	-0.8332	0.3016	-2.4128	0.0010	-1.5796	0.0035	-0.2494	0.8692	0.3159	0.6303	0.5653	0.3646
adp	adipose	1624637_at	-0.2616	0.4115	0.5279	0.0590	0.5104	0.0107	0.0651	0.9345	-0.3062	0.1875	-0.3713	0.0770	0.1281	0.9142	0.4826	0.2334	0.3545	0.4053
CG4365	CG4365	1624638_a_at	-0.3285	0.0546	0.4421	0.0843	0.7086	0.0038	0.1034	0.8327	-0.6422	0.0050	-0.7456	0.0016	-0.0988	0.8609	0.1261	0.6043	0.2249	0.3360
CG1420	CG1420	1624639_at	0.0019	0.9944	0.1892	0.5791	-0.3229	0.1339	-0.1373	0.8068	0.4156	0.0658	0.5529	0.0134	0.2588	0.7707	0.4581	0.2123	0.1994	0.6260
---	---	1624640_x_at	-0.1497	0.4838	-0.0919	0.5296	-0.0062	0.9780	0.0196	0.9777	0.0708	0.7399	0.0512	0.8013	-0.2063	0.7726	0.0317	0.9492	0.2380	0.4495
---	---	1624641_at	-0.5312	0.0175	-0.0389	0.7275	-0.1111	0.5705	-0.0736	0.9322	-0.4702	0.0706	-0.3966	0.0855	-0.0924	0.8846	-0.0574	0.8657	0.0350	0.9168
CG18420	CG18420	1624642_at	-0.0004	0.9983	-0.0840	0.6183	-0.0317	0.8964	0.1356	0.8025	0.0917	0.7157	-0.0438	0.8623	0.0495	0.9309	-0.0073	0.9838	-0.0568	0.8019
AlCR2	Drosophila allatos	1624643_a_at	0.1668	0.2687	-0.1498	0.5042	0.3024	0.0568	0.1249	0.8512	0.0408	0.9029	-0.0842	0.7498	0.0378	0.9657	-0.0076	0.9873	-0.0454	0.8884
CG5514	CG5514	1624644_a_at	-0.0655	0.8956	-0.1176	0.5965	-0.2170	0.2491	0.1264	0.8864	0.2520	0.4143	0.1257	0.6893	0.1464	0.9142	0.1161	0.8611	-0.0303	0.9659
---	---	1624645_at	0.0056	0.9772	0.2204	0.3046	0.2264	0.2538	0.1332	0.8792	-0.0520	0.8990	-0.1852	0.5242	0.1943	0.7284	0.0855	0.7707	-0.1089	0.6817
---	---	1624646_at	0.3899	0.1975	0.0670	0.5252	0.1454	0.4561	0.1709	0.6916	0.3254	0.1023	0.1545	0.4061	-0.0041	0.9978	-0.0793	0.8754	-0.0751	0.8745
---	---	1624647_at	0.1583	0.5285	-0.2220	0.3769	0.3744	0.1405	0.1726	0.6338	0.0423	0.8541	-0.1303	0.4349	-0.1814	0.8480	-0.0996	0.8546	0.0818	0.8786
---	---	1624648_at	-0.0029	0.9904	-0.3552	0.0448	-0.0871	0.7535	0.2067	0.6823	0.3803	0.1049	0.1736	0.4321	-0.1126	0.8270	-0.0545	0.8514	0.0581	0.8282
rogdi	rogdi	1624649_a_at	0.6984	0.0401	-0.7578	0.0212	-0.3217	0.3126	0.0872	0.9507	0.8371	0.0386	0.7499	0.0384	-0.1215	0.9142	-0.6262	0.1197	-0.5048	0.2199
CG33087	LDLR-like	1624650_at	0.2241	0.6577	0.1624	0.5742	0.0499	0.8218	-0.3138	0.5478	-0.5498	0.0473	-0.2360	0.3404	-0.1485	0.9522	-0.5338	0.4829	-0.3853	0.6312
---	---	1624651_at	0.1860	0.4173	-0.1664	0.4088	-0.5608	0.0118	-0.0651	0.9413	0.4970	0.0547	0.5621	0.0214	0.1513	0.7644	-0.0757	0.7692	-0.2270	0.2966
---	---	1624652_at	0.0630	0.8061	0.0031	0.9810	0.1295	0.5137	0.0775	0.9015	-0.0382	0.8897	-0.1157	0.5649	-0.0198	0.9898	0.0148	0.9797	0.0345	0.9387
Tsp42A	Tetraspanin 42A	1624653_at	0.1397	0.5034	0.0351	0.8384	0.0969	0.6792	-0.1572	0.7380	-0.0266	0.9276	0.1307	0.5170	-0.1124	0.8940	-0.0036	0.9961	0.1088	0.7682
CG13613	CG13613	1624654_at	0.0027	0.9896	-0.0666	0.6488	-0.1784	0.2705	-0.0980	0.8535	-0.1290	0.5283	-0.0310	0.8926	-0.0004	0.9998	-0.0714	0.8248	-0.0710	0.8172
CG6472	CG6472	1624655_at	0.0808	0.6938	-0.1987	0.1529	0.2241	0.2237	0.0437	0.9343	0.0905	0.5940	0.0468	0.7880	-0.1109	0.8608	-0.1243	0.6536	-0.0134	0.9727
CG4293	CG4293	1624656_s_at	1.5136	0.0021	1.2347	0.0321	1.5143	0.0001	-0.0851	0.9042	0.2065	0.3802	0.2916	0.1552	-0.1955	0.8692	0.1541	0.7971	0.3496	0.4758
---	---	1624657_at	0.0408	0.8314	0.1174	0.4549	-0.1131	0.4994	-0.3322	0.4440	-0.1172	0.6613	0.2149	0.3183	-0.0366	0.9653	-0.0502	0.8764	-0.0135	0.9686
CG33158 /// DmirCG33158	CG33158	1624658_at	0.0096	0.9616	-0.2178	0.6775	-0.1637	0.6447	0.2093	0.6176	0.8425	0.0026	0.6332	0.0054	0.3054	0.8160	0.7242	0.1694	0.4188	0.4557
CG32741	CG32741	1624659_at	0.0723	0.7068	-0.1475	0.4878	-0.3568	0.3113	-0.2464	0.7753	0.4173	0.2496	0.6637	0.0457	-0.1614	0.7768	0.1470	0.5715	0.3083	0.2263
Ptp10D	Protein tyrosine pl	1624660_at	-2.0944	0.0043	-0.8446	0.0065	-1.5485	0.0026	-0.4919	0.4822	-1.2327	0.0062	-0.7408	0.0341	0.1263	0.9467	-0.0859	0.9233	-0.2122	0.7484
---	---	1624661_at	0.2771	0.2560	0.4045	0.2163	0.1706	0.2743	-0.2098	0.6473	-0.0986	0.6999	0.1112	0.6231	-0.0105	0.9916	0.0727	0.8123	0.0832	0.7607
CG3106	CG3106	1624662_at	0.2161	0.5312	0.2879	0.1609	0.1315	0.4460	-0.0717	0.9314	-0.0356	0.9178	0.0361	0.9034	-0.0369	0.9848	-0.0599	0.9314	-0.0231	0.9727
vis	vismay	1624663_a_at	-0.4374	0.0398	-0.5115	0.2496	-0.5314	0.0370	0.1675	0.6513	0.4684	0.0171	0.3009	0.0636	0.0214	0.9914	0.1333	0.8153	0.1118	0.8419
snk	snake	1624664_at	-0.0793	0.6458	-0.0835	0.4658	-0.1176	0.4864	0.0170	0.9774	0.0769	0.6417	0.0599	0.6995	0.0398	0.9589	0.0540	0.8594	0.0142	0.9658
CG9257	CG9257	1624665_at	-0.1056	0.7266	0.1412	0.6145	0.6717	0.0704	-0.0042	0.9956	-0.1033	0.6562	-0.0991	0.6384	-0.4841	0.7423	0.1869	0.8174	0.6711	0.2960
CG10092	CG10092	1624666_at	0.1073	0.5521	-0.1483	0.6422	0.2608	0.1420	0.2942	0.4191	0.6215	0.0084	0.3273	0.0719	-0.1665	0.8510	0.3700	0.3000	0.5364	0.1665
CG31760	CG31760	1624667_at	0.0146	0.9570	-0.3396	0.0775	-0.1809	0.3174	0.0768	0.9339	0.4104	0.1293	0.3336	0.1684	-0.2428	0.7370	-0.1301	0.7089	0.1127	0.7504
CG9422	CG9422	1624668_s_at	0.4616	0.0489	0.6207	0.2724	0.4827	0.1440	-0.0967	0.9116	-0.3156	0.2543	-0.2189	0.3935	-0.0123	0.9962	-0.1080	0.8980	-0.0956	0.9020
CG7009	CG7009	1624669_at	-0.4894	0.0678	-0.4977	0.0466	-0.0377	0.8677	0.1153	0.7230	0.0060	0.9790	-0.1093	0.4272	-0.3131	0.6897	-0.0640	0.8910	0.2491	0.4472
CG7589	CG7589	1624670_at	-4.3906	0.0007	-5.0116	0.0101	-5.1775	0.0000	0.1406	0.8578	-0.2447	0.3960	-0.3853	0.1234	0.3264	0.9238	-0.8670	0.4452	-1.1934	0.3045
CG17782	CG17782	1624671_at	0.2009	0.2579	0.1742	0.1881	0.1452	0.5581	0.0901	0.8871	0.0890	0.7214	-0.0010	0.9972	-0.1677	0.7953	-0.2268	0.3976	-0.0591	0.8699
---	---	1624672_at	-0.0833	0.5697	0.0653	0.5774	-0.1160	0.4278	-0.1538	0.6954	-0.2560	0.1561	-0.1021	0.5700	0.1529	0.7440	0.0742	0.7509	-0.0787	0.7266
CG5899	CG5899	1624673_at	0.4554	0.0802	0.2201	0.5850	0.0200	0.9258	0.0012	0.9991	0.4914	0.0656	0.4902	0.0435	0.2393	0.8202	0.3851	0.3650	0.1459	0.7749
CRMP	dihydropyrimidine	1624674_at	0.1538	0.4131	-0.0402	0.7537	0.0994	0.5439	0.0747	0.8610	0.0785	0.6484	0.0038	0.9849	-0.0109	0.9916	-0.0355	0.9274	-0.0247	0.9431
UBL3	Ubiquitin-like 3	1624675_at	-0.3226	0.1926	-0.6945	0.0957	-1.0568	0.0071	-0.1370	0.7155	0.1554	0.3748	0.2925	0.0602	0.2131	0.8943	-0.1621	0.8418	-0.3752	0.5564
CG13331	CG13331	1624676_a_at	0.1227	0.4725	0.0997	0.3313	-0.1978	0.4878	-0.1795	0.7888	0.0272	0.9469	0.2066	0.4219	0.2061	0.7307	0.1042	0.7251	-0.1019	0.7291

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
tnc	tenectin	1624677_at	-0.2351	0.3181	0.0826	0.4307	0.1352	0.6595	-0.0890	0.8844	-0.4864	0.0275	-0.3974	0.0397	0.0406	0.9764	0.0494	0.9233	0.0088	0.9868
CG9570	CG9570	1624678_at	0.1552	0.3295	0.0061	0.9845	-0.0414	0.8172	0.0344	0.9705	0.1126	0.6790	0.0782	0.7653	0.1850	0.7418	-0.0063	0.9905	-0.1914	0.4308
CG33253	CG33253	1624679_at	-0.0376	0.8229	0.0173	0.8691	0.1336	0.3876	0.1628	0.5664	0.1021	0.5060	-0.0607	0.6913	-0.0353	0.9742	0.0069	0.9911	0.0423	0.9099
RhoGEF4	RhoGEF4	1624680_a_at	0.1723	0.6913	-0.0208	0.9515	-0.2511	0.4073	-0.1177	0.9030	0.2923	0.3611	0.4100	0.1422	-0.1107	0.9514	-0.2096	0.7424	-0.0989	0.8942
---	---	1624681_at	0.2469	0.2937	-0.0100	0.9330	0.2015	0.4200	0.0618	0.9435	0.0610	0.8484	-0.0008	0.9981	-0.1277	0.8903	-0.0968	0.8356	0.0309	0.9494
---	---	1624682_at	-0.0094	0.9629	-0.0282	0.8307	0.0099	0.9713	-0.0448	0.9404	-0.0248	0.9183	0.0200	0.9247	0.1558	0.7823	0.1217	0.6458	-0.0341	0.9194
CG17680	CG17680	1624683_at	-0.1724	0.5045	-0.0668	0.8347	-0.1675	0.3633	0.0398	0.9610	-0.3899	0.0748	-0.4297	0.0339	0.1810	0.8521	-0.1240	0.8119	-0.3050	0.4627
exo84	exo84	1624684_at	-0.0799	0.7528	0.1420	0.5305	0.3237	0.0878	0.0707	0.9186	0.0528	0.8484	-0.0179	0.9473	-0.0541	0.9682	0.2574	0.4812	0.3116	0.3921
NUCB1	NUCB1	1624685_at	0.2592	0.5506	0.0980	0.7919	0.4458	0.0414	0.3068	0.6325	0.8901	0.0120	0.5834	0.0438	-0.0615	0.9816	0.7234	0.2123	0.7849	0.2095
dnc	dunce	1624686_a_at	-1.5926	0.0025	-1.8434	0.0029	-2.3992	0.0001	-0.5309	0.0654	-0.2216	0.1768	0.3092	0.0423	0.0436	0.9870	-0.3606	0.5778	-0.4042	0.5300
CG15594	CG15594	1624687_a_at	0.0132	0.9628	-0.0537	0.7562	0.4457	0.0792	0.1439	0.8554	0.0554	0.8855	-0.0886	0.7805	-0.1503	0.8554	0.0499	0.9263	0.2002	0.5789
CG9360 /// DsimCG9360	CG9360	1624688_at	-2.6020	0.0008	-0.7260	0.2670	-3.0054	0.0005	-1.9321	0.0492	-2.7439	0.0011	-0.8118	0.1064	0.3833	0.8202	-0.7875	0.2430	-1.1709	0.1271
---	---	1624689_at	-0.0069	0.9761	0.1750	0.7052	0.0031	0.9910	-0.1607	0.7230	0.0687	0.7817	0.2294	0.2097	-0.0609	0.9755	0.1728	0.7698	0.2337	0.6569
CG6878	CG6878	1624690_at	-0.3035	0.1827	0.0308	0.9471	0.1586	0.6361	0.0505	0.9649	-0.4047	0.1855	-0.4553	0.0973	-0.1474	0.9296	-0.0609	0.9450	0.0864	0.9080
---	---	1624691_at	0.1411	0.4732	0.0153	0.8833	-0.0432	0.7971	-0.0294	0.9637	0.0593	0.7727	0.0887	0.6069	0.1341	0.8292	0.0720	0.8370	-0.0621	0.8545
CG6830	CG6830	1624692_at	0.7501	0.4498	1.3977	0.1301	0.8805	0.0476	-0.0474	0.9909	-0.7890	0.3988	-0.7416	0.3791	0.4370	0.8439	-0.0831	0.9557	-0.5200	0.5961
Arf102F	ADP ribosylation f	1624693_at	0.5879	0.0137	1.1730	0.0026	1.2898	0.0001	0.0240	0.9714	-0.0421	0.8472	-0.0661	0.7129	-0.0565	0.9447	0.5165	0.0571	0.5731	0.0559
CG13597	CG13597	1624694_at	-0.0415	0.8430	-0.0549	0.6028	-0.0338	0.8960	0.0156	0.9909	0.0165	0.9715	0.0009	0.9981	0.0442	0.9457	-0.0498	0.8546	-0.0940	0.6600
CG3999	CG3999	1624695_at	2.3647	0.0014	0.8684	0.3022	2.7007	0.0001	0.9132	0.3553	0.5021	0.3856	-0.4111	0.4391	-0.9518	0.6592	-0.7054	0.4212	0.2464	0.8237
CG12769	CG12769	1624696_a_at	-0.2041	0.1905	-0.0107	0.9581	0.0336	0.8625	-0.0116	0.9874	-0.0115	0.9669	0.0001	0.9995	-0.0155	0.9900	0.1032	0.7453	0.1187	0.6914
CG14983	CG14983	1624697_at	0.1513	0.4657	0.0622	0.6261	0.1622	0.4250	0.0394	0.9540	-0.0067	0.9809	-0.0461	0.8293	-0.0223	0.9816	-0.0689	0.8061	-0.0465	0.8746
CG31397	CG31397	1624698_at	0.1926	0.3120	0.0771	0.6535	0.1575	0.5355	-0.0033	0.9962	0.0640	0.8308	0.0673	0.7990	-0.1266	0.8202	-0.0053	0.9924	0.1213	0.6232
Bx	rhombotin	1624699_s_at	0.0218	0.9317	-0.6283	0.2935	-0.1748	0.7505	0.0098	0.9956	0.6439	0.2006	0.6340	0.1593	-0.4800	0.6999	-0.3748	0.4833	0.1052	0.8834
CG10175	CG10175	1624700_a_at	-1.0139	0.0710	-3.3384	0.0087	-2.1479	0.0010	0.5203	0.5975	0.9891	0.0518	0.4687	0.2990	-0.5925	0.7464	-1.1653	0.1321	-0.5728	0.4740
CG14050	CG14050	1624701_at	0.3647	0.2393	0.0087	0.9536	0.1397	0.5421	0.4364	0.3381	0.3692	0.1581	-0.0672	0.8218	0.1285	0.8609	0.0701	0.8662	-0.0584	0.8861
dpr10	dpr10	1624702_s_at	-0.7325	0.1739	-0.0167	0.8799	-0.0873	0.6090	-0.1637	0.9067	-0.6104	0.1724	-0.4467	0.2709	-0.0488	0.9441	-0.0500	0.8657	-0.0012	0.9979
pb	proposcipedia	1624703_at	0.1705	0.3433	0.1012	0.3490	0.2182	0.2281	0.2193	0.4571	0.0229	0.9187	-0.1964	0.1686	0.2147	0.6868	-0.0654	0.8174	-0.2801	0.2144
CG4741	CG4741	1624704_at	-0.0072	0.9747	-0.0354	0.8553	-0.0285	0.8952	-0.0495	0.9311	0.0343	0.8782	0.0838	0.6240	-0.1360	0.7826	-0.0900	0.7101	0.0461	0.8727
fax	failed axon conne	1624705_a_at	-1.9606	0.0087	-2.1843	0.0045	-2.5632	0.0000	-0.3926	0.5019	-0.5392	0.0836	-0.1466	0.6466	0.0740	0.9764	-0.7659	0.1897	-0.8399	0.1883
CG34349	CG11819	1624706_at	-1.0556	0.0051	-0.9352	0.0949	-1.0501	0.0372	0.1558	0.8337	0.1740	0.5665	0.0182	0.9591	0.1150	0.9725	0.2937	0.7754	0.1787	0.8746
---	---	1624707_at	-0.0509	0.8067	-0.0148	0.8885	-0.1671	0.2599	-0.0087	0.9934	0.1837	0.3832	0.1924	0.3030	-0.0206	0.9844	0.0027	0.9961	0.0233	0.9434
CG15769	CG15769	1624708_at	0.3485	0.2368	0.1442	0.3450	0.0728	0.6792	0.1998	0.7159	0.0007	0.9985	-0.1991	0.3916	0.1822	0.7917	-0.1621	0.6032	-0.3442	0.2557
---	---	1624709_at	0.0799	0.6051	0.0719	0.5898	0.1975	0.4199	0.2070	0.5008	0.1427	0.3933	-0.0643	0.7093	0.0771	0.9168	-0.0387	0.9229	-0.1158	0.6817
---	---	1624710_at	-0.2069	0.2457	-0.0292	0.8459	0.0795	0.5835	0.1108	0.8076	-0.1100	0.5745	-0.2209	0.1723	0.0006	0.9998	-0.0834	0.7383	-0.0840	0.7307
CG31740	CG31740	1624711_at	0.2003	0.3496	0.0839	0.4984	0.1750	0.4813	-0.0676	0.9500	-0.1022	0.7774	-0.0347	0.9248	-0.0536	0.9653	-0.1161	0.7728	-0.0626	0.8904
---	---	1624712_at	0.0667	0.7444	0.0892	0.4829	-0.1965	0.3834	-0.0886	0.8217	-0.1231	0.4383	-0.0345	0.8456	0.1216	0.9108	-0.0661	0.9111	-0.1877	0.6528
CG34371	CG13548	1624713_at	0.1715	0.4462	0.1229	0.5350	0.1921	0.3947	0.1041	0.9036	-0.0261	0.9505	-0.1302	0.6519	0.0252	0.9852	-0.0547	0.9085	-0.0798	0.8411
woc	without children	1624714_at	0.0934	0.8061	-0.3747	0.4727	-0.0903	0.6045	0.1196	0.8028	0.3879	0.0486	0.2682	0.1168	-0.1644	0.9405	-0.0694	0.9499	0.0950	0.9194
mtacp1	NADH-ubiquinone	1624715_at	0.0388	0.9302	0.7988	0.0534	0.6562	0.0062	-0.1712	0.7031	-1.0258	0.0011	-0.8546	0.0014	-0.0525	0.9816	-0.1522	0.8066	-0.0997	0.8800
---	---	1624716_at	-0.0123	0.9518	0.0931	0.6301	0.0961	0.6826	0.2013	0.7117	0.1270	0.6476	-0.0743	0.7878	0.0575	0.9277	0.0851	0.7200	0.0276	0.9231
CG32647	CG32647	1624717_s_at	0.7462	0.1174	-0.7920	0.3147	0.3960	0.4453	0.4104	0.1634	0.3676	0.0452	-0.0428	0.8338	-0.8077	0.7633	-1.2726	0.2404	-0.4649	0.7135
CG32584	CG32584	1624718_a_at	-0.0285	0.9484	-0.0153	0.9517	-0.0541	0.8021	-0.0039	0.9956	-0.0520	0.8495	-0.0482	0.8435	0.0998	0.9589	0.0307	0.9739	-0.0692	0.9273
CG10777 /// DmirCG10777	CG10777	1624719_at	-0.5693	0.2450	-0.2849	0.2583	-0.5573	0.0453	0.0250	0.9854	0.3314	0.2861	0.3064	0.2709	0.1446	0.9445	0.4020	0.5398	0.2574	0.7156
CG6043	CG6043	1624720_s_at	-1.6279	0.0024	-1.0796	0.0131	-2.0508	0.0008	-0.5087	0.1163	-0.5566	0.0118	-0.0479	0.8296	0.4312	0.8049	0.0176	0.9921	-0.4136	0.5988
vimar	visceral mesoderm	1624721_at	-0.0095	0.9860	-0.1162	0.7454	-0.8673	0.0087	-0.1870	0.5933	0.5290	0.0097	0.7159	0.0015	0.4012	0.8192	0.3900	0.6111	-0.0111	0.9928
---	---	1624722_at	0.0661	0.7640	-0.0263	0.8777	0.0325	0.8973	0.0853	0.8732	0.1589	0.4049	0.0737	0.7076	-0.0045	0.9964	0.0039	0.9941	0.0084	0.9838
---	---	1624723_at	0.1349	0.4781	0.3348	0.1513	0.0913	0.7209	-0.0654	0.9377	0.0524	0.8720	0.1178	0.6394	-0.1125	0.8692	0.0395	0.9275	0.1520	0.6053
b6	b6	1624724_at	0.0991	0.6301	0.3752	0.2790	0.1795	0.2903	0.1946	0.6471	0.0041	0.9897	-0.1905	0.3083	0.2427	0.7215	0.1889	0.5264	-0.0538	0.8918
CG9626	CG9626	1624725_at	-2.3837	0.0012	-1.9092	0.0308	-2.9442	0.0001	-0.2156	0.7851	-0.4270	0.1881	-0.2114	0.5002	0.8209	0.6955	0.0532	0.9703	-0.7677	0.3828
CG7145	CG7145	1624726_s_at	-0.7717	0.0097	-0.7422	0.1363	-1.1146	0.0047	-0.3426	0.2101	-0.6695	0.0024	-0.3269	0.0333	-0.0882	0.9667	-0.6406	0.2385	-0.5523	0.3375
Ubp64E	Ubiquitin-specific	1624727_s_at	-1.3730	0.0073	-0.7304	0.0171	-1.1378	0.0002	-0.1559	0.7855	-0.5458	0.0284	-0.3900	0.0676	0.2014	0.8648	0.0209	0.9810	-0.1805	0.7408
---	---	1624728_at	-0.0264	0.9255	0.1688	0.3974	0.2988	0.1369	0.0347	0.9627	-0.0499	0.8449	-0.0846	0.6822	-0.1083	0.9142	0.0770	0.8790	0.1853	0.6261
lola	longitudinals abse	1624729_at	-0.5906	0.2382	-0.5609	0.3939	-0.5361	0.0509	-0.0425	0.9659	0.2736	0.3019	0.3161	0.1772	-0.0882	0.9816	0.3385	0.7183	0.4267	0.6294
---	---	1624730_at	-0.1609	0.3459	-0.2169	0.1332	-0.2546	0.1914	-0.0218	0.9770	-0.0159	0.9538	0.0059	0.9799	-0.0528	0.9659	-0.0584	0.9085	-0.0056	0.9924
CG33205	CG33205	1624731_s_at	-2.1395	0.0069	-2.1491	0.0280	-3.3745	0.0001	-0.8699	0.0125	-0.7123	0.0024	0.1576	0.3223	0.3546	0.9011	-0.8032	0.4409	-1.1577	0.2830
GstE5	Glutathione S tran	1624732_at	0.0289	0.9371	-0.0154	0.9380	-0.0864	0.6525	-0.1168	0.8738	-0.3680	0.1484	-0.2511	0.2773	0.0787	0.9644	-0.3616	0.4479	-0.4404	0.3679
---	---	1624733_at	0.2075	0.3051	0.0242	0.8859	-0.1818	0.4495	-0.1179	0.7647	0.1557	0.3647	0.2736	0.0712	-0.1088	0.8745	-0.0854	0.8061	0.0234	0.9512
---	---	1624734_at	0.0857	0.5519	0.0123	0.9403	0.0152	0.9299	-0.1679	0.5744	-0.0131	0.9542	0.1548	0.2622	-0.0174	0.9862	-0.0824	0.7727	-0.0649	0.8247
---	---	1624735_at	0.0717	0.7119	-0.1391	0.3706	0.2384	0.1513	0.0725	0.8971	-0.0647	0.7712	-0.1372	0.4287	-0.3167	0.5352	-0.2581	0.2672	0.0585	0.8509
CG5819	CG5819	1624736_a_at	2.3827	0.0135	1.5422	0.0917	2.3318	0.0004	0.4659	0.6699	0.2350	0.6934	-0.2309	0.6686	-0.2273	0.9400	-0.4341	0.6716	-0.2068	0.8664
CG31001	CG31001	1624737_at	-0.0204	0.9149	-0.2776	0.1178	-0.4249	0.0499	-0.1261	0.7625	0.2281	0.2010	0.3542	0.0341	0.0463	0.9742	-0.0605	0.9125	-0.1068	0.8087
CG11386 /// CG33222	CG11386 /// CG3	1624738_at	0.5157	0.0233	0.3937	0.1826	0.5335	0.0058	0.0599	0.9186	0.0153	0.9546	-0.0447	0.8282	0.1990	0.8157	0.0454	0.9354	-0.1536	0.6964
CG3081	CG3081	1624739_at	0.2404	0.3410	0.2870	0.1652	0.1960	0.2612	-0.0788	0.9329	-0.1051	0.7486	-0.0263	0.9392	-0.0069	0.9939	-0.0738	0.7745	-0.0669	0.7918
---	---	1624740_at	0.0604	0.7541	0.1365	0.2576	0.1339	0.4684	0.0108	0.9900	-0.0713	0.7574	-0.0821	0.6851	0.0168	0.9898	-0.0060	0.9925	-0.0228	0.9525
CG14464	CG14464	1624741_at	0.4383	0.0586	-0.1375	0.4016	0.0281	0.9046	0.1280	0.8500	0.3950	0.1083	0.2670	0.2267	-0.1105	0.8860	-0.1919	0.5234	-0.0814	0.8229
Elongin-C	d-elongin C	1624742_a_at	-0.1889	0.2860	0.1470	0.5299	0.3250	0.0573	-0.0254	0.9640	-0.6122	0.0024	-0.5868	0.0017	-0.0824	0.9365	-0.1755	0.6162	-0.0930	0.8199
PpD3	protein phosphata	1624743_s_at	0.0679	0.7482	-0.0802	0.8107	-0.1134	0.5813	-0.1122	0.8244	0.2598	0.1823	0.3720	0.0401	-0.0549	0.9717	0.1708	0.7019	0.2257	0.5944
nuf	nuclear-fallout	1624744_a_at	-0.7100	0.0080	-1.4983	0.0126	-1.6904	0.0009	0.1059	0.8164	0.2809	0.1158	0.1751	0.2794	0.3484	0.7758	-0.4320	0.4094	-0.7804	0.1651
Ilp5	Drosophila insulin	1624745_at	0.1264	0.5444	-0.0037	0.9897	0.2875	0.1338	0.0845	0.8914	0.1191	0.6002	0.0346	0.8891	-0.0331	0.9707	-0.1041	0.6820	-0.0710	0.7964

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
betaTub85D	beta-tubulin	1624746_at	0.1549	0.4338	-0.0143	0.8893	0.1632	0.2771	0.0980	0.8640	0.2868	0.1562	0.1887	0.3063	-0.0271	0.9717	0.0516	0.8445	0.0787	0.7221
Ts	Ts	1624747_at	0.2038	0.5946	0.0815	0.9203	-0.2916	0.4848	-0.2527	0.6669	0.3512	0.2077	0.6039	0.0239	-0.1425	0.9726	0.1326	0.9366	0.2751	0.8350
Mnn1	Menin 1	1624748_a_at	-0.1099	0.7855	0.6818	0.0764	0.4912	0.0178	0.0453	0.9760	-0.5843	0.1148	-0.6297	0.0618	0.0800	0.9405	0.3040	0.3446	0.2240	0.5156
CG32079	CG32079	1624749_at	0.1428	0.3655	-0.3904	0.1509	0.0134	0.9443	0.3784	0.2129	0.4701	0.0183	0.0917	0.6220	-0.0954	0.8904	-0.1348	0.6319	-0.0394	0.9122
CG8545	CG8545	1624750_at	0.4664	0.3323	-0.3930	0.3398	-0.1428	0.6592	0.4039	0.6457	1.3705	0.0069	0.9666	0.0194	0.1646	0.9341	0.4299	0.5141	0.2653	0.7128
CG10478	CG10478	1624751_at	-0.1174	0.5488	-0.1908	0.3758	-0.2845	0.2011	0.0654	0.9165	0.1589	0.4287	0.0935	0.6384	0.0999	0.8940	0.0324	0.9430	-0.0675	0.8533
Prestin	Prestin	1624752_at	-0.4711	0.0288	0.1692	0.7004	0.1770	0.7171	0.1448	0.7929	0.2058	0.3694	0.0609	0.8091	0.1987	0.9101	0.7577	0.2118	0.5590	0.3828
CG13136	CG13136	1624753_at	0.1044	0.6566	0.0833	0.6693	0.1068	0.5357	-0.1451	0.7362	-0.2290	0.2303	-0.0839	0.6743	0.0129	0.9914	0.0600	0.8794	0.0472	0.9023
---	---	1624754_at	0.0280	0.9071	-0.0256	0.8547	-0.0977	0.5813	0.0483	0.9435	0.1474	0.4643	0.0990	0.6117	0.0722	0.9238	-0.0244	0.9508	-0.0966	0.7351
flw	Protein phosphata	1624755_a_at	-0.2863	0.0960	-0.3149	0.3492	-0.1401	0.3541	0.0243	0.9658	0.1842	0.2173	0.1599	0.2332	-0.1007	0.9309	0.2427	0.5363	0.3433	0.3748
CG5009	CG5009	1624756_at	-0.0040	0.9916	-0.5328	0.0204	-0.4410	0.0181	-0.0688	0.8979	-0.0099	0.9711	0.0589	0.7595	-0.1550	0.8510	-0.5802	0.0968	-0.4251	0.2282
CG10202	CG10202	1624757_at	0.0913	0.7272	0.1383	0.4619	0.0150	0.9539	-0.0761	0.8943	-0.0442	0.8596	0.0319	0.8890	-0.1984	0.8122	-0.0772	0.8764	0.1213	0.7631
CG4318	CG4318	1624758_at	-0.1588	0.4060	0.1797	0.4557	-0.1450	0.4000	-0.1676	0.6591	-0.1745	0.3476	-0.0069	0.9761	0.2191	0.7423	0.2428	0.3778	0.0237	0.9523
Antp	Sex comb extra	1624759_s_at	-0.2202	0.7160	-0.2310	0.2940	-0.4171	0.1277	-0.1029	0.9486	-0.3129	0.5065	-0.2100	0.6481	-0.0386	0.9841	-0.4962	0.2430	-0.4576	0.3089
CG8306	CG8306	1624760_at	-0.4140	0.3952	-1.0559	0.0999	-0.1791	0.3110	0.1854	0.7327	0.0508	0.8739	-0.1345	0.5769	-0.6872	0.7095	-0.6844	0.3669	0.0028	0.9987
CG14535	CG14535	1624761_at	0.0065	0.9861	-0.0135	0.9127	-0.2572	0.2040	-0.1253	0.8355	0.1369	0.5809	0.2622	0.1980	0.0540	0.9604	0.0036	0.9959	-0.0504	0.9056
CG14254	CG14254	1624762_at	0.5067	0.1690	0.0845	0.7257	0.0587	0.7927	0.0213	0.9860	0.2507	0.3982	0.2294	0.3930	0.1903	0.8521	0.0140	0.9870	-0.1763	0.7115
CG2556 /// DsimCG2556	CG2556	1624763_at	-2.3620	0.0068	-4.1135	0.0045	-3.3582	0.0001	0.3625	0.7664	1.3231	0.0192	0.9606	0.0446	-0.4663	0.8042	-0.3917	0.6447	0.0746	0.9460
CG40265	CG40265	1624764_a_at	0.1248	0.5150	-0.0706	0.5564	-0.0381	0.8765	0.1272	0.7488	0.0763	0.7021	-0.0510	0.7926	0.0826	0.9046	-0.0707	0.8304	-0.1533	0.5612
mRpL36	mitochondrial ribo	1624765_at	0.0081	0.9804	0.5700	0.1896	0.6591	0.0231	0.0177	0.9823	-0.3653	0.0561	-0.3830	0.0300	-0.0419	0.9860	0.1737	0.7985	0.2157	0.7240
Dsp1	Dorsal switch prot	1624766_s_at	-0.3994	0.1155	-0.0479	0.9337	-0.3566	0.1752	-0.2993	0.5067	-0.2858	0.2308	0.0135	0.9645	0.0526	0.9816	0.1270	0.8655	0.0744	0.9195
---	---	1624767_at	0.2695	0.1744	0.1688	0.4452	0.0697	0.6838	0.0292	0.9744	-0.0321	0.9167	-0.0613	0.8039	0.0272	0.9737	-0.1112	0.6075	-0.1384	0.5120
CG31407	CG31407	1624768_at	0.0310	0.8596	0.0137	0.8976	0.0660	0.7445	0.0102	0.9883	0.1042	0.5566	0.0941	0.5633	0.0555	0.9348	0.0188	0.9587	-0.0367	0.9026
Tl	toll	1624769_s_at	0.2670	0.2526	0.1681	0.3780	0.0307	0.8931	-0.2497	0.5283	-0.1795	0.3983	0.0702	0.7564	-0.0421	0.9717	-0.0145	0.9784	0.0276	0.9474
CG17929	CG17929	1624770_at	0.0774	0.6934	0.0264	0.7925	0.0600	0.8000	0.0398	0.9592	-0.0210	0.9438	-0.0608	0.7889	0.0874	0.8906	0.1053	0.6961	0.0179	0.9589
Nipped-A	Nipped	1624771_at	0.0456	0.8666	0.0664	0.5014	0.2695	0.2463	0.1341	0.8336	0.0600	0.8484	-0.0741	0.7835	0.1892	0.7215	0.0665	0.8185	-0.1226	0.6094
CG14186	CG14186	1624772_at	-0.4336	0.0351	-0.1554	0.5666	-0.8223	0.0207	-0.3475	0.4333	-0.1191	0.6663	0.2284	0.3003	0.0368	0.9732	0.0009	0.9996	-0.0360	0.9231
---	---	1624773_at	-0.0371	0.8695	0.0333	0.7424	0.1642	0.3045	0.0866	0.8735	0.0326	0.8999	-0.0540	0.7990	-0.0676	0.8943	0.0225	0.9411	0.0901	0.6630
Fas2	Fasciclin II	1624774_a_at	-0.2551	0.2013	-0.3837	0.5936	-0.3806	0.0722	0.1321	0.8196	0.1335	0.5892	0.0014	0.9963	0.1016	0.9672	-0.0643	0.9507	-0.1659	0.8414
CG10628	CG10628	1624775_at	-0.4672	0.1062	-0.0528	0.8866	0.3276	0.1439	0.1730	0.7368	-0.5867	0.0184	-0.7597	0.0035	-0.2218	0.8966	-0.0758	0.9409	0.1460	0.8613
mtacp1	NADH-ubiquinone	1624776_a_at	0.0146	0.9700	0.4740	0.1298	0.1939	0.3846	-0.1734	0.7053	-0.7795	0.0037	-0.6061	0.0068	0.1507	0.9011	-0.2665	0.5692	-0.4173	0.3622
CG11113	CG11113	1624777_at	0.2740	0.1477	0.1729	0.1993	0.1368	0.3495	-0.0957	0.8174	-0.0271	0.9040	0.0686	0.6877	0.0400	0.9816	0.0999	0.8420	0.0599	0.9067
pcm	Pacman	1624778_at	0.4083	0.3342	-0.0518	0.9323	-0.0052	0.9848	0.0544	0.9149	0.5816	0.0039	0.5273	0.0036	0.0148	0.9964	0.1409	0.8941	0.1261	0.8982
CG11583 /// DyakCG11583	CG11583	1624779_at	0.2438	0.5598	0.5403	0.3742	0.9251	0.0098	0.3836	0.6506	0.4438	0.2813	0.0602	0.9028	-0.0689	0.9816	0.5324	0.4359	0.6014	0.3898
CG2995 /// DereCG2995 ///	CG2995	1624780_at	0.3299	0.3422	0.8885	0.0843	0.2854	0.1081	-0.4310	0.5301	0.3735	0.3014	0.8045	0.0202	0.2284	0.8062	0.7212	0.0778	0.4929	0.2210
---	---	1624781_at	0.0261	0.9057	-0.1614	0.2675	-0.0151	0.9609	0.1223	0.8462	0.3063	0.1836	0.1840	0.3926	0.0027	0.9981	-0.0075	0.9877	-0.0102	0.9789
Uev1A	Uev1A	1624782_a_at	-0.0334	0.8693	0.3712	0.1322	0.3014	0.0990	-0.1948	0.5930	-0.2978	0.1058	-0.1030	0.5752	-0.2036	0.7697	0.1150	0.7412	0.3186	0.2960
CG14961	CG14961	1624783_at	-0.0483	0.8009	-0.0818	0.6323	0.1335	0.4314	0.3597	0.2851	0.1122	0.6054	-0.2475	0.1607	0.0367	0.9636	-0.0467	0.8824	-0.0834	0.7419
---	---	1624784_at	0.0897	0.5530	0.0380	0.7192	0.1277	0.5759	0.0350	0.9562	0.1108	0.5493	0.0758	0.6722	-0.0545	0.9457	-0.0757	0.8084	-0.0213	0.9504
---	---	1624785_at	-0.0340	0.8816	-0.0128	0.9101	0.1120	0.4684	-0.0315	0.9562	-0.0113	0.9612	0.0202	0.9151	-0.1528	0.8283	0.1218	0.7184	0.2746	0.3645
---	---	1624786_at	0.1926	0.2819	0.1221	0.5006	0.4000	0.0235	-0.0200	0.9772	-0.0116	0.9644	0.0084	0.9689	-0.0940	0.8461	-0.0421	0.8850	0.0519	0.8393
CG16959	CG16959	1624787_s_at	-0.1653	0.5080	-0.5152	0.0332	0.0687	0.7327	0.5810	0.0492	0.4987	0.0100	-0.0823	0.6333	-0.2525	0.7464	-0.0677	0.8879	0.1848	0.6038
CG3773	anon-fast-evolving	1624788_at	0.1469	0.4674	0.0244	0.8696	0.2610	0.1212	0.1933	0.5732	0.4069	0.0278	0.2137	0.1714	0.0260	0.9816	0.2939	0.2357	0.2679	0.3061
---	---	1624789_at	0.1325	0.6036	0.0149	0.9161	0.0309	0.8905	-0.0189	0.9863	-0.0884	0.7828	-0.0695	0.8163	-0.0030	0.9984	-0.2079	0.4677	-0.2050	0.4835
Toll-6	Toll-6	1624790_at	0.2158	0.3005	0.1202	0.2470	0.0282	0.8802	-0.0032	0.9956	0.0338	0.8834	0.0370	0.8529	0.0387	0.9589	-0.0675	0.8018	-0.1062	0.6389
---	---	1624791_at	0.1195	0.5441	0.0908	0.5008	0.1312	0.4157	0.0298	0.9712	0.0192	0.9515	-0.0106	0.9679	-0.0363	0.9589	-0.0541	0.8445	-0.0177	0.9508
RpS28b	Ribosomal protein	1624792_at	0.3221	0.0566	0.5508	0.0415	0.3584	0.0400	-0.0281	0.9598	-0.1640	0.2629	-0.1360	0.3047	0.1460	0.7893	-0.0508	0.8791	-0.1969	0.4067
GstD7	Glutathione S tran	1624793_at	1.0482	0.2450	0.7023	0.0614	1.6805	0.0002	-0.0200	0.9861	0.0373	0.9226	0.0574	0.8582	-0.9416	0.6749	-0.2520	0.8394	0.6896	0.4719
CG10845	CG10845	1624794_at	0.1225	0.4963	0.0927	0.4465	0.2177	0.1368	0.0077	0.9934	0.0083	0.9764	0.0006	0.9981	0.0160	0.9875	-0.0035	0.9942	-0.0195	0.9520
CG1105	CG1105	1624795_at	-1.2848	0.0035	-0.4202	0.0993	-0.3468	0.1971	-0.0074	0.9944	-0.6385	0.0123	-0.6311	0.0079	0.0434	0.9831	0.2346	0.6519	0.1912	0.7255
CG2930	CG2930	1624796_at	0.1546	0.4173	0.1286	0.3881	0.4317	0.0608	0.1791	0.5128	-0.0338	0.8583	-0.2128	0.0991	0.0073	0.9938	0.1259	0.5607	0.1186	0.5906
CG1637	CG1637	1624797_at	0.3779	0.4674	-0.4704	0.2876	0.1380	0.6208	0.2271	0.7647	0.3041	0.3574	0.0770	0.8358	-0.3938	0.8236	-0.5934	0.4190	-0.1996	0.8299
Galpha49B	G-protein-alpha[[q	1624798_s_at	-0.2541	0.3350	0.7478	0.0258	-0.1025	0.6458	-0.2570	0.5619	0.0513	0.8639	0.3084	0.1274	0.6113	0.5434	1.0091	0.0527	0.3978	0.4051
---	---	1624799_at	-0.1108	0.6062	-0.0236	0.8715	-0.0398	0.8625	-0.0616	0.8942	0.0097	0.9678	0.0713	0.6547	-0.1296	0.8806	0.1031	0.8082	0.2327	0.4957
CG5224	CG5224	1624800_at	0.4175	0.2738	1.0338	0.0067	1.3230	0.0001	-0.0420	0.9426	-0.4844	0.0105	-0.4425	0.0096	-0.3291	0.8015	0.1189	0.8792	0.4480	0.4204
CG1138	CG1138	1624801_at	0.1004	0.5065	-0.0313	0.8450	-0.0207	0.9231	0.0939	0.7937	0.1092	0.4760	0.0153	0.9328	-0.0351	0.9545	-0.0380	0.8841	-0.0029	0.9924
CG31686	CG31686	1624802_at	1.4619	0.0005	1.5545	0.0027	3.6372	0.0000	1.9978	0.0033	1.1752	0.0029	-0.8226	0.0084	0.1483	0.7305	1.3583	0.0005	1.2100	0.0023
CG34013	CG34013	1624803_at	-0.4887	0.1175	0.0133	0.9293	-0.9036	0.0036	-1.1175	0.0189	-0.5747	0.0295	0.5428	0.0236	0.1177	0.8202	0.1165	0.6106	-0.0011	0.9979
---	---	1624804_at	0.1561	0.4326	-0.0348	0.8744	-0.0275	0.9127	0.0252	0.9649	0.0887	0.5940	0.0635	0.6918	0.0602	0.9530	0.0423	0.9306	-0.0179	0.9696
CG31041	CG31041	1624805_at	-0.0257	0.9240	0.2083	0.2748	-0.0163	0.9373	-0.2672	0.4455	-0.1748	0.3696	0.0924	0.6363	0.1292	0.8963	0.1761	0.6658	0.0470	0.9271
yellow-g	yellow-g	1624806_at	-2.0746	0.0255	-3.3877	0.0027	-3.4599	0.0000	0.0119	0.9927	0.5843	0.0335	0.5724	0.0231	0.0339	0.9914	-0.6927	0.3340	-0.7265	0.3354
CG5629	CG5629	1624807_at	-1.2856	0.0177	-0.0186	0.9606	-1.0071	0.0017	-0.7667	0.1109	-0.9973	0.0047	-0.2306	0.3834	0.2735	0.8270	0.2599	0.6449	-0.0136	0.9870
CheA75a	CheA75a	1624808_at	-0.2188	0.4062	0.0188	0.8527	-0.0894	0.5780	-0.1502	0.7351	-0.2613	0.1832	-0.1111	0.5704	-0.0928	0.9056	-0.0862	0.8131	0.0066	0.9880
---	---	1624809_s_at	0.3328	0.1794	-0.0148	0.9471	0.0474	0.8354	0.0928	0.8794	0.3460	0.0983	0.2531	0.1749	0.0086	0.9950	-0.1518	0.6626	-0.1603	0.6389
CG32774	CG32774	1624810_at	-0.3063	0.1565	0.4403	0.3659	0.0138	0.9585	-0.3945	0.5680	-0.3107	0.3906	0.0838	0.8358	0.1851	0.8972	0.4938	0.3424	0.3087	0.5874
---	---	1624811_at	-0.1118	0.4509	0.0807	0.4701	-0.0249	0.9299	-0.1383	0.8379	-0.1937	0.4643	-0.0554	0.8506	0.0048	0.9964	-0.0132	0.9745	-0.0180	0.9592
CG11070	CG11070	1624812_at	0.1259	0.5562	0.3266	0.3433	0.3329	0.0944	-0.1066	0.8817	0.2354	0.3447	0.3420	0.1170	-0.0577	0.9588	0.4532	0.1395	0.5109	0.1303
CG34008	CG34008	1624813_s_at	0.4530	0.0188	0.1956	0.5514	0.3988	0.0694	-0.3022	0.2789	-0.0673	0.7265	0.2349	0.1096	-0.3643	0.6832	-0.2943	0.4212	0.0700	0.8900
---	---	1624814_at	0.3028	0.1132	0.0348	0.8032	0.1649	0.4149	-0.0906	0.8498	-0.0909	0.6422	-0.0003	0.9989	-0.2493	0.7036	-0.3617	0.1718	-0.1124	0.7154

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	**Gene Title**	**Affy ID**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**
Ssrp	Chorion factor	1624815_at	0.1307	0.6343	0.4668	0.1782	0.7668	0.0304	-0.0896	0.8605	-0.0401	0.8684	0.0495	0.8106	-0.4432	0.7981	0.4031	0.6019	0.8463	0.2592
CG30283	CG30283	1624816_at	-0.1011	0.7302	0.1797	0.5252	0.1963	0.3946	-0.0390	0.9603	-0.1095	0.6431	-0.0705	0.7597	-0.1183	0.9163	0.2422	0.5523	0.3605	0.3680
Sr-CIV	Scavenger recept	1624817_at	-0.3517	0.2725	-0.2879	0.2381	-0.2489	0.2384	0.0333	0.9763	-0.0493	0.8965	-0.0826	0.7905	-0.1176	0.8655	-0.1664	0.5679	-0.0488	0.8984
Sply	Sphingosine-1-ph	1624818_s_at	0.8762	0.0127	1.0810	0.0051	1.5478	0.0001	0.1221	0.7658	0.3469	0.0522	0.2248	0.1496	-0.3947	0.7116	0.4262	0.3259	0.8209	0.1011
---	---	1624819_s_at	1.6962	0.0039	0.8626	0.0285	1.0105	0.0120	0.2607	0.8053	1.0130	0.0243	0.7523	0.0506	-0.0088	0.9959	0.0623	0.9194	0.0711	0.8957
CG8800	CG8800	1624820_at	0.4572	0.1001	0.0751	0.7968	0.3438	0.1380	0.3209	0.5681	0.5930	0.0457	0.2721	0.2987	-0.1935	0.7726	-0.0830	0.8258	0.1105	0.7390
CG8586 /// DyakCG8586	CG8586	1624821_at	-0.3425	0.1995	-1.6901	0.0054	-0.9637	0.0017	0.3750	0.3481	0.8784	0.0032	0.5034	0.0225	-0.2899	0.6749	-0.3946	0.1579	-0.1047	0.7520
CG15133	CG15133	1624822_at	0.0892	0.5743	0.0015	0.9922	0.1209	0.4146	0.2425	0.4770	0.2190	0.2309	-0.0235	0.9162	0.0927	0.8571	0.0932	0.6935	0.0005	0.9992
---	---	1624823_at	-0.0018	0.9961	-0.0028	0.9823	0.1508	0.3528	0.2763	0.4420	0.1698	0.3992	-0.1065	0.5869	0.1089	0.8815	0.0103	0.9848	-0.0986	0.7614
Jon74E	Jonah 74E	1624824_at	1.0290	0.1977	0.2148	0.4759	0.4502	0.0931	0.0341	0.9637	0.1420	0.4939	0.1078	0.5851	-0.1509	0.9687	-0.6985	0.4991	-0.5477	0.6136
---	---	1624825_s_at	0.2146	0.1908	0.1311	0.4808	0.0529	0.7965	-0.1084	0.8385	-0.0445	0.8656	0.0639	0.7722	0.0373	0.9521	0.0383	0.8874	0.0010	0.9979
CG8552	CG8552	1624826_at	0.4980	0.1019	0.5195	0.0750	0.6768	0.0016	0.1680	0.7519	0.6052	0.0168	0.4372	0.0402	-0.0387	0.9816	0.5497	0.1421	0.5885	0.1502
CG11698	CG11698	1624827_at	0.0312	0.8892	0.0540	0.7793	0.0579	0.8008	0.0188	0.9838	-0.1655	0.4617	-0.1843	0.3517	0.0850	0.8609	0.0738	0.7506	-0.0113	0.9702
Obp50a	Odorant-binding p	1624828_at	0.0075	0.9719	-0.0402	0.7658	0.1343	0.4812	-0.0107	0.9922	0.0841	0.7533	0.0949	0.6873	-0.1157	0.8680	0.0204	0.9646	0.1360	0.6469
CG33288	CG33288	1624829_at	-0.0753	0.7348	-0.2174	0.1538	-0.0645	0.7282	0.1312	0.8196	0.2982	0.1839	0.1671	0.4304	-0.0528	0.9156	-0.0046	0.9908	0.0482	0.8247
Act79B	79B Actin	1624830_at	-0.0041	0.9915	-0.1266	0.4987	0.1336	0.6106	0.2569	0.6070	0.1795	0.4958	-0.0774	0.7817	-0.0143	0.9939	0.0441	0.9483	0.0584	0.9194
CG31360	CG31360	1624831_at	-0.2004	0.3834	0.2339	0.3942	0.3635	0.0564	0.0734	0.9373	-0.4647	0.0870	-0.5382	0.0335	-0.1795	0.7779	0.0009	0.9996	0.1804	0.5360
---	---	1624832_s_at	-0.0403	0.8714	-0.1997	0.1574	-0.0911	0.8004	-0.0453	0.9665	0.0731	0.8413	0.1183	0.6899	-0.3686	0.3517	-0.3039	0.1418	0.0648	0.8003
mAcR-60C	muscarinic recept	1624833_a_at	-1.3761	0.0011	-3.5740	0.0039	-2.7891	0.0007	0.9248	0.4602	2.1936	0.0067	1.2689	0.0428	0.0343	0.9848	0.2230	0.6139	0.1888	0.6766
CG14015	CG14015	1624834_at	-0.1403	0.6044	0.0075	0.9491	-0.0828	0.6644	0.0150	0.9863	-0.2292	0.2728	-0.2443	0.1885	0.1032	0.8825	-0.1187	0.6918	-0.2219	0.4119
CG10869	CG10869	1624835_at	-0.0525	0.8407	0.1559	0.4733	0.0683	0.8122	-0.0075	0.9917	-0.1488	0.3297	-0.1414	0.3003	0.1909	0.8258	0.0919	0.8519	-0.0991	0.8256
CG13795	CG13795	1624836_at	2.9824	0.0036	0.3680	0.5707	1.6475	0.0018	0.1115	0.9649	0.9571	0.1560	0.8456	0.1616	-1.0763	0.3712	-1.5615	0.0397	-0.4851	0.4764
CG7506	CG7506	1624837_at	0.1856	0.4763	-0.2291	0.4266	-0.1619	0.4150	0.0885	0.8732	0.1938	0.3199	0.1053	0.5837	0.0707	0.9636	-0.1503	0.7693	-0.2211	0.6268
sNPF	short neuropeptid	1624838_a_at	-1.9179	0.0723	-0.0766	0.4731	-0.2783	0.2706	-0.2168	0.9380	-1.7355	0.0407	-1.5187	0.0442	-0.1843	0.7726	-0.0689	0.8551	0.1154	0.7095
h	hairy	1624839_at	-0.5114	0.0364	-0.5367	0.0323	-0.6738	0.0017	-0.2788	0.2401	0.3683	0.0170	0.6471	0.0009	-0.0470	0.9717	0.4780	0.1450	0.5250	0.1443
Acp29AB	accessory gland p	1624840_at	0.0434	0.7638	-0.1878	0.1281	0.1380	0.3748	0.3627	0.1412	0.3131	0.0423	-0.0496	0.7614	0.0048	0.9952	0.0888	0.6683	0.0840	0.6861
CG13585 /// DyakCG13585	CG13585	1624841_s_at	0.8983	0.0020	0.7786	0.0077	0.6899	0.0146	0.1991	0.5128	0.2260	0.1575	0.0269	0.8883	0.3437	0.7697	0.1194	0.8642	-0.2243	0.6853
koi	CG18584	1624842_at	-0.5668	0.0238	0.1683	0.6586	0.0393	0.8280	0.1353	0.9130	-0.1103	0.8210	-0.2456	0.5165	0.2430	0.7561	0.5479	0.1049	0.3049	0.3695
---	---	1624843_at	-0.0466	0.7755	-0.0766	0.6025	0.0273	0.9003	0.1280	0.8155	0.1758	0.4305	0.0478	0.8483	-0.0617	0.9330	-0.0738	0.8068	-0.0121	0.9739
---	---	1624844_x_at	0.1775	0.3334	0.0205	0.8399	0.0650	0.7770	-0.1400	0.8463	0.2084	0.4495	0.3484	0.1402	-0.2249	0.7266	-0.0093	0.9870	0.2156	0.4520
sun	stunted	1624845_a_at	-0.0996	0.5800	-0.1226	0.4551	-0.0886	0.7159	-0.1092	0.8908	-0.2940	0.2742	-0.1848	0.4666	-0.2730	0.7392	-0.1417	0.7240	0.1312	0.7442
CG10764	CG10764	1624846_at	-0.0234	0.9370	0.7289	0.0655	0.3057	0.3084	-1.0119	0.0383	-0.3886	0.1478	0.6233	0.0178	-0.4057	0.7395	0.5453	0.2764	0.9510	0.1013
CG11985	CG11985	1624847_at	0.0701	0.6934	1.3415	0.0126	0.6186	0.0105	-0.1220	0.8009	-0.3971	0.0469	-0.2751	0.1129	0.4830	0.5978	0.7757	0.0708	0.2927	0.4824
Prp18	Prp18	1624848_at	0.5475	0.0089	-0.1683	0.6047	-0.0938	0.5814	-0.0543	0.9375	0.4537	0.0319	0.5079	0.0125	-0.0638	0.9535	-0.0344	0.9482	0.0294	0.9486
Prosap	Prosap	1624849_at	-0.6238	0.2777	0.0708	0.9189	-0.9316	0.0065	-0.6201	0.1446	-0.1937	0.4703	0.4264	0.0663	0.2922	0.9246	0.4491	0.7010	0.1569	0.9121
CG3556	CG3556	1624850_at	-0.6228	0.1359	-0.4329	0.0723	-0.7622	0.0044	-0.3019	0.5120	-0.3113	0.1983	-0.0094	0.9754	-0.1342	0.8744	-0.4888	0.1392	-0.3546	0.2993
DsimCG32629 /// Tango13	CG32629 /// Trans	1624851_at	2.2210	0.0318	2.0704	0.0176	2.7803	0.0010	-0.2398	0.7196	-0.2682	0.3868	-0.0284	0.9397	-0.2328	0.9571	0.3768	0.8132	0.6095	0.6457
CG9817	CG9817	1624852_at	0.0918	0.7047	0.1332	0.1934	0.3922	0.1030	0.0550	0.9247	0.0576	0.7858	0.0027	0.9904	-0.0096	0.9948	0.2456	0.4794	0.2552	0.4683
CG11635	CG11635	1624853_at	0.1220	0.6590	-0.1496	0.4966	0.2785	0.3338	0.1500	0.8543	0.0318	0.9402	-0.1182	0.7035	-0.0627	0.9698	-0.1218	0.8292	-0.0590	0.9197
Lis-1	Paf-AHbeta	1624854_s_at	-0.6278	0.0350	-0.9986	0.0057	-1.3059	0.0001	-0.2226	0.5633	-0.0965	0.6679	0.1261	0.5087	0.0152	0.9928	-0.4461	0.2487	-0.4613	0.2660
CG31988	CG31988	1624855_at	-0.2570	0.2103	-0.3630	0.1094	-0.1428	0.5105	0.0249	0.9761	0.3357	0.1013	0.3108	0.0900	-0.2097	0.7997	0.1037	0.8168	0.3134	0.3762
c(2)M	crossover suppres	1624856_at	-0.1910	0.3534	-0.7558	0.0845	-0.6200	0.2120	0.0375	0.9766	0.7820	0.0210	0.7446	0.0161	-0.2907	0.8655	0.2168	0.8053	0.5075	0.4746
CG17510	CG17510	1624857_at	-0.0122	0.9783	0.2764	0.4212	-0.2443	0.2219	-0.3208	0.6041	-0.3850	0.2147	-0.0642	0.8599	0.2469	0.7726	0.0484	0.9347	-0.1985	0.6166
CG6693	CG6693	1624858_at	-0.5665	0.0131	-0.2036	0.2181	-0.2372	0.1876	-0.0265	0.9637	-0.1290	0.4170	-0.1024	0.4861	-0.0850	0.9030	0.0250	0.9508	0.1100	0.6978
rpr	reaper	1624859_at	0.2045	0.4703	0.4424	0.3424	0.7070	0.0665	0.1686	0.8209	-0.0489	0.9029	-0.2175	0.4251	0.0346	0.9901	0.3668	0.5620	0.3322	0.6094
CG31804	CG31804	1624860_at	-0.0645	0.7022	0.1006	0.4593	0.1735	0.3489	-0.0665	0.9068	-0.0251	0.9220	0.0414	0.8444	-0.0902	0.8494	0.1158	0.5800	0.2060	0.3081
AlkB	AlkB	1624861_at	0.5287	0.0182	0.2333	0.4058	0.2402	0.1622	-0.0931	0.8905	0.2556	0.2611	0.3488	0.0859	-0.1646	0.8554	0.0105	0.9898	0.1751	0.6630
---	---	1624862_at	0.3787	0.0421	0.1832	0.3965	0.1138	0.7421	-0.0044	0.9962	0.2173	0.5091	0.2217	0.4505	-0.0562	0.9630	0.0433	0.9353	0.0994	0.8073
CG8042	CG8042	1624863_a_at	-0.5496	0.0881	0.8133	0.1107	1.3959	0.0002	0.2805	0.4455	-0.4259	0.0399	-0.7064	0.0027	-0.3222	0.7768	0.7902	0.1137	1.1123	0.0607
dock	dreadlocks	1624864_s_at	-0.7805	0.0583	0.5210	0.1084	0.4305	0.1235	-0.0050	0.9956	-0.6455	0.0045	-0.6404	0.0027	-0.1044	0.9588	0.4332	0.4406	0.5375	0.3538
---	---	1624865_at	0.1484	0.3594	0.0234	0.8634	-0.0153	0.9496	-0.0667	0.9004	-0.0913	0.6361	-0.0247	0.9058	-0.0009	0.9998	-0.1363	0.7199	-0.1354	0.7171
sif	still life type 1	1624866_at	0.1315	0.6843	-0.0098	0.9304	-0.0163	0.9506	0.2751	0.5357	0.2914	0.2072	0.0162	0.9565	-0.0255	0.9841	-0.1573	0.6122	-0.1318	0.6800
CG13964	CG13964	1624867_at	-0.2844	0.2529	0.0497	0.6303	-0.2095	0.4662	-0.1029	0.9011	-0.4287	0.1090	-0.3259	0.1737	0.0296	0.9774	0.0583	0.8665	0.0286	0.9342
Cap-D3	Chromosome ass	1624868_at	0.0536	0.7539	-0.0428	0.7091	-0.1688	0.2733	0.0571	0.9036	0.1036	0.5242	0.0464	0.7860	0.0260	0.9742	-0.1420	0.4864	-0.1680	0.4078
CG12511	CG12511	1624869_at	0.3062	0.1025	0.0572	0.6502	0.1506	0.3391	0.0060	0.9937	-0.0388	0.8383	-0.0449	0.7865	0.0697	0.9298	-0.1203	0.6714	-0.1900	0.4742
CG16853	CG16853	1624870_at	-0.1880	0.2585	0.0398	0.7108	0.0709	0.7362	-0.2468	0.5145	-0.3047	0.1265	-0.0579	0.7972	-0.0586	0.9221	0.0539	0.8407	0.1125	0.5995
---	---	1624871_s_at	0.1332	0.4212	0.1761	0.2123	0.3439	0.1799	0.1529	0.7432	-0.0867	0.7160	-0.2396	0.1911	0.0940	0.9004	0.1170	0.7078	0.0230	0.9525
---	---	1624872_at	0.0588	0.7656	0.0524	0.6286	0.2611	0.1409	-0.0312	0.9602	0.0118	0.9619	0.0431	0.8172	-0.1044	0.8215	0.0429	0.8760	0.1472	0.4573
kar	karmoisin	1624873_at	0.9124	0.1410	-0.0272	0.9413	1.1592	0.0030	-0.6925	0.1032	-0.3742	0.1350	0.3184	0.1552	-1.8296	0.2652	-1.4934	0.0843	0.3362	0.7223
CG32271	CG32271	1624874_at	0.6977	0.0200	0.4354	0.0258	0.5932	0.0452	-0.1163	0.8422	-0.0093	0.9780	0.1069	0.6288	0.0806	0.9340	-0.0826	0.8445	-0.1632	0.6263
CG3650	CG3650	1624875_at	0.3459	0.0342	0.1652	0.3194	0.3031	0.1146	0.2032	0.5070	0.1023	0.5592	-0.1008	0.5236	0.0510	0.9683	0.0428	0.9372	-0.0082	0.9875
Ect4	Ect4	1624876_at	0.0220	0.9533	0.0536	0.7950	-0.8653	0.0487	-0.4427	0.6120	-0.0989	0.8637	0.3437	0.3984	0.3063	0.7423	0.1010	0.8511	-0.2053	0.6294
CG1561	CG1561	1624877_at	0.0789	0.7706	0.1171	0.5341	0.2605	0.0945	0.0072	0.9943	-0.1363	0.5483	-0.1435	0.4755	-0.0247	0.9806	-0.0854	0.7424	-0.0607	0.8259
---	---	1624878_at	0.1164	0.5917	0.0241	0.8098	0.0729	0.6769	-0.0060	0.9956	0.0461	0.8811	0.0521	0.8454	0.0257	0.9773	-0.0062	0.9894	-0.0318	0.9128
Awh	arrowhead	1624879_at	0.2496	0.3117	0.0288	0.7672	0.1473	0.4742	-0.0953	0.8449	0.1013	0.6077	0.1966	0.2213	-0.0278	0.9848	0.0135	0.9829	0.0414	0.9282
CG8813	CG8813	1624880_at	0.1228	0.5638	0.2589	0.0946	0.0778	0.6346	-0.0984	0.8498	-0.0797	0.7168	0.0187	0.9364	0.0642	0.9421	-0.0444	0.9167	-0.1085	0.7280
CG11448	CG11448	1624881_at	-0.8077	0.0894	-1.6292	0.0159	-0.9348	0.0195	0.8250	0.1787	1.4834	0.0027	0.6584	0.0519	0.1371	0.9495	0.6174	0.3259	0.4804	0.4699
CG4066	CG4066	1624882_at	0.0440	0.8389	-0.0855	0.3812	0.0353	0.8175	0.0592	0.9228	0.1237	0.5288	0.0644	0.7465	-0.0085	0.9923	0.0287	0.9277	0.0372	0.8925
Nedd8	Nedd8	1624883_at	0.3896	0.0739	0.8243	0.0323	0.7797	0.0008	0.1373	0.6969	-0.2378	0.1411	-0.3751	0.0178	0.1895	0.8000	0.2164	0.4997	0.0269	0.9512

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG32712 /// DsimCG32712	CG32712	1624884_at	0.1333	0.3683	0.1195	0.4690	-0.0130	0.9614	-0.0446	0.9482	-0.1096	0.6024	-0.0650	0.7566	0.1008	0.9095	-0.0193	0.9709	-0.1200	0.7426
CG32645	CG32645	1624885_at	0.3481	0.2670	-0.5053	0.1018	-0.7195	0.0148	-0.1390	0.8800	0.0054	0.9903	0.1443	0.6585	0.2649	0.8141	-0.6903	0.1377	-0.9552	0.0772
---	---	1624886_s_at	0.2246	0.3555	-0.0314	0.7530	0.0481	0.8530	0.0148	0.9905	0.3734	0.1713	0.3586	0.1423	0.0588	0.9340	0.0850	0.7504	0.0262	0.9342
CG8223	CG8223	1624887_at	0.0605	0.8816	0.4896	0.2053	0.4825	0.0303	0.1520	0.8572	-0.1510	0.6587	-0.3030	0.2671	0.2465	0.8091	0.3794	0.3620	0.1329	0.7947
CG7081	CG7081	1624888_at	0.6832	0.0184	0.3846	0.1422	0.5920	0.0044	-0.0294	0.9777	0.1353	0.6452	0.1647	0.5157	-0.2272	0.6898	-0.0494	0.8824	0.1779	0.4582
l(2)k16918	lethal (2) k16918	1624889_a_at	-1.4695	0.0260	-0.9445	0.0092	-1.3821	0.0019	-0.3560	0.6908	0.3312	0.4413	0.6872	0.0670	-0.0419	0.9860	0.5917	0.2507	0.6336	0.2528
CG12393	CG12393	1624890_s_at	-0.6499	0.0049	-0.2886	0.1301	-0.2633	0.0827	-0.0627	0.8815	-0.2452	0.0878	-0.1825	0.1529	-0.1111	0.8465	0.0451	0.9003	0.1562	0.5332
CG13183	CG13183	1624891_at	-1.5756	0.0098	-4.4804	0.0019	-3.0799	0.0007	1.3661	0.2511	3.5278	0.0011	2.1616	0.0053	-0.0749	0.9618	0.3774	0.3778	0.4524	0.3097
CG11892 /// DyakCG11892	CG11892	1624892_s_at	-1.1343	0.0054	-0.5372	0.5295	-1.7281	0.0044	-0.7058	0.1682	-0.5640	0.0729	0.1418	0.6604	0.3654	0.8940	-0.0082	0.9980	-0.3736	0.7516
CG8852	CG8852	1624893_at	-1.5784	0.0020	-1.0094	0.0395	-1.7059	0.0000	0.0686	0.9094	-0.2781	0.1450	-0.3468	0.0482	0.0890	0.9717	-0.0189	0.9887	-0.1078	0.9023
CSN8	Drosophila COP9	1624894_s_at	0.5668	0.0113	0.5733	0.0411	0.3249	0.0636	-0.0826	0.8915	0.3084	0.1285	0.3910	0.0384	0.1192	0.8331	0.2951	0.1957	0.1758	0.4703
CG7529	CG7529	1624895_at	0.7442	0.3322	1.0974	0.1098	0.2667	0.3495	-0.3679	0.7924	-1.1135	0.0548	-0.7456	0.1409	0.4236	0.8692	-0.6882	0.5140	-1.1118	0.2891
---	---	1624896_at	-0.0868	0.7466	0.0374	0.8724	0.1901	0.2038	-0.0802	0.8987	-0.0025	0.9935	0.0777	0.7236	0.0088	0.9942	0.1355	0.6512	0.1267	0.6746
l(1)G0237	lethal (1) G0237	1624897_at	0.0737	0.8335	-0.7076	0.0088	-0.4982	0.0230	0.2870	0.5453	0.8746	0.0044	0.5877	0.0156	-0.0632	0.9577	0.0091	0.9905	0.0723	0.8729
Ark	Apaf-1 related kill	1624898_a_at	0.1591	0.4612	-0.0179	0.8983	-0.1064	0.5138	0.1075	0.8384	0.2429	0.2171	0.1355	0.4702	0.1886	0.7707	0.0109	0.9838	-0.1777	0.5403
Ras85D	Ras GTPase	1624899_at	0.0432	0.8252	-0.0293	0.8928	-0.2921	0.0687	-0.2254	0.4420	0.0223	0.9211	0.2478	0.0841	0.0154	0.9885	0.0342	0.9264	0.0188	0.9548
CG31051	CG31051	1624900_at	-1.1650	0.0435	-0.1468	0.3050	-0.7555	0.0126	-0.1990	0.8844	-0.8665	0.0668	-0.6675	0.1090	0.0629	0.9589	0.0465	0.9325	-0.0164	0.9761
CG9272	CG9272	1624901_at	-0.4453	0.0407	-0.6969	0.0397	-1.0571	0.0157	-0.5375	0.1476	0.2180	0.3393	0.7555	0.0033	-0.1820	0.8825	-0.3079	0.5271	-0.1258	0.8310
PH4alphaNE2	prolyl-4-hydroxyla	1624902_at	0.3111	0.1818	0.1627	0.3191	0.1688	0.3521	-0.0318	0.9649	0.0490	0.8432	0.0808	0.6873	-0.0731	0.9341	-0.0235	0.9600	0.0496	0.8984
ppk21	pickpocket 21	1624903_at	0.3172	0.0990	0.0259	0.8605	-0.1191	0.7840	-0.0274	0.9838	0.1957	0.5577	0.2231	0.4444	0.0584	0.9816	-0.1846	0.7873	-0.2430	0.6881
---	---	1624904_at	0.3000	0.1869	0.2162	0.5117	0.1318	0.5994	-0.0194	0.9862	0.1199	0.6956	0.1393	0.6038	0.0212	0.9862	0.0028	0.9970	-0.0184	0.9640
CG9000	CG9000	1624905_at	0.3386	0.1473	0.1602	0.6348	0.2927	0.0601	0.0595	0.9042	0.3295	0.0460	0.2700	0.0638	-0.0539	0.9742	0.2298	0.6059	0.2836	0.5145
RpS27A	male female steril	1624906_at	0.1814	0.2201	0.8969	0.0150	0.9564	0.0029	-0.0128	0.9912	-0.6860	0.0140	-0.6732	0.0093	0.1102	0.8425	0.0025	0.9962	-0.1077	0.6630
Tsp42Ee	tetraspanin 42E	1624907_at	-0.3661	0.0591	-0.0207	0.9212	-0.6242	0.0032	-0.1396	0.7005	-0.1404	0.4139	-0.0008	0.9972	0.4169	0.3738	0.2538	0.3132	-0.1631	0.5499
CG15287 /// CG33090	CG15287 /// CG3	1624908_at	0.1710	0.5026	0.0336	0.8396	-0.0019	0.9942	-0.0209	0.9777	-0.0164	0.9531	0.0046	0.9850	0.0265	0.9816	-0.0622	0.8586	-0.0887	0.7615
CG32778	CG32778	1624909_at	0.1992	0.4148	0.0006	0.9987	-0.2205	0.3486	-0.0465	0.9345	0.1742	0.3019	0.2207	0.1381	0.0130	0.9926	-0.0273	0.9590	-0.0403	0.9273
CG7653	CG7653	1624910_at	0.2943	0.3759	0.0008	0.9994	0.0069	0.9865	0.1193	0.8701	0.3851	0.1305	0.2658	0.2467	0.1822	0.8513	-0.0001	1.0000	-0.1824	0.6817
Cyp28d2	Cyp28d2	1624911_at	-1.4112	0.0023	-0.6214	0.2423	-1.1695	0.0090	-0.3184	0.6041	-0.7037	0.0302	-0.3853	0.1621	0.1013	0.9705	-0.0962	0.9310	-0.1975	0.8205
CG15256	CG15256	1624912_at	-0.0147	0.9506	-0.0932	0.5873	0.0767	0.6249	0.1076	0.7539	0.1032	0.5099	-0.0044	0.9811	-0.0352	0.9742	0.0650	0.8640	0.1002	0.7492
oaf	transcript-near-de	1624913_s_at	0.4014	0.1831	0.6757	0.0648	0.9484	0.0008	0.0128	0.9898	-0.0619	0.8308	-0.0747	0.7653	-0.2227	0.8157	0.1422	0.7656	0.3648	0.3678
CG8690	CG8690	1624914_at	0.1327	0.5357	-0.0977	0.6645	-0.2041	0.3637	-0.0256	0.9803	0.1444	0.6060	0.1700	0.4848	-0.1222	0.8909	-0.2986	0.3667	-0.1764	0.6261
---	---	1624915_s_at	0.3451	0.0528	-0.0947	0.3902	-0.0733	0.7408	0.0835	0.8667	0.3694	0.0416	0.2859	0.0708	-0.0806	0.9238	-0.1637	0.5776	-0.0831	0.8073
CG33293	CG33293	1624916_a_at	0.1988	0.2853	0.1742	0.4388	0.0088	0.9729	0.0570	0.9311	0.0368	0.8882	-0.0202	0.9328	0.0668	0.9690	0.0138	0.9885	-0.0530	0.9337
CG12900	CG12900	1624917_at	-0.0599	0.7206	0.0168	0.8688	0.0282	0.8705	-0.0880	0.7952	-0.0981	0.5010	-0.0101	0.9544	-0.0789	0.8960	-0.1015	0.6864	-0.0225	0.9433
---	---	1624918_at	0.1376	0.4681	-0.0090	0.9337	0.0750	0.6377	-0.0052	0.9951	0.0311	0.8942	0.0363	0.8560	0.0088	0.9914	-0.0253	0.9326	-0.0341	0.8947
CG8525	CG8525	1624919_at	0.7917	0.0254	0.4541	0.2035	0.9545	0.0010	0.3120	0.3454	0.1857	0.3376	-0.1263	0.4883	-0.2920	0.8016	-0.2181	0.6873	0.0739	0.9121
---	---	1624920_at	0.0322	0.8620	0.0371	0.7072	0.0318	0.9048	-0.1392	0.6861	-0.0976	0.5727	0.0416	0.8185	0.0903	0.9047	0.0422	0.9207	-0.0481	0.8982
CG14459	CG14459	1624921_at	0.1242	0.4817	0.0167	0.9507	0.2433	0.1653	0.0627	0.9171	0.1199	0.5504	0.0572	0.7837	-0.1853	0.8215	-0.0862	0.8543	0.0991	0.8153
---	---	1624922_at	0.0695	0.7159	0.0807	0.6049	0.0105	0.9625	-0.1077	0.8550	-0.0422	0.8821	0.0655	0.7826	0.0348	0.9677	0.0730	0.7985	0.0382	0.9025
---	---	1624923_s_at	0.0082	0.9785	0.4474	0.0888	0.2834	0.1260	0.2524	0.4455	0.1390	0.4596	-0.1134	0.5164	0.2811	0.7196	0.5523	0.0968	0.2712	0.4114
CG31182	CG31182	1624924_at	-0.4860	0.4548	-1.4265	0.0884	-1.0051	0.0059	-0.0931	0.9518	0.7552	0.0803	0.8483	0.0343	-0.4221	0.8609	-0.1580	0.9167	0.2642	0.8289
CG14446	CG14446	1624925_at	0.2921	0.1674	0.0598	0.5962	-0.0595	0.7707	-0.0225	0.9760	0.0062	0.9818	0.0287	0.8972	0.0904	0.8571	-0.0460	0.8754	-0.1364	0.5257
CG13557	CG13557	1624926_at	0.1352	0.5636	-0.1119	0.5007	0.3201	0.0674	-0.0989	0.8604	-0.1265	0.5603	-0.0276	0.9100	-0.3540	0.3400	-0.1026	0.6314	0.2515	0.2202
---	---	1624927_at	0.0810	0.6462	0.1373	0.3599	-0.1318	0.5004	-0.1713	0.5507	-0.0679	0.6904	0.1034	0.4619	0.0067	0.9942	-0.0410	0.8967	-0.0477	0.8641
TORC	TORC	1624928_at	-0.1193	0.7201	0.0579	0.8908	0.2165	0.2869	0.0884	0.8402	0.0736	0.6957	-0.0148	0.9422	0.0097	0.9964	0.2108	0.7525	0.2011	0.7568
CG31710	CG31710	1624929_at	-0.3746	0.0528	-0.2676	0.4356	-0.9183	0.0065	-0.1478	0.7598	0.1266	0.5724	0.2745	0.1377	0.0630	0.9611	0.0697	0.8981	0.0067	0.9919
---	---	1624930_at	-0.1600	0.4085	0.0463	0.6181	0.0079	0.9753	-0.1988	0.6844	-0.2664	0.2415	-0.0676	0.7904	-0.0556	0.9390	0.0999	0.6915	0.1556	0.5041
cib	ciboulot	1624931_at	-1.0517	0.0027	-0.2051	0.5348	-1.2854	0.0011	-0.1470	0.7929	-0.3189	0.1613	-0.1718	0.4224	0.8534	0.2884	0.6646	0.1077	-0.1888	0.6689
Obp49a	Odorant-binding p	1624932_at	0.3603	0.1898	0.0769	0.8122	-0.0821	0.7499	0.0739	0.9098	0.2246	0.2805	0.1507	0.4386	-0.0195	0.9923	-0.0661	0.9277	-0.0466	0.9431
---	---	1624933_s_at	0.1343	0.5777	0.0319	0.7968	0.0196	0.9182	0.0037	0.9956	-0.0655	0.7051	-0.0693	0.6552	0.1367	0.8850	-0.0354	0.9507	-0.1721	0.6574
olf413	olf413	1624934_at	-0.2220	0.5153	-0.2882	0.1109	-0.1132	0.6758	0.1570	0.8865	-0.0226	0.9689	-0.1796	0.6382	-0.0824	0.9365	0.0431	0.9345	0.1255	0.7397
CG9601	CG9601	1624935_at	-0.0563	0.7659	0.0838	0.5147	0.2960	0.0889	-0.3451	0.3040	-0.4153	0.0417	-0.0702	0.7412	-0.3350	0.5126	-0.0512	0.8830	0.2838	0.2557
---	---	1624936_at	-0.0096	0.9791	0.1479	0.3225	-0.0205	0.9501	0.0298	0.9838	-0.1044	0.8004	-0.1342	0.7064	0.2023	0.8016	0.2031	0.5647	0.0008	0.9992
CG12288	CG12288	1624937_at	0.9407	0.1514	-0.0734	0.9230	-0.1772	0.7441	0.0400	0.9857	2.0152	0.0027	1.9752	0.0017	0.1009	0.9831	0.8046	0.4753	0.7037	0.5481
CG14128	CG14128	1624938_at	0.4133	0.0426	0.1461	0.3756	0.2318	0.1315	-0.0113	0.9865	0.0404	0.8478	0.0517	0.7733	0.0059	0.9943	-0.0679	0.7710	-0.0738	0.7389
Cyp6a21	Cyp6a21	1624939_at	-1.7734	0.0231	-3.0890	0.0060	-2.7150	0.0000	-0.0002	0.9999	-0.0392	0.9162	-0.0391	0.9027	-0.2019	0.9467	-1.2867	0.1512	-1.0848	0.2561
dmrt11E	doublesex-Mab re	1624940_at	0.0222	0.9010	0.0167	0.9017	-0.2386	0.1731	-0.1010	0.8251	0.0438	0.8488	0.1448	0.3773	0.0641	0.9342	-0.0664	0.8450	-0.1304	0.6294
CG30169	CG30169	1624941_at	0.3181	0.2315	0.1537	0.5804	-0.1825	0.2248	-0.2154	0.5948	0.2807	0.1693	0.4960	0.0149	0.1353	0.8692	0.1707	0.6312	0.0354	0.9387
---	---	1624942_at	0.0821	0.6044	0.0185	0.8599	-0.0493	0.8137	0.0084	0.9921	0.1059	0.5706	0.0975	0.5695	-0.0009	0.9997	-0.0328	0.9231	-0.0319	0.9161
CG1636	CG1636	1624943_at	-0.1389	0.7915	1.1298	0.0946	1.0273	0.0022	-0.0307	0.9894	-0.6179	0.2492	-0.5872	0.2199	0.2544	0.8553	0.6932	0.2116	0.4389	0.4582
---	---	1624944_at	0.1368	0.4333	0.0872	0.3938	0.1259	0.6125	-0.0006	0.9994	-0.0329	0.9026	-0.0322	0.8899	-0.0878	0.8867	-0.0068	0.9890	0.0810	0.7622
CG10910	CG10910	1624945_a_at	0.0942	0.6803	-0.0516	0.8399	-1.5465	0.0061	-1.4826	0.0657	0.1912	0.7234	1.6738	0.0024	0.0040	0.9984	-0.0555	0.9326	-0.0595	0.9168
CG6790	CG6790	1624946_at	0.1289	0.5571	-0.1078	0.4551	-0.0380	0.8453	0.0803	0.9061	0.1255	0.6016	0.0452	0.8599	-0.0158	0.9913	-0.1752	0.5726	-0.1593	0.6154
---	---	1624947_x_at	0.4254	0.1828	0.2047	0.1559	0.2808	0.1265	-0.0928	0.8798	-0.1553	0.4801	-0.0625	0.7907	-0.0538	0.9545	-0.1370	0.6483	-0.0832	0.8056
CG12948	CG12948	1624948_at	-0.5054	0.0843	-0.6110	0.0281	-0.8267	0.0135	-0.1137	0.7432	0.2464	0.1058	0.3602	0.0165	-0.0363	0.9862	0.0572	0.9425	0.0936	0.8897
CG31546	CG31546	1624949_at	-0.0008	0.9965	0.0172	0.9591	0.2346	0.3762	0.1230	0.8589	0.1312	0.6377	0.0082	0.9790	-0.0123	0.9950	0.0251	0.9739	0.0374	0.9525
CG14823	CG14823	1624950_a_at	2.3991	0.0015	0.7025	0.2881	2.0838	0.0004	0.2120	0.8463	-0.0736	0.8924	-0.2856	0.4511	-1.1502	0.3660	-1.5608	0.0423	-0.4105	0.5676
CG13862	CG13862	1624951_at	0.1717	0.3715	-0.0039	0.9781	0.0604	0.7665	-0.0147	0.9863	0.0368	0.8965	0.0514	0.8276	-0.0667	0.9345	0.0045	0.9940	0.0712	0.8287
Gr47b	Gustatory recepto	1624952_at	-0.0061	0.9812	0.0067	0.9522	-0.0098	0.9797	-0.1314	0.8545	-0.1706	0.5381	-0.0393	0.8997	0.1279	0.9055	0.1102	0.8292	-0.0177	0.9769

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Cyp12c1	Cyp12c1	1624953_at	1.8531	0.0324	0.5302	0.5461	0.8699	0.0142	-0.2186	0.7949	-0.0954	0.8246	0.1232	0.7384	-0.4770	0.8732	-1.1724	0.3053	-0.6954	0.5789
---	---	1624954_at	0.0571	0.7505	-0.1495	0.3155	-0.0175	0.9276	0.0987	0.8405	0.1228	0.5293	0.0241	0.9142	-0.0993	0.8886	-0.2114	0.4280	-0.1121	0.7096
---	---	1624955_at	0.2160	0.3485	-0.0661	0.7097	-0.1873	0.3267	0.0668	0.9345	0.4599	0.0587	0.3931	0.0688	0.1421	0.8202	0.0708	0.8386	-0.0713	0.8256
tst	twister	1624956_at	-0.0974	0.6351	0.0966	0.5032	0.2285	0.2639	-0.0779	0.9356	-0.2117	0.4761	-0.1338	0.6477	-0.2847	0.6749	-0.1173	0.7056	0.1673	0.5653
Tequila	Tequila	1624957_a_at	2.9117	0.0020	1.1140	0.0816	2.7549	0.0000	0.6703	0.4706	0.8819	0.0797	0.2116	0.6870	-0.8712	0.4443	-0.8656	0.1403	0.0056	0.9956
CG8851	CG8851	1624958_a_at	0.0243	0.9177	0.0101	0.9447	0.0909	0.6631	0.0753	0.9182	0.0664	0.8116	-0.0090	0.9745	0.0435	0.9598	0.0615	0.8551	0.0179	0.9598
---	---	1624959_at	0.0084	0.9739	0.1222	0.2415	0.1333	0.3994	-0.1675	0.6421	-0.1224	0.5097	0.0451	0.8212	-0.0261	0.9800	0.0223	0.9522	0.0484	0.8779
mam	mastermind	1624960_at	0.1282	0.5096	0.4315	0.1354	0.1246	0.6775	-0.1687	0.6908	-0.1355	0.5153	0.0332	0.8872	0.1313	0.9168	0.2100	0.6605	0.0787	0.8949
CG6380	CG6380	1624961_at	0.1316	0.5027	0.0211	0.9186	0.0014	0.9945	-0.0078	0.9937	-0.1070	0.6052	-0.0992	0.6004	0.0677	0.9426	-0.0005	0.9998	-0.0682	0.8570
star1	Drosophila allatos	1624962_at	0.0726	0.8097	0.0267	0.8321	0.1461	0.4457	-0.0158	0.9852	-0.2922	0.1273	-0.2764	0.1070	-0.0665	0.9451	-0.3107	0.2714	-0.2442	0.4114
CG17802	CG17802	1624963_at	0.1479	0.5981	-0.2034	0.4637	-0.4187	0.0180	-0.3284	0.6506	0.5368	0.1237	0.8653	0.0136	-0.1242	0.8814	0.1761	0.6028	0.3003	0.3572
CG17105	CG17105	1624964_at	0.0610	0.8002	-0.0401	0.8844	0.0669	0.7193	0.0170	0.9838	0.0307	0.9102	0.0137	0.9565	-0.0661	0.9357	-0.0813	0.8092	-0.0152	0.9697
---	---	1624965_at	0.0854	0.6471	0.0974	0.4242	0.0693	0.6537	-0.1372	0.6984	-0.1412	0.3974	-0.0040	0.9849	0.0135	0.9894	-0.0533	0.8550	-0.0668	0.7944
CG7423	CG7423	1624966_at	0.2349	0.3390	0.4601	0.0405	0.4110	0.0353	-0.0953	0.8738	-0.2893	0.1639	-0.1940	0.3065	0.1025	0.8331	0.1527	0.4531	0.0502	0.8472
CG1530	CG1530	1624967_at	-0.0657	0.6849	0.0911	0.3917	0.0413	0.8363	-0.1511	0.6506	-0.0535	0.7844	0.0976	0.5345	0.0761	0.8909	0.1182	0.5962	0.0421	0.8834
CG16899	CG16899	1624968_at	0.1329	0.4605	0.0456	0.7020	0.1423	0.4011	0.0421	0.9380	0.0664	0.7194	0.0243	0.8991	-0.0176	0.9862	0.0781	0.7818	0.0957	0.7097
CG17041	CG17041	1624969_s_at	-2.3887	0.0206	-1.4812	0.0411	-2.1964	0.0002	-0.7905	0.1495	-2.1860	0.0004	-1.3955	0.0013	0.0136	0.9978	-1.1324	0.3053	-1.1460	0.3243
gft	Cullin3	1624970_s_at	0.1892	0.4927	0.7199	0.0667	0.5785	0.0061	-0.0577	0.9461	-0.2367	0.3331	-0.1790	0.4284	0.0170	0.9924	0.3968	0.3261	0.3799	0.3723
CG5254	CG5254	1624971_at	-0.7253	0.0833	-0.4270	0.4879	-0.3798	0.3462	-0.4465	0.6013	-1.8217	0.0021	-1.3751	0.0042	-0.4292	0.8202	-1.5568	0.0655	-1.1276	0.1701
---	---	1624972_at	-0.1044	0.5413	-0.1539	0.3407	-0.2488	0.1549	0.0336	0.9518	0.0656	0.7130	0.0319	0.8589	-0.0675	0.9142	-0.0568	0.8500	0.0107	0.9757
---	---	1624973_at	0.1205	0.5312	-0.4279	0.0568	0.0131	0.9539	0.1852	0.5969	0.2894	0.1020	0.1042	0.5491	-0.1304	0.8202	-0.1632	0.5061	-0.0328	0.9198
---	---	1624974_at	-0.1093	0.4613	0.0243	0.8152	-0.1631	0.3923	-0.0650	0.9149	0.0212	0.9382	0.0862	0.6630	0.0088	0.9922	0.0023	0.9961	-0.0065	0.9847
CG17028	CG17028	1624975_at	0.6909	0.0304	1.0829	0.0106	1.2221	0.0022	-0.1376	0.8977	-0.5094	0.1458	-0.3717	0.2383	-0.0678	0.9592	0.0453	0.9409	0.1131	0.8108
CG32026	CG32026	1624976_at	0.0105	0.9599	0.2475	0.4870	0.1686	0.4704	0.0411	0.9705	-0.0656	0.8612	-0.1067	0.7267	0.0499	0.9773	0.1241	0.8165	0.0742	0.8949
---	---	1624977_at	0.0557	0.7479	0.0231	0.8757	0.0075	0.9723	0.2124	0.4979	0.0206	0.9297	-0.1918	0.1952	0.1101	0.8270	-0.1412	0.5172	-0.2513	0.2501
CG33309	CG33309	1624978_at	0.0956	0.7384	0.0691	0.7647	0.0801	0.5825	0.1330	0.6615	0.0732	0.6563	-0.0598	0.6962	0.0257	0.9862	-0.0156	0.9802	-0.0414	0.9316
CG18594 /// DyakCG18594	CG18594	1624979_at	0.2963	0.7553	-0.3085	0.2013	0.5765	0.1116	1.0252	0.0899	0.6794	0.0588	-0.3458	0.2789	0.0004	0.9999	-0.1826	0.9111	-0.1830	0.9020
---	---	1624980_at	-0.0452	0.8450	-0.1257	0.3413	-0.2948	0.1047	-0.0307	0.9558	0.2322	0.1162	0.2628	0.0521	-0.0051	0.9978	-0.0566	0.9313	-0.0515	0.9282
CG10576	CG10576	1624981_a_at	0.0135	0.9782	1.4638	0.0310	1.8327	0.0000	0.3159	0.3473	-0.8778	0.0015	-1.1937	0.0003	-0.2018	0.9011	0.4654	0.4336	0.6672	0.2784
CG5080	CG5080	1624982_s_at	-1.6716	0.0010	-1.8247	0.1395	-1.0617	0.0025	0.0411	0.9675	-1.0152	0.0032	-1.0563	0.0016	-0.6975	0.7726	-0.9261	0.3623	-0.2286	0.8680
RpS18	transcription unit E	1624983_s_at	0.5777	0.0084	0.8075	0.0372	0.8342	0.0154	0.2360	0.7328	-0.2267	0.4829	-0.4627	0.0934	0.3101	0.7230	-0.0306	0.9617	-0.3406	0.3808
CG8021	CG8021	1624984_at	-0.5614	0.1004	-0.0254	0.9640	0.2969	0.3225	-0.0515	0.9666	-0.7157	0.0340	-0.6642	0.0293	-0.3259	0.8205	-0.0797	0.9339	0.2461	0.7149
CG7886	CG7886	1624985_at	0.1579	0.3097	-0.1343	0.5091	-0.0675	0.7387	0.0849	0.8990	0.2345	0.2872	0.1497	0.4718	0.0945	0.8901	0.0312	0.9402	-0.0632	0.8474
CG9836	CG9836	1624986_at	-0.1336	0.5085	-0.1404	0.4733	0.0344	0.8765	-0.0935	0.8362	-0.3254	0.0607	-0.2319	0.1272	-0.2361	0.6955	-0.2708	0.2587	-0.0347	0.9194
CG3345	CG3345	1624987_at	0.0767	0.7151	0.0935	0.5799	0.0483	0.7789	-0.0949	0.8698	0.0775	0.7478	0.1724	0.3555	0.0415	0.9609	0.1551	0.5259	0.1136	0.6574
---	---	1624988_at	-0.0994	0.5188	0.0729	0.6579	0.0581	0.8347	0.0990	0.8288	0.1900	0.2829	0.0910	0.6083	0.0192	0.9914	0.1133	0.8185	0.0941	0.8467
Ef1alpha100E	elongation factor 1	1624989_s_at	-1.8032	0.0123	-1.8967	0.0118	-2.4644	0.0001	-0.5949	0.0737	-0.8067	0.0021	-0.2118	0.2177	0.0576	0.9870	-0.7337	0.3565	-0.7913	0.3437
CG12063	CG12063	1624990_at	-0.3368	0.2539	-0.5557	0.0815	-0.5786	0.0170	0.3930	0.3433	0.3536	0.1374	-0.0394	0.8906	0.0978	0.9165	-0.0888	0.8376	-0.1866	0.5872
Tim13	Tim13	1624991_at	-0.1137	0.4507	0.2084	0.1522	0.3420	0.0498	0.1819	0.6360	-0.2390	0.2039	-0.4209	0.0209	-0.0314	0.9726	0.0208	0.9557	0.0522	0.8635
---	---	1624992_at	0.2956	0.2257	0.0177	0.9176	-0.2851	0.0620	-0.0786	0.9311	0.1022	0.7478	0.1808	0.4787	-0.0099	0.9918	-0.1148	0.6382	-0.1049	0.6686
CG4377	CG4377	1624993_at	0.1085	0.8535	-0.0076	0.9460	0.7233	0.0584	-0.0049	0.9956	-0.0967	0.7640	-0.0919	0.7540	-0.6082	0.7588	-0.1432	0.9110	0.4651	0.6033
dup	Double-parked	1624994_at	0.4697	0.1310	-0.2482	0.6980	-0.4024	0.2717	-0.1997	0.6802	0.4986	0.0346	0.6983	0.0049	-0.0620	0.9868	-0.1867	0.8786	-0.1247	0.9161
CG12355	CG12355	1624995_at	-0.7496	0.3598	0.0929	0.3916	-0.3766	0.2102	-0.3888	0.6901	-1.4659	0.0066	-1.0771	0.0157	0.2016	0.9445	-0.6325	0.4793	-0.8341	0.3546
CG3368	CG3368	1624996_at	0.1463	0.4897	0.2697	0.3228	0.1856	0.1932	0.0790	0.8707	-0.1390	0.4293	-0.2181	0.1500	0.1523	0.8692	0.0435	0.9409	-0.1089	0.8088
5-Sep	septin	1624997_s_at	-0.1799	0.5576	-0.1203	0.5347	-0.1476	0.6973	-0.1321	0.8143	0.0213	0.9492	0.1534	0.4659	-0.3961	0.7230	-0.0799	0.9150	0.3162	0.5334
---	---	1624998_at	0.1947	0.4215	0.5906	0.1934	0.1561	0.4417	-0.1113	0.8822	0.1666	0.5460	0.2779	0.2261	0.3893	0.7644	0.6806	0.2010	0.2914	0.6228
---	---	1624999_s_at	0.0270	0.8933	0.0200	0.8796	0.1828	0.2870	0.0686	0.9036	-0.1176	0.5521	-0.1862	0.2599	0.0004	0.9999	0.0112	0.9829	0.0109	0.9812
CG32441	CG32441	1625000_a_at	-0.6639	0.1052	0.0165	0.9254	0.5708	0.0480	0.0362	0.9530	-0.7110	0.0021	-0.7472	0.0010	-0.5393	0.7188	-0.0385	0.9702	0.5008	0.4322
---	---	1625001_at	0.2815	0.1356	0.1326	0.4071	0.0368	0.8832	0.0318	0.9773	0.0357	0.9249	0.0039	0.9907	0.0226	0.9860	-0.1556	0.6132	-0.1782	0.5587
CG10038 /// DyakCG10038	CG10038	1625002_a_at	0.9273	0.0048	0.3506	0.2554	0.4882	0.0085	0.0204	0.9757	0.5046	0.0085	0.4842	0.0061	-0.1719	0.8655	0.0169	0.9828	0.1888	0.6694
ETHR	ETHR	1625003_at	0.1820	0.3075	0.3554	0.0912	0.2017	0.2466	-0.2844	0.3625	-0.1904	0.2886	0.0940	0.5988	-0.0922	0.8745	0.0754	0.7941	0.1676	0.4820
CG18811	CG18811	1625004_at	-0.1369	0.7482	0.2492	0.3222	0.3364	0.0828	0.0068	0.9940	-0.0730	0.7380	-0.0798	0.6805	-0.1170	0.9489	0.2883	0.6237	0.4053	0.4730
Ant2	Adenine nucleotid	1625005_s_at	0.3502	0.2119	-0.4666	0.0933	-0.4619	0.0141	0.1027	0.8572	0.7123	0.0048	0.6095	0.0058	0.0731	0.9335	-0.0788	0.8329	-0.1518	0.6150
Tango4	Transport and Gol	1625006_at	-0.5342	0.0504	-0.0891	0.8607	0.0528	0.8701	0.0165	0.9832	-0.2616	0.1423	-0.2782	0.0833	0.0232	0.9943	0.1056	0.9291	0.0823	0.9374
---	---	1625007_a_at	-0.1429	0.3805	-0.3697	0.1761	-0.4508	0.0246	0.1322	0.7167	0.1878	0.2591	0.0556	0.7592	0.0763	0.9390	-0.0009	0.9996	-0.0771	0.8444
CG33330	CG33330	1625008_at	-0.5836	0.0991	0.2267	0.6885	0.4205	0.0445	-0.2333	0.7335	-0.8216	0.0151	-0.5883	0.0378	-0.2968	0.8215	0.0833	0.9224	0.3801	0.5032
---	---	1625009_at	-0.1690	0.3284	-0.5537	0.0710	-0.4057	0.3941	-0.0637	0.9760	0.3971	0.4661	0.4608	0.3340	-0.2937	0.7500	0.0715	0.9058	0.3652	0.3671
Gr97a	Gustatory recepto	1625010_a_at	0.2484	0.1774	0.0516	0.6346	-0.0007	0.9977	0.0248	0.9817	0.2364	0.3659	0.2116	0.3705	0.0136	0.9901	0.0246	0.9499	0.0110	0.9769
CG11206	CG11206	1625011_at	-1.7185	0.0035	-1.1224	0.0372	-1.5656	0.0007	-0.4786	0.2939	-0.8055	0.0093	-0.3270	0.1714	0.0070	0.9978	-0.2349	0.7340	-0.2419	0.7184
vir-1	virus-induced 1	1625012_s_at	-1.6961	0.0347	-1.7250	0.0172	-1.9688	0.0092	0.2473	0.8315	1.3088	0.0100	1.0614	0.0155	0.4659	0.8875	1.3313	0.2627	0.8654	0.4978
---	---	1625013_at	-0.0121	0.9732	-0.0002	1.0000	0.4183	0.0413	0.2093	0.5054	0.0219	0.9252	-0.1874	0.2083	-0.1102	0.8940	0.0035	0.9961	0.1137	0.7484
Ilk	integrin-linked kin	1625014_at	-0.7346	0.0094	-0.2609	0.2962	-0.0975	0.6550	-0.0004	0.9994	-0.1573	0.2742	-0.1569	0.2211	-0.1245	0.9204	0.1794	0.7058	0.3039	0.4851
mthl12	Mth-like 12	1625015_at	0.1777	0.2868	-0.0687	0.6248	0.1138	0.5054	-0.0067	0.9937	0.1943	0.2805	0.2010	0.2093	-0.2252	0.6483	-0.2089	0.2735	0.0163	0.9533
CG31516	CG31516	1625016_at	0.0530	0.8968	0.0157	0.9642	0.0507	0.8070	0.2271	0.7293	0.4139	0.1569	0.1868	0.5098	0.1331	0.8166	0.1983	0.3972	0.0652	0.8256
CG3732	CG3732	1625017_at	-0.4775	0.0470	0.2173	0.2192	0.5325	0.1303	-0.0186	0.9860	-0.2490	0.3111	-0.2304	0.2963	-0.3140	0.7953	0.4316	0.3909	0.7456	0.1670
CG32827	CG32827	1625018_at	0.2889	0.4302	0.1502	0.8938	0.2981	0.3082	-0.0070	0.9956	0.5147	0.1073	0.5217	0.0711	-0.0335	0.9941	0.4812	0.6780	0.5147	0.6469
angel	angel	1625019_at	-0.3796	0.0493	-0.3673	0.1524	-0.5383	0.0064	-0.0326	0.9558	-0.1000	0.5606	-0.0674	0.6869	0.1621	0.7981	0.0324	0.9402	-0.1297	0.6491
CG13989	CG13989	1625020_at	0.1901	0.3255	0.0061	0.9828	0.3299	0.0929	0.0630	0.9060	0.0822	0.6755	0.0192	0.9274	-0.1141	0.8940	-0.0607	0.8995	0.0534	0.9031
---	---	1625021_at	0.0652	0.6601	-0.1328	0.4209	0.0291	0.8821	0.0190	0.9777	0.0854	0.6612	0.0664	0.7171	-0.0423	0.9635	-0.0117	0.9797	0.0307	0.9310

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG6125	CG6125	1625022_a_at	-0.6001	0.7896	-0.9867	0.2061	-2.3970	0.0005	-1.0451	0.6122	-1.1396	0.2728	-0.0945	0.9416	0.0529	0.9959	-1.6746	0.4843	-1.7274	0.4756
nAcRbeta-21C	Dbeta3	1625023_a_at	-3.0314	0.0041	-3.7564	0.0029	-3.9361	0.0000	-0.2025	0.6672	-0.8493	0.0033	-0.6468	0.0067	0.0511	0.9893	-1.4861	0.0756	-1.5373	0.0876
zwilch	zwilch	1625024_at	-0.7715	0.0194	-1.3690	0.1454	-1.8459	0.0019	-0.2811	0.6015	0.6199	0.0300	0.9010	0.0035	-0.0976	0.9831	0.0735	0.9670	0.1711	0.9089
CG31286	CG31286	1625025_a_at	0.0619	0.8054	0.1085	0.4637	-0.0048	0.9846	-0.0233	0.9755	-0.0521	0.8218	-0.0287	0.8974	-0.0846	0.9112	-0.0379	0.9291	0.0467	0.9004
CG3348	CG3348	1625026_at	-0.0603	0.9528	-0.6472	0.2926	-1.0299	0.0284	-0.1768	0.9554	1.5580	0.0703	1.7347	0.0303	0.2557	0.9057	0.8637	0.2488	0.6080	0.4463
GXIVsPLA2	GXIVsPLA2	1625027_a_at	-0.9416	0.0039	0.3951	0.1544	0.9776	0.0006	-0.1355	0.7327	-1.2202	0.0003	-1.0847	0.0003	-0.6822	0.1953	0.2384	0.3793	0.9206	0.0320
---	---	1625028_at	0.0976	0.6778	0.0402	0.7185	-0.0969	0.5975	-0.0129	0.9874	0.1170	0.5832	0.1299	0.4897	0.0941	0.8653	0.1172	0.6216	0.0231	0.9407
---	---	1625029_s_at	0.0022	0.9948	-0.0032	0.9796	-0.0789	0.7129	0.0574	0.9465	0.0384	0.9089	-0.0190	0.9503	0.0415	0.9635	-0.1168	0.6673	-0.1583	0.5430
CG6751	CG6751	1625030_at	0.4767	0.0648	-0.1229	0.6367	-0.0451	0.8026	0.2929	0.5220	0.8563	0.0043	0.5634	0.0168	0.0974	0.8999	0.1165	0.7202	0.0191	0.9637
CG7841	CG7841	1625031_at	-0.4008	0.3600	0.2510	0.4667	0.6637	0.1494	-0.3399	0.4679	-1.3972	0.0007	-1.0574	0.0013	-0.6410	0.7464	-0.6000	0.4861	0.0410	0.9757
mod(r)	modifier of rudime	1625032_s_at	-0.3390	0.2247	0.4966	0.0281	0.5709	0.0671	0.0027	0.9985	-0.7423	0.0236	-0.7450	0.0146	-0.0854	0.9306	0.1272	0.7279	0.2126	0.5206
CG16791	CG16791	1625033_at	-0.3766	0.3101	-0.2762	0.5492	-0.1330	0.4746	0.1304	0.8299	0.2292	0.3315	0.0988	0.6852	0.0875	0.9717	0.1716	0.8353	0.0840	0.9211
tacc	transforming acidi	1625034_s_at	-0.3567	0.0824	-0.2575	0.3526	-0.2231	0.1459	0.1120	0.8512	-0.2788	0.1982	-0.3908	0.0496	0.0070	0.9959	-0.0437	0.9288	-0.0507	0.9057
insc	Inscutable	1625035_at	0.5873	0.2340	-0.3522	0.5582	0.6330	0.1430	0.6600	0.2077	0.8053	0.0188	0.1454	0.6553	-0.4062	0.8586	-0.1881	0.8886	0.2182	0.8559
CG18281	CG18281	1625036_at	-0.0703	0.6866	0.1567	0.3219	0.1906	0.4355	0.0740	0.8723	-0.0158	0.9472	-0.0898	0.5724	-0.0005	0.9999	0.2985	0.4011	0.2990	0.4114
---	---	1625037_at	-0.0193	0.9440	0.1251	0.3823	0.1006	0.5652	-0.2080	0.6010	-0.1893	0.3537	0.0187	0.9400	-0.0137	0.9862	0.0083	0.9808	0.0220	0.9330
lok	maternal nuclear l	1625038_s_at	-0.4223	0.2071	-0.7283	0.1810	-0.3719	0.3344	0.0061	0.9956	-0.3149	0.2253	-0.3210	0.1663	-0.3405	0.8744	-0.5784	0.5075	-0.2380	0.8221
CG13582	CG13582	1625039_at	0.0666	0.7348	0.0328	0.7771	0.1934	0.4077	0.0773	0.8676	-0.0321	0.8842	-0.1094	0.4849	-0.0486	0.9742	-0.0281	0.9644	0.0205	0.9721
CG13502	CG13502	1625040_a_at	0.0099	0.9645	0.0019	0.9892	0.1139	0.5548	0.0159	0.9845	-0.0142	0.9593	-0.0301	0.8928	0.0528	0.9391	0.0961	0.6873	0.0432	0.8818
Oatp74D	Organic anion trar	1625041_at	0.4768	0.2579	2.3713	0.0476	2.1445	0.0017	-1.0321	0.1882	-2.3644	0.0011	-1.3322	0.0076	-0.8830	0.6960	-0.5418	0.5959	0.3412	0.7577
CG31288	CG31288	1625042_at	2.0244	0.1261	2.4923	0.0031	2.4781	0.0010	0.4087	0.9194	0.5139	0.7215	0.1052	0.9464	0.3792	0.7644	1.0889	0.0637	0.7096	0.2073
CG12436	CG12436	1625043_at	0.1947	0.3678	0.2214	0.1533	0.1443	0.4142	-0.2682	0.4459	-0.2921	0.1269	-0.0239	0.9193	-0.0935	0.8222	-0.0903	0.6288	0.0033	0.9914
CG8891	CG8891	1625044_at	-0.2205	0.4404	0.0417	0.8428	0.2066	0.4477	0.1634	0.6845	-0.0015	0.9955	-0.1649	0.3344	0.0683	0.9717	0.3455	0.4791	0.2772	0.5866
---	---	1625045_at	0.0703	0.7328	0.1727	0.2267	0.2289	0.1967	-0.1208	0.7880	-0.0369	0.8810	0.0839	0.6612	0.0317	0.9676	0.1496	0.4699	0.1179	0.5874
CG10648	CG10648	1625046_at	-0.3826	0.1983	-0.2327	0.4856	0.3094	0.2986	0.4235	0.4232	0.1960	0.5293	-0.2274	0.4036	-0.3119	0.7677	0.1837	0.7200	0.4956	0.2833
qkr58E-2	quaking related 5(	1625047_at	-0.1737	0.6109	-0.0680	0.8960	0.0795	0.6301	-0.0138	0.9854	-0.2235	0.1852	-0.2097	0.1658	-0.0717	0.9769	0.0129	0.9925	0.0846	0.9171
Nf1	neurofibromatosis	1625048_at	-0.9728	0.0191	-0.8084	0.1584	-1.0121	0.0012	-0.0831	0.9127	-0.1911	0.4419	-0.1080	0.6636	0.1472	0.9235	0.0292	0.9726	-0.1180	0.8577
CG5480	CG5480	1625049_at	-0.0672	0.8344	0.0043	0.9727	0.0019	0.9943	0.0556	0.9643	0.0546	0.9045	-0.0010	0.9981	0.1159	0.7997	0.0937	0.6512	-0.0222	0.9342
---	---	1625050_s_at	-0.5632	0.6778	-1.3462	0.0144	-0.0923	0.6600	1.4381	0.4001	0.8452	0.3890	-0.5930	0.5242	0.1871	0.8049	-0.2035	0.5333	-0.3906	0.2291
4EHP	4EHP	1625051_at	2.1323	0.0044	0.8879	0.0919	-0.3350	0.6144	-0.2628	0.6954	2.0241	0.0004	2.2869	0.0001	1.0040	0.7149	1.0239	0.3635	0.0199	0.9924
CG5955	CG5955	1625052_at	-0.5808	0.0665	0.1002	0.8379	0.6218	0.1449	0.1611	0.5720	-0.2078	0.1517	-0.3689	0.0122	-0.3327	0.8837	0.5145	0.5722	0.8472	0.3404
CG16898	CG16898	1625053_at	3.2676	0.0452	0.7123	0.6518	2.6102	0.0003	0.7930	0.8192	-0.5842	0.7101	-1.3772	0.2603	-1.1730	0.7614	-3.1313	0.0694	-1.9582	0.2409
---	---	1625054_at	0.1433	0.3264	0.1215	0.6156	0.3153	0.1074	-0.0242	0.9712	-0.2027	0.2331	-0.1785	0.2424	-0.1822	0.8042	-0.1132	0.7559	0.0689	0.8643
Ost48	Oligosaccharyltrar	1625055_at	0.7004	0.0305	0.5373	0.0201	0.7941	0.0017	0.1525	0.6775	0.6655	0.0030	0.5130	0.0058	-0.1107	0.9137	0.6224	0.0912	0.7331	0.0767
---	---	1625056_at	0.1796	0.2708	0.0650	0.7205	-0.0162	0.9421	0.0118	0.9882	0.0056	0.9838	-0.0062	0.9787	0.1092	0.8465	0.0518	0.8775	-0.0574	0.8491
RSG7	Regulator of G-pr	1625057_at	-0.0083	0.9715	0.0659	0.5423	0.3803	0.1063	0.0117	0.9872	-0.0335	0.8860	-0.0452	0.8181	-0.2094	0.8215	0.2442	0.5439	0.4536	0.2557
yellow-d	yellow-d	1625058_at	1.7084	0.0011	0.9480	0.0134	2.5385	0.0000	1.2774	0.0096	0.7214	0.0095	-0.5560	0.0180	-0.0252	0.9742	-0.0102	0.9763	0.0150	0.9577
Leucokinin	drosokinin	1625059_at	0.6688	0.0582	-0.0117	0.9712	0.1447	0.4571	0.0500	0.9621	0.1598	0.6023	0.1097	0.7121	-0.1146	0.9161	-0.4625	0.2063	-0.3478	0.3699
CG7974	CG7974	1625060_at	-0.2886	0.1970	0.2303	0.4141	0.4344	0.0927	-0.0494	0.9620	-0.0428	0.9105	0.0066	0.9850	-0.1810	0.8439	0.4566	0.2123	0.6376	0.1240
CG17298	CG17298	1625061_at	0.3415	0.1886	0.0388	0.7054	0.2636	0.1267	-0.0154	0.9903	0.1295	0.6882	0.1449	0.6111	-0.1797	0.7464	-0.1572	0.5198	0.0225	0.9460
CG32806	CG32806	1625062_a_at	0.0901	0.5368	0.0169	0.8678	0.2993	0.1254	0.0566	0.9254	0.1194	0.5403	0.0628	0.7511	-0.2435	0.6868	-0.1421	0.5872	0.1015	0.7154
CG31606	CG31606	1625063_a_at	0.4938	0.1671	0.1922	0.4979	-0.0291	0.8589	-0.1221	0.7530	0.0078	0.9762	0.1299	0.4066	0.0356	0.9841	-0.1030	0.8546	-0.1386	0.7728
CG34400	CG17368	1625064_at	0.1929	0.2368	0.0213	0.8773	0.3294	0.1148	0.2060	0.5633	0.0966	0.6371	-0.1094	0.5407	-0.0641	0.9016	-0.0902	0.6673	-0.0261	0.9205
SK	small conductanc	1625065_s_at	-3.0207	0.0019	-5.0087	0.0051	-5.2754	0.0001	0.2024	0.9017	1.4565	0.0140	1.2541	0.0165	0.4983	0.8395	-0.2389	0.8665	-0.7372	0.4815
CG13230	CG13230	1625066_at	0.3198	0.1033	-0.2088	0.4612	0.2677	0.1544	0.1662	0.7693	0.2638	0.2742	0.0976	0.6995	-0.3270	0.6264	-0.3230	0.2306	0.0039	0.9936
CG32260	CG32260	1625067_at	0.3450	0.1319	0.2869	0.3293	0.2162	0.2192	-0.0330	0.9711	0.0064	0.9854	0.0393	0.8862	-0.0444	0.9717	0.0276	0.9583	0.0719	0.8626
Rad23	Rad23	1625068_a_at	0.3285	0.0627	0.2373	0.3495	0.0515	0.7538	-0.0380	0.9470	0.4452	0.0137	0.4832	0.0057	0.2556	0.7070	0.4001	0.1498	0.1445	0.6385
---	---	1625069_at	0.0662	0.7497	0.0120	0.9122	-0.0210	0.9041	0.0052	0.9951	0.0209	0.9298	0.0157	0.9404	0.0841	0.8692	0.0163	0.9620	-0.0679	0.7788
---	---	1625070_at	0.3380	0.3617	-0.0230	0.9429	0.2022	0.3967	0.2276	0.7531	0.1947	0.5647	-0.0329	0.9319	-0.1120	0.8940	-0.1638	0.6280	-0.0518	0.9055
Mlc2	myosin regulatory	1625071_a_at	-2.3311	0.0005	-2.6267	0.0363	-3.9189	0.0003	-0.6797	0.0660	-0.5906	0.0135	0.0891	0.6914	0.5525	0.8493	-0.7595	0.5469	-1.3120	0.2921
Ac3	Ac3	1625072_s_at	-0.1989	0.2757	-0.1675	0.2668	-0.0038	0.9859	0.0235	0.9675	0.0668	0.7023	0.0433	0.7991	-0.1452	0.7644	-0.1015	0.6483	0.0437	0.8745
---	---	1625073_at	0.1644	0.3112	0.0000	1.0000	-0.0364	0.8658	-0.1494	0.7599	0.0963	0.6889	0.2457	0.1896	-0.0027	0.9976	-0.0551	0.8287	-0.0524	0.8287
CG12486	CG12486	1625074_at	-0.7630	0.0722	-0.5598	0.1463	-1.4694	0.0004	-0.7727	0.1526	-0.2538	0.4559	0.5189	0.0774	0.0729	0.9571	-0.2936	0.4470	-0.3665	0.3552
Nep2	Neprilysin 2	1625075_at	-0.6314	0.0411	-1.3133	0.0074	-0.7092	0.0110	0.1432	0.7959	0.2410	0.2888	0.0977	0.6783	-0.3949	0.6749	-0.3891	0.2973	0.0058	0.9931
aret	bruno	1625076_at	1.6878	0.1465	-1.1937	0.5437	-1.1951	0.0004	-0.6880	0.6197	1.8310	0.0161	2.5190	0.0023	-0.7253	0.8882	-0.9275	0.6614	-0.2022	0.9406
CG30265	CG30265	1625077_at	0.0866	0.6489	0.0940	0.6398	-0.1103	0.4915	-0.2335	0.4612	-0.0077	0.9773	0.2259	0.1399	-0.0155	0.9885	-0.0442	0.9009	-0.0288	0.9300
eIF-3p40	Eukaryotic initiatio	1625078_at	0.5494	0.0255	1.0464	0.0169	1.0881	0.0064	0.3028	0.6010	-0.3782	0.1923	-0.6809	0.0171	0.2914	0.7493	0.1572	0.7267	-0.1342	0.7681
CG34126	CG34126	1625079_at	0.1250	0.4968	0.4109	0.0726	0.4598	0.0556	0.0141	0.9860	0.0429	0.8658	0.0288	0.9008	-0.1375	0.8504	0.0752	0.8563	0.2127	0.4970
tral	trailer hitch	1625080_at	0.6315	0.1559	0.5386	0.2176	-0.1000	0.6515	-0.4453	0.2620	0.6294	0.0154	1.0747	0.0009	0.2367	0.8940	0.5445	0.4093	0.3078	0.6719
CG17260	CG17260	1625081_at	0.0873	0.7434	-0.4485	0.2623	-0.1834	0.2439	0.2757	0.3433	0.5065	0.0091	0.2309	0.1219	0.0014	0.9998	0.0394	0.9567	0.0380	0.9514
CG7882	CG7882	1625082_a_at	-0.4613	0.5286	-0.2274	0.2668	-0.2661	0.1541	0.1135	0.9603	-0.4579	0.4736	-0.5714	0.3028	-0.0731	0.9689	-0.2369	0.6605	-0.1637	0.7774
rhi	rhino	1625083_at	0.1930	0.5639	-0.8324	0.2477	-0.8744	0.1221	-0.6147	0.2121	1.4373	0.0012	2.0519	0.0002	-0.7428	0.8465	0.2191	0.9312	0.9619	0.5681
---	---	1625084_at	0.0592	0.7954	0.1113	0.4558	0.0068	0.9835	0.0577	0.9319	0.0982	0.6677	0.0405	0.8626	0.0336	0.9742	0.1040	0.7218	0.0704	0.8255
CG30182	CG30182	1625085_at	-0.0086	0.9599	0.0089	0.9372	0.0459	0.7955	0.0246	0.9641	-0.0679	0.6827	-0.0925	0.5103	0.1079	0.8610	0.0912	0.7612	-0.0167	0.9638
CG17450 /// CG32819 /// C	CG17450 /// CG3	1625086_s_at	0.1486	0.4042	0.2519	0.0798	0.0709	0.7561	0.0084	0.9932	-0.2370	0.2216	-0.2455	0.1566	0.1506	0.7823	-0.0319	0.9326	-0.1825	0.4541
CG32137	CG32137	1625087_a_at	-0.1233	0.5379	0.0400	0.8557	-0.0481	0.8522	0.2032	0.5363	0.1925	0.2643	-0.0107	0.9600	0.2219	0.8202	0.3876	0.3259	0.1656	0.7170
CG17996	CG17996	1625088_at	-0.4038	0.1668	0.1032	0.5985	-0.2547	0.2137	-0.0583	0.9223	-0.4146	0.0292	-0.3562	0.0342	0.1626	0.8461	-0.0558	0.9170	-0.2184	0.5511
Gr93b	Gustatory recepto	1625089_at	0.1496	0.4290	0.0310	0.9119	-0.2949	0.2139	-0.3126	0.4022	-0.0560	0.8337	0.2566	0.1676	-0.0839	0.9235	-0.2464	0.3820	-0.1625	0.5988
---	---	1625090_at	0.1304	0.4395	-0.0882	0.5431	0.0512	0.8226	0.1251	0.8233	0.1604	0.4816	0.0352	0.8925	-0.0285	0.9778	-0.0352	0.9276	-0.0066	0.9865

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11588	CG11588	1625091_at	0.1590	0.2561	0.0014	0.9947	0.3178	0.0807	0.0034	0.9956	0.1107	0.5605	0.1073	0.5334	0.0123	0.9901	0.0917	0.7086	0.0794	0.7503
mthl6	Mth-like 6	1625092_at	0.0241	0.9006	0.0558	0.5895	0.2453	0.1298	0.0512	0.9314	0.0363	0.8777	-0.0149	0.9471	-0.1163	0.7997	-0.1153	0.5647	0.0010	0.9979
---	---	1625093_s_at	0.3566	0.1652	0.2532	0.3106	0.1897	0.2822	-0.0799	0.9017	0.0425	0.8794	0.1223	0.5535	0.0124	0.9914	-0.0666	0.8537	-0.0790	0.8054
CG14574	CG14574	1625094_at	-0.0649	0.7186	0.0414	0.6960	0.0705	0.7327	-0.0461	0.9518	-0.0632	0.8096	-0.0171	0.9494	0.2251	0.7215	0.2451	0.3421	0.0200	0.9585
CG2791	CG2791	1625095_at	-0.3489	0.0462	-0.2096	0.3381	-0.0827	0.7043	0.0913	0.8244	0.2184	0.1680	0.1271	0.3908	0.0280	0.9816	0.3471	0.2053	0.3191	0.2757
CG9300	CG9300	1625096_at	0.5900	0.0232	-0.3684	0.4421	-0.0409	0.8772	0.0574	0.9586	0.8543	0.0102	0.7969	0.0085	-0.2635	0.8369	-0.1053	0.8919	0.1582	0.8061
---	---	1625097_at	0.0481	0.8305	-0.0045	0.9855	0.0572	0.7865	0.0936	0.8738	0.0049	0.9879	-0.0888	0.6793	0.0242	0.9816	0.0384	0.9194	0.0142	0.9697
---	---	1625098_at	0.2433	0.2787	-0.0715	0.4801	0.2433	0.1696	0.3903	0.1818	0.2621	0.1372	-0.1282	0.4416	-0.1402	0.8182	0.0012	0.9991	0.1414	0.5988
CG17999	CG17999	1625099_at	2.9358	0.2554	0.1118	0.3131	0.2294	0.4709	-0.1409	0.9297	-0.5807	0.2133	-0.4397	0.2999	-0.0808	0.9952	-3.3417	0.2974	-3.2609	0.3332
---	---	1625100_at	0.0282	0.8665	-0.0796	0.5268	0.1975	0.2553	0.0301	0.9620	0.0546	0.7899	0.0245	0.9034	-0.1029	0.8472	0.0606	0.8386	0.1634	0.4759
CcapR	Cardioaccelerator	1625101_at	0.0899	0.6846	-0.1940	0.3923	-0.2739	0.0875	-0.1797	0.5855	0.1646	0.3342	0.3443	0.0295	0.0226	0.9816	0.0512	0.8690	0.0286	0.9249
CG14299	CG14299	1625102_at	0.3130	0.1652	-0.0461	0.7936	0.1564	0.3003	-0.0277	0.9669	0.0681	0.7417	0.0958	0.5836	-0.2294	0.7215	-0.3003	0.2475	-0.0709	0.8354
---	---	1625103_at	0.2559	0.2931	-0.2532	0.1372	0.0789	0.6834	0.0303	0.9704	0.1885	0.3756	0.1583	0.4166	-0.2997	0.7305	-0.3253	0.3733	-0.0256	0.9626
CG6763	CG6763	1625104_at	-0.3222	0.1372	0.1751	0.4360	0.2897	0.1262	0.1163	0.7845	-0.2361	0.1757	-0.3525	0.0319	0.0475	0.9657	0.1703	0.5872	0.1228	0.7134
---	---	1625105_s_at	-0.0739	0.7121	0.1205	0.3071	0.2769	0.2417	-0.1205	0.8111	-0.3683	0.0684	-0.2478	0.1644	-0.1397	0.7953	0.1085	0.6633	0.2481	0.2869
---	---	1625106_at	0.1012	0.5266	-0.1028	0.4809	-0.0861	0.6657	0.1468	0.7678	0.3205	0.1282	0.1736	0.3739	0.0510	0.9581	-0.1200	0.6949	-0.1710	0.5511
brn	Gene B	1625107_at	-0.1830	0.4583	-0.0992	0.7341	-0.0345	0.8850	-0.0739	0.9136	-0.0952	0.7011	-0.0213	0.9364	-0.2709	0.7772	-0.1163	0.8346	0.1546	0.7482
Lim3	lethal(2)37Bd	1625108_a_at	-0.2772	0.0837	-0.0166	0.8731	0.0564	0.7383	0.0710	0.8822	-0.0719	0.7033	-0.1429	0.3406	-0.1265	0.8191	-0.0320	0.9286	0.0946	0.7105
Rrp1	strand transferase	1625109_at	-0.6002	0.3928	-1.3704	0.0330	-1.2840	0.0057	-0.2401	0.7028	0.4439	0.1237	0.6841	0.0164	-0.2597	0.9199	-0.3258	0.7543	-0.0661	0.9584
cpa	capping protein al	1625110_at	-0.2582	0.0819	-0.5272	0.0238	-0.5071	0.0239	-0.0498	0.9121	0.4714	0.0062	0.5213	0.0024	-0.0599	0.9503	0.2645	0.3404	0.3244	0.2716
CG13786	CG13786	1625111_at	0.1507	0.3738	-0.2004	0.2168	-0.0244	0.9190	0.2563	0.5096	0.2536	0.2165	-0.0027	0.9915	0.0178	0.9848	-0.1047	0.6587	-0.1226	0.5980
Snr1	snf5-related 1	1625112_at	-0.4855	0.0655	-0.1608	0.2345	0.0481	0.7840	0.1034	0.8028	-0.1149	0.5127	-0.2183	0.1397	-0.0047	0.9968	0.1339	0.6819	0.1386	0.6621
CG6509	CG6509	1625113_s_at	-0.0615	0.7979	0.1605	0.3439	-0.2268	0.2049	0.0369	0.9626	0.2090	0.3233	0.1720	0.3724	0.2176	0.7686	0.3416	0.2593	0.1241	0.7306
Cyp310a1	Cyp310a1	1625114_at	0.1000	0.5020	-0.0307	0.8120	-0.1396	0.5923	0.0389	0.9620	0.3178	0.1398	0.2789	0.1476	0.0889	0.8882	0.0339	0.9279	-0.0549	0.8608
rha	rha	1625115_at	-0.1585	0.6464	0.1464	0.3301	-0.0707	0.7529	-0.2412	0.6785	-0.3408	0.2091	-0.0996	0.7337	0.0420	0.9764	-0.1085	0.8038	-0.1505	0.6893
trol	Perlecan	1625116_at	-1.6433	0.0006	-3.1695	0.0014	-3.0032	0.0001	0.2983	0.6652	0.9020	0.0140	0.6036	0.0458	0.0192	0.9913	-0.6887	0.0761	-0.7079	0.0897
l(1)G0084	enhancer of yellov	1625117_at	-0.0153	0.9814	-0.2979	0.7957	-0.2947	0.3941	0.1201	0.8334	0.4411	0.0470	0.3210	0.0966	0.3328	0.9270	0.3201	0.8432	-0.0127	0.9953
CG32033	CG32033	1625118_at	0.2460	0.2070	0.0040	0.9731	0.1883	0.3161	0.0769	0.8796	0.1320	0.4673	0.0551	0.7748	-0.0605	0.9405	0.0629	0.8546	0.1234	0.6460
RhoGAP54D	RhoGAP54D	1625119_at	-0.0290	0.9226	0.0584	0.8523	-0.2843	0.2682	-0.2709	0.5626	0.1495	0.5633	0.4204	0.0542	0.1878	0.8628	0.2353	0.6156	0.0475	0.9382
CG14630 /// CG33082	Transcription unit	1625120_at	1.9431	0.0070	0.8141	0.3811	1.4968	0.0020	0.1850	0.7353	0.2049	0.4142	0.0199	0.9486	-0.4397	0.8815	-0.7224	0.5404	-0.2828	0.8456
---	---	1625121_at	0.2227	0.1743	0.4358	0.1332	0.4168	0.0317	-0.1716	0.6972	-0.1428	0.5056	0.0288	0.9069	-0.0063	0.9952	0.0111	0.9806	0.0174	0.9614
Obp99c	Odorant-binding p	1625122_at	1.9596	0.0018	1.0649	0.2231	2.6975	0.0002	0.9285	0.1787	-0.2385	0.6034	-1.1670	0.0079	-0.6388	0.7644	-1.1183	0.2042	-0.4794	0.6248
CG2962	CG2962	1625123_at	0.1115	0.6027	-0.1638	0.4741	-0.0067	0.9803	0.1837	0.6669	0.1476	0.4888	-0.0361	0.8806	-0.1559	0.8494	-0.1057	0.8117	0.0502	0.9158
AttA	attacin	1625124_at	1.8430	0.5068	-2.3819	0.2216	-1.7896	0.0801	0.1490	0.9755	6.9923	0.0005	6.8433	0.0004	-0.4329	0.9816	3.1482	0.4302	3.5810	0.3819
CG5599 /// DsimCG5599	CG5599	1625125_at	0.5526	0.0278	0.1517	0.4612	0.3124	0.1004	-0.1057	0.7605	-0.0820	0.6196	0.0236	0.8953	-0.3631	0.6741	-0.3398	0.3152	0.0233	0.9637
mnm	mini-me	1625126_at	0.0612	0.8721	0.4660	0.0429	0.4229	0.0612	0.1253	0.8678	0.0391	0.9167	-0.0862	0.7693	0.1706	0.8903	0.4434	0.3278	0.2728	0.5819
fs(1)h	rancor	1625127_at	-0.6467	0.4823	-0.0975	0.9484	-1.0240	0.0169	-0.0502	0.9518	0.0663	0.8161	0.1165	0.6178	0.8674	0.8243	0.5887	0.7652	-0.2786	0.9020
CG2249 /// DsimCG2249 ///	anon-fast-evolving	1625128_a_at	0.1972	0.5945	0.9871	0.0308	0.7365	0.0041	-0.2471	0.3863	-1.0452	0.0004	-0.7981	0.0006	-0.0401	0.9862	-0.3151	0.5860	-0.2751	0.6398
CG1946	CG1946	1625129_at	-0.1579	0.5690	-0.1114	0.5629	-0.1565	0.4775	0.0396	0.9452	0.0861	0.6408	0.0465	0.8013	-0.1019	0.9238	-0.1227	0.7790	-0.0208	0.9696
CG8170	CG8170	1625130_at	1.3056	0.1477	0.1728	0.1642	0.2511	0.2525	-0.1041	0.8405	0.0685	0.7675	0.1726	0.3280	-0.0293	0.9943	-0.7567	0.4365	-0.7273	0.4697
Vha55	vacuolar ATPase	1625131_s_at	-0.6535	0.0073	-0.7145	0.0525	-1.3002	0.0001	-0.2602	0.3921	-0.1681	0.3336	0.0920	0.5913	0.2571	0.7215	-0.2927	0.3197	-0.5498	0.1033
---	---	1625132_s_at	0.0813	0.5774	0.0525	0.6031	-0.0345	0.8521	-0.0664	0.9110	-0.0517	0.8282	0.0147	0.9508	0.0330	0.9729	0.0982	0.7248	0.0652	0.8307
CG14131	CG14131	1625133_at	1.3935	0.0010	0.5260	0.0880	0.8014	0.0026	0.2341	0.5744	0.7007	0.0063	0.4666	0.0229	-0.0388	0.9816	-0.1599	0.7034	-0.1212	0.7836
p53	p53-like regulator	1625134_at	0.4264	0.0401	0.0237	0.9078	0.3729	0.0460	0.0190	0.9753	0.1007	0.5281	0.0816	0.5849	-0.2277	0.7644	-0.3083	0.3259	-0.0805	0.8458
Cby	Chibby	1625135_at	0.2137	0.2538	0.3253	0.1915	0.0021	0.9944	-0.0504	0.9436	0.0367	0.8934	0.0871	0.6798	0.4283	0.5228	0.2611	0.4124	-0.1672	0.6296
mRpS18B	mitochondrial ribo	1625136_s_at	0.3387	0.2020	0.7959	0.0330	0.7714	0.0023	-0.0386	0.9467	-0.1425	0.3946	-0.1040	0.5097	0.0094	0.9959	0.2817	0.5199	0.2722	0.5416
CG3281	CG3281	1625137_at	-0.6728	0.0121	-0.2298	0.3786	-0.3731	0.0236	-0.0639	0.9251	0.0195	0.9492	0.0834	0.6969	0.0619	0.9449	0.3957	0.1397	0.3338	0.2338
CG13530	CG13530	1625138_at	0.1278	0.3993	0.1539	0.4063	0.0487	0.8411	-0.1163	0.8274	-0.2668	0.1910	-0.1506	0.4349	0.0725	0.9246	-0.1109	0.7036	-0.1834	0.4913
Gadd45	Gadd45	1625139_at	0.3907	0.1781	0.8305	0.2224	0.6810	0.0459	0.4007	0.3812	0.2978	0.2504	-0.1029	0.7065	0.7284	0.6749	0.9398	0.1750	0.2114	0.8112
NtR	NtR	1625140_at	-2.5374	0.0265	-4.5670	0.0088	-4.8229	0.0000	-0.3095	0.4459	0.0564	0.8442	0.3659	0.0677	-0.0247	0.9964	-1.9407	0.1512	-1.9160	0.1911
CG14715	CG14715	1625141_at	0.8429	0.0086	0.1039	0.5851	0.0885	0.7003	0.0507	0.9466	0.8146	0.0033	0.7639	0.0026	0.0881	0.9489	-0.0589	0.9277	-0.1469	0.7565
CG33268 /// CG7377	CG33268 /// CG7	1625142_s_at	0.1843	0.4402	0.3440	0.1619	0.2141	0.2787	0.0015	0.9988	-0.1368	0.6356	-0.1382	0.5924	0.0794	0.9330	-0.0583	0.8981	-0.1377	0.6787
Cpr65Ea	CG8640	1625143_at	-0.3148	0.2737	-0.0817	0.4903	0.2224	0.3757	0.1649	0.7303	-0.1417	0.5359	-0.3066	0.1081	-0.1144	0.8504	0.0559	0.8764	0.1704	0.5158
CG6758 /// DmirCG6758	CG6758	1625144_at	-0.5219	0.0479	-0.1135	0.7838	-0.0877	0.7214	0.0693	0.8791	-0.2052	0.1841	-0.2745	0.0525	-0.0313	0.9898	0.1489	0.8353	0.1802	0.7751
Anxb11	Annexin B11	1625145_a_at	-0.5362	0.0115	-0.1783	0.3102	-0.4500	0.0094	0.0250	0.9672	0.1779	0.2638	0.1529	0.2865	0.0371	0.9689	0.4321	0.0929	0.3949	0.1409
RN-tre	tre oncogene-relat	1625146_a_at	-0.7477	0.0369	0.9295	0.1096	0.6157	0.0698	-0.1017	0.9196	-1.0815	0.0047	-0.9797	0.0043	0.2812	0.8222	0.5916	0.2387	0.3104	0.5741
CG7866	CG7866	1625147_at	0.3762	0.0943	-0.0841	0.6736	-0.2340	0.2226	0.0530	0.9346	0.5881	0.0083	0.5351	0.0077	0.1455	0.8270	0.1295	0.6746	-0.0161	0.9701
14-3-3zeta	leonardo	1625148_s_at	-1.1231	0.0343	-0.5333	0.1298	-1.5508	0.0003	-0.2527	0.6999	-0.0449	0.9150	0.2078	0.4667	0.6908	0.6659	0.4163	0.5329	-0.2745	0.7030
CG7460	CG7460	1625149_at	-1.2677	0.0035	1.3648	0.0328	0.4517	0.1703	-0.6023	0.2830	-1.9577	0.0005	-1.3554	0.0014	0.3137	0.7681	0.6504	0.1418	0.3367	0.4702
Ef2b	Elongation factor 2	1625150_s_at	-0.0328	0.8433	-0.0108	0.9323	0.0631	0.6759	0.0038	0.9956	-0.0269	0.8910	-0.0307	0.8558	0.0059	0.9939	0.0226	0.9366	0.0167	0.9466
CG32770	CG32770	1625151_at	-0.0026	0.9915	-0.1023	0.4871	0.0697	0.7005	-0.0371	0.9562	0.0184	0.9456	0.0555	0.7860	-0.0257	0.9764	-0.0142	0.9670	0.0116	0.9714
CG17612	CG17612	1625152_s_at	-0.1290	0.4923	-0.5906	0.0812	-0.9588	0.0005	-0.0583	0.9524	0.5998	0.0338	0.6581	0.0146	0.1995	0.6607	0.0541	0.8230	-0.1454	0.4457
p24-2	p24-related-2	1625153_at	-0.2867	0.4137	0.4402	0.2652	0.8568	0.0367	-0.0140	0.9931	0.0425	0.9244	0.0565	0.8815	-0.4212	0.7707	0.8863	0.1418	1.3075	0.0710
CG2467	CG2467	1625154_s_at	-1.5055	0.0680	-0.5477	0.6468	0.2390	0.6526	0.3552	0.6175	-0.8007	0.0315	-1.1559	0.0039	-0.2747	0.9592	0.4216	0.8390	0.6963	0.6842
gammaCop	gamma-coatomer	1625155_a_at	0.7323	0.0197	0.5777	0.0269	0.6438	0.0064	0.1938	0.6010	0.7920	0.0020	0.5981	0.0041	0.1834	0.8521	0.6386	0.1209	0.4552	0.2798
CG8757	CG8757	1625156_at	0.0924	0.6711	-0.0413	0.7950	0.2273	0.2341	0.1225	0.8678	0.1249	0.6739	0.0025	0.9940	-0.0601	0.9542	0.0077	0.9914	0.0679	0.8680
---	---	1625157_at	0.0371	0.8141	-0.0881	0.4872	-0.0056	0.9792	-0.0762	0.8663	-0.0667	0.7194	0.0095	0.9622	-0.2499	0.6955	-0.0654	0.8614	0.1845	0.5093
Teh4	tipE homolog 4	1625158_at	-0.0404	0.8454	0.0594	0.5596	0.0437	0.8643	-0.1056	0.9086	-0.2296	0.4523	-0.1240	0.6873	-0.0438	0.9741	-0.0836	0.8546	-0.0398	0.9319
CG18561	CG18561	1625159_at	-0.3057	0.0611	0.0088	0.9372	-0.1329	0.4165	-0.1460	0.7327	-0.2857	0.1323	-0.1397	0.4349	-0.1117	0.7956	-0.0514	0.8313	0.0602	0.7787

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1625160_at	0.1170	0.5292	-0.0036	0.9827	0.0535	0.7529	0.0798	0.8564	0.0862	0.6246	0.0064	0.9738	0.0958	0.8395	0.0359	0.9042	-0.0600	0.7995
---	---	1625161_at	0.4666	0.0472	0.1674	0.4274	0.0473	0.8106	-0.0809	0.8877	-0.0109	0.9710	0.0700	0.7364	0.0965	0.9296	-0.1040	0.8220	-0.2004	0.5984
CG4611	CG4611	1625162_at	0.2067	0.2827	-0.1734	0.6915	0.0236	0.9349	0.0737	0.9313	0.6273	0.0212	0.5536	0.0224	-0.0984	0.9545	0.3418	0.5121	0.4402	0.3921
Surf1	Surfeit 1	1625163_at	-0.6205	0.0391	-0.4538	0.0218	0.1432	0.4031	-0.0019	0.9980	-0.3429	0.0632	-0.3411	0.0424	-0.6014	0.3162	-0.1909	0.5629	0.4105	0.2110
TotX	Turandot X	1625164_at	3.2158	0.0010	0.5131	0.6866	3.4217	0.0179	2.5582	0.2679	1.8999	0.1617	-0.6583	0.6381	-0.2559	0.9589	-0.7321	0.6321	-0.4762	0.7755
CG17187	CG17187	1625165_at	-0.0806	0.7442	0.1358	0.5926	0.2195	0.2538	0.0339	0.9755	0.2618	0.3358	0.2279	0.3540	-0.0381	0.9762	0.5581	0.0809	0.5962	0.0878
CG15456	CG15456	1625166_at	-0.6931	0.0142	0.0325	0.8079	-0.0353	0.8846	0.0274	0.9782	-0.6061	0.0257	-0.6335	0.0134	-0.1063	0.8882	-0.0770	0.8397	0.0293	0.9413
---	---	1625167_at	-0.0853	0.5565	0.2038	0.3599	0.4103	0.0181	0.2546	0.4037	-0.0454	0.8349	-0.3000	0.0531	0.0674	0.9253	0.1668	0.4999	0.0994	0.7148
CG30153	CG30153	1625168_at	0.2687	0.3457	0.1276	0.5614	-0.4771	0.1502	-0.2163	0.6885	0.2577	0.3073	0.4740	0.0404	0.3428	0.8049	0.0989	0.9111	-0.2439	0.7067
CG14667	CG14667	1625169_at	0.1679	0.2918	-0.2509	0.1754	-0.1608	0.3123	0.0209	0.9759	0.2441	0.1475	0.2232	0.1386	-0.1003	0.8427	-0.1981	0.3269	-0.0979	0.6646
CG8001	CG8001	1625170_at	0.0284	0.9457	0.4482	0.2314	0.6708	0.0077	0.0847	0.8987	-0.3364	0.1211	-0.4211	0.0374	-0.1804	0.9036	0.0825	0.9222	0.2630	0.6533
---	---	1625171_at	-0.0429	0.8881	-0.0399	0.6804	-0.0115	0.9573	-0.0128	0.9893	0.0653	0.8104	0.0781	0.7421	-0.0721	0.9317	0.0613	0.8751	0.1334	0.6486
CG2316	CG2316	1625172_s_at	-0.6753	0.0412	0.5621	0.0616	0.4221	0.0137	-0.3534	0.2939	-0.8529	0.0019	-0.4995	0.0124	-0.2621	0.7220	0.2484	0.4255	0.5105	0.1313
CG11652	CG11652	1625173_s_at	-0.1642	0.4685	0.5838	0.0395	0.6247	0.0058	0.1659	0.7556	-1.0545	0.0015	-1.2204	0.0005	0.2065	0.7893	-0.2782	0.3919	-0.4846	0.1651
CG15067	CG15067	1625174_at	2.0826	0.0022	0.9907	0.1567	1.7306	0.0045	0.4428	0.5376	0.3736	0.3262	-0.0693	0.8763	-0.1651	0.9400	-0.5873	0.3738	-0.4223	0.5523
CG7082	CG7082	1625175_s_at	-0.5010	0.0662	-0.7135	0.0990	-1.0517	0.0002	0.2170	0.6338	0.8855	0.0027	0.6685	0.0057	0.4215	0.6695	0.5806	0.1397	0.1591	0.7262
CG33174 /// DmirCG33174	CG33174	1625176_at	-0.1009	0.5320	-0.0925	0.4540	-0.0828	0.6511	0.1256	0.7857	0.0658	0.7743	-0.0598	0.7771	0.0693	0.9340	-0.0364	0.9320	-0.1057	0.7280
CG14182	CG14182	1625177_at	0.1333	0.6682	0.4120	0.1417	0.3965	0.0380	-0.0667	0.9515	-0.2117	0.5068	-0.1450	0.6399	0.0195	0.9898	0.1636	0.6501	0.1440	0.6973
CG31038	CG31038	1625178_at	-0.9412	0.2951	-0.2205	0.8484	-0.1521	0.7680	0.0938	0.9056	-0.6865	0.0151	-0.7803	0.0052	0.0201	0.9976	0.1361	0.9561	0.1160	0.9578
CG6000	CG6000	1625179_at	-0.0412	0.7928	0.4711	0.0395	0.2417	0.4139	-0.2657	0.4815	-0.7827	0.0030	-0.5170	0.0117	-0.1516	0.8694	0.0563	0.9222	0.2079	0.6006
---	---	1625180_at	0.1698	0.4386	0.1771	0.1272	0.1798	0.4780	0.1137	0.8632	0.0224	0.9494	-0.0913	0.7136	0.1341	0.8521	0.0646	0.8793	-0.0694	0.8579
CG5172	CG5172	1625181_at	-0.5335	0.1088	-0.4770	0.0328	-0.6830	0.0022	-0.1707	0.7558	0.0388	0.9058	0.2095	0.3335	0.0082	0.9926	0.0130	0.9689	0.0048	0.9877
---	---	1625182_at	-0.0069	0.9715	-0.1631	0.4403	-0.1955	0.2829	-0.1367	0.7695	0.0328	0.9044	0.1695	0.3501	0.0119	0.9913	-0.0636	0.8319	-0.0755	0.7764
smi35A	Dual specificity tyr	1625183_at	-1.4370	0.0012	-1.7828	0.0103	-1.8716	0.0003	-0.0037	0.9963	0.0615	0.8645	0.0652	0.8349	-0.0741	0.9689	-0.2514	0.6408	-0.1773	0.7568
CG15403	CG15403	1625184_at	-0.0464	0.7968	0.0434	0.7543	0.2984	0.0961	-0.2521	0.4317	-0.3712	0.0413	-0.1191	0.4776	-0.3840	0.5089	-0.0504	0.9049	0.3335	0.2409
CAH2	Carbonic anhydra	1625185_at	0.0384	0.9060	0.3196	0.2995	0.1092	0.5975	-0.1914	0.7349	-0.5995	0.0250	-0.4081	0.0714	0.0899	0.9421	-0.3028	0.4296	-0.3928	0.3171
CG9589	CG9589	1625186_at	0.1476	0.5519	0.2600	0.2552	0.3099	0.1671	0.0462	0.9558	-0.0113	0.9743	-0.0575	0.8207	0.0956	0.8940	0.0799	0.8175	-0.0156	0.9695
Sbf	SET domain bindi	1625187_a_at	0.0657	0.8112	0.1195	0.3906	-0.2556	0.0769	-0.1252	0.6908	0.4458	0.0086	0.5710	0.0017	0.2364	0.7118	0.4653	0.0917	0.2290	0.4009
gpp	grappa	1625188_at	-0.0494	0.7849	-0.0852	0.8107	0.1454	0.3912	-0.0154	0.9883	-0.0679	0.8314	-0.0525	0.8572	-0.1556	0.7644	-0.0508	0.8684	0.1047	0.6567
---	---	1625189_at	0.2517	0.2499	-0.1379	0.3651	-0.2607	0.3601	-0.0979	0.8817	0.4336	0.0580	0.5316	0.0163	-0.1188	0.9024	-0.0333	0.9531	0.0855	0.8514
CG31105	CG31105	1625190_at	0.2519	0.2381	0.0643	0.5629	0.0047	0.9868	-0.1110	0.7631	0.0892	0.6072	0.2002	0.1550	-0.0761	0.9400	-0.1456	0.6713	-0.0694	0.8656
CG5022	CG5022	1625191_at	-0.0782	0.6965	0.0000	1.0000	0.4134	0.0359	0.1718	0.6394	-0.1054	0.5892	-0.2771	0.0848	-0.0881	0.9239	0.0278	0.9558	0.1159	0.7442
CG12885	CG12885	1625192_at	0.1433	0.4580	-0.0224	0.8885	0.1204	0.4796	0.0315	0.9639	0.0487	0.8371	0.0171	0.9404	-0.1028	0.8608	-0.1334	0.5943	-0.0306	0.9246
CG31106	CG31106	1625193_at	-0.1988	0.8601	-0.1740	0.3881	0.0563	0.7205	0.2630	0.8667	-0.5310	0.3447	-0.7940	0.1070	-0.0875	0.9816	-0.7848	0.3158	-0.6973	0.3939
CG34113	CG34113	1625194_at	0.0732	0.7442	0.0459	0.7537	0.1314	0.4269	0.2770	0.5735	0.2603	0.3050	-0.0167	0.9583	0.0320	0.9816	0.0433	0.9345	0.0113	0.9838
shn	quo vadis	1625195_s_at	-0.6978	0.2738	-1.3502	0.1187	-1.3181	0.0010	0.0048	0.9966	0.1612	0.6965	0.1564	0.6776	-0.0522	0.9913	-0.4839	0.6594	-0.4316	0.6988
CG14995	Gandalf	1625196_a_at	-1.8962	0.0028	-2.3337	0.0235	-2.8352	0.0000	-0.3309	0.2988	-0.2359	0.2059	0.0950	0.6166	0.1219	0.9637	-0.7782	0.2682	-0.9001	0.2326
exba	krasavietz	1625197_at	0.0957	0.7453	-0.2157	0.3388	-0.2954	0.0707	0.1083	0.8707	0.6073	0.0172	0.4989	0.0246	0.2081	0.8133	0.3085	0.3909	0.1004	0.8256
ss	spineless-aristape	1625198_at	-0.2338	0.1743	0.0104	0.9278	0.0948	0.5857	-0.0443	0.9311	-0.0951	0.5582	-0.0508	0.7574	-0.1244	0.8786	0.0752	0.8637	0.1996	0.5481
dap	decapo	1625199_s_at	0.0301	0.9546	-0.3453	0.4045	-0.7126	0.0118	-0.3300	0.6814	-0.1762	0.6796	0.1538	0.6960	-0.1781	0.8882	-0.6885	0.1418	-0.5104	0.2984
CG30386	CG30386	1625200_at	0.0899	0.6481	0.1677	0.5009	0.0908	0.6572	-0.0710	0.9065	0.0655	0.7810	0.1366	0.4584	0.0628	0.9543	0.2058	0.5398	0.1430	0.6877
CG17230	CG17230	1625201_s_at	-1.2454	0.0126	-1.1660	0.0381	-1.3853	0.0011	0.1284	0.9171	0.0383	0.9460	-0.0901	0.8372	0.0230	0.9928	-0.2436	0.7228	-0.2666	0.6873
---	---	1625202_at	0.1565	0.4802	-0.0037	0.9755	0.0346	0.8469	-0.0650	0.9228	0.0346	0.9010	0.0996	0.6288	-0.0053	0.9964	-0.0503	0.8989	-0.0450	0.9019
CG9826	CG9826	1625203_at	3.3359	0.0005	3.3573	0.0011	3.7163	0.0000	1.0651	0.0221	0.8279	0.0059	-0.2372	0.2931	0.5046	0.5126	0.8553	0.0413	0.3507	0.3518
CG34134 /// CG8475	CG8475 /// CG34	1625204_a_at	-0.2004	0.2662	-0.2733	0.2728	0.1467	0.3063	-0.0447	0.9451	-0.5109	0.0135	-0.4662	0.0124	-0.6425	0.1628	-0.6833	0.0227	-0.0407	0.8947
CG31544	CG31544	1625205_at	0.0618	0.7160	-0.0608	0.5189	-0.0296	0.8837	-0.0042	0.9956	0.0490	0.8087	0.0533	0.7680	-0.0915	0.8971	-0.1006	0.7454	-0.0091	0.9838
---	---	1625206_at	0.2303	0.2245	0.2270	0.1683	-0.0855	0.6739	-0.1167	0.7815	-0.0119	0.9644	0.1048	0.5389	0.2092	0.6749	0.0388	0.8995	-0.1705	0.4090
CG32636	CG32636	1625207_at	0.3766	0.3098	-0.0918	0.7803	0.0767	0.7813	0.1208	0.8967	0.3543	0.2499	0.2334	0.4165	-0.0279	0.9914	-0.0376	0.9664	-0.0097	0.9921
CG15393 /// Slh	CG15393 /// SLY-	1625208_at	-0.0424	0.8139	0.2089	0.2445	0.2207	0.2406	-0.0450	0.9325	-0.0747	0.6781	-0.0297	0.8736	-0.0216	0.9816	0.0550	0.8478	0.0766	0.7520
CG40392	CG40392	1625209_at	0.2480	0.3331	0.1376	0.3798	0.0306	0.8968	-0.0444	0.9422	-0.0901	0.6460	-0.0457	0.8172	-0.2335	0.6584	-0.0700	0.7972	0.1636	0.4631
Msp-300	nesprin	1625210_a_at	-2.6676	0.0075	-3.5459	0.0077	-3.5266	0.0000	-0.0748	0.9386	0.5183	0.0714	0.5932	0.0276	-0.0898	0.9831	-0.3143	0.8042	-0.2245	0.8635
CG32810	CG32810	1625211_s_at	0.3642	0.0406	0.1864	0.1694	0.0800	0.6199	-0.1594	0.6354	0.1467	0.3855	0.3061	0.0440	-0.0103	0.9916	0.0229	0.9506	0.0332	0.9171
CG4653	CG4653	1625212_at	-0.1247	0.9673	0.0719	0.6320	0.1879	0.2363	0.0025	0.9997	-1.6849	0.2541	-1.6874	0.1995	-0.0827	0.9928	-1.2535	0.5856	-1.1708	0.6167
---	---	1625213_at	0.0250	0.8800	-0.0403	0.6984	0.2999	0.1426	0.0245	0.9727	0.1411	0.4462	0.1166	0.4961	-0.2758	0.6955	-0.0529	0.9075	0.2229	0.4683
Jupiter	Jupiter	1625214_at	-0.2406	0.1386	-0.2615	0.1865	-1.0999	0.0013	-0.1836	0.6615	0.1719	0.4051	0.3554	0.0523	0.0266	0.9742	0.0756	0.7573	0.0490	0.8540
nrv2	nervana2	1625215_s_at	-1.5233	0.0023	-0.3156	0.6453	-2.0175	0.0003	-1.0807	0.0935	-1.4910	0.0028	-0.4103	0.2252	0.5766	0.7307	-0.2572	0.7678	-0.8338	0.2614
CG13724	CG13724	1625216_at	-0.0352	0.8695	0.0258	0.8106	0.0630	0.8568	-0.1547	0.8050	-0.3037	0.2236	-0.1490	0.5428	0.0166	0.9923	0.0155	0.9841	-0.0012	0.9990
MED17	Mediator complex	1625217_at	-0.1772	0.3667	0.4856	0.0752	1.3145	0.0002	-0.0047	0.9956	-0.9758	0.0018	-0.9711	0.0011	-0.4314	0.3828	-0.1536	0.5988	0.2779	0.3205
CG11399 /// DmirCG11399	CG11399	1625218_at	-0.0185	0.9598	0.0155	0.9591	-0.3666	0.1004	-0.2646	0.5931	-0.0212	0.9548	0.2434	0.2832	0.0685	0.9545	0.0922	0.8476	0.0237	0.9636
ninaC	Droninac	1625219_at	0.2532	0.2690	-0.0714	0.5773	0.1743	0.3491	0.1045	0.8462	0.1894	0.3464	0.0849	0.6814	-0.0133	0.9893	-0.0552	0.8432	-0.0419	0.8813
---	---	1625220_at	0.1582	0.4758	0.1185	0.6093	0.0720	0.7177	0.0345	0.9529	0.0019	0.9939	-0.0326	0.8619	-0.0637	0.9646	0.0620	0.9172	0.1258	0.7836
CG7994	CG7994	1625221_at	0.1547	0.3799	-0.0111	0.9181	0.0959	0.5393	0.1029	0.7939	0.1538	0.3473	0.0509	0.7750	0.0241	0.9831	-0.1115	0.7087	-0.1356	0.6328
Gr22f	Gustatory recepto	1625222_at	0.1350	0.4547	0.0334	0.8995	-0.1134	0.5824	0.0759	0.9028	0.1033	0.6476	0.0274	0.9101	0.0561	0.9619	-0.0159	0.9784	-0.0721	0.8694
stck	PINCH	1625223_a_at	-0.9891	0.0030	-0.2346	0.3522	-0.4187	0.0419	-0.0932	0.8143	-0.4413	0.0123	-0.3481	0.0211	-0.0909	0.9417	0.0004	0.9999	0.0913	0.8547
---	---	1625224_at	0.0653	0.7713	0.1244	0.5281	0.1465	0.4250	0.0143	0.9863	0.1309	0.5477	0.1167	0.5594	-0.1064	0.8692	0.0701	0.8386	0.1765	0.5083
---	---	1625225_at	0.0690	0.6978	-0.0206	0.8399	0.0684	0.6511	0.1654	0.5948	0.0763	0.6708	-0.0890	0.5690	-0.0071	0.9943	-0.0224	0.9506	-0.0153	0.9630
---	---	1625226_at	0.0796	0.5985	0.0200	0.8605	0.0728	0.6851	-0.0673	0.9136	-0.1315	0.5287	-0.0642	0.7653	0.0930	0.8541	0.0603	0.8254	-0.0327	0.9095
CG30456	CG30456	1625227_at	0.7528	0.0088	0.1422	0.5583	0.1118	0.4274	-0.0948	0.8402	0.6563	0.0035	0.7512	0.0012	0.0629	0.9701	0.1058	0.8584	0.0429	0.9433
CG8394	CG8394	1625228_at	-0.0452	0.8041	0.1621	0.4402	-0.0501	0.7936	0.1766	0.7442	0.0380	0.9084	-0.1386	0.5535	0.1135	0.8689	-0.0223	0.9605	-0.1358	0.6419

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1625229_at	0.0056	0.9795	0.0538	0.8048	0.0121	0.9689	0.0234	0.9816	0.0310	0.9269	0.0076	0.9799	0.0079	0.9939	-0.0328	0.9270	-0.0408	0.8949
CG31738	CG31738	1625230_a_at	0.4895	0.4020	0.1535	0.8176	-0.0087	0.9833	0.1496	0.8640	1.3973	0.0016	1.2477	0.0015	0.3609	0.8815	1.0432	0.2430	0.6823	0.4773
U2af38	U2 small nuclear r	1625231_at	-0.1439	0.4968	0.4120	0.0509	0.6193	0.0131	0.0113	0.9937	-0.3956	0.1747	-0.4070	0.1190	-0.1912	0.7893	0.1836	0.5670	0.3748	0.2350
---	---	1625232_at	0.1544	0.5487	-0.0394	0.7992	0.1251	0.4347	-0.0588	0.9254	-0.0975	0.6460	-0.0387	0.8605	-0.1357	0.8795	-0.1325	0.7424	0.0031	0.9959
Pi3K68D	dPI 3-kinase	1625233_at	-0.5097	0.0650	-0.6253	0.0881	-0.7935	0.0014	-0.4003	0.3276	-0.0056	0.9879	0.3947	0.0657	-0.1232	0.8692	-0.0830	0.8349	0.0402	0.9215
---	---	1625234_at	0.3098	0.1619	0.1777	0.5114	0.0743	0.7630	0.0520	0.9483	0.1505	0.5307	0.0985	0.6752	0.0239	0.9725	-0.0531	0.8150	-0.0770	0.6896
CG13325	CG13325	1625235_at	0.1919	0.2988	-0.0872	0.3662	-0.1412	0.5579	0.0686	0.9314	0.1370	0.5964	0.0683	0.7976	-0.0232	0.9853	-0.1738	0.5659	-0.1506	0.6293
Prosap	Prosap	1625236_s_at	0.0180	0.9667	0.8272	0.0508	-0.0352	0.8949	-0.6038	0.2438	-0.5669	0.0682	0.0369	0.9222	0.2099	0.8461	0.2586	0.5824	0.0487	0.9372
CG1894	CG1894	1625237_at	0.1436	0.4595	0.0924	0.5617	0.0520	0.8251	-0.2455	0.4456	-0.0781	0.6996	0.1674	0.2931	0.2713	0.7152	0.0795	0.8518	-0.1918	0.5612
CG12003	CG12003	1625238_at	0.2762	0.1087	0.1279	0.4640	-0.0304	0.8927	-0.0393	0.9475	0.0607	0.7640	0.1000	0.5447	0.0995	0.8331	-0.0418	0.8831	-0.1413	0.4888
---	---	1625239_at	0.0879	0.6828	0.1357	0.4276	0.0915	0.6430	-0.1013	0.8155	-0.0593	0.7727	0.0421	0.8291	0.0258	0.9831	0.0373	0.9353	0.0115	0.9809
---	---	1625240_at	0.0249	0.8976	-0.0103	0.9678	0.2459	0.1921	0.1833	0.6869	0.1881	0.3848	0.0048	0.9856	-0.0754	0.9238	0.0906	0.7755	0.1660	0.5445
---	---	1625241_at	0.1829	0.4622	0.0618	0.5690	0.0593	0.7154	0.0360	0.9470	0.0594	0.7423	0.0234	0.8991	0.0135	0.9914	-0.0548	0.8903	-0.0683	0.8444
CG31249	CG31249	1625242_at	-0.2197	0.1760	-0.2406	0.0633	-0.0610	0.7509	-0.1042	0.7469	0.1701	0.2271	0.2743	0.0365	-0.1707	0.7220	0.1163	0.5957	0.2870	0.1883
E(bx)	Nucleosome remo	1625243_a_at	0.6602	0.0608	0.2886	0.5821	-0.4779	0.0453	-0.3722	0.3862	0.4492	0.0676	0.8214	0.0033	0.3641	0.7707	0.0576	0.9467	-0.3065	0.5889
CG15433	CG15433	1625244_at	0.0967	0.5933	0.5102	0.1804	0.9847	0.0020	0.3405	0.2792	0.0506	0.8307	-0.2900	0.0821	-0.1089	0.9409	0.4359	0.3259	0.5448	0.2497
CG3759	laccase-like	1625245_at	-1.1298	0.0247	0.1626	0.5939	0.1629	0.4163	-0.0532	0.9774	-1.8349	0.0027	-1.7817	0.0018	-0.0275	0.9862	-0.3558	0.3184	-0.3283	0.3810
Her	HES-related	1625246_at	0.1540	0.3361	0.1014	0.4290	0.1900	0.5261	0.1059	0.9068	0.0196	0.9651	-0.0863	0.7916	0.0863	0.9400	0.0307	0.9585	-0.0556	0.9089
dro6	drosomycin-l	1625247_at	0.0091	0.9791	0.0176	0.9312	0.1767	0.5745	0.1178	0.8336	-0.0459	0.8707	-0.1637	0.4097	0.2502	0.8215	0.0089	0.9935	-0.2413	0.6271
RpS29	Ribosomal protein	1625248_at	0.1263	0.6555	0.2916	0.4103	0.9562	0.0356	-0.1048	0.8837	-0.0421	0.9001	0.0627	0.8212	-0.4890	0.7500	0.0232	0.9857	0.5122	0.4475
ld14	forked end	1625249_at	-1.1262	0.0119	-1.9859	0.0029	-1.7209	0.0004	0.0942	0.9098	0.7603	0.0112	0.6662	0.0124	-0.0150	0.9914	-0.0529	0.9075	-0.0379	0.9265
CG5802	anon-fast-evolving	1625250_at	0.7169	0.0035	0.5130	0.0755	0.8570	0.0031	0.1366	0.7857	0.6228	0.0093	0.4862	0.0169	-0.1065	0.8825	0.5948	0.0520	0.7013	0.0457
---	---	1625251_at	-0.0055	0.9801	-0.0036	0.9767	0.1357	0.3672	-0.0080	0.9915	-0.1078	0.5175	-0.0999	0.5099	-0.1801	0.7220	-0.0382	0.9080	0.1418	0.5334
CG12158	CG12158	1625252_s_at	0.2500	0.3221	0.3679	0.0404	0.1363	0.6903	-0.0150	0.9860	-0.0942	0.6835	-0.0792	0.7129	0.1427	0.9168	-0.1330	0.8324	-0.2757	0.5832
POSH	Plenty of SH3s	1625253_at	0.3648	0.0922	-0.1756	0.2696	-0.2932	0.1422	0.1507	0.7400	0.9669	0.0012	0.8162	0.0014	0.2250	0.7215	0.3338	0.1957	0.1088	0.7184
---	---	1625254_at	-0.1364	0.5547	0.1765	0.2904	-0.0227	0.9181	0.0243	0.9774	0.0300	0.9197	0.0056	0.9835	0.0132	0.9870	0.0195	0.9485	0.0063	0.9840
CG11029	CG11029	1625255_at	-3.4218	0.0067	0.7015	0.1503	-1.8600	0.0158	-1.9716	0.1109	-5.2202	0.0003	-3.2485	0.0010	0.6113	0.8331	-0.9666	0.4203	-1.5780	0.2100
---	---	1625256_at	0.0844	0.6605	0.2490	0.4808	0.1283	0.6736	0.3590	0.4840	0.1442	0.6381	-0.2148	0.3993	0.1665	0.8494	0.0792	0.8792	-0.0873	0.8533
CG31523	CG31523	1625257_s_at	1.3431	0.0091	1.1158	0.0066	1.1805	0.0019	0.4482	0.4174	0.5906	0.0583	0.1424	0.6530	0.3011	0.7464	0.3394	0.3820	0.0384	0.9444
cid	centromere identif	1625258_at	0.0813	0.7521	-0.1331	0.4682	-0.0671	0.8067	-0.2802	0.3909	0.1056	0.5998	0.3858	0.0270	-0.3265	0.7681	-0.2531	0.6152	0.0734	0.9096
CG10265	CG10265	1625259_at	0.0940	0.6606	-0.5444	0.1674	0.0855	0.5882	0.6251	0.0312	0.9066	0.0006	0.2816	0.0518	-0.4634	0.5967	-0.2228	0.5855	0.2406	0.5557
CG11802	CG11802	1625260_at	-0.5964	0.0181	-0.0977	0.8988	-0.1262	0.6631	-0.0370	0.9451	-0.7149	0.0014	-0.6778	0.0010	0.0703	0.9831	-0.0390	0.9779	-0.1092	0.9189
CG13722	CG13722	1625261_x_at	0.8987	0.0376	0.4198	0.2513	0.9197	0.0047	0.1650	0.8640	0.2031	0.5849	0.0381	0.9274	-0.0807	0.9643	-0.2709	0.6019	-0.1902	0.7337
CG17786	CG17786	1625262_at	-0.7144	0.2068	-2.7367	0.0026	-2.3767	0.0058	0.2787	0.9102	2.0840	0.0159	1.8053	0.0181	-0.2128	0.8882	0.0989	0.9090	0.3117	0.6111
Nipped-A	Nipped	1625263_at	0.0985	0.7326	-0.2331	0.6677	-0.4627	0.0168	-0.2913	0.4232	0.2983	0.1362	0.5897	0.0062	-0.0243	0.9918	-0.0405	0.9647	-0.0161	0.9852
Cp1	Cathepsin L	1625264_s_at	-0.0851	0.5907	-0.0730	0.5779	0.1741	0.4295	-0.0041	0.9956	-0.4719	0.0080	-0.4678	0.0050	-0.2067	0.8000	-0.4517	0.1718	-0.2450	0.4865
CG9119	CG9119	1625265_at	1.5255	0.0107	-0.2386	0.8676	0.2370	0.5212	-0.0603	0.9263	-0.3748	0.0600	-0.3145	0.0745	-0.4852	0.9046	-2.1234	0.1400	-1.6381	0.2782
Ag5r	antigen 5-related	1625266_at	0.5839	0.7172	0.1003	0.4402	-0.7810	0.0863	-0.9701	0.0882	-0.3127	0.3542	0.6574	0.0337	-0.0443	0.9952	-1.0510	0.5972	-1.0067	0.6166
CG40463	CG40463	1625267_at	-0.0114	0.9478	-0.1036	0.6169	0.0354	0.8721	0.0917	0.8462	0.0667	0.7460	-0.0251	0.9046	0.0345	0.9689	-0.1091	0.6692	-0.1436	0.5585
CG11268	CG11268	1625268_at	0.0099	0.9717	-0.3279	0.1015	0.3679	0.0310	0.0605	0.9303	-0.1166	0.5991	-0.1771	0.3407	-0.5451	0.3166	-0.2783	0.3259	0.2668	0.3708
CG9004 /// DmirCG9004	CG9004	1625269_at	0.0964	0.7716	-0.4258	0.0323	-0.0794	0.7399	-0.0322	0.9705	0.7254	0.0066	0.7575	0.0032	-0.4163	0.6824	0.1658	0.7345	0.5821	0.1901
CG31259	CG31259	1625270_at	0.5551	0.0509	-0.0730	0.6550	0.6770	0.0015	0.2280	0.3619	0.2566	0.0741	0.0286	0.8630	-0.3530	0.7070	-0.1607	0.7220	0.1923	0.6498
CG15438	CG15438	1625271_at	0.3894	0.0712	1.3716	0.0333	1.5385	0.0004	0.0109	0.9941	-0.0045	0.9916	-0.0154	0.9679	-0.1295	0.9144	0.9802	0.0418	1.1098	0.0438
CG31030	CG31030	1625272_at	-0.9500	0.0584	0.1425	0.3191	-0.1114	0.5585	-0.2454	0.7749	-1.0407	0.0109	-0.7952	0.0215	-0.0346	0.9816	-0.1696	0.6314	-0.1350	0.7150
sc	scute	1625273_at	0.0425	0.7905	-0.0507	0.6734	0.0385	0.8336	-0.0749	0.8730	-0.0719	0.7011	0.0030	0.9881	-0.1397	0.8032	-0.0981	0.7105	0.0416	0.8969
---	---	1625274_at	0.1487	0.3897	-0.1187	0.3636	-0.1045	0.4840	-0.0078	0.9937	0.1976	0.3419	0.2053	0.2648	0.0677	0.8882	0.0703	0.7418	0.0026	0.9935
CG32037	CG32037	1625275_at	0.1181	0.7487	0.0793	0.7113	0.0671	0.7372	0.2167	0.7368	-0.0199	0.9636	-0.2366	0.3697	0.0962	0.9215	-0.2491	0.4450	-0.3454	0.3011
Eip71CD	methionine-S-sulfi	1625276_a_at	-0.5476	0.1425	-0.1993	0.8012	-0.8419	0.0231	0.2035	0.8076	1.0065	0.0091	0.8030	0.0151	0.7861	0.6903	1.3296	0.1087	0.5436	0.5247
CG32463	CG32463	1625277_at	0.1181	0.5607	0.0599	0.6187	0.0742	0.7687	0.0054	0.9956	-0.0692	0.7845	-0.0746	0.7393	-0.0158	0.9862	-0.0951	0.6949	-0.0793	0.7492
CG31004	CG31004	1625278_s_at	3.9453	0.0423	1.7227	0.1559	3.3701	0.0073	2.9438	0.2114	3.7191	0.0168	0.7753	0.5836	1.2338	0.7633	1.5056	0.3720	0.2718	0.9088
tup	islet	1625279_a_at	-0.2321	0.5150	0.0725	0.5221	0.1178	0.5749	-0.0613	0.9455	-0.3205	0.2072	-0.2593	0.2586	-0.0028	0.9990	0.0409	0.9380	0.0437	0.9227
amon	amontillado	1625280_at	-0.3914	0.1174	-0.0748	0.6913	-0.1901	0.3650	-0.2702	0.6338	-0.4614	0.0982	-0.1912	0.4707	-0.0340	0.9643	0.0561	0.8386	0.0901	0.6925
CG6191	CG6191	1625281_at	-0.0537	0.7483	0.0808	0.7879	0.0137	0.9677	0.0375	0.9696	-0.1893	0.4759	-0.2268	0.3282	0.1461	0.8940	0.0778	0.9009	-0.0684	0.9045
CG13722	CG13722	1625282_at	0.8014	0.0705	0.4531	0.1038	0.9961	0.0009	0.0598	0.9647	0.0951	0.8352	0.0353	0.9364	-0.1305	0.9357	-0.2060	0.7299	-0.0754	0.9161
mod(mdg4)	Modifier67.2	1625283_at	-0.4364	0.1896	-0.2836	0.4429	-0.3552	0.1567	-0.5093	0.3222	-0.3340	0.2668	0.1753	0.5513	-0.0469	0.9742	0.0214	0.9723	0.0683	0.8918
CG13605	CG13605	1625284_at	0.3500	0.0464	0.3658	0.0574	0.6881	0.0022	0.1184	0.7447	0.0673	0.7166	-0.0511	0.7720	-0.1711	0.7810	0.1464	0.6053	0.3174	0.2497
---	---	1625285_at	-0.0258	0.9456	-0.2628	0.3640	-0.3761	0.2304	-0.1822	0.7880	0.1769	0.5549	0.3591	0.1477	0.0169	0.9928	-0.0703	0.9158	-0.0872	0.8818
---	---	1625286_at	-0.0347	0.8964	0.0759	0.5275	0.0754	0.7371	0.0340	0.9603	-0.0092	0.9743	-0.0432	0.8378	-0.0455	0.9734	0.0674	0.8918	0.1128	0.7764
CG40178	CG40178	1625287_at	-0.8163	0.0985	0.0967	0.8217	0.1127	0.6169	-0.0400	0.9761	-0.8991	0.0146	-0.8592	0.0110	-0.0261	0.9914	-0.1136	0.8919	-0.0875	0.9109
CG33177	CG33177	1625288_at	0.0855	0.6430	0.1100	0.5205	0.2129	0.4576	0.1019	0.9034	-0.0325	0.9323	-0.1344	0.6276	-0.1175	0.9101	-0.0709	0.8997	0.0466	0.9282
---	---	1625289_at	-0.0770	0.6345	0.0581	0.7843	0.0910	0.6162	-0.0343	0.9592	-0.0551	0.7998	-0.0208	0.9240	-0.0248	0.9778	0.0068	0.9875	0.0316	0.9147
---	---	1625290_at	-0.0567	0.7171	-0.2585	0.2763	-0.4283	0.0129	-0.2001	0.5353	0.1901	0.2606	0.3902	0.0177	0.0674	0.9489	0.0056	0.9935	-0.0618	0.8864
CG18766	CG18766	1625291_s_at	-0.9545	0.0051	-1.1584	0.0074	-1.2174	0.0006	-0.1143	0.8776	0.0387	0.9134	0.1530	0.5385	-0.1290	0.8603	-0.2162	0.4696	-0.0872	0.8111
CG12200 /// DsimCG12200	CG12200	1625292_at	0.1238	0.5913	-0.0203	0.8573	0.0283	0.9122	0.2046	0.5455	0.2180	0.2128	0.0134	0.9517	0.0326	0.9717	-0.0271	0.9429	-0.0597	0.8419
---	---	1625293_at	-4.1956	0.0005	-4.9087	0.0007	-4.1165	0.0001	0.8536	0.3357	0.1930	0.7513	-0.6607	0.1477	-0.1241	0.9320	-0.3987	0.3893	-0.2746	0.5830
ng3	new glue 3	1625294_at	0.2923	0.2084	0.1154	0.3252	0.0392	0.8437	0.0314	0.9599	0.1236	0.4714	0.0922	0.5712	0.1799	0.7644	0.0349	0.9325	-0.1450	0.5991
ksr	Kinase suppresso	1625295_at	-0.8033	0.1737	0.1064	0.8833	0.0461	0.8610	0.0808	0.9166	-0.3767	0.1151	-0.4575	0.0395	0.2337	0.9342	0.4844	0.6191	0.2507	0.8256
CG9508	CG9508	1625296_at	0.2511	0.1923	0.2226	0.3582	0.2190	0.1592	0.0886	0.8817	0.0603	0.8104	-0.0282	0.9086	-0.0281	0.9742	0.0924	0.7072	0.1205	0.6065
CG8128 /// DmirCG8128	CG8128	1625297_at	0.4732	0.0575	1.0848	0.0114	1.5194	0.0001	0.0707	0.9136	-1.1036	0.0008	-1.1742	0.0004	-0.3159	0.6695	-0.3523	0.2212	-0.0364	0.9310

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG34406	NAT1	1625298_at	0.2338	0.4173	-0.0820	0.8954	0.0804	0.7655	-0.0518	0.9683	0.2738	0.4361	0.3255	0.2891	-0.2645	0.7779	-0.0279	0.9665	0.2366	0.5856
CG33515	CG33515	1625299_at	0.1312	0.3825	-0.0600	0.6033	0.1152	0.4998	0.1517	0.7140	0.1851	0.3347	0.0334	0.8814	-0.0487	0.9340	0.0279	0.9247	0.0766	0.7179
CG40166	CG40166	1625300_at	0.1728	0.4658	0.0494	0.7143	0.4830	0.0102	0.0749	0.8452	0.0861	0.5731	0.0112	0.9502	-0.2988	0.5134	-0.0160	0.9648	0.2828	0.2148
CG34109	CG34109	1625301_at	0.0617	0.7348	0.0368	0.8995	0.3496	0.0496	0.0466	0.9496	-0.2505	0.2214	-0.2971	0.1049	-0.0433	0.9717	-0.2125	0.5071	-0.1692	0.6130
Task6	Task6	1625302_at	-0.0326	0.8861	-0.0077	0.9749	0.0737	0.7241	-0.0253	0.9697	-0.1035	0.5767	-0.0782	0.6590	-0.1771	0.8076	-0.0281	0.9538	0.1490	0.6475
CG30054	CG30054	1625303_at	-0.0158	0.9250	-0.0264	0.9194	0.2067	0.2977	0.0308	0.9649	0.1644	0.3899	0.1336	0.4477	-0.1027	0.9168	0.1711	0.6425	0.2737	0.4322
---	---	1625304_s_at	-0.5287	0.1419	-1.4978	0.0111	-1.1342	0.0298	0.6802	0.2397	1.1494	0.0057	0.4692	0.1236	0.0971	0.9741	0.1376	0.9016	0.0405	0.9715
CG12663	CG12663	1625305_at	0.0159	0.9273	-0.0828	0.4664	0.0182	0.9180	0.1171	0.7134	0.1105	0.4657	-0.0066	0.9710	-0.0307	0.9779	-0.0667	0.8546	-0.0360	0.9211
CG17388	CG17388	1625306_at	0.1791	0.4892	0.1218	0.5998	0.1463	0.5418	-0.0716	0.9037	-0.1605	0.4195	-0.0889	0.6553	0.0397	0.9816	0.0215	0.9723	-0.0182	0.9752
sut2	sugar transporter	1625307_at	-2.9972	0.0017	-4.2178	0.0012	-3.5925	0.0002	0.7791	0.4639	1.6626	0.0110	0.8835	0.0864	0.2699	0.8628	0.3437	0.6093	0.0739	0.9337
---	---	1625308_at	0.0606	0.7915	-0.0012	0.9962	-0.2708	0.1378	-0.2273	0.5932	0.1161	0.6294	0.3434	0.0770	-0.0551	0.9421	-0.1342	0.5888	-0.0790	0.7749
---	---	1625309_at	-0.1447	0.3802	0.1403	0.4477	0.0449	0.8150	0.0880	0.8432	0.0417	0.8450	-0.0464	0.8027	0.1232	0.8744	0.3617	0.2217	0.2385	0.4518
CG5235	CG5235	1625310_at	-0.1955	0.4494	-0.3119	0.2671	-0.8408	0.0496	-0.7294	0.2018	0.3479	0.3181	1.0773	0.0047	-0.2301	0.8513	-0.0568	0.9451	0.1732	0.7728
---	---	1625311_at	0.0482	0.8337	0.0781	0.6302	0.0372	0.8642	-0.1118	0.8551	-0.1060	0.6744	0.0058	0.9836	0.0601	0.9320	-0.0051	0.9925	-0.0652	0.8178
CG10553	CG10553	1625312_at	-3.2338	0.0033	-2.1131	0.0562	-4.2236	0.0001	-1.4941	0.0965	-3.3349	0.0004	-1.8407	0.0028	0.5602	0.8465	-2.1852	0.0807	-2.7454	0.0600
CG12826	CG12826	1625313_at	0.5850	0.4455	-0.0480	0.7234	-0.0866	0.7936	0.1558	0.7941	0.2503	0.3086	0.0945	0.7144	0.0525	0.9914	-0.4668	0.6881	-0.5193	0.6419
CG2022	CG2022	1625314_at	-2.0549	0.0012	-1.7999	0.0083	-2.1319	0.0002	0.3626	0.5357	-0.2954	0.3425	-0.6580	0.0242	0.1841	0.8480	-0.2218	0.6028	-0.4059	0.3205
CG4858	CG4858	1625315_at	0.2217	0.3454	0.0137	0.9389	0.2013	0.2619	-0.1456	0.7663	-0.4344	0.0436	-0.2888	0.1204	-0.2497	0.7392	-0.4971	0.1224	-0.2474	0.4523
---	---	1625316_s_at	-0.6209	0.2708	0.1607	0.7272	0.2871	0.3856	-0.3117	0.6591	-0.6718	0.0526	-0.3601	0.2366	-0.3502	0.8444	0.1121	0.9221	0.4622	0.5499
CG3735	CG3735	1625317_at	0.1399	0.5903	0.1183	0.8166	0.1951	0.2924	0.1157	0.8327	0.3855	0.0673	0.2698	0.1459	-0.0361	0.9884	0.1861	0.7787	0.2222	0.7150
---	---	1625318_at	-0.0073	0.9729	0.1913	0.3910	0.4024	0.0347	-0.0268	0.9610	0.0082	0.9711	0.0349	0.8338	-0.0806	0.9400	0.0771	0.8678	0.1577	0.6577
---	---	1625319_at	0.0320	0.8558	0.0290	0.8537	-0.0191	0.9124	0.1271	0.8074	0.0886	0.7121	-0.0386	0.8753	0.1616	0.7152	0.0814	0.6955	-0.0802	0.6977
CG1738 /// DereCG1738 ///	CG1738	1625320_at	0.0640	0.7169	-0.0203	0.9659	-0.1818	0.3313	-0.1417	0.8342	0.0371	0.9178	0.1788	0.4618	-0.0101	0.9950	0.1293	0.7787	0.1393	0.7482
CG7549	CG7549	1625321_a_at	-2.4100	0.0015	-3.3119	0.0049	-2.5071	0.0004	0.6447	0.5098	0.8730	0.0924	0.2282	0.6720	-0.1415	0.9309	-0.0866	0.9158	0.0549	0.9412
sgg	Glycogen Synthas	1625322_at	-0.0467	0.7600	0.0000	1.0000	0.0948	0.6008	0.1160	0.7576	0.0314	0.8851	-0.0846	0.6027	-0.0951	0.8461	0.0125	0.9704	0.1075	0.6225
CG32397	CG32397	1625323_at	-1.1515	0.0075	0.3401	0.5385	-0.3286	0.2836	-0.7393	0.3053	-1.8724	0.0017	-1.1331	0.0095	-0.1636	0.8973	-0.3375	0.4854	-0.1739	0.7495
CG18549	CG18549	1625324_at	-0.4858	0.1756	0.1107	0.7802	-0.5178	0.0097	-0.4532	0.3016	-0.5908	0.0296	-0.1377	0.5944	0.2278	0.8609	-0.0157	0.9887	-0.2436	0.6719
simj	simjang	1625325_s_at	0.4749	0.2861	-0.2167	0.7597	-0.6048	0.0334	0.0755	0.9533	0.9174	0.0179	0.8418	0.0161	0.4736	0.8202	0.2398	0.8331	-0.2338	0.8268
CG14959	CG14959	1625326_a_at	0.4308	0.2479	2.5308	0.0039	0.6440	0.0982	-2.2269	0.0153	-3.1056	0.0004	-0.8787	0.0436	0.0249	0.9916	-0.7565	0.1553	-0.7814	0.1792
---	---	1625327_at	0.2546	0.1658	0.0000	1.0000	-0.0682	0.7445	-0.1259	0.7451	0.1409	0.4223	0.2668	0.0804	-0.0030	0.9978	-0.0856	0.7787	-0.0826	0.7774
---	---	1625328_s_at	0.1184	0.4510	0.0223	0.8257	-0.0140	0.9548	-0.0139	0.9863	0.1681	0.4034	0.1820	0.3061	-0.0035	0.9964	0.0439	0.8527	0.0474	0.8256
CG12667	CG12667	1625329_at	-0.0242	0.9116	-0.2965	0.4514	-0.2315	0.1483	-0.0191	0.9857	0.0646	0.8344	0.0837	0.7502	-0.0204	0.9816	-0.0800	0.7421	-0.0596	0.8173
CG12126	CG12126	1625330_at	-0.0741	0.7159	0.0471	0.6935	0.0863	0.6344	-0.1994	0.6338	-0.1649	0.4383	0.0344	0.8882	-0.0818	0.8941	-0.0199	0.9587	0.0619	0.8307
---	---	1625331_at	0.1033	0.5282	-0.0441	0.7275	0.0236	0.8915	0.1171	0.7121	0.2011	0.1613	0.0840	0.5539	-0.0866	0.8692	-0.0822	0.7371	0.0044	0.9902
CG14764	CG14764	1625332_at	-1.1823	0.0194	0.3417	0.5277	-0.7609	0.0179	-0.8966	0.1408	-0.4279	0.2464	0.4687	0.1536	0.2604	0.8689	0.9878	0.1178	0.7274	0.2597
CG33172	CG33172	1625333_at	0.1294	0.6030	-0.6730	0.0420	-0.2134	0.4695	0.0049	0.9956	0.9711	0.0030	0.9662	0.0018	-0.4947	0.6955	0.1500	0.8262	0.6447	0.2306
NPC2	Epididymal secret	1625334_at	-0.4423	0.2532	-0.7086	0.2298	-0.0268	0.9393	0.3069	0.5044	0.2820	0.2489	-0.0249	0.9344	-0.3264	0.8479	0.0145	0.9929	0.3409	0.6569
CG14408 /// DsimCG14408	CG14408	1625335_at	0.3568	0.1056	0.1449	0.3162	0.2877	0.1571	0.0761	0.9011	0.0707	0.7669	-0.0054	0.9831	0.0697	0.9340	-0.0021	0.9981	-0.0718	0.8323
---	---	1625336_s_at	2.6584	0.0161	1.1493	0.0067	4.7622	0.0000	3.1652	0.0417	1.2942	0.1361	-1.8710	0.0245	0.1317	0.8479	0.0972	0.7831	-0.0345	0.9316
RpL32	r-protein 49	1625337_s_at	0.5544	0.0149	2.7435	0.0056	2.4918	0.0005	0.0558	0.9696	-1.9047	0.0010	-1.9604	0.0005	0.1239	0.8869	0.0255	0.9621	-0.0983	0.8047
Cyp4d20	Cyp4d20	1625338_at	0.2897	0.1169	0.4429	0.5297	0.3883	0.4424	-0.3534	0.7161	-0.5563	0.2072	-0.2029	0.6581	-0.0808	0.9816	-0.4450	0.5897	-0.3642	0.6672
CG31677	CG31677	1625339_at	0.1210	0.6003	0.1346	0.3341	0.2448	0.2756	0.0062	0.9952	0.0506	0.8543	0.0444	0.8578	0.0639	0.9216	0.1939	0.3625	0.1300	0.5724
nonA	no-on-transient A	1625340_at	-0.6557	0.0128	-1.8287	0.0192	-1.8897	0.0004	-0.2818	0.6338	0.2293	0.4474	0.5111	0.0534	-0.0971	0.9589	-0.7103	0.1694	-0.6132	0.2662
FucTC	alpha1,3-fucosyltr	1625341_at	-0.0858	0.6258	-0.0687	0.4827	0.0651	0.7487	-0.0921	0.7973	-0.0811	0.6130	0.0109	0.9529	-0.0937	0.9063	-0.0059	0.9929	0.0878	0.7995
CG33282	CG33282	1625342_at	0.0539	0.9629	0.1057	0.6071	0.1432	0.5111	0.0092	0.9979	-0.4771	0.6251	-0.4863	0.5788	-0.0243	0.9928	-0.1818	0.8175	-0.1575	0.8376
CG5931	CG5931	1625343_at	0.3026	0.1209	0.6647	0.2065	1.0677	0.0021	0.4426	0.3732	0.2972	0.2936	-0.1454	0.6077	0.0178	0.9928	0.6148	0.1634	0.5970	0.2094
fu12	fu12	1625344_at	1.6371	0.0082	1.1213	0.0363	2.2697	0.0002	0.4444	0.5863	0.1399	0.7815	-0.3045	0.4365	-0.6404	0.6955	-0.2316	0.7804	0.4087	0.5676
CG18508	CG18508	1625345_at	0.4520	0.1215	0.4774	0.0726	0.3502	0.0379	-0.0581	0.9466	-0.2072	0.4117	-0.1491	0.5330	0.1224	0.8653	-0.0493	0.9111	-0.1717	0.5746
---	---	1625346_s_at	0.0071	0.9753	-0.5894	0.0833	-0.4681	0.0310	0.0081	0.9943	0.4996	0.0434	0.4915	0.0295	-0.0006	0.9998	0.0271	0.9276	0.0277	0.9158
CG14829	CG14829	1625347_at	0.5210	0.0554	0.3467	0.0721	0.3401	0.1213	0.0073	0.9943	-0.0369	0.9030	-0.0442	0.8629	0.0395	0.9592	-0.1916	0.3797	-0.2311	0.3089
CG8892	CG8892	1625348_s_at	-0.0261	0.9267	-0.1399	0.5699	-0.0527	0.8520	0.0285	0.9603	0.2061	0.1745	0.1776	0.1921	-0.0809	0.9568	0.1537	0.7667	0.2346	0.6105
beat-IIIc	beat-IIIc	1625349_at	-1.2619	0.0036	-1.7019	0.0114	-1.1747	0.0285	0.3575	0.7401	0.2313	0.6685	-0.1262	0.8135	-0.1427	0.8033	-0.1511	0.5434	-0.0084	0.9837
---	---	1625350_at	0.0603	0.7909	0.2482	0.1413	0.1713	0.5625	-0.1701	0.7515	-0.2810	0.2275	-0.1109	0.6436	-0.0308	0.9741	-0.0256	0.9460	0.0051	0.9891
CG15572	CG15572	1625351_at	0.0290	0.9042	0.0222	0.8230	0.0810	0.6631	-0.0510	0.9601	-0.1033	0.7434	-0.0523	0.8669	-0.0100	0.9922	-0.0714	0.8119	-0.0614	0.8334
---	---	1625352_s_at	-0.8434	0.0112	-0.0579	0.8815	0.3294	0.1633	0.0560	0.9314	-0.2552	0.1854	-0.3112	0.0744	-0.3462	0.7768	0.4444	0.3922	0.7905	0.1574
CG34127	CG34127	1625353_at	-0.1084	0.4867	0.3075	0.0966	0.4669	0.0210	-0.0289	0.9677	0.0098	0.9734	0.0387	0.8599	-0.1067	0.9056	0.1476	0.6820	0.2543	0.4444
alpha-Est3	fragment B	1625354_at	-0.0533	0.8666	-0.1103	0.4301	0.3002	0.2374	-0.0494	0.9423	-0.7144	0.0038	-0.6649	0.0031	-0.4586	0.6879	-0.7594	0.1089	-0.3008	0.5403
---	---	1625355_at	0.2067	0.3053	0.0422	0.6933	-0.0279	0.9338	-0.0979	0.9029	0.0004	0.9992	0.0983	0.7228	0.2275	0.7215	0.0741	0.8353	-0.1534	0.5906
---	---	1625356_at	-0.0422	0.7915	-0.0029	0.9828	0.1203	0.5764	-0.0511	0.9345	-0.0468	0.8437	0.0042	0.9856	-0.0241	0.9816	0.0535	0.8812	0.0776	0.7964
---	---	1625357_at	0.2204	0.1466	0.1850	0.3662	0.3047	0.0738	0.0402	0.9441	0.0350	0.8718	-0.0052	0.9799	0.0118	0.9914	0.0442	0.9008	0.0324	0.9197
szar	CG9326	1625358_s_at	-1.6822	0.0060	0.9172	0.1313	0.4403	0.3123	-0.1465	0.8817	-1.6419	0.0011	-1.4953	0.0010	0.3305	0.8999	0.9659	0.3052	0.6354	0.5335
CG31281	CG31281	1625359_at	0.2381	0.2863	0.1390	0.2819	0.2029	0.3318	0.0365	0.9626	0.0616	0.8092	0.0251	0.9209	-0.0626	0.9296	-0.1084	0.6686	-0.0458	0.8844
CG33458	CG33458	1625360_at	-0.9126	0.1437	-1.4386	0.0248	-1.5482	0.0001	-0.1455	0.9036	0.2005	0.6464	0.3460	0.3301	-0.0091	0.9959	-0.2384	0.5842	-0.2292	0.6040
---	---	1625361_at	0.3005	0.2052	0.2482	0.0818	0.3064	0.1638	-0.0374	0.9678	-0.1812	0.4778	-0.1438	0.5486	0.0021	0.9990	-0.1746	0.5266	-0.1766	0.5276
Gclc	Glutamate-cystein	1625362_at	0.7359	0.0111	0.0586	0.9181	1.0880	0.0022	0.3655	0.3772	-0.2907	0.2129	-0.6562	0.0077	-0.6421	0.6272	-0.9373	0.0972	-0.2952	0.6195
Obp56i	Odorant-binding p	1625363_at	0.0712	0.7849	0.1533	0.2476	0.1207	0.4796	0.0269	0.9715	-0.0766	0.7302	-0.1035	0.5849	0.0248	0.9816	-0.0189	0.9621	-0.0437	0.8949
yellow-k	yellow-k	1625364_at	0.1063	0.5918	0.1611	0.3294	0.0775	0.7178	-0.0026	0.9968	0.0549	0.8308	0.0576	0.7996	0.0449	0.9593	0.0754	0.8185	0.0306	0.9316
CG17005	CG17005	1625365_at	0.1214	0.6043	-0.0069	0.9569	0.2278	0.3119	0.4067	0.2289	0.2874	0.1510	-0.1193	0.5450	0.0047	0.9964	0.1285	0.6322	0.1238	0.6434
rst	irregular chiasm C	1625366_at	-0.7024	0.1856	-0.5880	0.2224	-1.4481	0.0004	-0.8646	0.3048	-0.7801	0.1142	0.0845	0.8859	-0.0220	0.9928	-0.6267	0.2430	-0.6047	0.2917

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9448	CG9448	1625367_at	-0.3206	0.4851	-0.3732	0.3217	-0.7946	0.0066	-0.2604	0.6338	0.0615	0.8616	0.3219	0.1794	0.1210	0.9527	-0.0838	0.9315	-0.2048	0.7759
---	---	1625368_at	0.0537	0.7739	-0.1504	0.4191	-0.1680	0.2977	-0.0077	0.9937	0.1518	0.4400	0.1595	0.3604	0.0786	0.9142	0.0067	0.9908	-0.0719	0.8233
CG10877	CG10877	1625369_at	0.7551	0.3530	-0.2611	0.1830	0.4279	0.0897	0.2074	0.5735	0.2596	0.1655	0.0522	0.8047	-0.4282	0.8655	-0.7080	0.4943	-0.2798	0.8247
gem	persephone	1625370_s_at	-0.6333	0.0562	-0.2031	0.5147	-0.7889	0.0155	-0.5628	0.1586	-0.6434	0.0159	-0.0806	0.7610	0.0799	0.9760	-0.2253	0.7730	-0.3051	0.6602
CG13048	CG13048	1625371_at	0.1786	0.3930	0.0736	0.5971	-0.1030	0.5301	0.0919	0.8280	0.1154	0.4998	0.0235	0.9042	0.2192	0.7500	-0.0306	0.9492	-0.2498	0.4037
Gr98b	Gustatory recepto	1625372_at	0.0128	0.9673	0.1682	0.2451	-0.1595	0.5191	-0.3017	0.5756	-0.3830	0.1635	-0.0813	0.7929	0.0615	0.9445	-0.1175	0.6962	-0.1790	0.5233
CG6966	CG6966	1625373_at	-0.1384	0.7261	0.2782	0.3972	-0.3334	0.1146	-0.2460	0.5933	-0.3315	0.1545	-0.0855	0.7337	0.3295	0.8049	0.0241	0.9819	-0.3055	0.6128
CG9466	CG9466	1625374_at	0.0364	0.9315	0.0603	0.8178	0.0024	0.9942	0.0848	0.9592	0.0445	0.9447	-0.0403	0.9404	-0.0727	0.9589	-0.0078	0.9935	0.0649	0.9057
CG32201	CG32201	1625375_at	0.0243	0.9250	-0.0801	0.4578	0.0462	0.8654	0.1103	0.8738	0.1332	0.6187	0.0229	0.9393	-0.2938	0.6660	0.0761	0.8367	0.3699	0.2028
---	---	1625376_at	0.0667	0.7833	-0.0193	0.8645	0.0225	0.9356	0.1887	0.7351	0.0308	0.9296	-0.1579	0.5096	0.0054	0.9963	-0.1016	0.7351	-0.1070	0.7134
CG5104	CG5104	1625377_at	-0.2230	0.3184	-0.1734	0.5105	-0.2317	0.1338	-0.0039	0.9956	-0.1363	0.4931	-0.1325	0.4606	-0.0362	0.9816	-0.1978	0.6011	-0.1616	0.6783
---	---	1625378_at	-0.0090	0.9732	0.1227	0.3640	-0.2091	0.4935	-0.0882	0.9255	0.0461	0.9073	0.1343	0.6469	0.1137	0.8362	0.1451	0.5404	0.0313	0.9198
CG40091	CG40091	1625379_a_at	0.1567	0.3923	0.0959	0.4935	0.0819	0.6106	0.0278	0.9610	0.0526	0.7751	0.0248	0.8919	0.0576	0.9324	0.0008	0.9994	-0.0568	0.8378
CG15712	CG15712	1625380_at	0.2201	0.1814	-0.1525	0.4272	-0.0287	0.8846	-0.0569	0.9028	0.0666	0.7015	0.1234	0.3760	-0.1720	0.7893	-0.2247	0.4094	-0.0527	0.8891
Eig71Ed	Gene IV	1625381_at	0.0373	0.8588	0.1049	0.4772	-0.0043	0.9882	0.0058	0.9956	-0.0491	0.8712	-0.0549	0.8338	-0.0596	0.9705	-0.0122	0.9887	0.0475	0.9342
Osi6	Osiris	1625382_at	0.2485	0.2953	-0.3538	0.0429	-0.3779	0.0350	-0.0313	0.9649	0.3813	0.0497	0.4126	0.0233	-0.0498	0.9522	-0.2840	0.2394	-0.2342	0.3621
---	---	1625383_at	-0.2514	0.1588	-0.1057	0.4464	-0.4566	0.0279	0.0827	0.8380	0.0972	0.5516	0.0145	0.9386	0.1406	0.8270	0.0325	0.9402	-0.1081	0.7221
---	---	1625384_at	0.2956	0.3525	0.1722	0.2910	0.2322	0.1931	-0.1334	0.7018	-0.1646	0.3076	-0.0312	0.8676	0.0789	0.9550	0.0351	0.9589	-0.0438	0.9392
ct	cut	1625385_at	-0.8458	0.0260	-1.3033	0.0138	-1.6137	0.0001	-0.3778	0.4279	0.3893	0.1366	0.7671	0.0063	-0.2328	0.7498	-0.2457	0.4294	-0.0129	0.9802
da	Daughterless	1625386_at	0.2693	0.6110	-0.0361	0.9468	-0.8150	0.0864	-0.1026	0.7690	0.6751	0.0015	0.7777	0.0005	0.6626	0.7855	0.3713	0.7707	-0.2913	0.8243
E2f	E2F transcription	1625387_s_at	0.6172	0.2130	0.1295	0.7167	-0.1763	0.4797	-0.0733	0.8705	0.4775	0.0089	0.5509	0.0028	0.4015	0.8270	0.1088	0.9282	-0.2927	0.7415
CG4893	CG4893	1625388_at	-0.6192	0.0121	-0.6511	0.0554	-0.7287	0.0035	0.0634	0.9080	0.0613	0.7722	-0.0021	0.9922	-0.0215	0.9852	-0.0464	0.9085	-0.0249	0.9466
CG8209	CG8209	1625389_at	-0.5001	0.1318	0.6049	0.0335	1.0427	0.0002	0.4328	0.1754	-0.4947	0.0188	-0.9275	0.0007	-0.0079	0.9952	0.5314	0.1009	0.5392	0.1192
CG40090	CG40090	1625390_at	-0.1921	0.3635	-0.1150	0.4810	-0.3483	0.0481	-0.0402	0.9586	0.1134	0.6149	0.1536	0.4219	-0.0730	0.9238	0.1402	0.6041	0.2132	0.4083
---	---	1625391_at	-0.1016	0.4810	-0.0113	0.9156	0.1906	0.2174	0.1020	0.8475	0.0374	0.8865	-0.0646	0.7614	0.0453	0.9421	0.0874	0.6844	0.0421	0.8699
mRpL42	mitochondrial ribo	1625392_at	0.0394	0.8831	-0.0998	0.7573	0.0849	0.6957	-0.0129	0.9880	0.1281	0.5542	0.1410	0.4588	-0.0658	0.9705	0.1642	0.7625	0.2300	0.6367
CG5274	CG5274	1625393_s_at	0.0551	0.8053	-0.3385	0.1601	-0.4715	0.0466	-0.0340	0.9632	0.3202	0.1014	0.3543	0.0480	0.1667	0.8653	0.0228	0.9725	-0.1439	0.7495
CG9922	CG9922	1625394_at	0.2186	0.2576	0.4834	0.2538	0.3524	0.1096	0.2012	0.6454	0.3732	0.0797	0.1720	0.3806	0.3561	0.7506	0.7234	0.1336	0.3673	0.4597
CG5115	CG5115	1625395_at	-0.0552	0.8106	0.1219	0.3055	0.1141	0.6628	-0.0698	0.8841	-0.1114	0.5207	-0.0416	0.8223	-0.0085	0.9939	0.1128	0.6949	0.1214	0.6609
---	---	1625396_at	0.1925	0.2668	0.0270	0.8994	0.1401	0.5018	0.2220	0.4509	0.1736	0.2804	-0.0485	0.7865	-0.0497	0.9706	-0.0019	0.9989	0.0477	0.9194
---	---	1625397_at	0.0652	0.7292	-0.1189	0.4096	0.2766	0.0994	0.0366	0.9518	0.1703	0.3168	0.1338	0.3926	-0.0827	0.8650	-0.0352	0.9058	0.0476	0.8487
Fcp26Ac	Follicle cell proteir	1625398_at	0.2662	0.1164	0.0356	0.7488	0.1704	0.2405	0.0417	0.9322	0.1424	0.3360	0.1007	0.4664	0.0391	0.9621	-0.0826	0.7667	-0.1217	0.6225
CG6978	CG6978	1625399_at	0.1384	0.4991	0.0172	0.9364	0.1229	0.6684	-0.0038	0.9956	-0.0348	0.8996	-0.0310	0.8978	-0.0108	0.9923	-0.0795	0.8117	-0.0686	0.8315
Cpr47Ea	CG9079	1625400_at	0.0705	0.6825	0.4669	0.0632	0.5090	0.0180	0.0227	0.9819	-0.1037	0.7033	-0.1263	0.5928	-0.0514	0.9340	0.0712	0.7656	0.1226	0.5584
CG13026	CG13026	1625401_at	0.0018	0.9958	-0.0310	0.7623	-0.0756	0.6726	-0.0235	0.9838	0.0019	0.9960	0.0254	0.9385	-0.1140	0.8810	0.1122	0.7424	0.2261	0.4504
Ugt36Bb	Ugt36Bb	1625402_at	0.1998	0.2381	-0.0596	0.5977	0.0616	0.7247	-0.0125	0.9864	0.3318	0.0652	0.3443	0.0370	-0.0900	0.8903	0.0032	0.9952	0.0932	0.7416
Or49b	Odorant receptor	1625403_at	0.0211	0.9258	-0.0447	0.6795	-0.0253	0.8812	0.0549	0.9300	0.0922	0.6556	0.0373	0.8612	-0.1022	0.8655	0.0171	0.9653	0.1193	0.6463
---	---	1625404_at	-0.0249	0.8976	-0.1053	0.4952	-0.1176	0.6458	0.0298	0.9620	0.0574	0.7727	0.0277	0.8884	0.0189	0.9862	-0.1257	0.6332	-0.1446	0.5781
yellow-c	yellow-c	1625405_at	-3.1873	0.0103	-4.7115	0.0047	-4.6077	0.0000	0.2920	0.6597	0.9507	0.0095	0.6587	0.0285	0.1197	0.9824	-0.6054	0.6601	-0.7250	0.5873
---	---	1625406_s_at	-0.1081	0.5674	0.2322	0.1599	0.3160	0.3825	-0.0183	0.9840	-0.2998	0.1577	-0.2815	0.1380	0.0236	0.9898	-0.1359	0.7824	-0.1595	0.7251
CG40336	CG40336	1625407_at	-0.0686	0.6638	0.0114	0.9441	0.1365	0.3573	-0.0009	0.9988	0.1179	0.4725	0.1188	0.4184	0.0427	0.9571	0.1828	0.4151	0.1402	0.5585
CG10148	CG10148	1625408_at	-0.0754	0.6547	-0.0129	0.9476	-0.0123	0.9492	-0.0850	0.8717	-0.2104	0.2524	-0.1254	0.4719	-0.1981	0.7464	-0.0992	0.7439	0.0989	0.7409
CG17086	CG17086	1625409_at	-0.4316	0.0506	-0.2081	0.4610	-0.0037	0.9855	-0.0559	0.9436	-0.4832	0.0388	-0.4273	0.0406	-0.0308	0.9717	-0.0303	0.9296	0.0004	0.9992
CG5147	CG5147	1625410_at	0.0319	0.9173	0.9517	0.0381	1.4366	0.0006	0.1418	0.7705	-0.3684	0.0754	-0.5102	0.0133	-0.2188	0.8326	0.5240	0.2102	0.7428	0.1190
tectonic	tectonic	1625411_at	-0.1781	0.3720	-0.1805	0.4346	-0.0763	0.8349	0.0557	0.9413	-0.1753	0.4358	-0.2310	0.2344	0.0751	0.9514	-0.1411	0.7439	-0.2162	0.5808
CG4769	CG4769	1625412_at	-0.5810	0.0334	-0.1655	0.4076	-0.2810	0.2253	-0.1163	0.7929	-0.6551	0.0040	-0.5388	0.0057	-0.0697	0.9555	-0.3166	0.3760	-0.2469	0.5167
CG33490	CG33490	1625413_s_at	-0.0309	0.8711	-0.0090	0.9494	0.0027	0.9910	0.0928	0.8738	0.0152	0.9604	-0.0776	0.7214	0.0917	0.9046	-0.0108	0.9841	-0.1024	0.7484
ifc	degenerative sper	1625414_at	-1.4745	0.0010	-1.5862	0.0136	-1.6335	0.0000	-0.0356	0.9688	-0.1635	0.5107	-0.1279	0.5864	0.0811	0.9400	-0.0980	0.8215	-0.1791	0.6166
CG14306	CG14306	1625415_at	0.0201	0.9230	0.1065	0.6190	0.2022	0.1790	-0.0393	0.9518	-0.0543	0.8094	-0.0150	0.9483	-0.0520	0.9405	0.1143	0.6218	0.1663	0.4564
grp	grapes	1625416_x_at	0.4234	0.0419	-0.1253	0.6930	-0.3139	0.2680	-0.0121	0.9863	0.6060	0.0041	0.6181	0.0022	0.0971	0.9530	0.0634	0.9353	-0.0338	0.9628
CG14374 /// CG14377	CG14374 /// CG1	1625417_s_at	0.2920	0.2232	-0.0432	0.6695	0.1948	0.3265	0.1320	0.8281	0.1634	0.5078	0.0315	0.9112	-0.1141	0.8689	-0.0884	0.7985	0.0257	0.9466
---	---	1625418_at	-0.0250	0.8940	0.3481	0.0376	0.2451	0.1246	-0.1818	0.6325	-0.1677	0.3781	0.0142	0.9522	-0.0520	0.9407	0.2820	0.1786	0.3339	0.1490
---	---	1625419_at	0.1333	0.5220	-0.2827	0.2992	0.0342	0.8398	0.0590	0.9339	0.3453	0.0987	0.2863	0.1239	-0.1570	0.8461	-0.1770	0.6243	-0.0200	0.9690
---	---	1625420_at	-0.0699	0.8440	-1.1451	0.0213	-1.2009	0.0223	-0.2393	0.8822	1.1253	0.0470	1.3647	0.0135	-0.1502	0.8656	-0.1013	0.8297	0.0489	0.9198
CG30377	CG30377	1625421_at	0.2560	0.6347	-0.3377	0.4587	-0.8043	0.0241	-0.6767	0.3817	0.3617	0.4235	1.0383	0.0160	-0.2736	0.8825	-0.2888	0.7221	-0.0152	0.9899
CG4925	CG4925	1625422_at	-0.3055	0.1543	0.0562	0.8250	0.4396	0.1887	0.0070	0.9922	-0.0454	0.8013	-0.0524	0.7409	-0.1503	0.8972	0.5677	0.1761	0.7180	0.1292
---	---	1625423_at	0.0622	0.7745	-0.0917	0.4761	0.0628	0.7290	0.1625	0.7695	0.0688	0.8128	-0.0937	0.7059	0.2466	0.6955	0.0547	0.8868	-0.1919	0.4863
CG17360	CG17360	1625424_at	0.3066	0.1381	-0.2383	0.2654	-0.2789	0.1737	-0.1328	0.8112	0.1057	0.6740	0.2385	0.2270	-0.0535	0.9514	-0.3788	0.1420	-0.3253	0.2332
RpS15	Ribosomal protein	1625425_a_at	0.1470	0.3461	0.4645	0.0413	0.6130	0.0121	0.1384	0.8074	-0.2778	0.2170	-0.4163	0.0449	0.1141	0.8386	0.0573	0.8569	-0.0568	0.8480
CG5846	CG5846	1625426_at	-0.0964	0.8012	0.1981	0.2292	0.2812	0.0807	-0.0653	0.9020	-0.6690	0.0030	-0.6037	0.0027	-0.1710	0.8541	-0.3907	0.2960	-0.2197	0.5930
CG12241	CG12241	1625427_at	0.0887	0.8489	1.1230	0.0200	1.2468	0.0004	0.1984	0.7721	-0.3483	0.2265	-0.5467	0.0403	0.0089	0.9964	0.6927	0.1468	0.6837	0.1865
CG13707	CG13707	1625428_at	-0.0626	0.9215	-0.1857	0.2754	-0.1984	0.2444	-0.3757	0.1332	-0.4214	0.0128	-0.0457	0.7868	-0.4636	0.7707	-0.6292	0.3421	-0.1655	0.8505
---	---	1625429_at	0.2330	0.3659	-0.0441	0.6984	0.3808	0.0458	-0.0241	0.9833	-0.0997	0.7482	-0.0757	0.7973	-0.2492	0.6749	-0.0629	0.8531	0.1862	0.4677
CG14022 /// DyakCG14022	CG14022	1625430_at	-0.9369	0.0588	0.5651	0.3490	-0.1522	0.8496	-0.7315	0.5633	-3.3177	0.0009	-2.5862	0.0016	0.2293	0.9474	-1.4689	0.1523	-1.6982	0.1389
repo	reverse polarity	1625431_at	-0.6551	0.2988	0.3919	0.3636	-0.5611	0.0406	-0.5593	0.6013	-1.3203	0.0228	-0.7610	0.1138	0.0246	0.9922	-0.4605	0.4158	-0.4850	0.4021
ths	thisbe	1625432_at	0.0461	0.8529	0.1014	0.4646	-0.0415	0.8305	-0.0732	0.8815	-0.0904	0.6274	-0.0172	0.9343	-0.0133	0.9905	0.0288	0.9404	0.0421	0.8982
Scp1	calcium biding pro	1625433_at	0.5162	0.3101	-0.3434	0.3993	0.3454	0.2327	0.2705	0.5680	0.2397	0.3294	-0.0308	0.9173	-0.6117	0.7772	-0.6071	0.5336	0.0046	0.9979
CG3270	CG3270	1625434_at	0.9799	0.2598	0.5968	0.1907	0.4514	0.2520	0.1477	0.9620	-0.9791	0.2340	-1.1267	0.1240	0.3949	0.8510	-1.7072	0.0676	-2.1020	0.0542
CG7920 /// DsimCG7920	CG7920	1625435_at	0.1733	0.4706	0.1785	0.2972	0.1501	0.4600	0.1196	0.8395	-0.0224	0.9453	-0.1420	0.5046	0.1670	0.7953	0.1685	0.5523	0.0015	0.9979

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Uro	Urate oxidase	1625436_at	-1.3527	0.4198	0.3199	0.7456	-0.2589	0.2538	0.2019	0.9728	-1.5573	0.2949	-1.7592	0.1806	0.8392	0.6898	0.4690	0.6125	-0.3702	0.7026
CG32164	CG32164	1625437_at	0.2395	0.1573	0.3458	0.3556	-0.1281	0.4731	-0.1004	0.8898	-0.0237	0.9478	0.0767	0.7754	0.3597	0.6557	0.0651	0.8919	-0.2946	0.3791
Cpr65Av	CG32405	1625438_at	0.0266	0.9066	0.3862	0.3882	-0.1436	0.6031	0.1184	0.9045	0.1947	0.5745	0.0763	0.8357	0.1468	0.8215	0.0962	0.7692	-0.0506	0.8918
Tsp29Fa	Tetraspanin 29Fa	1625439_at	2.3146	0.0045	1.3555	0.0082	2.4437	0.0000	0.8199	0.2989	0.8588	0.0674	0.0388	0.9473	-0.2959	0.7726	-0.0093	0.9928	0.2866	0.5416
CG15374	CG15374	1625440_at	0.0035	0.9886	0.3052	0.1571	0.2050	0.3071	-0.1579	0.6988	-0.2999	0.1083	-0.1420	0.4179	-0.0848	0.9168	-0.0902	0.8022	-0.0053	0.9914
CG12710	CG12710	1625441_at	0.3437	0.1267	0.4395	0.1340	0.1872	0.2457	-0.1183	0.7803	-0.0620	0.7695	0.0563	0.7725	0.1085	0.8889	0.1110	0.7475	0.0025	0.9963
shi	dynamin	1625442_a_at	-0.1540	0.3459	0.4355	0.3285	0.6905	0.0273	0.0413	0.9592	-0.2712	0.2017	-0.3125	0.1008	-0.2076	0.8564	0.2080	0.6937	0.4157	0.3859
CG33054	CG33054	1625443_a_at	-0.3112	0.2941	0.6644	0.0359	0.9851	0.0016	-0.1021	0.8479	-1.2786	0.0004	-1.1765	0.0003	-0.2592	0.7810	-0.1851	0.6812	0.0741	0.8934
CG13773	CG13773	1625444_at	0.3848	0.1675	0.0586	0.9247	0.2679	0.1043	0.2782	0.5496	0.4239	0.0795	0.1457	0.5345	-0.0259	0.9923	0.0831	0.9333	0.1090	0.8983
shn	quo vadis	1625445_s_at	0.2647	0.3297	-0.1340	0.7483	-0.6825	0.0090	-0.0718	0.8903	0.6994	0.0026	0.7712	0.0010	0.5183	0.6898	0.2729	0.6324	-0.2455	0.6706
l(1)G0045	lethal (1) G0045	1625446_at	-0.0751	0.6398	-0.1524	0.1746	0.1543	0.3609	0.1415	0.7368	0.0691	0.7562	-0.0724	0.7149	-0.0672	0.9333	0.1028	0.7299	0.1700	0.5295
CAP-D2	CAP-D2 condensi	1625447_at	0.2084	0.3527	-0.4946	0.0198	-0.4071	0.0663	0.0210	0.9774	0.7431	0.0023	0.7221	0.0016	-0.1200	0.8680	-0.0476	0.9145	0.0724	0.8434
---	---	1625448_at	0.2506	0.2302	-0.1942	0.3014	-0.0679	0.7257	0.1262	0.7711	0.3777	0.0450	0.2515	0.1229	-0.0407	0.9581	-0.0457	0.8838	-0.0050	0.9884
CG40339	CG40339	1625449_at	0.3374	0.0546	-0.0181	0.8638	-0.0598	0.7494	-0.0564	0.9245	0.2207	0.2135	0.2771	0.0827	-0.0779	0.9243	-0.0412	0.9231	0.0367	0.9229
mthl14	methuselah-like 1	1625450_at	-1.2309	0.0879	-1.5185	0.0820	-1.7900	0.0001	-0.0336	0.9777	0.0777	0.8461	0.1113	0.7378	0.1004	0.9811	-0.1863	0.8868	-0.2867	0.7925
CG5060	CG5060	1625451_at	-0.0256	0.8836	0.0357	0.7473	0.3405	0.1061	0.0342	0.9647	-0.0169	0.9557	-0.0511	0.8265	-0.0300	0.9717	0.1244	0.5755	0.1543	0.4779
CG17075 /// DereCG17075	CG17075	1625452_at	0.1625	0.5423	0.0030	0.9842	0.1839	0.3654	0.1122	0.8485	-0.0355	0.9045	-0.1477	0.4690	-0.0270	0.9860	-0.1659	0.6624	-0.1389	0.7240
Sh3beta	Sh3beta	1625453_a_at	0.4421	0.0228	0.9490	0.0121	1.0686	0.0033	0.0769	0.8902	-0.0461	0.8487	-0.1229	0.4912	0.0269	0.9862	0.4535	0.1948	0.4266	0.2543
Gbeta5	Gbeta5	1625454_at	-1.4961	0.0071	-0.6121	0.0315	-1.1360	0.0002	-0.0122	0.9901	-0.4066	0.0673	-0.3944	0.0496	0.0782	0.9544	-0.2112	0.6242	-0.2894	0.4851
---	---	1625455_at	0.0757	0.6608	0.1660	0.3823	0.1237	0.4595	-0.0527	0.9249	-0.0101	0.9693	0.0427	0.8228	0.0971	0.8655	0.0516	0.8749	-0.0454	0.8850
---	---	1625456_at	0.0470	0.7632	0.0543	0.6546	-0.0378	0.8661	0.0191	0.9803	0.0864	0.6920	0.0673	0.7424	0.1083	0.8320	-0.0080	0.9842	-0.1163	0.6124
CG10031	CG10031	1625457_at	1.9353	0.0060	1.1073	0.1791	1.6083	0.0012	0.0362	0.9852	-0.2743	0.5609	-0.3105	0.4525	-0.4132	0.8305	-1.0181	0.1942	-0.6049	0.4686
msk	moleskin	1625458_a_at	0.4697	0.0880	0.6792	0.0297	0.4639	0.0076	0.1421	0.7094	0.4227	0.0231	0.2806	0.0728	0.2849	0.7571	0.6526	0.1012	0.3677	0.3572
CG3520	CG3520	1625459_at	0.3158	0.1398	0.6376	0.1894	1.4659	0.0002	0.2241	0.6319	-0.3568	0.1174	-0.5809	0.0121	-0.6267	0.5126	-0.0824	0.9064	0.5443	0.2501
dpr	defective proboscı	1625460_s_at	0.1211	0.4597	0.0732	0.5518	0.0446	0.7919	-0.1174	0.7701	0.0240	0.9200	0.1414	0.3675	-0.0455	0.9514	-0.0353	0.9192	0.0103	0.9769
---	---	1625461_at	0.3960	0.0487	0.0919	0.6223	0.3014	0.2128	0.0661	0.9339	0.0981	0.7196	0.0319	0.9103	-0.1449	0.8780	-0.1376	0.7504	0.0074	0.9905
CG31221	CG31221	1625462_s_at	-0.1482	0.3815	0.0294	0.7950	0.0845	0.6842	-0.1894	0.7134	-0.5541	0.0260	-0.3646	0.0832	-0.0514	0.9499	-0.1509	0.5539	-0.0995	0.7184
ort	Histamine-gated c	1625463_at	0.0261	0.8918	0.1898	0.3078	0.1870	0.3348	0.0121	0.9893	-0.0510	0.8501	-0.0631	0.7865	-0.0460	0.9635	-0.0174	0.9692	0.0286	0.9409
Marf	Drosophila mitofu	1625464_at	-0.2973	0.3231	0.3609	0.2173	0.3845	0.1633	-0.2587	0.4238	-1.0275	0.0006	-0.7688	0.0012	-0.3572	0.7707	-0.4127	0.4300	-0.0555	0.9396
CG6028	CG6028	1625465_a_at	-0.1746	0.5612	0.4880	0.0531	0.4951	0.0533	-0.2897	0.3381	-0.6693	0.0031	-0.3796	0.0231	-0.2857	0.7644	-0.0444	0.9457	0.2413	0.5698
CG2656	CG2656	1625466_at	0.2028	0.3866	0.3724	0.0634	0.3219	0.0845	-0.0018	0.9987	-0.0886	0.7380	-0.0868	0.7165	-0.0669	0.9588	-0.0564	0.9193	0.0105	0.9852
CG5860	CG5860	1625467_at	-0.2704	0.2815	0.0632	0.6710	0.1159	0.6389	-0.0101	0.9937	0.0865	0.7793	0.0967	0.7216	-0.1220	0.8841	0.1376	0.7012	0.2596	0.4241
CG9500	CG9500	1625468_at	0.0512	0.7373	0.1655	0.3793	0.0596	0.6882	-0.0811	0.8794	-0.1917	0.2989	-0.1106	0.5353	0.1731	0.6955	0.0089	0.9784	-0.1643	0.3807
CG3323	CG3323	1625469_at	0.1316	0.4531	0.0109	0.9187	0.1746	0.3406	0.0394	0.9375	0.1207	0.4218	0.0813	0.5733	-0.0225	0.9862	0.0139	0.9789	0.0364	0.9282
WASp	WASP	1625470_s_at	0.1080	0.8085	0.7348	0.1389	0.8812	0.0140	-0.1913	0.6854	-0.2941	0.1775	-0.1028	0.6494	-0.1469	0.9514	0.3539	0.6483	0.5008	0.4978
CG4928 /// DsimCG4928	CG4928	1625471_s_at	-0.0736	0.7705	0.0021	0.9965	-2.5543	0.0107	-1.9267	0.0082	-1.6026	0.0015	0.3241	0.3120	0.6466	0.8270	-1.2476	0.2964	-1.8942	0.1490
CG14515	CG14515	1625472_at	-0.3572	0.0955	0.2490	0.1716	0.4116	0.0274	-0.1119	0.8393	-1.2442	0.0005	-1.1323	0.0004	0.0159	0.9862	0.0593	0.8346	0.0434	0.8803
CG4221	CG4221	1625473_at	-2.5005	0.0010	-2.0343	0.0209	-1.6480	0.0012	0.3427	0.7328	-0.7546	0.0926	-1.0973	0.0140	-0.2039	0.9011	-0.3915	0.5317	-0.1876	0.7964
CG1441	CG1441	1625474_s_at	-0.3831	0.5339	-2.6622	0.0051	-1.5683	0.0084	0.8828	0.1956	1.5762	0.0032	0.6934	0.0634	-0.2559	0.9038	-0.5385	0.4954	-0.2825	0.7495
drk	Enhancer of sever	1625475_s_at	-0.3416	0.1202	-0.5470	0.0993	-0.8918	0.0020	-0.1719	0.7134	0.2508	0.2393	0.4227	0.0336	0.0400	0.9776	-0.0548	0.9170	-0.0948	0.8243
CG1674	CG1674	1625476_a_at	-2.5805	0.0043	-1.8942	0.0781	-2.4977	0.0000	-0.6485	0.1216	-1.3613	0.0008	-0.7128	0.0075	0.1035	0.9816	-0.7221	0.4550	-0.8256	0.3995
CG4797	CG4797	1625477_a_at	1.9572	0.0077	1.5373	0.0488	2.9185	0.0004	1.2869	0.1119	0.9770	0.0487	-0.3099	0.5065	-0.3011	0.8692	0.3762	0.6337	0.6773	0.3672
Mnn1	Menin 1	1625478_at	0.1309	0.7145	0.1594	0.2259	0.3866	0.1633	0.0283	0.9755	0.0596	0.8372	0.0313	0.9089	-0.1669	0.8650	0.0987	0.8546	0.2656	0.5151
---	---	1625479_at	0.0830	0.6338	0.1030	0.6030	0.2191	0.2074	0.0411	0.9602	-0.0075	0.9816	-0.0486	0.8459	-0.0980	0.8609	-0.1433	0.5431	-0.0452	0.8840
CG10671	CG10671	1625480_at	-0.5985	0.0493	0.3017	0.1385	0.8096	0.0016	0.1103	0.7604	-0.6852	0.0017	-0.7955	0.0006	-0.3056	0.7305	0.2459	0.5336	0.5515	0.1702
retn	dead ringer	1625481_a_at	-0.0343	0.9508	-0.7269	0.0625	-0.6690	0.0097	0.0696	0.9311	0.6603	0.0132	0.5907	0.0132	0.0808	0.9764	0.0713	0.9463	-0.0096	0.9934
---	---	1625482_s_at	0.2278	0.3158	-0.0591	0.6031	0.2110	0.3348	0.0438	0.9532	0.1652	0.4378	0.1214	0.5469	-0.1223	0.8940	-0.1145	0.7883	0.0078	0.9885
CG31080	CG31080	1625483_at	0.2308	0.1283	-0.0558	0.6020	0.1678	0.4412	0.0898	0.8544	0.0827	0.6814	-0.0071	0.9740	-0.0264	0.9816	0.0776	0.8061	0.1040	0.7072
---	---	1625484_at	0.1320	0.3512	0.0000	1.0000	-0.0967	0.6075	-0.1200	0.8189	0.1136	0.6162	0.2335	0.2019	-0.0096	0.9922	-0.0078	0.9857	0.0018	0.9965
---	---	1625485_at	0.0812	0.6700	0.1478	0.4911	0.1468	0.4706	-0.0090	0.9942	-0.0280	0.9393	-0.0189	0.9534	0.1009	0.8541	0.1032	0.6818	0.0023	0.9953
PGRP-LA	Peptidoglycan rec	1625486_a_at	1.8390	0.0004	0.2642	0.4964	1.0655	0.0512	-0.1247	0.7526	-0.2103	0.2180	-0.0857	0.6234	-1.0314	0.6035	-1.8749	0.0514	-0.8435	0.3419
CG6839	CG6839	1625487_at	0.2353	0.2140	0.3136	0.3969	0.4101	0.0802	0.1321	0.7922	0.0867	0.7134	-0.0454	0.8476	-0.0262	0.9862	0.1806	0.6312	0.2067	0.5767
Rho1	rho1 kinase	1625488_s_at	-0.3524	0.0982	0.1425	0.2362	-0.1099	0.4711	-0.2429	0.3887	-0.3825	0.0243	-0.1397	0.3375	0.0725	0.8882	0.2142	0.2520	0.1417	0.4824
---	---	1625489_at	0.1453	0.3805	-0.1278	0.3489	0.1961	0.2094	0.2292	0.4455	0.2184	0.1800	-0.0108	0.9587	-0.0520	0.9238	-0.1724	0.3268	-0.1204	0.5257
CG3500	CG3500	1625490_at	-0.4133	0.2068	0.3428	0.0695	0.7451	0.0194	0.1991	0.5754	-0.5500	0.0091	-0.7491	0.0014	-0.1789	0.8806	0.2291	0.6425	0.4079	0.3813
bwa	brain washing	1625491_at	0.8308	0.0230	-0.0934	0.7610	0.3596	0.4302	-0.2486	0.7447	-0.4230	0.2055	-0.1745	0.6061	-0.7711	0.6903	-1.5207	0.0717	-0.7495	0.3621
---	---	1625492_at	-0.0807	0.6193	0.0211	0.8449	-0.1597	0.2788	-0.1219	0.7906	0.1044	0.6131	0.2263	0.1756	-0.0257	0.9767	0.1552	0.4655	0.1809	0.3973
HLHm7	Enhancer of split	1625493_at	-0.0728	0.9046	0.2109	0.3216	0.2347	0.2471	0.0978	0.9480	-0.3363	0.4413	-0.4341	0.2509	0.0352	0.9779	-0.0613	0.8903	-0.0965	0.7918
CG4282	CG4282	1625494_at	0.2368	0.3448	0.3245	0.0947	0.2933	0.0786	-0.1500	0.7605	0.0131	0.9684	0.1631	0.4099	-0.0534	0.9457	0.0599	0.8546	0.1132	0.6604
CG12807	CG12807	1625495_at	0.1617	0.3522	-0.2276	0.1015	0.1531	0.4347	0.2075	0.4908	0.3295	0.0458	0.1220	0.4166	0.0670	0.9467	0.1132	0.7550	0.0462	0.9121
Ide	Insulin degrading	1625496_at	-0.0168	0.9249	-0.3004	0.4201	-0.4986	0.0071	-0.1004	0.7929	0.3063	0.0545	0.4066	0.0107	0.1007	0.9201	0.1394	0.7200	0.0387	0.9350
CG33280	CG33280	1625497_at	0.0913	0.5831	0.1157	0.3776	0.3042	0.3355	-0.0383	0.9602	-0.1524	0.4698	-0.1141	0.5685	-0.2228	0.7997	-0.0649	0.9080	0.1579	0.6993
CG13590	CG13590	1625498_at	-0.0783	0.6703	0.0171	0.9643	0.0362	0.8599	-0.0139	0.9857	0.0810	0.6861	0.0949	0.5880	-0.0236	0.9895	0.0210	0.9739	0.0447	0.9330
CG3809	CG3809	1625499_at	-0.1275	0.5419	0.1667	0.3646	0.2711	0.1093	0.0134	0.9857	-0.0986	0.5942	-0.1120	0.4914	0.0188	0.9878	0.3325	0.2126	0.3137	0.2718
Mlh1	Mlh1	1625500_at	-0.0218	0.9231	-0.0969	0.4476	0.0008	0.9972	-0.3105	0.3521	0.1277	0.5319	0.4382	0.0185	-0.2131	0.7464	-0.0156	0.9739	0.1974	0.4967
---	---	1625501_at	0.0912	0.6404	-0.0890	0.5424	-0.0116	0.9614	0.0798	0.8856	0.1308	0.5107	0.0510	0.8099	0.0048	0.9964	0.0326	0.9425	0.0278	0.9433
Ntf-2r	Drosophila nuclea	1625502_at	-0.0043	0.9844	-0.0960	0.4953	-0.3491	0.0518	-0.0048	0.9956	0.0136	0.9698	0.0184	0.9499	0.1035	0.8874	-0.0591	0.8804	-0.1626	0.5767
CG32425	CG32425	1625503_at	2.0995	0.0013	1.4432	0.0091	2.2648	0.0003	0.7526	0.1329	0.6405	0.0399	-0.1120	0.7278	-0.2190	0.8841	-0.1266	0.8765	0.0925	0.9065
---	---	1625504_at	0.0492	0.7785	-0.0350	0.7700	0.3165	0.1036	0.1875	0.5913	0.1392	0.4495	-0.0484	0.8078	-0.0898	0.8465	-0.0193	0.9505	0.0705	0.7514

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13896	CG13896	1625505_at	0.0279	0.9432	0.5257	0.0365	0.2598	0.1464	-0.2181	0.6084	-0.1162	0.6244	0.1019	0.6421	0.0412	0.9816	0.3094	0.4424	0.2682	0.5270
---	---	1625506_at	-0.0057	0.9801	-0.0127	0.9047	-0.0099	0.9715	-0.0847	0.8791	-0.0003	0.9991	0.0844	0.6690	-0.0600	0.9342	-0.1093	0.6723	-0.0494	0.8759
---	---	1625507_at	-0.0635	0.7578	0.0255	0.8310	0.0971	0.5335	0.0584	0.9149	0.0431	0.8459	-0.0154	0.9425	-0.0508	0.9564	-0.0016	0.9989	0.0492	0.8933
CG34353	CG12274	1625508_at	0.1638	0.3289	0.0301	0.8386	-0.2261	0.1283	-0.0828	0.9034	0.0041	0.9899	0.0869	0.7108	0.1374	0.8215	-0.1334	0.6246	-0.2708	0.2964
---	---	1625509_at	0.3398	0.0927	0.0721	0.4653	0.1591	0.3140	0.0510	0.9263	0.0382	0.8607	-0.0128	0.9515	-0.0448	0.9588	-0.1452	0.5640	-0.1003	0.7112
Osi1	Osiris	1625510_at	-0.0080	0.9732	-0.0445	0.7848	-0.0729	0.7509	0.0694	0.8897	-0.0746	0.6994	-0.1440	0.3494	0.0172	0.9860	-0.0774	0.7734	-0.0947	0.7007
CG34383	CG14365	1625511_at	-1.1661	0.0113	-1.5570	0.0793	-1.6481	0.0013	0.2690	0.6457	0.2294	0.4353	-0.0396	0.9084	0.3628	0.8815	-0.0663	0.9652	-0.4291	0.6740
CG34002	CG34002	1625512_s_at	0.1486	0.4279	-0.5158	0.1813	-0.9578	0.0120	-0.1933	0.8452	0.5304	0.1443	0.7236	0.0337	-0.0773	0.9142	-0.0508	0.8893	0.0265	0.9397
CG31297	CG31297	1625513_at	-0.0123	0.9435	0.0638	0.6761	0.1157	0.5283	-0.0216	0.9704	0.0660	0.6987	0.0875	0.5460	-0.0514	0.9404	-0.0058	0.9903	0.0456	0.8721
olf186-F	lethal (2) k11505	1625514_s_at	-1.1072	0.0023	-1.4248	0.0112	-1.3762	0.0011	0.2157	0.8161	0.5290	0.1443	0.3133	0.3485	0.1352	0.8461	0.2551	0.3686	0.1199	0.7134
sd	scalloped	1625515_a_at	-0.4889	0.0403	-0.0888	0.8580	-0.4109	0.0142	0.0671	0.9353	0.2789	0.2558	0.2118	0.3452	0.3431	0.7392	0.7058	0.1126	0.3626	0.4148
RhoGAP88C	---	1625516_at	0.0500	0.8365	-0.0510	0.7800	-0.2015	0.2684	0.0561	0.9255	0.1671	0.3660	0.1110	0.5294	0.1853	0.7953	-0.0339	0.9449	-0.2193	0.4822
Max	Max	1625517_at	0.4403	0.0130	0.5511	0.1058	0.4382	0.0239	0.1506	0.6854	-0.0036	0.9892	-0.1542	0.3282	0.2042	0.7305	0.0989	0.7393	-0.1053	0.7134
---	---	1625518_at	0.0626	0.7640	0.1132	0.4575	0.1661	0.3418	0.1055	0.8192	-0.0584	0.7923	-0.1639	0.3202	0.0630	0.9405	-0.0197	0.9634	-0.0827	0.7917
CG6453	CG6453	1625519_at	0.5776	0.0407	0.6736	0.0411	1.0572	0.0007	0.0652	0.9036	0.3832	0.0349	0.3180	0.0467	-0.2129	0.8284	0.6116	0.1374	0.8245	0.0828
CG34372	CG13552	1625520_at	0.1749	0.4516	0.1455	0.4102	0.0547	0.7638	-0.1039	0.8856	-0.1331	0.6247	-0.0293	0.9226	-0.1396	0.8875	-0.2220	0.5683	-0.0825	0.8668
CG30355	CG30355	1625521_at	0.0326	0.8846	0.0270	0.8079	0.3481	0.1428	0.0074	0.9952	-0.1162	0.6797	-0.1236	0.6230	-0.0249	0.9814	-0.0737	0.7968	-0.0488	0.8729
CG32551	CG32551	1625522_at	0.0855	0.6171	0.0581	0.5952	0.0881	0.7255	0.0447	0.9688	0.0285	0.9491	-0.0162	0.9658	-0.0198	0.9831	0.0959	0.6891	0.1157	0.6164
fd64A	forkhead domain (	1625523_at	-0.4599	0.0782	-0.0510	0.6424	0.1399	0.4674	0.0627	0.9559	-0.5715	0.0658	-0.6342	0.0285	-0.0175	0.9848	0.0307	0.9238	0.0482	0.8570
CG3165	CG3165	1625524_at	-0.0024	0.9922	-0.0557	0.7533	0.1672	0.5844	0.5494	0.1570	0.5743	0.0228	0.0249	0.9313	0.2515	0.7733	0.4216	0.2491	0.1701	0.6853
---	---	1625525_at	-0.0388	0.8218	0.0670	0.5393	0.3949	0.0508	0.1799	0.5780	-0.1443	0.3961	-0.3243	0.0356	-0.1278	0.8609	-0.0869	0.8208	0.0409	0.9195
CG17375	CG17375	1625526_at	0.0639	0.8659	0.1116	0.4154	0.2142	0.2560	-0.1461	0.8663	-0.2812	0.3691	-0.1351	0.6717	-0.1097	0.9246	-0.2618	0.5131	-0.1520	0.7320
CG5265	CG5265	1625527_at	0.3440	0.1533	0.1476	0.3755	0.3173	0.1859	-0.1081	0.8194	-0.0653	0.7696	0.0428	0.8414	-0.2059	0.7644	-0.0627	0.8794	0.1432	0.6457
CG17593	CG17593	1625528_at	0.1384	0.3670	0.0469	0.8309	-0.1035	0.4924	-0.1363	0.6321	0.4520	0.0066	0.5884	0.0012	0.1167	0.8903	0.4092	0.1857	0.2924	0.3711
CG32044	CG32044	1625529_at	-0.1057	0.7350	0.0850	0.5990	0.3168	0.1826	-0.0009	0.9994	-0.0291	0.9338	-0.0282	0.9262	-0.2241	0.7951	0.1150	0.8041	0.3391	0.3621
CG10462	CG10462	1625530_at	-0.2738	0.1043	-0.8229	0.0162	-0.5344	0.0121	-0.1407	0.7043	0.1037	0.5731	0.2444	0.1078	-0.2646	0.7506	-0.3246	0.3527	-0.0600	0.9031
Obp18a	Odorant-binding p	1625531_at	2.1248	0.0014	0.9738	0.2353	2.0430	0.0001	0.5343	0.2602	0.3374	0.2299	-0.1969	0.4600	-0.5288	0.7758	-0.8940	0.2488	-0.3652	0.6795
cv	twisted gastrulatio	1625532_at	-1.5299	0.0012	-0.5855	0.0406	-0.8698	0.0085	-0.1447	0.8350	-1.2727	0.0012	-1.1280	0.0012	0.2240	0.7697	-0.2794	0.3820	-0.5033	0.1490
bsk	Jun kinase	1625533_at	-0.0639	0.7255	-0.0372	0.9323	-0.3530	0.0362	-0.3359	0.2291	-0.0996	0.5811	0.2363	0.1106	-0.0318	0.9852	-0.0640	0.9156	-0.0323	0.9526
---	---	1625534_at	0.2606	0.2982	0.0318	0.7516	0.2367	0.3278	0.0256	0.9860	0.3358	0.3292	0.3102	0.3156	-0.1467	0.8141	-0.0448	0.9096	0.1019	0.7305
H15	neuromancer1	1625535_at	-1.4236	0.0035	-1.9881	0.0126	-2.3335	0.0002	-0.3042	0.6886	0.2666	0.4683	0.5708	0.0719	-0.1415	0.8564	-0.3415	0.2691	-0.2000	0.5555
CG15730	CG15730	1625536_at	0.0008	0.9973	0.0911	0.5408	-0.1428	0.4307	-0.0230	0.9777	-0.0979	0.6764	-0.0749	0.7353	0.1239	0.8076	-0.0215	0.9504	-0.1453	0.5100
CG3655	CG3655	1625537_at	-3.7122	0.0006	-1.3728	0.2625	-3.0646	0.0011	-1.2939	0.2117	-1.9911	0.0071	-0.6972	0.2111	0.3128	0.9342	0.2158	0.9084	-0.0970	0.9556
Jon66Ci	Jonah 66C	1625538_at	0.2975	0.2427	0.3455	0.0911	0.0253	0.9087	-0.0891	0.8869	-0.1754	0.4233	-0.0863	0.6988	0.0692	0.9499	-0.1539	0.6697	-0.2231	0.5123
---	---	1625539_at	-0.0420	0.8667	0.0765	0.6029	-0.0455	0.8140	-0.0684	0.9244	0.0201	0.9505	0.0885	0.6962	0.1116	0.8284	-0.0053	0.9916	-0.1169	0.6153
---	---	1625540_at	0.0827	0.6033	-0.1225	0.4197	0.0211	0.9206	-0.0890	0.8427	0.0514	0.8010	0.1404	0.3589	-0.1241	0.8097	-0.1628	0.4479	-0.0387	0.8949
Gr64b	Gustatory recepto	1625541_at	-0.0427	0.7805	0.2067	0.3695	-0.0392	0.8725	-0.2525	0.5125	-0.2436	0.2308	0.0090	0.9716	0.0245	0.9831	0.1055	0.7362	0.0810	0.8044
Tm1	Tropomyosin	1625542_a_at	-0.9567	0.0409	0.9429	0.0953	0.0376	0.9036	-0.3674	0.5038	-1.3153	0.0016	-0.9479	0.0039	0.5042	0.7220	0.6707	0.2468	0.1665	0.8247
CYLD	CYLD	1625543_s_at	-0.8524	0.0296	0.4399	0.2366	0.1619	0.2956	-0.3479	0.5515	-1.2265	0.0025	-0.8785	0.0065	-0.0859	0.9495	0.2345	0.5848	0.3204	0.4411
par-1	Par-1 kinase	1625544_s_at	-0.9624	0.0164	0.3294	0.4402	0.1702	0.4178	-0.0472	0.9637	-0.9809	0.0042	-0.9337	0.0031	-0.0645	0.9748	0.1798	0.7712	0.2444	0.6571
CG15816	CG15816	1625545_at	0.1700	0.2648	-0.0752	0.7261	0.0747	0.6289	0.0702	0.8642	0.1130	0.4569	0.0427	0.7942	-0.1571	0.8298	-0.1903	0.5523	-0.0332	0.9382
CG4557 /// DmirCG4557	CG4557	1625546_at	-0.8365	0.2734	0.5564	0.5811	0.6235	0.0176	0.0385	0.9562	-0.8137	0.0019	-0.8522	0.0010	0.1946	0.9710	0.8674	0.5373	0.6728	0.6423
---	---	1625547_at	-0.1362	0.4858	0.0371	0.8432	-0.2248	0.5655	-0.0949	0.9138	-0.1384	0.6577	-0.0435	0.8959	0.0372	0.9717	0.0858	0.7984	0.0486	0.8934
CG6661	CG6661	1625548_at	0.2927	0.1574	0.1461	0.3632	0.2044	0.2737	-0.1165	0.8470	-0.0422	0.8877	0.0743	0.7589	0.0288	0.9848	-0.1108	0.7964	-0.1396	0.7155
CG5970	CG5970	1625549_at	-0.0012	0.9966	0.5477	0.0924	0.8070	0.0007	-0.0133	0.9883	-0.4660	0.0380	-0.4527	0.0273	-0.1811	0.8395	0.2503	0.5105	0.4313	0.2574
CR33294	CR33294	1625550_at	1.3261	0.0009	1.3394	0.0502	1.7981	0.0001	0.3637	0.3863	0.3297	0.1629	-0.0339	0.9058	0.0562	0.9816	0.0374	0.9650	-0.0188	0.9835
Jhl-1	Juvenile hormone	1625551_at	0.1539	0.5363	0.7304	0.0440	0.6606	0.0387	-0.1521	0.7795	-0.2760	0.2202	-0.1238	0.5825	-0.0588	0.9717	0.3481	0.4073	0.4069	0.3508
CG14820	CG14820	1625552_at	-0.3559	0.8987	-0.2784	0.2070	-0.0545	0.7743	0.2246	0.9704	-1.3831	0.3797	-1.6077	0.2450	-0.1547	0.9894	-1.3978	0.5900	-1.2431	0.6377
CG33482	DS07721.6-like pr	1625553_at	0.1718	0.4494	-0.0875	0.5348	0.2150	0.2274	0.0923	0.8469	0.1869	0.2893	0.0947	0.5885	-0.1901	0.8092	-0.0763	0.8685	0.1139	0.7647
obst-F	CG7306	1625554_at	0.0510	0.8480	0.3412	0.1069	0.1002	0.5028	0.1064	0.7857	-0.0269	0.9044	-0.1333	0.3727	0.1434	0.8465	0.0735	0.8642	-0.0700	0.8633
---	---	1625555_at	0.3366	0.0967	-0.0472	0.8292	0.2430	0.2114	0.1517	0.6936	0.2403	0.1750	0.0886	0.6250	-0.3732	0.6123	-0.2343	0.4593	0.1389	0.6892
ari-2	ariadne-2	1625556_at	0.4437	0.0766	-0.1880	0.3097	-0.8157	0.0079	-0.3482	0.5128	0.6090	0.0370	0.9572	0.0032	0.2968	0.7062	0.0278	0.9590	-0.2690	0.4204
---	---	1625557_at	0.2223	0.3892	0.2752	0.1388	0.0898	0.6263	-0.0777	0.8660	-0.2455	0.1294	-0.1678	0.2502	0.1596	0.8141	-0.1320	0.6704	-0.2917	0.3089
CG17290	CG17290	1625558_a_at	-0.0176	0.9413	-0.0316	0.8470	0.0739	0.6189	0.0307	0.9610	0.0160	0.9496	-0.0147	0.9445	-0.1185	0.8846	-0.0557	0.9064	0.0628	0.8813
CG30217	CG30217	1625559_a_at	-1.9400	0.0057	-1.6141	0.0019	-1.6935	0.0001	0.0248	0.9774	-0.7253	0.0054	-0.7501	0.0027	-0.0347	0.9898	-0.5218	0.3534	-0.4872	0.4055
---	---	1625560_at	0.2591	0.2067	0.0810	0.5147	0.1600	0.4643	-0.0874	0.8732	-0.0055	0.9845	0.0819	0.6808	0.0769	0.9393	-0.0623	0.8918	-0.1392	0.6860
par-1	Par-1 kinase	1625561_s_at	-0.0314	0.9234	-0.0086	0.9377	-0.2148	0.2165	-0.1331	0.6629	0.0765	0.6409	0.2096	0.1079	0.0508	0.9589	0.0458	0.9157	-0.0050	0.9921
CG4959 /// DyakCG4959	CG4959	1625562_at	-0.0396	0.8546	0.0234	0.8184	0.1598	0.4035	0.0235	0.9777	-0.0955	0.6892	-0.1191	0.5634	0.0213	0.9808	0.0438	0.8727	0.0225	0.9338
CG14253	CG14253	1625563_s_at	-0.6020	0.0557	-0.1782	0.4636	-0.8415	0.0176	0.4770	0.1658	0.8096	0.0030	0.3326	0.0767	1.1278	0.2884	1.3320	0.0378	0.2041	0.7422
Coq2	CG9613	1625564_at	-0.3664	0.2427	-0.5025	0.0775	-0.2398	0.1329	0.0310	0.9620	0.0510	0.8113	0.0200	0.9252	-0.2028	0.8202	-0.1104	0.8172	0.0924	0.8444
Nmnat	Nicotinamide mon	1625565_at	0.0326	0.8892	-0.0695	0.6064	-0.0412	0.8102	0.0684	0.8898	0.1983	0.2294	0.1299	0.3980	-0.0372	0.9677	-0.0367	0.9229	0.0005	0.9992
CG9293	CG9293	1625566_a_at	0.0524	0.7967	0.1237	0.4486	0.2278	0.2951	0.0085	0.9922	-0.0452	0.8436	-0.0536	0.7863	-0.0973	0.9246	0.1206	0.7760	0.2179	0.5508
CG18223	CG18223	1625567_a_at	0.0294	0.9060	0.2071	0.2413	0.0808	0.7398	0.0391	0.9586	-0.0434	0.8684	-0.0826	0.6914	0.1740	0.8395	0.0526	0.9248	-0.1215	0.7692
CG15111	CG15111	1625568_a_at	-0.1002	0.5710	0.1611	0.4629	0.2000	0.1657	-0.0090	0.9893	0.0619	0.7316	0.0709	0.6565	0.0598	0.9589	0.3950	0.2168	0.3352	0.3207
CG7565	CG7565	1625569_at	-0.2601	0.3880	0.0150	0.9855	1.2314	0.0008	0.2001	0.8413	-0.0795	0.8720	-0.2796	0.4225	-1.0781	0.3712	0.1821	0.8346	1.2602	0.0879
Cpr64Ac	CG15008	1625570_at	0.3789	0.0268	0.1114	0.4847	0.1643	0.3391	-0.0508	0.9435	0.0926	0.6906	0.1435	0.4543	0.0112	0.9939	-0.0925	0.8252	-0.1037	0.7839
CG5385	CG5385	1625571_at	0.6959	0.0064	1.5589	0.0082	1.4461	0.0002	0.1161	0.8544	-0.0371	0.9068	-0.1531	0.4819	-0.0341	0.9826	0.5462	0.1297	0.5803	0.1358
Dbi	diazepam binding	1625572_s_at	1.1773	0.0023	0.8820	0.2543	0.4914	0.0548	0.1246	0.8162	-0.2368	0.2622	-0.3614	0.0586	0.4017	0.8461	-0.6669	0.4355	-1.0686	0.2282
Tim9a	Tim9a	1625573_at	-0.1797	0.4203	-0.0572	0.9028	-0.3524	0.0606	0.0007	0.9994	0.3246	0.0814	0.3240	0.0550	0.3151	0.7970	0.3614	0.4932	0.0463	0.9490

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Nos	nitric oixide synth	1625574_at	-0.0245	0.9178	0.0447	0.8886	-0.1482	0.6344	-0.0189	0.9803	0.1436	0.4569	0.1625	0.3385	0.0158	0.9928	0.2162	0.6157	0.2004	0.6434
CG34402	CG17793	1625575_at	0.0278	0.8769	-0.0314	0.7524	0.1575	0.3923	0.1639	0.7093	0.0044	0.9888	-0.1596	0.3916	-0.0925	0.8461	-0.1317	0.5179	-0.0392	0.8846
---	---	1625576_at	-0.2511	0.2453	0.1305	0.2949	0.0577	0.7744	-0.0963	0.8698	-0.1178	0.6015	-0.0215	0.9328	0.0784	0.9340	0.0680	0.8749	-0.0104	0.9838
CCKLR-17D3	CCK-like receptor	1625577_at	0.0430	0.9077	0.1344	0.5854	0.0116	0.9713	0.0430	0.9612	-0.0196	0.9555	-0.0626	0.8146	0.2327	0.7628	0.2325	0.4794	-0.0002	0.9997
CG33097	CG33097	1625578_at	0.3618	0.6034	0.5619	0.1878	0.2239	0.3538	0.0215	0.9873	0.0682	0.8732	0.0467	0.9042	0.4020	0.8682	0.6795	0.4881	0.2774	0.8168
---	---	1625579_at	0.0301	0.8823	0.0273	0.8808	-0.0706	0.7496	0.1373	0.7838	0.0740	0.7650	-0.0633	0.7831	0.0937	0.9098	-0.0675	0.8725	-0.1613	0.6128
---	---	1625580_at	0.1804	0.2802	-0.0361	0.7228	-0.1766	0.4240	0.0748	0.9086	0.2241	0.2845	0.1494	0.4461	0.1824	0.7387	-0.0441	0.8999	-0.2265	0.3424
CG10011	CG10011	1625581_at	0.2799	0.7694	-0.0046	0.9969	-0.9770	0.0207	0.2711	0.7929	0.9929	0.0262	0.7217	0.0591	1.1133	0.6881	0.8880	0.4307	-0.2253	0.8846
CG32187	CG32187	1625582_at	-0.0914	0.7662	0.1436	0.4417	0.2245	0.2181	-0.0251	0.9761	-0.2142	0.3068	-0.1891	0.3170	-0.0068	0.9964	0.0052	0.9941	0.0121	0.9837
CG16926	CG16926	1625583_at	2.3975	0.0010	2.2097	0.0041	2.4627	0.0037	0.3977	0.7949	-0.3573	0.6006	-0.7550	0.1748	0.0580	0.9851	-0.7106	0.3018	-0.7686	0.2924
CG8654	CG8654	1625584_at	1.9820	0.0058	-0.0685	0.9591	1.1166	0.0339	0.0133	0.9943	0.6465	0.1034	0.6332	0.0765	-1.1137	0.7070	-1.3719	0.2518	-0.2582	0.8785
CG6481	CG6481	1625585_at	0.0021	0.9924	-0.0741	0.5162	0.0668	0.7605	0.1268	0.8366	0.1751	0.4695	0.0484	0.8579	-0.1338	0.8202	0.0344	0.9289	0.1682	0.5081
---	---	1625586_at	0.2038	0.2613	0.0748	0.5980	0.2819	0.2230	0.0665	0.9060	-0.1330	0.4848	-0.1996	0.2198	-0.0359	0.9735	-0.0837	0.8070	-0.0478	0.8957
Jheh3	Juvenile hormone	1625587_at	4.5975	0.0009	3.4265	0.0014	6.2309	0.0000	2.5702	0.0189	1.3125	0.0300	-1.2577	0.0228	-0.3904	0.5134	0.1129	0.7371	0.5033	0.1156
CG6199	CG6199	1625588_s_at	-0.1743	0.6142	-0.9311	0.1119	0.0026	0.9956	-0.4117	0.3571	-0.5922	0.0280	-0.1804	0.4548	-1.2880	0.3712	-1.3875	0.0890	-0.0995	0.9275
CG4840	CG4840	1625589_at	-0.0505	0.8547	-0.1071	0.7329	-0.0767	0.7072	0.1454	0.8612	0.5490	0.0686	0.4036	0.1284	0.0651	0.9611	0.4025	0.2634	0.3374	0.3791
Ate1	Ate1	1625590_a_at	0.6665	0.0260	1.1294	0.0136	1.5436	0.0000	-0.1211	0.8182	-0.6396	0.0081	-0.5185	0.0126	-0.6383	0.2884	-0.3686	0.2217	0.2696	0.3967
CG4582	CG4582	1625591_at	-0.0060	0.9859	0.3296	0.0729	-0.1172	0.4738	-0.1978	0.5357	-0.1893	0.2578	0.0085	0.9679	0.0252	0.9835	0.0256	0.9557	0.0003	0.9994
CG11307	CG11307	1625592_s_at	0.3766	0.2030	0.5781	0.1098	1.0184	0.0058	0.1193	0.8973	-0.0919	0.8058	-0.2112	0.4637	-0.3132	0.7475	0.1401	0.7842	0.4533	0.2858
---	---	1625593_at	0.0380	0.9030	0.0476	0.6431	0.0284	0.8826	0.1438	0.7401	0.0028	0.9916	-0.1410	0.4319	0.0006	0.9999	0.0083	0.9924	0.0078	0.9891
Pka-R2	cAMP-dependent	1625594_s_at	-0.9695	0.0881	0.2737	0.5054	-0.9070	0.0115	-0.7796	0.1956	-0.5686	0.1172	0.2110	0.5576	0.3469	0.8534	0.4875	0.5457	0.1406	0.8947
DsimCG9245 /// Pis	CG9245	1625595_a_at	-0.0241	0.9388	1.2571	0.0334	1.2783	0.0003	-0.2161	0.7887	-1.3623	0.0023	-1.1462	0.0029	-0.1722	0.7743	-0.1351	0.6310	0.0371	0.9176
Osbp	oxysterol binding	1625596_at	-0.2602	0.5484	0.1422	0.8457	0.3115	0.1362	0.0773	0.9314	-0.1130	0.7157	-0.1903	0.4510	-0.1517	0.9611	0.3689	0.7103	0.5206	0.5767
CG6441	CG6441	1625597_a_at	0.0303	0.8840	0.0078	0.9803	-0.0291	0.8614	-0.0789	0.8524	-0.1499	0.3373	-0.0711	0.6550	0.0602	0.9677	-0.0079	0.9933	-0.0680	0.9007
CG31326	CG31326	1625598_at	2.5263	0.0017	1.2092	0.1868	2.3017	0.0000	0.6912	0.4137	0.7675	0.1013	0.0763	0.8919	-0.4576	0.8292	-0.6408	0.4794	-0.1832	0.8794
rumi	rumi	1625599_at	0.1825	0.4440	0.2352	0.4867	0.6836	0.0071	0.0993	0.8817	0.0409	0.8953	-0.0584	0.8215	-0.3611	0.7392	0.1131	0.8595	0.4742	0.3141
Asph	Aspartyl beta-hyd	1625600_a_at	-0.3036	0.4290	-0.0100	0.9674	-0.9391	0.0015	-0.8969	0.0573	-0.0170	0.9679	0.8800	0.0039	0.1093	0.9486	0.0930	0.9049	-0.0163	0.9847
resilin	resilin	1625601_at	0.0678	0.7102	0.3103	0.0781	0.2439	0.5027	0.0591	0.9441	-0.1000	0.7224	-0.1591	0.4917	0.1517	0.7644	0.2291	0.2691	0.0774	0.7543
---	---	1625602_at	0.0633	0.7632	0.1143	0.4116	0.1227	0.5806	0.0594	0.9270	0.0382	0.8835	-0.0212	0.9288	0.1044	0.8953	0.0864	0.8244	-0.0180	0.9679
CG13604	CG13604	1625603_at	0.2672	0.3747	-0.2880	0.3755	-0.3742	0.2014	-0.0851	0.9495	0.6087	0.1032	0.6938	0.0440	0.0010	0.9998	0.0342	0.9601	0.0332	0.9548
CG14259	CG14259	1625604_at	0.8467	0.3232	0.1948	0.3040	0.3693	0.1452	-0.1300	0.7278	-0.3613	0.0375	-0.2313	0.1208	-0.1833	0.9666	-0.9999	0.3802	-0.8166	0.4978
CG11251	CG11251	1625605_at	0.3514	0.0932	0.0204	0.8936	0.3415	0.0528	0.0252	0.9733	0.0265	0.9188	0.0014	0.9955	-0.0275	0.9775	-0.0873	0.7502	-0.0598	0.8394
Ocho	Ocho	1625606_at	0.1182	0.5880	0.2485	0.3982	0.5056	0.0060	0.0850	0.9080	-0.1378	0.5911	-0.2228	0.2967	-0.0752	0.9457	0.0992	0.8185	0.1744	0.6263
Syx5	syntaxin	1625607_s_at	0.1992	0.2432	0.4352	0.0521	0.4981	0.0218	0.1026	0.8140	0.2986	0.0831	0.1960	0.2008	-0.0016	0.9994	0.5542	0.0415	0.5558	0.0559
CG8840	CG8840	1625608_at	0.2458	0.2024	0.2461	0.3122	0.3131	0.1296	0.0711	0.9351	0.0479	0.8906	-0.0231	0.9425	-0.0041	0.9967	0.0251	0.9508	0.0292	0.9347
CG32450	CG32450	1625609_at	-0.0107	0.9619	-0.1483	0.4836	0.0541	0.8142	0.0203	0.9860	0.0920	0.7847	0.0718	0.8194	-0.1214	0.7953	-0.0066	0.9869	0.1148	0.5830
CG40376	CG40376	1625610_at	-0.1034	0.5312	0.0074	0.9478	-0.1174	0.5687	-0.0648	0.9094	-0.0445	0.8495	0.0203	0.9268	-0.0140	0.9914	0.0642	0.8673	0.0782	0.8178
CG33115	CG33115	1625611_at	-2.2772	0.0243	-3.9287	0.0062	-3.8687	0.0000	-0.2823	0.5511	-0.0586	0.8534	0.2237	0.3157	-0.1245	0.9816	-1.6438	0.1528	-1.5193	0.2165
---	---	1625612_at	0.1593	0.2845	0.1011	0.5852	-0.0321	0.8806	0.0310	0.9567	-0.0270	0.8981	-0.0579	0.7240	0.2256	0.8091	-0.1099	0.8309	-0.3355	0.3953
CG11899 /// DsimCG11899	CG11899	1625613_at	0.6439	0.0587	0.2061	0.8142	1.0479	0.0008	-0.0052	0.9943	-0.4594	0.0105	-0.4542	0.0067	-0.9271	0.6749	-0.8909	0.3176	0.0363	0.9812
---	---	1625614_at	0.1770	0.2143	0.2647	0.1578	0.4547	0.0154	-0.0161	0.9838	-0.2725	0.1289	-0.2565	0.1102	-0.0983	0.8647	-0.0004	0.9998	0.0979	0.7040
CG31179	CG31179	1625615_at	-0.0332	0.8284	-0.0200	0.9290	0.1713	0.4716	0.2391	0.5238	0.1614	0.4276	-0.0778	0.7086	0.0304	0.9831	0.0312	0.9528	0.0008	0.9992
CG14566	CG14566	1625616_at	0.0850	0.6687	-0.3626	0.0711	-0.1441	0.6786	0.1844	0.8402	0.3214	0.3555	0.1370	0.7036	-0.0972	0.9238	0.0281	0.9600	0.1254	0.7492
Roc2	Roc2	1625617_at	0.1380	0.5743	-0.2491	0.2311	-0.1088	0.6408	0.0626	0.9036	0.1032	0.5696	0.0406	0.8326	-0.1458	0.9031	-0.4418	0.2985	-0.2960	0.5156
CG1824	CG1824	1625618_at	-0.4026	0.2725	-0.1144	0.8980	-0.0741	0.7687	-0.0869	0.9068	-0.2569	0.2867	-0.1700	0.4518	0.0165	0.9963	0.0201	0.9911	0.0036	0.9979
pip	pipe	1625619_at	0.1687	0.5576	0.0290	0.7847	0.1939	0.2672	0.0997	0.8280	-0.0413	0.8576	-0.1410	0.3889	-0.1333	0.8875	-0.0697	0.8947	0.0636	0.8951
---	---	1625620_a_at	-0.4660	0.2437	-0.4728	0.3364	-0.5557	0.0294	-0.1312	0.9027	0.0884	0.8442	0.2196	0.5134	-0.2429	0.8661	-0.2240	0.7400	0.0189	0.9846
f	forked	1625621_s_at	-4.3142	0.0004	-5.3417	0.0014	-5.0988	0.0001	0.1559	0.9506	1.0982	0.1132	0.9423	0.1278	-0.1566	0.8846	-0.0185	0.9806	0.1381	0.7749
---	---	1625622_at	-0.1139	0.5327	0.0615	0.5821	-0.0690	0.6772	0.0238	0.9715	0.0360	0.8722	0.0122	0.9549	-0.0869	0.8628	0.0377	0.9027	0.1246	0.5615
Nrx-1	Neurexin 1	1625623_at	-1.0579	0.0124	-0.8016	0.0185	-1.3934	0.0011	-0.0464	0.9441	0.1342	0.5015	0.1806	0.2930	0.2157	0.7602	0.0829	0.8343	-0.1328	0.6855
CG14105 /// DyakCG14105	CG14105	1625624_at	0.2367	0.3965	-0.0681	0.7722	0.0705	0.6692	0.1762	0.7128	0.1243	0.6025	-0.0519	0.8353	-0.0878	0.9324	-0.0928	0.8353	-0.0049	0.9933
ImpE1	Y1-like	1625625_at	0.2264	0.1332	-0.2974	0.0814	-0.0912	0.5859	0.0754	0.9068	0.3097	0.1338	0.2343	0.2059	-0.1325	0.7395	-0.3053	0.0831	-0.1728	0.3175
CG4438	CG4438	1625626_at	0.1053	0.4817	-0.0194	0.8523	0.0909	0.5219	0.2004	0.5756	0.1740	0.3517	-0.0264	0.9051	-0.1018	0.8202	-0.1253	0.5131	-0.0235	0.9271
---	---	1625627_at	-0.0388	0.8411	-0.1477	0.4471	0.1948	0.4593	0.2081	0.5680	0.1277	0.5195	-0.0805	0.6802	-0.0983	0.9144	0.0691	0.8807	0.1674	0.6294
---	---	1625628_at	0.1693	0.4496	0.0093	0.9578	0.1207	0.4403	-0.0877	0.8485	-0.1030	0.5702	-0.0153	0.9418	-0.1599	0.7979	-0.0783	0.8175	0.0816	0.7955
alpha-Est6	fragment G	1625629_at	-0.0155	0.9440	0.4276	0.2451	0.0458	0.8233	-0.3033	0.4586	-0.3600	0.1038	-0.0567	0.8218	0.1422	0.8965	0.1827	0.6883	0.0406	0.9433
---	---	1625630_at	0.0625	0.8141	-0.1369	0.6223	0.0265	0.9015	0.0491	0.9494	0.1119	0.6438	0.0629	0.7949	-0.0493	0.9741	-0.0591	0.9185	-0.0099	0.9868
CG15180	CG15180	1625631_at	0.2338	0.2373	0.0111	0.9234	0.0238	0.9008	0.1683	0.6122	0.3304	0.0525	0.1621	0.2825	0.1116	0.8513	-0.0256	0.9494	-0.1372	0.6052
robl	roadblock	1625632_at	-0.4600	0.0350	0.1513	0.4994	-0.1903	0.2827	0.0527	0.9314	0.0420	0.8610	-0.0106	0.9632	0.4391	0.5754	0.6352	0.0807	0.1961	0.6017
Argk	insect intestinal m	1625633_a_at	-0.6069	0.0094	0.6736	0.0109	0.1659	0.6756	-0.1571	0.8432	-0.8902	0.0090	-0.7331	0.0131	0.2865	0.8049	0.3427	0.4863	0.0562	0.9341
Dbp45A	ATP-dependent R	1625634_at	-0.1778	0.3148	-0.2839	0.4542	-0.3161	0.0464	-0.0068	0.9931	-0.0367	0.8530	-0.0298	0.8674	-0.0401	0.9816	-0.1166	0.8257	-0.0764	0.8905
CG9316	CG9316	1625635_at	0.0171	0.9344	-0.0463	0.6715	0.0500	0.8054	0.0793	0.8908	0.0361	0.8906	-0.0432	0.8477	0.0976	0.8692	-0.0234	0.9505	-0.1210	0.6317
---	---	1625636_s_at	-0.1066	0.4947	-0.3373	0.5318	-0.3969	0.1019	0.0735	0.9300	0.5117	0.0434	0.4382	0.0507	0.1358	0.9095	0.2991	0.4915	0.1633	0.7389
Start1	Start1	1625637_at	0.4142	0.0327	0.5424	0.1308	0.5727	0.0028	-0.1247	0.7185	0.2542	0.1042	0.3789	0.0149	-0.1019	0.9221	0.3226	0.3397	0.4245	0.2352
Hex-A	hexokinase A	1625638_a_at	1.0518	0.0119	0.9427	0.0081	0.0736	0.8123	-0.5452	0.2172	0.1941	0.4852	0.7392	0.0079	0.4812	0.6898	0.2055	0.7161	-0.2757	0.6043
CG6737	CG6737	1625639_at	-0.0021	0.9934	0.1164	0.3028	0.0650	0.7201	-0.0052	0.9956	0.0145	0.9611	0.0197	0.9362	0.0561	0.9238	0.0955	0.6516	0.0394	0.8822
Tsp97E	Tetraspanin 97E	1625640_a_at	1.0051	0.0071	0.2006	0.5113	0.3629	0.0592	0.0129	0.9909	0.8385	0.0054	0.8256	0.0034	-0.2567	0.7478	-0.0507	0.9238	0.2061	0.5657
CG9650 /// DsimCG9650	CG9650	1625641_s_at	0.3808	0.0892	0.2529	0.1937	0.7122	0.0126	0.1853	0.7121	0.1179	0.6463	-0.0673	0.7929	-0.1876	0.7440	0.0865	0.7692	0.2741	0.2751
CG11509	CG11509	1625642_at	-0.1717	0.2882	-1.4559	0.0069	-1.2546	0.0162	0.5858	0.5199	1.6983	0.0044	1.1125	0.0175	-0.1115	0.8382	-0.0995	0.6949	0.0120	0.9726

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1625643_s_at	-0.0485	0.8784	-0.1436	0.6734	-0.0674	0.8437	0.0009	0.9994	0.4273	0.2688	0.4264	0.2153	-0.2936	0.7726	0.1844	0.7116	0.4779	0.2918
CG2641	CG2641	1625644_at	-0.4450	0.0260	-0.1752	0.2126	-0.2436	0.1962	-0.1590	0.7857	-0.0897	0.7562	0.0694	0.7986	0.0564	0.9238	0.2652	0.1599	0.2087	0.2964
CG31822	CG31822	1625645_at	0.0923	0.5636	0.0155	0.9055	0.0709	0.7372	-0.0892	0.8794	-0.0905	0.6949	-0.0013	0.9958	-0.1331	0.8625	-0.2353	0.4479	-0.1023	0.7798
CG11148	CG11148	1625646_at	0.4552	0.0305	0.0028	0.9956	-0.5927	0.0289	-0.2434	0.5504	0.8942	0.0020	1.1376	0.0005	0.3768	0.7062	0.4216	0.2984	0.0448	0.9387
CG14519	CG14519	1625647_at	0.2523	0.2487	0.0111	0.9169	0.1768	0.2297	0.0636	0.9228	0.1871	0.3564	0.1235	0.5237	0.0073	0.9939	0.0730	0.7787	0.0658	0.7964
CG3868	CG3868	1625648_at	-0.0589	0.9700	-0.0715	0.6036	-0.3232	0.3461	0.2267	0.9410	-0.4637	0.6365	-0.6904	0.3973	0.1878	0.9675	-0.6824	0.5967	-0.8701	0.4865
---	---	1625649_s_at	-0.0105	0.9672	-0.1214	0.4574	0.0465	0.7713	0.0826	0.9036	0.1380	0.5630	0.0553	0.8261	-0.1626	0.7070	0.0266	0.9264	0.1892	0.3079
CG1161	CG1161	1625650_at	-0.1570	0.3964	-0.0760	0.5458	-0.0241	0.8949	0.0754	0.8698	-0.2864	0.0776	-0.3618	0.0203	-0.0691	0.9296	-0.2549	0.3010	-0.1858	0.4793
CG18317	CG18317	1625651_at	0.1805	0.6422	0.7980	0.3110	-0.2095	0.5541	-0.3605	0.4455	-0.0364	0.9200	0.3240	0.1586	0.6714	0.7506	0.5487	0.5634	-0.1228	0.9215
CG12428	CG12428	1625652_s_at	0.7289	0.0260	0.8530	0.2035	1.4186	0.0002	0.1915	0.6086	0.1950	0.3043	0.0035	0.9880	-0.4081	0.8049	0.1999	0.8247	0.6080	0.3865
---	---	1625653_at	-0.0119	0.9738	0.0067	0.9686	-0.0395	0.8521	-0.0651	0.9042	-0.0226	0.9269	0.0424	0.8332	0.0973	0.9221	0.0562	0.9120	-0.0411	0.9289
GluRIIB	glutamate recepto	1625654_at	-1.7428	0.0005	-0.5100	0.1505	-1.6221	0.0004	-0.8213	0.0657	-1.0874	0.0020	-0.2661	0.2472	0.0922	0.9540	-0.2710	0.5863	-0.3632	0.4557
CG3214	CG3214	1625655_at	0.0519	0.8239	0.4427	0.2580	0.2783	0.2036	0.1452	0.7576	-0.6814	0.0056	-0.8266	0.0014	0.2260	0.8424	-0.1849	0.7339	-0.4109	0.3878
CG5625	CG5625	1625656_s_at	-0.0316	0.9071	0.3769	0.2155	0.4522	0.0236	0.0960	0.8074	-0.0100	0.9676	-0.1060	0.4777	0.1323	0.8967	0.4372	0.2346	0.3049	0.4336
CG32335	CG32335	1625657_at	0.2211	0.5413	0.0004	0.9997	-0.3699	0.2303	-0.0865	0.9260	-0.0213	0.9595	0.0652	0.8404	0.2487	0.8564	-0.0687	0.9402	-0.3174	0.5995
CG5181	CG5181	1625658_at	-0.4766	0.0806	-0.2578	0.1448	-0.3409	0.1092	-0.1675	0.7469	-0.1029	0.6929	0.0646	0.7996	-0.0122	0.9922	0.2356	0.3991	0.2478	0.3898
Obp50d	Odorant-binding p	1625659_at	0.1817	0.1966	0.2137	0.2918	0.1022	0.6887	-0.0179	0.9796	-0.0376	0.8681	-0.0197	0.9252	-0.0430	0.9752	0.0054	0.9941	0.0485	0.9181
CG4151	CG4151	1625660_at	0.1572	0.5400	0.0277	0.8079	0.3818	0.0674	0.1823	0.8053	-0.0273	0.9509	-0.2096	0.4511	0.0716	0.9550	-0.0777	0.8850	-0.1494	0.7239
CG17702	CG17702	1625661_at	0.1530	0.4295	0.1419	0.5887	0.0192	0.9273	0.0868	0.9026	0.0573	0.8472	-0.0295	0.9161	0.1637	0.7506	-0.0049	0.9925	-0.1686	0.4597
CG13287	CG13287	1625662_at	-0.0226	0.9117	-0.0082	0.9403	0.0282	0.8729	0.0947	0.8516	-0.1685	0.3701	-0.2631	0.1091	0.0239	0.9869	-0.0725	0.8793	-0.0964	0.8173
Kul	Kuzbanian-like	1625663_at	0.2396	0.2185	0.1902	0.2499	-0.0110	0.9543	-0.0223	0.9761	0.0787	0.7121	0.1010	0.5832	-0.0839	0.8882	-0.0427	0.8995	0.0412	0.8933
CG14059	CG14059	1625664_at	-1.1630	0.0399	-1.3736	0.0041	-0.7109	0.0947	0.5045	0.3744	0.1227	0.7478	-0.3819	0.1828	-0.0489	0.9756	-0.0426	0.9462	0.0063	0.9924
mRpL39	mitochondrial ribo	1625665_at	-0.2680	0.4170	0.0390	0.8782	0.5924	0.0186	0.2540	0.3016	-0.6358	0.0018	-0.8898	0.0003	-0.2115	0.8599	-0.2165	0.6881	-0.0050	0.9953
---	---	1625666_at	0.8373	0.0819	0.0419	0.8313	0.1030	0.7042	0.0299	0.9731	0.4190	0.0602	0.3891	0.0518	0.1999	0.8725	-0.2069	0.7105	-0.4068	0.4153
CG13561	CG13561	1625667_at	0.1741	0.4452	0.0470	0.6870	0.0664	0.7515	0.3424	0.3909	0.2445	0.2797	-0.0979	0.6776	0.0121	0.9933	-0.0044	0.9951	-0.0165	0.9752
CG33277	CG33277	1625668_at	0.1547	0.4075	0.1695	0.1695	0.2596	0.1551	-0.1334	0.7409	-0.1183	0.5331	0.0150	0.9468	-0.0522	0.9499	0.1498	0.5635	0.2020	0.4243
Tektin-A	TEKTIN A	1625669_at	0.1853	0.5019	0.1704	0.3716	0.1563	0.5793	0.2566	0.6321	0.2075	0.4462	-0.0491	0.8741	0.1744	0.8692	-0.0544	0.9353	-0.2288	0.6140
CG1387	CG1387	1625670_at	0.1315	0.4166	-0.0950	0.3665	0.1927	0.3208	0.1876	0.5998	0.1342	0.4760	-0.0535	0.7908	-0.0770	0.8655	-0.0787	0.7017	-0.0016	0.9960
---	---	1625671_at	2.4256	0.0039	1.8954	0.0303	3.7079	0.0001	0.4186	0.8444	-1.2927	0.1002	-1.7113	0.0232	0.4490	0.8446	-0.8158	0.3768	-1.2649	0.2029
---	---	1625672_s_at	-0.6224	0.2999	0.4519	0.6800	0.2580	0.3228	-0.2838	0.8124	-0.9062	0.0594	-0.6224	0.1394	-0.1362	0.9802	0.1883	0.9208	0.3244	0.8323
CG12983 /// DmirCG12983	CG12983	1625673_at	-0.0453	0.8155	-0.2990	0.2603	-0.2400	0.2629	0.2277	0.6716	0.3631	0.1517	0.1353	0.5954	0.0163	0.9913	0.0212	0.9655	0.0049	0.9924
CG12516	CG12516	1625674_at	0.2100	0.2143	0.1235	0.6596	-0.0450	0.8315	-0.1606	0.7138	-0.0801	0.7322	0.0805	0.7023	-0.0423	0.9717	-0.2191	0.4753	-0.1768	0.5819
CG4853	CG4853	1625675_a_at	-0.0636	0.7754	0.3819	0.3693	0.2440	0.1377	0.0717	0.8887	0.1216	0.5027	0.0499	0.7967	0.1974	0.8564	0.5051	0.2408	0.3077	0.5061
---	---	1625676_s_at	-0.0114	0.9548	0.1307	0.3365	0.2572	0.2833	0.0993	0.8132	0.0399	0.8528	-0.0594	0.7393	-0.1121	0.8988	-0.0246	0.9634	0.0875	0.8309
---	---	1625677_at	0.1770	0.2721	0.2093	0.2176	0.2119	0.1292	-0.0665	0.9060	-0.0305	0.9031	0.0360	0.8650	-0.0019	0.9990	0.0787	0.8018	0.0806	0.7815
Pi3K59F	dPI 3-kinase	1625678_at	-0.7006	0.1148	0.0155	0.9894	0.3575	0.0918	0.1510	0.7556	-0.7642	0.0039	-0.9152	0.0011	-0.2600	0.9296	-0.0809	0.9590	0.1790	0.8915
CG15386	CG15386	1625679_at	0.3062	0.1983	0.2322	0.3274	0.0534	0.8091	-0.0661	0.9375	-0.0010	0.9979	0.0651	0.8135	0.1311	0.8795	-0.1095	0.7925	-0.2406	0.4841
CG5792	CG5792	1625680_a_at	-0.3374	0.2315	0.2699	0.3769	-0.0427	0.8067	-0.3317	0.3767	-0.5110	0.0241	-0.1794	0.3579	-0.0425	0.9767	-0.0740	0.8823	-0.0316	0.9486
Gga	Gga	1625681_at	0.0562	0.8186	0.3917	0.0393	0.3462	0.0460	0.1589	0.7138	0.2472	0.2090	0.0883	0.6666	0.1057	0.9142	0.4013	0.2254	0.2956	0.3969
---	---	1625682_at	0.0202	0.9019	-0.0664	0.6973	-0.0029	0.9885	0.0805	0.8650	0.0514	0.8027	-0.0290	0.8839	0.2518	0.7092	0.1107	0.7371	-0.1411	0.6419
CG5112	CG5112	1625683_at	-0.9411	0.0077	0.2483	0.3238	0.7327	0.0223	0.0635	0.9253	-0.7219	0.0044	-0.7854	0.0018	-0.4139	0.6955	0.4654	0.2789	0.8793	0.0836
CG11523	CG11523	1625684_at	-0.3954	0.2172	-0.5690	0.2937	-0.4624	0.0124	0.0367	0.9728	0.8532	0.0073	0.8165	0.0053	0.0896	0.9698	0.8070	0.1720	0.7174	0.2557
---	---	1625685_at	1.3009	0.0462	1.1057	0.2914	1.0671	0.0024	0.0144	0.9883	-0.6074	0.0164	-0.6219	0.0091	0.1303	0.9772	-0.8170	0.4471	-0.9473	0.3878
---	---	1625686_s_at	0.7271	0.2259	1.6350	0.2227	2.3384	0.0001	0.6726	0.2501	0.0506	0.9173	-0.6219	0.0504	-0.0982	0.9884	1.0161	0.4972	1.1144	0.4597
perd	NG2-like	1625687_at	-1.5288	0.0046	-1.5020	0.0534	-1.8811	0.0001	-0.3256	0.5039	0.3352	0.1923	0.6608	0.0117	0.1086	0.9657	0.3122	0.6822	0.2035	0.8088
CG6293	CG6293	1625688_at	0.1123	0.9551	-2.1309	0.1012	-1.3666	0.1062	0.4524	0.3190	1.8326	0.0003	1.3802	0.0005	-0.4783	0.9581	-0.5265	0.8862	-0.0482	0.9914
---	---	1625689_at	0.0092	0.9596	-0.0384	0.7915	0.0498	0.7733	-0.0131	0.9857	0.0178	0.9422	0.0310	0.8776	0.0685	0.8846	0.0645	0.7656	-0.0039	0.9891
CG15485	CG15485	1625690_at	0.0992	0.8061	0.0233	0.8609	0.0024	0.9943	-0.1230	0.8844	-0.0414	0.9185	0.0816	0.8012	0.0724	0.9545	-0.0463	0.9390	-0.1188	0.7894
Tre1	trapped in endode	1625691_at	-1.3414	0.0774	-2.2310	0.0014	-1.7389	0.0071	0.2515	0.7005	0.8833	0.0102	0.6318	0.0267	-0.1533	0.9740	0.1116	0.9519	0.2648	0.8611
CG9865	CG9865	1625692_s_at	-0.1320	0.4597	0.1163	0.7668	-0.2139	0.1846	-0.3189	0.2729	-0.2455	0.1505	0.0734	0.6828	0.0769	0.9506	0.0658	0.9075	-0.0111	0.9849
---	---	1625693_at	-0.0255	0.8973	0.1513	0.2227	0.2095	0.2580	0.1142	0.8589	0.0590	0.8438	-0.0553	0.8350	-0.0013	0.9996	0.1361	0.6939	0.1375	0.6861
---	---	1625694_at	0.0531	0.7642	0.1355	0.4186	0.0984	0.5396	0.1544	0.7135	0.0999	0.6422	-0.0546	0.7996	0.0704	0.9405	0.1939	0.5183	0.1235	0.7040
Ets98B	Ets at 98B	1625695_at	-0.0833	0.7722	0.4020	0.1554	0.5772	0.0186	0.0194	0.9807	-0.3522	0.0738	-0.3716	0.0397	0.0932	0.9467	0.1945	0.6834	0.1013	0.8555
---	---	1625696_at	0.1378	0.3503	0.2282	0.4026	-0.0474	0.8592	-0.1085	0.8068	-0.0477	0.8320	0.0608	0.7514	0.1688	0.8222	-0.0526	0.9125	-0.2214	0.4924
Cyp6g1	CYP6-like	1625697_at	2.4230	0.0051	1.7202	0.0906	2.6722	0.0001	0.5552	0.5415	-0.3323	0.5060	-0.8875	0.0421	-0.4326	0.8354	-0.9790	0.2404	-0.5464	0.5488
CG6639	CG6639	1625698_at	0.4169	0.2961	0.5917	0.7672	0.1838	0.7610	0.8604	0.0840	1.1222	0.0029	0.2617	0.3182	1.0484	0.8425	0.9803	0.6847	-0.0681	0.9848
---	---	1625699_at	0.1315	0.5274	0.1858	0.3005	0.1172	0.5142	-0.0417	0.9538	-0.0740	0.7518	-0.0323	0.8907	0.0216	0.9841	-0.0404	0.9157	-0.0620	0.8444
cbx	male sterile(2)46C	1625700_a_at	-0.0508	0.7994	-0.3694	0.1230	-0.5481	0.0038	-0.0894	0.8033	0.3393	0.0260	0.4287	0.0058	0.0512	0.9589	0.0368	0.9358	-0.0145	0.9745
---	---	1625701_at	0.2869	0.2679	0.2100	0.3217	0.2536	0.3607	0.1960	0.6932	0.1883	0.4276	-0.0076	0.9788	0.2010	0.7644	0.2736	0.3259	0.0725	0.8426
Taf12	TBP-associated fa	1625702_s_at	-0.1673	0.4026	-0.1014	0.6324	0.0858	0.6616	0.1511	0.7293	-0.1135	0.5940	-0.2647	0.1283	0.0077	0.9953	0.0332	0.9504	0.0255	0.9574
---	---	1625703_at	0.1358	0.4657	0.2271	0.2385	-0.0065	0.9807	-0.1304	0.7507	-0.1630	0.3705	-0.0326	0.8776	0.0087	0.9946	0.0039	0.9954	-0.0049	0.9924
CG16879	CG16879	1625704_at	0.1897	0.3276	-0.0413	0.8324	0.1367	0.3704	0.0736	0.8967	0.1806	0.3427	0.1070	0.5631	0.0346	0.9764	0.0336	0.9408	-0.0010	0.9987
E(Pc)	transcript group III	1625705_s_at	0.1759	0.2748	0.2430	0.2985	-0.0083	0.9764	-0.3418	0.2500	-0.1891	0.2868	0.1527	0.3447	0.0934	0.9294	0.0722	0.8830	-0.0212	0.9666
Pros28.1B	20s proteasome	1625706_at	0.1150	0.6049	-0.0806	0.5427	0.2120	0.4683	0.1095	0.8640	0.0505	0.8658	-0.0590	0.8185	-0.1994	0.8023	-0.1437	0.7017	0.0556	0.9031
Ccp84Aa /// Ccp84Ab	cuticle cluster 8 ///	1625707_s_at	0.2418	0.1169	0.0651	0.6190	0.0639	0.7217	0.0278	0.9691	0.0675	0.7618	0.0397	0.8541	-0.0191	0.9862	-0.0843	0.7932	-0.0652	0.8418
---	---	1625708_at	0.1685	0.3808	0.1006	0.4555	0.2643	0.1743	0.0566	0.9422	-0.1499	0.5271	-0.2065	0.3065	-0.1417	0.8395	-0.1232	0.7043	0.0185	0.9659
CG34377	CG13859	1625709_at	0.0960	0.5179	0.0678	0.6110	0.0235	0.9036	-0.0728	0.8822	-0.0219	0.9253	0.0508	0.7868	0.0435	0.9589	0.0238	0.9505	-0.0197	0.9525
CG31717	CG31717	1625710_at	0.8275	0.0555	1.2234	0.0262	1.5781	0.0001	0.1135	0.7949	-0.4941	0.0132	-0.6075	0.0031	-0.3438	0.8065	-0.0624	0.9478	0.2814	0.6577
CG31530	CG31530	1625711_at	1.8760	0.3184	0.1160	0.3190	-0.1981	0.3481	-0.2654	0.3837	-0.1406	0.4299	0.1248	0.4403	0.1623	0.9872	-1.7677	0.4448	-1.9300	0.4114

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14780	CG14780	1625712_at	0.0668	0.6456	0.1100	0.5054	-0.1544	0.5477	-0.1292	0.8034	-0.0109	0.9733	0.1184	0.5652	0.1427	0.7893	0.0165	0.9648	-0.1261	0.6060
CG11263	CG11263	1625713_at	-0.0849	0.6142	-0.3095	0.0925	-0.3459	0.0531	0.1941	0.6086	0.5034	0.0164	0.3093	0.0717	-0.0337	0.9816	0.1100	0.7985	0.1437	0.7068
CG30423	CG30423	1625714_a_at	0.1534	0.5401	0.2212	0.2748	0.1607	0.4262	0.0310	0.9603	-0.0828	0.6577	-0.1138	0.4696	0.0499	0.9725	0.0605	0.9145	0.0106	0.9852
CG11462	CG11462	1625715_at	0.2995	0.2882	0.4094	0.0685	0.3939	0.0464	0.1535	0.7536	0.1025	0.6699	-0.0510	0.8326	0.0169	0.9923	0.0155	0.9842	-0.0013	0.9988
---	---	1625716_at	0.0462	0.7842	-0.0650	0.6893	0.0139	0.9396	-0.0234	0.9727	0.0554	0.7892	0.0788	0.6528	-0.0700	0.9057	-0.0389	0.9049	0.0311	0.9161
CG9350	CG9350	1625717_s_at	-0.4926	0.1250	-0.4951	0.1016	-0.1013	0.5940	-0.0315	0.9603	-0.5762	0.0053	-0.5447	0.0040	-0.2594	0.8141	-0.4694	0.2879	-0.2101	0.6731
---	---	1625718_at	0.1153	0.4534	0.0921	0.6110	-0.0523	0.7587	-0.1212	0.8076	-0.0793	0.7340	0.0420	0.8560	0.1340	0.7633	-0.0248	0.9341	-0.1588	0.4021
Atpalpha	Na,K ATPase alpl	1625719_at	-0.9293	0.0463	-1.5382	0.0186	-2.3893	0.0003	-0.1602	0.7608	0.7229	0.0072	0.8832	0.0018	0.6543	0.7220	0.0309	0.9838	-0.6234	0.4370
CG12259	CG12259	1625720_at	-0.0913	0.8099	1.4484	0.0263	1.3095	0.0003	-0.2041	0.7380	-0.7839	0.0107	-0.5798	0.0244	-0.1420	0.9201	0.7104	0.1374	0.8524	0.1114
Drl-2	Derailed 2	1625721_s_at	0.1137	0.4590	0.1327	0.4274	0.2053	0.2974	0.0739	0.8912	0.1070	0.5877	0.0331	0.8776	-0.2083	0.7196	-0.1064	0.6949	0.1018	0.7069
CG2774	CG2774	1625722_at	-0.1734	0.3904	0.9038	0.0193	0.7686	0.0028	-0.2252	0.5832	-0.5720	0.0141	-0.3468	0.0668	-0.1244	0.8561	0.3931	0.1504	0.5175	0.1011
CG14710	CG14710	1625723_at	0.0250	0.9383	-0.3005	0.0763	-0.6326	0.0109	-0.1564	0.7018	0.5332	0.0118	0.6896	0.0022	0.0680	0.9509	0.1820	0.6015	0.1140	0.7641
Pde1c	Phosphodiesteras	1625724_at	-1.2071	0.0074	-2.2480	0.0047	-2.0300	0.0006	0.0091	0.9937	0.7635	0.0054	0.7543	0.0034	-0.2204	0.7220	-0.2038	0.4418	0.0166	0.9666
CG30290	CG30290	1625725_at	-0.2086	0.4494	-0.2889	0.0852	-0.3133	0.0809	-0.1158	0.7777	-0.0185	0.9402	0.0973	0.5659	-0.0946	0.9340	-0.2094	0.5880	-0.1148	0.7936
---	---	1625726_at	0.2533	0.1398	0.0510	0.8722	-0.3511	0.1644	0.0099	0.9937	0.5127	0.0452	0.5028	0.0314	0.4506	0.6695	0.2309	0.6129	-0.2197	0.6324
CG32006	CG32006	1625727_at	-0.0377	0.8382	-0.2053	0.1100	-0.1439	0.5733	0.0271	0.9771	0.1676	0.4871	0.1405	0.5312	-0.0813	0.9156	-0.0005	0.9998	0.0809	0.8044
CG8866	CG8866	1625728_at	0.1752	0.4062	0.4237	0.0730	0.3641	0.0603	-0.1681	0.6721	-0.3855	0.0461	-0.2174	0.1955	-0.2025	0.7707	-0.1668	0.5973	0.0357	0.9318
---	---	1625729_at	-0.0235	0.9304	-0.0945	0.4675	-0.0619	0.7386	-0.0143	0.9890	0.0295	0.9298	0.0438	0.8776	-0.0132	0.9943	-0.1699	0.7248	-0.1567	0.7454
---	---	1625730_at	0.1956	0.4338	-0.0344	0.7216	0.0060	0.9855	0.0863	0.8590	0.1937	0.2701	0.1075	0.5282	0.0353	0.9640	-0.0458	0.8799	-0.0811	0.7408
CG13114	CG13114	1625731_at	0.1241	0.6800	-0.0739	0.7090	-0.3108	0.0863	-0.1528	0.8281	0.2728	0.3192	0.4256	0.0800	0.0295	0.9816	0.0340	0.9404	0.0046	0.9924
dpr8	dpr8	1625732_at	-0.1123	0.6264	-0.3240	0.0688	-0.4458	0.1014	0.0562	0.9562	0.3476	0.2069	0.2914	0.2396	-0.1317	0.8283	-0.0566	0.8764	0.0751	0.8119
CG31142	CG31142	1625733_at	0.1359	0.5511	0.4692	0.1268	0.4967	0.0990	0.1298	0.8024	0.0132	0.9676	-0.1166	0.5719	0.0244	0.9898	0.2695	0.5378	0.2450	0.5834
CG12184	CG12184	1625734_at	-0.0262	0.8994	-0.0607	0.6286	0.2743	0.2909	0.1099	0.7924	-0.0189	0.9380	-0.1288	0.4175	0.0049	0.9972	0.0286	0.9590	0.0237	0.9614
---	---	1625735_at	0.2918	0.1782	0.0523	0.7310	-0.0010	0.9972	0.0747	0.8942	0.3251	0.0838	0.2504	0.1333	0.0060	0.9964	0.0308	0.9521	0.0247	0.9574
---	---	1625736_at	0.1372	0.4649	-0.4492	0.0534	-0.8372	0.0148	-0.1069	0.9277	0.8546	0.0229	0.9615	0.0084	0.0191	0.9881	0.1166	0.7169	0.0975	0.7663
CG12133 /// DereCG12133	CG12133	1625737_at	0.5550	0.1316	0.0208	0.8400	0.0895	0.6280	0.2943	0.3837	0.4297	0.0319	0.1354	0.4532	0.0686	0.9405	0.0096	0.9868	-0.0590	0.8782
CG8420	CG8420	1625738_at	0.2469	0.4735	-0.1090	0.5406	-0.3254	0.0674	0.0190	0.9852	0.3050	0.1836	0.2860	0.1643	-0.0473	0.9726	-0.0279	0.9617	0.0194	0.9709
CG3292	CG3292	1625739_at	0.1653	0.4384	0.1204	0.4096	0.2214	0.2795	-0.0694	0.9242	-0.1173	0.6301	-0.0478	0.8503	0.0401	0.9589	0.0351	0.9198	-0.0050	0.9889
CG7466	CG7466	1625740_at	-0.5038	0.1658	-1.4792	0.0108	-1.5901	0.0007	-0.1053	0.9118	0.7828	0.0179	0.8881	0.0062	0.0822	0.9514	-0.0858	0.8831	-0.1680	0.7134
---	---	1625741_s_at	0.1784	0.3611	-0.1061	0.4445	0.0428	0.8549	0.2631	0.5842	0.4146	0.0916	0.1514	0.5254	-0.0066	0.9964	-0.0124	0.9854	-0.0058	0.9924
usnp	ubisnap	1625742_at	0.2956	0.0862	0.2149	0.1107	0.0026	0.9932	-0.1350	0.7795	0.1789	0.3826	0.3139	0.0810	0.1160	0.8326	0.0612	0.8432	-0.0548	0.8527
ninaB	beta-carotene dio	1625743_at	0.1860	0.2917	0.1064	0.5879	-0.1247	0.4888	-0.0618	0.9013	-0.0920	0.6037	-0.0302	0.8744	0.1644	0.7726	-0.0933	0.7454	-0.2576	0.3027
GstE6	Glutathione S tran	1625744_at	-0.2684	0.7964	0.7513	0.0154	1.0104	0.0622	-0.2110	0.8044	-0.3475	0.3115	-0.1366	0.7035	-0.4812	0.8930	0.7154	0.6164	1.1967	0.3810
CG31643	CG31643	1625745_at	-0.0665	0.6710	-0.2565	0.4354	-0.0048	0.9845	-0.0295	0.9620	0.0296	0.8943	0.0591	0.7375	-0.3092	0.7196	-0.1689	0.6686	0.1403	0.7323
CG3386	CG3386	1625746_at	-0.4848	0.0350	-0.7132	0.0152	-0.3683	0.0713	0.0399	0.9603	-0.2196	0.3131	-0.2596	0.1774	-0.3210	0.6660	-0.5529	0.0782	-0.2320	0.4524
RhoGAP92B	RhoGAP92B	1625747_at	-0.0397	0.9392	0.3622	0.1160	-0.0407	0.8469	-0.0401	0.9609	0.1509	0.5129	0.1910	0.3383	0.1608	0.9142	0.4912	0.3340	0.3304	0.5488
par-6	par-6	1625748_s_at	-0.5949	0.0271	-0.6125	0.0688	-0.6590	0.0046	-0.1075	0.8134	-0.1298	0.4908	-0.0223	0.9193	0.0218	0.9875	0.0031	0.9972	-0.0187	0.9692
CG14937	CG14937	1625749_at	0.0441	0.7972	0.2276	0.2734	0.4233	0.0182	-0.0029	0.9956	-0.1046	0.5601	-0.1017	0.5307	-0.1977	0.7131	-0.1102	0.6512	0.0875	0.7342
CG17272	CG17272	1625750_at	-0.0263	0.9375	0.0676	0.8895	0.2207	0.2610	-0.0706	0.8800	0.0590	0.7581	0.1297	0.3826	-0.2223	0.8692	0.2693	0.6425	0.4916	0.3711
Tim17b1	Tim17b1	1625751_at	-0.0252	0.8836	0.0348	0.7359	0.0799	0.6460	0.0929	0.8550	-0.0049	0.9859	-0.0978	0.5946	-0.0786	0.8903	0.0455	0.8830	0.1240	0.5868
TpnC41C	Troponin C at 41C	1625752_at	-2.2876	0.0402	-0.0761	0.5185	-1.7115	0.0152	-1.4711	0.2729	-2.8366	0.0042	-1.3655	0.0576	0.0643	0.9852	-0.7936	0.3006	-0.8579	0.2917
---	---	1625753_at	0.1962	0.2973	0.0952	0.4658	0.0294	0.8649	-0.0708	0.8676	-0.1080	0.4920	-0.0372	0.8279	0.0992	0.8461	0.0918	0.7001	-0.0074	0.9837
---	---	1625754_s_at	0.0544	0.8060	0.0000	1.0000	-0.0763	0.6911	-0.0860	0.8470	0.0887	0.6248	0.1748	0.2322	0.0538	0.9499	0.0618	0.8619	0.0080	0.9849
CG1291	CG1291	1625755_at	0.4467	0.1055	0.5356	0.1592	0.3838	0.0282	-0.0033	0.9956	0.4368	0.0172	0.4401	0.0102	0.1564	0.9096	0.5757	0.2254	0.4194	0.4020
CG6656	CG6656	1625756_at	-0.0312	0.9466	-0.2817	0.6515	-0.4686	0.0217	-0.4147	0.1950	-0.5025	0.0176	-0.0878	0.6615	-0.1732	0.9405	-0.5881	0.4073	-0.4148	0.5868
gfzf	GST-containing FI	1625757_at	-0.5754	0.0223	0.1995	0.2039	-0.3333	0.0531	-0.4324	0.2017	-0.9339	0.0015	-0.5015	0.0139	0.2770	0.6587	-0.1466	0.5896	-0.4236	0.1300
pit	pitchoune	1625758_s_at	0.5113	0.1825	0.6796	0.3634	1.1287	0.0016	0.4666	0.3596	0.6902	0.0255	0.2236	0.4097	-0.0036	0.9994	0.8048	0.2455	0.8084	0.2769
Ku80	Ku80	1625759_at	-0.2548	0.5157	-0.1742	0.4841	0.0403	0.9108	0.3280	0.3328	0.6962	0.0044	0.3682	0.0417	0.0510	0.9873	0.6837	0.3350	0.6327	0.3930
CG30272	CG30272	1625760_at	0.4845	0.2024	0.0759	0.5170	0.5797	0.0089	-0.0005	0.9994	-0.0114	0.9619	-0.0108	0.9573	-0.1140	0.9168	0.0085	0.9925	0.1225	0.7850
Mmp1	Matrix metalloprot	1625761_a_at	-0.3604	0.5245	0.0995	0.9432	0.8760	0.1036	0.7957	0.4741	0.8019	0.1772	0.0061	0.9934	0.1225	0.9831	1.1978	0.3667	1.0753	0.4370
CG32188 /// CG32189	CG32189 /// CG3	1625762_s_at	0.3864	0.1991	0.2829	0.2112	0.0626	0.7137	-0.1354	0.6506	0.0349	0.8521	0.1703	0.1857	0.0531	0.9734	0.0092	0.9924	-0.0440	0.9362
CG2789	CG2789	1625763_at	-0.7280	0.0383	-0.4282	0.0359	-0.3879	0.0548	0.0287	0.9592	-0.7326	0.0011	-0.7614	0.0006	-0.0692	0.9705	-0.5692	0.2168	-0.5000	0.3047
CG17572	CG17572	1625764_at	-0.0749	0.7857	0.0181	0.8767	-0.1218	0.4592	-0.1085	0.7845	-0.1246	0.4661	-0.0160	0.9370	-0.0480	0.9623	-0.0032	0.9961	0.0449	0.9086
---	---	1625765_at	0.0627	0.8067	0.6227	0.0574	-0.2162	0.2363	-0.3109	0.5068	-0.3828	0.1215	-0.0719	0.7968	0.4843	0.4443	0.2795	0.3914	-0.2049	0.5587
CG15030	CG15030	1625766_at	0.0133	0.9450	-0.1178	0.3103	-0.3454	0.2041	0.0114	0.9937	0.1531	0.6188	0.1417	0.6146	0.0808	0.9077	-0.0780	0.8061	-0.1588	0.5444
CG30056	CG30056	1625767_at	0.0747	0.7070	0.0015	0.9905	-0.0131	0.9677	-0.2666	0.5175	-0.1155	0.6339	0.1511	0.4632	-0.0191	0.9831	0.0757	0.7588	0.0948	0.6748
lola	longitudinals abse	1625768_s_at	0.1714	0.8547	-0.7639	0.2391	-1.6984	0.0152	-0.7307	0.5020	1.2823	0.0337	2.0130	0.0028	0.1321	0.9816	0.2890	0.8698	0.1569	0.9277
CG1113	CG1113	1625769_at	0.1648	0.4781	-0.0082	0.9403	0.2606	0.3297	-0.0913	0.8950	-0.0729	0.7963	0.0185	0.9495	-0.2089	0.6832	-0.1936	0.3465	0.0153	0.9603
CG15011	CG15011	1625770_at	0.7488	0.0056	0.3230	0.3947	0.8329	0.0009	0.3496	0.1332	0.7210	0.0009	0.3714	0.0100	0.0142	0.9943	0.4438	0.3098	0.4296	0.3538
Dll	Distalless	1625771_at	0.4783	0.1185	0.1293	0.2623	0.2076	0.4713	-0.0110	0.9922	-0.0094	0.9790	0.0015	0.9958	0.1818	0.8049	-0.1302	0.7088	-0.3120	0.3210
---	---	1625772_at	0.1326	0.4560	-0.1702	0.3197	-0.0598	0.7981	0.2684	0.6202	0.3864	0.1504	0.1180	0.6760	0.0930	0.8825	0.0912	0.7469	-0.0018	0.9965
CG6289	CG6289	1625773_at	0.0344	0.8922	0.0782	0.4330	0.1086	0.4921	-0.0331	0.9592	-0.1999	0.2393	-0.1668	0.2769	0.0088	0.9950	-0.1073	0.7922	-0.1161	0.7567
CG32838	CG32838	1625774_at	-0.0225	0.9049	-0.8465	0.0163	-0.9698	0.0282	0.0550	0.9744	0.7478	0.0759	0.6928	0.0670	-0.0950	0.8461	-0.0923	0.6823	0.0027	0.9940
csw	Enhancer of sever	1625775_a_at	-0.7462	0.0397	0.2073	0.6518	-1.0126	0.0009	-0.8117	0.0657	-1.1137	0.0018	-0.3020	0.1814	0.4405	0.7464	-0.1334	0.8716	-0.5739	0.3324
CG2291	CG2291	1625776_at	0.2132	0.3732	0.0917	0.4195	0.3937	0.0375	0.0013	0.9988	-0.0952	0.6872	-0.0965	0.6507	-0.2447	0.7187	-0.1273	0.6859	0.1174	0.7128
CG12692	CG12692	1625777_at	0.0986	0.6318	-0.0489	0.8048	-0.0036	0.9855	0.0805	0.8508	0.0334	0.8718	-0.0471	0.7872	0.1073	0.8692	0.0695	0.8422	-0.0379	0.9159
CG4562 /// DwilCG4562	CG4562	1625778_at	0.0175	0.9622	-0.0100	0.9230	0.1369	0.4985	0.0942	0.9034	-0.1754	0.5093	-0.2696	0.2312	0.0074	0.9948	-0.0130	0.9782	-0.0204	0.9570
fd59A	forkhead domain !	1625779_at	0.0028	0.9883	0.0939	0.3996	0.1246	0.4814	0.0348	0.9452	-0.0765	0.6369	-0.1113	0.4101	0.0215	0.9823	0.0328	0.9265	0.0113	0.9745
CG9812	CG9812	1625780_a_at	-2.9724	0.0066	0.3769	0.3385	-1.7862	0.0106	-2.2107	0.0660	-4.8608	0.0003	-2.6501	0.0019	0.0446	0.9916	-1.2878	0.1772	-1.3324	0.1993

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14194	CG14194	1625781_at	-0.4091	0.2008	-0.5599	0.0257	-0.5530	0.0154	0.0712	0.9400	0.4619	0.0918	0.3907	0.1090	0.1085	0.9365	0.2242	0.6310	0.1156	0.8310
CG14010	CG14010	1625782_at	0.0924	0.6991	0.1942	0.3264	0.0460	0.8437	-0.0655	0.9247	0.0262	0.9286	0.0918	0.6710	0.0836	0.9400	0.2589	0.4488	0.1753	0.6328
CG9445	CG9445	1625783_at	0.0516	0.7911	-0.1383	0.3766	0.1429	0.4178	0.2702	0.5217	0.3439	0.1186	0.0736	0.7613	-0.0562	0.9366	0.0474	0.8831	0.1036	0.6719
---	---	1625784_s_at	-0.1733	0.2529	0.0643	0.6550	-0.0636	0.7220	-0.1564	0.6823	-0.2747	0.1212	-0.1183	0.4869	0.0602	0.9390	-0.0620	0.8511	-0.1222	0.6389
l(1)G0196	lethal (1) G0196	1625785_at	0.3830	0.3030	-0.1798	0.4558	0.4139	0.3085	0.0325	0.9873	0.3992	0.4327	0.3666	0.4257	-0.5379	0.6898	-0.0793	0.9275	0.4586	0.4114
---	---	1625786_at	0.2187	0.2962	0.2594	0.1609	0.2385	0.3042	-0.1217	0.8164	0.0079	0.9801	0.1296	0.5157	-0.0042	0.9952	-0.0243	0.9296	-0.0201	0.9343
---	---	1625787_at	0.1176	0.4288	-0.0661	0.5431	0.0311	0.9178	0.1928	0.5896	0.2404	0.1825	0.0476	0.8166	-0.0173	0.9848	0.1089	0.6237	0.1262	0.5637
---	---	1625788_at	0.0794	0.6775	0.0744	0.4981	0.2678	0.1301	0.1144	0.7680	-0.0648	0.7390	-0.1793	0.2249	0.0232	0.9848	0.0589	0.8794	0.0357	0.9237
CG5790	CDC 7-related	1625789_at	0.0957	0.6087	-0.1114	0.5179	-0.1619	0.4646	0.2589	0.4162	0.3520	0.0511	0.0931	0.5994	0.0593	0.9503	-0.0243	0.9588	-0.0835	0.8112
CG11585	CG11585	1625790_at	-0.2022	0.4725	0.1080	0.6346	-0.0327	0.8790	-0.1006	0.8827	-0.1330	0.6016	-0.0324	0.9086	0.1557	0.8673	0.0623	0.9129	-0.0934	0.8435
---	---	1625791_s_at	-0.2475	0.2637	-1.0775	0.0395	-1.2446	0.0138	-0.0618	0.9777	0.8834	0.1090	0.9451	0.0597	-0.0222	0.9816	0.0170	0.9634	0.0392	0.8991
l(3)neo26	lethal (3) neo26	1625792_s_at	0.2986	0.5692	-0.5948	0.4215	-0.7002	0.1838	-0.8173	0.0347	0.6511	0.0087	1.4684	0.0002	-0.6946	0.8156	-0.2030	0.9152	0.4917	0.7251
Dref	DNA replication-re	1625793_at	0.2078	0.2286	0.1376	0.7709	-0.2699	0.1914	-0.2179	0.5197	0.4617	0.0177	0.6796	0.0019	0.2317	0.8379	0.4499	0.3236	0.2182	0.6677
CG5830	CG5830	1625794_at	-1.0542	0.0021	0.1404	0.7123	-0.1912	0.3817	-0.4547	0.1032	-1.1460	0.0003	-0.6913	0.0013	-0.1826	0.8837	0.0686	0.9274	0.2513	0.6225
CG3394	CG3394	1625795_a_at	1.9526	0.0038	-0.1066	0.8437	1.3750	0.0001	0.1240	0.9380	0.6282	0.1764	0.5042	0.2278	-1.6296	0.1902	-1.5801	0.0391	0.0495	0.9582
CG14231	CG14231	1625796_at	-0.0564	0.7829	-0.1180	0.8542	-0.2451	0.3248	-0.2016	0.6597	-0.2320	0.2958	-0.0303	0.9093	-0.0225	0.9946	-0.3570	0.6762	-0.3345	0.6973
CG11063	CG11063	1625797_at	-0.2393	0.6335	0.2228	0.8309	0.5014	0.1715	0.2654	0.7193	-0.1205	0.7640	-0.3858	0.1963	-0.0085	0.9990	0.2601	0.8546	0.2687	0.8375
---	---	1625798_at	0.0898	0.6564	-0.0418	0.8180	-0.0156	0.9441	-0.0488	0.9304	0.0741	0.6941	0.1229	0.4231	0.0282	0.9717	-0.0248	0.9399	-0.0530	0.8348
CG7196	CG7196	1625799_a_at	-0.0886	0.6483	0.0445	0.8617	0.0684	0.7802	-0.0795	0.8841	-0.1682	0.3767	-0.0886	0.6431	-0.0710	0.9474	-0.0289	0.9577	0.0421	0.9260
CG4074	CG4074	1625800_at	-0.7541	0.0049	0.4908	0.0945	0.2395	0.2235	-0.1955	0.7759	-0.7176	0.0207	-0.5222	0.0474	-0.0442	0.9530	0.5578	0.0371	0.6019	0.0422
skpA	skpA	1625801_s_at	-0.1115	0.4947	0.1239	0.2796	0.0469	0.7973	-0.0474	0.9434	-0.0778	0.7230	-0.0304	0.8929	0.0412	0.9611	0.0799	0.7860	0.0387	0.9057
CG13135	CG13135	1625802_a_at	0.1715	0.3929	0.1622	0.3133	0.0610	0.8128	0.2771	0.6101	0.0004	0.9992	-0.2767	0.2597	0.1576	0.8049	-0.0419	0.9191	-0.1994	0.4677
---	---	1625803_at	0.0952	0.6053	0.1644	0.3604	0.2032	0.2787	0.1186	0.8085	0.0791	0.7284	-0.0395	0.8619	0.0541	0.9467	0.1164	0.6675	0.0623	0.8428
CG17082	CG17082	1625804_s_at	-0.1185	0.5179	-0.4862	0.1097	-0.6907	0.0040	-0.1761	0.5932	0.1733	0.3045	0.3494	0.0273	0.1402	0.8882	-0.0991	0.8450	-0.2393	0.5454
---	---	1625805_at	0.1100	0.5620	-0.1701	0.4074	0.0068	0.9846	0.1907	0.6958	0.1109	0.6643	-0.0798	0.7420	0.0404	0.9677	-0.1804	0.5071	-0.2208	0.4093
Pp1alpha-96A	Protein phosphata	1625806_at	0.2129	0.4396	0.6167	0.0098	0.7894	0.0040	0.0454	0.9473	-0.5352	0.0136	-0.5807	0.0057	-0.0037	0.9984	-0.1424	0.7424	-0.1387	0.7442
---	---	1625807_at	0.1366	0.6462	0.1258	0.4282	0.0938	0.5826	0.0859	0.8584	0.0017	0.9945	-0.0842	0.6354	0.0410	0.9738	-0.0620	0.8903	-0.1031	0.7764
---	---	1625808_at	0.2124	0.4305	0.0323	0.8685	0.3113	0.1535	0.1079	0.9191	-0.0163	0.9743	-0.1242	0.7205	-0.1325	0.8202	-0.1765	0.4740	-0.0440	0.8949
---	---	1625809_at	0.1276	0.6049	0.1077	0.3300	-0.0837	0.6701	-0.0960	0.7975	-0.0723	0.6757	0.0236	0.8972	0.1629	0.8141	0.0185	0.9689	-0.1444	0.6394
CG13937	CG13937	1625810_s_at	-1.0884	0.0109	-0.8079	0.0637	-1.3344	0.0004	-0.2159	0.5948	-0.3077	0.1337	-0.0919	0.6672	0.3748	0.5126	-0.0591	0.8792	-0.4339	0.1392
---	---	1625811_at	0.1542	0.4494	0.0597	0.7218	0.1955	0.2819	-0.0058	0.9956	0.0512	0.8667	0.0570	0.8287	0.0611	0.9361	0.0594	0.8592	-0.0018	0.9970
---	---	1625812_at	0.1695	0.2775	-0.0497	0.8120	0.1438	0.3822	0.1350	0.7138	0.1513	0.3781	0.0163	0.9377	-0.0611	0.9421	-0.1152	0.6928	-0.0540	0.8779
CG8290	CG8290	1625813_at	0.2164	0.5223	-0.5773	0.0945	-0.4396	0.0485	-0.1253	0.8521	0.6613	0.0146	0.7866	0.0040	-0.1577	0.8960	-0.0353	0.9621	0.1224	0.8289
CG8230	CG8230	1625814_at	0.4604	0.0694	1.1234	0.0088	1.0538	0.0007	0.1247	0.7672	-0.1673	0.3605	-0.2921	0.0709	0.1529	0.8802	0.4228	0.2584	0.2698	0.5032
---	---	1625815_at	0.0685	0.6435	-0.0464	0.7494	0.0688	0.6584	0.1591	0.6372	0.1041	0.5602	-0.0550	0.7614	-0.0274	0.9717	0.0008	0.9994	0.0282	0.9171
CG4781	CG4781	1625816_at	0.0886	0.8827	0.1326	0.2194	0.0514	0.7710	0.1255	0.9376	-0.1473	0.7979	-0.2728	0.5540	-0.0369	0.9869	-0.3016	0.5776	-0.2646	0.6312
CG14868	CG14868	1625817_at	0.5215	0.3101	0.1030	0.7290	0.6142	0.0299	0.4645	0.5311	1.4096	0.0039	0.9452	0.0143	-0.0592	0.9862	0.9668	0.2102	1.0260	0.2136
CG14421	CG14421	1625818_at	-0.0147	0.9472	-0.0566	0.7647	-0.1994	0.3089	-0.1466	0.7143	-0.2552	0.1607	-0.1086	0.5441	-0.0141	0.9898	-0.1335	0.6093	-0.1193	0.6498
sno	Strawberry Notch	1625819_at	0.6289	0.0598	0.6049	0.1792	0.4131	0.1957	-0.1413	0.8974	0.1670	0.6856	0.3083	0.3517	0.1801	0.9092	0.2212	0.7339	0.0411	0.9596
---	---	1625820_s_at	0.0248	0.8816	-0.0141	0.8943	0.1219	0.5205	0.1018	0.8676	0.0247	0.9356	-0.0771	0.7403	-0.0361	0.9677	0.0115	0.9784	0.0477	0.8835
---	---	1625821_at	0.2223	0.4840	0.1279	0.3492	0.3065	0.0813	0.0794	0.9295	0.1270	0.6717	0.0476	0.8786	-0.0055	0.9966	0.0503	0.9225	0.0558	0.9029
CG32719	CG32719	1625822_at	-0.0617	0.7524	-0.2700	0.3787	-0.2458	0.1746	0.2394	0.6506	0.4211	0.0987	0.1817	0.4501	0.0539	0.9526	0.1500	0.6042	0.0961	0.7585
CG1075	CG1075	1625823_at	0.2632	0.1684	0.0621	0.5608	0.2150	0.2100	0.0298	0.9633	0.0406	0.8555	0.0108	0.9597	-0.0788	0.8815	-0.0506	0.8553	0.0282	0.9197
CG9779 /// DsimCG9779	CG9779	1625824_at	-1.0042	0.0047	-0.5327	0.0659	-0.4168	0.0800	0.0830	0.8244	-0.2010	0.1635	-0.2840	0.0355	-0.0191	0.9923	0.3999	0.3667	0.4190	0.3645
BM-40-SPARC	BM-40-SPARC	1625825_at	-0.8731	0.0088	-0.9826	0.2331	-0.7104	0.0117	-0.0329	0.9688	-0.3300	0.1301	-0.2971	0.1272	-0.3878	0.8461	-0.8810	0.2673	-0.4932	0.5725
CG7514	CG7514	1625826_at	0.1754	0.3328	0.0190	0.8992	0.2476	0.1464	-0.0645	0.8942	-0.0501	0.7998	0.0144	0.9431	-0.0826	0.8878	0.0485	0.8773	0.1312	0.5746
gw	gawky	1625827_s_at	-0.2027	0.1805	-0.1155	0.6581	-0.4078	0.0492	-0.1802	0.6409	0.0789	0.7194	0.2590	0.1229	0.1087	0.9030	0.1644	0.6410	0.0557	0.9020
CG11841	CG11841	1625828_at	2.6013	0.0014	1.6371	0.0996	2.1765	0.0002	0.6352	0.3019	0.1706	0.6794	-0.4646	0.1499	0.0452	0.9914	-0.8012	0.3471	-0.8464	0.3473
CG6784	CG6784	1625829_a_at	0.1356	0.5573	0.0125	0.9472	-0.0555	0.8166	0.0158	0.9834	-0.1205	0.5106	-0.1363	0.3981	-0.1182	0.9095	-0.2356	0.5411	-0.1174	0.7915
slim	scruin like at the n	1625830_a_at	-0.1068	0.6053	-0.3648	0.2226	-0.5389	0.0261	-0.0270	0.9744	0.1973	0.3370	0.2243	0.2150	0.1223	0.9137	-0.0602	0.9221	-0.1826	0.6717
---	---	1625831_at	-0.0034	0.9883	-0.0312	0.7964	-0.0759	0.6552	0.0167	0.9812	0.1033	0.5769	0.0866	0.6176	0.0049	0.9952	-0.0271	0.9353	-0.0321	0.9109
---	---	1625832_at	0.1491	0.4345	0.1276	0.4108	0.2745	0.1632	0.0261	0.9703	-0.0359	0.8801	-0.0620	0.7481	-0.1744	0.7726	-0.0814	0.8083	0.0930	0.7568
CG9896	CG9896	1625833_at	2.4237	0.0048	1.2964	0.1478	1.2368	0.0001	0.0179	0.9937	0.6924	0.1184	0.6745	0.0903	-0.0204	0.9948	-0.4343	0.5654	-0.4139	0.5917
---	---	1625834_at	-0.1041	0.5466	-0.5720	0.1005	-0.5711	0.0380	0.1369	0.8392	0.4776	0.0622	0.3407	0.1294	-0.0363	0.9611	-0.0202	0.9519	0.0161	0.9575
yellow-f2	yellow-f2	1625835_at	0.2286	0.2544	-0.5852	0.2496	0.0826	0.7577	0.0683	0.9376	-0.3522	0.1638	-0.4205	0.0674	-0.6518	0.6272	-1.2301	0.0509	-0.5783	0.3141
CG4036	CG4036	1625836_at	0.1115	0.6307	-0.1966	0.6838	0.2637	0.1027	0.2023	0.6041	0.0617	0.7980	-0.1406	0.4475	-0.1029	0.9514	-0.1059	0.8836	-0.0030	0.9979
CG17244 /// DyakCG17244	CG17244	1625837_at	2.6208	0.0014	1.2063	0.1792	2.5591	0.0001	0.7237	0.1902	0.6288	0.0637	-0.0949	0.8007	-0.6513	0.7644	-0.7988	0.3789	-0.1475	0.9083
Rab3	Rab-protein 3	1625838_at	-0.7139	0.2123	-0.1339	0.4422	-0.5729	0.0707	-0.4662	0.6175	-0.9772	0.0422	-0.5111	0.2202	-0.0264	0.9874	-0.2746	0.4753	-0.2481	0.5334
CG3267 /// X11Lbeta	CG3267 /// X11Lb	1625839_at	-0.0411	0.8424	0.7283	0.0574	0.9652	0.0064	-0.0426	0.9436	-0.5763	0.0058	-0.5337	0.0049	-0.2702	0.8292	0.1865	0.7703	0.4567	0.3921
CG12001	CG12001	1625840_at	-0.2171	0.5640	-0.1363	0.7362	-0.1686	0.5106	-0.0816	0.8816	-0.0388	0.8759	0.0428	0.8427	-0.0966	0.9684	-0.1358	0.8836	-0.0392	0.9678
CG11342 /// DyakCG11342	CG11342	1625841_at	0.0465	0.8137	0.0369	0.8437	0.0325	0.9049	-0.0573	0.9346	0.0384	0.8906	0.0958	0.6536	-0.0856	0.9092	0.0042	0.9941	0.0898	0.7749
---	---	1625842_at	0.1130	0.5647	-0.0232	0.8266	0.1021	0.5581	0.0941	0.8449	0.1217	0.5181	0.0276	0.8972	-0.0063	0.9950	-0.0443	0.8963	-0.0381	0.9031
Pif1A /// Pif1B	PFTAIRE-interacti	1625843_s_at	-1.4207	0.0057	-1.4815	0.0457	-1.9089	0.0000	-0.3851	0.1787	-0.1574	0.3750	0.2277	0.1411	0.0500	0.9862	-0.2129	0.8058	-0.2629	0.7346
apt	tracheae defective	1625844_s_at	0.3634	0.0416	0.6516	0.1896	0.3670	0.0321	-0.0851	0.9067	-0.0146	0.9679	0.0705	0.7872	0.2489	0.7953	0.2413	0.5668	-0.0075	0.9915
D1	Chromosomal pro	1625845_s_at	-0.0045	0.9891	-0.0870	0.5572	-0.2515	0.1460	0.1423	0.7688	0.5071	0.0218	0.3648	0.0516	0.2888	0.6999	0.2681	0.3872	-0.0207	0.9646
CG12439	CG12439	1625846_at	-0.1331	0.5179	-0.0328	0.8132	0.0055	0.9801	-0.0713	0.8776	-0.1096	0.5152	-0.0383	0.8332	-0.0297	0.9768	-0.0470	0.8980	-0.0173	0.9614
CG9034	CG9034	1625847_at	-0.1888	0.4463	0.4349	0.3466	0.2694	0.2716	-0.1321	0.8578	-0.7091	0.0158	-0.5770	0.0235	0.0075	0.9976	-0.1291	0.8749	-0.1367	0.8537
---	---	1625848_at	-0.4338	0.0710	-0.0666	0.5548	-0.6960	0.0531	-0.5605	0.0760	-0.1451	0.4436	0.4154	0.0191	-0.1086	0.9457	0.1999	0.7166	0.3084	0.5448
REG	REG	1625849_at	0.1378	0.4591	0.5864	0.0836	0.2227	0.2047	0.1276	0.6908	-0.0726	0.6676	-0.2002	0.1303	0.2762	0.6389	0.3711	0.1253	0.0949	0.7306

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	**Gene Title**	**Affy ID**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**
odd	Odd-skipped	1625850_at	-3.2609	0.0011	-1.0264	0.3495	-3.8765	0.0010	-1.9970	0.1462	-2.9480	0.0044	-0.9511	0.1993	0.7408	0.7893	-0.7944	0.5161	-1.5352	0.2130
Pp2A-29B	regulatory subunit	1625851_s_at	-0.4744	0.0448	0.2833	0.3703	0.3455	0.0881	-0.0930	0.8578	-0.6078	0.0065	-0.5147	0.0083	-0.1802	0.8427	0.0983	0.8468	0.2785	0.4710
DmirCG32227 /// gogo	golden goal /// CG	1625852_at	-0.1568	0.7350	-0.1762	0.6026	0.0870	0.6044	0.2020	0.6506	0.0643	0.8094	-0.1377	0.5080	-0.1037	0.9701	-0.0593	0.9593	0.0444	0.9665
CG15124	CG15124	1625853_at	-0.0606	0.7867	-0.0479	0.7759	0.0703	0.7091	-0.0628	0.9527	-0.1449	0.6613	-0.0821	0.8030	-0.0842	0.9029	0.0450	0.9068	0.1292	0.6328
CG3925	CG3925	1625854_at	-0.3424	0.0987	0.0606	0.6540	-0.1707	0.3669	-0.1846	0.6122	-0.4004	0.0355	-0.2158	0.1862	-0.0598	0.9550	-0.0421	0.9325	0.0177	0.9703
MED30	Mediator complex	1625855_at	-0.0316	0.8550	-0.1187	0.7019	0.1907	0.2986	0.1994	0.5735	0.4750	0.0170	0.2756	0.0896	-0.1154	0.9063	0.3048	0.3760	0.4202	0.2497
dm	diminutive	1625856_at	0.6534	0.0535	0.7122	0.1738	1.2678	0.0001	-0.1910	0.6350	-0.3913	0.0523	-0.2002	0.2599	-0.6841	0.4224	-0.1695	0.7541	0.5146	0.2743
CG8451	CG8451	1625857_at	1.9180	0.0078	1.8703	0.0108	2.2669	0.0000	-0.0635	0.9641	-0.1493	0.7336	-0.0858	0.8422	-0.4883	0.6749	-0.1951	0.7175	0.2931	0.5579
CG31785	CG31785	1625858_at	0.1227	0.5661	0.2030	0.2583	0.3662	0.0707	0.1155	0.7732	-0.0422	0.8464	-0.1577	0.3012	-0.0227	0.9884	0.0009	0.9996	0.0236	0.9622
Cct2	CTP:phosphochol	1625859_at	0.2651	0.6853	0.1963	0.2377	0.2548	0.1755	0.2078	0.6705	0.1824	0.4491	-0.0253	0.9288	0.2270	0.9092	0.2231	0.8053	-0.0039	0.9979
CG18542 /// CG32939	CG18542 /// CG3	1625860_s_at	-0.5187	0.1688	0.0527	0.8886	-0.2261	0.1479	-0.4160	0.0976	-0.4530	0.0084	-0.0370	0.8252	-0.1002	0.9589	0.0341	0.9704	0.1343	0.8514
CG13012	CG13012	1625861_at	-1.2033	0.0044	-2.7208	0.0034	-2.4060	0.0011	0.7219	0.4690	1.4971	0.0130	0.7752	0.1058	0.2006	0.8609	-0.1457	0.8068	-0.3463	0.4699
CG40178	CG40178	1625862_x_at	-0.8429	0.0395	0.1935	0.4756	0.2690	0.4181	-0.0776	0.9314	-1.1469	0.0019	-1.0693	0.0015	-0.1527	0.9246	-0.1144	0.8831	0.0383	0.9614
CG15577	CG15577	1625863_at	0.2443	0.2934	0.1129	0.5926	0.4326	0.0254	0.0547	0.9313	-0.0247	0.9250	-0.0794	0.6862	-0.2012	0.8062	-0.2408	0.4939	-0.0397	0.9343
CG32105	CG32105	1625864_at	-0.0262	0.9155	-0.2587	0.1424	0.0574	0.7585	0.1255	0.8085	0.0582	0.8212	-0.0673	0.7669	-0.1622	0.7506	-0.1891	0.3774	-0.0268	0.9299
CG14454	CG14454	1625865_at	0.0109	0.9628	-0.0027	0.9836	-0.0321	0.8610	-0.0565	0.9110	-0.0332	0.8758	0.0232	0.9039	0.0101	0.9915	-0.0678	0.8127	-0.0779	0.7607
Lcp65Aa	Lcp65Aa	1625866_at	0.1306	0.6392	0.4644	0.3104	0.2667	0.2621	-0.1159	0.9090	-0.4453	0.1657	-0.3294	0.2569	0.1248	0.8784	0.0316	0.9505	-0.0932	0.8111
---	---	1625867_at	-0.0505	0.8478	0.0430	0.7025	0.2449	0.1185	-0.0172	0.9807	-0.1335	0.4590	-0.1163	0.4810	-0.1582	0.8243	0.0288	0.9519	0.1870	0.5515
---	---	1625868_at	-0.0008	0.9966	0.1187	0.4347	0.3403	0.0895	0.3104	0.4602	0.1463	0.5506	-0.1641	0.4444	0.1376	0.8192	0.1826	0.4631	0.0450	0.8943
Mst87F	Male-specific RNA	1625869_at	0.0114	0.9705	-0.2432	0.2199	-0.2941	0.1372	-0.0333	0.9649	0.1100	0.6253	0.1433	0.4544	-0.1698	0.7811	-0.1529	0.5855	0.0169	0.9659
CG14223	CG14223	1625870_at	0.4948	0.0124	0.7189	0.0836	0.8799	0.0243	0.1084	0.8660	0.0115	0.9741	-0.0969	0.6832	0.1336	0.9467	0.3447	0.5922	0.2111	0.7647
Gr22c	Gustatory recepto	1625871_at	0.1772	0.3655	0.0336	0.7543	-0.3733	0.2102	-0.2065	0.8000	0.1038	0.7966	0.3102	0.2973	0.1644	0.7215	0.0286	0.9265	-0.1358	0.5021
CG34021	CG34021	1625872_at	0.0395	0.8789	0.0138	0.9375	-0.0976	0.5993	-0.0465	0.9564	0.0001	0.9997	0.0466	0.8618	-0.0022	0.9984	-0.1107	0.6450	-0.1085	0.6486
CG15661	CG15661	1625873_at	-0.1401	0.7273	-0.0991	0.8399	-0.7665	0.0343	-1.0919	0.1390	-1.2223	0.0140	-0.1305	0.7943	-0.4324	0.7691	-1.2198	0.0689	-0.7874	0.2235
CG6761	CG6761	1625874_at	0.0842	0.6838	0.1361	0.2376	-0.0166	0.9591	-0.2344	0.5080	-0.2573	0.1681	-0.0229	0.9209	0.0682	0.9039	0.0091	0.9810	-0.0591	0.8177
CG10635	CG10635	1625875_at	-0.9880	0.0015	-1.1185	0.0359	-1.0381	0.0006	0.1750	0.6615	0.5201	0.0146	0.3451	0.0493	0.1630	0.8425	0.5592	0.1047	0.3962	0.2557
AP-1sigma	AP-1sigma	1625876_at	0.4742	0.0114	0.5445	0.0929	0.4344	0.0171	-0.0741	0.8707	0.2206	0.1640	0.2947	0.0444	-0.0332	0.9752	0.1970	0.4593	0.2302	0.3921
CG40084 /// CG40271 /// C	CG40084 /// CG4(	1625877_s_at	-0.1579	0.5637	-1.2919	0.2145	-1.5348	0.0040	-0.3564	0.4415	1.4255	0.0007	1.7819	0.0002	0.0520	0.9914	0.4469	0.7059	0.3949	0.7440
Aats-cys	Cysteinyl-tRNA sy	1625878_at	0.5036	0.0647	1.0075	0.0412	0.8385	0.0006	0.1612	0.6854	0.2730	0.1379	0.1118	0.5353	0.2938	0.7633	0.8300	0.0637	0.5362	0.2095
Pomp	Pomp	1625879_at	0.2606	0.3958	0.6603	0.1849	0.2504	0.0998	-0.0748	0.9451	-0.2472	0.4429	-0.1724	0.5769	0.2242	0.7277	0.1060	0.7454	-0.1182	0.7040
CG17618	CG17618	1625880_at	-0.0274	0.9091	-0.2106	0.4550	-0.1521	0.4165	-0.1127	0.8350	0.1219	0.5833	0.2346	0.1993	-0.2618	0.6955	-0.0653	0.8642	0.1965	0.4865
iav	inactive	1625881_at	0.0493	0.7820	0.1425	0.5554	0.1532	0.4705	-0.0067	0.9953	-0.1752	0.4631	-0.1685	0.4332	0.0541	0.9467	0.0462	0.9009	-0.0079	0.9847
---	---	1625882_at	-0.0429	0.7844	0.1249	0.6088	0.0150	0.9292	0.0560	0.9110	0.0290	0.8931	-0.0270	0.8862	0.2444	0.6955	0.1923	0.4703	-0.0522	0.8849
CG13253	CG13253	1625883_at	0.1294	0.6097	-0.2390	0.1416	0.0241	0.9079	0.0061	0.9956	0.3132	0.2235	0.3071	0.1815	0.0245	0.9848	0.1857	0.5283	0.1612	0.5977
CG13059	CG13059	1625884_at	0.2146	0.3477	0.3926	0.0592	0.0675	0.8405	-0.0373	0.9704	-0.3061	0.2337	-0.2687	0.2445	0.2341	0.7536	-0.0377	0.9409	-0.2718	0.4011
betaTub97EF	beta tubulin	1625885_at	0.1086	0.5244	-0.1027	0.7673	0.2531	0.1085	0.3473	0.1899	0.1687	0.2972	-0.1785	0.2132	0.0215	0.9898	-0.0234	0.9682	-0.0449	0.9274
CG34028	CG34028	1625886_s_at	1.4999	0.0326	1.3259	0.4241	1.7343	0.0005	0.2479	0.5978	0.4251	0.0756	0.1772	0.4258	0.0219	0.9978	0.3039	0.9117	0.2820	0.9089
CG10732 /// DmirCG10732	CG10732	1625887_at	0.3111	0.5121	0.3358	0.6650	0.6884	0.0501	0.2435	0.7531	0.5447	0.1033	0.3012	0.3245	-0.0233	0.9952	0.5920	0.5488	0.6152	0.5361
gry	gryzun	1625888_a_at	-0.2540	0.3013	-0.2962	0.2371	-0.3571	0.0623	0.0231	0.9727	0.4570	0.0145	0.4339	0.0113	0.0985	0.9405	0.4720	0.2329	0.3735	0.3737
RpL3	Ribosomal protein	1625889_at	1.2018	0.0115	1.9317	0.0142	2.1921	0.0000	0.3217	0.5068	-0.1660	0.5479	-0.4876	0.0401	0.0858	0.9737	0.4945	0.4409	0.4087	0.5476
CG14479	CG14479	1625890_at	-0.0187	0.9311	0.0087	0.9451	0.0812	0.6384	0.1527	0.7168	0.1561	0.4317	0.0034	0.9887	-0.0198	0.9893	0.0260	0.9590	0.0458	0.9151
CG17111	CG17111	1625891_at	-1.3449	0.0124	-1.3537	0.0885	-1.2245	0.0005	0.2801	0.6936	0.0305	0.9509	-0.2496	0.4172	0.0927	0.9742	0.0079	0.9961	-0.0848	0.9323
CG17097	CG17097	1625892_a_at	0.0907	0.6812	0.1407	0.3690	0.1629	0.4644	0.0271	0.9649	0.1364	0.4188	0.1093	0.4835	0.0262	0.9853	0.0737	0.8752	0.0475	0.9168
Nep5	Neprilysin 5	1625893_a_at	-0.1985	0.2916	0.0000	1.0000	-0.0735	0.7153	-0.0903	0.8546	-0.0512	0.8171	0.0392	0.8495	0.0493	0.9503	0.2191	0.3397	0.1698	0.4854
CG11370	CG11370	1625894_at	0.2667	0.0870	0.0545	0.8071	0.1016	0.6769	0.1586	0.7906	0.0623	0.8437	-0.0963	0.7092	-0.0951	0.9112	-0.0553	0.9053	0.0398	0.9236
CG31211	CG31211	1625895_a_at	-0.8829	0.1279	-0.3849	0.6041	-0.3677	0.1150	-0.0535	0.9436	-0.2851	0.1836	-0.2316	0.2304	-0.0367	0.9916	0.0740	0.9557	0.1108	0.9215
---	---	1625896_at	0.0187	0.9232	0.1414	0.4402	0.2798	0.1130	-0.0945	0.8554	-0.2144	0.2542	-0.1198	0.5076	-0.2579	0.7149	-0.1336	0.6847	0.1244	0.7072
---	---	1625897_s_at	-3.6512	0.0012	-4.5576	0.0047	-4.9619	0.0000	-0.0829	0.8908	0.7342	0.0040	0.8171	0.0015	0.4307	0.8270	-0.2306	0.8307	-0.6612	0.4275
Lnk	Lnk	1625898_s_at	-0.2138	0.3488	0.2713	0.4547	0.4848	0.0204	-0.1612	0.7857	-0.8353	0.0053	-0.6741	0.0083	-0.2748	0.7779	-0.1688	0.7319	0.1060	0.8453
---	---	1625899_at	0.3167	0.1655	0.1075	0.5939	0.0392	0.8821	-0.0638	0.9247	0.0542	0.8344	0.1181	0.5541	0.2397	0.7899	0.0928	0.8625	-0.1469	0.7409
CG12729	CG12729	1625900_at	0.0786	0.6327	0.0365	0.7701	0.2628	0.1553	-0.0282	0.9627	-0.0623	0.7424	-0.0341	0.8550	-0.1304	0.8513	-0.0657	0.8730	0.0648	0.8646
---	---	1625901_s_at	-2.1945	0.1278	-1.8975	0.0417	-2.5440	0.0002	-0.6798	0.4529	-1.9252	0.0029	-1.2454	0.0122	-0.0856	0.9914	-1.7600	0.3261	-1.6743	0.3764
CG11655	CG11655	1625902_at	0.1457	0.6031	0.6483	0.1648	0.8384	0.0005	-0.1053	0.8054	-0.7951	0.0014	-0.6898	0.0015	-0.2737	0.8049	-0.3575	0.4379	-0.0838	0.8949
CG10101	CG10101	1625903_at	0.0156	0.9487	0.0713	0.6936	0.2489	0.1309	0.1297	0.7478	0.0425	0.8526	-0.0872	0.6262	-0.1041	0.9246	-0.0977	0.8444	0.0064	0.9924
CG3557	CG3557	1625904_at	0.1235	0.6399	0.0443	0.6625	-0.0035	0.9886	0.0038	0.9956	0.1187	0.4604	0.1149	0.4271	0.0749	0.9426	0.1241	0.7424	0.0492	0.9115
CG13926	CG13926	1625905_at	0.0779	0.8723	0.5203	0.1574	0.6576	0.0102	0.0471	0.9745	-0.3374	0.3629	-0.3845	0.2376	0.0213	0.9912	0.2402	0.5544	0.2189	0.6009
CG11105 /// DsimCG11105	CG11105	1625906_a_at	-0.7692	0.0086	-0.1565	0.5248	-0.3132	0.2415	-0.0929	0.9037	-0.8420	0.0055	-0.7491	0.0056	0.1550	0.8541	-0.2244	0.5336	-0.3794	0.2879
CG5591	CG5591	1625907_at	0.0770	0.9155	-0.3969	0.0628	-0.1163	0.5857	0.1392	0.8578	0.6597	0.0261	0.5204	0.0435	0.0477	0.9898	0.2856	0.7753	0.2379	0.8153
TppII	tripeptidyl peptida	1625908_a_at	0.4618	0.1378	0.3208	0.1009	0.1849	0.3373	0.0363	0.9553	0.5284	0.0091	0.4921	0.0076	0.1872	0.8554	0.4350	0.2877	0.2478	0.5811
---	---	1625909_at	0.1009	0.5141	0.0416	0.6763	0.0628	0.7146	-0.0972	0.8153	-0.0599	0.7593	0.0373	0.8433	-0.0531	0.9330	-0.0521	0.8501	0.0010	0.9979
CG1773	CG1773	1625910_at	1.9339	0.0023	0.6522	0.0881	1.7777	0.0001	0.2334	0.6533	0.2430	0.3398	0.0096	0.9757	-0.4154	0.7070	-0.7861	0.0965	-0.3707	0.4325
CG15760	CG15760	1625911_at	0.1767	0.3373	0.1248	0.2451	0.1278	0.4587	0.0576	0.9247	-0.0692	0.7490	-0.1268	0.4630	0.0146	0.9862	0.0231	0.9402	0.0085	0.9777
CG40290	CG40290	1625912_at	0.1035	0.5210	0.2766	0.1431	0.2011	0.3196	-0.0733	0.9168	-0.0625	0.8174	0.0108	0.9688	-0.0317	0.9816	0.0941	0.8083	0.1258	0.7114
CG9970	CG9970	1625913_at	0.1809	0.4762	-0.0061	0.9605	-0.0823	0.7006	-0.0867	0.8815	0.2047	0.3064	0.2913	0.0996	0.0626	0.9555	-0.0311	0.9528	-0.0938	0.8178
Or69a	Odorant receptor	1625914_at	-0.0348	0.8842	0.1094	0.4977	0.3435	0.0681	0.0186	0.9857	-0.1579	0.5325	-0.1765	0.4270	-0.0371	0.9637	0.0434	0.8960	0.0805	0.7526
CG4375	CG4375	1625915_at	0.0963	0.5110	0.0725	0.6177	-0.0313	0.8713	-0.1556	0.6818	-0.0945	0.6308	0.0611	0.7504	-0.0686	0.8960	-0.0627	0.8021	0.0059	0.9849
---	---	1625916_at	0.0051	0.9791	-0.1086	0.4262	-0.3399	0.1629	0.0611	0.8856	0.0810	0.6095	0.0199	0.9093	0.1320	0.8655	-0.1695	0.6111	-0.3014	0.3475
---	---	1625917_s_at	0.2217	0.1822	0.1700	0.3114	0.3019	0.1210	-0.0190	0.9852	-0.1227	0.6339	-0.1036	0.6652	-0.0207	0.9823	0.0723	0.7897	0.0930	0.7011
CG14346	CG14346	1625918_a_at	0.1587	0.6257	0.1545	0.2149	0.1283	0.4787	0.1178	0.8545	0.1363	0.5891	0.0185	0.9498	0.1058	0.9092	-0.0110	0.9869	-0.1168	0.7573

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG12936	CG12936	1625919_at	0.0602	0.7272	0.1888	0.5230	-0.0627	0.7890	-0.0612	0.9154	0.1161	0.5504	0.1772	0.2777	-0.0375	0.9816	0.0631	0.9144	0.1007	0.8334
---	---	1625920_at	0.2767	0.3060	-0.0396	0.8433	-0.0752	0.6769	0.1023	0.8405	0.2802	0.1385	0.1779	0.3019	0.0095	0.9950	-0.0620	0.9058	-0.0715	0.8774
---	---	1625921_at	0.2184	0.2305	0.1158	0.3718	-0.2263	0.1465	-0.1325	0.7618	0.0699	0.7553	0.2024	0.2291	0.2014	0.7120	0.1685	0.4660	-0.0329	0.9171
CG10508	CG10508	1625922_a_at	-1.6668	0.0014	-0.7984	0.0208	-1.0706	0.0025	-0.3195	0.5515	-0.6888	0.0213	-0.3694	0.1347	-0.0301	0.9913	0.0786	0.9278	0.1087	0.8862
Flo-2	flotillin 2	1625923_s_at	-1.0505	0.0034	-0.6625	0.1015	-0.8885	0.0069	-0.2781	0.5932	-0.1615	0.5711	0.1165	0.6717	-0.0370	0.9816	0.1836	0.6382	0.2205	0.5634
CG12661	CG12661	1625924_at	0.9491	0.0058	0.3323	0.2164	0.4753	0.0293	-0.1122	0.7409	-0.0166	0.9382	0.0956	0.5071	0.0691	0.9717	-0.4159	0.3856	-0.4851	0.3307
His2A:CG31618 /// His2A:C	His2A:CG31618 //	1625925_at	1.1213	0.3060	-1.2647	0.4299	-0.5344	0.5211	0.0953	0.9647	1.9908	0.0047	1.8955	0.0035	-0.5658	0.9390	-0.3493	0.9221	0.2165	0.9463
---	---	1625926_at	0.1185	0.6807	-0.1996	0.3168	-0.4939	0.0903	-0.2144	0.7658	0.1977	0.5495	0.4121	0.1315	0.0110	0.9939	-0.2822	0.3651	-0.2933	0.3678
CG3198	CG3198	1625927_at	-0.1191	0.5022	-0.5910	0.0595	-0.5670	0.0095	0.0267	0.9757	0.3935	0.0694	0.3667	0.0595	0.3166	0.6955	0.2175	0.5259	-0.0991	0.8071
HtrA2	HtrA2	1625928_at	0.2269	0.5947	0.3849	0.2011	1.0399	0.0005	0.1212	0.8544	-0.1382	0.5942	-0.2594	0.2240	-0.2596	0.6749	0.1213	0.6626	0.3809	0.1573
CG9059 /// DmirCG9059	CG9059	1625929_a_at	0.0297	0.9405	0.2430	0.1086	0.0064	0.9868	-0.1429	0.8403	-0.3172	0.2305	-0.1743	0.4922	0.3497	0.7506	0.2037	0.7007	-0.1461	0.7965
amn	cheap date	1625930_at	-0.0698	0.6324	-0.1383	0.4513	0.3157	0.0871	0.2516	0.3837	0.1754	0.2867	-0.0762	0.6518	-0.2259	0.7215	0.0686	0.8485	0.2945	0.2798
dbo	diablo	1625931_at	-0.0795	0.8827	-0.6618	0.0521	-1.1139	0.0004	-0.1561	0.7618	0.6335	0.0115	0.7896	0.0026	0.3565	0.8143	0.1979	0.8061	-0.1586	0.8435
Samuel	middleman of sev	1625932_at	0.1537	0.8389	1.0792	0.2246	0.3081	0.3833	-0.3741	0.3744	-0.3723	0.1164	0.0018	0.9953	0.4619	0.8906	0.6533	0.6324	0.1914	0.9121
CG14731	CG14731	1625933_at	0.0351	0.8993	0.0291	0.8593	0.0945	0.5994	-0.1039	0.8611	-0.0797	0.7540	0.0243	0.9257	-0.1234	0.8890	-0.1494	0.6889	-0.0260	0.9566
p	CG9770	1625934_at	0.7149	0.0482	0.5434	0.0302	0.7535	0.0013	0.2019	0.5735	0.3396	0.0671	0.1377	0.4224	0.0376	0.9812	0.2276	0.5187	0.1900	0.6040
Trim9	Trim9	1625935_at	0.9991	0.1314	-0.1415	0.4959	-0.0159	0.9797	-0.2412	0.8369	0.7893	0.0758	1.0305	0.0171	-0.3755	0.8553	-0.5326	0.5457	-0.1571	0.8921
---	---	1625936_at	0.1944	0.2035	0.1899	0.3636	-0.1245	0.4787	-0.1244	0.8182	-0.0261	0.9298	0.0983	0.6536	0.1396	0.8145	-0.0193	0.9621	-0.1589	0.5403
CG18646	CG18646	1625937_at	0.0915	0.6212	-0.1324	0.3347	0.0287	0.8710	0.0642	0.8882	0.1305	0.4095	0.0663	0.6800	-0.1141	0.8236	-0.0531	0.8569	0.0610	0.8187
---	---	1625938_s_at	-0.0517	0.8852	0.1545	0.3930	-0.0076	0.9688	0.0526	0.9351	0.1306	0.5181	0.0780	0.6965	0.2023	0.6955	0.3406	0.1199	0.1383	0.5460
---	---	1625939_at	0.0256	0.8986	0.0897	0.6166	0.1624	0.4294	-0.1789	0.6338	-0.2256	0.2204	-0.0467	0.8239	-0.1521	0.8461	-0.0359	0.9454	0.1161	0.7554
sim	singleminded	1625940_s_at	-2.9202	0.0008	-1.1288	0.1847	-3.1519	0.0002	-1.6237	0.0674	-2.4229	0.0013	-0.7991	0.0835	0.1496	0.9238	-0.6190	0.2181	-0.7686	0.1645
CG9782	CG9782	1625941_at	0.1449	0.4380	0.4160	0.0222	0.5000	0.0580	-0.0549	0.9376	-0.3241	0.1127	-0.2692	0.1404	-0.0551	0.9677	0.0794	0.8765	0.1345	0.7442
dia	diaphanous	1625942_s_at	-0.3944	0.2258	-0.1521	0.6550	-0.7266	0.0043	-0.2034	0.6494	0.3568	0.0987	0.5602	0.0110	0.3445	0.7925	0.5157	0.3393	0.1712	0.7982
RpS30	Ribosomal protein	1625943_at	0.6091	0.0966	0.3228	0.1247	-0.0254	0.9356	-0.0678	0.9186	0.2530	0.2147	0.3209	0.0806	0.3394	0.7464	0.0297	0.9678	-0.3097	0.5007
CG16838	CG16838	1625944_a_at	-0.0445	0.9090	0.0634	0.7814	-0.1463	0.5388	-0.1217	0.8518	0.4482	0.0622	0.5699	0.0152	0.0235	0.9914	0.4355	0.3456	0.4120	0.3921
CG1233	CG1233	1625945_a_at	0.3236	0.2982	0.4555	0.1471	0.3627	0.0881	-0.0913	0.8578	-0.1548	0.4119	-0.0635	0.7498	0.0337	0.9869	0.0215	0.9799	-0.0122	0.9873
---	---	1625946_at	-0.0457	0.7833	-0.1989	0.1073	0.2532	0.2447	0.2296	0.3766	0.1987	0.1740	-0.0309	0.8567	-0.3019	0.7230	-0.1903	0.6314	0.1116	0.8039
GLaz	Glial Lazarillo	1625947_at	0.8080	0.0310	-0.1751	0.2355	0.4850	0.0654	0.0497	0.9517	-0.2352	0.3030	-0.2849	0.1577	-0.7078	0.4809	-1.3400	0.0290	-0.6322	0.2199
CG17633	CG17633	1625948_at	0.6796	0.4586	-0.3560	0.1559	0.0346	0.8453	0.5026	0.1526	0.5494	0.0180	0.0468	0.8493	-0.1716	0.9677	-0.7701	0.5055	-0.5985	0.6195
Gpdh	5' gene	1625949_at	0.5656	0.0423	-0.3224	0.3094	0.1212	0.5764	0.5790	0.1054	0.9718	0.0015	0.3927	0.0442	0.1472	0.8480	0.1083	0.7862	-0.0389	0.9313
CG7777 /// DyakCG7777	CG7777	1625950_a_at	-0.3644	0.0940	0.1771	0.6085	-0.0791	0.6586	-0.4265	0.1327	-0.6373	0.0035	-0.2108	0.1656	-0.2527	0.8049	-0.1565	0.7587	0.0962	0.8644
CG17778	CG17778	1625951_at	-0.3665	0.1428	-0.0319	0.7531	0.1740	0.2584	0.2301	0.5617	-0.3688	0.0725	-0.5988	0.0061	-0.1330	0.8429	-0.1027	0.7544	0.0303	0.9372
---	---	1625952_at	-0.0519	0.7507	0.0250	0.8113	0.0080	0.9689	-0.1024	0.7939	-0.0531	0.7845	0.0493	0.7821	-0.0044	0.9959	0.0242	0.9402	0.0286	0.9169
CG13463	CG13463	1625953_at	-0.2069	0.2553	-0.2336	0.2072	-0.1120	0.4707	-0.0276	0.9704	-0.0347	0.8933	-0.0070	0.9766	0.0160	0.9878	-0.0100	0.9821	-0.0260	0.9371
CG2681	CG2681	1625954_at	-0.0840	0.8892	0.1368	0.2012	-0.1678	0.6387	-0.3291	0.7604	-0.3502	0.4699	-0.0211	0.9710	0.0008	0.9998	0.0207	0.9647	0.0199	0.9614
CG32000	CG32000	1625955_a_at	-0.5780	0.0633	-0.5652	0.1250	-1.3099	0.0011	-0.5906	0.2501	0.2430	0.4418	0.8336	0.0085	-0.0653	0.9689	0.0445	0.9504	0.1097	0.8461
CG32547	CG32547	1625956_at	-0.0909	0.5429	0.2014	0.3035	0.1798	0.4164	0.0662	0.9112	-0.1351	0.4932	-0.2013	0.2315	0.0954	0.9016	0.1103	0.7395	0.0149	0.9727
CG10051	CG10051	1625957_at	0.8039	0.1072	0.4614	0.2075	0.7841	0.0018	-0.0132	0.9866	0.1435	0.4744	0.1567	0.3773	-0.2849	0.8472	0.0951	0.9213	0.3801	0.5612
---	---	1625958_at	-0.1347	0.6605	-0.0150	0.9399	-0.0156	0.9558	-0.1008	0.9297	-0.1628	0.6687	-0.0620	0.8760	-0.0408	0.9695	-0.0263	0.9528	0.0145	0.9730
CG4372	CG4372	1625959_at	0.0795	0.6515	0.0290	0.7775	-0.0682	0.7260	0.0022	0.9974	0.0238	0.9289	0.0215	0.9258	-0.1402	0.8270	-0.0540	0.8884	0.0863	0.7836
noc	Scutoid	1625960_at	-0.8256	0.1785	-1.5283	0.0783	-1.9092	0.0033	0.2581	0.3604	0.5209	0.0067	0.2628	0.0702	0.6635	0.8192	-0.0520	0.9814	-0.7154	0.5698
---	---	1625961_at	0.1251	0.3933	0.0314	0.7541	0.1653	0.4103	-0.1241	0.6955	-0.1021	0.5093	0.0220	0.9004	-0.0244	0.9816	-0.0172	0.9666	0.0072	0.9856
RpII140	RNA polymerase	1625962_at	0.1885	0.3902	0.4865	0.2371	0.3255	0.0422	-0.1001	0.8076	0.1208	0.4815	0.2210	0.1299	0.0371	0.9848	0.3519	0.4317	0.3148	0.4980
CG1896	CG1896	1625963_at	0.2269	0.1279	0.3085	0.3055	0.1674	0.3914	-0.2260	0.5096	-0.2989	0.0994	-0.0728	0.7036	0.0402	0.9814	-0.1507	0.7200	-0.1910	0.6294
rec	recombination-def	1625964_at	-0.0050	0.9785	-0.1880	0.3173	-0.2358	0.4762	-0.0823	0.9502	0.1745	0.6748	0.2568	0.4572	-0.0997	0.8510	-0.1454	0.5228	-0.0458	0.8779
Osi2	Osiris	1625965_a_at	0.0133	0.9713	0.1536	0.4150	0.0626	0.8143	-0.0229	0.9824	-0.0764	0.7986	-0.0534	0.8502	0.1981	0.8270	0.1646	0.6969	-0.0335	0.9492
CG5500	CG5500	1625966_at	0.0814	0.6404	0.0271	0.9048	0.0750	0.6661	0.0747	0.8589	-0.2614	0.0851	-0.3361	0.0212	0.1730	0.8012	-0.0301	0.9487	-0.2031	0.4931
ps	pasilla	1625967_s_at	-0.2537	0.3565	-0.5436	0.0248	-0.7692	0.0150	0.0532	0.9441	0.2288	0.2954	0.1756	0.3818	0.3162	0.7953	-0.0415	0.9613	-0.3577	0.4980
CG11227 /// CG13630	CG13630 /// CG1	1625968_at	-0.2052	0.3609	-0.0688	0.7127	0.2635	0.2231	0.1824	0.6998	0.0503	0.8595	-0.1321	0.5332	-0.2577	0.7823	0.1614	0.7304	0.4191	0.3089
---	---	1625969_at	0.0288	0.8807	0.2160	0.2415	0.0791	0.6348	-0.2165	0.4860	-0.0802	0.6711	0.1363	0.3735	0.0432	0.9530	0.0719	0.7943	0.0286	0.9243
ex	expanded	1625970_at	0.8315	0.0360	1.7131	0.0076	1.1402	0.0016	-0.0857	0.9215	-0.1982	0.4695	-0.1125	0.6819	0.4865	0.6749	0.7574	0.1203	0.2710	0.6017
---	---	1625971_at	0.2449	0.4578	0.0323	0.7373	0.1252	0.4363	0.0623	0.8936	0.0744	0.6739	0.0120	0.9511	-0.0507	0.9717	-0.0871	0.8605	-0.0365	0.9422
CG31999	fibrillin-like	1625972_at	-2.1155	0.0035	-3.0905	0.0252	-3.7910	0.0000	-0.5249	0.0765	-0.4244	0.0215	0.1004	0.5507	0.1724	0.9643	-1.3866	0.1694	-1.5590	0.1577
CG3038 /// DsimCG3038	CG3038	1625973_a_at	-1.4870	0.0898	-2.1925	0.0097	-1.7468	0.0054	-0.0380	0.9847	0.4351	0.3418	0.4731	0.2409	-0.5496	0.8331	-0.4612	0.7078	0.0885	0.9542
ird5	LPS-responsive ki	1625974_at	-0.1652	0.4528	-0.4113	0.1040	-0.4286	0.0169	-0.1464	0.7278	0.0208	0.9385	0.1673	0.3314	-0.1358	0.8374	-0.2760	0.2976	-0.1401	0.6324
Tango12	Transport and Gol	1625975_at	3.1212	0.0088	0.9216	0.2799	1.9086	0.0002	0.2873	0.7349	0.4761	0.2048	0.1888	0.6227	-0.5341	0.8572	-1.7135	0.1489	-1.1794	0.3444
---	---	1625976_at	0.1033	0.4540	0.2511	0.2254	0.0214	0.9246	-0.2430	0.5633	-0.1989	0.3668	0.0442	0.8605	0.0880	0.8940	0.1835	0.4601	0.0955	0.7369
---	---	1625977_at	0.0149	0.9356	-0.1055	0.4071	0.0012	0.9961	0.0210	0.9726	0.0662	0.7061	0.0451	0.7908	-0.0903	0.8906	-0.0682	0.8376	0.0221	0.9492
CG15141	CG15141	1625978_at	0.0997	0.7020	0.2022	0.4457	0.1148	0.6600	-0.0132	0.9917	0.3834	0.1489	0.3966	0.0959	0.0892	0.9426	0.5310	0.1514	0.4419	0.2622
---	---	1625979_s_at	0.0764	0.7141	0.0788	0.5876	0.2274	0.3355	0.0249	0.9695	-0.0953	0.6038	-0.1202	0.4467	-0.1163	0.8331	0.0442	0.8988	0.1605	0.5007
CG11835	CG11835	1625980_at	-0.7137	0.0035	-0.6605	0.0573	-0.6570	0.0153	-0.0298	0.9610	-0.1056	0.5464	-0.0758	0.6522	0.1333	0.8953	-0.1446	0.7442	-0.2779	0.4764
rab3-GEF	rab3-GEF	1625981_at	-0.1669	0.5936	-0.1361	0.4953	-0.3151	0.1064	-0.2729	0.6578	-0.1413	0.6751	0.1316	0.6690	-0.2033	0.8262	-0.3014	0.4301	-0.0981	0.8411
SP2353	agrin-like	1625982_at	0.2448	0.3366	-0.0902	0.7165	-0.1340	0.4347	-0.0415	0.9518	0.1140	0.5812	0.1555	0.3773	-0.1519	0.8999	-0.3597	0.4144	-0.2078	0.6670
CG18666	CG18666	1625983_at	-0.0509	0.7799	0.0250	0.8207	0.2642	0.1202	0.1249	0.7082	-0.0243	0.9087	-0.1492	0.2785	-0.0845	0.8692	0.1296	0.5426	0.2141	0.3057
---	---	1625984_at	0.0202	0.9409	0.0265	0.7883	0.0218	0.9111	0.0028	0.9961	-0.1442	0.4499	-0.1471	0.3883	0.0783	0.9117	0.0191	0.9631	-0.0592	0.8555
CG31446	CG31446	1625985_at	0.0616	0.7486	0.0078	0.9702	-0.1858	0.3239	-0.1668	0.7735	0.0379	0.9107	0.2047	0.3603	-0.0654	0.9174	-0.0854	0.7381	-0.0200	0.9479
CG31826	CG31826	1625986_at	-0.8156	0.0108	0.0878	0.8389	-0.3699	0.2734	-0.5812	0.3381	-2.2971	0.0004	-1.7159	0.0006	0.0991	0.9421	-1.1724	0.0280	-1.2715	0.0333
Eig71Ee	Gene VII	1625987_at	0.2022	0.1936	0.0492	0.6895	0.1405	0.4526	-0.1969	0.6120	-0.0083	0.9781	0.1886	0.2834	-0.1582	0.8016	-0.1841	0.4954	-0.0259	0.9435

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Dys	Dystrophin-like pr	1625988_a_at	0.2640	0.7136	1.4277	0.0450	1.6309	0.0017	-0.3808	0.7304	-0.9594	0.0567	-0.5786	0.1896	-0.3872	0.8731	0.3933	0.7196	0.7806	0.4235
CG6144	CG6144	1625989_at	-0.4331	0.0379	-0.0397	0.6827	-0.0163	0.9277	0.2184	0.5067	-0.0937	0.6288	-0.3121	0.0499	0.1425	0.7953	0.2333	0.3000	0.0909	0.7312
CG11367 /// CG32454 /// D	CG11367 /// CG3	1625990_at	0.9949	0.0034	0.9520	0.0164	1.1807	0.0001	0.0745	0.8807	0.4733	0.0128	0.3988	0.0165	-0.0715	0.9449	0.5078	0.1108	0.5793	0.1003
RpL28	Ribosomal protein	1625991_s_at	0.2525	0.1216	0.7020	0.0213	0.7628	0.0044	0.0488	0.9436	-0.4348	0.0348	-0.4835	0.0141	0.0378	0.9683	-0.0446	0.9075	-0.0825	0.7815
CG34413	CG32836	1625992_s_at	-0.0464	0.8994	0.8632	0.0902	1.0753	0.0014	-0.2883	0.4022	-0.7718	0.0026	-0.4835	0.0126	-0.2569	0.8534	0.2704	0.6686	0.5272	0.3701
---	---	1625993_at	0.1518	0.4276	-0.0328	0.7433	0.2313	0.1620	-0.0191	0.9803	-0.0800	0.7114	-0.0609	0.7675	-0.3287	0.3517	-0.2147	0.2389	0.1140	0.5684
mRpS11	mitochondrial ribo	1625994_a_at	-0.5685	0.0460	-0.2899	0.3046	-0.4425	0.0430	0.0229	0.9777	-0.2828	0.1671	-0.3057	0.0963	0.0457	0.9816	-0.0569	0.9347	-0.1027	0.8554
Zw	glucose-6-phosph	1625995_a_at	0.2871	0.4492	0.8615	0.0768	1.6728	0.0002	0.5462	0.2941	-0.3476	0.2578	-0.8938	0.0063	-0.3272	0.7682	0.0774	0.9157	0.4046	0.3986
BG642312	putative noncodin	1625996_s_at	0.0344	0.9250	0.0127	0.9377	-0.1664	0.3854	-0.0187	0.9838	-0.2910	0.1759	-0.2722	0.1577	0.1731	0.8461	-0.3388	0.3414	-0.5119	0.1867
---	---	1625997_s_at	-0.4170	0.0680	-0.5594	0.3055	-1.0201	0.0019	-0.2370	0.6202	0.2965	0.2121	0.5334	0.0205	0.1958	0.8874	0.1984	0.7440	0.0026	0.9979
---	---	1625998_s_at	0.3555	0.2020	0.1127	0.4756	0.2371	0.2497	0.0556	0.9314	0.0752	0.7396	0.0195	0.9344	-0.1706	0.8692	-0.0792	0.8997	0.0913	0.8693
CG2919	CG2919	1625999_at	-0.1262	0.7600	-0.7468	0.0312	-1.1085	0.0012	-0.2529	0.4821	0.5599	0.0107	0.8128	0.0013	-0.1156	0.9409	-0.0642	0.9341	0.0513	0.9394
CG6508	CG6508	1626000_at	0.0908	0.6041	0.2354	0.3218	-0.0365	0.8234	-0.2512	0.5376	-0.2332	0.2759	0.0181	0.9467	-0.0242	0.9775	0.0076	0.9842	0.0318	0.9095
Nrg	neuroglian	1626001_at	-0.5011	0.0144	-0.4498	0.2024	-0.4616	0.0265	0.1547	0.6558	0.2493	0.1339	0.0946	0.5674	-0.0177	0.9914	0.2299	0.5679	0.2477	0.5403
Roe1	Roe1	1626002_at	-0.0099	0.9578	0.2355	0.3659	0.0995	0.4949	-0.0950	0.8507	-0.1681	0.3701	-0.0731	0.7065	0.0401	0.9683	0.1595	0.5679	0.1194	0.6850
CG30461 /// CG9002	CG9002 /// CG30	1626003_s_at	0.1260	0.4042	0.0325	0.7938	0.0931	0.6695	0.0699	0.8815	0.0505	0.7965	-0.0194	0.9213	0.0849	0.9260	0.0369	0.9402	-0.0479	0.9092
---	---	1626004_at	-0.0395	0.8623	0.1037	0.5943	0.0481	0.7872	-0.1064	0.8034	-0.0859	0.6564	0.0205	0.9221	-0.1261	0.8427	0.0905	0.7784	0.2166	0.4148
CG40188	CG40188	1626005_at	0.0403	0.8231	-0.1168	0.4836	-0.1236	0.4872	0.0129	0.9917	0.0799	0.8026	0.0670	0.8207	0.0046	0.9964	-0.0511	0.8749	-0.0557	0.8481
Dsor1	Downstream of ra	1626006_at	-0.0169	0.9280	-0.2363	0.1310	0.1023	0.5344	0.1398	0.6712	0.2363	0.1288	0.0965	0.5250	-0.0969	0.8841	0.0009	0.9994	0.0979	0.7355
CG32650	CG32650	1626007_at	-0.0676	0.8333	0.0235	0.8699	0.0523	0.8287	-0.0111	0.9883	-0.0985	0.6196	-0.0875	0.6331	-0.0038	0.9974	-0.1327	0.6497	-0.1288	0.6579
---	---	1626008_at	-0.0934	0.5258	-0.0703	0.5290	-0.1366	0.4898	-0.1697	0.6338	-0.0305	0.8985	0.1392	0.3919	0.0238	0.9831	0.0434	0.9144	0.0197	0.9585
CG30427	CG30427	1626009_s_at	0.3480	0.0499	0.1774	0.3179	0.2044	0.4347	-0.1203	0.8628	0.0923	0.7550	0.2126	0.3501	0.0516	0.9589	-0.0307	0.9460	-0.0823	0.8123
CG4960	CG4960	1626010_at	0.4367	0.0869	0.1276	0.4201	0.4363	0.0235	-0.0163	0.9854	0.1235	0.5707	0.1398	0.4645	-0.0635	0.9462	0.0137	0.9787	0.0772	0.8299
CG17834	CG17834	1626011_at	-0.3411	0.0725	-0.3348	0.3822	-0.9687	0.0007	-0.3845	0.2172	0.1398	0.4747	0.5243	0.0077	0.4015	0.7018	0.2196	0.6447	-0.1818	0.7135
CG32846	CG32846	1626012_at	0.2409	0.3467	0.3049	0.0705	0.0971	0.7039	-0.1006	0.9136	-0.0775	0.8364	0.0230	0.9503	0.0210	0.9816	-0.0545	0.8481	-0.0754	0.7536
CG11192	CG11192	1626013_at	0.3332	0.6725	-0.0268	0.9146	-0.6010	0.0393	-0.3010	0.6703	0.1515	0.6946	0.4525	0.1298	0.1801	0.9588	-0.4420	0.6827	-0.6221	0.5438
---	---	1626014_at	0.0962	0.5146	0.0289	0.8634	0.2838	0.0950	0.0955	0.8816	-0.0595	0.8307	-0.1551	0.4528	-0.0986	0.8395	-0.0640	0.8031	0.0347	0.9006
CG32776	CG32776	1626015_s_at	-0.0688	0.6622	-0.0031	0.9823	0.0821	0.5780	0.1244	0.6888	0.0197	0.9220	-0.1047	0.4397	0.0074	0.9928	-0.0010	0.9991	-0.0084	0.9791
tmod	Tropomodulin	1626016_s_at	-1.8396	0.0028	-1.0698	0.1069	-1.3804	0.0001	-0.0735	0.9255	-0.6507	0.0138	-0.5772	0.0144	0.3846	0.7999	0.2532	0.7300	-0.1315	0.8782
---	---	1626017_at	0.0865	0.6682	0.0290	0.8274	-0.0938	0.6022	-0.0765	0.8836	0.0724	0.7279	0.1489	0.3637	0.0525	0.9555	0.0664	0.8626	0.0139	0.9746
crol	crooked legs	1626018_a_at	-0.7778	0.3094	-0.0970	0.7883	-0.8684	0.0154	-0.1551	0.7918	-0.1314	0.6200	0.0238	0.9362	0.5631	0.8283	0.5293	0.6576	-0.0339	0.9849
---	---	1626019_at	-0.0777	0.7235	0.0528	0.8347	-0.0252	0.8923	0.0065	0.9943	-0.0476	0.8463	-0.0541	0.7991	0.0708	0.9193	0.1320	0.6032	0.0611	0.8407
---	---	1626020_at	0.0126	0.9442	0.2495	0.2554	0.1362	0.4064	0.2379	0.5680	0.1699	0.4435	-0.0681	0.7728	0.1538	0.8202	0.1786	0.5404	0.0248	0.9494
CG32055	CG32055	1626021_at	0.1884	0.4342	-0.0134	0.9002	0.3356	0.0842	0.2568	0.6010	0.1879	0.4675	-0.0689	0.8046	-0.1367	0.8680	-0.0167	0.9769	0.1200	0.7492
Cyp12e1	Cyp12e1	1626022_at	-0.2134	0.5598	-0.4433	0.2200	-0.7424	0.0047	-0.7898	0.1683	-1.3790	0.0028	-0.5893	0.0623	-0.7126	0.3587	-1.7667	0.0092	-1.0541	0.0457
CG14932	CG14932	1626023_at	0.3961	0.0883	0.1377	0.5039	-0.0243	0.8972	-0.1252	0.8611	-0.0867	0.7801	0.0385	0.9006	0.1884	0.6483	-0.0786	0.6686	-0.2670	0.1389
CG14329	CG14329	1626024_at	-0.0307	0.8452	0.0784	0.4499	-0.0437	0.8764	-0.1112	0.7969	-0.2761	0.1145	-0.1648	0.3003	0.1013	0.9006	-0.0181	0.9708	-0.1194	0.7243
---	---	1626025_at	0.1830	0.5309	0.2003	0.3859	0.0888	0.6772	-0.0649	0.9345	-0.1617	0.5098	-0.0968	0.6914	0.0286	0.9869	0.0024	0.9989	-0.0263	0.9646
CG5342	CG5342	1626026_at	-0.0283	0.8979	0.0237	0.9533	-0.0988	0.6599	-0.0315	0.9819	-0.0088	0.9859	0.0228	0.9568	0.1172	0.8650	0.1322	0.6605	0.0150	0.9717
Sr-CII	Scavenger Recep	1626027_at	-0.0182	0.9441	-0.0153	0.9326	-0.0126	0.9490	0.0911	0.8462	0.0943	0.6225	0.0032	0.9880	-0.0354	0.9663	-0.0764	0.7824	-0.0410	0.8935
CG4783	CG4783	1626028_at	0.1777	0.7221	1.4822	0.0564	1.2152	0.0443	-0.3998	0.8038	-1.0909	0.0911	-0.6910	0.2307	-0.1334	0.9611	0.1945	0.8539	0.3278	0.7027
CG9094	CG9094	1626029_at	-0.0378	0.8249	0.1527	0.4378	0.1119	0.5368	-0.0176	0.9838	-0.2549	0.1976	-0.2372	0.1804	0.0282	0.9829	0.0449	0.9248	0.0167	0.9715
---	---	1626030_at	0.1475	0.4379	0.0374	0.8050	0.0375	0.8423	0.0681	0.8883	-0.0729	0.6982	-0.1409	0.3464	0.3116	0.6695	0.0647	0.8782	-0.2469	0.4093
CG12539	CG12539	1626031_at	-0.3102	0.0642	0.2553	0.1926	0.1748	0.3825	-0.0982	0.8211	-0.4414	0.0174	-0.3433	0.0311	-0.0417	0.9689	0.1912	0.5100	0.2329	0.4142
schuy	schumacher-levy	1626032_at	0.1339	0.4720	-0.0476	0.7722	0.0545	0.7340	0.0872	0.8746	0.1280	0.5335	0.0408	0.8557	0.0590	0.9365	0.0069	0.9889	-0.0521	0.8674
Atg9	Autophagy-specifi	1626033_at	0.1949	0.4069	0.3186	0.2669	0.3994	0.1350	-0.1423	0.6699	-0.5456	0.0051	-0.4034	0.0120	-0.2532	0.8270	-0.4146	0.3802	-0.1614	0.7764
---	---	1626034_at	-0.0119	0.9629	0.0614	0.7992	0.0006	0.9984	-0.0200	0.9767	-0.0032	0.9899	0.0168	0.9362	0.0633	0.9589	0.0816	0.8680	0.0183	0.9730
---	---	1626035_x_at	-0.2095	0.2944	-0.1054	0.4099	-0.0305	0.8991	0.0334	0.9665	-0.0547	0.8383	-0.0882	0.6868	-0.0784	0.9142	-0.0328	0.9361	0.0456	0.8969
---	---	1626036_at	0.0476	0.8043	-0.0319	0.7927	0.0036	0.9919	0.1951	0.7760	0.2660	0.3637	0.0710	0.8284	-0.0225	0.9831	0.1500	0.5685	0.1725	0.5069
CG14380	CG14380	1626037_at	-0.0273	0.9293	0.1563	0.2723	0.1476	0.3139	-0.1315	0.8385	-0.0089	0.9806	0.1225	0.6176	-0.2617	0.5126	0.0571	0.8187	0.3189	0.1289
frc	UDP-sugar transp	1626038_at	-0.4556	0.1815	-0.7592	0.0387	-0.7510	0.0032	-0.2672	0.4279	-0.3262	0.0806	-0.0590	0.7742	-0.2833	0.8059	-0.6381	0.1718	-0.3548	0.4758
DyakCG4413 /// trem	CG4413	1626039_at	0.2926	0.3815	0.3883	0.0713	0.6473	0.0120	-0.0694	0.9282	0.1784	0.4515	0.2478	0.2245	-0.2254	0.8016	0.2255	0.5654	0.4509	0.2454
---	---	1626040_at	0.1323	0.5045	0.3852	0.0441	-0.0276	0.9207	-0.2021	0.5213	-0.2106	0.2014	-0.0084	0.9679	0.0408	0.9746	-0.0759	0.8660	-0.1167	0.7516
CG33453	CG33453	1626041_at	-0.2562	0.3193	0.1650	0.5869	0.1274	0.4758	-0.0938	0.8293	-0.4507	0.0149	-0.3569	0.0249	-0.1241	0.8608	-0.0625	0.8781	0.0616	0.8717
---	---	1626042_at	-0.0393	0.8779	0.0537	0.7717	0.0998	0.6951	-0.0694	0.9120	-0.1252	0.5608	-0.0558	0.8039	-0.0649	0.9514	0.0742	0.8685	0.1391	0.6949
---	---	1626043_at	-0.0574	0.7184	-0.0237	0.8702	-0.0302	0.8950	0.0421	0.9441	0.0257	0.9147	-0.0164	0.9391	-0.0239	0.9775	-0.0659	0.7951	-0.0420	0.8779
CG7628	CG7628	1626044_at	-0.8600	0.4347	-2.6893	0.0083	-2.0272	0.0068	0.5148	0.5913	1.4019	0.0111	0.8871	0.0467	-0.3321	0.9400	-0.2228	0.9157	0.1093	0.9548
Rpb4	Rpb4	1626045_at	-0.0087	0.9732	0.3377	0.1374	0.2174	0.1668	-0.1939	0.5651	-0.2529	0.1422	-0.0590	0.7549	-0.0345	0.9717	0.1858	0.4521	0.2203	0.3825
CG5585 /// CG6434	CG5585 /// CG64	1626046_s_at	0.0661	0.7849	0.6716	0.1499	1.2622	0.0043	-0.0611	0.9376	-0.4020	0.0796	-0.3409	0.0943	-0.3191	0.7677	0.2456	0.6165	0.5648	0.2336
Ssl2	Strictosidine synth	1626047_at	0.2079	0.6187	0.0801	0.7055	-0.3027	0.2985	-0.2024	0.8293	0.0326	0.9517	0.2350	0.4973	0.0822	0.9515	-0.0715	0.9085	-0.1538	0.7439
HLHmgamma	split locus enhanc	1626048_at	0.1017	0.8776	0.0261	0.7968	0.0651	0.6940	-0.1597	0.9220	-0.2305	0.6813	-0.0707	0.9051	-0.1381	0.8666	-0.2333	0.4856	-0.0952	0.8154
CG13089 /// DyakCG13089	CG13089	1626049_at	-0.3044	0.2748	-0.1999	0.5460	-0.0361	0.8676	-0.0778	0.8894	0.0254	0.9226	0.1032	0.5804	-0.2141	0.8480	0.1907	0.7200	0.4048	0.3921
oa2	octopamine recep	1626050_at	0.2047	0.1881	0.0821	0.6706	0.0681	0.7297	0.0769	0.8671	0.1175	0.4881	0.0407	0.8252	0.1522	0.8425	0.0680	0.8801	-0.0842	0.8310
CG13339 /// DyakCG13339	CG13339	1626051_at	-0.0281	0.9008	-0.0386	0.8823	-0.1841	0.5284	-0.1460	0.7488	-0.3223	0.1038	-0.1762	0.3314	0.0785	0.9503	-0.3469	0.3416	-0.4254	0.2729
MP1	Melanization Prot	1626052_at	1.1885	0.0050	1.3653	0.0095	1.5569	0.0004	-0.1787	0.7235	-0.6662	0.0105	-0.4875	0.0248	-0.2568	0.8016	-0.5058	0.2198	-0.2490	0.5838
robo3	robo3	1626053_at	0.0739	0.7870	0.0113	0.9383	0.0678	0.7751	0.0950	0.8914	0.0055	0.9879	-0.0895	0.7189	0.0956	0.9137	0.0118	0.9841	-0.0838	0.8289
shakB	shaking B	1626054_at	0.3404	0.1818	0.0627	0.8500	0.0476	0.8361	-0.1334	0.8350	-0.1362	0.6108	-0.0027	0.9928	-0.0036	0.9990	-0.2816	0.5384	-0.2780	0.5488
sphinx1	CG32383	1626055_at	-0.0800	0.7019	-0.2471	0.3205	-0.1975	0.3330	0.1073	0.8571	0.3398	0.1106	0.2325	0.2241	-0.2645	0.6955	-0.0832	0.8216	0.1812	0.5433
eco	Drosophila eco1	1626056_a_at	-0.0591	0.7313	0.0366	0.8651	-0.0756	0.6177	-0.0044	0.9956	0.1887	0.2531	0.1930	0.1890	0.1573	0.8157	0.4074	0.1395	0.2501	0.3812

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9368	CG9368	1626057_a_at	-0.0967	0.5452	0.0734	0.7317	0.0556	0.7841	-0.2224	0.3744	-0.2681	0.0620	-0.0457	0.7698	-0.1548	0.8023	-0.1611	0.5501	-0.0063	0.9875
lbl	ladybird	1626058_at	3.0321	0.0009	3.3695	0.0048	3.8058	0.0002	0.6493	0.5556	-0.2282	0.7322	-0.8774	0.0852	0.3331	0.7387	0.2626	0.5531	-0.0704	0.9045
knrl	knirps-related	1626059_at	-4.8710	0.0004	-5.3592	0.0007	-5.2909	0.0000	0.1545	0.9056	0.2058	0.6668	0.0513	0.9209	-0.0550	0.9774	-0.5762	0.1925	-0.5213	0.2697
---	---	1626060_at	0.2777	0.1511	-0.0193	0.8600	-0.0965	0.6027	0.0035	0.9956	0.1662	0.3712	0.1627	0.3278	0.0411	0.9742	-0.0092	0.9894	-0.0504	0.9089
CG34123	CG34123	1626061_at	0.4215	0.3282	-0.2476	0.5745	0.1774	0.3958	-0.0936	0.9201	0.5789	0.0475	0.6725	0.0164	-0.6538	0.7070	-0.2580	0.7709	0.3958	0.6140
CG34353	CG12274	1626062_at	-0.0951	0.7256	-0.1973	0.3772	-0.0791	0.7194	0.1390	0.8084	0.0595	0.8395	-0.0795	0.7478	-0.1350	0.8968	-0.1014	0.8478	0.0336	0.9510
---	---	1626063_at	0.3024	0.0744	0.4926	0.1238	0.4640	0.0102	-0.0565	0.9011	-0.1110	0.4712	-0.0545	0.7295	0.0206	0.9848	0.0129	0.9763	-0.0077	0.9849
CG4648	CG4648	1626064_at	0.1149	0.4838	0.0909	0.4209	0.3925	0.0400	-0.0730	0.8732	0.1199	0.4694	0.1928	0.1728	-0.3734	0.3712	0.1023	0.6894	0.4757	0.0741
CG12564	CG12564	1626065_at	0.0935	0.5470	0.0163	0.8731	0.0039	0.9898	0.1828	0.7203	0.1937	0.4162	0.0109	0.9700	0.0866	0.9174	0.0418	0.9275	-0.0448	0.9107
CG18748	CR18748	1626066_at	0.1779	0.3203	0.2035	0.3445	0.3125	0.1898	0.0572	0.9449	-0.0092	0.9800	-0.0664	0.8030	-0.1688	0.8193	0.0051	0.9941	0.1739	0.5917
CG7311	CG7311	1626067_a_at	0.1669	0.4736	0.1489	0.6135	0.2512	0.3187	0.0276	0.9715	0.0196	0.9459	-0.0080	0.9738	-0.0592	0.9737	0.1185	0.8432	0.1777	0.7243
lectin-46Cb	lectin-46Cb	1626068_at	-0.2686	0.2917	0.2285	0.1088	0.1752	0.3607	-0.0802	0.9311	-0.3540	0.1889	-0.2738	0.2616	0.1344	0.8386	0.3626	0.1694	0.2282	0.4093
---	---	1626069_at	0.0969	0.6015	0.0247	0.8444	-0.0587	0.7872	-0.2380	0.6650	0.1842	0.5060	0.4223	0.0736	-0.0580	0.9474	-0.0076	0.9898	0.0504	0.8915
---	---	1626070_at	0.0262	0.8973	0.0309	0.8895	0.2097	0.1979	0.1263	0.7336	0.1167	0.5037	-0.0096	0.9632	0.0207	0.9852	-0.0600	0.8680	-0.0807	0.7936
---	---	1626071_at	0.1365	0.4329	0.2008	0.3526	0.1660	0.3174	-0.1385	0.7929	-0.1381	0.5504	0.0005	0.9988	0.0839	0.8692	0.0899	0.6954	0.0059	0.9852
Unc-89	PROJECTIN-like	1626072_at	0.2362	0.3492	0.1014	0.5698	-0.0538	0.8024	-0.0650	0.9300	0.0338	0.9111	0.0988	0.6613	0.0429	0.9467	-0.1104	0.5936	-0.1533	0.4449
mud	mushroom body d	1626073_a_at	0.1967	0.1746	-0.0028	0.9825	-0.0121	0.9629	0.0737	0.8716	0.1014	0.5539	0.0277	0.8832	0.0274	0.9773	-0.0743	0.7932	-0.1016	0.6839
CG34018	CG34018	1626074_at	2.0313	0.0363	0.9570	0.2594	2.0116	0.0028	0.5313	0.5046	0.3126	0.4795	-0.2187	0.6083	-0.2726	0.9477	-0.4080	0.7985	-0.1355	0.9387
CG11664	CG11664	1626075_at	0.1952	0.3911	-0.1836	0.4953	-0.0146	0.9673	0.0837	0.9422	0.2369	0.4918	0.1533	0.6519	-0.1073	0.9246	-0.1803	0.6658	-0.0729	0.8885
CG8149	CG8149	1626076_at	0.3513	0.0668	0.0533	0.8525	-0.2688	0.2683	-0.0456	0.9471	0.6672	0.0051	0.7128	0.0023	0.1197	0.8940	0.2121	0.5414	0.0925	0.8247
CG11870	CG11870	1626077_s_at	-1.8034	0.0049	-1.3906	0.1305	-1.7208	0.0027	0.2253	0.5283	0.0263	0.9201	-0.1990	0.2341	0.5858	0.8243	0.5207	0.6676	-0.0651	0.9694
---	---	1626078_at	0.1922	0.2607	0.2915	0.3409	0.3098	0.0833	-0.0585	0.9477	-0.0144	0.9705	0.0441	0.8818	-0.1340	0.8643	0.1222	0.7393	0.2562	0.4235
CG2201	CG2201	1626079_a_at	-1.3248	0.0007	0.2781	0.4543	-0.5453	0.1225	-0.1950	0.6368	-0.4424	0.0359	-0.2474	0.1714	0.6359	0.6749	1.2595	0.0655	0.6236	0.3323
---	---	1626080_at	0.8374	0.1353	0.0878	0.5726	0.2006	0.4420	0.5633	0.1883	1.1029	0.0020	0.5396	0.0276	0.1964	0.9238	0.0741	0.9462	-0.1223	0.8951
CG8837	CG8837	1626081_at	-0.7034	0.1097	0.0933	0.6897	-0.4396	0.0573	-0.3775	0.5125	-0.7806	0.0185	-0.4031	0.1368	0.2810	0.7685	0.0266	0.9682	-0.2544	0.5518
CG11882	CG11882	1626082_at	0.2564	0.2467	-0.0400	0.7288	0.2459	0.2774	0.3854	0.1601	0.1472	0.3875	-0.2382	0.1081	0.0553	0.9742	-0.1060	0.8557	-0.1612	0.7424
CG14137	CG14137	1626083_at	0.2111	0.3891	0.1174	0.4467	-0.2739	0.2907	-0.1732	0.8107	-0.1421	0.6621	0.0310	0.9295	0.1645	0.8330	-0.2119	0.5283	-0.3764	0.2614
---	---	1626084_at	-0.1490	0.5469	0.0725	0.4857	-0.0188	0.9098	-0.2161	0.4279	-0.3090	0.0454	-0.0929	0.5250	0.0106	0.9914	-0.0236	0.9485	-0.0342	0.9118
---	---	1626085_at	0.1592	0.3560	0.0794	0.6190	0.3923	0.1459	-0.0400	0.9517	-0.0737	0.7290	-0.0337	0.8759	-0.2615	0.7633	-0.0913	0.8550	0.1702	0.6672
CG10621	selenocysteine me	1626086_at	2.3899	0.0108	0.9568	0.3147	2.5647	0.0001	0.6479	0.6062	0.4597	0.4876	-0.1882	0.7900	-1.0540	0.6557	-0.8557	0.3669	0.1983	0.8800
stan	flamingo	1626087_at	-0.1458	0.8600	-0.1003	0.4843	-0.0815	0.7038	0.0042	0.9988	-0.3193	0.6377	-0.3234	0.5935	-0.1986	0.8454	-0.2747	0.5259	-0.0761	0.8947
CG2177 /// DyakCG2177	CG2177	1626088_at	0.5012	0.0110	0.9652	0.0948	0.8077	0.0017	-0.2799	0.2753	-0.4920	0.0065	-0.2120	0.1156	-0.0001	0.9999	0.0026	0.9991	0.0027	0.9979
CG32234	CG32234	1626089_at	-1.8793	0.0067	0.1196	0.3862	-0.7630	0.0115	-0.5725	0.4568	-1.7392	0.0022	-1.1668	0.0077	0.2139	0.7677	0.1786	0.5836	-0.0354	0.9344
CG3973 /// DsimCG3973	CG3973	1626090_at	-0.4174	0.4886	-1.6017	0.0234	-2.1565	0.0009	0.1032	0.9254	0.9893	0.0095	0.8862	0.0094	0.3580	0.8814	-0.1691	0.9037	-0.5271	0.5887
---	---	1626091_at	0.4134	0.2650	0.2431	0.3506	0.3363	0.2200	0.0360	0.9643	0.2606	0.2204	0.2247	0.2400	-0.0429	0.9666	0.1332	0.6544	0.1761	0.5403
CG32690	CG32690	1626092_at	-0.0298	0.8705	-0.0131	0.9524	0.0187	0.9369	0.0504	0.9345	-0.1005	0.6131	-0.1508	0.3634	0.0875	0.9063	0.1484	0.6012	0.0609	0.8618
Jon65Aiii	Jonah 65A	1626093_at	1.6609	0.4143	-0.6082	0.1577	-0.0243	0.9447	0.5525	0.4861	0.6879	0.1039	0.1353	0.7725	-0.2541	0.9794	-1.6161	0.4812	-1.3620	0.5676
CG1941	CG1941	1626094_at	-0.0615	0.9297	1.5841	0.0755	0.7469	0.1078	-0.8865	0.3996	-1.6043	0.0144	-0.7178	0.1724	-0.0408	0.9916	0.1024	0.9451	0.1432	0.9095
CG18004	CG18004	1626095_at	0.1129	0.4751	-0.0416	0.8259	0.0104	0.9572	0.1229	0.7018	0.0427	0.8147	-0.0801	0.5864	-0.0686	0.9342	-0.1433	0.6157	-0.0746	0.8237
CG5220 /// DyakCG5220	CG5220	1626096_at	0.8303	0.0166	0.3940	0.4126	0.2605	0.1503	0.1210	0.8389	0.6192	0.0132	0.4982	0.0207	0.1843	0.9071	0.2025	0.7668	0.0182	0.9847
CG14754	CG14754	1626097_at	0.0653	0.7504	-0.0399	0.6912	0.0474	0.8312	0.0684	0.9259	0.0739	0.7832	0.0055	0.9847	0.0375	0.9552	0.0091	0.9796	-0.0284	0.9162
CG11453	CG11453	1626098_at	-0.1535	0.8267	-0.0256	0.7940	-0.2841	0.0876	-0.1075	0.9603	-0.1740	0.8031	-0.0665	0.9240	0.0466	0.9816	-0.2704	0.5929	-0.3170	0.5266
fray	Ste20-like protein	1626099_s_at	0.1501	0.5208	-0.3723	0.1162	-0.6848	0.0123	0.0073	0.9956	0.7333	0.0145	0.7260	0.0092	0.2687	0.7187	0.1779	0.5878	-0.0907	0.8137
CG16723	CG16723	1626100_at	0.1328	0.5596	-0.0733	0.6752	0.0604	0.7263	0.0106	0.9937	0.0153	0.9705	0.0048	0.9886	-0.0788	0.8756	-0.0359	0.9054	0.0429	0.8708
Scamp	anon-fast-evolving	1626101_at	0.2917	0.1577	0.1169	0.3955	0.0371	0.9049	0.0292	0.9687	-0.0025	0.9928	-0.0317	0.8907	0.1386	0.9246	-0.0615	0.9387	-0.2001	0.7189
---	---	1626102_at	0.0283	0.8618	0.0062	0.9564	0.1168	0.6139	-0.1200	0.7149	-0.0693	0.6865	0.0507	0.7568	-0.0926	0.8736	0.0270	0.9411	0.1196	0.6279
Cp38	Shell-38	1626103_at	0.3506	0.1103	0.4241	0.0780	-0.0079	0.9770	-0.0863	0.8640	-0.1642	0.3696	-0.0779	0.6776	0.3102	0.6749	-0.0688	0.8760	-0.3789	0.2336
---	---	1626104_at	-0.0911	0.5511	0.2150	0.5360	0.2182	0.1897	0.0082	0.9937	-0.1718	0.4191	-0.1800	0.3399	-0.0739	0.8940	0.0781	0.7504	0.1520	0.4795
---	---	1626105_at	0.2267	0.1267	0.1953	0.1400	0.2793	0.2775	0.0614	0.9314	0.0193	0.9517	-0.0421	0.8656	-0.0248	0.9855	0.0242	0.9624	0.0489	0.9094
CG7139 /// DmirCG7139 ///	CG7139	1626106_a_at	-0.0517	0.8027	0.0143	0.9022	-0.0473	0.8343	-0.0230	0.9777	-0.0567	0.8342	-0.0337	0.8948	-0.0599	0.9467	0.0071	0.9911	0.0671	0.8500
---	---	1626107_at	0.0590	0.7223	-0.0852	0.4984	0.0040	0.9881	-0.0291	0.9744	-0.0766	0.7689	-0.0475	0.8502	-0.0830	0.9032	-0.1764	0.4879	-0.0934	0.7445
CG15177	CG15177	1626108_at	0.1558	0.4578	0.0999	0.5046	0.1697	0.3796	-0.0423	0.9497	0.0131	0.9634	0.0553	0.7911	-0.1063	0.8814	-0.1002	0.7576	0.0060	0.9890
Drip	aquaporin	1626109_a_at	-1.1546	0.0068	-0.5316	0.4567	-0.5217	0.0251	-0.4267	0.3744	-1.0553	0.0030	-0.6286	0.0177	-0.4969	0.7768	-0.4442	0.5800	0.0528	0.9622
CG9053	CG9053	1626110_a_at	1.1909	0.0028	1.4665	0.0074	1.7409	0.0002	0.3230	0.2535	0.5262	0.0078	0.2032	0.1759	0.0941	0.9467	0.8037	0.0741	0.7096	0.1277
CG14651	CG14651	1626111_at	0.0901	0.6341	0.0185	0.8546	0.1351	0.4397	0.0849	0.8601	0.0230	0.9241	-0.0619	0.7393	-0.0153	0.9913	0.0369	0.9353	0.0522	0.8933
alpha-Man-II	alpha-mannosidas	1626112_at	-0.7241	0.0387	-0.0061	0.9821	0.2107	0.2432	-0.0070	0.9943	-0.2817	0.1842	-0.2747	0.1484	-0.2894	0.7307	0.2858	0.4280	0.5753	0.1393
CG4080	CG4080	1626113_at	-0.1641	0.5028	-0.9675	0.0786	-0.8468	0.0813	0.1402	0.9329	0.6407	0.1910	0.5005	0.2591	-0.0706	0.9717	-0.1509	0.8169	-0.0804	0.9078
---	---	1626114_at	0.1844	0.3419	-0.0287	0.8413	0.0039	0.9843	-0.0230	0.9778	-0.0041	0.9899	0.0188	0.9438	0.0793	0.9088	-0.0656	0.8433	-0.1449	0.5789
CG2702	CG2702	1626115_at	0.0125	0.9649	-0.2406	0.4674	-0.0578	0.8546	-0.0716	0.9518	0.0642	0.8851	0.1358	0.6952	-0.1830	0.8157	-0.0572	0.9083	0.1258	0.7355
Sp212	CG33329	1626116_at	0.0701	0.8754	-0.4149	0.6161	0.0341	0.9591	0.4397	0.6673	1.1426	0.0269	0.7028	0.1061	-0.2350	0.9487	0.3829	0.7741	0.6179	0.5988
CG5720	CG5720	1626117_at	0.0564	0.8226	0.0204	0.8840	-0.0234	0.9091	-0.0801	0.8875	0.1544	0.4355	0.2344	0.1686	0.0790	0.8940	0.0831	0.7490	0.0040	0.9920
---	---	1626118_at	-0.0658	0.7488	-0.0074	0.9551	0.2430	0.1799	0.2029	0.5067	0.1542	0.3491	-0.0487	0.7878	-0.1269	0.8494	0.0486	0.9085	0.1755	0.5499
---	---	1626119_s_at	0.1563	0.3483	0.1234	0.5059	-0.0420	0.8672	0.0469	0.9507	0.1020	0.6692	0.0551	0.8168	0.1339	0.8446	0.0959	0.7824	-0.0380	0.9215
CG9266	CG9266	1626120_at	-0.2137	0.5594	-0.6326	0.3742	-0.2680	0.2785	-0.3382	0.3749	0.0161	0.9604	0.3543	0.0689	-0.4643	0.7726	-0.2310	0.7897	0.2332	0.7750
---	---	1626121_at	0.1431	0.3120	0.1015	0.4524	0.1995	0.2151	0.0785	0.8667	0.0218	0.9244	-0.0567	0.7524	-0.0700	0.9063	-0.0305	0.9277	0.0394	0.8926
---	---	1626122_at	0.0467	0.8558	-0.0196	0.8459	0.1764	0.3276	-0.0190	0.9823	-0.0168	0.9554	0.0022	0.9928	-0.2211	0.7220	-0.0095	0.9854	0.2116	0.4325
---	---	1626123_at	0.3871	0.1412	0.6055	0.0863	0.7954	0.0133	-0.1311	0.9036	-0.4423	0.2054	-0.3113	0.3310	-0.1557	0.8608	-0.0652	0.9053	0.0905	0.8430
CG9717	CG9717	1626124_at	-1.6981	0.0365	-0.0072	0.9840	-1.9189	0.0070	-1.9375	0.0855	-2.0261	0.0083	-0.0885	0.9111	-0.0717	0.9816	-0.3465	0.6655	-0.2748	0.7434
gammaTub23C	gamma tubulin	1626125_at	-0.8566	0.0489	-0.5732	0.1336	-0.9928	0.0020	-0.3714	0.5933	-0.3905	0.2712	-0.0191	0.9661	-0.0853	0.9588	-0.0439	0.9522	0.0414	0.9486

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18368	CG18368	1626126_at	-0.0113	0.9684	0.0536	0.5813	0.0783	0.6683	0.0339	0.9558	0.0521	0.7966	0.0182	0.9281	0.0554	0.9589	0.1004	0.7910	0.0450	0.9147
---	---	1626127_at	-0.0069	0.9794	-0.0249	0.8160	0.1140	0.4717	0.3369	0.2596	0.2410	0.1717	-0.0959	0.5885	-0.0876	0.9056	-0.0302	0.9440	0.0573	0.8721
---	---	1626128_at	0.3289	0.1118	0.0335	0.8709	0.0015	0.9951	0.1471	0.7380	0.0956	0.6649	-0.0515	0.8142	0.2390	0.7202	-0.0483	0.9086	-0.2873	0.3089
5-HT1B	Serotonin recepto	1626129_at	0.2474	0.0898	-0.0312	0.7960	0.2211	0.3042	0.0461	0.9311	0.1485	0.3500	0.1024	0.4945	-0.2495	0.6955	-0.0495	0.9011	0.2000	0.4620
---	---	1626130_s_at	0.0538	0.9035	-1.1798	0.0199	-1.5557	0.0168	-0.2192	0.9158	0.9885	0.1257	1.2077	0.0433	-0.0831	0.9677	-0.1610	0.8206	-0.0779	0.9174
CG11362	CG11362	1626131_at	0.0709	0.6807	0.0052	0.9645	0.1618	0.2680	0.2160	0.4140	0.2074	0.1556	-0.0086	0.9629	-0.0122	0.9867	0.0202	0.9404	0.0324	0.8880
---	---	1626132_at	0.0099	0.9801	-0.0234	0.9146	0.2518	0.1420	0.0106	0.9931	-0.0803	0.7811	-0.0909	0.7204	0.0089	0.9946	0.0567	0.9035	0.0478	0.9096
---	---	1626133_s_at	1.6772	0.0044	2.0485	0.0026	2.7168	0.0000	0.5279	0.2936	-0.1082	0.7595	-0.6361	0.0241	-0.3225	0.7230	0.1332	0.7843	0.4557	0.2614
---	---	1626134_s_at	0.1960	0.4185	-0.0482	0.6362	0.2354	0.2987	0.2236	0.6615	0.0384	0.9102	-0.1853	0.4185	0.0365	0.9589	-0.0063	0.9878	-0.0428	0.8755
CG14476	glucosidase II	1626135_s_at	-0.2811	0.1688	0.0916	0.5378	0.4115	0.0222	0.1662	0.6338	0.0444	0.8405	-0.1218	0.4511	-0.1512	0.8191	0.4250	0.1199	0.5762	0.0708
GstD6	Glutathione S tran	1626136_at	0.1892	0.2725	0.0003	1.0000	0.1527	0.4613	0.0196	0.9819	0.0716	0.7714	0.0520	0.8231	-0.2163	0.7464	-0.1098	0.7383	0.1066	0.7423
---	---	1626137_at	0.1540	0.4381	0.0151	0.9575	0.0478	0.8214	0.0241	0.9766	-0.0056	0.9851	-0.0297	0.9028	-0.0127	0.9916	-0.0228	0.9614	-0.0101	0.9835
Eip63E	Pftaire	1626138_s_at	0.2004	0.2640	1.0270	0.0264	0.2512	0.4022	0.0573	0.9218	0.0253	0.9183	-0.0320	0.8776	0.7934	0.3800	0.9307	0.0757	0.1373	0.8255
CG13131	CG13131	1626139_at	0.1280	0.3849	-0.1160	0.3801	-0.0288	0.8819	-0.0334	0.9610	0.1941	0.2888	0.2275	0.1598	-0.2413	0.5651	-0.1081	0.5906	0.1332	0.4978
---	---	1626140_at	-0.1197	0.5440	-0.1085	0.4333	0.0869	0.6680	0.1210	0.8544	-0.0624	0.8383	-0.1833	0.4086	-0.1163	0.8609	0.0537	0.8918	0.1699	0.5488
CG34027	CG34027	1626141_at	0.1518	0.3876	-0.0058	0.9603	-0.0898	0.6664	-0.1501	0.7349	-0.0102	0.9735	0.1399	0.4518	-0.0096	0.9923	-0.1212	0.6312	-0.1115	0.6602
Lcp65Af	Lcp65Af	1626142_at	-0.0672	0.6507	-0.0586	0.5981	0.2484	0.1415	0.0738	0.8676	0.0166	0.9417	-0.0572	0.7341	-0.2765	0.6749	-0.0274	0.9505	0.2492	0.3764
CG32138 /// DmirCG32138	CG32138	1626143_a_at	-1.1946	0.0622	-0.2064	0.6008	-0.1477	0.5620	0.2099	0.8234	-0.4086	0.2658	-0.6185	0.0620	0.1595	0.9503	0.5029	0.5246	0.3435	0.6839
CG7763	CG7763	1626144_at	-0.1770	0.6937	2.1905	0.0395	1.7574	0.0129	-0.0992	0.9474	0.3960	0.3606	0.4952	0.1911	0.2300	0.9317	2.8298	0.0204	2.5997	0.0333
---	---	1626145_at	-0.0613	0.7100	0.1180	0.4410	0.0347	0.8286	-0.0396	0.9549	-0.1488	0.4518	-0.1092	0.5612	0.0066	0.9943	0.1466	0.5141	0.1400	0.5411
Obp46a	Odorant-binding p	1626146_at	0.1990	0.1896	0.4957	0.0676	0.3329	0.1064	0.0100	0.9903	0.0361	0.8831	0.0261	0.9060	0.2081	0.7506	0.1631	0.5844	-0.0450	0.9080
CPTI	mitochondrial carr	1626147_s_at	0.0632	0.8239	0.2612	0.6563	0.6492	0.0905	-0.4055	0.3576	-1.6031	0.0004	-1.1977	0.0008	-0.8420	0.6483	-1.4271	0.0714	-0.5851	0.4457
CG32110	CG32110	1626148_at	1.1274	0.0169	-0.1338	0.5338	-0.1018	0.6688	0.5130	0.0898	1.1095	0.0004	0.5965	0.0033	0.3773	0.8122	-0.0374	0.9734	-0.4147	0.5506
CG14341	CG14341	1626149_at	-0.0014	0.9958	0.0226	0.8262	0.0871	0.7481	0.0048	0.9956	-0.0754	0.7963	-0.0802	0.7581	-0.0114	0.9928	0.0282	0.9534	0.0396	0.9237
tsh	aeroplane	1626150_at	-0.2425	0.6097	0.3395	0.3202	0.0739	0.7889	-0.4240	0.6082	-0.6461	0.1177	-0.2221	0.5927	0.0800	0.9309	-0.0279	0.9538	-0.1079	0.7548
Gr32a	Gustatory recepto	1626151_at	-0.0426	0.8558	-0.0514	0.7322	-0.1649	0.4823	0.0035	0.9978	-0.0448	0.9197	-0.0483	0.8991	-0.0307	0.9816	0.0377	0.9387	0.0684	0.8636
Ror	Ror	1626152_at	0.8797	0.0591	0.5466	0.1746	0.8362	0.0026	-0.0802	0.9353	-0.1504	0.6471	-0.0702	0.8338	-0.2550	0.8292	-0.3325	0.5172	-0.0775	0.9096
Su(var)3-9	Enhancer of varie	1626153_at	0.5698	0.1076	1.1026	0.0723	1.0855	0.0080	-0.2343	0.6360	0.0193	0.9572	0.2536	0.2456	-0.0282	0.9914	0.6949	0.2386	0.7231	0.2528
CG12081 /// DsimCG12081	CG12081	1626154_at	-0.0416	0.9142	0.4890	0.2933	0.9886	0.0031	0.0897	0.9039	-1.3473	0.0007	-1.4369	0.0003	-0.3643	0.8122	-0.6971	0.2518	-0.3328	0.6257
CG14579	CG14579	1626155_at	-0.0688	0.7998	0.0040	0.9752	0.0065	0.9781	0.1339	0.7080	0.0148	0.9517	-0.1192	0.4372	0.0160	0.9913	0.0260	0.9590	0.0100	0.9846
DyakCG17347 /// l(2)37Ce	CG17347 /// lethal	1626156_at	-0.6742	0.0278	-0.5648	0.0372	-0.5349	0.0202	-0.0718	0.8716	-0.0047	0.9837	0.0671	0.6787	-0.0592	0.9716	0.0284	0.9664	0.0876	0.8745
CG13672	CG13672	1626157_at	0.2366	0.2662	0.0025	0.9845	0.2780	0.1265	0.1265	0.7760	0.0537	0.8157	-0.0727	0.7129	-0.0234	0.9862	-0.0446	0.9276	-0.0213	0.9628
nes	nessy	1626158_s_at	0.4094	0.1055	0.5809	0.0815	0.3181	0.2049	-0.4323	0.3353	-0.5912	0.0299	-0.1588	0.5294	-0.2906	0.7154	-0.4962	0.1362	-0.2056	0.5612
---	---	1626159_at	0.0863	0.6605	0.0409	0.7749	0.1999	0.2641	0.0677	0.8816	0.0756	0.6678	0.0079	0.9679	-0.0611	0.9495	0.1770	0.5576	0.2381	0.4153
---	---	1626160_s_at	-0.1312	0.4119	0.0553	0.7086	-0.0313	0.8627	0.2140	0.5756	-0.0690	0.7695	-0.2830	0.1056	0.1348	0.8454	0.0855	0.8166	-0.0493	0.8998
---	---	1626161_at	-0.1443	0.5202	-0.1650	0.3878	0.0901	0.7445	0.0893	0.9263	0.0617	0.8725	-0.0276	0.9393	-0.1240	0.8889	-0.0528	0.9189	0.0711	0.8745
alph	alphabet	1626162_s_at	-0.9981	0.0485	-0.1903	0.7713	-0.4623	0.0942	-0.2216	0.8671	-0.3039	0.5464	-0.0823	0.8818	0.0262	0.9914	0.5386	0.3320	0.5124	0.3807
Act5C	5C actin	1626163_s_at	-0.6325	0.0598	0.2639	0.3835	0.9502	0.0003	0.3184	0.5128	0.1274	0.6612	-0.1909	0.4284	-0.3336	0.6955	0.9640	0.0290	1.2975	0.0236
CG31380	CG31380	1626164_at	1.4705	0.0113	0.4801	0.4019	0.4603	0.0264	-0.0248	0.9673	0.0382	0.8531	0.0630	0.7086	-0.1001	0.9677	-0.8263	0.1884	-0.7262	0.2779
CG3754	CG3754	1626165_at	-0.0922	0.8385	0.5069	0.4670	0.2704	0.0817	-0.3685	0.3348	-0.8260	0.0035	-0.4574	0.0281	-0.0944	0.9816	-0.2231	0.8563	-0.1287	0.9173
CG13082	CG13082	1626166_at	0.3858	0.1660	-0.1669	0.1964	-0.0178	0.9453	0.0992	0.7902	0.1423	0.3598	0.0431	0.8012	-0.0652	0.9657	-0.4150	0.2973	-0.3498	0.4007
CG11125	CG11125	1626167_at	-1.2581	0.0406	-0.4311	0.1074	-0.8465	0.0130	-0.0171	0.9872	-0.9135	0.0043	-0.8964	0.0027	0.3321	0.7500	-0.3379	0.4487	-0.6700	0.1573
CG7211	CG7211	1626168_at	0.0754	0.8493	0.2237	0.2870	0.1702	0.3038	0.0140	0.9883	0.0589	0.8396	0.0450	0.8643	0.1342	0.8953	0.1668	0.6969	0.0326	0.9514
CG2701	transcript II	1626169_at	-0.4222	0.3526	-0.3538	0.4689	-0.1887	0.3669	0.1375	0.8156	0.1259	0.6249	-0.0116	0.9685	0.1083	0.9677	0.4037	0.5855	0.2954	0.7067
CG18445	O-acyl transferase	1626170_at	-0.8188	0.0054	-0.2520	0.3038	-0.5264	0.0092	0.0326	0.9626	-0.1911	0.3051	-0.2237	0.1744	0.3345	0.6898	0.3091	0.3617	-0.0254	0.9596
---	---	1626171_at	-2.5338	0.0060	-4.4323	0.0130	-4.2013	0.0001	0.1413	0.9314	1.6021	0.0064	1.4608	0.0058	0.0196	0.9964	-0.1524	0.9237	-0.1721	0.9025
CG12187	CG12187	1626172_at	-0.0471	0.8663	-0.0118	0.9101	-0.3416	0.1077	-0.0604	0.9363	-0.1053	0.6761	-0.0449	0.8616	0.0235	0.9816	-0.0926	0.7439	-0.1161	0.6567
SmB	polypeptide B	1626173_at	0.4215	0.0798	0.9774	0.0332	0.9909	0.0057	0.0992	0.8578	0.3261	0.1019	0.2269	0.2027	0.0066	0.9977	0.7110	0.1512	0.7044	0.1911
jdp	jdp	1626174_a_at	-0.2645	0.6917	-0.9186	0.0409	-0.8102	0.0023	0.2301	0.7293	0.5342	0.0752	0.3042	0.2561	0.0616	0.9862	-0.1262	0.9192	-0.1877	0.8575
CG11534	CG11534	1626175_at	-0.0168	0.9533	0.1141	0.3492	0.2687	0.1013	-0.0037	0.9956	-0.3426	0.0697	-0.3389	0.0478	-0.1541	0.7810	-0.1952	0.4097	-0.0411	0.9018
---	---	1626176_at	0.0506	0.7642	0.2642	0.0497	0.0183	0.9347	-0.2728	0.2977	-0.2792	0.0725	-0.0064	0.9732	0.0621	0.9238	-0.0695	0.8018	-0.1316	0.5670
Gint3	GDI interacting pr	1626177_s_at	-0.2403	0.1621	0.0730	0.7655	-0.0511	0.7824	-0.0004	0.9994	-0.0756	0.6505	-0.0751	0.6170	0.0795	0.9296	0.2119	0.4743	0.1324	0.6800
---	---	1626178_s_at	0.7258	0.0345	0.2794	0.1890	0.2945	0.2335	0.2984	0.6354	0.9669	0.0079	0.6685	0.0239	0.2516	0.7697	0.5384	0.1382	0.2868	0.4457
Hsp60C	Hsp60C	1626179_s_at	-0.1386	0.4559	0.0159	0.9099	0.1350	0.3370	0.2463	0.4637	0.0159	0.9531	-0.2305	0.1553	-0.0463	0.9374	-0.0156	0.9591	0.0308	0.9045
Gmer	GDP-4-keto-6-dec	1626180_at	0.8558	0.0034	0.5704	0.1821	0.4233	0.0304	-0.1277	0.7869	0.6613	0.0054	0.7890	0.0015	0.0800	0.9503	0.3563	0.3397	0.2763	0.4852
CG1969	CG1969	1626181_at	0.0675	0.6783	0.0555	0.6425	0.1838	0.2020	0.0297	0.9558	0.0196	0.9228	-0.0101	0.9567	-0.0302	0.9717	-0.0286	0.9341	0.0016	0.9966
CG5026	CG5026	1626182_a_at	-0.1941	0.3932	0.4498	0.0932	0.2883	0.0749	-0.1985	0.7091	-0.2312	0.3485	-0.0327	0.9111	-0.0921	0.9246	0.2939	0.3612	0.3860	0.2557
CG2083	CG2083	1626183_at	0.2097	0.1973	0.0790	0.6624	0.1207	0.5847	0.0361	0.9570	0.1478	0.4277	0.1117	0.5239	0.0196	0.9893	0.0608	0.8903	0.0412	0.9211
CG16978 /// DyakCG16978	CG16978	1626184_at	0.0906	0.6635	0.0412	0.6885	0.2000	0.2224	-0.0035	0.9956	-0.1227	0.5720	-0.1192	0.5451	-0.1513	0.8513	-0.1827	0.6093	-0.0314	0.9466
---	---	1626185_at	0.0076	0.9732	0.0789	0.6844	0.0918	0.6074	0.0585	0.9039	0.1281	0.4339	0.0696	0.6721	0.0217	0.9848	0.1257	0.6585	0.1040	0.7240
CG18193 /// DmauCG18193	CG18193	1626186_at	-0.0077	0.9708	-0.0605	0.5827	-0.2029	0.3327	-0.0624	0.9471	0.0370	0.9213	0.0994	0.7262	-0.0094	0.9928	0.0276	0.9440	0.0371	0.9118
Brf	Brf	1626187_at	0.2742	0.2222	-0.0203	0.8771	0.0182	0.9459	-0.0049	0.9956	0.3890	0.0435	0.3939	0.0264	0.0309	0.9831	0.2029	0.5684	0.1720	0.6375
---	---	1626188_at	0.2608	0.2368	0.0192	0.9515	0.0731	0.7193	0.1020	0.8841	0.2324	0.3373	0.1305	0.5836	-0.0500	0.9689	0.0344	0.9497	0.0844	0.8435
---	---	1626189_s_at	0.0930	0.6327	-0.1941	0.1743	-0.0039	0.9881	0.2327	0.3583	0.3613	0.0207	0.1286	0.3304	-0.1441	0.7726	0.0077	0.9854	0.1518	0.4984
CG10825	CG10825	1626190_a_at	-0.1328	0.6934	0.2547	0.1061	0.6583	0.0112	0.0193	0.9883	-0.1900	0.5647	-0.2093	0.4710	-0.3383	0.3712	0.1469	0.4870	0.4852	0.0559
CG10600	CG10600	1626191_at	-0.4552	0.2983	1.0914	0.0621	0.6215	0.0080	-0.3260	0.7190	-0.9962	0.0234	-0.6702	0.0704	0.1370	0.9291	0.4844	0.3169	0.3473	0.5003
CG32709	CG32709	1626192_at	0.2217	0.7145	-1.1432	0.0846	-0.6747	0.2874	-0.4907	0.5255	0.6686	0.0990	1.1593	0.0070	-0.8645	0.7644	-0.5920	0.6571	0.2724	0.8655
---	---	1626193_at	0.0314	0.8816	-0.0187	0.8768	0.1135	0.5975	0.0880	0.8732	0.0646	0.7803	-0.0233	0.9209	-0.0739	0.9324	-0.0849	0.8153	-0.0110	0.9816
melt	melted	1626194_at	-0.0267	0.9288	0.7014	0.0523	0.5845	0.0224	0.0433	0.9586	-0.3458	0.1154	-0.3891	0.0528	0.1959	0.8016	0.2589	0.4258	0.0629	0.8884

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG12620	CG12620	1626195_at	0.2746	0.2908	0.1789	0.3900	0.2337	0.1333	0.0523	0.9325	-0.0592	0.7924	-0.1115	0.5335	-0.0203	0.9894	-0.0567	0.9075	-0.0364	0.9349
CaBP1	CaBP1	1626196_at	1.0646	0.0017	0.8778	0.0330	1.4897	0.0004	0.2934	0.6041	1.1615	0.0024	0.8682	0.0053	-0.2155	0.7220	1.0480	0.0105	1.2635	0.0050
CG31391	CG31391	1626197_at	0.0867	0.5898	-0.0928	0.5697	0.1444	0.3983	-0.0002	0.9999	0.0117	0.9693	0.0118	0.9611	-0.1587	0.7644	-0.1081	0.6616	0.0507	0.8646
Cyp4d8	Cytochrome P450	1626198_at	-0.6161	0.4663	-0.3386	0.6545	-0.1153	0.7078	-0.6842	0.2390	-0.9537	0.0132	-0.2695	0.3990	-0.8993	0.7644	-0.8975	0.4954	0.0017	0.9992
CG12104	CG12104	1626199_at	0.0195	0.9264	0.5547	0.0321	0.6658	0.0032	-0.0003	0.9997	-0.3229	0.0988	-0.3226	0.0679	-0.0508	0.9401	0.2080	0.3042	0.2588	0.2292
tipE	temperature-induc	1626200_s_at	0.1161	0.5949	-0.0426	0.7992	0.1076	0.5569	-0.0568	0.9295	0.1520	0.4360	0.2088	0.2149	-0.0731	0.9411	-0.1193	0.7403	-0.0462	0.9125
CG31091	CG31091	1626201_at	0.1496	0.4016	0.0000	1.0000	-0.0232	0.9083	-0.0953	0.8837	-0.1898	0.4049	-0.0945	0.6842	-0.0156	0.9862	-0.2239	0.2571	-0.2083	0.3205
---	---	1626202_at	0.1854	0.4383	0.3528	0.1506	-0.1190	0.4718	-0.1738	0.5842	-0.0312	0.8863	0.1426	0.3354	0.1791	0.7677	0.1466	0.5909	-0.0325	0.9274
DyakCG17768	CG17768	1626203_at	-0.1279	0.6319	0.4570	0.0755	0.3338	0.0755	0.0772	0.8671	-0.0632	0.7421	-0.1404	0.3468	0.2005	0.8032	0.3446	0.2894	0.1440	0.7006
---	---	1626204_at	0.7682	0.0543	1.8398	0.0061	1.4962	0.0203	0.0115	0.9956	-0.1784	0.7844	-0.1900	0.7428	0.3835	0.7953	0.8933	0.1420	0.5099	0.4215
---	---	1626205_s_at	-0.5563	0.5776	-2.2634	0.0935	-2.1233	0.0014	-0.0675	0.9777	1.6988	0.0137	1.7663	0.0070	-0.1418	0.9829	-0.0295	0.9935	0.1124	0.9596
CG6947 /// DmirCG6947	CG6947	1626206_at	0.1568	0.5820	0.0902	0.4365	0.1314	0.5381	0.0757	0.8606	-0.0101	0.9666	-0.0858	0.5780	0.1372	0.8609	0.0266	0.9593	-0.1106	0.7647
---	---	1626207_at	0.2399	0.3164	-0.0928	0.4917	0.4081	0.0537	0.3268	0.4162	0.4593	0.0450	0.1326	0.5435	-0.2314	0.6749	-0.0324	0.9272	0.1990	0.3835
Exp6	Exportin 6	1626208_at	0.8708	0.0350	0.4330	0.1353	0.1684	0.4189	0.0143	0.9883	0.6866	0.0088	0.6723	0.0058	0.1476	0.9076	0.0880	0.8964	-0.0596	0.9234
RpL35	Ribosomal protein	1626209_a_at	0.1421	0.3227	0.1415	0.1988	0.1496	0.2771	0.0077	0.9909	-0.0622	0.7096	-0.0699	0.6373	-0.0236	0.9759	-0.0884	0.6729	-0.0648	0.7703
---	---	1626210_at	0.1005	0.6964	-0.0500	0.6072	-0.1031	0.5517	0.0929	0.8074	0.2380	0.1151	0.1452	0.2903	0.0069	0.9952	-0.1405	0.6387	-0.1474	0.6195
CG34351	CG11928	1626211_at	0.1740	0.4601	0.5056	0.0661	0.0766	0.8003	-0.0011	0.9991	-0.1028	0.7246	-0.1017	0.6996	0.1241	0.9112	0.1422	0.7667	0.0181	0.9769
---	---	1626212_at	0.0480	0.8601	0.2088	0.2358	0.1496	0.4278	-0.1691	0.7121	-0.1986	0.3478	-0.0296	0.9065	0.0888	0.8725	0.0533	0.8610	-0.0354	0.9065
Pcmt	protein D-aspartyl	1626213_a_at	-0.9300	0.0033	-1.4889	0.0145	-1.1987	0.0002	0.2782	0.3817	0.2003	0.2669	-0.0779	0.6784	0.0901	0.9460	-0.3115	0.4391	-0.4016	0.3300
CG40354 /// Eb1	Eb1 /// CG40354	1626214_s_at	-0.4212	0.2594	0.2105	0.4242	0.4876	0.0776	0.0425	0.9777	-0.3838	0.3111	-0.4263	0.2032	0.0395	0.9848	0.4374	0.3417	0.3979	0.4078
CG2202	CG2202	1626215_at	-0.6141	0.2848	0.3094	0.7026	0.6374	0.0083	0.1539	0.8102	-0.6033	0.0257	-0.7572	0.0058	-0.1738	0.9610	0.2661	0.8427	0.4399	0.6877
---	---	1626216_at	0.0365	0.8478	0.0769	0.5883	0.1796	0.4910	0.0549	0.9413	-0.0409	0.8877	-0.0958	0.6661	0.0064	0.9964	0.0740	0.8684	0.0676	0.8745
Ipp	inositol polyphosp	1626217_at	-1.5494	0.0008	-1.5688	0.0386	-1.8407	0.0001	-0.3090	0.3386	-0.4420	0.0256	-0.1330	0.4492	0.0527	0.9775	-0.3533	0.4285	-0.4061	0.3764
oaf	transcript-near-de	1626218_s_at	1.2376	0.0201	0.8100	0.0775	0.1882	0.5038	-0.3737	0.7014	0.5919	0.1835	0.9656	0.0239	0.2331	0.8270	0.2032	0.6812	-0.0299	0.9636
Ogt	O-glycosyltransfer	1626219_s_at	0.0947	0.6356	-0.1655	0.7157	-0.3579	0.0272	-0.2101	0.5008	0.4839	0.0105	0.6940	0.0013	0.0392	0.9831	0.3248	0.4446	0.2856	0.5202
CG3448	CG3448	1626220_at	0.0108	0.9829	0.1086	0.5331	-0.1482	0.6258	-0.0542	0.9643	0.5205	0.1048	0.5747	0.0505	0.1276	0.9514	0.7083	0.2430	0.5807	0.3680
Glycogenin	Glycogenin	1626221_at	-2.5249	0.0027	-1.3766	0.1061	-2.4178	0.0034	-0.4628	0.4197	-1.6426	0.0009	-1.1797	0.0022	0.5898	0.8465	-0.4672	0.7573	-1.0570	0.4114
Ser8	Serine Protease 3	1626222_at	0.2884	0.4108	1.4450	0.1331	1.0603	0.0049	-0.8576	0.3196	0.3266	0.5459	1.1842	0.0165	-0.5988	0.7752	1.2726	0.1498	1.8714	0.0755
CG2543 /// DyakCG2543	CG2543	1626223_at	0.3597	0.2024	0.0268	0.9522	0.4198	0.1831	0.4572	0.3855	0.2867	0.3442	-0.1705	0.5629	0.0917	0.9634	0.0200	0.9848	-0.0717	0.9234
fws	Drosophila Cog5	1626224_s_at	0.2409	0.4012	1.1218	0.0456	1.2929	0.0002	0.0927	0.8640	-0.2262	0.2398	-0.3189	0.0669	-0.1206	0.9246	0.6830	0.1151	0.8035	0.0947
---	---	1626225_at	0.2046	0.4089	-0.1354	0.4744	-0.4565	0.0310	-0.0371	0.9603	0.2364	0.2297	0.2735	0.1193	0.1549	0.8461	0.0403	0.9402	-0.1146	0.7651
---	---	1626226_at	0.2720	0.0824	0.1521	0.3241	0.0777	0.6566	0.1007	0.7604	0.1555	0.2765	0.0547	0.7174	0.0870	0.9046	0.0232	0.9572	-0.0638	0.8504
CG5337	CG5337	1626227_at	-0.4901	0.0808	-0.3701	0.1051	-0.3520	0.0821	-0.0910	0.8162	-0.0958	0.5637	-0.0047	0.9802	-0.0256	0.9901	0.2116	0.6673	0.2373	0.6218
CG6967	CG6967	1626228_a_at	-0.6763	0.0916	-1.5202	0.0302	-1.6132	0.0094	-0.1346	0.9177	0.8713	0.0366	1.0059	0.0128	-0.3419	0.8930	0.0253	0.9906	0.3672	0.7362
---	---	1626229_at	-0.0940	0.5639	-0.0210	0.8696	0.2139	0.1379	0.1507	0.6090	-0.0299	0.8796	-0.1806	0.1714	0.0608	0.9466	0.2741	0.3030	0.2133	0.4503
CG31358	CG31358	1626230_s_at	-0.1770	0.4178	0.1731	0.4307	0.2203	0.3085	0.0270	0.9633	-0.0275	0.8973	-0.0545	0.7460	-0.0501	0.9734	0.3023	0.4093	0.3523	0.3543
CG4704	CG4704	1626231_at	0.1328	0.5147	0.0211	0.8447	0.0693	0.6667	0.1966	0.5531	0.1436	0.4170	-0.0531	0.7809	-0.0007	0.9998	-0.0479	0.9046	-0.0471	0.8949
Camta	Calmodulin-bindin	1626232_at	0.7346	0.0822	0.4075	0.2278	-0.3030	0.5190	-0.5272	0.1358	0.3552	0.0964	0.8824	0.0014	0.4510	0.8331	0.3032	0.7852	-0.1478	0.9054
CG8965	CG8965	1626233_at	0.3544	0.4107	-0.4040	0.4733	0.3721	0.5180	0.8504	0.3240	1.8440	0.0038	0.9936	0.0338	0.1230	0.9734	1.1101	0.2053	0.9871	0.2897
---	---	1626234_at	0.0334	0.9011	-0.0280	0.7968	0.2168	0.1564	0.1178	0.8066	0.0719	0.7522	-0.0459	0.8338	-0.1609	0.7707	-0.0455	0.8981	0.1154	0.6498
m	miniature	1626235_at	0.0686	0.7112	-0.2690	0.1497	0.1019	0.6362	0.2645	0.4630	0.2990	0.1235	0.0345	0.8811	-0.2552	0.6898	-0.1615	0.5591	0.0937	0.7578
CG10343	CG10343	1626236_at	0.1443	0.3251	0.0901	0.7642	0.0636	0.7225	0.0285	0.9584	0.5397	0.0034	0.5112	0.0026	0.1003	0.9238	0.5555	0.1179	0.4552	0.2130
CG12734 /// DmirCG12734	CG12734	1626237_at	-0.1293	0.4494	-0.2831	0.0613	-0.5996	0.0093	-0.0518	0.9387	0.3544	0.0749	0.4062	0.0289	0.1609	0.7506	0.1309	0.5654	-0.0300	0.9198
CG31690	CG31690	1626238_at	-0.0548	0.8181	-0.0658	0.6290	0.0301	0.8614	-0.0658	0.9060	-0.2937	0.1020	-0.2279	0.1551	0.0293	0.9814	0.0092	0.9865	-0.0201	0.9585
bon	bonus	1626239_at	-0.7270	0.0461	-1.0927	0.0230	-1.3150	0.0002	-0.1006	0.8017	0.4385	0.0143	0.5391	0.0034	0.0606	0.9742	0.0381	0.9600	-0.0225	0.9756
CG6236	CG6236	1626240_s_at	-0.0640	0.8046	0.0350	0.9119	-0.1062	0.5250	-0.0373	0.9445	0.0516	0.7811	0.0889	0.5569	0.1000	0.9235	0.0997	0.8248	-0.0004	0.9995
Cyt-c-d	cytochrome c	1626241_at	0.4770	0.0544	-0.0082	0.9444	0.2761	0.2178	0.1827	0.7321	0.1237	0.6431	-0.0590	0.8282	-0.1046	0.9239	-0.1946	0.6209	-0.0900	0.8485
CG14795	CG14795	1626242_at	-2.5416	0.0049	0.0682	0.8754	-0.7801	0.0060	-0.6803	0.2397	-2.4776	0.0003	-1.7973	0.0005	0.2181	0.9238	-0.2890	0.7466	-0.5071	0.5257
CG11626	CG11626	1626243_at	0.2313	0.2752	0.0325	0.8386	-0.0065	0.9724	0.1557	0.6354	0.1930	0.2306	0.0374	0.8407	0.1748	0.7322	0.0551	0.8546	-0.1197	0.6121
---	---	1626244_at	-0.0079	0.9751	0.0714	0.6791	0.0704	0.7158	0.1222	0.7990	0.1256	0.5477	0.0034	0.9889	0.1239	0.8903	0.0857	0.8527	-0.0381	0.9352
---	---	1626245_at	0.2396	0.2626	0.1669	0.5456	0.0808	0.7172	0.0920	0.8738	-0.0300	0.9158	-0.1220	0.5373	0.1703	0.8472	-0.0310	0.9590	-0.2013	0.6112
vig	vasa intronic gene	1626246_s_at	0.2381	0.6635	0.0356	0.8951	-0.6164	0.1176	-0.3379	0.4455	0.7917	0.0063	1.1296	0.0008	0.2926	0.8960	0.5614	0.5161	0.2688	0.7869
CG4785	CG4785	1626247_at	0.0797	0.8601	0.1960	0.3725	0.4301	0.0165	0.1782	0.6589	0.0376	0.8856	-0.1406	0.4407	-0.1087	0.8930	0.1018	0.7858	0.2105	0.4997
CG18547	CG18547	1626248_at	1.5353	0.0031	2.4838	0.0075	2.3729	0.0000	0.3620	0.4645	-0.2540	0.3517	-0.6160	0.0182	0.4775	0.6955	0.6613	0.1868	0.1838	0.7574
CycE	cycline E	1626249_s_at	-0.1771	0.8141	-0.8393	0.3322	-2.1328	0.0057	-0.7964	0.3744	1.0458	0.0450	1.8421	0.0024	0.1600	0.9775	0.0640	0.9792	-0.0960	0.9614
CG14282	CG14282	1626250_at	-0.1449	0.4471	0.1427	0.3973	-0.1134	0.6231	-0.0938	0.9068	-0.0894	0.7731	0.0044	0.9892	0.0190	0.9852	0.1086	0.6762	0.0897	0.7409
Rh4	rhodopsin	1626251_at	0.2956	0.3115	0.5557	0.0328	0.5627	0.0226	0.2392	0.6122	-0.2356	0.3272	-0.4748	0.0325	0.1313	0.9125	-0.0749	0.9075	-0.2062	0.6484
---	---	1626252_at	0.2250	0.2838	0.2761	0.2602	0.2634	0.0994	-0.0800	0.9048	-0.0367	0.9026	0.0433	0.8640	-0.1457	0.8480	-0.0410	0.9353	0.1047	0.7794
GstD4	Glutathione S tran	1626253_at	0.0501	0.7708	0.0757	0.7400	0.3682	0.0522	0.2247	0.5444	0.2093	0.2787	-0.0154	0.9496	-0.0712	0.9427	0.0283	0.9540	0.0995	0.7836
CG34370	CG13499	1626254_s_at	-0.0534	0.7245	0.1670	0.4061	0.0773	0.7571	-0.1053	0.7884	-0.1834	0.2558	-0.0782	0.6352	0.0165	0.9893	0.0551	0.8792	0.0386	0.9121
Gr61a	Gustatory recepto	1626255_at	0.1204	0.6227	0.1008	0.4490	0.1676	0.5772	-0.0431	0.9668	-0.1380	0.6515	-0.0949	0.7469	0.1846	0.7707	-0.1759	0.5321	-0.3605	0.2055
CG13836	CG13836	1626256_at	-1.4117	0.0649	0.3231	0.3944	-0.8540	0.0013	-0.6743	0.4333	-1.3713	0.0108	-0.6970	0.0984	0.3634	0.7893	0.1077	0.9022	-0.2557	0.6881
CG13656	CG13656	1626257_at	0.1229	0.5411	0.0646	0.7114	0.1467	0.6199	-0.0982	0.9251	-0.0375	0.9321	0.0607	0.8682	-0.1513	0.7495	-0.1186	0.5800	0.0327	0.9067
---	---	1626258_at	0.1002	0.4803	0.4659	0.0944	0.5703	0.0436	0.0745	0.9015	-0.1946	0.3262	-0.2691	0.1217	0.0489	0.9677	0.2049	0.5374	0.1560	0.6491
su(s)	suppressor of sab	1626259_at	-0.1982	0.5554	0.3562	0.3859	0.2061	0.3402	0.0305	0.9603	-0.4912	0.0093	-0.5217	0.0042	0.1283	0.9498	0.0865	0.9277	-0.0418	0.9624
CG11778	CG11778	1626260_at	0.1278	0.6741	0.3293	0.0719	0.1343	0.4527	-0.1529	0.7753	-0.2554	0.2566	-0.1025	0.6590	0.1438	0.8326	0.0493	0.9088	-0.0945	0.7764
CG8029 /// DyakCG8029	CG8029	1626261_s_at	-0.7587	0.0040	-0.6379	0.0440	-0.8599	0.0006	-0.1633	0.5681	-0.2456	0.0949	-0.0823	0.5726	-0.0358	0.9816	-0.1973	0.5647	-0.1615	0.6464
CG3104	CG3104	1626262_s_at	-0.7699	0.0267	0.2557	0.3900	-0.0617	0.8272	-0.0223	0.9863	-0.6753	0.0358	-0.6530	0.0263	0.1904	0.8443	0.3954	0.3066	0.2050	0.6326
CG7510	CG7510	1626263_at	-0.6222	0.0349	-1.9905	0.0042	-1.4847	0.0044	0.2335	0.8078	1.0088	0.0165	0.7753	0.0310	-0.1731	0.8692	-0.3306	0.4317	-0.1575	0.7461

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9883	CG9883	1626264_at	0.0118	0.9812	-0.4500	0.2554	-0.5714	0.1927	-0.0004	0.9997	0.6746	0.0102	0.6750	0.0061	-0.1014	0.9816	0.1137	0.9404	0.2150	0.8618
rdx	roadkill	1626265_at	0.3982	0.1526	0.8383	0.1233	0.3438	0.1058	-0.2376	0.6010	-0.3401	0.1356	-0.1026	0.6659	0.2871	0.7770	0.2433	0.6041	-0.0438	0.9433
MAGE	MAGE	1626266_at	0.5307	0.0244	0.2066	0.5348	0.1907	0.2405	-0.0338	0.9627	0.4851	0.0200	0.5189	0.0091	-0.0310	0.9852	0.1986	0.6288	0.2296	0.5698
---	---	1626267_at	0.0711	0.6617	0.0443	0.8638	-0.0006	0.9978	0.0507	0.9375	0.2127	0.2605	0.1620	0.3486	0.1708	0.7810	0.0465	0.9085	-0.1243	0.6675
Klp54D	Kinesin-like protei	1626268_at	-0.0022	0.9936	0.0829	0.7248	-0.2872	0.0722	-0.1419	0.8086	-0.0662	0.8202	0.0758	0.7686	0.0642	0.9514	0.1254	0.7308	0.0612	0.8875
---	---	1626269_at	-0.0014	0.9957	0.0000	1.0000	0.0461	0.8286	0.0147	0.9840	0.0056	0.9823	-0.0091	0.9669	-0.0823	0.8692	0.0043	0.9925	0.0866	0.6982
---	---	1626270_s_at	0.1775	0.4477	0.2486	0.2331	0.1554	0.3522	-0.0009	0.9991	-0.1077	0.6246	-0.1068	0.5898	0.0936	0.9063	-0.0280	0.9508	-0.1216	0.7035
CG9634	CG9634	1626271_at	0.3904	0.3132	0.0434	0.9238	0.4505	0.0879	-0.5470	0.0687	-0.9037	0.0009	-0.3567	0.0298	-0.9104	0.5754	-1.3084	0.0871	-0.3979	0.6152
Sp7	Melanization Prot	1626272_s_at	-0.0837	0.8517	-0.3156	0.3697	0.4460	0.1376	0.2962	0.5008	0.4931	0.0404	0.1969	0.3572	-0.5338	0.7387	0.1809	0.8444	0.7147	0.3042
CG6232	CG6232	1626273_at	0.3022	0.1063	0.2212	0.3144	0.4203	0.0768	-0.0896	0.8889	-0.3762	0.0816	-0.2866	0.1345	-0.0729	0.9210	-0.2295	0.3397	-0.1566	0.5488
---	---	1626274_at	0.0777	0.7142	0.1773	0.3120	0.3746	0.0851	0.1560	0.7845	-0.0290	0.9298	-0.1850	0.3975	-0.0202	0.9906	0.0621	0.9111	0.0823	0.8635
---	---	1626275_at	-0.3038	0.1716	-1.8469	0.0247	-1.3597	0.0273	0.9329	0.4420	2.0439	0.0083	1.1110	0.0640	-0.0476	0.9653	-0.0106	0.9854	0.0371	0.9278
CG7967	CG7967	1626276_at	-0.6301	0.0202	-0.6625	0.0136	-0.5998	0.0552	0.3149	0.3580	0.2627	0.1768	-0.0522	0.8130	0.2210	0.8446	0.1662	0.7676	-0.0548	0.9330
---	---	1626277_at	0.2341	0.3188	0.3201	0.1390	0.1530	0.4015	-0.2100	0.6763	-0.1577	0.5324	0.0524	0.8484	0.1284	0.8222	0.0597	0.8550	-0.0688	0.8167
CG6180	CG6180	1626278_at	-0.0799	0.6327	0.2917	0.1450	0.3623	0.1308	-0.0910	0.9028	-0.3238	0.1770	-0.2328	0.2863	-0.0774	0.9168	0.0640	0.8554	0.1414	0.6084
CG33087	LDLR-like	1626279_at	0.1596	0.4597	0.0073	0.9642	-0.1114	0.5793	0.0837	0.8901	0.1291	0.5543	0.0454	0.8469	0.1054	0.8903	-0.0707	0.8587	-0.1761	0.5629
---	---	1626280_a_at	-0.3139	0.4201	-0.3694	0.2837	-0.7165	0.0053	-0.2241	0.6823	0.3037	0.2338	0.5278	0.0284	0.1158	0.9514	0.2216	0.7406	0.1059	0.8923
---	---	1626281_at	0.0454	0.8997	-0.0706	0.6811	-0.0454	0.7946	0.1465	0.8556	-0.0029	0.9945	-0.1494	0.6111	0.0122	0.9901	-0.1184	0.6080	-0.1306	0.5684
---	---	1626282_at	0.0582	0.6849	0.0872	0.4841	0.0800	0.7358	0.1975	0.7432	0.0481	0.8941	-0.1494	0.5704	0.1423	0.7779	0.0482	0.8782	-0.0941	0.7013
---	---	1626283_at	-0.2865	0.3047	0.1524	0.4873	0.0294	0.9036	-0.0946	0.8611	-0.2537	0.1882	-0.1591	0.3746	-0.0930	0.9407	-0.0767	0.8933	0.0162	0.9796
CG13500	CG13500	1626284_at	0.1141	0.4954	0.0191	0.8502	0.1629	0.4028	-0.0571	0.9314	0.1355	0.5167	0.1925	0.2791	0.0815	0.9164	0.0740	0.8376	-0.0075	0.9859
CG13806	CG13806	1626285_at	-0.0078	0.9811	0.2597	0.2543	0.1911	0.3581	-0.0573	0.9422	-0.2163	0.3447	-0.1590	0.4548	0.1162	0.8810	0.0791	0.8438	-0.0371	0.9282
CG11788 /// DsecCG11788	CG11788	1626286_at	-0.1119	0.6569	-0.3038	0.0722	0.0206	0.9438	-0.4704	0.2975	0.0803	0.8053	0.5508	0.0280	-0.7169	0.5186	-0.2203	0.7165	0.4966	0.3629
GV1	GV1	1626287_at	-0.0702	0.8830	-0.4113	0.4686	0.2477	0.4286	0.5138	0.4962	0.2289	0.6015	-0.2850	0.4505	-0.0604	0.9862	-0.1237	0.9192	-0.0633	0.9548
---	---	1626288_at	0.0676	0.8649	-0.1176	0.3380	-0.1992	0.2836	0.0253	0.9774	0.1896	0.3977	0.1643	0.4219	-0.0215	0.9913	-0.1244	0.8176	-0.1029	0.8470
Dgkepsilon	Diacyl glycerol kin	1626289_at	-0.1150	0.6462	-0.1061	0.5419	0.0719	0.6889	0.0656	0.9117	0.0219	0.9341	-0.0437	0.8407	-0.0725	0.9445	0.1234	0.7378	0.1959	0.5579
CG6752	CG6752	1626290_at	-0.0803	0.6165	-0.0275	0.7876	-0.1276	0.6115	-0.1478	0.7168	0.0845	0.6918	0.2323	0.1603	-0.1470	0.8427	-0.0244	0.9617	0.1226	0.7224
Cpr56F /// DsecCG9036 ///	CG9036	1626291_at	-1.7243	0.0253	1.1452	0.0432	-0.4769	0.3474	-1.0058	0.4420	-4.2334	0.0005	-3.2276	0.0009	0.7187	0.6749	-1.3038	0.0787	-2.0225	0.0368
CG4439	CG4439	1626292_at	-0.0199	0.9554	0.0989	0.4561	0.2419	0.2424	0.1760	0.7432	-0.0604	0.8436	-0.2364	0.2672	0.0593	0.9577	0.1617	0.6345	0.1024	0.7838
---	---	1626293_at	0.0764	0.5943	0.0772	0.6071	0.1262	0.3900	0.0088	0.9922	-0.1317	0.5055	-0.1404	0.4231	-0.0324	0.9652	-0.0111	0.9738	0.0212	0.9390
stai	D-stathmin	1626294_a_at	-1.2581	0.1175	-1.3960	0.1082	-1.7867	0.0273	-0.2146	0.8124	-0.2850	0.4419	-0.0705	0.8657	0.1830	0.9752	-0.3279	0.8749	-0.5109	0.7614
---	---	1626295_at	0.2455	0.2699	-0.0204	0.8645	0.1715	0.4036	0.0479	0.9464	0.0073	0.9809	-0.0406	0.8621	-0.1516	0.8049	-0.1815	0.4864	-0.0299	0.9338
CG31303	CG31303	1626296_at	-0.9742	0.0127	-0.2093	0.4753	-0.6348	0.0098	-0.4957	0.3193	-0.6007	0.0444	-0.1051	0.7363	-0.0099	0.9964	0.3363	0.5158	0.3462	0.5042
---	---	1626297_at	0.2485	0.1739	0.0784	0.5089	-0.1465	0.3925	-0.1498	0.6818	0.0472	0.8249	0.1971	0.1941	0.0442	0.9420	-0.0476	0.8527	-0.0918	0.6476
CG14958	CG14958	1626298_at	-0.7881	0.0322	0.1770	0.3278	-0.3374	0.2458	-0.3088	0.7271	-1.2129	0.0085	-0.9042	0.0189	0.1372	0.8823	-0.2088	0.5767	-0.3459	0.3406
yl	YP receptor	1626299_at	0.7923	0.4781	-1.6582	0.1422	-0.8343	0.1426	0.4837	0.2438	2.1608	0.0001	1.6771	0.0002	-0.3849	0.9514	-0.1001	0.9758	0.2848	0.9121
CG17068	CG17068	1626300_at	-0.1113	0.8705	-0.0902	0.8976	-0.3368	0.3733	-0.0699	0.9112	0.2807	0.1555	0.3506	0.0527	0.2229	0.9514	0.2992	0.8353	0.0764	0.9614
ldgf3	Imaginal disc grov	1626301_at	-0.4767	0.0214	-1.3609	0.0426	-1.1907	0.0007	0.2358	0.4979	0.9231	0.0010	0.6874	0.0021	-0.1260	0.8814	-0.0569	0.9084	0.0691	0.8739
beat-IIIb	beat-IIIb	1626302_at	0.2164	0.1799	0.0479	0.6977	-0.1276	0.6558	-0.0324	0.9592	-0.0175	0.9418	0.0149	0.9425	0.1326	0.8965	-0.0201	0.9759	-0.1527	0.7239
CG32194	CG32194	1626303_s_at	-1.4982	0.0006	-1.6099	0.0075	-1.6130	0.0001	-0.0958	0.8327	0.0050	0.9844	0.1008	0.5517	-0.1105	0.9357	-0.0316	0.9648	0.0789	0.8949
CG32446	CG32446	1626304_at	-0.0370	0.8478	0.0182	0.9174	0.0780	0.6858	-0.1327	0.7409	0.0818	0.6860	0.2145	0.1741	-0.0786	0.9238	0.2425	0.3606	0.3210	0.2527
---	---	1626305_at	-0.1787	0.2482	-0.0522	0.6355	0.1152	0.5199	-0.0057	0.9951	-0.2094	0.2656	-0.2037	0.2246	-0.0660	0.8909	0.0181	0.9505	0.0841	0.6711
rok	RHO kinase	1626306_at	-0.6309	0.1116	-0.0514	0.9561	0.7070	0.0050	0.1570	0.8633	-0.5698	0.0803	-0.7268	0.0204	-0.5486	0.7697	0.1360	0.9115	0.6846	0.3994
CG8958	CG8958	1626307_at	0.1031	0.5648	-0.0842	0.4524	0.0769	0.7056	-0.0559	0.9314	0.1361	0.5056	0.1920	0.2715	-0.1010	0.8875	-0.0326	0.9402	0.0684	0.8418
CG4019	CG4019	1626308_s_at	2.7975	0.0013	1.3163	0.2008	3.3645	0.0001	1.1167	0.1495	0.5957	0.2016	-0.5210	0.2136	-0.9222	0.7305	-0.7351	0.5397	0.1871	0.9057
CG6004	CG6004	1626309_at	0.6114	0.1651	-0.1208	0.4134	0.6135	0.1142	0.2957	0.3837	0.3229	0.0933	0.0271	0.9074	-0.2137	0.9101	-0.3333	0.6483	-0.1197	0.8972
---	---	1626310_at	-0.1555	0.6186	0.0451	0.8270	0.0230	0.9180	0.0075	0.9954	0.0159	0.9687	0.0084	0.9798	0.0153	0.9862	-0.0033	0.9941	-0.0186	0.9486
CG15088	CG15088	1626311_at	0.4223	0.1522	0.0067	0.9738	-0.0719	0.7558	-0.1082	0.8856	0.0716	0.8226	0.1797	0.4566	-0.0340	0.9869	-0.4346	0.3485	-0.4006	0.4078
---	---	1626312_at	0.1392	0.4041	0.0104	0.9218	0.2561	0.1344	0.2391	0.4141	0.1823	0.2644	-0.0568	0.7469	-0.2002	0.7215	-0.1408	0.5697	0.0594	0.8435
crl	courtless	1626313_at	-0.1242	0.4180	0.7518	0.0097	0.8618	0.0017	0.1299	0.7031	-0.5965	0.0033	-0.7264	0.0008	-0.1030	0.8446	0.1375	0.5415	0.2405	0.2816
vib	vibrator	1626314_at	1.6918	0.0017	0.8073	0.0761	0.4099	0.0655	0.1031	0.8469	0.9885	0.0010	0.8854	0.0009	0.5736	0.6695	0.2680	0.6497	-0.3056	0.6005
l(1)G0003	lethal (1) G0003	1626315_at	-1.1152	0.0033	-0.4098	0.5132	0.0014	0.9972	0.0292	0.9749	-0.9097	0.0027	-0.9389	0.0014	-0.2537	0.8903	-0.0500	0.9648	0.2037	0.8114
CG34348	CG11757	1626316_at	0.0343	0.9413	1.2286	0.0318	0.5430	0.0915	-0.8548	0.1883	-0.9642	0.0226	-0.1093	0.8047	-0.0166	0.9929	0.2874	0.5338	0.3041	0.5117
---	---	1626317_at	-0.2144	0.2748	0.0600	0.6354	0.2721	0.2548	0.1854	0.6822	-0.1813	0.4025	-0.3667	0.0557	-0.2729	0.5681	-0.0694	0.8000	0.2035	0.3583
---	---	1626318_at	0.0451	0.8721	0.0581	0.6229	-0.0928	0.6300	-0.0229	0.9744	-0.1316	0.4652	-0.1087	0.5155	0.0011	0.9996	-0.2155	0.3760	-0.2166	0.3921
IM10	Immune induced r	1626319_a_at	0.3599	0.5431	-1.2943	0.0265	-0.8015	0.1185	0.6067	0.5362	2.4197	0.0013	1.8130	0.0027	-0.0448	0.9913	0.4964	0.5808	0.5412	0.5457
CG31344	CG31344	1626320_at	0.0287	0.9045	-0.5467	0.0277	-0.4080	0.3354	0.2577	0.5317	1.3293	0.0004	1.0717	0.0006	0.0526	0.9829	0.6372	0.2424	0.5845	0.3097
---	---	1626321_at	0.1903	0.2052	0.0041	0.9722	-0.0626	0.7050	0.0273	0.9610	0.3865	0.0181	0.3592	0.0154	0.1798	0.7190	0.1384	0.5191	-0.0414	0.8844
---	---	1626322_at	0.3007	0.1970	-0.0498	0.7713	-0.1065	0.5104	-0.1370	0.7764	0.0856	0.7143	0.2226	0.2177	-0.0867	0.8903	-0.2613	0.2507	-0.1746	0.4746
CG17255	CG17255	1626323_s_at	-0.3262	0.7911	-0.1682	0.9253	-1.0110	0.0085	-0.2775	0.5988	0.6649	0.0207	0.9424	0.0027	0.7294	0.9030	0.8518	0.7383	0.1225	0.9709
Cyp309a1	Cyp309a1	1626324_at	2.2320	0.0169	0.6424	0.5015	1.4387	0.0466	0.6028	0.5068	0.2786	0.5963	-0.3242	0.4805	-0.0588	0.9928	-1.2467	0.4172	-1.1879	0.4569
CG7849 /// DmirCG7849	CG7849	1626325_a_at	-0.6705	0.0138	-0.3094	0.3823	-0.5704	0.0511	-0.0561	0.9436	-0.1151	0.6496	-0.0590	0.8169	0.0118	0.9952	-0.0326	0.9652	-0.0444	0.9443
CG31436	CG31436	1626326_at	0.8537	0.0111	-0.1392	0.3968	0.0864	0.6384	0.1538	0.6168	0.3220	0.0422	0.1682	0.2207	-0.0469	0.9771	-0.5314	0.1553	-0.4844	0.2237
---	---	1626327_s_at	0.1284	0.3634	0.1564	0.4529	-0.0557	0.7657	-0.0136	0.9838	0.0322	0.8733	0.0458	0.7880	0.1960	0.7506	0.1527	0.5855	-0.0433	0.9057
CG32259	CG32259	1626328_at	0.0641	0.7573	0.1869	0.1691	-0.0531	0.7662	-0.3269	0.2876	-0.1982	0.2757	0.1287	0.4513	0.0025	0.9990	0.0012	0.9994	-0.0013	0.9981
Octbeta3R	Octbeta3R	1626329_at	0.0152	0.9700	0.0278	0.8880	-0.0757	0.6958	-0.0450	0.9393	0.0505	0.8128	0.0955	0.5779	-0.1279	0.8479	-0.1393	0.6429	-0.0114	0.9804
---	---	1626330_at	0.2965	0.2408	-0.0981	0.4394	-0.0401	0.8372	0.2495	0.4245	0.2600	0.1307	0.0105	0.9611	0.1507	0.8378	0.0022	0.9986	-0.1486	0.6476
CG7712	CG7712	1626331_at	-0.1647	0.5381	0.1926	0.5054	0.3171	0.1140	-0.0379	0.9613	-0.6732	0.0069	-0.6353	0.0053	-0.1034	0.9416	-0.3491	0.4214	-0.2458	0.6000
CG11486	polyA-nuclease re	1626332_s_at	1.1042	0.0202	0.6718	0.1893	0.4914	0.1757	-0.2025	0.6822	-0.0239	0.9436	0.1786	0.4053	0.0491	0.9898	-0.3397	0.7151	-0.3888	0.6604

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1626333_at	0.0904	0.6658	0.1882	0.2124	0.1589	0.4672	-0.0298	0.9602	-0.0725	0.6869	-0.0426	0.8091	0.2298	0.7893	0.3285	0.3578	0.0987	0.8299
CG6628	CG6628	1626334_at	0.0569	0.7505	0.1156	0.2926	0.0599	0.7032	-0.1079	0.8038	-0.1317	0.4680	-0.0238	0.9099	-0.0354	0.9640	-0.0249	0.9420	0.0105	0.9752
---	---	1626335_at	-0.0426	0.7970	0.0999	0.4598	0.1375	0.4484	0.0527	0.9496	-0.0773	0.7841	-0.1300	0.5733	-0.0553	0.9467	0.0365	0.9277	0.0918	0.7537
Dnr1	Defense represso	1626336_at	-1.6407	0.1103	0.5890	0.5303	-0.0740	0.8522	-0.5977	0.1341	-2.1869	0.0001	-1.5893	0.0002	0.1268	0.9853	0.0796	0.9778	-0.0472	0.9849
Eap	Exu-associated pr	1626337_at	-0.0274	0.9583	0.4026	0.3401	0.4847	0.0796	0.2763	0.7409	0.1687	0.6896	-0.1076	0.7946	0.0609	0.9816	0.5889	0.3009	0.5280	0.3803
---	---	1626338_s_at	-0.0404	0.7829	-0.0796	0.5040	-0.1244	0.4508	0.1342	0.7857	0.0002	0.9996	-0.1341	0.4926	0.0367	0.9514	0.0000	1.0000	-0.0367	0.8788
garz	gartenzwerg	1626339_at	0.5403	0.2988	0.9406	0.0486	1.1141	0.0034	0.2607	0.6387	0.2485	0.3705	-0.0123	0.9711	0.0600	0.9831	0.7157	0.2518	0.6557	0.3227
betaCop	beta-coatomer prc	1626340_at	0.7678	0.1654	0.5578	0.2236	0.8006	0.0018	0.3460	0.4861	0.7214	0.0146	0.3753	0.1131	0.0823	0.9816	0.6831	0.3963	0.6009	0.4757
CG14122	CG14122	1626341_at	-1.9830	0.0040	-0.7237	0.0265	-1.2799	0.0020	-0.2123	0.7082	-0.8994	0.0048	-0.6871	0.0098	0.1907	0.8603	0.0394	0.9557	-0.1513	0.7654
CG12115	CG12115	1626342_at	0.1900	0.4262	0.1708	0.2423	0.1792	0.2647	0.0404	0.9620	-0.1897	0.4144	-0.2301	0.2563	0.0184	0.9848	-0.1265	0.5929	-0.1449	0.5374
CG15874	CG15874	1626343_at	-0.1434	0.5590	-0.0597	0.6811	0.1789	0.5961	-0.0696	0.9396	-0.0981	0.7585	-0.0284	0.9308	-0.0080	0.9963	0.0439	0.9420	0.0519	0.9197
CG14980	CG14980	1626344_at	-0.2015	0.5610	-0.0298	0.9524	0.0836	0.6470	0.0665	0.9185	-0.3126	0.1162	-0.3791	0.0401	-0.1815	0.8749	-0.0762	0.9130	0.1053	0.8591
IM4	Immune induced r	1626345_at	1.9370	0.0139	0.6931	0.4841	1.5186	0.0036	0.6296	0.5060	0.7443	0.1362	0.1146	0.8431	0.0398	0.9936	-0.2417	0.8792	-0.2815	0.8432
---	---	1626346_at	0.3827	0.0635	0.4533	0.0745	0.0935	0.5663	-0.0685	0.8791	-0.0490	0.7948	0.0195	0.9175	0.1538	0.8283	0.0218	0.9647	-0.1320	0.6877
CG15877 /// DyakCG15877	CG15877	1626347_at	-0.1114	0.6390	-0.3003	0.4248	-0.0233	0.9188	0.1045	0.8981	0.3964	0.1372	0.2918	0.2239	-0.0716	0.9568	0.1917	0.6404	0.2633	0.5011
---	---	1626348_at	0.1396	0.5039	0.0171	0.9081	0.1417	0.5841	0.0738	0.9325	0.1942	0.4723	0.1204	0.6519	-0.0254	0.9816	0.1074	0.6794	0.1328	0.5972
alpha-Spec	alpha-Spectrin	1626349_at	0.0552	0.9152	-0.1091	0.8592	-0.3871	0.0758	0.3275	0.5454	0.7912	0.0123	0.4637	0.0676	0.7079	0.7196	0.6869	0.3951	-0.0210	0.9874
alpha-Est10	fragment K	1626350_at	1.0700	0.0461	1.4057	0.0554	2.6027	0.0012	0.2732	0.6312	-0.3797	0.1710	-0.6528	0.0169	-0.8524	0.7387	0.0510	0.9796	0.9034	0.4114
Tsp42Ep	tetraspanin 42E	1626351_at	0.3259	0.1561	0.2604	0.4006	0.4376	0.1057	0.1711	0.6380	0.1687	0.3528	-0.0024	0.9913	0.0324	0.9848	0.0313	0.9617	-0.0011	0.9990
---	---	1626352_at	0.0194	0.9732	-0.5260	0.3113	-0.7868	0.0478	-0.1222	0.9110	0.4892	0.1564	0.6113	0.0530	0.1711	0.9467	0.0333	0.9821	-0.1377	0.8984
CG15738	CG15738	1626353_at	0.2052	0.3188	-0.0480	0.7239	0.0545	0.7445	-0.0724	0.8792	0.0172	0.9418	0.0895	0.5804	-0.1296	0.8122	-0.1789	0.4280	-0.0494	0.8699
---	---	1626354_at	0.1630	0.2585	0.0836	0.5811	0.0733	0.7441	0.1547	0.6577	0.1630	0.3386	0.0084	0.9687	0.2047	0.7266	0.1081	0.7056	-0.0966	0.7411
DNApol-epsilon	DNA polymerase	1626355_at	0.2224	0.6470	-1.0171	0.1252	-1.1243	0.0425	-0.4414	0.5106	1.2324	0.0048	1.6738	0.0008	-0.4535	0.8454	-0.1621	0.9120	0.2914	0.8023
CG15316 /// DsimCG15316	CG15316	1626356_s_at	-1.7463	0.0538	-2.6011	0.0878	-1.7496	0.0001	-0.2795	0.6864	0.3753	0.2442	0.6549	0.0303	-1.1181	0.7370	-0.4785	0.7866	0.6396	0.6829
---	---	1626357_at	0.1304	0.3609	0.3024	0.2278	0.3228	0.2306	0.0649	0.9220	-0.0095	0.9755	-0.0744	0.7276	0.0830	0.9092	0.0495	0.8992	-0.0335	0.9250
Fer1	48 related 1	1626358_at	0.0028	0.9898	0.2604	0.2769	0.0690	0.7050	-0.0584	0.9293	-0.0867	0.6956	-0.0283	0.9028	0.1130	0.8815	0.0427	0.9263	-0.0703	0.8512
CG11403 /// DereCG11403	CG11403	1626359_at	-0.3902	0.0701	-0.8225	0.0431	-0.6877	0.0034	0.2308	0.4568	0.7098	0.0021	0.4790	0.0071	-0.1532	0.8564	-0.0228	0.9682	0.1304	0.7442
CG4936	CG4936	1626360_at	0.0997	0.5139	0.0622	0.7160	0.5265	0.0086	-0.0395	0.9436	-0.1511	0.3539	-0.1116	0.4616	-0.2841	0.6145	0.0236	0.9508	0.3077	0.2158
bol	boule	1626361_at	-0.0681	0.6896	-0.0092	0.9318	-0.0110	0.9733	0.0019	0.9981	0.0748	0.7322	0.0730	0.7129	-0.1442	0.8732	-0.0155	0.9814	0.1287	0.7531
mio	missing oocyte	1626362_at	0.4656	0.0609	0.7670	0.0154	0.8192	0.0070	-0.0336	0.9518	-0.3660	0.0251	-0.3324	0.0235	-0.1465	0.8609	-0.1650	0.6520	-0.0185	0.9716
---	---	1626363_at	0.2702	0.3553	-0.5090	0.1462	-1.1319	0.0159	-0.4068	0.5848	0.5138	0.1737	0.9206	0.0147	-0.1079	0.9527	-0.2891	0.6209	-0.1812	0.7774
CG13000	CG13000	1626364_at	0.2684	0.1808	0.0538	0.6084	0.0541	0.7840	-0.2054	0.7042	-0.1252	0.6556	0.0803	0.7701	-0.0670	0.8960	-0.2703	0.1489	-0.2033	0.2993
CG33322	CG33322	1626365_at	0.1880	0.2024	0.1696	0.4072	0.1734	0.2228	0.0075	0.9937	0.0584	0.8031	0.0509	0.8134	0.0959	0.8603	0.1254	0.5890	0.0295	0.9220
ppl	pumpless	1626366_at	1.5894	0.0016	0.6269	0.2783	1.4625	0.0001	0.4650	0.2208	0.2504	0.2733	-0.2146	0.2979	-0.3844	0.7810	-0.6716	0.2430	-0.2872	0.6569
Sgt	small glutamine-ri	1626367_at	0.1300	0.5205	0.5915	0.0344	0.8429	0.0070	0.2380	0.6704	-0.1318	0.6610	-0.3698	0.1181	-0.0182	0.9893	0.3085	0.2518	0.3267	0.2597
lace	Serine palmitoyltra	1626368_at	0.0911	0.7507	-0.0805	0.7863	-0.2114	0.2701	0.1840	0.6993	0.7355	0.0056	0.5515	0.0124	0.4334	0.6749	0.6065	0.1395	0.1731	0.7128
CG12269	CG12269	1626369_at	0.2082	0.2024	0.0661	0.5422	0.2352	0.1773	0.0281	0.9633	0.0666	0.7247	0.0385	0.8358	-0.0296	0.9774	-0.0565	0.8721	-0.0269	0.9377
CG31120	CG31120	1626370_at	0.1634	0.4135	0.5954	0.1133	0.9533	0.0005	-0.0849	0.9060	-0.5704	0.0221	-0.4856	0.0271	-0.2718	0.6904	-0.0115	0.9829	0.2603	0.3683
CG8507	CG8507	1626371_at	-0.3232	0.0942	-0.8581	0.0409	-1.1361	0.0045	-0.0334	0.9602	0.5734	0.0063	0.6068	0.0029	0.2504	0.8424	0.0096	0.9935	-0.2408	0.6677
---	---	1626372_at	-0.0025	0.9936	0.0618	0.6039	0.0450	0.7986	0.0214	0.9735	0.0681	0.7101	0.0467	0.7926	0.0175	0.9884	-0.0371	0.9279	-0.0545	0.8779
CG32833	CG32833	1626373_at	1.2213	0.0196	0.9168	0.0861	2.0544	0.0000	0.2533	0.5008	0.1776	0.3828	-0.0758	0.7204	-0.6615	0.7230	-0.1715	0.8830	0.4900	0.5670
---	---	1626374_at	0.2428	0.1489	0.0817	0.5629	0.0594	0.7630	-0.1583	0.7556	-0.0968	0.7021	0.0615	0.8039	0.0533	0.9333	-0.0398	0.8963	-0.0931	0.6779
Usf	Usf	1626375_a_at	-0.3450	0.1743	-0.4547	0.1068	-0.1347	0.4924	-0.0084	0.9900	0.1869	0.2142	0.1953	0.1463	-0.2732	0.8222	0.2796	0.6058	0.5528	0.2904
CG32133	CG32133	1626376_at	-0.1587	0.4848	0.5636	0.1451	0.6131	0.0190	-0.1664	0.8350	-0.4217	0.1615	-0.2552	0.3591	-0.0965	0.9503	0.4258	0.3416	0.5223	0.2727
CG8358	CG8358	1626377_at	2.7002	0.0010	0.9502	0.1642	2.5708	0.0004	0.8989	0.4239	0.9016	0.1453	0.0027	0.9974	-0.6700	0.6898	-0.8049	0.2254	-0.1348	0.8870
---	---	1626378_at	-0.1995	0.2392	0.1779	0.3382	0.0299	0.8865	-0.0959	0.7979	-0.1986	0.1883	-0.1027	0.4776	-0.0017	0.9992	0.0631	0.8432	0.0647	0.8261
CG17490	CG17490	1626379_a_at	0.4868	0.2130	0.6515	0.2509	0.2121	0.2535	-0.6738	0.2654	-0.3157	0.3898	0.3581	0.2672	-0.0823	0.9683	-0.0963	0.9083	-0.0140	0.9874
---	---	1626380_s_at	0.0932	0.5576	0.0728	0.6452	0.1743	0.4829	0.0352	0.9603	-0.0020	0.9942	-0.0372	0.8639	-0.0835	0.8760	-0.0647	0.8118	0.0188	0.9491
---	---	1626381_at	0.1216	0.4783	-0.0285	0.7767	-0.0051	0.9846	0.0711	0.8942	0.1122	0.5584	0.0410	0.8414	0.0042	0.9967	-0.0711	0.8380	-0.0753	0.8154
CG13887	CG13887	1626382_s_at	0.1862	0.3876	0.0254	0.8459	0.3876	0.0755	0.3148	0.3127	0.4998	0.0140	0.1849	0.2603	-0.1229	0.8270	0.3709	0.1168	0.4938	0.0710
CG4096	CG4096	1626383_at	-0.0907	0.7703	0.0506	0.9209	-0.1061	0.6823	0.0859	0.9413	0.2164	0.5479	0.1305	0.7135	0.0225	0.9914	0.1572	0.7978	0.1348	0.8247
CG12182	CG12182	1626384_at	0.2921	0.0672	0.3270	0.3693	0.2940	0.0933	0.2632	0.5633	0.5436	0.0278	0.2804	0.1789	0.3306	0.6557	0.3268	0.2642	-0.0038	0.9945
wdb	widerborst	1626385_s_at	-0.4538	0.0989	0.2219	0.2905	-0.4550	0.0423	-0.3165	0.1949	-0.2659	0.0740	0.0506	0.7532	0.3103	0.7215	0.3568	0.3173	0.0465	0.9271
Cpr57A	CG18066	1626386_at	-2.4422	0.0194	-0.1144	0.5254	-1.5023	0.0047	-0.9069	0.2592	-3.1678	0.0004	-2.2609	0.0008	0.3323	0.9101	-0.9345	0.3669	-1.2668	0.2454
mei-217 /// mei-218	mei-217 /// transcr	1626387_s_at	-0.2466	0.1966	0.0581	0.8185	0.0924	0.5825	-0.1366	0.7588	-0.0486	0.8442	0.0881	0.6559	-0.0819	0.9126	0.2453	0.3375	0.3272	0.2282
Ctr1B	Copper transporte	1626388_at	0.5566	0.1882	0.3393	0.1587	0.4979	0.0348	-0.0540	0.9592	-0.2619	0.3528	-0.2078	0.4225	-0.0369	0.9862	-0.4100	0.4110	-0.3730	0.4742
---	---	1626389_at	0.0841	0.5774	-0.1064	0.4512	0.1184	0.4172	0.0798	0.8507	0.0671	0.7061	-0.0127	0.9483	-0.1100	0.8461	-0.0869	0.7561	0.0232	0.9434
Ced-12	Ced-12	1626390_at	-0.8086	0.0020	-1.0657	0.0461	-1.1928	0.0005	-0.0048	0.9955	0.2652	0.1151	0.2700	0.0754	0.1653	0.8744	-0.0065	0.9941	-0.1718	0.7120
---	---	1626391_at	0.4091	0.0554	0.1452	0.5329	0.0887	0.7261	-0.0059	0.9951	0.1373	0.4994	0.1432	0.4290	-0.0940	0.9238	-0.1748	0.6144	-0.0808	0.8459
---	---	1626392_s_at	-1.0952	0.0032	-1.6911	0.0065	-1.3431	0.0009	0.0251	0.9838	1.0496	0.0035	1.0245	0.0023	-0.3592	0.6955	0.3781	0.3016	0.7373	0.0859
CG8078	CG8078	1626393_at	0.2703	0.3146	0.1316	0.7331	-0.0233	0.9286	-0.1697	0.7289	0.3624	0.0980	0.5322	0.0146	0.1597	0.8999	0.3064	0.5264	0.1468	0.7940
Jon99Fi	Jonah 99F	1626394_at	1.1293	0.5306	-1.1327	0.0171	-1.9445	0.0091	-1.0227	0.4722	1.0068	0.1858	2.0294	0.0099	-0.3347	0.9701	-1.4820	0.5329	-1.1472	0.6409
---	---	1626395_at	0.1838	0.3853	-0.0459	0.7379	0.0931	0.5323	0.0653	0.9343	0.1003	0.7087	0.0350	0.9003	-0.0664	0.9309	-0.0006	0.9996	0.0657	0.8348
Menl-1 /// Menl-2	Menl-1 /// Menl-2	1626396_s_at	0.3308	0.2016	-0.0122	0.9524	0.1881	0.3304	0.5005	0.2484	0.4279	0.0961	-0.0726	0.8015	0.0074	0.9933	0.0063	0.9877	-0.0011	0.9979
bt	Projectin	1626397_at	-2.1189	0.0111	-2.1163	0.1232	-2.6743	0.0004	-0.3394	0.3863	-0.2929	0.1843	0.0465	0.8572	0.3471	0.9441	-0.2325	0.9207	-0.5796	0.7371
---	---	1626398_at	0.1881	0.1854	0.0367	0.7194	0.0799	0.6766	-0.0065	0.9942	0.0225	0.9309	0.0290	0.8963	0.0529	0.9309	0.0458	0.8718	-0.0072	0.9835
CG5024	CG5024	1626399_at	0.0615	0.8369	-0.0985	0.5985	0.1136	0.6325	0.0252	0.9647	0.0461	0.8031	0.0209	0.9087	-0.2354	0.8097	0.0198	0.9784	0.2552	0.5516
CG30049	CG30049	1626400_at	0.1930	0.4041	0.0091	0.9387	0.0363	0.8218	0.0359	0.9620	0.0784	0.7396	0.0425	0.8549	-0.0854	0.8692	-0.1092	0.6266	-0.0238	0.9347
Cyp6a2	cytochrome P450-	1626401_at	0.7596	0.4592	0.8195	0.6031	1.5559	0.0046	0.5169	0.6468	-1.7059	0.0079	-2.2228	0.0015	-0.1610	0.9852	-1.7570	0.3685	-1.5960	0.4335

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31427	CG31427	1626402_at	0.2045	0.1933	-0.0142	0.9636	-0.0060	0.9841	0.0414	0.9596	0.0557	0.8405	0.0143	0.9582	0.1313	0.8427	-0.0861	0.8053	-0.2175	0.4370
CG40384	CG40384	1626403_at	-0.0016	0.9961	-0.3244	0.1578	-0.1614	0.2719	0.2784	0.2870	0.4221	0.0139	0.1437	0.3028	-0.0166	0.9884	0.0352	0.9279	0.0518	0.8775
bsk	Jun kinase	1626404_a_at	-0.3581	0.2482	1.1992	0.0652	0.9422	0.0006	-0.4191	0.3056	-1.3835	0.0006	-0.9644	0.0016	-0.1873	0.8956	0.2458	0.6772	0.4331	0.4272
Rep1	Rep1	1626405_at	-0.5626	0.3401	-0.0799	0.6709	-0.9134	0.0067	-0.3152	0.5633	0.0996	0.7670	0.4148	0.0974	0.6633	0.6955	0.8684	0.2116	0.2051	0.8183
---	---	1626406_at	0.8087	0.0108	0.6727	0.0775	0.3494	0.0841	0.0381	0.9712	0.3729	0.1619	0.3348	0.1612	-0.0013	0.9998	-0.0314	0.9682	-0.0301	0.9661
CG6523	CG6523	1626407_at	-0.0913	0.5568	0.7550	0.0186	0.8441	0.0034	0.0022	0.9962	-0.6144	0.0023	-0.6166	0.0014	-0.0945	0.9401	0.1353	0.7760	0.2298	0.5781
---	---	1626408_at	0.0631	0.6961	0.0725	0.7519	0.2052	0.2726	0.1087	0.8339	0.0984	0.6576	-0.0103	0.9673	0.0695	0.9514	0.1741	0.6382	0.1046	0.8007
CG11997	CG11997	1626409_at	0.0234	0.9052	0.0219	0.9220	0.0217	0.9043	-0.0186	0.9852	-0.0303	0.9232	-0.0117	0.9677	0.0007	0.9998	0.0211	0.9504	0.0204	0.9440
---	---	1626410_at	-0.1024	0.6235	0.3003	0.3247	-0.0616	0.7759	-0.1046	0.8090	-0.1507	0.3938	-0.0462	0.8124	0.3015	0.6557	0.2480	0.3500	-0.0536	0.8864
l(1)G0193	lethal (1) G0193	1626411_at	-0.0643	0.7028	-0.6904	0.0435	-0.6097	0.0095	-0.2030	0.6354	-0.0686	0.7899	0.1345	0.5079	-0.3164	0.6497	-0.6175	0.0509	-0.3011	0.2982
CG8549	CG8549	1626412_at	0.2095	0.2861	0.0282	0.7831	0.4657	0.0086	0.1332	0.8102	-0.1465	0.5335	-0.2796	0.1544	-0.1640	0.8141	-0.0769	0.8438	0.0871	0.8044
Rbp1-like	Rbp1-like	1626413_at	-0.2061	0.5073	-0.0874	0.8811	-0.2140	0.2151	0.1993	0.6338	0.5459	0.0159	0.3466	0.0622	0.1540	0.9317	0.4393	0.4512	0.2854	0.6484
Jheh1	JH-epoxide hydrol	1626414_at	-0.2483	0.5694	-0.8427	0.0270	-0.4022	0.1381	0.4507	0.1216	-0.4771	0.0145	-0.9278	0.0005	-0.0615	0.9826	-1.2891	0.0661	-1.2275	0.0934
---	---	1626415_at	-0.0399	0.8901	-0.0382	0.7870	0.0308	0.9057	-0.0965	0.8707	0.2884	0.1627	0.3849	0.0440	-0.1362	0.8331	0.1155	0.7040	0.2517	0.3626
l(2)efl	lethal (2) essentia	1626416_a_at	-2.4238	0.0307	-0.6038	0.2596	-0.9943	0.0501	0.0745	0.9704	-1.1173	0.0355	-1.1917	0.0171	0.5615	0.8461	0.8697	0.4696	0.3082	0.8375
Cht3	chitinase-3	1626417_at	-1.0816	0.0041	-1.2619	0.0110	-1.5448	0.0034	-0.2436	0.7511	0.1154	0.7741	0.3589	0.2292	-0.1770	0.8191	-0.1202	0.7459	0.0568	0.8951
CG14691	CG14691	1626418_a_at	-0.0241	0.8875	0.0506	0.7505	0.1645	0.4145	-0.0464	0.9255	-0.0747	0.6586	-0.0283	0.8725	-0.1155	0.9095	0.0283	0.9623	0.1438	0.7287
---	---	1626419_at	0.0701	0.7171	0.0760	0.5339	0.2646	0.1032	0.1325	0.6886	0.0469	0.8030	-0.0856	0.5751	-0.0062	0.9950	0.0371	0.9170	0.0433	0.8915
CG8563	CG8563	1626420_at	0.1345	0.5207	-0.0424	0.7088	0.0223	0.9098	0.0633	0.8830	-0.1039	0.5010	-0.1672	0.2012	-0.0549	0.9426	-0.1985	0.3922	-0.1436	0.5649
CG4662	CG4662	1626421_at	0.3758	0.2674	0.9515	0.1009	0.5510	0.0074	-0.2980	0.6673	-0.6984	0.0410	-0.4005	0.1746	-0.0339	0.9862	-0.0904	0.8893	-0.0565	0.9271
---	---	1626422_at	0.0197	0.9218	0.0901	0.4675	-0.0769	0.7005	-0.1150	0.8233	0.0818	0.7262	0.1967	0.2765	-0.0784	0.9011	-0.0540	0.8684	0.0244	0.9401
---	---	1626423_at	0.2223	0.1866	-0.0139	0.9119	0.0010	0.9968	0.0047	0.9949	-0.1238	0.4190	-0.1284	0.3449	-0.0436	0.9445	-0.0815	0.7027	-0.0378	0.8837
---	---	1626424_at	0.0805	0.7034	0.1883	0.3881	0.1775	0.3253	0.0560	0.9435	-0.0057	0.9869	-0.0617	0.8056	0.0767	0.8991	0.0626	0.8343	-0.0141	0.9661
2-Sep	septin	1626425_at	-0.4477	0.0217	0.3636	0.1960	0.4823	0.0455	0.1980	0.6338	-0.1796	0.3906	-0.3776	0.0443	-0.0494	0.9768	0.5465	0.1643	0.5959	0.1651
CG11155	CG11155	1626426_at	-1.1663	0.0086	-0.9931	0.0122	-1.4151	0.0001	-0.3789	0.1832	-0.3045	0.0792	0.0744	0.6798	0.0992	0.9445	-0.2327	0.6137	-0.3319	0.4558
Tequila	Tequila	1626427_a_at	0.0301	0.8926	0.0675	0.6960	0.1608	0.5444	-0.0019	0.9988	-0.0141	0.9756	-0.0122	0.9736	0.0043	0.9977	-0.1195	0.7634	-0.1238	0.7442
---	---	1626428_s_at	-0.0488	0.8377	0.0168	0.8785	-0.1955	0.4145	-0.1560	0.6720	0.0448	0.8432	0.2008	0.1977	0.1116	0.8541	-0.0015	0.9989	-0.1131	0.6811
Lsp1gamma	Larval serum prot	1626429_at	-0.0030	0.9934	-0.1161	0.7448	-0.4226	0.0212	-0.1100	0.9247	0.0637	0.8931	0.1737	0.6250	0.2942	0.7464	-0.1284	0.7877	-0.4227	0.2833
---	---	1626430_at	0.1919	0.2487	0.1467	0.3972	0.0981	0.6214	0.1154	0.7487	0.0212	0.9228	-0.0942	0.5389	0.2584	0.6955	0.0207	0.9642	-0.2376	0.4016
CG10681	CG10681	1626431_at	0.1892	0.2908	-0.0428	0.8593	-0.2467	0.1816	-0.0465	0.9558	0.3041	0.1752	0.3506	0.0832	0.1809	0.8192	0.0215	0.9682	-0.1594	0.6476
CG10839 /// DereCG10839	CG10839	1626432_at	0.3721	0.0890	0.2648	0.3272	0.3030	0.1764	0.0835	0.9326	-0.0764	0.8395	-0.1600	0.5846	-0.0087	0.9928	-0.0950	0.7000	-0.0863	0.7291
mRpS16	mitochondrial ribo	1626433_at	-0.0057	0.9895	0.3940	0.3083	0.2830	0.2230	-0.0409	0.9518	-0.2005	0.2901	-0.1597	0.3554	0.1890	0.9128	0.2686	0.6945	0.0796	0.9237
HMS-Beaglepol	pol	1626434_s_at	0.1160	0.6042	-0.1066	0.6135	0.0877	0.7191	-0.1058	0.8639	-0.0087	0.9795	0.0971	0.6697	-0.0095	0.9952	-0.0785	0.8824	-0.0690	0.8918
CG15263	CG15263	1626435_at	-0.0182	0.9288	-0.0567	0.6733	0.1019	0.6049	0.0290	0.9730	0.0511	0.8529	0.0221	0.9332	-0.0224	0.9829	0.0060	0.9914	0.0284	0.9330
CG3570	CG3570	1626436_at	0.1174	0.5045	0.4756	0.0369	0.4972	0.0076	-0.0155	0.9825	-0.1710	0.3039	-0.1556	0.2975	0.1381	0.8049	0.2212	0.3309	0.0831	0.7577
lectin-22C	Lectin22C	1626437_at	-2.7984	0.0009	-3.6394	0.0011	-3.2902	0.0000	0.3008	0.7028	0.7610	0.0422	0.4602	0.1548	-0.0742	0.9390	-0.3341	0.2460	-0.2599	0.3921
---	---	1626438_at	-0.2477	0.3686	-0.8965	0.0223	-0.8011	0.0019	-0.0158	0.9879	0.6608	0.0147	0.6766	0.0081	-0.1601	0.7768	0.0433	0.9077	0.2035	0.4114
CG15353	CG15353	1626439_at	-0.2553	0.6404	0.1454	0.8617	-0.6570	0.1691	-0.5737	0.2420	-1.1126	0.0031	-0.5389	0.0437	0.1891	0.9677	-0.7728	0.5439	-0.9619	0.4446
---	---	1626440_at	-0.0493	0.7953	0.0500	0.7428	0.1206	0.5293	0.0864	0.8470	0.0385	0.8576	-0.0479	0.7946	-0.0395	0.9677	-0.0646	0.8550	-0.0251	0.9440
CG5139	CG5139	1626441_at	-0.1703	0.2789	0.0910	0.4959	0.0936	0.6249	0.1156	0.8869	-0.1085	0.7373	-0.2241	0.3776	-0.1471	0.8270	0.0174	0.9698	0.1645	0.5778
CG7224 /// DyakCG7224	CG7224	1626442_at	-0.2777	0.6274	-1.4225	0.0299	-1.5071	0.0296	-0.2656	0.8837	-0.3929	0.5549	-0.1273	0.8599	-0.0249	0.9952	-1.4857	0.1320	-1.4608	0.1612
CG13364	CG13364	1626443_at	0.0540	0.7920	0.5385	0.1051	0.2134	0.2178	0.0315	0.9549	0.0123	0.9562	-0.0192	0.9172	0.2861	0.7215	0.3793	0.2408	0.0932	0.8246
Fib	Fibrillarin	1626444_at	0.3571	0.4704	-0.2070	0.4327	0.2981	0.5101	0.7029	0.3925	1.4107	0.0088	0.7078	0.0878	0.2474	0.8905	0.8471	0.1957	0.5998	0.3855
---	---	1626445_at	0.2211	0.3244	0.0414	0.7020	0.0822	0.6320	0.0479	0.9461	0.0610	0.8038	0.0131	0.9589	-0.0715	0.8937	0.0031	0.9941	0.0745	0.7442
Mlc1	myosin light chain	1626446_a_at	-1.4187	0.0735	-1.2204	0.1069	-2.0654	0.0018	-0.3413	0.5361	-1.1393	0.0028	-0.7980	0.0081	0.5653	0.8521	-1.0185	0.4131	-1.5839	0.2230
CG10508	CG10508	1626447_at	-0.2355	0.3463	0.0286	0.9233	-0.1191	0.5249	-0.1599	0.8132	-0.1724	0.5479	-0.0126	0.9700	0.2455	0.7628	0.3494	0.2889	0.1039	0.8009
Spt5	Spt5	1626448_at	0.4154	0.0958	0.7091	0.0884	0.6550	0.0029	0.2026	0.6010	0.3547	0.0723	0.1520	0.4036	0.2164	0.8292	0.6067	0.1418	0.3903	0.3671
exba	krasavietz	1626449_s_at	0.0713	0.7570	-0.1597	0.5841	0.0829	0.6530	0.0549	0.9216	0.2584	0.1221	0.2036	0.1744	-0.0437	0.9754	0.1455	0.7127	0.1893	0.6141
---	---	1626450_at	-0.0547	0.7832	0.1944	0.3510	0.1749	0.2769	-0.0363	0.9592	-0.1568	0.4118	-0.1205	0.4987	0.0566	0.9479	0.1189	0.6839	0.0624	0.8545
CG13847	CG13847	1626451_at	0.2609	0.1558	0.0978	0.5451	0.1441	0.4367	0.0116	0.9880	0.1003	0.6144	0.0887	0.6300	0.0204	0.9860	0.0462	0.9075	0.0258	0.9433
CG2121	CG2121	1626452_at	3.8308	0.0012	2.4336	0.0203	3.1738	0.0013	0.5401	0.3298	1.1187	0.0048	0.5786	0.0488	-0.0555	0.9923	0.0183	0.9949	0.0738	0.9712
---	---	1626453_x_at	-0.6086	0.0920	-0.8287	0.1312	-1.1547	0.0014	-0.2783	0.6785	0.4597	0.1418	0.7380	0.0167	-0.0210	0.9914	0.1751	0.7413	0.1961	0.6970
CycB3	cyclin B3	1626454_at	0.0683	0.9508	-0.8894	0.4762	-1.7666	0.0738	-0.3113	0.7415	1.6263	0.0029	1.9377	0.0008	0.1089	0.9914	0.2363	0.9486	0.1274	0.9704
CG34005	CG34005	1626455_s_at	0.1258	0.8892	0.0849	0.5189	-0.0395	0.8148	-0.1198	0.9336	-0.2374	0.6100	-0.1176	0.8046	-0.1136	0.9672	-0.2385	0.7943	-0.1249	0.9017
l(1)G0320	lethal (1) G0320	1626456_at	1.9799	0.0035	3.4176	0.0035	2.9544	0.0001	0.3309	0.5680	0.6529	0.0351	0.3219	0.2267	0.7678	0.6955	1.9566	0.0391	1.1888	0.1701
CG31163	CG31163	1626457_s_at	0.3732	0.4799	-0.2234	0.3876	-0.5517	0.0129	-0.3438	0.6247	0.2249	0.5480	0.5687	0.0702	-0.1766	0.9084	-0.6796	0.1960	-0.5030	0.3678
Sirt2	Sirt2	1626458_at	-0.2648	0.1005	-0.0393	0.9067	0.4418	0.0154	0.1226	0.7556	-0.0876	0.6468	-0.2102	0.1658	-0.3627	0.6256	0.1406	0.6834	0.5033	0.1326
---	---	1626459_at	0.0540	0.8314	0.0893	0.4258	0.1395	0.4764	-0.0031	0.9959	-0.0204	0.9385	-0.0174	0.9395	0.0140	0.9923	0.0754	0.8737	0.0614	0.8926
CG2658	paraplegin_DROM	1626460_at	-0.4163	0.1603	-0.1873	0.7672	0.2489	0.2792	0.3125	0.5311	0.0586	0.8646	-0.2539	0.2807	-0.1173	0.9619	0.2630	0.7439	0.3803	0.6052
Ncc69	sodium chloride c	1626461_at	-0.5799	0.3210	0.5700	0.0499	0.5430	0.0541	-0.1985	0.7647	-1.2889	0.0015	-1.0904	0.0019	-0.2511	0.9071	-0.2140	0.8352	0.0371	0.9760
CG7691	CG7691	1626462_at	0.0967	0.6187	0.1751	0.2443	-0.0994	0.6138	-0.0698	0.8817	-0.0409	0.8437	0.0289	0.8791	0.0911	0.9138	0.1397	0.6673	0.0485	0.9055
---	---	1626463_at	0.0020	0.9955	0.0704	0.5373	0.3388	0.1638	0.0505	0.9424	-0.1161	0.5926	-0.1666	0.3639	-0.0918	0.9412	-0.0806	0.8852	0.0113	0.9852
---	---	1626464_at	-0.0799	0.6758	0.0798	0.6025	0.0667	0.6695	-0.0518	0.9311	-0.1198	0.5207	-0.0680	0.7151	0.0587	0.9340	0.1090	0.6604	0.0504	0.8667
hep	hemipterous	1626465_a_at	-0.3281	0.5320	0.0995	0.9119	0.1744	0.4839	0.1254	0.8289	0.0414	0.8907	-0.0840	0.7236	0.0148	0.9970	0.4739	0.6548	0.4591	0.6630
---	---	1626466_at	0.2380	0.1604	0.0751	0.6028	0.2869	0.0892	0.0626	0.8844	0.0003	0.9987	-0.0623	0.6854	-0.1449	0.7681	-0.1726	0.4032	-0.0277	0.9231
Obp19c	Odorant-binding p	1626467_a_at	-0.0513	0.8061	0.3770	0.0879	0.0407	0.9042	-0.1658	0.8268	-0.2072	0.5010	-0.0414	0.9070	0.0384	0.9812	-0.0297	0.9590	-0.0681	0.8870
KCNQ	KCNQ potassium	1626468_a_at	-1.8495	0.0025	-1.9805	0.0185	-1.3580	0.0005	0.2526	0.7023	-0.1681	0.6162	-0.4207	0.1196	-0.3261	0.8202	-0.1484	0.8546	0.1778	0.8044
CG8235	CG8235	1626469_at	0.0585	0.7442	0.3556	0.1930	0.4606	0.0571	-0.0590	0.9098	-0.1685	0.3125	-0.1096	0.4884	-0.0887	0.9467	0.2154	0.6209	0.3041	0.4683
CG10440	CG10440	1626470_at	-1.8387	0.0087	-0.4115	0.2212	-0.7670	0.0736	-0.3002	0.5931	-1.1225	0.0028	-0.8223	0.0067	0.2511	0.9142	0.3361	0.7200	0.0850	0.9411

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG34353	CG12274	1626471_at	-0.0005	0.9977	0.0881	0.5561	0.2483	0.2129	0.0876	0.8776	-0.2272	0.2457	-0.3148	0.0734	0.0967	0.9025	0.0717	0.8561	-0.0250	0.9510
CG30340	CG30340	1626472_at	0.0704	0.6682	-0.1561	0.3377	0.2067	0.3224	0.1737	0.7596	0.0704	0.8157	-0.1033	0.6857	-0.1712	0.7893	-0.1963	0.4794	-0.0251	0.9473
fon	CG15825	1626473_a_at	1.9487	0.0021	0.4132	0.5448	1.8216	0.0011	0.7865	0.2802	1.3427	0.0078	0.5562	0.1460	-0.6186	0.7230	-0.3311	0.6969	0.2875	0.7419
---	---	1626474_at	0.3596	0.0679	-0.1125	0.4759	-0.1324	0.4592	0.1283	0.7949	0.3440	0.0882	0.2157	0.2314	-0.1001	0.8889	-0.1173	0.7007	-0.0172	0.9660
CG6729	CG6729	1626475_at	0.0237	0.9441	0.0333	0.8982	-0.0104	0.9693	0.2297	0.5317	0.2878	0.1323	0.0581	0.7868	0.2343	0.8661	0.2069	0.7541	-0.0274	0.9754
CG31457	CG31457	1626476_at	-0.0098	0.9560	-0.0690	0.5763	-0.2853	0.1479	0.1905	0.5334	0.2247	0.1575	0.0342	0.8540	0.2216	0.6955	0.0987	0.7158	-0.1229	0.6311
CG7236	CG7236	1626477_at	-0.3362	0.0989	0.1496	0.3384	0.0954	0.7099	-0.0079	0.9956	-0.4380	0.1407	-0.4301	0.1063	0.1207	0.8202	0.1243	0.5917	0.0037	0.9924
CG12340	CG12340	1626478_at	-0.1848	0.6258	0.5940	0.1141	0.2807	0.1032	-0.2198	0.6564	0.2614	0.2733	0.4812	0.0310	0.1808	0.8999	1.0128	0.0714	0.8320	0.1438
Rapgap1	Rapgap1	1626479_at	-0.8272	0.0103	-2.5105	0.0030	-2.4339	0.0002	-0.0619	0.9610	1.7707	0.0009	1.8326	0.0005	0.0322	0.9778	0.0506	0.9046	0.0185	0.9637
---	---	1626480_s_at	0.1520	0.3251	0.2422	0.1012	0.0287	0.8986	-0.1949	0.5415	-0.1492	0.3792	0.0456	0.8064	0.0324	0.9774	0.0232	0.9587	-0.0093	0.9838
tafazzin	tafazzins	1626481_a_at	0.5224	0.0752	0.8811	0.0225	0.9499	0.0060	-0.0769	0.9011	-0.7865	0.0029	-0.7096	0.0027	-0.0763	0.9459	-0.2102	0.5542	-0.1339	0.7307
CG11227	CG11227	1626482_a_at	-0.0623	0.7865	0.0546	0.6226	0.1356	0.6479	-0.2684	0.6338	-0.3319	0.2300	-0.0635	0.8425	-0.1545	0.8270	-0.0415	0.9279	0.1129	0.7371
Gl	dynactin	1626483_at	-0.4294	0.1493	-0.2242	0.4135	0.0786	0.5994	0.2467	0.5311	0.1585	0.4582	-0.0882	0.6814	-0.0596	0.9710	0.3469	0.3941	0.4065	0.3360
---	---	1626484_at	0.1461	0.3101	0.0142	0.8936	0.3942	0.0723	0.0597	0.9072	-0.0841	0.6507	-0.1438	0.3418	-0.0415	0.9672	-0.0031	0.9961	0.0384	0.9168
sad	shadow	1626485_at	0.1846	0.2840	0.0777	0.6029	0.2152	0.4042	0.0534	0.9218	-0.0251	0.9127	-0.0786	0.6418	-0.0201	0.9848	-0.0118	0.9784	0.0083	0.9841
CG31815	CG31815	1626486_at	0.2233	0.2725	-0.0193	0.8874	0.1691	0.3096	0.3067	0.2870	0.3637	0.0382	0.0569	0.7583	0.0169	0.9870	0.0042	0.9935	-0.0127	0.9726
---	---	1626487_at	0.0195	0.9102	-0.2426	0.2724	-0.1798	0.2145	0.0309	0.9586	0.3320	0.0408	0.3011	0.0385	0.0083	0.9935	-0.0246	0.9482	-0.0330	0.9181
Rpn9	Nobody	1626488_s_at	0.0923	0.6091	0.5161	0.0929	0.9290	0.0071	-0.0100	0.9922	-0.2031	0.3531	-0.1931	0.3244	-0.3202	0.7220	0.3968	0.2833	0.7170	0.0959
CG7433	CG7433	1626489_s_at	-0.0262	0.9226	-0.0882	0.5036	-0.0271	0.8883	0.0534	0.9098	-0.0420	0.8226	-0.0954	0.5098	-0.0300	0.9806	-0.1793	0.5192	-0.1493	0.6052
CG14608	CG14608	1626490_s_at	-0.0344	0.8542	0.0441	0.7423	0.1088	0.4795	0.1026	0.7582	0.0658	0.6883	-0.0368	0.8212	-0.0237	0.9852	0.0534	0.9058	0.0772	0.8362
---	---	1626491_at	0.0839	0.7026	-0.0761	0.6769	-0.1497	0.3254	-0.2178	0.5932	0.1982	0.3466	0.4161	0.0319	-0.0848	0.9514	0.0092	0.9925	0.0940	0.8621
Sop2	Suppressor of pro	1626492_s_at	-0.8953	0.0041	-0.1291	0.7891	0.2633	0.6029	0.3152	0.3706	-0.1447	0.4881	-0.4599	0.0179	-0.0725	0.9848	0.6492	0.4614	0.7216	0.4167
---	---	1626493_at	0.2148	0.3785	0.1270	0.4025	-0.0005	0.9982	0.0406	0.9435	0.0078	0.9750	-0.0328	0.8619	0.0269	0.9814	0.0304	0.9390	0.0035	0.9935
DNApol-eta	DNApol-eta	1626494_at	-0.1615	0.8294	-0.7624	0.2913	-1.3143	0.0146	-0.4129	0.4767	0.7992	0.0185	1.2121	0.0018	0.1362	0.9775	0.1645	0.9279	0.0283	0.9875
---	---	1626495_at	-0.0317	0.9120	0.0364	0.8120	-0.2536	0.2985	0.0697	0.9461	0.5124	0.0798	0.4427	0.0891	0.2495	0.7697	0.3707	0.2938	0.1212	0.7767
CG10348	CG10348	1626496_at	0.1663	0.3122	0.0891	0.3986	0.1810	0.2564	-0.0757	0.9215	-0.1410	0.5729	-0.0653	0.8019	0.0316	0.9717	-0.0458	0.8875	-0.0775	0.7663
mRpS6	mitochondrial ribo	1626497_at	-0.0008	0.9966	0.5426	0.0440	0.4129	0.1065	-0.0629	0.9036	-0.2857	0.0913	-0.2228	0.1383	0.1742	0.8222	0.2900	0.3612	0.1158	0.7560
CG12612	CG12612	1626498_at	0.1123	0.5932	0.3115	0.0581	-0.0357	0.8304	-0.1010	0.8000	-0.1916	0.2324	-0.0906	0.5693	0.2016	0.7220	0.0978	0.7248	-0.1037	0.7007
GATAe	GATAe	1626499_at	-0.0213	0.9838	0.1795	0.1546	-0.0475	0.7992	-0.1410	0.9403	-0.3125	0.5977	-0.1715	0.7751	0.2500	0.8804	-0.0747	0.9417	-0.3247	0.6350
CTCF	CTCF	1626500_a_at	0.3903	0.2439	0.0684	0.6122	-0.0948	0.6395	0.1001	0.8327	0.2820	0.1161	0.1819	0.2620	0.0926	0.9309	0.0654	0.9009	-0.0272	0.9565
Or85a	Odorant receptor	1626501_at	-0.0033	0.9895	0.0294	0.7699	-0.1890	0.2721	-0.0668	0.9182	-0.0845	0.7157	-0.0177	0.9438	0.1151	0.8842	0.0238	0.9620	-0.0914	0.8025
bcd /// CG14578	CG14578 /// Bicoi	1626502_s_at	0.1783	0.4474	-0.7366	0.0804	-0.6891	0.0427	0.5959	0.4166	1.4892	0.0039	0.8933	0.0223	0.3070	0.6695	0.3437	0.2205	0.0367	0.9273
CG2254 /// DyakCG2254	CG2254	1626503_at	1.5960	0.0586	1.8726	0.1894	2.3220	0.0010	-0.1318	0.9397	-1.1575	0.0296	-1.0257	0.0308	-0.6595	0.8650	-1.0353	0.5176	-0.3758	0.8507
prom	CG30164	1626504_at	-0.0832	0.6330	0.2668	0.2322	0.1310	0.4678	-0.1454	0.7768	-0.2709	0.2046	-0.1255	0.5509	-0.0117	0.9914	-0.0139	0.9719	-0.0022	0.9956
---	---	1626505_at	0.3522	0.2759	-0.0242	0.9428	0.1902	0.2416	0.0831	0.8903	0.0026	0.9929	-0.0805	0.7064	0.2355	0.7644	0.1284	0.7439	-0.1071	0.7891
Syx18	syntaxin	1626506_at	0.8465	0.0141	0.9169	0.0132	0.8141	0.0008	0.1786	0.6493	0.8646	0.0015	0.6861	0.0024	0.3584	0.6706	1.0473	0.0222	0.6889	0.0810
CG7556	CG7556	1626507_at	0.2512	0.4873	1.0190	0.1230	1.3058	0.0002	0.2330	0.4786	-0.4407	0.0207	-0.6738	0.0019	-0.1267	0.9637	0.3960	0.6279	0.5227	0.5058
CG32428	CG32428	1626508_at	-0.0194	0.9040	0.5021	0.1026	0.4115	0.1015	-0.0640	0.9255	-0.7837	0.0033	-0.7197	0.0028	0.0407	0.9796	-0.2515	0.4972	-0.2922	0.4296
18w	18 Wheeler	1626509_at	-0.0183	0.9761	-1.6176	0.0075	-1.1921	0.0051	0.3278	0.6856	1.1790	0.0078	0.8511	0.0197	-0.0372	0.9898	-0.5006	0.4280	-0.4634	0.4799
---	---	1626510_at	0.0910	0.5744	0.0605	0.5432	0.0342	0.8690	-0.0314	0.9695	0.0718	0.7795	0.1032	0.6338	0.0639	0.9016	0.0431	0.8736	-0.0208	0.9372
CG32305	CG32305	1626511_at	0.1159	0.5045	0.1002	0.4025	-0.0313	0.8979	-0.1495	0.6932	-0.0229	0.9252	0.1266	0.4420	-0.0050	0.9952	-0.1195	0.5972	-0.1145	0.6166
---	---	1626512_at	0.0550	0.8054	0.0471	0.6545	0.1305	0.5592	-0.0255	0.9649	-0.0353	0.8610	-0.0098	0.9594	-0.1058	0.8609	0.0278	0.9445	0.1336	0.6130
MESR6	Misexpression su	1626513_at	-0.0319	0.9144	0.2290	0.4055	0.6306	0.0135	0.1494	0.7850	0.1533	0.5238	0.0039	0.9890	-0.1269	0.8768	0.4467	0.1504	0.5736	0.1083
CG5013	CG5013	1626514_at	0.1198	0.5175	0.0869	0.5185	0.1117	0.6590	0.0507	0.9412	0.1371	0.5097	0.0863	0.6742	-0.1358	0.8076	0.0357	0.9216	0.1715	0.4755
---	---	1626515_at	0.1036	0.6808	-0.0111	0.9507	-0.0738	0.7375	0.0245	0.9761	0.0914	0.6965	0.0669	0.7653	0.0426	0.9616	-0.0580	0.8678	-0.1007	0.7189
---	---	1626516_at	0.2122	0.5148	0.0578	0.6135	0.2094	0.4178	0.1914	0.7021	-0.0080	0.9816	-0.1994	0.3440	-0.1748	0.8075	-0.1278	0.7052	0.0471	0.9093
---	---	1626517_at	0.1499	0.4328	0.1217	0.4284	0.1728	0.2976	0.1251	0.8385	0.0670	0.8139	-0.0581	0.8244	-0.1282	0.8846	0.0722	0.8804	0.2004	0.5724
seq	sequoia	1626518_at	-0.2968	0.2119	0.0630	0.7596	-0.0306	0.8468	0.0364	0.9567	-0.1609	0.3828	-0.1972	0.2213	0.0587	0.9430	0.1232	0.6537	0.0644	0.8413
CG15526	CG15526	1626519_at	0.0444	0.8061	0.0970	0.6701	0.0113	0.9552	-0.1302	0.7143	-0.0477	0.8119	0.0824	0.6163	0.0436	0.9611	0.1131	0.6839	0.0696	0.8237
Rad17	Rad17	1626520_at	0.5179	0.1166	0.2366	0.1575	-0.0938	0.7122	-0.3288	0.6015	0.5624	0.0788	0.8912	0.0077	-0.0879	0.9306	0.1509	0.6765	0.2388	0.4795
CG10953	CG10953	1626521_x_at	0.2515	0.1973	0.2909	0.1611	0.6354	0.0137	0.1369	0.7949	0.0410	0.8877	-0.0959	0.6671	-0.0889	0.8991	0.0506	0.8960	0.1395	0.6166
Caf1	Nucleosome remc	1626522_at	0.3626	0.2269	1.1307	0.0520	1.2313	0.0001	-0.0270	0.9727	-0.2937	0.1336	-0.2667	0.1272	-0.0661	0.9734	0.4043	0.4037	0.4704	0.3501
Aats-glupro	Glutamyl-tRNA sy	1626523_a_at	0.6470	0.0158	0.8766	0.0509	0.9991	0.0016	0.3733	0.2989	0.4738	0.0321	0.1006	0.6429	0.3725	0.7220	0.7976	0.0841	0.4251	0.3523
CG16996	CG16996	1626524_at	-0.0909	0.7814	0.1505	0.5120	0.5116	0.0227	-0.0056	0.9956	-0.4122	0.1186	-0.4066	0.0868	-0.0475	0.9742	-0.0272	0.9647	0.0203	0.9716
---	---	1626525_at	0.0801	0.6204	-0.0462	0.8418	0.0528	0.7366	0.2891	0.4861	0.2723	0.2188	-0.0169	0.9523	-0.0466	0.9340	-0.0255	0.9279	0.0211	0.9337
TotF	Turandot F	1626526_at	-0.0028	0.9936	0.1435	0.4514	0.3435	0.1021	0.0181	0.9852	-0.2294	0.3006	-0.2475	0.2070	0.0028	0.9990	0.0222	0.9723	0.0194	0.9743
hh	hedgehog	1626527_at	-0.3901	0.3418	-1.4090	0.0070	-1.7461	0.0027	0.1442	0.9251	0.4216	0.3693	0.2774	0.5361	0.3510	0.7741	-0.4145	0.4333	-0.7655	0.1729
CG40042	CG40042	1626528_at	0.1296	0.6849	0.0828	0.8030	-0.0772	0.6021	-0.1762	0.5423	0.1804	0.2299	0.3566	0.0161	-0.0393	0.9862	0.0817	0.9193	0.1211	0.8582
CG12977	CG12977	1626529_at	0.1589	0.3255	0.2791	0.2216	0.1743	0.2678	-0.1298	0.7397	-0.2315	0.1763	-0.1017	0.5469	-0.1000	0.9068	-0.0782	0.8518	0.0218	0.9613
CecB	Cecropin	1626530_at	-0.0495	0.8769	-2.2577	0.0526	-0.7480	0.3697	1.6699	0.1119	2.6544	0.0021	0.9845	0.0805	0.2437	0.9507	0.4918	0.7106	0.2481	0.8753
Marcal1	dsmarcal1	1626531_at	-0.0580	0.7920	-0.3899	0.1763	-0.2073	0.2292	-0.1235	0.7863	0.1342	0.4975	0.2577	0.1237	-0.3910	0.3738	-0.1747	0.4802	0.2163	0.3828
CG40409	CG40409	1626532_at	-0.0543	0.8091	0.2144	0.2514	0.1237	0.4704	-0.1722	0.7312	-0.2033	0.3740	-0.0311	0.9086	-0.0570	0.9365	0.1163	0.6344	0.1733	0.4589
CG11727 /// DsimCG11727	CG11727	1626533_a_at	-0.4836	0.3528	0.9291	0.1916	1.2465	0.0010	0.1730	0.9263	-0.0430	0.9592	-0.2160	0.7183	-0.2866	0.9000	1.0983	0.1777	1.3849	0.1303
tko	mitochondrial ribo	1626534_at	-0.0157	0.9405	0.6168	0.0817	0.3127	0.1180	0.0101	0.9861	-0.3224	0.0278	-0.3325	0.0155	0.2493	0.7945	0.2279	0.5929	-0.0214	0.9726
---	---	1626535_at	0.2997	0.2743	0.1044	0.4274	-0.0240	0.9489	0.1763	0.7605	0.1931	0.4561	0.0169	0.9573	0.0404	0.9742	0.0365	0.9417	-0.0039	0.9945
CG6776	CG6776	1626536_at	-0.0158	0.9732	0.7060	0.1088	0.3561	0.1053	-0.4160	0.2977	-0.8163	0.0048	-0.4003	0.0605	0.0032	0.9994	-0.0779	0.9290	-0.0811	0.9161
CG17321	CG17321	1626537_at	0.5852	0.1675	0.1482	0.6659	0.6166	0.1074	-0.0733	0.9351	-0.0856	0.7925	-0.0123	0.9708	-0.1849	0.9309	-0.2620	0.7498	-0.0770	0.9369
---	---	1626538_at	0.2879	0.1834	0.2594	0.0703	0.2995	0.1350	-0.0622	0.9067	-0.0744	0.7061	-0.0121	0.9557	0.0582	0.9168	-0.0115	0.9716	-0.0697	0.7527
---	---	1626539_at	0.1369	0.5077	0.1111	0.4676	0.1471	0.3126	-0.0526	0.9542	-0.0354	0.9200	0.0172	0.9573	-0.1970	0.7588	-0.0297	0.9460	0.1674	0.5581

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14076	CG14076	1626540_at	0.0787	0.8025	-0.0806	0.5978	0.0934	0.6458	0.0606	0.9494	0.1638	0.5691	0.1032	0.7151	-0.0724	0.9467	0.0000	1.0000	0.0725	0.8680
---	---	1626541_at	0.0484	0.7839	-0.0901	0.6064	0.0819	0.6758	0.2012	0.5311	0.3101	0.0676	0.1089	0.4995	-0.0199	0.9862	-0.0136	0.9763	0.0063	0.9877
---	---	1626542_at	-0.0128	0.9422	-0.1391	0.3859	0.0992	0.5056	0.0753	0.8844	0.0769	0.7058	0.0016	0.9942	-0.2718	0.6272	-0.1720	0.4697	0.0998	0.7040
CG7066	CG7066	1626543_at	0.2054	0.4256	-0.2745	0.0875	-0.2254	0.1426	-0.1012	0.8605	0.2837	0.1675	0.3849	0.0434	-0.1386	0.8439	-0.2244	0.4363	-0.0858	0.8098
CG14818	CG14818	1626544_at	-1.2504	0.0022	-0.8553	0.0775	-1.4582	0.0008	-0.2941	0.6324	-0.8572	0.0117	-0.5631	0.0424	0.2846	0.7644	-0.4959	0.1975	-0.7805	0.0859
pHCl	pHCl	1626545_at	-0.8717	0.0043	-0.3013	0.1419	-0.5270	0.0115	-0.3824	0.2455	-0.6563	0.0056	-0.2739	0.1149	-0.0340	0.9816	-0.0857	0.8584	-0.0517	0.9145
Tsp74F	Tetraspanin 74F	1626546_s_at	-2.4036	0.0085	-0.5764	0.2730	-1.8936	0.0008	-1.2517	0.0359	-2.6013	0.0002	-1.3496	0.0014	0.1174	0.9774	-0.7539	0.4392	-0.8712	0.3828
---	---	1626547_at	0.1873	0.4833	-0.8071	0.0401	-1.2209	0.0015	-0.2833	0.6080	1.0650	0.0031	1.3482	0.0007	0.0172	0.9928	0.0425	0.9538	0.0254	0.9714
---	---	1626548_s_at	-0.0085	0.9653	-0.0418	0.7350	0.0821	0.7058	0.1062	0.7909	0.1285	0.4491	0.0223	0.9100	-0.0282	0.9816	0.0679	0.8500	0.0961	0.7513
CG7262	CG7262	1626549_at	-0.1092	0.4949	0.3872	0.1818	0.5760	0.0256	-0.1549	0.7929	-0.3184	0.1816	-0.1635	0.4733	-0.1792	0.8270	0.2351	0.5020	0.4143	0.2392
CG32705	CG32705	1626550_at	0.0047	0.9915	-0.1759	0.8612	-0.5131	0.1273	-0.1504	0.7872	0.3173	0.1643	0.4677	0.0302	0.3654	0.8806	0.2483	0.8437	-0.1171	0.9275
---	---	1626551_at	0.0432	0.8294	0.1322	0.4134	0.1590	0.2726	0.0084	0.9922	-0.0779	0.7095	-0.0863	0.6423	-0.0078	0.9923	-0.0214	0.9429	-0.0136	0.9596
CG3568	CG3568	1626552_at	-1.2113	0.0157	-2.1487	0.0041	-2.1131	0.0002	-0.0762	0.9421	0.5969	0.0505	0.6731	0.0198	-0.4046	0.6955	-0.2677	0.5591	0.1369	0.7954
CG10376	CG10376	1626553_at	-0.3831	0.0527	-0.6604	0.0109	-0.8360	0.0011	-0.0751	0.8924	0.1767	0.3503	0.2518	0.1279	0.0857	0.8882	-0.1000	0.7002	-0.1856	0.4308
CG16777	CG16777	1626554_at	-0.4166	0.2337	0.0235	0.9781	-0.9525	0.0688	-0.4156	0.5913	-0.1592	0.7290	0.2565	0.4965	0.5195	0.8215	0.1930	0.8899	-0.3265	0.7710
---	---	1626555_at	0.0959	0.6565	-0.2086	0.3364	-0.2388	0.2332	0.0100	0.9914	0.3848	0.0558	0.3747	0.0401	0.0492	0.9646	-0.0716	0.8642	-0.1208	0.7223
Arpc3A	Arpc3A	1626556_at	-0.3047	0.1341	0.2680	0.2654	0.6933	0.0206	0.0858	0.8642	-0.3264	0.0700	-0.4122	0.0178	0.0651	0.9689	0.5809	0.1730	0.5158	0.2561
NitFhit	NitFlit	1626557_at	0.5668	0.0226	0.9224	0.0091	1.2142	0.0022	-0.3339	0.3102	-1.2846	0.0004	-0.9507	0.0006	-0.5537	0.5228	-0.9036	0.0512	-0.3498	0.4076
CG32241	CG32241	1626558_at	0.0401	0.8747	0.0958	0.6612	-0.1339	0.5483	0.0466	0.9626	0.0217	0.9557	-0.0249	0.9394	0.0015	0.9994	-0.1255	0.6146	-0.1270	0.6129
CG13148	CG13148	1626559_s_at	0.1554	0.3531	0.0554	0.7010	-0.0424	0.8614	-0.0787	0.8863	0.0124	0.9656	0.0911	0.6322	0.0699	0.9234	0.0038	0.9941	-0.0661	0.8256
AQP /// blw	aquaporin /// mitoc	1626560_at	-2.6038	0.0055	-2.6218	0.0027	-2.9053	0.0000	-0.1235	0.8507	-0.7245	0.0089	-0.6010	0.0125	0.1396	0.9618	-0.9322	0.2361	-1.0718	0.2079
CG7692	CG7692	1626561_at	0.3658	0.3300	0.4407	0.1001	0.2032	0.3043	-0.1241	0.7825	0.3983	0.0386	0.5224	0.0076	0.0595	0.9773	0.3373	0.5060	0.2777	0.5988
---	---	1626562_at	0.2468	0.1752	0.1368	0.3488	0.2538	0.2367	0.1361	0.6988	0.1974	0.2202	0.0613	0.7214	0.0385	0.9742	0.1446	0.6514	0.1061	0.7544
Taf6	Suppressor of Ra	1626563_at	0.2137	0.3164	0.1972	0.3925	0.5856	0.0080	0.0968	0.8676	-0.4649	0.0307	-0.5617	0.0085	-0.1677	0.8461	-0.3203	0.3556	-0.1527	0.6993
---	---	1626564_at	0.0299	0.8759	0.0171	0.8895	0.0853	0.6521	-0.0640	0.9228	0.0178	0.9520	0.0818	0.6957	-0.1632	0.7826	-0.0380	0.9253	0.1252	0.6498
CG2790	CG2790	1626565_at	-0.3550	0.1257	0.3208	0.2783	0.5834	0.0791	0.0106	0.9915	-0.3258	0.1165	-0.3364	0.0731	-0.2885	0.8156	0.2749	0.6152	0.5634	0.2842
CG12262	CG12262	1626566_at	1.0116	0.0079	0.6641	0.2424	0.9362	0.0008	-0.0791	0.9149	-0.1861	0.4383	-0.1070	0.6553	-0.3288	0.8016	-0.5641	0.2859	-0.2352	0.6991
beat-VI	beat-VI	1626567_at	0.0888	0.7960	0.2413	0.1498	-0.2529	0.2500	0.0403	0.9659	0.1485	0.5826	0.1082	0.6776	0.3063	0.7057	0.0961	0.8297	-0.2101	0.5571
CG7357	CG7357	1626568_at	-0.1074	0.4514	0.0247	0.9127	0.3847	0.0636	-0.0385	0.9436	-0.5010	0.0066	-0.4625	0.0056	-0.3603	0.7070	-0.2622	0.5286	0.0980	0.8498
---	---	1626569_at	0.2264	0.1439	-0.0940	0.5985	-0.0486	0.7814	-0.0597	0.9039	0.0588	0.7606	0.1185	0.4333	-0.0277	0.9816	-0.1290	0.6456	-0.1013	0.7325
SNF4Agamma	loechrig	1626570_s_at	-0.9408	0.0327	-0.0120	0.9882	-0.6730	0.0144	-0.0942	0.8166	-0.4913	0.0084	-0.3972	0.0132	0.5833	0.7628	0.2280	0.8309	-0.3553	0.6893
CG7942	CG7942	1626571_at	0.0407	0.7870	0.1408	0.6137	0.2959	0.0833	-0.1116	0.8234	0.0199	0.9445	0.1315	0.4772	-0.2217	0.7768	0.2157	0.5414	0.4374	0.2136
CG10822	CG10822	1626572_at	0.0500	0.8223	0.1330	0.4924	0.1478	0.3226	-0.0060	0.9942	-0.1534	0.3767	-0.1474	0.3439	-0.0399	0.9718	0.0387	0.9328	0.0786	0.8282
CG13579	CG13579	1626573_at	0.0543	0.7632	0.1256	0.6888	0.2173	0.3142	0.1059	0.8676	0.0041	0.9899	-0.1017	0.6615	0.0955	0.9092	0.1027	0.7784	0.0072	0.9875
CG9935	CG9935	1626574_at	-0.4438	0.0461	-0.8210	0.1133	-1.1908	0.0149	0.0219	0.9923	0.6537	0.1678	0.6318	0.1367	0.0071	0.9963	0.1491	0.6823	0.1419	0.6969
CG1657 /// DmirCG1657	CG1657	1626575_at	-0.9952	0.0630	-0.1586	0.8113	-0.0487	0.8396	-0.1100	0.8446	-0.5888	0.0118	-0.4788	0.0177	-0.1684	0.9467	0.1067	0.9292	0.2751	0.7537
---	---	1626576_at	0.0867	0.6710	-0.0006	0.9994	0.1065	0.5140	-0.0542	0.9311	0.1675	0.3730	0.2217	0.1772	-0.0709	0.9238	0.0721	0.8248	0.1430	0.5889
CG15556	CG15556	1626577_at	-3.3522	0.0005	-3.4111	0.0075	-3.9630	0.0000	-0.4638	0.2753	-0.0339	0.9228	0.4299	0.0591	0.0208	0.9928	-0.0982	0.9077	-0.1190	0.8729
CG5195	CG5195	1626578_at	-0.0382	0.8327	-0.0139	0.9453	-0.0391	0.8625	0.0725	0.8732	0.0519	0.7870	-0.0206	0.9148	-0.0464	0.9657	0.0734	0.8548	0.1198	0.7154
CG3040 /// DyakCG3040	CG3040	1626579_at	-0.5739	0.0084	-0.5328	0.0638	-0.5504	0.0133	0.2135	0.5680	0.1301	0.5229	-0.0834	0.6774	0.2314	0.7230	0.2196	0.4355	-0.0118	0.9798
aft	adrift	1626580_at	0.5484	0.0374	-0.0727	0.8001	-0.4478	0.1664	-0.3526	0.2642	0.2308	0.2149	0.5834	0.0047	-0.0309	0.9889	-0.3111	0.5246	-0.2802	0.5767
CG1607	CG1607	1626581_s_at	-0.1298	0.6441	-1.4472	0.0093	-0.6639	0.0102	0.4701	0.1345	0.5398	0.0118	0.0698	0.7318	-0.3537	0.6896	-0.7027	0.0684	-0.3490	0.3458
---	---	1626582_x_at	0.2567	0.2001	0.2644	0.0925	0.1280	0.4745	0.0402	0.9436	0.1561	0.3462	0.1159	0.4509	0.1363	0.7686	0.1223	0.5523	-0.0140	0.9614
Or94b	Odorant receptor	1626583_at	0.2118	0.2947	0.2129	0.1129	0.2806	0.2498	0.0840	0.8640	0.0198	0.9361	-0.0642	0.7307	-0.0055	0.9952	-0.0661	0.8132	-0.0606	0.8225
SMSr	SMSr	1626584_a_at	-0.2157	0.1652	0.4359	0.0521	0.5946	0.0069	-0.0567	0.9410	-0.5292	0.0239	-0.4725	0.0241	-0.1575	0.8157	0.1369	0.6516	0.2944	0.3027
CG4589	CG4589	1626585_s_at	-0.1608	0.4729	0.4955	0.1739	0.5834	0.0070	-0.3088	0.4631	-0.8227	0.0041	-0.5140	0.0198	-0.2878	0.7215	0.0119	0.9858	0.2998	0.3859
Obp57a	Odorant-binding p	1626586_at	-1.1748	0.0041	-0.2285	0.7710	-1.6908	0.0004	-1.1984	0.0741	-1.5663	0.0025	-0.3679	0.2952	0.1284	0.9589	-0.7083	0.3011	-0.8367	0.2527
---	---	1626587_s_at	0.2079	0.3855	0.2164	0.1217	0.0619	0.8233	0.0074	0.9934	-0.0368	0.8732	-0.0442	0.8229	0.0557	0.9659	-0.0901	0.8511	-0.1458	0.7134
CG5325	Peroxisomal farne	1626588_a_at	-0.9116	0.0020	0.0271	0.8423	0.5061	0.1120	0.0803	0.9345	-1.0301	0.0043	-1.1104	0.0018	-0.2919	0.7506	-0.2318	0.5769	0.0602	0.9115
---	---	1626589_at	0.0893	0.6791	0.1294	0.4467	0.0153	0.9661	-0.0004	0.9995	-0.0020	0.9942	-0.0016	0.9947	0.0482	0.9746	0.1840	0.6573	0.1358	0.7555
MCPH1	Microcephalin	1626590_a_at	-0.5495	0.1443	-1.3371	0.0144	-0.9358	0.0019	0.3064	0.5932	0.9970	0.0051	0.6907	0.0160	-0.0079	0.9952	0.2996	0.3236	0.3075	0.3343
siz	schizo	1626591_at	0.2423	0.6091	0.0887	0.9122	-0.5373	0.0197	0.0297	0.9622	0.4344	0.0173	0.4047	0.0146	0.8098	0.6955	0.4354	0.6380	-0.3743	0.6923
---	---	1626592_at	-0.0681	0.8161	0.2573	0.3021	0.2676	0.1621	-0.0466	0.9649	-0.0603	0.8723	-0.0138	0.9689	0.0796	0.9309	0.1555	0.6281	0.0759	0.8418
CG14693	CG14693	1626593_at	-0.0084	0.9675	0.0252	0.8546	0.2687	0.0802	-0.0873	0.8211	-0.1240	0.4260	-0.0368	0.8300	-0.1727	0.7307	0.0281	0.9353	0.2008	0.3641
svp	seven-up	1626594_s_at	1.5766	0.1562	1.5009	0.1463	1.1274	0.0394	0.0758	0.8735	0.1013	0.5711	0.0254	0.8987	0.4759	0.9238	0.1653	0.9505	-0.3106	0.8909
---	---	1626595_s_at	0.2184	0.3098	0.3107	0.0927	0.1443	0.4355	-0.1796	0.5832	-0.2746	0.0993	-0.0950	0.5646	0.1858	0.7893	0.0054	0.9935	-0.1804	0.5669
CG32437	CG32437	1626596_at	0.0801	0.6880	0.0295	0.8777	-0.3268	0.1071	-0.1299	0.7705	-0.0717	0.7482	0.0582	0.7810	0.1991	0.6955	-0.0034	0.9941	-0.2025	0.3601
CG9521	CG9521	1626597_at	-0.4292	0.1112	-1.3414	0.0246	-1.3161	0.0013	0.2186	0.8379	1.1716	0.0105	0.9530	0.0159	0.0806	0.9246	-0.0262	0.9551	-0.1068	0.7456
---	---	1626598_at	0.0512	0.7693	-0.0713	0.5460	0.2933	0.1501	0.1280	0.7243	0.0187	0.9355	-0.1093	0.4838	-0.2222	0.7588	-0.0665	0.8785	0.1556	0.6331
CG17050	CG17050	1626599_at	0.0190	0.9566	0.2234	0.4124	-0.1450	0.5529	-0.1519	0.8439	-0.1697	0.5832	-0.0178	0.9599	0.2229	0.7843	0.1098	0.8079	-0.1130	0.7865
CG30016	CG30016	1626600_at	-0.2236	0.6315	-0.0751	0.5516	-0.0676	0.6878	0.1689	0.9036	-0.3544	0.4502	-0.5234	0.1939	-0.0308	0.9776	-0.0909	0.7780	-0.0601	0.8614
CG8629	CG8629	1626601_at	0.1705	0.2928	0.2239	0.1688	0.1021	0.4808	-0.0270	0.9649	0.0383	0.8583	0.0653	0.7065	0.2908	0.4415	0.2767	0.1498	-0.0141	0.9613
Rack1	Guanine nucleotid	1626602_at	0.1918	0.1761	1.0763	0.0030	1.1693	0.0002	0.1250	0.8034	-0.6258	0.0080	-0.7508	0.0021	-0.0122	0.9885	0.0464	0.8550	0.0586	0.7918
zormin	D-Titin	1626603_at	0.4022	0.1319	0.0133	0.9523	-1.1144	0.0013	-0.3383	0.3979	0.7670	0.0052	1.1053	0.0006	0.7312	0.4415	0.1693	0.7804	-0.5618	0.2717
CG30363	CG30363	1626604_at	0.2806	0.0670	0.1597	0.4131	0.2195	0.2327	0.1728	0.6122	0.2118	0.2150	0.0390	0.8435	-0.0282	0.9819	0.0358	0.9402	0.0640	0.8694
CG14075	CG14075	1626605_at	0.0460	0.8601	0.1419	0.3735	0.5328	0.0140	-0.1247	0.8294	-0.2276	0.3103	-0.1029	0.6540	-0.1618	0.7726	-0.0555	0.8725	0.1063	0.6941
CG10630	CG10630	1626606_at	1.0898	0.0206	0.4217	0.5031	0.3164	0.2022	0.3658	0.5495	0.7865	0.0210	0.4207	0.1347	0.2801	0.9014	0.2273	0.8382	-0.0527	0.9660
sec63	sec63	1626607_at	0.9572	0.0108	0.7943	0.0314	1.0736	0.0001	0.1742	0.7386	0.7385	0.0071	0.5642	0.0141	-0.1314	0.9076	0.5650	0.1498	0.6963	0.1184
---	---	1626608_at	0.0828	0.5880	0.0996	0.4713	-0.0707	0.6962	-0.1269	0.7604	0.0328	0.8924	0.1598	0.3285	0.1336	0.8193	0.1324	0.6049	-0.0011	0.9979

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG10006	CG10006	1626609_at	0.8501	0.0238	1.5216	0.1434	2.6552	0.0000	0.1510	0.8128	-1.3869	0.0007	-1.5380	0.0003	-0.5620	0.7726	-0.3674	0.6949	0.1946	0.8577
---	---	1626610_at	-0.1063	0.5738	0.1510	0.2906	0.3165	0.0905	0.1296	0.7658	-0.1464	0.4478	-0.2760	0.0959	-0.0110	0.9898	-0.0018	0.9972	0.0092	0.9756
---	---	1626611_at	0.1345	0.6034	0.0713	0.6301	0.1121	0.4954	-0.1161	0.7929	-0.1542	0.4034	-0.0381	0.8555	-0.1935	0.7440	-0.1280	0.6370	0.0655	0.8364
---	---	1626612_at	-0.3156	0.0818	0.1601	0.6307	0.0773	0.7473	-0.2341	0.6688	-0.4760	0.0703	-0.2419	0.3063	0.0874	0.9449	0.0707	0.9041	-0.0167	0.9783
CG11236 /// DsimCG11236	CG11236	1626613_at	3.0729	0.0012	1.5385	0.0434	3.4370	0.0001	1.1329	0.1316	0.7096	0.1143	-0.4233	0.3003	-0.8365	0.6749	-0.9162	0.2378	-0.0797	0.9433
---	---	1626614_at	0.0638	0.8287	0.0312	0.7877	0.0569	0.7986	-0.0240	0.9834	0.0267	0.9455	0.0507	0.8705	-0.0757	0.9092	-0.0412	0.9085	0.0345	0.9160
CG31156	CG31156	1626615_at	-0.8616	0.0072	0.0386	0.8102	0.1289	0.5887	-0.2404	0.7352	-0.3692	0.2438	-0.1288	0.6988	-0.1005	0.9046	0.6385	0.0567	0.7390	0.0527
CG7465	CG7465	1626616_at	0.4009	0.0737	0.3784	0.1247	0.4861	0.0251	0.0864	0.9300	0.1168	0.7299	0.0304	0.9319	-0.1208	0.8609	0.0602	0.8807	0.1810	0.5399
dx	deltex	1626617_at	-0.0461	0.9152	-0.3761	0.3887	-0.4361	0.0191	0.0944	0.8493	0.5146	0.0119	0.4202	0.0176	0.0801	0.9742	0.2181	0.7755	0.1379	0.8679
CG34342	CG10629	1626618_at	0.0838	0.7096	-0.0588	0.6595	0.1534	0.3373	0.1242	0.8000	0.0013	0.9963	-0.1229	0.5227	-0.0778	0.9011	-0.1016	0.6930	-0.0238	0.9410
CG9919	CG9919	1626619_at	-0.3075	0.3425	-0.2928	0.1997	-0.5022	0.0189	-0.1615	0.8247	-0.2895	0.3100	-0.1280	0.6620	-0.0646	0.9335	-0.2727	0.2518	-0.2080	0.4086
MRG15	MRG15	1626620_at	-0.2647	0.1013	0.0805	0.6286	0.0339	0.8363	-0.1214	0.8090	0.0128	0.9674	0.1342	0.4818	0.0001	0.9999	0.3679	0.1151	0.3678	0.1390
CG1407	CG1407	1626621_at	0.2824	0.2514	0.5241	0.1503	0.3305	0.0839	-0.2727	0.5492	-0.0849	0.7637	0.1878	0.3906	0.0320	0.9816	0.2918	0.3075	0.2599	0.3877
CG10750	CG10750	1626622_at	0.0980	0.8546	-0.4991	0.3288	-1.0144	0.0081	-0.4271	0.5039	0.3075	0.3747	0.7346	0.0236	0.0538	0.9869	0.1558	0.8852	0.1020	0.9211
CG11670	CG11670	1626623_at	-0.1487	0.7560	0.2427	0.3991	-0.1366	0.5691	-0.0631	0.9232	0.0083	0.9786	0.0714	0.7370	0.2086	0.8837	0.0677	0.9387	-0.1410	0.8374
Pax	paxillin	1626624_s_at	0.3363	0.3146	-0.0609	0.7080	-0.3321	0.2427	0.1439	0.8863	0.2666	0.4519	0.1227	0.7378	0.5193	0.6955	0.0444	0.9617	-0.4750	0.3995
---	---	1626625_at	-0.3093	0.3929	0.2818	0.6190	-0.2405	0.2693	-0.5062	0.3665	-0.2490	0.4524	0.2572	0.3834	-0.0221	0.9923	0.2857	0.6124	0.3078	0.5834
CG13079	CG13079	1626626_at	0.1083	0.7555	0.0752	0.4388	0.1308	0.4321	0.0044	0.9956	-0.0535	0.8041	-0.0578	0.7639	0.0151	0.9913	-0.0729	0.8500	-0.0879	0.7962
CG11718	CG11718	1626627_at	0.3171	0.1456	0.2331	0.4006	0.1748	0.4893	-0.0859	0.8671	-0.1173	0.5479	-0.0314	0.8839	-0.0779	0.9528	-0.1536	0.7386	-0.0757	0.8886
---	---	1626628_at	0.1575	0.3130	-0.1149	0.4133	0.2487	0.2566	0.1587	0.7451	0.2266	0.2944	0.0679	0.7749	-0.1000	0.8571	-0.0128	0.9735	0.0872	0.7391
---	---	1626629_s_at	0.0244	0.9059	0.1637	0.5023	0.2359	0.3323	-0.2316	0.6869	-0.2453	0.3687	-0.0137	0.9679	-0.1659	0.8097	0.0205	0.9650	0.1864	0.5367
Sgs4	salivary gland sec	1626630_at	0.0543	0.8506	0.1481	0.2860	0.1713	0.3233	0.0051	0.9956	-0.1390	0.5535	-0.1441	0.4892	0.0240	0.9853	0.0390	0.9353	0.0150	0.9749
---	---	1626631_at	0.1107	0.6853	0.2882	0.3010	0.2413	0.0966	-0.0323	0.9755	0.0052	0.9896	0.0374	0.9035	-0.0242	0.9764	0.0031	0.9941	0.0273	0.9197
---	---	1626632_at	-0.1510	0.7171	-0.2462	0.3923	-0.6333	0.0153	-0.2458	0.7130	0.3107	0.3115	0.5566	0.0462	-0.1939	0.8608	-0.0028	0.9989	0.1911	0.6995
CG32559	CG32559	1626633_at	0.2110	0.4049	0.1389	0.4196	-0.0151	0.9477	-0.2230	0.5478	0.0304	0.9091	0.2534	0.1383	0.1046	0.8990	0.0016	0.9991	-0.1030	0.7744
---	---	1626634_at	0.3570	0.1752	0.0565	0.6522	0.2333	0.1333	0.1464	0.6886	0.2073	0.2202	0.0609	0.7385	-0.0405	0.9717	-0.1349	0.6746	-0.0944	0.7834
dpr3	dpr3	1626635_at	0.0245	0.9144	0.2312	0.3331	0.1436	0.3574	0.0165	0.9860	0.0053	0.9876	-0.0112	0.9679	0.1128	0.9056	0.1995	0.5808	0.0867	0.8422
---	---	1626636_at	-0.0998	0.6295	0.1620	0.2947	-0.0176	0.9231	-0.1426	0.7815	-0.2055	0.3433	-0.0629	0.7923	0.1685	0.7230	0.1326	0.5374	-0.0358	0.9000
bif	bifocal	1626637_a_at	-1.4738	0.0037	-2.3352	0.0077	-2.1274	0.0001	0.0731	0.8836	0.2238	0.1850	0.1507	0.3316	-0.1770	0.9342	-0.4137	0.5679	-0.2367	0.7695
escl	esc-like	1626638_at	0.0130	0.9704	-0.2594	0.3274	-0.2613	0.1418	-0.0959	0.9343	0.2982	0.3956	0.3941	0.1967	-0.0402	0.9778	0.0046	0.9951	0.0448	0.9246
CG4655	CG4655	1626639_a_at	-0.1449	0.6637	-1.3990	0.0053	-2.0500	0.0037	-0.1080	0.9558	1.1961	0.0299	1.3041	0.0132	-0.0909	0.9239	-0.2370	0.4601	-0.1461	0.6761
CG4565	CG4565	1626640_at	-0.1851	0.3664	0.0000	1.0000	-0.1134	0.6289	-0.0702	0.9037	0.2507	0.1818	0.3210	0.0611	-0.1490	0.8655	0.2769	0.4285	0.4260	0.2398
glob1	Hemoglobin	1626641_s_at	0.1688	0.6687	0.0609	0.9394	-0.6410	0.0340	-1.0389	0.1119	-1.1225	0.0112	-0.0836	0.8538	-0.4040	0.8141	-1.3012	0.0795	-0.8972	0.2230
CG6486	CG6486	1626642_at	-0.2092	0.4042	-0.0442	0.7505	-0.0610	0.7635	0.0508	0.9311	0.0520	0.8071	0.0011	0.9959	0.0400	0.9721	-0.0769	0.8445	-0.1168	0.7222
CG32160	CG32160	1626643_at	0.0833	0.5469	0.0388	0.6944	0.2818	0.1377	-0.0145	0.9819	-0.0560	0.7551	-0.0415	0.8056	-0.0437	0.9503	0.0069	0.9871	0.0506	0.8514
CG33062	CG33062	1626644_at	0.0474	0.7585	0.1124	0.4058	0.1891	0.3796	-0.0894	0.9037	-0.1045	0.7061	-0.0152	0.9597	0.0907	0.9238	-0.0103	0.9872	-0.1011	0.7884
CG8596	CG8596	1626645_at	-0.5803	0.0122	-0.2549	0.4189	-0.2125	0.4841	0.0672	0.9098	-0.4494	0.0251	-0.5165	0.0085	0.0311	0.9898	-0.1308	0.8569	-0.1619	0.8002
CG4729	CG4729	1626646_s_at	0.8011	0.0087	0.4748	0.0223	0.7950	0.0012	0.2218	0.5766	0.0387	0.8885	-0.1831	0.3246	0.0625	0.9514	-0.1620	0.6259	-0.2245	0.4820
Mcm5	Minichromosome	1626647_at	-0.1632	0.8191	-0.9914	0.4537	-1.5045	0.0992	-0.4767	0.3328	1.0949	0.0031	1.5715	0.0004	-0.2527	0.9780	0.0183	0.9980	0.2709	0.9295
CG12897	CG12897	1626648_at	0.1749	0.5225	0.0289	0.9441	0.1518	0.2891	-0.0657	0.9627	-0.0387	0.9436	0.0270	0.9547	-0.0673	0.8882	0.0512	0.8302	0.1185	0.5319
koko	kokopelli	1626649_s_at	0.2902	0.2234	-0.0067	0.9674	-0.1468	0.3782	0.2650	0.4573	0.8522	0.0017	0.5872	0.0053	0.3008	0.6749	0.4657	0.1188	0.1649	0.6030
mod(mdg4)	Modifier67.2	1626650_at	-0.0182	0.9432	0.1109	0.6629	-0.1128	0.4567	-0.1480	0.6533	-0.1547	0.3378	-0.0068	0.9727	-0.0724	0.9587	-0.0868	0.8754	-0.0144	0.9832
c-cup	calcutta cup	1626651_at	0.1119	0.4886	-0.1790	0.1782	-0.0738	0.7336	-0.0976	0.8676	0.1058	0.6500	0.2034	0.2773	-0.1562	0.8202	-0.1625	0.5902	-0.0063	0.9891
Orc1	Origin recognition	1626652_at	0.8435	0.0032	0.3904	0.3264	-0.2615	0.4973	-0.5138	0.2472	1.1425	0.0018	1.6563	0.0003	0.3460	0.8292	0.8151	0.2114	0.4691	0.5007
ferrochelatase	ferrochelatase	1626653_a_at	-0.1660	0.4895	0.7614	0.0236	1.1683	0.0012	0.1212	0.8436	-0.8085	0.0044	-0.9296	0.0014	-0.1573	0.8461	0.1280	0.7439	0.2853	0.4004
CG11791	CG11791	1626654_a_at	-0.5469	0.0392	-0.4846	0.0699	-0.7395	0.0071	0.2377	0.4442	0.3999	0.0260	0.1623	0.2898	0.4030	0.6145	0.4565	0.1682	0.0536	0.9095
bnk	bottleneck	1626655_at	0.2209	0.1472	0.1326	0.3769	0.2794	0.1848	0.2654	0.4356	0.3571	0.0598	0.0917	0.6303	0.2022	0.8362	0.3641	0.3571	0.1619	0.7245
---	---	1626656_at	0.0999	0.5952	0.1436	0.5015	0.1152	0.4872	0.0633	0.8987	0.0507	0.7998	-0.0126	0.9511	0.1005	0.8541	0.0676	0.8168	-0.0329	0.9161
wfs1	wolfram syndrome	1626657_a_at	-0.0296	0.9182	-0.3225	0.1073	-0.3220	0.0737	0.2114	0.5947	0.6305	0.0078	0.4191	0.0278	0.1754	0.8145	0.1914	0.5539	0.0160	0.9735
---	---	1626658_at	0.0875	0.5367	0.3245	0.3553	0.5034	0.0254	-0.1100	0.7825	-0.2886	0.0812	-0.1786	0.2260	-0.1371	0.8740	0.0465	0.9320	0.1836	0.6161
CG4980	CG4980	1626659_at	0.1126	0.5419	0.1825	0.4288	0.1448	0.3304	-0.0411	0.9319	0.0523	0.7613	0.0934	0.4983	-0.0615	0.9449	0.1080	0.7299	0.1696	0.5515
CG15024	CG15024	1626660_at	0.3342	0.0916	0.0783	0.6118	0.1047	0.5525	0.0646	0.9149	0.1511	0.4419	0.0865	0.6590	0.0346	0.9657	0.0115	0.9760	-0.0231	0.9395
CG31355	CG31355	1626661_at	0.1454	0.5462	-0.0327	0.7456	0.0892	0.5677	0.2432	0.6122	0.2107	0.3944	-0.0325	0.9111	0.1646	0.8427	-0.0442	0.9353	-0.2087	0.5637
CG12309	CG12309	1626662_at	0.3691	0.0617	0.0372	0.7379	0.1978	0.3339	0.0766	0.9185	0.0665	0.8157	-0.0101	0.9718	0.0009	0.9994	-0.1356	0.4683	-0.1365	0.4742
Eb1	Eb1	1626663_s_at	-0.5390	0.0368	0.1592	0.3822	0.0220	0.9036	-0.0963	0.8470	-0.3078	0.0930	-0.2115	0.1952	0.0638	0.9495	0.3330	0.2501	0.2691	0.3808
CG3285	CG3285	1626664_at	-0.4235	0.4481	0.0948	0.5951	0.0801	0.7368	0.1094	0.9491	-0.4491	0.3583	-0.5586	0.1918	0.0924	0.9390	0.1204	0.7989	0.0279	0.9588
CG7408	CG7408	1626665_at	-0.9413	0.0591	-1.4114	0.0611	-2.1240	0.0002	-0.1621	0.7250	-0.1271	0.5701	0.0350	0.8873	-0.1819	0.8097	-0.4874	0.1245	-0.3054	0.3457
RhoGAP16F	RhoGAP16F	1626666_s_at	-1.1141	0.0230	-2.4090	0.0083	-2.8304	0.0000	-0.0755	0.9339	1.0510	0.0029	1.1265	0.0013	0.1210	0.9467	-0.3387	0.5613	-0.4596	0.4153
miple	miple	1626667_at	-0.4569	0.1138	-0.0617	0.6311	-0.2191	0.2966	-0.2014	0.6650	-0.4354	0.0545	-0.2340	0.2396	0.0233	0.9841	-0.0155	0.9723	-0.0389	0.9158
---	---	1626668_at	0.0951	0.4875	0.0266	0.7901	0.1611	0.3780	-0.0186	0.9767	0.1384	0.3898	0.1570	0.2672	-0.0845	0.8609	0.0433	0.8765	0.1279	0.5345
---	---	1626669_at	0.0627	0.7708	-0.1163	0.4053	-0.0832	0.6966	0.0284	0.9630	0.0343	0.8736	0.0059	0.9770	0.0434	0.9588	-0.0662	0.8310	-0.1096	0.6668
CG13744	CG13744	1626670_at	0.0713	0.7982	0.3360	0.0755	0.2803	0.1454	0.2167	0.5842	0.0517	0.8432	-0.1649	0.3773	0.3558	0.6955	0.3893	0.2924	0.0334	0.9502
---	---	1626671_at	0.0028	0.9907	-0.0117	0.9596	-0.1418	0.3375	0.0023	0.9972	0.1131	0.5648	0.1108	0.5327	0.0402	0.9550	0.0084	0.9831	-0.0318	0.9118
---	---	1626672_at	0.1387	0.3781	0.0904	0.3946	0.1202	0.5342	-0.0578	0.9313	-0.1632	0.4251	-0.1054	0.5946	0.0042	0.9963	-0.0150	0.9623	-0.0192	0.9433
CG31716	CG31716	1626673_s_at	-0.1894	0.3140	-0.3096	0.4427	-0.5571	0.0144	-0.1684	0.7135	0.2039	0.3371	0.3723	0.0510	0.1399	0.9246	0.1913	0.7424	0.0514	0.9412
---	---	1626674_at	-0.0530	0.7713	0.0472	0.7168	-0.0395	0.9036	-0.0175	0.9838	-0.0448	0.8626	-0.0272	0.9089	0.1582	0.8541	0.1083	0.8132	-0.0499	0.9189
XRCC1	XRCC1	1626675_at	-0.6492	0.2662	-0.3889	0.2442	-0.5888	0.0048	0.0710	0.9451	0.4891	0.0960	0.4181	0.1102	0.0866	0.9783	0.6951	0.3454	0.6084	0.4322
plexB	Plexin-B	1626676_at	-0.2219	0.3971	-0.1044	0.8617	-0.4581	0.0528	-0.2514	0.6627	-0.0154	0.9711	0.2360	0.3517	0.0355	0.9892	0.0119	0.9925	-0.0236	0.9784
---	---	1626677_s_at	0.0233	0.9043	0.1621	0.2997	0.2254	0.1373	-0.0723	0.8875	-0.1403	0.4324	-0.0680	0.7100	-0.1352	0.8202	-0.0200	0.9613	0.1153	0.6662

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG7135	CG7135	1626678_at	0.0686	0.7160	-0.0153	0.8805	0.2431	0.1912	0.2085	0.5255	0.2124	0.2156	0.0039	0.9856	-0.1309	0.7953	0.0001	1.0000	0.1309	0.5612
CG17327 /// DyakCG17327	CG17327	1626679_at	-0.4977	0.0888	0.3714	0.2117	0.7380	0.0043	-0.0428	0.9375	-0.9155	0.0006	-0.8726	0.0004	-0.2792	0.7990	0.1084	0.8658	0.3876	0.4044
CG9305	CG9305	1626680_at	-0.3568	0.1668	-0.3516	0.1110	-0.1968	0.2544	0.2119	0.5735	0.2418	0.2087	0.0300	0.8974	0.0595	0.9547	0.2960	0.3236	0.2365	0.4568
---	---	1626681_s_at	0.2612	0.1855	-0.0835	0.7186	0.0959	0.7268	0.1608	0.7929	0.1084	0.7061	-0.0524	0.8567	-0.0807	0.9441	-0.2162	0.5531	-0.1355	0.7353
CG9135	CG9135	1626682_s_at	-0.3327	0.1734	0.4097	0.0911	0.6947	0.0407	-0.3105	0.4791	-0.6363	0.0147	-0.3258	0.1200	-0.5885	0.5126	0.1955	0.6783	0.7841	0.1003
CG1514 /// DsimCG1514	CG1514	1626683_at	-0.5975	0.1087	0.1067	0.5353	0.0157	0.9431	-0.3653	0.4962	-0.5944	0.0430	-0.2292	0.3852	-0.0359	0.9884	0.3210	0.5706	0.3569	0.5270
CG7607	CG7607	1626684_at	-0.7310	0.0459	-0.9699	0.0370	-0.6953	0.0035	0.4608	0.1138	1.3159	0.0002	0.8552	0.0006	0.1875	0.8882	1.0507	0.0577	0.8632	0.1190
CG14418	CG14418	1626685_at	0.1286	0.5320	0.3472	0.1411	-0.0722	0.7379	-0.4541	0.2438	-0.1522	0.5390	0.3019	0.1427	0.2156	0.7961	0.0487	0.9306	-0.1670	0.6607
---	---	1626686_at	0.2204	0.3454	0.0926	0.5052	0.0695	0.7813	0.0123	0.9865	-0.0291	0.9045	-0.0414	0.8378	0.0885	0.9152	-0.1156	0.7331	-0.2041	0.4978
CG10956	CG10956	1626687_at	-0.0337	0.9150	-0.4584	0.0755	-0.0412	0.8146	0.0302	0.9696	0.0888	0.7056	0.0586	0.7978	-0.1139	0.8472	-0.1051	0.7050	0.0088	0.9835
CG13004	CG13004	1626688_at	-1.1977	0.0233	-1.3707	0.0487	-0.9646	0.0530	-0.1478	0.9314	-0.0758	0.9155	0.0720	0.9074	-0.5400	0.7220	-0.2911	0.6844	0.2489	0.7362
Cyp6t1	Cyp6t1	1626689_at	2.7609	0.0136	1.3035	0.1309	2.8254	0.0001	0.7082	0.6854	0.0174	0.9890	-0.6908	0.3548	-0.8905	0.6724	-1.3824	0.1095	-0.4919	0.5870
---	---	1626690_at	0.4517	0.0357	0.3232	0.1347	0.2403	0.3925	-0.0058	0.9953	0.0957	0.6739	0.1016	0.6153	0.0411	0.9637	0.0448	0.9054	0.0037	0.9933
---	---	1626691_at	-0.0046	0.9857	-0.1929	0.4705	0.0539	0.7436	0.2878	0.4455	0.2040	0.3279	-0.0839	0.6982	0.0129	0.9916	0.0511	0.9075	0.0381	0.9231
CG14573	CG14573	1626692_at	0.1584	0.2929	0.1577	0.4799	0.2870	0.1019	-0.0619	0.9393	-0.0657	0.8254	-0.0038	0.9901	0.0131	0.9914	0.1694	0.5299	0.1563	0.5709
RpS29	Ribosomal protein	1626693_at	0.6576	0.0908	1.5966	0.0149	0.8716	0.0160	-0.1279	0.7028	-0.5111	0.0060	-0.3832	0.0132	0.4992	0.7644	0.3058	0.6975	-0.1934	0.8255
CG1674	CG1674	1626694_at	-1.2357	0.0054	-0.3038	0.2477	-1.0186	0.0039	-0.4016	0.4481	-0.9131	0.0074	-0.5116	0.0516	0.0834	0.9514	-0.0498	0.9402	-0.1333	0.7838
CG9518	CG9518	1626695_at	0.1374	0.6434	-0.2714	0.1387	-0.4456	0.0913	-0.1719	0.8628	0.2080	0.5890	0.3800	0.2308	-0.0305	0.9848	-0.1276	0.7730	-0.0970	0.8310
E(z)	Enhancer-of-zeste	1626696_at	-0.9081	0.0130	-1.8623	0.0089	-1.3335	0.0023	0.4449	0.6122	0.9126	0.0444	0.4677	0.2363	-0.0494	0.9653	0.1015	0.7903	0.1509	0.6457
---	---	1626697_at	0.0190	0.9362	0.0914	0.4659	0.2616	0.1296	-0.0207	0.9777	-0.1975	0.2794	-0.1769	0.2809	-0.0111	0.9898	0.0332	0.9054	0.0443	0.8505
CG15543	CG15543	1626698_at	0.1982	0.4420	0.0528	0.6870	0.0700	0.7948	0.2747	0.5731	0.2030	0.4306	-0.0717	0.7978	-0.0730	0.9421	-0.2231	0.4794	-0.1500	0.6533
CG34002	CG34002	1626699_at	0.0349	0.8810	-0.1936	0.2025	-0.2936	0.0867	0.1120	0.7857	0.3668	0.0378	0.2548	0.0941	0.1973	0.8145	0.0775	0.8764	-0.1199	0.7689
Act88F	arthrin	1626700_at	-0.3559	0.5774	-0.1857	0.3276	-0.2732	0.1350	-0.1505	0.8826	-0.5021	0.1471	-0.3516	0.2616	-0.1450	0.9499	-0.3840	0.5995	-0.2390	0.7647
CG4743	CG4743	1626701_at	-0.0548	0.7108	0.0644	0.7296	0.3707	0.1629	0.0807	0.8707	0.2075	0.2310	0.1268	0.4368	-0.2321	0.8236	0.2656	0.5597	0.4977	0.2677
fz3	Dfrizzled-3	1626702_at	0.2424	0.3082	0.7858	0.2785	0.2111	0.4991	-0.3920	0.3514	-0.5415	0.0321	-0.1495	0.5242	0.0432	0.9901	-0.2054	0.8432	-0.2486	0.7849
---	---	1626703_at	0.1147	0.4716	0.1484	0.3755	0.0446	0.7952	-0.1960	0.5455	-0.1862	0.2702	0.0098	0.9632	0.0405	0.9665	-0.0313	0.9402	-0.0718	0.8237
CG7248	CG7248	1626704_at	0.2493	0.3098	-0.0818	0.5103	-0.0051	0.9807	0.2032	0.7067	0.2780	0.2626	0.0748	0.7868	-0.0216	0.9848	-0.0488	0.8962	-0.0272	0.9380
CG14063	CG14063	1626705_at	-0.0696	0.6643	-0.0413	0.7086	0.0525	0.7691	0.1290	0.7130	-0.0443	0.8249	-0.1733	0.2252	0.0654	0.9126	0.1033	0.6503	0.0379	0.8949
katanin-60	katanin 60	1626706_at	-0.1762	0.4718	-0.2494	0.1831	-0.4959	0.0287	-0.1350	0.7138	-0.0342	0.8788	0.1008	0.5391	0.1754	0.8541	0.0193	0.9784	-0.1561	0.7284
AnnIX	Annexin IX	1626707_a_at	-0.5000	0.0303	0.3447	0.2202	0.3583	0.0830	0.0905	0.8738	-0.1471	0.4795	-0.2376	0.1804	0.0938	0.9309	0.5432	0.1249	0.4493	0.2177
Cbl	Cbl	1626708_at	0.2836	0.2247	-0.2957	0.2141	-0.6052	0.0475	-0.2125	0.6916	0.4707	0.0615	0.6833	0.0083	0.1250	0.9152	-0.0230	0.9739	-0.1480	0.7541
CG30179	CG30179	1626709_at	-0.6225	0.0192	-0.0588	0.6712	-0.4411	0.0456	-0.1904	0.5690	-0.5005	0.0106	-0.3102	0.0483	0.1948	0.7550	-0.0896	0.7860	-0.2844	0.2964
SA	Stromalin	1626710_at	-0.3632	0.1088	-0.1003	0.4716	0.2844	0.0911	0.0006	0.9994	-0.1896	0.4164	-0.1902	0.3603	-0.2130	0.7142	0.2176	0.3615	0.4306	0.1116
---	---	1626711_at	-0.0487	0.8319	0.0779	0.8292	0.2030	0.3368	0.2159	0.7234	-0.0683	0.8482	-0.2842	0.2501	-0.0260	0.9816	-0.0971	0.7343	-0.0711	0.8166
---	---	1626712_at	0.3538	0.0655	-0.1094	0.4664	-0.0222	0.9054	0.1103	0.8000	0.0895	0.6505	-0.0209	0.9230	-0.0253	0.9816	-0.1164	0.6473	-0.0911	0.7351
CG6629	CG6629	1626713_at	0.2023	0.5340	0.0585	0.6915	0.0403	0.8525	0.0363	0.9819	0.0965	0.8437	0.0602	0.8947	-0.0848	0.8768	0.1119	0.6266	0.1967	0.3693
---	---	1626714_s_at	0.0081	0.9788	0.0837	0.4550	0.1875	0.2248	-0.1643	0.7579	-0.2251	0.3331	-0.0608	0.8140	-0.0104	0.9952	-0.0255	0.9697	-0.0151	0.9832
CG5084	CG5084	1626715_at	0.0663	0.6535	0.2194	0.4103	0.2866	0.2032	0.0839	0.9186	0.0556	0.8681	-0.0283	0.9272	-0.0199	0.9871	0.1531	0.6120	0.1731	0.5632
---	---	1626716_at	-0.0223	0.9435	-0.1574	0.1630	-0.0398	0.8643	0.0754	0.9005	0.0577	0.8112	-0.0177	0.9422	-0.0973	0.8736	0.0315	0.9350	0.1288	0.6189
CG14612	CG14612	1626717_at	0.3696	0.0387	0.3515	0.1268	0.2285	0.2820	-0.1703	0.6937	-0.1324	0.5361	0.0379	0.8725	0.0644	0.9309	0.0342	0.9286	-0.0301	0.9287
CG33127	CG33127	1626718_at	-0.1467	0.9538	-0.0064	0.9553	0.0281	0.9301	-0.0848	0.9883	-1.2551	0.3591	-1.1703	0.3410	-0.0185	0.9984	-0.9724	0.6098	-0.9539	0.6190
wls	wntless	1626719_a_at	-0.8387	0.0049	-1.1025	0.0112	-1.0203	0.0005	-0.0550	0.9196	0.2137	0.1976	0.2687	0.0733	-0.0713	0.9328	0.0223	0.9605	0.0936	0.7674
CG34049	CG34049	1626720_at	0.0707	0.6828	0.0331	0.7971	-0.1362	0.4298	-0.0486	0.9445	0.0236	0.9330	0.0722	0.7369	0.1738	0.7241	0.0482	0.8749	-0.1256	0.5808
CG11762	CG11762	1626721_at	0.0577	0.7764	-0.0094	0.9496	0.2598	0.1455	0.1319	0.7438	-0.0332	0.8902	-0.1651	0.3053	-0.0611	0.9467	-0.1082	0.7406	-0.0472	0.9025
CG14087	CG14087	1626722_at	0.1613	0.6497	0.1186	0.6516	-0.0120	0.9643	-0.1604	0.6013	-0.0025	0.9910	0.1579	0.2571	-0.1224	0.9360	-0.1405	0.8247	-0.0181	0.9832
Takr86C	Neuropeptide rece	1626723_at	-0.0183	0.9436	0.0222	0.8264	-0.0010	0.9972	-0.0361	0.9641	-0.1044	0.6670	-0.0682	0.7725	-0.0646	0.9308	-0.0793	0.7925	-0.0147	0.9678
CG32687	CG32687	1626724_at	1.1428	0.0625	1.2553	0.1864	1.8548	0.0016	0.2518	0.7556	-0.4576	0.1870	-0.7094	0.0305	-0.3662	0.9011	-0.3916	0.7642	-0.0253	0.9881
CG2970	CG2970	1626725_at	-0.3404	0.1176	0.2639	0.0478	0.3693	0.0473	0.1303	0.7031	-0.3958	0.0202	-0.5261	0.0034	0.0574	0.9408	0.1616	0.5131	0.1042	0.6955
---	---	1626726_at	-0.3160	0.0526	-0.2361	0.1624	-0.3532	0.0798	-0.0917	0.8716	0.0728	0.7582	0.1645	0.3675	-0.1008	0.8978	0.0345	0.9411	0.1353	0.6717
Mct1	Monocarboxylate	1626727_at	1.1961	0.2024	-0.0935	0.9222	0.1877	0.6772	0.0657	0.9559	0.5360	0.0956	0.4703	0.1011	-0.0656	0.9914	-0.6016	0.7012	-0.5360	0.7377
CG34150	CG34150	1626728_at	-0.0865	0.6467	0.0294	0.8038	-0.0400	0.8287	-0.0842	0.8872	-0.0667	0.7844	0.0175	0.9445	0.0142	0.9914	0.0553	0.9009	0.0412	0.9189
CG3803	CG3803	1626729_at	-0.3715	0.1251	-0.1306	0.6840	-0.2549	0.4314	-0.3119	0.2500	-0.0508	0.7965	0.2611	0.0716	-0.2428	0.8504	0.1552	0.8244	0.3980	0.4697
tna	tonalli	1626730_s_at	-1.3120	0.0568	-1.2727	0.1688	-1.2443	0.0069	0.0795	0.9130	0.1663	0.4895	0.0868	0.7228	0.0397	0.9939	0.0460	0.9841	0.0064	0.9979
Klc	kinesin	1626731_at	-0.5192	0.0578	-0.6569	0.0804	-0.5370	0.0499	0.1356	0.8493	0.3701	0.1529	0.2345	0.3234	0.1015	0.9174	0.3322	0.3134	0.2307	0.5137
CG30217	CG30217	1626732_at	0.0651	0.7529	0.1104	0.4311	0.0875	0.5923	-0.2492	0.5328	-0.1044	0.6574	0.1448	0.4638	-0.0552	0.9342	0.0556	0.8500	0.1108	0.6318
Ski6	Ski6	1626733_at	0.2334	0.3469	-0.0071	0.9917	0.0947	0.7487	0.3955	0.4167	0.8882	0.0062	0.4927	0.0469	0.1521	0.9412	0.5849	0.3486	0.4327	0.5173
CG6484	CG6484	1626734_at	-0.9308	0.0218	-0.0097	0.9779	-0.4352	0.1129	-0.8723	0.2500	-2.3348	0.0009	-1.4625	0.0039	-0.3675	0.6749	-1.2881	0.0158	-0.9206	0.0441
CG13551	CG13551	1626735_at	0.2998	0.1560	0.1127	0.3761	0.4041	0.1174	0.1475	0.7271	0.0208	0.9393	-0.1267	0.4834	0.0408	0.9831	-0.0455	0.9505	-0.0864	0.8897
ph-d	Polyhomeotic	1626736_at	0.0589	0.7858	0.1642	0.3871	0.3439	0.1875	0.0773	0.9102	0.0055	0.9870	-0.0718	0.7630	-0.2354	0.7644	0.0678	0.8879	0.3032	0.3719
Gr89a	Gr89a	1626737_at	0.2308	0.2482	0.0792	0.6081	0.1419	0.3920	-0.1742	0.6763	-0.1615	0.4260	0.0127	0.9593	-0.0950	0.8954	-0.0486	0.9064	0.0463	0.9016
CG4644	CG4644	1626738_at	-0.1604	0.5495	-0.2320	0.4853	-0.2265	0.2585	-0.1614	0.6513	-0.2861	0.0977	-0.1247	0.4427	-0.1398	0.9032	-0.2957	0.4899	-0.1560	0.7468
GluRIIC	glutamate recepto	1626739_s_at	-2.0281	0.0021	-2.7910	0.0106	-3.2355	0.0000	0.0016	0.9988	0.2170	0.3758	0.2154	0.3249	0.4382	0.7812	-0.5108	0.4572	-0.9490	0.1901
Cpr72Ec	CG4784	1626740_at	0.1556	0.4259	0.1152	0.4410	0.1737	0.3538	0.0495	0.9314	-0.1780	0.3010	-0.2275	0.1337	0.1421	0.8388	0.1444	0.6437	0.0022	0.9965
CG12683	CG12683	1626741_at	0.1086	0.5596	0.1856	0.1728	0.3676	0.0370	-0.0401	0.9518	-0.1132	0.5696	-0.0731	0.7076	-0.1899	0.7633	-0.0198	0.9639	0.1701	0.5420
CG8964	CG8964	1626742_at	-0.4300	0.2102	0.2294	0.1584	0.3382	0.1483	-0.2922	0.6202	-0.9488	0.0068	-0.6566	0.0207	-0.3455	0.7220	-0.1741	0.7090	0.1715	0.7128
CG8543	CG8543	1626743_at	0.4369	0.0714	0.2643	0.1951	0.3607	0.1175	0.0181	0.9860	0.0670	0.8314	0.0489	0.8654	-0.0026	0.9990	-0.0309	0.9528	-0.0283	0.9508
---	---	1626744_at	0.2098	0.2850	0.1019	0.4367	0.2330	0.1096	0.0816	0.8837	0.1416	0.4762	0.0600	0.7758	-0.0546	0.9467	0.0047	0.9935	0.0593	0.8558
Sucb	Sucb	1626745_at	0.1960	0.3800	0.1047	0.7201	-0.0328	0.8710	0.1258	0.7298	0.2337	0.1497	0.1079	0.4888	0.1520	0.8558	0.1728	0.6425	0.0207	0.9686
---	---	1626746_at	0.1833	0.2534	-0.0374	0.8090	0.1459	0.4815	0.0152	0.9819	0.0050	0.9827	-0.0101	0.9597	0.0449	0.9752	-0.0146	0.9829	-0.0595	0.9033

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Pros28.1A	20s proteasome	1626747_at	0.0127	0.9535	-0.0383	0.7554	-0.0859	0.6656	0.0567	0.9255	0.0593	0.7913	0.0026	0.9910	0.0053	0.9952	0.0166	0.9620	0.0113	0.9726
dpr9	dpr9	1626748_at	0.2509	0.2054	0.0680	0.7918	0.3430	0.0747	0.1650	0.6936	0.0579	0.8091	-0.1072	0.5787	-0.2032	0.7768	-0.2041	0.5246	-0.0009	0.9990
CG18135	CG18135	1626749_a_at	0.0445	0.9247	0.3763	0.1896	-0.7465	0.0071	-0.7888	0.1140	-0.5309	0.0786	0.2580	0.3487	0.3494	0.7686	-0.1393	0.8372	-0.4888	0.3468
CG30479	CG30479	1626750_at	0.3068	0.3301	0.0874	0.6241	0.3299	0.0727	-0.0097	0.9917	0.0264	0.9226	0.0360	0.8758	-0.0256	0.9848	0.0257	0.9599	0.0514	0.9045
CG4603	CG4603	1626751_at	0.0280	0.9009	0.1139	0.2762	0.3146	0.1246	0.0724	0.8698	-0.1068	0.5156	-0.1792	0.1951	-0.0606	0.9515	0.0619	0.8867	0.1225	0.7185
mTTF	mitochondrial tran	1626752_at	1.0553	0.0035	1.6803	0.0162	1.4365	0.0001	0.0847	0.8899	0.2365	0.2498	0.1517	0.4305	0.3457	0.7506	0.8400	0.0834	0.4944	0.3011
CG12947	CG12947	1626753_at	-1.1813	0.0019	0.0556	0.7248	0.3904	0.1225	0.0960	0.8817	-1.2145	0.0007	-1.3105	0.0003	-0.2879	0.8235	-0.0756	0.9289	0.2123	0.7294
---	---	1626754_at	-0.0632	0.7195	0.0954	0.5485	-0.2382	0.4115	-0.2130	0.6327	0.0349	0.9084	0.2479	0.2040	0.1163	0.8655	-0.0017	0.9989	-0.1180	0.7007
CG30085	CG30085	1626755_at	0.1904	0.5674	-0.2801	0.7157	-0.2766	0.5505	-0.0661	0.9117	0.4537	0.0247	0.5198	0.0084	-0.1239	0.9814	0.0259	0.9924	0.1498	0.9231
RhoGEF3	RhoGEF3	1626756_a_at	-0.7883	0.1155	-2.5410	0.0022	-2.6087	0.0003	0.1478	0.9295	1.2775	0.0172	1.1297	0.0179	0.2226	0.9075	-0.5256	0.4392	-0.7482	0.2862
---	---	1626757_at	0.0383	0.8301	0.0267	0.8026	0.2943	0.1376	0.1752	0.7803	0.0359	0.9220	-0.1393	0.5935	-0.0309	0.9771	0.1199	0.6686	0.1508	0.5768
---	---	1626758_at	0.2036	0.2883	-0.1273	0.4558	-0.0210	0.9143	0.2123	0.5832	0.1713	0.3965	-0.0410	0.8585	-0.0363	0.9677	-0.1112	0.6673	-0.0749	0.7895
CG34136	CG34136	1626759_at	1.4408	0.0117	-0.1174	0.9096	0.1324	0.7889	0.4859	0.6360	0.5122	0.3150	0.0263	0.9674	0.2057	0.9499	-1.0356	0.2722	-1.2414	0.2199
CG2974	CG2974	1626760_at	0.5998	0.0649	0.4714	0.1001	0.2253	0.1606	-0.2668	0.5948	0.0156	0.9686	0.2824	0.2149	0.0733	0.9503	-0.0892	0.8519	-0.1626	0.6689
---	---	1626761_at	0.1362	0.4386	-0.1225	0.2793	0.1296	0.5585	0.1636	0.6578	0.0964	0.6247	-0.0672	0.7220	-0.0956	0.8270	-0.1281	0.4879	-0.0325	0.8951
cin	cinnamon	1626762_s_at	-0.3004	0.1405	0.2021	0.3638	-0.0139	0.9623	-0.1987	0.6990	-0.6084	0.0182	-0.4098	0.0563	0.0298	0.9816	-0.1128	0.7200	-0.1426	0.6300
veli	veli	1626763_a_at	-0.3093	0.1493	-0.3427	0.2311	-0.6679	0.0078	-0.1285	0.7608	0.2008	0.2717	0.3294	0.0481	0.2369	0.7953	0.1832	0.6656	-0.0537	0.9194
CG10182	CG10182	1626764_at	0.0440	0.8544	-0.0441	0.6653	0.0156	0.9466	0.1298	0.8281	0.2981	0.1927	0.1683	0.4365	0.0115	0.9884	-0.0367	0.8812	-0.0482	0.8216
CG6478 /// DyakCG6478	CG6478	1626765_at	0.0987	0.5255	0.2087	0.2139	0.3637	0.0597	0.0599	0.9017	-0.0584	0.7562	-0.1183	0.4225	-0.0735	0.9171	0.0261	0.9477	0.0995	0.7192
Alp23B	dActivin2	1626766_s_at	1.7609	0.0196	1.5568	0.0121	2.3399	0.0009	0.2138	0.8671	0.4167	0.3646	0.2029	0.6636	-0.4790	0.8049	0.3481	0.7027	0.8271	0.3163
Tsp2A	Tetraspanin 2A	1626767_at	0.0596	0.9748	-0.0159	0.9713	0.1335	0.7206	-0.0608	0.9893	-0.6422	0.5663	-0.5814	0.5724	0.0504	0.9924	-0.6210	0.6416	-0.6714	0.6130
CG7044	CG7044	1626768_at	0.5392	0.1138	0.2898	0.1801	0.4416	0.0379	0.1891	0.6970	0.4489	0.0498	0.2598	0.1930	0.0767	0.9589	0.2998	0.4810	0.2231	0.6190
CG1553	CG1553	1626769_s_at	-0.4055	0.1427	-0.3206	0.1909	-1.0502	0.0224	-0.3982	0.4442	-1.1483	0.0025	-0.7500	0.0100	0.0771	0.9775	-1.1323	0.0890	-1.2094	0.0953
---	---	1626770_at	0.2477	0.2012	-0.0266	0.8719	-0.0773	0.7865	0.0798	0.8949	0.2422	0.2273	0.1624	0.3833	0.0121	0.9916	0.0866	0.8008	0.0745	0.8255
spri	Sprint	1626771_at	0.1489	0.3946	0.3727	0.2979	0.3549	0.0572	-0.0217	0.9734	-0.1329	0.4223	-0.1113	0.4637	0.0561	0.9727	0.1859	0.6850	0.1298	0.7931
---	---	1626772_at	0.0667	0.7425	0.0000	1.0000	0.2230	0.1394	0.2589	0.5008	0.0576	0.8202	-0.2014	0.2718	0.1389	0.8076	-0.0581	0.8611	-0.1970	0.4148
---	---	1626773_s_at	-0.9775	0.3850	0.0307	0.7596	-0.2408	0.1430	-0.1077	0.9777	-0.8691	0.3940	-0.7614	0.4123	-0.0466	0.9860	-0.1858	0.8084	-0.1392	0.8577
robo	Roundabout 1	1626774_s_at	-0.1681	0.6229	-0.0015	0.9949	-0.2017	0.3702	-0.0778	0.9380	0.0935	0.7961	0.1713	0.5545	-0.0295	0.9869	0.1802	0.6961	0.2097	0.6344
CG41447	CG41447	1626775_at	0.1918	0.3098	0.1393	0.2788	0.1007	0.6677	-0.0317	0.9610	-0.0436	0.8470	-0.0119	0.9573	0.0186	0.9913	-0.1245	0.7640	-0.1431	0.7128
CG10474	CG10474	1626776_at	0.1343	0.5634	0.1771	0.2623	0.1247	0.4444	-0.2068	0.6988	-0.1357	0.6173	0.0711	0.7968	-0.0214	0.9742	0.0388	0.8678	0.0602	0.7514
---	---	1626777_at	0.1707	0.2849	-0.3797	0.0597	-0.2316	0.2083	0.1347	0.7743	0.4805	0.0228	0.3458	0.0536	-0.0214	0.9772	-0.0773	0.6964	-0.0560	0.7915
---	---	1626778_at	0.1868	0.3952	0.0011	0.9932	0.2456	0.1341	-0.0008	0.9994	0.0928	0.6920	0.0936	0.6578	-0.1733	0.7726	-0.1706	0.5234	0.0027	0.9953
---	---	1626779_at	0.0413	0.7930	0.0378	0.7049	0.0344	0.9273	0.1047	0.8794	0.1162	0.6642	0.0115	0.9689	0.1675	0.8086	0.0372	0.9353	-0.1302	0.6817
CG4678	CG4678	1626780_s_at	-0.5324	0.0369	-0.4342	0.0223	-0.1625	0.6049	0.2231	0.7217	0.0281	0.9466	-0.1950	0.4670	-0.2131	0.8202	-0.1731	0.6861	0.0400	0.9410
Pp2B-14D	calcineurin	1626781_at	-0.3012	0.4185	-0.1141	0.8470	0.1408	0.5975	0.0261	0.9639	-0.1323	0.3988	-0.1583	0.2480	-0.0740	0.9831	0.0581	0.9650	0.1321	0.9057
CG17331	CG17331	1626782_at	0.3298	0.2332	0.4743	0.0843	0.5146	0.0106	0.0313	0.9649	0.0500	0.8372	0.0187	0.9364	-0.0220	0.9914	0.2337	0.6109	0.2556	0.5762
CG16756 /// DyakCG16756	CG16756	1626783_at	2.9643	0.0026	1.3708	0.1798	3.2114	0.0005	0.6228	0.6560	-0.4841	0.4929	-1.1069	0.0685	-0.6666	0.7997	-1.5839	0.1410	-0.9173	0.4093
---	---	1626784_at	0.0950	0.6610	0.1507	0.3678	-0.0034	0.9872	-0.1793	0.5593	-0.2219	0.1578	-0.0426	0.8106	0.0079	0.9950	0.0062	0.9925	-0.0017	0.9979
Nap1	Nucleosome asse	1626785_at	0.1562	0.4204	1.0623	0.0070	0.8029	0.0494	0.0717	0.9482	-0.8015	0.0178	-0.8732	0.0074	0.2619	0.7953	0.0358	0.9590	-0.2261	0.6205
CG8494	CG8494	1626786_at	-0.2930	0.1427	-0.0347	0.8777	0.1696	0.3900	0.0685	0.9037	-0.1799	0.3360	-0.2484	0.1295	-0.0630	0.9568	0.2147	0.5342	0.2777	0.4114
CG17556 /// CG3678	CG17556 /// CG3(	1626787_s_at	0.0796	0.6713	0.6010	0.0929	0.3208	0.0440	-0.2349	0.3801	-0.0076	0.9743	0.2273	0.0929	0.1191	0.8991	0.5596	0.1124	0.4404	0.2202
CstF-50	CstF-50	1626788_a_at	0.4685	0.0182	0.2735	0.6000	0.8784	0.0029	0.2360	0.5552	0.3459	0.0945	0.1100	0.5953	-0.3481	0.7768	0.1182	0.8764	0.4664	0.3899
CG13618 /// DyakCG13618	CG13618	1626789_at	0.1112	0.9632	-1.4865	0.0186	-0.6263	0.0832	0.9192	0.2426	0.0841	0.8943	-0.8352	0.0495	0.0123	0.9994	-1.7176	0.4875	-1.7300	0.4889
Fak56D	Focal Adhesion Ki	1626790_s_at	0.3138	0.2746	0.3623	0.4424	0.2711	0.1599	-0.0564	0.9436	0.2384	0.2972	0.2948	0.1445	0.0726	0.9734	0.3164	0.5776	0.2438	0.6798
CG11406	CG11406	1626791_at	0.1330	0.4662	-0.0247	0.8480	-0.1700	0.2998	0.0653	0.9075	0.0866	0.6743	0.0213	0.9235	0.2593	0.6695	0.0628	0.8518	-0.1966	0.4353
---	---	1626792_at	-0.1207	0.4457	-0.2903	0.3381	-0.2339	0.1260	0.2852	0.5213	0.4647	0.0515	0.1794	0.4108	0.1120	0.8049	0.2024	0.2664	0.0904	0.6581
CG30427	CG30427	1626793_at	0.3136	0.2252	0.0494	0.8114	0.0530	0.8057	0.1620	0.7281	0.2028	0.3392	0.0408	0.8672	0.0373	0.9816	-0.1702	0.6834	-0.2075	0.6055
CG40486	CG40486	1626794_at	-0.0874	0.9027	-0.1476	0.6000	-0.4812	0.0306	-0.3901	0.7253	-0.5304	0.2922	-0.1403	0.8030	-0.1594	0.8997	-0.5285	0.2395	-0.3691	0.4399
---	---	1626795_at	-0.0699	0.6991	-0.0388	0.7030	-0.0131	0.9454	0.1065	0.8281	0.0687	0.7617	-0.0379	0.8632	0.0605	0.9309	0.0545	0.8625	-0.0060	0.9868
---	---	1626796_at	0.1569	0.4948	0.0610	0.7280	-0.0032	0.9910	-0.1213	0.7929	0.0036	0.9899	0.1249	0.4870	-0.0791	0.9515	-0.2106	0.6162	-0.1315	0.7764
Ntl	CG7075	1626797_a_at	-0.4593	0.0587	-0.6286	0.0959	-0.8530	0.0078	0.0162	0.9921	0.4173	0.2080	0.4011	0.1765	-0.0355	0.9672	-0.1074	0.6682	-0.0719	0.7923
Bsg25D	Blastoderm-specif	1626798_a_at	-0.4642	0.2315	-0.1164	0.3679	-0.9262	0.0819	-0.3180	0.6114	-0.2147	0.5194	0.1033	0.7640	0.5167	0.7726	0.1548	0.8926	-0.3619	0.6689
CG16758	CG16758	1626799_a_at	-0.4787	0.0819	-0.4442	0.3636	-0.6913	0.0170	-0.0005	0.9996	-0.0759	0.7864	-0.0754	0.7656	-0.0570	0.9816	-0.1972	0.7587	-0.1402	0.8372
CG11630	CG11630	1626800_at	0.2531	0.0964	-0.0230	0.8851	-0.0830	0.6484	-0.0507	0.9339	0.0866	0.6715	0.1373	0.4124	0.1039	0.8446	-0.0573	0.8468	-0.1612	0.4747
CG10638	CG10638	1626801_at	-0.0890	0.5327	-0.3429	0.0323	-0.4023	0.1287	0.0076	0.9943	0.6841	0.0084	0.6765	0.0053	-0.0722	0.9342	0.3160	0.2424	0.3883	0.1901
Sh	Shaker	1626802_a_at	-1.3006	0.0168	-1.0265	0.0791	-1.7219	0.0006	-0.3140	0.7733	0.3065	0.5296	0.6205	0.1274	-0.0221	0.9916	-0.0378	0.9629	-0.0156	0.9847
---	---	1626803_at	0.2432	0.2894	0.0508	0.8227	0.2101	0.2537	0.1501	0.7854	0.1651	0.4881	0.0150	0.9586	0.0079	0.9952	-0.0587	0.8963	-0.0666	0.8674
CG5379	CG5379	1626804_at	5.5112	0.0016	4.3017	0.0022	7.9120	0.0000	5.6398	0.0031	2.9264	0.0040	-2.7134	0.0033	1.6961	0.3482	1.4424	0.1321	-0.2537	0.8336
gish	gilgamesh	1626805_s_at	0.3460	0.1574	0.7746	0.0433	0.5062	0.0137	-0.0053	0.9951	0.0837	0.6685	0.0890	0.6083	0.3007	0.7628	0.5612	0.1633	0.2605	0.5518
tra	transformer	1626806_at	-0.0728	0.7096	-0.2387	0.1658	-0.3306	0.1804	0.1093	0.7838	0.2626	0.1092	0.1533	0.3045	0.1569	0.8270	0.0151	0.9778	-0.1419	0.6621
Nhe3	Nhe3	1626807_a_at	-0.2023	0.5240	0.1459	0.3610	-0.4418	0.0768	-0.5556	0.1515	-0.7179	0.0085	-0.1623	0.4645	-0.0195	0.9914	-0.3435	0.3993	-0.3240	0.4449
Rho1	rho1 kinase	1626808_s_at	-0.1186	0.5179	0.3918	0.0349	0.0822	0.6406	0.0756	0.8761	0.0087	0.9735	-0.0670	0.7046	0.3764	0.5191	0.5159	0.0782	0.1396	0.6414
CG33980	CG33980	1626809_at	0.1196	0.5968	0.0557	0.7167	0.0869	0.7058	0.0188	0.9838	0.0370	0.9015	0.0182	0.9473	0.1243	0.8049	0.0842	0.7275	-0.0400	0.8893
CG4400 /// DyakCG4400	CG4400	1626810_at	-0.3382	0.1622	0.2156	0.1716	0.1338	0.3407	0.1526	0.7618	-0.1689	0.4535	-0.3215	0.0963	0.1228	0.8379	0.3085	0.1962	0.1857	0.4677
CG5510	CG5510	1626811_at	-0.0502	0.8070	-0.2046	0.4927	-0.0297	0.8606	0.0928	0.8507	0.3501	0.0554	0.2573	0.1074	-0.1533	0.8331	0.1932	0.5418	0.3465	0.2707
CG14132	CG14132	1626812_at	-0.4321	0.1658	-1.8950	0.0080	-1.1346	0.0024	0.7355	0.3005	1.1397	0.0142	0.4042	0.2840	0.0462	0.9790	-0.1423	0.7741	-0.1885	0.6706
Smn	survival motor neu	1626813_at	-0.0044	0.9897	-0.4783	0.0721	-0.2205	0.2115	0.1132	0.8064	0.6190	0.0056	0.5058	0.0083	-0.1825	0.8454	0.0497	0.9353	0.2322	0.5688
CG14625	CG14625	1626814_at	-0.1221	0.5340	0.1888	0.2092	0.3194	0.1631	-0.0497	0.9558	-0.3528	0.1447	-0.3031	0.1618	-0.0620	0.9400	0.0704	0.8353	0.1325	0.6280
CG9264	CG9264	1626815_a_at	-0.6655	0.1087	-0.1317	0.4063	-0.1688	0.2846	-0.1853	0.8705	-0.7863	0.0525	-0.6010	0.0906	0.0842	0.9552	-0.4181	0.3268	-0.5023	0.2674

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1626816_at	0.0652	0.7428	0.2694	0.3363	0.4441	0.0150	0.1523	0.8422	-0.0665	0.8581	-0.2188	0.4097	0.0422	0.9717	0.1271	0.7064	0.0849	0.8199
Rrp45	Rrp45	1626817_at	0.6050	0.0767	0.0003	1.0000	0.5232	0.0144	0.0897	0.9042	0.4742	0.0554	0.3845	0.0779	-0.4019	0.6749	-0.1683	0.7041	0.2336	0.5755
CG13097	CG13097	1626818_at	0.4248	0.1672	0.1240	0.7746	0.6880	0.0052	0.3684	0.4532	0.6063	0.0304	0.2379	0.3283	-0.2234	0.8379	0.2925	0.5312	0.5159	0.2674
CG11552	CG11552	1626819_s_at	0.0172	0.9535	0.1100	0.3769	0.3520	0.0704	0.1907	0.6890	-0.0342	0.9114	-0.2249	0.2583	-0.0713	0.9309	0.0795	0.8197	0.1508	0.6005
CG5169	CG5169	1626820_at	-0.2223	0.6072	-0.0309	0.8560	-0.1098	0.6099	0.1608	0.7513	0.2392	0.2805	0.0784	0.7413	0.2141	0.8689	0.4680	0.3486	0.2539	0.6452
Hsp70Ba /// Hsp70Bb /// Hs	Hsp70Bb /// heat	1626821_s_at	-0.9128	0.8143	-1.6143	0.0574	-1.3912	0.6125	1.8750	0.8485	3.3995	0.3526	1.5245	0.6854	1.7639	0.8439	2.7548	0.4556	0.9910	0.8306
---	---	1626822_s_at	-0.0153	0.9296	0.0177	0.8647	0.0617	0.7690	-0.0300	0.9666	-0.0907	0.6718	-0.0607	0.7701	-0.0130	0.9872	0.0347	0.9019	0.0477	0.8401
CG7945	CG7945	1626823_a_at	-0.4780	0.0443	-0.0082	0.9810	0.0817	0.5975	0.0832	0.8791	-0.3373	0.0729	-0.4206	0.0198	-0.0289	0.9831	0.1353	0.7043	0.1642	0.6294
---	---	1626824_at	0.0429	0.8510	0.2231	0.4417	0.1959	0.1857	-0.1021	0.8578	-0.0699	0.7783	0.0321	0.8966	0.0155	0.9862	0.1019	0.6567	0.0864	0.7148
---	---	1626825_at	-0.0076	0.9742	-0.0037	0.9813	0.0655	0.8409	0.1618	0.7278	0.2889	0.1629	0.1271	0.5312	-0.0155	0.9922	0.1424	0.7350	0.1579	0.6941
ng1 /// ng2	new glue 2 /// nes	1626826_s_at	0.0247	0.8892	0.0656	0.6044	-0.0834	0.6600	0.0353	0.9484	0.1196	0.4524	0.0842	0.5809	0.2025	0.7116	0.1533	0.5181	-0.0491	0.8736
CG12480	CG12480	1626827_at	0.1648	0.5820	0.0707	0.6122	0.1016	0.6160	0.0018	0.9983	0.0349	0.8918	0.0331	0.8823	-0.0956	0.8906	-0.0115	0.9814	0.0841	0.7873
---	---	1626828_at	-0.1011	0.6265	-0.2192	0.1086	-0.0537	0.7499	0.0969	0.8578	0.1179	0.5745	0.0210	0.9295	-0.0727	0.9238	-0.0539	0.8830	0.0188	0.9595
CG3494	CG3494	1626829_s_at	0.1591	0.4935	0.3868	0.1648	0.0499	0.8003	-0.2736	0.5140	-0.3284	0.1366	-0.0548	0.8284	0.3117	0.6272	0.1534	0.5897	-0.1583	0.5792
Lcp1	cuticle p. l	1626830_at	-0.0395	0.8513	0.0411	0.7511	0.1074	0.6120	0.1722	0.7441	0.0160	0.9638	-0.1562	0.4796	0.1530	0.7423	0.0832	0.7087	-0.0698	0.7578
CG6579	CG6579	1626831_at	-0.5313	0.0217	0.3840	0.0751	-0.4916	0.0898	-0.0471	0.9266	-0.2027	0.1841	-0.1556	0.2598	0.8247	0.3362	0.7014	0.1248	-0.1233	0.8287
CG31343	CG31343	1626832_at	-0.6235	0.8800	0.3412	0.6794	-4.1016	0.0001	-3.6688	0.2941	-3.3759	0.1005	0.2928	0.9063	0.6927	0.9590	-2.1815	0.5989	-2.8742	0.4755
CG11839	CG11839	1626833_at	0.2021	0.3238	0.6767	0.1709	0.3061	0.0693	-0.3612	0.5175	-0.2972	0.3185	0.0641	0.8513	0.0190	0.9913	0.2468	0.4997	0.2277	0.5445
eIF4E-5	eIF4E-5	1626834_at	0.2145	0.1865	0.0717	0.6130	0.0271	0.8795	-0.0526	0.9223	0.0364	0.8656	0.0890	0.5839	-0.1236	0.8283	-0.1190	0.6464	0.0046	0.9914
Syb	synaptobrevin	1626835_at	0.1890	0.3655	1.1439	0.0091	0.8624	0.0033	-0.0127	0.9893	-0.5295	0.0219	-0.5167	0.0152	0.2625	0.7506	0.4697	0.1718	0.2072	0.5821
nub	CG15488	1626836_at	0.1248	0.5842	0.3786	0.1314	0.2524	0.0801	-0.1991	0.6122	-0.2720	0.1661	-0.0729	0.7305	0.0708	0.9391	-0.0141	0.9782	-0.0849	0.8098
CG32245	CG32245	1626837_a_at	-1.3449	0.0032	-0.1532	0.4405	-0.4749	0.0736	-0.2879	0.6919	-1.4057	0.0022	-1.1178	0.0035	-0.0319	0.9722	-0.0206	0.9567	0.0113	0.9758
---	---	1626838_at	-0.0559	0.7509	0.2072	0.1668	0.1831	0.3846	0.0258	0.9744	-0.1712	0.3899	-0.1970	0.2599	0.1098	0.8473	0.1653	0.4981	0.0555	0.8577
blot	bloated tubules	1626839_s_at	0.7979	0.0372	0.1320	0.3733	0.2228	0.3530	0.4501	0.3016	1.0223	0.0027	0.5722	0.0207	0.3107	0.8000	0.3878	0.4495	0.0771	0.9125
opa1-like	optic atrophy 1-lik	1626840_a_at	-0.4525	0.0554	-0.0683	0.8030	0.1009	0.6737	-0.0942	0.9020	-0.3814	0.1231	-0.2872	0.1952	-0.1320	0.8270	0.0555	0.8764	0.1875	0.4677
ck	myosin VIIa	1626841_s_at	-0.5544	0.0400	-0.8311	0.0562	-1.1622	0.0006	-0.2416	0.7143	0.4320	0.1498	0.6736	0.0207	0.0294	0.9822	0.1085	0.7710	0.0791	0.8393
l(3)82Fd	late puff gene at 8	1626842_a_at	-1.5191	0.0226	-0.4691	0.6772	-1.0478	0.0134	-0.4788	0.5492	-0.7592	0.0691	-0.2804	0.4757	0.0336	0.9946	0.2928	0.8550	0.2592	0.8673
---	---	1626843_at	0.2784	0.1905	-0.0253	0.8363	0.1128	0.4348	-0.1813	0.5633	-0.0780	0.6709	0.1033	0.5060	-0.1538	0.7506	-0.2330	0.2437	-0.0792	0.7411
Hsf	Heat shock factor	1626844_at	-0.1039	0.5870	-0.1807	0.4072	-0.2608	0.1127	0.0068	0.9941	0.2387	0.2053	0.2319	0.1690	0.1314	0.8270	0.2700	0.2593	0.1386	0.6043
CG8351	CG8351	1626845_at	-0.1680	0.6443	0.0606	0.7680	0.1628	0.5312	0.5043	0.4386	0.6272	0.0797	0.1228	0.7514	0.3934	0.6695	0.8123	0.0509	0.4189	0.2757
CG5973	CG5973	1626846_s_at	-0.8280	0.1745	1.1140	0.1650	-0.1761	0.6325	-0.8993	0.1772	-2.1056	0.0009	-1.2063	0.0057	0.5371	0.8222	0.1114	0.9450	-0.4257	0.7010
---	---	1626847_at	0.3651	0.1799	-0.1586	0.3103	0.0192	0.9263	0.2547	0.4908	0.4685	0.0255	0.2138	0.2258	0.0038	0.9969	-0.1848	0.4391	-0.1886	0.4446
---	---	1626848_at	0.2177	0.3111	0.0194	0.9233	-0.0136	0.9518	0.0089	0.9888	-0.0632	0.7110	-0.0721	0.6322	0.0813	0.9432	-0.2125	0.5618	-0.2938	0.4080
CG7251	CG7251	1626849_at	0.0870	0.5707	0.1747	0.1368	0.2613	0.1946	0.0031	0.9962	-0.0534	0.8482	-0.0565	0.8165	0.0265	0.9721	0.0430	0.8764	0.0164	0.9517
CG31953 /// gkt	CG31953 /// glaiki	1626850_s_at	0.2464	0.3715	0.4242	0.0675	0.8389	0.0016	0.3116	0.3222	0.3839	0.0423	0.0724	0.7092	-0.0480	0.9717	0.5210	0.1287	0.5690	0.1271
CG9460	CG9460	1626851_at	0.1678	0.4954	-0.1764	0.2670	0.2531	0.3219	0.1511	0.8441	0.3517	0.2138	0.2006	0.4543	-0.1734	0.8617	-0.0926	0.8721	0.0809	0.8837
CG40241	CG40241	1626852_at	0.1095	0.4968	-0.0301	0.8020	0.2597	0.1804	0.0993	0.7931	0.1594	0.3072	0.0602	0.7138	-0.2347	0.6955	-0.0862	0.7787	0.1485	0.5767
CG14017	CG14017	1626853_a_at	0.0328	0.8284	0.1857	0.4335	0.1011	0.6600	-0.0613	0.9319	0.0080	0.9804	0.0692	0.7689	-0.0085	0.9943	-0.0104	0.9848	-0.0019	0.9974
CG32148	CG32148	1626854_at	-0.0341	0.9135	-0.0827	0.5651	-0.2185	0.3363	-0.0629	0.9393	0.0035	0.9918	0.0663	0.8059	-0.0217	0.9816	-0.0452	0.8782	-0.0235	0.9348
CG31263	CG31263	1626855_at	0.0660	0.6381	0.0188	0.9350	0.3176	0.0578	0.1834	0.7138	-0.0133	0.9702	-0.1966	0.3474	-0.1144	0.8425	0.0900	0.7456	0.2044	0.3953
CG12560	CG12560	1626856_at	0.6545	0.0333	0.4687	0.0599	0.7722	0.0033	0.3256	0.3723	0.7341	0.0044	0.4085	0.0338	-0.0232	0.9898	0.4559	0.2298	0.4791	0.2391
CG4408	CG4408	1626857_at	2.9537	0.0025	1.2909	0.1641	2.6513	0.0001	0.4912	0.3512	0.9423	0.0079	0.4511	0.0947	-0.7726	0.7628	-0.5081	0.6664	0.2645	0.8472
CG5071	CG5071	1626858_a_at	0.2674	0.3194	-0.1384	0.5032	0.1074	0.4844	0.1278	0.7857	0.2619	0.1748	0.1341	0.4660	-0.2445	0.7193	-0.2190	0.4367	0.0255	0.9486
CG32374	CG32374	1626859_at	0.1581	0.5028	0.0284	0.7731	0.3968	0.0296	-0.0841	0.8883	-0.0297	0.9166	0.0544	0.8106	-0.2626	0.6903	-0.1293	0.6670	0.1334	0.6486
---	---	1626860_at	0.1223	0.6166	0.0744	0.7060	0.0458	0.8234	0.0202	0.9838	-0.0309	0.9220	-0.0512	0.8443	0.0144	0.9898	-0.1040	0.7019	-0.1184	0.6475
Octbeta3R	Octbeta3R	1626861_at	0.1718	0.2087	0.0526	0.6960	0.0033	0.9892	0.0807	0.8550	-0.0125	0.9580	-0.0931	0.5534	0.0649	0.9156	0.0346	0.9170	-0.0304	0.9180
Su(var)3-7	Suppressor of var	1626862_at	0.0272	0.9540	-0.4928	0.1470	-0.8162	0.0145	-0.1473	0.8217	0.5080	0.0512	0.6553	0.0114	0.2543	0.8331	0.0399	0.9620	-0.2145	0.7040
---	---	1626863_at	0.1915	0.2442	-0.1216	0.4684	-0.1275	0.4264	0.0264	0.9695	0.4438	0.0209	0.4174	0.0168	0.0348	0.9672	0.0512	0.8724	0.0164	0.9596
Pmm45A	Phosphomannom	1626864_at	-0.2933	0.2604	0.3745	0.0786	0.6690	0.0065	0.4467	0.3056	-0.0708	0.8272	-0.5174	0.0316	0.1281	0.8425	0.5615	0.0558	0.4334	0.1313
---	---	1626865_at	0.3321	0.1539	0.1194	0.3709	0.1128	0.7155	0.0476	0.9603	0.1638	0.5495	0.1163	0.6590	0.1089	0.8875	-0.0550	0.8999	-0.1639	0.5984
---	---	1626866_at	-0.0216	0.9311	0.0344	0.7967	0.2337	0.1331	0.0919	0.8362	0.0300	0.8957	-0.0619	0.7305	-0.0730	0.9238	0.0129	0.9772	0.0860	0.7741
His3:CG31613 /// His3:CG	His3:CG31613 ///	1626867_at	-0.4766	0.6063	-2.7440	0.0405	-1.8665	0.0087	0.1756	0.9182	1.6415	0.0079	1.4660	0.0079	-0.7839	0.8292	-0.6706	0.6952	0.1133	0.9592
CG17687 /// DmirCG17687	CG17687	1626868_at	0.1809	0.3206	0.0948	0.4570	0.1884	0.2911	-0.0479	0.9305	-0.0639	0.7364	-0.0159	0.9364	-0.1006	0.8534	-0.0139	0.9701	0.0867	0.7371
CG31619	CG31619	1626869_a_at	-0.6497	0.0383	0.0209	0.8453	0.0718	0.6853	-0.0854	0.8899	-0.7587	0.0038	-0.6732	0.0039	0.0100	0.9913	0.1149	0.5609	0.1049	0.6045
---	---	1626870_at	0.1129	0.6562	0.0591	0.6579	0.2068	0.1794	0.0211	0.9733	0.0272	0.8985	0.0061	0.9750	-0.1512	0.8276	-0.1178	0.7259	0.0334	0.9347
icln	icln	1626871_at	-0.1259	0.5493	0.3820	0.2016	0.2433	0.5504	-0.2283	0.8046	0.4074	0.2709	0.6357	0.0576	-0.2251	0.8940	0.9366	0.1345	1.1618	0.1000
CG14867	CG14867	1626872_at	0.1460	0.3671	0.1707	0.2343	0.2239	0.1951	-0.1454	0.7492	-0.1637	0.4240	-0.0182	0.9408	0.0379	0.9589	0.0985	0.6686	0.0606	0.8159
CG12448	CG12448	1626873_a_at	0.1653	0.5452	0.0921	0.4824	0.2966	0.0855	0.1835	0.7409	0.0610	0.8471	-0.1225	0.6220	0.0269	0.9775	0.0592	0.8478	0.0323	0.9176
Eip63F-1	Ecdysone-induce	1626874_at	-1.2371	0.0030	-0.0218	0.9666	-1.5742	0.0009	-0.9660	0.0460	-1.3625	0.0011	-0.3965	0.1064	0.5870	0.6390	-0.0511	0.9505	-0.6381	0.2282
CG13758 /// DanaCG1375	CG13758	1626875_at	0.8399	0.1054	1.0208	0.0793	2.0720	0.0001	0.1388	0.8146	-0.3951	0.0908	-0.5339	0.0185	-0.6273	0.7215	-0.1323	0.9075	0.4950	0.5266
CG40119	CG40119	1626876_x_at	-0.1398	0.3189	-0.2210	0.1761	-0.1260	0.4759	0.1175	0.7451	0.2729	0.0866	0.1554	0.2785	-0.0480	0.9677	0.0277	0.9563	0.0757	0.8495
---	---	1626877_at	0.1146	0.5477	0.1262	0.3938	0.1297	0.5975	-0.0117	0.9922	-0.0249	0.9459	-0.0133	0.9666	-0.0690	0.8885	-0.0298	0.9175	0.0392	0.8761
CG13811	CG13811	1626878_at	0.0393	0.8512	0.0806	0.7242	0.1005	0.6031	0.0367	0.9688	-0.0471	0.8859	-0.0837	0.7528	0.0105	0.9928	0.0056	0.9929	-0.0049	0.9921
CG15198 /// DyakCG15198	CG15198	1626879_at	0.1651	0.4180	0.2543	0.2866	0.3544	0.0624	-0.0150	0.9854	-0.1291	0.5102	-0.1140	0.5278	0.0397	0.9421	0.0176	0.9492	-0.0221	0.9242
CG1972	CG1972	1626880_at	-0.0215	0.8973	0.2734	0.1075	0.4847	0.0119	0.0531	0.9375	-0.3066	0.1171	-0.3597	0.0461	-0.1511	0.7826	0.0866	0.7606	0.2377	0.3271
CG8907	CG8907	1626881_at	0.8092	0.4225	-0.3840	0.5478	-0.0232	0.9448	0.2233	0.9409	0.0451	0.9733	-0.1782	0.8598	-0.2061	0.9365	-1.0875	0.1718	-0.8814	0.2960
lush	lush	1626882_at	0.5076	0.4036	0.2237	0.4974	0.0684	0.7422	-0.2189	0.7880	-0.1193	0.7663	0.0996	0.7900	-0.1361	0.9647	-0.3490	0.7098	-0.2129	0.8407
CG1620	CG1620	1626883_at	-0.0230	0.9293	0.4249	0.3376	0.4978	0.0126	-0.0518	0.9311	-0.0896	0.6514	-0.0378	0.8534	-0.2211	0.8424	0.1759	0.7424	0.3971	0.3921
CG32062	CG32062	1626884_a_at	0.1412	0.7994	0.1865	0.3423	-0.4615	0.2227	-0.0674	0.9220	0.0219	0.9422	0.0894	0.6814	0.4632	0.8222	0.0826	0.9521	-0.3806	0.6883

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11752 /// DyakCG11752	CG11752	1626885_at	-0.0947	0.7755	0.3711	0.3198	0.2779	0.0833	-0.0669	0.9225	-0.6688	0.0068	-0.6019	0.0065	0.0895	0.9611	-0.2335	0.6848	-0.3229	0.5533
CG3663	CG3663	1626886_at	0.4672	0.0375	0.1287	0.4845	0.1123	0.5281	0.0341	0.9672	0.2494	0.2534	0.2154	0.2736	0.1073	0.8940	-0.0298	0.9506	-0.1371	0.6774
---	---	1626887_at	0.1966	0.2529	0.1311	0.4534	0.0988	0.7554	-0.0940	0.8507	0.1301	0.4953	0.2241	0.1664	0.1265	0.9221	0.0752	0.9085	-0.0513	0.9324
CG9426	CG9426	1626888_at	-0.4304	0.2236	0.5067	0.0429	0.2393	0.1623	-0.1363	0.7605	-0.5267	0.0141	-0.3904	0.0309	0.1287	0.9296	0.4290	0.3497	0.3003	0.5457
CG16710	CG16710	1626889_at	-0.0104	0.9551	0.1690	0.3376	0.1966	0.2350	-0.0276	0.9629	-0.0719	0.6924	-0.0443	0.8039	0.0075	0.9928	0.2376	0.2154	0.2302	0.2614
CG11294	CG11294	1626890_at	-0.0018	0.9958	0.1614	0.3579	-0.1067	0.5672	-0.1862	0.7857	-0.1268	0.6968	0.0594	0.8572	0.0929	0.9154	-0.1230	0.7272	-0.2159	0.4944
CG15414	CG15414	1626891_at	2.1880	0.0019	3.5222	0.0025	4.0476	0.0001	0.6654	0.5008	-1.4699	0.0126	-2.1352	0.0015	0.0040	0.9994	0.0564	0.9519	0.0525	0.9491
del	CG9252	1626892_at	-1.3205	0.0463	-2.7053	0.0818	-3.0414	0.0018	0.0111	0.9889	1.6431	0.0001	1.6320	0.0001	0.5090	0.9106	0.4331	0.8431	-0.0759	0.9766
Gasp	Gasp	1626893_at	-1.2881	0.0298	0.2251	0.7047	-1.0315	0.0166	-1.2403	0.0898	-2.7375	0.0004	-1.4972	0.0027	0.1811	0.9445	-1.0780	0.1583	-1.2591	0.1390
CG13165	CG13165	1626894_at	-0.7849	0.0060	-0.2003	0.1961	-0.7343	0.0090	-0.2354	0.6035	-0.9352	0.0024	-0.6998	0.0051	-0.0310	0.9821	-0.2936	0.3496	-0.2626	0.4256
CG40313	CG40313	1626895_at	0.0647	0.6591	-0.0186	0.8573	0.1983	0.1978	0.0789	0.8281	-0.0244	0.8985	-0.1033	0.4318	-0.1014	0.8283	-0.0486	0.8575	0.0528	0.8307
CG16918	CG16918	1626896_at	-1.2167	0.0441	-0.5638	0.0111	-0.9315	0.0002	0.0176	0.9852	-0.3372	0.1089	-0.3548	0.0634	0.0329	0.9862	-0.2830	0.5336	-0.3159	0.4852
HisCl1	Histamine-gated c	1626897_at	0.2899	0.3431	0.0011	0.9936	-0.0961	0.7109	0.0037	0.9961	0.3263	0.1661	0.3226	0.1261	-0.0794	0.9032	-0.0239	0.9505	0.0555	0.8578
CG4751	CG4751	1626898_at	-0.3216	0.3358	-0.2493	0.5928	-0.1783	0.2834	0.0624	0.9351	0.3034	0.1775	0.2410	0.2342	-0.0031	0.9996	0.3524	0.6430	0.3555	0.6388
GATAd	GATAd	1626899_at	-0.0465	0.9025	0.0880	0.6967	-0.6531	0.0333	-0.2646	0.5008	0.5216	0.0205	0.7863	0.0020	0.4615	0.7215	0.5553	0.2959	0.0938	0.9016
Mef2	Complementation	1626900_a_at	0.3961	0.4148	0.9990	0.0613	1.2951	0.0040	-0.0263	0.9803	-0.2127	0.4296	-0.1864	0.4474	0.0446	0.9912	0.6610	0.4138	0.6164	0.4648
CG15890	CG15890	1626901_at	1.4489	0.0144	3.4382	0.0019	3.2361	0.0009	-0.4655	0.7679	-1.9718	0.0102	-1.5063	0.0202	-0.1837	0.9168	0.0940	0.9214	0.2777	0.6787
Rab11	lethal(3)93Bi	1626902_a_at	0.2481	0.1281	0.6150	0.0394	0.2498	0.1367	0.0533	0.9017	0.0757	0.6261	0.0224	0.8937	0.3416	0.5259	0.4333	0.1020	0.0917	0.7585
CG3355	CG3355	1626903_at	-0.5829	0.1163	-1.1640	0.0082	-1.0668	0.0071	0.5569	0.3752	1.2373	0.0048	0.6804	0.0388	0.1600	0.8306	0.1234	0.7356	-0.0366	0.9340
CG2112	CG2112	1626904_at	0.0137	0.9380	0.1580	0.4064	0.1327	0.4931	0.0966	0.8571	0.0313	0.9068	-0.0653	0.7561	-0.0088	0.9929	0.0080	0.9868	0.0168	0.9614
Cyp317a1	Cyp317a1	1626905_at	-1.1998	0.0045	0.1328	0.6049	0.4638	0.0165	-0.2506	0.6354	-1.2418	0.0012	-0.9912	0.0019	-0.3984	0.6955	0.2868	0.5140	0.6851	0.1361
CG17030	CG17030	1626906_at	0.0325	0.8618	0.0250	0.8056	0.1347	0.3983	0.0518	0.9186	0.0093	0.9695	-0.0425	0.8079	-0.0962	0.8465	0.0466	0.8749	0.1428	0.5045
---	---	1626907_at	0.2211	0.1942	0.0889	0.7449	0.1572	0.5456	0.1100	0.8663	-0.0023	0.9945	-0.1123	0.6335	0.0006	0.9999	-0.0820	0.8872	-0.0826	0.8788
CG8066	CG8066	1626908_at	-1.7884	0.0008	-1.5626	0.0115	-2.2649	0.0002	-0.0366	0.9723	-0.4340	0.0992	-0.3974	0.0913	0.3856	0.7436	-0.3285	0.5243	-0.7141	0.1752
CG10931	CG10931	1626909_at	0.2071	0.4515	0.0304	0.7647	-0.0602	0.7868	-0.0159	0.9860	-0.0981	0.6827	-0.0821	0.7131	-0.0847	0.9161	-0.0764	0.8382	0.0083	0.9852
CG15282	CG15282	1626910_at	0.1084	0.7663	-0.0475	0.6899	-0.1294	0.5297	-0.1625	0.8162	-0.0995	0.7625	0.0629	0.8426	-0.0475	0.9589	-0.1308	0.6401	-0.0834	0.7853
CG6495	CG6495	1626911_at	1.2881	0.3512	0.3763	0.7602	1.3451	0.0009	0.6530	0.0770	0.5449	0.0190	-0.1081	0.6183	-0.1335	0.9893	-0.2843	0.9325	-0.1508	0.9609
CG17381	CG17381	1626912_at	0.2108	0.4157	0.0000	1.0000	0.2728	0.1417	0.0648	0.9218	0.0515	0.8411	-0.0133	0.9580	-0.0455	0.9645	-0.0007	0.9997	0.0448	0.9061
CG32175	CG32175	1626913_at	0.2365	0.3352	0.0799	0.5140	0.1646	0.3748	0.0528	0.9297	0.1399	0.4414	0.0871	0.6253	0.0444	0.9514	0.0371	0.9119	-0.0073	0.9842
---	---	1626914_at	-0.1399	0.6245	0.0732	0.7331	-0.0539	0.7950	0.0941	0.9300	-0.0157	0.9743	-0.1098	0.7476	0.0504	0.9589	0.0327	0.9411	-0.0177	0.9659
---	---	1626915_at	0.1426	0.4060	0.0327	0.7561	0.0651	0.7375	-0.0888	0.8914	0.0447	0.8778	0.1335	0.5306	0.0218	0.9898	-0.0638	0.9115	-0.0857	0.8618
CG33998	CG33998	1626916_at	0.0388	0.8675	-0.0038	0.9771	0.2062	0.2612	0.0741	0.8589	-0.0023	0.9915	-0.0764	0.6151	-0.1356	0.7997	0.0155	0.9661	0.1511	0.5149
CG31337	CG31337	1626917_at	-0.0367	0.8836	0.0168	0.9245	-0.3343	0.1633	-0.2242	0.5000	-0.2415	0.1669	-0.0173	0.9364	0.0563	0.9674	-0.1141	0.8061	-0.1704	0.6655
CG13879	CG13879	1626918_at	-0.3666	0.2007	-0.2239	0.4753	-0.3989	0.0833	0.1473	0.8479	0.0962	0.7768	-0.0510	0.8782	0.1936	0.7848	0.0751	0.8591	-0.1185	0.7352
---	---	1626919_at	0.0513	0.7972	0.0266	0.8101	0.1049	0.5266	0.1005	0.8234	0.1498	0.4049	0.0493	0.8026	-0.0543	0.9357	0.0895	0.7124	0.1437	0.5202
---	---	1626920_at	-0.1190	0.4973	-0.0665	0.5588	-0.0169	0.9331	0.0784	0.8844	-0.0479	0.8395	-0.1263	0.4696	0.0884	0.8857	0.2026	0.3733	0.1142	0.6471
---	---	1626921_at	0.0365	0.8579	-0.0294	0.8313	0.0317	0.8517	0.1680	0.6954	0.0856	0.7077	-0.0824	0.6918	0.0442	0.9521	0.0001	1.0000	-0.0441	0.8846
Cpr30F	CG31876	1626922_at	0.2409	0.2391	0.3934	0.1563	0.5534	0.0293	0.0301	0.9814	-0.3882	0.2074	-0.4183	0.1284	0.0975	0.8905	-0.0049	0.9935	-0.1024	0.7392
CG1924	CG1924	1626923_at	-0.0595	0.7754	0.0908	0.4751	-0.0055	0.9783	-0.0361	0.9507	-0.0077	0.9755	0.0283	0.8822	0.1662	0.7307	0.1924	0.3410	0.0262	0.9274
RpS6	ribosomal protein	1626924_a_at	-0.0037	0.9883	0.5365	0.0124	1.2280	0.0180	0.5562	0.5012	-0.4589	0.3007	-1.0151	0.0178	-0.0529	0.9342	-0.0316	0.9221	0.0214	0.9413
Gyc76C	receptor-type guai	1626925_at	-0.7002	0.0255	-0.4222	0.0350	-0.6329	0.0130	-0.1948	0.6802	0.0892	0.7299	0.2840	0.1484	0.0639	0.9589	0.2824	0.4246	0.2185	0.5612
---	---	1626926_at	-0.0185	0.9261	0.0533	0.7960	0.2144	0.4179	-0.0251	0.9744	-0.0361	0.8877	-0.0111	0.9631	0.0161	0.9869	-0.0820	0.7567	-0.0981	0.6892
---	---	1626927_at	-0.4302	0.3459	-0.2954	0.2972	-0.9095	0.0032	-0.2286	0.5515	0.1007	0.6522	0.3292	0.0666	0.2375	0.8236	0.0680	0.9221	-0.1695	0.7417
CG13190	CG13190	1626928_at	0.2474	0.2213	0.0634	0.7750	0.1460	0.3538	-0.2247	0.4504	0.0056	0.9816	0.2303	0.1102	-0.1586	0.8395	-0.1462	0.6854	0.0125	0.9821
CG9357	CG9357	1626929_at	0.2337	0.3384	0.1837	0.3710	0.1544	0.3466	-0.0467	0.9251	-0.0154	0.9438	0.0313	0.8567	-0.1624	0.8202	-0.1364	0.6781	0.0260	0.9503
CG6230	CG6230	1626930_at	0.4778	0.0125	0.4967	0.2054	0.4870	0.0355	-0.1030	0.8865	0.4370	0.0760	0.5400	0.0216	-0.1159	0.9117	0.4637	0.1957	0.5796	0.1443
RpS8	Ribosomal protein	1626931_a_at	0.3290	0.0695	1.5220	0.0048	1.4180	0.0001	0.0118	0.9922	-0.9026	0.0039	-0.9144	0.0022	0.1126	0.8689	0.0202	0.9639	-0.0924	0.7704
CG15194	CG15194	1626932_at	0.0424	0.8075	0.1349	0.3566	0.1207	0.5211	0.0358	0.9518	-0.1153	0.5082	-0.1511	0.3142	0.1202	0.8326	0.1167	0.6497	-0.0035	0.9933
CG2469	CG2469	1626933_s_at	-0.7414	0.0130	0.2460	0.1730	0.6636	0.0465	0.0741	0.8979	-0.5214	0.0138	-0.5955	0.0045	-0.3254	0.7475	0.4102	0.3259	0.7356	0.1190
wb	laminin alpha1,2	1626934_a_at	0.4834	0.5152	1.0202	0.0581	0.6868	0.0206	-0.6885	0.1532	-0.7727	0.0162	-0.0842	0.7975	-0.1790	0.9581	-0.1099	0.9429	0.0691	0.9603
Lcp65Ab2	Lcp65Ab2	1626935_at	0.2058	0.2033	-0.0456	0.6896	-0.0164	0.9350	0.0267	0.9603	0.1810	0.2079	0.1543	0.2324	-0.0298	0.9751	-0.0362	0.9217	-0.0064	0.9866
---	---	1626936_at	-0.0801	0.6676	-0.0604	0.5822	0.2081	0.2091	0.3030	0.3596	0.1221	0.5473	-0.1808	0.2897	-0.0875	0.8700	0.0951	0.6915	0.1826	0.4015
trpgamma	trpgamma	1626937_s_at	0.0301	0.8897	0.2450	0.1635	0.2737	0.2002	-0.0957	0.8817	0.0367	0.9038	0.1324	0.5376	-0.0202	0.9862	0.0912	0.7769	0.1114	0.7045
---	---	1626938_at	0.2125	0.2520	-0.0352	0.7780	0.1720	0.3156	-0.1219	0.7743	-0.0315	0.8971	0.0904	0.6165	-0.1308	0.8097	-0.1454	0.5363	-0.0146	0.9658
CG14313	CG14313	1626939_at	0.0432	0.7849	0.1575	0.2769	-0.0269	0.8846	-0.0043	0.9956	0.1284	0.4642	0.1327	0.3960	0.2534	0.6660	0.1034	0.6977	-0.1501	0.5485
nito	One Twenty Two	1626940_at	0.0794	0.7507	0.0277	0.8799	-0.0766	0.7386	-0.2893	0.2789	0.0537	0.7777	0.3430	0.0230	-0.1845	0.8439	-0.0190	0.9783	0.1655	0.7007
Best4	Bestrophin 4	1626941_at	0.1371	0.6061	-0.0415	0.6932	0.0012	0.9956	0.0629	0.9186	0.0973	0.6474	0.0344	0.8779	0.0374	0.9742	-0.0422	0.9259	-0.0797	0.8243
CG9996	CG9996	1626942_at	0.0089	0.9783	-0.3271	0.0451	-0.4181	0.1917	-0.0977	0.9329	0.4634	0.1747	0.5612	0.0699	-0.0141	0.9943	0.0893	0.8995	0.1034	0.8679
CG14609	CG14609	1626943_at	0.2455	0.3748	-0.2272	0.4374	-0.1769	0.4103	-0.2042	0.7048	-0.0114	0.9764	0.1928	0.4033	-0.1850	0.8513	-0.1670	0.7213	0.0180	0.9778
---	---	1626944_at	0.1635	0.4040	-0.0696	0.6390	-0.0197	0.9368	-0.0908	0.8836	0.1571	0.4765	0.2478	0.1883	-0.1641	0.8379	-0.0631	0.8991	0.1010	0.8007
Snx6	Snx6	1626945_at	-0.2065	0.1473	0.1377	0.3686	0.1101	0.5563	-0.0777	0.8730	-0.4628	0.0140	-0.3850	0.0189	-0.0419	0.9534	0.0111	0.9763	0.0530	0.8461
oc	orthodenticle	1626946_at	0.0253	0.9043	-0.1590	0.4560	-0.0373	0.8522	0.1745	0.7503	0.3061	0.1965	0.1316	0.5794	0.0932	0.9201	0.0670	0.8824	-0.0262	0.9526
pho	pleiohomeotic	1626947_s_at	-0.1325	0.6217	0.2673	0.2256	-0.1792	0.3545	-0.1773	0.7022	-0.2108	0.3286	-0.0335	0.8956	0.2075	0.7946	0.2613	0.4335	0.0538	0.9072
fd96Cb	forkhead domain (	1626948_at	-2.3631	0.0007	-3.2185	0.0011	-2.5275	0.0001	0.3287	0.7039	0.5095	0.1983	0.1808	0.6613	-0.0067	0.9949	-0.0716	0.8185	-0.0649	0.8301
CG11158	CG11158	1626949_at	0.6467	0.0330	0.5674	0.0216	-0.0987	0.6180	-0.4012	0.2502	0.2608	0.2077	0.6620	0.0042	0.1722	0.8202	0.0971	0.8121	-0.0751	0.8540
---	---	1626950_at	-0.0772	0.6699	-0.0300	0.7740	-0.1442	0.4429	0.1384	0.7293	0.0015	0.9951	-0.1368	0.4125	0.1317	0.8097	-0.0785	0.7787	-0.2102	0.3680
---	---	1626951_at	-0.0739	0.7410	-0.0101	0.9315	0.1230	0.5814	0.0361	0.9610	0.0886	0.6956	0.0525	0.8132	-0.0365	0.9717	0.0170	0.9688	0.0535	0.8812
pip	pipe	1626952_at	0.0508	0.8601	0.1151	0.2730	0.1354	0.4470	0.0296	0.9728	0.0028	0.9931	-0.0268	0.9193	-0.0079	0.9943	0.0972	0.7459	0.1051	0.7154
Fmo-1	Flavin-containing	1626953_at	1.7372	0.0072	0.2107	0.4126	0.5197	0.0312	0.0619	0.9562	-0.2867	0.3562	-0.3486	0.2002	-0.2535	0.8878	-1.5151	0.0461	-1.2617	0.0964

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1626954_at	0.1474	0.5601	0.0017	0.9902	0.2353	0.1343	0.0132	0.9857	0.0256	0.9145	0.0124	0.9544	-0.1417	0.8744	-0.0164	0.9796	0.1253	0.7577
CG12014	CG12014	1626955_at	-0.8151	0.0320	-0.0272	0.8673	-0.6437	0.0176	-0.3504	0.3710	-0.8966	0.0025	-0.5461	0.0137	0.1670	0.8236	-0.0328	0.9488	-0.1998	0.5450
---	---	1626956_s_at	0.1815	0.3083	-0.0348	0.7750	-0.1066	0.5524	-0.1441	0.7850	0.1831	0.4148	0.3272	0.0927	-0.0047	0.9952	0.0273	0.9301	0.0320	0.9057
CG15450	CG15450	1626957_at	0.1796	0.2728	0.1640	0.3837	0.3917	0.0768	0.0318	0.9602	0.0067	0.9793	-0.0250	0.9007	-0.0150	0.9923	-0.0029	0.9980	0.0121	0.9839
---	---	1626958_at	0.0551	0.7514	-0.0373	0.7806	0.1030	0.5504	0.1149	0.7204	-0.0116	0.9569	-0.1265	0.3415	-0.0607	0.9412	-0.0582	0.8731	0.0025	0.9954
Obp58b	Odorant-binding p	1626959_at	0.2248	0.3191	0.0926	0.7411	0.0477	0.7961	0.0439	0.9551	0.0120	0.9708	-0.0319	0.9000	0.2189	0.7726	0.0116	0.9854	-0.2072	0.5457
---	---	1626960_at	0.1686	0.2860	0.0121	0.9090	0.1798	0.2523	-0.0581	0.9154	0.0872	0.6514	0.1452	0.3565	-0.0547	0.9108	0.0557	0.8018	0.1103	0.5481
---	---	1626961_s_at	0.2220	0.2214	0.0870	0.5167	0.2014	0.2010	-0.0141	0.9863	0.0260	0.9252	0.0401	0.8620	0.0458	0.9588	-0.0064	0.9911	-0.0522	0.8761
CG32652	CG32652	1626962_x_at	0.1707	0.3505	-0.2227	0.1311	0.0328	0.8720	0.1476	0.7658	0.1500	0.5010	0.0023	0.9928	-0.0769	0.9142	-0.1904	0.4466	-0.1134	0.6800
CG8746	CG8746	1626963_at	-0.1696	0.3294	0.0684	0.6756	-0.0338	0.8821	0.2000	0.6822	0.0447	0.8836	-0.1553	0.4724	0.1091	0.8400	0.0640	0.8285	-0.0451	0.8829
RpL39	Ribosomal protein	1626964_at	0.1558	0.4236	0.4946	0.0323	0.4055	0.0239	0.0520	0.9351	-0.1885	0.3240	-0.2405	0.1523	0.0576	0.9226	0.0357	0.9058	-0.0219	0.9367
CG31743	CG31743	1626965_at	1.3050	0.0091	1.1042	0.0959	1.7409	0.0012	0.0403	0.9761	-0.8075	0.0231	-0.8479	0.0118	-0.6156	0.7436	-0.9917	0.1962	-0.3761	0.6621
---	---	1626966_s_at	-0.8753	0.0387	-0.2771	0.2270	-1.0332	0.0496	-0.4908	0.3452	-0.9735	0.0066	-0.4827	0.0734	0.2537	0.8904	-0.4698	0.5121	-0.7235	0.3079
CalpB	Calpain B	1626967_at	-0.4640	0.1514	0.3771	0.4443	-0.0226	0.9330	0.1402	0.6999	-0.0730	0.7050	-0.2132	0.1521	0.5525	0.7369	0.7384	0.2746	0.1859	0.8334
CG7207	CG7207	1626968_at	0.3162	0.2422	0.3468	0.1821	0.3902	0.0275	-0.0340	0.9518	0.0862	0.6191	0.1203	0.4112	0.0005	0.9999	0.1184	0.8133	0.1179	0.8032
CG11231	CG11231	1626969_at	-0.0070	0.9732	-0.0260	0.8807	-0.0617	0.7362	0.1414	0.8085	0.2362	0.3145	0.0948	0.6983	0.1726	0.7644	0.2720	0.2501	0.0994	0.7205
Ugt35b	UDP-glycosyltrans	1626970_at	-0.2667	0.4588	0.5526	0.3322	0.2724	0.3650	-0.1295	0.9303	-1.5477	0.0044	-1.4181	0.0038	0.0288	0.9862	-0.5738	0.1382	-0.6025	0.1479
---	---	1626971_at	-0.0540	0.8238	-0.0123	0.9189	0.0534	0.8302	-0.0004	0.9996	0.0078	0.9816	0.0082	0.9772	-0.1317	0.8875	0.0116	0.9870	0.1434	0.7156
CG11874	CG11874	1626972_at	-0.0841	0.7878	-0.2569	0.0624	-0.4871	0.0151	-0.1209	0.8140	0.4099	0.0480	0.5308	0.0103	0.0649	0.9535	0.3403	0.2879	0.2753	0.4114
---	---	1626973_at	0.1643	0.3835	0.0653	0.6512	-0.0959	0.7193	0.1502	0.7658	0.0892	0.7220	-0.0610	0.8009	0.1362	0.8276	0.0281	0.9476	-0.1081	0.7154
CG15060	CG15060	1626974_at	0.3232	0.1446	0.3019	0.2568	0.2632	0.1341	-0.0624	0.9496	-0.0159	0.9703	0.0465	0.8870	-0.0258	0.9816	-0.0636	0.8494	-0.0379	0.9109
CG13882	CG13882	1626975_at	0.1400	0.3669	0.0113	0.9158	0.0692	0.7588	0.1356	0.7293	0.3100	0.0793	0.1743	0.2712	-0.0027	0.9982	-0.0214	0.9583	-0.0187	0.9578
---	---	1626976_at	0.5812	0.0292	0.0524	0.6277	0.2278	0.1644	0.1560	0.7631	0.2735	0.2165	0.1175	0.5975	0.1093	0.8049	-0.0122	0.9682	-0.1214	0.5299
Gyc76C	receptor-type guai	1626977_at	0.0019	0.9944	0.1855	0.3761	0.3017	0.0603	-0.1036	0.8118	-0.1858	0.2840	-0.0823	0.6419	-0.0939	0.9309	0.0571	0.9170	0.1510	0.7026
CG31873	CG31873	1626978_at	0.0134	0.9590	-0.0212	0.9287	-0.2056	0.3627	0.0157	0.9869	0.1933	0.4170	0.1776	0.4098	0.0203	0.9916	0.0890	0.8963	0.0686	0.9127
Gr92a	Gustatory recepto	1626979_at	-0.0456	0.8741	-0.0094	0.9509	-0.0286	0.9032	0.0165	0.9857	0.1980	0.3598	0.1816	0.3509	-0.0202	0.9884	0.1421	0.6698	0.1623	0.6171
Pli	pellino	1626980_at	0.0641	0.8897	0.1762	0.3704	-0.0756	0.7872	-0.0770	0.9046	0.4060	0.0562	0.4830	0.0177	0.1381	0.9407	0.5002	0.3740	0.3622	0.5501
CG8979	CG8979	1626981_s_at	-0.4366	0.0824	0.1707	0.3113	-0.0615	0.7535	-0.2029	0.5564	-0.4057	0.0301	-0.2029	0.2017	0.1348	0.7779	0.0204	0.9508	-0.1144	0.6094
---	---	1626982_at	0.1835	0.4609	0.0590	0.5842	0.3002	0.0683	0.0640	0.9005	0.0515	0.7998	-0.0124	0.9532	-0.3013	0.6749	-0.1485	0.6483	0.1527	0.6366
---	---	1626983_at	0.0719	0.7641	-0.1454	0.2546	-0.1005	0.7185	0.0244	0.9688	0.0615	0.7485	0.0371	0.8426	-0.1887	0.8465	-0.2312	0.5897	-0.0425	0.9402
Gld	glucose oxidase	1626984_at	0.6677	0.6326	-0.9779	0.0197	-2.5903	0.0006	-2.1395	0.0010	0.5080	0.0387	2.6476	0.0000	-0.3439	0.9514	-0.9239	0.6053	-0.5800	0.7663
CG7263	CG7263	1626985_a_at	0.2716	0.2728	0.4607	0.1393	0.5914	0.0266	-0.0644	0.9311	0.3523	0.1070	0.4166	0.0399	-0.1536	0.8930	0.4706	0.2507	0.6242	0.1655
CG5516	CG5516	1626986_at	-0.3766	0.1601	-0.2142	0.2055	-0.4632	0.0903	-0.0128	0.9863	0.1360	0.4627	0.1488	0.3628	0.0878	0.9487	0.2195	0.6157	0.1317	0.7868
CG15734	CG15734	1626987_at	0.0418	0.8558	0.0312	0.9119	0.0400	0.8418	0.0231	0.9673	0.1743	0.2398	0.1512	0.2565	0.0662	0.9491	0.1021	0.7927	0.0359	0.9342
CG6805	CG6805	1626988_at	-0.7364	0.0096	-0.7982	0.0096	-0.6465	0.0046	0.0852	0.8471	0.1742	0.2856	0.0890	0.5820	-0.1837	0.7464	0.0396	0.9148	0.2233	0.3627
CG16886	CG16886	1626989_at	0.2183	0.3821	0.0714	0.8158	-0.3484	0.0396	-0.3265	0.5008	0.0658	0.8432	0.3922	0.0878	0.0358	0.9816	0.0264	0.9634	-0.0094	0.9868
Mipp1	Multiple inositol po	1626990_s_at	-0.5363	0.3035	-0.2143	0.6688	-0.8387	0.0033	-0.2746	0.4611	-1.3208	0.0004	-1.0462	0.0005	0.1992	0.9342	-0.9076	0.2269	-1.1068	0.1806
CG11374	CG11374	1626991_at	0.0005	0.9985	-0.3340	0.3996	-0.1243	0.7099	-0.0032	0.9963	0.2646	0.2518	0.2677	0.1927	-0.3347	0.8049	-0.2331	0.7175	0.1016	0.8949
---	---	1626992_at	0.1653	0.4236	0.1206	0.6030	0.1908	0.3206	-0.1408	0.8441	0.0167	0.9677	0.1574	0.5428	0.0508	0.9675	0.1039	0.8045	0.0531	0.9065
Sra-1	specifically Rac1-a	1626993_at	0.2246	0.1182	0.0726	0.7073	-0.5858	0.0196	-0.0803	0.8776	0.4919	0.0138	0.5722	0.0041	0.5765	0.3828	0.3525	0.3225	-0.2240	0.5630
CG33492	CG33492	1626994_at	0.0272	0.8827	-0.0697	0.7538	-0.1044	0.5268	-0.1904	0.6085	0.0279	0.9148	0.2183	0.1896	-0.1433	0.8298	-0.1042	0.7544	0.0392	0.9194
CSN1a	Drosophila COP9	1626995_at	0.1887	0.2951	0.2224	0.4935	0.1770	0.2497	0.3231	0.3428	0.2307	0.2396	-0.0924	0.6469	0.0057	0.9967	-0.1793	0.6312	-0.1850	0.6189
br	broad complex	1626996_s_at	0.2113	0.4439	-0.5337	0.0344	-0.6640	0.0034	0.2820	0.4770	0.7856	0.0037	0.5035	0.0163	0.0683	0.9514	0.1014	0.8125	0.0331	0.9431
CG6425	CG6425	1626997_at	-0.0092	0.9768	-0.4434	0.1104	-0.6747	0.1383	-0.1931	0.7432	0.4398	0.0904	0.6329	0.0142	-0.0426	0.9898	-0.0133	0.9935	0.0293	0.9809
CG8159	CG8159	1626998_at	0.3616	0.0298	0.8019	0.0208	0.6247	0.0052	-0.0346	0.9475	-0.0441	0.8096	-0.0095	0.9593	0.1165	0.8609	0.3038	0.2404	0.1873	0.4998
---	---	1626999_at	0.2854	0.3175	0.1909	0.4138	0.1479	0.4004	-0.1966	0.6718	0.1741	0.4459	0.3707	0.0628	0.0161	0.9914	0.2531	0.4709	0.2370	0.5095
CG6231	CG6231	1627000_s_at	-0.9289	0.1125	-1.2342	0.0235	-1.2639	0.0008	0.2064	0.8794	0.8205	0.0795	0.6141	0.1384	0.1905	0.8825	0.5070	0.2880	0.3165	0.5421
---	---	1627001_at	-0.2201	0.3229	-0.0411	0.8034	0.3085	0.1002	0.2411	0.5363	-0.0623	0.8031	-0.3034	0.0955	-0.1045	0.8368	-0.0231	0.9460	0.0814	0.7345
CG17446	CG17446	1627002_at	-0.0963	0.7600	0.0337	0.8113	-0.0095	0.9687	-0.0459	0.9567	-0.1396	0.5698	-0.0937	0.6950	0.0892	0.9474	0.0155	0.9842	-0.0738	0.8957
MED27	Mediator complex	1627003_at	-0.0884	0.8218	-0.0090	0.9845	0.1880	0.2652	0.0523	0.9186	0.1184	0.4724	0.0661	0.6896	-0.3243	0.8091	0.1133	0.8890	0.4376	0.4457
CG7632	CG7632	1627004_at	0.0602	0.7292	-0.0932	0.6725	-0.3801	0.1320	-0.0526	0.9153	0.1520	0.3334	0.2047	0.1377	0.1566	0.8954	-0.0712	0.9175	-0.2278	0.6331
Reg-5	Drosophila Rhythr	1627005_at	-1.0674	0.0058	-0.5349	0.1606	-0.5621	0.0241	-0.0637	0.9325	-0.9597	0.0019	-0.8960	0.0015	0.0146	0.9943	-0.4554	0.3259	-0.4700	0.3354
Vha13	Vacuolar H[+] ATF	1627006_at	-0.9099	0.0397	-0.5246	0.0330	-0.8764	0.0008	-0.1360	0.6308	-0.3741	0.0149	-0.2381	0.0588	0.1385	0.9030	-0.2673	0.5337	-0.4059	0.3388
CG4306	CG4306	1627007_at	1.8010	0.0016	0.9862	0.1634	1.3753	0.0003	0.1154	0.8612	-0.0507	0.8718	-0.1661	0.4583	-0.1910	0.9272	-0.7845	0.2378	-0.5935	0.3954
CG15034	CG15034	1627008_at	0.2227	0.1493	0.1665	0.4199	0.1915	0.2404	0.2122	0.6580	0.1469	0.5539	-0.0652	0.8012	0.2825	0.6749	0.0820	0.8247	-0.2005	0.4916
stil	STAND STILL	1627009_at	0.4405	0.2272	0.0620	0.7957	0.0465	0.8812	-0.0446	0.9484	0.1566	0.4343	0.2012	0.2457	0.1049	0.9459	-0.0991	0.8831	-0.2040	0.6956
CG13021	CG13021	1627010_s_at	0.0494	0.7426	0.0570	0.5378	-0.0378	0.8121	-0.1682	0.5729	-0.0793	0.6427	0.0889	0.5531	-0.0039	0.9963	-0.0496	0.8386	-0.0458	0.8443
---	---	1627011_at	0.0930	0.6947	-0.0104	0.9573	0.0910	0.5879	-0.0522	0.9297	-0.0150	0.9534	0.0372	0.8534	-0.1424	0.8206	-0.1934	0.4631	-0.0510	0.8874
---	---	1627012_at	0.2276	0.2203	-0.0185	0.8611	-0.0258	0.8954	-0.1991	0.6577	0.0744	0.7750	0.2735	0.1551	0.0966	0.8424	-0.0295	0.9250	-0.1262	0.5488
CG1597	CG1597	1627013_at	0.2131	0.4190	0.1620	0.6625	0.3440	0.0802	-0.2776	0.7131	-0.1276	0.7553	0.1499	0.6766	-0.3842	0.5905	-0.0672	0.8830	0.3170	0.3271
CG6783	CG6783	1627014_a_at	0.7848	0.0590	0.0952	0.7709	0.1840	0.2443	0.2699	0.5735	0.4273	0.0833	0.1574	0.5067	0.0599	0.9781	-0.3953	0.4495	-0.4553	0.3921
sni	Sniffer	1627015_at	-1.0030	0.0035	-0.9044	0.0170	-0.4013	0.0261	0.2127	0.6389	-0.4029	0.0723	-0.6157	0.0081	-0.2070	0.7644	-0.2490	0.3893	-0.0420	0.9174
---	---	1627016_at	0.0274	0.8826	-0.6792	0.1698	-0.6600	0.0537	-0.1248	0.8902	0.6671	0.0346	0.7919	0.0105	-0.1831	0.9168	-0.1316	0.8797	0.0515	0.9520
Dlc90F	dynein light chain;	1627017_at	-0.3238	0.0758	-0.4620	0.0618	-0.5475	0.0389	0.2976	0.3147	0.4723	0.0142	0.1747	0.2619	0.3422	0.6955	0.4753	0.1786	0.1332	0.7514
ATPsyn-gamma	F1ATP synthase	1627018_s_at	-0.2840	0.2848	0.4388	0.1006	0.5472	0.0972	-0.0486	0.9659	-1.2778	0.0020	-1.2292	0.0014	-0.0199	0.9922	-0.3133	0.5131	-0.2934	0.5488
CG33472	CG33472	1627019_s_at	0.0358	0.8558	0.1132	0.4784	0.2566	0.1400	0.1963	0.6075	0.0085	0.9775	-0.1878	0.2782	0.0673	0.9296	0.0977	0.7344	0.0304	0.9299
CG11110	CG11110	1627020_at	-0.5130	0.0818	0.3547	0.4836	1.0631	0.0044	0.2990	0.4532	-0.8305	0.0031	-1.1295	0.0005	-0.4530	0.7726	0.0805	0.9407	0.5335	0.4405
CG8202	CG8202	1627021_at	-0.3804	0.0860	0.3167	0.3843	0.4955	0.0335	-0.1707	0.7576	-0.7595	0.0071	-0.5889	0.0134	-0.3848	0.7230	0.0525	0.9449	0.4373	0.3616
CG9582	CG9582	1627022_at	0.2225	0.1619	0.1342	0.5514	0.1114	0.6008	0.0819	0.9247	0.2213	0.4087	0.1394	0.5924	-0.0047	0.9964	-0.0377	0.9231	-0.0330	0.9234

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG4884	CG4884	1627023_at	0.3135	0.0855	0.0930	0.6424	-0.0224	0.9049	-0.3882	0.1505	0.2244	0.1671	0.6126	0.0021	-0.3417	0.6824	-0.1161	0.7873	0.2256	0.5340
CG9164	CG9164	1627024_s_at	0.0006	0.9973	-0.2380	0.2450	-0.3025	0.2181	0.0232	0.9854	0.1071	0.7530	0.0839	0.7932	-0.0210	0.9852	-0.2374	0.3486	-0.2164	0.4114
---	---	1627025_at	0.1027	0.5241	0.0937	0.5081	0.0020	0.9946	-0.1789	0.7753	-0.1175	0.6974	0.0614	0.8392	-0.0730	0.8732	-0.0316	0.9085	0.0415	0.8608
Edg78E	Ecdysone-depend	1627026_at	0.1939	0.2067	-0.0448	0.6843	0.0510	0.8271	0.1446	0.6852	0.2231	0.1756	0.0785	0.6449	-0.1163	0.8510	-0.1558	0.5618	-0.0395	0.9109
CG15168	CG15168	1627027_at	-0.4785	0.0118	-0.1549	0.5725	-0.1291	0.5262	0.2207	0.4117	0.2170	0.1437	-0.0037	0.9849	0.2637	0.7726	0.5520	0.1512	0.2883	0.4823
CG9030	CG9030	1627028_at	0.1577	0.3467	-0.3532	0.2223	0.0421	0.8438	0.2199	0.5626	0.3988	0.0461	0.1789	0.3122	-0.1629	0.8480	-0.0053	0.9942	0.1576	0.6875
---	---	1627029_at	-0.0362	0.8519	0.0721	0.6248	0.2344	0.2002	-0.1003	0.8315	-0.0486	0.8318	0.0517	0.7981	0.2318	0.6955	0.1922	0.4394	-0.0397	0.9084
---	---	1627030_s_at	0.0489	0.7780	0.3330	0.1780	0.2108	0.6008	-0.3204	0.3828	-0.3415	0.0993	-0.0211	0.9344	0.0491	0.9778	0.1606	0.7456	0.1115	0.8341
---	---	1627031_at	0.0471	0.7972	0.0548	0.6097	-0.0294	0.9082	0.2380	0.5735	0.1921	0.3873	-0.0459	0.8557	0.2417	0.6955	0.2050	0.4139	-0.0368	0.9164
CG8611	CG8611	1627032_a_at	0.6201	0.0333	0.1694	0.3201	0.9319	0.0268	0.1647	0.8479	0.4193	0.1796	0.2546	0.3818	-0.4601	0.7215	0.0571	0.9487	0.5173	0.3552
CG11048 /// DsecCG11048	CG11048	1627033_at	0.2008	0.3886	0.0116	0.9515	0.0478	0.7936	0.0610	0.9229	0.0849	0.6997	0.0239	0.9189	0.0798	0.9142	0.1199	0.6794	0.0401	0.9110
CG9410 /// DyakCG9410	CG9410	1627034_a_at	0.0731	0.7320	0.3415	0.0485	0.4019	0.0430	-0.1869	0.5332	-0.3741	0.0249	-0.1871	0.1801	-0.2875	0.6955	-0.0936	0.8150	0.1938	0.5511
---	---	1627035_at	-0.1062	0.7145	0.0424	0.6830	0.1721	0.3923	0.2071	0.6354	-0.0367	0.9011	-0.2438	0.2032	-0.1912	0.7215	0.0128	0.9734	0.2040	0.3765
Ory	Occludin-Related	1627036_at	0.0604	0.7098	0.0799	0.4894	-0.1501	0.3313	0.0116	0.9857	0.0728	0.6604	0.0612	0.6905	0.0798	0.8609	0.0442	0.8642	-0.0356	0.8884
---	---	1627037_s_at	-0.0345	0.8816	0.0533	0.8308	0.0020	0.9935	0.0211	0.9744	-0.0169	0.9422	-0.0379	0.8359	0.1428	0.8331	-0.0284	0.9505	-0.1713	0.5689
CG13992	CG13992	1627038_at	-0.1983	0.5081	-0.3976	0.0352	-0.1715	0.2517	0.3890	0.5444	0.2723	0.4287	-0.1167	0.7457	0.1829	0.7215	0.0454	0.8841	-0.1375	0.5457
---	---	1627039_at	0.1080	0.5952	-0.0746	0.6847	-0.0222	0.9041	0.0307	0.9689	0.0964	0.6800	0.0657	0.7721	-0.0981	0.8609	-0.0371	0.9157	0.0610	0.8307
CG1344	CG1344	1627040_at	-0.6277	0.0124	-0.1201	0.5729	-0.0945	0.6481	-0.1374	0.7711	-0.4908	0.0220	-0.3534	0.0518	-0.2352	0.7644	0.0235	0.9656	0.2587	0.4523
ine	inebriated	1627041_s_at	-0.1005	0.9231	0.0762	0.9029	-0.7348	0.0071	-0.1262	0.7610	-0.8337	0.0014	-0.7075	0.0016	0.6623	0.8270	-0.6523	0.6332	-1.3146	0.3089
---	---	1627042_at	0.2148	0.2549	-0.0561	0.6011	0.1265	0.5629	0.0485	0.9339	0.0618	0.7640	0.0133	0.9511	-0.1213	0.8846	-0.1597	0.6425	-0.0383	0.9316
CG4674	CG4674	1627043_at	-0.2389	0.5256	-0.0281	0.9315	-0.4954	0.2107	-0.3725	0.3749	0.3462	0.1436	0.7187	0.0054	-0.0448	0.9860	0.3227	0.5972	0.3675	0.5444
Ca-P60A	organellar-type Ca	1627044_s_at	-1.1432	0.0132	-0.8997	0.0638	-0.7574	0.0011	-0.1602	0.6506	-0.3082	0.0736	-0.1480	0.3444	-0.3978	0.7685	-0.1813	0.8053	0.2164	0.7437
CG14877	CG14877	1627045_at	0.2174	0.3856	0.0553	0.7587	-0.0159	0.9592	-0.0870	0.9295	0.1027	0.7665	0.1898	0.4923	-0.0353	0.9514	-0.0537	0.8117	-0.0184	0.9396
Ntf-2	Nuclear transport	1627046_at	0.2122	0.3193	-0.2904	0.2832	-0.5004	0.0048	0.0380	0.9639	0.6048	0.0143	0.5668	0.0117	0.1872	0.8270	0.1189	0.7917	-0.0683	0.8902
CG3987	CG3987	1627047_at	0.2469	0.9244	-0.1687	0.5306	-0.7891	0.0677	-0.4816	0.9218	-0.8513	0.5994	-0.3698	0.8265	-0.0339	0.9964	-1.3367	0.4729	-1.3028	0.4921
CG17293	CG17293	1627048_at	0.3544	0.0797	0.0978	0.7189	0.1354	0.4186	0.1023	0.7608	0.3077	0.0399	0.2054	0.1112	0.0719	0.9467	0.0373	0.9438	-0.0346	0.9394
---	---	1627049_at	0.0062	0.9795	0.0663	0.6739	0.1946	0.2247	-0.0839	0.9036	-0.2367	0.2963	-0.1528	0.4748	-0.0868	0.9092	0.0657	0.8611	0.1526	0.5988
CG9249	CG9249	1627050_at	0.2899	0.3004	0.1210	0.4944	0.2988	0.1797	-0.2257	0.6122	-0.0297	0.9240	0.1960	0.3371	-0.4930	0.5754	-0.2629	0.5214	0.2302	0.5856
Lcp9	Larval cuticle prot	1627051_at	0.3477	0.0490	0.1678	0.3285	0.2382	0.2491	0.0402	0.9448	0.1190	0.4941	0.0788	0.6437	-0.0138	0.9900	-0.0118	0.9784	0.0021	0.9963
CG4617	CG4617	1627052_at	0.1450	0.6676	1.1741	0.0083	1.1367	0.0024	0.0824	0.9247	-0.4694	0.0747	-0.5518	0.0262	0.1022	0.9467	0.4472	0.3259	0.3450	0.4755
slp2	sloppy-paired	1627053_at	-0.3006	0.4074	-0.2856	0.2748	-0.1262	0.6512	0.0535	0.9705	-0.2775	0.4729	-0.3310	0.3268	-0.1345	0.8744	-0.1675	0.6447	-0.0330	0.9434
osp	arc-like wing	1627054_at	-0.9019	0.1180	0.1489	0.4670	-1.1774	0.0013	-0.3248	0.5008	-0.4802	0.0642	-0.1554	0.5343	1.0974	0.4415	0.7065	0.3312	-0.3909	0.6274
CG6709	CG6709	1627055_at	0.1215	0.6097	0.1394	0.4523	0.2735	0.3074	0.3055	0.4068	0.3568	0.0828	0.0513	0.8258	0.1881	0.8110	0.2802	0.3802	0.0922	0.8209
l(2)05714	lethal (2) 05714	1627056_a_at	0.4393	0.0774	0.4576	0.2423	0.2159	0.3236	-0.0738	0.8950	0.3776	0.0475	0.4514	0.0146	0.1859	0.8332	0.3730	0.3000	0.1871	0.6388
---	---	1627057_at	0.1881	0.3609	0.2136	0.5697	0.2129	0.2309	0.0200	0.9860	0.0945	0.7689	0.0745	0.8047	0.1014	0.9056	0.1015	0.7921	0.0002	0.9997
Caf1-105	Caf1-105	1627058_at	0.3704	0.2629	0.0084	0.9882	-0.1878	0.2785	-0.2197	0.6189	0.3011	0.1686	0.5208	0.0162	-0.0776	0.9635	0.0069	0.9942	0.0845	0.8941
und	uninitiated	1627059_at	-0.2830	0.0899	-0.0580	0.7886	0.1602	0.4503	-0.0447	0.9518	-0.3552	0.0836	-0.3105	0.0897	-0.1901	0.7506	-0.1408	0.6085	0.0493	0.8891
Gel	gelsolin	1627060_s_at	-0.4886	0.0264	0.1659	0.6164	0.6216	0.0051	0.0184	0.9761	-0.9583	0.0004	-0.9768	0.0002	-0.4946	0.6389	-0.2505	0.5827	0.2441	0.5984
---	---	1627061_at	0.4894	0.0135	-0.0009	0.9960	0.0384	0.8854	0.1046	0.9036	0.2520	0.3783	0.1474	0.6001	-0.0572	0.9340	-0.1931	0.3713	-0.1359	0.5608
CG14414 /// DsimCG14414	CG14414	1627062_s_at	-0.0633	0.8161	-0.2624	0.1576	-0.2192	0.1523	0.0510	0.9477	0.3889	0.0773	0.3379	0.0847	0.1023	0.8909	0.2656	0.3322	0.1633	0.5862
---	---	1627063_at	0.0644	0.6590	0.1628	0.3822	0.2100	0.2166	0.1162	0.8578	-0.0271	0.9338	-0.1433	0.5258	0.0439	0.9710	0.0172	0.9738	-0.0268	0.9502
aly	always early	1627064_at	0.1402	0.3854	-0.1729	0.1655	0.0235	0.9049	0.0571	0.9043	0.1313	0.4093	0.0742	0.6397	-0.1368	0.8206	-0.1288	0.6344	0.0080	0.9847
---	---	1627065_at	0.1429	0.6457	-0.1668	0.7934	-0.6062	0.2581	-0.9950	0.1787	0.5711	0.1998	1.5660	0.0028	-0.3275	0.9168	0.2524	0.8685	0.5799	0.6189
capt	act up	1627066_at	0.0871	0.6080	-0.0967	0.7265	0.0031	0.9886	0.2990	0.4238	0.3231	0.1166	0.0240	0.9246	0.0092	0.9914	0.0139	0.9647	0.0047	0.9874
CanB2	Calcineurin suppr	1627067_at	-0.2751	0.1132	0.0094	0.9738	-0.0975	0.5456	-0.1163	0.7507	-0.2410	0.1279	-0.1247	0.3988	-0.0511	0.9677	-0.0336	0.9501	0.0175	0.9727
Cyp4aa1	Cyp4aa1	1627068_at	0.3574	0.6003	0.2831	0.3486	0.4512	0.0802	0.1680	0.7966	-0.1155	0.7056	-0.2835	0.2317	-0.0823	0.9742	-0.0388	0.9704	0.0435	0.9624
CG7191	CG7191	1627069_at	0.1973	0.3763	0.0104	0.9244	0.1359	0.4087	-0.1746	0.6457	-0.0320	0.8990	0.1426	0.4066	-0.1301	0.8480	-0.1000	0.7716	0.0302	0.9394
CG15040	CG15040	1627070_x_at	-0.0009	0.9964	-0.0201	0.8744	0.0487	0.7704	0.0543	0.9154	-0.0140	0.9531	-0.0683	0.6832	0.0341	0.9535	0.0674	0.7411	0.0333	0.8891
dpr	defective proboscl	1627071_at	0.0193	0.9405	0.2250	0.1706	0.1368	0.5038	0.0112	0.9860	-0.1046	0.4998	-0.1158	0.3976	-0.0136	0.9923	-0.0830	0.8500	-0.0694	0.8726
---	---	1627072_at	0.0738	0.7085	0.2009	0.2949	0.1203	0.5287	-0.1291	0.8000	-0.1892	0.3677	-0.0601	0.7943	-0.0972	0.8689	-0.0782	0.7883	0.0190	0.9539
CG10126	CG10126	1627073_a_at	-0.2878	0.0929	-0.0944	0.6021	-0.2178	0.3269	0.1863	0.5735	0.5711	0.0055	0.3848	0.0191	0.2964	0.7337	0.9614	0.0335	0.6650	0.1110
CG10185	CG10185	1627074_at	0.2180	0.2169	-0.0129	0.9637	0.1046	0.5315	0.2585	0.2977	0.2604	0.0765	0.0020	0.9915	-0.0630	0.9400	-0.0633	0.8584	-0.0004	0.9992
Gr22e	Gustatory recepto	1627075_at	0.1727	0.2632	-0.1493	0.4174	-0.0800	0.7069	0.0694	0.9011	0.1654	0.3705	0.0960	0.5968	0.0321	0.9822	-0.0719	0.8807	-0.1040	0.7964
CG34109	CG34109	1627076_at	-0.0215	0.9261	-0.0181	0.9146	-0.0509	0.8142	0.0700	0.9098	0.0605	0.8014	-0.0095	0.9690	0.0731	0.9317	0.0590	0.8821	-0.0142	0.9739
CG12917	CG12917	1627077_at	0.3266	0.1216	0.1760	0.1196	0.0399	0.8088	-0.1854	0.6086	-0.1220	0.5292	0.0633	0.7481	0.0212	0.9816	-0.0910	0.6793	-0.1122	0.5983
Oatp58Dc	Organic anion trar	1627078_at	-1.2167	0.3061	-0.1344	0.7014	-0.1138	0.5199	0.2595	0.9311	-1.6657	0.0637	-1.9252	0.0233	0.0441	0.9914	-0.6132	0.5175	-0.6572	0.4867
---	---	1627079_at	0.1055	0.6099	-0.0109	0.9337	0.1830	0.2875	-0.1689	0.6673	-0.0647	0.7744	0.1042	0.5767	-0.1425	0.8564	-0.1067	0.7916	0.0357	0.9369
CG17752	CG17752	1627080_at	-1.0182	0.2009	-0.1830	0.7657	-0.9209	0.0895	-0.0846	0.9857	-0.6568	0.5726	-0.5722	0.5961	0.5080	0.7266	0.1344	0.8816	-0.3736	0.5758
robl37BC	robl37BC	1627081_at	0.0839	0.6601	0.3410	0.0321	0.7402	0.0182	0.1565	0.6229	-0.0418	0.8368	-0.1982	0.1595	-0.1035	0.9095	0.0812	0.8551	0.1847	0.5964
CG15465	CG15465	1627082_at	-0.0788	0.8158	-0.8491	0.1642	-0.6089	0.0414	0.8639	0.1469	1.3317	0.0037	0.4679	0.1452	0.4230	0.7780	0.5133	0.4303	0.0904	0.9197
---	---	1627083_at	0.0274	0.8838	-0.0535	0.8162	0.0217	0.8983	0.0714	0.9029	0.1074	0.6067	0.0359	0.8725	-0.0175	0.9816	-0.0161	0.9563	0.0014	0.9965
SIP2	Syntaxin Interactir	1627084_s_at	0.0749	0.8119	0.1106	0.4605	0.0383	0.8480	-0.0421	0.9563	-0.4412	0.0400	-0.3991	0.0383	0.0226	0.9869	-0.2186	0.4976	-0.2412	0.4569
CG33087	LDLR-like	1627085_a_at	-0.1397	0.4409	-0.1996	0.2910	-0.0930	0.7301	0.0422	0.9598	-0.0242	0.9385	-0.0664	0.7862	-0.0941	0.8794	-0.0798	0.7886	0.0143	0.9686
CG14871	CG14871	1627086_at	0.0739	0.6908	0.0562	0.6063	0.0777	0.7369	0.1256	0.7564	0.0039	0.9883	-0.1216	0.4615	0.0028	0.9978	-0.0711	0.8151	-0.0739	0.7931
CG14805	CG14805	1627087_at	0.3752	0.0521	0.4291	0.0964	0.5978	0.0042	-0.0454	0.9376	0.1535	0.3756	0.1989	0.1886	-0.1450	0.8049	0.1359	0.6025	0.2809	0.2668
fit	female-specific ind	1627088_at	2.7106	0.0039	1.6483	0.3751	4.9050	0.0002	3.0142	0.0093	2.2448	0.0027	-0.7694	0.1259	-0.3401	0.9599	1.1047	0.5880	1.4448	0.4683
---	---	1627089_at	0.1509	0.5232	0.0914	0.4751	0.4118	0.1058	0.0384	0.9705	0.0276	0.9434	-0.0108	0.9736	-0.2396	0.7215	0.0609	0.8819	0.3005	0.3026
Orc6	Origin recognition	1627090_at	-0.0435	0.8774	-0.3338	0.1993	-0.2950	0.1442	-0.1875	0.5917	0.1380	0.4548	0.3255	0.0457	-0.3577	0.6557	-0.0954	0.8215	0.2623	0.4285
CG30287	CG30287	1627091_at	0.1038	0.5811	-0.2172	0.0953	0.1042	0.5678	0.3237	0.5680	0.2658	0.3695	-0.0580	0.8637	-0.0306	0.9816	-0.0103	0.9865	0.0203	0.9636

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31894	CG31894	1627092_at	0.1498	0.4198	0.1103	0.6299	-0.0563	0.7277	-0.0013	0.9988	0.0448	0.8477	0.0462	0.8213	0.1008	0.9092	-0.0237	0.9633	-0.1245	0.7264
CG9284 /// DsimCG9284	CG9284	1627093_at	-0.1176	0.6462	-0.0391	0.7117	0.1674	0.4975	0.1263	0.7760	0.0170	0.9517	-0.1094	0.5489	-0.0201	0.9869	0.1369	0.6512	0.1571	0.5995
---	---	1627094_at	0.1032	0.4981	0.1948	0.2390	0.1343	0.3923	-0.0293	0.9586	-0.0182	0.9304	0.0111	0.9534	0.0401	0.9689	0.0612	0.8730	0.0212	0.9562
---	---	1627095_at	0.0135	0.9522	0.1513	0.4796	-0.0687	0.6827	-0.0097	0.9922	0.0704	0.7795	0.0800	0.7171	0.1678	0.8062	0.2025	0.4875	0.0347	0.9312
Tkr	BTB-protein-III	1627096_s_at	2.7303	0.0026	1.8955	0.0229	2.9030	0.0000	0.1628	0.7444	-0.0180	0.9557	-0.1808	0.3724	-0.0702	0.9848	-0.1258	0.9233	-0.0556	0.9640
CG31858	CG31858	1627097_at	-0.0458	0.8030	-0.0935	0.5432	0.2346	0.1523	0.0398	0.9387	-0.1192	0.4503	-0.1590	0.2435	-0.1296	0.8461	-0.0340	0.9402	0.0956	0.7673
---	---	1627098_s_at	-0.1120	0.4552	0.0209	0.8458	-0.0918	0.7290	0.1438	0.7507	0.1804	0.3693	0.0366	0.8752	0.1305	0.8738	0.2846	0.3695	0.1541	0.6601
CG14391	CG14391	1627099_at	-0.0863	0.6703	0.0587	0.5912	0.1341	0.4195	0.0520	0.9295	-0.0520	0.8057	-0.1040	0.5353	-0.1117	0.8968	0.0910	0.8307	0.2027	0.5499
drm	drumstick	1627100_s_at	-3.7611	0.0006	-0.4889	0.6889	-3.3863	0.0020	-2.2331	0.0414	-3.5507	0.0006	-1.3177	0.0236	0.5585	0.8882	-0.3804	0.8535	-0.9389	0.5525
scyl	scylla	1627101_at	0.0893	0.8769	0.2550	0.5252	-1.1614	0.0519	-0.2940	0.5726	0.0023	0.9951	0.2963	0.2132	1.0803	0.6903	0.1392	0.9390	-0.9411	0.4083
DsimCG9245 /// Pis	CG9245	1627102_at	-0.3106	0.1014	0.0198	0.8573	0.0074	0.9786	-0.1076	0.8671	-0.5551	0.0228	-0.4475	0.0347	-0.0676	0.9226	-0.1823	0.4285	-0.1147	0.6441
---	---	1627103_at	0.1569	0.4840	0.1370	0.2094	-0.1564	0.3561	-0.3063	0.3743	-0.1758	0.3771	0.1305	0.4827	-0.0049	0.9976	-0.2183	0.5764	-0.2134	0.5876
CG8064	CG8064	1627104_at	-0.0023	0.9944	0.0257	0.9568	0.7421	0.0633	0.4897	0.2876	0.3698	0.1700	-0.1199	0.6705	-0.1847	0.9291	0.2777	0.7173	0.4624	0.5063
TrxT	thioredoxin	1627105_at	0.0394	0.8371	-0.1778	0.1828	0.0191	0.9094	0.1603	0.5766	0.1011	0.5173	-0.0591	0.7035	-0.1037	0.9238	-0.1501	0.7166	-0.0463	0.9263
CG10237	CG10237	1627106_a_at	-2.2858	0.0112	-2.1870	0.0077	-2.5481	0.0002	-0.4816	0.5067	-1.1795	0.0084	-0.6980	0.0474	-0.0715	0.9862	-1.0724	0.2386	-1.0009	0.2994
CG12821	CG12821	1627107_at	0.3060	0.1176	0.8561	0.1479	0.5889	0.0325	-0.3270	0.4861	-0.6796	0.0147	-0.3526	0.1149	0.0063	0.9964	0.0495	0.9326	0.0432	0.9330
---	---	1627108_at	-0.0190	0.9192	0.0742	0.7439	-0.0879	0.6019	-0.1350	0.7409	-0.1384	0.4604	-0.0035	0.9875	0.0218	0.9816	0.0448	0.8850	0.0230	0.9387
Hsc70-5	Heat shock protei	1627109_at	0.0417	0.8385	-0.2673	0.0930	-0.2781	0.1469	0.0317	0.9610	0.2748	0.1130	0.2431	0.1166	-0.0194	0.9913	-0.0025	0.9988	0.0169	0.9769
StlP	StlP	1627110_at	0.1722	0.5407	0.8066	0.1764	1.6673	0.0004	0.0008	0.9994	-0.5408	0.0252	-0.5416	0.0157	-0.5354	0.6957	0.1007	0.9093	0.6361	0.2865
NHP2	NHP2	1627111_at	0.7745	0.0868	0.0485	0.9181	0.3552	0.1082	0.3550	0.6817	0.9143	0.0306	0.5593	0.1191	0.0424	0.9848	0.1198	0.8634	0.0774	0.9105
---	---	1627112_s_at	-0.1579	0.7342	-3.0871	0.0052	-2.8666	0.0010	0.4037	0.8432	3.0763	0.0024	2.6726	0.0027	0.2618	0.8646	0.1420	0.8695	-0.1198	0.8870
---	---	1627113_at	0.3919	0.1569	0.1336	0.3506	-0.1369	0.5300	0.0919	0.9353	0.2122	0.5562	0.1203	0.7384	0.1209	0.8553	-0.0871	0.8018	-0.2080	0.4597
CG3793	CG3793	1627114_at	-0.5243	0.0091	-0.0610	0.5248	-0.0600	0.8284	-0.1081	0.8535	-0.2770	0.1884	-0.1689	0.3909	-0.0539	0.9306	0.1160	0.5847	0.1699	0.4042
---	---	1627115_at	0.0472	0.7870	0.2696	0.4206	0.0521	0.8641	-0.0711	0.9376	-0.0840	0.7966	-0.0129	0.9691	0.1236	0.9092	0.0727	0.9005	-0.0509	0.9231
CG18316	CG18316	1627116_at	0.6449	0.1094	0.4983	0.1959	0.0685	0.8642	-0.4771	0.3674	-0.6443	0.0385	-0.1672	0.5763	-0.0840	0.9705	-0.9407	0.1088	-0.8567	0.1612
CG3301	CG3301	1627117_a_at	0.5726	0.1362	-0.0970	0.7812	0.3292	0.1074	0.0728	0.9603	-0.6135	0.1151	-0.6863	0.0538	-0.3382	0.8222	-1.2720	0.0637	-0.9339	0.1591
HPS	HPS	1627118_at	0.7748	0.0082	0.2408	0.3133	0.2901	0.2336	0.1415	0.8033	0.4851	0.0393	0.3436	0.0910	0.1352	0.8882	0.0547	0.9225	-0.0805	0.8660
CG8329	CG8329	1627119_at	0.1219	0.6618	3.8100	0.0291	4.1204	0.0002	0.9513	0.0441	0.9876	0.0037	0.0363	0.9105	0.6567	0.8202	4.4733	0.0134	3.8166	0.0320
robl62A	robl62A	1627120_at	0.1670	0.4463	-0.1043	0.3805	0.0878	0.6458	0.0910	0.8837	0.1684	0.4446	0.0774	0.7346	-0.0276	0.9755	-0.0754	0.7810	-0.0478	0.8709
par-1	Par-1 kinase	1627121_s_at	0.0797	0.6110	0.0808	0.7724	-0.1540	0.4773	-0.1180	0.8073	0.0271	0.9200	0.1451	0.4201	0.1810	0.8395	0.0082	0.9925	-0.1728	0.6668
CG32767	CG32767	1627122_at	-0.0890	0.8681	0.0118	0.9171	-0.6154	0.0261	-0.5167	0.4333	0.3585	0.3286	0.8752	0.0149	0.1765	0.8802	0.4092	0.3494	0.2327	0.6310
CG5828	CG5828	1627123_at	-0.1402	0.4218	-0.0328	0.7644	-0.0374	0.8568	-0.0533	0.9413	-0.0499	0.8529	0.0034	0.9897	-0.0501	0.9495	0.0265	0.9450	0.0766	0.7839
CG7326	CG7326	1627124_at	0.1615	0.7449	0.5672	0.0181	-0.1214	0.6320	-0.2535	0.6818	0.5874	0.0466	0.8409	0.0063	0.2689	0.8521	0.7269	0.2101	0.4580	0.4584
Gr36c	Gustatory recepto	1627125_at	0.2976	0.3108	-0.0335	0.7992	0.3226	0.0722	0.0340	0.9627	0.0357	0.8920	0.0017	0.9945	-0.1164	0.8202	-0.0008	0.9994	0.1156	0.6113
---	---	1627126_at	0.1303	0.3926	-0.0448	0.6559	0.0686	0.7445	0.1207	0.7922	0.1199	0.5495	-0.0008	0.9974	-0.1630	0.8012	-0.1084	0.7287	0.0547	0.8818
CG13609	CG13609	1627127_at	-0.0608	0.7278	0.0387	0.7805	0.0678	0.6826	-0.0253	0.9745	-0.0005	0.9986	0.0249	0.9157	-0.1072	0.8446	0.0795	0.7719	0.1866	0.4132
ppk6	pickpocket 6	1627128_at	0.3737	0.0971	0.1086	0.5177	0.0114	0.9584	-0.0040	0.9956	0.1948	0.3030	0.1988	0.2361	0.0452	0.9743	0.0413	0.9421	-0.0039	0.9952
---	---	1627129_at	-1.5789	0.0141	-1.4042	0.1016	-2.4965	0.0001	-0.3078	0.2870	0.2364	0.1622	0.5442	0.0041	0.8607	0.7116	0.4839	0.6472	-0.3768	0.7368
CG7568	CG7568	1627130_at	0.1759	0.4085	0.0336	0.8525	0.3611	0.0253	0.1318	0.7303	-0.0433	0.8446	-0.1751	0.2546	-0.0303	0.9816	-0.1792	0.5441	-0.1489	0.6260
spt4	spt4	1627131_at	0.5842	0.0397	0.5584	0.1448	0.2947	0.2509	0.0478	0.9672	0.7770	0.0191	0.7293	0.0156	0.3041	0.7822	0.7365	0.1248	0.4324	0.3764
CG15073	CG15073	1627132_at	-0.3934	0.2217	-0.3671	0.0443	-0.6974	0.0196	0.0576	0.8971	0.2282	0.1184	0.1706	0.1931	0.2918	0.7768	0.2108	0.6675	-0.0811	0.8945
Vha100-3	Vha100-3	1627133_at	-0.0539	0.8509	0.1058	0.3709	0.0153	0.9368	-0.1425	0.7556	-0.0751	0.7485	0.0674	0.7550	0.1555	0.8541	0.1233	0.7707	-0.0322	0.9471
ApepP	Aminopeptidase F	1627134_at	0.2456	0.3251	0.5813	0.1402	0.3333	0.0432	-0.0304	0.9730	0.1612	0.4901	0.1916	0.3489	0.1750	0.8326	0.5147	0.1345	0.3397	0.3357
bnl	fibroblast growth f	1627135_at	-1.6082	0.0901	-1.8189	0.0052	-2.2942	0.0012	-0.2894	0.7425	-0.0501	0.9265	0.2392	0.5259	0.3422	0.9310	-0.1067	0.9600	-0.4489	0.7647
Khc-73	kinesin-73	1627136_at	-1.3410	0.0291	-2.0854	0.0111	-2.5025	0.0001	0.0612	0.9745	0.6533	0.1626	0.5921	0.1580	0.2528	0.8461	-0.2159	0.7299	-0.4688	0.3921
---	---	1627137_at	0.1087	0.4178	-0.0318	0.7522	-0.0282	0.8793	0.0067	0.9922	-0.0232	0.9062	-0.0299	0.8567	0.0408	0.9589	0.0349	0.9217	-0.0058	0.9874
CG4271	CG4271	1627138_at	1.5484	0.0008	0.4085	0.3414	1.6630	0.0017	0.4249	0.1927	0.6212	0.0076	0.1964	0.2701	-0.7741	0.6660	-0.2489	0.7799	0.5252	0.4839
CG8734	CG8734	1627139_at	-0.4229	0.1741	-0.0227	0.8378	0.4227	0.0252	-0.1896	0.5771	-0.6016	0.0050	-0.4120	0.0163	-0.5098	0.3828	-0.2776	0.3813	0.2322	0.4869
CG31883	CG31883	1627140_at	0.0488	0.8231	0.0584	0.6873	0.2406	0.2711	0.0720	0.8865	0.0572	0.7820	-0.0147	0.9458	-0.0849	0.9305	0.0812	0.8500	0.1661	0.6233
CG18041	CG18041	1627141_at	-0.0813	0.5692	-0.4497	0.0409	-0.3459	0.0812	-0.0372	0.9518	0.0190	0.9380	0.0562	0.7628	-0.1446	0.8023	-0.1642	0.5100	-0.0196	0.9544
CG2943 /// DyakCG2943	CG2943	1627142_at	-0.2614	0.4050	0.3228	0.1644	0.5455	0.0045	0.2479	0.5376	-0.0613	0.8134	-0.3092	0.0991	0.0588	0.9742	0.6174	0.1634	0.5586	0.2360
CG30034	CG30034	1627143_at	0.1616	0.3377	0.1556	0.2494	0.1499	0.3796	-0.0910	0.8096	0.0971	0.5479	0.1881	0.1601	0.0306	0.9762	0.0749	0.8151	0.0443	0.8951
CG10560	CG10560	1627144_at	-0.9227	0.5532	0.1292	0.5508	-0.1893	0.2002	0.0731	0.9893	-1.5395	0.2138	-1.6125	0.1452	0.2422	0.9031	-0.3179	0.6969	-0.5601	0.4554
CG14457	CG14457	1627145_at	-0.0154	0.9358	0.1335	0.3661	0.2042	0.2406	0.0569	0.9339	0.0106	0.9732	-0.0463	0.8424	-0.0329	0.9717	-0.0694	0.8166	-0.0366	0.9086
CG41456	CG41456	1627146_at	0.2471	0.1208	0.0307	0.8343	-0.0211	0.9122	-0.0033	0.9956	0.0468	0.8442	0.0501	0.8087	-0.0131	0.9901	-0.1602	0.4954	-0.1471	0.5421
CG14644	CG14644	1627147_at	0.0251	0.9366	0.4123	0.0314	0.6628	0.0053	-0.1352	0.7380	-0.5030	0.0123	-0.3678	0.0289	0.0843	0.9340	0.2115	0.5302	0.1272	0.7333
Aats-arg	Arginyl-tRNA synt	1627148_at	0.4733	0.0548	0.9379	0.1632	0.9260	0.0023	0.0582	0.9300	0.2581	0.1803	0.1999	0.2503	0.0763	0.9780	0.7723	0.2259	0.6960	0.3039
---	---	1627149_at	0.0191	0.9245	0.2268	0.4280	0.1515	0.4337	-0.0324	0.9580	-0.1369	0.4131	-0.1046	0.5031	-0.1191	0.8814	0.0265	0.9587	0.1455	0.6604
CG3056	CG3056	1627150_at	-0.0969	0.7942	-0.2626	0.0713	-0.4072	0.0817	-0.0190	0.9857	0.2655	0.2644	0.2845	0.1783	0.1397	0.8995	0.2718	0.5192	0.1321	0.7847
CG4612	CG4612	1627151_at	-0.4599	0.2815	0.0244	0.9440	-0.1139	0.5468	-0.2618	0.5756	0.0158	0.9659	0.2776	0.1946	-0.0742	0.9816	0.4221	0.5539	0.4963	0.4825
CG12918	CG12918	1627152_at	1.9961	0.0004	1.5942	0.0062	2.5585	0.0000	0.7773	0.0260	1.2546	0.0004	0.4773	0.0124	-0.1132	0.9277	0.7467	0.0795	0.8599	0.0718
CG30157	CG30157	1627153_at	0.8035	0.0312	0.6062	0.0260	1.4944	0.0000	0.4943	0.2331	0.5338	0.0369	0.0395	0.8936	-0.2208	0.8305	0.5157	0.2168	0.7365	0.1192
CG10904	CG10904	1627154_at	-0.3466	0.2136	0.3729	0.1037	0.6866	0.0074	0.2175	0.6623	-0.6493	0.0145	-0.8668	0.0024	-0.0418	0.9793	0.1318	0.7692	0.1736	0.6656
CG9106	CG9106	1627155_s_at	0.2629	0.1632	-0.1613	0.5707	0.1116	0.6499	0.1633	0.7482	0.1584	0.4966	-0.0048	0.9862	-0.1694	0.7751	-0.1597	0.5523	0.0098	0.9824
CG33225	CG33225	1627156_at	0.3985	0.0660	0.0767	0.5621	0.0523	0.7840	-0.0506	0.9528	-0.0076	0.9826	0.0430	0.8776	-0.0235	0.9875	-0.0613	0.9064	-0.0378	0.9367
CG1126	CG1126	1627157_at	0.2908	0.1730	0.4152	0.1663	0.2231	0.1762	-0.1261	0.8162	-0.1361	0.5531	-0.0100	0.9702	0.0716	0.9557	0.0502	0.9339	-0.0213	0.9704
---	---	1627158_at	0.0723	0.7007	0.0608	0.7277	0.0828	0.6737	-0.0343	0.9500	-0.0789	0.6434	-0.0445	0.7933	0.0170	0.9914	0.0204	0.9714	0.0034	0.9953
CG13427	CG13427	1627159_at	-0.3815	0.6024	0.0515	0.7723	0.2799	0.2098	0.0058	0.9985	-0.5362	0.4446	-0.5420	0.3873	-0.1028	0.8956	-0.1570	0.6159	-0.0542	0.8926
---	---	1627160_at	0.1257	0.5989	0.0528	0.7916	0.1739	0.2771	-0.1659	0.7531	-0.0757	0.7847	0.0902	0.7090	-0.3316	0.7062	-0.1273	0.7778	0.2043	0.6040

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1627161_at	0.1430	0.4440	-0.0729	0.5285	0.2294	0.2040	0.1921	0.5156	0.2627	0.0930	0.0706	0.6630	-0.2085	0.6749	-0.1175	0.5882	0.0910	0.6877
CG31681	CG31681	1627162_at	2.2280	0.0101	0.8331	0.0070	4.3631	0.0000	3.1963	0.0144	1.3793	0.0432	-1.8170	0.0084	-0.0664	0.8956	0.0394	0.8852	0.1058	0.6028
---	---	1627163_at	0.3272	0.0388	0.3385	0.3485	0.2757	0.1714	0.0448	0.9436	0.0609	0.7811	0.0160	0.9438	0.0801	0.8689	0.0150	0.9621	-0.0651	0.7736
CG4616	CG4616	1627164_at	-0.0091	0.9613	-0.1229	0.4183	-0.5708	0.0691	-0.2408	0.6615	0.1980	0.4722	0.4388	0.0656	0.0583	0.9467	-0.0360	0.9325	-0.0943	0.7578
Nmnat	Nicotinamide mon	1627165_a_at	-0.2973	0.2845	0.6649	0.1107	0.9051	0.0009	-0.0156	0.9910	-0.6035	0.0507	-0.5879	0.0363	-0.1724	0.8326	0.3427	0.3000	0.5152	0.1548
CG17287	CG17287	1627166_at	0.0835	0.7193	-0.0251	0.8639	0.1093	0.5746	0.0969	0.9004	0.0106	0.9785	-0.0863	0.7524	-0.0992	0.9277	-0.1757	0.6512	-0.0765	0.8737
CG40485	CG40485	1627167_a_at	0.0872	0.6989	0.1548	0.4337	0.1490	0.5641	-0.0599	0.9496	-0.2405	0.3728	-0.1806	0.4724	0.1334	0.8541	0.0674	0.8745	-0.0659	0.8674
CG13476	CG13476	1627168_at	-0.0128	0.9655	-0.1411	0.3194	-0.1830	0.3647	0.0939	0.8248	0.4785	0.0106	0.3846	0.0169	-0.0881	0.9246	0.0863	0.8370	0.1744	0.6040
CG3264	CG3264	1627169_at	-0.0211	0.9315	-0.0576	0.7108	0.0436	0.8625	0.0929	0.9098	0.0212	0.9560	-0.0716	0.8061	0.0485	0.9515	-0.0060	0.9915	-0.0545	0.8635
---	---	1627170_at	0.2040	0.1740	-0.0118	0.9096	0.0986	0.6427	-0.1491	0.7497	0.0236	0.9341	0.1728	0.3549	-0.0841	0.9090	-0.1134	0.7000	-0.0293	0.9360
l(3)87Df	lethal (3) 87Df	1627171_at	-0.3766	0.0893	-0.7591	0.0615	-0.4644	0.0560	0.1474	0.8300	0.0330	0.9289	-0.1144	0.6776	-0.1530	0.8760	-0.3863	0.3002	-0.2333	0.5657
CG10254	CG10254	1627172_s_at	0.6759	0.0612	0.2466	0.0737	0.0714	0.7588	0.1067	0.8470	1.0681	0.0008	0.9615	0.0008	0.2743	0.7751	0.5866	0.1489	0.3123	0.4679
CG33224 /// CG4040	CG4040 /// CG33	1627173_at	1.1634	0.3060	0.5911	0.5034	-1.1194	0.0711	-0.3780	0.8164	1.4005	0.0358	1.7785	0.0081	1.3873	0.7215	0.7177	0.6931	-0.6696	0.7137
CG18823	CG18823	1627174_at	0.4009	0.0898	0.2066	0.0931	0.2469	0.1274	0.0431	0.9598	0.0348	0.9112	-0.0083	0.9766	0.1176	0.8023	0.0757	0.7424	-0.0419	0.8743
CG15324	CG15324	1627175_at	0.0328	0.8741	0.0390	0.7668	0.2475	0.1651	-0.0158	0.9808	-0.0652	0.7273	-0.0494	0.7818	-0.0408	0.9487	0.1467	0.4394	0.1875	0.3360
CG31431	CG31431	1627176_at	0.5571	0.1746	1.0920	0.0357	0.8113	0.0582	-0.5932	0.3237	-0.6840	0.0550	-0.0908	0.8185	-0.1102	0.9449	-0.1877	0.7400	-0.0775	0.9073
---	---	1627177_x_at	0.2234	0.2848	0.0158	0.8775	0.0371	0.8209	0.0647	0.9200	0.1311	0.5321	0.0665	0.7570	-0.0561	0.9238	-0.0582	0.8187	-0.0021	0.9953
Acp62F	Accessory gland p	1627178_at	-0.4769	0.0276	-0.1505	0.4143	-1.0340	0.0122	-1.0354	0.2401	-0.3001	0.5986	0.7354	0.1136	-0.0863	0.9342	-0.0189	0.9738	0.0673	0.8844
CG32758	CG32758	1627179_s_at	-0.5963	0.0689	-0.0660	0.8123	0.0044	0.9854	-0.1715	0.7531	-0.6324	0.0153	-0.4609	0.0358	-0.0422	0.9657	0.1010	0.7462	0.1432	0.6166
Cyp4d14	Cyp4d14	1627180_at	0.4447	0.0680	0.1804	0.5011	0.2873	0.1411	0.0212	0.9863	-0.2209	0.4798	-0.2422	0.3819	0.0118	0.9914	-0.3241	0.1925	-0.3359	0.2095
RpS7	Ribosomal protein	1627181_at	0.0894	0.5803	0.1587	0.4744	0.1223	0.6458	-0.0729	0.9254	0.0213	0.9517	0.0943	0.7034	-0.0492	0.9514	0.1054	0.7002	0.1546	0.5457
CG13299	CG13299	1627182_at	-1.6726	0.0375	0.1797	0.7891	-3.4272	0.0036	-2.6427	0.0510	-3.9940	0.0009	-1.3512	0.0564	1.0162	0.7576	-1.8755	0.1637	-2.8917	0.0741
CG14906	open reading fram	1627183_at	-0.0941	0.6135	0.0275	0.8079	-0.1077	0.7813	0.0596	0.9295	0.3770	0.0598	0.3174	0.0735	0.1736	0.8692	0.4873	0.2289	0.3137	0.4697
scaf6	scaf6	1627184_at	-0.1823	0.3954	0.0825	0.8401	0.3660	0.0833	0.0670	0.9380	-0.1052	0.7214	-0.1722	0.4749	-0.2934	0.7632	0.2118	0.6288	0.5052	0.2287
---	---	1627185_at	0.1213	0.5605	0.1939	0.3450	0.3835	0.0934	-0.0445	0.9445	-0.0730	0.7342	-0.0285	0.8972	-0.0263	0.9831	0.0303	0.9492	0.0566	0.8880
Gr23a	Gustatory recepto	1627186_at	-0.0798	0.8443	-0.0423	0.7397	-0.0020	0.9952	-0.0432	0.9777	-0.0384	0.9485	0.0048	0.9921	-0.0788	0.9095	-0.0126	0.9771	0.0662	0.8310
---	---	1627187_at	-0.0952	0.5500	-0.0470	0.8521	-0.0757	0.8123	-0.0185	0.9875	0.0238	0.9550	0.0423	0.9009	0.0408	0.9742	0.0536	0.9111	0.0129	0.9804
phm	phantom	1627188_at	0.4602	0.0240	0.1327	0.4922	0.0118	0.9615	0.1983	0.5836	0.2903	0.1141	0.0920	0.6240	0.1649	0.8202	-0.0076	0.9914	-0.1724	0.5921
CG12289	CG12289	1627189_a_at	0.0849	0.8369	0.0376	0.7090	0.0402	0.8832	-0.1220	0.9023	-0.1487	0.6872	-0.0267	0.9479	0.0591	0.9676	0.0077	0.9931	-0.0513	0.9220
CG13228	CG13228	1627190_at	-0.1090	0.5884	-0.0554	0.6702	0.1308	0.5836	0.1301	0.8632	-0.0681	0.8446	-0.1982	0.4318	-0.0007	0.9998	0.0113	0.9828	0.0120	0.9777
ena	enabled	1627191_a_at	0.3313	0.5759	0.0075	0.9985	-1.0644	0.0511	0.1048	0.9375	1.8448	0.0012	1.7401	0.0010	1.1306	0.7423	1.3403	0.3438	0.2097	0.9170
---	---	1627192_at	0.2756	0.1665	0.1486	0.3938	-0.1370	0.5258	-0.0634	0.9518	-0.0079	0.9859	0.0556	0.8720	0.0546	0.9457	-0.0825	0.7837	-0.1371	0.5990
---	---	1627193_at	-0.0430	0.8528	0.1245	0.3860	0.1790	0.3219	0.1166	0.7820	-0.0539	0.8010	-0.1706	0.2791	0.2228	0.7062	0.2405	0.3149	0.0177	0.9609
l(1)sc	Lethal of Scute	1627194_at	0.0546	0.8083	-0.0412	0.6732	-0.4120	0.0641	-0.3433	0.3793	0.0511	0.8595	0.3944	0.0504	-0.0913	0.8692	-0.0632	0.8294	0.0280	0.9273
CG32642	CG32642	1627195_at	0.0897	0.7178	-0.0174	0.8741	-0.0451	0.8713	-0.1658	0.7929	-0.1170	0.6913	0.0488	0.8718	-0.0785	0.9095	-0.1322	0.6154	-0.0537	0.8699
---	---	1627196_at	0.1096	0.5289	-0.0275	0.8105	0.2399	0.1032	0.1565	0.6331	0.1968	0.2194	0.0404	0.8253	-0.0761	0.8756	-0.0280	0.9263	0.0480	0.8436
Rop	Ras opposite	1627197_at	-0.5844	0.0230	-0.3868	0.1931	-0.3576	0.1488	0.1966	0.6041	0.2535	0.1817	0.0570	0.7894	0.2376	0.8284	0.5363	0.2265	0.2987	0.5379
CG33928	CG33928	1627198_at	-0.3707	0.1746	-0.8349	0.0779	-0.9578	0.0043	-0.0071	0.9956	0.3833	0.2174	0.3904	0.1601	-0.1308	0.8382	-0.1854	0.4943	-0.0546	0.8788
---	---	1627199_at	0.0298	0.8620	0.0817	0.4771	0.2512	0.3000	0.1186	0.7947	-0.0458	0.8489	-0.1643	0.3329	0.0563	0.9460	-0.0223	0.9573	-0.0786	0.7962
---	---	1627200_at	0.0374	0.8324	0.1216	0.4973	-0.0109	0.9613	0.0449	0.9380	-0.0513	0.8065	-0.0962	0.5683	0.0559	0.9516	0.0434	0.9198	-0.0125	0.9777
---	---	1627201_at	0.2721	0.3097	-0.0135	0.9610	-0.0741	0.8139	-0.1322	0.8908	-0.0361	0.9378	0.0961	0.7899	0.0740	0.9506	-0.1533	0.7020	-0.2273	0.5444
AP-2sigma	AP-2sigma	1627202_at	-0.3008	0.2843	-0.0214	0.8760	-0.0583	0.7946	-0.1517	0.5766	-0.2196	0.1141	-0.0678	0.6343	-0.1497	0.8680	0.0268	0.9639	0.1765	0.6460
CG17150 /// DmirCG17150	CG17150	1627203_at	0.3384	0.1794	0.0000	1.0000	0.4122	0.0205	0.1184	0.8189	0.1885	0.3610	0.0701	0.7498	-0.1223	0.8837	-0.2604	0.4091	-0.1381	0.6970
CG4749	CG4749	1627204_at	-0.2511	0.2477	-0.5675	0.0447	-0.3008	0.0792	0.2372	0.5008	0.4533	0.0235	0.2160	0.1956	-0.0502	0.9555	0.0187	0.9652	0.0689	0.8364
---	---	1627205_at	0.2619	0.1006	0.2662	0.0641	0.2954	0.2835	0.0727	0.8676	-0.0025	0.9907	-0.0753	0.6328	-0.0400	0.9816	0.0118	0.9886	0.0518	0.9234
RpL35	Ribosomal protein	1627206_at	0.6744	0.1172	1.3577	0.0179	1.0407	0.0029	-0.0636	0.9659	-0.7573	0.0587	-0.6936	0.0533	0.1835	0.8609	0.3386	0.4245	0.1551	0.7514
CG14352	CG14352	1627207_at	-0.7448	0.0217	0.4075	0.2884	0.7315	0.0393	-0.1160	0.9196	-0.6820	0.0569	-0.5660	0.0737	-0.1378	0.9011	0.5415	0.1694	0.6793	0.1271
CG18418	CG18418	1627208_at	0.2527	0.1751	-0.5614	0.0488	-0.2890	0.1638	0.1585	0.6257	0.4988	0.0081	0.3403	0.0261	-0.0765	0.9032	-0.1911	0.3999	-0.1146	0.6419
mRNA-capping-enzyme	mRNA-capping-er	1627209_at	-0.1045	0.5521	0.5921	0.0762	0.5777	0.0057	0.0296	0.9672	0.0053	0.9851	-0.0243	0.9157	0.0947	0.9075	0.6304	0.0512	0.5358	0.1003
MstProx	MstProx	1627210_at	-0.0117	0.9551	0.0197	0.8466	0.0650	0.6973	0.0570	0.9108	-0.0117	0.9615	-0.0686	0.6833	-0.0254	0.9831	0.0536	0.8993	0.0790	0.8218
CG7202	CG7202	1627211_at	-0.0990	0.4961	-0.0038	0.9771	0.1842	0.1962	0.2673	0.3851	-0.0244	0.9204	-0.2917	0.0642	0.1654	0.7230	0.1457	0.4791	-0.0197	0.9440
ref(2)P	refractaire	1627212_at	-0.8895	0.0039	-0.9995	0.0465	-0.4121	0.1027	0.2139	0.5917	-0.2776	0.1672	-0.4915	0.0146	-0.4068	0.7220	-0.4169	0.3832	-0.0101	0.9902
---	---	1627213_at	0.1432	0.3560	0.0539	0.6913	0.0526	0.8067	0.0753	0.8768	0.1825	0.2770	0.1072	0.5056	0.1374	0.8427	0.1584	0.6019	0.0210	0.9596
CG7650	CG7650	1627214_s_at	0.1443	0.4897	0.0473	0.7020	0.6846	0.0480	0.0647	0.9300	-0.2305	0.2900	-0.2952	0.1247	-0.4689	0.6496	-0.1684	0.7269	0.3005	0.4865
CG14310	CG14310	1627215_at	0.1658	0.3751	0.2273	0.0872	-0.0782	0.6463	0.0373	0.9467	0.0930	0.5885	0.0557	0.7441	0.1882	0.7644	0.1368	0.6382	-0.0515	0.8889
Zyx102EF	Zyx102EF	1627216_s_at	0.2666	0.2782	-0.1908	0.4607	-0.4069	0.0585	-0.3095	0.4128	0.2514	0.2298	0.5609	0.0095	-0.1372	0.9070	-0.1634	0.7399	-0.0262	0.9666
Glut3	Glucose transport	1627217_at	0.1126	0.5329	0.0108	0.9182	0.2630	0.0805	-0.0331	0.9672	-0.0390	0.8910	-0.0060	0.9822	-0.0590	0.9208	-0.0497	0.8563	0.0093	0.9769
CG7646	CG7646	1627218_a_at	-1.2834	0.0075	-0.4725	0.5450	-1.4635	0.0002	-0.5378	0.4568	-0.7118	0.0720	-0.1741	0.6691	0.5460	0.7464	0.0855	0.9406	-0.4605	0.5411
nAcRbeta-64B	AChR protein of D	1627219_at	-0.0112	0.9802	-0.3182	0.2542	-0.2637	0.2907	-0.0002	0.9999	0.2926	0.3701	0.2928	0.3146	-0.1854	0.8521	0.0050	0.9957	0.1904	0.6719
ND23	NADH:ubiquinone	1627220_at	-0.2341	0.3968	-0.1353	0.6000	-0.1781	0.4645	-0.1287	0.8498	-0.6202	0.0205	-0.4915	0.0335	-0.0111	0.9952	-0.4765	0.2880	-0.4654	0.3253
cdm	cadmus	1627221_at	-0.2921	0.1620	0.0153	0.8819	-0.0386	0.8616	0.0412	0.9441	-0.3352	0.0508	-0.3764	0.0204	0.1432	0.8270	0.1356	0.6497	-0.0076	0.9860
---	---	1627222_at	0.1808	0.3439	0.3231	0.2244	0.0872	0.7405	-0.1488	0.7596	-0.1574	0.4714	-0.0086	0.9736	0.1310	0.8833	0.0585	0.9110	-0.0725	0.8744
---	---	1627223_at	-0.0180	0.9192	-0.0258	0.8061	0.1605	0.4804	-0.0096	0.9893	-0.0001	0.9996	0.0095	0.9629	-0.1719	0.8062	0.0188	0.9689	0.1906	0.5361
CG11692	CG11692	1627224_at	-0.0773	0.6139	0.0641	0.6019	-0.0990	0.6173	-0.0538	0.9491	-0.0503	0.8718	0.0035	0.9906	0.0387	0.9717	0.1024	0.7573	0.0637	0.8618
Tsp26A	Tetraspanin 26A	1627225_at	0.2837	0.2453	0.0929	0.5226	0.2093	0.2853	0.0882	0.9064	-0.0186	0.9589	-0.1068	0.6709	0.0421	0.9585	-0.0241	0.9476	-0.0662	0.8120
---	---	1627226_at	0.1636	0.4360	0.1603	0.3956	0.1599	0.2599	-0.1180	0.8121	-0.1118	0.6034	0.0062	0.9799	0.0991	0.8657	0.0129	0.9741	-0.0862	0.7495
Pglym78	Phosphoglycerom	1627227_at	-0.0117	0.9555	-0.4418	0.1012	-0.3108	0.0578	-0.0664	0.8942	-0.0223	0.9231	0.0441	0.8135	-0.2273	0.7500	-0.4558	0.1345	-0.2285	0.4683
CG15309	CG15309	1627228_at	0.2300	0.4892	0.3973	0.1328	0.2550	0.5843	-0.2456	0.6493	-0.7941	0.0088	-0.5485	0.0267	-0.1139	0.9672	-0.6187	0.3912	-0.5048	0.5058
CG34364	CG12927	1627229_at	0.3369	0.1140	0.1098	0.4622	0.3344	0.0685	0.2504	0.5770	0.2469	0.2871	-0.0035	0.9905	0.0050	0.9964	-0.0381	0.9325	-0.0431	0.9109

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1627230_at	-0.0371	0.9046	0.0379	0.7167	-0.0053	0.9806	-0.0037	0.9956	-0.0108	0.9699	-0.0071	0.9755	0.1418	0.8882	0.0754	0.8920	-0.0663	0.8984
CG40342	CG40342	1627231_at	0.0084	0.9682	-0.0162	0.8964	0.0286	0.8850	0.0038	0.9956	-0.0110	0.9640	-0.0148	0.9404	0.0022	0.9984	0.0031	0.9950	0.0009	0.9985
---	---	1627232_at	0.2418	0.3807	0.1083	0.4914	0.0035	0.9918	-0.1866	0.6202	-0.1860	0.3278	0.0006	0.9981	0.0074	0.9964	-0.2709	0.4833	-0.2783	0.4771
CG14047	CG14047	1627233_at	0.2896	0.1057	0.1235	0.6564	-0.0206	0.9211	-0.0890	0.8528	0.1911	0.2738	0.2801	0.0734	0.1291	0.8609	0.0874	0.8220	-0.0417	0.9194
---	---	1627234_at	0.0169	0.9375	-0.0008	0.9956	0.0701	0.6816	-0.0112	0.9900	0.1263	0.5609	0.1375	0.4730	0.0436	0.9488	0.0981	0.6605	0.0544	0.8310
CG17207	CG17207	1627235_at	0.1505	0.4193	0.1000	0.5128	0.0299	0.8785	-0.1645	0.8119	-0.0215	0.9578	0.1431	0.6039	-0.0798	0.9238	-0.1371	0.6512	-0.0573	0.8800
---	---	1627236_s_at	1.1094	0.3092	1.7205	0.0958	0.5053	0.6023	0.0077	0.9931	0.6734	0.0031	0.6657	0.0019	1.2296	0.7707	1.9995	0.2501	0.7699	0.7027
CG15564	CG15564	1627237_at	0.2200	0.4422	0.2746	0.2584	0.1142	0.5698	0.2265	0.7235	0.1010	0.7713	-0.1255	0.6776	0.3119	0.7215	0.0654	0.9076	-0.2466	0.5230
CG6520	CG6520	1627238_at	0.1200	0.7022	0.0158	0.9002	0.0296	0.8703	0.0104	0.9942	-0.1482	0.6534	-0.1585	0.5885	-0.0844	0.8637	-0.1745	0.3672	-0.0901	0.6766
CG17034	CG17034	1627239_s_at	-0.8813	0.0041	-1.1546	0.0172	-0.8193	0.0109	-0.2476	0.5531	0.1197	0.6157	0.3672	0.0607	-0.5365	0.5228	-0.1750	0.7002	0.3615	0.3807
CG13039	CG13039	1627240_at	0.0862	0.5597	0.0286	0.8403	0.3918	0.0296	0.0107	0.9873	-0.0476	0.8112	-0.0583	0.7369	-0.2405	0.6749	-0.0821	0.7741	0.1584	0.5172
CG32696	CG32696	1627241_at	0.2939	0.2271	0.1953	0.1605	0.3712	0.1443	-0.0378	0.9413	0.0120	0.9561	0.0498	0.7570	-0.1955	0.7644	-0.0416	0.9225	0.1539	0.6017
l(2)efl	lethal (2) essentia	1627242_at	-1.4252	0.0590	-1.4526	0.0117	-2.2828	0.0001	-0.1259	0.7995	0.3821	0.0602	0.5080	0.0121	0.5448	0.8049	0.4811	0.6264	-0.0637	0.9622
---	---	1627243_at	0.1484	0.3832	-0.0129	0.9029	0.1864	0.2896	-0.0769	0.8671	0.0261	0.9077	0.1030	0.5098	-0.1255	0.8270	-0.0841	0.7707	0.0414	0.8993
CG18870	CG18870	1627244_at	0.0797	0.6702	-0.5378	0.0311	-0.6487	0.0258	-0.2240	0.5859	0.5173	0.0218	0.7413	0.0027	-0.0503	0.9698	-0.1516	0.6937	-0.1012	0.8111
CG5746	CG5746	1627245_s_at	0.4759	0.1647	0.3287	0.2388	0.0425	0.8832	-0.2640	0.6247	0.1268	0.6774	0.3907	0.1014	0.0326	0.9895	0.0764	0.9263	0.0439	0.9523
Pcaf	Pcaf	1627246_at	0.1709	0.2801	-0.0918	0.7456	0.1933	0.3018	0.1449	0.6513	0.3532	0.0301	0.2083	0.1307	-0.0535	0.9657	0.0946	0.8319	0.1480	0.6892
Oseg1	Oseg1	1627247_at	-0.5810	0.0452	-0.8336	0.1058	-1.1211	0.0008	-0.1836	0.7432	0.0323	0.9252	0.2158	0.3390	-0.3073	0.8202	-0.2144	0.7424	0.0929	0.9032
mura	murashka	1627248_at	0.3632	0.2014	0.4069	0.3919	0.3188	0.0804	-0.1749	0.7539	-0.2652	0.2733	-0.0903	0.7254	-0.0482	0.9824	-0.2356	0.6744	-0.1874	0.7484
l(2)35Bg	transcription unit E	1627249_a_at	0.2035	0.2915	0.1400	0.2123	0.9294	0.0107	0.3780	0.1290	0.4277	0.0115	0.0497	0.7617	-0.1758	0.8400	0.4167	0.2288	0.5925	0.1300
Msp-300	nesprin	1627250_at	-1.1386	0.1500	-1.0030	0.0896	-2.8045	0.0000	-1.1143	0.0741	0.2578	0.4957	1.3721	0.0019	0.2822	0.9092	0.0128	0.9943	-0.2694	0.8006
CG12565	CG12565	1627251_s_at	0.0340	0.8689	-0.0203	0.8997	0.1335	0.3746	0.0556	0.9138	0.0555	0.7768	-0.0001	0.9995	0.0158	0.9884	0.0030	0.9955	-0.0129	0.9717
---	---	1627252_at	-0.0175	0.9161	0.1044	0.5327	0.2392	0.1223	0.0352	0.9586	-0.0370	0.8765	-0.0722	0.7019	-0.0252	0.9717	0.0394	0.8792	0.0646	0.7568
cutlet	gilead	1627253_at	-0.0131	0.9844	-0.6561	0.2048	-0.6629	0.0452	-0.0849	0.9314	0.7506	0.0186	0.8355	0.0071	0.0432	0.9914	0.5017	0.5878	0.4585	0.6261
CG7427	CG7427	1627254_at	-0.1023	0.5339	-0.0205	0.9564	-0.0005	0.9980	0.0666	0.8853	0.1188	0.4681	0.0522	0.7612	-0.0178	0.9913	0.1253	0.7563	0.1431	0.7067
CG12625	CG12625	1627255_at	0.1393	0.5079	0.0216	0.8374	0.1216	0.5389	0.0133	0.9893	-0.0134	0.9698	-0.0267	0.9235	0.0003	0.9999	0.0005	0.9998	0.0002	0.9997
ATPsyn-d /// CG14642	CG14642 /// ATPa	1627256_s_at	-0.1669	0.4535	0.9007	0.0725	1.0079	0.0012	-0.0720	0.9287	-1.3206	0.0007	-1.2486	0.0005	-0.2137	0.8222	-0.2615	0.5246	-0.0478	0.9316
CG4433	CG4433	1627257_at	-0.8279	0.0509	0.8535	0.0335	0.8789	0.0047	0.0784	0.9346	-0.8020	0.0119	-0.8805	0.0047	0.1438	0.8909	0.8897	0.0461	0.7459	0.0947
---	---	1627258_at	-0.1465	0.6003	-0.0592	0.6058	-0.2649	0.1946	-0.3781	0.2876	-0.1340	0.5522	0.2441	0.1911	-0.0465	0.9589	-0.1324	0.6381	-0.0859	0.7788
ovo	shaven baby	1627259_a_at	-1.7523	0.0075	-2.5500	0.0152	-2.8590	0.0000	-0.3209	0.4979	0.4846	0.0577	0.8055	0.0041	-0.1653	0.9409	-0.3619	0.6265	-0.1966	0.8188
CG11693	CG11693	1627260_at	0.1718	0.4865	0.2589	0.0764	-0.0846	0.7150	-0.1289	0.7803	-0.1515	0.4469	-0.0227	0.9235	0.1476	0.8235	-0.0068	0.9916	-0.1544	0.6024
Cnx99A	calnexin	1627261_s_at	0.1592	0.5934	0.3310	0.0949	0.3328	0.0759	0.1469	0.6679	0.6567	0.0026	0.5098	0.0047	-0.1041	0.9391	0.6966	0.1047	0.8006	0.0908
CG3599	CG3599	1627262_at	0.0973	0.8186	0.1365	0.4522	-0.5306	0.0312	-0.4817	0.2359	-0.3326	0.1652	0.1490	0.5250	0.0703	0.9742	-0.2903	0.6184	-0.3606	0.5274
CG31601	CG31601	1627263_at	0.0523	0.7717	0.0659	0.6806	0.1082	0.5877	0.0460	0.9300	0.1357	0.3873	0.0898	0.5509	-0.0642	0.9549	0.2098	0.5455	0.2740	0.4175
---	---	1627264_at	-0.0069	0.9725	-0.1002	0.4628	0.1116	0.5991	0.0800	0.8628	0.0985	0.5818	0.0185	0.9267	0.0066	0.9963	0.0471	0.9213	0.0405	0.9231
CG9139	CG9139	1627265_at	-0.1153	0.5407	-0.4263	0.1174	-0.3444	0.1336	0.2420	0.4907	0.5631	0.0092	0.3210	0.0588	0.1522	0.8472	0.2692	0.4094	0.1169	0.7578
---	---	1627266_at	0.0880	0.6528	-0.0918	0.4567	0.0342	0.8762	-0.0452	0.9639	0.1564	0.5842	0.2016	0.4126	-0.1139	0.8608	-0.0043	0.9941	0.1097	0.7083
MESR3	Misexpression sup	1627267_at	-0.9546	0.0011	-1.1825	0.0564	-1.8152	0.0000	-0.0145	0.9842	0.2604	0.1221	0.2749	0.0716	0.5036	0.6557	0.0559	0.9402	-0.4477	0.3406
CG13386	CG13386	1627268_at	0.1487	0.4701	-0.0365	0.8050	0.0793	0.7352	0.1582	0.6954	0.2947	0.1130	0.1366	0.4359	0.1955	0.8016	0.2642	0.4121	0.0687	0.8759
---	---	1627269_at	-0.0746	0.6396	-0.0294	0.9189	-0.1607	0.4214	0.0584	0.9371	0.0594	0.8263	0.0010	0.9973	0.0596	0.9330	-0.0621	0.8386	-0.1217	0.6189
CG15893	CG15893	1627270_at	-1.2808	0.0013	-0.8819	0.0577	-1.4203	0.0002	-0.3847	0.1391	-0.3588	0.0299	0.0259	0.8907	-0.0055	0.9982	0.0339	0.9664	0.0394	0.9551
CG13422	CG13422	1627271_at	0.2999	0.3522	1.1496	0.1011	0.9648	0.1058	0.1023	0.9603	-0.4294	0.4474	-0.5317	0.2791	0.3466	0.8553	0.3776	0.6596	0.0310	0.9811
CG30380	CG30380	1627272_at	-2.2120	0.0070	-1.9450	0.0129	-2.5048	0.0000	-0.1221	0.9086	-0.3104	0.3705	-0.1883	0.5780	0.1371	0.9445	-0.3603	0.5636	-0.4974	0.4114
CG12035	CG12035	1627273_at	0.1096	0.6435	0.1182	0.5348	0.1106	0.6900	-0.0833	0.9225	-0.2322	0.3787	-0.1489	0.5593	-0.0364	0.9787	-0.0377	0.9409	-0.0014	0.9982
CG31044	CG31044	1627274_at	0.3377	0.0654	0.1432	0.5842	0.3558	0.0768	-0.0566	0.9413	-0.0039	0.9902	0.0527	0.8340	-0.2292	0.7464	-0.1031	0.7787	0.1260	0.7075
CG9902	CG9902	1627275_at	0.2929	0.1587	-0.0019	0.9974	-0.2707	0.3043	-0.3734	0.2753	0.3640	0.0742	0.7374	0.0023	0.1612	0.9095	0.1068	0.8836	-0.0545	0.9387
CG40450	CG40450	1627276_a_at	0.1586	0.3776	-0.0097	0.9269	0.1630	0.3592	0.1041	0.8473	0.0499	0.8461	-0.0543	0.8075	0.0518	0.9467	0.0158	0.9670	-0.0360	0.9109
CG2930	CG2930	1627277_s_at	0.2629	0.2147	1.4309	0.0230	1.3580	0.0012	-0.0077	0.9952	-0.8694	0.0058	-0.8617	0.0035	-0.1237	0.9112	0.1389	0.7730	0.2626	0.5257
CG14619 /// DyakCG1461	CG14619	1627278_s_at	-0.4543	0.4173	-0.0346	0.9476	-1.1003	0.0146	-0.2068	0.8096	0.4626	0.1729	0.6694	0.0352	0.9091	0.6749	0.7571	0.3922	-0.1520	0.9025
nwk	nervous wreck	1627279_at	0.1347	0.3413	0.0648	0.5774	-0.0290	0.8923	0.0979	0.8336	0.0675	0.7475	-0.0304	0.8843	-0.0533	0.9305	-0.0774	0.7368	-0.0241	0.9304
pnut	septin	1627280_s_at	-0.4800	0.1470	0.0555	0.9645	-0.5490	0.0241	0.0292	0.9695	-0.0243	0.9298	-0.0535	0.8091	0.5310	0.8331	0.4471	0.7072	-0.0839	0.9553
CG3117	CG3117	1627281_at	0.1335	0.4074	-0.0968	0.7228	0.0947	0.6507	-0.0066	0.9956	0.1080	0.7135	0.1147	0.6634	-0.1113	0.8967	0.0063	0.9935	0.1177	0.7492
CG15548	CG15548	1627282_at	0.1541	0.5880	0.0672	0.5431	0.3534	0.0533	-0.1533	0.8244	0.0795	0.8099	0.2329	0.3424	-0.3400	0.5461	-0.1073	0.7200	0.2327	0.3844
---	---	1627283_at	0.0368	0.8347	-0.1572	0.6273	0.0521	0.7659	0.1521	0.8190	0.2264	0.3967	0.0744	0.7991	-0.0595	0.9514	0.0598	0.8857	0.1193	0.7135
PH4alphaPV	prolyl-4-hydroxyla	1627284_at	-0.1625	0.4761	-0.1407	0.2376	-0.3333	0.0401	0.0544	0.9476	0.1914	0.4275	0.1371	0.5495	-0.0753	0.9449	-0.1056	0.8031	-0.0303	0.9479
CG6118	CG6118	1627285_at	0.2969	0.1591	0.3298	0.0466	0.2794	0.0990	-0.0137	0.9852	-0.0548	0.7851	-0.0411	0.8276	0.0165	0.9884	-0.0414	0.9139	-0.0579	0.8577
CG11369	CG11369	1627286_at	0.0498	0.8223	0.1569	0.4047	0.3219	0.1197	0.1106	0.8891	-0.2148	0.4360	-0.3254	0.1702	-0.0546	0.9523	-0.1095	0.7311	-0.0549	0.8842
CG5966	CG5966	1627287_at	1.9029	0.1404	0.3480	0.8260	2.0073	0.0009	0.7948	0.6517	-0.1809	0.8725	-0.9757	0.1989	-0.6247	0.9016	-1.3992	0.4514	-0.7745	0.7118
Cpn	calcium-binding pr	1627288_a_at	0.3240	0.1888	-0.0424	0.7087	0.2964	0.1170	0.1782	0.7613	0.2653	0.2979	0.0871	0.7515	-0.1779	0.7707	-0.1460	0.6010	0.0319	0.9310
---	---	1627289_at	-0.1751	0.3414	0.0278	0.8731	-0.1584	0.4592	-0.1507	0.7874	-0.0600	0.8395	0.0906	0.7099	0.0457	0.9626	0.0698	0.8500	0.0241	0.9492
CG13042	CG13042	1627290_at	0.1928	0.5811	0.2198	0.0718	0.2637	0.1528	0.0043	0.9961	-0.1212	0.7060	-0.1255	0.6635	0.0561	0.9672	0.0136	0.9842	-0.0425	0.9330
inx6	pas related protein	1627291_at	0.0417	0.8236	0.1038	0.3575	0.1856	0.3184	-0.1390	0.7777	-0.3214	0.1164	-0.1824	0.3316	-0.0102	0.9915	-0.0343	0.9216	-0.0241	0.9386
CG2898	CG2898	1627292_at	0.3226	0.0977	0.2065	0.2594	0.1735	0.3473	0.0037	0.9960	-0.0454	0.8856	-0.0491	0.8572	0.0343	0.9742	-0.0796	0.8153	-0.1139	0.6974
CG14439	CG14439	1627293_at	1.1797	0.0043	2.2682	0.0512	3.2340	0.0000	0.4351	0.6120	-0.1609	0.7535	-0.5961	0.1196	-0.1334	0.9657	0.9849	0.2236	1.1183	0.2051
---	---	1627294_at	0.2543	0.0933	0.0538	0.8263	-0.1503	0.4910	-0.1431	0.7579	0.0470	0.8568	0.1901	0.2931	0.1094	0.8572	-0.1011	0.7228	-0.2105	0.4047
CycD	cyclin D	1627295_s_at	0.5742	0.0444	0.8452	0.1141	0.3529	0.0720	-0.2932	0.4317	-0.0153	0.9604	0.2779	0.1272	0.1302	0.9342	0.1777	0.7730	0.0475	0.9466
---	---	1627296_at	0.2403	0.3005	-0.0517	0.7058	0.3047	0.0963	0.2180	0.5332	0.1739	0.3493	-0.0441	0.8327	0.0956	0.9056	0.0230	0.9620	-0.0726	0.8444
CG15270	Axs-like	1627297_at	-0.1559	0.3734	-0.0374	0.7612	-0.3437	0.0836	-0.3136	0.4815	-0.1972	0.4242	0.1164	0.6334	0.0000	0.9999	-0.0761	0.7933	-0.0761	0.7813
---	---	1627298_at	0.1473	0.5050	-0.0193	0.8973	0.1913	0.2913	-0.0665	0.9251	-0.0299	0.9200	0.0366	0.8843	0.0291	0.9816	-0.0010	0.9994	-0.0302	0.9371

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13216 /// DmirCG13216	CG13216	1627299_at	-0.0433	0.8554	-0.3047	0.2916	0.0351	0.8765	0.1501	0.8057	0.1555	0.5525	0.0055	0.9858	-0.3184	0.6538	-0.2873	0.3005	0.0312	0.9375
para	paralyzed	1627300_at	0.1855	0.4969	0.2681	0.3887	-0.0157	0.9440	-0.1403	0.7140	-0.2369	0.1732	-0.0965	0.5795	-0.0545	0.9717	-0.1175	0.8132	-0.0631	0.9056
CG7208	CG7208	1627301_s_at	0.5086	0.5179	-0.7573	0.5394	-0.9300	0.2251	-0.8941	0.0144	1.7121	0.0001	2.6062	0.0000	-0.9844	0.8236	0.1845	0.9505	1.1689	0.5488
Fmo-2	Flavin-containing	1627302_at	-0.4314	0.0591	-0.2774	0.6931	-0.3721	0.2243	-0.2012	0.5008	-0.2721	0.0867	-0.0709	0.6662	-0.1129	0.9677	-0.1160	0.9196	-0.0031	0.9982
---	---	1627303_at	0.3222	0.1318	-0.0266	0.8101	0.0446	0.8217	-0.0651	0.9220	0.1264	0.5591	0.1915	0.2921	0.0165	0.9916	-0.0786	0.8814	-0.0950	0.8380
CG13073	CG13073	1627304_at	0.0863	0.6628	0.0437	0.7695	0.2489	0.1032	0.0248	0.9755	0.0373	0.8884	0.0125	0.9597	-0.1670	0.7810	-0.0326	0.9382	0.1344	0.6311
CG5398	CG5398	1627305_at	0.2064	0.3589	-0.1832	0.1131	0.1322	0.4607	0.4069	0.2100	0.3545	0.0719	-0.0525	0.8113	-0.0411	0.9522	0.0313	0.9223	0.0724	0.7615
---	---	1627306_at	0.2075	0.2865	0.0874	0.5842	-0.1730	0.5975	-0.1045	0.9171	0.1211	0.7414	0.2257	0.4402	0.1355	0.9076	0.1056	0.8537	-0.0300	0.9609
CG5681	CG5681	1627307_at	-0.1666	0.2986	-0.1751	0.3771	0.0575	0.7682	0.0671	0.8975	0.0923	0.6282	0.0252	0.9027	-0.1108	0.8494	-0.0051	0.9925	0.1057	0.6958
RpL15	ribosomal L15	1627308_s_at	-0.0356	0.8240	0.3686	0.0515	0.3285	0.1181	-0.0634	0.8836	-0.2928	0.0518	-0.2294	0.0829	-0.0204	0.9831	0.0423	0.9013	0.0627	0.8243
---	---	1627309_at	0.0347	0.8440	0.4205	0.2701	0.1985	0.3957	-0.3136	0.4741	-0.3937	0.0950	-0.0800	0.7561	0.1525	0.8585	-0.0598	0.9105	-0.2123	0.5634
---	---	1627310_at	0.1207	0.4950	0.0214	0.8318	0.1230	0.6423	0.1222	0.8605	0.0803	0.7932	-0.0420	0.8880	0.0939	0.8814	-0.0101	0.9823	-0.1041	0.6978
CG3829	CG3829	1627311_at	0.1113	0.6061	-0.0121	0.9666	0.2443	0.2285	0.1579	0.8132	0.0841	0.7947	-0.0737	0.8039	0.0534	0.9420	0.0443	0.8953	-0.0090	0.9810
RpL12	Ribosomal protein	1627312_at	0.3695	0.4338	0.3783	0.3433	0.5230	0.0375	-0.1104	0.9098	-0.3392	0.2752	-0.2288	0.4310	-0.0652	0.9846	-0.2512	0.7891	-0.1861	0.8470
CG8260	CG8260	1627313_at	0.2300	0.1851	-0.0364	0.8114	-0.0635	0.7868	-0.1348	0.7830	0.1605	0.4446	0.2952	0.1017	-0.0435	0.9571	-0.0945	0.7200	-0.0510	0.8672
CG31864 /// escl	CG31864 /// esc-li	1627314_s_at	0.2692	0.1446	-0.3359	0.0345	-0.2877	0.1451	0.0274	0.9649	0.8442	0.0009	0.8169	0.0006	-0.0620	0.9331	0.2647	0.2468	0.3267	0.1911
Pld	Phospholipase D	1627315_s_at	1.2533	0.0005	0.6244	0.0889	0.5067	0.0079	-0.0179	0.9796	0.6108	0.0043	0.6288	0.0022	0.0544	0.9503	-0.0524	0.8893	-0.1068	0.7148
---	---	1627316_at	0.1579	0.5309	0.0685	0.5235	0.1016	0.6525	-0.0425	0.9672	-0.1140	0.7168	-0.0715	0.8151	0.0076	0.9939	-0.0155	0.9678	-0.0230	0.9433
Tbh	tyramine-beta-hyd	1627317_a_at	0.0178	0.9508	0.0117	0.9564	0.1317	0.3819	0.0287	0.9603	0.0286	0.8906	-0.0001	0.9995	-0.0810	0.9558	0.1016	0.8643	0.1826	0.7013
CG31117	CG31117	1627318_at	-0.1135	0.5456	-0.0343	0.7382	-0.2052	0.2649	-0.0383	0.9466	-0.0633	0.7397	-0.0250	0.8976	0.1336	0.8236	-0.0302	0.9404	-0.1638	0.5340
CG14430	CG14430	1627319_at	-1.2046	0.0171	-1.3976	0.0122	-1.1796	0.0004	-0.0584	0.9029	-0.6378	0.0024	-0.5793	0.0021	-0.2568	0.8533	-0.8834	0.1250	-0.6266	0.2869
zfh2	Zn finger homeod	1627320_at	-0.1835	0.3466	-0.0583	0.6754	-0.1548	0.4477	-0.4516	0.1366	-0.3585	0.0554	0.0932	0.6165	-0.1434	0.8235	-0.1356	0.6397	0.0078	0.9852
GS	Glutathione Synth	1627321_x_at	0.2492	0.1093	0.0392	0.7473	0.2241	0.1646	-0.0807	0.8316	0.0288	0.8814	0.1096	0.4184	0.0369	0.9653	0.0746	0.7963	0.0377	0.9055
CG14300	CG14300	1627322_at	0.2406	0.2181	0.1748	0.4442	0.2255	0.1530	0.2297	0.5744	0.0507	0.8525	-0.1790	0.3528	0.1548	0.8157	0.0388	0.9286	-0.1160	0.7068
CG8388	CG8388	1627323_at	0.1163	0.5619	-0.1645	0.5004	-0.2094	0.2606	-0.2745	0.5008	0.1340	0.5649	0.4084	0.0405	-0.0090	0.9963	-0.0385	0.9557	-0.0295	0.9622
lola	longitudinals abse	1627324_at	-0.4282	0.3335	-1.1846	0.2276	-0.9821	0.0037	0.0669	0.9412	0.8125	0.0078	0.7456	0.0069	-0.1679	0.9647	0.0771	0.9634	0.2450	0.8533
aur	Aurora-A	1627325_at	0.1126	0.6976	-0.4961	0.1357	-0.4314	0.1611	-0.0763	0.9223	0.8260	0.0047	0.9023	0.0019	-0.2941	0.8331	0.0342	0.9720	0.3284	0.5988
CG6860	CG6860	1627326_a_at	-0.9290	0.0219	0.4157	0.2227	-0.6530	0.0089	-0.5379	0.0999	-0.5724	0.0098	-0.0345	0.8802	0.4119	0.7686	0.6374	0.2668	0.2255	0.7427
dro5	Drosomycin B	1627327_at	0.5823	0.0116	0.9617	0.2311	0.7670	0.0076	0.3150	0.5880	-0.1182	0.7334	-0.4332	0.1008	0.2673	0.9174	0.1248	0.9299	-0.1425	0.9084
scw	screw	1627328_at	0.0565	0.7183	0.2541	0.4733	0.0864	0.7737	-0.1203	0.8498	-0.1207	0.6422	-0.0005	0.9988	0.2525	0.8222	0.2150	0.6784	-0.0375	0.9548
CG33558	anon-fast-evolving	1627329_at	0.5790	0.3060	0.1813	0.4115	0.8034	0.0050	0.0354	0.9777	-0.0116	0.9809	-0.0469	0.9034	-0.3845	0.8461	0.0125	0.9941	0.3970	0.6533
CG10189	CG10189	1627330_at	-0.0837	0.6676	0.2810	0.1770	0.8794	0.0010	0.2323	0.3837	-0.5381	0.0043	-0.7705	0.0006	-0.2765	0.7220	-0.1404	0.7078	0.1361	0.7154
Taf5	TBP-associated fa	1627331_at	0.1050	0.7325	0.6129	0.1463	1.1533	0.0006	0.1463	0.7906	-0.4833	0.0382	-0.6296	0.0077	-0.2480	0.8349	0.2739	0.6032	0.5219	0.3027
---	---	1627332_at	0.0802	0.6940	-0.3723	0.2554	-0.0660	0.6942	0.2681	0.5777	0.3649	0.1375	0.0967	0.7121	-0.1043	0.9113	-0.0387	0.9425	0.0657	0.8861
CG11226	CG11226	1627333_at	-4.2679	0.0004	-3.6467	0.0017	-3.2823	0.0001	0.0611	0.9639	-1.1923	0.0058	-1.2534	0.0027	-0.0593	0.9657	-0.2325	0.5483	-0.1732	0.6677
Tsp39D	Tetraspanin 39D	1627334_at	-1.3986	0.0318	-1.0143	0.0313	-1.6568	0.0004	-0.2386	0.7010	0.2364	0.4231	0.4750	0.0659	0.1880	0.9092	0.3735	0.5411	0.1855	0.7923
---	---	1627335_at	0.0501	0.7507	-0.1399	0.5127	0.1235	0.5090	0.1269	0.8470	0.1376	0.6025	0.0107	0.9715	-0.0474	0.9467	-0.0958	0.6949	-0.0483	0.8668
---	---	1627336_at	0.1406	0.5580	0.2025	0.2218	0.0244	0.8914	0.1643	0.6441	0.3515	0.0473	0.1872	0.2262	0.2439	0.6898	0.1868	0.4563	-0.0571	0.8618
CG11555	CG11555	1627337_at	0.3692	0.0868	0.4721	0.3561	0.0963	0.7120	0.1868	0.7164	0.3824	0.1021	0.1956	0.3648	0.4022	0.7266	0.3811	0.4418	-0.0211	0.9791
---	---	1627338_at	0.0257	0.9125	-0.0644	0.5789	-0.0720	0.7060	0.1387	0.7562	0.0223	0.9343	-0.1164	0.5361	0.0953	0.8586	-0.0239	0.9460	-0.1192	0.6096
CG32298	CG32298	1627339_at	0.2122	0.4199	0.1446	0.3344	0.0826	0.7739	0.0050	0.9956	-0.1168	0.5638	-0.1218	0.4994	0.0530	0.9707	-0.0785	0.8807	-0.1315	0.7545
pigeon	lionette	1627340_at	-0.3028	0.3508	-0.2972	0.1953	-0.1530	0.4456	0.3020	0.6416	0.7959	0.0202	0.4939	0.0823	-0.0327	0.9848	0.5917	0.1243	0.6244	0.1313
CG31556	CG31556	1627341_at	0.1371	0.4152	-0.1252	0.3311	0.2274	0.1535	0.2364	0.5859	0.2773	0.2083	0.0409	0.8762	-0.1499	0.7644	0.0087	0.9829	0.1586	0.4704
CG15251	CG15251	1627342_at	-0.1151	0.6724	0.2190	0.1104	-0.0126	0.9619	-0.3669	0.4073	-0.9312	0.0033	-0.5642	0.0185	0.0545	0.9741	-0.3067	0.4600	-0.3612	0.3907
CG5535	CG5535	1627343_a_at	0.8853	0.0406	1.4431	0.0627	1.4019	0.0009	0.7228	0.1074	0.5499	0.0447	-0.1729	0.4993	0.7267	0.7046	1.1059	0.1528	0.3792	0.6607
---	---	1627344_at	0.0095	0.9736	-0.0925	0.4736	-0.1848	0.4950	0.1731	0.7734	0.0819	0.7905	-0.0911	0.7369	0.2300	0.8122	-0.0706	0.9085	-0.3006	0.4686
HDC15381	HDC15381	1627345_at	0.0314	0.9269	-0.0593	0.5480	0.0897	0.7757	0.0053	0.9956	0.0815	0.8272	0.0762	0.8204	-0.2018	0.8270	-0.1457	0.7455	0.0561	0.9158
---	---	1627346_at	0.3164	0.1024	0.1736	0.1251	-0.0159	0.9477	-0.0521	0.9232	0.0429	0.8382	0.0949	0.5542	0.1728	0.7726	0.1575	0.5591	-0.0153	0.9691
---	---	1627347_at	0.2145	0.3602	0.2982	0.0813	0.1273	0.6180	-0.0188	0.9803	-0.0369	0.8810	-0.0182	0.9367	0.1798	0.8461	0.1075	0.8331	-0.0723	0.8907
CG5285	CG5285	1627348_at	0.7811	0.0051	1.1187	0.0140	1.5057	0.0001	0.1124	0.8132	0.1708	0.3726	0.0584	0.7794	-0.3844	0.5186	0.4043	0.1471	0.7887	0.0410
Bka	missing imaginal	1627349_at	0.1195	0.7349	0.3204	0.4524	0.3456	0.0510	0.1571	0.8067	0.2190	0.4034	0.0619	0.8312	0.0190	0.9928	0.3127	0.5284	0.2937	0.5630
CG12420	CG12420	1627350_at	-0.0208	0.9267	-0.1860	0.3656	0.0618	0.7672	0.1322	0.7469	0.1415	0.4468	0.0093	0.9674	-0.2344	0.7230	-0.0493	0.9111	0.1852	0.5409
CG15335	CG15335	1627351_at	0.2072	0.2788	0.0590	0.6452	0.2866	0.2179	0.0989	0.8154	0.2480	0.1343	0.1491	0.3258	-0.0015	0.9994	0.0878	0.7897	0.0892	0.7734
Oat	Ornithine aminotra	1627352_at	-0.3233	0.3834	-0.4286	0.5548	0.6566	0.0268	-0.0654	0.9435	-0.8396	0.0069	-0.7742	0.0059	-1.0457	0.6145	-1.0457	0.2212	0.0000	1.0000
Gr59f	Gustatory recepto	1627353_at	0.0157	0.9501	0.0692	0.7400	0.1766	0.4061	0.1042	0.8880	-0.0846	0.7795	-0.1888	0.4184	-0.1286	0.8235	-0.0551	0.8721	0.0735	0.8011
CG11779 /// CG5835 /// Da	CG11779 /// CG5	1627354_at	1.3438	0.0092	0.9461	0.1661	2.1639	0.0051	1.3056	0.0262	1.7363	0.0007	0.4307	0.1294	0.0691	0.9898	1.2682	0.2430	1.1992	0.2993
---	---	1627355_at	-0.0234	0.8821	0.1440	0.3382	0.1430	0.3375	-0.1791	0.5617	-0.0638	0.7304	0.1153	0.4386	-0.0413	0.9514	0.0806	0.7269	0.1220	0.5649
---	---	1627356_at	0.1171	0.5879	0.2109	0.3288	0.2750	0.2710	0.0544	0.9345	-0.1657	0.4103	-0.2201	0.2075	-0.1719	0.8743	0.0026	0.9989	0.1745	0.7170
---	---	1627357_at	-0.1020	0.6277	-0.0314	0.7568	0.0429	0.8982	-0.0229	0.9803	-0.0613	0.8286	-0.0385	0.8846	-0.0462	0.9599	0.0479	0.9049	0.0942	0.7544
---	---	1627358_at	0.0171	0.9523	0.2624	0.2176	0.2488	0.2054	-0.0067	0.9937	-0.1770	0.3270	-0.1704	0.2906	0.0176	0.9894	0.1080	0.7416	0.0904	0.7847
CG12398	CG12398	1627359_at	0.0844	0.7229	0.0310	0.8944	0.0811	0.6165	0.1172	0.8473	0.2044	0.3695	0.0872	0.7129	0.0309	0.9816	0.0413	0.9277	0.0105	0.9835
pyd	tamou	1627360_at	0.6156	0.2936	0.4818	0.4455	0.5250	0.0061	-0.1429	0.7023	0.0710	0.7229	0.2139	0.1628	0.0420	0.9913	0.2653	0.7943	0.2233	0.8256
---	---	1627361_at	0.0811	0.6131	-0.0244	0.8236	-0.1622	0.4141	0.1375	0.6511	0.1918	0.1890	0.0544	0.7295	0.2600	0.6557	0.2440	0.2912	-0.0159	0.9637
Su(var)2-10	zinc finger-contair	1627362_a_at	-0.1247	0.6681	-0.1605	0.6498	-0.2671	0.4070	-0.4479	0.6108	0.1926	0.7053	0.6405	0.1036	-0.1833	0.8810	0.2501	0.6170	0.4334	0.3671
---	---	1627363_at	-0.0818	0.6272	-0.1312	0.4918	-0.0518	0.8128	-0.0415	0.9346	-0.0319	0.8724	0.0096	0.9593	-0.1120	0.8744	-0.1075	0.7424	0.0045	0.9924
RpS12	40S ribosomal prc	1627364_s_at	0.2442	0.1514	0.6197	0.0114	0.6775	0.0025	0.0636	0.9117	-0.2494	0.1695	-0.3130	0.0590	0.0515	0.9407	-0.0064	0.9886	-0.0580	0.8301
---	---	1627365_at	0.1557	0.5478	0.1797	0.3000	0.0138	0.9619	-0.1281	0.7140	0.0510	0.7940	0.1792	0.2091	0.1707	0.8802	0.1448	0.7897	-0.0259	0.9690
---	---	1627366_at	-0.0885	0.7292	0.0191	0.8504	0.1271	0.5513	0.2397	0.5842	0.1560	0.5091	-0.0837	0.7255	0.1077	0.8751	0.1167	0.7010	0.0090	0.9843
CG5653	CG5653	1627367_at	0.1505	0.4927	-0.4417	0.3499	-0.3423	0.1535	0.1859	0.8631	0.6263	0.1020	0.4404	0.1980	0.0207	0.9894	-0.1688	0.6341	-0.1895	0.5876

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15497	CG15497	1627368_at	-0.4256	0.1376	0.3640	0.2338	0.1247	0.4711	-0.0155	0.9921	-0.7909	0.0211	-0.7754	0.0143	0.0886	0.9238	0.1369	0.6915	0.0483	0.9095
CG14581	CG14581	1627369_at	0.1851	0.3178	0.2189	0.3664	0.0762	0.6715	-0.1210	0.7929	-0.1002	0.6325	0.0208	0.9276	-0.0206	0.9853	0.0282	0.9462	0.0488	0.8904
CG7835	CG7835	1627370_at	0.0958	0.7022	0.1954	0.2660	0.3599	0.0632	-0.3710	0.3016	-0.2353	0.2669	0.1357	0.5041	-0.0110	0.9948	0.0949	0.8619	0.1059	0.8299
CG4744	CG4744	1627371_at	0.1001	0.6457	0.1058	0.3555	0.2333	0.2391	0.0558	0.9154	-0.0161	0.9478	-0.0719	0.6764	0.0184	0.9898	0.1117	0.7554	0.0933	0.7980
lig	transcription unit /	1627372_a_at	0.3049	0.6811	-0.2995	0.7173	-0.5065	0.0729	0.0846	0.8578	0.8228	0.0012	0.7383	0.0011	0.3995	0.9092	0.1994	0.9175	-0.2001	0.9073
Tsp42Eg	tetraspanin 42E	1627373_at	-0.9961	0.0167	-0.3405	0.4025	-1.6067	0.0027	-0.4016	0.1658	-0.1075	0.5711	0.2941	0.0647	0.7394	0.7062	0.4925	0.5647	-0.2469	0.8044
---	---	1627374_at	-0.0434	0.7795	-0.0690	0.7757	-0.0573	0.7950	0.0112	0.9931	0.0568	0.8656	0.0456	0.8801	-0.0282	0.9848	-0.0189	0.9727	0.0093	0.9856
---	---	1627375_at	0.2346	0.2193	-0.0592	0.7777	-0.0064	0.9835	0.1073	0.8103	0.2719	0.1243	0.1646	0.3075	-0.0203	0.9885	-0.1333	0.7002	-0.1130	0.7492
Rel	relish	1627376_at	-0.0484	0.8409	-1.7353	0.0039	-1.4573	0.0089	0.5156	0.4220	1.8581	0.0009	1.3425	0.0020	0.3136	0.8270	0.3157	0.6236	0.0021	0.9987
Grip75	Grip75	1627377_at	-0.8864	0.0544	-0.8461	0.0136	-0.7584	0.0012	-0.1605	0.8251	0.1112	0.7377	0.2717	0.2863	-0.2357	0.7415	0.0439	0.9269	0.2796	0.3671
CG31395	CG31395	1627378_at	0.2640	0.1623	-0.0981	0.5029	-0.2253	0.3647	-0.2237	0.7149	0.1779	0.5545	0.4016	0.1073	0.0286	0.9810	-0.0416	0.9170	-0.0702	0.8287
zormin	D-Titin	1627379_at	-0.0796	0.6940	0.2548	0.2508	0.0242	0.9432	-0.2745	0.5808	-0.3267	0.1963	-0.0522	0.8599	0.0662	0.9520	0.1003	0.8153	0.0340	0.9414
RpA-70	Drosophila Replic	1627380_at	-0.0125	0.9629	-0.4071	0.0815	-0.5136	0.0606	-0.1015	0.7857	0.6798	0.0018	0.7813	0.0006	0.0338	0.9871	0.2964	0.5531	0.2626	0.6109
CG40158	CG40158	1627381_at	-0.0555	0.7573	-0.0283	0.7957	0.0724	0.6685	0.1341	0.7312	0.0547	0.7982	-0.0794	0.6604	0.0541	0.9400	0.0615	0.8352	0.0074	0.9842
Sec61beta	Sec61beta	1627382_at	0.9306	0.0015	1.0905	0.0047	1.1506	0.0002	0.0222	0.9704	0.1933	0.2025	0.1711	0.2090	-0.0710	0.9238	0.4590	0.0767	0.5300	0.0702
asp	abnormal spindle	1627383_at	0.3501	0.3215	0.0279	0.9372	-0.9547	0.0434	-0.7545	0.0300	0.6158	0.0066	1.3703	0.0002	0.0230	0.9939	0.1852	0.8323	0.1622	0.8474
CG6225	CG6225	1627384_at	-4.2257	0.0006	-4.5065	0.0026	-4.0999	0.0001	0.3568	0.7604	0.1152	0.8612	-0.2416	0.6390	-0.2171	0.8689	-0.3207	0.5543	-0.1036	0.8844
Pbprp1	Pheromone-bindir	1627385_at	0.7553	0.1746	0.2570	0.4705	0.4807	0.0283	-0.3693	0.4861	-0.0450	0.9089	0.3243	0.2000	-0.3608	0.8235	-0.2551	0.7466	0.1057	0.9095
CG34017	CG34017	1627386_at	0.0592	0.7230	0.1501	0.4543	0.1454	0.3483	0.0360	0.9584	0.0019	0.9942	-0.0341	0.8738	0.0002	0.9999	0.0119	0.9784	0.0117	0.9761
---	---	1627387_at	-0.0088	0.9678	0.0872	0.4443	0.0233	0.9200	-0.0660	0.9037	-0.1096	0.5669	-0.0436	0.8293	0.1172	0.8182	0.0863	0.7183	-0.0309	0.9152
Lcp65Ag3	Larval cuticle prot	1627388_at	-0.0447	0.7780	-0.2132	0.2311	-0.1503	0.3255	-0.0155	0.9777	0.0479	0.7828	0.0634	0.6704	-0.0795	0.9101	-0.1235	0.6503	-0.0440	0.8984
---	---	1627389_at	0.0979	0.7241	-0.0830	0.6395	0.0213	0.9292	-0.0155	0.9860	0.0589	0.8232	0.0744	0.7433	0.0372	0.9751	-0.0873	0.8208	-0.1245	0.7071
CG2926	CG2926	1627390_at	-0.0414	0.9578	0.6656	0.3296	0.5762	0.0547	-0.0628	0.9649	-0.0155	0.9790	0.0473	0.9189	-0.0050	0.9994	0.7471	0.4368	0.7521	0.4475
---	---	1627391_s_at	0.1430	0.3603	0.0276	0.8056	0.6606	0.0115	0.1728	0.5765	0.1087	0.5184	-0.0640	0.7017	-0.3016	0.6584	0.0368	0.9341	0.3384	0.2398
---	---	1627392_at	0.3128	0.1769	0.1336	0.6358	0.3468	0.0344	0.2079	0.7327	0.2370	0.3949	0.0291	0.9305	0.0274	0.9798	0.0715	0.8218	0.0441	0.8949
CG9029	CG9029	1627393_at	0.2730	0.3071	-0.0593	0.5836	-0.0004	0.9984	0.0611	0.9346	-0.0196	0.9526	-0.0807	0.7330	0.1324	0.8890	-0.1990	0.6019	-0.3314	0.3673
aop	pokkuri	1627394_s_at	-0.1169	0.7386	-0.6374	0.0561	-1.1666	0.0002	0.0217	0.9782	0.6057	0.0095	0.5840	0.0068	0.4507	0.6045	-0.0218	0.9727	-0.4725	0.2253
CG34123	CG34123	1627395_a_at	0.2070	0.5320	0.1629	0.3736	0.6190	0.0070	-0.1453	0.7749	-0.0648	0.8042	0.0805	0.7224	-0.4715	0.7142	0.0623	0.9425	0.5339	0.3307
su(f)	lethal3Des	1627396_a_at	0.2498	0.5130	-0.0791	0.8799	0.3268	0.1440	-0.2896	0.7123	-0.4416	0.2155	-0.1520	0.6851	-0.5440	0.7131	-0.7898	0.1875	-0.2457	0.7269
CG33515	CG33515	1627397_at	0.0111	0.9601	0.0605	0.6031	-0.0083	0.9713	-0.2165	0.5068	-0.0587	0.7795	0.1578	0.3167	-0.0407	0.9459	0.0788	0.7007	0.1195	0.5322
Gbeta13F	G-protein beta 13	1627398_a_at	-0.2032	0.4915	0.0072	0.9864	-0.0032	0.9916	0.1593	0.6533	0.2661	0.1176	0.1068	0.5193	0.1995	0.8811	0.4458	0.3726	0.2463	0.6545
CG17266	CG17266	1627399_at	0.6210	0.0303	0.7502	0.0290	0.5237	0.1106	-0.2015	0.7327	-0.2979	0.2590	-0.0965	0.7323	0.1123	0.9081	0.0834	0.8625	-0.0289	0.9525
CG5613	CG5613	1627400_a_at	-1.8899	0.0016	-0.9674	0.0500	-0.8511	0.0422	-0.2343	0.7805	-1.2239	0.0047	-0.9896	0.0073	-0.3600	0.7743	-0.5191	0.3268	-0.1591	0.8138
CG6789	CG6789	1627401_at	-0.1888	0.6838	-1.1519	0.0616	-1.0436	0.0577	0.1521	0.9353	1.1989	0.0369	1.0468	0.0406	0.0449	0.9860	0.2226	0.7400	0.1777	0.7967
CG7630	CG7630	1627402_a_at	0.0757	0.8211	1.5149	0.0340	1.1984	0.0008	-0.1149	0.8717	-1.5835	0.0004	-1.4686	0.0003	0.1306	0.9467	-0.2356	0.7346	-0.3662	0.5629
Paf-AHalpha	platelet-activating	1627403_s_at	0.2404	0.1777	-0.2427	0.4743	-0.6311	0.0078	-0.0205	0.9749	0.8777	0.0007	0.8982	0.0004	0.2309	0.8384	0.3117	0.5186	0.0808	0.9004
---	---	1627404_at	0.1766	0.3287	-0.0181	0.8573	-0.0163	0.9467	0.2146	0.4464	0.0736	0.6746	-0.1410	0.3135	0.0150	0.9914	-0.0588	0.8940	-0.0738	0.8467
CG14966	CG14966	1627405_at	-0.5886	0.0158	0.5182	0.2216	1.1782	0.0008	0.0425	0.9641	-0.9802	0.0029	-1.0227	0.0015	-0.6008	0.5126	0.1802	0.7202	0.7810	0.1116
CG3009	CG3009	1627406_at	0.0623	0.7722	-0.1248	0.2185	-0.1941	0.2628	-0.0813	0.8534	-0.0302	0.8884	0.0511	0.7713	0.0058	0.9963	-0.0179	0.9684	-0.0236	0.9512
SNCF	SoxNeuro Co-Fac	1627407_at	0.1191	0.4180	0.2534	0.2273	0.1400	0.4079	0.0497	0.9436	-0.0589	0.8128	-0.1086	0.5890	0.0346	0.9650	0.0487	0.8718	0.0140	0.9640
CG32107	CG32107	1627408_at	-0.0135	0.9700	-0.1069	0.4267	-0.2276	0.2718	-0.0266	0.9777	-0.0017	0.9960	0.0250	0.9307	-0.0781	0.9142	-0.0870	0.7801	-0.0088	0.9837
CG15727	CG15727	1627409_at	-0.1960	0.2790	0.3798	0.1101	0.2292	0.1348	-0.1519	0.7143	-0.3470	0.0700	-0.1951	0.2518	-0.0641	0.9409	0.2237	0.3976	0.2878	0.2960
Amph	amphiphysin	1627410_at	-1.2466	0.0025	-1.0156	0.1110	-2.1763	0.0001	-0.5792	0.1951	-1.0138	0.0035	-0.4346	0.0739	0.3739	0.7953	-0.8768	0.1405	-1.2507	0.0750
CG14427	CG14427	1627411_at	0.1845	0.3758	0.0449	0.8699	0.2792	0.2140	0.2245	0.6096	0.1217	0.6184	-0.1028	0.6520	0.1436	0.8806	0.1321	0.7642	-0.0115	0.9849
---	---	1627412_at	-0.1742	0.2748	-0.0992	0.4493	-0.0883	0.7185	-0.1204	0.7388	-0.0512	0.7932	0.0692	0.6778	-0.2014	0.7707	-0.0552	0.9022	0.1462	0.6457
CG34383	CG14365	1627413_at	0.2064	0.4893	0.1682	0.4886	0.1398	0.4615	0.0253	0.9639	-0.0593	0.7316	-0.0846	0.5622	0.0612	0.9775	0.0177	0.9865	-0.0434	0.9525
---	---	1627414_at	-0.0320	0.8832	0.1056	0.3091	-0.1875	0.2769	-0.0616	0.8959	-0.0686	0.7021	-0.0070	0.9711	0.1152	0.8270	-0.0486	0.8782	-0.1638	0.4741
CG9986	CG9986	1627415_at	0.4248	0.0589	0.0062	0.9891	-0.0424	0.8028	-0.0307	0.9761	0.7832	0.0081	0.8139	0.0040	-0.0470	0.9716	0.2815	0.3802	0.3285	0.3262
CG32182	CG32182	1627416_at	0.0258	0.8808	0.1148	0.3955	0.2426	0.1508	0.2684	0.3166	0.0604	0.7439	-0.2080	0.1352	-0.0145	0.9898	0.0750	0.8119	0.0895	0.7492
CG12038	CG12038	1627417_s_at	-0.0201	0.9112	0.1253	0.4479	-0.0751	0.7616	0.0000	0.9999	-0.1017	0.7064	-0.1017	0.6776	0.0563	0.9309	0.0645	0.8132	0.0082	0.9819
---	---	1627418_at	0.1399	0.3108	-0.0925	0.4771	-0.1574	0.2822	0.0685	0.9136	0.3058	0.1224	0.2373	0.1809	-0.0110	0.9898	-0.1266	0.5042	-0.1156	0.5515
His4r	Histone H4 replac	1627419_at	-0.2092	0.3390	-0.0636	0.8835	-0.5080	0.0324	-0.0509	0.9695	-0.3012	0.3904	-0.2503	0.4364	0.3887	0.7266	-0.0008	0.9998	-0.3895	0.4285
CG7702	connectin-like	1627420_s_at	0.2671	0.7933	-0.1529	0.8273	0.4482	0.3767	0.4519	0.8172	0.2977	0.7425	-0.1542	0.8629	-0.1496	0.9735	-0.0439	0.9838	0.1056	0.9462
CG18292	CG18292	1627421_at	-0.3463	0.4582	-0.3353	0.4873	-0.8104	0.1333	-0.0413	0.9620	0.1260	0.6210	0.1673	0.4389	0.4833	0.8521	0.1640	0.9221	-0.3193	0.8095
---	---	1627422_at	0.0933	0.5614	0.0144	0.9233	0.2215	0.1492	0.1246	0.7409	0.1237	0.4762	-0.0009	0.9966	-0.0902	0.8504	0.1026	0.6312	0.1928	0.3439
CG6178	CG6178	1627423_at	1.6134	0.0084	0.7027	0.0249	1.1495	0.0001	0.0607	0.9518	0.6190	0.0339	0.5583	0.0328	-0.2996	0.7498	-0.3299	0.4067	-0.0303	0.9584
---	---	1627424_at	0.1062	0.5759	0.3633	0.0467	0.0019	0.9959	-0.1854	0.5838	-0.2676	0.1187	-0.0822	0.6423	0.1582	0.8270	-0.0218	0.9648	-0.1800	0.5709
CG30465	CG30465	1627425_at	-0.0727	0.7054	0.0449	0.6988	0.0485	0.7652	0.0410	0.9436	0.0028	0.9903	-0.0382	0.8407	-0.0880	0.8692	0.1144	0.6183	0.2024	0.3578
CG3403	CG3403	1627426_at	-0.2760	0.0934	0.2552	0.1518	-0.0417	0.8135	-0.1187	0.7929	-0.4491	0.0230	-0.3304	0.0497	0.1173	0.8305	0.0381	0.9149	-0.0792	0.7674
CG9002	CG9002	1627427_s_at	0.1688	0.2305	-0.1786	0.3322	0.1546	0.3285	0.1582	0.7409	0.2676	0.2030	0.1095	0.6075	-0.1050	0.8284	-0.1299	0.5404	-0.0249	0.9300
---	---	1627428_at	-0.0945	0.5435	0.0463	0.7341	-0.0109	0.9625	-0.1369	0.7143	-0.1590	0.3616	-0.0220	0.9157	-0.1150	0.8461	-0.1285	0.6264	-0.0135	0.9716
---	---	1627429_at	-0.0604	0.7642	0.2533	0.0974	0.1458	0.4603	-0.1961	0.6970	-0.2428	0.2995	-0.0467	0.8626	-0.0702	0.9342	0.0410	0.9237	0.1112	0.7201
CG8888	CG8888	1627430_at	-3.4379	0.0162	-4.8029	0.0157	-5.0094	0.0000	0.0235	0.9817	0.0352	0.9175	0.0117	0.9692	0.3699	0.9459	-1.2339	0.4566	-1.6037	0.3451
CG8155	CG8155	1627431_at	0.3840	0.1827	0.2213	0.2176	-0.0306	0.8583	-0.0154	0.9798	0.2028	0.1665	0.2181	0.0974	0.3861	0.6557	0.0939	0.8439	-0.2922	0.4132
CG6041	CG6041	1627432_at	-0.2245	0.8890	-0.2109	0.3103	-0.8292	0.0014	-0.3526	0.8623	-0.6665	0.3684	-0.3139	0.6789	0.0055	0.9994	-0.8766	0.4523	-0.8821	0.4643
CG3163	CG3163	1627433_at	0.1929	0.3094	-0.4451	0.1895	-0.4027	0.0332	0.1135	0.7145	0.5271	0.0036	0.4136	0.0063	-0.0586	0.9677	-0.2117	0.6032	-0.1532	0.7245
CG5741	CG5741	1627434_at	-1.1492	0.0049	-0.4577	0.3704	-1.5368	0.0000	-0.0841	0.9004	0.5010	0.0291	0.5851	0.0093	1.1524	0.2884	1.3902	0.0391	0.2378	0.7113
clumsy	clumsy	1627435_at	-2.6923	0.0014	-2.7661	0.0083	-3.3043	0.0000	-0.2377	0.5539	-0.2024	0.3373	0.0353	0.8862	0.0713	0.9806	-0.4097	0.5401	-0.4810	0.4683
Pdp1	PAR-Domain Prot	1627436_s_at	-0.7205	0.0694	-0.6132	0.3455	-1.8454	0.0023	-0.3823	0.2654	-0.0935	0.6875	0.2888	0.1106	0.6976	0.7506	0.0699	0.9642	-0.6277	0.5239

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
ref2	ref2	1627437_at	0.1719	0.4304	-0.0310	0.9095	0.2014	0.3791	0.0229	0.9745	0.0935	0.6307	0.0706	0.7013	-0.0939	0.9461	-0.0684	0.9167	0.0255	0.9684
CG31084	CG31084	1627438_at	0.2825	0.2853	-0.0746	0.8439	0.1870	0.2288	0.1156	0.8267	0.2975	0.1426	0.1819	0.3282	0.0138	0.9939	-0.0630	0.9232	-0.0768	0.8934
CG14840	CG14840	1627439_at	-0.0235	0.8979	0.1223	0.3364	-0.1621	0.4568	-0.2593	0.4586	-0.3041	0.1073	-0.0448	0.8363	0.2015	0.6898	0.0236	0.9443	-0.1779	0.4008
---	---	1627440_at	-0.0125	0.9548	-0.0084	0.9642	-0.1352	0.4542	0.0385	0.9387	0.0149	0.9453	-0.0236	0.8930	0.1235	0.8331	0.0062	0.9911	-0.1172	0.6583
TyrR	honoka	1627441_at	-0.1884	0.2589	-0.0711	0.6522	-0.0958	0.5976	-0.0193	0.9753	-0.0202	0.9243	-0.0009	0.9964	-0.0493	0.9523	0.0348	0.9291	0.0842	0.7734
---	---	1627442_at	0.1229	0.4134	-0.2002	0.2827	-0.0894	0.7663	-0.0117	0.9909	0.4835	0.0347	0.4952	0.0199	0.0258	0.9854	0.1604	0.6483	0.1345	0.7134
CG5941	CG5941	1627443_at	0.2123	0.4771	0.5046	0.1990	0.3241	0.1756	-0.0936	0.9375	-0.3674	0.2931	-0.2738	0.3953	0.0318	0.9857	-0.0830	0.8858	-0.1149	0.8178
CG31819	CG31819	1627444_at	0.2159	0.3210	-0.0246	0.8900	-0.0235	0.9117	0.2825	0.5008	0.2641	0.2364	-0.0185	0.9479	0.0927	0.8940	0.0939	0.7667	0.0012	0.9979
en	spermatheca	1627445_s_at	-2.4026	0.0007	-1.1642	0.1165	-3.0447	0.0001	-1.4511	0.1009	-1.5301	0.0104	-0.0790	0.9000	0.0393	0.9852	-0.2765	0.5931	-0.3158	0.5391
net	net	1627446_at	-0.0823	0.8729	-0.3959	0.1636	-0.5908	0.1009	-0.1665	0.9228	0.0977	0.8889	0.2642	0.6136	-0.2363	0.6898	-0.4224	0.0900	-0.1861	0.4563
CG14759	CG14759	1627447_at	0.0666	0.6896	0.0457	0.6688	0.2383	0.1489	0.1254	0.6988	0.0083	0.9714	-0.1171	0.3972	0.0391	0.9588	-0.0096	0.9805	-0.0487	0.8602
Updo	Updo	1627448_at	-0.2926	0.2708	-0.4684	0.0314	-0.1377	0.4367	0.1210	0.7351	0.2344	0.1389	0.1134	0.4494	-0.1906	0.8202	0.0589	0.9111	0.2495	0.4841
CG8111	CG8111	1627449_at	-0.4643	0.0175	-0.4821	0.0163	-0.3041	0.2057	0.0666	0.9029	0.3416	0.0579	0.2749	0.0833	0.0350	0.9657	0.3202	0.1397	0.2852	0.2130
CG40282	CG40282	1627450_at	0.1815	0.3020	0.2820	0.1500	0.1969	0.4091	-0.1470	0.7970	-0.1183	0.6505	0.0287	0.9195	0.0582	0.9226	0.0717	0.7656	0.0135	0.9630
CG14538	CG14538	1627451_at	0.1448	0.5511	0.0135	0.8979	0.1013	0.5052	0.0933	0.8350	0.1156	0.5214	0.0222	0.9143	0.0721	0.9343	0.1513	0.6157	0.0793	0.8223
l(3)mbt	tumor-supressor	1627452_a_at	-0.7739	0.0482	-0.6419	0.2592	-0.0405	0.9046	0.1502	0.8792	-0.2218	0.5396	-0.3720	0.2177	-0.3645	0.8062	0.0216	0.9864	0.3860	0.5579
spi	spitz	1627453_s_at	-0.7549	0.0039	-0.8914	0.0500	-0.9334	0.0017	-0.0538	0.9110	-0.0912	0.5818	-0.0374	0.8296	0.0095	0.9952	-0.2882	0.4796	-0.2977	0.4710
cora	Coracle	1627454_a_at	-1.2465	0.0975	-0.5745	0.5227	-0.9335	0.0019	0.0971	0.8794	-0.2083	0.3491	-0.3054	0.1171	0.3984	0.9137	0.5669	0.6949	0.1685	0.9237
CG4448	CG4448	1627455_at	0.1216	0.5576	0.1465	0.5588	-0.1749	0.3109	-0.1658	0.7512	0.2279	0.3221	0.3936	0.0567	0.2083	0.7726	0.2468	0.4197	0.0386	0.9279
mub	mushroom bodies	1627456_s_at	-0.0704	0.8882	-0.0356	0.9337	-0.5321	0.1358	-0.0757	0.9038	0.1120	0.6196	0.1878	0.3108	0.3606	0.8608	0.0280	0.9868	-0.3326	0.7228
CG10793	CG10793	1627457_at	0.0585	0.8225	0.0443	0.6715	-0.0816	0.7269	-0.2417	0.5842	-0.1337	0.5847	0.1080	0.6402	-0.0202	0.9884	-0.1366	0.6827	-0.1163	0.7355
Catsup	catsup	1627458_at	0.3838	0.0424	1.5354	0.0109	1.7431	0.0000	-0.0002	0.9997	-0.5675	0.0033	-0.5673	0.0020	-0.3011	0.6955	0.4624	0.1460	0.7636	0.0559
CG9154	CG9154	1627459_at	-0.1571	0.6183	0.4146	0.2035	0.6059	0.0222	0.2816	0.4779	-0.0182	0.9542	-0.2998	0.1140	0.1341	0.9342	0.5453	0.2814	0.4112	0.4450
CG18343	CG18343	1627460_at	0.3770	0.0466	0.4687	0.1095	0.4653	0.0133	-0.0117	0.9893	0.0531	0.8365	0.0648	0.7696	0.0315	0.9816	0.0355	0.9402	0.0041	0.9937
bsf	bicoid stability fac	1627461_at	-0.7378	0.0220	-0.2120	0.3424	-0.1433	0.5478	0.0342	0.9672	-0.3695	0.0914	-0.4036	0.0444	-0.0535	0.9717	0.2237	0.5835	0.2772	0.4865
CG4662	CG4662	1627462_a_at	-1.0524	0.0044	-0.5379	0.2427	-0.4485	0.0328	-0.0930	0.8791	-0.7119	0.0052	-0.6189	0.0058	-0.1507	0.8928	-0.1710	0.7200	-0.0203	0.9758
Damm	Death associated	1627463_at	-0.6915	0.7642	-0.0579	0.6709	-0.1167	0.5142	-0.0823	0.9921	-1.2856	0.4667	-1.2032	0.4511	-0.0713	0.9916	-0.9226	0.5824	-0.8513	0.6171
LvpD	hdl cuticle gene cl	1627464_at	0.0226	0.9958	-0.1665	0.4549	-0.6883	0.0667	-0.3841	0.9610	-1.6546	0.4490	-1.2705	0.5357	0.1281	0.9923	-1.8768	0.5641	-2.0049	0.5403
Mes4	NF-YC-like	1627465_at	-0.3752	0.1169	-0.3593	0.4522	-0.4478	0.0105	-0.4537	0.4001	-0.5570	0.0681	-0.1033	0.7540	-0.2382	0.7644	-0.5064	0.1353	-0.2682	0.4424
---	---	1627466_at	-0.1718	0.2339	0.1381	0.2604	0.1893	0.2210	-0.0457	0.9314	-0.2784	0.0774	-0.2327	0.0961	0.0056	0.9943	0.1234	0.5146	0.1177	0.5420
CG13949	CG13949	1627467_at	0.2845	0.1313	0.0573	0.6445	0.0559	0.7281	-0.1487	0.6932	-0.0419	0.8528	0.1068	0.5293	0.0042	0.9964	-0.1440	0.5042	-0.1481	0.4931
Gr43b	Gr43b	1627468_at	-0.1567	0.6270	-0.9479	0.0346	-0.4759	0.0502	0.5874	0.3625	0.7479	0.0472	0.1605	0.6763	0.1097	0.9016	-0.1488	0.6820	-0.2585	0.4399
CG16736	CG16736	1627469_at	0.1046	0.5831	0.1129	0.4543	0.1202	0.5357	-0.0892	0.8053	-0.0531	0.7580	0.0360	0.8264	-0.1234	0.8379	-0.0233	0.9533	0.1002	0.7228
phl	Pole hole	1627470_at	-0.0057	0.9795	0.3413	0.1110	0.2446	0.1166	0.0916	0.8424	-0.0677	0.7372	-0.1593	0.3053	0.1856	0.7726	0.3157	0.2400	0.1300	0.6673
dmrt99B	doublesex-Mab re	1627471_at	0.5103	0.0362	0.1245	0.5847	0.0780	0.7193	0.0956	0.9046	0.1956	0.4710	0.1000	0.7172	0.1707	0.8069	-0.0694	0.8649	-0.2401	0.4175
AnnX	Annexin X	1627472_at	-0.8020	0.0149	-0.6639	0.3409	-0.3959	0.0966	0.1325	0.7568	0.2563	0.1617	0.1239	0.4805	-0.2060	0.9246	0.2698	0.7572	0.4757	0.5369
CG8090	CG8090	1627473_at	0.5423	0.0560	-0.0177	0.9771	-0.1072	0.5799	-0.1954	0.5633	0.6245	0.0041	0.8199	0.0008	-0.0570	0.9816	0.2035	0.7551	0.2605	0.6607
---	---	1627474_at	0.1563	0.4690	0.1739	0.3879	0.1311	0.5005	0.0443	0.9584	0.0783	0.7709	0.0340	0.8994	0.0698	0.9412	0.0653	0.8764	-0.0044	0.9931
---	---	1627475_at	0.3153	0.2444	0.0446	0.9016	-0.1078	0.6704	0.1134	0.8388	0.1718	0.4266	0.0584	0.8039	0.1868	0.8472	-0.0975	0.8625	-0.2843	0.4975
BBS8	BBS8	1627476_at	0.1758	0.2936	0.0000	1.0000	0.1718	0.4270	0.2092	0.5808	0.2883	0.1348	0.0791	0.6976	-0.0504	0.9449	0.0640	0.8258	0.1144	0.6318
---	---	1627477_at	0.0552	0.8099	-0.0119	0.9113	-0.1361	0.3959	-0.1812	0.6615	0.0863	0.7060	0.2675	0.1294	0.1244	0.8276	0.0261	0.9462	-0.0983	0.7156
CG33232	villin-like	1627478_at	-0.8093	0.0529	-0.4696	0.5244	-0.3508	0.2077	-0.0427	0.9603	-0.5962	0.0164	-0.5536	0.0140	-0.2268	0.9330	-0.3613	0.7127	-0.1345	0.9100
---	---	1627479_at	0.1487	0.3302	0.0024	0.9912	0.1207	0.4375	0.0450	0.9353	0.0884	0.6281	0.0434	0.8145	-0.2039	0.7387	-0.2433	0.3366	-0.0393	0.9121
---	---	1627480_at	-0.2137	0.3098	0.2187	0.2183	0.1760	0.3347	-0.0492	0.9376	-0.2351	0.2003	-0.1858	0.2629	0.0117	0.9913	0.1612	0.4670	0.1496	0.5115
CG15615	CG15615	1627481_at	0.1899	0.2937	0.1598	0.3566	0.2680	0.1427	0.2461	0.5179	0.0538	0.8340	-0.1922	0.2903	0.1578	0.8270	0.0242	0.9617	-0.1337	0.6883
---	---	1627482_at	0.1829	0.4976	0.0401	0.7304	0.1595	0.3140	0.0937	0.8164	0.0323	0.8789	-0.0614	0.7165	0.0128	0.9928	-0.0726	0.8787	-0.0854	0.8404
mod(mdg4)	Modifier67.2	1627483_at	-0.2040	0.3928	0.0405	0.9266	-0.0161	0.9345	-0.1842	0.7243	-0.3685	0.1158	-0.1843	0.3997	-0.0748	0.9717	-0.0626	0.9407	0.0122	0.9882
CG13694	CG13694	1627484_at	-0.0544	0.7978	0.1286	0.4139	0.0803	0.7269	0.1447	0.6933	0.0544	0.7932	-0.0903	0.5945	-0.0850	0.9340	-0.0059	0.9935	0.0791	0.8524
---	---	1627485_at	0.0469	0.7828	0.3686	0.1226	0.1323	0.4643	-0.0978	0.7982	-0.2001	0.1947	-0.1023	0.4910	0.2191	0.7893	0.1551	0.6928	-0.0640	0.8934
---	---	1627486_at	-0.0055	0.9769	0.0446	0.6728	0.0692	0.8071	0.0326	0.9745	-0.0146	0.9701	-0.0472	0.8729	-0.0040	0.9964	0.1024	0.6584	0.1063	0.6392
CG34379	CG13942	1627487_at	0.2208	0.2356	0.2381	0.2752	0.4764	0.0182	0.1132	0.8885	-0.0730	0.8331	-0.1862	0.4701	-0.1052	0.9294	0.0045	0.9960	0.1097	0.8178
twf	twinfillin	1627488_at	-0.7421	0.0113	-0.0112	0.9545	-0.2841	0.1369	-0.1276	0.7380	-0.2337	0.1637	-0.1061	0.5165	0.1554	0.8940	0.3457	0.4234	0.1903	0.6918
CG10433	CG10433	1627489_a_at	-0.8394	0.0033	-0.1300	0.4189	-0.7017	0.0177	-0.6192	0.1663	-1.0626	0.0029	-0.4434	0.0706	-0.0557	0.9589	-0.3839	0.1957	-0.3282	0.2964
CG30027	CG30027	1627490_at	0.1644	0.3105	0.1225	0.2496	0.0944	0.6760	-0.1043	0.8892	-0.2268	0.3764	-0.1225	0.6330	-0.0610	0.8963	-0.0738	0.7086	-0.0128	0.9596
G-ialpha65A	G protein alpha o	1627491_at	0.0138	0.9535	-0.7860	0.0915	-0.8684	0.1222	0.0805	0.9314	1.1074	0.0027	1.0269	0.0022	0.0819	0.9831	0.2842	0.8061	0.2023	0.8654
CG5018	CG5018	1627492_at	0.5158	0.2649	-0.5040	0.4402	-0.2628	0.3916	0.2013	0.8837	1.2756	0.0145	1.0743	0.0187	-0.0875	0.9742	0.2314	0.7819	0.3188	0.6644
CG5958	CG5958	1627493_at	1.9155	0.0011	1.2380	0.0152	1.9591	0.0001	0.3476	0.5680	0.2461	0.4489	-0.1015	0.7686	-0.3508	0.6898	-0.5264	0.1400	-0.1756	0.6534
oaf	transcript-near-de	1627494_s_at	0.0831	0.7974	0.3513	0.0700	0.0241	0.8934	-0.2498	0.6086	-0.4376	0.0765	-0.1878	0.4126	0.1332	0.8802	-0.1685	0.6458	-0.3018	0.3825
CG18234	CG18234	1627495_at	-6.1745	0.0005	-4.5258	0.0020	-5.2171	0.0000	-0.6431	0.1883	-1.4459	0.0011	-0.8028	0.0088	-0.1267	0.9557	-0.0562	0.9590	0.0704	0.9396
Myo95E	Myosin 95E	1627496_at	0.1053	0.6187	0.0343	0.8278	0.3258	0.0537	0.1130	0.8121	-0.1282	0.5214	-0.2411	0.1501	-0.1581	0.8284	-0.1828	0.5711	-0.0247	0.9548
Txl	Thioredoxin-like	1627497_at	-0.4049	0.0612	0.1085	0.6211	0.0993	0.5266	0.0996	0.7762	-0.0765	0.6422	-0.1761	0.1782	0.0120	0.9939	0.3082	0.3698	0.2962	0.4055
Synd	Syndapin	1627498_at	-0.3127	0.2944	0.4843	0.1088	0.3245	0.0860	-0.1902	0.6291	-0.4354	0.0318	-0.2453	0.1552	-0.0735	0.9514	0.4065	0.2430	0.4800	0.2056
CG2016	CG2016	1627499_at	-0.7963	0.1414	-2.7945	0.0028	-1.7285	0.0026	0.9256	0.2487	1.3553	0.0117	0.4297	0.3216	0.0992	0.9498	-0.4014	0.3820	-0.5006	0.2977
CG30345	CG30345	1627500_at	0.3403	0.2908	1.5696	0.0406	1.9972	0.0002	-0.4009	0.4926	-1.6527	0.0008	-1.2517	0.0015	-0.8809	0.4809	-0.4457	0.4802	0.4351	0.4978
B4	B4	1627501_s_at	0.1330	0.5623	-0.0818	0.7043	-0.1778	0.4997	0.0393	0.9627	0.5852	0.0172	0.5459	0.0144	0.1791	0.8971	0.3361	0.5289	0.1570	0.8024
CG9304	CG9304	1627502_at	1.0381	0.0245	0.8393	0.0162	0.9048	0.0079	-0.2184	0.7380	0.2467	0.4059	0.4650	0.0735	-0.1224	0.9333	0.1239	0.8455	0.2464	0.6260
CG31878	CG31878	1627503_at	0.0345	0.8453	-0.0709	0.6467	0.0671	0.7431	0.1303	0.7777	0.0844	0.7039	-0.0459	0.8351	-0.1275	0.8243	-0.0066	0.9903	0.1209	0.6389
CG32440	CG32440	1627504_at	0.1013	0.6258	0.1790	0.3650	0.1490	0.4910	-0.0422	0.9518	-0.1387	0.4953	-0.0965	0.6246	-0.0289	0.9816	0.0981	0.7594	0.1270	0.6619
CG12159	CG12159	1627505_at	0.4282	0.0508	1.0380	0.0055	1.1935	0.0005	-0.1996	0.6038	-1.2204	0.0004	-1.0208	0.0005	-0.2382	0.7485	-0.4434	0.1498	-0.2052	0.5367

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
ed	echinoid	1627506_at	-1.0916	0.0037	-0.9715	0.0083	-1.0571	0.0096	-0.1374	0.7313	0.1162	0.5460	0.2536	0.1114	-0.1437	0.9246	0.0494	0.9519	0.1931	0.7425
CG5704	CG5704	1627507_at	-0.1701	0.4359	-1.0588	0.0111	-0.9119	0.0035	0.2410	0.4310	0.6715	0.0027	0.4305	0.0116	0.0730	0.9689	-0.2058	0.7105	-0.2788	0.5964
CG31887	CG31887	1627508_at	-0.0342	0.8698	-0.1857	0.3398	-0.0969	0.6572	0.0642	0.9065	0.1326	0.4707	0.0685	0.7136	-0.1010	0.9135	0.0124	0.9841	0.1134	0.7640
hoe1	hoepel	1627509_s_at	-0.2858	0.1787	-0.0975	0.7505	-0.0546	0.7453	0.0637	0.8987	0.0187	0.9359	-0.0450	0.8066	0.0421	0.9742	0.2200	0.5071	0.1780	0.6060
CG12842	CG12842	1627510_at	0.0460	0.8558	0.0197	0.8802	-0.1018	0.5697	0.1321	0.7191	0.1453	0.3951	0.0132	0.9503	0.1869	0.7769	0.0492	0.9111	-0.1377	0.6567
chrb	charybdis	1627511_at	-0.3164	0.4728	-0.1104	0.2786	-1.2005	0.0085	-0.4093	0.3793	0.3225	0.2197	0.7318	0.0080	0.2857	0.8510	0.3665	0.5825	0.0808	0.9259
CG31915	CG31915	1627512_at	-0.3110	0.3808	-0.2100	0.7791	0.4207	0.0752	-0.0390	0.9672	-0.0221	0.9528	0.0170	0.9573	-0.6164	0.7324	0.2384	0.8123	0.8548	0.2842
---	---	1627513_at	-0.1156	0.5538	-0.4188	0.1893	-0.2561	0.3241	0.2849	0.6122	0.5607	0.0515	0.2757	0.2789	-0.1009	0.8461	0.0087	0.9829	0.1097	0.6366
CG14294 /// DereCG14294	CG14294	1627514_at	0.1859	0.3974	0.0874	0.7894	0.2869	0.1498	0.1953	0.7118	0.1930	0.4378	-0.0024	0.9937	0.0045	0.9976	0.0595	0.9085	0.0550	0.9067
CG10641	CG10641	1627515_at	-0.7333	0.0138	0.0816	0.8433	-0.1714	0.6651	-0.1480	0.8069	-0.9048	0.0034	-0.7568	0.0045	-0.0262	0.9929	-0.1072	0.9225	-0.0811	0.9347
CG6834	CG6834	1627516_at	1.9790	0.0104	1.9089	0.0061	2.4816	0.0066	-0.3397	0.8732	-0.9932	0.1771	-0.6535	0.3340	-0.8186	0.7215	-1.2101	0.1962	-0.3915	0.7222
---	---	1627517_at	-0.0062	0.9752	0.0450	0.7806	0.1195	0.4650	0.0259	0.9790	0.0122	0.9743	-0.0137	0.9648	-0.1011	0.9296	-0.1399	0.7490	-0.0388	0.9406
CG10219 /// DyakCG10219	CG10219	1627518_at	-0.6720	0.1241	-0.7014	0.0302	-0.6844	0.0055	-0.0916	0.8658	-0.3059	0.1089	-0.2143	0.2104	-0.0608	0.9816	-0.3531	0.5860	-0.2923	0.6602
---	---	1627519_at	0.1766	0.4438	0.2994	0.1528	0.1201	0.4716	-0.0365	0.9532	-0.0475	0.8247	-0.0111	0.9592	0.1931	0.7215	0.1065	0.6757	-0.0866	0.7434
---	---	1627520_at	0.0595	0.8231	-0.1491	0.6121	0.2578	0.1641	0.2990	0.5680	0.2200	0.4286	-0.0790	0.7929	-0.0958	0.9361	0.0483	0.9363	0.1441	0.7417
Ero1L	Ero1L	1627525_a_at	0.6608	0.0098	0.7117	0.0263	1.2384	0.0004	0.2207	0.6823	0.3898	0.1193	0.1690	0.4806	-0.1467	0.7644	0.5941	0.0276	0.7408	0.0238
CG8778 /// DmirCG8778	CG8778	1627526_at	-0.2553	0.2598	0.2602	0.1317	0.2894	0.1619	-0.3641	0.4440	-0.8333	0.0069	-0.4692	0.0480	-0.3592	0.6272	-0.3190	0.2877	0.0402	0.9246
CG9992	CG9992	1627527_s_at	0.5663	0.0277	0.8934	0.1152	0.4837	0.0218	-0.2437	0.5735	-0.1358	0.5709	0.1079	0.6347	0.2802	0.8097	0.2160	0.6909	-0.0642	0.9231
CG8257	CG8257	1627528_at	0.0979	0.5460	0.1186	0.5608	0.2069	0.2250	-0.0536	0.9345	-0.2231	0.2463	-0.1695	0.3344	-0.2850	0.7230	-0.0092	0.9906	0.2758	0.4353
CG32726	CG32726	1627529_at	0.1072	0.5115	0.0470	0.6760	-0.1226	0.4714	0.1161	0.7747	0.0925	0.6243	-0.0237	0.9084	-0.0511	0.9246	-0.0544	0.8174	-0.0033	0.9921
CG6700	CG6700	1627530_at	0.1701	0.4345	0.2197	0.6020	0.2070	0.4814	-0.1302	0.8545	-0.2618	0.3118	-0.1316	0.6117	-0.2047	0.8666	-0.2869	0.5756	-0.0823	0.9025
---	---	1627531_s_at	0.0977	0.5706	0.0446	0.6598	-0.1725	0.3179	0.0557	0.9262	0.2216	0.2189	0.1659	0.3129	0.1712	0.7644	0.0768	0.8061	-0.0944	0.7369
Rpb4	Rpb4	1627532_at	0.1840	0.3929	0.0152	0.9702	-0.0272	0.8710	-0.2224	0.5507	-0.0066	0.9816	0.2157	0.2105	-0.0627	0.9515	0.0535	0.9101	0.1162	0.7442
stj	straightjacket	1627533_at	-0.9137	0.0172	-3.5359	0.0014	-2.8773	0.0046	0.6153	0.7683	2.3409	0.0164	1.7256	0.0364	-0.1487	0.8400	-0.2477	0.4101	-0.0990	0.7836
---	---	1627534_at	-0.0573	0.7830	-0.0143	0.8893	0.1754	0.3766	0.1153	0.7915	-0.0987	0.6146	-0.2140	0.1770	-0.4376	0.5228	-0.2389	0.4729	0.1986	0.5676
CG34016	CG34016	1627535_at	-0.1840	0.3271	-0.0718	0.5977	0.0314	0.8429	0.0264	0.9649	0.0497	0.7998	0.0233	0.9034	0.0008	0.9998	0.0862	0.7439	0.0854	0.7418
LpR1	yolkless-like	1627536_at	0.0070	0.9764	-0.0643	0.6579	-0.1305	0.4736	0.0516	0.9540	0.1959	0.4403	0.1444	0.5485	-0.0496	0.9438	0.0212	0.9506	0.0708	0.7785
CG3085 /// DereCG3085	CG3085	1627537_at	-0.0106	0.9672	-0.0830	0.6431	0.0594	0.7561	0.0187	0.9819	-0.0593	0.8060	-0.0779	0.7072	-0.1116	0.9025	-0.0081	0.9917	0.1034	0.7964
CG6770 /// DyakCG6770	CG6770	1627538_at	-0.2094	0.2045	0.0948	0.5123	-0.3292	0.2101	-0.5784	0.0584	-1.1309	0.0004	-0.5526	0.0045	-0.1797	0.8446	-0.7548	0.0654	-0.5751	0.1524
NiPp1	Nuclear inhibitor o	1627539_at	-0.3009	0.1367	0.3341	0.1078	0.2735	0.1255	0.0061	0.9951	-0.4112	0.0458	-0.4173	0.0277	0.0241	0.9816	0.2256	0.3315	0.2016	0.4061
---	---	1627540_at	0.0906	0.6733	0.3126	0.1063	0.0780	0.6783	-0.0286	0.9629	-0.0098	0.9694	0.0189	0.9252	0.2494	0.7070	0.3000	0.2599	0.0505	0.8941
CG32855	CG32855	1627541_at	-0.1307	0.4806	-0.0977	0.4457	0.0450	0.8053	0.1002	0.7688	0.1424	0.3319	0.0422	0.7949	-0.1132	0.8611	0.0931	0.7707	0.2063	0.4457
CG17012	CG17012	1627542_at	2.1910	0.0233	0.9562	0.0065	4.7931	0.0000	3.3141	0.0241	1.3637	0.0869	-1.9504	0.0139	0.0338	0.9677	0.0462	0.8850	0.0124	0.9709
CG34370	CG13499	1627543_at	0.2406	0.2164	0.1649	0.4417	-0.0504	0.8218	-0.2750	0.5910	-0.2552	0.3323	0.0198	0.9519	-0.0497	0.9717	-0.3197	0.3571	-0.2700	0.4615
CG14797	CG14797	1627544_at	-0.0199	0.9168	0.2914	0.0606	0.1116	0.6064	0.0539	0.9553	-0.0207	0.9569	-0.0746	0.7981	0.1272	0.8692	0.0165	0.9758	-0.1108	0.7562
CG13319	CG13319	1627545_at	0.7463	0.0071	0.8294	0.0236	0.9843	0.0004	0.1206	0.7539	-0.2600	0.1164	-0.3806	0.0186	-0.0987	0.9246	-0.1215	0.7771	-0.0228	0.9651
CTCF	CTCF	1627546_at	-0.0015	0.9972	0.3511	0.0470	0.6473	0.0078	0.1150	0.7656	0.1154	0.5062	0.0005	0.9982	-0.1788	0.8744	0.4476	0.3006	0.6264	0.1839
CG31746	CG31746	1627547_at	-0.0082	0.9782	0.0311	0.7861	0.0030	0.9896	-0.2072	0.5799	-0.1250	0.5438	0.0823	0.6814	-0.0921	0.8968	0.0042	0.9941	0.0963	0.7520
CG12541	CG12541	1627548_at	-0.9604	0.0380	-0.9578	0.0598	-0.8161	0.0055	0.1637	0.8632	-0.0352	0.9434	-0.1989	0.5488	0.0840	0.9056	0.1365	0.6155	0.0525	0.8788
CG34114	CG34114	1627549_at	-0.0403	0.8589	0.1393	0.3242	0.0029	0.9922	-0.1586	0.6999	-0.1884	0.3257	-0.0298	0.8955	0.0551	0.9421	0.0800	0.7820	0.0249	0.9394
l(1)G0168	lethal (1) G0168	1627550_a_at	0.0595	0.7604	0.0123	0.9447	0.2821	0.2283	0.1171	0.8544	0.0500	0.8701	-0.0672	0.7949	-0.0610	0.9342	0.0352	0.9259	0.0962	0.7224
AttA /// AttB	attacin	1627551_s_at	1.0283	0.6878	-2.3334	0.2421	-1.9197	0.0367	-0.1757	0.9733	5.6920	0.0018	5.8677	0.0009	-0.5843	0.9619	2.4733	0.4785	3.0577	0.3819
---	---	1627552_at	0.2956	0.5440	-0.3620	0.0613	-0.0704	0.6973	0.5284	0.0657	0.6831	0.0023	0.1547	0.3030	0.0046	0.9990	-0.2807	0.6673	-0.2853	0.6567
fan	farinelli	1627553_at	0.0780	0.7438	0.0384	0.8549	-0.0790	0.6783	-0.0572	0.8968	0.1111	0.4644	0.1682	0.1949	0.2028	0.7633	0.2958	0.2760	0.0929	0.7791
CG15057	CG15057	1627554_at	0.1204	0.4913	0.4132	0.0444	0.3168	0.1241	-0.0999	0.7833	-0.2074	0.1644	-0.1075	0.4474	0.1600	0.7893	0.1509	0.5770	-0.0091	0.9835
---	---	1627555_s_at	0.3278	0.4464	-0.0986	0.7919	-0.1706	0.5089	-0.2753	0.7539	-0.0340	0.9520	0.2413	0.5103	-0.2429	0.8089	-0.5822	0.1512	-0.3393	0.4275
CG15639	CG15639	1627556_at	-0.1132	0.6097	0.1491	0.2813	0.2728	0.1189	-0.0366	0.9672	-0.2974	0.2011	-0.2608	0.2118	-0.2031	0.7707	-0.0224	0.9634	0.1807	0.5653
CG9632	CG9632	1627557_a_at	0.0877	0.6403	0.0554	0.7397	0.4302	0.0298	0.0476	0.9314	0.0841	0.6474	0.0365	0.8475	-0.2196	0.7628	-0.0830	0.8383	0.1365	0.6841
Sas-4	lethal (3) s2214	1627558_at	-0.1637	0.5831	-0.5453	0.2938	-0.2661	0.2272	-0.5238	0.3604	0.1171	0.7646	0.6409	0.0351	-0.7449	0.3162	-0.1983	0.6382	0.5466	0.1849
---	---	1627559_at	0.0514	0.8382	0.0114	0.9207	0.1660	0.3221	0.0867	0.9088	0.0242	0.9460	-0.0626	0.8192	0.0502	0.9445	0.0058	0.9905	-0.0444	0.8810
CG31308	CG31308	1627560_at	0.1923	0.3857	0.0670	0.5528	-0.0775	0.7111	0.0625	0.9276	0.0768	0.7530	0.0142	0.9568	0.0728	0.9516	-0.0305	0.9590	-0.1033	0.8189
---	---	1627561_at	-0.0279	0.9163	0.1847	0.3093	0.0915	0.6134	-0.0291	0.9693	-0.0634	0.7908	-0.0342	0.8820	0.0316	0.9683	0.1540	0.4694	0.1224	0.5834
Irp-1A	cytosolic aconitas	1627562_at	0.0157	0.9422	0.1860	0.2614	0.4913	0.0529	0.1077	0.7680	-0.0052	0.9822	-0.1129	0.4392	-0.0999	0.9296	0.2456	0.5146	0.3455	0.3568
CG7320	CG7320	1627563_at	0.1098	0.4386	0.0839	0.6411	0.2432	0.1013	0.0723	0.8658	0.1445	0.3475	0.0722	0.6418	-0.0542	0.9168	-0.0546	0.8145	-0.0004	0.9992
Smg5	Smg5	1627564_s_at	0.9708	0.0199	0.0732	0.7191	-0.2994	0.4491	-0.0186	0.9838	0.7642	0.0041	0.7828	0.0022	0.3255	0.8424	-0.2441	0.7618	-0.5696	0.4037
CG14435 /// DsimCG14435	CG14435	1627565_at	-0.4877	0.2171	1.0055	0.1045	0.8203	0.0061	-0.1871	0.7906	-1.1204	0.0029	-0.9333	0.0039	0.1144	0.9588	0.4495	0.4699	0.3351	0.6128
CG12163 /// CG33293 /// D	CG12163 /// CG3	1627566_a_at	0.2261	0.2219	-0.1626	0.1853	-0.1196	0.4916	-0.0438	0.9441	0.0844	0.6792	0.1281	0.4477	-0.1387	0.7726	-0.2993	0.1395	-0.1606	0.4465
Myb	Myb oncogene-lik	1627567_at	-0.3396	0.0771	0.4770	0.0846	0.4340	0.0223	-0.0518	0.9263	-0.6817	0.0025	-0.6298	0.0021	-0.0343	0.9777	0.1160	0.7341	0.1502	0.6329
CG14109	CG14109	1627568_at	-2.5923	0.0050	-2.9281	0.0116	-3.4287	0.0000	-0.2266	0.4898	-0.1877	0.2865	0.0389	0.8478	0.2854	0.9125	-0.5232	0.5896	-0.8086	0.3865
CG33541	CG33541	1627569_at	-0.0212	0.9056	-0.0237	0.8931	-0.0621	0.7885	0.0337	0.9603	0.0223	0.9298	-0.0114	0.9597	0.0649	0.9516	0.0229	0.9652	-0.0419	0.9246
CG8765	CG8765	1627570_a_at	-0.0362	0.8574	0.1324	0.5779	0.2172	0.1918	-0.0373	0.9649	-0.1200	0.6322	-0.0827	0.7314	-0.0052	0.9964	0.1150	0.7460	0.1202	0.7271
CrebA	Box B-binding fac	1627571_at	0.7376	0.0298	0.9741	0.1205	0.6255	0.0264	-0.0070	0.9937	0.3666	0.0464	0.3736	0.0276	0.3379	0.8270	0.5860	0.3465	0.2481	0.7353
CG31219	CG31219	1627572_at	0.0623	0.8346	0.0595	0.6248	0.0252	0.8986	0.0409	0.9633	-0.0346	0.9167	-0.0755	0.7720	-0.0416	0.9770	-0.0998	0.8231	-0.0582	0.9023
---	---	1627573_at	0.1240	0.4941	0.0249	0.9096	0.0042	0.9898	0.0088	0.9936	0.0234	0.9380	0.0146	0.9568	0.0357	0.9743	-0.0263	0.9531	-0.0620	0.8673
CG4623	CG4623	1627574_at	0.3909	0.8542	-0.4524	0.0942	-0.8917	0.0023	-0.1193	0.8611	-0.6892	0.0123	-0.5699	0.0172	0.2351	0.9816	-1.6093	0.4687	-1.8444	0.4076
---	---	1627575_at	0.3166	0.2769	0.0290	0.8496	0.1815	0.3217	0.0581	0.9339	0.0607	0.8112	0.0026	0.9918	-0.0631	0.9421	-0.1502	0.6011	-0.0871	0.7853
CG8247	CG8247	1627576_at	0.0311	0.9405	-0.0405	0.6889	0.1444	0.4663	-0.5699	0.2541	0.2808	0.3535	0.8506	0.0070	-0.5067	0.3539	0.2510	0.3812	0.7578	0.0441
CG7409	CG7409	1627577_at	-2.4779	0.0831	-1.4319	0.1628	-2.8244	0.0029	-0.5629	0.8473	-0.3288	0.8026	0.2340	0.8506	0.7821	0.8472	0.8032	0.6673	0.0210	0.9945
CG7255	CG7255	1627578_s_at	0.9508	0.0330	2.2711	0.0193	2.0103	0.0002	0.0517	0.9620	-0.3034	0.2975	-0.3551	0.1678	0.3229	0.8744	0.7566	0.3346	0.4336	0.6168

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Grd	GABA-and-glycine	1627579_at	0.0717	0.7130	0.0465	0.6888	0.1664	0.3259	0.0235	0.9749	-0.0220	0.9319	-0.0454	0.8275	0.0161	0.9869	-0.0272	0.9402	-0.0433	0.8884
Dcr-1	Dicer1	1627580_at	0.7162	0.0915	-0.0330	0.9227	0.1266	0.5384	0.3063	0.4751	0.9994	0.0018	0.6932	0.0056	0.1176	0.9514	0.3005	0.6253	0.1828	0.7883
CG8080	CG8080	1627581_at	1.0513	0.0090	0.9194	0.0696	1.2914	0.0002	-0.1023	0.8770	-0.4984	0.0351	-0.3961	0.0552	-0.4358	0.7220	-0.3723	0.4848	0.0635	0.9311
CG30035 /// DmirCG30035	CG30035	1627582_a_at	1.0695	0.0085	1.0622	0.0226	1.0789	0.0003	0.2722	0.6402	-0.3555	0.2100	-0.6277	0.0219	0.1618	0.8049	-0.3841	0.1498	-0.5459	0.0832
CG15213	CG15213	1627583_at	0.2928	0.6063	0.0628	0.6441	0.2332	0.2744	0.2862	0.7093	-0.0743	0.8725	-0.3605	0.2509	0.2850	0.8890	-0.1517	0.8948	-0.4367	0.5984
---	---	1627584_at	0.0512	0.7993	0.0000	1.0000	0.1397	0.4064	0.0146	0.9863	0.0998	0.6705	0.0852	0.6940	-0.1321	0.8283	-0.0172	0.9672	0.1149	0.6840
ro	Rough	1627585_a_at	0.0274	0.9036	0.0209	0.8528	0.2222	0.1590	0.0441	0.9486	0.1330	0.5104	0.0889	0.6528	-0.0360	0.9666	0.1170	0.6380	0.1530	0.5239
---	---	1627586_at	0.4185	0.0398	0.0338	0.8003	-0.1224	0.4559	0.1058	0.8730	0.3268	0.1525	0.2211	0.2873	0.2478	0.7644	0.0066	0.9935	-0.2412	0.5093
Epac	Epac	1627587_at	-0.1256	0.6757	0.3134	0.0459	0.1600	0.4421	-0.0106	0.9886	-0.1904	0.2745	-0.1798	0.2481	0.0608	0.9495	0.2012	0.4915	0.1404	0.6476
CG12483	CG12483	1627588_at	0.1755	0.3528	0.0864	0.4858	-0.1067	0.5279	-0.2153	0.4073	-0.0703	0.6687	0.1450	0.2658	0.1011	0.8541	-0.0151	0.9678	-0.1163	0.6328
CG9520	CG9520	1627589_s_at	0.4724	0.0497	0.6030	0.3598	0.4822	0.1562	-0.4043	0.3353	-0.3166	0.1884	0.0877	0.7363	-0.0831	0.9798	-0.1587	0.8807	-0.0756	0.9421
CG9336 /// DyakCG9336	CG9336	1627590_at	-0.8198	0.0350	0.0705	0.7055	-0.1261	0.5512	-0.1149	0.8358	-0.8023	0.0032	-0.6874	0.0038	0.0691	0.9717	0.1329	0.8393	0.0638	0.9243
---	---	1627591_at	0.0365	0.8056	0.0983	0.6336	0.3570	0.0732	0.0275	0.9637	0.0057	0.9809	-0.0217	0.9101	-0.0468	0.9589	0.2308	0.3571	0.2776	0.2923
CG3295	CG3295	1627592_at	-0.1703	0.4216	0.6630	0.0400	0.7781	0.0149	-0.1469	0.7010	-1.2827	0.0003	-1.1358	0.0002	-0.3099	0.7480	-0.4467	0.2566	-0.1368	0.7764
CG2678	transcript B	1627593_at	0.1199	0.6849	-0.7831	0.2205	-0.2763	0.4550	0.0340	0.9666	0.4648	0.0356	0.4308	0.0309	-0.5360	0.8145	-0.5509	0.5825	-0.0148	0.9926
---	---	1627594_at	0.1619	0.4914	0.1089	0.4567	0.3313	0.0590	0.0887	0.9247	0.0305	0.9398	-0.0582	0.8586	-0.1435	0.7772	0.0143	0.9688	0.1578	0.4888
CG11314	CG11314	1627595_at	2.3213	0.0008	1.4583	0.1302	2.8712	0.0000	0.7931	0.1787	0.1015	0.8201	-0.6916	0.0373	-0.3512	0.8461	-0.6013	0.4210	-0.2501	0.7774
CG4446	CG4446	1627596_at	-1.7714	0.0053	-3.6298	0.0020	-2.2923	0.0132	1.1965	0.4245	2.1334	0.0183	0.9369	0.2057	-0.0118	0.9939	-0.0063	0.9935	0.0055	0.9924
CG10694	CG10694	1627597_at	0.1505	0.3369	0.2696	0.1652	-0.0154	0.9607	-0.2474	0.4612	-0.0523	0.8171	0.1951	0.2313	-0.0090	0.9939	0.1320	0.6573	0.1411	0.6299
CG11490	CG11490	1627598_at	0.7740	0.0107	0.6788	0.0121	0.5806	0.0142	-0.0656	0.9255	0.0557	0.8382	0.1213	0.5629	0.0637	0.9590	-0.0660	0.9040	-0.1296	0.7527
Ptp4E	Protein tyrosine pl	1627599_at	-0.7196	0.0436	-0.6279	0.1580	-1.6454	0.0000	-0.5289	0.2783	0.1339	0.6829	0.6628	0.0180	0.2277	0.8023	0.0052	0.9949	-0.2225	0.5817
CG4770	CG4770	1627600_at	0.2193	0.4170	-0.1077	0.4539	0.0081	0.9666	0.0866	0.8930	0.2082	0.3402	0.1216	0.5680	-0.1774	0.8097	-0.1779	0.5836	-0.0005	0.9992
---	---	1627601_at	0.1591	0.5278	0.1828	0.2579	0.0183	0.9223	0.0001	0.9999	-0.0496	0.8112	-0.0497	0.7913	0.1136	0.9076	0.0116	0.9870	-0.1021	0.8138
tutl	turtle	1627602_at	0.2118	0.2662	0.0620	0.5190	0.1268	0.5797	0.0343	0.9664	0.1081	0.6525	0.0738	0.7504	0.1494	0.8403	0.0688	0.8754	-0.0806	0.8354
mod(mdg4)	Modifier67.2	1627603_at	-0.1020	0.6141	0.2797	0.3233	0.4644	0.0605	-0.0962	0.9042	-0.4341	0.0955	-0.3379	0.1451	-0.3387	0.6749	-0.1102	0.7863	0.2285	0.4985
CG34109	CG34109	1627604_at	0.5181	0.2645	-0.7687	0.0755	0.1652	0.3503	1.0487	0.0069	1.0206	0.0007	-0.0281	0.8991	0.1966	0.8960	-0.1542	0.8380	-0.3508	0.5554
E2f2	E2F transcription	1627605_at	0.0093	0.9699	0.1316	0.4924	-0.2719	0.1773	-0.2174	0.3985	0.0971	0.5283	0.3145	0.0231	0.0108	0.9943	0.1995	0.5855	0.1887	0.6129
---	---	1627606_at	0.2286	0.1844	-0.0672	0.5813	0.0634	0.6990	0.0283	0.9610	0.1837	0.2369	0.1554	0.2670	-0.0843	0.9238	-0.0976	0.7896	-0.0133	0.9769
CG4229	CG4229	1627607_at	0.2418	0.1819	0.0626	0.6453	0.3719	0.1345	0.0784	0.9228	0.0472	0.8857	-0.0312	0.9166	-0.0022	0.9990	-0.0010	0.9994	0.0012	0.9979
CG32048	CG32048	1627608_s_at	-0.1042	0.5430	-0.0148	0.8930	-0.1676	0.3255	-0.0127	0.9893	-0.0665	0.8008	-0.0538	0.8257	0.1014	0.8384	-0.0497	0.8619	-0.1510	0.4758
CG9742	CG9742	1627609_at	0.1558	0.5060	0.2355	0.4209	0.3278	0.2297	-0.0014	0.9988	-0.0503	0.8452	-0.0488	0.8296	-0.1165	0.9421	0.0950	0.8991	0.2114	0.7040
---	---	1627610_at	0.2613	0.2479	0.0972	0.4048	0.1756	0.3906	-0.1516	0.6292	-0.0326	0.8732	0.1189	0.4091	-0.0444	0.9742	-0.1472	0.7050	-0.1028	0.8079
CG6596	CG6596	1627611_at	-0.0003	0.9994	-0.0299	0.8051	-0.3377	0.1778	-0.1102	0.8550	-0.0565	0.8407	0.0536	0.8293	0.0030	0.9989	-0.1380	0.7188	-0.1409	0.7073
Crag	Calmodulin-bindin	1627612_a_at	0.3260	0.4918	0.0800	0.9228	-0.0141	0.9644	0.0151	0.9889	0.4284	0.0897	0.4133	0.0690	0.0333	0.9939	0.1966	0.8864	0.1634	0.9018
Mtk	metchnikowin	1627613_at	1.7247	0.4198	-1.7516	0.5664	-1.7608	0.3686	0.4722	0.8735	5.0063	0.0012	4.5342	0.0011	0.4942	0.9816	1.3511	0.8244	0.8569	0.8926
chn	charlatan	1627614_at	-0.3648	0.5341	-1.3268	0.0460	-1.9164	0.0002	-0.2479	0.5539	1.1187	0.0009	1.3666	0.0002	0.0919	0.9698	-0.1873	0.8174	-0.2792	0.6861
---	---	1627615_at	-0.2748	0.1129	-0.0049	0.9817	-0.0339	0.8780	-0.1562	0.7161	-0.0798	0.7271	0.0764	0.7129	-0.1356	0.8454	0.0796	0.8349	0.2152	0.4657
c12.1	c12.1	1627616_at	0.2612	0.2790	0.7094	0.0200	0.7272	0.0099	0.1322	0.7855	-0.4049	0.0484	-0.5371	0.0094	-0.0212	0.9898	-0.0364	0.9506	-0.0152	0.9801
Best1	Bestrophin 1	1627617_at	-1.4247	0.0014	-1.7479	0.0164	-2.1723	0.0001	-0.0701	0.9008	0.4027	0.0338	0.4728	0.0108	0.3775	0.7633	0.1660	0.8053	-0.2115	0.7228
CG15455	CG15455	1627618_at	0.1403	0.5484	0.0976	0.4248	0.1226	0.5078	0.0488	0.9154	0.0230	0.9084	-0.0258	0.8800	-0.0345	0.9816	0.1551	0.6830	0.1896	0.6049
CG15332	CG15332	1627619_at	0.1090	0.6049	-0.1140	0.5492	0.0737	0.7492	0.2059	0.5650	0.2942	0.1082	0.0883	0.6391	-0.0289	0.9816	-0.0337	0.9393	-0.0048	0.9923
CG14034	CG14034	1627620_at	-0.0049	0.9886	0.0693	0.4741	0.0528	0.8127	-0.1259	0.8180	-0.2087	0.3379	-0.0828	0.7165	0.0172	0.9898	-0.0478	0.9167	-0.0651	0.8681
---	---	1627621_at	-0.1887	0.3912	0.0145	0.9230	-0.2515	0.2925	-0.1167	0.8611	-0.1900	0.4444	-0.0733	0.7834	0.0705	0.9393	-0.0476	0.9141	-0.1180	0.7135
---	---	1627622_at	0.0301	0.8743	-0.0352	0.8335	-0.1928	0.2480	-0.0290	0.9637	0.0840	0.6585	0.1130	0.4810	0.0624	0.9467	-0.1113	0.7406	-0.1737	0.5672
bic	Enhancer of Bicau	1627623_s_at	-0.0628	0.7447	-0.4382	0.0226	-0.5207	0.0116	0.0106	0.9883	0.4111	0.0240	0.4005	0.0167	0.0153	0.9914	0.0235	0.9650	0.0083	0.9874
---	---	1627624_s_at	0.2933	0.2670	0.2689	0.1433	0.4179	0.0681	0.1067	0.8667	-0.0086	0.9801	-0.1153	0.6111	-0.1053	0.9142	-0.0882	0.8500	0.0171	0.9749
---	---	1627625_at	0.1979	0.2648	-0.1162	0.5580	0.1666	0.3994	0.0646	0.9194	0.0756	0.7436	0.0110	0.9647	-0.1928	0.8049	-0.2257	0.4983	-0.0330	0.9430
---	---	1627626_at	-0.0396	0.8337	-0.0071	0.9560	0.1626	0.2970	0.0264	0.9664	-0.0204	0.9294	-0.0468	0.7993	-0.0197	0.9852	-0.0141	0.9724	0.0056	0.9884
CG10466	CG10466	1627627_at	0.1777	0.3533	-0.0332	0.8091	0.4204	0.1121	0.0863	0.9242	0.1852	0.5219	0.0988	0.7355	-0.2621	0.7215	0.0224	0.9649	0.2846	0.3716
---	---	1627628_at	0.0610	0.7312	0.0763	0.6229	0.0769	0.7124	-0.0562	0.9451	-0.0029	0.9933	0.0533	0.8427	-0.1210	0.8999	0.1195	0.7870	0.2406	0.5154
CG3446 /// DyakCG3446	CG3446	1627629_at	-0.0382	0.8517	-0.2260	0.3736	-0.4261	0.1199	0.1117	0.8794	-0.1755	0.5092	-0.2872	0.2008	0.2504	0.7822	-0.3158	0.4177	-0.5662	0.1725
CG9650 /// DsimCG9650	CG9650	1627630_s_at	0.3026	0.2465	0.4958	0.1093	0.3268	0.3421	-0.1095	0.8707	-0.2274	0.3427	-0.1179	0.6231	0.2129	0.8846	-0.0684	0.9402	-0.2813	0.6389
His4r	Histone H4 replac	1627631_s_at	-0.1844	0.2288	0.0355	0.8437	-0.0200	0.9176	-0.0142	0.9838	-0.1898	0.2368	-0.1756	0.2213	0.0229	0.9816	-0.0207	0.9572	-0.0437	0.8919
CG17777	CG17777	1627632_at	0.3086	0.2282	0.4942	0.0260	0.3717	0.1653	-0.2176	0.6541	-0.2796	0.2316	-0.0621	0.8141	0.0423	0.9775	0.1877	0.6209	0.1454	0.7154
CG8932	CG8932	1627633_at	0.0464	0.8758	0.0584	0.6661	0.0573	0.7185	0.0814	0.9017	-0.0239	0.9380	-0.1053	0.6293	0.0438	0.9467	-0.0392	0.8962	-0.0830	0.7142
CG15098	CG15098	1627634_at	-0.0890	0.7613	0.9720	0.0101	1.1349	0.0024	-0.0513	0.9540	-1.1432	0.0013	-1.0919	0.0010	-0.1772	0.8692	-0.1632	0.7466	0.0140	0.9846
---	---	1627635_at	0.2806	0.1895	0.1412	0.4759	0.1658	0.4885	-0.0065	0.9956	0.0485	0.8940	0.0550	0.8599	-0.0204	0.9865	-0.1912	0.5071	-0.1708	0.5634
CG14636	CG14636	1627636_at	-0.5400	0.0279	-0.4789	0.0747	-0.5225	0.0173	0.5166	0.1641	0.4610	0.0461	-0.0556	0.8284	0.1701	0.7997	-0.0159	0.9734	-0.1860	0.5252
d	myosin 29D	1627637_at	-0.3270	0.4359	-0.1160	0.3078	-0.4868	0.0168	-0.0790	0.9467	-0.1713	0.6490	-0.0923	0.8056	0.1368	0.8283	-0.1285	0.6576	-0.2653	0.3271
T-cp1	Tcp1-like	1627638_s_at	-0.2911	0.3765	-0.3022	0.1786	-0.0194	0.9751	0.3985	0.5317	0.8363	0.0202	0.4379	0.1396	0.2017	0.9309	0.9121	0.2148	0.7104	0.3627
CG32802	CG32802	1627639_at	0.0274	0.8908	0.2722	0.0624	0.1066	0.5515	0.0442	0.9558	-0.0359	0.9045	-0.0801	0.7291	0.2208	0.7436	0.1441	0.6408	-0.0768	0.8306
---	---	1627640_at	-0.0724	0.7502	0.2016	0.3201	0.2350	0.2405	-0.1361	0.7539	-0.5621	0.0094	-0.4260	0.0193	-0.0253	0.9848	-0.1595	0.6216	-0.1341	0.6872
CG17494 /// DyakCG17494	CG17494	1627641_s_at	0.2367	0.3412	0.1808	0.6381	-0.1013	0.6978	-0.2410	0.6591	0.1113	0.7143	0.3523	0.1314	0.0595	0.9742	0.0803	0.9075	0.0208	0.9769
CG14974	CG14974	1627642_at	-0.0001	0.9994	0.0063	0.9562	0.0389	0.8604	-0.0626	0.9228	-0.1451	0.4757	-0.0826	0.6857	-0.1345	0.8461	0.0411	0.9275	0.1756	0.5612
CG7546 /// Mgstl	CG7546 /// Micros	1627643_at	0.5493	0.1929	1.0045	0.0106	1.2254	0.0019	0.0801	0.9422	-0.7328	0.0273	-0.8129	0.0110	-0.0551	0.9816	-0.3427	0.5374	-0.2876	0.6166
CG3588	CG3588	1627644_a_at	0.3091	0.0765	0.2998	0.0626	0.5044	0.0156	-0.0485	0.9387	-0.3025	0.1014	-0.2540	0.1225	-0.0608	0.9365	-0.0239	0.9507	0.0370	0.9113
CG9427	CG9427	1627645_at	-0.4612	0.6916	-0.4946	0.1821	-0.1859	0.3014	-0.0292	0.9602	-0.7969	0.0009	-0.7678	0.0006	-0.2632	0.9589	-1.1285	0.4391	-0.8654	0.5781
MED9	Mediator complex	1627646_at	-0.2991	0.1140	0.6196	0.0775	0.7309	0.0114	0.1723	0.8164	-0.7695	0.0165	-0.9419	0.0040	0.0007	0.9998	0.0142	0.9682	0.0135	0.9667
CG3940	CG3940	1627647_at	2.9098	0.0046	0.9231	0.2105	1.8194	0.0051	0.4375	0.7028	0.6718	0.2010	0.2343	0.6693	-0.5357	0.8517	-1.3843	0.2259	-0.8486	0.4888

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13070	CG13070	1627648_at	-0.0778	0.6699	0.0518	0.6514	0.1458	0.5264	0.0384	0.9666	-0.1749	0.4877	-0.2132	0.3319	0.1034	0.8215	0.1621	0.3856	0.0587	0.7983
Neu3	Meltrin-like	1627649_at	-1.5813	0.0025	-1.6175	0.0091	-1.1843	0.0218	0.1256	0.9249	-0.0029	0.9960	-0.1285	0.7711	-0.0852	0.9460	-0.1993	0.6280	-0.1141	0.8078
CG32431	CG32431	1627650_at	0.1832	0.2057	-0.0039	0.9781	0.0923	0.5592	0.1045	0.8601	0.1992	0.3583	0.0947	0.6681	-0.1184	0.7770	-0.0179	0.9507	0.1005	0.6058
loco	locomotion defect	1627651_a_at	-0.6930	0.0840	-0.6765	0.0421	-0.8224	0.0011	0.1655	0.6844	0.5670	0.0094	0.4015	0.0257	0.2384	0.8270	0.3838	0.3878	0.1455	0.7865
CG17761	CG17761	1627652_at	-0.0151	0.9561	0.2051	0.3277	-0.0144	0.9534	-0.1386	0.8251	-0.1561	0.5506	-0.0175	0.9549	0.2910	0.6749	0.2327	0.4073	-0.0583	0.8800
l(1)G0469	lethal (1) G0469	1627653_at	0.3305	0.4181	2.3312	0.0061	0.9096	0.0536	-0.5985	0.1956	-1.1812	0.0021	-0.5827	0.0278	0.7445	0.6955	0.7038	0.3623	-0.0408	0.9734
---	---	1627654_at	0.2863	0.0825	0.1060	0.4906	0.1125	0.5799	-0.0332	0.9603	0.0056	0.9826	0.0388	0.8492	0.0576	0.9142	-0.0776	0.7196	-0.1352	0.4869
CG30410	CG30410	1627655_at	0.9649	0.0028	1.0865	0.0727	1.9325	0.0006	0.0056	0.9956	-0.8616	0.0061	-0.8672	0.0035	-0.4016	0.7220	-0.4626	0.3265	-0.0610	0.9277
grk	gurken	1627656_at	-2.4506	0.0011	-2.3426	0.0223	-2.7378	0.0002	-0.2558	0.6615	-0.3871	0.1643	-0.1313	0.6489	0.1274	0.9670	-0.3991	0.6512	-0.5266	0.5391
CG16749	CG16749	1627657_at	1.8235	0.2156	1.0964	0.1717	1.8846	0.0215	0.6087	0.8285	-0.6282	0.5938	-1.2369	0.1997	-0.1652	0.9816	-1.2697	0.4542	-1.1046	0.5355
CG4995	CG4995	1627658_s_at	2.7349	0.0012	1.4818	0.1169	2.7443	0.0006	0.4794	0.5545	0.3932	0.3579	-0.0862	0.8599	-0.7224	0.7423	-0.8802	0.3306	-0.1578	0.9025
CG32510	CG32510	1627659_at	0.4456	0.3043	-0.7606	0.0793	-0.8100	0.0240	0.0512	0.9819	0.9958	0.0645	0.9446	0.0515	0.2613	0.7823	-0.0024	0.9989	-0.2637	0.5415
Plip	MKP-like	1627660_a_at	0.1888	0.4402	-0.1647	0.3741	-0.0062	0.9748	0.1020	0.8009	0.1567	0.3440	0.0547	0.7584	-0.1088	0.9030	-0.1413	0.7002	-0.0325	0.9433
Dph5	Diphthamide meth	1627661_at	0.1756	0.6142	0.3663	0.0713	1.0011	0.0118	0.5210	0.1609	0.1397	0.5656	-0.3813	0.0620	-0.0436	0.9848	0.4124	0.4329	0.4560	0.3953
Ugt35a	UDP-glycosyltrans	1627662_at	-0.4657	0.3198	-0.8442	0.0834	-1.2402	0.0033	-0.1815	0.8827	-0.8736	0.0435	-0.6921	0.0680	0.1767	0.9411	-1.3590	0.0795	-1.5358	0.0746
---	---	1627663_at	0.0357	0.8774	0.0466	0.7023	0.1685	0.3853	-0.0227	0.9759	-0.1818	0.3328	-0.1591	0.3487	-0.0124	0.9928	0.0479	0.9233	0.0604	0.8907
CG11178	CG11178	1627664_a_at	0.2152	0.7028	0.3269	0.6611	0.2534	0.2681	0.1421	0.7161	0.0965	0.6253	-0.0456	0.8216	0.2612	0.9296	0.2676	0.8351	0.0064	0.9974
CG17486	CG17486	1627665_at	0.1993	0.3856	0.5297	0.1486	0.8459	0.0008	-0.1476	0.7326	-0.4559	0.0257	-0.3084	0.0747	-0.2836	0.7753	-0.1610	0.7520	0.1226	0.8187
CG10979	CG10979	1627666_at	-0.5643	0.2289	0.4782	0.1893	0.4479	0.0601	0.0621	0.9562	-0.2507	0.4277	-0.3129	0.2552	0.2652	0.8571	0.7863	0.1770	0.5211	0.3941
CG16753	CG16753	1627667_at	-0.1907	0.5424	-1.1306	0.0246	-0.8679	0.0145	0.2768	0.6908	0.9675	0.0093	0.6907	0.0247	0.1397	0.8999	0.1398	0.7813	0.0001	0.9999
bbx	bobby sox	1627668_s_at	-0.1812	0.3616	0.5633	0.0207	0.6673	0.0079	-0.0368	0.9538	-0.6486	0.0036	-0.6118	0.0027	-0.0984	0.8940	0.0523	0.9003	0.1506	0.6124
CG1791	CG1791	1627669_at	0.7772	0.0602	0.3259	0.3168	0.6313	0.1646	-0.2489	0.6402	-0.5043	0.0570	-0.2554	0.2774	-0.5054	0.7387	-0.7950	0.2010	-0.2895	0.6839
CG18788	CG18788	1627670_at	-1.0025	0.0710	-1.1011	0.0077	-1.2787	0.0010	0.0877	0.9559	0.0520	0.9313	-0.0356	0.9478	-0.0063	0.9976	-0.2722	0.5848	-0.2660	0.5988
---	---	1627671_s_at	0.1038	0.4986	0.0186	0.8574	0.1079	0.5583	-0.1339	0.7732	-0.1125	0.5969	0.0214	0.9283	-0.0643	0.9405	-0.0531	0.8924	0.0113	0.9793
CG33970	CG33970	1627672_at	0.5276	0.0969	0.0742	0.7300	-0.7430	0.0437	-0.1770	0.7000	0.5801	0.0138	0.7572	0.0025	0.4982	0.6749	-0.0211	0.9814	-0.5193	0.2960
sbr	locus 1	1627673_at	0.3174	0.1127	-0.0485	0.8986	0.0196	0.9369	-0.2242	0.4420	-0.2844	0.0759	-0.0602	0.7220	-0.1384	0.8999	-0.4600	0.2395	-0.3216	0.4398
---	---	1627674_at	0.0817	0.6035	0.0003	1.0000	0.0387	0.8055	0.2295	0.4728	0.1035	0.5793	-0.1260	0.4356	0.0226	0.9851	-0.1006	0.7604	-0.1233	0.6861
---	---	1627675_at	0.1909	0.2747	0.2091	0.3118	0.0555	0.7879	-0.0978	0.8544	-0.0121	0.9678	0.0858	0.6671	0.0096	0.9940	0.1245	0.7093	0.1150	0.7346
CG1074	CG1074	1627676_at	0.6545	0.0173	0.4802	0.3756	0.6230	0.0037	0.0585	0.9255	0.3818	0.0487	0.3233	0.0596	-0.0890	0.9558	0.2789	0.5707	0.3679	0.4448
CG10743	CG10743	1627677_s_at	-0.2037	0.3971	-0.1814	0.1648	-0.3776	0.0645	0.0613	0.9110	0.1187	0.5205	0.0573	0.7637	0.2760	0.7149	0.1607	0.6384	-0.1153	0.7513
CG18302	CG18302	1627678_at	-0.0352	0.8796	-0.0366	0.8373	0.0852	0.7038	0.0842	0.9149	0.0290	0.9338	-0.0552	0.8472	-0.1668	0.8461	0.0521	0.9265	0.2189	0.5615
CG12470	CG12470	1627679_at	-1.0578	0.1118	-1.3978	0.0077	-2.1411	0.0002	-0.7721	0.3645	-0.0591	0.9307	0.7130	0.0995	-0.1655	0.9365	-0.6190	0.3342	-0.4535	0.5067
CG8777	CG8777	1627680_at	-0.2957	0.1986	-0.1564	0.7863	-0.1533	0.3659	-0.1211	0.7890	-0.0125	0.9652	0.1087	0.5509	-0.2115	0.8815	0.0640	0.9414	0.2755	0.6385
Cpr35B	CG3474	1627681_at	0.0303	0.9155	0.2558	0.2281	0.0839	0.6464	0.0262	0.9774	-0.1896	0.4166	-0.2158	0.2921	0.2113	0.7230	0.0854	0.7897	-0.1258	0.6475
CG14949	CG14949	1627682_at	2.2047	0.3856	-2.8513	0.0226	0.1213	0.5934	3.4013	0.0018	2.8676	0.0004	-0.5337	0.1655	0.2725	0.9845	-2.2506	0.4807	-2.5230	0.4318
---	---	1627683_at	0.0375	0.8844	-0.0294	0.8059	-0.0613	0.7684	-0.1727	0.6580	-0.0733	0.7399	0.0993	0.5975	-0.0277	0.9848	-0.0679	0.8852	-0.0402	0.9295
lrrk	CG5483	1627684_at	-0.0127	0.9862	0.2436	0.4637	0.4659	0.0492	-0.0946	0.9448	-0.0042	0.9942	0.0904	0.8409	-0.1955	0.9030	0.3391	0.5827	0.5346	0.3716
His3.3A	histone H3.3A	1627685_a_at	-0.5165	0.0090	0.6039	0.0207	0.7567	0.0244	0.0461	0.9687	-1.3470	0.0017	-1.3932	0.0009	-0.0850	0.8926	-0.1544	0.5259	-0.0695	0.8112
---	---	1627686_at	0.1458	0.4491	0.3273	0.1942	0.2751	0.1158	0.0295	0.9612	0.0051	0.9827	-0.0243	0.9007	0.0184	0.9848	0.0000	1.0000	-0.0184	0.9512
CG14101	CG14101	1627687_at	0.0828	0.6311	0.1492	0.2773	-0.0674	0.6646	-0.0638	0.9057	0.0733	0.7161	0.1371	0.3965	0.2339	0.6749	0.2760	0.2198	0.0421	0.8949
CG4989	CG4989	1627688_at	-0.7670	0.0622	-0.4855	0.2050	-0.5987	0.3498	-0.2415	0.8539	0.4770	0.3193	0.7185	0.0897	-0.2117	0.9238	0.5798	0.4258	0.7915	0.2927
CG17010	CG17010	1627689_at	-0.0106	0.9588	-0.0321	0.8270	-0.3035	0.1234	-0.0997	0.8107	0.0824	0.6587	0.1821	0.2180	0.1975	0.7506	0.0528	0.8929	-0.1447	0.6140
---	---	1627690_at	0.1268	0.5351	0.0881	0.6241	0.1210	0.4599	0.0150	0.9814	0.1689	0.2743	0.1539	0.2671	0.0972	0.9061	0.1632	0.6041	0.0661	0.8655
dnk	deoxyribonucleosi	1627691_at	-0.1373	0.6911	-0.2118	0.5009	-0.6280	0.0380	-0.2429	0.4012	0.3067	0.0618	0.5496	0.0032	0.0194	0.9943	0.1031	0.9170	0.0837	0.9237
---	---	1627692_at	0.0818	0.6577	-0.1075	0.4514	-0.2134	0.3397	-0.1399	0.7774	0.0623	0.8066	0.2022	0.2786	0.0068	0.9952	-0.2326	0.3856	-0.2394	0.3898
CG1965	CG1965	1627693_at	-0.2562	0.2996	-0.1121	0.8600	-0.2767	0.1273	-0.2391	0.5012	-0.1234	0.5430	0.1156	0.5300	-0.1303	0.9361	-0.0151	0.9890	0.1152	0.8655
CG9715 /// DereCG9715 ///	CG9715	1627694_at	0.2956	0.1886	-0.1098	0.7069	0.0318	0.9064	0.1242	0.7511	0.4299	0.0198	0.3057	0.0495	-0.0256	0.9898	0.0071	0.9941	0.0327	0.9584
---	---	1627695_at	0.0374	0.8993	0.3276	0.2713	0.1853	0.3228	0.0126	0.9897	-0.0039	0.9902	-0.0165	0.9535	0.1285	0.8754	0.2119	0.5246	0.0834	0.8379
---	---	1627696_at	0.2523	0.2600	-0.0422	0.7237	0.1349	0.4160	0.0332	0.9603	0.1184	0.5270	0.0853	0.6354	-0.1132	0.8740	-0.1330	0.6675	-0.0198	0.9619
rho-7	rhomboid-7	1627697_at	-0.2094	0.3929	0.3510	0.2176	0.7719	0.0063	0.0612	0.9149	-0.4103	0.0295	-0.4715	0.0102	-0.2421	0.7196	0.0994	0.7699	0.3415	0.2365
CCS	CCS	1627698_at	-0.0110	0.9629	-0.1693	0.3910	-0.3804	0.0487	-0.1685	0.7604	0.1611	0.5239	0.3297	0.1193	0.0687	0.9225	0.0415	0.9077	-0.0272	0.9342
CG14070	CG14070	1627699_at	-0.1730	0.3526	-0.0661	0.7448	0.2857	0.3072	-0.0751	0.8671	-0.2116	0.1795	-0.1365	0.3481	-0.2914	0.8016	-0.1392	0.8272	0.1522	0.7940
CG6420	CG6420	1627700_at	-0.4447	0.3547	0.0806	0.9233	0.0476	0.7884	-0.1654	0.8930	-0.6820	0.0964	-0.5166	0.1576	0.0990	0.9781	0.1119	0.9353	0.0129	0.9932
---	---	1627701_at	0.1226	0.4892	0.1247	0.4219	0.4000	0.0306	0.2471	0.5128	0.0121	0.9690	-0.2350	0.1870	0.0204	0.9862	-0.0094	0.9868	-0.0298	0.9387
---	---	1627702_at	-0.0769	0.8693	-0.1073	0.8666	-0.1541	0.4724	0.1340	0.8640	0.1057	0.7485	-0.0283	0.9344	0.2150	0.9275	0.1593	0.8867	-0.0557	0.9609
eys	eyes shut	1627703_at	-0.1734	0.3229	0.0201	0.9494	0.0182	0.9640	-0.0136	0.9917	0.0475	0.8983	0.0611	0.8457	-0.1073	0.8889	0.1240	0.7048	0.2313	0.4367
Traf1	TNF-receptor-ass	1627704_a_at	-1.9541	0.0103	-1.6562	0.0074	-2.4958	0.0001	-1.1472	0.0666	-1.3186	0.0039	-0.1714	0.6451	-0.1356	0.9474	-1.0115	0.1119	-0.8759	0.1875
---	---	1627705_at	0.0692	0.8028	-0.0100	0.9261	0.1721	0.6172	-0.0691	0.9445	0.0156	0.9718	0.0846	0.7899	-0.1427	0.8037	-0.0551	0.8712	0.0876	0.7503
CG33174 /// DmirCG33174	CG33174	1627706_a_at	-0.9370	0.0590	0.0530	0.7392	-0.1652	0.5606	-0.0515	0.9610	-0.4880	0.0856	-0.4364	0.0851	-0.1752	0.8846	0.2201	0.6614	0.3953	0.3964
CG3313	CG3313	1627707_at	-0.2260	0.3697	0.5288	0.0655	-0.0360	0.8450	-0.3141	0.3596	-0.4860	0.0211	-0.1719	0.3362	0.0343	0.9742	-0.0787	0.8168	-0.1130	0.6985
---	---	1627708_at	0.2017	0.4107	0.0539	0.6302	-0.0057	0.9865	-0.0386	0.9673	-0.0469	0.8887	-0.0083	0.9787	0.0127	0.9922	-0.0728	0.8551	-0.0856	0.8129
CG30371	CG30371	1627709_at	3.1023	0.0016	1.1792	0.0086	4.8067	0.0000	3.4911	0.0104	2.2573	0.0059	-1.2338	0.0475	0.1296	0.8004	0.3016	0.1489	0.1720	0.4322
---	---	1627710_at	0.0723	0.6558	0.0096	0.9596	-0.0029	0.9916	-0.0095	0.9934	0.0215	0.9486	0.0310	0.9084	0.0456	0.9549	-0.0173	0.9647	-0.0629	0.8323
CdsA	CDP-DAG syntha	1627711_at	-0.5249	0.0166	-0.8212	0.0498	-0.8007	0.0018	-0.0412	0.9496	0.2292	0.2052	0.2704	0.0953	-0.0751	0.9425	0.0248	0.9628	0.0999	0.7968
CG34369	CG13489	1627712_at	0.1279	0.6139	0.0891	0.4827	-0.0457	0.8311	-0.1558	0.7982	-0.0594	0.8529	0.0965	0.7129	0.0169	0.9831	0.0251	0.9315	0.0081	0.9777
CG8289	CG8289	1627713_at	0.1155	0.5240	0.1039	0.7620	0.0633	0.7045	0.1122	0.7579	0.3090	0.0532	0.1968	0.1588	0.2052	0.8270	0.3448	0.3737	0.1396	0.7593
CG3748	CG3748	1627714_at	-0.0545	0.8258	-0.0220	0.8954	0.0039	0.9881	-0.0242	0.9777	-0.0587	0.8340	-0.0345	0.8956	0.0778	0.9365	-0.0430	0.9296	-0.1208	0.7323
---	---	1627715_at	-0.0598	0.8529	0.0021	0.9940	-0.0786	0.6781	0.0856	0.9116	0.1492	0.5727	0.0636	0.8187	0.0251	0.9831	0.0817	0.8225	0.0566	0.8813
CG11598	CG11598	1627716_at	0.1843	0.2763	0.1050	0.6149	0.1099	0.5491	0.1551	0.6081	0.1266	0.4188	-0.0285	0.8738	0.1595	0.8270	0.2016	0.5246	0.0421	0.9198

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1627717_at	0.0832	0.6043	-0.0264	0.8240	0.0030	0.9922	0.0286	0.9722	0.0623	0.8048	0.0337	0.8907	0.0526	0.9391	0.0217	0.9499	-0.0309	0.9164
---	---	1627718_at	0.0031	0.9892	0.2301	0.3492	-0.0576	0.8710	0.0583	0.9299	-0.0864	0.6991	-0.1447	0.4260	0.1333	0.8608	0.0561	0.9046	-0.0772	0.8434
gol	goliath	1627719_at	1.1821	0.0088	1.3922	0.0238	1.3003	0.0011	0.1042	0.8906	-0.4164	0.1009	-0.5207	0.0294	0.2125	0.8768	-0.2753	0.6337	-0.4878	0.3743
CG14301	CG14301	1627720_at	-2.9254	0.0019	-3.5970	0.0153	-3.3811	0.0001	0.2685	0.7647	-0.4310	0.2629	-0.6994	0.0467	0.1410	0.9677	-0.5213	0.5936	-0.6623	0.4865
CG12880	CG12880	1627721_at	-2.9686	0.0022	-1.0252	0.2535	-3.0587	0.0049	-1.5807	0.0627	-2.5797	0.0008	-0.9989	0.0293	0.5146	0.9011	-0.5438	0.7672	-1.0584	0.5003
---	---	1627722_at	0.2615	0.3098	0.1778	0.2035	0.1317	0.4732	-0.0997	0.8076	-0.0977	0.5817	0.0020	0.9919	0.1470	0.8465	-0.0202	0.9690	-0.1672	0.6225
CG14701	CG14701	1627723_at	-0.0429	0.8764	0.3123	0.2748	0.3635	0.2974	0.3650	0.5242	0.3300	0.2740	-0.0350	0.9243	0.2048	0.8889	0.6036	0.2561	0.3989	0.4852
---	---	1627724_at	0.0161	0.9253	-0.0920	0.4728	0.0289	0.8656	-0.0366	0.9471	0.0399	0.8435	0.0764	0.6313	-0.1048	0.8302	-0.0660	0.7998	0.0388	0.8905
CG5213	CG5213	1627725_at	0.3213	0.1572	0.1465	0.4476	0.3199	0.1168	0.2072	0.6558	-0.0120	0.9729	-0.2192	0.2782	0.2253	0.7823	0.0366	0.9494	-0.1888	0.6196
CG6404	CG6404	1627726_a_at	0.0977	0.7305	0.2180	0.2947	0.2109	0.4549	-0.1099	0.7931	0.2480	0.1459	0.3579	0.0273	-0.2000	0.8650	0.2851	0.5618	0.4851	0.3097
---	---	1627727_at	-0.1383	0.5047	0.0622	0.5702	-0.0872	0.7592	-0.1491	0.7075	-0.2529	0.1615	-0.1038	0.5631	0.0057	0.9969	0.0159	0.9812	0.0102	0.9856
CG31099	CG31099	1627728_at	0.1676	0.3942	0.1474	0.3973	0.2778	0.1442	0.1442	0.6990	0.0723	0.7158	-0.0719	0.6895	0.1909	0.7628	0.1366	0.6314	-0.0542	0.8800
pUf68	half-pint	1627729_s_at	-0.6958	0.1142	-0.0760	0.9323	0.2413	0.1629	-0.1427	0.6597	-0.7747	0.0011	-0.6320	0.0015	-0.3692	0.8609	0.1757	0.8864	0.5448	0.5444
Rgk3	Rgk3	1627730_at	0.1028	0.5044	0.2701	0.2605	-0.0868	0.7744	0.0249	0.9716	0.0677	0.7478	0.0428	0.8326	0.4683	0.5421	0.2389	0.5199	-0.2294	0.5448
CG17438	CG17438	1627731_at	-0.0506	0.7855	0.0878	0.4646	0.1161	0.4369	-0.0103	0.9895	-0.1984	0.2747	-0.1881	0.2467	-0.0264	0.9717	0.0505	0.8394	0.0769	0.7138
CG30415	CG30415	1627732_s_at	-0.1821	0.5152	0.0595	0.8608	-0.0715	0.7325	-0.0536	0.9297	-0.4050	0.0303	-0.3514	0.0343	0.0631	0.9721	-0.3205	0.4954	-0.3836	0.4095
CG12347 /// DyakCG12347	CG12347	1627733_at	-0.0396	0.8377	-0.1722	0.6213	-0.1831	0.4881	0.1735	0.8043	-0.0048	0.9903	-0.1783	0.5090	0.1624	0.9032	-0.0600	0.9401	-0.2224	0.6773
nau	myogenic determi	1627734_at	0.1277	0.3744	0.0868	0.4867	0.2881	0.3153	-0.0843	0.8594	0.1252	0.4856	0.2095	0.1703	-0.1129	0.9011	-0.0520	0.9198	0.0609	0.8938
CG13908 /// DmirCG13908	CG13908	1627735_at	0.1726	0.4903	-0.2332	0.2343	-0.1218	0.5015	0.1540	0.7017	0.4360	0.0259	0.2820	0.0875	-0.0089	0.9939	-0.0801	0.8228	-0.0712	0.8373
activin-beta	dActivin	1627736_at	-0.0722	0.6741	0.0170	0.8886	0.1494	0.4745	0.0106	0.9922	-0.0902	0.7377	-0.1007	0.6719	-0.2220	0.7633	-0.0839	0.8393	0.1381	0.6854
CG13008	CG13008	1627737_at	0.1614	0.3367	-0.0347	0.7583	0.3123	0.2651	0.1852	0.5311	0.0271	0.8980	-0.1581	0.2549	0.0277	0.9683	-0.1144	0.5523	-0.1421	0.4558
msl-3	male specifc letha	1627738_a_at	0.2416	0.3000	0.5105	0.1302	0.8415	0.0016	0.1428	0.7556	-0.1422	0.4940	-0.2850	0.1067	-0.1467	0.8940	0.1836	0.6876	0.3303	0.4301
SC35	SC35	1627739_at	0.3028	0.1454	0.2602	0.0847	0.3273	0.0632	0.1705	0.6947	0.4525	0.0315	0.2820	0.1146	0.0589	0.9330	0.3523	0.1199	0.2935	0.2095
unc-13-4A	unc-13-4A	1627740_s_at	-0.1332	0.4605	0.0355	0.7775	0.3397	0.1670	-0.0116	0.9903	-0.0911	0.7100	-0.0796	0.7253	-0.1441	0.8009	0.0817	0.7802	0.2259	0.3564
CG13086	CG13086	1627741_at	1.8455	0.0108	1.3956	0.1015	2.1401	0.0005	0.2203	0.7760	-0.2487	0.4614	-0.4690	0.1046	-0.6130	0.8023	-0.6386	0.5505	-0.0255	0.9874
Bmcp	Bmcp	1627742_at	0.1427	0.6701	-0.5446	0.0411	-0.5223	0.0098	-0.1798	0.7205	-0.0271	0.9325	0.1527	0.4810	-0.2707	0.7561	-0.8216	0.0476	-0.5508	0.1591
CG5585	CG5585	1627743_at	0.1594	0.2907	-0.0771	0.7881	0.1919	0.3742	0.0299	0.9641	0.4151	0.0261	0.3852	0.0226	-0.3925	0.6824	0.0082	0.9928	0.4008	0.3205
CG15209	CG15209	1627744_at	-0.5290	0.4842	-0.6746	0.1977	-1.1012	0.0798	0.2423	0.8578	-0.2631	0.6253	-0.5054	0.2457	0.5898	0.8378	-0.4411	0.7544	-1.0308	0.3921
---	---	1627745_s_at	-0.6909	0.0035	-1.5919	0.1388	-1.7458	0.0006	-0.1081	0.8828	0.6191	0.0215	0.7272	0.0065	-0.0597	0.9884	-0.1750	0.8939	-0.1154	0.9245
CG6124 /// eater	CG6124 /// eater	1627746_at	-2.9253	0.0107	-4.5690	0.0149	-4.7709	0.0000	-0.6454	0.0852	0.0690	0.7963	0.7144	0.0038	-0.5328	0.8890	-1.6022	0.2489	-1.0694	0.4730
CG34017	CG34017	1627747_s_at	0.2620	0.1820	-0.0968	0.4920	0.1286	0.5202	0.1281	0.7850	0.1595	0.4259	0.0314	0.8924	0.0600	0.9225	-0.1324	0.5330	-0.1923	0.3601
CG9236	CG9236	1627748_at	-0.0449	0.8831	-0.0481	0.6506	-0.0308	0.8771	0.2303	0.4979	0.1650	0.3695	-0.0653	0.7361	0.0819	0.9011	0.1228	0.6394	0.0409	0.9025
CG6412	CG6412	1627749_at	-0.1686	0.3335	-0.0288	0.9293	-0.2201	0.1904	0.0234	0.9705	-0.0505	0.7998	-0.0739	0.6607	0.1945	0.8065	0.0184	0.9739	-0.1760	0.6222
CG15184	CG15184	1627750_at	0.3271	0.2529	0.1242	0.3636	0.3002	0.0926	0.0869	0.8485	0.0411	0.8488	-0.0458	0.8061	-0.1281	0.8744	-0.1842	0.5855	-0.0562	0.8982
CG40121	CG40121	1627751_at	0.1866	0.3766	0.0143	0.8912	-0.0752	0.7997	-0.1161	0.8449	0.1027	0.6806	0.2188	0.2674	0.0243	0.9831	-0.1023	0.7439	-0.1266	0.6607
SK	small conductanc	1627752_s_at	-2.7455	0.0012	-4.6934	0.0084	-4.4172	0.0001	0.0675	0.9803	1.4467	0.0371	1.3792	0.0287	-0.3092	0.9090	-0.4420	0.6771	-0.1328	0.9197
---	---	1627753_at	-0.0297	0.8827	-0.1206	0.5041	0.3070	0.1170	0.2163	0.5932	0.1796	0.3938	-0.0367	0.8802	-0.1657	0.8379	0.0144	0.9819	0.1801	0.6166
---	---	1627754_at	0.2342	0.1654	0.1785	0.5039	0.1220	0.7042	0.0464	0.9633	-0.0063	0.9879	-0.0526	0.8662	0.0951	0.9342	0.1469	0.7341	0.0518	0.9194
CG14185	CG14185	1627755_at	0.4136	0.0760	0.3218	0.2344	0.2590	0.2886	-0.2340	0.6854	-0.2270	0.4120	0.0070	0.9840	0.0517	0.9535	0.0255	0.9519	-0.0262	0.9433
CG6006	CG6006	1627756_at	-0.3262	0.1141	-0.0390	0.7803	-0.0349	0.8420	0.1208	0.8384	-0.4362	0.0541	-0.5570	0.0127	-0.0667	0.9174	-0.1888	0.3909	-0.1222	0.6120
Hmx	H6-like-homeobo	1627757_at	0.2200	0.3138	0.2049	0.3388	0.1568	0.4704	0.0979	0.8830	0.2251	0.3275	0.1271	0.5720	0.0685	0.9503	0.1534	0.6699	0.0849	0.8373
CG13407	CG13407	1627758_at	-0.0171	0.9264	0.1348	0.2840	0.3531	0.1380	0.0938	0.8923	-0.0453	0.8846	-0.1391	0.5393	-0.1861	0.8541	0.0833	0.8924	0.2694	0.5412
CG30080	CG30080	1627759_at	0.4113	0.1547	0.1438	0.5632	0.3135	0.0934	0.1697	0.7538	-0.3128	0.1779	-0.4825	0.0286	0.0242	0.9872	-0.3956	0.2351	-0.4197	0.2378
---	---	1627760_s_at	0.1518	0.4437	0.0067	0.9752	0.0625	0.7526	0.0856	0.9116	0.2727	0.2647	0.1872	0.4106	-0.0493	0.9588	0.0650	0.8591	0.1143	0.6980
fat2	fat-like	1627761_at	-0.3603	0.4893	0.3733	0.6096	0.1588	0.6423	-0.3942	0.5280	-0.8568	0.0169	-0.4626	0.1119	-0.1199	0.9742	-0.0833	0.9561	0.0366	0.9816
---	---	1627762_at	0.0757	0.7870	0.1256	0.4443	0.0555	0.8435	-0.1260	0.8085	-0.0605	0.8134	0.0655	0.7751	-0.1048	0.9235	-0.0215	0.9716	0.0833	0.8599
CG31048	CG31048	1627763_at	-0.8822	0.0824	0.2002	0.4577	0.2234	0.3516	-0.0958	0.9445	-0.9126	0.0277	-0.8169	0.0277	-0.0668	0.9812	0.1138	0.9011	0.1806	0.8082
CG6194	CG6194	1627764_at	-0.0603	0.8113	-0.0646	0.5615	-0.0933	0.5499	-0.0743	0.8959	-0.2114	0.2625	-0.1371	0.4374	-0.0708	0.9420	-0.2245	0.4548	-0.1537	0.6328
endoB	endophilin	1627765_a_at	-0.0839	0.6588	-0.4986	0.0833	-0.6207	0.0326	-0.2764	0.6553	0.2018	0.5243	0.4782	0.0743	-0.1539	0.8400	-0.1045	0.7916	0.0494	0.9095
CG40245	CG40245	1627766_at	-0.1746	0.2757	0.1068	0.4875	0.0825	0.5766	-0.0561	0.9040	-0.1932	0.1998	-0.1372	0.3202	0.0143	0.9875	0.1170	0.5907	0.1027	0.6414
---	---	1627767_at	0.2986	0.2060	0.2376	0.2228	0.0466	0.8659	0.0421	0.9602	0.1328	0.5816	0.0906	0.6976	0.1229	0.8689	0.0580	0.8949	-0.0650	0.8684
CG12384 /// DyakCG12384	CG12384	1627768_at	0.6546	0.0204	1.4913	0.0086	0.9707	0.0019	-0.1449	0.7680	-0.9134	0.0018	-0.7685	0.0022	0.1368	0.9029	-0.1677	0.7200	-0.3045	0.4699
---	---	1627769_at	-0.1954	0.3255	-0.2467	0.0945	-0.0489	0.8009	0.2061	0.6338	0.3305	0.1186	0.1245	0.5516	-0.0596	0.9342	0.0839	0.7640	0.1435	0.5587
Cks30A	Cyclin-dependent	1627770_at	0.3115	0.6217	-0.4665	0.5431	-0.8764	0.1861	-0.7417	0.1403	0.7832	0.0183	1.5249	0.0006	-0.6112	0.8744	-0.0215	0.9947	0.5898	0.7354
CG13075	CG13075	1627771_at	0.4386	0.4900	-0.0664	0.7074	-1.2791	0.0231	-1.2038	0.2506	0.0620	0.9468	1.2658	0.0309	0.0463	0.9913	-0.4957	0.6026	-0.5420	0.5669
---	---	1627772_at	0.1669	0.5278	-0.1907	0.1589	-0.0197	0.9142	0.1480	0.7929	0.1379	0.5812	-0.0102	0.9716	-0.0026	0.9984	-0.0881	0.7910	-0.0855	0.7865
Jheh2	Juvenile hormone	1627773_a_at	-1.9795	0.0024	-3.5055	0.0021	-2.1247	0.0001	0.3038	0.6063	-0.0638	0.8712	-0.3676	0.1642	-1.0063	0.2740	-1.5396	0.0204	-0.5333	0.2729
---	---	1627774_a_at	0.0778	0.5934	0.0477	0.6538	0.1215	0.3988	0.0330	0.9482	-0.0494	0.7729	-0.0824	0.5573	0.0367	0.9545	-0.0227	0.9404	-0.0594	0.7936
ect	Interferon-like pro	1627775_s_at	1.4293	0.2476	0.3411	0.3635	0.5218	0.0551	0.0306	0.9598	-0.2698	0.0911	-0.3003	0.0412	0.0105	0.9990	-1.3248	0.3002	-1.3352	0.3205
polybromo	polybromo	1627776_at	0.1543	0.6735	0.4068	0.1821	0.5599	0.0765	0.1963	0.8518	0.0527	0.9219	-0.1437	0.7248	0.0812	0.9588	0.4003	0.3456	0.3191	0.4793
---	---	1627777_at	0.0829	0.7799	-0.3414	0.1931	-0.0350	0.9030	-0.1459	0.8717	0.4708	0.1337	0.6168	0.0356	-0.4331	0.6557	-0.1254	0.8022	0.3077	0.4457
Art3	Arginine methyltra	1627778_a_at	0.3258	0.3440	0.2818	0.6556	0.6616	0.0099	0.2333	0.7023	0.5291	0.0629	0.2957	0.2396	-0.1455	0.9418	0.4558	0.4619	0.6013	0.3406
CG13295	CG13295	1627779_at	-0.3712	0.2645	-0.2150	0.1201	-0.7233	0.0084	-0.4829	0.1749	-0.6514	0.0091	-0.1685	0.4014	-0.0320	0.9884	-0.5238	0.2509	-0.4918	0.3089
---	---	1627780_at	0.0790	0.7283	0.1145	0.4419	0.1186	0.4994	-0.1471	0.7031	-0.1183	0.5335	0.0288	0.8924	-0.0385	0.9640	0.1110	0.6614	0.1495	0.5391
Ucp4B	Ucp4B	1627781_at	-0.0538	0.8667	0.0589	0.5349	-0.0177	0.9348	-0.0228	0.9807	-0.1903	0.4229	-0.1675	0.4377	0.0740	0.9340	-0.0096	0.9868	-0.0837	0.8138
CG13751	CG13751	1627782_at	-0.4632	0.0197	-0.2271	0.3761	-0.3822	0.1183	0.0210	0.9773	0.1781	0.3257	0.1572	0.3362	0.2103	0.8203	0.3996	0.2839	0.1892	0.6476
CG18599	CG18599	1627783_at	0.1624	0.5596	0.2318	0.1625	0.6967	0.0040	-0.0408	0.9529	-0.1159	0.5732	-0.0751	0.7101	-0.1599	0.7768	0.1548	0.5418	0.3146	0.2136
---	---	1627784_at	-0.8792	0.1673	-0.5868	0.5413	-0.8620	0.0083	-0.4620	0.5370	-0.0183	0.9760	0.4437	0.2037	-0.1082	0.9831	0.1933	0.9169	0.3015	0.8435
Tsp42En	tetraspanin 42E	1627785_at	-1.6871	0.0138	-2.5667	0.0064	-2.8888	0.0000	0.0630	0.9592	-0.1638	0.6514	-0.2268	0.4568	0.4016	0.8270	-0.9443	0.1986	-1.3459	0.1121

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1627786_s_at	0.0399	0.8892	0.1623	0.6802	-0.0530	0.8592	-0.2432	0.7117	-0.5113	0.0878	-0.2681	0.3253	0.0348	0.9875	-0.3670	0.4729	-0.4018	0.4370
---	---	1627787_at	-0.0500	0.8526	-0.0992	0.4658	-0.0546	0.8206	0.0020	0.9985	0.0664	0.8128	0.0643	0.8007	0.0102	0.9944	0.0138	0.9829	0.0036	0.9953
CG14722	CG14722	1627788_at	-0.2238	0.2123	0.6085	0.1287	0.6843	0.0048	0.2038	0.5552	-0.2807	0.1145	-0.4844	0.0088	0.0978	0.9457	0.4154	0.3259	0.3176	0.4793
CG11444	CG11444	1627789_at	0.0069	0.9892	1.1390	0.0247	1.0816	0.0073	0.3835	0.5471	-0.2658	0.4371	-0.6493	0.0348	0.4118	0.6749	0.5706	0.1512	0.1589	0.7355
CG31461	CG31461	1627790_at	0.2040	0.3799	0.0517	0.7339	-0.0834	0.6181	0.1249	0.8189	0.0856	0.7322	-0.0393	0.8767	0.1474	0.8222	-0.1544	0.5957	-0.3018	0.2841
CG34351	CG11928	1627791_at	0.1335	0.6406	0.0492	0.6748	0.3309	0.0455	0.1159	0.8897	0.0415	0.9154	-0.0744	0.8135	-0.0081	0.9950	0.0279	0.9531	0.0359	0.9316
---	---	1627792_at	0.0355	0.8846	0.1025	0.4735	0.0346	0.8823	-0.0496	0.9436	-0.0100	0.9743	0.0396	0.8651	-0.0636	0.9503	-0.0313	0.9496	0.0323	0.9394
CG9134	CG9134	1627793_s_at	-3.5555	0.0011	-2.3976	0.0500	-3.0714	0.0000	-0.4685	0.4840	-1.5355	0.0019	-1.0670	0.0057	0.1732	0.9598	-0.4311	0.6952	-0.6043	0.5610
Trax	Translin associate	1627794_a_at	-0.4353	0.0527	0.1689	0.5053	0.1125	0.6563	-0.1280	0.8817	-0.2566	0.3946	-0.1286	0.6752	-0.0380	0.9774	0.3359	0.2713	0.3739	0.2536
Ir	Drosophila inward	1627795_at	-1.8512	0.0157	-1.1708	0.0367	-1.9238	0.0001	-0.2243	0.8045	-0.3450	0.3486	-0.1208	0.7605	0.6731	0.6898	0.3900	0.5943	-0.2831	0.7154
gpp	grappa	1627796_s_at	0.8220	0.1851	0.1841	0.6399	-1.3726	0.0266	-0.6609	0.3723	1.4003	0.0057	2.0611	0.0006	0.7628	0.7697	0.5944	0.6156	-0.1684	0.9121
CG10019	CG10019	1627797_at	-0.1012	0.7150	0.0548	0.8098	0.0096	0.9643	-0.0052	0.9956	-0.0172	0.9550	-0.0120	0.9629	-0.1437	0.9013	-0.0320	0.9634	0.1117	0.8357
CG6971	CG6971	1627798_at	0.3049	0.1215	0.0624	0.5182	0.4415	0.0716	0.2379	0.6010	0.1705	0.4762	-0.0674	0.7929	-0.1980	0.7686	-0.1124	0.7361	0.0856	0.8069
CG12161	CG12161	1627799_at	0.3034	0.2478	0.1164	0.2893	-0.0087	0.9737	0.0368	0.9465	0.1560	0.3195	0.1191	0.4099	0.1282	0.9087	-0.1115	0.8319	-0.2397	0.5670
---	---	1627800_at	0.1156	0.4934	0.0967	0.4966	0.1133	0.4859	-0.0126	0.9838	0.0435	0.8094	0.0562	0.7175	0.0799	0.9081	0.1377	0.5994	0.0578	0.8572
CG9497	CG9497	1627801_at	0.2096	0.4279	0.5310	0.1502	1.1077	0.0020	-0.0139	0.9937	-0.5260	0.1182	-0.5121	0.0903	-0.3344	0.7644	0.0063	0.9950	0.3407	0.4852
Sur-8	Sur-8	1627802_s_at	-0.2592	0.6027	-0.5563	0.0854	-1.0780	0.0016	-0.1646	0.8279	0.4959	0.0889	0.6605	0.0192	0.2225	0.9024	0.1737	0.8472	-0.0489	0.9596
CG11875	CG11875	1627803_at	0.2995	0.2545	0.2359	0.1972	0.3299	0.1242	-0.2050	0.5317	-0.0258	0.9142	0.1792	0.2436	-0.3199	0.7070	0.0247	0.9664	0.3445	0.3485
---	---	1627804_at	0.2484	0.1888	-0.0224	0.8372	-0.0871	0.5692	0.1760	0.6379	0.1963	0.2882	0.0202	0.9281	0.0033	0.9976	-0.1730	0.4976	-0.1763	0.4925
CG40225	CG40225	1627805_at	0.0359	0.8657	0.0920	0.3957	-0.0216	0.9404	-0.0564	0.9225	-0.0872	0.6587	-0.0308	0.8815	0.0609	0.9589	-0.0049	0.9942	-0.0659	0.8849
CG11133	CG11133	1627806_at	0.0438	0.8495	-0.0747	0.8730	-0.3290	0.2609	-0.3195	0.5068	0.3794	0.1356	0.6989	0.0085	-0.0510	0.9831	0.2719	0.6576	0.3229	0.5884
w-cup	world cup	1627807_at	0.3266	0.0527	0.0242	0.8251	0.0524	0.8179	0.0806	0.8470	0.1293	0.4129	0.0487	0.7737	-0.0143	0.9898	-0.2379	0.3040	-0.2236	0.3616
CG13011 /// DyakCG13011	CG13011	1627808_at	0.2220	0.4138	-0.7709	0.0491	0.2918	0.4544	0.0944	0.9266	0.5916	0.0585	0.4972	0.0725	-1.0301	0.2488	-0.6285	0.1489	0.4016	0.3798
CG17047	CG17047	1627809_at	0.0292	0.8803	-0.0403	0.8106	0.0552	0.8170	0.0805	0.9037	0.2104	0.3385	0.1299	0.5389	-0.0131	0.9901	0.0946	0.7200	0.1077	0.6670
---	---	1627810_at	0.1391	0.4344	-0.1125	0.4966	0.2289	0.3395	0.2538	0.4800	0.2450	0.2017	-0.0088	0.9710	-0.1408	0.8331	-0.0121	0.9817	0.1288	0.6729
---	---	1627811_at	0.1227	0.5634	-0.2868	0.1301	-0.1647	0.3581	0.0699	0.9101	0.3067	0.1170	0.2368	0.1769	-0.1131	0.8609	-0.1007	0.7424	0.0124	0.9762
Rbp1	RNA-binding prote	1627812_s_at	0.1873	0.3412	0.2244	0.4717	0.1269	0.6148	0.0404	0.9564	0.2963	0.1356	0.2559	0.1497	0.1174	0.9174	0.3410	0.3743	0.2236	0.5945
wa-cup	walker cup	1627813_at	0.1215	0.4947	0.0024	0.9902	0.1876	0.3407	0.0956	0.8707	0.1178	0.5999	0.0223	0.9293	-0.0778	0.9137	-0.0480	0.8991	0.0298	0.9330
dpr2	dpr2	1627814_at	0.0858	0.6558	0.0421	0.8258	0.1278	0.4106	0.0141	0.9852	0.0309	0.8948	0.0168	0.9372	-0.0548	0.9296	0.0547	0.8390	0.1095	0.6129
CG13996	CG13996	1627815_at	-0.0407	0.7920	0.0463	0.7165	-0.0359	0.8773	-0.0683	0.8899	0.0331	0.8809	0.1014	0.5295	0.0045	0.9964	0.1107	0.7019	0.1061	0.7134
CG32264	CG32264	1627816_a_at	-0.3803	0.5045	-0.6989	0.2109	-1.1380	0.0066	-0.2182	0.5789	0.7266	0.0041	0.9448	0.0008	0.2906	0.9221	0.3021	0.8132	0.0115	0.9947
CG33109	CG33109	1627817_at	0.1760	0.3961	0.0318	0.8190	0.1822	0.3194	-0.0257	0.9745	0.0806	0.7250	0.1064	0.5884	-0.0079	0.9928	-0.0590	0.8212	-0.0511	0.8412
Gad1	Glutamic acid dec	1627818_s_at	0.1862	0.3363	-0.2026	0.2018	0.0593	0.7441	0.0279	0.9755	0.2416	0.2801	0.2137	0.2885	-0.2444	0.7485	-0.1943	0.5679	0.0501	0.9108
Or13a	Odorant receptor	1627819_at	0.1143	0.5674	-0.2711	0.0959	0.1303	0.5218	0.2793	0.4589	0.3166	0.1196	0.0373	0.8776	-0.1594	0.7768	-0.0240	0.9507	0.1354	0.6049
CG30096	CG30096	1627820_at	0.0694	0.8374	0.3366	0.0405	0.2813	0.1365	-0.1159	0.8578	-0.1804	0.4547	-0.0644	0.8047	-0.0713	0.9390	0.1417	0.6496	0.2130	0.4730
Syt7	Synaptotagmin VI	1627821_s_at	-1.0623	0.1037	-1.4887	0.0317	-1.7694	0.0007	0.0855	0.9463	0.5577	0.1182	0.4722	0.1393	0.3363	0.8940	0.1204	0.9354	-0.2159	0.8602
CG2938	CG2938	1627822_at	-0.8615	0.0290	0.4348	0.1528	0.7365	0.0024	-0.1727	0.7230	-1.1126	0.0010	-0.9399	0.0012	-0.2636	0.7697	0.3379	0.3685	0.6015	0.1433
mRpL35	mitochondrial ribo	1627823_at	0.3712	0.1095	0.7658	0.0613	0.3013	0.2385	0.0001	0.9999	-0.2635	0.2475	-0.2637	0.1939	0.2907	0.7728	0.0579	0.9341	-0.2329	0.6216
CG33635	CG33635	1627824_at	0.7261	0.0507	0.9179	0.0946	0.9945	0.0004	0.0640	0.9228	0.1906	0.3476	0.1265	0.5110	0.0136	0.9952	0.2606	0.6601	0.2471	0.6764
CG13305	CG13305	1627825_at	-0.0007	0.9974	0.0851	0.6176	0.1811	0.2309	0.0988	0.8868	0.1857	0.4459	0.0870	0.7289	-0.0138	0.9894	0.1010	0.6773	0.1148	0.6263
CG12744	CG12744	1627826_s_at	0.6348	0.1359	-0.5514	0.1206	-0.0959	0.7655	0.8133	0.1857	0.9472	0.0191	0.1339	0.7379	0.2955	0.8076	-0.1449	0.8284	-0.4404	0.3921
RpII33	RNA polymerase	1627827_a_at	0.1438	0.5320	0.3983	0.2761	0.3174	0.0681	0.0258	0.9777	0.0893	0.7372	0.0635	0.8030	0.1091	0.9174	0.2458	0.5146	0.1367	0.7448
Iat	Origin recognition	1627828_s_at	0.1274	0.6266	-0.3729	0.1820	-0.1780	0.4678	-0.2621	0.4553	0.3078	0.1048	0.5699	0.0057	-0.2682	0.7893	-0.1272	0.8185	0.1410	0.7788
Rh3	rhodopsin	1627829_at	0.2830	0.1013	0.2888	0.1704	0.0441	0.8516	-0.1098	0.8714	-0.0175	0.9604	0.0922	0.7135	0.2347	0.7116	0.0730	0.8372	-0.1616	0.5644
rpk	ripped pocket	1627830_at	0.8596	0.4751	-2.0897	0.1813	-1.9977	0.0612	-0.4139	0.2478	2.8396	0.0000	3.2535	0.0000	-0.4275	0.9611	-0.2701	0.9462	0.1575	0.9659
CG4891	CG4891	1627831_at	0.0220	0.9274	0.1287	0.2099	0.1969	0.4009	-0.2163	0.5735	-0.1012	0.6466	0.1151	0.5509	-0.0198	0.9873	0.1715	0.5635	0.1913	0.5175
scat	scattered	1627832_at	-0.4038	0.1117	-0.3575	0.2358	-0.2078	0.3685	0.2372	0.6354	0.4198	0.0884	0.1826	0.4284	0.0996	0.9409	0.4118	0.3087	0.3121	0.4683
Cralbp	Cellular retinaldeh	1627833_at	-0.2295	0.4027	0.1355	0.2280	-0.0060	0.9839	-0.1687	0.8218	-0.3007	0.3079	-0.1320	0.6630	-0.3426	0.5461	-0.1435	0.6135	0.1992	0.4683
aret	bruno	1627834_a_at	1.2987	0.1260	-1.2055	0.4063	-0.8531	0.0154	-0.3330	0.8189	1.6338	0.0113	1.9668	0.0029	-0.5959	0.8807	-0.7137	0.6704	-0.1177	0.9571
CG31973	CG31973	1627835_a_at	-1.1032	0.0574	-3.2386	0.0019	-2.3273	0.0046	0.9949	0.4456	2.0701	0.0107	1.0752	0.0911	-0.0299	0.9816	-0.1401	0.6483	-0.1102	0.7342
CG13801	CG13801	1627836_at	0.1710	0.4701	0.1312	0.4371	0.1256	0.4875	0.1700	0.7795	0.0345	0.9220	-0.1354	0.5912	-0.0291	0.9737	0.0210	0.9519	0.0501	0.8602
CG5527	CG5527	1627837_at	1.5280	0.0109	0.7503	0.1896	2.2751	0.0001	0.9609	0.1321	1.7096	0.0017	0.7487	0.0360	-0.7564	0.6272	0.8431	0.1800	1.5995	0.0518
uzip	zipper	1627838_at	-1.3909	0.0112	-0.2440	0.5799	-1.0067	0.0048	-0.5401	0.4509	-1.4940	0.0031	-0.9539	0.0140	0.2473	0.8714	-0.1667	0.8379	-0.4139	0.5125
CG2107	CG2107	1627839_at	0.6701	0.0097	0.5909	0.0463	0.8752	0.0027	0.0005	0.9994	-0.1112	0.5549	-0.1118	0.5081	-0.2040	0.7589	-0.1529	0.6093	0.0511	0.8938
zip	myosin II	1627840_a_at	0.1839	0.6945	1.1354	0.0625	0.4249	0.0711	0.2313	0.6936	0.3295	0.2234	0.0982	0.7363	0.9308	0.5258	1.2348	0.0890	0.3040	0.6978
CG11170	CG11170	1627841_at	-1.4474	0.0020	-1.6786	0.0207	-1.3030	0.0001	0.0727	0.9257	-0.4681	0.0514	-0.5408	0.0182	-0.3259	0.7458	-0.6449	0.1287	-0.3189	0.4649
CG14898	CG14898	1627842_at	-0.6468	0.0123	-0.4648	0.1283	-0.5511	0.0792	-0.2105	0.7576	-0.1709	0.5926	0.0396	0.9111	-0.1618	0.8718	0.0244	0.9704	0.1862	0.6668
CG12253 /// DyakCG12253	CG12253	1627843_at	-0.2528	0.2063	-1.2310	0.0239	-1.2579	0.0013	0.0283	0.9766	0.9136	0.0031	0.8852	0.0021	-0.1126	0.9247	-0.2079	0.6314	-0.0952	0.8552
Cyp4e2	Cytochrome P450	1627844_at	0.9327	0.0614	-0.0783	0.8546	-0.1085	0.5770	-0.0620	0.8967	0.0851	0.6268	0.1471	0.3032	-0.1205	0.9710	-1.0041	0.2145	-0.8836	0.3019
CG6673	CG6673	1627845_at	-0.0914	0.7505	0.2088	0.3734	0.1870	0.2373	-0.2045	0.4850	-0.3707	0.0256	-0.1662	0.2350	-0.1166	0.9301	-0.1243	0.8256	-0.0076	0.9921
---	---	1627846_at	-0.0046	0.9861	0.1030	0.4687	-0.0009	0.9964	-0.1045	0.8383	-0.1577	0.4269	-0.0532	0.8049	0.0215	0.9870	-0.0595	0.8946	-0.0810	0.8310
yip3	yippee interacting	1627847_s_at	0.1599	0.4432	0.1236	0.5360	0.1089	0.5655	-0.0506	0.9445	0.0340	0.9058	0.0846	0.6988	-0.0436	0.9742	-0.0450	0.9341	-0.0015	0.9982
---	---	1627848_at	0.2127	0.3844	0.0930	0.5696	-0.0353	0.8212	-0.0520	0.9375	-0.0408	0.8735	0.0113	0.9632	0.0798	0.9142	-0.1602	0.5543	-0.2400	0.3678
CG11456	CG11456	1627849_at	0.0076	0.9818	-0.0220	0.9220	-0.2226	0.1231	-0.2393	0.6304	0.0829	0.7790	0.3222	0.1379	-0.1211	0.8400	0.0921	0.7545	0.2132	0.3972
CG3689	CG3689	1627850_at	0.2999	0.2144	0.1164	0.6688	-0.0335	0.8394	-0.2266	0.5014	0.1697	0.3517	0.3963	0.0215	0.0949	0.9409	0.0556	0.9291	-0.0393	0.9434
CG10914	CG10914	1627851_at	-0.0717	0.6734	0.4967	0.0460	0.5647	0.0972	0.1342	0.7117	-0.0640	0.7422	-0.1982	0.1776	-0.1432	0.9092	0.2813	0.5452	0.4245	0.3545
CG14655	CG14655	1627852_at	-2.1445	0.0007	-2.1769	0.0027	-2.4617	0.0000	-0.2176	0.6120	-0.8151	0.0033	-0.5975	0.0080	0.3271	0.7726	-0.5712	0.2259	-0.8983	0.1013
---	---	1627853_at	0.1530	0.3718	0.1063	0.5345	0.0532	0.7524	-0.0723	0.8940	-0.0744	0.7234	-0.0021	0.9927	0.1783	0.7464	0.0264	0.9438	-0.1519	0.5334
CG9914	Enoyl-CoA hydrati	1627854_at	2.9390	0.0017	1.3353	0.3097	2.5593	0.0004	0.2827	0.6506	-0.8010	0.0160	-1.0837	0.0025	-0.9952	0.7606	-2.4301	0.0871	-1.4349	0.3056

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
yellow-e	yellow-e	1627855_at	0.1218	0.6748	0.1701	0.5617	0.1769	0.3359	0.0447	0.9302	-0.0325	0.8698	-0.0773	0.6098	-0.0251	0.9885	-0.0642	0.9115	-0.0392	0.9411
Hus1-like	Hus1-like	1627856_at	0.0469	0.9264	-0.2355	0.6677	-0.4370	0.0293	0.0155	0.9873	0.4521	0.0569	0.4366	0.0425	0.2503	0.8999	0.2420	0.7923	-0.0083	0.9950
---	---	1627857_at	0.0100	0.9633	0.1002	0.3453	0.0754	0.7389	-0.0118	0.9874	-0.0760	0.7146	-0.0642	0.7393	-0.0780	0.8999	0.0757	0.7914	0.1537	0.5194
---	---	1627858_at	-0.0299	0.8654	0.0850	0.4478	-0.1123	0.5305	-0.1266	0.7488	-0.2089	0.2239	-0.0823	0.6413	-0.0184	0.9816	-0.0365	0.8889	-0.0181	0.9431
CG31646	CG31646	1627859_at	-0.0256	0.9011	-0.1486	0.3972	0.1906	0.3528	-0.0168	0.9777	0.0013	0.9951	0.0181	0.9228	-0.1727	0.8042	-0.0961	0.7912	0.0766	0.8348
Or59c	Odorant receptor	1627860_at	0.0853	0.6386	0.0634	0.6411	0.3160	0.0461	-0.0348	0.9488	0.0504	0.7838	0.0852	0.5701	-0.1630	0.7500	0.0127	0.9724	0.1758	0.4327
Aldh-III	Fatty aldehyde de	1627861_s_at	-0.5789	0.0611	-0.1880	0.2283	-0.2449	0.3924	-0.0359	0.9622	-0.4554	0.0315	-0.4194	0.0280	-0.0791	0.9467	-0.1121	0.8123	-0.0330	0.9486
CG15267	CG15267	1627862_at	-0.2360	0.3352	-0.8509	0.0054	-0.4383	0.0405	0.2440	0.6999	0.4729	0.1033	0.2289	0.3975	-0.0901	0.9012	0.0594	0.8760	0.1495	0.6016
CG9947	CG9947	1627863_at	-0.4850	0.1331	0.3976	0.0919	1.0215	0.0004	0.1890	0.6202	-0.5636	0.0098	-0.7526	0.0016	-0.3481	0.6897	0.3527	0.3055	0.7007	0.0836
l(2)k07433	lethal (2) k07433	1627864_at	-0.0846	0.7471	0.1617	0.6772	-0.1646	0.5359	-0.1552	0.8028	0.2103	0.4130	0.3655	0.1005	0.1360	0.9359	0.5145	0.3259	0.3786	0.4955
---	---	1627865_a_at	0.3829	0.6824	0.4591	0.1095	-0.4787	0.4559	-0.2595	0.3619	0.3832	0.0259	0.6427	0.0016	0.8339	0.8049	0.7158	0.6394	-0.1181	0.9525
Top3alpha	Topoisomerase III	1627866_at	-0.4077	0.1932	-0.0985	0.8087	-0.0953	0.6538	-0.1023	0.8994	-0.4391	0.0967	-0.3367	0.1527	0.0678	0.9734	-0.0178	0.9855	-0.0856	0.9020
CG12438	CG12438	1627867_at	-0.0671	0.6932	0.0609	0.5757	0.0829	0.7236	0.1381	0.8034	0.1240	0.6128	-0.0141	0.9597	0.2077	0.7230	0.1551	0.5613	-0.0527	0.8788
Prat	transcript D	1627868_at	-0.3223	0.1830	-0.0545	0.6809	0.3705	0.1519	0.1597	0.8028	-0.2239	0.3944	-0.3836	0.0939	-0.2993	0.7363	0.1153	0.8147	0.4147	0.2869
Jon25Bi	Jonah 25B	1627869_at	0.4684	0.3664	0.0036	0.9836	-0.0123	0.9719	0.1604	0.7828	0.1366	0.6049	-0.0238	0.9363	0.0746	0.9721	-0.2156	0.7307	-0.2902	0.6175
CG31900	CG31900	1627870_at	-1.9042	0.0307	-0.1602	0.7565	-2.3276	0.0059	-1.9861	0.0492	-2.9195	0.0010	-0.9334	0.0729	0.0951	0.9860	-1.0540	0.3738	-1.1491	0.3552
Abd-B	Abdominal-B	1627871_at	0.2840	0.2035	0.2356	0.1493	0.0461	0.8643	-0.0087	0.9935	0.1187	0.5977	0.1274	0.5250	0.1292	0.8554	0.0515	0.9080	-0.0777	0.8321
CG3770	CG3770	1627872_at	-1.2670	0.0114	-0.7817	0.0264	-0.4890	0.1033	0.2136	0.7023	0.9252	0.0041	0.7115	0.0081	-0.2842	0.8331	1.3076	0.0415	1.5919	0.0385
CG5641	Interleukin enhanc	1627873_at	0.0653	0.6841	0.2907	0.0993	0.7773	0.0026	0.3271	0.2596	-0.1510	0.3922	-0.4781	0.0080	-0.1981	0.7220	0.0389	0.9157	0.2370	0.3276
xmas-1	transcript g	1627874_at	0.2222	0.4144	0.1895	0.2026	0.3500	0.0744	0.0788	0.8676	0.0209	0.9294	-0.0580	0.7499	0.1054	0.8889	0.1487	0.6280	0.0433	0.9121
CG12814	CG12814	1627875_at	0.2188	0.3410	0.5908	0.0454	0.2262	0.2115	-0.0832	0.8794	-0.1794	0.3478	-0.0962	0.6125	0.3629	0.6955	0.2749	0.4794	-0.0880	0.8602
CG5556	CG5556	1627876_at	0.0556	0.7727	0.1522	0.2646	0.1453	0.5438	-0.1705	0.6705	-0.2394	0.2069	-0.0689	0.7363	-0.0966	0.8960	-0.0029	0.9969	0.0937	0.7734
---	---	1627877_at	-0.0847	0.6236	0.0527	0.6058	0.2158	0.1401	-0.0877	0.8512	-0.1305	0.4627	-0.0428	0.8254	-0.1884	0.7220	0.0682	0.8165	0.2567	0.2731
---	---	1627878_at	-0.0823	0.7022	0.1316	0.3996	0.1424	0.3358	0.0569	0.9295	-0.1438	0.4643	-0.2007	0.2337	0.0156	0.9860	0.0054	0.9903	-0.0102	0.9739
---	---	1627879_at	0.0649	0.7408	0.0028	0.9818	-0.0797	0.6944	-0.0270	0.9626	0.0401	0.8375	0.0671	0.6722	0.0578	0.9342	0.0508	0.8749	-0.0069	0.9849
CG30334	CG30334	1627880_at	0.3539	0.1752	0.4013	0.1653	0.7436	0.0470	0.2235	0.7010	0.1827	0.5200	-0.0408	0.8992	0.0481	0.9816	0.2915	0.5715	0.2434	0.6434
salm	spalt	1627881_at	-0.1892	0.6708	-0.3492	0.2916	0.2316	0.3387	0.4456	0.5617	0.0696	0.8972	-0.3761	0.2916	-0.2196	0.8331	-0.2960	0.5076	-0.0764	0.8982
dlg1	Discs large	1627882_at	-2.8751	0.0023	-2.8921	0.0014	-3.0099	0.0001	0.1657	0.9319	0.4856	0.4148	0.3199	0.5782	-0.0803	0.9589	0.2340	0.6154	0.3143	0.4851
CG40230	CG40230	1627883_at	0.0558	0.8305	0.1433	0.2084	0.1736	0.3274	0.0030	0.9956	-0.0779	0.6821	-0.0808	0.6359	0.0261	0.9848	-0.1037	0.7921	-0.1298	0.7137
CG5281	CG5281	1627884_at	-0.1148	0.6197	-0.1050	0.4457	-0.4003	0.0367	-0.1125	0.7743	-0.1825	0.2690	-0.0700	0.6844	0.0941	0.9168	0.0725	0.8685	-0.0216	0.9622
CG12316	CG12316	1627885_s_at	0.1591	0.5710	0.0597	0.8667	-0.3001	0.0909	-0.1550	0.5931	0.0816	0.6148	0.2366	0.0736	0.0970	0.9342	-0.1011	0.8439	-0.1981	0.6282
Atg1	Autophagy-specifi	1627886_at	0.1613	0.8186	1.3486	0.0125	1.4780	0.0001	0.2753	0.7608	-0.1541	0.7372	-0.4295	0.2179	0.2353	0.8973	1.0365	0.1297	0.8012	0.2557
---	---	1627887_at	-0.1248	0.4613	-0.0214	0.8868	-0.1230	0.5829	-0.0848	0.9068	-0.0532	0.8627	0.0317	0.9109	-0.0647	0.9400	0.0348	0.9341	0.0996	0.7437
CG6672	CG6672	1627888_at	0.3853	0.0597	0.7284	0.0354	0.8372	0.0013	0.0050	0.9956	0.2782	0.1362	0.2733	0.1023	0.0084	0.9952	0.6746	0.0568	0.6662	0.0743
CG17826	CG17826	1627889_at	0.0796	0.7865	0.2199	0.3524	0.0710	0.7850	-0.1685	0.7647	-0.1037	0.7083	0.0648	0.8109	0.2806	0.7663	0.2031	0.6408	-0.0775	0.8884
GstD10	Glutathione S tran	1627890_at	-0.7545	0.0833	-0.9653	0.0426	-0.1227	0.6584	0.2630	0.6041	0.4372	0.0880	0.1743	0.4736	-0.5940	0.6483	0.0981	0.9019	0.6920	0.2075
CG12725	CG12725	1627891_at	0.0774	0.7096	0.0013	0.9928	0.0621	0.7058	0.1090	0.8485	0.2428	0.2458	0.1337	0.5067	0.0128	0.9914	0.0758	0.8353	0.0630	0.8602
---	---	1627892_at	0.2315	0.1635	0.1172	0.3958	-0.0802	0.6292	-0.0136	0.9857	0.0341	0.8837	0.0477	0.8066	0.0654	0.9152	-0.0830	0.7424	-0.1484	0.5050
CG18177	CG18177	1627893_at	0.0727	0.6875	0.0274	0.9203	-0.2695	0.1394	-0.2579	0.5128	0.1206	0.5953	0.3785	0.0474	-0.1194	0.8814	-0.0749	0.8602	0.0445	0.9164
Chi	Chip	1627894_s_at	-0.4521	0.1104	-0.0962	0.8861	-0.1368	0.6824	0.1709	0.6268	-0.1564	0.3781	-0.3273	0.0412	-0.1013	0.9776	0.0858	0.9506	0.1871	0.8717
CG18404	CG18404	1627895_at	0.0339	0.9226	0.0222	0.8266	-0.1190	0.5290	0.0190	0.9863	0.0562	0.8720	0.0372	0.9071	0.1114	0.9168	-0.0370	0.9505	-0.1484	0.7245
CG32275	CG32275	1627896_at	0.0297	0.8758	-0.0403	0.7309	0.1707	0.4546	0.0635	0.9393	0.0365	0.9145	-0.0270	0.9274	-0.1813	0.7644	-0.0830	0.7948	0.0983	0.7362
CG12265	CG12265	1627897_at	0.0689	0.8160	-0.3564	0.1229	-0.2798	0.3978	-0.0240	0.9777	0.4595	0.0430	0.4835	0.0223	-0.2811	0.8000	0.0126	0.9904	0.2937	0.5488
Sce	dRing1	1627898_at	-0.5296	0.1018	-0.1868	0.3492	-0.1713	0.4701	0.1584	0.8138	0.0738	0.8244	-0.0846	0.7725	0.1542	0.8903	0.4015	0.3259	0.2473	0.5803
---	---	1627899_at	0.2027	0.5101	-0.1498	0.6840	-0.6994	0.0195	-0.1418	0.8907	0.5895	0.0888	0.7313	0.0260	0.0864	0.9652	0.1168	0.8776	0.0304	0.9702
CG3058	CG3058	1627900_at	-0.7823	0.0144	-0.5664	0.1160	-0.6850	0.0026	-0.1254	0.8014	-0.1166	0.5932	0.0088	0.9718	-0.0282	0.9831	0.0739	0.8658	0.1021	0.7820
CG9692	CG9692	1627901_a_at	0.2861	0.2632	0.1390	0.2074	-0.0600	0.7753	-0.0785	0.8676	-0.0251	0.9142	0.0534	0.7715	0.3097	0.5461	-0.0146	0.9708	-0.3243	0.1990
CG41434	CG41434	1627902_at	0.0873	0.6261	-0.0753	0.6659	0.1266	0.4359	-0.0267	0.9671	0.0951	0.6092	0.1218	0.4439	-0.2342	0.7822	-0.0529	0.9275	0.1813	0.6457
CG9040	CG9040	1627903_at	-0.0302	0.8519	0.1271	0.4436	0.1056	0.4731	-0.0604	0.9011	-0.0247	0.9103	0.0358	0.8441	0.0217	0.9813	0.1260	0.5404	0.1043	0.6231
CG15375	CG15375	1627904_at	0.0618	0.7574	0.0086	0.9446	0.2366	0.1369	-0.0145	0.9873	-0.0476	0.8696	-0.0331	0.9000	-0.0597	0.9309	0.0142	0.9690	0.0739	0.7777
---	---	1627905_at	-0.0617	0.8326	0.0452	0.6908	0.0924	0.5485	0.0947	0.8409	0.0515	0.8128	-0.0432	0.8296	0.1520	0.8472	0.1130	0.7788	-0.0389	0.9330
---	---	1627906_at	-0.0158	0.9634	0.1667	0.3853	0.1475	0.5578	-0.0109	0.9943	0.0508	0.9122	0.0617	0.8747	-0.0035	0.9971	0.0769	0.7996	0.0804	0.7749
---	---	1627907_at	0.0931	0.5260	0.1683	0.3490	0.1300	0.3770	-0.1105	0.7243	-0.1740	0.2163	-0.0635	0.6643	-0.0207	0.9816	-0.0275	0.9347	-0.0069	0.9846
CG14561 /// DmojCG14561	CG14561	1627908_at	0.1780	0.5025	0.1786	0.3347	0.0807	0.6274	-0.1574	0.7967	-0.0226	0.9517	0.1348	0.5869	0.0374	0.9819	0.0297	0.9629	-0.0077	0.9907
---	---	1627909_at	0.0079	0.9758	0.0800	0.7681	0.0709	0.6772	0.0979	0.8463	0.0347	0.8902	-0.0631	0.7552	0.1682	0.8885	0.1180	0.8481	-0.0502	0.9369
CG8036	CG8036	1627910_s_at	0.7717	0.0230	1.0529	0.0395	1.8655	0.0001	0.5551	0.0564	-0.8762	0.0009	-1.4313	0.0001	-0.1829	0.8875	-0.5710	0.2224	-0.3881	0.4364
CG14286	CG14286	1627911_at	0.5903	0.0246	0.1952	0.5664	0.0617	0.7358	0.1868	0.7169	0.3833	0.1021	0.1964	0.3634	0.1888	0.8650	-0.0105	0.9923	-0.1993	0.6843
CG5493	CG5493	1627912_at	2.0720	0.0075	-0.2135	0.9126	1.2955	0.4207	1.0451	0.6607	1.6283	0.1517	0.5833	0.6137	-0.2430	0.9822	-0.4943	0.8960	-0.2513	0.9433
---	---	1627913_at	0.1911	0.3107	0.0716	0.5432	0.1997	0.3254	0.0612	0.8852	0.1305	0.3742	0.0694	0.6382	0.0139	0.9895	0.0150	0.9670	0.0011	0.9979
---	---	1627914_at	0.0631	0.8367	0.0273	0.9329	-0.0333	0.8733	0.1005	0.8603	-0.0043	0.9891	-0.1048	0.6128	0.0806	0.9552	-0.0542	0.9353	-0.1348	0.7868
Gapdh2	G3-P dehydrogen	1627915_a_at	-0.0387	0.8262	-0.1727	0.4617	-0.0974	0.5111	-0.1583	0.6927	-0.2574	0.1625	-0.0991	0.5931	-0.1494	0.7997	-0.3367	0.1579	-0.1874	0.4597
ras	inosine monophos	1627916_s_at	0.8140	0.0058	0.2041	0.2345	0.4641	0.0248	0.1966	0.6013	0.4478	0.0261	0.2512	0.1373	-0.0280	0.9816	-0.1107	0.7439	-0.0827	0.8178
dpr4	dpr4	1627917_at	0.2619	0.1936	0.3908	0.0314	0.5135	0.0107	0.0410	0.9375	-0.1369	0.3781	-0.1779	0.1896	-0.0888	0.9087	-0.1529	0.6009	-0.0640	0.8577
CG15776	CG15776	1627918_at	-0.0526	0.7600	0.1669	0.4354	-0.0888	0.6139	0.0232	0.9767	-0.0565	0.8157	-0.0796	0.6988	0.0475	0.9514	-0.0560	0.8631	-0.1035	0.6877
CG2713	CG2713	1627919_at	-0.4599	0.1427	0.1361	0.4056	0.8956	0.0035	0.2788	0.6506	-0.3534	0.2349	-0.6322	0.0257	-0.4704	0.6531	0.2167	0.6337	0.6871	0.1313
CG17758 /// DmirCG17758	CG17758	1627920_at	0.0379	0.8097	-0.0766	0.5017	0.0459	0.8224	0.1216	0.7514	0.0673	0.7303	-0.0543	0.7675	-0.1214	0.8331	-0.0547	0.8731	0.0667	0.8247
CG33681	CG33681	1627921_at	0.1262	0.5676	0.1398	0.2965	0.1566	0.3923	0.0874	0.8841	-0.1645	0.4369	-0.2519	0.1664	-0.0175	0.9898	-0.0244	0.9590	-0.0069	0.9878
CG32544	CG32544	1627922_at	-0.3089	0.0888	0.0667	0.7903	-0.0465	0.8060	0.1366	0.8655	-0.0737	0.8415	-0.2103	0.4326	0.2547	0.7215	0.2448	0.4107	-0.0099	0.9846
CG12950	CG12950	1627923_at	0.2257	0.4247	0.1557	0.2198	0.2713	0.1081	-0.1418	0.7906	-0.1111	0.6506	0.0307	0.9075	-0.0742	0.9246	-0.0254	0.9519	0.0488	0.8921

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Rpl135	RNA polymerase	1627924_at	0.7188	0.1054	-0.0117	0.9882	0.6031	0.0605	0.4074	0.6513	1.0792	0.0215	0.6718	0.0852	-0.2415	0.9016	0.3700	0.6325	0.6115	0.4017
spict	spichthyin	1627925_at	0.0026	0.9896	-0.6392	0.0484	-0.1604	0.4684	0.1235	0.8469	0.3127	0.1793	0.1893	0.3832	-0.3471	0.5614	-0.3417	0.1948	0.0054	0.9907
CG8315	CG8315	1627926_at	0.6678	0.0486	0.1917	0.3385	-0.0266	0.8788	-0.1168	0.7432	0.4863	0.0081	0.6031	0.0019	0.2451	0.7975	0.1422	0.7699	-0.1029	0.8402
CG8443	CG8443	1627927_at	0.2196	0.2748	-0.1292	0.8025	0.0301	0.9049	0.0541	0.9132	0.3607	0.0291	0.3065	0.0357	-0.2336	0.8740	-0.0915	0.9188	0.1421	0.8484
CG12229 /// CG14101	CG14101 /// CG1	1627928_at	0.0349	0.8744	0.0419	0.7018	0.1558	0.3575	0.1059	0.8738	0.1111	0.6757	0.0052	0.9858	0.0162	0.9881	0.1480	0.5523	0.1318	0.6056
CG14632	CG14632	1627929_at	-0.1540	0.3728	0.0361	0.7293	0.0566	0.7389	0.0637	0.8858	-0.0137	0.9517	-0.0773	0.6111	-0.0314	0.9717	0.0962	0.6977	0.1276	0.5865
CG32280	CG32280	1627930_s_at	-0.4082	0.1266	-0.7637	0.0159	-0.8581	0.0009	-0.0313	0.9649	0.2000	0.2983	0.2313	0.1740	0.0349	0.9816	-0.2323	0.4829	-0.2672	0.4175
---	---	1627931_at	0.0338	0.8827	-0.0140	0.9318	-0.1458	0.4728	-0.0885	0.8138	0.1088	0.4828	0.1973	0.1344	0.0474	0.9775	0.0085	0.9925	-0.0389	0.9435
how	struthio	1627932_at	0.0733	0.8141	-0.0538	0.6223	-0.1115	0.5214	0.0565	0.9371	0.0941	0.6942	0.0375	0.8791	0.0467	0.9677	-0.1403	0.6762	-0.1870	0.5605
mXr	mXr	1627933_a_at	-0.0654	0.7037	-0.0779	0.6367	-0.0251	0.9077	-0.0441	0.9436	0.1489	0.4195	0.1930	0.2274	-0.0721	0.9099	0.0233	0.9499	0.0954	0.7087
CG11723	CG11723	1627934_at	0.0827	0.6936	0.2638	0.1648	0.3334	0.0427	0.0091	0.9921	-0.0609	0.7887	-0.0700	0.7245	-0.0562	0.9460	0.1627	0.5336	0.2189	0.3921
Obp59a	Odorant-binding p	1627935_a_at	0.0491	0.7836	0.0860	0.5884	0.2124	0.2737	-0.0426	0.9558	-0.1033	0.6613	-0.0607	0.7955	-0.0373	0.9653	0.0176	0.9629	0.0549	0.8534
---	---	1627936_s_at	0.1222	0.7490	-1.6263	0.0387	-1.1058	0.0239	0.4006	0.7202	2.0963	0.0023	1.6956	0.0034	-0.0061	0.9978	0.2048	0.7128	0.2109	0.7004
CG32413	CG32413	1627937_at	0.0856	0.5266	0.0410	0.7938	0.0372	0.8517	0.0311	0.9672	-0.0318	0.9080	-0.0629	0.7742	-0.0201	0.9852	0.0263	0.9482	0.0464	0.8918
CG17840	CG17840	1627938_at	-0.2482	0.2867	-0.5975	0.1573	-0.1023	0.6147	0.2471	0.4908	0.4463	0.0272	0.1991	0.2455	-0.2181	0.8619	0.0882	0.9093	0.3063	0.5691
ferrochelatase	ferrochelatase	1627939_a_at	0.0322	0.8527	0.2187	0.3756	0.5572	0.0126	0.1517	0.7447	-0.3396	0.0972	-0.4913	0.0153	-0.2794	0.6984	-0.1093	0.7668	0.1701	0.6030
---	---	1627940_at	-0.0264	0.8693	-0.0709	0.5297	-0.1319	0.4916	-0.0366	0.9532	0.1284	0.4736	0.1649	0.2875	-0.0486	0.9514	-0.0696	0.8247	-0.0209	0.9505
CG12984	CG12984	1627941_at	0.0723	0.6513	0.0251	0.8209	0.0702	0.7499	-0.1339	0.6965	-0.1546	0.3359	-0.0207	0.9143	-0.0965	0.8903	-0.0636	0.8619	0.0329	0.9279
CG17186	CG17186	1627942_at	-0.1232	0.6868	0.3340	0.0906	0.5126	0.0498	-0.0421	0.9592	-0.4469	0.0446	-0.4048	0.0424	-0.1694	0.8680	0.1044	0.8499	0.2738	0.5154
Ilp3	Drosophila insulin	1627943_at	0.0828	0.7640	0.1222	0.5825	0.1759	0.4814	0.2366	0.6471	0.3137	0.2083	0.0772	0.7820	-0.0174	0.9893	0.0297	0.9460	0.0471	0.8985
CG9018	CG9018	1627944_a_at	-0.1214	0.6009	0.5567	0.0417	0.5876	0.0255	0.1879	0.5564	-0.4417	0.0154	-0.6296	0.0019	0.1469	0.8742	0.3898	0.2673	0.2430	0.5257
Fdxh	Ferredoxin	1627945_at	-0.2546	0.1267	-0.0914	0.6882	0.1987	0.3035	-0.0333	0.9558	-0.2816	0.0813	-0.2484	0.0849	-0.2265	0.7783	-0.1493	0.7058	0.0772	0.8682
CG12068	CG12068	1627946_at	0.1808	0.3042	0.1944	0.5528	0.2039	0.3057	0.0242	0.9722	-0.0418	0.8529	-0.0661	0.7216	0.0712	0.9030	0.1054	0.6483	0.0343	0.9070
---	---	1627947_at	-0.0193	0.9092	0.1342	0.3650	-0.1083	0.5004	-0.1007	0.8033	-0.0401	0.8484	0.0606	0.7258	0.1751	0.8202	0.0761	0.8634	-0.0990	0.7964
aPKC	pschur	1627948_at	-0.6729	0.1438	-0.2802	0.5514	-0.9144	0.0021	-0.3547	0.4828	-0.4438	0.1005	-0.0891	0.7637	0.2390	0.8882	0.0123	0.9935	-0.2266	0.7577
---	---	1627949_at	-0.0342	0.8494	-0.0160	0.9078	0.0185	0.9183	-0.0501	0.9289	-0.0534	0.7916	-0.0033	0.9874	-0.0370	0.9624	-0.0972	0.6912	-0.0602	0.8256
CG14237	CG14237	1627950_at	0.0117	0.9652	0.0722	0.5827	-0.0083	0.9830	-0.2672	0.6041	-0.2290	0.3887	0.0382	0.9027	-0.0588	0.9467	-0.1045	0.7411	-0.0457	0.9025
CG33289	CG33289	1627951_at	0.0020	0.9948	0.0767	0.5835	0.0366	0.9019	-0.0847	0.8937	-0.1371	0.5459	-0.0524	0.8277	0.1054	0.9227	0.0143	0.9837	-0.0911	0.8434
CG8102	CG8102	1627952_a_at	0.1634	0.3634	0.0196	0.9700	0.0343	0.8453	0.0336	0.9770	0.1549	0.6247	0.1213	0.6850	0.1349	0.8427	0.1349	0.6601	0.0001	0.9999
mod(mdg4)	Modifier67.2	1627953_at	-0.4257	0.3511	-0.0146	0.9753	-0.6383	0.0887	-0.4491	0.2842	-0.2552	0.3054	0.1938	0.3985	-0.0103	0.9977	-0.1036	0.9353	-0.0933	0.9340
CG9184 /// DyakCG9184	CG9184	1627954_a_at	-0.2457	0.2237	-0.2160	0.2402	-0.6754	0.0149	-0.0837	0.9381	0.2376	0.4736	0.3213	0.2602	-0.0656	0.9589	0.0338	0.9531	0.0993	0.8247
Asator	Asator	1627955_a_at	-2.0114	0.0008	-1.8245	0.0584	-2.3199	0.0001	-0.4454	0.3228	-0.6935	0.0162	-0.2481	0.2960	0.0580	0.9849	-0.3879	0.6053	-0.4459	0.5488
CG14216	CG14216	1627956_x_at	-0.1024	0.5661	-0.3712	0.3158	-0.6656	0.0126	-0.2297	0.6149	-0.0449	0.8838	0.1848	0.3834	-0.0653	0.9514	-0.3884	0.2102	-0.3232	0.3221
CG1638	CG1638	1627957_at	-0.0951	0.5683	0.0934	0.5406	0.5413	0.0269	0.2036	0.5540	-0.1770	0.3267	-0.3806	0.0248	-0.1008	0.8062	0.0189	0.9460	0.1198	0.5003
CG30269	CG30269	1627958_at	0.3974	0.0342	0.2122	0.5643	0.4696	0.1429	0.0350	0.9649	0.2019	0.3514	0.1670	0.3990	-0.0802	0.9717	0.0752	0.9345	0.1555	0.8287
dj	don juan	1627959_a_at	-0.0283	0.9226	-0.1801	0.1178	0.0372	0.8884	0.1065	0.8771	0.1246	0.6378	0.0182	0.9517	-0.2254	0.7633	-0.0025	0.9981	0.2229	0.4925
---	---	1627960_at	0.2547	0.2998	-0.0353	0.9127	-0.0364	0.8938	0.0889	0.8623	0.2775	0.1278	0.1886	0.2507	0.1507	0.9011	0.1746	0.7361	0.0240	0.9722
CG18507	CG18507	1627961_a_at	-2.2415	0.0006	-0.8925	0.1865	-0.8319	0.0067	-0.2843	0.4568	-1.9616	0.0001	-1.6773	0.0001	-0.2603	0.8910	-0.5124	0.4848	-0.2521	0.7619
CG31749	CG31749	1627962_at	-0.4082	0.2503	-0.3759	0.0425	-0.7109	0.0595	0.4424	0.2646	0.7747	0.0061	0.3323	0.1117	0.6947	0.6749	0.8149	0.2101	0.1202	0.8955
Hsc70-1	Heat shock protei	1627963_s_at	-1.5532	0.0022	-1.1409	0.1080	-1.4185	0.0002	-0.0629	0.9466	-0.3848	0.1461	-0.3219	0.1749	0.0408	0.9812	-0.0880	0.8657	-0.1288	0.7649
CG7888	CG7888	1627964_s_at	-1.0278	0.5982	-0.9640	0.2221	-2.8831	0.0004	-1.8843	0.0417	-2.3357	0.0016	-0.4514	0.3411	0.0430	0.9964	-2.3841	0.3120	-2.4271	0.3271
---	---	1627965_s_at	-0.1512	0.4853	-0.7506	0.0169	-1.4790	0.0150	-0.0076	0.9956	1.2470	0.0082	1.2546	0.0048	0.0651	0.9441	0.0299	0.9477	-0.0351	0.9282
CG6406	CG6406	1627966_s_at	-0.3616	0.0819	-0.5299	0.0534	-0.8278	0.0038	-0.2602	0.5008	0.3751	0.0692	0.6353	0.0048	-0.0081	0.9958	0.2586	0.4697	0.2667	0.4648
mthl4	mth-like 4	1627967_a_at	-0.6584	0.0105	-1.6271	0.0290	-1.5125	0.0004	0.1923	0.7631	0.8637	0.0075	0.6714	0.0139	0.0656	0.9677	0.0139	0.9873	-0.0517	0.9318
---	---	1627968_at	0.1498	0.3740	0.0133	0.9095	-0.2081	0.2440	-0.1974	0.5259	0.1461	0.3812	0.3434	0.0266	0.0559	0.9225	-0.0247	0.9353	-0.0807	0.7060
CG9289	glutactin-like	1627969_at	0.0665	0.7404	0.0556	0.6081	0.1780	0.3865	0.0326	0.9637	-0.0432	0.8618	-0.0758	0.7039	0.0501	0.9398	0.0254	0.9388	-0.0248	0.9310
Hel89B	89B helicase	1627970_at	-0.3617	0.1285	-0.5500	0.0450	-0.5312	0.0521	0.4877	0.3040	0.9313	0.0058	0.4436	0.0768	0.4625	0.5461	0.7344	0.0613	0.2719	0.4597
sano	serrano	1627971_s_at	0.0861	0.8987	0.0705	0.8665	-0.7436	0.0301	-0.3300	0.4567	-0.4755	0.0524	-0.1454	0.5349	0.3859	0.8692	-0.5606	0.5640	-0.9465	0.3175
CG9822	CG9822	1627972_at	0.1444	0.4492	0.1657	0.2771	0.2171	0.2128	-0.0488	0.9413	-0.0147	0.9585	0.0341	0.8800	-0.2606	0.7046	-0.2171	0.4493	0.0435	0.9121
---	---	1627973_s_at	-0.5178	0.0723	-0.2657	0.4048	0.0526	0.8110	0.1363	0.7524	-0.2097	0.2644	-0.3461	0.0444	-0.2393	0.8042	0.0167	0.9828	0.2561	0.5457
Cyp6t3	Cyp6t3	1627974_at	0.4915	0.0335	0.2522	0.3250	0.9764	0.0084	0.1330	0.6984	0.0214	0.9226	-0.1116	0.4535	-0.4008	0.7070	-0.1753	0.7357	0.2255	0.6389
TotB	Turandot B	1627975_at	0.6317	0.0365	0.1875	0.4672	0.2191	0.1921	0.2210	0.5478	0.0320	0.9029	-0.1890	0.2709	0.0860	0.9572	-0.2261	0.6479	-0.3121	0.5099
Gr39a	Gustatory recepto	1627976_at	0.0287	0.8783	-0.1072	0.5524	0.0867	0.5932	0.0309	0.9639	0.1411	0.4526	0.1102	0.5318	-0.0416	0.9589	0.0303	0.9353	0.0718	0.7978
CG1671	CG1671	1627977_at	0.5692	0.1219	-0.4400	0.4185	-0.0461	0.7850	0.3671	0.5869	0.9020	0.0170	0.5348	0.0833	-0.0879	0.9643	-0.0566	0.9477	0.0313	0.9695
CG13394	CG13394	1627978_at	-0.1418	0.4107	-0.0431	0.7954	-0.1697	0.5183	-0.0376	0.9757	-0.1024	0.7784	-0.0648	0.8528	0.1093	0.8890	-0.1452	0.6512	-0.2546	0.3972
Wnt6	Wnt6	1627979_at	-1.5352	0.0036	-0.5227	0.2650	-1.1851	0.0047	-0.3672	0.5159	-0.0132	0.9775	0.3541	0.1838	0.1262	0.9095	0.4420	0.2466	0.3158	0.4367
---	---	1627980_at	-0.3204	0.0678	-0.1042	0.4845	0.0499	0.7802	0.1592	0.7149	-0.0744	0.7535	-0.2336	0.1880	0.0351	0.9611	0.0259	0.9363	-0.0093	0.9769
CG32238	CG32238	1627981_at	0.3673	0.1603	0.1171	0.2536	0.3848	0.0492	0.1074	0.8492	0.0383	0.8906	-0.0691	0.7592	-0.0777	0.9342	-0.1842	0.5622	-0.1065	0.7620
CG31077	CG31077	1627982_at	0.2119	0.3223	-0.0114	0.9152	0.0525	0.7890	0.0947	0.8714	-0.0259	0.9282	-0.1206	0.5471	-0.1130	0.8292	-0.3330	0.1308	-0.2200	0.3271
---	---	1627983_at	0.2160	0.3447	0.5294	0.0436	0.5622	0.0160	0.0233	0.9733	-0.1749	0.3122	-0.1982	0.1951	0.1023	0.9239	0.1733	0.6597	0.0710	0.8857
CG13917 /// DmirCG13917	CG13917	1627984_at	-0.9098	0.1319	-0.9231	0.2184	-1.3599	0.0019	0.0434	0.9688	0.2168	0.4714	0.1734	0.5375	0.4257	0.8690	0.2227	0.8790	-0.2030	0.8849
CG31870	CG31870	1627985_a_at	-0.0573	0.7941	0.0228	0.8207	-0.0185	0.9373	-0.0514	0.9436	0.2441	0.2370	0.2956	0.1082	0.0060	0.9964	0.1292	0.7317	0.1232	0.7422
PGRP-SC1a /// PGRP-SC1	PGRP-SC1a /// Pt	1627986_s_at	0.1902	0.4717	0.2920	0.0763	0.1868	0.4325	-0.0899	0.9282	-0.1844	0.5626	-0.0944	0.7723	0.0870	0.9032	0.0383	0.9264	-0.0487	0.8918
---	---	1627987_s_at	-0.1212	0.4569	0.2207	0.2634	0.2976	0.1511	0.0047	0.9956	-0.1710	0.3467	-0.1757	0.2765	0.0185	0.9913	0.1801	0.6218	0.1616	0.6607
RpL4	Ribosomal protein	1627988_at	0.0308	0.8382	0.1055	0.6096	0.2999	0.0670	0.1623	0.6202	-0.0802	0.6651	-0.2425	0.0961	-0.0237	0.9831	-0.0878	0.7903	-0.0641	0.8508
---	---	1627989_at	-0.1198	0.5636	-0.0781	0.5552	0.1043	0.6395	-0.0004	0.9994	-0.0087	0.9734	-0.0083	0.9689	0.0521	0.9672	0.2255	0.5120	0.1734	0.6291
Acp53Ea	Accessory gland p	1627990_at	0.0902	0.5441	0.0407	0.8387	0.2037	0.3620	0.0616	0.9117	-0.0041	0.9880	-0.0657	0.7278	-0.0350	0.9737	-0.0847	0.7971	-0.0497	0.8897
---	---	1627991_at	0.1557	0.4291	-0.0036	0.9788	0.1908	0.2578	0.1029	0.7845	-0.0038	0.9875	-0.1067	0.4687	-0.1048	0.8671	-0.0145	0.9724	0.0904	0.7526
---	---	1627992_at	0.2421	0.3377	-0.0771	0.5643	0.0197	0.9131	0.1099	0.8594	0.1950	0.3944	0.0851	0.7204	0.0346	0.9775	0.0748	0.8500	0.0402	0.9197

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11396	CG11396	1627993_at	0.3196	0.0926	-0.0356	0.9445	-0.3468	0.0829	-0.1408	0.7937	0.9002	0.0024	1.0410	0.0008	0.1198	0.9238	0.5138	0.1975	0.3940	0.3539
CG33296	CG33296	1627994_at	0.1336	0.4472	-0.0660	0.6780	0.0167	0.9398	0.0426	0.9451	0.0779	0.7021	0.0353	0.8637	-0.2203	0.7633	-0.2062	0.5141	0.0140	0.9769
Art8	Arginine methyltra	1627995_at	-0.6923	0.0554	-1.2795	0.0859	-0.6075	0.0316	0.3987	0.4757	0.7633	0.0193	0.3646	0.1738	-0.2712	0.8846	0.2297	0.7978	0.5009	0.4944
Pi4KIIalpha	Pi4KIIalpha	1627996_a_at	-0.4132	0.1635	0.2806	0.4825	0.2989	0.1790	0.0407	0.9603	-0.2940	0.1732	-0.3346	0.0852	-0.0104	0.9964	0.4204	0.4433	0.4308	0.4457
eyc	eyes closed	1627997_at	-0.0044	0.9859	0.0000	1.0000	-0.0056	0.9827	0.0255	0.9715	-0.0203	0.9377	-0.0458	0.8234	0.1726	0.7979	0.0339	0.9404	-0.1387	0.6475
CG2225	CG2225	1627998_s_at	-0.0507	0.9245	0.1748	0.7193	-0.0838	0.8123	0.0257	0.9777	-0.1083	0.6724	-0.1340	0.5428	0.2749	0.8903	0.2625	0.7747	-0.0124	0.9924
barr	Barren	1627999_s_at	0.0755	0.7304	-0.3738	0.1176	-0.4062	0.0410	-0.1856	0.7227	0.3265	0.1638	0.5120	0.0233	-0.1901	0.7726	-0.1414	0.6479	0.0487	0.9020
CG33489	CG33489	1628000_s_at	0.0169	0.9425	-0.1369	0.3313	0.1113	0.4994	0.2299	0.5388	0.2659	0.1710	0.0360	0.8767	-0.0473	0.9589	-0.0442	0.9111	0.0031	0.9946
CG10492	transcription unit 3	1628001_at	0.4420	0.0559	-0.0170	0.9647	0.0183	0.9306	-0.1251	0.8034	0.3392	0.0926	0.4644	0.0179	-0.1101	0.9117	-0.1089	0.8117	0.0012	0.9988
CG33986	CG33986	1628002_at	0.0188	0.9130	0.1172	0.4524	-0.0631	0.7487	-0.0144	0.9834	-0.0682	0.7102	-0.0538	0.7565	0.1845	0.6955	0.1445	0.4702	-0.0400	0.8818
---	---	1628003_s_at	0.1653	0.4339	0.0920	0.3450	0.2411	0.2193	0.0118	0.9865	0.0412	0.8521	0.0294	0.8840	-0.0643	0.9522	0.0215	0.9679	0.0858	0.8326
---	---	1628004_at	0.1730	0.5589	-0.0558	0.5990	0.0078	0.9837	-0.0702	0.9380	0.0211	0.9569	0.0914	0.7523	0.0537	0.9774	-0.0150	0.9868	-0.0687	0.9108
CG31666 /// chinmo	CG31666 /// chror	1628005_at	-0.0341	0.9081	-0.1841	0.3857	-0.2394	0.5106	-0.3042	0.7678	0.0304	0.9638	0.3346	0.4139	-0.2131	0.7726	-0.1926	0.5629	0.0204	0.9659
DebB	Developmental en	1628006_at	0.5380	0.1015	0.5711	0.0875	0.7710	0.0372	0.1072	0.8416	0.1801	0.3776	0.0728	0.7345	-0.0815	0.9735	0.2600	0.7040	0.3415	0.5995
---	---	1628007_at	0.0739	0.6476	0.0745	0.5103	-0.0933	0.6669	0.0271	0.9665	0.1367	0.4353	0.1096	0.4999	-0.0376	0.9781	-0.0554	0.9121	-0.0178	0.9717
TER94	Complementation	1628008_a_at	0.3037	0.3735	0.4149	0.0505	0.5566	0.0062	0.1861	0.5528	0.4070	0.0203	0.2209	0.1258	0.0367	0.9848	0.4908	0.2329	0.4541	0.2984
fas	faint sausage	1628009_at	0.2102	0.3512	-0.0289	0.8489	0.0892	0.5351	-0.0296	0.9687	0.1355	0.5112	0.1651	0.3575	-0.0381	0.9717	-0.0297	0.9455	0.0084	0.9848
CG34386	CG14496	1628010_at	0.1276	0.4785	0.1315	0.6130	0.1784	0.4901	0.0247	0.9838	0.0551	0.8811	0.0304	0.9281	0.3120	0.7070	0.1202	0.7787	-0.1918	0.6087
CG12877	CG12877	1628011_a_at	-0.7575	0.1779	-0.4077	0.4402	-0.1319	0.4340	0.1768	0.7929	0.1001	0.7613	-0.0767	0.8041	-0.1282	0.9589	0.1995	0.8352	0.3277	0.6761
---	---	1628012_at	-0.0489	0.8097	0.0653	0.5428	-0.1068	0.5060	-0.1568	0.7138	-0.2312	0.2360	-0.0743	0.7204	0.1367	0.8235	0.0787	0.8123	-0.0580	0.8635
CG3597	CG3597	1628013_at	2.7461	0.0009	1.5825	0.1026	3.2138	0.0003	0.9843	0.2355	0.5979	0.2235	-0.3864	0.3988	-0.9268	0.6898	-0.6989	0.4714	0.2279	0.8542
CG5703 /// DyakCG5703	CG5703	1628014_at	-0.0318	0.9405	0.0781	0.7922	-0.0428	0.8502	-0.1396	0.7962	-0.5433	0.0217	-0.4037	0.0456	-0.0208	0.9926	-0.4179	0.4095	-0.3971	0.4524
cngl	CNG channel-like	1628015_at	0.1151	0.5054	0.1135	0.5574	0.0759	0.6387	-0.0226	0.9790	0.0430	0.8803	0.0656	0.7831	0.1298	0.8494	0.0274	0.9521	-0.1025	0.7541
ari-1	Ariadne-1	1628016_s_at	-0.7790	0.2264	0.0943	0.4895	-0.6381	0.0310	-0.1343	0.8500	-0.4103	0.1117	-0.2761	0.2334	0.5806	0.7324	0.4533	0.5523	-0.1273	0.8990
me31B	maternal expressi	1628017_a_at	0.3879	0.3301	-0.5771	0.2569	-0.5553	0.1165	-0.1427	0.7937	1.3085	0.0006	1.4512	0.0002	-0.1723	0.9450	0.2914	0.7439	0.4638	0.5637
Hml	hemolectin	1628018_at	-2.5312	0.0074	-4.4672	0.0077	-4.5659	0.0000	-0.1728	0.5388	0.1674	0.2555	0.3402	0.0176	-0.0014	0.9999	-1.7655	0.1120	-1.7641	0.1362
spas	Dspastin	1628019_a_at	-0.2264	0.5982	0.8685	0.0713	1.3036	0.0029	0.0334	0.9774	-0.4809	0.0965	-0.5143	0.0513	-0.3129	0.8326	0.6013	0.3152	0.9141	0.1612
CG15358	CG15358	1628020_at	-0.1404	0.7948	-0.2315	0.8599	0.9094	0.0936	1.5307	0.0674	0.6691	0.1650	-0.8616	0.0514	-0.0246	0.9970	0.3395	0.8804	0.3641	0.8600
CG17364	CG17364	1628021_at	0.0928	0.6094	0.0427	0.6973	0.1078	0.4484	0.1234	0.7535	0.0313	0.8920	-0.0921	0.5874	0.1672	0.7707	0.0911	0.7544	-0.0761	0.7964
---	---	1628022_at	0.0373	0.8836	0.0475	0.7492	-0.0208	0.9344	-0.0361	0.9749	0.1095	0.7414	0.1456	0.6082	0.0744	0.9195	0.1881	0.4531	0.1137	0.6795
---	---	1628023_at	0.1148	0.6036	-0.0696	0.7188	0.0911	0.6075	0.0004	0.9996	0.1125	0.6523	0.1121	0.6184	-0.0058	0.9946	-0.0186	0.9528	-0.0128	0.9646
---	---	1628024_at	0.0677	0.6735	0.2061	0.1632	0.1252	0.5677	-0.1276	0.8417	-0.2179	0.3695	-0.0903	0.7214	-0.0129	0.9914	0.0358	0.9289	0.0487	0.8893
CG13055	CG13055	1628025_at	0.2982	0.2789	0.2904	0.0811	0.0769	0.8000	-0.1215	0.7658	-0.1135	0.5440	0.0080	0.9708	0.0961	0.9167	0.0087	0.9903	-0.0873	0.8256
---	---	1628026_at	0.3611	0.1427	0.2007	0.4106	0.1708	0.4464	-0.0091	0.9943	0.0164	0.9699	0.0255	0.9413	0.0180	0.9914	0.0142	0.9848	-0.0038	0.9955
PyK	pyruvate kinase	1628027_a_at	-0.3219	0.0564	0.3079	0.4360	0.8029	0.0006	0.1185	0.8025	-0.8550	0.0016	-0.9735	0.0006	-0.4353	0.6557	-0.2728	0.4976	0.1625	0.7134
RpL17	Ribosomal protein	1628028_s_at	0.2972	0.0830	0.5012	0.0398	0.7419	0.0042	0.2165	0.5932	-0.1647	0.4400	-0.3812	0.0440	-0.0030	0.9977	-0.0635	0.8382	-0.0605	0.8372
CG14480	CG14480	1628029_at	0.2348	0.3206	0.0784	0.7938	0.0558	0.8024	0.1167	0.8578	0.3810	0.1035	0.2643	0.2072	0.0909	0.9309	0.1394	0.7220	0.0485	0.9181
CG18808	CG18808	1628030_at	0.1516	0.4927	0.0480	0.8058	0.4316	0.0632	0.1734	0.6584	0.0686	0.7613	-0.1048	0.5769	-0.2149	0.7181	-0.1595	0.5338	0.0554	0.8653
---	---	1628031_at	0.1239	0.5375	0.0107	0.9515	-0.1106	0.6008	-0.1842	0.7143	-0.1372	0.5825	0.0470	0.8605	0.1321	0.8193	-0.0206	0.9587	-0.1527	0.5416
LpR2	a2M-receptor-like	1628032_a_at	0.5427	0.2986	-0.1390	0.6935	-0.8814	0.1622	-0.4297	0.2959	0.3368	0.1613	0.7665	0.0043	0.1927	0.9619	-0.3542	0.8065	-0.5470	0.6533
---	---	1628033_at	0.2156	0.1589	0.0909	0.5833	0.1021	0.4814	0.0883	0.8405	0.0821	0.6568	-0.0062	0.9761	-0.0402	0.9589	-0.0186	0.9590	0.0216	0.9435
CG3162	CG3162	1628034_at	0.2434	0.1251	0.2089	0.2168	0.0722	0.7282	0.0612	0.9017	0.0328	0.8786	-0.0284	0.8815	0.0538	0.9515	0.1141	0.7046	0.0603	0.8636
CG18536	CG18536	1628035_at	0.2014	0.2136	0.1911	0.4274	-0.1955	0.2935	-0.2492	0.5933	0.0214	0.9517	0.2706	0.2008	0.0968	0.8909	-0.0847	0.8037	-0.1815	0.5123
hdm	hold'em	1628036_at	0.1192	0.5604	0.0097	0.9659	0.1443	0.4596	0.0251	0.9677	0.0717	0.7034	0.0466	0.7992	0.0297	0.9737	-0.0039	0.9938	-0.0336	0.9121
CG32533	CG32533	1628037_at	-0.4523	0.4762	0.1595	0.8537	0.1589	0.5121	0.2594	0.6496	-0.0475	0.8999	-0.3069	0.2132	0.2022	0.9653	0.5569	0.6894	0.3547	0.8190
CG13577	CG13577	1628038_at	-0.0047	0.9816	0.1957	0.3430	0.2318	0.1235	0.0501	0.9313	-0.0618	0.7639	-0.1118	0.4970	0.0140	0.9862	0.0529	0.8331	0.0389	0.8788
---	---	1628039_at	0.1349	0.4188	-0.0146	0.9091	0.0325	0.8973	0.0802	0.8903	0.1140	0.5907	0.0338	0.8834	-0.0556	0.9503	-0.0085	0.9875	0.0472	0.8972
CG10993	CG10993	1628040_at	-0.6152	0.0665	-0.6167	0.1166	-0.4512	0.0243	0.1517	0.7431	0.1089	0.6309	-0.0428	0.8569	-0.1419	0.8461	-0.1130	0.7509	0.0290	0.9449
Dek	Dek	1628041_a_at	0.3704	0.1174	0.7187	0.0458	0.2780	0.0966	-0.2044	0.6010	0.1026	0.6424	0.3070	0.0825	0.1068	0.9238	0.4071	0.2571	0.3003	0.4323
CG32488	CG32488	1628042_at	0.1020	0.6901	0.3359	0.0716	0.3463	0.0423	-0.0878	0.9036	-0.1378	0.5901	-0.0499	0.8552	0.0132	0.9923	0.1225	0.7424	0.1094	0.7674
---	---	1628043_s_at	-0.0904	0.5778	-0.1075	0.7339	0.1570	0.4165	0.0081	0.9934	0.1471	0.4630	0.1390	0.4420	-0.1779	0.8215	-0.0326	0.9506	0.1454	0.6881
CG10214	CG10214	1628044_at	0.3681	0.1656	0.2968	0.6023	0.1073	0.6900	0.0598	0.9518	0.1602	0.5955	0.1004	0.7360	0.1813	0.9238	0.1054	0.9139	-0.0759	0.9316
DmirCG32227 /// gogo	golden goal /// CG	1628045_s_at	0.0146	0.9699	0.1417	0.5189	0.0466	0.7776	0.0059	0.9956	0.0835	0.7399	0.0776	0.7355	0.0850	0.9491	0.0922	0.8690	0.0072	0.9921
CG15828	CG15828	1628049_at	2.4992	0.0025	1.1031	0.0765	2.4080	0.0005	0.7420	0.2536	1.6955	0.0018	0.9535	0.0136	-0.6241	0.7305	0.2124	0.8384	0.8365	0.2960
---	---	1628050_at	0.1244	0.4164	0.1069	0.4827	0.1035	0.6763	0.1142	0.8033	0.0150	0.9560	-0.0992	0.5888	-0.0985	0.8940	-0.0661	0.8628	0.0324	0.9330
CG30376	CG30376	1628051_at	0.4150	0.2991	0.3024	0.1584	0.1935	0.3968	0.0383	0.9603	-0.0321	0.9096	-0.0704	0.7500	0.0088	0.9952	0.0707	0.8979	0.0618	0.9025
Cyp6a17	Cyp6a17	1628052_at	-0.6033	0.1801	-0.9097	0.0596	-0.2687	0.1398	-0.5240	0.0884	-1.4297	0.0002	-0.9057	0.0006	-1.1226	0.3362	-1.7141	0.0285	-0.5915	0.3552
sav	salvador/shar-pei	1628053_at	-0.5454	0.1433	-0.3336	0.6492	-0.1663	0.4121	0.2216	0.4139	0.0583	0.7424	-0.1632	0.2252	-0.0284	0.9939	0.1023	0.9404	0.1306	0.9113
---	---	1628054_at	0.1909	0.3181	-0.0201	0.8582	0.1604	0.5135	0.0306	0.9603	0.0144	0.9531	-0.0162	0.9366	-0.1745	0.8541	-0.1575	0.7263	0.0170	0.9781
Gr58b	Gustatory recepto	1628055_at	0.0669	0.6861	-0.1240	0.3446	0.1371	0.4674	-0.0044	0.9956	0.0603	0.8128	0.0646	0.7772	-0.1198	0.8270	-0.1056	0.6793	0.0142	0.9673
---	---	1628056_at	0.1321	0.4893	0.3054	0.0435	0.2223	0.2154	0.1084	0.7576	0.0735	0.6685	-0.0350	0.8407	0.2243	0.7749	0.0547	0.9170	-0.1697	0.6419
CG18616	CG18616	1628057_at	-0.2478	0.2025	-0.2090	0.5613	-0.1308	0.4914	-0.0126	0.9838	-0.0771	0.6367	-0.0645	0.6710	-0.0734	0.9677	-0.1416	0.8218	-0.0682	0.9182
---	---	1628058_at	0.3297	0.1024	0.1606	0.5895	0.2788	0.0982	-0.0712	0.9165	0.0381	0.8944	0.1093	0.6083	-0.0986	0.8861	-0.1216	0.6711	-0.0230	0.9493
CG31270	CG31270	1628059_at	0.0239	0.9043	0.0361	0.8431	0.0946	0.6183	0.0387	0.9635	0.0972	0.7061	0.0586	0.8172	-0.0706	0.9155	-0.0664	0.8287	0.0042	0.9921
---	---	1628060_at	0.0810	0.7171	0.0689	0.6202	0.0566	0.7269	0.1441	0.7202	0.0339	0.8906	-0.1103	0.5345	0.1726	0.8191	0.0320	0.9494	-0.1406	0.6777
RpL37a	Ribosomal protein	1628061_at	0.5713	0.0444	1.0819	0.0231	1.2354	0.0019	0.1427	0.7760	-0.5020	0.0248	-0.6447	0.0051	0.1999	0.8270	-0.0428	0.9451	-0.2427	0.5499
CG34048	CG34048	1628062_at	-0.1612	0.4681	-0.0325	0.8763	-0.2947	0.0651	-0.0266	0.9745	-0.0039	0.9899	0.0227	0.9279	0.1038	0.8270	0.0041	0.9925	-0.0997	0.6366
dpr6	dpr6	1628063_at	0.5969	0.0107	0.4237	0.2116	0.2888	0.1969	-0.1012	0.8942	0.1412	0.6126	0.2423	0.2898	-0.0469	0.9611	-0.0051	0.9935	0.0418	0.9109
---	---	1628064_at	0.1170	0.5240	0.0259	0.8069	0.0380	0.8625	0.0627	0.8915	0.1369	0.3870	0.0741	0.6399	0.0064	0.9963	0.0364	0.9402	0.0301	0.9431

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Ravus	Ravus	1628065_at	-0.1642	0.4133	-0.5096	0.0854	0.2368	0.2589	0.1955	0.6627	0.1863	0.3980	-0.0093	0.9724	-0.4710	0.3985	-0.1668	0.6079	0.3041	0.3277
---	---	1628066_at	0.1666	0.3109	0.0297	0.8287	0.0700	0.6552	0.1185	0.7425	0.2072	0.1893	0.0887	0.5738	-0.0056	0.9946	-0.0521	0.8380	-0.0465	0.8488
dyl	dusky-like	1628067_s_at	0.1989	0.2636	-0.2155	0.3735	-0.0152	0.9408	0.1108	0.7845	0.1757	0.2973	0.0649	0.7144	-0.0699	0.9467	-0.1829	0.5896	-0.1130	0.7595
---	---	1628068_s_at	0.1789	0.2482	-0.1092	0.5040	0.2605	0.0962	0.1348	0.7011	0.2421	0.1319	0.1073	0.4860	-0.1530	0.7779	-0.0607	0.8501	0.0923	0.7333
---	---	1628069_at	0.2190	0.2482	-0.1108	0.6084	0.1007	0.5799	0.1563	0.7589	0.3164	0.1463	0.1601	0.4365	-0.0736	0.9306	-0.1119	0.7200	-0.0383	0.9192
CG14541	CG14541	1628070_at	0.1730	0.3193	0.0408	0.8086	-0.0409	0.8254	0.1204	0.8422	0.1089	0.6703	-0.0115	0.9679	0.0965	0.8825	0.0259	0.9492	-0.0706	0.8204
---	---	1628071_at	0.1753	0.2520	-0.0446	0.7762	-0.0376	0.8730	0.0658	0.9380	0.0569	0.8616	-0.0090	0.9772	-0.0786	0.8999	-0.0713	0.8117	0.0073	0.9847
CG12993	CG12993	1628072_at	0.2080	0.3413	-0.0104	0.9220	0.0806	0.6803	0.0317	0.9732	0.0930	0.7345	0.0613	0.8165	0.1854	0.7100	0.0293	0.9291	-0.1561	0.4686
---	---	1628073_s_at	0.1967	0.2585	0.0158	0.8915	-0.0637	0.8219	0.1911	0.6888	0.3858	0.0826	0.1947	0.3362	0.1362	0.8903	0.0327	0.9578	-0.1035	0.8243
---	---	1628074_at	-0.0089	0.9625	0.0993	0.4330	0.0651	0.7271	0.0804	0.8405	0.0304	0.8786	-0.0500	0.7595	0.0602	0.9405	0.0892	0.7692	0.0290	0.9342
olf186-M	olf186-M	1628075_at	1.1691	0.0024	-0.1375	0.4818	0.2465	0.1439	0.3270	0.2806	1.1970	0.0004	0.8700	0.0007	0.0152	0.9923	-0.0876	0.8629	-0.1028	0.8209
Lip3	Lip3	1628076_at	0.1065	0.5048	0.3641	0.0478	0.0390	0.8302	-0.1033	0.7658	-0.2013	0.1710	-0.0980	0.4881	0.1547	0.7956	0.0526	0.8850	-0.1021	0.7202
---	---	1628077_at	-0.0257	0.9717	-0.1103	0.7570	-0.7537	0.0222	-0.2003	0.7081	0.2595	0.2921	0.4598	0.0414	0.3203	0.8905	0.1459	0.9144	-0.1744	0.8846
CG2256	CG2256	1628078_at	0.0237	0.8982	0.0948	0.5057	0.2436	0.1667	0.1949	0.5541	0.0356	0.8747	-0.1592	0.3028	0.0114	0.9884	0.0555	0.7972	0.0441	0.8393
CG5458	CG5458	1628079_at	0.1981	0.4860	-0.1281	0.6295	0.3397	0.2373	0.3857	0.5117	0.3274	0.2960	-0.0583	0.8738	-0.1581	0.8814	0.0627	0.9205	0.2209	0.6124
GIIIspla2	CG1583	1628080_at	0.7109	0.0329	0.1771	0.7901	0.3974	0.1883	-0.0576	0.9422	0.3882	0.0865	0.4457	0.0342	-0.1621	0.9514	-0.1521	0.8963	0.0099	0.9945
CG7530	CG7530	1628081_s_at	0.2379	0.3091	-0.1728	0.6104	0.0956	0.8127	0.0324	0.9807	0.2861	0.3937	0.2537	0.4052	-0.2741	0.8305	-0.0123	0.9924	0.2617	0.6498
---	---	1628082_at	0.0763	0.7171	0.1785	0.2260	0.3309	0.0818	0.1006	0.8803	-0.1881	0.4260	-0.2886	0.1563	0.0230	0.9848	0.0928	0.7808	0.0698	0.8394
CG5036	CG5036	1628083_at	-2.4627	0.0008	-2.9890	0.0027	-3.2222	0.0000	-0.2901	0.7031	0.2285	0.5441	0.5186	0.0972	-0.0336	0.9848	-0.3752	0.3259	-0.3416	0.3921
BobA	Brother of Bearde	1628084_at	0.0400	0.9267	0.2289	0.3586	-0.1239	0.4706	-0.0922	0.9257	-0.1144	0.7465	-0.0222	0.9534	0.2140	0.7644	0.0867	0.8280	-0.1273	0.7075
CG15756	CG15756	1628085_at	-0.0103	0.9667	0.1185	0.4034	0.3663	0.0224	0.0302	0.9688	-0.0031	0.9913	-0.0334	0.8891	-0.0966	0.8609	0.0896	0.7266	0.1862	0.4093
CG34367	CG13141	1628086_at	0.2067	0.2669	0.1963	0.2724	-0.1174	0.5305	-0.1308	0.6936	-0.0914	0.5829	0.0394	0.8212	0.1967	0.7576	0.0589	0.8782	-0.1378	0.6328
CG10200 /// DsimCG10200	CG10200	1628087_s_at	-2.1890	0.0514	-1.0107	0.0126	-2.5409	0.0010	-1.1920	0.1293	-2.0038	0.0020	-0.8118	0.0585	-0.0105	0.9990	-1.1236	0.3651	-1.1130	0.3878
snRNP69D	small nuclear ribo	1628088_at	0.1731	0.3970	0.4818	0.1526	0.5893	0.0583	-0.1351	0.7146	-0.3290	0.0556	-0.1939	0.1974	-0.1689	0.9142	0.2308	0.7127	0.3998	0.4848
ham	hamlet	1628089_at	-1.5753	0.0031	-0.2617	0.3638	-1.2053	0.0005	-0.6799	0.1341	-1.1025	0.0026	-0.4225	0.0856	-0.0033	0.9984	-0.3439	0.2650	-0.3407	0.2993
---	---	1628090_at	0.0773	0.7099	-0.0659	0.5608	0.0302	0.9180	-0.1886	0.6070	-0.0829	0.6952	0.1056	0.5617	-0.0522	0.9619	0.0073	0.9923	0.0595	0.8877
CG8060	CG8060	1628091_at	-0.2523	0.3199	0.7294	0.0672	1.2021	0.0070	0.0053	0.9956	-0.4763	0.0621	-0.4816	0.0388	-0.2664	0.8454	0.5094	0.3519	0.7758	0.1910
Tfb4	Tfb4	1628092_at	0.0927	0.4969	0.6465	0.0426	0.7176	0.0069	0.0116	0.9922	-0.2415	0.3278	-0.2531	0.2466	0.0166	0.9898	0.2310	0.3931	0.2144	0.4471
CG14395	CG14395	1628093_at	0.3784	0.3324	0.2517	0.1742	0.7311	0.0014	-0.0408	0.9570	-0.1035	0.6507	-0.0627	0.7815	-0.4202	0.7510	-0.3889	0.5075	0.0313	0.9716
CG17639	CG17639	1628094_at	0.4742	0.3244	-0.1242	0.5751	-0.1408	0.6965	-0.0516	0.9740	0.2933	0.4647	0.3449	0.3254	0.1972	0.9156	-0.2713	0.7127	-0.4685	0.4852
CG15370	CG15370	1628095_at	-0.1773	0.3420	0.1633	0.3173	0.0079	0.9806	0.1315	0.7970	0.1003	0.6714	-0.0312	0.9003	0.1291	0.8882	0.2507	0.4732	0.1216	0.7597
CG3995	CG3995	1628096_at	0.2618	0.1798	0.2026	0.3874	0.2317	0.1343	0.0581	0.9204	0.1845	0.3014	0.1265	0.4482	-0.0295	0.9816	0.1031	0.7843	0.1325	0.6954
elk	eag-like K[+] chan	1628097_at	0.3605	0.1827	0.1546	0.2516	0.0992	0.6412	-0.1223	0.8356	-0.0256	0.9352	0.0967	0.6778	-0.0123	0.9933	-0.1442	0.7067	-0.1319	0.7352
---	---	1628098_at	0.0303	0.8956	-0.0619	0.7647	-0.1100	0.5919	-0.0679	0.9037	-0.0124	0.9644	0.0554	0.7860	-0.0820	0.9064	-0.0268	0.9476	0.0552	0.8680
bor	belphegor	1628099_at	0.5597	0.0493	-0.2884	0.1889	-0.0091	0.9796	0.1606	0.8231	0.8375	0.0093	0.6769	0.0146	-0.1617	0.9016	0.0392	0.9609	0.2008	0.7099
CG8957	CG8957	1628100_at	-0.1312	0.7145	0.1549	0.5696	0.2110	0.1497	-0.0304	0.9810	-0.2978	0.3435	-0.2674	0.3457	-0.0099	0.9929	-0.0331	0.9389	-0.0232	0.9505
CG32581 /// CG8974	CG32581 /// CG8	1628101_s_at	-0.5151	0.1788	1.5234	0.0089	1.4350	0.0017	0.0193	0.9926	-1.8820	0.0018	-1.9013	0.0010	0.0455	0.9831	0.0417	0.9590	-0.0039	0.9965
CG32573	CG32573	1628102_a_at	0.1833	0.3242	-0.0644	0.7636	0.2033	0.3831	0.1916	0.5854	0.2966	0.0977	0.1050	0.5504	0.0312	0.9848	0.1633	0.6876	0.1321	0.7524
CG31496	CG31496	1628103_at	-0.6204	0.0255	-1.0609	0.0073	-0.8920	0.0030	0.0784	0.9314	0.3416	0.2068	0.2632	0.2834	0.0107	0.9914	0.0031	0.9944	-0.0076	0.9835
---	---	1628104_at	0.1139	0.5755	0.0207	0.8480	-0.1033	0.6126	0.0419	0.9580	0.1221	0.5942	0.0802	0.7204	0.3265	0.6960	0.1566	0.6949	-0.1698	0.6580
Prm	paramyosin	1628105_a_at	-1.8005	0.0134	-1.9295	0.0057	-2.6390	0.0001	-0.2204	0.5628	-0.4851	0.0213	-0.2646	0.1282	0.4764	0.8283	-0.5862	0.5416	-1.0627	0.2655
CG12034	CG12034	1628106_at	0.6826	0.0099	0.5360	0.1004	0.7383	0.0019	-0.0115	0.9866	0.1773	0.2943	0.1888	0.2075	-0.1703	0.8379	0.0903	0.8461	0.2606	0.4620
CG3513	CG3513	1628107_at	0.3915	0.0327	0.1841	0.4979	0.3524	0.1210	0.1913	0.7130	-0.2838	0.2300	-0.4752	0.0322	0.0267	0.9862	-0.2244	0.5531	-0.2511	0.5041
CG32114	CG32114	1628108_at	0.0720	0.6641	-0.0542	0.6983	0.0794	0.7009	0.0679	0.8836	0.1361	0.4019	0.0682	0.6798	0.0260	0.9774	-0.0762	0.7722	-0.1022	0.6621
CG1737	CG1737	1628109_at	0.4702	0.1290	0.0696	0.8444	-0.3152	0.3009	-0.0844	0.9117	0.7669	0.0074	0.8513	0.0027	0.3695	0.7697	0.2744	0.6380	-0.0951	0.8977
E(Pc)	transcript group III	1628110_s_at	-0.0106	0.9708	-0.1036	0.4811	0.4484	0.0954	-0.1506	0.7016	-0.0058	0.9826	0.1448	0.3870	-0.3684	0.6955	-0.0387	0.9505	0.3297	0.4016
Dip1	Dorsal interacting	1628111_at	0.3511	0.2011	0.6950	0.0586	0.7378	0.0025	0.1772	0.7341	0.1257	0.6259	-0.0515	0.8483	-0.0681	0.9174	0.3811	0.1029	0.4492	0.0852
SMC1	SMC1	1628112_at	-0.1076	0.6871	-0.1887	0.3822	-0.0818	0.6866	0.0210	0.9826	0.3692	0.1009	0.3482	0.0844	-0.0888	0.9235	0.2390	0.4311	0.3278	0.2960
Fili	CG13487	1628113_at	0.0990	0.6587	0.1314	0.4120	0.0598	0.8574	0.0966	0.8764	-0.1118	0.6416	-0.2084	0.2809	0.1512	0.8833	-0.0943	0.8631	-0.2456	0.5535
Pten	Pten	1628114_s_at	-0.3567	0.2908	0.1672	0.6047	-0.2378	0.2354	-0.1325	0.8571	-0.0500	0.8887	0.0826	0.7781	0.1286	0.9221	0.4019	0.3452	0.2733	0.5540
CG6793	CG6793	1628115_at	0.2299	0.2763	-0.0566	0.6494	0.2733	0.1489	0.1831	0.7621	0.0702	0.8320	-0.1129	0.6776	-0.0122	0.9939	-0.1952	0.6200	-0.1830	0.6423
CG14007	CG14007	1628116_at	0.1289	0.4903	-0.0160	0.8776	0.2172	0.1962	0.1357	0.7015	0.0529	0.7901	-0.0828	0.6147	-0.2650	0.6145	-0.0818	0.7603	0.1832	0.4226
Hsp27	heat shock proteir	1628117_at	-0.5098	0.6408	-2.0266	0.0223	-1.8612	0.0334	0.4963	0.8738	2.5900	0.0253	2.0937	0.0381	0.3872	0.9142	1.0731	0.3897	0.6859	0.6150
---	---	1628118_at	0.1238	0.6514	-0.1149	0.4523	0.0649	0.7556	0.0054	0.9956	0.0904	0.7553	0.0850	0.7469	-0.1955	0.7293	-0.0798	0.7932	0.1157	0.6601
---	---	1628119_at	0.1417	0.5054	-0.0236	0.8768	-0.1169	0.6400	-0.0186	0.9790	0.0976	0.6135	0.1161	0.4881	-0.0492	0.9742	-0.1263	0.7898	-0.0772	0.8812
CG18416	CG18416	1628120_at	0.1496	0.3175	-0.1138	0.4129	-0.1331	0.5467	-0.0861	0.8507	-0.0131	0.9578	0.0730	0.6746	-0.1365	0.8283	-0.0670	0.8539	0.0695	0.8354
Osi14	Osiris	1628121_at	-0.9896	0.0198	-1.3690	0.0074	-0.7226	0.0058	0.8519	0.1162	0.8516	0.0174	-0.0003	0.9995	0.0790	0.9142	-0.0608	0.8674	-0.1397	0.6150
CG32631	CG32631	1628122_at	0.0205	0.9346	0.0674	0.6698	0.1148	0.5104	0.0732	0.8856	-0.0098	0.9705	-0.0830	0.6386	-0.0436	0.9604	0.0327	0.9341	0.0763	0.7983
CG4332	CG4332	1628123_at	0.3291	0.1051	0.3684	0.3496	0.2086	0.2530	-0.1432	0.6936	0.1624	0.3426	0.3056	0.0475	0.0852	0.9588	0.2696	0.5778	0.1844	0.7229
CG3032	CG3032	1628124_at	-0.0669	0.7790	-0.3520	0.1546	-0.2569	0.2151	0.1533	0.7556	0.5495	0.0179	0.3963	0.0431	0.1200	0.8700	0.2431	0.4021	0.1231	0.7087
Doc2	Dorsocross2	1628125_at	0.0564	0.8125	-0.0829	0.6399	-0.0222	0.9025	0.0298	0.9715	-0.0656	0.7999	-0.0954	0.6630	-0.0499	0.9646	-0.1172	0.7424	-0.0673	0.8672
CG31202	CG31202	1628126_at	0.2214	0.4338	0.0859	0.6302	0.5120	0.0344	0.1082	0.8337	0.0999	0.6502	-0.0083	0.9727	-0.1780	0.8541	-0.0121	0.9885	0.1658	0.7134
CG9926 /// DyakCG9926	CG9926	1628127_at	0.3111	0.1799	-0.1385	0.7168	-0.1637	0.7973	-0.1504	0.8485	0.4054	0.1570	0.5558	0.0374	-0.2743	0.8740	0.1045	0.9216	0.3788	0.6043
CG6333	CG6333	1628128_at	0.2563	0.1002	0.0028	0.9904	0.4026	0.0461	0.0932	0.9196	0.1015	0.7646	0.0083	0.9812	-0.1959	0.7547	-0.1861	0.4958	0.0097	0.9835
Sdic3 /// Sdic4 /// sw	dynein /// Sdic:CG	1628129_a_at	-0.6829	0.1370	-0.2393	0.7122	-0.4577	0.1516	-0.0719	0.9311	-0.4220	0.0857	-0.3501	0.1086	0.1842	0.9449	0.1145	0.9279	-0.0697	0.9512
CG31771	CG31771	1628130_at	0.3776	0.0344	0.0788	0.5322	0.3212	0.0493	0.0789	0.8707	0.0315	0.8907	-0.0474	0.8039	-0.2444	0.6868	-0.2500	0.3000	-0.0055	0.9889
RpIII128	RNA polymerase	1628131_at	0.3946	0.0401	-0.0877	0.8615	0.2183	0.3099	0.0749	0.9413	0.6097	0.0430	0.5347	0.0461	-0.1925	0.8692	0.1419	0.8127	0.3343	0.4850
CG7099	CG7099	1628132_at	0.4311	0.0230	0.2706	0.3798	0.5840	0.0128	0.3573	0.2851	0.3533	0.0748	-0.0040	0.9872	-0.0401	0.9761	0.2588	0.4198	0.2989	0.3671
CG5204	CG5204	1628133_at	0.2571	0.1620	0.1421	0.4992	-0.1546	0.4297	0.1577	0.7121	0.2093	0.2843	0.0516	0.8135	0.2495	0.7220	0.0513	0.9109	-0.1983	0.5257

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Cad96Cb	Cad96Cb	1628134_at	-0.1658	0.3255	0.1102	0.3969	0.1749	0.3646	-0.0616	0.9113	-0.1208	0.5139	-0.0593	0.7565	-0.1300	0.8714	0.1110	0.7769	0.2410	0.4648
pck	megatrachea	1628135_s_at	-0.6655	0.0081	-0.4666	0.1085	-0.7722	0.0026	-0.0035	0.9956	-0.1657	0.3050	-0.1623	0.2599	0.2742	0.7464	0.1405	0.7354	-0.1337	0.7448
CG15904	CG15904	1628136_at	0.2150	0.2094	0.0049	0.9730	-0.0332	0.9055	-0.1174	0.8550	-0.0935	0.7306	0.0239	0.9341	-0.0137	0.9926	-0.1890	0.5992	-0.1753	0.6299
---	---	1628137_at	0.0364	0.8444	-0.0166	0.8881	0.1231	0.4810	-0.0733	0.8959	-0.1002	0.6265	-0.0269	0.9039	-0.1167	0.8192	-0.0869	0.7165	0.0298	0.9176
CG18166 /// CG18273	CG18166 /// CG1	1628138_s_at	-0.6281	0.2814	0.3753	0.2121	1.2229	0.0030	0.0094	0.9951	-1.0097	0.0062	-1.0190	0.0035	-0.2965	0.7644	0.3061	0.4743	0.6026	0.1764
CG30385	CG30385	1628139_at	-0.7867	0.0166	-2.5306	0.0097	-2.5277	0.0027	0.2579	0.8999	1.3503	0.0480	1.0923	0.0694	0.0184	0.9898	-0.2508	0.3999	-0.2692	0.3820
---	---	1628140_at	0.1571	0.5424	-0.3344	0.2667	-0.2190	0.1549	0.3070	0.6257	0.5542	0.0771	0.2472	0.3930	0.1304	0.8814	-0.0316	0.9531	-0.1620	0.6534
rhea	tendrils	1628141_at	-0.4973	0.0899	-0.8954	0.0295	-1.0966	0.0002	0.0749	0.9030	0.3750	0.0635	0.3002	0.0921	0.2525	0.7741	-0.0127	0.9870	-0.2653	0.5030
CG12986	CG12986	1628142_at	0.2454	0.2811	0.2580	0.6492	0.3805	0.0472	0.0382	0.9777	0.0464	0.9214	0.0082	0.9849	-0.0048	0.9970	0.1095	0.7730	0.1143	0.7495
rg	Drosophila A kina	1628143_a_at	0.3001	0.7044	-1.6981	0.0109	-1.5449	0.0229	0.4830	0.7492	2.7415	0.0022	2.2585	0.0030	0.4030	0.8882	0.6972	0.5363	0.2942	0.8287
CG14273	CG14273	1628144_at	-0.0788	0.7667	-0.1292	0.4227	-0.1394	0.4095	0.0964	0.8578	0.0185	0.9492	-0.0779	0.7025	-0.0757	0.9246	-0.2039	0.4475	-0.1283	0.6589
CG12361	CG12361	1628145_at	-1.6236	0.0008	0.0400	0.7793	-0.1959	0.5485	-0.4161	0.4338	-1.6078	0.0007	-1.1917	0.0015	-0.0960	0.9076	-0.1509	0.6361	-0.0549	0.8919
crb	Crumbs	1628146_at	-1.0796	0.0136	-1.2431	0.0195	-1.5496	0.0008	0.1342	0.9011	0.5726	0.1021	0.4384	0.1608	0.4302	0.7230	0.3458	0.5246	-0.0844	0.9070
neuroligin	neuroligin	1628147_at	0.0334	0.8418	-0.0610	0.5579	0.0232	0.9205	-0.0112	0.9873	0.0955	0.6015	0.1067	0.5079	-0.1326	0.8395	-0.0795	0.8216	0.0531	0.8864
unc-119	unc-119	1628148_at	0.2200	0.1719	0.2793	0.3923	0.3653	0.1455	-0.0316	0.9760	0.1034	0.7345	0.1350	0.6062	0.1065	0.9260	0.2081	0.6093	0.1016	0.8323
Pak3	Pak3	1628149_a_at	0.4620	0.0313	-0.0513	0.8219	-0.1172	0.6436	0.2281	0.5453	1.0155	0.0009	0.7875	0.0015	0.2962	0.7644	0.5216	0.1978	0.2253	0.6171
CG9449	CG9449	1628150_a_at	1.6958	0.0041	0.8777	0.0959	2.1671	0.0000	-0.0773	0.9518	-0.4533	0.1954	-0.3760	0.2326	-0.8244	0.6483	-0.7252	0.3066	0.0992	0.9211
---	---	1628151_at	-0.1450	0.4976	-0.1024	0.4737	-0.0394	0.8761	0.0809	0.8889	0.0389	0.8809	-0.0420	0.8525	-0.0572	0.9521	0.0489	0.9111	0.1061	0.7484
Clc	clathrin	1628152_at	-0.2534	0.1511	0.1928	0.2429	0.0982	0.5419	0.0795	0.8544	-0.0525	0.7820	-0.1320	0.3589	0.1483	0.7997	0.3540	0.1397	0.2057	0.4057
---	---	1628153_at	0.0934	0.6965	0.0007	0.9987	0.1032	0.6408	0.1076	0.8959	0.1206	0.7008	0.0129	0.9697	-0.0583	0.9374	-0.0826	0.7707	-0.0243	0.9410
CG33492	CG33492	1628154_at	0.0936	0.7484	0.2008	0.1255	0.2902	0.0603	-0.2199	0.6755	-0.2264	0.3695	-0.0065	0.9836	-0.1514	0.8202	0.0024	0.9978	0.1538	0.6040
klar	marbles	1628155_at	-0.2335	0.6576	-0.8944	0.0751	-0.9981	0.0022	0.1938	0.7906	0.5180	0.0830	0.3242	0.2240	0.0815	0.9814	0.0124	0.9941	-0.0691	0.9471
---	---	1628156_at	0.1521	0.3120	-0.2369	0.3705	-0.3475	0.2167	0.0489	0.9666	0.5081	0.0962	0.4592	0.0923	0.0004	0.9999	-0.0489	0.9291	-0.0494	0.9183
---	---	1628157_at	0.0054	0.9801	-0.1077	0.4901	-0.0207	0.9231	0.1091	0.8467	0.2303	0.2674	0.1212	0.5506	-0.0030	0.9967	0.0645	0.7678	0.0675	0.7447
---	---	1628158_at	0.1561	0.4371	-0.0160	0.9228	0.0024	0.9943	0.1296	0.8479	0.0204	0.9556	-0.1092	0.6717	0.0481	0.9538	-0.0482	0.8954	-0.0964	0.7323
CG32206	CG32206	1628159_a_at	-0.1930	0.1922	-0.0448	0.6599	-0.1041	0.5470	-0.1350	0.7647	-0.0488	0.8449	0.0862	0.6666	-0.0375	0.9619	0.0384	0.9085	0.0760	0.7663
dom	domino	1628160_a_at	0.6325	0.2747	0.4551	0.3822	0.2493	0.1636	-0.0584	0.9120	0.5867	0.0047	0.6451	0.0019	0.1293	0.9689	0.4012	0.6789	0.2719	0.7963
Nup62	Nucleoporin	1628161_at	0.7699	0.0134	0.2702	0.3609	0.3963	0.1367	0.0683	0.9036	0.1384	0.4692	0.0701	0.7186	0.1471	0.9296	-0.1193	0.8775	-0.2663	0.6471
CG9757	CG9757	1628162_at	0.1675	0.3149	0.1637	0.5413	0.2075	0.3620	0.0098	0.9917	0.1276	0.5455	0.1178	0.5388	0.1306	0.8814	0.1964	0.5778	0.0658	0.8861
---	---	1628163_at	0.1297	0.6503	0.1099	0.5347	0.1949	0.3294	0.1062	0.8244	0.0727	0.7396	-0.0336	0.8786	0.0846	0.9400	0.1093	0.8055	0.0247	0.9613
Pp1-13C	Protein phosphata	1628164_at	-0.1686	0.3138	-0.0341	0.7194	-0.0587	0.7799	-0.0364	0.9641	-0.1076	0.6569	-0.0711	0.7621	-0.0492	0.9365	0.1210	0.5534	0.1702	0.3921
---	---	1628165_at	0.1972	0.3736	0.0209	0.8458	0.1883	0.2460	0.1047	0.7351	0.0224	0.9068	-0.0823	0.5402	0.0955	0.9024	-0.1412	0.6478	-0.2367	0.4114
DmsR-2	Dromyosuppressi	1628166_a_at	-0.1194	0.6035	-0.0157	0.8799	0.0911	0.5857	-0.1169	0.7278	-0.2795	0.0664	-0.1625	0.2275	-0.2002	0.7305	-0.2892	0.2289	-0.0890	0.7567
CG6597	CG6597	1628167_s_at	-0.3482	0.1116	-0.5209	0.0216	-0.7644	0.0024	-0.0682	0.8841	-0.0575	0.7639	0.0107	0.9582	0.0826	0.9491	-0.2027	0.6270	-0.2854	0.4755
CG5665	CG5665	1628168_at	-0.0058	0.9841	0.1510	0.4542	0.1651	0.4367	-0.0043	0.9956	-0.1915	0.4747	-0.1871	0.4371	0.0465	0.9514	0.0998	0.6977	0.0533	0.8577
mRpS22	mitochondrial ribo	1628169_at	-0.4715	0.0916	0.5632	0.0498	0.6991	0.0012	0.0463	0.9468	-1.0261	0.0009	-1.0724	0.0004	-0.1727	0.8461	-0.0279	0.9631	0.1448	0.7287
---	---	1628170_at	0.1448	0.4748	0.1104	0.5258	0.0025	0.9927	-0.0626	0.8791	-0.1338	0.3490	-0.0712	0.6170	0.1038	0.9056	-0.1757	0.6003	-0.2796	0.3838
CG13504	CG13504	1628171_at	0.0751	0.7061	0.2640	0.2911	0.2557	0.1530	0.0248	0.9777	-0.2131	0.3616	-0.2379	0.2467	-0.0278	0.9862	0.0155	0.9821	0.0433	0.9329
msps	MINISPINDLES	1628172_at	0.0344	0.8589	-0.7910	0.0120	-0.8178	0.0039	-0.1210	0.8299	0.5215	0.0238	0.6425	0.0058	-0.1360	0.8049	-0.2992	0.1798	-0.1632	0.4934
CG17189	CG17189	1628173_at	0.4997	0.5130	-0.1889	0.2072	-0.2132	0.2047	-0.1391	0.8640	0.0241	0.9553	0.1632	0.5687	-0.1705	0.9514	-0.6099	0.4698	-0.4395	0.6233
CG33119	CG33119	1628174_at	-1.0139	0.0452	-0.7611	0.2138	-0.5351	0.0391	-0.2735	0.5150	-0.3705	0.0955	-0.0970	0.6761	-0.7181	0.7095	-0.3140	0.7403	0.4041	0.6414
CG6325	CG6325	1628175_at	1.1011	0.0245	0.4222	0.0615	0.4197	0.0367	-0.1914	0.6662	0.3831	0.0729	0.5745	0.0088	0.0327	0.9884	-0.1106	0.8748	-0.1432	0.8138
---	---	1628176_at	-0.0042	0.9868	-0.0664	0.7072	0.0494	0.7748	-0.1505	0.7401	-0.1014	0.6521	0.0491	0.8295	-0.2375	0.7386	-0.1741	0.5855	0.0634	0.8769
jhamt	juvenile hormone	1628177_at	-3.6483	0.0505	-4.1849	0.0116	-3.6215	0.0001	0.0963	0.9873	2.8695	0.0514	2.7732	0.0380	-0.2197	0.9644	2.6649	0.0654	2.8846	0.0679
CG4793	CG4793	1628178_a_at	0.0310	0.9023	-0.0346	0.7306	0.0642	0.7834	0.1394	0.8049	0.0480	0.8723	-0.0914	0.6999	-0.0559	0.9457	-0.0455	0.9046	0.0104	0.9794
CG12589	CG12589	1628179_at	0.1997	0.1781	0.0989	0.5227	0.0332	0.8785	0.1155	0.7647	0.1628	0.3297	0.0474	0.7981	0.2734	0.6927	0.1256	0.6949	-0.1479	0.6296
CG12909	CG12909	1628180_at	0.1789	0.7145	-0.0763	0.8571	0.5745	0.1787	0.3549	0.7121	0.5103	0.2445	0.1554	0.7417	-0.2547	0.8909	0.3345	0.6655	0.5892	0.4090
---	---	1628181_at	0.0169	0.9298	0.0344	0.8465	0.0617	0.7354	0.1094	0.7915	-0.0368	0.8691	-0.1461	0.3441	-0.0972	0.8464	-0.1020	0.6512	-0.0049	0.9880
CG13785	CG13785	1628182_at	0.1856	0.3992	0.1413	0.3887	-0.0686	0.6751	-0.0633	0.9297	-0.1253	0.5829	-0.0620	0.7926	0.1215	0.8202	-0.0540	0.8614	-0.1755	0.4413
---	---	1628183_at	-0.2212	0.4123	-0.2622	0.5292	-0.5526	0.1077	-0.2095	0.8791	0.2398	0.6505	0.4494	0.2912	0.0573	0.9779	0.2129	0.7032	0.1556	0.7940
Tsp3A	Tetraspanin 3A	1628184_at	-0.0317	0.9482	0.5428	0.1133	1.3055	0.0023	0.0351	0.9790	-0.5777	0.0823	-0.6129	0.0443	-0.2887	0.8342	0.2322	0.7260	0.5209	0.3766
CG32121	CG32121	1628185_at	-0.2272	0.2299	-0.0745	0.6592	0.0131	0.9688	-0.2329	0.6506	-0.3864	0.1183	-0.1534	0.5255	-0.1383	0.8464	-0.0899	0.8153	0.0485	0.9057
CG11165	CG11165	1628186_at	0.0667	0.9155	-0.9225	0.3503	-1.0703	0.0729	-0.1434	0.9455	1.0421	0.0821	1.1856	0.0335	-0.0145	0.9978	0.0585	0.9788	0.0730	0.9688
CG9691	CG9691	1628187_s_at	0.3004	0.5314	0.7617	0.3490	0.5260	0.1748	-0.9318	0.1194	-2.2189	0.0004	-1.2871	0.0026	-0.4863	0.8650	-1.6113	0.1489	-1.1250	0.3359
---	---	1628188_at	1.0977	0.1642	0.0435	0.8828	0.2308	0.4389	0.2777	0.8942	1.4607	0.0418	1.1831	0.0606	0.1439	0.9340	0.3018	0.6105	0.1579	0.8190
Rep2	Rep2	1628189_at	0.0818	0.5998	-0.1283	0.4935	-0.0697	0.7629	0.1262	0.7018	0.2335	0.1211	0.1073	0.4525	0.0166	0.9900	-0.0056	0.9935	-0.0222	0.9575
---	---	1628190_at	-0.9929	0.0021	-0.2381	0.2499	-0.4339	0.2348	-0.1607	0.8385	-0.3933	0.1811	-0.2326	0.3978	-0.1174	0.9400	0.0184	0.9841	0.1358	0.8196
CG17199	CG17199	1628191_at	0.0853	0.7247	0.0257	0.8777	0.1370	0.4314	0.2984	0.3357	0.1069	0.5825	-0.1915	0.2321	0.1554	0.8243	0.0926	0.8045	-0.0628	0.8729
CG34113	CG34113	1628192_at	0.0923	0.5936	0.1446	0.2894	0.1266	0.4037	0.0743	0.8594	0.0272	0.8948	-0.0471	0.7785	0.0255	0.9764	0.0265	0.9363	0.0011	0.9979
CG34047	CG34047	1628193_s_at	-0.0397	0.8374	-0.1742	0.2053	0.0767	0.6647	0.0941	0.8084	0.1518	0.3327	0.0577	0.7262	-0.1042	0.8488	0.0359	0.9192	0.1401	0.5615
CG2918	CG2918	1628194_at	1.3736	0.0049	1.5315	0.0100	1.3115	0.0004	0.0922	0.9182	0.8755	0.0073	0.7834	0.0072	0.1378	0.9137	1.0446	0.0405	0.9069	0.0778
---	---	1628195_at	0.1939	0.3389	0.0009	0.9994	0.2094	0.3007	0.1273	0.8633	0.1276	0.6691	0.0004	0.9994	-0.1382	0.8215	-0.1155	0.6834	0.0227	0.9491
---	---	1628196_at	0.1848	0.2607	0.0046	0.9885	0.2912	0.1631	0.2638	0.3490	0.0027	0.9903	-0.2611	0.0732	0.0243	0.9831	0.0007	0.9996	-0.0236	0.9507
---	---	1628197_at	1.4297	0.4201	-0.1103	0.3774	0.0589	0.8348	0.3729	0.3147	0.2484	0.2529	-0.1245	0.5612	0.1646	0.9884	-1.2300	0.6446	-1.3945	0.5972
okr	okra	1628198_at	-0.3709	0.1378	-0.4569	0.0550	-0.4860	0.0125	-0.1546	0.6823	0.1859	0.2953	0.3404	0.0373	-0.1844	0.8076	0.0269	0.9590	0.2112	0.5233
CG15345	CG15345	1628199_at	-0.2344	0.3589	-0.2853	0.2157	-0.4462	0.0654	-0.4255	0.4567	0.2310	0.4754	0.6565	0.0258	-0.3194	0.6955	0.0580	0.9111	0.3774	0.2798
---	---	1628200_at	-0.0195	0.9152	0.0828	0.5041	0.2344	0.1303	0.0684	0.8939	-0.1257	0.4810	-0.1941	0.2022	0.0409	0.9667	0.0127	0.9784	-0.0282	0.9382
CG4061	CG4061	1628201_at	-0.4298	0.1018	0.2105	0.3128	0.7920	0.0161	0.3696	0.3166	-0.2055	0.3484	-0.5751	0.0096	-0.0793	0.9515	0.4408	0.2430	0.5201	0.2056
Syx17	syntaxin	1628202_at	-0.1718	0.3335	-0.0671	0.7190	-0.0399	0.8477	-0.1151	0.7785	-0.5582	0.0063	-0.4431	0.0108	-0.1619	0.8740	-0.4765	0.2181	-0.3146	0.4457

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15476	CG15476	1628203_at	0.0943	0.6599	0.1963	0.2462	0.0908	0.6127	-0.0316	0.9753	-0.0270	0.9398	0.0046	0.9880	0.1391	0.8042	0.1143	0.6519	-0.0248	0.9394
CG8270	CG8270	1628204_at	0.0371	0.9196	0.0877	0.8400	0.1031	0.5266	-0.0814	0.8855	0.0033	0.9903	0.0847	0.6709	-0.1001	0.9441	0.1495	0.7780	0.2496	0.5869
---	---	1628205_at	0.0750	0.7149	0.2525	0.3003	0.0658	0.8039	0.0507	0.9502	0.0649	0.8160	0.0142	0.9597	0.1751	0.8292	0.1830	0.6154	0.0078	0.9884
---	---	1628206_at	0.1719	0.3217	0.5113	0.1056	0.1893	0.3429	-0.1988	0.6869	-0.1077	0.6800	0.0911	0.7065	0.0711	0.9365	0.2006	0.4861	0.1295	0.6746
CG5065	CG5065	1628207_at	0.5158	0.6778	-0.3426	0.5742	-0.5635	0.0156	0.3599	0.6493	0.3019	0.4469	-0.0580	0.8997	0.2752	0.9657	-0.8141	0.6673	-1.0893	0.5488
CG7603	CG7603	1628208_at	-0.2138	0.6261	-0.2343	0.4190	-0.1980	0.2237	-0.0165	0.9860	-0.4099	0.0727	-0.3933	0.0558	-0.1914	0.8940	-0.4902	0.3472	-0.2988	0.6040
resilin /// Vkor	resilin /// Vitamin-I	1628209_at	0.0727	0.7509	-0.0464	0.8318	0.2579	0.1422	0.1553	0.7446	-0.0480	0.8612	-0.2033	0.2822	-0.0635	0.8940	0.0701	0.7349	0.1335	0.4677
gcl	germ cell-less	1628210_at	-0.2907	0.1822	-0.2354	0.0909	-0.5276	0.0443	-0.0314	0.9674	-0.0507	0.8463	-0.0193	0.9386	0.3164	0.6832	0.1024	0.8018	-0.2140	0.5232
---	---	1628211_s_at	0.1829	0.5658	0.0050	0.9854	0.2538	0.1381	0.3026	0.4690	0.1302	0.5964	-0.1724	0.4134	0.1059	0.9405	0.0130	0.9887	-0.0929	0.8752
CG17319	CG17319	1628212_at	0.0135	0.9413	0.0255	0.7950	0.1585	0.6020	-0.0921	0.8822	-0.0474	0.8656	0.0447	0.8567	-0.0279	0.9816	0.0927	0.7901	0.1206	0.6978
---	---	1628213_s_at	0.6407	0.0829	1.0887	0.0152	1.0168	0.0016	0.0188	0.9883	-0.3343	0.2677	-0.3531	0.1876	0.0258	0.9848	-0.0184	0.9710	-0.0442	0.9164
CG5989	SAM methionine r	1628214_at	-0.1861	0.3391	-0.1792	0.4276	0.0835	0.6476	0.1754	0.6197	-0.1114	0.5567	-0.2868	0.0708	-0.1067	0.8954	-0.1367	0.6862	-0.0300	0.9434
kuz	lethal (2) c00136	1628215_s_at	-1.1462	0.0089	-0.1364	0.5557	-1.7187	0.0007	0.0896	0.8967	0.3437	0.1294	0.2541	0.2116	1.5180	0.1902	1.3507	0.0454	-0.1673	0.8235
ppan	Peter Pan	1628216_at	-0.0712	0.8769	-0.2212	0.5451	-0.4121	0.0653	0.1847	0.8424	0.5323	0.1211	0.3476	0.2631	0.1733	0.9056	0.2251	0.7040	0.0518	0.9434
---	---	1628217_at	0.0536	0.8095	0.0017	0.9899	0.2976	0.0735	0.0451	0.9584	0.0527	0.8599	0.0077	0.9787	-0.1581	0.8072	-0.1428	0.6231	0.0153	0.9704
---	---	1628218_at	0.0073	0.9717	-0.0106	0.9415	-0.0167	0.9231	-0.1009	0.8172	-0.0551	0.7928	0.0457	0.8130	0.0226	0.9816	0.0637	0.8275	0.0412	0.8924
CG32050	CG32050	1628219_at	-0.0136	0.9358	0.0541	0.6072	0.0689	0.7295	0.0296	0.9704	-0.0269	0.9244	-0.0565	0.8046	0.0182	0.9816	0.0230	0.9341	0.0048	0.9859
---	---	1628220_at	-0.0647	0.8065	0.1209	0.5322	-0.1099	0.5336	-0.0126	0.9860	-0.0624	0.7550	-0.0498	0.7908	0.1584	0.8284	0.0525	0.9111	-0.1059	0.7676
CG17036	CG17036	1628221_at	0.0598	0.7404	-0.1188	0.3973	0.4487	0.0802	0.1216	0.8798	0.2985	0.2816	0.1769	0.5056	-0.2905	0.7220	-0.0212	0.9708	0.2692	0.4542
CG30106	CG30106	1628222_at	-1.0914	0.0371	0.0122	0.9564	-0.1008	0.8008	0.0352	0.9774	-1.0397	0.0054	-1.0749	0.0027	0.1491	0.8774	-0.0337	0.9562	-0.1828	0.6522
---	---	1628223_at	0.3528	0.3086	-0.1479	0.5364	0.1122	0.6797	-0.1189	0.9254	0.2901	0.4667	0.4090	0.2315	-0.0613	0.9816	-0.1117	0.9021	-0.0504	0.9523
E23	Early gene at 23	1628224_a_at	-0.6958	0.1062	-0.8886	0.0258	-1.1436	0.0052	-0.1900	0.7960	0.5662	0.0621	0.7562	0.0123	0.0431	0.9848	0.2958	0.5800	0.2527	0.6410
CG32391	CG32391	1628225_at	0.1428	0.3448	0.0022	0.9891	-0.0106	0.9680	0.0094	0.9922	0.0583	0.8147	0.0489	0.8305	0.0297	0.9717	-0.1171	0.5964	-0.1468	0.4944
CG8539 /// DsimCG8539	CG8539	1628226_at	2.1652	0.0057	2.1663	0.0321	2.4523	0.0022	-0.3135	0.8768	-0.7418	0.2900	-0.4283	0.5268	-0.5913	0.7810	-0.7236	0.4300	-0.1323	0.9171
---	---	1628227_at	-0.0442	0.8283	-0.0955	0.4217	-0.2334	0.1954	-0.1550	0.7753	0.0984	0.7077	0.2534	0.2129	-0.0670	0.9405	-0.0324	0.9420	0.0346	0.9282
CG4218	CG4218	1628228_at	0.0815	0.7096	0.0510	0.6226	0.0826	0.5755	-0.0223	0.9777	-0.0128	0.9670	0.0095	0.9692	0.0906	0.9246	0.0334	0.9492	-0.0572	0.8972
Spn27A	serpin27A	1628229_at	-0.6987	0.0558	0.0901	0.4474	0.0783	0.8354	-0.4408	0.5610	-0.8002	0.0454	-0.3594	0.3094	-0.4348	0.6749	-0.0086	0.9935	0.4262	0.3360
CG31001	CG31001	1628230_x_at	-0.1754	0.4527	-0.9015	0.1801	-0.7041	0.0107	-0.0136	0.9956	0.7725	0.1100	0.7861	0.0719	0.0482	0.9589	-0.0137	0.9763	-0.0619	0.8570
Ac76E	adenylyl cyclase	1628231_at	-1.8531	0.0017	-1.2415	0.0188	-2.1650	0.0006	-1.1429	0.0437	-1.2212	0.0032	-0.0782	0.8300	-0.3507	0.8009	-0.6936	0.2160	-0.3429	0.5781
CG30046	CG30046	1628232_at	-0.0543	0.9059	0.8063	0.2972	1.2364	0.0090	0.0697	0.8950	-0.8818	0.0010	-0.9515	0.0005	-0.3406	0.8940	0.0998	0.9492	0.4403	0.6748
CG31355	CG31355	1628233_at	0.1434	0.6655	-0.1090	0.4209	0.1295	0.5194	0.1630	0.7443	0.0879	0.7328	-0.0751	0.7540	-0.0238	0.9889	-0.1783	0.6503	-0.1545	0.7030
burs	furled wing	1628234_at	0.2511	0.3964	0.1486	0.3957	0.4830	0.0236	0.2684	0.5465	0.0468	0.8789	-0.2215	0.2888	-0.1273	0.9221	-0.0558	0.9353	0.0715	0.9054
CG7203	CG7203	1628235_at	1.1667	0.5715	-0.3358	0.4402	-0.2464	0.3010	-0.0607	0.9866	-0.5438	0.5651	-0.4831	0.5803	0.1444	0.9898	-2.2508	0.3225	-2.3953	0.3149
wntD	wnt inhibitor of Do	1628236_at	0.0183	0.9284	0.1161	0.4901	0.2458	0.1361	0.0359	0.9612	-0.2433	0.2170	-0.2792	0.1129	-0.0178	0.9943	-0.1567	0.8349	-0.1389	0.8471
CG10602	CG10602	1628237_s_at	-0.2243	0.2291	-0.0003	1.0000	-0.2244	0.2352	-0.1873	0.6457	0.0441	0.8656	0.2314	0.1894	-0.0294	0.9869	0.2621	0.5411	0.2915	0.4932
Tektin-C	TEKTIN C	1628238_at	-0.0599	0.7094	0.0597	0.6616	-0.0461	0.7983	0.0006	0.9994	-0.1633	0.3644	-0.1639	0.3061	0.1047	0.8283	0.0430	0.8824	-0.0617	0.8017
---	---	1628239_at	0.5769	0.0220	0.6296	0.3740	0.3160	0.2938	-0.0862	0.9158	-0.1330	0.6422	-0.0468	0.8773	0.3738	0.8215	0.1374	0.8918	-0.2364	0.7683
CG4825	Phosphatidylserin	1628240_at	0.4710	0.3347	1.0761	0.0123	1.5610	0.0084	0.4613	0.2988	0.8901	0.0053	0.4288	0.0688	-0.0345	0.9928	1.5543	0.0957	1.5889	0.1156
CG2614	CG2614	1628241_at	0.2969	0.1885	-0.4628	0.2343	0.0909	0.5354	-0.1424	0.7018	0.3830	0.0321	0.5254	0.0050	-0.5583	0.3771	-0.2535	0.4729	0.3048	0.3898
---	---	1628242_s_at	-0.1453	0.3342	0.0317	0.9113	0.0004	0.9984	-0.0447	0.9346	-0.2015	0.2090	-0.1568	0.2809	0.0482	0.9734	0.0188	0.9760	-0.0294	0.9530
dbr	debra	1628243_at	-0.2097	0.4277	0.4788	0.1651	0.1883	0.3262	-0.1409	0.8057	-0.4289	0.0637	-0.2880	0.1571	0.0563	0.9755	0.1879	0.7124	0.1316	0.8138
---	---	1628244_at	0.1637	0.3308	0.0337	0.7530	0.2534	0.2516	0.0709	0.8817	-0.0987	0.5743	-0.1696	0.2454	-0.2584	0.6832	-0.1109	0.7069	0.1475	0.5995
sens-2	senseless-2	1628245_at	-0.5073	0.2560	-0.0198	0.9207	-0.2617	0.2237	-0.0750	0.9715	-0.8430	0.1115	-0.7680	0.1054	-0.0323	0.9860	-0.1277	0.8117	-0.0954	0.8600
CG14372	CG14372	1628246_at	0.1334	0.4082	-0.0571	0.6801	0.1356	0.5140	0.1528	0.7507	0.0899	0.7096	-0.0629	0.7878	0.0003	0.9999	-0.0003	0.9999	-0.0006	0.9992
Him	Him	1628247_at	-0.1335	0.5334	0.0663	0.5280	0.1470	0.3921	-0.2163	0.5735	-0.3828	0.0557	-0.1665	0.3557	-0.1554	0.7823	-0.1331	0.6093	0.0223	0.9479
Scm	Sex combs on mic	1628248_at	0.1819	0.6088	0.1556	0.2240	-0.1275	0.4106	-0.0437	0.9373	0.2857	0.0780	0.3293	0.0298	0.3936	0.7062	0.3873	0.3651	-0.0063	0.9935
CG14145	CG14145	1628249_at	-0.1725	0.5168	0.3543	0.3800	-0.0600	0.8277	0.0473	0.9665	0.0672	0.8624	0.0199	0.9573	0.3749	0.7726	0.5949	0.2729	0.2200	0.7333
---	---	1628250_at	0.0116	0.9610	0.4153	0.0288	0.5903	0.0098	-0.0748	0.8841	-0.3685	0.0415	-0.2937	0.0637	-0.1139	0.8775	0.2070	0.4796	0.3209	0.2795
CG33054	CG33054	1628251_at	-0.2233	0.4127	-0.1615	0.5723	-0.1234	0.5336	-0.0954	0.8906	-0.3524	0.1278	-0.2570	0.2161	-0.1876	0.8461	-0.2692	0.5129	-0.0816	0.8800
CG17919	CG17919	1628252_at	-1.1430	0.0085	-2.1425	0.0016	-2.4019	0.0001	0.1299	0.7576	0.0216	0.9323	-0.1083	0.5401	0.4596	0.7215	-0.7888	0.1389	-1.2484	0.0559
---	---	1628253_at	0.0055	0.9820	0.1478	0.2266	0.0233	0.8845	-0.0062	0.9951	-0.0815	0.7303	-0.0753	0.7273	-0.0202	0.9816	0.0556	0.8335	0.0758	0.7403
CG31847	CG31847	1628254_at	0.1023	0.9059	0.8399	0.1770	0.6263	0.0120	-0.2143	0.8446	-0.8459	0.0421	-0.6316	0.0809	0.2610	0.9087	0.4914	0.5597	0.2304	0.8178
---	---	1628255_at	0.1372	0.5424	-0.0449	0.7023	0.3097	0.0766	-0.0026	0.9974	0.1211	0.6263	0.1238	0.5790	-0.2319	0.7215	-0.0484	0.9074	0.1835	0.5154
Mst77F	Male-specific tran	1628256_at	-0.2746	0.2340	-0.1815	0.2425	-0.1925	0.3176	0.0538	0.9491	-0.0489	0.8759	-0.1027	0.6767	0.0129	0.9916	0.0321	0.9450	0.0192	0.9636
PHGPx	PHGPx	1628257_s_at	0.1456	0.4066	0.1696	0.5047	0.7152	0.0022	0.3797	0.3431	-0.0701	0.8044	-0.4498	0.0360	-0.2210	0.7779	-0.3242	0.3284	-0.1032	0.8043
CG14526	CG14526	1628258_at	0.6458	0.0495	0.0056	0.9937	-0.0344	0.8690	-0.2728	0.6035	-0.0380	0.9183	0.2348	0.3312	-0.2013	0.8960	-0.6901	0.2156	-0.4888	0.4037
---	---	1628259_at	0.2017	0.2223	0.2428	0.1727	0.1751	0.3635	-0.0210	0.9777	0.0147	0.9569	0.0357	0.8682	-0.0023	0.9987	0.0126	0.9783	0.0149	0.9696
CG13186	CG13186	1628260_at	-0.0084	0.9791	-0.0496	0.6033	0.0990	0.5220	0.2870	0.4599	0.1011	0.6739	-0.1860	0.3340	0.0521	0.9515	-0.0260	0.9504	-0.0781	0.8044
CG13067	CG13067	1628261_at	0.5762	0.1969	0.6223	0.0763	-0.2547	0.2821	-0.8599	0.1210	-0.6719	0.0497	0.1880	0.5712	0.1120	0.9543	-0.5147	0.3605	-0.6268	0.2879
zfh1	Zn finger homeodi	1628262_a_at	-2.5189	0.0035	-2.2000	0.0194	-2.7166	0.0003	-0.4057	0.4553	-0.1484	0.6554	0.2573	0.3400	0.0103	0.9980	0.1249	0.9312	0.1146	0.9273
CG15236	CG15236	1628263_s_at	0.1411	0.3678	-0.0384	0.7504	-0.1269	0.5317	-0.0438	0.9421	0.2064	0.2275	0.2502	0.1017	-0.0004	0.9998	-0.0819	0.7027	-0.0815	0.7010
RecQ5	Drosophila homol	1628264_a_at	-0.3371	0.3390	-0.0438	0.9046	-0.5407	0.0104	-0.2199	0.7138	-0.1071	0.7396	0.1128	0.6941	0.2218	0.8465	-0.0227	0.9789	-0.2445	0.6328
CG32245	CG32245	1628265_a_at	-0.1175	0.4109	-0.0221	0.8269	-0.1172	0.4360	0.0962	0.8402	-0.0103	0.9703	-0.1065	0.5409	-0.0084	0.9914	0.0237	0.9277	0.0321	0.8880
---	---	1628266_at	0.0910	0.6933	-0.1257	0.5287	-0.1645	0.3449	0.1547	0.6537	0.1169	0.5057	-0.0378	0.8441	-0.0269	0.9816	-0.2286	0.3862	-0.2017	0.4677
CG15167	CG15167	1628267_at	0.2043	0.3805	0.1610	0.4054	0.2466	0.1119	0.0626	0.9101	0.0220	0.9292	-0.0406	0.8414	-0.0217	0.9862	-0.0656	0.8685	-0.0439	0.9096
CG18249	CG18249	1628268_at	1.0445	0.0045	-0.3295	0.5492	0.4445	0.0833	0.0476	0.9371	0.4294	0.0225	0.3818	0.0230	-0.7472	0.6660	-0.9057	0.1806	-0.1584	0.8636
Dox-A2	phenoloxidase	1628269_at	0.1855	0.2025	0.5288	0.0705	0.9940	0.0004	0.2294	0.4333	-0.1487	0.3649	-0.3781	0.0166	-0.1983	0.7116	0.1763	0.4303	0.3746	0.1271
CG30321	CG30321	1628270_at	0.0719	0.7169	0.0649	0.6587	0.1345	0.3686	0.0507	0.9110	0.0642	0.6982	0.0135	0.9395	-0.0962	0.8940	0.0738	0.8386	0.1701	0.5512
CG3609	CG3609	1628271_at	0.0658	0.6853	0.7630	0.0119	1.3375	0.0001	0.1846	0.4690	-1.0503	0.0002	-1.2350	0.0001	-0.2500	0.5461	-0.1126	0.5824	0.1374	0.4913

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG40486	CG40486	1628272_a_at	-0.4811	0.3366	-0.6447	0.0411	-0.8700	0.0031	-0.5598	0.1140	-0.8883	0.0022	-0.3286	0.0844	-0.2301	0.9057	-0.9982	0.1432	-0.7681	0.2855
---	---	1628273_at	-0.2380	0.1142	0.1257	0.4143	-0.0987	0.4994	-0.2534	0.4675	-0.1307	0.5112	0.1227	0.4974	0.0113	0.9914	0.0035	0.9942	-0.0078	0.9842
CG31924	CG31924	1628274_a_at	0.1532	0.6352	0.0563	0.7833	0.3608	0.1061	0.1236	0.8068	0.2077	0.3054	0.0841	0.6897	-0.1518	0.8611	0.1298	0.7579	0.2816	0.4370
Trl	GAGA factor	1628275_at	0.1601	0.7578	-0.0723	0.5746	-0.1245	0.7282	-0.0386	0.9860	0.0168	0.9822	0.0554	0.9285	-0.0078	0.9964	-0.2323	0.5800	-0.2245	0.5984
Bsg	Basigin	1628276_s_at	-0.5326	0.0634	-0.2057	0.3577	-0.6701	0.0370	-0.0898	0.8671	-0.0679	0.7654	0.0219	0.9252	0.3808	0.7644	0.2266	0.7105	-0.1542	0.8182
CG14384	CG14384	1628277_at	0.1030	0.6127	0.0656	0.6951	0.2919	0.1535	0.0650	0.9110	0.1100	0.5825	0.0450	0.8300	-0.1030	0.9101	0.0503	0.9213	0.1533	0.6649
CG10711	CG10711	1628278_at	-0.1216	0.5523	0.0773	0.5374	0.3523	0.0476	0.0527	0.9353	-0.2034	0.2931	-0.2561	0.1334	-0.1180	0.8586	0.0384	0.9277	0.1564	0.5838
CG11076	CG11076	1628279_s_at	0.2218	0.3953	0.4358	0.3433	0.4265	0.0315	0.1496	0.8249	0.1896	0.4900	0.0400	0.8991	0.2514	0.8284	0.5821	0.2168	0.3307	0.5154
Tim17b2	Translocase inner	1628280_at	0.0882	0.6412	0.0349	0.8128	0.3219	0.1361	0.1129	0.8409	0.0074	0.9811	-0.1055	0.6215	-0.0786	0.9174	-0.0245	0.9538	0.0541	0.8800
Mo25	Mo25	1628281_at	-0.1455	0.3439	0.2940	0.2474	-0.0509	0.7840	-0.0954	0.8155	0.1136	0.5025	0.2090	0.1443	0.1467	0.8461	0.4634	0.1355	0.3167	0.3186
CG11896	CG11896	1628282_at	-0.2075	0.7122	0.7125	0.1822	0.7254	0.0012	-0.2400	0.5785	-0.6034	0.0143	-0.3634	0.0689	-0.3813	0.8374	0.3227	0.7090	0.7040	0.3695
CG1575	CG1575	1628283_at	0.4085	0.2445	0.5909	0.2221	1.1397	0.0006	0.2423	0.7649	0.1812	0.6409	-0.0611	0.8815	-0.1938	0.8692	0.4785	0.2958	0.6723	0.1766
CG3690	CG3690	1628284_at	-0.1863	0.8109	-0.0074	0.9855	0.2027	0.4966	0.2528	0.8671	-0.4427	0.4225	-0.6955	0.1437	0.1482	0.8651	-0.1863	0.6170	-0.3345	0.3503
Rga	Regena	1628285_a_at	-0.0839	0.8212	0.3204	0.6277	0.7542	0.0050	0.0403	0.9603	-0.3572	0.0957	-0.3975	0.0439	-0.3926	0.8202	0.1761	0.8567	0.5688	0.4335
---	---	1628286_at	0.1337	0.5968	-0.1504	0.4482	0.0568	0.7966	0.1372	0.8248	0.1858	0.4573	0.0486	0.8621	-0.1284	0.8823	0.0672	0.8893	0.1956	0.5789
---	---	1628287_at	-0.0725	0.6952	0.0321	0.8638	0.0224	0.9206	0.1333	0.6645	-0.1012	0.5150	-0.2345	0.0756	0.1862	0.8270	0.0201	0.9736	-0.1661	0.6683
Nep4	Neprilysin 4	1628288_s_at	-0.1540	0.7546	-0.5121	0.3531	0.4506	0.4135	0.9130	0.2008	0.4601	0.2890	-0.4529	0.2407	-0.1843	0.9589	-0.0711	0.9664	0.1132	0.9372
ems	empty spiracles	1628289_at	-2.3292	0.0036	-0.0311	0.7653	-1.5578	0.0045	-1.2527	0.1171	-2.1863	0.0015	-0.9336	0.0371	0.0796	0.9142	-0.0722	0.8349	-0.1519	0.5814
CG17292	CG17292	1628290_s_at	0.8629	0.0139	0.2339	0.2176	0.6901	0.0038	0.1881	0.7349	0.1178	0.6756	-0.0703	0.7997	-0.3295	0.6593	-0.5036	0.1055	-0.1741	0.5954
CG4553	CG4553	1628291_at	-0.3735	0.2024	-0.1029	0.6192	0.2151	0.4094	-0.0629	0.9311	-0.3671	0.0876	-0.3041	0.1111	-0.2527	0.7628	-0.0435	0.9390	0.2093	0.5731
HLH106	Sterol regulatory e	1628292_s_at	-0.1871	0.6219	0.6231	0.1004	0.5133	0.0329	0.4486	0.2996	0.0775	0.8041	-0.3711	0.1019	0.6052	0.6483	1.0745	0.0654	0.4693	0.3898
bw	Plum	1628293_a_at	0.1020	0.5634	0.2076	0.2127	-0.0660	0.7314	0.0691	0.9059	0.0973	0.6461	0.0282	0.9007	0.2708	0.7230	0.1631	0.6489	-0.1077	0.7830
---	---	1628294_at	0.0072	0.9852	0.0000	1.0000	0.1473	0.5368	0.0826	0.9254	-0.0217	0.9559	-0.1043	0.7108	-0.0959	0.8943	-0.0043	0.9941	0.0917	0.7745
CG17300	CG17300	1628295_at	0.0477	0.8093	0.1622	0.2164	0.0381	0.8153	-0.1069	0.7492	-0.0635	0.7056	0.0434	0.7898	0.1274	0.8395	0.1125	0.7003	-0.0149	0.9702
pgant4	polypeptide GalN/	1628296_at	0.1705	0.3589	-0.1697	0.3950	-0.0628	0.8204	0.1848	0.8241	0.3859	0.2305	0.2011	0.5203	-0.0392	0.9589	-0.0299	0.9315	0.0093	0.9791
CG8478	CG8478	1628297_a_at	0.2433	0.5832	-0.9054	0.0291	-0.7744	0.0366	-0.1545	0.6756	1.1107	0.0004	1.2652	0.0002	-0.4644	0.7697	-0.1639	0.8619	0.3004	0.6877
Dox-A3	prophenoloxidase	1628298_at	-0.0589	0.7104	-0.1089	0.4876	-0.2439	0.2263	0.0614	0.8776	0.1041	0.4664	0.0426	0.7794	0.0594	0.9589	-0.1376	0.7124	-0.1971	0.5724
CG5339	CG5339	1628299_at	0.2689	0.1056	0.2650	0.0947	0.5425	0.0321	0.1148	0.7880	0.0859	0.6662	-0.0288	0.8907	-0.0677	0.9457	0.0804	0.8431	0.1481	0.6464
CG34030	CG34030	1628300_s_at	-0.2189	0.3251	-0.0668	0.6679	0.0120	0.9505	0.1437	0.7303	-0.1103	0.5861	-0.2540	0.1258	-0.0807	0.9092	0.0172	0.9664	0.0979	0.7328
CG32159	CG32159	1628301_at	-1.1536	0.0283	-0.4641	0.0437	-0.5690	0.0055	-0.0947	0.9507	-0.3931	0.3658	-0.2985	0.4586	-0.1173	0.8906	-0.0430	0.9340	0.0743	0.8602
CG13748	CG13748	1628302_at	-0.1597	0.5115	-0.1958	0.1463	-0.3901	0.3219	0.1436	0.9042	0.4086	0.2972	0.2651	0.4731	-0.1810	0.7681	-0.0854	0.7943	0.0957	0.7495
CG33231	CG33231	1628303_at	-0.1381	0.6839	0.0703	0.8406	-0.1323	0.5955	-0.2411	0.5978	-0.0884	0.7469	0.1528	0.4892	-0.0531	0.9710	0.0304	0.9600	0.0835	0.8634
---	---	1628304_at	-0.0319	0.8738	0.1174	0.3972	0.1645	0.3855	-0.0290	0.9722	-0.0603	0.8145	-0.0313	0.9003	-0.0164	0.9859	0.0883	0.7054	0.1047	0.6389
---	---	1628305_at	0.1449	0.4178	0.0506	0.6318	0.4113	0.0327	0.1424	0.7618	-0.1032	0.6500	-0.2456	0.1732	-0.1017	0.8861	-0.0775	0.8250	0.0242	0.9486
trc	tricorner	1628306_at	-0.3117	0.2477	-0.2594	0.1212	-0.4278	0.0347	-0.0584	0.9303	0.2557	0.1897	0.3141	0.0748	0.1842	0.8479	0.3371	0.3926	0.1529	0.7409
CG32388	CG32388	1628307_at	0.0675	0.6828	0.0286	0.8447	0.2150	0.3590	0.1435	0.8033	0.1404	0.5731	-0.0031	0.9913	-0.0967	0.8837	0.0033	0.9952	0.1000	0.7269
CG12714	CG12714	1628308_at	0.2299	0.2883	-0.0071	0.9579	0.0677	0.6896	-0.0300	0.9603	0.1062	0.5288	0.1362	0.3484	-0.1252	0.8091	-0.0170	0.9620	0.1082	0.6399
CG5126	CG5126	1628309_at	1.0035	0.0030	0.7281	0.0403	0.5940	0.0069	-0.0229	0.9777	0.2933	0.1518	0.3162	0.0860	-0.0868	0.9243	-0.0570	0.9022	0.0299	0.9435
---	---	1628310_at	0.1046	0.4396	-0.0039	0.9739	0.0235	0.9292	-0.1588	0.6637	0.0665	0.7483	0.2253	0.1478	0.0378	0.9611	0.0049	0.9925	-0.0329	0.9122
Rrp42	Rrp42	1628311_at	0.4726	0.0847	0.0966	0.8484	0.4432	0.0793	0.2163	0.5680	0.5367	0.0135	0.3204	0.0678	-0.1808	0.9246	0.1195	0.9018	0.3003	0.6646
CG15425	CG15425	1628312_at	0.0326	0.8973	0.0796	0.5297	0.2073	0.1544	-0.0716	0.9011	-0.1587	0.4087	-0.0870	0.6517	-0.1251	0.8374	0.0245	0.9512	0.1496	0.5744
CG41452	CG41452	1628313_at	0.0242	0.9160	-0.0908	0.6494	-0.4032	0.0664	-0.0005	0.9996	0.3576	0.1213	0.3581	0.0849	0.2257	0.7324	0.2642	0.3371	0.0386	0.9211
CG6891	CG6891	1628314_a_at	-1.4591	0.0004	-0.8957	0.2533	-1.2413	0.0748	0.0262	0.9715	0.6750	0.0039	0.6488	0.0028	0.3608	0.9342	1.2307	0.3720	0.8699	0.5587
CG17264	CG17264	1628315_at	-1.0806	0.2357	-1.8297	0.0752	-1.9667	0.0001	-0.0190	0.9863	0.9447	0.0044	0.9637	0.0024	0.1678	0.9717	0.0908	0.9634	-0.0770	0.9659
CG7670 /// DsimCG7670	CG7670	1628316_at	0.4854	0.4344	-0.6063	0.6379	-1.1800	0.1842	-0.7510	0.5604	1.5756	0.0253	2.3265	0.0027	-0.5752	0.9221	0.1441	0.9647	0.7194	0.7495
CG14410	CG14410	1628317_at	-0.0333	0.8926	0.0584	0.5757	-0.3061	0.1749	-0.1347	0.7138	-0.0108	0.9672	0.1239	0.4312	0.1054	0.9394	0.0447	0.9487	-0.0607	0.9176
CG11451	CG11451	1628318_at	-0.2675	0.4121	-0.0380	0.8851	-0.4298	0.2236	-0.5811	0.2172	-0.0095	0.9819	0.5716	0.0313	-0.4203	0.8023	-0.0207	0.9888	0.3995	0.5954
sdk	sidekick	1628319_s_at	-0.2069	0.6185	0.1686	0.6107	0.4126	0.1009	0.0475	0.9647	-0.1503	0.6367	-0.1978	0.4628	-0.0617	0.9816	0.2317	0.7299	0.2934	0.6388
CG9471 /// DyakCG9471	biliverdin reductas	1628320_at	-1.1209	0.0079	0.0474	0.9130	-0.3074	0.4677	-0.5706	0.3953	-1.5774	0.0022	-1.0068	0.0096	-0.2103	0.8967	-0.4431	0.4696	-0.2328	0.7381
CG4788	CG4788	1628321_at	-0.6016	0.0750	-1.0492	0.0089	-1.1195	0.0008	0.0022	0.9978	0.4455	0.0354	0.4432	0.0230	0.0290	0.9872	-0.0684	0.9152	-0.0975	0.8570
dah	Apodystrophin	1628322_at	0.1770	0.6209	-0.4661	0.2610	-0.2878	0.2071	-0.0719	0.8609	0.4300	0.0105	0.5018	0.0031	0.0098	0.9976	0.1643	0.8754	0.1545	0.8761
ogre	Innexin 1	1628323_s_at	-1.2489	0.0025	-1.4577	0.0097	-1.2883	0.0037	-0.2745	0.6844	-0.9464	0.0092	-0.6719	0.0248	-0.4874	0.6660	-0.8832	0.0694	-0.3958	0.3921
CG14442	CG14442	1628324_at	-0.5503	0.2352	-0.4837	0.5555	-0.3686	0.0488	-0.1485	0.7330	-0.2640	0.1717	-0.1155	0.5435	-0.3339	0.8940	-0.1003	0.9470	0.2336	0.8444
Krn	spitz2	1628325_at	0.2854	0.2596	-0.6329	0.0355	-0.2054	0.4666	0.4579	0.1223	0.6615	0.0037	0.2036	0.2019	0.1444	0.9142	-0.2793	0.5755	-0.4237	0.3807
---	---	1628326_at	-0.0287	0.9101	-0.0509	0.7165	-0.0220	0.9258	0.0667	0.9218	0.0479	0.8583	-0.0188	0.9412	-0.0039	0.9984	-0.0899	0.8664	-0.0860	0.8636
---	---	1628327_at	-0.0150	0.9516	0.1071	0.4713	0.1533	0.5627	-0.2237	0.5932	-0.5814	0.0140	-0.3577	0.0630	0.0479	0.9643	0.0244	0.9590	-0.0235	0.9542
GstE10	Glutathione S tran	1628328_at	-0.1280	0.8478	0.0592	0.5993	0.1304	0.4706	0.1520	0.9351	-0.3244	0.5878	-0.4764	0.3453	0.3145	0.7187	0.0887	0.8602	-0.2258	0.5566
Gr22b	Gustatory recepto	1628329_at	0.0729	0.7349	0.0998	0.4887	0.0654	0.7395	-0.0347	0.9622	-0.0159	0.9562	0.0189	0.9380	0.0928	0.8940	0.0775	0.8187	-0.0153	0.9695
CG34116	CG34116	1628330_at	0.0078	0.9758	0.2577	0.0613	0.3700	0.1231	-0.2159	0.6338	-0.5013	0.0318	-0.2854	0.1508	-0.2359	0.7436	-0.1300	0.7053	0.1059	0.7647
Ca-beta	calcium channel b	1628331_at	-0.2505	0.4274	-1.0531	0.0297	-1.7648	0.0001	-0.4403	0.6391	0.8665	0.0629	1.3068	0.0072	0.1251	0.9243	-0.0842	0.8988	-0.2094	0.6603
CG14990	CG14990	1628332_at	0.7425	0.1756	0.5398	0.0114	1.1188	0.0082	0.1802	0.7206	-0.0058	0.9869	-0.1860	0.3775	-0.1048	0.9689	-0.1874	0.8433	-0.0826	0.9331
CG4655	CG4655	1628333_at	-0.4446	0.2063	-1.3684	0.0050	-2.4457	0.0004	-0.3868	0.7104	0.7832	0.0985	1.1700	0.0136	0.0378	0.9814	-0.3356	0.3167	-0.3734	0.2926
m4	E(spl) region trans	1628334_at	-0.0072	0.9801	-0.0001	1.0000	0.1986	0.1918	0.2020	0.4902	0.0186	0.9323	-0.1833	0.1876	-0.0756	0.8919	-0.0311	0.9247	0.0445	0.8745
CG5222	CG5222	1628335_at	0.1702	0.3799	-0.1577	0.4499	0.0661	0.6772	0.1084	0.8211	0.5719	0.0086	0.4635	0.0134	-0.1518	0.8597	0.2067	0.5707	0.3585	0.3097
Lac	lachesin	1628336_a_at	-0.8622	0.0027	-0.8970	0.0277	-1.0036	0.0005	0.1761	0.6096	0.3336	0.0583	0.1574	0.3186	0.2746	0.6955	0.2880	0.3045	0.0133	0.9769
CG30354	CG30354	1628337_at	0.5811	0.0113	0.2366	0.3943	0.2642	0.3153	-0.1188	0.8140	0.2537	0.2006	0.3725	0.0426	-0.1531	0.8305	-0.1259	0.7105	0.0272	0.9486
CG32756	CG32756	1628338_at	-0.0023	0.9953	-0.4134	0.1398	-0.6078	0.0135	0.0863	0.8869	0.7421	0.0040	0.6557	0.0042	0.1019	0.9309	0.2294	0.5651	0.1275	0.7764
CG7920 /// DsimCG7920	CG7920	1628339_a_at	-0.0434	0.9349	0.7541	0.0418	1.1701	0.0054	0.1509	0.9218	-1.1955	0.0182	-1.3464	0.0065	-0.2107	0.9038	-0.1916	0.8158	0.0192	0.9849
lark	lark	1628340_s_at	-0.0024	0.9936	0.6279	0.1353	0.3774	0.0266	0.1065	0.8589	-0.1714	0.4435	-0.2779	0.1478	0.3426	0.7464	0.4985	0.2468	0.1559	0.7638

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1628341_at	-0.1379	0.8900	-0.9355	0.1389	-1.4065	0.0111	-0.5917	0.6099	0.9963	0.0878	1.5880	0.0086	-0.2140	0.9330	-0.0323	0.9837	0.1816	0.8667
ball	ballchen	1628342_s_at	0.0762	0.7195	0.0325	0.7891	0.2737	0.2409	-0.1001	0.8865	-0.0983	0.7222	0.0018	0.9953	-0.4568	0.5126	-0.0839	0.8550	0.3729	0.2798
---	---	1628343_at	0.1811	0.3043	-0.0034	0.9781	0.0889	0.5630	0.1018	0.8097	0.0593	0.7692	-0.0426	0.8239	0.0202	0.9816	-0.0415	0.8927	-0.0618	0.8111
CkIIbeta	casein kinase 2	1628344_at	-0.2189	0.3287	0.3842	0.3862	0.1055	0.6754	-0.2916	0.3433	-0.3997	0.0315	-0.1081	0.5321	0.2384	0.8692	0.4400	0.4475	0.2016	0.7641
Cyp6a9	cytochrome P450	1628345_at	-1.4200	0.0019	-0.4272	0.4796	-1.2339	0.0062	-0.6438	0.3842	-1.6708	0.0027	-1.0270	0.0140	0.0776	0.9781	-0.7245	0.2655	-0.8021	0.2525
---	---	1628346_at	-0.0444	0.8044	0.0606	0.6106	0.2696	0.3518	0.0547	0.9125	-0.0474	0.8072	-0.1021	0.5002	-0.0948	0.9514	0.1866	0.7269	0.2814	0.5659
---	---	1628347_s_at	0.0701	0.6970	-0.0651	0.5764	-0.0314	0.8571	0.1143	0.8671	0.0210	0.9531	-0.0933	0.7165	0.0856	0.9238	-0.0498	0.9129	-0.1354	0.6761
CG32081	CG32081	1628348_at	0.1348	0.4007	-0.0432	0.8470	0.0620	0.7878	0.2348	0.6854	0.2801	0.3032	0.0454	0.8878	-0.0954	0.8692	-0.0640	0.8376	0.0314	0.9216
CG14507	CG14507	1628349_a_at	0.1789	0.2857	-0.0629	0.5977	-0.0961	0.6124	0.0185	0.9803	0.2401	0.1864	0.2216	0.1743	-0.1027	0.8795	-0.0574	0.8773	0.0452	0.8998
CG40368	CG40368	1628350_a_at	-0.0574	0.7591	0.0287	0.8216	0.0008	0.9977	0.1268	0.8190	0.1189	0.6217	-0.0079	0.9770	0.1089	0.8141	0.0710	0.7554	-0.0379	0.8858
CHORD	CHORD containin	1628351_at	0.0364	0.9173	0.1322	0.6561	0.0303	0.8753	0.0632	0.9488	0.4369	0.1112	0.3737	0.1269	0.1430	0.8692	0.3816	0.2507	0.2386	0.5041
CG8798	CG8798	1628352_a_at	0.0985	0.5722	-0.4150	0.0230	-0.4565	0.1843	-0.1219	0.9203	0.4656	0.2083	0.5876	0.0783	-0.0020	0.9994	0.1596	0.6952	0.1616	0.6870
GstD1	CG10045 Glutathi	1628353_at	0.0022	0.9915	0.4705	0.0308	0.8125	0.0044	0.0932	0.9008	-0.3023	0.2092	-0.3954	0.0698	-0.2223	0.7337	0.1678	0.5684	0.3902	0.1891
---	---	1628354_at	-0.1112	0.5973	-0.0518	0.8495	-0.0155	0.9473	0.0972	0.8421	0.1142	0.5585	0.0170	0.9395	-0.0039	0.9974	0.0634	0.8725	0.0673	0.8509
CG32379	CG32379	1628355_at	-0.1327	0.9410	0.0490	0.6639	-0.3105	0.0833	-0.0427	0.9922	-0.9842	0.2857	-0.9414	0.2522	0.0515	0.9928	-0.6420	0.6512	-0.6935	0.6225
---	---	1628356_at	0.0217	0.9143	0.0469	0.6679	-0.0497	0.7535	-0.0066	0.9943	-0.0861	0.7061	-0.0795	0.7032	0.0553	0.9238	-0.0206	0.9477	-0.0759	0.7320
CG6142	CG6142	1628357_at	-4.0701	0.0008	-5.8323	0.0014	-4.8923	0.0000	0.9096	0.3514	1.4071	0.0202	0.4975	0.3309	-0.0415	0.9862	-0.2986	0.6059	-0.2571	0.6617
Tob	Tob	1628358_at	-0.1627	0.7840	-0.3821	0.3760	-1.4301	0.0530	-0.6207	0.5688	0.1487	0.8356	0.7694	0.1222	0.4324	0.8768	-0.0545	0.9779	-0.4869	0.6857
---	---	1628359_at	0.0132	0.9478	0.1012	0.5292	0.0002	0.9994	-0.1579	0.6916	-0.2686	0.1433	-0.1107	0.5385	0.0046	0.9970	-0.0822	0.8439	-0.0869	0.8218
CG17150 /// DmirCG17150	CG17150	1628360_at	0.0187	0.9566	0.1341	0.4625	0.0328	0.8574	0.1188	0.7622	-0.0542	0.7932	-0.1730	0.2540	0.2210	0.7440	0.0864	0.8153	-0.1345	0.6641
CG32986	CG32986	1628361_at	0.0185	0.9105	-0.0805	0.7014	0.0058	0.9859	-0.0416	0.9672	0.0250	0.9500	0.0666	0.8253	-0.0468	0.9751	-0.1551	0.7127	-0.1083	0.8147
CG2941 /// CG32783 /// CC	CG2941 /// CG32	1628362_s_at	-0.3779	0.2810	0.1095	0.5572	-0.2609	0.1497	-0.4096	0.3596	-0.3786	0.1360	0.0310	0.9210	-0.0860	0.9587	0.1009	0.8782	0.1869	0.7194
inx3	innexin 3	1628363_at	1.3159	0.0030	1.1904	0.0435	1.2672	0.0070	-0.2378	0.5376	-0.1964	0.3377	0.0414	0.8599	-0.3686	0.8023	-0.1748	0.8303	0.1938	0.7933
CG7968 /// DyakCG7968	CG7968	1628364_at	0.0445	0.8874	0.0750	0.5697	0.6131	0.0091	0.1776	0.6970	-0.0582	0.8257	-0.2358	0.2092	-0.5149	0.3712	-0.2481	0.4317	0.2668	0.4075
CG5538	CG5538	1628365_at	0.8364	0.0107	0.0225	0.9184	0.3286	0.1225	-0.0086	0.9943	0.2080	0.4148	0.2166	0.3388	-0.2456	0.7215	-0.3933	0.1644	-0.1477	0.6385
CG31065	CG31065	1628366_at	0.1915	0.3221	0.1528	0.4449	0.0224	0.9218	-0.0290	0.9749	0.0943	0.7202	0.1233	0.5851	0.0773	0.9216	0.0052	0.9933	-0.0720	0.8260
---	---	1628367_at	0.2328	0.2732	0.4611	0.0623	0.0847	0.7007	-0.1229	0.7121	-0.0427	0.8212	0.0802	0.5975	0.1081	0.9365	0.0087	0.9931	-0.0994	0.8600
Sir2	Sir2	1628368_at	0.2536	0.3425	0.0871	0.4524	-0.2126	0.2628	-0.0817	0.8902	0.2150	0.2805	0.2967	0.0939	0.3342	0.6749	0.0628	0.8979	-0.2714	0.4114
boss	bride of sevenless	1628369_at	0.3665	0.0546	0.1067	0.4850	0.0961	0.6772	0.0984	0.9036	0.1502	0.6027	0.0519	0.8662	0.2201	0.7220	0.0675	0.8500	-0.1526	0.5838
---	---	1628370_at	0.0831	0.5466	0.1237	0.4135	0.1498	0.3906	0.0017	0.9974	-0.0203	0.9211	-0.0220	0.9000	0.0465	0.9653	0.1026	0.7692	0.0561	0.8874
HLH4C	Helix loop helix pr	1628371_at	0.2340	0.2024	0.0483	0.6497	0.0242	0.9161	-0.1484	0.6854	0.0750	0.7021	0.2234	0.1406	0.0036	0.9964	-0.0035	0.9935	-0.0071	0.9819
DNApol-alpha73	DNA polymerase `	1628372_a_at	0.2094	0.3594	0.2018	0.7857	-0.0589	0.9075	-0.4605	0.3016	0.0935	0.7664	0.5540	0.0263	0.0054	0.9994	0.0928	0.9487	0.0873	0.9434
CG32085	CG32085	1628373_at	-0.3980	0.4641	-0.8024	0.0996	-0.6161	0.0293	-0.6973	0.1731	-0.7049	0.0305	-0.0077	0.9849	-0.8392	0.3800	-1.1541	0.0512	-0.3149	0.5765
Gr22a	Gustatory recepto	1628374_at	0.0115	0.9468	0.1412	0.3485	0.3018	0.1338	0.1049	0.8342	0.0770	0.7297	-0.0280	0.9028	0.0488	0.9640	0.1411	0.6605	0.0923	0.7935
CG11786	CG11786	1628375_at	0.1529	0.3266	0.1493	0.1547	0.1090	0.5094	-0.0803	0.8862	0.0113	0.9694	0.0916	0.6381	0.0892	0.8657	0.0351	0.9145	-0.0541	0.8409
CG13947	CG13947	1628376_x_at	0.0547	0.8526	0.1336	0.5943	-0.0869	0.8399	0.0221	0.9894	-0.0458	0.9326	-0.0679	0.8815	0.1653	0.8692	0.1727	0.7007	0.0073	0.9920
AP-50	AP-50	1628377_s_at	-0.4798	0.0319	-0.1549	0.3931	-0.1973	0.3015	0.1239	0.6988	0.0339	0.8605	-0.0900	0.5306	-0.0239	0.9835	0.2584	0.3217	0.2823	0.3027
---	---	1628378_at	-0.0191	0.9459	0.0326	0.9028	0.0976	0.6126	0.0685	0.9345	-0.0135	0.9717	-0.0819	0.7604	0.1315	0.8141	0.0381	0.9156	-0.0934	0.7222
CG10749	CG10749	1628379_at	0.0239	0.9448	0.0217	0.8412	0.1022	0.6025	0.0735	0.8736	-0.0749	0.6827	-0.1485	0.3092	-0.0297	0.9841	-0.0663	0.8949	-0.0366	0.9381
CG13647	CG13647	1628380_at	0.1639	0.4145	0.1202	0.3347	-0.0610	0.7661	0.0209	0.9803	0.0383	0.8919	0.0174	0.9473	0.1906	0.7358	-0.0229	0.9528	-0.2136	0.3855
CG3579	CG3579	1628381_at	-0.0374	0.8739	0.1391	0.4470	0.3571	0.0724	0.1602	0.6673	-0.2529	0.1529	-0.4131	0.0175	-0.0642	0.9296	-0.0675	0.8284	-0.0033	0.9937
Srp54	Srp54	1628382_at	-0.0948	0.5939	0.2345	0.1383	0.0692	0.7163	0.0849	0.8479	-0.2627	0.1040	-0.3475	0.0246	0.2862	0.6749	0.0857	0.8172	-0.2005	0.4978
CG3330	CG3330	1628383_at	0.0251	0.9226	0.0073	0.9604	-0.0654	0.7235	0.0462	0.9584	0.1132	0.6692	0.0670	0.7981	0.0404	0.9657	0.1021	0.7275	0.0617	0.8507
---	---	1628384_at	-0.1970	0.2491	0.1708	0.4402	-0.0555	0.8104	-0.3529	0.4455	-0.4234	0.0934	-0.0705	0.8039	0.0179	0.9884	0.0488	0.9048	0.0308	0.9342
Mef2	Complementation	1628385_a_at	1.4386	0.0409	1.4620	0.0026	1.6911	0.0001	0.3093	0.2693	0.5387	0.0065	0.2294	0.1195	0.0335	0.9924	0.5738	0.4958	0.5403	0.5320
---	---	1628386_at	-1.0229	0.0287	-1.4374	0.0163	-1.4367	0.0087	0.1863	0.9260	0.9119	0.1296	0.7256	0.1780	-0.2544	0.6700	0.0269	0.9451	0.2813	0.2597
CG30080	CG30080	1628387_s_at	0.5339	0.1093	0.8456	0.0929	0.6901	0.0095	0.0015	0.9988	-0.5522	0.0436	-0.5537	0.0276	0.1494	0.9246	-0.4134	0.4335	-0.5629	0.2993
CG3823	CG3823	1628388_at	-1.3630	0.0255	2.2509	0.0410	1.2890	0.0156	-1.2524	0.2063	-5.2057	0.0001	-3.9533	0.0002	-0.0459	0.9914	-1.1994	0.1925	-1.1536	0.2398
CG4225	CG4225	1628389_at	-0.3456	0.1058	0.2064	0.5442	0.4609	0.0167	-0.0789	0.8611	-0.8044	0.0011	-0.7255	0.0010	-0.5355	0.5540	-0.0927	0.8765	0.4428	0.3006
CG6151	CG6151	1628390_at	0.5770	0.0462	0.9788	0.0508	0.7703	0.0011	-0.1067	0.8485	-0.0497	0.8511	0.0570	0.8039	0.0121	0.9943	0.2385	0.5657	0.2263	0.5944
---	---	1628391_at	0.1450	0.4658	0.4160	0.2141	0.6845	0.0033	0.2086	0.5249	-0.1035	0.5785	-0.3121	0.0475	0.1280	0.8689	0.2069	0.5141	0.0789	0.8401
---	---	1628392_x_at	-0.0159	0.9275	-0.0720	0.7037	0.0495	0.7799	0.0336	0.9550	0.0304	0.8894	-0.0032	0.9874	0.1047	0.8779	0.0875	0.7914	-0.0171	0.9655
CG10039	CG10039	1628393_at	0.0488	0.8831	0.0365	0.7142	0.1296	0.5872	0.2392	0.5586	0.1296	0.5685	-0.1096	0.6057	0.0042	0.9984	-0.1185	0.8161	-0.1227	0.7953
---	---	1628394_at	-0.0187	0.9482	-0.0891	0.4476	0.0986	0.6045	0.2182	0.5161	0.1990	0.2642	-0.0192	0.9296	-0.1038	0.8439	0.0193	0.9567	0.1231	0.5972
---	---	1628395_at	0.0326	0.8661	0.0876	0.4384	0.2967	0.1101	0.1021	0.8699	-0.0121	0.9715	-0.1142	0.6038	0.0221	0.9816	0.0196	0.9528	-0.0025	0.9946
CG7397	CG7397	1628396_at	0.0983	0.5838	-0.0848	0.5992	-0.0152	0.9486	0.1261	0.8067	-0.0169	0.9556	-0.1430	0.4607	0.1538	0.8023	-0.0291	0.9445	-0.1829	0.4907
CG8707 /// DyakCG8707	CG8707	1628397_at	0.0165	0.9251	0.3798	0.0751	0.2259	0.3108	-0.0377	0.9622	-0.1709	0.4354	-0.1331	0.5154	0.1561	0.8378	0.1625	0.6312	0.0064	0.9903
CG5853	CG5853	1628398_at	1.8279	0.0230	0.9019	0.1018	2.2404	0.0000	0.5957	0.6391	0.1923	0.8008	-0.4034	0.4997	-0.5996	0.6955	-0.6471	0.3000	-0.0475	0.9595
B52	Serine/arginine ric	1628399_s_at	0.1942	0.4069	0.3649	0.1015	0.1352	0.3308	0.1854	0.5156	0.5060	0.0056	0.3206	0.0255	0.2791	0.7215	0.5052	0.1275	0.2261	0.5092
CG40367	CG40367	1628400_at	0.3840	0.2928	0.0117	0.9108	-0.0148	0.9369	0.0436	0.9518	0.0770	0.7453	0.0335	0.8887	-0.0383	0.9816	-0.1820	0.6380	-0.1437	0.7223
CG31952	CG31952	1628401_at	0.0958	0.5473	0.2359	0.2082	0.0130	0.9386	-0.0615	0.8990	-0.0829	0.6416	-0.0213	0.9112	0.2045	0.7357	0.1570	0.5618	-0.0475	0.8933
GlcAT-S	GlcAT-S	1628402_a_at	-1.4760	0.0014	-0.7744	0.0555	-1.1819	0.0317	-0.2062	0.5633	0.2428	0.1819	0.4490	0.0139	0.1188	0.9701	0.7243	0.3647	0.6054	0.4704
---	---	1628403_at	0.0532	0.8185	-0.0346	0.7322	0.0360	0.8560	0.0257	0.9777	0.2301	0.3140	0.2043	0.3223	-0.1616	0.7726	0.0334	0.9313	0.1950	0.4336
CG2767	CG2767	1628404_at	0.0119	0.9538	0.2521	0.2002	0.3758	0.0387	-0.0872	0.8605	-0.0760	0.7123	0.0112	0.9597	-0.2790	0.6955	0.1550	0.6256	0.4339	0.1651
CG7298	CG7298	1628405_at	0.3537	0.5727	0.0371	0.8266	-0.2065	0.4029	-0.5064	0.3749	-0.2789	0.4012	0.2275	0.4568	-0.0995	0.9775	-0.6866	0.4037	-0.5871	0.4944
---	---	1628406_s_at	-0.3799	0.2671	-1.8168	0.1592	-1.4212	0.0129	0.1747	0.8605	1.3324	0.0035	1.1577	0.0039	-0.0919	0.9848	0.0601	0.9739	0.1520	0.9189
CG17042	CG17042	1628407_at	-0.0389	0.8287	0.1187	0.3143	0.2527	0.1879	0.1289	0.7551	-0.0808	0.6926	-0.2097	0.1867	-0.0160	0.9892	0.0988	0.7269	0.1148	0.6668
CG34041	CG34041	1628408_at	0.1346	0.5470	0.1493	0.3496	0.0413	0.8293	-0.1168	0.8578	-0.0853	0.7620	0.0315	0.9116	0.1506	0.7893	0.0416	0.9101	-0.1090	0.6806
CG32576	CG32576	1628409_at	0.4018	0.1517	0.0300	0.9148	-0.2266	0.1401	-0.0746	0.8636	0.5009	0.0065	0.5755	0.0021	0.1592	0.8628	0.2600	0.4954	0.1007	0.8293

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1628410_at	0.0802	0.6415	0.1047	0.4465	0.0281	0.8708	0.0431	0.9413	0.0938	0.6125	0.0507	0.7871	0.0065	0.9946	-0.0134	0.9719	-0.0199	0.9503
CG17390	CG17390	1628411_at	-0.0257	0.9222	-0.0776	0.5127	-0.1923	0.2226	-0.0293	0.9592	0.0550	0.7618	0.0844	0.5760	-0.0088	0.9935	-0.0039	0.9941	0.0049	0.9907
Ac78C	Adenylyl cyclase	1628412_at	-2.5057	0.0032	0.2209	0.8649	-1.3647	0.0162	-1.4054	0.1401	-2.3126	0.0026	-0.9071	0.0796	0.1034	0.9841	0.4268	0.7642	0.3234	0.8267
---	---	1628413_at	-0.0244	0.9192	0.3450	0.2020	0.1908	0.2396	0.0499	0.9310	0.0539	0.7935	0.0041	0.9850	-0.0348	0.9647	0.2641	0.1974	0.2989	0.1829
CG4955	CG4955	1628414_at	0.0164	0.9528	0.1050	0.6488	0.2339	0.2652	-0.0600	0.9345	-0.1504	0.5064	-0.0904	0.6878	-0.1610	0.7848	0.0068	0.9905	0.1678	0.5298
---	---	1628415_at	-0.1140	0.5981	0.2351	0.1718	-0.1120	0.5677	-0.0360	0.9668	-0.0051	0.9883	0.0309	0.9101	0.3476	0.5421	0.3450	0.1804	-0.0026	0.9956
Dp1	Drosophila dodeca	1628416_s_at	0.4479	0.1648	0.6301	0.0818	0.7294	0.0830	0.0335	0.9834	0.2180	0.5887	0.1845	0.6248	0.1408	0.9238	0.4909	0.2984	0.3501	0.4851
CG14324	CG14324	1628417_at	-0.0704	0.6913	-0.1186	0.4072	-0.0088	0.9756	0.0353	0.9603	0.1782	0.3520	0.1429	0.4166	-0.1417	0.7768	-0.0035	0.9941	0.1381	0.5450
abo	abnormal oocyte	1628418_at	0.0515	0.7911	0.2004	0.3599	0.4967	0.0072	0.1817	0.5516	0.0322	0.8786	-0.1495	0.2964	-0.1019	0.8692	0.1697	0.4997	0.2715	0.2830
CG8909	yolkless-like	1628419_a_at	-0.1862	0.3115	-0.2105	0.2748	-0.1351	0.7261	0.0412	0.9734	0.1877	0.5686	0.1465	0.6390	-0.0386	0.9751	-0.0021	0.9988	0.0365	0.9323
CG6923	CG6923	1628420_s_at	-0.1895	0.5337	-0.8045	0.0920	-0.7123	0.0100	0.2165	0.8028	0.8579	0.0216	0.6414	0.0444	0.0216	0.9914	0.1849	0.7341	0.1633	0.7638
lola	longitudinals abse	1628421_at	-0.5928	0.0775	-0.3282	0.1937	-0.7572	0.0087	0.0317	0.9777	0.4714	0.0907	0.4398	0.0782	0.5413	0.5754	0.7680	0.0890	0.2267	0.6306
CG31650	CG31650	1628422_s_at	-1.0393	0.0011	-0.6900	0.0953	-0.7898	0.0005	-0.0059	0.9943	0.1212	0.5060	0.1271	0.4331	0.1417	0.8714	0.5000	0.1418	0.3583	0.3128
Bap170	Brahma associate	1628423_at	0.3617	0.2613	-0.1314	0.5329	-0.0339	0.8661	0.0877	0.9110	0.7001	0.0123	0.6125	0.0137	0.1128	0.9309	0.3021	0.4864	0.1894	0.6870
CG5913	CG5913	1628424_at	-0.2674	0.4903	-0.1951	0.1163	-0.3465	0.0659	-0.0210	0.9745	0.4179	0.0183	0.4389	0.0091	0.0162	0.9923	0.3015	0.4301	0.2853	0.4710
CG31898	CG31898	1628425_at	0.2067	0.3309	-0.1966	0.1098	-0.4115	0.0405	-0.1154	0.8409	0.2680	0.2099	0.3834	0.0504	-0.0222	0.9898	-0.2433	0.5321	-0.2211	0.5795
CG8740	CG8740	1628426_s_at	-0.6606	0.4999	-0.0507	0.9800	0.2506	0.5796	0.3085	0.6663	-0.5048	0.1371	-0.8133	0.0157	-0.0471	0.9952	-0.0209	0.9965	0.0262	0.9941
Rab40	Rab40	1628427_a_at	-0.6439	0.1740	1.0829	0.1296	0.2149	0.3820	-0.1112	0.9023	-1.1115	0.0033	-1.0003	0.0032	0.7676	0.6898	0.6061	0.4418	-0.1615	0.8803
Fpps	Farnesyl diphospl	1628428_at	0.1099	0.7560	-0.1005	0.7505	0.0851	0.7723	0.2693	0.5724	-0.2141	0.3913	-0.4834	0.0336	-0.0506	0.9816	-0.4302	0.3802	-0.3796	0.4631
Sec61gamma	Sec61gamma	1628429_at	0.2782	0.1261	1.2976	0.0080	1.1248	0.0005	0.0412	0.9371	0.0150	0.9469	-0.0262	0.8842	0.1283	0.8650	0.9272	0.0222	0.7989	0.0422
CG15704	CG15704	1628430_at	0.3462	0.0706	-0.0615	0.5896	0.1497	0.3723	0.0418	0.9647	-0.0647	0.8415	-0.1064	0.6851	0.0972	0.8680	-0.1268	0.6080	-0.2240	0.3467
---	---	1628431_at	0.0441	0.8552	-0.0491	0.8605	-0.0581	0.7684	0.0845	0.9326	0.1410	0.6764	0.0565	0.8716	0.0273	0.9771	-0.0818	0.7604	-0.1090	0.6519
CG5590	CG5590	1628432_at	0.1296	0.3912	0.2204	0.4735	0.6604	0.0058	-0.0575	0.9343	-0.6408	0.0079	-0.5833	0.0072	-0.4431	0.6897	-0.4669	0.2854	-0.0238	0.9727
---	---	1628433_at	-0.1096	0.6490	0.0657	0.5657	-0.1077	0.6878	-0.1787	0.6354	-0.1453	0.4499	0.0334	0.8790	0.0615	0.9514	0.0041	0.9950	-0.0574	0.8901
CG33220	CG33220	1628434_at	0.0421	0.8472	0.4804	0.1331	-0.0540	0.8642	-0.1026	0.8578	-0.1830	0.3874	-0.0805	0.7136	0.1633	0.8374	0.2841	0.3756	0.1208	0.7492
sqd	RNA-binding prote	1628435_at	1.4633	0.0142	1.0984	0.0052	-0.2061	0.5468	-0.4605	0.2697	0.2041	0.4233	0.6646	0.0091	0.7238	0.6557	0.4105	0.5522	-0.3133	0.6607
CG13203	CG13203	1628436_at	0.0309	0.9013	0.2603	0.1807	0.4332	0.0303	-0.0206	0.9757	-0.1863	0.2627	-0.1657	0.2671	0.0290	0.9816	0.1830	0.5296	0.1540	0.6111
CG12783	CG12783	1628437_at	-0.0562	0.6888	0.0160	0.8982	-0.0224	0.9038	0.2792	0.5140	0.1075	0.6767	-0.1717	0.4174	0.1470	0.7726	0.0238	0.9473	-0.1233	0.6040
CG9044	CG9044	1628438_at	0.8399	0.0069	0.4390	0.1038	0.3722	0.3227	-0.0270	0.9757	-0.0566	0.8395	-0.0296	0.9100	0.0514	0.9848	-0.2833	0.6820	-0.3347	0.6166
---	---	1628439_s_at	-1.8482	0.0038	-1.6512	0.0028	-2.1470	0.0005	-0.0103	0.9956	-0.2658	0.6289	-0.2556	0.6084	0.0027	0.9970	0.0078	0.9801	0.0051	0.9849
---	---	1628440_at	0.0921	0.6736	-0.1489	0.2996	0.0014	0.9956	0.1553	0.7130	0.0802	0.7195	-0.0750	0.7129	-0.1074	0.8461	-0.1186	0.6288	-0.0113	0.9752
CG40138	CG40138	1628441_at	0.1493	0.5180	-0.2299	0.1608	-0.0954	0.6145	-0.0378	0.9603	0.3413	0.0938	0.3791	0.0433	-0.1502	0.8141	-0.1414	0.6181	0.0088	0.9843
---	---	1628442_at	0.2030	0.5411	0.1028	0.5961	0.3761	0.1015	0.0462	0.9479	0.0270	0.9226	-0.0192	0.9379	-0.0152	0.9928	-0.0384	0.9505	-0.0232	0.9686
---	---	1628443_at	0.0440	0.7697	0.1419	0.4243	-0.0754	0.6639	-0.0887	0.8649	-0.0550	0.8099	0.0337	0.8776	0.1293	0.7644	0.0502	0.8370	-0.0791	0.6963
---	---	1628444_at	0.1691	0.3953	0.4238	0.4645	0.0233	0.9332	-0.3683	0.2873	-0.5392	0.0162	-0.1709	0.3620	-0.0800	0.9618	-0.3190	0.5066	-0.2390	0.6328
CG33213	CG33213	1628445_at	0.1050	0.7161	-0.6641	0.0727	-0.5248	0.0667	-0.1693	0.8660	0.7547	0.0395	0.9240	0.0107	-0.3238	0.6557	-0.0086	0.9894	0.3152	0.2980
MtnC	Metallothionein C	1628446_at	-0.5061	0.8784	0.2180	0.5693	-0.1749	0.3014	-0.2886	0.9715	-2.0328	0.3271	-1.7442	0.3528	0.2198	0.9848	-1.3914	0.6259	-1.6113	0.5653
CG33993	CG33993	1628447_at	0.1734	0.3651	0.1156	0.4829	-0.2875	0.2396	-0.0294	0.9671	-0.2044	0.2742	-0.1750	0.2995	0.4282	0.6955	0.1069	0.8642	-0.3213	0.4869
CG31473	CG31473	1628448_at	-0.2163	0.4981	0.0973	0.4145	0.2735	0.2761	-0.1423	0.7278	-0.6990	0.0032	-0.5568	0.0053	-0.2065	0.8283	-0.2532	0.5422	-0.0467	0.9338
Ugt86Dc	Ugt86Dc	1628449_at	0.1594	0.7851	-0.2156	0.2634	-0.1374	0.7269	-0.0601	0.9422	-0.3522	0.1344	-0.2920	0.1656	-0.0446	0.9893	-0.8198	0.2168	-0.7752	0.2757
Nplp1	neuropeptide-like	1628450_at	0.3546	0.0319	0.0073	0.9817	0.3638	0.0610	0.1509	0.7031	0.0801	0.7033	-0.0708	0.7149	-0.2540	0.6695	-0.2047	0.3865	0.0493	0.8800
SuUR	Suppressor of Un	1628451_at	0.2517	0.3609	0.5569	0.1450	-0.0594	0.7330	-0.1973	0.6517	0.4722	0.0325	0.6695	0.0043	0.3876	0.7576	0.8188	0.1248	0.4313	0.4258
---	---	1628452_at	-0.1539	0.3636	0.1094	0.4189	-0.0711	0.7340	-0.0302	0.9602	-0.1457	0.3695	-0.1155	0.4397	0.1682	0.7772	0.1585	0.5587	-0.0097	0.9832
l(1)G0289	lethal (1) G0289	1628453_at	1.7695	0.0098	1.1269	0.0956	0.3968	0.4568	-0.7600	0.2461	0.0172	0.9775	0.7772	0.0324	-0.0518	0.9914	-0.6514	0.5333	-0.5996	0.5756
ppk12	pickpocket 12	1628454_at	0.2069	0.3341	-0.0209	0.8608	0.0575	0.7362	0.0788	0.8894	0.1231	0.5460	0.0444	0.8390	0.0185	0.9875	-0.0165	0.9701	-0.0350	0.9236
---	---	1628455_at	0.2531	0.2251	0.3074	0.1463	0.0311	0.9111	-0.1019	0.8545	-0.0893	0.6994	0.0126	0.9598	0.3080	0.7436	0.1183	0.8187	-0.1898	0.6601
Spt3	Spt3	1628456_at	-0.4116	0.0401	0.0408	0.8318	0.2405	0.1986	0.3008	0.3016	-0.2375	0.1633	-0.5383	0.0045	-0.0269	0.9848	-0.0361	0.9431	-0.0093	0.9852
---	---	1628457_at	0.0952	0.5477	0.1757	0.1503	0.0886	0.5991	0.0029	0.9956	0.1134	0.5153	0.1105	0.4814	-0.0533	0.9365	0.0570	0.8428	0.1102	0.6299
milt	Milton	1628458_at	-0.6840	0.0140	-1.1387	0.0335	-1.2944	0.0001	0.0313	0.9649	0.3804	0.0507	0.3491	0.0458	0.1973	0.8331	-0.0949	0.8615	-0.2922	0.4686
Acp36DE	36DE accessory g	1628459_at	0.1007	0.6063	-0.0205	0.8668	0.0225	0.9484	-0.1089	0.8738	-0.0377	0.9096	0.0712	0.7902	-0.0306	0.9778	-0.0314	0.9404	-0.0009	0.9988
CG13160	CG13160	1628460_at	0.2808	0.7941	0.1652	0.4324	-0.5140	0.1592	-0.7736	0.3106	-0.4730	0.2944	0.3006	0.4809	0.0936	0.9824	-0.3294	0.7883	-0.4230	0.6993
yki	yorkie	1628461_a_at	-0.5038	0.0346	0.1340	0.4526	-0.0009	0.9973	0.0789	0.8921	-0.3839	0.0537	-0.4628	0.0160	0.2580	0.7324	0.1160	0.7679	-0.1420	0.6930
---	---	1628462_s_at	-0.1879	0.3516	0.0373	0.9188	-0.3486	0.0976	-0.3284	0.4568	-0.0236	0.9471	0.3048	0.1528	0.0478	0.9705	0.2257	0.5055	0.1779	0.6150
CG7242	lethal (3) c00064	1628463_at	0.4443	0.3240	-0.4465	0.6963	0.0379	0.9522	-0.0442	0.9664	1.0175	0.0037	1.0617	0.0018	-0.4969	0.8846	0.2880	0.8764	0.7849	0.5676
---	---	1628464_s_at	0.1786	0.2185	0.1973	0.4271	0.5265	0.0132	0.2451	0.6577	0.0788	0.8104	-0.1663	0.5186	-0.2602	0.6749	0.0321	0.9387	0.2923	0.2717
CG34417	CG3950	1628465_a_at	-1.0958	0.0038	-1.1953	0.0236	-1.7721	0.0000	-0.0898	0.8881	0.1315	0.5710	0.2213	0.2516	0.4095	0.5434	0.0273	0.9587	-0.3822	0.2398
Rpn6	Proteasome p44.5	1628466_s_at	0.3521	0.0560	0.4633	0.0485	0.4175	0.0161	0.0194	0.9759	0.0948	0.5811	0.0754	0.6428	0.1049	0.8692	0.2523	0.3075	0.1473	0.5869
CG14492	CG14492	1628467_at	0.1475	0.5692	0.0007	0.9986	0.3696	0.0491	-0.0223	0.9763	-0.0950	0.6507	-0.0727	0.7135	-0.1671	0.7826	-0.1029	0.7371	0.0642	0.8491
CG8738	CG8738	1628468_at	-1.7120	0.0082	-3.1520	0.0153	-1.8429	0.0021	1.2258	0.1140	1.3222	0.0118	0.0964	0.8588	-0.1338	0.9701	-0.0170	0.9935	0.1169	0.9263
CG32529	CG32529	1628469_a_at	-0.8304	0.1201	-0.9618	0.1648	-1.2331	0.0030	-0.0062	0.9956	0.2933	0.3275	0.2995	0.2599	0.2967	0.8979	0.1958	0.8734	-0.1009	0.9337
---	---	1628470_s_at	-0.1068	0.5936	0.0414	0.6829	-0.0290	0.9022	-0.1807	0.6681	-0.2074	0.3068	-0.0266	0.9132	0.0289	0.9734	-0.0565	0.8450	-0.0854	0.7253
---	---	1628471_at	0.0102	0.9547	0.1344	0.3102	0.0819	0.6695	-0.0169	0.9816	-0.0552	0.7963	-0.0383	0.8495	0.0984	0.8270	0.1093	0.5800	0.0109	0.9696
deltaCOP	delta-coatomer pr	1628472_at	0.5084	0.0211	0.7021	0.0220	0.9959	0.0004	0.1628	0.6490	0.0369	0.8720	-0.1259	0.4384	-0.2108	0.7330	0.3064	0.2319	0.5172	0.0859
ng2	new glue 2	1628473_at	0.1290	0.6085	-0.0577	0.7396	0.1564	0.2978	0.0300	0.9603	0.0520	0.7930	0.0220	0.9101	-0.0695	0.9309	-0.0336	0.9353	0.0360	0.9197
CG16712	CG16712	1628474_at	1.3947	0.0067	0.4881	0.5583	1.4299	0.0004	0.2741	0.5376	0.0250	0.9407	-0.2491	0.2310	-0.5900	0.7726	-0.8864	0.2990	-0.2965	0.7744
CG8678	CG8678	1628475_at	-0.4043	0.3209	-0.2151	0.4735	-0.2048	0.5264	-0.0082	0.9943	-0.4181	0.0853	-0.4099	0.0618	-0.0211	0.9943	-0.2609	0.7439	-0.2398	0.7607
PP2A-B	PP2A B' subunit	1628476_at	0.0066	0.9812	0.0165	0.9665	0.5040	0.0428	0.0484	0.9471	-0.5029	0.0230	-0.5513	0.0095	-0.4631	0.6584	-0.4138	0.3199	0.0493	0.9344
mab-2	mab-2	1628477_s_at	0.3619	0.3271	0.1651	0.3071	0.1448	0.4581	0.0389	0.9521	0.1146	0.5543	0.0757	0.6888	0.0502	0.9672	-0.0863	0.8420	-0.1364	0.7040
CG34415	CG33455	1628478_at	0.0949	0.7207	0.4371	0.3167	-0.0544	0.7908	-0.3305	0.4521	0.2644	0.2701	0.5949	0.0131	0.1870	0.8540	0.5588	0.1682	0.3718	0.3831

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15625	CG15625	1628479_at	0.2121	0.3551	-0.0963	0.4736	-0.1405	0.4979	-0.1367	0.7735	0.1680	0.4138	0.3047	0.0877	0.0320	0.9816	-0.0454	0.9265	-0.0774	0.8453
Gr98d	Gustatory recepto	1628480_at	0.1760	0.3366	-0.0532	0.7863	-0.0668	0.8127	-0.1069	0.8724	0.1600	0.5138	0.2669	0.1965	-0.0618	0.9238	-0.1171	0.6122	-0.0553	0.8412
mRpL1	mitochondrial ribo	1628481_at	-0.0242	0.9311	0.2733	0.2230	0.2839	0.2187	-0.1112	0.7975	-0.3507	0.0510	-0.2395	0.1259	-0.1706	0.8692	-0.0435	0.9484	0.1271	0.7983
Rab39	Rab39	1628482_at	0.2342	0.3327	0.0623	0.6664	0.2460	0.3081	0.0989	0.8676	-0.4679	0.0331	-0.5667	0.0091	0.0495	0.9737	-0.5911	0.1160	-0.6406	0.1184
CG3004	CG3004	1628483_at	0.2606	0.1331	0.3660	0.1503	0.4787	0.0220	0.2121	0.5498	0.1378	0.4722	-0.0743	0.7007	0.0806	0.9066	0.0891	0.7642	0.0084	0.9839
---	---	1628484_at	0.2525	0.2882	-0.0086	0.9507	0.1062	0.4706	-0.0172	0.9840	-0.1368	0.5216	-0.1196	0.5428	-0.0233	0.9860	-0.2683	0.3579	-0.2451	0.4192
CG13564	CG13564	1628485_at	0.0759	0.6453	0.0466	0.7186	0.4192	0.0821	0.0089	0.9937	0.0344	0.9077	0.0255	0.9228	-0.2192	0.7770	0.0665	0.8939	0.2857	0.4037
d4	d4	1628486_a_at	0.1435	0.3934	-0.2724	0.2711	-0.3654	0.0997	-0.0172	0.9803	0.3986	0.0274	0.4158	0.0145	0.1795	0.8002	0.1704	0.5860	-0.0091	0.9849
mri	mrityu	1628487_s_at	0.7512	0.0094	0.1766	0.1923	0.5344	0.0057	0.0335	0.9626	-0.1174	0.5661	-0.1509	0.3916	-0.2060	0.7513	-0.5408	0.0694	-0.3348	0.2474
CG6051	CG6051	1628488_at	-0.2060	0.5706	0.2219	0.5873	-0.1449	0.5672	-0.1309	0.8073	-0.4271	0.0501	-0.2962	0.1189	0.1182	0.9506	0.0005	1.0000	-0.1177	0.8783
dally	division abnormall	1628489_at	0.4397	0.0297	-0.1530	0.6920	-2.0358	0.0067	-1.1088	0.0144	0.8787	0.0032	1.9875	0.0001	0.8574	0.7215	0.3392	0.7883	-0.5182	0.6328
bigmax	bigmax	1628490_at	-0.2880	0.1049	0.0810	0.6732	0.2713	0.1634	-0.1144	0.8564	-1.0148	0.0017	-0.9004	0.0016	-0.3269	0.5765	-0.5262	0.0664	-0.1993	0.4683
---	---	1628491_at	-0.0638	0.7455	0.0252	0.8267	0.1623	0.4601	-0.0602	0.9375	-0.0743	0.7844	-0.0141	0.9597	-0.0603	0.9306	0.0245	0.9462	0.0848	0.7431
Aprt	Adenine phosphor	1628492_at	0.4360	0.0346	0.9982	0.0199	0.9321	0.0008	-0.0686	0.8903	-0.0748	0.6962	-0.0063	0.9761	-0.0700	0.9515	0.3225	0.3383	0.3926	0.2729
kal-1	Kallmann	1628493_at	-0.6382	0.2740	-0.3762	0.5237	-0.6817	0.0093	-0.1153	0.9302	0.0817	0.8732	0.1970	0.6145	0.2913	0.8953	0.3544	0.7042	0.0631	0.9578
Picot	Picot	1628494_a_at	-0.4700	0.0245	-0.1858	0.6579	-0.5759	0.0058	0.0040	0.9956	0.3321	0.0873	0.3281	0.0617	0.3791	0.7070	0.6628	0.1199	0.2837	0.5185
CG9220	CG9220	1628495_at	-0.1185	0.8604	2.3983	0.0301	1.7827	0.0011	-0.3933	0.6144	-1.7812	0.0015	-1.3879	0.0027	0.0908	0.9816	0.6303	0.4856	0.5394	0.5649
dpr16	dpr16	1628496_at	-0.0736	0.8192	-0.1922	0.3669	0.0281	0.8812	0.2134	0.6908	0.1356	0.6235	-0.0778	0.7781	-0.0799	0.9066	-0.0110	0.9808	0.0689	0.8209
CG7420	CG7420	1628497_at	-0.2899	0.1215	-0.4206	0.0962	-0.6007	0.0042	-0.2733	0.3909	-0.1391	0.4571	0.1343	0.4254	-0.1357	0.8202	-0.4205	0.0953	-0.2848	0.2614
obst-E	CG11142	1628498_at	-2.9731	0.0022	-1.2529	0.0087	-2.0453	0.0010	-0.6141	0.4171	-2.5942	0.0004	-1.9801	0.0008	0.2805	0.8672	-0.7514	0.2427	-1.0318	0.1479
CG14331	CG14331	1628499_at	-0.0605	0.7712	0.0878	0.5932	0.1804	0.3555	-0.0071	0.9956	-0.2552	0.3447	-0.2481	0.3029	-0.0963	0.8650	-0.0917	0.7245	0.0046	0.9903
CG32483	CG32483	1628500_at	0.4889	0.1467	0.3040	0.0925	0.3088	0.0702	-0.0582	0.9463	-0.0034	0.9922	0.0548	0.8460	-0.1427	0.8320	-0.1376	0.6512	0.0051	0.9920
---	---	1628501_at	0.2549	0.2708	0.2260	0.4962	0.0109	0.9666	-0.0620	0.9120	0.1139	0.5532	0.1759	0.2751	0.0996	0.9156	0.1463	0.6903	0.0468	0.9176
---	---	1628502_at	-0.0193	0.9150	-0.0440	0.7434	-0.1321	0.4164	0.0458	0.9507	0.1298	0.5582	0.0841	0.6977	0.1686	0.7485	0.1778	0.4274	0.0092	0.9806
CG30424	CG30424	1628503_at	0.9871	0.0184	0.6743	0.0809	1.2047	0.0014	-0.0055	0.9956	-0.3829	0.1902	-0.3774	0.1497	-0.1434	0.9342	-0.3875	0.5024	-0.2441	0.6965
CG16852	CG16852	1628504_at	-0.0072	0.9784	0.0495	0.7319	0.0982	0.6551	-0.1193	0.8304	-0.1276	0.5817	-0.0083	0.9754	-0.0431	0.9589	0.0354	0.9263	0.0786	0.7836
CG11964	CG11964	1628505_at	0.1558	0.3001	-0.0087	0.9763	-0.0658	0.6584	0.0499	0.9255	0.2340	0.1439	0.1840	0.1997	0.0883	0.9016	0.1037	0.7344	0.0154	0.9695
---	---	1628506_a_at	0.1598	0.3591	0.1111	0.4914	0.2148	0.3274	0.1260	0.8189	0.0550	0.8436	-0.0710	0.7636	0.0090	0.9946	-0.0237	0.9628	-0.0327	0.9394
CG10053	CG10053	1628507_at	0.2392	0.2013	0.2880	0.3481	0.6040	0.0072	0.1531	0.7028	0.1282	0.5114	-0.0249	0.9111	0.0001	0.9999	0.2043	0.5162	0.2042	0.5209
CG30062	CG30062	1628508_at	0.3106	0.0978	0.0764	0.5608	-0.0349	0.8675	-0.1350	0.7845	0.0622	0.8031	0.1972	0.2839	0.0382	0.9717	-0.1274	0.6673	-0.1656	0.5612
CG7364	CG7364	1628509_at	0.4750	0.0237	0.5170	0.1123	0.7365	0.0024	-0.1262	0.7658	0.2102	0.2452	0.3364	0.0433	-0.2581	0.7707	0.2492	0.5254	0.5073	0.2037
---	---	1628510_at	0.0483	0.7981	0.0220	0.8274	0.0979	0.5248	0.0253	0.9643	-0.0227	0.9138	-0.0480	0.7724	0.0240	0.9816	-0.0528	0.8684	-0.0768	0.7733
---	---	1628511_at	0.1728	0.3039	0.2752	0.1904	0.2387	0.2230	0.0392	0.9562	-0.0492	0.8437	-0.0884	0.6585	0.1310	0.8122	0.0048	0.9925	-0.1262	0.6053
CG7441 /// DmirCG7441 ///	CG7441	1628512_at	0.2681	0.5044	0.3875	0.1949	0.8893	0.0069	-0.1817	0.7617	-0.2794	0.2809	-0.0977	0.7219	-0.3621	0.7266	-0.0001	1.0000	0.3620	0.4301
---	---	1628513_at	0.3633	0.4757	0.2554	0.0529	0.5281	0.0295	0.1372	0.8358	0.1175	0.6800	-0.0197	0.9503	-0.0713	0.9619	0.0985	0.8667	0.1698	0.7189
Ipk2	Ipk2	1628514_at	-0.4397	0.1008	-0.5770	0.0596	-0.7602	0.0367	0.0020	0.9981	-0.0160	0.9567	-0.0180	0.9422	0.1931	0.9046	-0.1400	0.8631	-0.3330	0.5929
---	---	1628515_at	0.1161	0.6033	-0.0275	0.8102	-0.0121	0.9571	0.1913	0.6557	0.1216	0.5881	-0.0698	0.7570	0.1737	0.8133	0.0170	0.9739	-0.1567	0.6328
CG15432	CG15432	1628516_at	-0.0691	0.7642	-0.2642	0.0621	-0.0196	0.9322	0.1384	0.7575	-0.0111	0.9703	-0.1495	0.4048	-0.1798	0.8193	-0.2111	0.5283	-0.0313	0.9444
---	---	1628517_at	0.2036	0.2882	0.0508	0.6371	0.1245	0.5854	0.0574	0.9308	0.0537	0.8247	-0.0036	0.9878	-0.0135	0.9875	-0.0545	0.8326	-0.0410	0.8751
---	---	1628518_at	0.3094	0.1343	-0.0780	0.4953	-0.2333	0.1676	-0.3914	0.3558	0.1977	0.4307	0.5892	0.0146	-0.1382	0.8480	-0.1319	0.6969	0.0063	0.9902
CG15741	CG15741	1628519_at	0.0999	0.5568	-0.0638	0.6768	-0.0025	0.9935	0.0256	0.9819	0.0409	0.9126	0.0153	0.9638	-0.0383	0.9653	-0.1639	0.4982	-0.1255	0.6209
---	---	1628520_at	0.0811	0.5660	0.2302	0.2086	0.1685	0.2432	-0.1109	0.7351	-0.1567	0.2861	-0.0457	0.7775	0.0601	0.9457	0.0902	0.7860	0.0301	0.9356
Gmd	GDP-mannose 4,(	1628521_at	0.5795	0.0274	1.0211	0.0562	1.2741	0.0001	0.1046	0.8701	0.2896	0.1938	0.1850	0.3711	-0.1354	0.9056	0.6461	0.1199	0.7816	0.0906
---	---	1628522_at	-0.1109	0.4157	0.0134	0.9370	0.1982	0.2973	0.0926	0.8636	0.0566	0.8112	-0.0360	0.8735	-0.0966	0.9142	0.0747	0.8665	0.1713	0.6150
---	---	1628523_at	0.0252	0.8981	0.0260	0.8408	0.0805	0.6025	0.0377	0.9488	0.0065	0.9795	-0.0312	0.8716	-0.0681	0.9246	-0.0376	0.9204	0.0305	0.9277
CG31008	CG31008	1628524_at	0.2256	0.3457	-0.0088	0.9365	0.1107	0.5038	0.0841	0.9133	0.1773	0.4877	0.0932	0.7198	0.0965	0.9142	0.1049	0.7883	0.0083	0.9868
CG6405	CG6405	1628525_at	0.0655	0.8135	-0.1704	0.2796	0.1816	0.5658	0.1659	0.8068	0.0866	0.7932	-0.0793	0.7931	-0.1075	0.9330	0.0180	0.9812	0.1255	0.8003
CG15337	CG15337	1628526_at	-0.1575	0.3880	0.3758	0.1015	0.2047	0.3743	-0.1419	0.7604	-0.2681	0.1779	-0.1262	0.5151	0.1007	0.9095	0.2432	0.4409	0.1426	0.6833
---	---	1628527_at	0.1327	0.5805	0.0225	0.9033	0.2249	0.1600	-0.0200	0.9854	-0.0133	0.9727	0.0068	0.9829	-0.0873	0.8599	-0.0458	0.8721	0.0415	0.8779
---	---	1628528_at	0.0920	0.5851	0.0135	0.8976	0.0635	0.7050	-0.0850	0.8605	-0.0852	0.6658	-0.0002	0.9995	-0.1284	0.7997	-0.0720	0.7822	0.0564	0.8310
---	---	1628529_a_at	0.0275	0.9079	0.5341	0.0911	0.6813	0.0016	0.0926	0.8686	-0.3982	0.0490	-0.4909	0.0132	-0.0331	0.9816	0.1784	0.6024	0.2115	0.5314
CG10584	CG10584	1628530_at	-1.0434	0.0096	0.0061	0.9882	0.2845	0.2223	-0.0489	0.9518	-1.0698	0.0012	-1.0209	0.0009	-0.2413	0.8465	0.0631	0.9402	0.3043	0.5816
Rpn11	yippee interacting	1628531_at	0.2981	0.1744	0.4980	0.1133	0.8444	0.0009	0.1668	0.6475	-0.0953	0.6277	-0.2621	0.0965	-0.1396	0.8655	0.1553	0.6695	0.2949	0.3819
CG15768	CG15768	1628532_at	0.1299	0.6039	0.0287	0.7775	-0.0724	0.7158	-0.0411	0.9580	0.1653	0.4396	0.2064	0.2671	0.0282	0.9816	0.1545	0.6189	0.1262	0.6956
CG13578	CG13578	1628533_at	-0.1078	0.5183	0.1026	0.5779	0.0135	0.9534	0.0596	0.8940	-0.2160	0.1445	-0.2756	0.0440	0.1874	0.7780	0.1987	0.5064	0.0113	0.9819
CG15059	CG15059	1628534_at	-0.0482	0.8091	0.0609	0.6751	0.2899	0.1205	0.1185	0.8671	0.0001	0.9997	-0.1184	0.6446	-0.0513	0.9467	0.1085	0.6762	0.1599	0.5114
CG33771	CG33771	1628535_at	0.1726	0.3453	0.0917	0.4744	0.2050	0.2628	-0.1014	0.8701	-0.0655	0.8066	0.0359	0.8902	-0.1685	0.7644	-0.0264	0.9451	0.1422	0.5700
CG11880	CG11880	1628536_s_at	1.3323	0.0107	1.2466	0.0264	1.5956	0.0017	0.8754	0.2506	1.5703	0.0050	0.6949	0.0873	0.4590	0.7070	1.7209	0.0194	1.2620	0.0505
CG8368	RNase H-like	1628537_s_at	0.4350	0.1032	0.4120	0.3309	0.5455	0.0173	0.2449	0.5771	0.3343	0.1361	0.0894	0.7076	0.0019	0.9997	0.2232	0.6876	0.2212	0.6872
CG34398	CG15867	1628538_at	0.1759	0.4957	0.1969	0.1811	0.0090	0.9674	-0.2201	0.4611	-0.1448	0.3814	0.0753	0.6527	0.0746	0.9445	-0.0988	0.8132	-0.1735	0.6186
CG31174	CG31174	1628539_at	-0.4149	0.0423	-0.3615	0.4120	-0.1991	0.2836	-0.2987	0.3862	-0.2062	0.2943	0.0925	0.6442	0.0470	0.9816	0.2421	0.6256	0.1950	0.7055
---	---	1628540_s_at	-0.0751	0.6684	-0.0121	0.9549	-0.0837	0.7005	0.0080	0.9937	-0.1448	0.4728	-0.1527	0.3941	-0.0453	0.9616	-0.1051	0.7299	-0.0598	0.8626
Takl1	Tak1-like 1	1628541_at	0.1394	0.8861	0.0577	0.5827	0.3161	0.2977	-0.1649	0.9375	-0.3853	0.5572	-0.2204	0.7366	-0.0717	0.9831	-0.1760	0.8760	-0.1043	0.9231
---	---	1628542_at	0.3996	0.1264	0.2477	0.3334	0.1246	0.7350	-0.3543	0.5255	-0.2810	0.3450	0.0733	0.8255	-0.0405	0.9838	-0.0390	0.9590	0.0016	0.9987
Fas3	Fasciclin III	1628543_a_at	-1.0272	0.0302	0.1116	0.8630	0.6089	0.1826	0.6959	0.2905	-0.1177	0.8046	-0.8136	0.0268	0.3712	0.8740	1.2395	0.1655	0.8683	0.3592
CG33642	CG33642	1628544_at	0.0364	0.8163	0.0019	0.9948	0.0840	0.5935	0.0706	0.9182	0.0151	0.9623	-0.0555	0.8165	-0.0014	0.9990	-0.0115	0.9704	-0.0101	0.9723
CG13195	CG13195	1628545_at	0.2107	0.3226	0.2260	0.1292	0.2182	0.3828	0.0615	0.9425	0.0088	0.9809	-0.0527	0.8516	-0.1529	0.8012	0.0468	0.9036	0.1997	0.4424
CG31210	CG31210	1628546_at	0.1467	0.4148	0.0892	0.5564	0.0825	0.6441	-0.0210	0.9745	-0.0187	0.9350	0.0023	0.9911	0.0945	0.8963	-0.1243	0.6803	-0.2188	0.4308
CG5167	CG5167	1628547_at	-0.2554	0.5302	0.4282	0.1214	1.5285	0.0001	0.2524	0.6890	-1.1649	0.0027	-1.4173	0.0007	-0.7743	0.4128	-0.3056	0.5647	0.4688	0.3678

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Apc	adenomatous poly	1628548_at	0.7584	0.0597	0.1264	0.7992	-0.2298	0.2603	-0.0627	0.9308	0.7456	0.0045	0.8083	0.0019	0.3431	0.8326	0.1919	0.8304	-0.1512	0.8655
---	---	1628549_x_at	-0.1720	0.4726	-0.1186	0.4665	-0.2169	0.1637	0.1159	0.8857	0.0092	0.9818	-0.1066	0.7136	0.1138	0.8714	-0.0788	0.8319	-0.1925	0.5059
Pmm45A	Phosphomannomi	1628550_a_at	-0.5012	0.0598	-0.0812	0.5135	0.1554	0.4203	0.4193	0.3361	0.0890	0.7683	-0.3303	0.1409	0.1516	0.8016	0.5471	0.0514	0.3954	0.1438
Cyp303a1	lethal(2)35Fb	1628551_at	0.1286	0.5972	0.0031	0.9805	0.3189	0.0455	0.1248	0.7970	-0.0534	0.8315	-0.1782	0.3214	-0.1154	0.8202	-0.1861	0.3685	-0.0707	0.7761
CG7156	CG7156	1628552_at	-0.9870	0.0184	0.1295	0.3705	0.3213	0.1080	-0.1886	0.7680	-1.2214	0.0016	-1.0328	0.0019	-0.2915	0.7464	0.1277	0.7899	0.4191	0.2869
---	---	1628553_at	0.0530	0.7807	0.0954	0.4932	0.2770	0.0952	0.0474	0.9413	-0.0055	0.9840	-0.0528	0.7989	-0.0474	0.9467	-0.0711	0.7932	-0.0237	0.9372
---	---	1628554_at	-0.0402	0.8294	-0.2913	0.3041	-0.1710	0.3539	0.1045	0.8908	0.3395	0.1807	0.2351	0.3107	-0.0235	0.9788	0.0283	0.9318	0.0518	0.8444
Osi19	Osiris	1628555_at	0.1784	0.3591	0.0125	0.9153	0.1200	0.4111	0.1797	0.5539	0.1305	0.4209	-0.0492	0.7779	0.0550	0.9101	-0.0108	0.9701	-0.0659	0.7434
Hr39	Hormone receptor	1628556_s_at	1.8330	0.0193	1.9692	0.0321	2.2267	0.0009	-0.3533	0.4397	-0.0455	0.8932	0.3077	0.1669	-0.2644	0.9112	0.2411	0.8287	0.5054	0.5722
CG2063	CG2063	1628557_at	0.2093	0.1864	0.1212	0.4521	0.0345	0.8277	0.0105	0.9866	0.2787	0.0728	0.2682	0.0554	0.0296	0.9767	0.2037	0.3941	0.1741	0.4865
CG30022	CG30022	1628558_at	0.2899	0.1389	0.4148	0.0689	0.7135	0.0015	0.2187	0.5008	0.2038	0.2376	-0.0149	0.9458	-0.0627	0.9309	0.2440	0.2849	0.3067	0.2111
CG14876	CG14876	1628559_at	-0.0805	0.5872	-0.0010	0.9954	0.1297	0.4900	0.1895	0.7596	0.2067	0.4550	0.0172	0.9592	0.2007	0.7118	0.2573	0.2408	0.0566	0.8461
CG9344 /// DyakCG9344	CG9344	1628560_at	0.6520	0.0267	0.9820	0.0207	0.9611	0.0022	-0.1167	0.8009	-0.2762	0.1365	-0.1595	0.3510	-0.0451	0.9778	0.1126	0.8247	0.1576	0.7190
CG31515	CG31515	1628561_at	0.0320	0.8676	0.0791	0.5407	0.2017	0.1925	0.0992	0.8281	0.0872	0.6615	-0.0120	0.9573	-0.1784	0.7070	0.0424	0.8830	0.2208	0.2833
mod(mdg4)	Modifier67.2	1628562_s_at	0.4256	0.0333	0.6783	0.0675	0.6274	0.0053	-0.0755	0.8791	0.2718	0.1063	0.3472	0.0290	-0.0002	0.9999	0.4882	0.1175	0.4884	0.1409
CG33181	CG33181	1628563_at	-3.9030	0.0004	-0.8551	0.3198	-3.2821	0.0009	-1.9826	0.0154	-2.9270	0.0003	-0.9443	0.0212	0.2762	0.9309	-0.0973	0.9531	-0.3735	0.7527
Spn43Ad	Spn43Ad	1628564_at	-0.4659	0.0944	0.3847	0.4517	0.0769	0.8377	-0.1565	0.9018	-1.3730	0.0058	-1.2165	0.0060	0.2066	0.8878	-0.4127	0.4541	-0.6193	0.2757
CG11454	CG11454	1628565_at	-0.0873	0.7692	0.2074	0.2593	0.6833	0.0022	0.2422	0.5311	-0.2835	0.1581	-0.5257	0.0107	-0.2379	0.7493	0.0312	0.9508	0.2691	0.4054
---	---	1628566_at	0.0884	0.6601	-0.2647	0.2820	-0.5629	0.0591	0.0871	0.9376	0.3979	0.2180	0.3108	0.2895	0.0656	0.9365	-0.1349	0.6314	-0.2006	0.4568
CG13565	CG13565	1628567_at	0.2416	0.3730	0.0290	0.7740	-0.0502	0.8209	0.0093	0.9922	0.0984	0.6664	0.0891	0.6701	0.0894	0.9095	-0.0574	0.8880	-0.1468	0.6279
---	---	1628568_at	0.1967	0.4588	0.1191	0.4819	0.3831	0.0505	0.0402	0.9666	0.0809	0.7932	0.0408	0.8926	-0.1650	0.7726	0.0306	0.9390	0.1956	0.4388
---	---	1628569_at	0.0084	0.9705	-0.0664	0.5681	-0.4462	0.0204	-0.2271	0.6641	0.2836	0.2584	0.5107	0.0299	-0.0873	0.8541	-0.0425	0.8792	0.0447	0.8618
CG4021	CG4021	1628570_at	0.2228	0.1436	0.0483	0.6913	-0.0241	0.9083	-0.0595	0.9023	-0.1392	0.3882	-0.0796	0.6178	0.1359	0.7953	-0.1868	0.3899	-0.3227	0.1659
---	---	1628571_at	-0.0003	0.9994	0.0624	0.7861	0.0890	0.6582	0.0546	0.9451	-0.0393	0.8990	-0.0938	0.6925	0.0308	0.9816	0.0498	0.9085	0.0190	0.9638
su(rdgB)69	suppressor (rdgB)	1628572_s_at	-0.6078	0.0265	-0.2487	0.0571	-0.4131	0.0782	-0.1527	0.8249	-0.0719	0.8334	0.0808	0.7869	-0.2465	0.6898	-0.0391	0.9207	0.2074	0.4182
ATPsyn-b	Fo-ATP synthase	1628573_a_at	-0.3628	0.0797	-0.1386	0.5785	-0.1553	0.4430	-0.1598	0.7028	-0.5622	0.0106	-0.4024	0.0273	-0.0718	0.9449	-0.3930	0.1957	-0.3211	0.3170
CG14402	CG14402	1628574_at	0.0655	0.6803	0.1562	0.1518	0.1150	0.5219	-0.0362	0.9648	-0.2029	0.3598	-0.1667	0.4109	-0.0231	0.9776	-0.0613	0.8101	-0.0382	0.8880
---	---	1628575_at	-0.1453	0.6117	0.1000	0.4282	0.2864	0.2355	-0.1265	0.7503	-0.1343	0.4547	-0.0078	0.9711	0.0145	0.9914	0.1235	0.7100	0.1090	0.7482
CG10883	CG10883	1628576_at	0.0490	0.8159	0.0946	0.5917	0.1360	0.5513	0.0023	0.9983	0.0187	0.9582	0.0165	0.9573	0.1023	0.9128	0.1431	0.7000	0.0408	0.9289
---	---	1628577_at	0.0956	0.6027	-0.1539	0.3887	-0.0338	0.8867	0.3351	0.2899	0.5425	0.0106	0.2074	0.2131	0.1005	0.8909	-0.0049	0.9935	-0.1054	0.7410
CG10651	CG10651	1628578_at	-0.0469	0.7688	0.0450	0.6681	0.1279	0.4477	0.0758	0.8942	-0.1196	0.5592	-0.1953	0.2519	-0.1309	0.8141	-0.0426	0.9037	0.0883	0.7398
---	---	1628579_at	0.3385	0.2981	-0.5421	0.4606	-0.3282	0.2436	0.0994	0.9410	0.3102	0.4358	0.2108	0.5823	-0.2554	0.8940	-0.6471	0.3571	-0.3917	0.6129
Cdep	Cdep	1628580_a_at	-0.1133	0.7196	-0.2926	0.0947	-0.2201	0.4106	0.0109	0.9912	-0.1474	0.5153	-0.1583	0.4305	-0.0659	0.9742	-0.2067	0.7260	-0.1408	0.8256
CG3884	CG3884	1628581_at	0.7434	0.6913	-0.1821	0.8477	0.5816	0.3611	0.9733	0.7608	0.8931	0.5469	-0.0802	0.9632	0.3000	0.9610	-0.0765	0.9808	-0.3764	0.8686
CG10738	CG10738	1628582_at	0.1401	0.4437	0.1630	0.4612	0.2143	0.2537	-0.1183	0.8474	-0.0791	0.7705	0.0393	0.8830	-0.1644	0.8023	-0.1429	0.6281	0.0215	0.9563
CG31688	CG31688	1628583_at	-0.0897	0.8981	1.1988	0.2271	0.2514	0.4343	-0.0128	0.9909	-0.4683	0.0557	-0.4555	0.0402	0.9044	0.7464	0.7198	0.5632	-0.1846	0.9101
Cyp305a1	Cyp305a1	1628584_at	4.3106	0.0008	2.5624	0.0025	4.7107	0.0000	1.8310	0.0300	2.1227	0.0015	0.2917	0.5123	-0.3267	0.7980	0.4969	0.3397	0.8236	0.1491
Gyc-89Da	Guanylyl cyclase	1628585_at	-0.0565	0.7343	-0.0569	0.8433	0.1617	0.5147	0.1335	0.7556	-0.0035	0.9899	-0.1370	0.4289	-0.0368	0.9816	-0.1253	0.7829	-0.0884	0.8524
CG15642	CG15642	1628586_at	0.1380	0.5146	0.3717	0.1178	0.4305	0.0286	-0.0421	0.9507	-0.1968	0.2998	-0.1547	0.3735	-0.1621	0.8270	-0.0385	0.9387	0.1236	0.7247
CG8607	CG8607	1628587_at	-0.7676	0.1718	-0.0983	0.7273	0.2662	0.1180	0.0759	0.8771	-0.1388	0.4267	-0.2147	0.1533	-0.1699	0.9390	0.4388	0.5386	0.6087	0.3838
Not1	Not1	1628588_at	0.0205	0.9481	-0.0332	0.8311	0.2057	0.2727	0.1325	0.8479	0.0167	0.9659	-0.1158	0.6586	-0.2269	0.8049	-0.0364	0.9528	0.1905	0.6429
CG5010	CG5010	1628589_at	0.6285	0.0769	-0.0191	0.9747	-0.1017	0.5746	0.1842	0.6325	0.6615	0.0046	0.4773	0.0122	0.3866	0.7908	0.2060	0.7906	-0.1806	0.8152
CG12920	CG12920	1628590_at	0.0147	0.9650	0.0285	0.8730	0.1518	0.3466	0.0388	0.9674	-0.0938	0.7519	-0.1326	0.5956	-0.1703	0.8609	-0.0359	0.9550	0.1344	0.7694
Cyp313a3	Cyp313a3	1628591_at	-1.9884	0.0023	1.5427	0.0250	-1.4420	0.0323	-2.8039	0.0082	-4.0310	0.0002	-1.2271	0.0153	0.1799	0.9469	-0.6922	0.3981	-0.8721	0.3033
CG15539	CG15539	1628592_at	0.1362	0.6123	0.0089	0.9475	-0.0438	0.7811	0.1366	0.7117	0.2006	0.2306	0.0640	0.7199	0.0813	0.9309	-0.1171	0.7424	-0.1984	0.5341
CG33252	CG33252	1628593_at	-0.1319	0.5154	-0.1919	0.3081	-0.1741	0.2122	0.1097	0.7929	0.0432	0.8442	-0.0665	0.7129	0.1932	0.8157	-0.0072	0.9925	-0.2004	0.5830
rut	rutabaga	1628594_at	-3.3461	0.0025	-1.0434	0.0948	-2.8128	0.0003	-0.2788	0.8350	-0.2899	0.6015	-0.0111	0.9861	1.3527	0.4440	1.8343	0.0661	0.4816	0.6286
UbcD4	Ubiquitin conjugat	1628595_at	0.0505	0.7667	0.0842	0.5396	0.3100	0.1861	0.1059	0.8671	0.0391	0.8979	-0.0668	0.7884	-0.1279	0.8513	0.0303	0.9476	0.1582	0.6024
CG4610	CG4610	1628596_at	-0.1233	0.5710	-0.2766	0.3887	-0.4454	0.1400	-0.0026	0.9982	-0.0051	0.9898	-0.0025	0.9938	0.2566	0.8016	-0.1472	0.7787	-0.4038	0.3572
---	---	1628597_at	0.0707	0.6443	0.0798	0.4109	0.0581	0.7056	-0.2410	0.4814	-0.1093	0.5825	0.1317	0.4456	0.0344	0.9514	0.0856	0.6387	0.0512	0.8025
sec23	sec23	1628598_s_at	0.4773	0.3444	0.5348	0.1768	0.9188	0.0007	0.2587	0.4910	0.3554	0.0780	0.0967	0.6380	0.0246	0.9939	0.6674	0.3543	0.6429	0.3921
Psf2	Psf2	1628599_at	1.4197	0.0022	0.5370	0.3973	0.7804	0.0015	0.2022	0.5688	0.4440	0.0230	0.2418	0.1360	0.0144	0.9963	-0.3376	0.6380	-0.3520	0.6211
CG12448	CG12448	1628600_at	0.0427	0.8167	-0.0110	0.9664	-0.0843	0.7325	-0.1244	0.7743	-0.1635	0.3784	-0.0392	0.8527	0.0630	0.9467	-0.1655	0.5855	-0.2286	0.4374
Klp64D	Kinesin-like protei	1628601_at	-1.0256	0.0081	0.3566	0.1795	0.6832	0.0142	0.0970	0.9004	-0.9465	0.0034	-1.0434	0.0013	-0.3132	0.7196	0.2843	0.4302	0.5975	0.1300
CG9248	CG9248	1628602_a_at	0.3691	0.1318	0.5364	0.1510	0.9453	0.0004	0.1338	0.6539	0.0713	0.6619	-0.0625	0.6776	-0.1651	0.8744	0.1579	0.7439	0.3230	0.4457
RpL26	Ribosomal protein	1628603_at	0.2877	0.0764	2.4595	0.0041	2.6439	0.0000	0.2063	0.7271	-2.2490	0.0001	-2.4553	0.0000	-0.0583	0.9466	-0.0323	0.9402	0.0260	0.9434
D	fish-hook	1628604_at	-2.3910	0.0009	-0.8005	0.2789	-2.2829	0.0021	-1.3322	0.1615	-2.3854	0.0023	-1.0532	0.0469	-0.0942	0.9647	-0.8774	0.1287	-0.7833	0.1972
CG5662	CG5662	1628605_at	0.1185	0.6310	-0.0411	0.7381	0.3731	0.1095	0.1780	0.7604	0.0548	0.8693	-0.1232	0.6314	-0.1087	0.8875	-0.0107	0.9844	0.0979	0.7734
---	---	1628606_at	0.3008	0.2185	-0.0297	0.8520	0.1796	0.3462	0.1782	0.6154	0.2236	0.2064	0.0454	0.8216	0.1495	0.7791	0.0234	0.9501	-0.1261	0.6133
---	---	1628607_at	-0.1325	0.5127	0.0612	0.5572	0.0328	0.8608	0.0658	0.9247	-0.1559	0.4749	-0.2218	0.2356	-0.0195	0.9831	-0.1219	0.5972	-0.1024	0.6630
CG17475	CG17475	1628608_at	-0.9493	0.8201	0.1901	0.4148	-0.6761	0.0350	-0.8825	0.9070	-3.9610	0.1032	-3.0785	0.1560	0.0200	0.9994	-2.8615	0.4372	-2.8815	0.4475
CG15709	CG15709	1628609_at	0.2502	0.1916	0.0311	0.7556	-0.0233	0.9263	-0.1003	0.8189	-0.0007	0.9979	0.0996	0.5542	0.0103	0.9943	0.0627	0.9004	0.0523	0.9088
CG7380 /// DyakCG7380	CG7380	1628610_at	-0.3767	0.1569	-0.4547	0.0144	-0.2889	0.2099	0.1318	0.7795	0.4548	0.0277	0.3230	0.0675	-0.0333	0.9860	0.4648	0.2520	0.4981	0.2557
CG11241	CG11241	1628611_at	3.5595	0.0008	2.0293	0.0136	3.6283	0.0000	0.8624	0.2747	0.4916	0.2944	-0.3708	0.3909	-0.8575	0.5754	-1.1438	0.1037	-0.2864	0.7120
CG31860	CG31860	1628612_at	0.1013	0.6640	-0.2144	0.1529	-0.0640	0.7890	0.2297	0.6720	0.4195	0.1023	0.1898	0.4318	-0.1820	0.7493	-0.1939	0.4242	-0.0119	0.9756
CG30492	CG30492	1628613_a_at	-0.6749	0.0884	-0.3422	0.4543	-0.5923	0.0057	-0.4715	0.3317	-0.8076	0.0112	-0.3361	0.1792	-0.1684	0.9101	-0.4177	0.4308	-0.2493	0.6668
CG14014	CG14014	1628614_at	0.3469	0.0665	0.0040	0.9905	0.1229	0.5502	-0.0463	0.9381	0.1236	0.5046	0.1699	0.2840	-0.0913	0.9445	-0.2081	0.6265	-0.1169	0.8119
CG10375	CG10375	1628615_at	-0.1613	0.4463	0.6280	0.0297	0.6284	0.0062	-0.0038	0.9956	-0.9166	0.0007	-0.9129	0.0004	-0.0623	0.9421	-0.1861	0.4951	-0.1237	0.6683
CG8419	CG8419	1628616_at	3.9115	0.0004	2.5801	0.0080	4.4417	0.0000	0.8633	0.1138	0.6044	0.0671	-0.2588	0.3934	-0.1038	0.9296	-0.0733	0.8981	0.0305	0.9549

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
dro4	drosomycin-F	1628617_at	0.5949	0.4962	0.3925	0.1163	0.3530	0.0618	-0.0796	0.9538	-0.2687	0.4944	-0.1892	0.6187	-0.0309	0.9950	-0.5934	0.6199	-0.5626	0.6391
CG9240	CG9240	1628618_at	-0.6400	0.0075	-0.1538	0.3640	0.2079	0.3127	0.1894	0.6473	-0.4270	0.0396	-0.6164	0.0050	-0.1282	0.8680	0.0691	0.8738	0.1973	0.5391
---	---	1628619_at	0.0175	0.9450	-0.0100	0.9635	-0.2762	0.0879	-0.1906	0.5528	-0.0634	0.7475	0.1272	0.4123	0.0096	0.9928	-0.1060	0.7054	-0.1156	0.6683
CG32806	CG32806	1628620_at	0.1775	0.4915	0.0341	0.7401	-0.1008	0.5949	-0.1972	0.7168	-0.0597	0.8527	0.1375	0.5769	-0.0409	0.9522	-0.1078	0.6264	-0.0669	0.7836
mRpS18C	mitochondrial ribo	1628621_at	0.2432	0.2362	0.0756	0.7936	-0.3600	0.0426	0.1494	0.6533	0.6289	0.0029	0.4795	0.0057	0.5002	0.6272	0.4070	0.3285	-0.0932	0.8707
CG3409	CG3409	1628622_s_at	0.1662	0.3244	0.0904	0.4888	0.3850	0.0913	0.0214	0.9810	0.0227	0.9432	0.0013	0.9963	-0.1078	0.8740	-0.0193	0.9649	0.0885	0.7788
CG9904	CG9904	1628623_at	0.4618	0.3407	0.1571	0.7334	0.7385	0.0064	-0.0766	0.9599	-0.0495	0.9298	0.0271	0.9573	-0.5768	0.6955	-0.2515	0.7267	0.3253	0.6288
---	---	1628624_s_at	0.1096	0.5598	0.0219	0.8546	0.4840	0.0711	0.3572	0.2654	0.1129	0.5833	-0.2443	0.1482	-0.0600	0.9174	-0.0221	0.9460	0.0380	0.8905
CG12926	CG12926	1628625_at	-2.6633	0.0006	-0.8528	0.1074	-0.7385	0.0651	-0.2812	0.6506	-1.2105	0.0025	-0.9292	0.0049	-0.3370	0.8534	0.4182	0.6014	0.7552	0.3253
CG32537	CG32537	1628626_at	-0.4291	0.4379	0.4676	0.4598	0.7996	0.0210	0.0949	0.8950	-0.5843	0.0228	-0.6792	0.0072	0.1752	0.9672	0.3854	0.7810	0.2102	0.8918
CG11807	CG11807	1628627_at	-0.1661	0.4168	-0.1263	0.5196	-0.4437	0.0139	-0.1898	0.6425	0.1109	0.6139	0.3006	0.0913	0.1642	0.7726	0.1928	0.4294	0.0285	0.9341
CalpA	Calpain A	1628628_at	-1.2866	0.0015	-0.6036	0.0395	-0.5172	0.0266	-0.1010	0.8189	-0.7383	0.0019	-0.6373	0.0022	-0.0822	0.9421	-0.0521	0.9246	0.0301	0.9512
CG30038	CG30038	1628629_at	-0.1507	0.5568	-0.8368	0.0471	-0.2819	0.1091	0.2217	0.7461	0.6478	0.0373	0.4261	0.1108	-0.3133	0.6496	0.0095	0.9865	0.3228	0.2622
Taf13	TBP-associated fa	1628630_at	0.1370	0.4063	0.4008	0.1547	0.6846	0.0050	0.0240	0.9777	-0.2923	0.1759	-0.3163	0.1025	-0.2194	0.7070	-0.1072	0.6977	0.1122	0.6763
---	---	1628631_at	-0.0101	0.9728	0.0479	0.6478	0.3022	0.1374	0.0643	0.9435	-0.0615	0.8530	-0.1258	0.6308	-0.2318	0.7149	0.0745	0.8329	0.3063	0.2622
Paip2	polyA-binding prot	1628632_at	0.6159	0.1282	-0.1509	0.7266	-0.3612	0.5719	-0.3552	0.6514	-0.2750	0.4900	0.0802	0.8563	-0.0251	0.9952	-1.0318	0.2680	-1.0067	0.3081
Pig1	gland specific gen	1628633_at	0.2643	0.1454	0.0658	0.8545	0.0626	0.7399	0.0180	0.9760	0.0243	0.9052	0.0063	0.9727	0.1410	0.8837	0.0266	0.9648	-0.1144	0.7962
Papst2	CG7853	1628634_at	0.1126	0.6559	0.2309	0.5237	0.4239	0.0162	-0.0870	0.8405	0.0884	0.6208	0.1754	0.2248	-0.2270	0.8400	0.2468	0.6209	0.4738	0.3175
Osi9	Osiris	1628635_at	0.2775	0.1608	-0.0566	0.5852	-0.0323	0.8616	-0.0516	0.9265	0.1328	0.4469	0.1844	0.2202	-0.0160	0.9892	-0.2149	0.3706	-0.1989	0.4268
CG11943 /// DmirCG11943	CG11943	1628636_a_at	0.4229	0.4396	-0.0225	0.9810	0.1959	0.3767	0.0254	0.9777	0.6037	0.0145	0.5782	0.0108	-0.2143	0.9514	0.1687	0.9158	0.3830	0.7484
ubl	ubiquitin like	1628637_at	-0.2907	0.2235	0.7589	0.0818	0.2609	0.2073	-0.3803	0.4167	-0.9482	0.0039	-0.5679	0.0228	0.1707	0.8609	0.2125	0.6156	0.0418	0.9398
CG7200	CG7200	1628638_at	0.7966	0.0220	0.2253	0.4911	0.0032	0.9898	-0.0937	0.8699	0.6227	0.0081	0.7164	0.0026	0.1077	0.9409	0.0067	0.9942	-0.1011	0.8652
CG5791	CG5791	1628639_at	2.1102	0.0160	0.9123	0.3433	1.4557	0.0169	0.2392	0.8498	0.7353	0.1091	0.4961	0.2287	-0.5305	0.8609	-0.5962	0.6573	-0.0658	0.9725
CG14627	CG14627	1628640_at	0.1159	0.5393	0.0048	0.9858	-0.0147	0.9727	-0.0460	0.9608	0.0064	0.9869	0.0524	0.8569	0.2150	0.8480	0.0773	0.9152	-0.1377	0.8123
CG14470	CG14470	1628641_at	0.0150	0.9418	-0.0140	0.9218	0.0592	0.7690	0.1148	0.8350	0.0997	0.6750	-0.0151	0.9547	-0.0669	0.8940	-0.0162	0.9586	0.0506	0.8287
CG9895 /// DlitCG9895 /// [	CG9895 /// GA22	1628642_at	0.0778	0.6947	0.2158	0.2884	0.2127	0.2248	-0.0463	0.9436	-0.1364	0.4826	-0.0901	0.6352	0.1121	0.8609	0.0397	0.9220	-0.0724	0.8237
CG31267	CG31267	1628643_at	-0.5777	0.8885	-0.0577	0.5985	-0.0322	0.8993	-0.2932	0.9745	-2.9111	0.1960	-2.6179	0.1946	-0.2126	0.9894	-2.2723	0.5140	-2.0597	0.5634
CG17350	CG17350	1628644_at	-4.4365	0.0005	-5.0899	0.0041	-5.1139	0.0000	0.4443	0.6506	1.3236	0.0129	0.8793	0.0440	0.2488	0.8940	0.3931	0.5923	0.1444	0.8779
CG6018	CG6018	1628645_at	-0.4342	0.1278	0.0026	0.9844	0.0671	0.7382	0.0725	0.9300	-0.5518	0.0299	-0.6243	0.0112	-0.0367	0.9816	-0.1586	0.6896	-0.1218	0.7691
CG18507	CG18507	1628646_at	-1.9559	0.0054	-0.8881	0.0503	-0.7853	0.0017	-0.1525	0.7857	-1.5213	0.0004	-1.3688	0.0003	-0.1565	0.9257	-0.5636	0.3002	-0.4070	0.4824
a5	antennal protein 5	1628647_at	0.1848	0.2367	0.0955	0.5524	0.2810	0.2663	0.1709	0.7767	0.0756	0.8080	-0.0953	0.7226	-0.0332	0.9589	0.0129	0.9660	0.0461	0.8460
CG3909	CG3909	1628648_at	0.1446	0.5485	0.2722	0.3101	0.3858	0.0243	-0.1107	0.7693	0.1042	0.5451	0.2149	0.1311	-0.1037	0.9309	0.2859	0.4689	0.3896	0.3271
Mst57Db	male specific tran	1628649_at	-0.0491	0.7905	-0.0151	0.9388	-0.0850	0.6821	0.0119	0.9922	0.0444	0.8973	0.0325	0.9156	0.0603	0.9390	0.0865	0.7706	0.0261	0.9387
CG31538	CG31538	1628650_at	0.0102	0.9634	-0.0182	0.8634	-0.0088	0.9715	-0.0348	0.9630	0.0468	0.8578	0.0816	0.6979	-0.0296	0.9816	-0.0144	0.9758	0.0152	0.9709
Nf1	neurofibromatosis	1628651_a_at	-0.3604	0.1517	-0.1501	0.5757	-0.0295	0.8884	-0.0569	0.9339	-0.0797	0.7378	-0.0228	0.9263	-0.1124	0.9239	0.0871	0.8771	0.1995	0.6388
Myo10A	Myosin XV	1628652_at	-1.1367	0.1885	-1.5558	0.2111	-2.0610	0.0006	0.4059	0.6402	0.9821	0.0283	0.5762	0.1271	0.7341	0.8331	0.5522	0.7466	-0.1820	0.9282
CG31238	CG31238	1628653_at	0.0539	0.7953	-0.1919	0.3052	0.1340	0.3685	0.2186	0.4259	0.0549	0.7627	-0.1637	0.2276	-0.0880	0.8874	-0.1528	0.5259	-0.0649	0.8243
---	---	1628654_at	-0.0215	0.9170	0.0759	0.5401	0.1833	0.1963	0.2432	0.5156	0.1535	0.4559	-0.0898	0.6615	0.0681	0.9309	0.1546	0.5618	0.0865	0.7725
CG4476	CG4476	1628655_at	0.1034	0.8477	-1.8160	0.1203	-1.3762	0.1427	0.0921	0.9434	1.8354	0.0010	1.7434	0.0008	-0.4123	0.9467	0.0465	0.9914	0.4588	0.8474
CG34352	LDLR-like	1628656_at	-0.1309	0.7122	-0.3497	0.2091	-0.4272	0.0725	-0.0372	0.9586	0.5100	0.0149	0.5471	0.0066	-0.0831	0.9478	-0.0362	0.9531	0.0469	0.9310
GstE9	Glutathione S tran	1628657_at	2.0241	0.0060	0.8972	0.4397	2.3469	0.0001	1.0332	0.0532	0.5444	0.0780	-0.4888	0.0765	-0.3677	0.9130	-0.5383	0.6827	-0.1707	0.9169
gbb	glass bottom boat	1628658_at	-0.6161	0.0964	-0.2374	0.5016	0.1214	0.5732	0.3286	0.2442	-0.4246	0.0194	-0.7532	0.0010	-0.0294	0.9898	-0.0479	0.9531	-0.0185	0.9835
Mdr49	Multi drug resistar	1628659_at	-1.8355	0.1207	-3.3952	0.0386	-2.6677	0.0032	0.6209	0.2501	1.0343	0.0068	0.4135	0.1482	-0.0353	0.9964	-0.5454	0.8228	-0.5102	0.8268
CG7130	CG7130	1628660_at	0.0912	0.8509	-0.0490	0.6733	-0.0834	0.7441	0.2210	0.6673	0.6717	0.0140	0.4506	0.0455	0.1614	0.8943	0.3763	0.4041	0.2149	0.6636
CG34393	CG15405	1628661_at	-0.0024	0.9898	0.1022	0.4972	0.3085	0.1794	0.0764	0.9254	-0.0731	0.8094	-0.1496	0.5318	-0.0764	0.8940	0.0542	0.8535	0.1306	0.5634
CG13531	CG13531	1628662_at	0.6746	0.0323	0.0616	0.6912	0.0846	0.6616	-0.0717	0.8645	0.5092	0.0053	0.5810	0.0018	-0.0583	0.9677	-0.0489	0.9358	0.0094	0.9874
CG18259 /// CG6961	CG6961 /// CG18	1628663_s_at	0.0617	0.7857	-0.0020	0.9907	0.1277	0.4746	-0.0754	0.9079	0.2465	0.2381	0.3219	0.0854	-0.1011	0.8828	0.2106	0.4192	0.3117	0.2524
CG12484	CG12484	1628664_at	-2.4086	0.0010	-0.7175	0.0935	-1.3499	0.0004	-0.2977	0.5419	-1.1759	0.0014	-0.8783	0.0029	0.0198	0.9929	-0.0933	0.9085	-0.1131	0.8751
CG6841 /// DmirCG6841	CG6841	1628665_at	0.2382	0.3056	0.1684	0.4236	0.4251	0.0319	0.2657	0.4861	0.2799	0.1670	0.0142	0.9567	-0.0865	0.9291	0.2400	0.4448	0.3265	0.3089
CG3328	CG3328	1628666_at	-2.4227	0.0082	-3.0432	0.0384	-3.5607	0.0000	0.1108	0.8856	0.5591	0.0388	0.4483	0.0590	0.6523	0.8193	-0.1083	0.9541	-0.7606	0.5379
CG6689	CG6689	1628667_at	-1.0099	0.0019	-0.4504	0.1387	-0.7704	0.0071	-0.2535	0.5150	-0.7930	0.0031	-0.5395	0.0104	-0.0748	0.9296	-0.3229	0.2217	-0.2481	0.3775
---	---	1628668_at	0.2316	0.2766	-0.0504	0.7334	0.0450	0.8449	0.0773	0.9314	0.1367	0.6476	0.0594	0.8472	-0.0746	0.8882	-0.1797	0.3594	-0.1051	0.6260
CG9754	CG9754	1628669_at	-0.2219	0.2025	0.0092	0.9635	-0.1553	0.3689	-0.1356	0.7007	-0.0132	0.9560	0.1224	0.4195	-0.0636	0.9309	0.1645	0.4998	0.2281	0.3503
CG18577	CG18577	1628670_at	0.0842	0.7150	-0.0197	0.8807	0.0651	0.7552	-0.0582	0.9153	-0.0126	0.9606	0.0456	0.8134	-0.1383	0.8480	-0.1341	0.6913	0.0042	0.9937
sls	D-Titin	1628671_a_at	-0.8451	0.1514	-1.5920	0.0242	-3.1986	0.0005	-0.3423	0.2654	0.9038	0.0011	1.2461	0.0002	1.3832	0.5821	0.2362	0.8850	-1.1470	0.3205
pio	PioPio	1628672_at	-1.2619	0.0078	-0.1358	0.4633	-1.0926	0.0010	-0.7832	0.1641	-0.6974	0.0465	0.0858	0.8252	-0.0645	0.9235	0.0249	0.9449	0.0894	0.7220
hk	hook	1628673_at	-0.2677	0.1012	-0.3427	0.3107	-0.1499	0.3514	-0.0019	0.9973	0.3356	0.0351	0.3375	0.0217	-0.2244	0.7707	0.2378	0.4794	0.4622	0.1865
---	---	1628674_at	0.2488	0.2520	0.0160	0.9771	0.3597	0.0466	0.1027	0.8841	-0.1256	0.6413	-0.2283	0.2943	-0.0481	0.9521	-0.2223	0.3419	-0.1743	0.4835
Caps	Calcium activated	1628675_at	0.2221	0.4897	0.1807	0.6004	-0.1837	0.5600	-0.0872	0.9345	0.3371	0.2846	0.4243	0.1276	0.2246	0.8909	0.2803	0.6839	0.0557	0.9487
CG30089	CG30089	1628676_at	0.1316	0.3605	0.2413	0.0848	0.0746	0.7050	-0.0453	0.9319	-0.2176	0.1662	-0.1723	0.2237	0.2291	0.7464	-0.0971	0.7984	-0.3262	0.2917
Pkcdelta	protein kinase C	1628677_at	-1.8024	0.0027	-1.6186	0.0506	-2.3820	0.0004	-0.4639	0.3317	-0.2609	0.3536	0.2030	0.4339	0.2803	0.9088	-0.0741	0.9590	-0.3544	0.7168
CG13920	CG13920	1628678_at	-2.6873	0.0208	-2.7105	0.0148	-3.6696	0.0000	-0.9420	0.0657	-1.2436	0.0021	-0.3016	0.2555	0.2548	0.9514	-0.9918	0.4203	-1.2466	0.3246
Snm1	mutagen-sensitive	1628679_at	-0.8466	0.0489	-1.1219	0.0140	-0.7839	0.0053	-0.0473	0.9576	0.6471	0.0141	0.6945	0.0062	-0.3479	0.7070	0.1960	0.6414	0.5439	0.1867
CG13581	CG13581	1628680_at	0.1529	0.5554	0.1344	0.3480	-0.2297	0.1585	-0.1578	0.6840	0.0266	0.9162	0.1845	0.2553	0.1545	0.8611	-0.0305	0.9590	-0.1850	0.6318
CG31128	CG31128	1628681_at	0.2341	0.1173	0.1499	0.1776	0.2299	0.1767	-0.0590	0.9013	0.2306	0.1341	0.2897	0.0423	-0.0235	0.9764	-0.0244	0.9363	-0.0009	0.9979
CG5577	CG5577	1628682_at	0.3938	0.0403	0.8687	0.0111	0.8404	0.0013	-0.0736	0.8631	-0.7765	0.0009	-0.7029	0.0008	-0.1644	0.8122	-0.3246	0.2394	-0.1602	0.6018
CG6272	CG6272	1628683_at	-0.4354	0.1199	-0.4274	0.1642	-0.2956	0.1574	0.1104	0.8815	0.1112	0.7021	0.0008	0.9981	0.0638	0.9511	0.1245	0.7248	0.0607	0.8858
Teh1	tipE homolog 1	1628684_at	-0.2003	0.2988	-0.1371	0.4824	-0.0108	0.9575	0.0498	0.9436	-0.0409	0.8780	-0.0907	0.6615	-0.0913	0.8804	0.0679	0.8230	0.1592	0.5092
---	---	1628685_at	-0.2002	0.3174	0.0987	0.4129	0.1801	0.2611	-0.0544	0.9518	-0.2770	0.2665	-0.2227	0.3254	-0.0129	0.9898	0.0652	0.8153	0.0782	0.7524

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
su(f)	lethal3Des	1628686_a_at	0.1986	0.2543	0.0481	0.9130	0.2859	0.1321	0.0024	0.9961	-0.2255	0.1522	-0.2279	0.1063	-0.0657	0.9552	-0.3008	0.3698	-0.2351	0.5073
CG13602	CG13602	1628687_at	-0.0031	0.9877	0.1449	0.2610	0.0204	0.9017	-0.0843	0.8219	-0.1328	0.3730	-0.0486	0.7613	0.2729	0.6272	0.1996	0.3884	-0.0732	0.7964
CG12054 /// DyakCG12054	CG12054	1628688_at	0.0892	0.9297	-0.1241	0.4549	-0.6712	0.0207	-0.2235	0.7929	0.6994	0.0486	0.9229	0.0096	0.4608	0.7337	0.4598	0.4270	-0.0010	0.9992
CG17211	CG17211	1628689_at	-0.0342	0.9080	0.2960	0.2097	0.5648	0.0105	-0.0156	0.9838	-0.3024	0.0826	-0.2869	0.0674	-0.0873	0.9351	0.1811	0.6231	0.2684	0.4463
CG1503	CG1503	1628690_at	0.0907	0.6311	0.1701	0.4829	0.0344	0.8275	-0.0327	0.9649	-0.1121	0.6047	-0.0794	0.7031	0.0020	0.9994	-0.1527	0.6447	-0.1546	0.6386
CG17514	CG17514	1628691_at	0.3030	0.1033	0.2084	0.6424	-0.0058	0.9860	-0.1517	0.6903	0.2077	0.2394	0.3594	0.0302	0.0955	0.9589	0.1257	0.8678	0.0302	0.9706
---	---	1628692_at	0.0369	0.8705	-0.0642	0.7700	0.0945	0.6445	-0.0050	0.9956	0.0421	0.8905	0.0470	0.8577	0.0097	0.9939	-0.0887	0.8176	-0.0984	0.7777
CG15429	CG15429	1628693_at	0.0340	0.8389	-0.0256	0.9401	-0.0071	0.9734	-0.0532	0.9068	-0.1071	0.4854	-0.0539	0.7311	-0.0568	0.9525	0.0703	0.8591	0.1271	0.6877
14-3-3epsilon	Suppressor of Ra	1628694_a_at	0.0817	0.6658	0.0979	0.5173	-0.0755	0.6395	-0.0209	0.9758	0.0630	0.7541	0.0839	0.6250	0.1181	0.8705	0.1906	0.5246	0.0725	0.8444
CG33224	CG33224	1628695_at	0.1994	0.2378	0.2765	0.2183	0.2322	0.1862	-0.0555	0.9436	-0.2197	0.3304	-0.1641	0.4324	-0.0805	0.9299	-0.1268	0.7052	-0.0463	0.9096
CG12643 /// DyakCG12643	CG12643	1628696_at	0.6555	0.3375	-0.3365	0.4737	-0.4005	0.5241	-0.5094	0.5301	-0.5124	0.2253	-0.0029	0.9957	-0.3320	0.9324	-1.3950	0.2505	-1.0631	0.4061
CG14952	CG14952	1628697_at	0.2833	0.3108	0.3378	0.0944	0.3971	0.0456	-0.0631	0.9154	-0.1887	0.3156	-0.1256	0.4789	0.0078	0.9962	-0.0019	0.9989	-0.0097	0.9863
---	---	1628698_at	0.2216	0.3111	-0.0948	0.3756	-0.1948	0.2720	-0.0696	0.8841	0.1248	0.4622	0.1944	0.1804	-0.1353	0.8059	-0.1271	0.6041	0.0082	0.9835
Lsp1beta	Larval serum prot	1628699_at	2.0293	0.1784	1.1640	0.5965	1.8325	0.0344	1.1490	0.3581	1.0339	0.1457	-0.1151	0.8928	0.5286	0.9611	0.3890	0.9363	-0.1395	0.9769
CG15472	CG15472	1628700_at	-0.1134	0.4226	-0.0586	0.6213	0.0039	0.9849	-0.0042	0.9956	-0.0247	0.9149	-0.0205	0.9195	0.1546	0.7953	0.2078	0.4028	0.0533	0.8750
CG14798	CG14798	1628701_a_at	-0.2272	0.5339	-0.2265	0.1081	-0.0897	0.6217	0.1703	0.6988	0.1394	0.5151	-0.0309	0.8991	-0.1251	0.9309	-0.1427	0.8153	-0.0177	0.9831
CG32580	CG32580	1628702_at	0.0735	0.6694	0.0365	0.8588	0.0733	0.7174	-0.1352	0.7850	-0.1825	0.3827	-0.0472	0.8412	-0.0672	0.9092	0.0047	0.9925	0.0718	0.7696
---	---	1628703_at	-0.0294	0.8892	0.0000	1.0000	0.2158	0.2235	0.0856	0.8589	0.0999	0.5977	0.0143	0.9482	-0.0039	0.9964	0.0615	0.8343	0.0654	0.8088
Or69a	Odorant receptor	1628704_at	0.0808	0.6818	-0.0335	0.7342	0.1227	0.6216	0.0728	0.8841	0.1726	0.3153	0.0998	0.5505	-0.0372	0.9761	0.0318	0.9478	0.0690	0.8604
PH4alphaEFB	prolyl-4-hydroxyla	1628705_at	1.6304	0.0008	1.0389	0.0137	1.6201	0.0000	0.2087	0.6195	0.6084	0.0107	0.3996	0.0392	-0.2507	0.7152	0.1007	0.7741	0.3514	0.2363
CG17982	CG17982	1628706_at	-0.1314	0.5265	0.6613	0.0310	0.4659	0.1287	0.1248	0.9050	-0.5666	0.0952	-0.6914	0.0302	0.1609	0.8903	0.3782	0.3802	0.2173	0.6457
CG12452	CG12452	1628707_at	0.3483	0.1318	0.3023	0.0671	0.0240	0.9408	-0.1762	0.6891	-0.0073	0.9809	0.1689	0.3707	0.1900	0.7853	-0.1143	0.7456	-0.3042	0.3205
CG30044	CG30044	1628708_at	-0.6576	0.0064	0.2169	0.4478	-0.0465	0.8647	-0.2704	0.7067	-0.8034	0.0228	-0.5330	0.0720	-0.0058	0.9964	0.1163	0.7339	0.1221	0.7128
---	---	1628709_at	-0.1460	0.9405	1.0089	0.2154	-0.1542	0.7242	-1.2476	0.6988	-0.6497	0.7034	0.5978	0.7017	-0.0851	0.9776	0.3315	0.6820	0.4166	0.5908
Grasp65	Grasp65	1628710_at	0.3481	0.5459	1.3799	0.0400	1.7233	0.0000	0.4122	0.4532	-0.1939	0.5460	-0.6061	0.0307	0.0380	0.9914	0.9004	0.2404	0.8624	0.2923
CG12307	CG12307	1628711_at	-0.0739	0.7350	0.0662	0.5148	0.1328	0.5178	0.0774	0.8781	-0.1988	0.2508	-0.2762	0.0754	0.1820	0.8298	0.0784	0.8773	-0.1036	0.8147
CG17754	CG17754	1628712_at	0.2761	0.5648	0.2043	0.6802	-0.4217	0.2117	-0.2546	0.7556	-0.0355	0.9457	0.2191	0.5235	0.1874	0.9373	-0.1785	0.8644	-0.3659	0.6495
Cpr65Aw	CG32404	1628713_at	-0.2742	0.1700	0.0055	0.9660	0.0068	0.9816	0.0098	0.9937	-0.1041	0.7228	-0.1139	0.6627	0.1885	0.7062	0.1922	0.3442	0.0037	0.9923
CG12129	CG12129	1628714_at	0.6570	0.0578	0.3649	0.4159	0.6326	0.0028	0.0280	0.9672	0.3132	0.0814	0.2852	0.0763	-0.0784	0.9657	0.1225	0.8550	0.2009	0.7154
Karl	Karl	1628715_a_at	-2.8633	0.0132	-3.3454	0.0212	-4.0717	0.0000	-0.5262	0.1956	-1.0327	0.0021	-0.5065	0.0291	0.2849	0.9515	-1.3004	0.3421	-1.5853	0.2757
CG13283	CG13283	1628716_at	-0.2208	0.6207	-1.5844	0.0744	-1.5401	0.0117	-0.4679	0.4335	-0.2337	0.4985	0.2342	0.4489	-0.6857	0.7707	-1.7005	0.0979	-1.0148	0.3226
mud	mushroom body d	1628717_a_at	0.1770	0.3855	-0.0562	0.5437	0.1132	0.5572	0.0194	0.9852	0.2217	0.3491	0.2024	0.3429	-0.1344	0.8160	-0.0431	0.9064	0.0913	0.7410
CG11289	CG11289	1628718_at	-5.1898	0.0004	-5.7240	0.0035	-5.7203	0.0000	-0.1669	0.8578	-0.2686	0.4378	-0.1017	0.7854	-0.1668	0.9457	-0.7801	0.2753	-0.6133	0.4148
cry	crybaby	1628719_at	-0.3015	0.0932	0.6683	0.0256	1.2075	0.0002	-0.2411	0.5290	-1.4580	0.0003	-1.2169	0.0003	-0.5911	0.2884	-0.2847	0.3059	0.3064	0.2964
CG4440	CG4440	1628720_at	-0.1174	0.7145	-0.0894	0.6036	-0.0116	0.9598	-0.0144	0.9894	0.0368	0.9167	0.0512	0.8608	-0.0608	0.9550	0.0256	0.9613	0.0864	0.8270
CG9864 /// DsecCG9864 ///	CG9864	1628721_at	0.4854	0.3806	-1.6394	0.0109	-0.6365	0.0200	-0.2221	0.8350	0.7399	0.0699	0.9620	0.0158	-1.3249	0.2652	-1.3404	0.0483	-0.0155	0.9874
Task7	Task7	1628722_at	0.2165	0.2331	0.0747	0.6606	0.1078	0.5076	0.2182	0.6677	0.1510	0.5633	-0.0672	0.8047	-0.0948	0.8846	-0.0886	0.7684	0.0061	0.9874
CG4038	CG4038	1628723_at	0.5368	0.3777	0.2397	0.6585	-0.3106	0.2778	-0.2530	0.7293	0.7886	0.0242	1.0417	0.0043	0.3681	0.8680	0.6922	0.4287	0.3241	0.7492
---	---	1628724_at	0.1667	0.2670	0.1231	0.3914	0.1309	0.4115	0.0777	0.8705	0.1479	0.3858	0.0703	0.6877	0.0408	0.9535	-0.0018	0.9979	-0.0425	0.8800
CG14127	CG14127	1628725_at	-0.0130	0.9597	-0.0240	0.8463	0.0978	0.5744	0.0333	0.9599	0.1717	0.3297	0.1384	0.3915	-0.0918	0.8940	0.1869	0.4710	0.2786	0.2891
Rgl	Ral guanine nucle	1628726_s_at	-1.3079	0.0094	-1.2969	0.0734	-1.6118	0.0005	-0.0841	0.9518	0.3930	0.3155	0.4771	0.1674	0.1927	0.9257	0.4830	0.4947	0.2903	0.7072
CG15599	CG15599	1628727_at	0.2960	0.0983	0.1705	0.4402	0.2488	0.1185	0.0471	0.9563	0.1227	0.6373	0.0755	0.7679	-0.0762	0.9199	0.0070	0.9905	0.0832	0.7849
CG6163	CG6163	1628728_at	-0.2036	0.2668	-0.0025	0.9846	-0.0645	0.7657	-0.0562	0.9228	-0.1569	0.3811	-0.1007	0.5606	0.0677	0.8940	0.0360	0.9004	-0.0317	0.9033
---	---	1628729_at	-0.7930	0.0649	-0.2398	0.3437	-0.7503	0.0749	-0.3819	0.6371	-0.6298	0.1120	-0.2479	0.5204	-0.0841	0.9555	-0.1908	0.7010	-0.1067	0.8509
rdgA	DAG kinase	1628730_at	-0.5791	0.0604	-0.9080	0.0308	-0.7869	0.0051	0.1809	0.8408	0.6059	0.0740	0.4249	0.1561	0.0205	0.9853	0.0987	0.7434	0.0782	0.8017
CG13741 /// CG8080 /// Dy	CG13741 /// CG8	1628731_s_at	1.7486	0.0124	-0.0226	0.9788	-0.0402	0.8952	-0.5627	0.4221	1.5741	0.0025	2.1368	0.0004	-0.5341	0.7779	-0.2066	0.8550	0.3275	0.7269
pon	Partner of Numb	1628732_at	-0.2465	0.4649	0.1411	0.7248	-0.1534	0.4419	0.0371	0.9744	-0.0565	0.8806	-0.0936	0.7612	0.1178	0.9441	-0.0076	0.9942	-0.1254	0.8480
Oseg6	Oseg6	1628733_at	0.1268	0.5535	-0.0971	0.6888	-0.0800	0.7169	0.1228	0.8546	0.4446	0.0694	0.3218	0.1357	-0.0025	0.9994	0.2720	0.5321	0.2745	0.5340
CG11286	CG11286	1628734_at	0.1976	0.1931	-0.0843	0.5954	0.0075	0.9762	0.0585	0.9139	0.2521	0.1373	0.1937	0.2032	0.1232	0.8825	0.0237	0.9643	-0.0995	0.7954
---	---	1628735_at	0.2121	0.2290	0.0649	0.6648	0.1467	0.5325	-0.0240	0.9704	0.0152	0.9513	0.0391	0.8358	0.0364	0.9775	0.0181	0.9721	-0.0183	0.9691
CG1137	CG1137	1628736_at	0.2095	0.2623	0.1571	0.3424	0.2381	0.1356	-0.0072	0.9946	0.0246	0.9398	0.0318	0.9075	-0.0827	0.8743	-0.0312	0.9225	0.0515	0.8444
CG12912	CG12912	1628737_at	-0.1440	0.5328	-0.0109	0.9523	-0.1124	0.5569	0.1130	0.8315	-0.1219	0.5787	-0.2349	0.1937	0.0833	0.9032	-0.0212	0.9590	-0.1045	0.7106
CG31251	CG31251	1628738_at	-0.4066	0.0245	0.2487	0.1142	0.3963	0.0224	-0.0441	0.9380	-0.3132	0.0627	-0.2691	0.0718	-0.1132	0.8439	0.3927	0.1038	0.5059	0.0702
CG33528	CG33528	1628739_at	-0.2173	0.8389	0.1507	0.8510	-1.3090	0.0080	-0.8939	0.2378	-2.0134	0.0016	-1.1195	0.0130	0.4249	0.9076	-1.4678	0.2430	-1.8927	0.1700
---	---	1628740_at	-0.1466	0.6015	0.1849	0.2875	0.0383	0.8486	-0.2063	0.5680	-0.1958	0.2944	0.0104	0.9645	-0.0407	0.9775	0.1368	0.7300	0.1775	0.6309
CG7364	CG7364	1628741_s_at	0.1387	0.3600	0.0866	0.5608	-0.2109	0.3204	-0.0747	0.8932	0.1109	0.5822	0.1856	0.2673	0.0569	0.9514	0.0878	0.8045	0.0309	0.9362
mld	molting defective	1628742_at	-0.5214	0.0806	-1.9708	0.0263	-1.7972	0.0199	0.3957	0.8402	1.6515	0.0315	1.2558	0.0590	-0.0796	0.9717	0.0010	0.9998	0.0806	0.9183
glec	Gliolectin	1628743_at	-0.8105	0.0129	-0.9919	0.0269	-1.5591	0.0001	0.0111	0.9931	0.3155	0.2107	0.3045	0.1774	0.5597	0.4562	0.1770	0.6819	-0.3827	0.3342
CG17597 /// ScpX	CG17597 /// stero	1628744_s_at	0.4305	0.0669	0.9158	0.0348	1.3842	0.0001	0.1019	0.8738	-0.8449	0.0034	-0.9468	0.0012	-0.3169	0.6898	-0.2507	0.4365	0.0662	0.8800
CG18313	CG18313	1628745_at	-0.0547	0.7342	0.1435	0.4342	0.1666	0.3560	-0.0413	0.9586	-0.0784	0.7535	-0.0371	0.8809	0.0819	0.8870	0.1065	0.6500	0.0246	0.9350
CG8851	CG8851	1628746_at	-0.0112	0.9596	0.0423	0.8702	0.0034	0.9886	0.0401	0.9602	-0.0676	0.7940	-0.1077	0.6146	0.0020	0.9982	-0.1234	0.5084	-0.1254	0.5041
---	---	1628747_at	0.2946	0.3494	0.1948	0.1964	0.1298	0.3698	0.0035	0.9956	0.1143	0.4456	0.1108	0.4108	-0.0034	0.9971	-0.0584	0.8539	-0.0550	0.8542
---	---	1628748_at	0.0940	0.6434	-0.1850	0.1452	-0.3607	0.0324	0.1511	0.7749	0.4265	0.0581	0.2753	0.1627	0.0213	0.9862	-0.1372	0.6633	-0.1585	0.6071
CG17599	CG17599	1628749_at	0.2895	0.1149	0.1413	0.5185	0.0380	0.8832	0.0099	0.9908	0.1793	0.3435	0.1694	0.3179	0.0942	0.8909	0.0783	0.8165	-0.0159	0.9675
CG33999	CG33999	1628750_at	-0.0933	0.5934	0.1722	0.4185	0.2746	0.2265	0.0666	0.9308	-0.1843	0.4276	-0.2509	0.2114	-0.0835	0.9137	-0.0417	0.9208	0.0418	0.9107
CG8728	CG8728	1628751_at	-0.1785	0.4208	0.1663	0.4935	0.5718	0.0423	0.0748	0.8990	-0.2309	0.2305	-0.3056	0.0777	-0.2173	0.8571	0.3303	0.5172	0.5476	0.2819
CG17085	CG17085	1628752_at	0.0565	0.7442	-0.1659	0.4537	0.2877	0.1658	0.1254	0.8915	0.0941	0.8043	-0.0314	0.9347	0.0131	0.9914	0.0757	0.8232	0.0626	0.8515
---	---	1628753_at	-0.3538	0.4201	-0.4034	0.3801	-0.5592	0.0495	-0.4792	0.5492	0.1080	0.8395	0.5872	0.1112	-0.1694	0.9101	-0.1933	0.7656	-0.0239	0.9777
Ggamma30A	G protein gamma	1628754_s_at	0.0693	0.7063	-0.0676	0.6925	0.1719	0.4910	0.0890	0.9011	0.0828	0.7664	-0.0061	0.9836	-0.1733	0.7628	-0.1123	0.6711	0.0610	0.8417

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Gfr	Golgi GDP-fucose	1628755_at	0.5407	0.1313	0.1615	0.6453	0.4657	0.0266	-0.0003	0.9997	0.5022	0.0130	0.5025	0.0079	-0.2620	0.8494	0.1604	0.8343	0.4225	0.4759
---	---	1628756_at	0.1408	0.4671	-0.0394	0.8879	-0.0148	0.9635	0.0679	0.9377	0.2079	0.4298	0.1400	0.5809	0.1164	0.8692	-0.0561	0.8910	-0.1725	0.5612
Glut1	glucose transporte	1628757_at	-0.3252	0.2333	-1.1277	0.1107	-1.1517	0.0011	-0.1253	0.8735	0.2764	0.3176	0.4017	0.1000	-0.1226	0.9543	-0.4335	0.5020	-0.3109	0.6450
CG1774	CG1774	1628758_at	0.2725	0.3363	0.0978	0.8556	0.8255	0.0019	0.1504	0.7604	-0.6311	0.0096	-0.7815	0.0022	-0.6432	0.5352	-0.9175	0.0758	-0.2743	0.6033
CG14073	Nirvana	1628759_a_at	-0.2050	0.7502	-0.3505	0.3248	-1.0528	0.0030	-0.1192	0.9022	0.7180	0.0299	0.8372	0.0097	0.3766	0.8270	0.4203	0.5835	0.0436	0.9686
CG8517	thioredoxin	1628760_at	0.2977	0.3710	0.0888	0.4384	0.3167	0.2678	0.1504	0.8385	0.0962	0.7724	-0.0541	0.8671	-0.0015	0.9994	0.0583	0.8959	0.0598	0.8834
CG12207	CG12207	1628761_s_at	-0.7170	0.0079	-1.5147	0.0179	-1.3655	0.0146	0.2514	0.8544	0.6298	0.2011	0.3784	0.4134	0.1777	0.8806	-0.0869	0.8987	-0.2646	0.5829
---	---	1628762_at	0.0853	0.6042	0.1233	0.3848	0.1677	0.4284	0.1052	0.8076	-0.0228	0.9244	-0.1280	0.4280	0.0790	0.9277	0.1058	0.7549	0.0268	0.9466
Ptpmeg	split central compl	1628763_at	1.1631	0.0486	0.5251	0.3599	0.4004	0.1479	0.0620	0.9314	0.3087	0.1492	0.2466	0.1987	0.1197	0.9737	-0.3546	0.7340	-0.4744	0.6225
l(3)s1921	lethal (3) s1921	1628764_at	0.2771	0.3355	0.1339	0.6363	0.0826	0.7408	0.1170	0.8852	0.4968	0.0737	0.3797	0.1212	0.1514	0.9066	0.4346	0.3346	0.2832	0.5634
---	---	1628765_at	-0.0398	0.8250	-0.0703	0.7192	0.0485	0.8155	0.1556	0.7167	0.1434	0.4830	-0.0123	0.9597	-0.0084	0.9940	0.1482	0.5917	0.1565	0.5709
CG4984	CG4984	1628766_at	-2.1796	0.0113	-1.1741	0.1007	-2.1945	0.0003	-0.6419	0.4191	-1.9855	0.0016	-1.3437	0.0052	0.1175	0.9742	-1.0157	0.2507	-1.1333	0.2326
MED26	Mediator complex	1628767_s_at	0.3397	0.1494	0.3069	0.3958	-0.5224	0.1288	-0.2628	0.3855	0.4655	0.0145	0.7283	0.0012	0.5736	0.7230	0.2970	0.7064	-0.2766	0.7287
CG4676	CG4676	1628768_at	-0.9926	0.0366	-0.8023	0.0097	-1.2209	0.0194	-0.0481	0.9774	0.1429	0.7777	0.1910	0.6595	-0.2432	0.8243	-0.1280	0.8341	0.1152	0.8435
---	---	1628769_at	-0.0029	0.9911	0.0204	0.9308	0.2795	0.1492	0.0380	0.9441	-0.0562	0.7613	-0.0942	0.5320	-0.0099	0.9939	0.1379	0.6743	0.1478	0.6419
---	---	1628770_at	-0.0157	0.9416	0.0153	0.9399	0.2161	0.3077	-0.1335	0.7725	-0.1699	0.3937	-0.0364	0.8738	-0.2273	0.6376	-0.0682	0.7787	0.1591	0.4322
CG32700 /// DsimCG32700	CG32700	1628771_s_at	5.1724	0.0018	3.2810	0.0024	6.2468	0.0000	3.0092	0.0403	2.4990	0.0083	-0.5102	0.5164	-0.0284	0.9848	0.4864	0.1410	0.5148	0.1512
---	---	1628772_at	-0.0055	0.9812	-0.1080	0.4315	0.0418	0.8022	0.1420	0.7854	0.2442	0.2578	0.1022	0.6439	0.0013	0.9994	0.0927	0.7430	0.0913	0.7422
---	---	1628773_at	-0.0301	0.8674	-0.0229	0.8817	0.1362	0.4814	0.1263	0.7279	0.0595	0.7604	-0.0669	0.6959	-0.0795	0.9142	0.0602	0.8685	0.1397	0.6156
boi	CG32796	1628774_a_at	-0.5967	0.0463	-0.6757	0.1318	-1.0824	0.0483	-0.3594	0.4861	0.3545	0.1963	0.7138	0.0110	-0.6642	0.7810	-0.0849	0.9605	0.5793	0.6029
RpL8	Ribosomal protein	1628775_s_at	-0.0161	0.9226	-0.0484	0.6598	0.0785	0.6268	0.1070	0.7507	0.0758	0.6448	-0.0312	0.8545	0.1013	0.8379	0.0708	0.7779	-0.0305	0.9139
---	---	1628776_at	0.2161	0.3013	0.0434	0.9046	0.1167	0.5439	0.0790	0.9023	0.1427	0.5200	0.0637	0.7846	-0.0175	0.9832	-0.0729	0.7541	-0.0554	0.8207
CG12140	CG12140	1628777_at	0.8103	0.0159	0.4327	0.2579	0.7869	0.0026	-0.1046	0.7815	-0.2127	0.1714	-0.1081	0.4660	-0.5393	0.6898	-0.5569	0.3022	-0.0176	0.9848
itp	ion transport pepti	1628778_at	-2.2576	0.0083	-4.0382	0.0055	-4.8456	0.0001	-0.1252	0.9413	1.7708	0.0039	1.8960	0.0018	0.6925	0.7485	0.0225	0.9917	-0.6700	0.4799
svp	seven-up	1628779_a_at	1.1024	0.1665	1.1457	0.2255	1.4111	0.0005	0.0955	0.9528	-0.4499	0.3171	-0.5454	0.1693	-0.1437	0.9764	-0.3492	0.8177	-0.2055	0.8982
CG11582	CG11582	1628780_at	0.2221	0.3255	0.0535	0.7364	0.0658	0.6836	-0.1203	0.8495	-0.1056	0.6902	0.0147	0.9597	-0.0502	0.9665	-0.1705	0.6157	-0.1203	0.7431
CG30482	CG30482	1628781_at	0.2242	0.1928	0.2970	0.1162	-0.0479	0.7766	-0.1396	0.7115	-0.0091	0.9732	0.1305	0.4151	0.1177	0.8122	-0.0485	0.8660	-0.1662	0.4275
CG4341	CG4341	1628782_at	-0.0847	0.5912	0.0145	0.9096	-0.0531	0.7550	-0.1343	0.6954	-0.2319	0.1410	-0.0977	0.5255	0.0642	0.9238	0.0220	0.9507	-0.0422	0.8912
Rfx	Regulatory factor	1628783_at	-0.3385	0.3034	0.2924	0.2482	-0.0424	0.8256	-0.2510	0.6513	-0.5291	0.0535	-0.2781	0.2484	0.0458	0.9611	0.1671	0.5415	0.1212	0.6728
CG16817	CG16817	1628784_at	0.4795	0.0203	0.4194	0.1100	-0.1821	0.4095	-0.0435	0.9436	0.5267	0.0093	0.5701	0.0039	0.6223	0.3734	0.4583	0.2172	-0.1640	0.7034
l(3)04053	lethal (3) 04053	1628785_at	-0.3380	0.3121	-0.3009	0.4992	-0.4800	0.0197	-0.1413	0.7293	0.0496	0.8308	0.1909	0.2440	0.0679	0.9805	0.0282	0.9797	-0.0397	0.9650
---	---	1628786_at	-0.0197	0.9099	-0.0967	0.4658	-0.1679	0.4535	-0.1235	0.8200	0.1394	0.5418	0.2628	0.1658	-0.0747	0.9101	-0.0417	0.9084	0.0330	0.9194
CG31238	CG31238	1628787_s_at	0.1424	0.5600	0.0788	0.6702	0.1496	0.3340	-0.0223	0.9753	0.0078	0.9775	0.0301	0.8850	-0.0925	0.9056	-0.0827	0.8212	0.0098	0.9837
dsh	dishevelled	1628788_a_at	-0.5112	0.2062	-0.5573	0.0144	-0.5559	0.0134	-0.1078	0.8676	0.3005	0.1862	0.4082	0.0506	-0.1713	0.8768	0.1712	0.7330	0.3425	0.4389
---	---	1628789_x_at	0.1430	0.4755	0.1117	0.3736	0.1645	0.2724	0.1255	0.7131	0.0419	0.8326	-0.0836	0.5907	0.1008	0.8999	0.0539	0.9052	-0.0469	0.9084
CG6345	CG6345	1628790_at	0.2485	0.1607	-0.0133	0.9536	0.1148	0.5690	0.1623	0.7689	0.2672	0.2555	0.1049	0.6661	0.0116	0.9914	0.0098	0.9816	-0.0018	0.9965
CG13933	CG13933	1628791_at	-0.1947	0.7022	0.1582	0.2084	-0.3220	0.3596	-0.4536	0.4442	-0.7545	0.0273	-0.3008	0.3051	-0.0670	0.9848	-0.2061	0.8501	-0.1391	0.8998
Obp56g	Odorant-binding p	1628792_at	0.1109	0.6644	0.2505	0.2477	0.1864	0.2123	-0.2299	0.4850	0.0365	0.8759	0.2664	0.0901	0.0462	0.9589	0.1411	0.6043	0.0949	0.7482
CG6106	CG6106	1628793_at	-2.0807	0.0005	-1.0868	0.0308	-2.0739	0.0002	-0.3997	0.3521	-0.5741	0.0275	-0.1744	0.4535	0.0598	0.9514	-0.3913	0.1657	-0.4511	0.1479
---	---	1628794_at	0.2773	0.3611	-0.1468	0.2995	0.1914	0.3176	0.2073	0.6823	0.2487	0.2958	0.0414	0.8819	-0.0431	0.9816	0.0045	0.9962	0.0476	0.9344
CG9175	CG9175	1628795_s_at	0.6911	0.0287	0.8926	0.1093	0.7256	0.0163	-0.0165	0.9863	0.4033	0.0826	0.4198	0.0477	0.3525	0.8122	0.6496	0.2729	0.2971	0.6545
CG6353	CG6353	1628796_at	0.2901	0.3748	0.1152	0.6426	0.6663	0.0505	0.1194	0.8594	-0.1337	0.6183	-0.2531	0.2449	-0.4269	0.7492	-0.3347	0.5800	0.0923	0.9067
CG40260	CG40260	1628797_at	-0.0350	0.8342	-0.0045	0.9853	-0.0075	0.9729	-0.0705	0.8906	0.0657	0.7469	0.1362	0.3932	-0.1127	0.8215	0.0613	0.8212	0.1740	0.4093
lin-52	lin-52	1628798_at	-0.1080	0.5929	-0.3512	0.3189	-0.5214	0.0400	0.1945	0.6338	0.5507	0.0140	0.3561	0.0518	0.1097	0.9343	0.1262	0.8220	0.0165	0.9821
zormin	D-Titin	1628799_at	-0.0372	0.8558	0.0578	0.5727	0.0110	0.9645	-0.2583	0.5735	-0.2966	0.2048	-0.0383	0.8916	0.0431	0.9742	0.0550	0.9148	0.0119	0.9835
Jon65Aiv	Jonah 65A	1628800_at	1.4732	0.4716	-1.0486	0.0395	0.0855	0.8669	0.8366	0.1899	0.8873	0.0293	0.0507	0.9142	-0.5659	0.9499	-1.9054	0.4875	-1.3394	0.6426
kto	khotalo	1628801_at	-0.1276	0.8695	-0.1624	0.8485	-0.1158	0.6460	0.0645	0.9558	0.3112	0.3308	0.2467	0.4007	0.0955	0.9831	0.2838	0.8438	0.1884	0.8977
CG32554	CG32554	1628802_at	0.0148	0.9376	-0.0602	0.8680	-0.1342	0.5477	0.1786	0.7672	0.5184	0.0487	0.3399	0.1370	0.0509	0.9742	0.4388	0.2507	0.3880	0.3406
CG31462	CG31462	1628803_at	0.0508	0.8165	0.1219	0.4356	-0.0482	0.7822	-0.0144	0.9854	0.0534	0.8052	0.0678	0.7174	0.1060	0.8586	0.1588	0.5248	0.0528	0.8694
CG11419	CG11419	1628804_at	0.0183	0.9528	-0.0701	0.6682	-0.0250	0.9190	0.0285	0.9777	0.1085	0.7077	0.0800	0.7727	0.0082	0.9950	-0.0328	0.9462	-0.0410	0.9211
---	---	1628805_s_at	0.0198	0.9485	0.0213	0.9059	-0.0040	0.9916	-0.0250	0.9819	0.1041	0.7372	0.1291	0.6313	-0.0614	0.9514	-0.1057	0.7709	-0.0443	0.9151
Rh5	Rhodopsin5	1628806_at	0.1610	0.3758	0.3166	0.0842	0.3551	0.1016	0.0238	0.9760	-0.2014	0.3054	-0.2252	0.1948	0.0503	0.9672	-0.0996	0.8123	-0.1499	0.6708
CG5205	CG5205	1628807_at	0.8878	0.0067	0.3831	0.1197	0.3577	0.2146	0.0364	0.9758	1.0728	0.0039	1.0364	0.0027	0.0358	0.9831	0.4700	0.2151	0.4342	0.2830
---	---	1628808_at	0.0843	0.6702	0.0877	0.5142	0.0038	0.9889	0.0796	0.8594	0.0214	0.9245	-0.0582	0.7377	0.0192	0.9868	-0.0781	0.8212	-0.0973	0.7492
D19B	D19B	1628809_at	-0.5402	0.3071	0.0483	0.9517	0.3010	0.3357	-0.0089	0.9951	-0.3851	0.1854	-0.3761	0.1492	-0.1948	0.9457	0.2746	0.8039	0.4695	0.6140
CG15597	CG15597	1628810_at	-0.1161	0.6097	-0.4456	0.0632	-0.2467	0.3340	0.4900	0.3985	0.7827	0.0242	0.2927	0.3244	0.0707	0.8956	0.0635	0.8061	-0.0073	0.9835
CG14546	CG14546	1628811_at	0.0139	0.9584	0.0310	0.8691	-0.0174	0.9348	0.0085	0.9941	-0.1300	0.6226	-0.1385	0.5540	0.0144	0.9916	-0.0016	0.9991	-0.0160	0.9745
---	---	1628812_at	0.2137	0.3944	-0.0044	0.9923	0.0002	0.9995	-0.1072	0.9048	-0.0559	0.8869	0.0513	0.8815	0.0665	0.9460	-0.0720	0.8624	-0.1386	0.6690
CG31381	CG31381	1628813_at	0.5056	0.2315	1.0776	0.0454	1.0075	0.0031	-0.0146	0.9907	-0.3650	0.1776	-0.3505	0.1487	0.1116	0.9689	0.5425	0.4789	0.4310	0.5908
CG9894 /// DyakCG9894	CG9894	1628814_s_at	0.7235	0.0233	0.3744	0.0440	0.2371	0.3253	0.0230	0.9777	0.1878	0.3706	0.1648	0.3878	0.1755	0.8814	-0.3123	0.4985	-0.4877	0.2923
sl	phospholipase Cg	1628815_at	0.2730	0.1569	0.0163	0.9601	-0.1676	0.4274	-0.0681	0.9314	0.4844	0.0444	0.5524	0.0164	0.1670	0.8465	0.2483	0.5000	0.0814	0.8635
---	---	1628816_at	0.1436	0.4158	0.0496	0.6232	-0.0947	0.6770	-0.0104	0.9876	0.1491	0.3566	0.1594	0.2648	0.1223	0.8425	0.1257	0.6465	0.0033	0.9941
---	---	1628817_at	-0.0631	0.7191	0.0688	0.5185	0.0941	0.5396	0.0100	0.9883	-0.0072	0.9772	-0.0172	0.9307	0.0078	0.9928	0.0553	0.8356	0.0475	0.8545
CG17562	CG17562	1628818_at	0.5880	0.7330	-2.6985	0.3769	1.3230	0.2135	3.2496	0.1162	2.1705	0.0828	-1.0791	0.3452	-0.8733	0.9340	-1.1153	0.7922	-0.2420	0.9612
---	---	1628819_at	-0.0632	0.8472	0.0015	0.9913	-0.0611	0.7473	0.0776	0.9435	0.0574	0.8906	-0.0202	0.9588	0.1296	0.8270	0.1461	0.5715	0.0165	0.9636
CG32521	CG32521	1628820_at	-0.1697	0.3671	0.0500	0.8397	-0.0093	0.9756	0.0620	0.9515	-0.2599	0.3662	-0.3219	0.2003	-0.0118	0.9914	-0.1781	0.4345	-0.1664	0.4822
CG17027	CG17027	1628821_at	-2.0003	0.0013	-1.6046	0.0063	-1.5565	0.0027	0.0019	0.9988	-0.7391	0.0134	-0.7410	0.0081	-0.1471	0.9467	-0.4182	0.5503	-0.2711	0.7214
CG30271	CG30271	1628822_at	4.2818	0.0023	3.3406	0.0381	4.8485	0.0000	0.5391	0.6327	0.1414	0.8435	-0.3977	0.4455	-1.1724	0.7233	-0.9025	0.5479	0.2698	0.8918
CG14120	CG14120	1628823_at	0.1540	0.9535	-0.1238	0.5168	-0.3235	0.1687	-0.0466	0.9917	-0.7252	0.4586	-0.6786	0.4427	0.0171	0.9990	-1.3669	0.5618	-1.3840	0.5612

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
l(1)G0060	lethal (1) G0060	1628824_s_at	0.3340	0.4651	-0.0082	0.9785	-0.2324	0.2689	-0.0418	0.9627	0.6013	0.0201	0.6431	0.0092	0.3653	0.8042	0.3872	0.5447	0.0219	0.9835
CG8092	CG8092	1628825_at	0.6747	0.0230	0.1569	0.5796	0.2962	0.1181	-0.2340	0.5568	0.3987	0.0566	0.6327	0.0050	-0.2712	0.7726	0.0727	0.9068	0.3439	0.4047
---	---	1628826_at	4.6243	0.0007	3.0208	0.0033	4.5118	0.0000	1.7178	0.0221	1.6495	0.0024	-0.0683	0.8864	0.1461	0.9095	0.0287	0.9698	-0.1174	0.8410
CG40500	CG40500	1628827_s_at	0.0715	0.7366	-0.1293	0.4009	0.1238	0.4986	0.2052	0.6013	0.1266	0.5530	-0.0786	0.7076	-0.1021	0.8439	-0.0788	0.7563	0.0233	0.9373
Oscillin	Oscillin	1628828_s_at	0.2610	0.6351	0.4254	0.1223	0.6682	0.0359	-0.0335	0.9766	0.2422	0.4025	0.2757	0.2785	-0.1644	0.9340	0.4034	0.5405	0.5678	0.3807
CG4624	CG4624	1628829_at	-0.1340	0.6598	-0.3420	0.0556	-0.2355	0.2175	0.0769	0.8856	0.2478	0.1692	0.1709	0.2982	-0.1504	0.8940	0.0571	0.9325	0.2076	0.6471
l(3)S147910	lethal (3) S14791(	1628830_at	-0.0401	0.9167	-0.1081	0.5756	-0.1812	0.4618	-0.0967	0.8650	0.1404	0.5098	0.2371	0.1876	-0.0689	0.9657	-0.1344	0.8117	-0.0655	0.9124
torp4a	torp4a	1628831_at	-0.2434	0.2030	0.1811	0.3492	0.4119	0.0592	-0.0687	0.8776	-0.3443	0.0318	-0.2756	0.0491	-0.2468	0.7826	0.1422	0.7591	0.3890	0.3271
---	---	1628832_s_at	0.0723	0.7368	0.0924	0.4737	0.4376	0.0383	0.0724	0.9037	0.0081	0.9789	-0.0644	0.7628	0.0343	0.9826	0.1048	0.8255	0.0705	0.8877
CG7311	CG7311	1628833_at	-0.0754	0.6971	-0.1495	0.4685	-0.1935	0.4536	0.0873	0.9339	0.3138	0.3157	0.2265	0.4364	-0.0414	0.9732	-0.0231	0.9633	0.0182	0.9687
SelD	selD-like	1628834_a_at	-0.1239	0.6257	0.8795	0.0253	0.6651	0.0029	-0.0664	0.8717	-0.5922	0.0023	-0.5258	0.0023	0.0865	0.9499	0.3582	0.3733	0.2716	0.5274
CG14534	CG14534	1628835_at	-0.2121	0.2025	-0.6629	0.2776	-0.9755	0.0018	-0.0352	0.9808	0.2350	0.5368	0.2701	0.4166	-0.1143	0.8097	-0.2165	0.2540	-0.1022	0.6313
---	---	1628836_at	0.3787	0.2033	0.2122	0.2176	0.2351	0.1330	0.0020	0.9985	0.0834	0.7640	0.0814	0.7452	-0.0001	0.9999	-0.0887	0.8123	-0.0886	0.8003
---	---	1628837_at	0.0353	0.8324	0.0668	0.5260	0.0978	0.6207	0.0472	0.9247	0.0043	0.9846	-0.0429	0.7981	0.0418	0.9653	0.0490	0.9002	0.0072	0.9860
CG15286	CG15286	1628838_at	0.3179	0.1074	0.0392	0.7758	0.3210	0.1225	0.1570	0.6945	0.1293	0.5073	-0.0277	0.9006	-0.1167	0.9093	-0.0451	0.9395	0.0716	0.8864
CG17110	CG17110	1628839_at	-0.1090	0.6476	0.0266	0.8608	0.1617	0.2808	0.1804	0.6888	-0.0897	0.7101	-0.2701	0.1470	-0.0804	0.8875	-0.1132	0.6209	-0.0328	0.9109
ken	ken barbie	1628840_at	-0.0719	0.7468	-0.3275	0.1998	-0.4380	0.0546	0.0589	0.9599	0.1880	0.5706	0.1291	0.6888	0.1861	0.8431	-0.0219	0.9735	-0.2080	0.6169
CG31522	CG31522	1628841_at	-0.0408	0.8061	0.0828	0.5003	0.0948	0.6031	-0.0030	0.9962	-0.1629	0.4316	-0.1599	0.3892	0.0148	0.9898	0.1825	0.4785	0.1677	0.5268
cals	calsyntenin-1	1628842_s_at	-0.1183	0.5349	-0.0088	0.9854	-0.5759	0.0256	-0.3974	0.2067	-0.1633	0.4015	0.2342	0.1655	0.2746	0.7644	0.0637	0.9151	-0.2108	0.6116
Ipod	Interaction partne	1628843_at	0.0813	0.7644	0.1243	0.3423	-0.0285	0.8846	-0.0648	0.9441	-0.2366	0.3813	-0.1719	0.4987	0.0406	0.9619	-0.0366	0.9216	-0.0772	0.7847
WDY	CG40449	1628844_at	0.1032	0.5648	0.1639	0.2579	0.0726	0.6708	-0.0910	0.8281	-0.0723	0.6965	0.0187	0.9248	0.1101	0.8400	0.0806	0.7692	-0.0295	0.9242
---	---	1628845_at	-0.1296	0.5703	-0.0958	0.4056	-0.1284	0.5473	-0.0044	0.9956	0.3079	0.1963	0.3123	0.1432	-0.1657	0.7953	0.0157	0.9723	0.1814	0.5166
CG17806	CG17806	1628846_at	0.4582	0.1283	0.0107	0.9575	0.2422	0.1842	0.0379	0.9649	0.4932	0.0381	0.4552	0.0336	-0.2225	0.8049	0.0157	0.9828	0.2382	0.5490
CG17167	CG40146	1628847_a_at	0.3075	0.1566	0.2372	0.3376	0.2127	0.2572	-0.3303	0.4455	-0.1635	0.5120	0.1668	0.4543	-0.1033	0.8910	-0.1875	0.5251	-0.0842	0.8112
CG17083	CG17083	1628848_at	0.0330	0.8904	0.3073	0.0416	0.3201	0.0503	0.0791	0.8810	-0.1995	0.2701	-0.2786	0.0844	0.0216	0.9816	0.0724	0.7763	0.0508	0.8491
par-1	Par-1 kinase	1628849_at	0.0849	0.7105	0.1823	0.6641	-0.1050	0.5543	-0.4695	0.2299	0.0603	0.8409	0.5298	0.0191	-0.1975	0.8882	0.3120	0.5756	0.5096	0.3475
CG7056	CG7056	1628850_at	0.1005	0.6094	0.1980	0.1353	0.1169	0.5205	-0.1474	0.6854	-0.1447	0.4059	0.0027	0.9898	0.0921	0.8973	0.0424	0.9178	-0.0497	0.8918
---	---	1628851_at	0.2942	0.2376	0.0710	0.6267	0.0596	0.7499	-0.0165	0.9838	-0.0390	0.8782	-0.0224	0.9235	-0.0142	0.9894	-0.0611	0.8386	-0.0470	0.8774
CG14262	CG14262	1628852_at	0.1067	0.5516	0.0418	0.7209	-0.0618	0.7046	-0.3245	0.4529	-0.1036	0.7018	0.2209	0.2989	-0.0432	0.9672	-0.0940	0.7860	-0.0508	0.8941
CG8915 /// DmirCG8915	CG8915	1628853_at	0.6168	0.0428	0.1427	0.6198	-0.0989	0.5961	-0.2775	0.4873	0.7081	0.0060	0.9856	0.0009	0.0458	0.9793	0.2749	0.5110	0.2291	0.5988
CG32252	CG32252	1628854_at	0.1043	0.5739	-0.0122	0.9166	-0.0339	0.8880	-0.0263	0.9677	-0.0384	0.8646	-0.0121	0.9556	-0.1831	0.8076	-0.1358	0.7017	0.0472	0.9125
---	---	1628855_s_at	0.3274	0.0787	0.0974	0.6069	0.2486	0.1375	0.1505	0.6998	0.0510	0.8194	-0.0995	0.5769	0.0522	0.9515	0.0615	0.8657	0.0093	0.9835
---	---	1628856_at	-0.2020	0.6908	-0.2905	0.3662	-0.7218	0.0218	-0.3616	0.7023	0.1554	0.7656	0.5170	0.1822	-0.0609	0.9816	0.0959	0.9157	0.1568	0.8307
---	---	1628857_at	-0.0668	0.7187	-0.1274	0.4522	0.0521	0.7492	0.0130	0.9865	-0.0292	0.9107	-0.0422	0.8456	-0.0549	0.9306	-0.0473	0.8687	0.0076	0.9832
---	---	1628858_at	0.0569	0.7091	0.2317	0.3600	0.2368	0.2619	0.1017	0.9182	0.1789	0.5890	0.0772	0.8234	-0.0303	0.9742	0.0885	0.7439	0.1188	0.6317
shi	dynamin	1628859_at	0.8417	0.1452	1.0203	0.0100	0.5221	0.2048	-0.0725	0.9339	0.2583	0.3201	0.3308	0.1481	0.4672	0.8202	0.5684	0.5183	0.1013	0.9326
---	---	1628860_at	0.1524	0.5568	0.1336	0.5255	0.0635	0.7457	-0.1648	0.6884	-0.2054	0.2844	-0.0406	0.8545	0.0464	0.9717	0.1144	0.7738	0.0680	0.8779
CG7120	CG7120	1628861_at	-2.1776	0.0043	-0.9523	0.0077	-1.6031	0.0001	-0.7000	0.1119	-1.7681	0.0004	-1.0682	0.0014	-0.0864	0.9589	-0.4524	0.3259	-0.3660	0.4546
tankyrase	tankyrase	1628862_at	0.2610	0.5050	0.3382	0.5289	0.5187	0.0372	-0.0469	0.9538	0.1068	0.6729	0.1537	0.4657	-0.0970	0.9735	0.2192	0.8144	0.3162	0.6918
---	---	1628863_at	0.1879	0.4331	0.1310	0.2671	0.0170	0.9399	-0.0374	0.9518	-0.0377	0.8673	-0.0003	0.9991	-0.0566	0.9412	-0.0719	0.8165	-0.0153	0.9659
Gr39b	Gustatory recepto	1628864_at	0.1414	0.4804	-0.1139	0.4097	-0.0102	0.9756	0.1330	0.8508	0.3052	0.2367	0.1722	0.4839	-0.1017	0.9142	-0.0742	0.8751	0.0275	0.9525
opa	odd-paired	1628865_at	0.1014	0.7041	-0.0366	0.7141	0.1041	0.5092	0.0598	0.9422	-0.0175	0.9604	-0.0773	0.7642	-0.1226	0.8016	-0.1690	0.4015	-0.0464	0.8626
---	---	1628866_at	0.1981	0.5491	0.8199	0.0833	0.5634	0.0073	0.0401	0.9518	0.4740	0.0176	0.4340	0.0159	0.3210	0.8185	1.0120	0.0881	0.6910	0.2437
CG33334 /// CG5740	CG5740 /// CG33	1628867_s_at	-2.1228	0.0014	-1.9444	0.0226	-2.7545	0.0000	-0.1447	0.8014	-0.4316	0.0652	-0.2869	0.1640	0.7346	0.6272	-0.2926	0.6754	-1.0272	0.1313
CG9778	CG9778	1628868_at	0.0980	0.5201	-0.1326	0.4320	0.0699	0.6482	0.0378	0.9538	0.0548	0.8026	0.0169	0.9393	-0.0698	0.9059	-0.1646	0.4361	-0.0949	0.6857
CG11504	CG11504	1628869_s_at	-0.3102	0.3325	0.3031	0.1839	0.1672	0.4634	-0.0393	0.9603	-0.3710	0.0797	-0.3317	0.0796	0.0711	0.9689	0.2673	0.5971	0.1962	0.7138
---	---	1628870_at	0.2602	0.2477	-0.0742	0.6755	0.0468	0.8110	0.0884	0.8921	0.2747	0.2070	0.1863	0.3528	0.0026	0.9984	0.0040	0.9943	0.0013	0.9979
CG11145	CG11145	1628871_at	0.1021	0.5152	-0.0059	0.9588	0.0075	0.9795	0.0389	0.9624	-0.0363	0.9045	-0.0752	0.7536	-0.0078	0.9939	-0.0625	0.8439	-0.0547	0.8577
CG32351	CG32351	1628872_at	0.1086	0.6347	-0.0521	0.6449	-0.1709	0.2553	-0.0449	0.9325	-0.0214	0.9223	0.0235	0.9009	0.0310	0.9779	-0.0903	0.7874	-0.1213	0.6817
Mio	Mlx interactor	1628873_s_at	0.8870	0.0090	0.3688	0.0621	0.5929	0.0249	0.1318	0.7759	0.3614	0.0654	0.2296	0.1838	-0.1050	0.9342	-0.0908	0.8764	0.0142	0.9838
CG15200	CG15200	1628874_at	0.0165	0.9275	0.0771	0.6901	-0.1361	0.6051	-0.0232	0.9688	0.0127	0.9557	0.0359	0.8404	0.0482	0.9635	0.1026	0.7678	0.0544	0.8902
pain	painless	1628875_at	-2.1467	0.0005	-1.8036	0.0330	-2.2226	0.0001	0.0813	0.8914	0.0311	0.9108	-0.0502	0.8253	0.4272	0.7810	0.1393	0.8850	-0.2879	0.6978
---	---	1628876_at	0.0725	0.6947	0.0053	0.9826	0.2245	0.1847	0.0838	0.8578	0.0406	0.8527	-0.0432	0.8211	-0.2011	0.7196	-0.0571	0.8593	0.1440	0.5591
---	---	1628877_at	0.1104	0.5580	0.0428	0.6600	0.1164	0.5857	-0.1144	0.8663	0.0092	0.9801	0.1236	0.6102	-0.0150	0.9901	0.1883	0.4868	0.2033	0.4573
---	---	1628878_at	0.1419	0.3832	0.0488	0.6064	0.0632	0.6803	-0.0157	0.9860	-0.0116	0.9720	0.0041	0.9875	0.0581	0.9342	0.0214	0.9531	-0.0368	0.9067
CG13223	CG13223	1628879_at	0.1436	0.4492	0.0013	0.9980	-0.0547	0.8464	-0.1244	0.8507	0.1342	0.6131	0.2587	0.2304	-0.0687	0.9317	0.0242	0.9534	0.0929	0.7560
fau	fau	1628880_at	-0.0248	0.9102	-0.0759	0.5017	-0.1752	0.2935	-0.0061	0.9943	0.0673	0.7553	0.0734	0.7006	0.0237	0.9892	0.0891	0.8604	0.0654	0.8952
---	---	1628881_at	0.0529	0.7490	0.0068	0.9659	-0.0309	0.8596	-0.0644	0.9068	0.0578	0.7899	0.1222	0.4683	-0.1463	0.7899	-0.0752	0.8018	0.0712	0.8039
CG7378	MKP-like	1628882_at	-1.3895	0.0055	-1.3886	0.0925	-1.7622	0.0002	-0.2516	0.6367	-0.1203	0.6880	0.1313	0.6218	0.1472	0.9514	-0.0425	0.9716	-0.1897	0.8325
CG7484	CG7484	1628883_at	0.0469	0.8287	0.3600	0.1886	0.2677	0.0971	-0.1479	0.7923	0.1043	0.6905	0.2521	0.2171	-0.1279	0.8628	0.3292	0.2517	0.4571	0.1512
PGRP-SA	semmelweis	1628884_at	1.4450	0.0112	-0.4671	0.6802	0.2996	0.2014	0.3485	0.5461	1.9739	0.0004	1.6254	0.0005	-0.3688	0.9083	-0.0886	0.9622	0.2803	0.8474
CG32698	CG32698	1628885_at	1.9930	0.0118	1.2093	0.2084	2.1306	0.0008	0.1089	0.8498	-0.0207	0.9470	-0.1296	0.5246	-0.7087	0.7826	-0.5497	0.6497	0.1591	0.9176
CG4629	CG4629	1628886_at	-0.0722	0.7577	0.0269	0.8819	-0.0242	0.9082	-0.1166	0.8251	0.0080	0.9800	0.1246	0.5325	-0.0299	0.9742	-0.0310	0.9333	-0.0011	0.9979
CG32112	CG32112	1628887_a_at	-0.5172	0.0400	-0.3452	0.0354	-0.4826	0.0090	-0.4116	0.1363	-0.8537	0.0009	-0.4421	0.0099	-0.3807	0.5120	-0.6753	0.0383	-0.2946	0.2960
CG7180	CG7180	1628888_at	-2.6052	0.0032	-3.0088	0.0481	-4.4985	0.0000	-0.7776	0.0144	-0.4585	0.0123	0.3191	0.0353	0.8206	0.7893	-0.8256	0.5426	-1.6462	0.2221
---	---	1628889_s_at	0.2075	0.3692	0.0196	0.9136	-0.0223	0.9419	0.1511	0.7732	0.1141	0.6443	-0.0370	0.8878	0.0703	0.9462	-0.0108	0.9865	-0.0812	0.8408
CG11873	CG11873	1628890_at	-0.2211	0.1983	0.1164	0.7236	0.3261	0.2136	0.2072	0.5324	0.0080	0.9762	-0.1992	0.1979	-0.1476	0.8963	0.2679	0.5431	0.4155	0.3374
---	---	1628891_at	-0.0070	0.9896	-0.6547	0.1231	-0.5266	0.2969	0.0177	0.9952	0.7370	0.2189	0.7193	0.1797	-0.4056	0.7770	-0.1846	0.8196	0.2211	0.7577
CG10984	CG10984	1628892_s_at	-0.7886	0.0019	-0.2833	0.2768	-0.1838	0.3274	-0.0160	0.9852	-0.2879	0.1343	-0.2720	0.1138	-0.0483	0.9589	0.2358	0.3645	0.2841	0.2964

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15589	CG15589	1628893_at	-1.7021	0.0034	-0.8013	0.0314	-0.6731	0.0102	-0.0121	0.9931	-1.3747	0.0010	-1.3625	0.0006	-0.1701	0.8999	-0.5478	0.2518	-0.3777	0.4622
CG34340	CG10017	1628894_a_at	-0.0406	0.8483	-0.0736	0.6622	-0.0427	0.8904	0.0679	0.9204	-0.1398	0.5254	-0.2077	0.2651	0.0621	0.9635	0.0805	0.8791	0.0184	0.9751
ham	hamlet	1628895_at	0.1012	0.4540	0.1052	0.4950	-0.2746	0.2938	0.0002	0.9999	0.0774	0.7558	0.0771	0.7296	0.3761	0.6749	0.1471	0.7260	-0.2290	0.5514
CG4210	CG4210	1628896_a_at	-0.3637	0.5166	-1.0302	0.2377	-0.5534	0.0411	0.3916	0.2180	0.0998	0.6415	-0.2918	0.0877	0.0104	0.9986	-0.5374	0.6605	-0.5478	0.6476
---	---	1628897_at	0.1472	0.4351	0.1773	0.1688	0.0368	0.8536	-0.1700	0.7711	-0.0617	0.8471	0.1084	0.6740	0.0272	0.9831	-0.0347	0.9429	-0.0619	0.8797
---	---	1628898_at	-0.0647	0.7951	-0.1246	0.4561	0.0679	0.7451	0.1118	0.7973	0.2529	0.1505	0.1410	0.3905	-0.1253	0.8076	-0.0417	0.8966	0.0836	0.7342
CG13534	CG13534	1628899_at	0.1394	0.6317	0.0094	0.9305	0.0707	0.6878	-0.0234	0.9664	0.0351	0.8526	0.0585	0.7029	0.0120	0.9916	-0.0537	0.8903	-0.0658	0.8474
---	---	1628900_at	-0.1182	0.4133	0.0221	0.8273	0.1779	0.5161	0.1312	0.6958	0.1588	0.3100	0.0275	0.8806	-0.1636	0.8400	0.1316	0.7371	0.2952	0.3891
Asx	Additional sex cor	1628901_at	-0.3398	0.3206	0.1098	0.7138	-0.3593	0.1668	0.0137	0.9866	0.0450	0.8642	0.0314	0.8959	0.3510	0.8033	0.4669	0.4253	0.1159	0.8861
mRpS11	mitochondrial ribo	1628902_at	-0.7413	0.0453	0.2653	0.2199	0.5229	0.1804	0.0235	0.9863	-1.0733	0.0065	-1.0968	0.0035	0.0291	0.9914	0.0292	0.9782	0.0001	0.9999
Sin3A	Enhancer of GMR	1628903_s_at	0.2344	0.3193	0.3643	0.2987	0.0283	0.8696	-0.1417	0.8180	-0.0172	0.9623	0.1244	0.6096	0.1891	0.8235	0.2678	0.4483	0.0787	0.8666
---	---	1628904_at	-0.0249	0.9009	0.1732	0.3798	0.2337	0.1130	0.0765	0.8717	-0.0976	0.5866	-0.1740	0.2396	-0.0159	0.9901	-0.0563	0.8903	-0.0403	0.9169
---	---	1628905_at	0.4585	0.0198	0.0281	0.8258	-0.3303	0.4164	-0.1432	0.8460	0.4210	0.1182	0.5642	0.0276	-0.0832	0.9495	-0.1089	0.8353	-0.0257	0.9646
---	---	1628906_at	-0.0045	0.9809	-0.0951	0.5016	-0.0159	0.9389	-0.0830	0.8752	0.1227	0.5306	0.2057	0.2097	-0.0351	0.9779	0.0183	0.9711	0.0534	0.8982
T3dh	Type III alcohol de	1628907_at	-0.2875	0.4547	-0.2274	0.5041	0.1576	0.5857	0.0069	0.9956	-0.2627	0.4715	-0.2697	0.4074	-0.5153	0.6458	-0.3166	0.4947	0.1987	0.6909
CG5075	CG5075	1628908_at	0.0375	0.8585	0.0369	0.8393	0.1105	0.5054	0.0009	0.9988	0.0374	0.8732	0.0365	0.8599	-0.0374	0.9705	0.1173	0.6827	0.1546	0.5709
CG9686	CG9686	1628909_at	0.2340	0.3951	-0.0897	0.6592	0.2153	0.1724	0.2835	0.5311	0.1534	0.5467	-0.1301	0.5811	-0.0309	0.9775	-0.1343	0.6344	-0.1034	0.7291
---	---	1628910_at	0.0788	0.6735	0.1354	0.4276	0.1883	0.5238	0.0586	0.9538	0.0212	0.9593	-0.0374	0.9111	0.1112	0.8270	-0.0144	0.9672	-0.1255	0.5808
---	---	1628911_at	0.2835	0.1712	0.0642	0.6356	0.2534	0.1251	0.1591	0.7248	0.1520	0.4747	-0.0071	0.9778	-0.0745	0.9076	-0.0410	0.9068	0.0335	0.9160
CG12672	CG12672	1628912_at	-0.1716	0.3674	-0.8420	0.0126	-1.0325	0.0003	-0.0543	0.9380	0.5275	0.0180	0.5818	0.0071	0.0444	0.9653	-0.2532	0.3438	-0.2976	0.2926
CG32365	CG32365	1628913_at	0.7630	0.0164	0.2139	0.7562	0.4741	0.0813	0.0066	0.9922	0.2514	0.0769	0.2448	0.0564	-0.1156	0.9657	-0.1334	0.9049	-0.0178	0.9875
CG3194	CG3194	1628914_at	-0.5963	0.0962	0.2612	0.6559	0.6369	0.0125	0.3358	0.6720	-0.2229	0.5853	-0.5587	0.0960	0.1690	0.9324	0.7880	0.2042	0.6191	0.3476
Exn	CG3799	1628915_s_at	-0.4070	0.3679	-0.0207	0.9812	-0.1199	0.6619	-0.3908	0.2042	-0.8954	0.0012	-0.5046	0.0087	-0.2390	0.9244	-0.4540	0.6139	-0.2149	0.8412
---	---	1628916_at	0.3508	0.2430	0.1021	0.5247	0.0379	0.8821	0.1585	0.6615	0.5067	0.0105	0.3482	0.0319	0.2934	0.7485	0.2242	0.5880	-0.0692	0.8977
HDC02577	HDC02577	1628917_at	0.1395	0.3586	0.0590	0.5868	-0.1188	0.5854	-0.0989	0.8578	0.1124	0.6075	0.2113	0.2363	0.0694	0.9238	0.0632	0.8478	-0.0062	0.9874
CG17478	CG17478	1628918_at	0.0135	0.9683	-0.3209	0.1502	-0.1744	0.4371	-0.1038	0.8827	0.0958	0.7343	0.1996	0.3676	-0.2710	0.6955	-0.1962	0.5176	0.0748	0.8414
---	---	1628919_at	0.1251	0.5659	-0.0386	0.7132	0.2755	0.1681	0.0596	0.9493	0.0965	0.7574	0.0369	0.9074	-0.0712	0.9441	0.0403	0.9350	0.1115	0.7526
CG12237	CG12237	1628920_at	-0.8273	0.0040	-0.3569	0.6181	-0.0605	0.7788	0.0991	0.8578	-0.4954	0.0213	-0.5946	0.0058	-0.2099	0.9238	-0.1449	0.8939	0.0650	0.9494
mod(mdg4)	Modifier67.2	1628921_s_at	0.1286	0.5273	0.3608	0.1233	0.3576	0.0360	-0.0659	0.8942	0.0221	0.9232	0.0880	0.5941	-0.0513	0.9611	0.2081	0.4939	0.2593	0.3914
veg	vegetable	1628922_s_at	0.1610	0.3770	0.1000	0.6829	-0.0379	0.8737	0.1889	0.5511	0.4722	0.0111	0.2834	0.0570	0.2560	0.8016	0.4402	0.2854	0.1842	0.6978
form3	formin 3	1628923_at	-0.1268	0.6097	-0.9232	0.0312	-0.9978	0.0379	-0.0261	0.9915	0.7109	0.1677	0.7370	0.1107	-0.1064	0.8885	-0.1188	0.7162	-0.0124	0.9784
CG13567	CG13567	1628924_a_at	0.1785	0.4364	-0.1362	0.4737	-0.1609	0.2832	-0.0022	0.9970	0.3304	0.0605	0.3326	0.0384	-0.0110	0.9939	-0.0799	0.8634	-0.0689	0.8788
CG17625	CG17625	1628925_at	0.0298	0.9390	-0.0639	0.6591	0.2329	0.1281	0.2090	0.4518	0.2206	0.1421	0.0116	0.9517	-0.1270	0.9061	0.0396	0.9499	0.1666	0.6993
Pde11	Phosphodiesteras	1628926_at	-0.1507	0.5499	0.3103	0.1792	0.3789	0.0276	0.0651	0.8836	-0.0968	0.5530	-0.1619	0.2326	-0.0120	0.9939	0.2135	0.5647	0.2254	0.5461
yellow-b	yellow-b	1628927_at	-0.1897	0.8684	-1.2407	0.1638	-0.5725	0.4260	0.5336	0.3837	0.6512	0.0635	0.1176	0.7561	-0.0752	0.9924	-0.2273	0.9397	-0.1521	0.9530
RpL31	Ribosomal protein	1628928_s_at	0.2713	0.0845	1.2976	0.0069	1.7364	0.0114	0.5365	0.6839	-0.8188	0.1788	-1.3554	0.0215	0.0864	0.8909	0.0053	0.9925	-0.0811	0.7713
CG1902	CG1902	1628929_s_at	-0.6879	0.0094	-0.6503	0.0340	-0.6750	0.0021	-0.2345	0.3454	0.1662	0.2464	0.4007	0.0077	-0.0768	0.9063	0.2575	0.2518	0.3343	0.1767
swi2	swi2	1628930_at	-1.4176	0.3876	-1.5440	0.0632	-2.5071	0.0006	-0.7120	0.6718	-1.0232	0.1968	-0.3113	0.7129	0.0413	0.9964	-1.4613	0.5062	-1.5026	0.4952
CG3301	CG3301	1628931_at	0.7774	0.0339	-0.1205	0.5573	0.2099	0.2550	0.0361	0.9672	0.1501	0.5423	0.1140	0.6251	-0.2926	0.7768	-0.7498	0.1009	-0.4573	0.3153
mod(mdg4)	Modifier67.2	1628932_at	-0.5653	0.0620	0.0281	0.8634	0.0023	0.9950	-0.0955	0.9017	-0.5187	0.0447	-0.4232	0.0628	-0.1265	0.8882	0.1037	0.8131	0.2302	0.5141
CG40153 /// CG41138	CG40153 /// CG4	1628933_at	-0.0364	0.8167	0.0569	0.5944	0.0495	0.7944	0.0606	0.9271	-0.0099	0.9743	-0.0705	0.7408	0.0205	0.9776	0.0277	0.9184	0.0072	0.9801
---	---	1628934_at	-0.0641	0.6768	-0.0291	0.8835	0.1341	0.5135	0.0350	0.9603	0.0464	0.8464	0.0114	0.9606	-0.0602	0.9306	0.0470	0.8845	0.1072	0.6577
CG10962	CG10962	1628935_at	0.7437	0.0331	2.1569	0.0082	2.7326	0.0002	0.7376	0.0408	-0.7849	0.0029	-1.5225	0.0001	0.1316	0.9474	0.8977	0.1381	0.7661	0.2231
---	---	1628936_at	-0.0842	0.6485	-0.1117	0.4549	-0.0633	0.7276	-0.0613	0.9228	0.1015	0.6364	0.1628	0.3555	-0.0178	0.9839	0.1117	0.6012	0.1294	0.5403
rin	rasputin	1628937_s_at	1.2692	0.0525	1.2991	0.0223	1.0703	0.0078	-0.1539	0.8281	0.6117	0.0321	0.7656	0.0076	0.2771	0.9066	0.6591	0.4351	0.3820	0.6824
CG10882	CG10882	1628938_at	0.8549	0.1116	1.1168	0.0989	1.4876	0.0001	0.4641	0.2920	0.8929	0.0050	0.4287	0.0673	0.0391	0.9914	1.1142	0.1536	1.0751	0.2051
---	---	1628939_at	0.1944	0.4033	0.1774	0.2307	-0.0322	0.9086	-0.0758	0.9318	0.0980	0.7552	0.1738	0.4909	0.1304	0.8692	0.0108	0.9867	-0.1196	0.7439
---	---	1628940_at	0.1431	0.4180	0.1655	0.3577	-0.1793	0.5336	-0.1813	0.8293	-0.1240	0.7457	0.0573	0.8807	0.2027	0.8122	0.0872	0.8584	-0.1156	0.7822
CG32628	CG32628	1628941_at	-0.5729	0.0301	0.0102	0.9225	0.0184	0.9477	0.0388	0.9586	-0.6161	0.0078	-0.6549	0.0035	0.1701	0.8680	0.1367	0.7871	-0.0334	0.9533
CG15385	CG15385	1628942_at	0.7072	0.1991	1.0343	0.0289	0.8563	0.0036	0.0153	0.9863	0.5295	0.0230	0.5142	0.0163	0.2610	0.8861	0.8350	0.2093	0.5740	0.4096
Tsf2	Transferrin 2	1628943_at	-0.5750	0.0182	-0.1868	0.6842	-0.3109	0.1136	0.1749	0.6755	0.2640	0.1768	0.0890	0.6620	0.3008	0.8033	0.6069	0.2126	0.3061	0.5658
CG13634	CG13634	1628944_at	0.0408	0.9071	0.3211	0.0843	0.0925	0.6233	-0.0615	0.9376	-0.0078	0.9819	0.0537	0.8396	0.0604	0.9672	0.1437	0.7554	0.0833	0.8729
CG33275	CG33275	1628945_at	-0.0848	0.6969	-0.0869	0.6096	-0.0278	0.9051	-0.0668	0.9153	-0.0560	0.8190	0.0108	0.9657	-0.0751	0.9467	-0.1002	0.8250	-0.0251	0.9601
lola	longitudinals abse	1628946_at	-0.7989	0.0554	-0.7405	0.3996	-0.9402	0.0119	0.0863	0.9467	0.3383	0.3647	0.2520	0.4675	0.3247	0.8903	0.4905	0.6041	0.1659	0.8918
mnd	minidiscs	1628947_s_at	1.4152	0.0020	1.7769	0.0233	2.0154	0.0006	0.4002	0.3665	0.8061	0.0071	0.4058	0.0734	-0.0052	0.9990	1.0417	0.1293	1.0469	0.1512
CG14408 /// DsimCG1440	CG14408	1628948_a_at	0.5094	0.4396	1.1259	0.0816	0.4695	0.1058	-0.2479	0.5132	0.1915	0.3477	0.4394	0.0224	0.4900	0.8558	0.6511	0.5778	0.1611	0.9162
shn	quo vadis	1628949_at	0.1526	0.3920	0.1185	0.5940	0.1977	0.2903	-0.0324	0.9610	0.1020	0.5964	0.1344	0.4155	0.0876	0.9168	0.1515	0.6281	0.0639	0.8687
CG10178	CG10178	1628950_at	-0.5012	0.0318	1.1364	0.0074	-0.0956	0.8614	-1.3674	0.1140	-2.5289	0.0011	-1.1614	0.0204	0.0981	0.9390	-0.5732	0.1406	-0.6713	0.1210
CG13739	CG13739	1628951_at	0.1569	0.5328	0.1234	0.3125	0.3083	0.0690	-0.0332	0.9567	-0.2049	0.2122	-0.1717	0.2457	0.0241	0.9816	-0.0364	0.9193	-0.0605	0.8368
Fur1	Furin-like protease	1628952_s_at	-2.0009	0.0156	-1.6966	0.0608	-2.1674	0.0001	0.0862	0.9333	0.0697	0.8612	-0.0165	0.9658	0.6254	0.7726	0.4742	0.6413	-0.1513	0.9067
CG7131	CG7131	1628953_at	0.3761	0.0784	0.0842	0.4523	0.0981	0.7232	0.2086	0.6489	-0.0022	0.9945	-0.2107	0.2931	0.1796	0.8049	-0.1472	0.6596	-0.3268	0.2964
CG11898	CG11898	1628954_at	-0.0694	0.7632	0.4699	0.1008	0.2498	0.1258	-0.1597	0.8057	-0.3540	0.1687	-0.1943	0.4225	0.2772	0.6660	0.2976	0.2375	0.0204	0.9564
Ance-3	Ance-3	1628955_at	-1.4721	0.0564	1.3092	0.1494	1.3549	0.0272	-0.6254	0.6615	-2.3600	0.0050	-1.7346	0.0122	-0.5976	0.8077	0.3694	0.7660	0.9670	0.3592
BCL7-like	BCL7-like	1628956_at	-0.3652	0.3491	0.3763	0.4621	0.6682	0.2531	-0.1825	0.9028	-0.4339	0.3802	-0.2514	0.6069	-0.4811	0.8441	0.2283	0.8737	0.7094	0.4954
l(2)35Bd	lethal 9 in the blac	1628957_at	0.1493	0.5044	-0.0267	0.8896	0.1262	0.4818	0.0015	0.9986	0.1151	0.5490	0.1136	0.5098	-0.0485	0.9588	0.0718	0.8374	0.1204	0.6728
CG13442 /// DyakCG1344	CG13442	1628958_at	0.1006	0.6618	0.0688	0.6830	0.0496	0.7727	0.0670	0.9067	-0.0179	0.9485	-0.0849	0.6556	0.1374	0.8461	-0.0352	0.9404	-0.1726	0.5789
---	---	1628959_at	-0.4796	0.0411	0.3016	0.0758	0.1043	0.5569	-0.1748	0.5644	-0.5223	0.0057	-0.3475	0.0208	-0.1134	0.8881	0.0968	0.8016	0.2102	0.5010
CG17829	CG17829	1628960_s_at	0.5141	0.0114	0.2221	0.4870	0.3753	0.0688	-0.0788	0.9314	0.3643	0.1764	0.4432	0.0699	-0.3002	0.6557	-0.0104	0.9841	0.2897	0.3007
CG32494	CG32494	1628961_at	0.0483	0.7849	0.3445	0.1022	0.2450	0.2387	-0.0878	0.8396	-0.0310	0.8873	0.0567	0.7464	0.3409	0.6272	0.3359	0.2361	-0.0050	0.9924

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG12708	CG12708	1628962_at	-0.1085	0.6113	0.0690	0.5562	0.1680	0.4123	-0.0156	0.9838	-0.0248	0.9220	-0.0092	0.9679	-0.1653	0.7779	0.0742	0.8244	0.2395	0.3601
CG4716 /// DsimCG4716 ///	CG4716	1628963_at	2.2455	0.0077	0.7605	0.1326	1.8085	0.0018	0.5813	0.7217	0.4521	0.5696	-0.1291	0.8820	-0.3214	0.8206	-0.8280	0.1498	-0.5066	0.3982
CG4804	CG4804	1628964_at	-1.7892	0.0196	-1.8224	0.0269	-1.8795	0.0000	-0.2821	0.2965	-0.5870	0.0037	-0.3049	0.0384	-0.1219	0.9589	-0.4106	0.5597	-0.2886	0.7008
CG4194	CG4194	1628965_at	-0.6430	0.0627	-0.7836	0.0257	-0.4898	0.0097	0.5929	0.0899	0.6583	0.0069	0.0654	0.7725	0.0215	0.9914	0.2464	0.5798	0.2249	0.6186
---	---	1628966_at	0.0278	0.8997	0.2810	0.2731	0.1842	0.3527	0.0545	0.9592	-0.0175	0.9679	-0.0720	0.8214	0.0494	0.9701	0.0707	0.8850	0.0212	0.9665
Eig71Ec	Gene III	1628967_at	0.0998	0.5921	0.0356	0.8209	-0.0558	0.7975	-0.0183	0.9854	0.1022	0.6946	0.1205	0.5950	0.0339	0.9677	-0.0251	0.9445	-0.0590	0.8325
CG18469	CG18469	1628968_at	-0.0238	0.9241	-0.0143	0.8992	0.0462	0.8214	0.1162	0.8315	0.0724	0.7727	-0.0438	0.8567	0.0604	0.9409	-0.0196	0.9627	-0.0800	0.7925
CG8485	CG8485	1628969_s_at	0.6898	0.0412	0.1880	0.1252	0.0575	0.8248	-0.0718	0.8578	0.0113	0.9579	0.0830	0.5595	0.1708	0.8810	-0.4708	0.2625	-0.6416	0.1651
CG11249	CG11249	1628970_at	0.0456	0.8078	-0.0196	0.9282	-0.0623	0.6886	0.0133	0.9860	0.1719	0.3489	0.1586	0.3364	-0.0021	0.9990	0.0312	0.9451	0.0333	0.9323
CG5116	CG5116	1628971_at	0.2035	0.6805	-0.4900	0.2012	-0.1659	0.6662	0.2769	0.8164	0.6824	0.1437	0.4055	0.3453	-0.1028	0.9657	-0.2715	0.7105	-0.1687	0.8368
CG30488	CG30488	1628972_at	0.0526	0.8369	-0.0787	0.6385	0.1212	0.4790	-0.1829	0.7877	-0.0682	0.8506	0.1147	0.6961	-0.2224	0.6594	-0.1649	0.4214	0.0575	0.8240
ey	eyeless	1628973_a_at	0.2252	0.3607	0.0487	0.7218	-0.0651	0.6832	-0.0885	0.8865	0.0513	0.8506	0.1398	0.4856	0.0239	0.9862	-0.0171	0.9760	-0.0410	0.9271
CG32711 /// DsimCG32711	CG32711	1628974_at	-0.3798	0.0550	-0.2809	0.1100	-0.2379	0.3569	-0.0446	0.9491	-0.3753	0.0582	-0.3306	0.0613	-0.0667	0.9577	-0.1777	0.6437	-0.1111	0.7947
---	---	1628975_at	0.1972	0.3834	0.1128	0.5991	0.2967	0.0516	0.1108	0.7432	0.1271	0.4063	0.0163	0.9286	0.0062	0.9957	-0.0041	0.9941	-0.0103	0.9821
CG14222	CG14222	1628976_at	0.3640	0.0509	0.0746	0.7329	-0.4240	0.0503	-0.1176	0.7949	0.3251	0.0794	0.4426	0.0153	0.4507	0.5126	0.0271	0.9605	-0.4236	0.2140
CG12190	CG12190	1628977_at	-0.3784	0.1071	-0.2520	0.4987	-0.4508	0.0117	0.0782	0.8939	-0.1765	0.3756	-0.2547	0.1425	0.1464	0.8972	0.0006	0.9998	-0.1458	0.7674
CG5360	CG5360	1628978_at	0.1985	0.3313	0.2020	0.3686	-0.0085	0.9687	-0.0666	0.9288	0.4502	0.0456	0.5168	0.0165	0.3107	0.6749	0.4129	0.1619	0.1022	0.7754
CG32396	CG32396	1628979_at	-0.4348	0.2567	0.3162	0.3278	0.0719	0.7973	-0.0839	0.9375	-0.2605	0.4163	-0.1766	0.5657	0.1667	0.8283	0.3030	0.3342	0.1364	0.7060
CG6707	CG6707	1628980_s_at	-0.1007	0.6705	-0.1507	0.3453	-0.3862	0.0512	-0.1738	0.6518	0.1858	0.3247	0.3596	0.0373	0.0078	0.9950	0.0879	0.8154	0.0801	0.8256
CG16849	CG16849	1628981_at	-0.0423	0.8515	0.0949	0.4845	0.2201	0.2748	0.1236	0.8124	-0.0098	0.9756	-0.1334	0.5002	-0.0365	0.9717	-0.0108	0.9823	0.0257	0.9433
NPFR1	neuropeptide F re	1628982_at	-1.4691	0.0060	-0.0137	0.9543	-0.1401	0.5391	-0.3921	0.3890	-1.6040	0.0004	-1.2118	0.0008	-0.0237	0.9871	-0.1275	0.7455	-0.1039	0.7967
---	---	1628983_at	0.1335	0.4597	0.1491	0.3491	0.0630	0.7285	-0.0359	0.9592	0.0538	0.8144	0.0897	0.6362	0.1032	0.8513	0.0689	0.8161	-0.0344	0.9124
HmgZ	HMG protein Z	1628984_s_at	-0.2542	0.3796	-0.5586	0.0227	-0.2704	0.1996	0.3015	0.5870	-0.0846	0.8087	-0.3862	0.1250	0.1008	0.9400	-0.4077	0.3087	-0.5085	0.2331
Caki	Camguk	1628985_a_at	-1.5992	0.0291	-2.1831	0.0307	-2.3730	0.0002	-0.0688	0.9540	0.4916	0.1314	0.5604	0.0592	0.2326	0.9460	0.0806	0.9627	-0.1520	0.9161
gp210	gp210	1628986_at	0.5739	0.0185	0.5008	0.4437	0.7595	0.0449	-0.0813	0.9023	0.3753	0.0817	0.4566	0.0255	-0.2256	0.9296	0.4327	0.6259	0.6583	0.4356
CG5869	CG5869	1628987_at	0.4491	0.0610	1.1821	0.0057	0.8367	0.0047	-0.0044	0.9956	0.0030	0.9903	0.0073	0.9735	0.2641	0.7726	0.6609	0.1042	0.3968	0.3293
---	---	1628988_at	0.3067	0.2275	0.0914	0.3981	0.3207	0.0706	0.0164	0.9794	0.0238	0.9129	0.0074	0.9697	-0.0067	0.9958	0.0880	0.8174	0.0947	0.7858
---	---	1628989_at	-0.1085	0.6336	-0.5715	0.0764	-0.5617	0.0325	0.3033	0.5008	0.7175	0.0093	0.4143	0.0566	0.0691	0.9390	-0.0856	0.8123	-0.1547	0.6052
Hmgcr	columbus	1628990_at	-1.0801	0.0149	0.2072	0.6910	0.5150	0.0878	-0.1301	0.8666	-1.0087	0.0037	-0.8786	0.0040	-0.3311	0.8292	0.3949	0.5589	0.7260	0.2757
colt	congested-like tra	1628991_at	0.6398	0.0148	0.6427	0.1709	0.9274	0.0012	0.1522	0.7777	-0.1592	0.5019	-0.3113	0.1201	-0.0741	0.9611	-0.2291	0.6173	-0.1550	0.7529
---	---	1628992_at	-1.2943	0.0067	-0.8949	0.0257	-0.7212	0.0047	0.0499	0.9425	-0.2109	0.2879	-0.2607	0.1370	-0.2415	0.8099	0.1066	0.8522	0.3481	0.4114
beat-IIb	beat-IIb	1628993_at	-0.1445	0.4393	0.0833	0.6286	0.0456	0.8371	-0.0210	0.9824	-0.3205	0.1487	-0.2995	0.1310	0.0835	0.8874	0.0227	0.9496	-0.0608	0.8256
---	---	1628994_at	0.3075	0.1713	0.0745	0.4770	0.0562	0.7980	0.0069	0.9937	-0.0039	0.9891	-0.0108	0.9629	0.1245	0.8802	0.0343	0.9467	-0.0902	0.8184
CG8006	CG8006	1628995_at	0.0410	0.8545	-0.0157	0.8896	0.1636	0.4648	0.2347	0.5628	0.1048	0.6555	-0.1299	0.5199	0.0554	0.9589	-0.0914	0.8165	-0.1468	0.6534
---	---	1628996_at	-0.0187	0.9555	-0.3541	0.2478	-0.1442	0.5905	0.1948	0.6641	0.3455	0.1061	0.1508	0.4575	0.1938	0.7893	-0.0747	0.8625	-0.2686	0.3939
CG14763	CG14763	1628997_at	0.0618	0.7291	0.0712	0.5547	0.1913	0.5604	0.2207	0.7409	0.1589	0.6264	-0.0618	0.8567	0.2405	0.7070	0.2440	0.3469	0.0035	0.9942
CG31907	CG31907	1628998_at	-0.0614	0.6947	0.0591	0.5897	0.2401	0.2310	0.0151	0.9834	-0.0983	0.5837	-0.1133	0.4703	-0.0473	0.9599	0.0245	0.9553	0.0718	0.8310
---	---	1628999_at	0.1245	0.4578	0.0992	0.5849	0.1385	0.3998	0.1957	0.6589	0.1582	0.4776	-0.0375	0.8815	0.1549	0.8449	0.0789	0.8619	-0.0760	0.8577
CG14117	CG14117	1629000_at	-0.4118	0.1663	-0.1919	0.2768	-0.3551	0.0424	-0.0960	0.8875	-0.0072	0.9827	0.0888	0.7153	0.0128	0.9939	0.0910	0.8625	0.0782	0.8785
CG5787	CG5787	1629001_at	-0.0884	0.7805	-0.2019	0.2209	0.1607	0.3994	0.1984	0.6705	0.2948	0.1807	0.0964	0.6775	-0.2709	0.7493	0.2311	0.5405	0.5020	0.1904
---	---	1629002_at	0.0261	0.8797	0.1975	0.1387	-0.0472	0.8338	-0.2250	0.5125	-0.0849	0.6813	0.1401	0.4075	0.0140	0.9862	0.0570	0.8061	0.0430	0.8549
CG6208	CG6208	1629003_at	0.2391	0.2845	0.0713	0.6231	0.1835	0.4072	0.0684	0.9096	-0.0260	0.9226	-0.0945	0.6281	-0.0156	0.9922	-0.0067	0.9935	0.0089	0.9874
---	---	1629004_at	0.1378	0.4587	0.1766	0.1570	-0.0544	0.8113	-0.0375	0.9603	0.0176	0.9531	0.0551	0.8062	0.0451	0.9520	0.1295	0.5860	0.0845	0.7442
CG3209	CG3209	1629005_a_at	-0.1838	0.5243	0.8539	0.0361	1.0360	0.0002	0.0144	0.9847	-0.7829	0.0015	-0.7973	0.0008	-0.1717	0.8825	0.1803	0.7248	0.3520	0.4407
CG30127	CG30127	1629006_at	0.2133	0.3678	0.0699	0.5460	0.0757	0.6558	0.1528	0.7539	0.0659	0.7986	-0.0869	0.6944	0.0555	0.9653	-0.1978	0.5872	-0.2533	0.4764
CG8952	CG8952	1629007_at	-0.2978	0.9226	-0.0374	0.8104	-1.1361	0.0456	-1.1541	0.7550	-1.4199	0.3838	-0.2657	0.8890	-0.0595	0.9952	-1.0433	0.6789	-0.9838	0.6973
CG3817	CG3817	1629008_at	-0.1669	0.4492	0.2750	0.3800	0.3539	0.2311	0.3212	0.4779	0.1692	0.5097	-0.1519	0.5191	0.2329	0.8472	0.5530	0.2522	0.3200	0.5476
Cyp28a5	Cyp28a5	1629009_at	-0.1620	0.6987	0.3375	0.1652	0.4107	0.1336	-0.1702	0.8645	-0.6887	0.0562	-0.5185	0.1009	-0.2130	0.8609	-0.2005	0.7234	0.0125	0.9874
rho-5	rhomboid-5	1629010_at	-0.1314	0.5759	-0.1327	0.4241	-0.1639	0.5356	0.0537	0.9540	0.0359	0.9202	-0.0178	0.9568	-0.0684	0.9093	-0.0989	0.6765	-0.0305	0.9176
CG5476	CG5476	1629011_at	0.0697	0.7578	0.1584	0.3761	0.4946	0.0558	-0.1164	0.8738	-0.1049	0.7255	0.0115	0.9711	0.1188	0.8917	0.2856	0.3740	0.1668	0.6366
capt	act up	1629012_at	-0.6593	0.0390	-0.4144	0.1835	-0.4706	0.0535	0.0726	0.8913	0.2500	0.1595	0.1774	0.2709	0.1373	0.9309	0.6516	0.1886	0.5143	0.3248
CG11436	CG11436	1629013_at	-0.0683	0.8078	-0.1876	0.3704	0.0032	0.9912	-0.0043	0.9956	0.5220	0.0293	0.5262	0.0177	-0.2086	0.8191	0.3058	0.4073	0.5144	0.1911
CG2604	CG2604	1629014_s_at	0.2096	0.4364	0.2637	0.6202	0.6783	0.0059	0.3394	0.2753	-0.5739	0.0078	-0.9133	0.0006	-0.0047	0.9990	-0.4245	0.5131	-0.4198	0.5235
DAAM	Dishevelled Assoc	1629015_a_at	-0.9064	0.0147	-1.7331	0.0297	-2.0167	0.0010	-0.0309	0.9823	0.8301	0.0195	0.8611	0.0102	0.2744	0.8270	0.1567	0.8169	-0.1177	0.8640
unc-13-4A	unc-13-4A	1629016_s_at	0.3608	0.2084	0.1654	0.3894	-0.1435	0.6045	-0.1441	0.8578	-0.0530	0.8924	0.0911	0.7765	0.1619	0.8689	-0.2123	0.6080	-0.3742	0.3475
yemalpha	yemanuclein-alph	1629017_s_at	-0.0963	0.7974	-0.2662	0.5765	-0.5117	0.0210	-0.2604	0.4711	0.1913	0.3327	0.4517	0.0175	-0.0332	0.9906	0.0515	0.9587	0.0846	0.9180
CG8372	CG8372	1629018_a_at	0.0927	0.6840	-0.0231	0.9233	0.2706	0.0983	0.2146	0.4633	-0.1166	0.4823	-0.3311	0.0273	-0.0699	0.9545	-0.2176	0.5647	-0.1477	0.7160
CG13551	CG13551	1629019_s_at	0.3258	0.0543	0.5495	0.0270	0.3568	0.2122	0.0775	0.9016	-0.3802	0.0645	-0.4577	0.0199	0.1712	0.8202	-0.2593	0.4004	-0.4305	0.1911
arm	Beta-catenin	1629020_s_at	-0.3012	0.4964	0.3207	0.5200	0.1068	0.6781	0.2108	0.6005	-0.0788	0.7417	-0.2896	0.1088	0.2823	0.8818	0.4305	0.5751	0.1483	0.8810
CG3356	CG3356	1629021_at	0.0145	0.9629	0.0132	0.9728	0.2896	0.1164	0.0333	0.9592	0.0380	0.8656	0.0047	0.9826	-0.1880	0.8446	0.1628	0.7183	0.3508	0.3835
CG13991	CG13991	1629022_at	0.0458	0.8024	0.2011	0.3324	-0.0517	0.7538	-0.2501	0.3955	-0.1945	0.2404	0.0555	0.7592	0.0813	0.8740	0.0117	0.9724	-0.0695	0.7674
CG40191	CG40191	1629023_s_at	-0.5642	0.0401	-0.7158	0.0880	-0.6131	0.0097	-0.0864	0.8605	0.2414	0.1686	0.3278	0.0437	-0.2029	0.8608	0.1084	0.8685	0.3113	0.5230
ppk25	pickpocket 25	1629024_at	0.1304	0.4333	-0.0443	0.6682	0.1756	0.5305	-0.0672	0.8923	-0.0042	0.9869	0.0630	0.7220	-0.0847	0.9449	-0.0806	0.8792	0.0041	0.9951
PFE	Pray For Elves	1629025_at	0.1924	0.2303	0.1958	0.1411	0.4770	0.0132	0.0312	0.9584	-0.1206	0.4627	-0.1518	0.2865	-0.2205	0.7464	-0.0697	0.8629	0.1508	0.6293
Aats-asp	lethal(2)49Db	1629026_at	0.3450	0.0360	0.3682	0.0323	0.4251	0.0242	0.1436	0.6457	0.4527	0.0095	0.3091	0.0302	0.1828	0.7644	0.5630	0.0515	0.3802	0.1659
CG14662	CG14662	1629027_at	-0.8766	0.0114	-1.4429	0.0077	-2.3063	0.0002	-0.1559	0.8280	0.8861	0.0069	1.0421	0.0020	0.0308	0.9852	0.0358	0.9531	0.0050	0.9941
---	---	1629028_at	0.2035	0.2349	0.1642	0.5666	0.0360	0.8647	0.0734	0.9009	0.1858	0.3377	0.1123	0.5487	0.1153	0.8870	0.0314	0.9496	-0.0839	0.8256
CG5160	CG5160	1629029_at	-0.0499	0.8926	-0.0622	0.8000	-0.1278	0.6840	-0.0922	0.8721	0.0148	0.9604	0.1070	0.5917	-0.1312	0.8861	-0.1564	0.6842	-0.0252	0.9597
CG18659	CG18659	1629030_a_at	-0.5065	0.2126	-0.1098	0.8607	-0.2828	0.2932	0.0748	0.9018	-0.0168	0.9542	-0.0916	0.6528	0.2401	0.9142	0.3980	0.6407	0.1579	0.8834

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629031_s_at	0.2060	0.3698	0.1827	0.3155	0.0948	0.6562	-0.2304	0.7067	-0.1548	0.6173	0.0755	0.8109	0.1922	0.7743	0.0157	0.9752	-0.1766	0.5669
---	---	1629032_at	0.0553	0.7722	0.2137	0.1853	0.3896	0.0259	0.0209	0.9757	-0.1026	0.5696	-0.1235	0.4316	-0.1102	0.8461	-0.0265	0.9447	0.0837	0.7568
CG2277	CG2277	1629033_at	-0.8877	0.0186	-1.8363	0.0048	-1.6191	0.0000	0.2321	0.4707	0.8514	0.0011	0.6193	0.0026	0.1753	0.8454	0.0528	0.9277	-0.1225	0.7785
CG10253 /// DmirCG10253	CG10253	1629034_at	0.0569	0.9732	-0.0321	0.7870	-1.0711	0.0060	-0.9310	0.1170	-1.2611	0.0046	-0.3301	0.3108	-0.0324	0.9963	-1.5685	0.2812	-1.5361	0.3171
CG12290	CG12290	1629035_at	-1.1321	0.0263	-0.2544	0.5753	-0.7614	0.0301	-0.0597	0.9313	0.1667	0.4302	0.2263	0.2153	0.4363	0.8033	0.9047	0.1995	0.4684	0.5433
amx	almondex	1629036_at	-0.0742	0.7071	-0.0155	0.9620	-0.1518	0.4484	0.0343	0.9612	0.0554	0.8128	0.0210	0.9274	0.1745	0.7726	-0.1162	0.6879	-0.2906	0.2778
CG9650 /// DsimCG9650	CG9650	1629037_at	0.2005	0.3650	0.4105	0.0408	-0.0537	0.8390	-0.1183	0.7303	-0.0182	0.9330	0.1002	0.4970	0.2705	0.8157	0.1096	0.8725	-0.1610	0.7756
CG31545	CG31545	1629038_at	0.1081	0.6730	0.0433	0.6698	0.0680	0.6892	-0.0229	0.9814	0.0290	0.9298	0.0519	0.8489	-0.0634	0.9449	-0.0053	0.9935	0.0580	0.8779
asrij	asrij	1629039_at	-0.9559	0.0101	-0.0274	0.9337	0.6459	0.0266	0.2364	0.4455	-1.0707	0.0004	-1.3071	0.0001	-0.5086	0.6898	-0.1757	0.7856	0.3329	0.5439
CG3476	CG3476	1629040_at	-0.5482	0.0333	-0.9714	0.0126	-0.6704	0.1077	0.2730	0.6122	0.6953	0.0184	0.4222	0.0823	-0.0499	0.9845	0.2113	0.7573	0.2611	0.6782
CG15357	CG15357	1629041_at	-0.0703	0.6658	0.0429	0.8009	0.0885	0.6395	0.2261	0.5311	0.0080	0.9784	-0.2181	0.1948	0.0808	0.9138	0.1558	0.5711	0.0750	0.8178
CG15674	CG15674	1629042_at	-0.0471	0.8023	-0.0922	0.4886	-0.3237	0.1607	0.0568	0.9154	0.0950	0.6052	0.0382	0.8434	-0.0392	0.9741	-0.0580	0.8962	-0.0188	0.9661
Gr93d	Gustatory recepto	1629043_at	-0.0693	0.7519	-0.0046	0.9882	0.0330	0.8931	-0.0792	0.9019	0.0577	0.8263	0.1368	0.4943	-0.1640	0.8016	-0.1090	0.7299	0.0550	0.8826
trbl	tribbles	1629044_at	0.0637	0.8609	1.1481	0.0077	1.3048	0.0009	0.1095	0.9218	-1.3562	0.0026	-1.4657	0.0011	0.0102	0.9946	-0.1403	0.7341	-0.1505	0.7053
CG32452	CG32452	1629045_s_at	0.3508	0.2483	0.5636	0.1163	0.6105	0.0242	0.0335	0.9761	0.1852	0.5309	0.1517	0.5835	0.1025	0.9400	0.2980	0.4868	0.1956	0.6680
CG4716 /// DsimCG4716 ///	CG4716	1629046_a_at	2.6569	0.0027	1.1801	0.1015	2.5202	0.0007	0.7292	0.5633	0.3104	0.6741	-0.4188	0.5009	-0.4831	0.7726	-1.0652	0.1376	-0.5821	0.4301
---	---	1629047_at	0.2905	0.1596	0.0153	0.8835	0.1330	0.4760	0.0665	0.9029	0.2491	0.1573	0.1826	0.2503	0.0099	0.9923	0.0155	0.9684	0.0056	0.9879
CG9674 /// DmirCG9674	CG9674	1629048_s_at	-0.7916	0.0209	-0.1691	0.6316	-0.2982	0.0873	-0.4382	0.2501	-1.4223	0.0004	-0.9841	0.0011	-0.2802	0.7576	-0.7914	0.0609	-0.5111	0.2049
CG34370	CG13499	1629049_at	0.1215	0.6080	0.0595	0.6473	-0.0463	0.8558	0.1766	0.7255	0.2026	0.3807	0.0260	0.9252	0.1449	0.8461	0.0459	0.9240	-0.0990	0.7865
CG10477	NF-YB-like	1629050_at	0.1658	0.6608	-0.2490	0.1544	0.0335	0.9052	0.4324	0.1599	0.2963	0.1104	-0.1360	0.4368	0.0795	0.9647	-0.0941	0.8967	-0.1736	0.7558
---	---	1629051_at	0.2368	0.3224	0.0036	0.9878	0.0855	0.6569	-0.0332	0.9542	-0.0267	0.9027	0.0065	0.9736	-0.1398	0.8603	-0.1392	0.6997	0.0006	0.9992
---	---	1629052_at	0.1431	0.4363	-0.0471	0.7168	0.4949	0.0582	0.1570	0.7003	-0.0431	0.8611	-0.2000	0.2330	-0.3056	0.7095	-0.2032	0.5750	0.1024	0.8081
CG8370	CG8370	1629053_at	-0.1092	0.7993	-0.8473	0.0185	-0.6888	0.0192	0.2454	0.5906	0.7908	0.0052	0.5454	0.0165	0.0708	0.9689	0.0246	0.9769	-0.0463	0.9444
CG40143	CG40143	1629054_at	0.1747	0.3285	-0.0405	0.7559	-0.0293	0.8744	0.0188	0.9753	0.0390	0.8368	0.0201	0.9100	-0.0179	0.9874	-0.1441	0.5987	-0.1261	0.6476
Got2	Glutamate oxaloa	1629055_a_at	0.7163	0.0044	0.6017	0.0883	0.6237	0.0152	-0.2050	0.6584	-0.5299	0.0255	-0.3250	0.1023	-0.1732	0.8400	-0.6860	0.0700	-0.5128	0.1729
Eig71Ea	Gene I	1629056_at	0.1519	0.3173	-0.1462	0.3798	0.1428	0.4086	0.2393	0.5290	0.4559	0.0293	0.2166	0.2232	-0.1115	0.9092	-0.0702	0.8893	0.0413	0.9323
---	---	1629057_at	0.1812	0.2218	0.0360	0.7668	0.0253	0.9213	-0.1600	0.6084	0.0579	0.7572	0.2179	0.1187	-0.0797	0.8903	-0.0357	0.9152	0.0440	0.8813
Tsp47F	Tetraspanin 47F	1629058_a_at	-0.2356	0.2429	-0.0856	0.5002	0.3144	0.2306	0.2607	0.5586	0.0226	0.9470	-0.2381	0.2475	-0.1595	0.8736	0.0639	0.9167	0.2234	0.5984
---	---	1629059_at	0.4591	0.1116	0.1804	0.1388	0.1315	0.4724	0.0338	0.9761	0.2608	0.3607	0.2270	0.3806	0.0082	0.9939	-0.0317	0.9328	-0.0398	0.9025
CG40139 /// CG40163	CG40139 /// CG4(	1629060_s_at	-0.0627	0.8216	-0.9769	0.0089	-0.9281	0.0863	0.0877	0.9639	1.0632	0.0442	0.9755	0.0399	-0.2658	0.7691	-0.0765	0.8929	0.1892	0.6476
---	---	1629061_s_at	0.0075	0.9973	-1.7615	0.0426	-1.4298	0.4209	1.5319	0.7556	3.7245	0.0801	2.1926	0.2486	1.1235	0.8292	1.7619	0.4167	0.6383	0.8138
CG13252	CG13252	1629062_at	-0.1096	0.5243	0.0623	0.9017	-0.4857	0.0521	-0.2282	0.5455	0.2514	0.1969	0.4796	0.0140	0.2885	0.8097	0.4618	0.3416	0.1733	0.7651
desert	desert	1629063_at	-0.0091	0.9732	-0.3006	0.3735	-0.2382	0.3606	0.1054	0.8844	0.3558	0.1483	0.2505	0.2603	0.0798	0.9514	0.0099	0.9911	-0.0699	0.8949
ewg	erect wing	1629064_at	0.3814	0.4135	-0.0067	0.9916	-0.0264	0.9490	-0.2894	0.7699	-0.2630	0.5602	0.0264	0.9597	-0.1195	0.9543	-0.5185	0.3899	-0.3989	0.5358
Vha26	E subunit	1629065_s_at	-0.5869	0.0093	-0.8727	0.0414	-1.0965	0.0003	-0.2334	0.3628	-0.1641	0.2638	0.0693	0.6446	-0.0618	0.9643	-0.2993	0.4201	-0.2374	0.5464
GRHRII	Drosophila Corazc	1629066_at	-0.7702	0.4131	-1.0621	0.0936	0.1819	0.7615	0.6483	0.7401	0.0492	0.9705	-0.5992	0.4628	-0.5831	0.8379	-0.3811	0.7971	0.2020	0.9012
mthl7	Mth-like 7	1629067_at	0.3248	0.0664	0.0260	0.7946	0.2259	0.1734	0.0477	0.9233	0.1321	0.3882	0.0845	0.5686	-0.1106	0.8465	0.0123	0.9766	0.1228	0.6317
---	---	1629068_at	0.0087	0.9700	0.1806	0.4108	-0.0283	0.8927	-0.1329	0.8251	-0.3072	0.1849	-0.1743	0.4249	0.1083	0.8609	-0.0769	0.8123	-0.1852	0.4740
CG7504	CG7504	1629069_at	0.0682	0.8892	-0.1269	0.4513	-0.3173	0.2494	-0.2166	0.8069	0.6226	0.0791	0.8392	0.0158	-0.1368	0.9309	0.2742	0.6191	0.4110	0.4367
CG1208	CG1208	1629070_at	-0.5762	0.6553	-2.1588	0.1268	-2.5973	0.0000	-0.6292	0.1627	-0.2015	0.4810	0.4277	0.0812	-0.1925	0.9816	-1.5639	0.4287	-1.3714	0.5042
---	---	1629071_at	-0.0587	0.7097	0.1195	0.4088	-0.0649	0.7171	-0.1381	0.6808	-0.1901	0.2237	-0.0520	0.7617	0.1439	0.7506	0.0538	0.8353	-0.0900	0.6701
su(w[a])	suppressor of whi	1629072_at	-0.6600	0.0229	-0.8221	0.2056	-0.6996	0.0076	-0.1254	0.7708	-0.1009	0.6149	0.0245	0.9112	-0.4031	0.7707	-0.3550	0.5650	0.0482	0.9537
CG32715	CG32715	1629073_at	0.1616	0.5263	-0.2073	0.1419	-0.1711	0.4325	0.1692	0.6144	0.4601	0.0141	0.2909	0.0573	0.1332	0.8882	-0.1028	0.8258	-0.2361	0.5281
nxf2	nuclear RNA expc	1629074_a_at	-0.2986	0.6064	-0.5134	0.1630	-0.8560	0.0630	-0.3222	0.6015	0.5945	0.0604	0.9167	0.0061	-0.3640	0.8564	-0.0079	0.9973	0.3561	0.6985
---	---	1629075_at	-0.1060	0.6916	0.0202	0.8507	0.0141	0.9561	-0.0169	0.9807	-0.1047	0.5695	-0.0878	0.6103	-0.0370	0.9706	-0.1901	0.4592	-0.1530	0.5720
His2B:CG40461	His2B:CG40461	1629076_at	0.2111	0.1671	-0.0325	0.7512	-0.0560	0.7660	0.0069	0.9937	0.0926	0.6468	0.0856	0.6428	-0.2520	0.6999	-0.3182	0.2311	-0.0662	0.8524
CG1021	CG1021	1629077_s_at	0.3209	0.3631	1.1769	0.2029	0.9350	0.0077	-0.0440	0.9553	-0.4670	0.0347	-0.4230	0.0330	0.2004	0.9495	0.2121	0.8750	0.0117	0.9945
Pk61C	Phosphoinositide	1629078_s_at	0.0389	0.9258	-0.1466	0.5861	-0.6560	0.0131	-0.2047	0.6175	0.7563	0.0037	0.9611	0.0008	0.3462	0.7810	0.6232	0.2289	0.2770	0.6318
CG15517	CG15517	1629079_at	0.2975	0.4225	0.0001	1.0000	0.0263	0.8836	-0.0824	0.8899	0.0516	0.8408	0.1340	0.4805	-0.0814	0.9588	-0.1892	0.7027	-0.1079	0.8482
CG41101	CG41101	1629080_at	0.3303	0.0494	0.4803	0.0586	0.1673	0.2899	-0.0318	0.9532	-0.0066	0.9775	0.0252	0.8871	0.1884	0.8427	0.1557	0.7317	-0.0327	0.9525
ord	orientation disrupt	1629081_at	0.0834	0.6775	-0.1266	0.4524	0.0306	0.9006	0.0273	0.9744	0.1957	0.3525	0.1684	0.3786	-0.1308	0.8486	0.0339	0.9409	0.1647	0.5881
CG14407	CG14407	1629082_at	-0.6566	0.0750	-0.2849	0.1368	-0.1584	0.4106	0.0350	0.9507	-0.6428	0.0024	-0.6778	0.0011	-0.1868	0.8795	-0.3768	0.4242	-0.1900	0.7236
CG16704	CG16704	1629083_at	3.2472	0.0033	1.5136	0.2960	2.4598	0.0004	0.8095	0.5808	-0.3814	0.6510	-1.1909	0.0769	0.0805	0.9895	-1.8506	0.1418	-1.9311	0.1574
betaggt-II	beta subunit of tyr	1629084_at	-0.2935	0.0833	0.0924	0.7992	0.2924	0.1483	0.1308	0.7021	-0.0566	0.7639	-0.1874	0.1806	-0.2068	0.8609	0.2611	0.6080	0.4679	0.3375
CG14227	CG14227	1629085_at	-0.0273	0.8816	0.0748	0.5932	0.0123	0.9569	-0.1129	0.8103	0.0655	0.7708	0.1784	0.2920	0.0164	0.9848	0.1860	0.3206	0.1695	0.3878
Scs-fp	succinate dehydrc	1629086_s_at	-0.7141	0.0112	-1.2345	0.0048	-0.9732	0.0005	-0.0347	0.9580	-0.1572	0.3777	-0.1225	0.4577	-0.4045	0.5126	-0.7857	0.0292	-0.3812	0.2132
---	---	1629087_s_at	0.1574	0.3255	-0.0044	0.9745	0.1302	0.4483	-0.1315	0.7857	-0.1059	0.6351	0.0256	0.9162	-0.0730	0.8956	-0.0866	0.7151	-0.0135	0.9654
CG3911	CG3911	1629088_at	0.3149	0.2471	0.8485	0.0688	1.1281	0.0004	0.1483	0.7929	-0.4316	0.0649	-0.5800	0.0126	-0.1463	0.8889	0.1682	0.7060	0.3144	0.4374
CG12963	CG12963	1629089_a_at	-0.2012	0.9008	0.0944	0.6159	-0.3182	0.1205	-0.4790	0.8481	-1.2577	0.1668	-0.7787	0.3540	0.0958	0.9862	-0.6416	0.6483	-0.7374	0.5964
---	---	1629090_at	0.0129	0.9612	-0.0696	0.4702	0.2695	0.2212	0.1231	0.8794	0.1303	0.6811	0.0072	0.9840	-0.1520	0.7707	0.0202	0.9556	0.1722	0.4533
CG5177	CG5177	1629091_at	-2.1627	0.0145	-2.5962	0.0263	-2.9700	0.0001	-0.3940	0.2613	-0.4525	0.0362	-0.0585	0.8026	-0.0669	0.9906	-0.7876	0.5266	-0.7207	0.5722
---	---	1629092_at	0.3213	0.2615	0.1065	0.6767	-0.0944	0.7205	-0.2555	0.6038	0.1443	0.5930	0.3997	0.0731	0.1169	0.8692	-0.0117	0.9828	-0.1286	0.6763
CG14960	CG14960	1629093_at	-0.0827	0.6760	-0.0664	0.5764	-0.2426	0.3554	-0.1713	0.7749	0.1406	0.6105	0.3119	0.1627	-0.0126	0.9898	0.0554	0.8476	0.0680	0.7873
---	---	1629094_at	0.1167	0.5228	-0.0984	0.4968	-0.0290	0.8732	0.0435	0.9356	0.2195	0.1642	0.1760	0.2150	-0.0364	0.9718	-0.0798	0.8153	-0.0433	0.9055
Men	malic enzyme	1629095_a_at	0.9994	0.0522	0.4001	0.4427	1.4230	0.0003	0.6445	0.0657	0.1977	0.3351	-0.4468	0.0215	-0.2607	0.9112	-0.3783	0.6824	-0.1176	0.9181
CG13984	CG13984	1629096_at	0.0696	0.6843	0.2859	0.0727	0.2724	0.2800	0.0635	0.9136	-0.0348	0.8894	-0.0983	0.5953	-0.0607	0.9409	0.0046	0.9935	0.0653	0.8393
kni	radius incompletu	1629097_at	-3.9146	0.0011	-5.0714	0.0014	-4.4610	0.0000	0.0089	0.9951	0.2211	0.4722	0.2122	0.4434	-0.1958	0.9274	-0.5561	0.4285	-0.3603	0.6342
CG7631	CG7631	1629098_at	-0.0568	0.7824	-0.1956	0.2333	-0.0403	0.8229	0.0928	0.8942	0.0932	0.7322	0.0003	0.9994	-0.0008	0.9994	-0.0175	0.9496	-0.0167	0.9435
CG13721	CG13721	1629099_at	-0.0332	0.8827	-0.0676	0.6686	-0.0232	0.8981	0.0964	0.8897	0.1354	0.5933	0.0390	0.8883	-0.0600	0.9467	0.0093	0.9865	0.0693	0.8414

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14634	CG14634	1629100_at	0.2329	0.2830	-0.1714	0.1494	-0.2022	0.1803	0.1566	0.7039	0.4142	0.0357	0.2577	0.1267	0.0608	0.9514	-0.1128	0.7490	-0.1736	0.5859
CG10395	CG10395	1629101_a_at	-0.4794	0.0944	-0.4449	0.1848	-0.3365	0.2727	0.0269	0.9825	0.2757	0.3452	0.2488	0.3448	-0.0701	0.9672	0.1396	0.8112	0.2098	0.6682
---	---	1629102_at	0.1695	0.2634	0.0452	0.7090	0.0150	0.9466	-0.0665	0.8908	0.0321	0.8831	0.0986	0.5340	-0.0933	0.8744	-0.0223	0.9525	0.0711	0.8017
---	---	1629103_at	0.2209	0.2640	-0.0891	0.5065	0.2192	0.2356	0.0457	0.9320	-0.0336	0.8736	-0.0793	0.6253	-0.0327	0.9774	-0.0466	0.9111	-0.0138	0.9743
CG2258 /// DmirCG2258	CG2258	1629104_at	-0.2640	0.4663	-0.4231	0.0846	-0.7473	0.0249	-0.1848	0.7939	0.3553	0.2171	0.5401	0.0426	0.1398	0.9296	0.3498	0.5084	0.2100	0.7159
CG16735	CG16735	1629105_at	0.0506	0.7896	-0.0175	0.9515	-0.0539	0.7385	0.0285	0.9749	0.1735	0.4526	0.1450	0.4969	0.0373	0.9651	0.0493	0.8799	0.0120	0.9727
CG2233	CG2233	1629106_at	0.9995	0.0067	0.4055	0.2801	1.5480	0.0006	0.7137	0.2438	0.2937	0.4350	-0.4200	0.1944	-0.3023	0.7436	-0.2674	0.5068	0.0348	0.9492
RpL7	Ribosomal protein	1629107_at	0.1778	0.2025	0.0255	0.8082	0.1661	0.2803	0.0492	0.9098	0.0187	0.9226	-0.0305	0.8484	-0.0160	0.9841	-0.0429	0.8659	-0.0270	0.9145
---	---	1629108_at	0.3108	0.0904	0.3228	0.0927	0.1479	0.3411	0.0241	0.9714	0.0322	0.8884	0.0081	0.9694	0.1740	0.7266	0.0284	0.9341	-0.1456	0.5163
CG7979	CG7979	1629109_at	-0.3032	0.1292	-0.0766	0.5779	-0.0537	0.7955	-0.1902	0.6650	-0.3151	0.1317	-0.1248	0.5433	-0.2645	0.7387	-0.0163	0.9784	0.2482	0.4755
CG4168	chaoptic-like	1629110_x_at	0.3038	0.3035	0.4897	0.0461	0.2845	0.3686	-0.0019	0.9988	-0.1327	0.6740	-0.1308	0.6460	0.2261	0.7049	0.0921	0.7559	-0.1340	0.6168
MED1	Mediator complex	1629111_at	-0.8224	0.1853	-0.1966	0.7076	-0.2651	0.2832	-0.0154	0.9883	0.0847	0.7778	0.1000	0.7020	-0.0421	0.9914	0.4698	0.6080	0.5120	0.5742
CG5078	CG5078	1629112_at	0.0785	0.6272	0.0500	0.6805	-0.0483	0.7989	-0.0794	0.8791	0.0313	0.8974	0.1107	0.5213	0.0120	0.9914	-0.0168	0.9657	-0.0288	0.9319
mus312	mutagen-sensitive	1629113_a_at	-0.0167	0.9715	-0.1368	0.7138	-0.1457	0.4100	-0.2476	0.5735	0.0127	0.9710	0.2602	0.1948	-0.1484	0.9034	-0.0813	0.9053	0.0671	0.9124
---	---	1629114_at	0.2066	0.3224	-0.0531	0.6079	0.1726	0.3409	0.1325	0.8009	0.2971	0.1581	0.1646	0.4035	-0.1667	0.8097	-0.0733	0.8527	0.0934	0.7840
scrib	scribble	1629115_s_at	-1.1369	0.0260	-1.6159	0.0618	-1.9353	0.0003	-0.4306	0.5533	0.0250	0.9656	0.4555	0.1740	-0.3564	0.8156	-0.5634	0.3631	-0.2069	0.7815
CG14545	CG14545	1629116_at	0.9010	0.4641	-1.6615	0.2184	-1.0057	0.2601	-0.0888	0.9314	2.7713	0.0001	2.8601	0.0000	-0.8495	0.8903	-0.0398	0.9940	0.8096	0.7625
GalNAc-T2	polypeptide GalN/	1629117_at	0.0333	0.9061	0.1635	0.4733	0.3429	0.0976	-0.2350	0.6371	-0.1493	0.5719	0.0857	0.7459	-0.2335	0.8270	0.2743	0.5538	0.5078	0.2677
PGRP-SC2	PGRP-SC2	1629118_at	1.3568	0.0254	2.5939	0.0059	3.8642	0.0000	1.2845	0.0035	1.8260	0.0001	0.5415	0.0081	-0.1277	0.9635	3.0863	0.0105	3.2140	0.0114
CG33221	CG33221	1629119_at	0.1075	0.4976	0.0033	0.9923	-0.0030	0.9881	0.0396	0.9470	0.1286	0.4635	0.0890	0.5975	0.1427	0.8192	0.1562	0.5597	0.0135	0.9727
alpha-Man-I /// CG32682	CG32682 /// alpha	1629120_at	-1.7272	0.0263	-4.4755	0.0080	-4.1099	0.0000	0.3761	0.7307	1.2840	0.0167	0.9079	0.0437	0.1353	0.9721	-1.3140	0.1682	-1.4493	0.1649
CG10337	CG10337	1629121_at	-0.3717	0.2490	-0.9988	0.0400	-1.1591	0.0025	0.2443	0.8074	1.4147	0.0044	1.1704	0.0062	0.3880	0.7338	0.6979	0.1446	0.3098	0.5488
CG11697	CG11697	1629122_at	0.1763	0.4311	0.0634	0.7053	0.3226	0.0876	0.0800	0.9345	0.1910	0.5305	0.1111	0.7138	-0.0213	0.9775	0.0664	0.7593	0.0877	0.6542
spas	Dspastin	1629123_at	0.3305	0.2181	-0.3118	0.1961	-0.2399	0.2770	0.0502	0.9425	0.7717	0.0029	0.7214	0.0023	0.0307	0.9781	0.0703	0.8467	0.0397	0.9151
CG17153	CG17153	1629124_at	-0.2884	0.1918	0.0074	0.9487	0.3521	0.0573	0.0655	0.8794	-0.0102	0.9639	-0.0757	0.6117	-0.1475	0.8439	0.3745	0.2099	0.5220	0.1211
gp210	gp210	1629125_at	-0.0737	0.7751	0.0438	0.6776	0.7200	0.0728	0.0873	0.8140	-0.1378	0.3556	-0.2251	0.0860	-0.4168	0.7187	0.0382	0.9620	0.4550	0.3569
---	---	1629126_at	0.0268	0.8981	-0.0492	0.7236	0.0839	0.5724	0.0107	0.9883	-0.1318	0.4605	-0.1425	0.3676	-0.0678	0.9235	-0.0487	0.8850	0.0191	0.9533
CG6996	CG6996	1629127_at	0.1136	0.6757	0.1899	0.4239	0.1242	0.5173	0.0097	0.9946	-0.0295	0.9479	-0.0392	0.9151	0.1485	0.8284	-0.0045	0.9941	-0.1530	0.6216
CG8701 /// DyakCG8701	CG8701	1629128_at	0.0535	0.8116	0.0856	0.5299	0.0088	0.9729	-0.0125	0.9883	-0.1620	0.4393	-0.1495	0.4306	0.0778	0.9401	-0.0617	0.8979	-0.1394	0.6941
Tie	tyrosine kinase	1629129_at	0.4361	0.2300	-0.4940	0.3313	-0.7736	0.0945	0.1819	0.8764	0.6358	0.1136	0.4539	0.2076	0.5411	0.7330	-0.1216	0.9075	-0.6627	0.3439
Pif1A	PFTAIRE-interacti	1629130_at	-0.0437	0.8223	0.0487	0.8082	-0.0046	0.9853	-0.0276	0.9610	-0.0218	0.9183	0.0058	0.9756	0.1165	0.8692	0.0527	0.9019	-0.0638	0.8636
---	---	1629131_at	0.0292	0.8709	-0.0060	0.9649	-0.0012	0.9959	0.1302	0.7760	0.1489	0.4548	0.0187	0.9372	-0.0175	0.9870	-0.0500	0.8886	-0.0325	0.9231
---	---	1629132_at	-0.1572	0.3550	0.0260	0.8106	-0.0084	0.9666	-0.0596	0.9228	-0.1676	0.3792	-0.1080	0.5571	-0.0345	0.9657	-0.0403	0.9017	-0.0058	0.9866
CG3815	CG3815	1629133_at	0.2527	0.5180	-0.2651	0.5862	-0.5916	0.0325	-0.0275	0.9745	0.8159	0.0031	0.8435	0.0016	0.1492	0.9400	0.0883	0.9269	-0.0608	0.9431
CG15373	CG15373	1629134_at	0.1377	0.6091	0.1020	0.4206	0.1112	0.5851	-0.0340	0.9586	0.0654	0.7497	0.0995	0.5600	0.0228	0.9829	0.1316	0.6104	0.1088	0.6839
---	---	1629135_at	0.1440	0.6091	0.1010	0.5594	-0.0023	0.9956	0.0796	0.8944	0.1474	0.4814	0.0677	0.7566	-0.1239	0.9342	0.0803	0.9144	0.2042	0.7060
capa	Drm-myotropin	1629136_at	0.3042	0.0912	0.2497	0.2224	0.1029	0.7430	0.0181	0.9796	0.0755	0.7046	0.0574	0.7612	0.1646	0.8202	-0.0374	0.9383	-0.2021	0.5163
---	---	1629137_at	0.3081	0.1387	0.1356	0.4658	0.1696	0.3692	0.0324	0.9586	0.0274	0.9040	-0.0050	0.9805	0.0317	0.9862	0.0217	0.9766	-0.0100	0.9879
---	---	1629138_at	0.0778	0.6870	0.1131	0.3801	0.0314	0.8872	-0.0356	0.9602	0.0576	0.8022	0.0931	0.6248	0.0887	0.8991	0.0516	0.8919	-0.0370	0.9173
---	---	1629139_at	0.1477	0.6015	-0.1708	0.3769	0.0415	0.8637	0.1603	0.8111	0.2922	0.2745	0.1319	0.6263	-0.1625	0.7953	-0.0605	0.8721	0.1020	0.7369
CG13058	CG13058	1629140_at	0.3209	0.1153	0.3377	0.2361	0.2064	0.1621	0.0887	0.9005	-0.0356	0.9114	-0.1243	0.5887	0.0491	0.9515	-0.0291	0.9404	-0.0783	0.7861
InR	insulin receptor	1629141_at	-0.3074	0.6050	0.1066	0.6518	-0.5163	0.0773	-0.3448	0.4279	0.0231	0.9489	0.3679	0.0864	0.2489	0.9142	0.1190	0.9263	-0.1299	0.9073
DopR	dopamine D1 rece	1629142_at	0.1468	0.5028	-0.1768	0.2623	-0.0834	0.7329	0.2152	0.5948	0.3991	0.0564	0.1839	0.3265	0.0495	0.9618	0.0218	0.9634	-0.0277	0.9449
CG31563	CG31563	1629143_at	0.0099	0.9561	-0.0638	0.6183	0.1917	0.3034	0.0381	0.9434	0.1404	0.3632	0.1023	0.4776	-0.2587	0.6584	-0.0360	0.9231	0.2227	0.3642
CG15440	CG15440	1629144_at	0.0028	0.9895	-0.0978	0.7077	0.1374	0.4156	0.1101	0.7929	0.0979	0.6016	-0.0122	0.9555	-0.1814	0.8202	-0.0054	0.9941	0.1760	0.6166
HERC2	HERC2	1629145_at	-1.2993	0.2469	-0.5282	0.7672	-1.3995	0.0128	0.0551	0.9603	0.0359	0.9298	-0.0193	0.9586	0.9963	0.8515	0.6424	0.8244	-0.3539	0.9077
CG31699	CG31699	1629146_at	0.1761	0.3389	0.0175	0.9002	0.4238	0.0276	0.1060	0.8609	0.1174	0.6265	0.0114	0.9673	-0.2970	0.6145	-0.0975	0.7413	0.1995	0.4399
mRpL37	mitochondrial ribo	1629147_at	0.1985	0.2757	0.0443	0.6821	-0.0514	0.8414	-0.0989	0.8057	-0.0422	0.8378	0.0567	0.7424	0.0475	0.9741	-0.2019	0.5986	-0.2494	0.5041
CG9776	CG9776	1629148_a_at	-0.5207	0.0729	-0.7623	0.0185	-0.4876	0.0102	0.0367	0.9506	0.0984	0.5803	0.0616	0.7244	-0.2229	0.8091	0.0254	0.9682	0.2482	0.5391
Fibp	Fibp	1629149_at	-0.1779	0.5460	0.9286	0.0269	0.9163	0.0008	0.1379	0.8302	-0.8389	0.0053	-0.9768	0.0016	0.0439	0.9781	0.2245	0.5800	0.1806	0.6630
CG17337 /// DvirCG17337	CG17337	1629150_at	-0.4859	0.1574	-0.8350	0.0230	-0.9967	0.0006	-0.3495	0.2495	0.2273	0.2046	0.5767	0.0041	-0.2107	0.8521	-0.0270	0.9727	0.1837	0.7318
spn-F	spn-F	1629151_at	0.0419	0.8067	0.1421	0.5894	-0.1927	0.4272	-0.0883	0.8589	0.0246	0.9218	0.1129	0.5129	0.0718	0.9514	0.0936	0.8450	0.0218	0.9673
CG31451	CG31451	1629152_at	-1.8269	0.0025	-2.4149	0.0017	-2.5902	0.0000	-0.1537	0.7425	0.0837	0.7279	0.2375	0.1963	-0.0296	0.9901	-0.5158	0.3097	-0.4862	0.3646
CG5991	CG5991	1629153_s_at	0.3494	0.3770	0.7372	0.0407	1.0445	0.0028	-0.1695	0.6506	-0.3532	0.0555	-0.1837	0.2591	-0.4569	0.7644	-0.1244	0.8989	0.3325	0.6334
Or43a	Olfactory receptor	1629154_at	0.1319	0.6846	0.2865	0.1530	0.2972	0.1217	-0.1287	0.8155	-0.2347	0.2813	-0.1060	0.6321	-0.0312	0.9829	-0.1103	0.7906	-0.0792	0.8543
CG4764 /// DereCG4764 ///	CG4764	1629155_at	-0.5131	0.0254	0.4734	0.0179	0.7532	0.0015	0.2634	0.4910	-0.4075	0.0507	-0.6709	0.0037	0.0647	0.9291	0.4636	0.0665	0.3989	0.1210
---	---	1629156_at	-0.2712	0.5664	0.7714	0.0221	0.4431	0.1351	-0.0642	0.9117	-0.3223	0.0811	-0.2581	0.1153	0.1679	0.9421	0.6296	0.3681	0.4617	0.5386
---	---	1629157_at	-0.0389	0.8335	0.1200	0.3700	0.0780	0.6486	-0.0377	0.9563	-0.0067	0.9809	0.0310	0.8890	0.0280	0.9826	0.0078	0.9908	-0.0202	0.9628
CG4946	CG4946	1629158_at	0.3947	0.0749	-0.0065	0.9873	-0.0722	0.8579	0.0703	0.9376	0.0801	0.8040	0.0097	0.9766	0.0945	0.9589	-0.2589	0.6536	-0.3534	0.5252
CG16865	CG16865	1629159_at	0.2922	0.1242	0.4861	0.0644	0.4557	0.0468	0.1411	0.7228	0.2365	0.1847	0.0953	0.5956	0.1316	0.8400	0.4233	0.1199	0.2917	0.2929
---	---	1629160_s_at	-0.4490	0.4694	0.5838	0.4165	-0.6007	0.0662	-0.6360	0.1879	-0.4724	0.1046	0.1637	0.5723	0.5467	0.8202	0.5572	0.5972	0.0105	0.9953
Obp56f	Odorant-binding p	1629161_at	0.0573	0.7312	-0.0077	0.9674	0.0487	0.8110	-0.1082	0.7891	-0.0038	0.9880	0.1044	0.5157	-0.1109	0.8270	-0.0687	0.7978	0.0422	0.8846
Gr63a	Gustatory recepto	1629162_at	-0.0713	0.7992	0.0579	0.5910	0.1242	0.5301	0.1563	0.7185	0.1342	0.5207	-0.0221	0.9268	0.0654	0.9589	0.0952	0.8445	0.0298	0.9525
CG17764	CG17764	1629163_at	0.1288	0.3598	0.1588	0.3010	0.0698	0.7942	0.1948	0.6122	0.1761	0.3717	-0.0188	0.9376	0.1748	0.8024	0.1198	0.7200	-0.0550	0.8915
CG9825	CG9825	1629164_at	-0.1190	0.5123	0.1361	0.4879	0.0735	0.7777	0.0462	0.9334	0.0450	0.8264	-0.0012	0.9955	0.1714	0.8692	0.3151	0.4446	0.1437	0.7641
CG4203	CG4203	1629165_at	0.0481	0.8954	0.5350	0.0230	0.6707	0.0028	0.0365	0.9688	-0.1945	0.4373	-0.2310	0.2910	-0.1551	0.8668	0.1039	0.8319	0.2590	0.4971
---	---	1629166_at	-0.0549	0.7979	0.3085	0.2272	0.2318	0.2483	-0.0628	0.9334	-0.2564	0.2481	-0.1935	0.3399	0.1490	0.8692	0.0670	0.9040	-0.0820	0.8641
CG11060	CG11060	1629167_at	0.2967	0.2140	-0.2537	0.4211	-0.2765	0.1743	0.1192	0.8449	0.3379	0.1314	0.2187	0.2814	-0.1409	0.8331	-0.4189	0.1345	-0.2779	0.3315
Menl-1	Menl-1	1629168_s_at	0.2194	0.4535	0.0059	0.9660	0.0604	0.7657	0.0804	0.8418	0.0599	0.7331	-0.0205	0.9100	-0.0754	0.9530	-0.2242	0.5800	-0.1488	0.7346

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629169_at	0.1853	0.4215	0.2947	0.2394	0.2150	0.1461	0.0146	0.9879	-0.1652	0.4844	-0.1798	0.3909	0.0361	0.9762	-0.0542	0.9027	-0.0903	0.7978
Xpd	Xeroderma pigme	1629170_s_at	0.1436	0.5440	0.4695	0.1314	0.8079	0.0087	-0.2350	0.6615	-0.1552	0.5785	0.0797	0.7809	-0.4671	0.6824	0.1915	0.7221	0.6586	0.1865
---	---	1629171_at	0.3581	0.0538	-0.0260	0.8173	0.1947	0.4214	0.1197	0.8073	0.1793	0.3683	0.0596	0.7845	-0.1662	0.8033	-0.1634	0.5783	0.0027	0.9954
rk	leucine-rich repea	1629172_s_at	1.2700	0.0266	-0.0487	0.9252	0.7093	0.0901	0.8318	0.2753	1.1337	0.0199	0.3019	0.4810	0.1005	0.9672	-0.1603	0.8561	-0.2607	0.7209
beat-Va	beat-Va	1629173_at	-0.0566	0.7346	0.1399	0.4327	0.4041	0.0380	-0.0636	0.9094	-0.0061	0.9816	0.0575	0.7694	-0.0940	0.9238	0.1548	0.6664	0.2488	0.4597
CG30264	CG30264	1629174_at	5.8479	0.0020	4.0195	0.0048	7.2754	0.0000	2.4863	0.1156	1.0243	0.2843	-1.4619	0.0859	-0.6890	0.7095	-0.6698	0.3775	0.0192	0.9874
---	---	1629175_at	0.3398	0.1349	-0.0779	0.6108	0.1383	0.4125	0.0126	0.9863	0.2370	0.1818	0.2244	0.1588	0.0021	0.9990	-0.1273	0.6930	-0.1295	0.6827
vls	Valois	1629176_at	0.0295	0.9000	0.1121	0.3555	0.3754	0.0263	0.2157	0.4968	0.1255	0.4754	-0.0902	0.5922	-0.1144	0.8692	0.2211	0.4212	0.3355	0.2398
---	---	1629177_at	-0.0889	0.6356	0.0057	0.9791	0.3263	0.1517	0.0462	0.9627	-0.0605	0.8612	-0.1066	0.7013	-0.2338	0.7230	-0.0466	0.9157	0.1872	0.5332
CG30181	CG30181	1629178_at	-0.0356	0.9188	-0.0532	0.6223	0.2357	0.1529	0.1029	0.8612	0.0885	0.7186	-0.0144	0.9573	-0.0797	0.9390	-0.0169	0.9761	0.0628	0.8874
CG9465	CG9465	1629179_at	0.0293	0.8692	-0.0034	0.9787	-0.0472	0.8429	-0.0332	0.9586	-0.0946	0.6112	-0.0614	0.7355	0.0293	0.9742	-0.0508	0.8735	-0.0802	0.7557
O-fut2	O-fucosyltransfera	1629180_at	-0.7876	0.3023	0.2229	0.8696	-0.7796	0.3179	-0.0333	0.9603	-0.7403	0.0021	-0.7070	0.0016	0.8612	0.8541	0.0453	0.9924	-0.8160	0.7063
CG33494	CG33494	1629181_at	0.9540	0.0614	-0.5371	0.5955	-0.7103	0.0185	-0.2011	0.8462	0.4987	0.1874	0.6998	0.0453	0.0164	0.9964	-1.0640	0.2442	-1.0804	0.2715
CG10140	CG10140	1629182_at	0.3038	0.2400	0.1560	0.2331	0.2120	0.4138	-0.0160	0.9833	0.0625	0.7640	0.0785	0.6630	-0.0634	0.9711	0.0444	0.9504	0.1078	0.8488
---	---	1629183_at	0.0858	0.8141	0.2169	0.3758	-0.0757	0.6882	-0.1864	0.8086	-0.1920	0.5613	-0.0056	0.9880	0.1366	0.8450	0.0015	0.9991	-0.1352	0.6646
CG3819	CG3819	1629184_at	0.4951	0.2934	-0.0487	0.7495	0.1311	0.4474	0.2553	0.3793	0.1818	0.2705	-0.0735	0.6662	0.0960	0.9606	-0.1892	0.7781	-0.2852	0.6280
CG6734	CG6734	1629185_at	0.2167	0.5085	0.9053	0.0351	0.8496	0.0021	0.2356	0.7118	0.0601	0.8759	-0.1755	0.5345	0.2626	0.7733	0.6831	0.0943	0.4204	0.3002
AlstR	Galanin Receptor	1629186_a_at	-0.0446	0.7758	-0.1764	0.2915	0.1518	0.2876	0.1909	0.5581	0.3287	0.0548	0.1379	0.3739	-0.1698	0.7953	0.0613	0.8764	0.2311	0.4086
wdb	widerborst	1629187_s_at	-0.0883	0.5777	0.2768	0.2486	0.0922	0.6415	-0.2301	0.5904	-0.2277	0.2979	0.0023	0.9929	0.0147	0.9914	0.2226	0.4601	0.2079	0.5033
CG12304	CG12304	1629188_at	0.1533	0.4669	0.0969	0.8184	-0.0519	0.7677	0.1103	0.7970	0.3245	0.0656	0.2142	0.1678	0.2412	0.8235	0.3461	0.4410	0.1049	0.8581
sec5	sec5	1629189_at	-1.0123	0.0057	-0.0005	1.0000	0.2159	0.3029	0.1558	0.6698	-0.7721	0.0017	-0.9278	0.0005	-0.0785	0.9725	0.3604	0.5471	0.4390	0.4597
---	---	1629190_at	0.1809	0.4831	0.4229	0.0599	0.1276	0.4430	-0.1750	0.6890	-0.3437	0.0911	-0.1687	0.3676	0.2195	0.7196	0.1548	0.5613	-0.0647	0.8418
CG6154	CG6154	1629191_s_at	-0.0026	0.9895	0.0263	0.7908	0.2700	0.0890	-0.0256	0.9637	-0.2038	0.1673	-0.1782	0.1784	-0.1149	0.8202	-0.0900	0.7050	0.0249	0.9330
---	---	1629192_at	0.1849	0.3432	0.2191	0.4144	0.2021	0.1519	0.0637	0.9067	-0.0146	0.9552	-0.0784	0.6661	0.0036	0.9964	-0.0176	0.9507	-0.0211	0.9329
---	---	1629193_at	0.3132	0.1132	0.1775	0.3653	0.0375	0.9119	0.0096	0.9903	0.0136	0.9585	0.0039	0.9858	0.1588	0.8202	0.0492	0.9121	-0.1096	0.7439
ade2	adenosine2	1629194_s_at	2.0553	0.0024	0.6379	0.4824	1.8310	0.0000	0.6247	0.0510	0.9954	0.0008	0.3707	0.0317	-0.6182	0.8049	-0.4388	0.7090	0.1794	0.9006
CG4851	CG4851	1629195_at	1.0565	0.0033	0.5627	0.1780	0.6256	0.0279	-0.2455	0.5854	-0.2805	0.2202	-0.0349	0.8992	-0.2934	0.8160	-0.8326	0.1156	-0.5392	0.3109
---	---	1629196_at	0.1145	0.4424	0.0767	0.5378	0.0798	0.6125	0.0379	0.9441	-0.0071	0.9764	-0.0450	0.7961	0.0907	0.8940	0.0390	0.9209	-0.0517	0.8801
CG14372	CG14372	1629197_at	0.1514	0.3586	0.1910	0.3655	0.2187	0.2089	0.0247	0.9688	-0.1015	0.5649	-0.1261	0.4080	0.0003	0.9999	-0.0348	0.9064	-0.0351	0.8948
l(1)G0095	lethal (1) G0095	1629198_at	0.0102	0.9891	-0.0350	0.9540	0.8933	0.0015	0.3795	0.3947	0.1878	0.4739	-0.1917	0.4139	-0.3883	0.8326	0.2642	0.7787	0.6524	0.4016
Ag5r2	Antigen 5-related	1629199_at	0.4102	0.1658	0.0866	0.4959	0.1433	0.4844	0.1802	0.7582	0.2787	0.2702	0.0985	0.7129	-0.0425	0.9807	-0.1768	0.6765	-0.1343	0.7615
CG40113	CG40113	1629200_at	-0.0993	0.5309	-0.1721	0.2993	-0.2276	0.1057	0.1603	0.6557	0.4617	0.0156	0.3014	0.0553	0.0898	0.8611	0.1012	0.6581	0.0115	0.9716
CG5550	Fibrinogen-like	1629201_at	0.2652	0.5523	0.1011	0.5789	0.4417	0.0237	0.2281	0.8164	-0.1360	0.7693	-0.3641	0.2982	0.0114	0.9946	-0.3242	0.4094	-0.3356	0.4058
CG7362	CG7362	1629202_at	-0.0826	0.5676	-0.1307	0.6029	-0.1782	0.4140	-0.0603	0.9482	0.0391	0.9144	0.0994	0.7197	0.0629	0.9550	0.0306	0.9531	-0.0323	0.9434
---	---	1629203_at	0.0580	0.8601	-0.0080	0.9496	0.2286	0.1547	0.0309	0.9641	-0.0103	0.9707	-0.0413	0.8443	-0.0798	0.8738	-0.0877	0.6926	-0.0079	0.9810
CG7705	CG7705	1629204_a_at	0.2641	0.3287	0.0820	0.6051	0.2677	0.1492	0.1473	0.8000	0.0788	0.7812	-0.0684	0.7946	-0.0281	0.9848	-0.0041	0.9955	0.0240	0.9596
CG32032	CG32032	1629205_at	0.2414	0.4219	1.7628	0.0729	2.2472	0.0004	0.2327	0.8164	-1.7266	0.0018	-1.9594	0.0006	-0.4039	0.8236	-0.1890	0.8553	0.2149	0.8200
---	---	1629206_at	0.1058	0.5923	-0.0797	0.4692	0.0055	0.9831	0.1612	0.7023	0.2369	0.2204	0.0757	0.7129	0.0276	0.9816	0.0322	0.9425	0.0046	0.9924
CG6287 /// DsimCG6287	CG6287	1629207_at	0.4825	0.0400	0.4516	0.1002	0.6347	0.0024	0.0819	0.8743	-0.3028	0.0941	-0.3847	0.0254	-0.1391	0.8718	-0.2940	0.3802	-0.1549	0.6811
Obp51a	Odorant-binding p	1629208_at	0.1395	0.4739	0.2256	0.3996	0.1448	0.3030	0.0637	0.9303	-0.0036	0.9903	-0.0673	0.7765	0.0805	0.9174	0.0017	0.9989	-0.0787	0.8145
CG9286	CG9286	1629209_at	0.1072	0.6934	0.1432	0.8101	0.3439	0.1145	0.3425	0.3810	0.2678	0.2243	-0.0746	0.7570	0.1310	0.9495	0.2201	0.7692	0.0891	0.9169
---	---	1629210_at	0.2736	0.2028	0.0169	0.9420	0.0770	0.7005	-0.0771	0.9040	0.0177	0.9542	0.0949	0.6601	-0.0784	0.9452	-0.0622	0.9064	0.0162	0.9767
---	---	1629211_at	0.1092	0.5266	0.0291	0.7960	-0.0226	0.8966	-0.0529	0.9029	-0.0493	0.7702	0.0036	0.9844	0.0326	0.9775	-0.1656	0.5635	-0.1981	0.4841
lox	lysyl oxidase-like	1629212_at	0.1513	0.5199	0.0010	0.9986	0.1007	0.6802	0.2104	0.6927	0.2488	0.3173	0.0384	0.8969	-0.0129	0.9922	0.0358	0.9404	0.0487	0.9057
---	---	1629213_at	0.3485	0.0916	0.0598	0.6779	0.1498	0.3304	0.0445	0.9376	0.0419	0.8463	-0.0027	0.9900	0.0795	0.8861	-0.1003	0.6614	-0.1798	0.3972
---	---	1629214_at	0.3239	0.1402	0.1957	0.3020	0.2252	0.3043	-0.0547	0.9191	0.0006	0.9979	0.0553	0.7600	-0.0326	0.9775	-0.1737	0.5455	-0.1411	0.6331
Kua	Kua	1629215_at	1.2532	0.0230	3.3868	0.0256	2.4950	0.0012	-0.0839	0.9592	-0.5289	0.2173	-0.4450	0.2489	0.7295	0.7506	1.4824	0.1321	0.7529	0.4582
CG32810	CG32810	1629216_at	-0.1119	0.7126	-0.1811	0.2929	-0.2375	0.1452	0.0324	0.9626	0.1652	0.3792	0.1327	0.4427	-0.0339	0.9816	-0.0402	0.9431	-0.0063	0.9921
Cenp-C	lethal (3) 85Aa	1629217_at	0.1588	0.4364	-0.2796	0.2785	-0.4075	0.0450	-0.3612	0.3433	0.6137	0.0128	0.9750	0.0010	-0.1638	0.8439	0.1994	0.5805	0.3632	0.2988
Top3beta	Topoisomerase 3	1629218_at	0.1375	0.7911	-0.1620	0.7448	-0.3628	0.0943	-0.1266	0.8796	0.5391	0.0636	0.6657	0.0175	0.0746	0.9816	0.3088	0.7202	0.2343	0.7969
---	---	1629219_at	0.2302	0.1849	-0.1321	0.4135	0.0362	0.9024	0.2180	0.5492	0.1340	0.4990	-0.0841	0.6670	0.0435	0.9725	-0.1440	0.6842	-0.1875	0.5789
Ilp2	Drosophila insulin	1629220_at	0.4271	0.1250	0.1435	0.3425	-0.0326	0.9130	-0.1986	0.5255	0.1273	0.4559	0.3259	0.0335	0.1299	0.9056	-0.0448	0.9438	-0.1747	0.6877
morgue	modifier of rpr and	1629221_at	-0.1265	0.4629	-0.2975	0.1559	0.1858	0.5515	0.1597	0.7664	0.2702	0.2392	0.1105	0.6384	-0.4135	0.6955	0.1046	0.8636	0.5181	0.2527
Ahcy13	AdoHcyase	1629222_at	0.4123	0.2061	1.0340	0.1167	1.5537	0.0007	0.4466	0.4615	-0.0430	0.9252	-0.4896	0.0953	-0.1095	0.9679	0.4493	0.5523	0.5588	0.4541
---	---	1629223_at	0.3045	0.0777	0.0289	0.7696	0.1827	0.2970	0.0730	0.8817	0.1566	0.3604	0.0836	0.6246	-0.1943	0.7464	-0.1078	0.7058	0.0865	0.7704
CG14420	CG14420	1629224_at	-0.0243	0.9017	0.0541	0.6302	-0.0792	0.7451	-0.0704	0.8942	0.0475	0.8320	0.1179	0.4821	0.0595	0.9246	0.0963	0.6807	0.0368	0.8985
CG31386	CG31386	1629225_s_at	0.0808	0.6039	0.0015	0.9926	0.0176	0.9432	-0.0219	0.9834	-0.1857	0.4614	-0.1638	0.4772	0.1308	0.7893	-0.0436	0.8850	-0.1744	0.4093
CG32507	CG32507	1629226_at	0.0235	0.8892	0.2342	0.2545	0.0281	0.8757	-0.0703	0.8823	-0.2101	0.1937	-0.1398	0.3487	0.2529	0.7464	0.1982	0.5647	-0.0547	0.9042
baz	Bazooka	1629227_at	0.3362	0.7098	1.8033	0.0912	1.1436	0.0017	-0.3480	0.6327	-0.8913	0.0208	-0.5432	0.0891	0.2613	0.9514	0.4754	0.7514	0.2142	0.9026
Hs3st-A	CG33147	1629228_at	-0.5138	0.0948	-0.3324	0.1850	-0.4563	0.0207	-0.2137	0.6650	-0.2632	0.2661	-0.0495	0.8569	-0.1223	0.8973	0.0655	0.9036	0.1877	0.6216
srp	A-box-binding fact	1629229_a_at	1.6142	0.0182	2.2176	0.0094	1.4991	0.0004	-0.1124	0.9011	-0.1512	0.6463	-0.0388	0.9132	0.5689	0.7485	0.4105	0.6109	-0.1584	0.8773
---	---	1629230_at	-0.0466	0.8106	0.1897	0.3428	0.0660	0.7600	-0.1516	0.7345	-0.1924	0.3410	-0.0408	0.8599	0.0185	0.9898	-0.0226	0.9634	-0.0411	0.9199
---	---	1629231_at	-0.0402	0.8402	0.0621	0.6625	-0.0346	0.8319	0.0201	0.9722	-0.0071	0.9746	-0.0272	0.8730	0.0411	0.9530	0.0344	0.9156	-0.0068	0.9846
CG12869	CG12869	1629232_at	-0.6433	0.0251	0.0210	0.9524	-0.0935	0.6398	-0.0988	0.9093	-0.7811	0.0120	-0.6823	0.0134	-0.1575	0.8270	-0.0473	0.9192	0.1102	0.7498
CG16708	CG16708	1629233_s_at	0.0816	0.7484	0.1662	0.7767	0.1900	0.5911	-0.0330	0.9641	0.1508	0.4573	0.1838	0.2982	0.1686	0.9405	0.2462	0.7709	0.0776	0.9363
eca	eclair	1629234_at	0.5226	0.0423	0.6363	0.0412	0.8498	0.0015	0.0943	0.9041	0.3419	0.1773	0.2475	0.2825	-0.0716	0.9309	0.5324	0.0661	0.6040	0.0607
wupA	Troponin I	1629235_s_at	-1.0376	0.0182	-1.6232	0.0304	-2.2828	0.0005	-0.1387	0.6338	-0.1826	0.1991	-0.0439	0.7827	0.4940	0.8215	-0.6771	0.4598	-1.1711	0.2140
---	---	1629236_at	0.0164	0.9328	0.0813	0.5560	0.1785	0.2798	-0.1201	0.7292	-0.0980	0.5573	0.0221	0.9060	-0.0068	0.9946	0.1801	0.4410	0.1869	0.4370
Mes2	Mes2	1629237_at	0.1179	0.5282	-0.0234	0.8156	0.0466	0.8013	0.2490	0.5008	0.4086	0.0431	0.1596	0.3786	-0.0782	0.8909	0.0688	0.8020	0.1470	0.5113

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13047	CG13047	1629238_at	0.0479	0.8785	0.1634	0.3822	0.0073	0.9859	-0.0526	0.9593	-0.2158	0.4448	-0.1632	0.5393	0.2115	0.8270	-0.0083	0.9929	-0.2197	0.6094
CG2246	CG2246	1629239_s_at	-0.2678	0.0943	0.2303	0.2723	-0.5467	0.0068	-0.0899	0.8578	-0.2262	0.2065	-0.1363	0.4175	0.5794	0.3162	0.1945	0.5374	-0.3849	0.2199
Mst84Dc	Male-specific RNA	1629240_at	0.2523	0.2003	0.0868	0.5647	0.2612	0.1261	0.1699	0.7749	0.1447	0.5932	-0.0252	0.9348	0.0627	0.9467	-0.0677	0.8660	-0.1304	0.6773
Ptpa	Phosphotyrosyl pl	1629241_at	-0.5512	0.0445	-0.0878	0.6733	0.1749	0.2344	0.1315	0.8671	-0.4598	0.0977	-0.5913	0.0253	-0.0555	0.9552	0.1798	0.5518	0.2353	0.4253
---	---	1629242_x_at	-0.2812	0.1128	-0.4094	0.1228	-0.5551	0.0178	0.0607	0.8937	0.3338	0.0342	0.2731	0.0483	0.1567	0.8706	0.1859	0.6586	0.0292	0.9582
Dab	disabled	1629243_at	-1.9318	0.0112	-1.3313	0.0575	-1.8790	0.0001	-0.6127	0.5680	-1.2225	0.0333	-0.6098	0.2149	-0.1041	0.9735	-0.7273	0.3340	-0.6232	0.4317
---	---	1629244_at	0.2066	0.2639	0.0317	0.7583	0.0325	0.8659	0.2867	0.2783	0.2795	0.0752	-0.0072	0.9708	0.1186	0.8603	-0.0039	0.9944	-0.1225	0.6817
Gr39a	Gustatory recepto	1629245_at	0.0427	0.8064	0.0240	0.9056	-0.0305	0.9042	0.0170	0.9834	0.0704	0.7530	0.0534	0.8002	-0.0650	0.9420	-0.0404	0.9263	0.0246	0.9492
CG12219	CG12219	1629246_at	-0.3804	0.1020	0.1626	0.7800	0.5290	0.0527	0.1866	0.7278	-0.6491	0.0148	-0.8358	0.0028	-0.0800	0.9816	0.1014	0.9306	0.1813	0.8467
---	---	1629247_at	0.2770	0.3801	-0.1560	0.4327	-0.1107	0.6050	-0.2060	0.7400	0.1372	0.6563	0.3431	0.1592	-0.1764	0.8431	-0.2613	0.4856	-0.0848	0.8599
yrt	lethal(3)87Ek	1629248_at	-0.2861	0.2871	-0.1110	0.4486	-0.4868	0.0112	-0.1167	0.7952	0.0555	0.8054	0.1722	0.3003	0.2241	0.7893	0.1965	0.6068	-0.0276	0.9575
---	---	1629249_at	0.0518	0.7764	-0.0150	0.9238	-0.1538	0.3373	-0.1489	0.6822	-0.0142	0.9550	0.1347	0.3916	0.1070	0.9092	-0.0213	0.9690	-0.1282	0.7389
CG10077	CG10077	1629250_at	-0.1742	0.6091	-0.3674	0.2647	-0.5427	0.0293	-0.0045	0.9956	0.1190	0.5911	0.1235	0.5340	0.2278	0.8882	-0.0484	0.9620	-0.2762	0.6800
Rpb7	Rpb7	1629251_at	0.0157	0.9598	-0.0135	0.9522	0.2970	0.2075	0.0189	0.9803	0.3699	0.0516	0.3510	0.0411	-0.3938	0.7046	0.3085	0.4856	0.7023	0.1336
CG14657	CG14657	1629252_at	0.0421	0.9629	0.4456	0.1655	0.3448	0.1547	0.1382	0.8594	0.4861	0.0847	0.3479	0.1648	0.2651	0.9126	0.7437	0.3733	0.4786	0.6001
RabX2	RabX2	1629253_at	0.0588	0.7392	0.1748	0.2652	0.1495	0.3011	-0.1068	0.7589	-0.1360	0.3730	-0.0292	0.8675	0.0046	0.9952	0.0399	0.8807	0.0354	0.8897
---	---	1629254_at	0.0377	0.8692	-0.1356	0.5889	-0.1536	0.5320	0.1400	0.8233	0.2315	0.3505	0.0915	0.7249	0.0222	0.9862	0.1119	0.7481	0.0897	0.8039
---	---	1629255_at	0.2581	0.1006	-0.0838	0.4060	-0.0854	0.6758	-0.0259	0.9688	0.0812	0.6792	0.1072	0.5218	-0.0205	0.9816	-0.0923	0.6747	-0.0718	0.7527
CG31547	CG31547	1629256_a_at	1.8729	0.4277	-0.7828	0.3110	-0.5934	0.0749	0.5875	0.5019	-0.1541	0.7846	-0.7416	0.0757	0.5123	0.9619	-2.8172	0.3349	-3.3295	0.2830
CG17715	CG17715	1629257_s_at	-0.4703	0.1561	-0.2963	0.3250	-0.7212	0.0043	-0.3825	0.4338	-0.1136	0.7153	0.2689	0.2668	0.0015	0.9997	-0.0693	0.8986	-0.0708	0.8868
nAcRalpha-7E	Dalpha3	1629258_at	0.0608	0.8472	-0.0908	0.5872	0.1657	0.4077	0.0239	0.9903	-0.1141	0.8390	-0.1381	0.7750	-0.0436	0.9653	-0.0293	0.9463	0.0143	0.9727
CG4186	CG4186	1629259_at	-0.0520	0.8735	0.4889	0.2034	-0.0489	0.7840	-0.4402	0.3743	-0.3504	0.2087	0.0898	0.7715	0.1389	0.8949	0.1910	0.6576	0.0521	0.9231
---	---	1629260_at	-0.1042	0.5934	0.1834	0.2223	0.2511	0.1500	-0.1998	0.5948	-0.4384	0.0291	-0.2386	0.1601	-0.1496	0.8122	-0.0672	0.8500	0.0824	0.7931
eIF-4B	Eukaryotic initiatio	1629261_at	-0.0592	0.7484	-0.2025	0.3201	0.0278	0.9162	0.2805	0.6354	0.6275	0.0373	0.3470	0.1801	0.1272	0.8760	0.3543	0.2489	0.2271	0.4900
CG30001	CG30001	1629262_at	0.1931	0.2738	0.1833	0.3598	0.2136	0.2579	-0.0324	0.9610	0.0002	0.9995	0.0325	0.8773	-0.0125	0.9939	-0.0699	0.8995	-0.0575	0.9095
CG13699	CG13699	1629263_at	0.0933	0.5782	-0.0723	0.7385	-0.0439	0.8678	0.1639	0.5765	0.2536	0.0926	0.0897	0.5428	-0.1056	0.9088	0.1483	0.6830	0.2539	0.4498
CG13025	CG13025	1629264_at	0.0888	0.6718	-0.1232	0.4513	-0.5702	0.0497	-0.2018	0.7857	0.2404	0.4512	0.4422	0.1061	-0.1094	0.8940	-0.0667	0.8790	0.0427	0.9194
dib	disembodied	1629265_at	-0.0176	0.9413	0.0932	0.4133	0.1792	0.5391	-0.2608	0.6656	-0.3353	0.2452	-0.0745	0.8194	-0.2054	0.7726	-0.0639	0.8850	0.1415	0.6719
CG13349	CG13349	1629266_s_at	0.3593	0.0270	0.2454	0.1097	0.4633	0.0097	0.1663	0.5823	0.0703	0.6909	-0.0960	0.5213	-0.0392	0.9589	0.0430	0.8939	0.0822	0.7433
---	---	1629267_at	0.0675	0.6931	0.1776	0.3011	-0.0127	0.9474	0.1018	0.8526	0.2209	0.2684	0.1190	0.5398	0.1904	0.7215	0.2301	0.2874	0.0397	0.8969
---	---	1629268_at	0.0274	0.8908	0.0233	0.8510	0.0713	0.6992	-0.0500	0.9491	0.0170	0.9583	0.0670	0.7837	0.0921	0.8480	0.0322	0.9170	-0.0598	0.8073
CG32204	CG32204	1629269_at	-0.2150	0.2017	-0.1466	0.2251	-0.3323	0.0562	-0.0970	0.8293	0.0651	0.7511	0.1622	0.3003	-0.1131	0.8490	-0.0475	0.8987	0.0656	0.8334
Sry-delta	serendipity delta	1629270_at	-0.1663	0.6429	1.0398	0.0570	1.1188	0.0004	0.1979	0.6923	-0.7901	0.0052	-0.9880	0.0012	0.0626	0.9816	0.2851	0.6673	0.2225	0.7487
CG10444 /// DsecCG10444	CG10444	1629271_at	0.7648	0.0115	-0.0335	0.8459	-0.0854	0.7571	-0.1548	0.7511	0.5417	0.0189	0.6965	0.0037	-0.2339	0.8033	-0.4250	0.2591	-0.1911	0.6498
CG14230	CG14230	1629272_at	0.2360	0.3830	0.2604	0.2543	0.8316	0.0141	0.3297	0.3126	0.1726	0.3767	-0.1571	0.3735	-0.0954	0.9405	0.2217	0.5988	0.3170	0.4336
---	---	1629273_at	0.2988	0.0960	0.0303	0.8266	0.0334	0.8989	0.0125	0.9860	0.1209	0.5013	0.1084	0.5103	0.1134	0.8517	-0.0167	0.9680	-0.1301	0.6311
---	---	1629274_at	-0.0620	0.6838	0.0181	0.8976	0.0537	0.7845	0.0616	0.9022	-0.1267	0.4537	-0.1882	0.1943	0.0503	0.9611	-0.0010	0.9994	-0.0513	0.8963
---	---	1629275_at	0.0299	0.8666	-0.0592	0.6273	-0.1510	0.3283	-0.0068	0.9941	0.0879	0.6857	0.0947	0.6244	0.1464	0.7689	-0.0819	0.7424	-0.2282	0.2960
---	---	1629276_at	0.1166	0.4136	0.0640	0.7891	0.0459	0.7728	-0.0797	0.8460	0.0985	0.5425	0.1782	0.1848	0.0479	0.9588	0.0910	0.7729	0.0431	0.9033
CG31475	CG31475	1629277_at	-1.6045	0.0073	-0.3053	0.3761	-0.8180	0.0013	-0.2660	0.7753	-1.5262	0.0029	-1.2601	0.0040	0.0116	0.9928	-0.3201	0.2814	-0.3316	0.2945
---	---	1629278_at	-0.5913	0.0332	-0.4826	0.1468	-1.1944	0.0061	-0.1110	0.9387	0.5022	0.2350	0.6132	0.1042	0.0652	0.9451	0.2318	0.4165	0.1667	0.5862
CG8211	CG8211	1629279_at	0.3908	0.3209	-0.1162	0.8313	-0.1600	0.4273	0.0849	0.8822	0.3558	0.0725	0.2708	0.1210	0.1297	0.9611	-0.0760	0.9499	-0.2057	0.8256
CG12344	CG12344	1629280_at	-0.2163	0.6683	-0.0330	0.7314	-0.2842	0.5656	-0.3698	0.7121	-0.3540	0.4555	0.0158	0.9781	-0.5406	0.7230	-0.4749	0.4743	0.0656	0.9431
Dhh1	DEAD/DEAH RNA	1629281_at	0.1989	0.5455	-0.0183	0.9659	-0.4078	0.0707	-0.3860	0.4299	0.5194	0.0577	0.9054	0.0033	-0.1526	0.9168	0.3025	0.5679	0.4552	0.3800
---	---	1629282_at	0.1764	0.4545	-0.0064	0.9620	0.2474	0.2140	0.0031	0.9961	0.0750	0.7529	0.0719	0.7380	-0.1670	0.7644	-0.1682	0.4803	-0.0012	0.9979
CG33298	CG33298	1629283_at	0.2812	0.1354	0.3308	0.0974	0.3265	0.0524	-0.0747	0.9060	-0.0684	0.7824	0.0063	0.9801	0.0796	0.9174	0.2434	0.3458	0.1638	0.5610
CG32387 /// DmirCG32387	CG32387	1629284_a_at	0.1086	0.5466	0.0696	0.5813	0.1387	0.4929	-0.0315	0.9647	-0.0316	0.9030	-0.0001	0.9995	0.0319	0.9798	0.0881	0.8111	0.0561	0.8857
CG9617 /// DereCG9617 ///	CG9617	1629285_at	-0.6436	0.0237	-0.0504	0.6675	-0.1628	0.6023	-0.0281	0.9649	-0.1955	0.2491	-0.1675	0.2730	0.0506	0.9793	0.2822	0.5455	0.2316	0.6311
Lag1	longevity	1629286_s_at	-0.3908	0.0634	-0.2295	0.2146	-0.0937	0.7655	0.1899	0.7111	0.2635	0.2594	0.0736	0.7754	-0.0873	0.9495	0.1974	0.6605	0.2848	0.5032
CG14130	CG14130	1629287_at	-0.0552	0.8667	0.3127	0.3800	0.4511	0.0455	-0.0080	0.9943	0.1668	0.4852	0.1747	0.4123	-0.2447	0.8202	0.4399	0.3091	0.6846	0.1498
CG12608 /// CG9123	CG12608 /// CG9	1629288_s_at	0.7241	0.0112	0.4904	0.0875	0.5394	0.0186	0.1544	0.6223	0.6587	0.0021	0.5043	0.0040	0.0318	0.9862	0.4657	0.2518	0.4338	0.3149
CG31921	CG31921	1629289_at	0.2756	0.1278	0.3019	0.3104	0.1830	0.4111	-0.1833	0.5799	-0.1120	0.5364	0.0713	0.6884	-0.0592	0.9737	-0.0752	0.9115	-0.0159	0.9835
Smox	Smad on X	1629290_at	-0.1391	0.7148	0.6211	0.0534	0.3044	0.1171	-0.1999	0.6580	-0.7302	0.0056	-0.5303	0.0142	-0.0777	0.9672	0.0938	0.9009	0.1716	0.7683
---	---	1629291_at	-0.0296	0.8967	0.0348	0.7908	0.0901	0.5739	-0.0851	0.8578	-0.0574	0.7801	0.0277	0.8919	-0.0029	0.9978	0.1023	0.7151	0.1053	0.7020
---	---	1629292_at	-0.0719	0.6314	0.1213	0.4733	0.1756	0.2346	-0.0497	0.9186	0.0728	0.6690	0.1225	0.3768	-0.2301	0.6692	0.0963	0.6915	0.3264	0.1562
CG32635	CG32635	1629293_at	-0.2896	0.1223	-0.2328	0.1388	-0.5843	0.0457	-0.2142	0.7093	0.0365	0.9213	0.2507	0.2903	0.1251	0.8472	0.1001	0.7554	-0.0250	0.9471
---	---	1629294_at	0.1111	0.5917	-0.0704	0.6639	0.1506	0.3907	-0.0154	0.9805	0.0487	0.7965	0.0640	0.6918	-0.1970	0.7506	-0.1844	0.4997	0.0126	0.9764
CHES-1-like	Checkpoint suppr	1629295_at	0.5323	0.4675	0.1271	0.7798	-1.2310	0.0067	-1.0908	0.2227	0.1566	0.8128	1.2474	0.0172	0.2254	0.9462	-0.1366	0.9328	-0.3620	0.7568
CG15882	CG15882	1629296_at	0.0516	0.8374	-0.0039	0.9745	-0.0672	0.8004	-0.0083	0.9922	-0.0902	0.6554	-0.0819	0.6586	0.0115	0.9914	-0.0800	0.7824	-0.0915	0.7338
---	---	1629297_at	0.2952	0.1519	-0.1145	0.4315	0.1145	0.4596	0.1574	0.6854	0.3226	0.0764	0.1652	0.3155	-0.1990	0.8202	-0.0531	0.9265	0.1459	0.7229
Gr94a	Gustatory recepto	1629298_at	0.0913	0.7900	-0.0247	0.8553	0.0440	0.8146	-0.0570	0.9494	0.0145	0.9702	0.0715	0.7991	-0.1354	0.8611	-0.1555	0.6483	-0.0201	0.9659
CG33524	CG33524	1629299_at	0.0812	0.7229	-0.0465	0.6594	0.1346	0.3957	0.0369	0.9568	0.1276	0.5099	0.0907	0.6268	-0.0111	0.9898	0.0593	0.8062	0.0705	0.7457
---	---	1629300_at	0.3941	0.0261	-0.0260	0.9022	0.0790	0.6339	0.0656	0.8860	0.2831	0.0708	0.2175	0.1156	0.0078	0.9950	-0.0476	0.9151	-0.0554	0.8884
---	---	1629301_at	0.1808	0.4432	0.0854	0.4748	0.2259	0.3022	0.1637	0.6937	0.1745	0.3742	0.0108	0.9648	0.0581	0.9342	0.0511	0.8745	-0.0070	0.9849
CG30447	CG30447	1629302_at	0.1338	0.3971	0.2045	0.2787	0.0468	0.8128	-0.0191	0.9792	0.1585	0.3938	0.1775	0.2780	-0.1510	0.7893	-0.0277	0.9429	0.1233	0.6328
CG32832	CG32832	1629303_at	0.0568	0.7261	-0.0385	0.7715	0.2103	0.2056	0.1075	0.8449	-0.0280	0.9215	-0.1355	0.4834	0.0151	0.9901	0.0129	0.9784	-0.0023	0.9965
Cpr50Cb	CG6305	1629304_at	-0.1067	0.6838	-0.2289	0.0884	0.0749	0.7475	0.2600	0.5634	0.3748	0.1042	0.1148	0.6259	-0.1201	0.8379	0.0458	0.9004	0.1659	0.5092
Nup214	Nup214	1629305_at	0.3935	0.0919	0.1656	0.3228	0.3491	0.2346	-0.0982	0.8958	0.4227	0.0891	0.5209	0.0268	-0.2608	0.7230	0.1607	0.6425	0.4215	0.2075
X11L	X11L	1629306_at	-0.6946	0.3224	0.1850	0.7071	-0.1172	0.6631	-0.0857	0.9030	-0.4558	0.0510	-0.3700	0.0722	0.1254	0.9742	0.4335	0.6895	0.3081	0.7915

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
hdc	Fusion-6	1629307_s_at	-0.0696	0.8979	-0.9876	0.1741	-0.2128	0.4547	0.0686	0.9461	0.4202	0.1408	0.3516	0.1688	-0.6585	0.7423	-0.4870	0.5860	0.1715	0.8813
CG17192	CG17192	1629308_at	0.2883	0.3316	0.1373	0.6176	0.2345	0.1347	-0.0468	0.9350	0.1843	0.2794	0.2312	0.1254	-0.0167	0.9928	0.0402	0.9531	0.0569	0.9231
CG7627	CG7627	1629309_at	-0.6418	0.0250	-0.2887	0.6720	-0.1755	0.5138	-0.0244	0.9838	-0.2302	0.4258	-0.2059	0.4329	-0.2310	0.8768	-0.0104	0.9935	0.2207	0.7422
CG11265 /// DsimCG11265	CG11265	1629310_at	1.1393	0.1436	0.8535	0.2415	0.8393	0.0344	-0.0259	0.9705	-0.0740	0.7233	-0.0480	0.8124	0.1530	0.9717	-0.2245	0.8858	-0.3775	0.7647
Fancd2	Fancd2	1629311_at	0.2045	0.2189	0.0645	0.5741	0.0681	0.7710	0.0359	0.9486	0.1365	0.3918	0.1006	0.5008	0.0521	0.9545	-0.0975	0.7623	-0.1497	0.6040
Obp57e	Odorant-binding p	1629312_at	0.0931	0.6017	-0.1360	0.2676	-0.1034	0.5455	0.1735	0.6247	0.2233	0.2022	0.0498	0.8009	-0.0619	0.9491	-0.1331	0.6794	-0.0712	0.8483
CG8661 /// DyakCG8661	CG8661	1629313_at	0.4289	0.0695	0.0130	0.9099	0.3035	0.0603	0.0141	0.9860	-0.0048	0.9870	-0.0190	0.9362	-0.1040	0.8882	-0.1558	0.5994	-0.0518	0.8921
CG12997	CG12997	1629314_at	-0.4777	0.0239	-0.0603	0.6619	-0.2664	0.1345	-0.1181	0.8507	-0.5673	0.0211	-0.4492	0.0346	0.1431	0.7726	-0.1243	0.5815	-0.2674	0.2235
---	---	1629315_at	0.0864	0.5918	0.1441	0.4274	0.1354	0.4713	-0.1130	0.7795	-0.0431	0.8428	0.0699	0.6914	-0.0548	0.9460	0.0264	0.9470	0.0812	0.7782
l-t	Inhibitor-t	1629316_at	-0.0618	0.7073	0.2157	0.3782	0.2045	0.2414	0.0087	0.9919	-0.0164	0.9509	-0.0251	0.9067	0.0565	0.9599	0.1644	0.6344	0.1078	0.7749
Nplp3	neuropeptide-like	1629317_at	0.9509	0.2905	0.0208	0.8406	-0.1525	0.3037	-0.1846	0.7855	-0.2560	0.3700	-0.0714	0.8219	-0.1047	0.9826	-1.0726	0.3178	-0.9679	0.3898
CG17104	CG17104	1629318_at	0.0258	0.9528	0.0352	0.8235	0.2316	0.3340	0.3266	0.5931	0.2291	0.4795	-0.0975	0.7767	0.0840	0.9238	0.1940	0.5246	0.1099	0.7456
CG10428	CG10428	1629319_at	0.1494	0.5620	-0.1358	0.4563	-0.2741	0.0923	-0.2184	0.5008	0.1055	0.5694	0.3239	0.0410	-0.1632	0.8141	-0.1735	0.5635	-0.0103	0.9835
---	---	1629320_at	-0.1380	0.4170	-0.2471	0.1973	0.0714	0.6754	0.0821	0.8470	0.0864	0.6164	0.0042	0.9831	-0.0916	0.9016	-0.1081	0.7327	-0.0165	0.9681
ash1	discs absent, sma	1629321_at	0.1373	0.7030	0.1490	0.6940	0.2756	0.1643	0.1491	0.8244	0.0174	0.9655	-0.1317	0.6176	0.1861	0.8647	0.2891	0.5214	0.1030	0.8555
Pde6	Phosphodiesteras	1629322_at	0.2287	0.3951	3.0528	0.0062	0.6148	0.1844	-0.7242	0.3280	-1.3033	0.0089	-0.5791	0.1292	1.6051	0.2388	1.3912	0.0625	-0.2139	0.7940
CG32644	CG32644	1629323_at	0.1273	0.5580	-0.0505	0.8243	0.0106	0.9659	0.0679	0.9306	0.0637	0.8240	-0.0043	0.9880	-0.0688	0.9506	-0.1075	0.7967	-0.0387	0.9337
CG6870	cytochrome B5	1629324_at	3.1681	0.0035	2.2748	0.1026	3.2501	0.0010	0.0211	0.9942	0.1923	0.7907	0.1712	0.7972	-1.0245	0.7644	-0.7880	0.6061	0.2366	0.9054
---	---	1629325_at	0.2003	0.4060	-0.0136	0.9713	-0.2126	0.3345	-0.0579	0.9482	0.4182	0.0946	0.4760	0.0396	0.1504	0.8603	0.3695	0.2643	0.2192	0.5457
eg	eagle	1629326_s_at	-4.3007	0.0009	-2.9148	0.0208	-4.1986	0.0000	-1.1153	0.1675	-1.3968	0.0118	-0.2814	0.5624	0.1490	0.9585	-0.0841	0.9484	-0.2331	0.8135
CG5693	CG5693	1629327_at	0.0219	0.9144	0.1415	0.4549	-0.0257	0.9218	-0.0185	0.9803	0.1690	0.3693	0.1875	0.2579	0.1380	0.8692	0.0843	0.8531	-0.0538	0.9064
---	---	1629328_at	-0.0030	0.9896	-0.0732	0.6155	-0.0610	0.7014	0.0363	0.9669	0.0327	0.9189	-0.0036	0.9901	0.0860	0.8479	0.0716	0.7439	-0.0143	0.9575
---	---	1629329_at	-0.0269	0.8911	-0.0098	0.9288	0.0783	0.6715	-0.0027	0.9956	-0.0632	0.7322	-0.0604	0.7183	-0.0756	0.9168	0.0006	0.9996	0.0762	0.8039
---	---	1629330_at	0.1461	0.6390	0.0045	0.9774	0.0797	0.7007	0.1508	0.7768	0.1598	0.4937	0.0089	0.9740	0.0679	0.9550	0.1342	0.7459	0.0663	0.8915
CG31468 /// DsimCG31468	CG31468	1629331_at	0.1270	0.4215	-0.1485	0.4882	0.1230	0.4536	0.2161	0.6338	0.2773	0.2111	0.0612	0.8063	0.0201	0.9852	0.0216	0.9583	0.0015	0.9979
CG9642	CG9642	1629332_at	0.0158	0.9535	0.0745	0.6214	0.2027	0.2942	-0.1095	0.8679	-0.1449	0.5648	-0.0354	0.8997	-0.0403	0.9653	0.0862	0.7787	0.1266	0.6389
---	---	1629333_at	0.3157	0.1619	-0.1102	0.4390	0.0373	0.8934	0.0866	0.9152	0.2046	0.4382	0.1180	0.6536	-0.0085	0.9923	-0.1945	0.3266	-0.1860	0.3738
CG12552	CG12552	1629334_at	0.1525	0.4893	-0.0029	0.9840	0.0043	0.9831	-0.0136	0.9854	0.1185	0.5018	0.1322	0.3959	0.2541	0.7611	0.3058	0.3733	0.0517	0.9155
---	---	1629335_at	0.1680	0.3667	0.1899	0.2935	0.2110	0.1652	-0.1408	0.7447	-0.0980	0.6460	0.0428	0.8456	-0.1700	0.8192	0.0508	0.9139	0.2208	0.4825
---	---	1629336_at	0.0433	0.8264	0.0789	0.4187	0.0646	0.8437	0.0441	0.9602	-0.0827	0.7661	-0.1268	0.5836	0.0753	0.9400	-0.0115	0.9844	-0.0868	0.8221
CG32721	CG32721	1629337_at	0.3972	0.0864	0.6222	0.1055	0.8583	0.0019	0.0171	0.9833	-0.1979	0.2930	-0.2150	0.1964	-0.3044	0.7628	0.1611	0.7439	0.4655	0.2833
---	---	1629338_at	0.4278	0.0305	0.1134	0.5244	0.1458	0.5305	0.2209	0.7271	0.3785	0.1793	0.1575	0.5769	-0.0399	0.9713	-0.0887	0.8023	-0.0488	0.8984
---	---	1629339_at	0.1426	0.3578	-0.0788	0.7530	0.0215	0.9118	0.1430	0.7228	0.1727	0.3504	0.0297	0.8913	-0.0749	0.9521	-0.1397	0.7541	-0.0648	0.9006
Lcp65Aa	Lcp65Aa	1629340_s_at	0.0025	0.9915	0.3211	0.0397	0.2502	0.1729	0.1591	0.6844	0.0067	0.9809	-0.1524	0.3623	0.1149	0.8461	0.2134	0.3706	0.0985	0.7204
Or22c	Olfactory receptor	1629341_at	-0.0815	0.6572	-0.0542	0.6581	0.0766	0.6417	0.1648	0.7518	-0.0612	0.8318	-0.2259	0.2675	0.0113	0.9916	0.0478	0.9009	0.0366	0.9169
CG1746 /// DyakCG1746	CG1746	1629342_s_at	-0.5823	0.0500	-0.1608	0.2408	-0.3620	0.0971	-0.3735	0.2438	-0.5507	0.0108	-0.1772	0.3030	-0.1917	0.8270	-0.2762	0.4513	-0.0846	0.8602
CG4090	CG4090	1629343_at	0.0896	0.7189	0.2305	0.3005	0.1996	0.3850	0.1107	0.8402	0.2048	0.3241	0.0941	0.6574	0.2458	0.7726	0.2405	0.5289	-0.0053	0.9935
CG4730	CG4730	1629344_at	-0.1259	0.5865	-0.0023	0.9854	-0.3413	0.0968	-0.5751	0.1057	-0.1027	0.6687	0.4725	0.0205	-0.0702	0.9284	0.0408	0.9170	0.1110	0.6941
CG6540	CG6540	1629345_at	0.0615	0.8426	0.1242	0.5248	0.2236	0.2120	0.0217	0.9745	0.0689	0.7250	0.0472	0.8030	-0.2019	0.8318	0.0557	0.9291	0.2576	0.5373
CG1443	CG1443	1629346_at	-0.8661	0.4463	-2.6142	0.0074	-2.4057	0.0001	0.2111	0.7135	0.8065	0.0083	0.5954	0.0191	0.0334	0.9952	-0.7747	0.6049	-0.8081	0.5892
klu	klumphuss	1629347_at	0.0914	0.7795	-0.0866	0.4843	-0.3764	0.4163	-0.2553	0.8073	0.0155	0.9804	0.2708	0.4980	-0.2251	0.8297	-0.1383	0.8056	0.0867	0.8857
ec	echinus	1629348_at	0.1051	0.5074	-0.0190	0.9022	-0.0371	0.8647	-0.0314	0.9529	0.0805	0.6201	0.1118	0.4145	-0.0571	0.9246	0.0079	0.9841	0.0650	0.7884
CG12682	CG12682	1629349_at	0.1982	0.2085	0.0250	0.8223	0.0310	0.8929	0.0616	0.9218	0.2552	0.1758	0.1936	0.2561	-0.1453	0.8465	-0.1184	0.7480	0.0269	0.9512
---	---	1629350_at	0.1990	0.3221	0.0601	0.6019	-0.0060	0.9886	0.0145	0.9937	-0.2474	0.5503	-0.2619	0.4748	0.0451	0.9467	-0.0899	0.7007	-0.1350	0.5361
---	---	1629351_at	-0.0130	0.9601	-0.0845	0.4808	-0.0667	0.6948	0.0711	0.8877	0.1050	0.5671	0.0339	0.8639	-0.0862	0.8972	-0.1066	0.7043	-0.0204	0.9535
sgl	UDP-glucose deh	1629352_at	0.8869	0.0511	1.6467	0.0151	1.5955	0.0002	0.2395	0.4270	0.0989	0.5839	-0.1406	0.3579	0.2122	0.9229	0.8768	0.2119	0.6646	0.3719
CG18210	CG18210	1629353_at	0.0072	0.9732	0.0811	0.6659	0.1684	0.4346	0.0140	0.9900	-0.0786	0.7988	-0.0926	0.7307	0.0482	0.9555	0.0406	0.9170	-0.0076	0.9849
---	---	1629354_at	0.2849	0.2596	0.2281	0.2012	0.0981	0.6184	-0.1219	0.7028	0.0890	0.5753	0.2109	0.1072	0.1643	0.8465	0.0388	0.9460	-0.1255	0.7577
CG17374	CG17374	1629355_at	0.0732	0.9729	-3.1686	0.4514	0.7807	0.4889	1.6035	0.5510	1.4325	0.3073	-0.1710	0.9201	-2.5250	0.8004	-2.0673	0.6474	0.4577	0.9372
Rpl1	RNA polymerase	1629356_at	0.3873	0.1419	0.0726	0.9048	0.3064	0.2376	0.3049	0.6424	0.8194	0.0187	0.5145	0.0739	0.0366	0.9898	0.4258	0.4870	0.3892	0.5379
SNF1A	G protein-coupled	1629357_s_at	-0.7712	0.1267	-0.4284	0.5980	-0.4298	0.2415	-0.1200	0.7225	-0.5638	0.0037	-0.4438	0.0064	-0.0424	0.9929	-0.2377	0.8852	-0.1953	0.9020
Tsp29Fb	Tetraspanin 29Fb	1629358_s_at	-0.2032	0.2091	-0.1407	0.4140	-0.0977	0.5388	0.0334	0.9586	-0.2289	0.1750	-0.2622	0.0847	-0.0002	0.9999	-0.1257	0.6918	-0.1254	0.6877
YL-1	YL-1	1629359_at	-0.0407	0.8352	0.1898	0.4522	0.2466	0.1597	-0.0453	0.9426	-0.0285	0.9089	0.0168	0.9397	-0.0467	0.9717	0.2420	0.4631	0.2888	0.3868
jdp	jdp	1629360_at	0.1903	0.2693	0.1492	0.4214	0.1392	0.4181	-0.0031	0.9956	0.1411	0.4517	0.1441	0.3891	0.0419	0.9653	0.0732	0.8307	0.0313	0.9313
CG14809	CG14809	1629361_at	0.0094	0.9606	0.1922	0.1868	-0.0038	0.9900	-0.1825	0.7425	-0.1081	0.7014	0.0744	0.7857	0.0598	0.9447	0.1814	0.4976	0.1216	0.6679
mthl8	Mth-like 8	1629362_at	-0.2875	0.2570	-0.4336	0.2103	-0.4830	0.1288	0.1861	0.8087	0.3251	0.2929	0.1390	0.6615	0.0952	0.9238	-0.0309	0.9533	-0.1261	0.7411
Acyp2	Acylphosphatase	1629363_at	-0.2214	0.3063	0.5856	0.0842	0.4040	0.0595	-0.1646	0.7230	-0.7930	0.0033	-0.6284	0.0056	0.0549	0.9693	0.0552	0.9248	0.0004	0.9995
---	---	1629364_at	0.2206	0.2782	0.0360	0.8120	0.2301	0.1869	0.1508	0.7031	0.1561	0.4046	0.0052	0.9816	-0.2653	0.6584	-0.1665	0.5100	0.0988	0.7223
CG30324	CG30324	1629365_at	-0.0130	0.9578	-0.0124	0.9167	0.2766	0.1299	-0.0929	0.8449	0.0039	0.9884	0.0968	0.5792	-0.0842	0.9205	0.1196	0.7106	0.2037	0.4886
CG6216	CG6216	1629366_at	0.1651	0.4183	0.0198	0.8467	0.1870	0.2647	-0.0531	0.9222	-0.0647	0.7380	-0.0116	0.9561	-0.2358	0.7154	-0.1472	0.6106	0.0885	0.7824
CG15534	CG15534	1629367_at	0.0116	0.9495	0.1120	0.5799	0.1334	0.5535	-0.0641	0.9011	-0.3199	0.0594	-0.2558	0.0869	-0.3252	0.7506	-0.1839	0.7120	0.1413	0.7869
slou	slouch	1629368_at	-0.1311	0.5802	0.0002	1.0000	0.1344	0.3844	0.1774	0.7576	0.0164	0.9656	-0.1610	0.4973	0.0545	0.9449	0.0890	0.7541	0.0345	0.9169
CG7457 /// DmirCG7457	CG7457	1629369_at	0.0686	0.8635	0.1119	0.5184	0.3574	0.1973	0.1544	0.8971	0.1640	0.7202	0.0096	0.9850	-0.1179	0.8903	0.3328	0.2835	0.4506	0.1849
---	---	1629370_at	0.0326	0.8614	-0.0710	0.4827	0.0144	0.9332	-0.1091	0.7507	0.1758	0.2386	0.2848	0.0391	0.0014	0.9994	0.0979	0.7787	0.0966	0.7725
Elo68beta	CG11801	1629371_at	0.0130	0.9500	0.0040	0.9730	0.0422	0.8568	0.0143	0.9857	0.0963	0.6294	0.0820	0.6587	0.0282	0.9725	-0.0345	0.9138	-0.0627	0.8029
sec71	sec71	1629372_a_at	0.1358	0.4494	0.3025	0.1285	0.2980	0.1393	0.0457	0.9558	0.2415	0.2801	0.1957	0.3357	0.0464	0.9611	0.3318	0.1975	0.2854	0.2964
CG5199	CG5199	1629373_at	-0.3454	0.2482	-0.0716	0.5325	-0.0069	0.9881	0.1605	0.8247	-0.2170	0.4576	-0.3775	0.1306	-0.1821	0.8395	-0.0808	0.8791	0.1013	0.8281
CG31739	CG31739	1629374_at	-0.0907	0.7649	-0.2321	0.4437	-0.4272	0.0599	0.0180	0.9852	0.3796	0.0822	0.3616	0.0661	0.2291	0.8434	0.2339	0.6512	0.0048	0.9953
CG11400	CG11400	1629375_at	1.3896	0.0025	1.1019	0.0793	0.6455	0.2178	-0.4298	0.7278	-0.6837	0.2152	-0.2539	0.6578	-0.0386	0.9898	-0.9545	0.1349	-0.9159	0.1762

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629376_at	-0.0629	0.8067	-0.0123	0.9437	0.0358	0.8418	-0.0599	0.9116	-0.0061	0.9809	0.0538	0.7758	-0.1675	0.8012	-0.0857	0.8117	0.0818	0.8112
CG32263	CG32263	1629377_at	0.0551	0.7861	0.0361	0.7267	0.1270	0.5412	-0.0115	0.9922	0.1094	0.6956	0.1208	0.6253	-0.0228	0.9816	-0.0015	0.9988	0.0213	0.9431
CG10752	CG10752	1629378_at	0.1093	0.5907	0.0159	0.9388	-0.0348	0.8617	-0.0768	0.8942	0.0267	0.9214	0.1035	0.5912	-0.0265	0.9816	0.0016	0.9989	0.0280	0.9387
---	---	1629379_at	-0.1162	0.6215	-0.2188	0.1718	0.0392	0.8302	0.3522	0.2438	0.2424	0.1740	-0.1098	0.5295	-0.0636	0.9400	0.0667	0.8511	0.1304	0.6411
Cp16	Chorion protein 1(	1629380_at	-0.0608	0.7693	0.1024	0.6586	0.1565	0.4429	-0.0135	0.9857	-0.1077	0.5663	-0.0942	0.5882	-0.0015	0.9994	-0.1813	0.5523	-0.1798	0.5612
---	---	1629381_at	0.1384	0.5530	0.1668	0.3250	0.2525	0.2226	-0.0320	0.9603	-0.1306	0.4592	-0.0987	0.5538	-0.1582	0.8270	-0.0594	0.8933	0.0989	0.7793
fal	falten	1629382_at	0.4809	0.0416	-0.1253	0.3286	-0.1714	0.5345	-0.1061	0.9029	0.5190	0.0684	0.6250	0.0213	-0.1733	0.8270	-0.1248	0.7518	0.0485	0.9164
CG6416	CG6416	1629383_a_at	-2.4618	0.0039	-1.6344	0.0924	-2.6598	0.0020	-0.4666	0.5531	-1.5606	0.0031	-1.0941	0.0092	0.5227	0.8677	-0.8260	0.5208	-1.3486	0.2923
Brd8	Brd8	1629384_at	0.0430	0.9369	-0.2893	0.6437	0.2077	0.5035	0.3581	0.2500	0.5136	0.0130	0.1555	0.3591	-0.3703	0.8714	0.0621	0.9666	0.4323	0.6607
---	---	1629385_s_at	-2.7238	0.0005	-2.4842	0.0202	-2.6861	0.0000	-0.3078	0.5735	-1.3481	0.0012	-1.0403	0.0021	0.2067	0.8906	-0.6661	0.2226	-0.8728	0.1498
---	---	1629386_at	0.0559	0.7751	-0.0295	0.7710	0.1756	0.2785	0.1041	0.8449	0.0310	0.9096	-0.0731	0.7273	-0.0344	0.9689	-0.0923	0.7305	-0.0580	0.8444
aru	arouser	1629387_s_at	-0.9485	0.0237	-0.8573	0.0886	-0.9281	0.0030	0.0587	0.9436	-0.0826	0.7724	-0.1413	0.5432	-0.0558	0.9816	-0.0857	0.9198	-0.0300	0.9717
Arp5	CG7950	1629388_s_at	0.3392	0.1809	0.2023	0.5563	0.2098	0.3236	0.0219	0.9803	0.4137	0.0576	0.3917	0.0462	0.0783	0.9610	0.3159	0.4881	0.2377	0.6210
CG32809	CG32809	1629389_a_at	-0.1869	0.4058	0.2236	0.2683	-0.0228	0.9148	-0.1550	0.8513	-0.1475	0.6664	0.0075	0.9849	0.2417	0.7541	0.2562	0.4341	0.0146	0.9777
---	---	1629390_at	0.1005	0.4706	0.2129	0.4001	0.1800	0.3725	0.0500	0.9436	-0.0505	0.8471	-0.1005	0.6253	0.0243	0.9831	0.1104	0.7177	0.0862	0.7862
lectin-46Ca	lectin-44Ca	1629391_at	0.0440	0.7880	0.3279	0.1088	0.4175	0.0505	0.0233	0.9777	-0.0803	0.7443	-0.1035	0.6250	0.0767	0.9309	0.1284	0.6889	0.0517	0.8949
---	---	1629392_at	-0.1341	0.4609	0.0397	0.7445	-0.0586	0.7235	0.0027	0.9956	-0.0128	0.9530	-0.0155	0.9307	0.0474	0.9588	0.1394	0.6100	0.0920	0.7548
CG34022	CG34022	1629393_s_at	0.0594	0.7942	-0.0874	0.6471	0.1628	0.4631	0.1944	0.6823	0.0965	0.7053	-0.0978	0.6705	-0.0755	0.9291	-0.0388	0.9286	0.0367	0.9230
Cpr64Ab	CG15007	1629394_at	0.2920	0.1614	0.0559	0.6790	0.0855	0.5935	0.1289	0.8058	-0.0244	0.9338	-0.1533	0.4345	0.0704	0.9250	-0.1727	0.5050	-0.2431	0.3467
CG9743	CG9743	1629395_at	-0.2951	0.3454	-0.0483	0.9010	0.2448	0.2398	0.2831	0.4786	-0.3574	0.0956	-0.6405	0.0057	-0.1179	0.9400	-0.1671	0.7784	-0.0491	0.9428
CG8486	CG8486	1629396_a_at	-0.2955	0.3306	1.7922	0.0177	0.7038	0.0065	0.0294	0.9798	-0.3201	0.2570	-0.3495	0.1638	0.8679	0.3587	1.7870	0.0158	0.9192	0.1039
---	---	1629397_at	-0.1285	0.4162	0.0283	0.8379	-0.0604	0.7727	0.0261	0.9649	-0.0119	0.9604	-0.0380	0.8350	0.0736	0.9405	0.1255	0.7199	0.0520	0.9023
CG10383	CG10383	1629398_at	-0.7726	0.0334	-0.1942	0.7409	-1.2129	0.0001	0.6073	0.0928	0.4713	0.0333	-0.1360	0.5058	1.6724	0.1628	1.1658	0.0761	-0.5066	0.4285
CG30282	CG30282	1629399_at	4.8768	0.0005	3.0414	0.0371	3.0480	0.0000	0.3354	0.3836	1.4977	0.0003	1.1623	0.0005	0.2560	0.9421	-0.0999	0.9548	-0.3558	0.7822
RpL18	Ribosomal protein	1629400_at	0.5543	0.0081	0.9659	0.0097	1.1380	0.0008	0.1712	0.7604	-0.1573	0.5452	-0.3285	0.1267	0.0107	0.9913	-0.0710	0.7784	-0.0816	0.7266
CG1814	CG1814	1629401_s_at	-0.9645	0.0025	-0.1050	0.6611	0.0590	0.8045	0.0048	0.9943	-0.9016	0.0004	-0.9064	0.0003	-0.2567	0.7215	-0.2090	0.5057	0.0476	0.9095
CG7024	CG7024	1629402_at	0.0463	0.8517	0.0958	0.4658	-0.1564	0.4490	-0.2502	0.4162	-0.0942	0.6178	0.1560	0.3157	0.0145	0.9913	-0.0678	0.8546	-0.0824	0.8007
CG31466	CG31466	1629403_at	0.0278	0.9104	-0.0645	0.5583	0.1801	0.4012	0.2791	0.5502	0.1887	0.4526	-0.0904	0.7258	-0.1196	0.8222	-0.0257	0.9425	0.0939	0.7060
---	---	1629404_at	0.0457	0.7573	0.0679	0.5459	0.0091	0.9700	0.0397	0.9567	0.0274	0.9213	-0.0123	0.9599	0.0272	0.9672	-0.0218	0.9387	-0.0490	0.8221
---	---	1629405_s_at	0.4453	0.3953	0.1800	0.6263	-0.6898	0.0234	-0.2620	0.6101	0.8886	0.0050	1.1506	0.0010	0.5991	0.7485	0.6330	0.4242	0.0339	0.9790
Slbp	Stem-loop binding	1629406_at	0.1832	0.3305	-0.2528	0.1765	-0.3490	0.0833	-0.0803	0.8794	0.3705	0.0461	0.4508	0.0131	-0.0629	0.9604	-0.1810	0.6408	-0.1180	0.7794
CG13113	CG13113	1629407_at	0.1353	0.6676	-0.1928	0.4133	-0.0663	0.8028	0.1726	0.7189	0.2862	0.1854	0.1136	0.6040	-0.1009	0.9296	-0.1707	0.6768	-0.0697	0.8910
CG30387	CG30387	1629408_s_at	-0.2807	0.3743	-0.6148	0.0808	-0.4695	0.0982	0.1446	0.8518	0.7961	0.0121	0.6516	0.0177	-0.0774	0.9689	0.3466	0.5228	0.4240	0.4287
---	---	1629409_s_at	-0.1587	0.5755	-0.3610	0.4406	-0.3928	0.0488	-0.3946	0.3837	-0.2111	0.4260	0.1836	0.4475	-0.4922	0.6955	-0.4877	0.3350	0.0045	0.9963
CG32572	CG32572	1629410_at	-0.1122	0.5576	-0.1294	0.3398	-0.1305	0.3949	-0.0656	0.8947	0.0459	0.8245	0.1115	0.4776	-0.0635	0.9142	-0.0340	0.9152	0.0295	0.9174
CG8444 /// DyakCG8444	CG8444	1629411_at	-1.2955	0.0155	-0.3673	0.1821	-0.2568	0.3223	-0.0652	0.9518	-1.1818	0.0027	-1.1166	0.0021	-0.2091	0.8603	-0.2435	0.6382	-0.0344	0.9609
CG40091	CG40091	1629412_at	0.1998	0.1951	0.0698	0.5998	0.1842	0.2693	0.1323	0.6856	0.1978	0.1906	0.0655	0.6803	0.0596	0.9357	0.0319	0.9325	-0.0277	0.9330
CG12703	CG12703	1629413_at	0.4085	0.5029	0.8390	0.2215	1.2944	0.0007	0.2798	0.5014	-0.1276	0.5965	-0.4074	0.0450	-0.2329	0.9411	0.3149	0.8000	0.5479	0.6026
---	---	1629414_at	0.1053	0.5611	-0.0222	0.8651	0.2541	0.2581	0.1721	0.6829	0.1346	0.5180	-0.0375	0.8713	-0.0819	0.9340	0.0199	0.9696	0.1018	0.7872
CG40441	CG40441	1629415_x_at	0.0280	0.8827	0.1925	0.2025	0.1501	0.3816	-0.0711	0.8894	-0.1219	0.4989	-0.0508	0.7911	0.0418	0.9611	0.1363	0.5897	0.0945	0.7287
---	---	1629416_s_at	0.0987	0.5392	-0.1664	0.4439	-0.0377	0.8642	-0.0978	0.8449	0.0403	0.8689	0.1381	0.4244	-0.1167	0.8903	-0.0068	0.9925	0.1099	0.7656
CG14853	CG14853	1629417_s_at	-0.1255	0.3690	-0.1226	0.4205	0.0005	0.9982	0.0582	0.9039	0.0613	0.7401	0.0031	0.9874	-0.1489	0.8192	-0.0206	0.9628	0.1283	0.6574
sei	seizure potassium	1629418_s_at	-1.5089	0.0105	-2.8831	0.0019	-2.5515	0.0000	-0.0272	0.9838	1.0157	0.0064	1.0429	0.0034	-0.2382	0.8446	-0.2586	0.6314	-0.0204	0.9807
---	---	1629419_at	0.1122	0.4501	0.2276	0.3640	0.4624	0.0462	0.0984	0.8490	-0.0957	0.6525	-0.1941	0.2522	0.0163	0.9831	0.0687	0.7481	0.0524	0.8173
CG12433	CG12433	1629420_at	0.1177	0.4043	0.0813	0.6020	-0.0979	0.6173	-0.0974	0.8578	0.0812	0.7211	0.1785	0.3149	0.1679	0.7387	0.2336	0.2566	0.0657	0.8003
CG16789	CG16789	1629421_at	-0.8437	0.0885	-0.1289	0.4879	-0.1265	0.7668	0.2449	0.8290	-0.4516	0.3041	-0.6965	0.0752	-0.0287	0.9762	0.0317	0.9322	0.0604	0.8374
---	---	1629422_at	0.0214	0.9229	0.1788	0.1836	0.1447	0.5045	0.1025	0.7764	-0.0016	0.9942	-0.1041	0.4675	0.0576	0.9514	0.1040	0.7549	0.0465	0.9046
ics	Ras suppressor-1	1629423_at	-0.3134	0.1587	0.3327	0.1292	0.3569	0.1221	0.1307	0.7205	-0.2156	0.1893	-0.3463	0.0270	0.0285	0.9877	0.2586	0.5523	0.2302	0.6056
CG32550	CG32550	1629424_at	0.4484	0.0530	0.2965	0.1124	0.3361	0.1293	0.0449	0.9466	0.1217	0.5504	0.0767	0.7007	0.2127	0.8049	0.1262	0.7741	-0.0866	0.8524
CG13157	CG13157	1629425_at	0.0910	0.6716	-0.1067	0.4465	0.4769	0.0400	0.1851	0.6327	0.0392	0.8765	-0.1459	0.4109	-0.2810	0.6955	-0.1858	0.5509	0.0952	0.7890
Trx-2	thioredoxin	1629426_s_at	-0.6719	0.0224	-0.3360	0.1046	-0.1992	0.4148	-0.0119	0.9863	-0.2568	0.1218	-0.2449	0.1000	-0.0281	0.9862	-0.0030	0.9981	0.0251	0.9640
CG5660	CG5660	1629427_at	0.2491	0.1581	-0.0393	0.7999	0.1550	0.5697	0.0814	0.8796	0.6620	0.0041	0.5806	0.0044	-0.1021	0.8882	0.1576	0.5847	0.2597	0.3543
CG7342	CG7342	1629428_at	-2.4170	0.0020	-3.3704	0.0014	-2.9289	0.0002	0.6255	0.5496	1.0891	0.0490	0.4637	0.3501	-0.0158	0.9914	-0.0586	0.9022	-0.0427	0.9211
CG31388	CG31388	1629429_at	-0.1537	0.6299	-0.5408	0.0705	-0.3248	0.0801	-0.0587	0.9375	0.1762	0.4342	0.2350	0.2271	-0.2438	0.8122	-0.2874	0.5143	-0.0435	0.9425
regucalcin	regucalcin	1629430_s_at	-2.3097	0.0016	-0.7051	0.0364	-0.8230	0.0364	-0.0010	0.9994	-1.5672	0.0026	-1.5662	0.0015	-0.1458	0.8486	-0.4653	0.1395	-0.3195	0.3271
CG18747	CG18747	1629431_at	-0.4150	0.4674	-0.2740	0.1181	-0.7153	0.0042	-0.2310	0.8640	-0.1067	0.8656	0.1243	0.8185	0.2060	0.7393	-0.1016	0.7424	-0.3076	0.2585
mRpL15	mitochondrial ribo	1629432_at	-0.0528	0.7329	0.0790	0.8117	-0.2808	0.1235	-0.0593	0.9186	-0.0087	0.9750	0.0506	0.8007	0.4009	0.6483	0.1448	0.7200	-0.2561	0.4825
CG12091 /// DyakCG12091	CG12091	1629433_at	-0.0485	0.7577	0.1193	0.8458	0.2446	0.2518	-0.0331	0.9610	-1.0187	0.0007	-0.9857	0.0004	-0.2824	0.8461	-0.8181	0.1599	-0.5357	0.3825
CG9380	CG9380	1629434_at	-0.2232	0.9494	-0.4523	0.2534	-2.0383	0.0011	-1.1863	0.7845	-1.1211	0.5609	0.0651	0.9770	0.2752	0.9800	-1.2834	0.6270	-1.5585	0.5476
CG7710	CG7710	1629435_at	0.1259	0.6858	0.2622	0.4254	0.1091	0.6821	-0.1321	0.8611	-0.3168	0.2428	-0.1847	0.4734	-0.1174	0.8904	-0.0986	0.8126	0.0189	0.9695
---	---	1629436_at	0.2186	0.1885	0.1646	0.3217	0.1184	0.4513	-0.0275	0.9745	-0.0044	0.9892	0.0231	0.9293	0.1689	0.7154	0.0721	0.7544	-0.0968	0.6419
---	---	1629437_at	0.0505	0.8043	-0.1099	0.4063	-0.0301	0.8786	0.1019	0.8074	0.1225	0.4815	0.0206	0.9193	0.0074	0.9929	-0.0569	0.8283	-0.0643	0.7849
---	---	1629438_at	0.6589	0.0296	0.5288	0.2131	0.0972	0.6935	-0.1069	0.8701	0.4477	0.0552	0.5545	0.0147	0.2116	0.8692	0.1895	0.7572	-0.0221	0.9788
CG10431	CG10431	1629439_at	0.1129	0.7490	0.1090	0.6681	0.2292	0.2265	0.1075	0.8738	0.0351	0.9150	-0.0723	0.7830	0.0996	0.9449	0.2115	0.6593	0.1119	0.8411
mRpS2	mitochondrial ribo	1629440_at	0.2110	0.4597	0.7100	0.0542	0.5443	0.0231	0.1577	0.7556	-0.3731	0.0878	-0.5308	0.0144	0.2055	0.8616	0.0653	0.9320	-0.1402	0.8131
Fit2	Fermitin 2	1629441_at	-1.6322	0.0012	-1.8806	0.0045	-2.1159	0.0000	0.0495	0.9257	0.3478	0.0366	0.2983	0.0433	0.1667	0.8400	0.0545	0.9175	-0.1122	0.7804
egr	Eiger	1629442_at	-1.9059	0.0038	-0.8750	0.0574	-1.6196	0.0001	-0.3488	0.6557	-0.5578	0.1365	-0.2090	0.5769	-0.0157	0.9916	-0.1597	0.6819	-0.1441	0.7148
CG12655	CG12655	1629443_at	-2.8725	0.0041	-1.4376	0.0191	-1.9808	0.0018	-0.1648	0.9185	-0.9421	0.0610	-0.7773	0.0805	0.1041	0.9690	0.0938	0.9341	-0.0103	0.9933
CG11756	CG11756	1629444_at	0.0297	0.8810	0.0723	0.6288	0.1780	0.4512	0.1890	0.6954	0.1441	0.5479	-0.0449	0.8626	0.0816	0.9235	0.2917	0.2822	0.2100	0.4683

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
mtrm	matrimony	1629445_at	0.7991	0.4863	-2.4780	0.2620	-1.9041	0.0991	0.1721	0.7225	3.6282	0.0000	3.4561	0.0000	-0.3867	0.9741	0.4119	0.9287	0.7986	0.8271
CG1136	CG1136	1629446_at	0.1900	0.5757	-0.2503	0.5352	-0.1484	0.4848	0.4341	0.3328	0.5354	0.0447	0.1013	0.7129	0.3467	0.7743	0.0881	0.9132	-0.2586	0.6469
CG32829	CG32829	1629447_at	0.2488	0.3681	0.3809	0.1630	0.0856	0.5793	0.0142	0.9860	-0.1044	0.6244	-0.1186	0.5250	0.1876	0.8548	0.1892	0.6883	0.0015	0.9987
CG14358	CG14358	1629448_at	0.1946	0.3341	-0.3099	0.1887	0.0149	0.9506	0.3103	0.5680	0.2835	0.3122	-0.0268	0.9381	0.0298	0.9816	-0.1378	0.6605	-0.1676	0.5816
---	---	1629449_at	0.1540	0.4791	-0.0036	0.9839	0.0496	0.7744	0.0780	0.8606	0.1643	0.3077	0.0862	0.5888	-0.1808	0.7500	-0.1099	0.6847	0.0709	0.8142
CG14453	CG14453	1629450_at	0.0687	0.6913	0.0087	0.9361	-0.0021	0.9927	0.2273	0.4751	0.1935	0.2594	-0.0338	0.8655	0.1437	0.7961	-0.0384	0.9157	-0.1821	0.4507
---	---	1629451_at	-0.0510	0.8052	0.0619	0.5763	0.1483	0.4174	0.1388	0.6998	0.0155	0.9505	-0.1234	0.4255	0.0615	0.9309	0.0578	0.8546	-0.0037	0.9924
CG5862	CG5862	1629452_at	0.1517	0.4661	1.0353	0.0073	1.2844	0.0002	-0.0406	0.9445	-0.3934	0.0261	-0.3527	0.0261	-0.2856	0.7116	0.4846	0.1345	0.7702	0.0559
FucTA	alpha1,3-fucosyltr	1629453_at	-0.2840	0.2307	-0.9227	0.0108	-0.6711	0.0235	0.3373	0.5621	0.7889	0.0163	0.4516	0.0905	-0.1899	0.7661	-0.0507	0.9022	0.1392	0.6328
---	---	1629454_at	0.1069	0.5648	-0.0432	0.8967	0.0845	0.7185	-0.0785	0.8507	0.0443	0.8139	0.1228	0.3832	-0.0617	0.9717	-0.0661	0.9217	-0.0044	0.9953
CG8925	CG8925	1629455_at	0.0634	0.8689	2.0831	0.0226	1.5595	0.0095	-0.7906	0.4440	-2.4112	0.0019	-1.6206	0.0068	-0.2593	0.8825	-0.3358	0.6447	-0.0765	0.9350
---	---	1629456_at	0.4196	0.0361	0.3011	0.1386	0.2169	0.1290	-0.0627	0.8908	-0.1104	0.4929	-0.0477	0.7802	0.1064	0.9174	-0.1693	0.6655	-0.2757	0.4523
---	---	1629457_at	-0.0459	0.7891	-0.2866	0.2603	-0.7817	0.0456	-0.3627	0.5332	0.3535	0.2455	0.7162	0.0165	0.0129	0.9943	0.1029	0.8625	0.0900	0.8759
CG9411	CG9411	1629458_at	-0.6007	0.0191	-1.0947	0.0080	-1.2557	0.0026	0.1658	0.7831	0.7339	0.0101	0.5680	0.0189	0.0050	0.9964	0.0156	0.9733	0.0106	0.9819
Doc3	Dorsocross3	1629459_at	0.2264	0.3387	0.0515	0.6751	0.0491	0.8832	0.0652	0.9236	0.0337	0.9052	-0.0315	0.8985	-0.3601	0.7204	-0.2697	0.5342	0.0904	0.8718
CG32679	CG32679	1629460_at	0.2145	0.2662	0.0872	0.4452	0.4621	0.0543	0.0249	0.9629	0.0357	0.8464	0.0109	0.9522	-0.2648	0.6898	-0.0154	0.9737	0.2494	0.3723
CG15216	CG15216	1629461_at	0.1503	0.3683	0.0964	0.5811	0.1435	0.5139	0.0184	0.9812	-0.0447	0.8548	-0.0631	0.7589	0.0760	0.9291	0.0167	0.9712	-0.0593	0.8729
CG30192	CG30192	1629462_at	0.0404	0.8386	0.0299	0.8360	0.0854	0.5742	0.0099	0.9900	0.0008	0.9976	-0.0090	0.9679	0.0602	0.9101	0.0212	0.9451	-0.0390	0.8801
---	---	1629463_at	0.0818	0.6518	-0.1129	0.3503	-0.0099	0.9619	0.1038	0.8550	0.1587	0.4576	0.0550	0.8127	-0.1394	0.7628	-0.1699	0.3698	-0.0305	0.9088
CG32473	CG32473	1629464_a_at	-0.1256	0.9302	0.2211	0.5023	0.1845	0.3158	0.0384	0.9931	-0.6876	0.4517	-0.7260	0.3707	0.0999	0.9829	-0.3250	0.8129	-0.4249	0.7243
---	---	1629465_at	0.1322	0.5917	-0.1839	0.3446	-0.0641	0.8022	0.3228	0.2975	0.5602	0.0084	0.2374	0.1432	-0.0126	0.9898	-0.0310	0.9225	-0.0183	0.9491
cdc16	cdc16	1629466_at	0.7199	0.0312	0.3264	0.0748	0.6573	0.0042	-0.0949	0.9135	0.2934	0.2906	0.3883	0.1134	-0.4053	0.5667	-0.0253	0.9620	0.3800	0.2525
pan	CG32005	1629467_at	0.2130	0.2085	0.0315	0.7815	0.2778	0.1629	0.0438	0.9375	0.0588	0.7621	0.0150	0.9405	-0.0493	0.9619	-0.0527	0.9049	-0.0034	0.9946
CG9372	CG9372	1629468_at	-0.4943	0.2444	-0.8308	0.0945	-0.6787	0.0057	-0.1992	0.6617	-0.1331	0.5727	0.0661	0.7868	-0.3591	0.8076	-0.4050	0.5248	-0.0459	0.9590
CG10960 /// DyakCG10960	CG10960	1629469_s_at	1.5434	0.0051	0.5512	0.0759	0.9141	0.0063	0.0047	0.9956	1.1728	0.0017	1.1680	0.0010	-0.4165	0.7220	0.1961	0.7354	0.6125	0.2338
HP1c	HP1c	1629470_at	0.0932	0.7317	0.1717	0.3910	0.2035	0.3725	-0.0535	0.9451	0.0053	0.9879	0.0588	0.8141	-0.0464	0.9775	0.2692	0.5064	0.3156	0.4322
CG10803	CG10803	1629471_at	-0.4706	0.2692	-0.2727	0.4629	-0.5326	0.0178	-0.2407	0.5790	0.2036	0.3694	0.4443	0.0325	-0.0459	0.9862	0.3375	0.6146	0.3834	0.5634
CG9356	CG9356	1629472_s_at	-0.1718	0.3924	0.6045	0.0461	0.8383	0.0014	0.2438	0.4908	-0.5588	0.0099	-0.8026	0.0012	0.0753	0.9447	0.2719	0.4022	0.1967	0.5722
cdi	center divider	1629473_at	-0.5091	0.5243	0.2600	0.3745	-0.1206	0.6842	-0.4283	0.4405	-0.5405	0.0766	-0.1122	0.7294	-0.0043	0.9994	0.0425	0.9789	0.0468	0.9730
CG11368	CG11368	1629474_at	-0.1070	0.6907	-0.1770	0.3372	-0.1500	0.3765	-0.0257	0.9838	-0.1232	0.7193	-0.0975	0.7623	-0.0483	0.9687	-0.2337	0.4753	-0.1854	0.5889
nudC	nudC	1629475_at	0.4528	0.0796	0.3196	0.4567	0.1570	0.6079	-0.0376	0.9603	0.5139	0.0193	0.5515	0.0086	0.2968	0.8202	0.5290	0.3097	0.2322	0.6975
CG7542 /// DyakCG7542	CG7542	1629476_at	0.6746	0.2714	0.1687	0.1639	-0.0155	0.9393	-0.2026	0.6013	-0.0909	0.6837	0.1117	0.5633	0.1762	0.9400	-0.3849	0.6154	-0.5611	0.4457
---	---	1629477_at	-0.0105	0.9620	0.0931	0.4558	0.1807	0.2370	-0.0590	0.8950	-0.0433	0.8139	0.0157	0.9314	-0.2377	0.7230	-0.0011	0.9994	0.2366	0.4279
mRpS18A	mitochondrial ribo	1629478_at	0.0819	0.7592	0.3525	0.3437	0.0315	0.9161	-0.1847	0.6755	0.0372	0.8957	0.2219	0.2327	0.1903	0.8909	0.2861	0.6093	0.0957	0.8941
fok	fledgling of Klp38l	1629479_a_at	1.0349	0.0035	1.1461	0.0033	1.4508	0.0007	0.1460	0.8189	-0.8117	0.0065	-0.9577	0.0019	-0.1301	0.9061	-0.7828	0.0664	-0.6527	0.1300
---	---	1629480_at	0.1156	0.5336	0.2525	0.2124	0.4556	0.1354	0.1338	0.8164	0.0711	0.7982	-0.0627	0.8054	-0.1128	0.9342	0.3440	0.4328	0.4568	0.3089
---	---	1629481_at	0.2201	0.3086	0.0436	0.7918	-0.0482	0.7577	0.0116	0.9863	0.0139	0.9538	0.0023	0.9907	-0.0068	0.9952	-0.0774	0.8218	-0.0706	0.8307
CG40001	CG40001	1629482_at	-0.1252	0.4902	0.0151	0.9122	0.1539	0.2987	0.1303	0.6823	0.0363	0.8503	-0.0940	0.5119	-0.0015	0.9994	0.0355	0.9330	0.0370	0.9194
---	---	1629483_at	0.0846	0.6828	0.0712	0.6770	0.3324	0.0964	0.1104	0.8000	-0.0512	0.8139	-0.1616	0.3122	-0.1102	0.8317	0.0418	0.8979	0.1520	0.4978
CG31666 /// chinmo	CG31666 /// chror	1629484_s_at	-0.0119	0.9535	-0.1996	0.1176	-0.1081	0.5026	0.0658	0.8679	0.1920	0.1674	0.1262	0.3228	-0.0306	0.9748	0.0259	0.9476	0.0565	0.8577
gd	Gastrulation Defe	1629485_at	-0.1236	0.4950	-0.0698	0.7809	0.0943	0.5836	0.1577	0.7695	0.0291	0.9278	-0.1286	0.5695	-0.2153	0.7997	-0.0472	0.9339	0.1680	0.6607
Ccp84Ag	cuticle cluster 2	1629486_at	0.0780	0.8135	-0.2185	0.3433	-0.1238	0.4064	0.2353	0.6558	0.2149	0.4114	-0.0204	0.9495	0.0326	0.9741	-0.1267	0.6345	-0.1593	0.5403
Cdk9	Cyclin-dependent	1629487_at	0.1221	0.4720	0.0678	0.6572	0.3827	0.0264	0.0059	0.9937	-0.1046	0.4971	-0.1105	0.4202	-0.1383	0.8202	-0.1411	0.5972	-0.0028	0.9951
---	---	1629488_at	0.3912	0.1336	0.0323	0.7799	-0.1260	0.4808	0.0652	0.9413	0.2654	0.2988	0.2002	0.3948	0.0475	0.9679	-0.0528	0.9125	-0.1003	0.7884
CG5111	CG5111	1629489_at	0.0571	0.7171	-0.3080	0.0410	-0.2406	0.3635	0.1048	0.8794	0.3893	0.0998	0.2845	0.1777	-0.1484	0.7644	-0.0991	0.6698	0.0493	0.8577
---	---	1629490_at	0.0743	0.7086	-0.0406	0.7248	0.0529	0.7503	0.1174	0.8281	0.0318	0.9127	-0.0856	0.6958	0.0197	0.9812	-0.0481	0.8407	-0.0678	0.7418
CG13318	CG13318	1629491_at	-1.6104	0.0045	-2.2909	0.0654	-1.8140	0.0027	0.1960	0.8449	0.4410	0.2314	0.2450	0.4874	-0.2347	0.9455	-0.1151	0.9454	0.1196	0.9344
CG31626	CG31626	1629492_s_at	-0.0396	0.8621	-0.1580	0.5572	0.0892	0.7472	-0.0038	0.9961	0.0439	0.8937	0.0477	0.8656	-0.2906	0.7464	-0.2382	0.5518	0.0524	0.9205
U26	U26	1629493_at	0.3017	0.0666	0.1752	0.4830	0.1161	0.4614	-0.2750	0.3533	0.0689	0.7284	0.3438	0.0309	-0.1059	0.9092	0.0799	0.8619	0.1858	0.5995
---	---	1629494_s_at	-0.1628	0.3154	-0.3259	0.1196	-0.2277	0.1587	0.0207	0.9777	0.1788	0.3550	0.1581	0.3663	-0.1059	0.8202	-0.0524	0.8397	0.0535	0.8251
---	---	1629495_at	0.1214	0.4910	0.1533	0.2568	0.1779	0.3280	-0.0331	0.9580	-0.0561	0.7801	-0.0230	0.9086	0.0624	0.9056	0.0192	0.9499	-0.0432	0.8622
l(3)L1231	lethal (3) L1231	1629496_at	1.4386	0.0432	0.6830	0.0775	-0.6857	0.3276	-0.6191	0.6073	1.1789	0.0557	1.7980	0.0058	0.8208	0.7733	0.4430	0.7678	-0.3778	0.8016
CG11323	CG11323	1629497_at	0.4507	0.0831	0.0800	0.6106	-0.0538	0.8036	-0.0121	0.9894	0.2051	0.3373	0.2172	0.2502	0.1869	0.8156	-0.1017	0.8123	-0.2886	0.3921
Gyc32E	Guanylate cyclase	1629498_at	0.0630	0.7730	0.4840	0.0301	0.2275	0.1770	-0.1705	0.6763	-0.1764	0.3686	-0.0059	0.9802	0.3220	0.7220	0.3992	0.2871	0.0772	0.8844
CG11915	CG11915	1629499_at	-2.5946	0.0016	-1.7301	0.0523	-2.7098	0.0000	-0.7327	0.1033	-1.5790	0.0005	-0.8463	0.0043	0.2402	0.9238	-0.7834	0.3352	-1.0236	0.2352
CG31609	CG31609	1629500_at	0.1421	0.3686	0.1488	0.4391	0.1493	0.4663	-0.0323	0.9610	-0.0218	0.9298	0.0106	0.9621	0.0816	0.9309	0.1269	0.7180	0.0454	0.9155
CG15601	CG15601	1629501_at	0.2141	0.4165	0.3257	0.4310	0.1263	0.4118	-0.2494	0.5515	0.1767	0.4307	0.4261	0.0349	-0.2836	0.7917	0.1688	0.7556	0.4524	0.3311
CG31358	CG31358	1629502_at	0.1707	0.4150	0.0990	0.3309	0.3830	0.1117	-0.0436	0.9598	-0.1173	0.6451	-0.0737	0.7681	-0.0103	0.9914	0.0702	0.8052	0.0805	0.7526
Pitslre	PITSLRE	1629503_s_at	0.3532	0.0754	-0.1605	0.5103	0.0179	0.9453	-0.0583	0.9441	0.0399	0.9029	0.0982	0.6940	-0.1005	0.8156	-0.0431	0.8619	0.0574	0.7864
---	---	1629504_at	0.1176	0.4437	0.1300	0.4262	0.0771	0.7362	-0.1370	0.8339	-0.1130	0.6912	0.0240	0.9379	0.0669	0.8940	-0.0304	0.9170	-0.0972	0.6317
---	---	1629505_at	0.1328	0.4578	0.0366	0.7194	0.0414	0.8453	-0.0962	0.8810	-0.0283	0.9267	0.0679	0.7805	-0.1178	0.8222	-0.0609	0.8347	0.0570	0.8366
pip	pipe	1629506_at	0.3894	0.0710	0.0616	0.6809	-0.1452	0.6719	-0.0051	0.9956	0.0879	0.7194	0.0930	0.6709	0.1860	0.8400	-0.1393	0.7591	-0.3253	0.4004
mle	no action potentia	1629507_a_at	0.1427	0.7781	0.4632	0.2368	0.3107	0.3432	-0.3087	0.6972	-0.0918	0.8463	0.2170	0.5444	-0.1645	0.9289	0.2595	0.7000	0.4240	0.4913
---	---	1629508_at	0.0909	0.6987	0.0667	0.8270	0.1824	0.2347	0.0417	0.9603	-0.0617	0.8241	-0.1034	0.6517	-0.0591	0.9611	-0.0344	0.9501	0.0248	0.9598
car	carnation	1629509_s_at	-0.7710	0.2150	-0.2078	0.6580	0.3082	0.0902	0.2548	0.5008	-0.5916	0.0100	-0.8465	0.0012	-0.2720	0.9246	-0.0467	0.9784	0.2253	0.8577
CG4294	CG4294	1629510_at	0.2073	0.4910	0.4181	0.1210	-0.0344	0.9080	-0.2915	0.5109	0.1537	0.5413	0.4451	0.0400	0.1048	0.9276	0.2504	0.5214	0.1456	0.7377
CG5326	CG5326	1629511_at	0.6858	0.3367	-1.6763	0.0025	-0.0166	0.9453	1.4653	0.0031	1.4471	0.0003	-0.0182	0.9458	-0.1098	0.9816	-0.6642	0.5246	-0.5544	0.6094
---	---	1629512_at	0.1504	0.4139	0.2878	0.0705	0.2967	0.2112	-0.0230	0.9838	-0.2758	0.2927	-0.2527	0.2813	0.1254	0.8917	-0.0110	0.9876	-0.1365	0.7307
---	---	1629513_at	0.0638	0.8257	-0.0280	0.7887	0.0438	0.7814	-0.0554	0.9435	-0.0303	0.9220	0.0251	0.9260	-0.0512	0.9538	-0.1763	0.5141	-0.1251	0.6588

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629514_at	0.0715	0.6916	-0.1538	0.2753	0.2239	0.1924	0.2230	0.5238	0.2028	0.2708	-0.0203	0.9279	-0.1052	0.8901	-0.0464	0.9164	0.0588	0.8794
Pdk	pyruvate dehydro	1629515_at	0.3732	0.1102	-0.5389	0.1881	-0.5303	0.1293	0.1800	0.7952	0.4528	0.1084	0.2728	0.2865	0.0036	0.9994	-0.5897	0.3458	-0.5934	0.3683
---	---	1629516_at	-0.0169	0.9461	-0.2872	0.1288	-0.4431	0.1065	-0.1466	0.8162	0.4058	0.1011	0.5524	0.0208	-0.0654	0.9445	0.0960	0.7854	0.1614	0.5964
CG5118	CG5118	1629517_at	0.3962	0.3234	-0.8022	0.0077	-0.3192	0.6045	0.5657	0.5929	1.4180	0.0163	0.8523	0.0769	0.1200	0.9618	0.2903	0.7162	0.1703	0.8504
CG2493	CG2493	1629518_at	-0.2821	0.3832	-1.0638	0.0776	-0.7611	0.0129	0.0717	0.8908	-0.0676	0.7437	-0.1393	0.3916	-0.2531	0.8885	-0.7899	0.2289	-0.5368	0.4441
CG11356	CG11356	1629519_at	0.1455	0.4185	0.1685	0.3015	0.2865	0.1212	0.0409	0.9403	-0.0254	0.9077	-0.0663	0.6882	-0.1526	0.8283	-0.0477	0.9170	0.1048	0.7578
CG8997 /// DsimCG8997	CG8997	1629520_at	-0.2247	0.7857	-0.2880	0.5109	-0.4055	0.2269	0.0194	0.9937	0.1152	0.8586	0.0958	0.8686	0.1805	0.9530	-0.0307	0.9869	-0.2112	0.8601
---	---	1629521_at	0.0046	0.9795	0.0052	0.9712	-0.1639	0.3014	0.0798	0.8839	0.2048	0.2738	0.1250	0.4818	-0.0369	0.9611	0.0418	0.8946	0.0787	0.7484
Osi3	Osiris	1629522_at	0.2391	0.2593	0.3349	0.1329	0.2298	0.3183	0.0952	0.9130	-0.2163	0.4494	-0.3115	0.2049	0.0948	0.9128	0.0275	0.9557	-0.0673	0.8677
lola	longitudinals abse	1629523_at	-0.5091	0.2648	-0.6186	0.4517	-0.2629	0.3089	0.2463	0.7093	0.3974	0.1842	0.1512	0.6217	-0.0882	0.9831	0.4176	0.7054	0.5058	0.6311
---	---	1629524_at	0.0652	0.7258	0.1834	0.2141	0.1326	0.3983	-0.1320	0.6994	-0.1900	0.2252	-0.0581	0.7295	-0.0098	0.9901	0.0336	0.8966	0.0434	0.8444
CG30152 /// DafsCG30152	CG30152 /// CG1(	1629525_at	-0.3657	0.1569	0.1801	0.1708	0.1805	0.5279	-0.1722	0.6678	-0.6974	0.0039	-0.5253	0.0084	-0.1859	0.8655	-0.1081	0.8584	0.0779	0.8972
---	---	1629526_at	0.1425	0.4220	0.0228	0.8369	0.1834	0.2896	0.0594	0.9185	-0.0110	0.9685	-0.0704	0.7095	-0.0295	0.9717	0.0478	0.8687	0.0774	0.7442
CG40159	CG40159	1629527_a_at	0.0059	0.9870	0.0914	0.4120	-0.0246	0.9005	-0.1455	0.7558	-0.0671	0.7845	0.0785	0.7144	-0.0166	0.9898	0.0808	0.8168	0.0974	0.7526
---	---	1629528_a_at	0.0370	0.8033	0.0919	0.5748	0.0267	0.8934	0.0710	0.8667	0.0630	0.7163	-0.0081	0.9660	0.1362	0.8202	0.1101	0.6890	-0.0260	0.9396
---	---	1629529_at	-0.2312	0.2467	0.1561	0.1927	0.0356	0.8562	0.0438	0.9471	-0.1146	0.5704	-0.1584	0.3567	0.0477	0.9515	0.1596	0.5129	0.1119	0.6612
IM23	Immune induced r	1629530_at	0.8427	0.6183	0.4764	0.7334	1.3180	0.0781	0.1165	0.9671	1.2907	0.0812	1.1742	0.0763	-0.5521	0.9411	1.1181	0.6601	1.6703	0.4856
CG6460	CG6460	1629531_at	0.2028	0.3378	0.1280	0.4882	0.0924	0.6935	0.0383	0.9540	0.0730	0.7317	0.0347	0.8713	0.1721	0.8609	0.0862	0.8792	-0.0860	0.8709
CG18157	CG18157	1629532_at	0.1475	0.5156	0.0814	0.6416	0.1084	0.6334	-0.0921	0.8757	-0.0149	0.9611	0.0772	0.7219	-0.0387	0.9742	-0.0198	0.9682	0.0189	0.9665
---	---	1629533_at	0.0110	0.9752	0.1125	0.4672	0.1008	0.5317	0.0282	0.9744	-0.0296	0.9214	-0.0578	0.8093	-0.0527	0.9499	-0.1748	0.4958	-0.1221	0.6498
CG18568	CG18568	1629534_at	-0.1158	0.6015	-0.0347	0.7560	-0.0001	0.9998	-0.0170	0.9777	0.0879	0.5979	0.1049	0.4694	-0.0241	0.9829	0.1211	0.6713	0.1452	0.6009
CG14861	CG14861	1629535_at	0.1532	0.2947	-0.0582	0.8048	0.1073	0.6248	0.2558	0.3907	0.3808	0.0308	0.1250	0.4274	-0.0625	0.9555	-0.0619	0.9003	0.0005	0.9992
CG14079	CG14079	1629536_at	-0.0560	0.8377	0.2236	0.1471	0.2693	0.1929	0.1507	0.6537	-0.0528	0.7899	-0.2035	0.1592	-0.0503	0.9710	0.1615	0.6762	0.2118	0.5658
CG30394	CG30394	1629537_s_at	0.2285	0.3113	0.7400	0.1889	0.6691	0.0060	-0.0162	0.9834	-0.0115	0.9672	0.0047	0.9835	0.0080	0.9974	0.4317	0.4452	0.4237	0.4683
---	---	1629538_s_at	-0.5030	0.6801	-0.8076	0.0097	-0.4958	0.0058	0.3215	0.4739	-0.7506	0.0080	-1.0722	0.0010	-0.0442	0.9939	-1.2054	0.3397	-1.1612	0.3812
CG14042 /// SP555	SP555 /// CG1404	1629539_s_at	0.2419	0.4139	0.1522	0.5692	0.1076	0.4875	-0.0955	0.8605	-0.0240	0.9299	0.0715	0.7327	0.0642	0.9589	0.0477	0.9341	-0.0165	0.9769
bw	Plum	1629540_a_at	0.0603	0.7668	0.0430	0.6822	-0.0175	0.9473	-0.0598	0.8915	-0.0642	0.7060	-0.0043	0.9812	0.0436	0.9589	-0.0298	0.9390	-0.0733	0.8007
CG13018	CG13018	1629541_at	0.2456	0.4494	0.1884	0.5004	0.0797	0.7619	-0.1700	0.7123	0.0009	0.9978	0.1709	0.3791	-0.1314	0.9467	-0.0809	0.9322	0.0505	0.9517
CG31821	CG31821	1629542_at	-1.4482	0.0018	-0.5100	0.0868	-2.5265	0.0002	-1.3650	0.0855	-0.4327	0.3598	0.9323	0.0317	0.3004	0.7485	0.0965	0.8619	-0.2039	0.6336
---	---	1629543_at	0.2564	0.2701	-0.4904	0.1546	-0.2833	0.2937	-0.0741	0.9380	0.3082	0.2715	0.3823	0.1236	-0.4931	0.6695	-0.3624	0.4376	0.1307	0.8237
---	---	1629544_at	0.1353	0.3339	-0.0381	0.7093	-0.2857	0.0653	-0.0732	0.8699	0.2365	0.1296	0.3096	0.0341	0.0867	0.8541	-0.0626	0.8000	-0.1493	0.4572
Yp1	yolk protein 1	1629545_at	0.5121	0.0514	0.0812	0.7373	0.4361	0.0560	0.2045	0.7031	0.3697	0.1336	0.1652	0.4839	-0.0714	0.9405	-0.0366	0.9393	0.0348	0.9331
CG7054	CG7054	1629546_at	0.0786	0.8520	0.7183	0.0336	0.9910	0.0004	0.2252	0.5138	-0.2245	0.2167	-0.4498	0.0136	-0.0643	0.9775	0.4896	0.3604	0.5540	0.3205
Rad51C	Rad51C	1629547_at	-0.1876	0.3036	0.0099	0.9416	0.0386	0.9026	-0.0548	0.9518	-0.1371	0.6239	-0.0823	0.7660	0.0225	0.9914	0.1910	0.7017	0.1685	0.7411
---	---	1629548_x_at	0.1008	0.6422	0.0240	0.8817	0.1536	0.3762	-0.0644	0.9185	0.0699	0.7634	0.1343	0.4639	0.0916	0.9142	0.1072	0.7656	0.0156	0.9730
CG40190 /// rl	MAP kinase /// CC	1629549_s_at	0.2580	0.4257	1.2454	0.0277	1.0867	0.0003	-0.3617	0.5008	-1.1505	0.0025	-0.7887	0.0081	-0.2275	0.8122	-0.1547	0.7383	0.0728	0.8932
CG14650	CG14650	1629550_at	0.4039	0.0893	0.8788	0.0607	0.8484	0.0031	0.0513	0.9603	-0.2112	0.4637	-0.2625	0.2943	0.0040	0.9984	0.2985	0.4564	0.2946	0.4753
CG12268	CG12268	1629551_s_at	-2.0491	0.0267	-3.8765	0.0100	-2.9160	0.0001	0.5160	0.6015	1.4894	0.0095	0.9734	0.0360	-0.3435	0.9095	-0.3165	0.8218	0.0271	0.9874
rdgBbeta	rdgBbeta	1629552_at	0.2735	0.2986	0.3691	0.0925	0.5950	0.0637	0.0077	0.9937	0.1677	0.4259	0.1600	0.3983	-0.1431	0.9246	0.2774	0.6108	0.4205	0.4172
CG14153	CG14153	1629553_at	0.2245	0.9049	-0.6054	0.1344	-1.8450	0.0001	-0.0778	0.9445	-1.3578	0.0020	-1.2800	0.0016	1.1151	0.8202	-2.1963	0.2593	-3.3114	0.1313
CG14928	CG14928	1629554_at	0.1305	0.6763	0.1861	0.3442	0.0414	0.8206	-0.0220	0.9803	-0.1427	0.5327	-0.1207	0.5701	0.0800	0.9101	0.0874	0.7780	0.0074	0.9852
---	---	1629555_at	-0.0631	0.8736	0.0150	0.9027	0.1721	0.3420	0.0333	0.9753	-0.2880	0.2728	-0.3214	0.1682	0.1167	0.8729	-0.0345	0.9404	-0.1512	0.6259
---	---	1629556_at	0.1147	0.5820	0.2515	0.1634	0.3344	0.0531	-0.1221	0.7294	-0.1323	0.4174	-0.0102	0.9592	0.0313	0.9793	0.1766	0.5404	0.1453	0.6260
---	---	1629557_at	0.2555	0.2482	0.2002	0.3862	0.4560	0.0386	0.1666	0.7350	0.1032	0.6792	-0.0634	0.7967	0.0276	0.9780	0.0203	0.9590	-0.0073	0.9849
CG15657	CG15657	1629558_at	0.1498	0.5105	-0.0499	0.6717	0.1664	0.3613	0.1272	0.8085	-0.0488	0.8583	-0.1760	0.3601	-0.0465	0.9619	-0.1845	0.5102	-0.1380	0.6371
Atet	ABC transporter e	1629559_s_at	1.1478	0.0031	1.8490	0.0058	0.9271	0.0003	-0.2630	0.6724	-0.1779	0.5773	0.0851	0.7973	0.7381	0.1628	0.5700	0.0568	-0.1681	0.5587
---	---	1629560_at	0.0700	0.7325	-0.0759	0.7456	0.1073	0.6113	0.0671	0.9375	0.1403	0.6081	0.0732	0.7931	-0.0669	0.9589	-0.0441	0.9402	0.0228	0.9665
---	---	1629561_at	-0.0406	0.8827	0.0788	0.5168	0.1528	0.3345	0.0178	0.9777	-0.1195	0.4720	-0.1373	0.3462	-0.0281	0.9816	-0.0506	0.8981	-0.0224	0.9514
---	---	1629562_at	0.1410	0.5657	0.1048	0.4519	0.1116	0.5504	0.0251	0.9705	0.0470	0.8318	0.0219	0.9178	0.0101	0.9940	0.0347	0.9451	0.0246	0.9562
CG17059	CG17059	1629563_at	-0.6300	0.0113	-0.4472	0.0870	-0.2965	0.0899	0.1475	0.7838	-0.2665	0.2289	-0.4139	0.0430	0.0090	0.9928	-0.1036	0.6952	-0.1126	0.6580
CG1504	CG1504	1629564_at	0.0247	0.9149	0.0997	0.3192	0.1429	0.4717	0.0804	0.8987	-0.0594	0.8171	-0.1398	0.4780	0.0702	0.9405	0.0594	0.8867	-0.0107	0.9832
CG30160 /// Tsp42Eb	CG30160 /// tetras	1629565_s_at	0.2968	0.8043	-0.8507	0.2408	-0.5102	0.0369	0.9422	0.3749	1.5141	0.0201	0.5720	0.2937	0.3428	0.9246	0.1680	0.9312	-0.1748	0.9173
CG8834	CG8834	1629566_at	-0.3964	0.0607	-0.0922	0.6099	-0.0774	0.6911	-0.0364	0.9518	-0.0828	0.6607	-0.0464	0.8043	-0.0528	0.9514	0.1026	0.7351	0.1554	0.5744
---	---	1629567_at	0.0714	0.7681	0.1027	0.7102	-0.0450	0.8420	-0.0813	0.8769	-0.0943	0.6415	-0.0130	0.9549	0.1873	0.8065	-0.0004	0.9999	-0.1877	0.5811
CG33276	CG33276	1629568_at	0.4954	0.0248	1.3934	0.0269	1.1608	0.0092	-0.0945	0.9040	-0.2128	0.4195	-0.1183	0.6536	0.1447	0.9401	0.5693	0.3259	0.4245	0.4889
Fbxl4	CG1839	1629569_at	0.5091	0.0506	0.5001	0.5431	0.3162	0.3686	-0.3117	0.4962	0.2247	0.3653	0.5364	0.0217	0.0378	0.9922	0.3414	0.7400	0.3036	0.7663
---	---	1629570_at	0.3410	0.2565	0.0413	0.7494	0.3385	0.1096	0.1966	0.7208	0.1083	0.7060	-0.0883	0.7417	0.1256	0.9101	0.0566	0.9278	-0.0690	0.9001
---	---	1629571_at	0.1553	0.4295	0.2154	0.2837	-0.0310	0.8850	-0.2194	0.4142	-0.0558	0.7540	0.1636	0.2208	0.0631	0.9662	0.0587	0.9263	-0.0044	0.9951
fat-spondin	fat-spondin	1629572_a_at	1.0174	0.0343	-0.0390	0.9672	0.7226	0.0114	0.1577	0.8663	0.2052	0.5648	0.0475	0.9050	-0.5250	0.7990	-0.8441	0.3132	-0.3191	0.7484
CG34360	CG12802	1629573_at	0.0381	0.8889	0.4399	0.2676	0.1631	0.2814	-0.0231	0.9774	-0.0337	0.9026	-0.0106	0.9669	0.3298	0.7464	0.4794	0.2471	0.1496	0.7647
Spn7	Serine protease ir	1629574_at	-0.0143	0.9483	-0.1013	0.5407	-0.1873	0.4884	-0.0427	0.9371	0.0687	0.7045	0.1114	0.4525	-0.0800	0.9523	-0.1010	0.8546	-0.0211	0.9730
---	---	1629575_at	0.2442	0.2604	0.1507	0.2748	0.1657	0.4008	0.0321	0.9767	0.0340	0.9289	0.0019	0.9958	0.0115	0.9914	-0.0054	0.9925	-0.0169	0.9622
Map60	Centrosome asso	1629576_at	-0.0517	0.8970	-0.2930	0.2977	-0.5287	0.0272	-0.0452	0.9666	0.5074	0.0756	0.5526	0.0360	0.0005	0.9999	0.0786	0.9157	0.0780	0.9057
CG5690	CG5690	1629577_at	0.1530	0.5339	0.2541	0.3158	0.3358	0.0306	0.0628	0.9255	0.0709	0.7717	0.0080	0.9748	0.1173	0.9174	0.3219	0.4089	0.2046	0.6294
Tango7	Transport and Gol	1629578_at	0.0510	0.8147	0.0113	0.9702	0.2123	0.2972	0.3563	0.3381	0.1283	0.5819	-0.2280	0.2355	0.1872	0.8122	0.0942	0.8256	-0.0930	0.8184
CG2478	CG2478	1629579_at	0.2234	0.3836	0.1705	0.4833	-0.0012	0.9954	-0.1215	0.7067	0.2206	0.1339	0.3421	0.0177	0.1038	0.9352	0.1526	0.7518	0.0488	0.9321
---	---	1629580_at	0.2179	0.2585	0.0371	0.9010	0.2053	0.1917	0.1971	0.7312	0.1141	0.6994	-0.0830	0.7689	0.0335	0.9840	-0.1321	0.7760	-0.1657	0.6941
CG12079	CG12079	1629581_at	-0.2048	0.6039	0.6097	0.0394	0.7769	0.0027	-0.0853	0.8815	-1.1543	0.0005	-1.0691	0.0004	-0.2332	0.8609	-0.3483	0.5329	-0.1151	0.8737
---	---	1629582_at	0.1543	0.4173	0.0910	0.5401	0.2433	0.2304	-0.0267	0.9729	-0.0982	0.6557	-0.0715	0.7330	-0.0045	0.9964	-0.0203	0.9590	-0.0158	0.9640

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629583_at	0.0686	0.6445	-0.0527	0.7927	0.0673	0.7616	0.1811	0.6429	0.1580	0.4217	-0.0231	0.9210	-0.0117	0.9922	-0.0368	0.9314	-0.0252	0.9466
CG11298	CG11298	1629584_at	0.1282	0.4464	0.1180	0.5514	0.1476	0.3476	-0.0658	0.8942	-0.0955	0.5940	-0.0297	0.8789	0.1414	0.8202	0.0564	0.8787	-0.0849	0.7810
---	---	1629585_at	-0.0679	0.6845	-0.6784	0.0127	-0.4115	0.2803	0.2049	0.7695	0.5115	0.0854	0.3066	0.2489	0.0470	0.9831	-0.0412	0.9617	-0.0882	0.9020
CG34126	CG34126	1629586_at	-0.0233	0.9708	0.1902	0.7194	0.3860	0.3094	0.4475	0.6354	-0.0432	0.9505	-0.4908	0.2374	0.1799	0.9530	0.2573	0.8317	0.0775	0.9517
CG34363	CG12904	1629587_at	-0.8131	0.0157	-0.0198	0.8564	-0.3048	0.1624	-0.4267	0.4110	-1.2677	0.0017	-0.8410	0.0063	0.0259	0.9816	-0.1386	0.5827	-0.1645	0.5050
CG14240	CG14240	1629588_at	0.0946	0.5879	0.1395	0.3697	0.3111	0.0556	0.1693	0.7243	0.0727	0.7812	-0.0966	0.6669	0.0099	0.9911	0.0432	0.8642	0.0332	0.8932
Gr43a	Gustatory recepto	1629589_at	-0.4434	0.0576	0.1484	0.3525	-0.1414	0.3967	-0.2502	0.5376	-0.5785	0.0142	-0.3283	0.0841	-0.0302	0.9743	0.0555	0.8678	0.0857	0.7527
Gr64c	Gustatory recepto	1629590_at	0.1058	0.5671	0.1162	0.4766	-0.0359	0.8883	-0.2288	0.6338	-0.1521	0.5492	0.0768	0.7679	-0.0379	0.9562	-0.0317	0.9180	0.0061	0.9849
kl-2	male fertility factor	1629591_at	0.0805	0.6142	-0.0703	0.6452	0.1158	0.5694	0.0493	0.9506	0.0944	0.7102	0.0450	0.8605	-0.0656	0.9333	-0.0011	0.9993	0.0645	0.8404
CG8931	CG8931	1629592_at	-0.1375	0.4699	-0.2332	0.4542	-0.3220	0.0511	-0.0877	0.8507	0.3259	0.0592	0.4136	0.0144	-0.1951	0.8270	0.1696	0.6839	0.3648	0.3424
CG10866	CG10866	1629593_at	0.1027	0.5698	0.5587	0.0470	0.8508	0.0121	-0.1175	0.8655	-0.7880	0.0069	-0.6705	0.0086	-0.1631	0.8378	-0.1684	0.6344	-0.0053	0.9924
---	---	1629594_s_at	0.1701	0.3094	0.0911	0.4710	0.1585	0.3153	0.1191	0.8008	0.3028	0.1099	0.1838	0.2846	-0.0100	0.9914	0.1394	0.5024	0.1494	0.4745
---	---	1629595_at	0.2527	0.1957	-0.0072	0.9542	0.0075	0.9689	0.0372	0.9567	0.0799	0.7076	0.0427	0.8410	0.1027	0.8629	-0.1461	0.5611	-0.2489	0.3089
Rh7	rhodopsin	1629596_s_at	0.2110	0.1938	0.2956	0.1654	0.2558	0.1159	0.0577	0.9201	-0.1050	0.5821	-0.1627	0.3079	0.0631	0.9225	0.0885	0.7199	0.0254	0.9330
Asph	Aspartyl beta-hydr	1629597_a_at	-2.2487	0.0025	-1.6085	0.0484	-2.0945	0.0003	-0.1174	0.7431	-0.2168	0.1663	-0.0994	0.5146	0.4196	0.8579	0.4737	0.6472	0.0541	0.9704
FK506-bp1	FK506-binding pro	1629598_at	0.7397	0.0170	0.0809	0.8533	0.4867	0.1176	0.3232	0.2870	0.9702	0.0007	0.6470	0.0024	-0.0232	0.9940	0.5169	0.4825	0.5401	0.4683
---	---	1629599_at	-0.0546	0.7431	0.2388	0.2481	-0.0343	0.9041	-0.3480	0.5311	-0.3071	0.2943	0.0409	0.9075	0.0040	0.9964	-0.0479	0.8845	-0.0518	0.8626
CG17660	CG17660	1629600_at	-0.2270	0.2311	0.1514	0.5322	0.6230	0.0774	0.0010	0.9991	-0.2635	0.2139	-0.2645	0.1630	-0.2828	0.7519	0.2139	0.6009	0.4967	0.2130
malpha	E(spl) region trans	1629601_at	-0.0173	0.9715	0.1018	0.5606	0.1027	0.6259	-0.0523	0.9455	-0.2685	0.2156	-0.2162	0.2704	0.0040	0.9976	0.1544	0.6275	0.1504	0.6366
CG14211	MKP-like	1629602_at	-0.3532	0.1188	-0.2791	0.1849	-0.4821	0.0085	0.0668	0.9154	0.1004	0.6505	0.0336	0.8859	-0.0396	0.9588	0.1311	0.5591	0.1708	0.4371
CG13127	CG13127	1629603_at	0.1199	0.5865	0.0127	0.9642	0.3415	0.0758	0.2263	0.7469	0.0788	0.8426	-0.1475	0.6390	-0.0297	0.9759	0.0619	0.8481	0.0916	0.7381
---	---	1629604_s_at	0.3688	0.2569	0.0059	0.9669	0.0147	0.9477	-0.0556	0.9436	0.0256	0.9354	0.0812	0.7348	-0.0333	0.9816	-0.2614	0.3998	-0.2280	0.4825
CG15056	CG15056	1629605_at	-0.0759	0.6327	0.1241	0.4371	0.2293	0.3472	-0.0853	0.9313	-0.2011	0.5151	-0.1158	0.7075	0.0319	0.9590	0.0864	0.6607	0.0544	0.8039
CG13890	CG13890	1629606_at	0.3394	0.1702	0.2421	0.6276	0.9388	0.0014	0.0300	0.9744	-0.7212	0.0071	-0.7512	0.0035	-0.5576	0.6898	-0.6102	0.2714	-0.0526	0.9479
CG6179	CG6179	1629607_at	-0.1990	0.5285	0.1590	0.4855	0.0633	0.7481	-0.0737	0.9108	-0.1580	0.4664	-0.0844	0.6999	0.1533	0.8609	0.2248	0.5433	0.0715	0.8831
Gyc88E	Guanylyl cyclase a	1629608_a_at	0.2140	0.2252	-0.1334	0.4209	-0.0333	0.8606	0.1264	0.8038	0.1926	0.3528	0.0662	0.7679	0.0008	0.9998	-0.1230	0.7567	-0.1238	0.7480
GRHR	hormone receptor	1629609_at	1.5973	0.0138	0.2426	0.1676	0.9595	0.0413	-0.2840	0.7631	-0.0350	0.9534	0.2490	0.5266	-0.6816	0.6824	-1.2141	0.0881	-0.5325	0.4533
Cyp28c1	Cyp28c1	1629610_at	0.0784	0.6063	0.0923	0.3458	0.0615	0.8244	-0.0755	0.9220	-0.1006	0.7077	-0.0252	0.9295	0.1528	0.7485	0.0005	0.9996	-0.1523	0.4683
---	---	1629611_at	0.1547	0.4044	0.0847	0.5673	0.0722	0.7704	0.1635	0.6041	0.0924	0.5939	-0.0711	0.6662	0.3234	0.5186	0.1593	0.5208	-0.1641	0.5099
---	---	1629612_at	0.0210	0.9297	0.0367	0.7060	0.1985	0.1706	-0.1313	0.7917	-0.2378	0.2407	-0.1065	0.6008	-0.0837	0.8888	-0.2015	0.3626	-0.1178	0.6280
omd	oocyte maintenan	1629613_at	0.1011	0.7292	0.0522	0.8518	0.0793	0.6867	0.0057	0.9941	0.1432	0.3781	0.1375	0.3457	0.0542	0.9717	0.1628	0.7080	0.1086	0.8209
CG31549	CG31549	1629614_at	0.3110	0.2125	0.4102	0.1507	0.4156	0.0318	-0.0253	0.9728	0.1035	0.6110	0.1288	0.4615	-0.0259	0.9870	0.1859	0.6385	0.2118	0.5862
aust	australin	1629615_at	0.2103	0.2802	0.1479	0.5015	0.0735	0.7252	-0.1999	0.4770	-0.0524	0.7729	0.1475	0.2791	0.0054	0.9963	-0.2165	0.3820	-0.2219	0.3895
---	---	1629616_at	0.0570	0.7894	0.0186	0.8543	0.0235	0.9268	-0.1216	0.8247	-0.1460	0.5176	-0.0243	0.9253	0.0481	0.9301	-0.0182	0.9499	-0.0663	0.7456
CG14893	CG14893	1629617_at	0.0346	0.8693	0.1406	0.4523	-0.0005	0.9977	-0.0898	0.8676	-0.0413	0.8703	0.0485	0.8225	-0.0734	0.8966	-0.0587	0.8349	0.0146	0.9622
CG12256	CG12256	1629618_at	0.3223	0.1885	0.5972	0.0569	0.1976	0.3067	-0.1960	0.6615	-0.0733	0.7776	0.1227	0.5631	0.1411	0.8423	0.0865	0.8185	-0.0545	0.8915
Aats-trp	Tryptophanyl-tRN	1629619_at	0.4049	0.3224	-0.1620	0.5329	-0.7357	0.0081	0.0426	0.9704	1.3046	0.0017	1.2619	0.0011	0.3955	0.7677	0.6460	0.2395	0.2505	0.6902
Hel25E	RNA helicase	1629620_s_at	0.4559	0.0234	0.4738	0.2059	-0.0352	0.8517	0.0733	0.8667	0.3477	0.0315	0.2743	0.0511	0.4647	0.6159	0.3833	0.3178	-0.0814	0.8791
Rab27	Rab27	1629621_a_at	0.2242	0.6142	0.0915	0.6736	-0.0911	0.6869	-0.5074	0.1968	-0.1462	0.5695	0.3612	0.0896	-0.1965	0.9076	-0.1098	0.9054	0.0867	0.9171
---	---	1629622_at	-0.0560	0.8138	-0.2028	0.1596	-0.1663	0.4724	-0.0383	0.9759	0.1210	0.7401	0.1592	0.6110	-0.1208	0.8467	0.0245	0.9531	0.1452	0.6036
CG15916	CG15916	1629623_at	-0.0071	0.9724	-0.0784	0.7623	-0.2722	0.2249	-0.1239	0.7924	-0.1849	0.3425	-0.0610	0.7728	0.0959	0.9426	-0.0530	0.9353	-0.1489	0.7527
---	---	1629624_at	0.0799	0.6041	0.0556	0.6431	0.1183	0.4632	0.0437	0.9387	0.1639	0.3318	0.1202	0.4427	-0.1583	0.8192	0.0746	0.8461	0.2328	0.4192
CG13211	CG13211	1629625_at	0.7037	0.0297	1.3371	0.0136	1.5782	0.0002	0.2611	0.6010	0.0257	0.9447	-0.2354	0.3017	-0.0591	0.9775	0.5719	0.2369	0.6310	0.2226
---	---	1629626_s_at	0.0820	0.7080	0.1875	0.4256	0.0516	0.7670	-0.1203	0.8337	-0.1088	0.6568	0.0114	0.9671	0.1269	0.9152	0.1801	0.7020	0.0532	0.9264
---	---	1629627_at	0.1038	0.5389	0.1174	0.4673	-0.0019	0.9945	-0.1752	0.6673	0.0596	0.8027	0.2348	0.1759	0.1989	0.7726	0.1946	0.5246	-0.0042	0.9935
---	---	1629628_at	0.0974	0.6257	0.1477	0.2349	0.3763	0.0280	0.0330	0.9608	-0.1418	0.4456	-0.1748	0.2794	-0.0639	0.9092	-0.0103	0.9765	0.0535	0.8307
Pdp	CG12151	1629629_at	-0.1190	0.4649	0.4650	0.0230	0.6255	0.0029	-0.1607	0.5556	-0.6828	0.0012	-0.5222	0.0022	-0.1920	0.7475	-0.0316	0.9388	0.1604	0.5488
---	---	1629630_at	-0.0016	0.9950	-0.0215	0.8384	0.0375	0.8377	0.0654	0.9223	0.2141	0.2938	0.1487	0.4337	0.0647	0.9333	0.1972	0.4242	0.1325	0.6189
CG10853	CG10853	1629631_at	0.0482	0.8247	-0.1649	0.1726	-0.1683	0.3293	0.0311	0.9704	0.1884	0.3899	0.1573	0.4337	-0.1331	0.7723	-0.0407	0.8857	0.0923	0.6647
CG8135	CG8135	1629632_at	-0.8398	0.0292	0.2366	0.5601	0.4629	0.0775	0.1044	0.9116	-0.5319	0.0768	-0.6363	0.0253	-0.0486	0.9831	0.5847	0.2518	0.6334	0.2499
CG15255	CG15255	1629633_at	-0.3597	0.9040	-0.4762	0.0836	-0.2396	0.1581	0.1757	0.9811	-1.3024	0.4874	-1.4781	0.3709	-0.1487	0.9872	-1.4543	0.5057	-1.3056	0.5612
CG1531	CG1531	1629634_at	0.0903	0.8342	1.0730	0.1228	1.1570	0.0015	-0.0931	0.9518	-0.4233	0.3286	-0.3303	0.4075	-0.3447	0.8192	0.3667	0.5750	0.7115	0.2666
CG2051	CG2051	1629635_at	-0.0599	0.7337	-0.0350	0.7585	0.0914	0.6179	0.0022	0.9962	0.0592	0.7354	0.0571	0.7187	-0.0480	0.9679	0.1801	0.5923	0.2281	0.4865
---	---	1629636_at	0.0481	0.8333	0.0555	0.6767	-0.0113	0.9569	-0.0712	0.9147	-0.0544	0.8391	0.0168	0.9494	0.0116	0.9898	0.0766	0.7274	0.0650	0.7710
CG13794	CG13794	1629637_at	1.0423	0.0629	0.7724	0.0778	0.4552	0.1676	-1.2359	0.1302	-0.4547	0.3638	0.7812	0.0764	-0.9897	0.3362	-0.8370	0.1280	0.1527	0.8237
Als	Acid Labile Subun	1629638_at	-0.0898	0.7618	-0.3350	0.0433	-0.5077	0.0965	-0.0196	0.9888	0.0148	0.9764	0.0344	0.9311	0.2185	0.7994	-0.0248	0.9664	-0.2433	0.5151
CG14369	CG14369	1629639_at	0.0957	0.6872	0.2673	0.3180	0.3516	0.0297	-0.1413	0.8738	-0.3660	0.2398	-0.2246	0.4429	-0.0552	0.9277	-0.1066	0.6166	-0.0514	0.8393
---	---	1629640_at	0.1096	0.6964	-0.5787	0.2829	-0.8007	0.0157	-0.0916	0.9380	1.1826	0.0052	1.2742	0.0022	0.0146	0.9958	0.3357	0.6259	0.3211	0.6398
---	---	1629641_s_at	-1.3148	0.0037	-1.3537	0.1518	-2.0068	0.0008	-0.3300	0.4913	0.1686	0.5380	0.4986	0.0360	0.1286	0.9734	-0.0036	0.9994	-0.1321	0.9197
CG17664	CG17664	1629642_a_at	0.5282	0.3654	0.5684	0.4533	0.9189	0.0235	-0.0261	0.9673	-0.2313	0.1651	-0.2052	0.1698	-0.6013	0.8331	-0.3984	0.7886	0.2030	0.9022
CG32186	CG32186	1629643_at	-0.0190	0.9374	0.1101	0.5399	-0.1329	0.5528	-0.1854	0.7042	0.0105	0.9761	0.1959	0.3411	0.0499	0.9611	0.1598	0.6035	0.1099	0.7417
Mtl	Mtl(Rac-like)	1629644_s_at	-1.0057	0.0046	0.1738	0.2916	0.0420	0.8556	-0.0431	0.9603	-0.8233	0.0045	-0.7802	0.0034	0.0090	0.9946	0.3613	0.2247	0.3523	0.2697
---	---	1629645_at	-0.0009	0.9964	0.0285	0.8930	-0.0306	0.9314	-0.1012	0.8402	0.0079	0.9785	0.1091	0.5535	-0.1750	0.8815	-0.0832	0.9037	0.0917	0.8811
Or22a	Olfactory receptor	1629646_at	0.0051	0.9858	0.0179	0.9543	0.0522	0.8221	0.0571	0.9375	0.3410	0.1050	0.2839	0.1306	-0.0340	0.9831	0.1187	0.8052	0.1527	0.7185
l(2)06225	lethal (2) 06225	1629647_at	-0.0897	0.7431	0.2160	0.6286	0.2047	0.3123	-0.0060	0.9956	-0.7148	0.0092	-0.7087	0.0057	-0.0257	0.9914	-0.3660	0.4926	-0.3403	0.5341
---	---	1629648_s_at	-0.7525	0.1072	-0.7339	0.3450	-0.7106	0.0116	0.0206	0.9770	0.0291	0.9044	0.0084	0.9692	0.2723	0.8882	0.2326	0.8018	-0.0397	0.9717
siz	schizo	1629649_at	0.3851	0.2650	0.3057	0.1463	-0.2142	0.3009	-0.1904	0.5744	-0.0908	0.6416	0.0996	0.5623	0.5358	0.3564	0.2598	0.3972	-0.2760	0.3838
Cog3	CG3248	1629650_at	-0.3846	0.3396	0.4011	0.3081	0.3504	0.0601	0.2524	0.5946	0.3033	0.2057	0.0509	0.8557	0.3725	0.7644	1.2621	0.0391	0.8896	0.1210
CG5386	CG5386	1629651_at	-0.1162	0.5260	0.1859	0.1265	0.1482	0.4141	0.0558	0.9260	-0.0456	0.8459	-0.1013	0.5724	0.0711	0.8940	0.1603	0.4211	0.0892	0.6875

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629652_at	0.1653	0.3736	0.0251	0.8051	0.5604	0.0155	0.1308	0.6940	0.0097	0.9677	-0.1211	0.3972	-0.2238	0.7478	-0.0221	0.9640	0.2017	0.5123
CG16782	CG16782	1629653_at	0.0691	0.7144	0.2614	0.2904	0.1681	0.3446	-0.1372	0.8281	-0.1449	0.5827	-0.0077	0.9799	0.2538	0.6749	0.3843	0.1297	0.1305	0.6311
---	---	1629654_at	-0.1432	0.5269	-0.0115	0.9515	-0.0898	0.7099	0.0281	0.9805	0.0397	0.9200	0.0115	0.9736	0.0041	0.9965	0.0798	0.7940	0.0757	0.7964
Aos1	Ubiquitin activatin	1629655_at	0.4362	0.0806	0.2482	0.4292	0.3995	0.0380	-0.0135	0.9808	0.3409	0.0207	0.3544	0.0108	-0.0728	0.9589	0.1690	0.7117	0.2417	0.5722
---	---	1629656_at	-0.2279	0.2445	0.0030	0.9919	-0.0422	0.8263	-0.2069	0.5756	-0.2490	0.1857	-0.0421	0.8478	0.0779	0.9296	0.2369	0.4031	0.1591	0.6051
l(1)19Ec	section 7	1629657_at	0.0803	0.7938	0.1603	0.2486	0.0049	0.9885	-0.1544	0.7904	-0.0810	0.7775	0.0734	0.7809	0.0499	0.9689	0.0445	0.9342	-0.0054	0.9924
CG33298	CG33298	1629658_at	-0.0953	0.8247	-0.1992	0.5472	-0.8067	0.0107	-0.1204	0.8943	0.4190	0.1634	0.5394	0.0506	0.5045	0.7196	0.3069	0.6259	-0.1977	0.7743
pAbp	doppio fuso	1629659_s_at	0.0437	0.9065	0.4171	0.1336	-0.1962	0.3659	-0.1152	0.8076	-0.0332	0.8974	0.0820	0.6776	0.4305	0.7196	0.2875	0.5855	-0.1431	0.8178
CG11379	CG11379	1629660_at	0.2748	0.2124	0.2271	0.3222	0.4535	0.0267	0.0538	0.9367	-0.0265	0.9231	-0.0803	0.6991	-0.0712	0.9441	-0.1025	0.7900	-0.0313	0.9428
CG13723	CG13723	1629661_at	0.1057	0.6916	0.0193	0.9204	0.1246	0.5854	0.0014	0.9988	-0.0887	0.7676	-0.0901	0.7379	-0.2333	0.7768	-0.2346	0.5248	-0.0013	0.9986
CG3527	CG3527	1629662_at	0.2828	0.2296	0.1214	0.5614	0.3896	0.0654	0.0741	0.8897	0.1246	0.5103	0.0504	0.8025	-0.2612	0.8270	0.0461	0.9541	0.3073	0.5612
CG1832	CG1832	1629663_at	0.3146	0.1151	0.2275	0.5032	-0.3338	0.1710	-0.0573	0.9182	0.3782	0.0342	0.4354	0.0120	0.5139	0.6557	0.3183	0.5114	-0.1956	0.7121
---	---	1629664_at	0.2409	0.2465	0.2536	0.1461	0.1374	0.4214	-0.1607	0.6844	0.0659	0.7648	0.2266	0.1666	0.1038	0.8825	0.1776	0.5244	0.0737	0.8264
---	---	1629665_s_at	0.0868	0.7560	0.2451	0.1268	0.1048	0.6427	-0.1912	0.6010	-0.2451	0.1807	-0.0538	0.7944	0.0091	0.9914	0.0262	0.9325	0.0170	0.9494
CG4865	CG4865	1629666_at	0.1928	0.4671	0.2598	0.2859	0.4746	0.1787	-0.4241	0.4815	-0.0255	0.9576	0.3986	0.1660	-0.3291	0.7230	0.0779	0.8964	0.4070	0.3205
CG14298 /// DereCG14298	CG14298 /// GA1	1629667_at	-3.6280	0.0008	-4.4358	0.0052	-4.2625	0.0000	0.1197	0.8738	0.2263	0.4025	0.1066	0.7005	-0.0461	0.9914	-0.6510	0.4802	-0.6048	0.5232
CG32586	CG32586	1629668_at	0.0176	0.9245	-0.0342	0.7880	0.0081	0.9747	0.0947	0.8076	-0.0177	0.9357	-0.1124	0.4408	-0.1246	0.8395	-0.0757	0.8185	0.0489	0.8888
---	---	1629669_x_at	-0.1201	0.5532	-0.2652	0.0915	-0.4619	0.0282	-0.1787	0.7031	0.0439	0.8788	0.2226	0.2504	-0.0493	0.9589	-0.2040	0.4520	-0.1547	0.5942
CG17376	CG17376	1629670_at	0.2181	0.4186	0.1248	0.4239	-0.0101	0.9729	0.0814	0.9300	-0.1623	0.5818	-0.2437	0.3251	0.1184	0.8692	-0.1262	0.6957	-0.2446	0.4017
CG14193	CG14193	1629671_at	-0.8198	0.0403	-2.8354	0.0048	-2.4332	0.0065	0.5901	0.7351	1.8123	0.0273	1.2223	0.0796	0.0305	0.9848	-0.3451	0.3281	-0.3756	0.3097
Obp93a	Odorant-binding p	1629672_at	-0.0200	0.9250	0.2352	0.1503	0.1450	0.5241	-0.2103	0.6338	-0.1796	0.4216	0.0307	0.9067	-0.1219	0.8461	-0.1351	0.6310	-0.0132	0.9743
CG15418	CG15418	1629673_at	0.7488	0.0083	0.3168	0.2138	0.3070	0.1665	-0.1424	0.8189	-0.0291	0.9321	0.1133	0.6519	-0.1733	0.8461	-0.4274	0.2263	-0.2541	0.5041
CG10999	CG10999	1629674_s_at	-1.3700	0.0098	-0.7063	0.0252	-0.4671	0.0662	0.5495	0.3193	0.3225	0.3222	-0.2271	0.4551	0.2308	0.8479	0.9855	0.0667	0.7547	0.1568
CG18814	CG18814	1629675_at	0.1513	0.3855	0.0202	0.8978	0.1549	0.3261	-0.0085	0.9940	0.0916	0.7372	0.1000	0.6797	-0.1284	0.8283	-0.0685	0.8365	0.0599	0.8514
Tbp-1	Proteasome 26S	1629676_at	0.1894	0.4356	-0.0307	0.8469	-0.1289	0.4807	-0.0634	0.9212	0.4496	0.0279	0.5130	0.0099	-0.0257	0.9846	0.2673	0.3571	0.2931	0.3343
Tim17a2	Tim17a2	1629677_at	-0.6373	0.0075	-0.2815	0.1651	-0.2029	0.4454	0.0640	0.9452	-0.2789	0.2983	-0.3429	0.1477	0.0043	0.9982	-0.0275	0.9657	-0.0318	0.9542
Aplip1	APP-like protein ir	1629678_a_at	-0.1895	0.2367	-0.0021	0.9912	0.0313	0.8960	-0.0716	0.8735	-0.1165	0.4760	-0.0449	0.7981	-0.0672	0.9113	0.0106	0.9782	0.0778	0.7526
---	---	1629679_at	0.0703	0.6397	-0.1417	0.2416	0.0581	0.8307	0.0344	0.9683	0.0713	0.7965	0.0369	0.8912	-0.1256	0.8427	-0.0801	0.8123	0.0455	0.8985
CG5984	Filamin-like	1629680_at	-0.6614	0.0096	0.0332	0.8215	-0.0774	0.6758	-0.0915	0.8034	-0.5914	0.0026	-0.4999	0.0032	0.0365	0.9545	0.0876	0.6754	0.0511	0.8295
CG4068	CG4068	1629681_x_at	-1.2095	0.2064	-1.0637	0.4918	-2.2078	0.0084	-0.1938	0.5760	0.2553	0.1492	0.4491	0.0124	0.9024	0.8692	0.4462	0.8868	-0.4562	0.8771
CG13542	CG13542	1629682_at	0.0911	0.7160	0.2459	0.1297	0.1834	0.2775	-0.0425	0.9603	-0.2288	0.3188	-0.1862	0.3743	0.1139	0.8706	0.0029	0.9967	-0.1110	0.7269
br	broad complex	1629683_at	0.0795	0.7196	-0.0175	0.8687	0.0193	0.9347	0.0717	0.8987	0.1626	0.3898	0.0910	0.6288	0.0831	0.8999	-0.0043	0.9935	-0.0875	0.7557
---	---	1629684_at	0.2963	0.2762	0.2308	0.2743	0.1534	0.3282	0.0611	0.9314	0.1246	0.5832	0.0635	0.7861	0.0244	0.9816	0.0718	0.8187	0.0474	0.8866
CG16986	CG16986	1629685_at	0.4035	0.0452	0.3683	0.1797	0.4562	0.0554	-0.2752	0.5140	-0.3578	0.1074	-0.0826	0.7294	-0.3197	0.7387	-0.4259	0.2799	-0.1062	0.8372
---	---	1629686_at	0.0903	0.6886	0.1068	0.4327	0.1207	0.4166	0.1013	0.8449	0.0371	0.8855	-0.0643	0.7581	-0.0119	0.9928	-0.0627	0.8857	-0.0508	0.9031
---	---	1629687_s_at	-0.0998	0.5619	0.0064	0.9568	0.1948	0.2496	0.1747	0.5757	-0.0581	0.7622	-0.2328	0.1032	-0.2289	0.6763	-0.1000	0.6985	0.1289	0.6001
Oseg4	Oseg4	1629688_at	-0.6703	0.0145	-0.8391	0.0283	-0.9105	0.0610	0.4353	0.6825	0.6868	0.1638	0.2515	0.6170	0.2031	0.7899	0.0005	0.9998	-0.2027	0.5571
CG14673	CG14673	1629689_at	-0.0058	0.9749	0.0648	0.5884	0.1439	0.4715	-0.1155	0.7488	-0.0257	0.9058	0.0898	0.5631	-0.1440	0.8049	-0.0165	0.9670	0.1275	0.6279
CG6791	CG6791	1629690_at	-0.7166	0.1049	0.5656	0.0527	0.6325	0.0410	-0.1868	0.7743	-1.0536	0.0031	-0.8668	0.0043	-0.2070	0.8846	0.3730	0.5027	0.5800	0.2964
CG5028	CG5028	1629691_at	-0.7953	0.0186	0.2342	0.4243	0.3403	0.1510	-0.1816	0.7931	-1.5817	0.0007	-1.4001	0.0006	-0.2549	0.8049	-0.4174	0.3185	-0.1625	0.7434
CG15653	CG15653	1629692_at	0.0184	0.9192	0.0320	0.9232	0.1135	0.6023	-0.0915	0.8498	-0.1758	0.3241	-0.0843	0.6402	0.0347	0.9816	0.0051	0.9942	-0.0296	0.9514
CG15080	CG15080	1629693_at	0.1840	0.3565	0.0681	0.7915	-0.2145	0.5523	-0.0296	0.9819	0.3783	0.2166	0.4079	0.1357	-0.0487	0.9677	0.1519	0.6658	0.2006	0.5516
Pabp2	polyA-binding prot	1629694_at	-0.1171	0.6686	0.0956	0.8384	0.0516	0.7577	-0.2808	0.5840	-0.0663	0.8446	0.2146	0.3741	-0.2059	0.8222	0.1329	0.7754	0.3387	0.3830
CG4686 /// DpseGA18356	CG4686 /// GA18	1629695_at	0.0463	0.7513	0.1353	0.3589	0.3553	0.0802	-0.0384	0.9518	-0.2816	0.1112	-0.2431	0.1235	-0.0988	0.8521	-0.0447	0.8886	0.0541	0.8477
CG30497	CG30497	1629696_a_at	-0.2886	0.4188	0.1165	0.6545	-0.4297	0.0195	-0.1102	0.8373	-0.0245	0.9309	0.0857	0.6851	0.4968	0.6749	0.3592	0.4757	-0.1376	0.8247
beat-VII	beat-VII	1629697_a_at	-2.2412	0.0014	-3.0911	0.0022	-2.4516	0.0002	0.6409	0.4566	0.8940	0.0593	0.2530	0.5872	-0.0224	0.9875	-0.0697	0.8807	-0.0473	0.9161
CG3515	CG3515	1629698_at	0.1224	0.5544	0.0645	0.5904	0.2973	0.1058	0.1819	0.7349	0.1794	0.4727	-0.0026	0.9930	-0.2141	0.7707	-0.2117	0.5143	0.0024	0.9965
CG15311	CG15311	1629699_at	0.0099	0.9678	0.1103	0.5871	0.0426	0.8941	0.0314	0.9759	-0.0746	0.8141	-0.1060	0.6940	0.0280	0.9862	-0.0913	0.8636	-0.1193	0.7953
---	---	1629700_at	0.4126	0.4004	-0.6128	0.0656	-1.6316	0.0253	-0.2790	0.7380	1.7460	0.0013	2.0249	0.0004	0.7713	0.7500	0.6258	0.5636	-0.1455	0.9189
CG15767	CG15767	1629701_at	-0.0160	0.9323	0.0670	0.6830	-0.1680	0.4527	-0.0282	0.9774	-0.0124	0.9743	0.0158	0.9592	0.1777	0.7810	0.0092	0.9870	-0.1686	0.5653
ab	clueless	1629702_a_at	-0.2873	0.7972	-0.0528	0.6025	-0.2739	0.1106	-0.0223	0.9956	-0.5399	0.5602	-0.5176	0.5376	0.2374	0.7758	-0.2208	0.5591	-0.4582	0.2199
npf	neuropeptide F	1629703_at	0.0223	0.9125	0.1307	0.4422	0.0937	0.6348	-0.2219	0.5109	-0.0835	0.6814	0.1384	0.4058	-0.0495	0.9449	0.0756	0.7760	0.1251	0.5881
---	---	1629704_at	-0.1334	0.4042	0.0180	0.8672	0.0210	0.9136	0.1207	0.7161	-0.0091	0.9690	-0.1298	0.3498	-0.0476	0.9505	-0.0034	0.9942	0.0442	0.8880
DyakCG6084	CG6084	1629705_at	-0.5413	0.3459	0.8419	0.0267	-0.2690	0.1303	0.1438	0.6886	0.1847	0.2692	0.0409	0.8293	1.3738	0.3712	1.6551	0.0664	0.2813	0.7754
Obp50b	Odorant-binding p	1629706_at	-0.2917	0.2461	0.0717	0.6028	-0.0785	0.6519	-0.1045	0.8939	-0.1784	0.5189	-0.0739	0.8007	0.0881	0.8465	0.1456	0.4393	0.0576	0.8022
---	---	1629707_at	0.1558	0.3491	0.0768	0.5874	0.2770	0.1385	0.1313	0.7351	-0.0142	0.9557	-0.1455	0.3595	-0.0568	0.9503	-0.1263	0.6747	-0.0695	0.8409
CG8219	CG8219	1629708_at	0.3936	0.0217	-0.0973	0.4622	-0.0586	0.8143	0.0627	0.9196	0.3490	0.0697	0.2863	0.0925	0.0146	0.9914	-0.0099	0.9873	-0.0245	0.9549
Meics	zinc-finger-motif-p	1629709_at	-0.0369	0.9298	0.3392	0.4056	-0.0020	0.9945	-0.1375	0.8660	-0.0236	0.9557	0.1139	0.7085	0.2512	0.8384	0.4048	0.4242	0.1537	0.8041
Lip1	Lipase 1	1629710_at	0.0904	0.5640	0.0962	0.4490	0.2974	0.1622	0.0338	0.9558	-0.0158	0.9492	-0.0497	0.7878	0.1055	0.8379	0.1687	0.4266	0.0632	0.8098
CG10597	CG10597	1629711_at	-0.0741	0.6392	0.1389	0.3450	0.1819	0.3594	0.1121	0.8233	-0.1013	0.6437	-0.2135	0.2214	0.0609	0.9421	0.1180	0.6839	0.0571	0.8692
---	---	1629712_at	0.2228	0.1830	-0.1138	0.5341	-0.0353	0.8715	0.1753	0.6096	0.4090	0.0256	0.2337	0.1274	-0.0101	0.9939	0.1205	0.7299	0.1306	0.6975
Psf1	Psf1	1629713_at	0.3324	0.0791	0.5382	0.0472	0.3962	0.0658	-0.0956	0.8312	0.0243	0.9197	0.1199	0.4593	0.0101	0.9916	0.2037	0.3458	0.1936	0.3921
bgcn	Putative nuclear b	1629714_at	0.1039	0.5557	-0.0530	0.6600	-0.1636	0.5844	-0.0383	0.9641	0.2261	0.3236	0.2644	0.1896	-0.0844	0.9230	-0.0545	0.9009	0.0300	0.9412
CG7891	CG7891	1629715_at	-0.0542	0.8517	0.3402	0.0911	0.5153	0.0134	0.2248	0.6247	0.0156	0.9643	-0.2092	0.3122	0.1033	0.8906	0.4243	0.1345	0.3210	0.2738
CG13284	CG13284	1629716_a_at	2.3557	0.0023	0.6682	0.1434	1.4255	0.0035	0.3574	0.2654	0.8857	0.0014	0.5283	0.0080	-0.4011	0.8454	-0.9055	0.2664	-0.5044	0.5744
CG33795	CG33795	1629717_at	0.1649	0.4578	0.3242	0.3711	0.0129	0.9470	-0.0231	0.9768	0.0677	0.7737	0.0908	0.6528	0.3251	0.7464	0.2566	0.5644	-0.0685	0.9068
CG16727	CG16727	1629718_at	0.0647	0.8922	0.0988	0.4315	0.0309	0.8706	0.1777	0.8807	-0.1403	0.7730	-0.3180	0.3977	0.2269	0.7916	0.0329	0.9557	-0.1940	0.6211
---	---	1629719_at	0.2394	0.3112	0.0858	0.6330	0.2344	0.2370	0.0445	0.9375	0.0335	0.8789	-0.0111	0.9577	-0.1583	0.8446	-0.1667	0.6437	-0.0084	0.9874
---	---	1629720_s_at	0.1259	0.3648	-0.0312	0.8320	-0.0012	0.9964	-0.2289	0.5756	-0.2869	0.1686	-0.0579	0.8054	-0.3502	0.3712	-0.4110	0.0714	-0.0608	0.8209

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1629721_at	0.0323	0.8620	-0.0970	0.5055	0.0436	0.8399	0.1335	0.8376	0.2331	0.3489	0.0996	0.6989	-0.0603	0.8973	-0.0479	0.8376	0.0124	0.9614
---	---	1629722_at	-0.0250	0.9124	-0.0166	0.8754	-0.0631	0.7529	0.1055	0.8140	0.1384	0.4518	0.0329	0.8753	0.1289	0.8680	0.0513	0.9141	-0.0776	0.8434
CG14714	CG14714	1629723_at	-1.8903	0.0112	1.7740	0.1597	0.3452	0.5416	-0.3937	0.7556	-1.9935	0.0039	-1.5998	0.0062	1.1717	0.6955	1.6850	0.1623	0.5134	0.7144
---	---	1629724_at	-0.2145	0.2678	-0.0218	0.8401	0.0019	0.9958	0.0042	0.9956	0.0780	0.7627	0.0738	0.7539	-0.0884	0.9092	0.0283	0.9496	0.1167	0.7053
CG33331	CG33331	1629725_at	-0.0618	0.7105	0.4284	0.1425	1.5722	0.0001	0.0485	0.9686	-0.8006	0.0205	-0.8491	0.0098	-0.8095	0.2884	0.0017	0.9994	0.8111	0.0755
CG10472 /// DyakCG10472	CG10472	1629726_at	-0.6322	0.3728	-0.3657	0.2050	-0.4472	0.0186	0.1281	0.8908	-0.6952	0.0335	-0.8233	0.0102	-0.1426	0.9589	-0.5634	0.4870	-0.4208	0.6225
CG14258	CG14258	1629727_at	0.1550	0.4858	-0.0898	0.5372	0.1321	0.3913	0.0369	0.9515	0.1213	0.4901	0.0844	0.6207	-0.2349	0.7215	-0.1740	0.5415	0.0609	0.8668
---	---	1629728_s_at	0.0314	0.8844	0.1876	0.2763	0.3329	0.1787	0.0528	0.9352	-0.1292	0.5271	-0.1819	0.2947	0.0005	0.9999	0.0117	0.9844	0.0111	0.9835
CG31019	CG31019	1629729_at	-0.0380	0.9102	0.0819	0.5474	0.0384	0.8647	-0.1611	0.8578	-0.2362	0.4830	-0.0751	0.8397	-0.0900	0.8768	-0.0370	0.9157	0.0529	0.8577
Tom7	Translocase of ou	1629730_at	0.1753	0.4651	0.5624	0.0251	0.7155	0.0037	0.0290	0.9675	-0.2734	0.1429	-0.3024	0.0730	-0.1280	0.8875	0.0520	0.9217	0.1800	0.6216
CG14565	CG14565	1629731_at	0.2466	0.1991	0.1429	0.1652	0.0661	0.6903	-0.1445	0.6196	-0.0257	0.8966	0.1188	0.3751	-0.1835	0.7230	-0.1060	0.6653	0.0775	0.7640
CG8791	CG8791	1629732_at	-0.1011	0.9624	0.2017	0.7173	-2.2580	0.0009	-1.9857	0.2101	-1.3217	0.1630	0.6640	0.4615	0.3739	0.9611	-0.8908	0.7166	-1.2647	0.5808
Lim1	---	1629733_at	0.5680	0.1981	-0.1709	0.7758	-0.2765	0.5613	0.1518	0.9371	0.7759	0.1626	0.6241	0.2114	0.2094	0.9142	0.0986	0.9269	-0.1107	0.9057
---	---	1629734_at	0.0687	0.6849	-0.1008	0.3480	0.3159	0.0734	0.1364	0.6591	0.1929	0.1919	0.0565	0.7212	-0.1283	0.7707	-0.0519	0.8372	0.0764	0.7200
CG40264	CG40264	1629735_s_at	0.1705	0.4520	0.1671	0.3216	0.1144	0.5396	-0.1175	0.8171	0.0610	0.8031	0.1785	0.3271	-0.1707	0.8157	-0.1182	0.7393	0.0525	0.9004
---	---	1629736_at	-0.0235	0.9056	0.1285	0.4827	0.1361	0.4636	-0.1957	0.5552	-0.0435	0.8446	0.1522	0.3311	-0.0870	0.8846	0.0422	0.9033	0.1293	0.5969
dpa	Drosophila prolife	1629737_at	0.4696	0.3609	-0.8031	0.3974	-0.6640	0.0827	0.0129	0.9943	1.5429	0.0029	1.5300	0.0018	-0.2430	0.9550	0.0244	0.9935	0.2674	0.8751
CG14957	CG14957	1629738_at	-0.3258	0.3377	0.0785	0.4429	0.0042	0.9896	0.0311	0.9824	-0.3133	0.3507	-0.3444	0.2449	0.1596	0.8461	0.1397	0.7180	-0.0199	0.9695
---	---	1629739_s_at	0.2088	0.2185	0.0440	0.6373	-0.1200	0.4435	-0.1392	0.7425	0.0342	0.8924	0.1734	0.3059	0.0390	0.9590	-0.1111	0.6394	-0.1501	0.5092
His1:CG31617 /// His1:CG3	His1:CG31617 ///	1629740_at	0.4962	0.6067	-1.5030	0.1476	-1.3272	0.0565	-0.2401	0.8281	2.7321	0.0004	2.9723	0.0002	-0.4655	0.9301	0.7482	0.7000	1.2137	0.4944
CG32666 /// DsimCG32666	CG32666	1629741_at	0.6004	0.2162	0.0478	0.8886	0.5375	0.0517	-0.2005	0.8096	-0.0979	0.8108	0.1026	0.7805	-0.6690	0.6749	-0.5716	0.3706	0.0974	0.9131
CG31058	CG31058	1629742_at	0.0090	0.9706	0.0626	0.5858	-0.0400	0.8487	-0.0485	0.9353	-0.0822	0.6834	-0.0337	0.8714	-0.0033	0.9970	-0.1674	0.4355	-0.1641	0.4597
CG11663	CG11663	1629743_at	0.0578	0.7911	0.0315	0.8285	-0.0479	0.7757	-0.0358	0.9610	-0.0353	0.8942	0.0005	0.9982	0.1720	0.8023	-0.0292	0.9499	-0.2013	0.4978
alpha-Man-I	alpha-mannosidas	1629744_a_at	-1.3292	0.0082	-1.5088	0.0091	-1.6202	0.0005	-0.1981	0.7845	0.5808	0.0563	0.7789	0.0107	-0.1837	0.8609	0.2657	0.5469	0.4494	0.2993
CG6439	Isocitrate dehydro	1629745_at	-0.9178	0.0105	0.2738	0.3012	0.1123	0.6521	-0.1076	0.8640	-1.3902	0.0004	-1.2827	0.0004	-0.0012	0.9998	-0.1296	0.8034	-0.1284	0.7932
CG4041 /// DmirCG4041	CG4041	1629746_at	-0.3437	0.3668	0.8525	0.1567	1.1194	0.0002	0.2224	0.7475	-0.8442	0.0127	-1.0667	0.0027	-0.0979	0.9717	0.4848	0.4875	0.5828	0.4017
Cpr49Ag	CG8511	1629747_at	0.3267	0.3005	0.0386	0.8429	0.0365	0.9044	0.0402	0.9639	-0.0561	0.8540	-0.0964	0.6968	0.2268	0.8589	-0.2242	0.7003	-0.4510	0.3921
CG2747	CG2747	1629748_s_at	-0.7693	0.0620	0.4440	0.4402	0.4697	0.0667	0.2282	0.8321	-0.2604	0.5546	-0.4885	0.1807	0.2457	0.8744	0.9972	0.1089	0.7515	0.2338
CG13870	CG13870	1629749_a_at	0.2830	0.2887	0.0127	0.9022	0.1507	0.4683	0.0871	0.8605	0.2139	0.2278	0.1268	0.4493	0.1502	0.8298	-0.0277	0.9531	-0.1779	0.5670
PpY-55A	protein phosphata	1629750_at	0.0815	0.6734	0.1472	0.2787	-0.0191	0.9330	0.0309	0.9744	-0.1006	0.7100	-0.1315	0.5712	0.1597	0.7673	0.0106	0.9797	-0.1491	0.5360
CG13693	CG13693	1629751_at	0.0822	0.6658	-0.0665	0.6888	-0.0586	0.7711	0.0175	0.9838	0.1926	0.3491	0.1751	0.3452	-0.0402	0.9689	-0.0548	0.8893	-0.0146	0.9721
CG5525	CG5525	1629752_at	-0.1703	0.4374	0.0570	0.6944	0.2804	0.3421	0.1816	0.7168	0.2364	0.3044	0.0548	0.8336	0.0725	0.9653	0.5719	0.1857	0.4994	0.2782
Kr	kruppel	1629753_at	0.1218	0.5251	-0.1095	0.3376	-0.1023	0.4759	0.0554	0.9012	0.0795	0.6204	0.0241	0.8893	0.1042	0.8740	-0.0734	0.8306	-0.1776	0.5088
Trl	GAGA factor	1629754_s_at	0.2186	0.5205	0.8114	0.0301	0.4424	0.0365	0.0314	0.9755	0.0190	0.9598	-0.0124	0.9688	0.3644	0.6531	0.5666	0.0917	0.2022	0.5585
---	---	1629755_at	0.2340	0.3164	0.2156	0.3602	0.3646	0.0710	0.0415	0.9715	0.1045	0.7666	0.0631	0.8533	-0.0060	0.9964	0.1568	0.6458	0.1628	0.6311
CG11092	CG11092	1629756_at	0.3219	0.0509	0.5165	0.1698	0.8195	0.0041	-0.0443	0.9518	-0.0835	0.7262	-0.0392	0.8704	-0.2531	0.7823	0.2167	0.6093	0.4697	0.2532
CG30327	CG30327	1629757_at	-0.1824	0.4755	0.0323	0.7537	-0.0481	0.8562	-0.1525	0.7608	-0.1700	0.4484	-0.0175	0.9491	-0.0430	0.9677	-0.0048	0.9940	0.0382	0.9211
Aats-pro	Prolyl-tRNA synth	1629758_at	-0.2755	0.1464	0.1712	0.5517	0.4466	0.0234	-0.0073	0.9952	-0.4847	0.0520	-0.4774	0.0358	-0.3174	0.6955	-0.1187	0.7778	0.1987	0.5856
CG18815	CG18815	1629759_at	0.0272	0.8794	0.5065	0.0270	0.4304	0.0243	-0.0414	0.9441	-0.2565	0.1271	-0.2151	0.1526	-0.0070	0.9946	0.1220	0.6382	0.1289	0.6166
---	---	1629760_s_at	-0.6376	0.0659	-1.2676	0.1784	-2.1091	0.0001	-0.1989	0.8190	0.9033	0.0160	1.1022	0.0040	0.2856	0.9138	-0.0478	0.9766	-0.3334	0.7534
CG17781	CG17781	1629761_at	0.6704	0.3579	0.1667	0.3672	0.1431	0.5408	-0.0098	0.9893	0.0048	0.9846	0.0146	0.9442	0.1029	0.9717	-0.3529	0.6605	-0.4557	0.5578
CG5148	CG5148	1629762_at	0.3941	0.3721	-0.1795	0.5069	-0.3600	0.1110	0.0711	0.9696	0.8322	0.0846	0.7611	0.0780	0.1938	0.8236	0.2489	0.5068	0.0551	0.9121
CG11262	CG11262	1629763_at	0.2382	0.1174	-0.0864	0.6546	-0.0744	0.6974	0.2522	0.6402	0.2902	0.2732	0.0381	0.9045	0.0590	0.9405	-0.0574	0.8675	-0.1164	0.6593
---	---	1629764_at	0.1824	0.2292	0.2403	0.3381	0.0383	0.8952	-0.1141	0.8281	-0.0072	0.9809	0.1068	0.5988	0.1056	0.9048	-0.0191	0.9714	-0.1247	0.7327
CG7715	CG7715	1629765_at	0.2408	0.1669	0.2507	0.1937	0.0113	0.9717	-0.0793	0.9149	-0.0147	0.9678	0.0647	0.8047	0.1971	0.7848	-0.0808	0.8485	-0.2779	0.3818
ca	claret	1629766_at	-0.4411	0.3332	0.0179	0.9874	0.6269	0.0447	0.0903	0.8915	-0.2749	0.2147	-0.3651	0.0688	-0.3623	0.8810	0.3436	0.7544	0.7059	0.4597
Src42A	Suppressor of poli	1629767_s_at	0.5323	0.0449	-0.1797	0.5125	-0.6421	0.0104	0.0273	0.9672	0.9768	0.0007	0.9495	0.0004	0.4537	0.6272	0.2872	0.4655	-0.1665	0.7022
CG40356	CG40356	1629768_at	0.1027	0.6807	0.1458	0.3836	0.0766	0.6842	0.0454	0.9507	0.0047	0.9881	-0.0407	0.8640	0.0431	0.9732	-0.0222	0.9658	-0.0653	0.8795
CG4763	CG4763	1629769_at	-0.1721	0.3001	0.0439	0.7545	0.0950	0.5590	0.0029	0.9962	-0.0815	0.7343	-0.0844	0.6941	-0.0441	0.9555	0.0136	0.9719	0.0578	0.8444
---	---	1629770_at	0.2905	0.2427	0.0435	0.6971	-0.0126	0.9477	-0.0412	0.9542	-0.0171	0.9531	0.0241	0.9208	0.1246	0.8760	-0.0921	0.8215	-0.2168	0.5034
Cyp12a5	Cyp12a5	1629771_at	0.5973	0.0859	0.7472	0.0523	1.6297	0.0006	-0.1071	0.8942	-1.5508	0.0006	-1.4437	0.0005	-0.6695	0.6211	-1.2220	0.0529	-0.5525	0.3458
CG9541	CG9541	1629772_at	1.7620	0.0397	2.3409	0.0170	2.9169	0.0000	0.7682	0.1140	0.7507	0.0182	-0.0175	0.9619	0.0573	0.9913	1.2888	0.2239	1.2315	0.2784
CG14518	CG14518	1629773_at	-1.2020	0.0560	-1.6709	0.0100	-1.5666	0.0038	0.3607	0.7604	0.5412	0.2921	0.1805	0.7420	0.0405	0.9764	0.0156	0.9793	-0.0250	0.9582
CG5357	CG5357	1629774_at	0.1009	0.6740	0.1271	0.3490	-0.1569	0.6126	-0.2018	0.6720	0.1529	0.5249	0.3546	0.0805	0.0747	0.9717	0.1513	0.8329	0.0766	0.9176
---	---	1629775_at	0.1516	0.4273	0.0430	0.6452	-0.1779	0.2560	0.0236	0.9704	0.0817	0.6557	0.0581	0.7398	0.1043	0.9063	-0.0378	0.9409	-0.1421	0.6861
CG6643	CG6643	1629776_a_at	-0.8080	0.0408	-0.7933	0.0348	-0.6451	0.0046	0.2855	0.5603	0.2061	0.4287	-0.0795	0.7757	0.1680	0.8427	0.1931	0.6032	0.0251	0.9609
CG9588	CG9588	1629777_at	-0.1905	0.3572	0.7448	0.0085	0.8842	0.0032	-0.0386	0.9592	-0.9854	0.0011	-0.9468	0.0008	-0.0755	0.9421	0.0731	0.8734	0.1486	0.6717
CG5130	CG5130	1629778_s_at	1.4246	0.0106	1.3072	0.0608	1.8143	0.0007	0.4110	0.6480	0.5519	0.2034	0.1410	0.7693	-0.1995	0.8653	0.2039	0.7003	0.4035	0.3981
Mcr	Macroglobulin con	1629779_at	-0.2780	0.2111	0.0415	0.9211	-0.4362	0.0377	0.0112	0.9893	0.2057	0.2867	0.1946	0.2595	0.4339	0.6817	0.5019	0.2329	0.0681	0.9096
CG13471	CG13471	1629780_at	0.1572	0.3138	0.1168	0.3849	0.2342	0.4005	0.0597	0.9136	0.1273	0.4845	0.0676	0.7140	-0.1151	0.8521	0.0271	0.9478	0.1422	0.6045
CG32334	CG32334	1629781_at	0.1171	0.6300	0.0338	0.7456	0.1358	0.4234	0.0968	0.8839	-0.1144	0.6530	-0.2112	0.3029	-0.0116	0.9898	0.0007	0.9994	0.0123	0.9667
---	---	1629782_at	0.1475	0.5596	0.0016	0.9979	0.1719	0.4276	0.1172	0.9311	-0.0296	0.9604	-0.1468	0.7307	-0.1511	0.8379	-0.0692	0.8736	0.0819	0.8307
---	---	1629783_s_at	-0.1127	0.7566	-0.1656	0.6176	-0.1713	0.3873	0.0159	0.9937	-0.0028	0.9960	-0.0187	0.9694	0.1498	0.8903	0.0011	0.9996	-0.1487	0.7520
Myo28B1	myosin VII 28B1-I	1629784_a_at	2.0008	0.0233	0.5730	0.1470	1.7782	0.0003	0.8564	0.2674	1.2734	0.0129	0.4169	0.3157	-0.0489	0.9862	0.2851	0.6912	0.3340	0.6282
Sp1	Sp1	1629785_a_at	-1.7316	0.0017	0.1230	0.2828	0.0978	0.7265	-0.4212	0.4895	-1.9066	0.0006	-1.4854	0.0009	-0.1929	0.7726	-0.0504	0.9081	0.1425	0.6419
CG31116	CG31116	1629786_a_at	0.1545	0.5102	0.7914	0.1039	0.3314	0.1894	-0.5450	0.1927	-0.1117	0.7019	0.4333	0.0595	0.0671	0.9741	0.5780	0.2378	0.5110	0.3227
CG31642	CG31642	1629787_at	0.0338	0.8766	-0.0681	0.6341	0.1827	0.2351	0.0502	0.9345	0.0276	0.9127	-0.0226	0.9187	-0.0654	0.8941	-0.0543	0.8218	0.0111	0.9686
CG33182	CG33182	1629788_at	0.7700	0.2670	0.6099	0.3669	-0.2358	0.7351	-0.4618	0.2352	0.3631	0.1145	0.8250	0.0025	0.4940	0.8842	0.2832	0.8779	-0.2109	0.9057
CG9649	CG9649	1629789_at	3.3308	0.0009	0.8486	0.3099	2.4554	0.0002	0.8580	0.4183	1.0599	0.0740	0.2019	0.7529	-0.7323	0.7046	-1.4405	0.0853	-0.7083	0.3891

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11267	CG11267	1629790_at	0.2573	0.3301	-0.0393	0.9124	0.0844	0.8349	0.0667	0.8908	0.1315	0.4380	0.0647	0.7085	0.0419	0.9869	-0.1303	0.8764	-0.1722	0.8119
CG17271	CG17271	1629791_at	1.0761	0.0029	0.9647	0.0285	1.1158	0.0008	0.1667	0.7030	0.6021	0.0095	0.4354	0.0237	0.0481	0.9812	0.4669	0.2660	0.4188	0.3493
CG17493 /// DyakCG17493	CG17493	1629792_at	-0.1027	0.6718	0.1534	0.5568	-0.1065	0.5672	-0.2294	0.6122	-0.2990	0.1858	-0.0696	0.7837	0.1021	0.9031	0.1099	0.7667	0.0078	0.9874
CG12772	CG12772	1629793_at	-0.9936	0.0238	0.2404	0.2671	0.2728	0.3426	-0.1394	0.8578	-0.8412	0.0092	-0.7019	0.0125	-0.0879	0.9545	0.1707	0.7501	0.2586	0.5956
iHog	interference Hedg	1629794_at	-0.8413	0.0233	-1.5620	0.0285	-1.8976	0.0001	-0.0310	0.9675	0.5145	0.0186	0.5456	0.0088	-0.0060	0.9984	-0.4348	0.4871	-0.4289	0.4984
PTP-ER	Protein tyrosine pl	1629795_at	-0.4434	0.3525	-0.3475	0.3363	-0.4738	0.0929	-0.3294	0.4507	-0.0369	0.9108	0.2925	0.1665	-0.0769	0.9741	-0.0602	0.9495	0.0167	0.9855
cin	cinnamon	1629796_s_at	0.2023	0.2364	0.2679	0.4856	0.1429	0.5742	-0.1355	0.7409	0.2405	0.1811	0.3760	0.0278	0.0031	0.9994	0.3121	0.5214	0.3090	0.5326
Rab6	warthog	1629797_at	0.4786	0.0183	0.2718	0.4659	0.0691	0.7525	0.0785	0.8955	1.0943	0.0007	1.0159	0.0006	0.0466	0.9778	0.7298	0.0795	0.6832	0.1192
CG34365	CG12958	1629798_at	-0.0706	0.8025	-0.2837	0.0811	-0.3211	0.0862	0.2022	0.6513	0.6145	0.0118	0.4123	0.0394	0.0895	0.8461	0.1434	0.4514	0.0539	0.8204
sphinx2	CG32382	1629799_at	0.1026	0.6808	0.1705	0.3716	0.1029	0.5740	-0.1781	0.7098	-0.1895	0.3967	-0.0114	0.9674	-0.2006	0.7588	-0.1600	0.5825	0.0406	0.9152
---	---	1629800_at	0.0268	0.8734	0.1536	0.1988	0.0383	0.8765	-0.0005	0.9994	0.0009	0.9979	0.0014	0.9958	0.0125	0.9898	0.0653	0.8028	0.0528	0.8393
CG13050	CG13050	1629801_at	0.1959	0.2269	0.0616	0.5742	0.0382	0.8381	-0.1969	0.5842	-0.1001	0.6234	0.0968	0.5996	0.0801	0.8513	0.0116	0.9682	-0.0685	0.7369
CG15484	CG15484	1629802_at	0.0721	0.6553	-0.4396	0.0302	-0.5362	0.0116	-0.0380	0.9637	0.3562	0.1045	0.3942	0.0500	0.1356	0.8215	-0.0477	0.8993	-0.1833	0.4786
scb	volado	1629803_a_at	0.0015	0.9965	-0.4820	0.1849	-0.2669	0.6660	0.6947	0.1372	1.7217	0.0005	1.0270	0.0024	0.4038	0.8692	0.9545	0.3160	0.5507	0.6005
CG6329	CG6329	1629804_s_at	-1.8890	0.0256	-1.8157	0.0427	-2.6887	0.0000	-0.2191	0.7795	-0.7422	0.0298	-0.5231	0.0735	0.0977	0.9742	-0.5671	0.4485	-0.6647	0.3838
SAK	SAK	1629805_at	-0.8406	0.0915	-0.7477	0.1162	-0.8870	0.0182	-0.3269	0.7815	-0.1385	0.8202	0.1884	0.7169	-0.2302	0.7810	-0.2632	0.4673	-0.0330	0.9471
CG32103	CG32103	1629806_a_at	0.2076	0.5924	0.2786	0.1146	0.2807	0.1375	0.0367	0.9777	0.2837	0.4027	0.2470	0.4245	0.0795	0.9552	0.4993	0.2123	0.4197	0.3205
---	---	1629807_at	0.2562	0.1520	0.1393	0.4967	0.1928	0.3174	0.0487	0.9471	0.0782	0.7518	0.0295	0.9068	-0.0572	0.9652	0.0511	0.9268	0.1082	0.7982
---	---	1629808_at	-0.0065	0.9857	0.4853	0.3450	0.4177	0.0715	0.1798	0.7435	0.0741	0.8027	-0.1057	0.6760	0.0549	0.9707	0.3302	0.3738	0.2753	0.4824
CG30491	CG30491	1629809_at	0.0062	0.9861	-0.3241	0.4151	-0.1521	0.4967	-0.0066	0.9937	0.2519	0.1295	0.2585	0.0839	-0.2831	0.7644	-0.0959	0.8632	0.1872	0.6668
CG14340	CG14340	1629810_at	-0.0906	0.6479	0.0219	0.8553	0.2439	0.3130	0.1283	0.7598	-0.1407	0.4532	-0.2690	0.0948	-0.0371	0.9831	0.1256	0.8151	0.1627	0.7307
CG9619	CG9619	1629811_at	0.1237	0.8675	-0.1797	0.6150	-0.3565	0.2675	0.1678	0.8822	-0.0299	0.9585	-0.1977	0.6136	0.0583	0.9906	-0.6465	0.5543	-0.7049	0.5194
CG14915 /// DyakCG14915	CG14915	1629812_at	0.0801	0.6988	-0.0038	0.9873	0.0775	0.7591	0.0967	0.8676	0.1535	0.4737	0.0568	0.8049	-0.0318	0.9816	-0.0056	0.9935	0.0262	0.9527
CG17691	CG17691	1629813_at	-0.5304	0.1924	-0.4712	0.0426	-0.7167	0.0016	-0.3911	0.2474	-0.5462	0.0143	-0.1551	0.4066	-0.2146	0.8736	-0.5254	0.3076	-0.3108	0.5814
RacGAP84C	RnRacGAP	1629814_at	-0.0502	0.7365	-0.0585	0.5913	0.1202	0.5808	-0.0199	0.9777	0.0172	0.9492	0.0371	0.8598	-0.2757	0.6749	-0.0357	0.9350	0.2400	0.3897
CG33679	CG33679	1629815_at	-0.0610	0.7583	0.0072	0.9579	-0.0134	0.9536	0.1781	0.6763	0.0116	0.9703	-0.1665	0.3682	0.0546	0.9400	0.1040	0.6714	0.0494	0.8668
---	---	1629816_at	-0.0399	0.8466	-0.0963	0.5657	-0.0036	0.9900	0.0894	0.8987	0.0087	0.9801	-0.0807	0.7452	-0.1101	0.8802	-0.0165	0.9720	0.0936	0.7764
CG3339	CG3339	1629817_at	0.1889	0.2845	0.1574	0.3657	0.0852	0.6423	-0.2151	0.4770	0.0254	0.9105	0.2405	0.0973	0.0114	0.9898	-0.0081	0.9828	-0.0194	0.9431
---	---	1629818_at	0.0983	0.7290	-0.0317	0.8835	0.2763	0.2417	0.0610	0.9584	0.1408	0.6905	0.0799	0.8191	-0.2328	0.7230	-0.1894	0.5191	0.0435	0.9115
Sema-2a	semiphorin	1629819_s_at	-2.3322	0.0069	-3.5246	0.0144	-4.4525	0.0000	-0.5424	0.2747	-0.6235	0.0394	-0.0811	0.8047	0.2916	0.9296	-1.8202	0.0954	-2.1119	0.0834
CG14353	CG14353	1629820_at	-0.2175	0.3638	0.5211	0.0248	0.6492	0.0264	-0.2605	0.4979	-0.3234	0.1108	-0.0630	0.7809	-0.3302	0.6749	0.3778	0.2366	0.7080	0.0702
CG30055	CG30055	1629821_at	-0.3963	0.1074	-0.1287	0.7246	-0.0421	0.8196	-0.2499	0.6360	-0.7201	0.0132	-0.4701	0.0480	-0.2223	0.8122	-0.3988	0.2912	-0.1765	0.6800
CG15506	CG15506	1629822_a_at	0.1607	0.7332	0.1686	0.4857	0.2607	0.2531	0.1923	0.7924	0.3547	0.2317	0.1624	0.5839	-0.0724	0.9734	0.2395	0.6883	0.3119	0.5833
CG4230	CG4230	1629823_at	-0.6178	0.0636	0.3570	0.3340	0.2617	0.1246	0.0514	0.9651	-0.8016	0.0189	-0.8530	0.0088	-0.2242	0.8205	0.1652	0.7275	0.3894	0.3565
CG13081	CG13081	1629824_at	-0.0152	0.9432	0.0009	0.9952	0.0879	0.6230	0.0172	0.9857	-0.1835	0.4084	-0.2008	0.3053	-0.0832	0.8875	-0.1408	0.5411	-0.0576	0.8372
CG1801	CG1801	1629825_a_at	0.2644	0.0816	0.0537	0.6276	0.2076	0.3070	0.0153	0.9850	-0.0126	0.9639	-0.0279	0.8991	-0.0768	0.8878	-0.0387	0.9004	0.0381	0.8918
---	---	1629826_at	0.0530	0.8958	0.0208	0.8437	0.1761	0.4346	0.0450	0.9664	0.0650	0.8593	0.0200	0.9554	-0.0678	0.9296	0.0040	0.9941	0.0718	0.8172
Hsp70Ba /// Hsp70Bb /// Hs	Hsp70Bbb /// 87C	1629827_s_at	0.0551	0.8517	-0.1974	0.2942	0.0912	0.7650	0.0686	0.9435	0.2994	0.2742	0.2309	0.3568	-0.0423	0.9841	0.1452	0.8161	0.1874	0.7341
CG13375	CG13375	1629828_at	-0.0142	0.9474	0.1622	0.3704	0.0876	0.6480	0.0299	0.9722	0.0229	0.9411	-0.0070	0.9793	0.2154	0.6749	0.3337	0.1199	0.1184	0.6040
CG13454	CG13454	1629829_at	0.1732	0.2800	0.1357	0.5894	0.1974	0.2236	-0.0378	0.9496	-0.0339	0.8789	0.0040	0.9851	-0.1385	0.8427	-0.0647	0.8750	0.0738	0.8407
Or7a	Odorant receptor	1629830_at	0.0443	0.8096	0.0087	0.9660	-0.0923	0.6301	0.0322	0.9600	-0.0076	0.9771	-0.0398	0.8368	0.0120	0.9895	0.0030	0.9941	-0.0090	0.9767
CG17801	CG17801	1629831_at	0.0386	0.8810	0.2794	0.1559	-0.0763	0.7051	-0.1162	0.8280	-0.1672	0.4317	-0.0511	0.8275	0.0763	0.9029	0.1205	0.6216	0.0442	0.8879
---	---	1629832_at	0.2167	0.2314	0.1166	0.3871	0.0090	0.9666	-0.0612	0.9252	-0.1022	0.6418	-0.0409	0.8576	0.0780	0.9467	-0.1649	0.6793	-0.2429	0.5163
Gen	XPG-like endonuc	1629833_at	-0.2473	0.4046	-0.2125	0.3147	-0.1829	0.4300	-0.0871	0.8875	-0.0441	0.8728	0.0431	0.8591	-0.2141	0.7689	-0.0050	0.9941	0.2091	0.5179
CG18628	CG18628	1629834_at	-3.7220	0.2815	-4.5400	0.1423	-3.3967	0.1262	0.0359	0.9435	0.0694	0.6685	0.0335	0.8378	-1.0860	0.9589	-1.5688	0.8500	-0.4828	0.9553
CG5048	CG5048	1629835_at	-0.0247	0.9196	-0.6635	0.0410	-0.3461	0.1480	0.5838	0.1560	0.7805	0.0078	0.1967	0.3960	-0.0875	0.9056	-0.0791	0.8185	0.0084	0.9847
Ugt36Ba	Ugt36Ba	1629836_at	-0.8788	0.0171	-0.4890	0.0312	-0.5359	0.0135	0.2108	0.6409	-0.9034	0.0024	-1.1141	0.0006	-0.1739	0.6999	-0.5467	0.0276	-0.3728	0.0852
CG2330	CG2330	1629837_at	-2.4389	0.0022	-3.1922	0.0106	-3.4426	0.0000	-0.0252	0.9722	0.4728	0.0169	0.4980	0.0082	0.2782	0.9076	-0.2094	0.8595	-0.4876	0.5942
CG15925	CG15925	1629838_at	0.3398	0.0917	0.1067	0.5355	-0.0087	0.9666	0.0745	0.8776	0.1545	0.3616	0.0800	0.6384	0.1452	0.8141	0.0275	0.9472	-0.1177	0.6734
Ggamma1	G protein gamma	1629839_a_at	-1.5578	0.0216	-0.2576	0.4702	-0.7579	0.0167	-0.0747	0.9558	-0.9435	0.0178	-0.8688	0.0157	0.1920	0.9400	0.1771	0.8749	-0.0149	0.9914
---	---	1629840_at	-0.0315	0.8505	0.2330	0.1078	0.1304	0.4475	-0.0545	0.8979	-0.1063	0.4663	-0.0517	0.7295	0.0792	0.9021	0.0438	0.9027	-0.0353	0.9128
CG32418	CG32418	1629841_at	0.2286	0.3598	0.1253	0.5822	0.1399	0.3835	0.1494	0.7426	0.5186	0.0178	0.3692	0.0449	0.1124	0.9370	0.4108	0.3469	0.2984	0.5258
Gap1	sextra	1629842_at	-0.5794	0.0525	-0.8695	0.0138	-1.0057	0.0020	0.3572	0.4102	0.6071	0.0199	0.2499	0.2471	0.4886	0.5519	0.3210	0.3909	-0.1676	0.6892
Ubc-E2H	Ubc-E2H	1629843_s_at	-0.1566	0.5573	0.1513	0.2123	0.3838	0.1708	-0.1799	0.6338	-0.7034	0.0033	-0.5235	0.0073	-0.2377	0.8122	-0.2302	0.6012	0.0075	0.9921
rap	fizzy-related	1629844_s_at	-0.0610	0.8672	0.3266	0.3636	0.2121	0.4306	-0.0104	0.9937	-0.4304	0.0922	-0.4200	0.0683	0.1438	0.9246	-0.0281	0.9741	-0.1719	0.7754
CG34375	CG13834	1629845_at	0.0443	0.8102	-0.0756	0.5280	-0.0784	0.6263	0.1246	0.8164	0.2713	0.1971	0.1467	0.4638	0.1366	0.7387	0.1720	0.3090	0.0354	0.8812
CG11619	CG11619	1629846_at	-0.0201	0.9365	0.1672	0.1939	0.3035	0.1018	-0.1237	0.7821	-0.2797	0.1294	-0.1561	0.3594	-0.1005	0.8882	0.0262	0.9508	0.1267	0.6623
Pxn	peroxidasin	1629847_at	-0.2961	0.1377	0.0368	0.6955	-0.1472	0.4991	0.0854	0.8942	-0.1932	0.3777	-0.2786	0.1446	0.1563	0.8292	0.0774	0.8527	-0.0789	0.8375
CG17184	CG17184	1629848_s_at	-0.2476	0.0959	-0.1106	0.7173	-0.3027	0.0849	-0.0742	0.8817	0.1291	0.4689	0.2034	0.1813	0.1147	0.8692	0.2392	0.3778	0.1245	0.6843
---	---	1629849_at	0.1799	0.2311	-0.2132	0.2553	0.1075	0.5560	0.3655	0.1605	0.2794	0.0787	-0.0861	0.5853	-0.0508	0.9646	-0.2321	0.4523	-0.1812	0.5808
---	---	1629850_at	0.0530	0.8071	0.3726	0.0463	0.2605	0.1286	-0.3180	0.4200	-0.2544	0.2454	0.0636	0.7958	-0.1590	0.7707	0.0654	0.8343	0.2244	0.3503
CG31998	CG31998	1629851_at	0.1286	0.7529	0.6261	0.3417	0.0785	0.6689	-0.5267	0.2935	0.0003	0.9994	0.5271	0.0502	0.0678	0.9823	0.5457	0.4410	0.4779	0.5192
CG9018	CG9018	1629852_at	0.4906	0.0362	-0.0170	0.9342	-0.0098	0.9695	0.0899	0.8327	0.5037	0.0079	0.4138	0.0116	0.1832	0.8202	0.0642	0.8981	-0.1190	0.7568
CG3699	CG3699	1629853_at	3.2452	0.0012	1.5524	0.3866	3.8210	0.0002	2.1187	0.0055	0.0917	0.8431	-2.0270	0.0004	-0.1417	0.9860	-1.6527	0.3446	-1.5110	0.4078
CG15743	CG15743	1629854_at	-0.9523	0.0084	-1.1551	0.0032	-0.5723	0.0553	-0.1083	0.9080	0.2817	0.3575	0.3901	0.1451	-0.6611	0.4377	0.0961	0.8797	0.7572	0.1192
---	---	1629855_at	0.0030	0.9915	0.0339	0.7948	0.0480	0.8486	-0.0024	0.9967	0.0948	0.6328	0.0971	0.5844	0.1177	0.8657	0.0551	0.8938	-0.0626	0.8648
---	---	1629856_at	0.0215	0.9211	0.0888	0.4550	-0.0499	0.7626	-0.0792	0.8663	-0.0993	0.5818	-0.0201	0.9214	0.0806	0.8571	-0.0523	0.8287	-0.1328	0.4824
CG10359	CG10359	1629857_at	-1.9292	0.0385	-4.4690	0.0025	-3.5730	0.0000	0.6375	0.2789	1.1792	0.0053	0.5418	0.0821	-0.2736	0.9365	-1.2649	0.2265	-0.9913	0.3728
CG2528	CG2528	1629858_at	0.2778	0.1938	0.1446	0.2320	-0.0319	0.8418	-0.1481	0.6084	-0.0376	0.8409	0.1106	0.4149	0.0222	0.9869	-0.1335	0.7000	-0.1557	0.6375

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
vap	vacuolar peduncle	1629859_s_at	-0.7280	0.0031	-1.2686	0.0344	-1.7629	0.0001	-0.2779	0.3431	0.2452	0.1442	0.5231	0.0048	0.1620	0.8894	-0.2393	0.6093	-0.4013	0.3719
CG6462	CG6462	1629860_at	0.2252	0.4726	0.2126	0.3890	0.3626	0.0822	0.1327	0.8747	-0.0343	0.9338	-0.1670	0.5646	0.0391	0.9816	0.0608	0.9175	0.0216	0.9704
---	---	1629861_at	0.1742	0.6043	0.0435	0.8190	0.3575	0.0500	0.1519	0.6257	0.0204	0.9243	-0.1314	0.3528	-0.1361	0.9015	-0.1195	0.8203	0.0166	0.9803
CG11815	CG11815	1629862_at	-0.1425	0.5092	0.0847	0.5869	-0.0787	0.7115	0.1017	0.8535	0.0366	0.8924	-0.0651	0.7663	0.1431	0.8069	0.1875	0.4421	0.0444	0.8949
Cad89D	Cad89D	1629863_at	-0.1108	0.6858	0.0650	0.5729	-0.0803	0.6342	0.0558	0.9116	0.0724	0.6913	0.0166	0.9327	0.1034	0.9011	0.1895	0.5523	0.0862	0.8209
CG2685	CG2685	1629864_at	-0.4306	0.2390	-0.0764	0.8687	0.4001	0.0606	0.3110	0.2466	-0.0823	0.6453	-0.3932	0.0135	-0.2220	0.8439	0.4018	0.3749	0.6238	0.1995
ric8a	ric8a	1629865_s_at	0.3495	0.0675	1.0212	0.0065	0.9186	0.0059	-0.0409	0.9777	-0.4110	0.2738	-0.3701	0.2728	0.1817	0.7230	0.3249	0.1389	0.1432	0.5391
Suv4-20	Suv4-20	1629866_at	-0.0367	0.9415	-0.0950	0.8095	-0.1112	0.5276	0.0103	0.9940	0.2662	0.3562	0.2559	0.3217	0.0720	0.9741	0.2050	0.7504	0.1330	0.8499
RpS29	Ribosomal protein	1629867_a_at	0.2968	0.1775	1.4487	0.0042	1.3181	0.0004	0.0651	0.9465	-0.8282	0.0090	-0.8933	0.0038	0.1012	0.8700	0.0707	0.8287	-0.0305	0.9295
CG3838	CG3838	1629868_s_at	0.3988	0.1250	1.4301	0.0057	1.3317	0.0002	-0.0515	0.9295	-1.0366	0.0005	-0.9851	0.0004	-0.0345	0.9831	0.0728	0.8954	0.1073	0.8167
Fatp	Fatty acid (long cl	1629869_s_at	0.3356	0.1047	-0.1180	0.5790	0.3347	0.1176	0.3834	0.3631	0.2255	0.3574	-0.1579	0.4932	-0.0056	0.9964	-0.2853	0.3404	-0.2797	0.3756
CG3687	CG3687	1629870_at	0.1022	0.5599	0.0824	0.5230	0.0271	0.8932	-0.0994	0.8761	-0.1256	0.6025	-0.0261	0.9230	0.0771	0.8558	-0.0489	0.8333	-0.1260	0.4846
CG11665 /// DyakCG11665	CG11665	1629871_at	-0.1890	0.4157	0.1472	0.7288	0.1122	0.4848	-0.1775	0.5944	-0.3850	0.0301	-0.2076	0.1684	-0.2053	0.8609	-0.0930	0.8953	0.1123	0.8552
---	---	1629872_at	0.1904	0.5228	0.2020	0.3803	-0.1452	0.5199	-0.1460	0.8856	0.1198	0.7724	0.2659	0.4078	0.1574	0.8270	0.1139	0.7472	-0.0435	0.9168
---	---	1629873_at	0.4379	0.0302	0.2647	0.2084	0.0285	0.8654	-0.1155	0.7941	0.0005	0.9985	0.1160	0.5053	0.1818	0.7485	-0.0708	0.8220	-0.2525	0.3056
Kif3C	Kif3C	1629874_at	0.1469	0.4215	-0.0606	0.5990	0.0411	0.8008	0.1782	0.6202	0.2488	0.1628	0.0706	0.7105	-0.0647	0.9011	-0.0362	0.9003	0.0285	0.9137
Cyp4d1	Cytochrome P450	1629875_a_at	-0.1856	0.4822	0.0512	0.8580	0.0991	0.6608	0.0476	0.9602	-0.1573	0.5609	-0.2049	0.3774	-0.0727	0.9549	0.0585	0.9204	0.1312	0.7641
Cks85A	Cyclin-dependent	1629876_at	-0.0939	0.7507	-0.1167	0.4737	-0.1016	0.6983	-0.2964	0.4612	-0.3363	0.1214	-0.0399	0.8776	-0.1036	0.9400	-0.0563	0.9341	0.0474	0.9362
CG31683	CG31683	1629877_at	0.0495	0.9364	0.1756	0.6356	0.3118	0.1789	-0.2320	0.8471	-0.7898	0.0758	-0.5578	0.1561	-0.3773	0.7628	-0.6449	0.2012	-0.2677	0.6328
---	---	1629878_at	-0.0694	0.7096	-0.0527	0.7968	0.0247	0.8917	0.0298	0.9592	0.0930	0.5775	0.0631	0.6964	-0.0018	0.9994	0.1317	0.6512	0.1335	0.6428
CG13434	CG13434	1629879_at	0.3568	0.3346	-0.3762	0.1184	-0.1691	0.2771	0.1459	0.7753	0.1720	0.4383	0.0261	0.9209	0.0658	0.9775	-0.4333	0.4355	-0.4991	0.3810
grp	grapes	1629880_at	0.4466	0.1416	-0.4425	0.5109	-0.5616	0.1288	0.2382	0.5802	0.8650	0.0028	0.6268	0.0071	-0.1023	0.9762	-0.1120	0.9325	-0.0098	0.9946
Nxt1	NTF2-related expo	1629881_at	0.1530	0.6061	-0.0409	0.8553	0.1593	0.5138	0.1007	0.8923	0.2570	0.3068	0.1564	0.5170	-0.0949	0.9499	0.0183	0.9829	0.1132	0.8444
CG6683	CG6683	1629882_at	-0.0358	0.8946	0.0821	0.8459	-0.2019	0.3062	-0.1818	0.8507	-0.0627	0.8957	0.1191	0.7564	0.1673	0.7588	0.1142	0.6458	-0.0531	0.8599
---	---	1629883_at	0.1028	0.6388	0.2878	0.3071	0.2195	0.3156	-0.1960	0.7937	-0.1830	0.5818	0.0129	0.9729	-0.0537	0.9727	-0.1893	0.6605	-0.1356	0.7677
CG34125	CG34125	1629884_at	0.0496	0.8295	-0.4243	0.1564	-0.2561	0.1779	0.2377	0.4386	0.4173	0.0212	0.1796	0.2322	0.0273	0.9837	-0.0183	0.9718	-0.0455	0.9151
CG30174	CG30174	1629885_at	-0.1443	0.4276	-0.0701	0.6594	-0.0520	0.7496	-0.1404	0.6817	-0.3048	0.0596	-0.1644	0.2502	-0.1289	0.8049	-0.1660	0.4479	-0.0371	0.9025
for	foraging	1629886_s_at	0.7963	0.0069	0.6239	0.1332	0.8333	0.0010	0.0018	0.9983	-0.3731	0.0535	-0.3749	0.0339	-0.2506	0.8202	-0.5795	0.1894	-0.3289	0.4851
CG8180	CG8180	1629887_at	-1.0678	0.0328	-0.3154	0.5220	-1.0570	0.0448	-0.2658	0.5323	-0.2064	0.3629	0.0593	0.8130	0.0629	0.9898	0.1141	0.9467	0.0513	0.9758
CG6220	CG6220	1629888_at	-0.1198	0.6386	0.3315	0.0792	0.3273	0.0785	-0.1504	0.7815	-0.3379	0.1313	-0.1875	0.3657	-0.2971	0.5754	0.0273	0.9438	0.3244	0.1990
regucalcin	regucalcin	1629889_s_at	-1.8110	0.0131	-2.9401	0.0033	-2.5237	0.0000	0.0477	0.9513	-0.2903	0.1754	-0.3380	0.0806	-0.2742	0.8999	-1.3212	0.1068	-1.0470	0.2110
CG6262	CG6262	1629890_a_at	0.2895	0.1818	-0.0044	0.9699	0.1280	0.4246	-0.0504	0.9293	0.0492	0.8108	0.0995	0.5401	-0.2563	0.6832	-0.1711	0.5246	0.0852	0.7813
---	---	1629891_at	0.1729	0.3035	0.0545	0.7793	0.0489	0.8146	0.0119	0.9909	0.1296	0.6024	0.1177	0.6079	-0.0747	0.9126	-0.0814	0.7837	-0.0067	0.9859
CG34377	CG13859	1629892_at	0.0606	0.8339	0.0950	0.4654	0.1788	0.3839	-0.0265	0.9798	-0.0467	0.8920	-0.0203	0.9498	-0.1065	0.8882	-0.0631	0.8778	0.0434	0.9121
---	---	1629893_s_at	0.0176	0.9215	-0.2147	0.1937	-0.2330	0.2174	0.0778	0.8967	0.3719	0.0635	0.2941	0.0957	0.0001	0.9999	0.1771	0.4177	0.1770	0.4322
CG3044	CG3044	1629894_at	-0.6036	0.0234	-0.3230	0.2799	-0.3366	0.0479	-0.1430	0.7229	-0.4098	0.0316	-0.2667	0.1006	-0.2057	0.7843	-0.1628	0.6425	0.0429	0.9231
---	---	1629895_at	-0.0083	0.9629	-0.0641	0.6935	0.1178	0.4686	0.0000	0.9999	-0.0846	0.6322	-0.0845	0.5942	-0.1225	0.8379	0.0479	0.8967	0.1704	0.5050
CG32845	CG32845	1629896_at	0.1216	0.3758	-0.4651	0.0711	0.0412	0.8540	0.3935	0.2101	0.5109	0.0147	0.1174	0.5224	-0.3363	0.6509	-0.1361	0.6842	0.2002	0.5249
CG6044	CG6044	1629897_a_at	-0.4765	0.1728	-0.2681	0.0481	-1.0397	0.0013	-0.2644	0.6591	-0.1723	0.5818	0.0921	0.7723	-0.1158	0.8553	-0.0786	0.8165	0.0371	0.9176
CG10737	Gene 1	1629898_s_at	-1.7095	0.0020	-1.1318	0.0834	-1.2099	0.0005	0.0330	0.9776	-0.4776	0.0970	-0.5106	0.0516	0.1549	0.9246	0.1015	0.9036	-0.0534	0.9440
cact	cactus	1629899_at	-1.0941	0.0016	-0.5773	0.0506	-1.7215	0.0001	-0.3383	0.4420	0.1696	0.5034	0.5080	0.0261	0.7399	0.2884	0.7133	0.0648	-0.0266	0.9588
---	---	1629900_at	0.0691	0.7529	-0.1078	0.3556	0.0374	0.8785	0.1499	0.6678	0.0614	0.7578	-0.0885	0.5956	-0.0945	0.8956	-0.0731	0.8394	0.0214	0.9551
CG11881	CG11881	1629901_at	-0.1439	0.4042	-0.2332	0.2317	-0.5205	0.0272	-0.2328	0.6919	0.2600	0.3473	0.4928	0.0478	-0.0353	0.9751	0.0144	0.9766	0.0497	0.8972
Cap-G	lethal (2) c00093	1629902_at	-0.0819	0.8175	0.2248	0.1208	0.4344	0.0414	0.1058	0.8667	-0.3583	0.1060	-0.4641	0.0274	-0.0271	0.9852	0.0870	0.8481	0.1142	0.7709
CG30359	CG30359	1629903_at	-0.2093	0.3309	-0.8050	0.1656	-0.6797	0.0398	-0.0806	0.9303	-0.6864	0.0193	-0.6058	0.0204	-0.1435	0.9460	-1.3900	0.0519	-1.2465	0.0878
fru	fru-satori	1629904_at	-0.0776	0.7219	-0.2188	0.4031	-0.1265	0.5126	-0.0486	0.9539	0.1272	0.6176	0.1758	0.4145	-0.0754	0.9246	-0.0470	0.9085	0.0284	0.9397
---	---	1629905_s_at	0.8857	0.1819	0.7519	0.1715	0.0208	0.9764	-0.2260	0.5407	0.2217	0.2503	0.4477	0.0173	0.4769	0.8461	0.0724	0.9650	-0.4045	0.7253
---	---	1629906_s_at	0.5495	0.1542	0.3401	0.6293	1.3381	0.0007	0.4746	0.1358	-0.0746	0.7424	-0.5492	0.0071	-0.4805	0.8192	-0.1169	0.9325	0.3636	0.7071
CG14989	CG14989	1629907_at	0.2401	0.3613	0.0224	0.9022	0.0968	0.5136	0.0567	0.9380	0.0038	0.9901	-0.0529	0.8279	-0.1040	0.8875	-0.1497	0.6122	-0.0457	0.9053
---	---	1629908_at	0.0073	0.9706	0.1206	0.3906	0.2499	0.2097	-0.2247	0.5445	-0.2123	0.2716	0.0124	0.9593	-0.1833	0.7794	0.1124	0.7340	0.2958	0.3081
---	---	1629909_at	0.0833	0.6384	0.1331	0.2828	0.0777	0.6951	-0.0150	0.9811	-0.0236	0.9116	-0.0086	0.9648	0.0783	0.9076	0.0188	0.9621	-0.0596	0.8467
CG11570	CG11570	1629910_at	0.2790	0.1377	-0.1332	0.3104	-0.1328	0.4694	0.0873	0.8294	0.2664	0.0875	0.1791	0.1967	-0.0044	0.9967	-0.1192	0.6952	-0.1148	0.7053
CG3222	CG3222	1629911_at	-0.1818	0.4183	0.0713	0.5382	-0.2830	0.2077	-0.1256	0.7857	0.0069	0.9809	0.1325	0.4640	0.1327	0.8386	0.1706	0.5422	0.0379	0.9187
---	---	1629912_at	0.0042	0.9865	-0.0108	0.9181	0.1507	0.4991	0.0726	0.8605	0.0793	0.6304	0.0068	0.9710	-0.0365	0.9677	-0.0089	0.9842	0.0276	0.9338
---	---	1629913_at	0.0489	0.7298	0.0128	0.9329	-0.0955	0.6046	0.0438	0.9528	0.0948	0.6844	0.0510	0.8263	-0.0359	0.9555	-0.0209	0.9451	0.0150	0.9549
CG14328	CG14328	1629914_at	-0.0645	0.8162	-0.0294	0.8387	-0.0725	0.7010	0.0854	0.9029	0.1561	0.5169	0.0707	0.7796	0.0453	0.9499	0.1522	0.4875	0.1069	0.6428
---	---	1629915_at	0.1116	0.6476	0.0098	0.9493	0.2430	0.1966	0.1126	0.8817	0.0337	0.9257	-0.0789	0.7853	-0.1752	0.7436	0.0049	0.9925	0.1800	0.4374
debcl	deborg	1629916_at	-1.1774	0.0012	-0.7528	0.0132	-0.2857	0.3274	0.1214	0.7293	-0.1701	0.2816	-0.2916	0.0438	-0.2132	0.8513	0.2722	0.5855	0.4854	0.3139
---	---	1629917_at	0.0303	0.8939	0.1246	0.2397	0.2183	0.1644	-0.0907	0.8705	-0.1021	0.6396	-0.0114	0.9630	0.0309	0.9792	0.0102	0.9842	-0.0206	0.9582
CG6197	CG6197	1629918_at	-0.0987	0.4918	0.7106	0.0321	0.8905	0.0030	0.1530	0.7018	-0.3001	0.1017	-0.4531	0.0136	-0.0819	0.9457	0.5492	0.1321	0.6311	0.1179
Cpr100A	CG12045	1629919_at	0.5918	0.2295	0.0478	0.8199	0.4072	0.1233	0.1748	0.8096	0.0702	0.8513	-0.1046	0.7361	0.0368	0.9885	-0.2656	0.6634	-0.3024	0.6134
Kap-alpha1	karyopherin-alpha	1629920_at	0.1978	0.2185	0.5513	0.1041	0.4353	0.0181	-0.0809	0.8873	-0.2248	0.2444	-0.1439	0.4258	0.0202	0.9829	0.1775	0.4068	0.1573	0.4822
Su(H)	suppressor of hair	1629921_a_at	0.8278	0.0242	0.2706	0.2293	0.0707	0.7206	-0.2378	0.5850	0.5973	0.0151	0.8351	0.0020	0.1408	0.8963	0.2367	0.5800	0.0959	0.8548
CG11409 /// DmirCG11409	CG11409	1629922_at	-4.4145	0.0005	-4.7933	0.0016	-5.1730	0.0000	0.3106	0.7658	0.3033	0.5216	-0.0073	0.9897	0.2654	0.7743	0.0280	0.9661	-0.2374	0.5786
CG7371	CG7371	1629923_at	-0.5301	0.0096	-0.4509	0.1387	-0.2967	0.0522	0.1292	0.7349	0.1306	0.4602	0.0013	0.9953	0.0167	0.9898	0.2641	0.3349	0.2474	0.3893
CG10238	CG10238	1629924_at	-0.5399	0.3094	-0.3294	0.1611	-0.9846	0.0007	-0.4194	0.1583	0.3049	0.0911	0.7243	0.0015	0.0942	0.9691	0.4553	0.4848	0.3611	0.5988
CG8007	CG8007	1629925_a_at	0.3130	0.1954	0.2881	0.0699	0.4382	0.0245	0.0931	0.8546	0.1537	0.4185	0.0606	0.7639	-0.0010	0.9994	0.0235	0.9451	0.0245	0.9342
CG1342	CG1342	1629926_at	-0.0727	0.7269	0.2649	0.1463	0.1000	0.5429	-0.0854	0.8942	-0.2578	0.2278	-0.1724	0.3852	0.0398	0.9683	0.0714	0.8402	0.0316	0.9321
alc	beta homologue	1629927_at	0.0868	0.7497	-0.0599	0.8089	-0.0512	0.8495	-0.2292	0.6506	-0.2570	0.2978	-0.0278	0.9260	-0.1508	0.8973	-0.1933	0.6939	-0.0425	0.9440

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG32141	CG32141	1629928_a_at	-0.0475	0.8068	0.1872	0.3756	0.0321	0.8909	-0.0909	0.8619	0.0312	0.9038	0.1221	0.4960	0.0924	0.8943	0.2086	0.4229	0.1162	0.6877
---	---	1629929_at	-0.0676	0.7700	-0.1384	0.4102	-0.0237	0.9189	-0.0908	0.9066	-0.1505	0.5769	-0.0597	0.8340	-0.0753	0.9168	0.0011	0.9994	0.0763	0.8015
Cct1	Phosphocholine c	1629930_s_at	-0.5698	0.4393	0.0734	0.9472	0.4047	0.0609	0.7377	0.3772	0.5775	0.2202	-0.1602	0.7564	0.4719	0.8692	0.9354	0.4093	0.4635	0.7205
---	---	1629931_at	0.1174	0.4218	0.2100	0.2158	0.1196	0.6114	0.1083	0.8856	0.0160	0.9678	-0.0923	0.7371	0.0831	0.9238	0.0986	0.7787	0.0156	0.9717
yellow-d2	yellow-d2	1629932_at	1.2938	0.0110	1.1577	0.0270	1.6577	0.0002	-0.0580	0.9218	-0.3231	0.0732	-0.2651	0.0965	-0.3209	0.8016	-0.2365	0.6929	0.0844	0.9088
Sas10	Sas10	1629933_at	0.7140	0.0291	0.3468	0.4962	0.5897	0.0155	0.3726	0.4861	1.0450	0.0039	0.6724	0.0166	0.0372	0.9884	0.6696	0.2155	0.6324	0.2738
CG14219	CG14219	1629934_at	0.1371	0.3971	0.2071	0.1813	0.3345	0.0605	-0.0104	0.9895	-0.1561	0.4059	-0.1457	0.3909	0.0088	0.9952	0.1802	0.6392	0.1714	0.6545
CG13312	CG13312	1629935_at	-0.0279	0.8836	0.2275	0.2634	0.1786	0.2891	0.0061	0.9937	-0.0919	0.6019	-0.0980	0.5325	-0.0632	0.9238	0.0300	0.9325	0.0932	0.7072
mRpL16	mitochondrial ribo	1629936_at	-0.1122	0.7009	0.1339	0.4183	0.3404	0.2484	0.2085	0.5836	-0.3229	0.0957	-0.5315	0.0084	0.0905	0.9499	0.1423	0.7910	0.0518	0.9316
CG34379	CG13942	1629937_a_at	0.1646	0.8661	0.8197	0.4786	1.8963	0.0001	0.6606	0.1317	-0.3431	0.1899	-1.0037	0.0019	-0.3118	0.9499	0.3123	0.8838	0.6241	0.7071
Ctr1A	Copper transporte	1629938_at	-1.1267	0.0088	-0.6965	0.0888	-0.6286	0.0054	-0.0152	0.9893	-0.2618	0.3201	-0.2467	0.2954	-0.0544	0.9781	0.1465	0.8119	0.2010	0.7060
CG1140	citric acid synthet	1629939_a_at	0.1875	0.3375	1.5535	0.0065	1.8703	0.0000	0.2693	0.4927	-0.9154	0.0018	-1.1847	0.0004	-0.0107	0.9914	0.3246	0.1391	0.3353	0.1548
CG13895	CG13895	1629940_at	-0.4758	0.0267	-1.0406	0.0208	-1.1843	0.0003	-0.0362	0.9592	0.4569	0.0226	0.4931	0.0100	0.1126	0.9142	-0.1365	0.7533	-0.2491	0.5137
CG8044	CG8044	1629941_a_at	-0.1161	0.5598	-0.1703	0.6036	-0.2001	0.3581	0.0280	0.9744	0.3170	0.1289	0.2890	0.1212	0.0049	0.9978	0.3063	0.4691	0.3014	0.4851
CG5292	CG5292	1629942_at	0.2491	0.4453	0.8187	0.0153	0.7246	0.0044	-0.1876	0.5539	-0.1574	0.3453	0.0302	0.8767	-0.0826	0.9611	0.4687	0.3087	0.5513	0.2611
CG7112	CG7112	1629943_at	-0.1235	0.7684	0.0894	0.8580	0.6892	0.0015	0.3513	0.5174	-0.2428	0.4101	-0.5941	0.0280	-0.3281	0.8202	-0.0462	0.9628	0.2819	0.6644
CG12814	CG12814	1629944_at	0.0847	0.8090	-0.1617	0.8098	-0.2805	0.1078	0.3209	0.5375	0.4916	0.0729	0.1708	0.5164	0.5668	0.7142	0.3256	0.6417	-0.2412	0.7456
CG32341	CG32341	1629945_at	0.4057	0.1746	-0.0332	0.8800	0.0766	0.7618	0.1000	0.8863	0.3910	0.0986	0.2910	0.1680	-0.2363	0.7336	-0.1888	0.5425	0.0475	0.9080
CG32118	CG32118	1629946_at	0.0096	0.9732	0.0503	0.6348	0.1984	0.2397	-0.1414	0.7221	-0.0695	0.7425	0.0719	0.7036	0.0366	0.9623	0.1124	0.6288	0.0757	0.7619
CG32812	CG32812	1629947_at	-0.4731	0.0320	-0.0533	0.6367	-0.2213	0.3756	0.0774	0.9247	-0.2182	0.3838	-0.2955	0.1760	0.1788	0.7633	0.0242	0.9516	-0.1545	0.5585
CG5194	CG5194	1629948_at	0.4236	0.3858	-0.0757	0.8437	-0.2050	0.5955	-0.3147	0.2686	0.4598	0.0140	0.7745	0.0009	-0.0668	0.9848	0.1460	0.9064	0.2128	0.8354
CG14647	CG14647	1629949_at	-0.2865	0.4062	-0.7679	0.0435	-0.5422	0.0320	-0.3342	0.3571	-0.0051	0.9875	0.3291	0.0772	-0.4768	0.6955	-0.3820	0.4523	0.0948	0.8918
CG1785	CG1785	1629950_at	0.5581	0.0769	0.2188	0.6660	0.3920	0.1151	0.2416	0.7327	0.6575	0.0434	0.4159	0.1390	0.0593	0.9751	0.3393	0.4785	0.2801	0.5744
CG9863	CG9863	1629951_at	0.1172	0.4800	0.0207	0.9229	0.1853	0.3804	0.2374	0.6096	0.1567	0.5288	-0.0807	0.7507	0.0200	0.9870	0.0657	0.8678	0.0458	0.9057
CG15227	CG15227	1629952_at	-0.0845	0.6842	-0.0218	0.9044	-0.0002	0.9991	-0.0157	0.9825	-0.0137	0.9560	0.0020	0.9923	-0.0492	0.9742	-0.1085	0.8255	-0.0593	0.9089
gt	giant	1629953_at	0.0112	0.9667	-0.1354	0.4276	-0.1036	0.6553	-0.0352	0.9666	0.1846	0.4164	0.2197	0.2679	-0.2476	0.7436	-0.1935	0.5635	0.0541	0.9025
CG10153	CG10153	1629954_at	-0.0543	0.8494	0.0838	0.7511	-0.1191	0.4548	-0.0063	0.9950	-0.0821	0.7311	-0.0757	0.7281	0.1401	0.8825	0.0849	0.8679	-0.0552	0.9121
CG13309 /// DyakCG1330	CG13309	1629955_at	-0.2255	0.7229	0.1267	0.5368	0.2114	0.1497	0.0588	0.9761	-0.6364	0.1900	-0.6952	0.1102	0.1756	0.8270	-0.2207	0.5282	-0.3962	0.2558
CG10098	CG10098	1629956_at	-1.0479	0.0280	-0.7038	0.0681	-0.5272	0.1174	0.4630	0.3301	0.6220	0.0312	0.1590	0.5586	0.1942	0.9142	1.0072	0.1151	0.8129	0.2136
---	---	1629957_at	0.2310	0.1541	0.3124	0.2215	0.2235	0.1764	-0.0041	0.9956	-0.0247	0.9045	-0.0206	0.9095	0.0245	0.9816	0.1188	0.6658	0.0943	0.7434
CG5316	CG5316	1629958_at	0.0785	0.7119	0.1681	0.1584	-0.0415	0.8277	-0.0026	0.9970	0.0174	0.9547	0.0200	0.9376	0.0203	0.9787	-0.0212	0.9407	-0.0416	0.8577
CG32569	CG32569	1629959_at	0.0921	0.6371	0.1361	0.3486	0.2631	0.2000	0.0123	0.9847	-0.0321	0.8675	-0.0444	0.7854	-0.1807	0.8236	-0.0102	0.9886	0.1705	0.6395
Rep3	Rep3	1629960_s_at	-0.0368	0.8275	0.1566	0.3377	0.0752	0.7453	-0.0825	0.8834	-0.0370	0.8858	0.0455	0.8358	0.1776	0.7707	0.1111	0.7058	-0.0664	0.8418
l(2)44DEa	Fatty acyl CoA syr	1629961_s_at	1.0298	0.0157	0.6458	0.1275	-0.1075	0.7712	0.0570	0.9228	0.2463	0.1625	0.1893	0.2328	0.9150	0.5126	-0.0202	0.9885	-0.9352	0.1829
CG8916	CG8916	1629962_at	-0.0349	0.8715	0.0431	0.8229	0.0978	0.6009	0.0386	0.9506	0.1482	0.4027	0.1095	0.5103	-0.0443	0.9713	0.2268	0.4667	0.2711	0.3880
sut1	sugar transporter	1629963_at	1.4957	0.0101	0.0583	0.8447	0.4757	0.0791	-0.0128	0.9937	0.5688	0.0725	0.5816	0.0440	-0.5558	0.6763	-0.8647	0.1233	-0.3088	0.6052
Nufip	Nufip	1629964_at	0.1256	0.6272	-0.0806	0.8607	-0.4503	0.0237	-0.2781	0.5330	0.3093	0.1816	0.5874	0.0124	0.1425	0.8930	0.1371	0.7779	-0.0054	0.9940
CG8262	CG8262	1629965_at	-0.0786	0.8231	0.1321	0.5661	-0.3200	0.1663	-0.1292	0.7929	-0.0527	0.8382	0.0765	0.7200	0.1006	0.9330	-0.0767	0.8919	-0.1774	0.6724
E(spl)	Enhancer of split r	1629966_at	0.0569	0.8587	0.1089	0.5256	0.1155	0.4791	0.1186	0.8956	-0.1067	0.7665	-0.2253	0.4219	0.1361	0.7768	0.0455	0.8786	-0.0906	0.6963
park	dParkin	1629967_s_at	-0.4357	0.0310	-0.0328	0.9069	-0.0252	0.8815	0.0465	0.9325	-0.4351	0.0157	-0.4816	0.0060	-0.0566	0.9526	-0.0929	0.7963	-0.0364	0.9264
CG17370	CG17370	1629968_s_at	0.2764	0.2843	0.3031	0.2260	0.2744	0.1083	-0.1365	0.7266	0.0346	0.8828	0.1711	0.2789	-0.0157	0.9928	0.1405	0.7694	0.1562	0.7287
CG11426	CG11426	1629969_at	0.7508	0.0869	2.8339	0.0283	3.7188	0.0000	1.6292	0.0049	1.3923	0.0009	-0.2370	0.3329	0.9679	0.6557	3.5759	0.0105	2.6080	0.0340
eIF-4E	eukaryotic initiatio	1629970_s_at	0.2711	0.1476	0.1518	0.5119	-0.1057	0.5888	-0.0544	0.9120	0.2257	0.1476	0.2801	0.0503	0.2277	0.7893	0.1406	0.7424	-0.0871	0.8534
---	---	1629971_at	0.0142	0.9413	0.3074	0.2408	-0.0129	0.9632	-0.1244	0.7243	-0.1632	0.3088	-0.0388	0.8300	0.0486	0.9742	0.0073	0.9935	-0.0413	0.9371
CG8188	CG8188	1629972_at	0.6899	0.0778	0.7801	0.0750	0.5701	0.0552	0.1037	0.8897	0.2845	0.2570	0.1807	0.4424	0.3422	0.8427	0.4239	0.5699	0.0817	0.9344
CG6218	CG6218	1629973_at	0.8274	0.0938	0.8329	0.0683	0.7909	0.0008	0.0936	0.9451	0.5156	0.1802	0.4220	0.2232	0.1937	0.8461	0.4896	0.2204	0.2959	0.4889
Taf2	TBP-associated fa	1629974_at	0.5614	0.0179	0.1232	0.6186	-0.2620	0.1623	-0.2868	0.5332	0.6901	0.0113	0.9769	0.0015	0.2392	0.7783	0.3946	0.2688	0.1554	0.7087
CG7406	CG7406	1629975_at	-1.4171	0.0147	-0.7062	0.2391	-0.9855	0.0025	-0.4103	0.5833	-0.5463	0.1497	-0.1360	0.7403	-0.1891	0.8270	-0.0481	0.9341	0.1410	0.7338
CG31424	CG31424	1629976_at	0.1928	0.2315	0.0068	0.9837	-0.0464	0.8292	0.0774	0.8939	0.1881	0.3370	0.1107	0.5614	0.0354	0.9816	0.0366	0.9500	0.0012	0.9987
CG9921	CG9921	1629977_at	-0.4139	0.2268	-0.1098	0.7007	0.1214	0.5643	-0.1566	0.6178	-0.8324	0.0008	-0.6758	0.0011	-0.3083	0.7893	-0.4181	0.3885	-0.1098	0.8665
CG12250	CG12250	1629978_at	0.0278	0.8827	-0.2017	0.4374	-0.2873	0.1498	0.0652	0.9090	0.3483	0.0594	0.2831	0.0826	0.0596	0.9589	-0.0876	0.8444	-0.1472	0.6861
---	---	1629979_at	0.1825	0.4173	0.0547	0.7968	0.1810	0.3300	0.1492	0.6970	0.0541	0.8026	-0.0951	0.5900	-0.0617	0.9717	-0.0676	0.9221	-0.0059	0.9941
---	---	1629980_at	0.1311	0.5652	0.0021	0.9877	-0.0427	0.7972	0.1240	0.8216	0.0695	0.7912	-0.0546	0.8227	0.0463	0.9589	-0.0022	0.9978	-0.0484	0.8896
LamC	lamin C	1629981_at	-0.7541	0.1944	-0.8840	0.0572	-0.9404	0.0006	0.4340	0.5680	0.6950	0.0763	0.2610	0.4821	0.5187	0.6955	0.5826	0.2829	0.0638	0.9350
---	---	1629982_at	0.1562	0.3888	0.1728	0.2713	0.1116	0.4889	0.0051	0.9956	0.0188	0.9492	0.0136	0.9568	-0.0453	0.9588	-0.0006	0.9996	0.0447	0.8938
CG1578	CG1578	1629983_at	-0.2428	0.6258	-0.2629	0.7302	-0.3813	0.0460	-0.0697	0.9149	-0.4170	0.0468	-0.3473	0.0611	0.0874	0.9816	-0.3993	0.6943	-0.4867	0.6166
RpL38	Enhancer on chro	1629984_s_at	0.0429	0.8054	0.5018	0.0348	0.5544	0.0089	0.0053	0.9943	-0.3541	0.0362	-0.3594	0.0216	-0.0092	0.9928	0.1148	0.6512	0.1240	0.6225
CG15614	CG15614	1629985_at	0.0888	0.6857	0.2396	0.2537	-0.0781	0.6531	0.1242	0.8190	0.0677	0.7951	-0.0565	0.8135	0.2204	0.7387	0.1315	0.6713	-0.0889	0.7918
---	---	1629986_at	0.3529	0.1030	0.1472	0.4119	0.1890	0.2044	0.0266	0.9774	-0.0141	0.9686	-0.0407	0.8807	-0.0553	0.9331	-0.0526	0.8561	0.0027	0.9945
---	---	1629987_s_at	-0.0107	0.9621	-0.0243	0.8531	-0.0829	0.7327	0.1165	0.8908	0.0601	0.8736	-0.0564	0.8656	0.0688	0.9467	0.0546	0.9085	-0.0142	0.9769
CG34380	CG13988	1629988_at	0.0621	0.7513	0.0040	0.9825	-0.0671	0.7689	-0.0379	0.9506	0.0248	0.9166	0.0627	0.7278	-0.0245	0.9862	-0.0372	0.9450	-0.0128	0.9821
---	---	1629989_at	-0.0631	0.7720	-0.0789	0.5514	0.0838	0.6245	0.0074	0.9943	0.0006	0.9984	-0.0067	0.9799	-0.0453	0.9514	-0.0389	0.9085	0.0064	0.9856
CG14040	CG14040	1629990_at	-0.1532	0.6635	0.8374	0.1128	0.8488	0.0018	-0.1678	0.6122	-0.5652	0.0053	-0.3974	0.0155	-0.1945	0.9016	0.3856	0.5184	0.5801	0.3256
CG31028	CG31028	1629991_a_at	-0.3400	0.2585	0.1414	0.4224	-0.1342	0.3761	-0.2358	0.6480	-0.5700	0.0300	-0.3343	0.1327	0.2336	0.7726	0.0898	0.8504	-0.1438	0.7154
---	---	1629992_at	0.0012	0.9954	0.0304	0.8399	-0.0023	0.9924	-0.1955	0.6122	-0.0420	0.8696	0.1534	0.3930	-0.0948	0.8945	0.0211	0.9620	0.1159	0.6981
CG41056	CG41056	1629993_at	0.0484	0.7953	-0.0805	0.4284	0.1101	0.6424	0.1542	0.6823	0.2199	0.2082	0.0657	0.7255	-0.1098	0.9038	-0.0576	0.9088	0.0523	0.9088
CG14521	CG14521	1629994_at	-0.1801	0.4074	0.0701	0.6430	0.2044	0.3509	0.0420	0.9540	0.0305	0.9130	-0.0115	0.9635	-0.2020	0.8049	0.0088	0.9911	0.2108	0.5612
CG32104	CG32104	1629995_at	0.6652	0.0201	0.4166	0.3011	0.4801	0.0275	-0.0118	0.9921	0.4592	0.0611	0.4710	0.0360	-0.0111	0.9950	0.2110	0.6391	0.2221	0.6189
CG11910	CG11910	1629996_at	0.0063	0.9812	0.0860	0.7213	-0.0278	0.8951	-0.1014	0.9116	-0.1344	0.6844	-0.0330	0.9257	0.0455	0.9653	0.0837	0.8180	0.0381	0.9211

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13858	CG13858	1629997_at	0.2748	0.3207	0.0719	0.6814	0.2214	0.3833	0.1283	0.8192	0.0736	0.7828	-0.0547	0.8276	0.1529	0.8878	-0.1853	0.6827	-0.3381	0.4119
---	---	1629998_at	0.5557	0.0566	0.2946	0.3202	0.2783	0.3565	0.0068	0.9956	-0.0643	0.8736	-0.0711	0.8403	-0.1897	0.8845	-0.3038	0.5576	-0.1141	0.8602
CG15770	CG15770	1629999_at	-0.7168	0.0084	-0.3625	0.2695	-0.3920	0.0927	0.1539	0.7937	-0.2793	0.2458	-0.4332	0.0488	-0.0621	0.9421	-0.3671	0.1524	-0.3050	0.2659
CG7358 /// DrepCG7358	CG7358	1630000_at	-0.6604	0.4969	0.3307	0.8341	0.1767	0.6458	0.1277	0.8564	-0.2530	0.3257	-0.3807	0.0932	0.4065	0.9492	0.7442	0.7383	0.3377	0.8976
CG12535	CG12535	1630001_at	0.2713	0.1302	0.0572	0.7655	0.0660	0.7050	-0.1682	0.7461	0.0900	0.7372	0.2581	0.2038	-0.0851	0.8861	-0.0965	0.7020	-0.0115	0.9739
Hr39	Hormone receptor	1630002_s_at	2.7685	0.0006	1.6904	0.0100	1.9032	0.0003	-0.0853	0.9500	0.4832	0.1984	0.5685	0.0920	-0.1794	0.8874	-0.4868	0.2912	-0.3074	0.5391
CG12825	CG12825	1630003_at	-0.4525	0.6606	-0.1281	0.7957	-1.0508	0.0006	-0.7974	0.4102	-0.8905	0.0975	-0.0931	0.8836	0.1369	0.9778	-0.6340	0.6146	-0.7709	0.5334
beat-Ib	beaten path Ib	1630004_at	-0.1466	0.4728	-0.0117	0.9125	0.1352	0.6778	-0.1073	0.8102	0.0128	0.9616	0.1201	0.4774	-0.2874	0.7423	0.1408	0.7455	0.4283	0.2615
Ubi-p5E	Ubiquitin-5E	1630005_at	-0.0119	0.9631	0.0218	0.9158	-0.1922	0.1955	-0.0732	0.9029	0.0115	0.9696	0.0847	0.6769	0.1557	0.7633	0.0832	0.7439	-0.0725	0.7764
CG5278	CG5278	1630006_at	0.1921	0.1904	-0.0403	0.7242	0.0638	0.7509	0.1809	0.6330	0.1846	0.3272	0.0037	0.9874	-0.1476	0.7997	-0.1176	0.6587	0.0299	0.9295
Plc21C	Phospholipase C	1630007_s_at	-0.2941	0.2354	-0.4838	0.1388	-1.1899	0.0010	-0.6264	0.1119	-0.0263	0.9380	0.6002	0.0118	0.1326	0.9063	-0.2852	0.4898	-0.4178	0.3101
CG18428	CG18428	1630008_at	0.1284	0.4115	0.4947	0.1132	0.4492	0.0603	0.0078	0.9922	-0.1766	0.2944	-0.1844	0.2172	0.0301	0.9848	0.1604	0.6912	0.1303	0.7527
CG12209	CG12209	1630009_at	-0.0646	0.7139	-0.0092	0.9828	0.1014	0.8143	0.0014	0.9988	-0.0561	0.8112	-0.0574	0.7869	0.1109	0.9400	0.1758	0.7411	0.0649	0.9173
pnt	pointed	1630010_a_at	-0.3066	0.1423	-0.2445	0.2050	-0.9295	0.0015	-0.6544	0.2401	-0.2222	0.5276	0.4323	0.1413	0.0838	0.9032	-0.0131	0.9763	-0.0969	0.7378
CG31659	CG31659	1630011_at	0.6468	0.0458	0.1942	0.4514	0.4321	0.1395	0.1625	0.8337	0.2848	0.3400	0.1224	0.6914	-0.1368	0.8982	-0.1381	0.7776	-0.0013	0.9988
CG31225	CG31225	1630012_at	-0.0807	0.6635	0.0384	0.6786	0.2577	0.1281	0.0874	0.8321	-0.0273	0.8986	-0.1147	0.4364	-0.2222	0.7136	0.1036	0.7200	0.3258	0.2136
Sps2	Selenophosphate	1630013_s_at	-0.1478	0.5697	0.0600	0.6203	-0.0218	0.9053	-0.2866	0.4529	-0.5697	0.0138	-0.2831	0.1249	-0.1078	0.8930	-0.3646	0.2086	-0.2568	0.3986
---	---	1630014_at	-2.2724	0.0129	-3.1404	0.0113	-3.1588	0.0057	0.4793	0.8621	0.7347	0.4754	0.2554	0.8191	0.1521	0.9721	-0.2934	0.8445	-0.4455	0.7224
---	---	1630015_at	0.1831	0.2965	0.1952	0.2941	0.1885	0.2894	0.1129	0.8078	0.0032	0.9903	-0.1097	0.5453	0.0256	0.9808	-0.1060	0.6789	-0.1316	0.5944
Vha16-2	Vha16-2	1630016_at	-0.0102	0.9624	-0.0316	0.8241	0.1451	0.5561	0.0971	0.8164	0.0383	0.8591	-0.0589	0.7398	-0.0565	0.9641	0.0267	0.9620	0.0833	0.8488
Ptp99A	Protein tyrosine pl	1630017_at	0.0162	0.9392	0.2920	0.0394	0.2910	0.0742	0.0359	0.9463	-0.0891	0.5857	-0.1250	0.3679	0.1024	0.8597	0.1827	0.4325	0.0802	0.7691
spn-B	Spindle-B	1630018_at	-0.0484	0.9010	-0.2639	0.3887	-0.2846	0.2346	-0.1713	0.6549	0.1626	0.3899	0.3339	0.0482	-0.1849	0.9365	0.0372	0.9773	0.2221	0.8039
CG12026	CG12026	1630019_s_at	-2.9343	0.0006	0.3333	0.8002	-1.6251	0.0188	-1.9127	0.1054	-3.1808	0.0015	-1.2682	0.0482	-0.1179	0.9841	0.1438	0.9476	0.2617	0.8874
CG13041	CG13041	1630020_at	-0.1000	0.6243	-0.0137	0.8936	0.0653	0.6988	0.1239	0.8103	0.0441	0.8712	-0.0797	0.7142	0.0554	0.9460	0.0360	0.9271	-0.0194	0.9564
Gr59a	Gustatory recepto	1630021_at	0.1205	0.6093	0.1075	0.5858	0.1253	0.4181	-0.0135	0.9860	0.0125	0.9648	0.0260	0.9074	-0.1026	0.8611	-0.0200	0.9593	0.0827	0.7656
CG40212	CG40212	1630022_at	0.0090	0.9734	0.0645	0.6081	0.1814	0.4347	0.0639	0.9255	0.0488	0.8554	-0.0150	0.9544	-0.0861	0.9246	0.0538	0.9084	0.1400	0.6761
DLP	Daxx-like protein	1630023_at	0.7122	0.0041	0.2978	0.3273	-0.0942	0.5477	-0.2315	0.5886	0.3451	0.1115	0.5766	0.0099	0.1082	0.8890	0.0388	0.9341	-0.0695	0.8549
MED24	Mediator complex	1630024_at	-0.1043	0.6473	0.5946	0.1636	0.7308	0.0151	-0.0311	0.9745	-0.4851	0.0470	-0.4539	0.0395	-0.1901	0.8825	0.3296	0.5176	0.5197	0.3023
CG2865	CG2865	1630025_at	-0.3859	0.4265	0.2197	0.5928	0.1117	0.5270	0.3398	0.5917	0.3152	0.3338	-0.0246	0.9517	0.3775	0.8037	0.8306	0.1749	0.4531	0.4876
dpp	shortvein	1630026_s_at	-0.8283	0.2414	-1.4657	0.0181	-2.5222	0.0004	-0.2652	0.6888	0.7548	0.0223	1.0200	0.0035	0.6805	0.7560	0.0601	0.9687	-0.6203	0.5232
Rtnl1	Rtnl1	1630027_s_at	0.1947	0.3805	0.5750	0.0599	0.1571	0.4865	0.0267	0.9744	-0.0498	0.8489	-0.0765	0.7219	0.4041	0.5228	0.3030	0.3034	-0.1011	0.7774
p53	p53-like regulator	1630028_at	-0.0506	0.7663	-0.1348	0.4327	-0.0383	0.8832	0.0732	0.9130	0.0675	0.7951	-0.0057	0.9831	0.0934	0.8940	0.1553	0.5715	0.0619	0.8549
---	---	1630029_at	-0.0884	0.7447	0.0503	0.6109	-0.0279	0.8864	0.0703	0.9110	0.0343	0.8999	-0.0360	0.8788	0.1504	0.7485	0.0340	0.9121	-0.1165	0.5862
CG7777 /// DyakCG7777	CG7777	1630030_at	-0.3619	0.2002	-0.7280	0.0912	0.1901	0.5545	0.2460	0.6597	0.3925	0.1419	0.1465	0.5846	-0.9128	0.3712	-0.1029	0.9039	0.8098	0.1701
RpS5a	Minute 1(o)	1630031_at	0.2320	0.1751	0.2367	0.1287	0.4100	0.0201	0.0591	0.9088	-0.1229	0.4747	-0.1820	0.2163	-0.0187	0.9848	-0.0368	0.9121	-0.0180	0.9526
---	---	1630032_at	0.1229	0.6077	0.1213	0.4911	0.0776	0.6441	0.0268	0.9672	0.0533	0.7997	0.0266	0.8974	0.1713	0.7726	0.0792	0.8117	-0.0921	0.7545
---	---	1630033_at	0.1554	0.4292	0.0853	0.6469	-0.0819	0.7261	-0.0916	0.8676	-0.1393	0.4953	-0.0477	0.8296	0.0660	0.9095	0.0377	0.9052	-0.0282	0.9211
CG12119	CG12119	1630034_at	-3.1138	0.0052	-3.9235	0.0074	-3.6919	0.0000	0.0658	0.9225	-0.8500	0.0022	-0.9158	0.0010	-0.1115	0.9816	-1.5890	0.1498	-1.4775	0.2107
---	---	1630035_at	-0.0725	0.7490	0.0283	0.7799	0.0192	0.9148	-0.2144	0.4631	-0.2022	0.1991	0.0122	0.9511	-0.1305	0.8076	-0.0411	0.9046	0.0894	0.7257
CG13594	CG13594	1630036_at	-0.9084	0.1215	0.0175	0.8687	-0.0446	0.9117	-0.0952	0.9627	-1.8439	0.0054	-1.7487	0.0041	0.0764	0.9589	-0.5914	0.1504	-0.6678	0.1422
Or59b	Odorant receptor	1630037_at	0.2182	0.2934	0.0210	0.8366	0.0387	0.8857	-0.0655	0.9436	0.0215	0.9560	0.0870	0.7637	0.0228	0.9816	0.0071	0.9877	-0.0157	0.9621
pyd3	pyd3	1630038_at	-1.2699	0.0008	-2.1173	0.0039	-2.0838	0.0000	-0.1095	0.8297	0.1207	0.5667	0.2302	0.1861	-0.1384	0.8692	-0.6846	0.0622	-0.5462	0.1313
fred	friend of echinoid	1630039_at	-0.0271	0.8834	-0.0731	0.5843	0.3315	0.0613	0.2648	0.4139	0.1002	0.6131	-0.1646	0.3142	-0.0287	0.9653	0.0409	0.8721	0.0696	0.7334
---	---	1630040_at	0.2134	0.1904	-0.0230	0.8777	0.0461	0.7748	-0.0227	0.9671	0.1303	0.3742	0.1530	0.2339	-0.0613	0.8972	-0.1080	0.5597	-0.0467	0.8347
CG10899	CG10899	1630041_at	0.1995	0.2050	0.1238	0.3214	0.0037	0.9889	-0.0739	0.8822	0.0550	0.7925	0.1289	0.4192	0.1246	0.8097	-0.0397	0.9039	-0.1643	0.4597
Ca-alpha1T	Ca-alpha1T	1630042_at	-0.1406	0.6803	0.2246	0.5015	0.2540	0.2154	-0.0338	0.9727	-0.3325	0.1730	-0.2987	0.1723	0.1908	0.8463	0.1911	0.6721	0.0002	0.9997
alpha-catenin-related	---	1630043_a_at	-1.6895	0.0026	-2.1556	0.0069	-2.1426	0.0001	0.2339	0.8403	0.8847	0.0468	0.6507	0.0931	0.0240	0.9829	0.0410	0.9178	0.0170	0.9643
CG6904	Glycogen synthas	1630044_s_at	0.7324	0.0149	0.7643	0.1162	1.0771	0.0029	-0.1595	0.8196	-1.2076	0.0017	-1.0480	0.0019	-0.4775	0.7116	-1.2464	0.0418	-0.7689	0.1783
Scgalpha	Sarcoglycan alpha	1630045_at	-1.8427	0.0006	0.0407	0.7253	-0.5438	0.1972	-0.5576	0.4998	-1.8368	0.0021	-1.2792	0.0062	-0.1734	0.7979	-0.2649	0.3366	-0.0915	0.7859
---	---	1630046_at	0.0982	0.5487	0.0005	0.9994	0.0400	0.8661	0.0372	0.9633	0.0110	0.9738	-0.0262	0.9209	-0.0610	0.9277	-0.0414	0.9015	0.0196	0.9492
GABA-B-R1	metabotropic GAE	1630047_at	0.2868	0.1082	0.0478	0.7891	-0.0228	0.8893	-0.0510	0.9181	0.0794	0.6423	0.1304	0.3510	-0.0846	0.9168	-0.1962	0.4969	-0.1116	0.7282
ppk	pickpocket1	1630048_at	0.1146	0.5946	-0.0856	0.4567	0.0352	0.8386	0.1236	0.8327	0.2408	0.2821	0.1172	0.6001	-0.1380	0.7707	-0.0323	0.9178	0.1057	0.6291
CG4461	CG4461	1630049_at	-0.6338	0.1541	0.0830	0.5418	-0.6287	0.0093	-0.3224	0.4183	-0.6241	0.0122	-0.3018	0.1248	0.3794	0.7220	0.0322	0.9655	-0.3472	0.4627
CG8569	CG8569	1630050_at	-0.0813	0.8034	0.0712	0.8062	-0.0207	0.9332	0.0433	0.9325	0.0640	0.7157	0.0208	0.9100	0.1527	0.9087	0.2595	0.6058	0.1067	0.8636
Tsp42Ec	tetraspanin 42E	1630051_at	0.1492	0.5490	0.0648	0.5773	0.2989	0.0551	0.4500	0.2920	0.2263	0.3781	-0.2237	0.3298	0.1199	0.8513	0.0129	0.9784	-0.1070	0.7221
CG8446	CG8446	1630052_s_at	-0.4345	0.1226	0.1493	0.6191	-0.1786	0.4494	-0.1181	0.8578	-0.4210	0.0780	-0.3029	0.1524	0.0701	0.9689	0.1455	0.8126	0.0754	0.9084
---	---	1630053_at	0.0553	0.7781	-0.0394	0.8523	-0.2442	0.1859	-0.0581	0.9300	0.0895	0.6862	0.1476	0.4149	0.0828	0.9246	0.0069	0.9924	-0.0759	0.8373
CG4465 /// DereCG4465 ///	CG4465	1630054_at	0.6123	0.3120	-0.0392	0.7362	-0.0450	0.8063	0.1566	0.8189	0.1578	0.5878	0.0012	0.9973	0.0363	0.9914	-0.6362	0.3630	-0.6725	0.3583
sec31	sec31	1630055_a_at	0.3146	0.5339	0.7798	0.1565	1.0161	0.0002	0.2727	0.4102	0.3785	0.0450	0.1058	0.5597	-0.0787	0.9816	0.7909	0.2960	0.8696	0.2784
CG32054	CG32054	1630056_at	0.0136	0.9861	-0.0022	0.9905	-0.1157	0.4944	-0.1313	0.9302	-0.3035	0.5139	-0.1723	0.7121	0.0541	0.9816	-0.2375	0.7054	-0.2916	0.6263
CG11753	CG11753	1630057_at	-0.1908	0.4847	-0.0233	0.8215	0.0279	0.8809	0.0312	0.9621	-0.2112	0.2258	-0.2424	0.1191	-0.0545	0.9644	-0.0691	0.8850	-0.0146	0.9777
---	---	1630058_at	0.1921	0.2391	0.1281	0.5396	0.2615	0.1721	0.0550	0.8950	0.0222	0.9083	-0.0328	0.8364	-0.0513	0.9677	0.0261	0.9623	0.0775	0.8602
CG34053	CG34053	1630059_at	-0.0529	0.7171	0.0003	1.0000	0.0018	0.9935	0.0210	0.9697	-0.1181	0.4235	-0.1391	0.2803	0.0576	0.9076	-0.0375	0.8830	-0.0951	0.6168
---	---	1630060_at	0.1023	0.6249	0.0041	0.9768	0.2816	0.2754	0.0657	0.9120	0.0019	0.9945	-0.0638	0.7570	-0.0863	0.9365	0.0978	0.8287	0.1841	0.6169
---	---	1630061_at	-0.0168	0.9413	-0.0929	0.5521	0.0235	0.8983	0.1125	0.7855	0.0932	0.6226	-0.0194	0.9260	-0.0606	0.8940	-0.0140	0.9605	0.0465	0.8287
l(2)35Di	lethal (2) 35Di	1630062_a_at	-0.3720	0.1410	0.5523	0.0795	0.9136	0.0150	0.0181	0.9909	-1.4197	0.0025	-1.4378	0.0014	-0.1869	0.8825	-0.4358	0.3571	-0.2489	0.6328
eIF-3p66	Eukaryotic initiatio	1630063_a_at	0.2789	0.2339	0.8122	0.0521	0.9564	0.0022	0.3691	0.4326	-0.1521	0.5877	-0.5212	0.0314	0.1385	0.8461	0.3533	0.2145	0.2148	0.4807
CG5639	N-like	1630064_at	0.8119	0.0593	0.2939	0.6087	0.7740	0.0447	0.1792	0.6948	0.2213	0.2985	0.0420	0.8640	-0.2232	0.9467	-0.3303	0.7944	-0.1071	0.9394
CG6912	CG6912	1630065_at	1.5508	0.0034	-0.7565	0.3916	0.6640	0.0294	0.9029	0.0546	1.4560	0.0008	0.5531	0.0295	-0.4207	0.8427	-0.8641	0.3073	-0.4433	0.6366

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG4302	CG4302	1630066_at	1.7790	0.0071	0.9528	0.1770	1.1082	0.0065	-0.4541	0.6506	-0.2559	0.6367	0.1982	0.6966	-0.6842	0.6999	-1.2796	0.0954	-0.5955	0.4370
TepII	Thiolester contain	1630067_a_at	1.4742	0.0239	-0.6285	0.2766	0.1366	0.6820	0.2667	0.7929	2.0435	0.0011	1.7768	0.0012	-0.5141	0.7726	-0.0280	0.9849	0.4862	0.5498
---	---	1630068_at	0.2014	0.3418	0.1508	0.2696	0.0252	0.9191	0.0134	0.9860	0.0699	0.7388	0.0565	0.7737	0.3277	0.5875	0.1258	0.6729	-0.2019	0.4683
CG13728	CG13728	1630069_at	0.1069	0.7383	0.0594	0.5827	0.1682	0.2938	-0.1155	0.7749	-0.2200	0.1906	-0.1045	0.5250	0.0485	0.9717	-0.0759	0.8765	-0.1244	0.7524
Or46a	Olfactory receptor	1630070_at	0.0920	0.6072	0.2136	0.1818	0.0509	0.7761	-0.0795	0.8650	-0.1357	0.4291	-0.0562	0.7566	0.0351	0.9611	-0.0482	0.8654	-0.0833	0.7154
---	---	1630071_at	0.1096	0.5634	0.3151	0.1272	0.0181	0.9487	-0.2466	0.4757	-0.2993	0.1063	-0.0527	0.8007	-0.0136	0.9914	0.0024	0.9980	0.0160	0.9721
---	---	1630072_a_at	1.9601	0.4298	-1.5365	0.1217	-2.0486	0.0028	0.2137	0.9711	1.3511	0.3768	1.1374	0.4163	0.6887	0.9331	-2.3381	0.3664	-3.0268	0.2661
CG12173	CG12173	1630073_at	-0.5451	0.2535	-0.2791	0.3389	-0.2088	0.2874	-0.1716	0.6081	-0.5497	0.0063	-0.3780	0.0200	-0.2503	0.8683	-0.2777	0.6757	-0.0274	0.9769
---	---	1630074_at	0.0546	0.8130	-0.0186	0.8535	0.2073	0.2376	0.3139	0.2401	0.2305	0.1436	-0.0834	0.5999	0.0635	0.9544	0.1536	0.6686	0.0900	0.8255
CG9699	CG9699	1630075_s_at	-0.4843	0.5054	-0.8269	0.3716	-0.9360	0.0166	-0.2831	0.5932	-0.5722	0.0396	-0.2891	0.2295	-0.3245	0.9306	-1.0942	0.3522	-0.7697	0.5450
Hop	Hsp70/Hsp90 orga	1630076_at	-0.0777	0.9130	-0.0084	0.9817	-0.2027	0.1518	0.1500	0.9363	0.4430	0.4366	0.2931	0.5942	0.2442	0.7810	0.4477	0.2217	0.2035	0.6189
Or2a	Olfactory receptor	1630077_at	-0.0020	0.9936	-0.0637	0.5534	-0.1942	0.1861	0.1507	0.7409	0.1828	0.3723	0.0320	0.8942	0.1802	0.6749	0.1144	0.5329	-0.0658	0.7465
CG32000	CG32000	1630078_a_at	-0.3200	0.1539	-0.4145	0.3385	-0.8819	0.0023	-0.3844	0.2697	0.0499	0.8520	0.4342	0.0263	0.1040	0.9460	-0.0338	0.9648	-0.1378	0.8112
Atu	Another transcript	1630079_at	-0.4180	0.2026	0.1692	0.6844	0.3502	0.0409	-0.0637	0.8877	-0.1712	0.2629	-0.1075	0.4552	-0.1197	0.9400	0.2411	0.6478	0.3608	0.4729
phol	pho-like	1630080_at	0.4621	0.0603	0.2607	0.1501	-0.0080	0.9682	-0.4523	0.1160	0.1054	0.5731	0.5577	0.0040	-0.0844	0.9056	0.0557	0.8790	0.1402	0.6111
---	---	1630081_at	0.1246	0.4785	0.0290	0.8258	-0.0793	0.7473	-0.0993	0.8102	-0.2128	0.1930	-0.1135	0.4667	0.0020	0.9989	-0.0697	0.8174	-0.0718	0.7983
---	---	1630082_at	0.1600	0.3641	0.2124	0.2671	0.1564	0.4870	0.0206	0.9722	-0.0046	0.9827	-0.0252	0.8853	0.1557	0.8215	0.1376	0.6626	-0.0181	0.9661
Rab8	Rab-protein 8	1630083_at	-0.1785	0.4169	0.3680	0.2924	-0.1781	0.4592	-0.3042	0.5735	-0.4944	0.0765	-0.1902	0.4697	0.2840	0.7513	0.2255	0.5800	-0.0586	0.9121
CG10205	CG10205	1630084_at	-0.0334	0.8561	-0.1724	0.4651	0.0122	0.9748	0.0462	0.9491	0.2994	0.1363	0.2531	0.1595	-0.1876	0.8608	0.1680	0.7381	0.3557	0.4148
Peritrophin-A	peritrophin A	1630085_s_at	-3.6251	0.0003	-3.7442	0.0014	-3.4043	0.0000	0.4260	0.4167	-0.1324	0.6918	-0.5584	0.0385	0.0417	0.9611	-0.1745	0.4753	-0.2162	0.3807
wuho	CG15897	1630086_at	-0.0066	0.9879	0.3103	0.3409	0.5096	0.0635	0.2341	0.6405	-0.1266	0.6453	-0.3607	0.0990	0.0440	0.9853	0.2433	0.6949	0.1993	0.7527
Tsp33B	Tetraspanin 33B	1630087_at	0.2651	0.2917	0.2089	0.3636	0.1038	0.6199	-0.1392	0.8366	-0.1395	0.6207	-0.0003	0.9995	0.0114	0.9916	0.0701	0.8393	0.0587	0.8626
CG16743 /// DyakCG16743	CG16743	1630088_at	1.7837	0.0013	1.5511	0.0217	2.4563	0.0003	0.3882	0.5251	0.7277	0.0304	0.3395	0.2363	-0.6673	0.6660	0.8346	0.1694	1.5018	0.0557
fzo	fuzzy onions	1630089_at	0.2403	0.2852	0.2399	0.2482	0.2131	0.2474	-0.2240	0.6034	-0.1191	0.6204	0.1049	0.6372	-0.0681	0.9467	0.0018	0.9989	0.0699	0.8666
mxc	lethal(1)malignant	1630090_at	-0.1477	0.5229	0.3834	0.2559	0.7232	0.0166	-0.0090	0.9951	-0.0723	0.8484	-0.0633	0.8520	-0.2750	0.7541	0.6167	0.1054	0.8917	0.0559
CG31296	CG31296	1630091_at	0.1979	0.1961	-0.2089	0.4256	0.1349	0.5603	0.4323	0.2782	0.3428	0.1424	-0.0894	0.7204	-0.1058	0.8454	-0.1450	0.5296	-0.0392	0.8982
CG13088	CG13088	1630092_at	0.3893	0.0466	0.6135	0.2158	0.4540	0.0573	-0.1556	0.6864	0.2210	0.2157	0.3766	0.0267	-0.0104	0.9952	0.3849	0.3620	0.3953	0.3705
RpS24	Ribosomal protein	1630093_at	0.3946	0.0466	0.3897	0.0641	0.5973	0.0246	0.1428	0.8299	-0.1038	0.7270	-0.2466	0.2840	0.0743	0.8999	-0.0787	0.7649	-0.1530	0.4978
l(1)G0269	lethal (1) G0269	1630094_at	0.1963	0.6219	-0.2701	0.6613	-0.5028	0.1649	-0.2742	0.5096	0.6580	0.0095	0.9322	0.0012	0.0456	0.9913	0.2338	0.8397	0.1882	0.8699
Drl-2	Derailed 2	1630095_a_at	-0.0587	0.7541	0.0694	0.5407	-0.1272	0.4651	-0.0104	0.9893	-0.0381	0.8698	-0.0277	0.8955	-0.0352	0.9829	-0.1707	0.6819	-0.1355	0.7524
CG2533	CG2533	1630096_at	0.1978	0.2357	0.1285	0.3103	0.2089	0.1633	-0.0351	0.9473	-0.0054	0.9809	0.0297	0.8654	0.1100	0.8207	0.0072	0.9855	-0.1028	0.6366
CG17834	CG17834	1630097_s_at	-2.3365	0.0024	-2.7410	0.0100	-2.8122	0.0000	-0.0637	0.9603	0.0131	0.9801	0.0767	0.8456	-0.0518	0.9841	-0.1546	0.8473	-0.1028	0.8993
CG6447	CG6447	1630098_a_at	0.1485	0.4504	-0.1770	0.3168	0.1737	0.3265	0.2018	0.7135	0.2556	0.3114	0.0538	0.8530	-0.1734	0.7769	-0.1028	0.7416	0.0706	0.8332
mip120	Myb-interacting pr	1630099_at	0.4089	0.1600	0.5698	0.0626	0.6295	0.0152	-0.0887	0.9194	0.2014	0.4735	0.2901	0.2277	-0.0204	0.9898	0.3683	0.2520	0.3888	0.2614
CG31607	CG31607	1630100_at	-3.2781	0.0025	-0.0645	0.9713	-3.3453	0.0021	-3.0192	0.0256	-4.5619	0.0004	-1.5428	0.0264	0.3978	0.9401	-1.2792	0.4317	-1.6769	0.3147
Smr	SMRTER	1630101_s_at	0.0692	0.9173	0.1456	0.8593	-1.1617	0.0023	-0.5084	0.5419	0.4389	0.3167	0.9473	0.0228	0.8592	0.7138	0.5473	0.5994	-0.3119	0.7898
CG12617	CG12617	1630102_at	0.0673	0.7150	0.0185	0.8546	-0.0144	0.9490	-0.0455	0.9346	-0.0825	0.6560	-0.0370	0.8455	0.0803	0.8882	-0.0185	0.9588	-0.0988	0.6749
mmd	Meltrin-like	1630103_at	0.1563	0.4062	0.0431	0.7008	0.0606	0.7420	0.1395	0.7764	0.1838	0.3816	0.0443	0.8526	0.0023	0.9979	0.0109	0.9739	0.0086	0.9778
---	---	1630104_at	0.1087	0.6681	-0.0864	0.4867	0.1570	0.3547	0.1245	0.7857	0.1512	0.4394	0.0268	0.9067	-0.0136	0.9914	-0.0358	0.9377	-0.0222	0.9565
CG3891	CG3891	1630105_at	-0.1668	0.5542	-0.0330	0.9478	-0.1194	0.4337	0.0111	0.9897	0.0851	0.7076	0.0740	0.7242	0.2199	0.8379	0.2748	0.5523	0.0549	0.9289
CG10362	CG10362	1630106_at	-0.0886	0.6268	-0.1468	0.4606	0.0827	0.6135	0.0639	0.9118	-0.1929	0.2979	-0.2569	0.1165	-0.0947	0.9168	-0.1258	0.7275	-0.0311	0.9433
CG15685	CG15685	1630107_at	0.0332	0.8964	0.0913	0.4543	0.1298	0.4206	0.0621	0.9263	-0.1062	0.6385	-0.1683	0.3661	-0.0042	0.9976	-0.0777	0.8584	-0.0735	0.8577
r-cup	ryder cup	1630108_a_at	0.0384	0.8644	-0.1988	0.3471	-0.0107	0.9691	0.1740	0.7138	0.2065	0.3478	0.0325	0.9004	-0.0242	0.9853	-0.1999	0.5251	-0.1757	0.5870
CG8628	CG8628	1630109_at	0.0776	0.9728	0.0749	0.7009	0.3458	0.1308	-0.0592	0.9909	-1.2903	0.2572	-1.2311	0.2258	0.0516	0.9928	-0.6915	0.6344	-0.7431	0.6081
CG12935	CG12935	1630110_at	0.1273	0.5642	0.1386	0.7297	0.1187	0.8043	-0.0019	0.9987	0.5061	0.0366	0.5080	0.0230	-0.1384	0.9626	0.2693	0.7925	0.4077	0.6419
---	---	1630111_at	-0.0214	0.9049	-0.0848	0.5083	0.1538	0.3641	0.1797	0.5639	0.0310	0.8856	-0.1487	0.3061	0.0091	0.9938	0.0730	0.8307	0.0638	0.8467
CG18731	CG18731	1630112_at	0.2363	0.2290	0.4795	0.1015	0.2892	0.0939	-0.1548	0.6864	-0.0180	0.9456	0.1369	0.4109	-0.0197	0.9862	0.1568	0.5697	0.1765	0.5193
CG10660	CG10660	1630113_at	-0.5230	0.0189	1.3375	0.0141	0.9040	0.0009	-0.4452	0.2941	-1.5413	0.0004	-1.0962	0.0011	0.0243	0.9848	0.4270	0.1382	0.4027	0.1897
---	---	1630114_at	-0.1081	0.5598	-0.0572	0.5939	-0.0678	0.7228	0.0422	0.9426	0.0006	0.9981	-0.0417	0.8265	0.1135	0.8086	0.0128	0.9682	-0.1007	0.6317
CG15602	CG15602	1630115_at	-0.2000	0.2155	-0.6416	0.0144	-0.6207	0.0607	0.0641	0.9011	0.1932	0.2502	0.1291	0.4097	0.0384	0.9823	-0.2667	0.5200	-0.3052	0.4617
CG13446	CG13446	1630116_at	0.0349	0.8558	0.0088	0.9415	0.3528	0.0718	0.2601	0.3773	0.0545	0.7888	-0.2056	0.1665	-0.0410	0.9499	0.0485	0.8551	0.0895	0.6728
CG34357	CG14652	1630117_at	-0.0114	0.9572	0.1843	0.3137	-0.0176	0.9591	-0.0362	0.9622	-0.2683	0.1842	-0.2321	0.2007	0.2227	0.8122	0.1055	0.8392	-0.1172	0.8041
Dyb	Dystrobrevin-like	1630118_s_at	-1.2926	0.0966	0.1911	0.7193	-0.6608	0.0160	-0.0686	0.9254	-0.1132	0.6476	-0.0445	0.8621	0.7986	0.7519	1.3663	0.1928	0.5677	0.6263
norpA	phospholipase C	1630119_s_at	-1.3011	0.0039	1.1134	0.1146	0.1469	0.5359	-0.4846	0.3477	-0.8538	0.0112	-0.3692	0.1630	0.5178	0.7241	1.5790	0.0378	1.0612	0.1275
CG32636	CG32636	1630120_s_at	-0.0296	0.8696	0.2823	0.1821	0.3672	0.0787	-0.2608	0.3837	-0.2349	0.1634	0.0259	0.8991	0.0657	0.9457	0.2657	0.3499	0.2000	0.5092
CG9392	CG9392	1630121_at	0.0948	0.6934	0.1673	0.3785	0.1535	0.2988	-0.2053	0.5680	-0.3489	0.0617	-0.1436	0.4019	0.0438	0.9700	-0.2375	0.4245	-0.2813	0.3590
CG13978	CG13978	1630122_at	-0.2431	0.4402	-0.0806	0.3983	-0.1798	0.3254	-0.1010	0.9080	-0.1013	0.7634	-0.0003	0.9995	-0.0048	0.9964	0.0185	0.9684	0.0234	0.9534
CG3099	CG3099	1630123_at	-1.0749	0.0487	-1.6147	0.0307	-1.1807	0.0412	0.3811	0.8005	1.4063	0.0283	1.0252	0.0626	0.0213	0.9943	0.9651	0.1542	0.9438	0.1995
CG5514	CG5514	1630124_at	-0.1070	0.7551	0.0571	0.7793	-0.6686	0.0263	-0.1036	0.8987	0.2786	0.3009	0.3822	0.1081	0.3754	0.7697	0.3996	0.4711	0.0243	0.9777
---	---	1630125_s_at	0.1776	0.4062	0.0491	0.6532	-0.0047	0.9826	-0.0343	0.9507	0.0545	0.7674	0.0888	0.5583	0.0483	0.9589	-0.0409	0.9220	-0.0892	0.7764
CG6452	CG6452	1630126_at	-0.0605	0.8589	0.1681	0.3650	0.1646	0.3647	-0.0883	0.9186	-0.2743	0.3068	-0.1860	0.4597	0.0336	0.9742	-0.0535	0.8850	-0.0872	0.7727
---	---	1630127_at	-1.5638	0.0060	-2.1596	0.0163	-2.0779	0.0001	0.1208	0.9039	0.5308	0.1028	0.4100	0.1582	0.0688	0.9698	-0.2694	0.5860	-0.3382	0.4848
---	---	1630128_at	-0.0261	0.9059	-0.0209	0.8802	0.0000	0.9998	-0.0037	0.9956	0.0209	0.9455	0.0246	0.9228	0.0886	0.8825	0.0676	0.8212	-0.0210	0.9478
CG32355	CG32355	1630129_at	-2.1859	0.0023	-2.5446	0.0014	-3.4253	0.0000	-0.3081	0.3743	-0.1247	0.5543	0.1834	0.3021	0.4636	0.7726	-0.5804	0.3978	-1.0440	0.1573
CG4404	CG4404	1630130_at	-1.0782	0.0041	-0.7468	0.0109	-0.8985	0.0566	-0.3097	0.5735	-1.0684	0.0033	-0.7587	0.0091	-0.1666	0.9390	-0.6935	0.2871	-0.5269	0.4457
CG6567	CG6567	1630131_at	0.6660	0.0146	0.9638	0.0111	0.9337	0.0006	0.0814	0.8667	-0.2704	0.1127	-0.3518	0.0290	0.0832	0.9306	0.0382	0.9395	-0.0450	0.9162
l(1)G0022	lethal (1) G0022	1630132_at	-0.2322	0.5475	-0.0623	0.5779	-0.1544	0.5683	0.4939	0.4333	0.6436	0.0659	0.1497	0.6789	0.4641	0.5757	0.7194	0.0714	0.2554	0.5141
zen2	zerknullt 2	1630133_at	0.0820	0.6987	0.1293	0.5687	0.1038	0.5137	0.0910	0.8405	-0.0656	0.7422	-0.1566	0.3070	-0.0045	0.9976	-0.0296	0.9577	-0.0251	0.9589
CG32233	CG32233	1630134_at	0.1994	0.4349	-0.0139	0.9229	0.1179	0.4496	0.1645	0.6235	0.2127	0.1986	0.0482	0.7965	-0.0107	0.9918	-0.0772	0.8038	-0.0665	0.8264

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1630135_at	0.2702	0.1269	0.0772	0.6882	0.0873	0.6058	0.1326	0.6578	0.3111	0.0372	0.1785	0.1642	-0.0946	0.8940	-0.0189	0.9650	0.0757	0.8178
---	---	1630136_at	0.0785	0.7896	0.1915	0.3359	-0.3184	0.2100	-0.1317	0.7104	-0.0546	0.7799	0.0771	0.6415	0.1353	0.8941	-0.0176	0.9797	-0.1530	0.7245
CG15696 /// DereCG15696	CG15696	1630137_at	-0.0788	0.7150	0.1294	0.4579	-0.1125	0.7201	0.0189	0.9838	-0.0986	0.6906	-0.1175	0.5862	0.1492	0.8331	0.2057	0.4976	0.0565	0.8893
Eig71Ej	Eig71Ej	1630138_at	-0.0757	0.6443	-0.2020	0.3650	0.0409	0.8181	0.1240	0.7916	0.0667	0.7717	-0.0573	0.7884	-0.0695	0.9095	-0.0756	0.7779	-0.0061	0.9859
---	---	1630139_at	0.0664	0.7722	0.0760	0.5195	-0.0483	0.8230	-0.0474	0.9592	0.2187	0.3812	0.2661	0.2239	0.0097	0.9943	0.0719	0.8690	0.0622	0.8834
CG14931	CG14931	1630140_at	0.2346	0.3721	-0.1329	0.2313	-0.3139	0.4125	-0.3821	0.2435	0.4220	0.0362	0.8041	0.0013	-0.1713	0.9174	0.0349	0.9709	0.2061	0.7549
CG2158	CG2158	1630141_at	0.3646	0.0849	0.1725	0.6302	0.3086	0.0763	0.0379	0.9507	0.5940	0.0048	0.5561	0.0038	-0.0698	0.9672	0.3510	0.4317	0.4208	0.3614
sog	short gastrulation	1630142_at	-0.1467	0.3750	0.0100	0.9604	-0.0907	0.6398	0.1539	0.7425	0.0705	0.7776	-0.0835	0.7010	0.2704	0.7506	0.2909	0.4223	0.0206	0.9706
---	---	1630143_at	0.1469	0.4494	-0.0783	0.4999	-0.0979	0.6704	0.0115	0.9895	0.0344	0.9019	0.0229	0.9262	-0.0844	0.9296	-0.1622	0.6265	-0.0778	0.8443
Cdk7	Cyclin-dependent	1630144_at	0.0946	0.6311	0.1173	0.4746	0.4492	0.0261	0.1141	0.8051	-0.0490	0.8359	-0.1631	0.3334	-0.2714	0.6994	-0.1024	0.7782	0.1689	0.5944
Tsp42Ea	tetraspanin 42E	1630145_s_at	-0.9491	0.0324	-0.3993	0.3945	-0.7503	0.0080	-0.2204	0.6134	-0.5579	0.0190	-0.3375	0.0853	0.1002	0.9677	-0.1307	0.8929	-0.2309	0.7620
CG12508	CG12508	1630146_at	-0.6377	0.1117	0.1800	0.7136	0.5055	0.0790	-0.1098	0.9023	-1.1213	0.0031	-1.0115	0.0028	-0.2937	0.8395	-0.0876	0.9263	0.2061	0.7668
Hex-C	Hexokinase	1630147_at	2.8590	0.0024	2.3433	0.0015	2.7996	0.0000	0.8230	0.4142	0.3574	0.5536	-0.4656	0.3676	0.1972	0.6903	-0.2561	0.1955	-0.4533	0.0625
---	---	1630148_at	0.0220	0.9239	0.3152	0.0336	0.1170	0.5709	0.0725	0.9311	-0.0347	0.9201	-0.1072	0.6775	0.2500	0.6702	0.1142	0.6605	-0.1357	0.5929
CG6290	CG6290	1630149_at	0.1788	0.3522	-0.1859	0.2176	-0.1393	0.5963	0.1945	0.6597	0.3888	0.0698	0.1943	0.3145	-0.1070	0.8541	-0.0563	0.8678	0.0507	0.8761
Cg25C	type IV collagen, a	1630150_s_at	-1.8814	0.0299	-1.4073	0.3822	-1.5307	0.0006	0.0809	0.9149	-0.5542	0.0273	-0.6351	0.0095	0.1465	0.9848	-0.1998	0.9438	-0.3463	0.8857
spn-D	spindle D	1630151_at	0.1071	0.4509	0.1006	0.3178	0.0191	0.9558	-0.1129	0.8938	0.0435	0.9114	0.1564	0.5765	-0.1046	0.8395	-0.1002	0.6694	0.0044	0.9902
dpr3	dpr3	1630152_at	0.0996	0.6763	0.2004	0.2916	0.0387	0.8785	-0.1432	0.7647	-0.0393	0.8869	0.1039	0.6162	0.2124	0.7464	0.1922	0.4976	-0.0201	0.9599
CG1988	CG1988	1630153_a_at	-0.1002	0.6768	-0.0992	0.3413	-0.1343	0.6007	0.0909	0.8266	0.2137	0.1795	0.1227	0.4120	-0.0431	0.9797	0.1718	0.6876	0.2149	0.6012
CG3437	CG3437	1630154_at	0.1009	0.7745	0.2388	0.1920	0.5461	0.0139	-0.0623	0.9435	0.1247	0.6559	0.1870	0.4207	-0.4090	0.6660	0.2328	0.5618	0.6418	0.1267
Obp19c	Odorant-binding p	1630155_at	0.1362	0.4759	0.1135	0.2554	0.2238	0.2287	0.0378	0.9436	0.0322	0.8723	-0.0056	0.9766	0.0176	0.9862	0.1134	0.6536	0.0958	0.7148
CG12859 /// DyakCG12859	anon-fast-evolving	1630156_at	-0.3890	0.0636	-0.0780	0.8431	-0.0280	0.9240	-0.1095	0.8794	-0.5083	0.0451	-0.3988	0.0727	-0.0336	0.9831	-0.1236	0.7922	-0.0900	0.8524
CG7453	CG7453	1630157_a_at	0.2562	0.2303	0.2914	0.1127	-0.0120	0.9555	-0.1635	0.6046	0.1366	0.4034	0.3001	0.0412	0.1856	0.8122	0.1631	0.6414	-0.0225	0.9622
CG30429	CG30429	1630158_at	0.2806	0.3034	0.1034	0.3599	0.1029	0.5716	-0.1247	0.7235	0.0631	0.7364	0.1878	0.1856	0.0613	0.9306	0.0707	0.8106	0.0093	0.9804
CG11672	CG11672	1630159_at	0.0106	0.9732	0.0949	0.4655	0.0377	0.8547	0.1395	0.7377	0.0838	0.6922	-0.0557	0.7868	0.1700	0.8305	0.1870	0.5979	0.0170	0.9740
Fas2	Fasciclin II	1630160_at	0.0414	0.8889	-0.6832	0.2541	-0.6863	0.0258	0.0639	0.9228	0.5427	0.0142	0.4788	0.0151	0.1713	0.9296	-0.1051	0.9138	-0.2764	0.6902
fau	fau	1630161_at	-1.8314	0.0113	-2.4681	0.0525	-3.5941	0.0005	-0.7751	0.1462	-0.1015	0.7978	0.6736	0.0275	0.6123	0.8521	-0.6888	0.6407	-1.3011	0.3519
CG14870	CG14870	1630162_at	-0.0003	0.9991	0.1150	0.5371	0.0610	0.7382	-0.1555	0.7912	-0.1585	0.5357	-0.0030	0.9919	0.0391	0.9516	0.0206	0.9483	-0.0185	0.9460
CG32373	CG32373	1630163_at	-0.7906	0.0345	-0.7767	0.0653	-1.6809	0.0006	-0.8232	0.1250	0.0698	0.8704	0.8930	0.0082	-0.1335	0.8657	-0.1852	0.5800	-0.0516	0.9057
---	---	1630164_at	0.1869	0.4038	-0.0794	0.4184	-0.2694	0.1902	0.0448	0.9379	0.1136	0.5259	0.0688	0.6974	0.0502	0.9571	-0.2061	0.4379	-0.2564	0.3488
Sin3A	Enhancer of GMR	1630165_s_at	0.5894	0.0967	0.4107	0.1196	0.1450	0.4075	0.0039	0.9956	0.4659	0.0318	0.4620	0.0208	0.3243	0.7305	0.3128	0.4379	-0.0114	0.9856
Bj1	CG18640 (Bj1), R	1630166_s_at	0.1772	0.5044	-0.1624	0.6679	0.1141	0.6789	-0.0613	0.9515	0.2467	0.3852	0.3079	0.2149	-0.3654	0.7677	-0.2391	0.6827	0.1263	0.8524
Pcd	pterin-4a-carbinola	1630167_at	-0.3038	0.1872	-0.7599	0.1988	-0.3982	0.0727	-0.0248	0.9744	-0.1121	0.5825	-0.0874	0.6527	-0.3045	0.8122	-0.5342	0.3013	-0.2297	0.6990
CG31636	CG31636	1630168_at	0.1633	0.3255	0.0217	0.8960	0.1788	0.2201	0.1102	0.8350	0.0315	0.9099	-0.0787	0.7119	-0.0454	0.9309	-0.0415	0.8593	0.0039	0.9884
Sap47	Synapse-associat	1630169_a_at	0.7776	0.0470	0.0496	0.9546	0.0326	0.9087	0.0945	0.8742	0.5844	0.0126	0.4898	0.0166	0.0918	0.9816	-0.1869	0.8830	-0.2787	0.7923
Cyp12b2	Cyp12b2	1630170_at	1.1733	0.0024	0.3672	0.0770	0.5092	0.0236	-0.2881	0.4796	-0.0094	0.9780	0.2786	0.1524	-0.2415	0.7062	-0.7314	0.0285	-0.4900	0.1011
Cpr97Ea	CG6131	1630171_at	0.3850	0.0625	0.3060	0.3662	0.2372	0.2510	0.1186	0.8180	-0.2345	0.2462	-0.3531	0.0548	0.0952	0.9101	-0.2408	0.4229	-0.3360	0.2800
CG10950	CG10950	1630172_at	-0.1723	0.2861	-0.0877	0.5994	-0.0506	0.8227	-0.1383	0.8501	0.0031	0.9939	0.1414	0.5972	0.0164	0.9852	-0.0088	0.9815	-0.0252	0.9282
CG15745	CG15745	1630173_a_at	2.0885	0.0014	1.7863	0.0077	1.4062	0.0002	-0.0738	0.9487	0.9548	0.0096	1.0286	0.0041	0.1209	0.9246	0.4094	0.3255	0.2884	0.5171
CG4908	CG4908	1630174_s_at	-0.4044	0.2673	-0.0788	0.7151	-0.0771	0.7005	-0.2808	0.4586	-0.4619	0.0318	-0.1811	0.3347	-0.3195	0.7726	-0.2774	0.5800	0.0421	0.9492
---	---	1630175_s_at	0.4022	0.1819	0.1150	0.5118	0.1013	0.6053	0.1415	0.8485	0.2913	0.2850	0.1498	0.5787	0.0472	0.9653	-0.1907	0.5251	-0.2379	0.4203
CG18005	CG18005	1630176_at	-0.2316	0.3068	0.5547	0.1440	0.8645	0.0020	-0.0102	0.9900	-0.7984	0.0018	-0.7882	0.0011	-0.1324	0.9168	0.2254	0.6366	0.3578	0.4273
Ranbp11	Ranbp11	1630177_at	-0.5137	0.3812	-0.0049	0.9986	0.1045	0.6000	0.2354	0.5179	-0.2864	0.1344	-0.5218	0.0088	0.2314	0.9426	0.3450	0.7770	0.1136	0.9347
CG12082	CG12082	1630178_at	0.2263	0.3635	0.9065	0.0780	1.2216	0.0005	0.3499	0.2282	-0.1758	0.3138	-0.5257	0.0053	-0.0279	0.9913	0.5010	0.3389	0.5289	0.3369
CG40341	CG40341	1630179_at	-0.2353	0.1543	0.1558	0.6176	0.0743	0.6552	-0.0842	0.9136	-0.1791	0.4855	-0.0949	0.7156	-0.0338	0.9777	0.2083	0.4794	0.2421	0.4093
CG4815	CG4815	1630180_at	-0.0229	0.8834	0.1302	0.4220	0.0439	0.8142	-0.3091	0.2905	-0.3473	0.0487	-0.0382	0.8482	-0.1487	0.7697	-0.0397	0.9046	0.1090	0.6392
CG9194	CG9194	1630181_at	-2.5966	0.0028	-1.9177	0.0129	-2.4920	0.0000	-0.4966	0.5407	-0.9539	0.0314	-0.4573	0.2269	-0.2031	0.8564	-0.4059	0.3680	-0.2029	0.6888
bin	Biniou	1630182_at	-0.1247	0.5153	0.0046	0.9700	0.1144	0.5619	0.1417	0.7949	-0.0191	0.9538	-0.1608	0.4416	-0.0307	0.9816	0.1045	0.7883	0.1352	0.6968
---	---	1630183_at	0.1360	0.5240	0.1673	0.2525	0.1017	0.5974	-0.2153	0.4510	-0.0816	0.6385	0.1337	0.3478	0.0035	0.9976	0.0195	0.9647	0.0160	0.9684
CG40115	CG40115	1630184_at	-0.3539	0.2097	-0.5508	0.0398	-1.1486	0.0398	-0.3145	0.8271	0.7186	0.1925	1.0331	0.0433	0.1041	0.8960	0.2660	0.3625	0.1618	0.6140
---	---	1630185_x_at	0.0894	0.6762	-0.9605	0.0450	-0.9447	0.0016	-0.0530	0.9578	0.8148	0.0085	0.8677	0.0038	0.1555	0.8903	0.1423	0.7872	-0.0131	0.9849
CG13743	CG13743	1630186_at	0.0752	0.6535	0.0695	0.6190	-0.1737	0.2540	-0.2468	0.4769	-0.1348	0.4900	0.1120	0.5381	0.0467	0.9545	-0.0559	0.8690	-0.1026	0.7040
CG6299	CG6299	1630187_a_at	0.6949	0.0811	0.8320	0.1278	1.3103	0.0001	0.3963	0.2943	-0.0968	0.7060	-0.4931	0.0213	-0.0271	0.9928	0.1627	0.8725	0.1898	0.8323
---	---	1630188_at	0.0423	0.8405	0.0280	0.8893	0.0785	0.6703	-0.0227	0.9744	-0.0068	0.9793	0.0159	0.9400	-0.0582	0.9405	-0.1357	0.5975	-0.0775	0.7875
---	---	1630189_at	0.0173	0.9435	0.1715	0.4761	-0.0534	0.8055	0.0053	0.9950	-0.0701	0.7203	-0.0754	0.6660	0.1012	0.9024	0.0636	0.8836	-0.0376	0.9282
---	---	1630190_at	0.1331	0.5369	0.0458	0.8540	0.1313	0.4916	-0.0725	0.9266	0.1883	0.4429	0.2609	0.2177	0.0864	0.9342	0.2521	0.4541	0.1657	0.6464
CG32436	CG32436	1630191_at	-0.0363	0.8855	0.0766	0.7046	0.2239	0.1988	0.2282	0.6144	-0.0197	0.9534	-0.2479	0.2230	-0.0030	0.9984	0.0297	0.9504	0.0327	0.9369
CG32762	CG32762	1630192_at	0.0967	0.6681	0.0158	0.8936	0.1481	0.5764	0.0698	0.9110	0.0451	0.8624	-0.0248	0.9187	0.0577	0.9409	-0.0024	0.9971	-0.0601	0.8461
CG34347	CG11339	1630193_at	-0.0199	0.9504	0.1753	0.2040	0.0289	0.9087	-0.1539	0.7556	-0.1442	0.5247	0.0097	0.9710	0.0332	0.9742	0.1393	0.6138	0.1061	0.7154
---	---	1630194_at	0.1475	0.5080	0.1813	0.2929	0.0513	0.8070	-0.0838	0.8817	-0.0198	0.9442	0.0639	0.7617	0.1258	0.8386	-0.0108	0.9821	-0.1367	0.6186
Or42b	Odorant receptor	1630195_at	0.0753	0.7632	0.0361	0.8056	0.0756	0.6045	0.1635	0.6823	0.1307	0.5060	-0.0328	0.8815	0.1038	0.8465	-0.0032	0.9947	-0.1070	0.6574
Bem46	Bem46	1630196_at	0.1103	0.6553	0.4048	0.3561	0.2790	0.1221	-0.2593	0.6338	-0.1972	0.4795	0.0620	0.8404	-0.1692	0.8903	0.1777	0.7425	0.3469	0.4673
---	---	1630197_at	0.1771	0.2747	-0.1807	0.3994	0.2771	0.2703	0.2393	0.4596	0.4465	0.0191	0.2072	0.1856	-0.0402	0.9679	0.0544	0.8867	0.0946	0.7564
Aly	Aly	1630198_at	-0.0129	0.9824	0.6848	0.0675	0.8190	0.0072	-0.0715	0.9610	-0.7021	0.0759	-0.6306	0.0741	-0.1729	0.8917	0.2012	0.7093	0.3741	0.4462
---	---	1630199_at	-0.0206	0.8971	-0.0019	0.9913	0.0188	0.9177	-0.0470	0.9343	0.0450	0.8335	0.0920	0.5824	-0.0334	0.9677	-0.0404	0.9040	-0.0070	0.9847
---	---	1630200_at	0.3497	0.2601	0.1403	0.5887	0.1048	0.6458	-0.1021	0.8336	0.0117	0.9676	0.1139	0.5250	-0.0780	0.9171	-0.0049	0.9935	0.0730	0.8221
CG11321	CG11321	1630201_a_at	-0.5279	0.1004	-1.2056	0.0031	-1.3292	0.0011	0.0056	0.9956	0.4755	0.1429	0.4699	0.1059	-0.0861	0.9259	-0.2416	0.4311	-0.1555	0.6389
CG8632	CG8632	1630202_a_at	-0.1160	0.5180	-0.1885	0.3636	-0.0260	0.8954	-0.2423	0.4455	0.0219	0.9282	0.2642	0.0881	-0.3981	0.6557	-0.0436	0.9407	0.3545	0.3458
Edem2	CG5682	1630203_at	1.1653	0.0833	0.8131	0.0902	0.2553	0.3884	-0.2525	0.8200	0.6863	0.1135	0.9388	0.0239	0.2503	0.9020	0.2544	0.7818	0.0041	0.9979

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG8136	CG8136	1630204_a_at	0.3898	0.1049	0.4870	0.0287	0.2774	0.1346	-0.1665	0.6321	-0.3253	0.0598	-0.1588	0.3034	0.1117	0.9057	-0.0777	0.8721	-0.1894	0.6036
---	---	1630205_at	0.0582	0.7230	-0.0012	0.9933	0.2088	0.1534	0.1221	0.6869	-0.0498	0.7681	-0.1720	0.1712	-0.0048	0.9952	-0.0757	0.7379	-0.0709	0.7502
CS-2	Chitin Synthase A	1630206_at	0.2370	0.6699	0.5513	0.5643	-1.6157	0.0004	-1.7043	0.0407	-0.0838	0.8940	1.6205	0.0027	0.3142	0.9277	-0.0121	0.9965	-0.3264	0.8177
CG16972	CG16972	1630207_at	0.4407	0.1192	0.2803	0.1660	0.3948	0.1220	0.1360	0.8738	0.2242	0.4710	0.0882	0.7910	0.0783	0.9168	0.1276	0.6517	0.0494	0.8905
ligatin	Ligatin	1630208_at	-0.1061	0.5624	0.1382	0.4552	0.3662	0.0420	0.2788	0.4694	-0.1339	0.5491	-0.4127	0.0335	-0.0371	0.9515	0.0690	0.7504	0.1061	0.5871
CG32445	CG32445	1630209_at	0.1222	0.4558	0.0425	0.7220	0.2285	0.1580	0.0511	0.9234	-0.1951	0.2227	-0.2463	0.0862	-0.0302	0.9725	-0.0605	0.8380	-0.0303	0.9199
---	---	1630210_at	0.2544	0.1748	0.0852	0.5964	0.3944	0.0876	0.0695	0.8663	0.0432	0.8104	-0.0263	0.8788	-0.1056	0.8722	-0.0122	0.9796	0.0934	0.7548
---	---	1630211_at	0.1204	0.4976	0.0386	0.8979	0.0327	0.8762	-0.1509	0.8196	-0.0526	0.8791	0.0983	0.7219	-0.1307	0.8093	-0.0546	0.8619	0.0761	0.7745
CG2065	CG2065	1630212_at	-1.1349	0.0015	-2.2579	0.0042	-2.0303	0.0002	0.6831	0.0978	1.4632	0.0005	0.7801	0.0041	0.4052	0.7366	0.4296	0.3973	0.0244	0.9764
CG1078	CG1078	1630213_at	-0.0293	0.9703	0.3048	0.5952	0.1587	0.6121	0.0114	0.9921	0.0999	0.7101	0.0885	0.7207	-0.0404	0.9923	0.4145	0.7069	0.4549	0.6683
CG7824	CG7824	1630214_a_at	-0.0793	0.7849	0.3371	0.1650	0.8197	0.0062	-0.1153	0.8402	-1.4177	0.0004	-1.3024	0.0003	-0.7195	0.3587	-0.9176	0.0509	-0.1981	0.6634
---	---	1630215_at	0.0869	0.6423	-0.0372	0.7156	0.2119	0.4174	0.1834	0.7031	-0.0213	0.9492	-0.2047	0.3092	-0.1057	0.8882	0.0181	0.9690	0.1238	0.6939
drosha	Drosha	1630216_at	-0.3094	0.1748	-0.4511	0.0479	-0.3992	0.1043	-0.0470	0.9558	0.5084	0.0337	0.5554	0.0147	-0.1338	0.8376	0.2852	0.2707	0.4190	0.1443
CG13650	CG13650	1630217_at	-1.7063	0.0048	-3.1061	0.0149	-3.4638	0.0000	-0.4811	0.4102	0.3667	0.2595	0.8478	0.0105	-0.2112	0.9342	-0.8782	0.2693	-0.6670	0.4301
CG13227	CG13227	1630218_at	0.2077	0.2885	0.0110	0.9167	0.0330	0.8872	0.1340	0.6654	0.0241	0.9062	-0.1099	0.4285	0.0112	0.9934	-0.1216	0.7378	-0.1328	0.7025
CG14837	CG14837	1630219_at	0.1629	0.4136	-0.1667	0.3556	-0.0802	0.7265	-0.0189	0.9834	0.2641	0.2036	0.2830	0.1272	-0.0518	0.9467	-0.1254	0.6209	-0.0736	0.7964
---	---	1630220_at	0.2682	0.2819	0.3451	0.1795	0.0476	0.7744	-0.1150	0.7432	-0.0046	0.9840	0.1105	0.4480	0.0752	0.9421	0.1598	0.6458	0.0846	0.8350
CG11600	CG11600	1630221_at	-0.0352	0.8464	0.0358	0.8254	0.2571	0.1820	0.0294	0.9599	-0.0415	0.8320	-0.0709	0.6552	-0.0940	0.9238	0.1077	0.7906	0.2017	0.5578
CG40317	CG40317	1630222_at	-0.0479	0.8429	0.0175	0.8638	-0.0603	0.7383	-0.1593	0.6533	-0.2039	0.2320	-0.0446	0.8178	-0.0006	0.9998	0.0517	0.9037	0.0523	0.8918
skl	Sickle	1630223_at	-0.1555	0.8533	0.3053	0.3698	0.5907	0.0461	0.0767	0.9766	-0.3808	0.5869	-0.4575	0.4521	-0.1269	0.9296	0.0363	0.9622	0.1632	0.7623
CG6410 /// DyakCG6410	CG6410	1630224_at	0.0734	0.7739	0.7707	0.0674	0.7250	0.0161	-0.0714	0.9380	-0.4654	0.0853	-0.3940	0.1014	0.1711	0.8653	0.3719	0.3424	0.2008	0.6423
---	---	1630225_at	0.3470	0.1571	0.0997	0.5244	0.2453	0.2658	-0.0715	0.9371	-0.0048	0.9899	0.0667	0.8234	-0.1677	0.8141	-0.0616	0.8852	0.1062	0.7549
CG31619	CG31619	1630226_at	-0.4401	0.0293	0.0226	0.8437	0.0025	0.9910	0.1648	0.6189	-0.1801	0.2780	-0.3449	0.0273	0.0311	0.9611	-0.0311	0.9095	-0.0621	0.7656
CG8176	CG8176	1630227_a_at	-0.4263	0.2085	0.1365	0.6639	-0.1707	0.3862	-0.1676	0.6354	-0.3094	0.0768	-0.1418	0.3773	0.2728	0.8270	0.3410	0.5259	0.0682	0.9237
---	---	1630228_at	0.1015	0.6725	0.0821	0.6546	0.0779	0.6125	-0.1257	0.8578	-0.0220	0.9532	0.1038	0.6964	0.0062	0.9952	0.0168	0.9652	0.0106	0.9775
CG9815	CG9815	1630229_at	-1.2218	0.0070	-2.4157	0.0038	-1.9790	0.0006	0.5368	0.6720	1.2335	0.0457	0.6967	0.1937	0.0909	0.9342	-0.1158	0.7955	-0.2067	0.5838
---	---	1630230_at	0.1533	0.4927	0.1341	0.2013	0.0671	0.6583	-0.1519	0.6496	-0.0520	0.7928	0.0999	0.5246	-0.0287	0.9816	-0.0380	0.9339	-0.0093	0.9846
fw	wrinkled	1630231_at	0.0491	0.9071	-0.7791	0.0902	-0.7403	0.0451	0.2269	0.8776	0.6147	0.2226	0.3878	0.4109	-0.0783	0.9630	-0.4402	0.3285	-0.3620	0.4475
---	---	1630232_at	0.0429	0.7807	-0.0533	0.7833	0.0892	0.6460	0.1020	0.8074	0.0296	0.8965	-0.0724	0.6776	0.1443	0.8461	0.0805	0.8478	-0.0638	0.8784
CG5399 /// DyakCG5399	CG5399	1630233_at	4.0661	0.0069	0.2882	0.8104	2.3731	0.0219	2.5720	0.1331	5.3763	0.0009	2.8043	0.0088	0.6544	0.8903	1.7401	0.3152	1.0857	0.5649
CG40497	CG40497	1630234_x_at	0.2091	0.4132	0.2955	0.1194	0.3216	0.0498	0.1358	0.7823	0.0106	0.9734	-0.1252	0.5270	-0.0944	0.8903	-0.0106	0.9829	0.0838	0.7827
---	---	1630235_at	0.0587	0.7676	0.1700	0.3444	0.2400	0.1881	-0.0995	0.7979	-0.1320	0.4169	-0.0325	0.8598	-0.1404	0.7697	-0.0195	0.9525	0.1208	0.5781
CG15093	CG15093	1630236_s_at	0.9632	0.2108	0.2496	0.3834	1.8829	0.0045	0.8627	0.3048	1.2631	0.0188	0.4004	0.3818	-0.7324	0.7485	0.5321	0.6080	1.2645	0.2110
Dll	Distalless	1630237_a_at	-2.3272	0.0041	-0.0907	0.6653	-1.9501	0.0089	-1.0623	0.3420	-1.8345	0.0117	-0.7722	0.1783	0.5791	0.6898	0.0463	0.9624	-0.5328	0.3825
CG11018 /// Lcp4	CG11018 /// cuticl	1630238_at	0.0057	0.9762	-0.0066	0.9730	0.0276	0.8793	0.0864	0.8589	0.0446	0.8437	-0.0417	0.8348	-0.0237	0.9816	-0.0049	0.9925	0.0189	0.9523
alphaTub67C	alpha-Tubulin	1630239_at	0.5847	0.4994	-0.7866	0.6479	-0.8269	0.3859	-0.2471	0.5564	2.3614	0.0001	2.6085	0.0000	-0.3366	0.9717	0.4938	0.8882	0.8304	0.7692
---	---	1630240_at	0.1540	0.4377	0.1875	0.1512	0.1885	0.4212	0.2065	0.6280	0.1702	0.4344	-0.0363	0.8839	0.1602	0.8427	0.2060	0.5523	0.0458	0.9198
lambdaTry	lambdaTry	1630241_at	-0.0304	0.9758	-0.0061	0.9639	-0.1498	0.6603	0.2401	0.8803	-0.2823	0.6450	-0.5224	0.2903	0.1074	0.9778	-0.2271	0.8588	-0.3344	0.7526
CG31510	CG31510	1630242_at	0.1259	0.6934	0.1866	0.7880	-0.2854	0.1757	-0.0865	0.9182	0.3363	0.1914	0.4227	0.0702	0.2929	0.8692	0.3764	0.6209	0.0835	0.9331
CG13685	CG13685	1630243_at	0.0137	0.9535	0.0964	0.7194	0.0574	0.8024	0.2914	0.6763	0.0288	0.9531	-0.2627	0.3887	0.2505	0.6695	0.0525	0.8765	-0.1980	0.4098
CG31809 /// CG31810	CG31810 /// CG3	1630244_s_at	0.8586	0.6686	-1.6806	0.0347	-1.0641	0.0009	1.1030	0.0657	1.6946	0.0011	0.5917	0.0590	0.1964	0.9852	-1.1626	0.6652	-1.3590	0.6040
shu	Shutdown	1630245_at	0.1338	0.7384	-0.4776	0.4856	-0.7024	0.0075	-0.2769	0.4599	0.6743	0.0059	0.9512	0.0008	-0.0564	0.9898	0.0139	0.9941	0.0703	0.9572
CG14199	CG14199	1630246_at	0.6797	0.0233	0.9240	0.0289	0.9771	0.0014	0.1302	0.7760	0.1313	0.5178	0.0011	0.9963	-0.0814	0.9589	0.3152	0.4833	0.3966	0.3803
CG4554	CG4554	1630247_at	0.5735	0.0737	0.0449	0.9602	0.5507	0.2256	0.5055	0.5075	1.3471	0.0059	0.8416	0.0278	0.0716	0.9841	0.7006	0.3828	0.6290	0.4569
CG5346	CG5346	1630248_at	-1.3389	0.0222	-1.2886	0.0597	-1.7978	0.0019	0.2661	0.8246	0.4065	0.3944	0.1404	0.7871	0.9068	0.5228	0.6017	0.3685	-0.3051	0.6857
CG31090	CG31090	1630249_at	-0.4229	0.3198	0.2245	0.3503	0.1724	0.6554	-0.0460	0.9777	-0.6422	0.1300	-0.5961	0.1161	0.1156	0.9306	0.0158	0.9848	-0.0998	0.8581
CG40115	CG40115	1630250_at	0.0077	0.9709	-0.1311	0.4435	-0.0285	0.8934	0.1805	0.6465	0.1289	0.5256	-0.0516	0.8102	0.1006	0.8386	0.0798	0.7374	-0.0208	0.9425
CG9624	CG9624	1630251_at	0.2299	0.2734	0.1074	0.6136	0.1095	0.6882	-0.1218	0.8217	-0.0650	0.8013	0.0568	0.8111	-0.1473	0.8909	-0.0960	0.8664	0.0513	0.9273
RpS11	Ribosomal protein	1630252_at	-0.1566	0.5235	0.2281	0.0710	-0.1267	0.3864	-0.0854	0.8908	-0.3917	0.0644	-0.3062	0.1009	0.1227	0.8609	-0.1472	0.6288	-0.2699	0.3538
---	---	1630253_at	-0.0396	0.8472	0.1898	0.3276	-0.0557	0.7437	-0.1281	0.7599	-0.1521	0.4129	-0.0239	0.9129	0.0466	0.9457	0.0639	0.8119	0.0174	0.9525
CG2964	CG2964	1630254_at	0.4427	0.1591	0.3200	0.0521	0.4956	0.0198	0.0786	0.9247	-0.1362	0.6234	-0.2148	0.3485	0.0140	0.9914	-0.0208	0.9655	-0.0348	0.9330
CG31676	CG31676	1630255_at	-1.7045	0.0387	-0.1661	0.8100	-1.6880	0.0064	-1.2570	0.0912	-2.4892	0.0006	-1.2322	0.0074	0.3008	0.9330	-1.2151	0.2729	-1.5159	0.2075
lectin-24Db	Lectin24Db	1630256_at	-2.6467	0.0364	-4.1848	0.0048	-4.6819	0.0000	-0.2705	0.7031	-0.4147	0.2030	-0.1442	0.6720	0.1670	0.9814	-1.8066	0.2195	-1.9736	0.2132
---	---	1630257_s_at	0.1796	0.3053	-0.0364	0.9078	-0.0882	0.6229	-0.0517	0.9314	0.0094	0.9733	0.0611	0.7539	0.0718	0.9340	-0.0217	0.9621	-0.0936	0.7745
GstD2	Glutathione S tran	1630258_at	1.2133	0.1883	0.1033	0.8459	1.0596	0.0598	1.0818	0.1156	2.3784	0.0006	1.2967	0.0046	0.1558	0.9781	1.3154	0.3203	1.1596	0.4016
CG31422	CG31422	1630259_at	-0.1283	0.3986	0.2183	0.3862	-0.0303	0.8977	-0.1297	0.6975	-0.0887	0.5964	0.0410	0.8130	0.1165	0.9109	0.1998	0.6144	0.0833	0.8646
CG31832	CG31832	1630260_at	0.0322	0.8621	-0.0702	0.5394	0.1520	0.3987	0.0890	0.8297	0.1154	0.4877	0.0264	0.8893	-0.1421	0.8164	-0.0077	0.9885	0.1344	0.6216
up	troponin T	1630261_a_at	-2.6020	0.0051	-1.6478	0.0361	-2.4356	0.0011	-0.5285	0.2289	-1.2459	0.0012	-0.7174	0.0084	0.2977	0.9330	-0.1887	0.9145	-0.4864	0.7015
---	---	1630262_s_at	-0.6187	0.0590	-0.2915	0.3672	-0.1537	0.6025	0.0115	0.9931	0.0204	0.9578	0.0088	0.9785	-0.1876	0.7893	0.1093	0.7595	0.2970	0.3307
CG31370	CG31370	1630263_at	0.8963	0.0190	0.2551	0.3626	0.2891	0.5109	-0.0622	0.9567	0.9143	0.0098	0.9765	0.0043	-0.0769	0.9775	0.3074	0.6694	0.3843	0.5804
RpL22-like	CG9871	1630264_at	-0.0800	0.7848	0.0575	0.6372	-0.0474	0.8052	0.0268	0.9777	-0.0851	0.7724	-0.1119	0.6586	-0.0156	0.9862	-0.0833	0.7200	-0.0677	0.7774
CG12426	CG12426	1630265_at	0.0649	0.8110	0.1291	0.5629	0.0676	0.7789	0.0012	0.9988	-0.1570	0.5184	-0.1581	0.4666	-0.0323	0.9862	0.0495	0.9425	0.0818	0.8887
Ilp4	Drosophila insulin	1630266_at	-0.1457	0.5044	-0.1320	0.4073	-0.1625	0.4125	-0.0679	0.9500	0.0928	0.8013	0.1607	0.5934	0.1896	0.7215	0.1209	0.6129	-0.0687	0.8000
CG10858	CG10858	1630267_at	-0.0330	0.8609	-0.0640	0.5805	0.2878	0.1018	0.1484	0.6844	0.1641	0.3415	0.0157	0.9404	-0.0330	0.9737	0.1753	0.4865	0.2082	0.4058
CG11781	CG11781	1630268_at	0.0431	0.8968	0.2862	0.1865	0.0919	0.6569	-0.1834	0.7749	-0.0206	0.9591	0.1628	0.5345	0.0367	0.9852	0.2994	0.5248	0.2627	0.5871
skd	blind spot	1630269_s_at	-0.0153	0.9751	-0.0893	0.6821	-0.6392	0.0685	-0.3735	0.6338	0.5973	0.1186	0.9707	0.0124	0.1372	0.9335	0.3684	0.4951	0.2313	0.6912
RpL3	Ribosomal protein	1630270_s_at	0.2906	0.0587	0.1186	0.2376	0.2043	0.1599	0.1492	0.6409	0.2092	0.1778	0.0600	0.7180	0.0687	0.8842	0.0040	0.9925	-0.0647	0.7531
Hem	kette	1630271_at	0.0757	0.7781	-0.0585	0.9244	-0.2382	0.2155	0.0402	0.9647	0.5209	0.0368	0.4807	0.0326	0.2928	0.8206	0.4455	0.4016	0.1527	0.8194
MED23	Mediator complex	1630272_at	0.3833	0.0395	0.3242	0.2397	0.4130	0.0217	-0.0597	0.9489	0.2209	0.4092	0.2806	0.2278	0.1167	0.8692	0.2990	0.2703	0.1823	0.5379

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15145	CG15145	1630273_at	-0.0005	0.9980	0.0477	0.6935	-0.1110	0.5081	0.1085	0.8475	0.1372	0.5345	0.0287	0.9086	0.0953	0.8744	0.0333	0.9291	-0.0621	0.8366
CG12099	CG12099	1630274_s_at	-0.0089	0.9783	0.9902	0.0488	0.4306	0.0830	-0.2171	0.5639	-0.2327	0.2280	-0.0156	0.9494	0.2429	0.8374	0.6721	0.1542	0.4291	0.3878
---	---	1630275_at	0.0923	0.6775	0.1285	0.4670	0.1061	0.5608	-0.0574	0.9314	0.0219	0.9385	0.0793	0.7046	0.0197	0.9851	0.0532	0.8767	0.0335	0.9194
CG5149	CG5149	1630276_at	-0.1037	0.7117	-0.0642	0.7831	-0.1398	0.5973	-0.2816	0.5357	0.1906	0.4383	0.4722	0.0333	-0.2817	0.7768	0.1924	0.6893	0.4741	0.2842
CG30048	CG30048	1630277_at	0.1367	0.6293	0.1917	0.1326	0.0037	0.9896	-0.0047	0.9956	0.0862	0.7262	0.0909	0.6805	0.1652	0.8202	0.0467	0.9193	-0.1185	0.7264
---	---	1630278_s_at	0.2163	0.5809	-0.5644	0.0846	-0.6966	0.0134	0.3467	0.5229	1.0608	0.0035	0.7141	0.0127	0.5485	0.6557	0.2771	0.6044	-0.2714	0.6152
CG40383	CG40383	1630279_at	-0.0830	0.6049	-0.0926	0.5068	-0.2674	0.1787	-0.1607	0.6424	-0.0284	0.9026	0.1323	0.3992	0.0381	0.9514	0.0225	0.9404	-0.0157	0.9523
CG9216	CG9216	1630280_s_at	-0.3358	0.3016	-0.3002	0.1152	-0.6282	0.0033	-0.3029	0.2501	-0.1536	0.3345	0.1493	0.2931	0.1099	0.9309	-0.1518	0.7554	-0.2617	0.5450
dpr12	dpr12	1630281_at	0.1908	0.4249	0.1014	0.6464	0.1057	0.5975	0.1181	0.8920	0.1211	0.7214	0.0030	0.9934	0.1587	0.7780	0.1850	0.4512	0.0263	0.9387
CG14852	CG14852	1630282_at	0.3039	0.1629	0.0427	0.6990	0.0051	0.9852	0.0682	0.9149	0.2003	0.3242	0.1321	0.4914	0.3843	0.6551	0.0597	0.9100	-0.3246	0.3601
Cdc37	cdc37 protein	1630283_at	-0.4528	0.1386	-0.0440	0.8668	0.0965	0.5764	0.0703	0.9497	0.0046	0.9916	-0.0657	0.8557	0.0415	0.9657	0.3833	0.1392	0.3417	0.2101
CG14082	CG14082	1630284_at	0.1927	0.3536	0.3830	0.1544	0.1246	0.5090	0.1246	0.7949	-0.0351	0.8953	-0.1597	0.3773	0.2439	0.6898	0.0258	0.9494	-0.2180	0.3921
RhoGAP100F	RhoGAP100F	1630285_at	-0.3208	0.1039	-0.2768	0.1238	-0.1261	0.5396	-0.0263	0.9745	-0.0161	0.9569	0.0102	0.9679	-0.2738	0.7726	0.0041	0.9970	0.2779	0.5152
AnnIX	Annexin IX	1630286_at	-0.4312	0.0543	-0.1898	0.2535	-0.4622	0.0197	-0.0536	0.9205	0.1354	0.4234	0.1890	0.1952	0.1859	0.7810	0.3245	0.2442	0.1386	0.6579
UbcD6	Ubiquitin conjugat	1630287_at	-0.1441	0.6287	0.9972	0.0235	0.9736	0.0005	-0.0531	0.9420	-0.8001	0.0031	-0.7470	0.0025	-0.1406	0.8714	0.2513	0.4729	0.3919	0.2721
CG3588	CG3588	1630288_a_at	-0.5164	0.1132	0.0099	0.9661	-0.3723	0.1047	-0.2928	0.5328	-0.7348	0.0094	-0.4420	0.0488	-0.1007	0.9421	-0.1799	0.7124	-0.0792	0.8932
CG6435	CG6435	1630289_at	-2.4796	0.0026	-0.1073	0.5886	-1.5209	0.0125	-1.6685	0.0925	-2.4960	0.0019	-0.8276	0.1102	-0.4063	0.7215	-0.3062	0.5415	0.1001	0.8779
mld	molting defective	1630290_at	-0.7014	0.0238	-2.5548	0.0014	-2.4192	0.0005	0.4318	0.6615	1.6651	0.0046	1.2334	0.0107	0.2454	0.7220	-0.2965	0.2981	-0.5419	0.1000
MtnB	Metallothionein B	1630291_at	-2.7239	0.0164	-4.7446	0.0020	-4.4743	0.0001	0.0502	0.9818	1.4077	0.0146	1.3575	0.0105	-0.0903	0.9816	-0.5123	0.5905	-0.4221	0.6656
dve	defective proventr	1630292_a_at	0.2351	0.2916	0.0446	0.6734	0.0947	0.5820	-0.0269	0.9759	-0.0315	0.9183	-0.0046	0.9870	-0.0683	0.9365	-0.0622	0.8724	0.0061	0.9890
CG32454	CG32454	1630293_at	0.0335	0.8884	-0.0021	0.9912	0.2679	0.1226	-0.0234	0.9819	-0.1061	0.7046	-0.0828	0.7548	-0.0740	0.9407	0.0228	0.9646	0.0968	0.7940
CG15733	CG15733	1630294_at	0.1093	0.5182	0.0794	0.4510	0.2530	0.1126	-0.0205	0.9803	-0.0365	0.8952	-0.0161	0.9503	-0.0731	0.9238	0.0884	0.7784	0.1615	0.5498
---	---	1630295_at	-0.0294	0.9432	-0.2002	0.2255	-0.1403	0.6369	-0.1752	0.8578	-0.0223	0.9676	0.1529	0.6776	-0.3561	0.5200	-0.3120	0.2217	0.0441	0.9040
---	---	1630296_at	-0.1939	0.3098	0.1176	0.6430	0.1107	0.6549	0.2543	0.6574	0.0133	0.9750	-0.2410	0.3392	-0.0384	0.9648	-0.0224	0.9526	0.0160	0.9628
---	---	1630297_at	0.2587	0.0833	0.2263	0.3281	0.2348	0.2717	-0.1321	0.7432	-0.1816	0.3107	-0.0495	0.8041	0.1094	0.8533	0.0598	0.8595	-0.0496	0.8812
CG30496	CG30496	1630298_at	0.1750	0.3107	0.6787	0.1026	0.7559	0.0007	0.0380	0.9627	-0.1033	0.6746	-0.1413	0.4956	-0.0175	0.9914	0.4605	0.1886	0.4781	0.2069
CG17325	CG17325	1630299_at	-0.3564	0.7642	-0.1346	0.7992	-1.8377	0.0027	-1.3852	0.1327	-1.6587	0.0094	-0.2735	0.6276	0.2679	0.9640	-1.2900	0.4203	-1.5579	0.3467
CG4706	CG4706	1630300_at	0.0716	0.7895	0.1340	0.2433	0.0322	0.9011	-0.1959	0.6558	-0.1012	0.6746	0.0948	0.6660	-0.0077	0.9943	0.0207	0.9590	0.0284	0.9344
CG12204	CG12204	1630301_at	-0.2863	0.2413	-0.4739	0.1110	-0.1677	0.3516	0.2677	0.4115	-0.2268	0.2077	-0.4945	0.0085	-0.0542	0.9717	-0.2701	0.4996	-0.2159	0.6053
CG8562	CG8562	1630302_at	0.0406	0.9535	0.0191	0.9173	-0.1223	0.5824	0.0803	0.8732	-0.0665	0.7479	-0.1468	0.3601	0.1355	0.9514	-0.1153	0.9105	-0.2508	0.7442
---	---	1630303_at	0.0781	0.6084	-0.0445	0.6960	0.0609	0.8028	0.0525	0.9436	0.0114	0.9727	-0.0411	0.8686	0.0370	0.9610	-0.0175	0.9599	-0.0545	0.8387
Dhc93AB	dynein-related hea	1630304_at	0.2007	0.3173	0.1391	0.4200	0.0871	0.5975	-0.0811	0.8732	-0.0304	0.8999	0.0506	0.7983	0.0823	0.9246	0.0446	0.9220	-0.0376	0.9252
fkh	Fork Head	1630305_at	0.8893	0.3047	-0.5070	0.0388	-0.5148	0.0720	0.0018	0.9988	0.1620	0.5694	0.1602	0.5336	-0.0120	0.9982	-0.8670	0.4258	-0.8550	0.4464
CG15012 /// DyakCG15012	CG15012	1630306_at	0.0657	0.7649	-0.0527	0.6296	-0.1382	0.4659	0.0266	0.9647	-0.0080	0.9743	-0.0346	0.8516	0.0001	0.9999	-0.1921	0.5711	-0.1922	0.5746
Cpr64Ad	CG1259	1630307_at	0.1432	0.6838	0.0190	0.8741	0.1835	0.3421	0.0121	0.9893	0.1640	0.4468	0.1519	0.4361	0.0629	0.9587	0.1033	0.8127	0.0404	0.9319
CG3726	CG3726	1630308_at	-1.3681	0.0113	-1.1841	0.2162	-1.1481	0.0073	0.4537	0.4861	0.4840	0.1626	0.0303	0.9451	0.4234	0.8699	0.6170	0.5708	0.1935	0.8918
---	---	1630309_s_at	0.0916	0.9014	0.1723	0.5887	-0.7013	0.2416	-0.4345	0.7121	-0.0744	0.9214	0.3601	0.4809	0.3839	0.8740	-0.0418	0.9814	-0.4256	0.6861
CG32751	CG32751	1630310_at	5.7386	0.0005	3.4483	0.0022	7.1485	0.0000	2.8619	0.0153	2.3404	0.0031	-0.5215	0.3498	-0.7271	0.3517	0.0637	0.9215	0.7908	0.0905
dpr16	dpr16	1630311_at	0.0361	0.8757	-0.0270	0.8012	0.0141	0.9474	-0.0038	0.9956	0.0199	0.9284	0.0237	0.9003	-0.0860	0.9108	-0.1134	0.7162	-0.0274	0.9431
Ard1	Ard1	1630312_s_at	0.0669	0.7869	0.6076	0.0794	0.5971	0.0082	0.0867	0.8743	-0.1580	0.4240	-0.2447	0.1509	0.1721	0.8541	0.3668	0.3292	0.1947	0.6394
CG15086	CG15086	1630313_a_at	0.0445	0.8800	-0.1127	0.6028	0.2691	0.1679	0.1330	0.8836	-0.0626	0.8793	-0.1956	0.5130	-0.0274	0.9829	-0.1176	0.7299	-0.0902	0.8003
CG31141	CG31141	1630314_at	0.2559	0.2045	-0.0411	0.6893	-0.2303	0.1737	-0.0789	0.8707	0.0152	0.9518	0.0941	0.5769	0.0048	0.9964	-0.1712	0.4696	-0.1760	0.4659
CG8214	CG8214	1630315_at	-0.1946	0.3230	-0.0125	0.9093	-0.1118	0.6288	0.0640	0.9311	-0.0454	0.8752	-0.1094	0.6215	-0.0416	0.9467	0.0281	0.9241	0.0697	0.7484
CG14814	CG14814	1630316_a_at	-0.2235	0.7573	-1.3426	0.2783	-1.8430	0.0142	-0.2576	0.7127	1.3458	0.0021	1.6035	0.0006	-0.0031	0.9998	0.1709	0.9463	0.1740	0.9369
CG5127	CG5127	1630317_at	-0.1426	0.5248	0.1723	0.3222	0.3555	0.0305	-0.0341	0.9666	-0.5191	0.0230	-0.4850	0.0191	-0.0473	0.9611	0.0007	0.9996	0.0480	0.8984
MED7	Mediator complex	1630318_at	0.0440	0.8365	0.0561	0.7599	-0.0666	0.7900	0.1181	0.7888	0.0799	0.7014	-0.0382	0.8560	0.1008	0.8870	0.1236	0.6760	0.0227	0.9514
---	---	1630319_at	0.4911	0.0255	-0.1936	0.3485	0.1233	0.4135	0.2479	0.5444	0.2067	0.3373	-0.0412	0.8682	-0.0853	0.9142	-0.2204	0.4258	-0.1352	0.6519
betaTry	betaTrypsin	1630320_at	1.7496	0.5023	-0.2873	0.1812	0.2360	0.7074	0.2707	0.6557	0.2019	0.5137	-0.0689	0.8372	-0.4601	0.9737	-2.0105	0.5800	-1.5504	0.6817
---	---	1630321_at	-0.0487	0.8161	0.1555	0.4523	-0.0598	0.6954	-0.0105	0.9915	-0.1594	0.4601	-0.1489	0.4453	-0.0408	0.9489	-0.0425	0.8760	-0.0017	0.9959
Apc2	Epithelial Adenom	1630322_at	0.4196	0.1996	-0.2858	0.2631	0.1289	0.3987	0.2891	0.6122	0.7241	0.0198	0.4350	0.0898	0.0322	0.9816	0.0567	0.9081	0.0245	0.9578
CG6914	CG6914	1630323_at	0.1397	0.3954	0.0533	0.8101	0.0774	0.6619	-0.1164	0.7780	0.0571	0.7847	0.1735	0.2631	-0.0718	0.9036	-0.0591	0.8380	0.0126	0.9695
l(2)k07824	lethal (2) k07824	1630324_at	0.6701	0.0434	-0.0653	0.9174	0.1339	0.6639	0.2600	0.6354	1.0074	0.0035	0.7474	0.0081	0.1030	0.9640	0.2913	0.6686	0.1884	0.8017
CG15262	CG15262	1630325_at	0.3339	0.0644	0.0590	0.6695	0.0302	0.8679	0.0331	0.9603	0.0982	0.6186	0.0651	0.7340	0.1917	0.8076	0.0435	0.9341	-0.1482	0.6817
Obp47b	Odorant-binding p	1630326_at	0.1605	0.4281	0.0716	0.7990	0.3816	0.0422	0.1562	0.7950	-0.0219	0.9523	-0.1781	0.4399	-0.0563	0.9330	-0.1526	0.4873	-0.0962	0.6857
CG14684	CG14684	1630327_at	0.1215	0.5948	-0.0303	0.7658	0.1126	0.4980	0.1251	0.7556	0.1930	0.2670	0.0679	0.7121	-0.1192	0.8700	-0.1498	0.6344	-0.0305	0.9405
---	---	1630328_at	0.1454	0.4954	0.1506	0.4044	0.0809	0.6248	-0.0413	0.9477	0.0212	0.9323	0.0625	0.7443	-0.0652	0.9224	0.0821	0.7562	0.1473	0.5271
CG1716 /// DrepCG1716	CG1716	1630329_at	0.0572	0.9082	-0.5543	0.4018	-0.5182	0.0049	-0.0237	0.9808	0.5480	0.0295	0.5717	0.0156	-0.1712	0.9474	-0.0946	0.9407	0.0767	0.9444
CG31112	CG31112	1630330_at	-0.0097	0.9735	0.0553	0.5991	0.0413	0.7968	-0.0395	0.9629	-0.0727	0.7932	-0.0332	0.9031	0.0095	0.9946	0.0222	0.9666	0.0127	0.9818
---	---	1630331_at	0.0383	0.8975	0.0596	0.6686	-0.0478	0.7509	-0.1699	0.6908	-0.1061	0.6308	0.0639	0.7705	-0.0422	0.9701	-0.0640	0.8765	-0.0219	0.9584
CG3652	CG3652	1630332_at	1.0740	0.0056	1.3098	0.0134	1.0046	0.0002	-0.0411	0.9463	0.1954	0.2581	0.2365	0.1229	0.2576	0.7644	0.3801	0.2753	0.1226	0.7734
Cpr50Ca	CG13338	1630333_at	1.8649	0.0015	1.3115	0.0815	1.4062	0.0100	-0.0920	0.8462	-0.0753	0.7073	0.0166	0.9389	-0.1480	0.9487	-0.3477	0.6414	-0.1997	0.8161
CG15406	CG15406	1630334_at	-0.4902	0.2311	0.0922	0.6053	0.4955	0.0148	0.2778	0.8028	-0.6901	0.1197	-0.9680	0.0233	-0.1222	0.8906	-0.0694	0.8858	0.0528	0.9093
CG7084	CG7084	1630335_s_at	-0.8952	0.3705	0.5986	0.2215	-0.5052	0.2199	-0.7211	0.5402	-2.1596	0.0046	-1.4385	0.0168	0.1596	0.9565	-0.5881	0.4958	-0.7477	0.3844
---	---	1630336_at	0.1955	0.4545	-0.0166	0.9536	-0.3924	0.0507	-0.1750	0.6041	0.2214	0.1902	0.3964	0.0172	-0.0058	0.9977	-0.0026	0.9989	0.0032	0.9973
CG15927	CG15927	1630337_at	0.1438	0.3193	-0.1071	0.5123	-0.0373	0.8423	0.0114	0.9883	0.1194	0.5480	0.1080	0.5527	-0.1131	0.8592	-0.0901	0.7747	0.0230	0.9492
---	---	1630338_at	0.0593	0.8314	0.1630	0.5965	0.1987	0.2646	-0.2266	0.4704	-0.1968	0.2455	0.0298	0.8815	-0.0522	0.9721	0.0748	0.8938	0.1270	0.7755
---	---	1630339_at	-0.3578	0.3144	0.1360	0.5760	0.0451	0.8508	-0.1206	0.9229	-0.2668	0.5025	-0.1463	0.7151	-0.0615	0.9657	0.0812	0.8846	0.1428	0.7492
CG11927	CG11927	1630340_at	-0.2647	0.3439	-0.2172	0.0986	-0.2215	0.3655	0.0679	0.8721	0.3345	0.0302	0.2666	0.0477	0.0407	0.9816	0.2320	0.5966	0.1913	0.6694
mbt	mushroom bodies	1630341_at	-0.6452	0.0476	-0.5420	0.2777	-0.4342	0.1869	-0.0090	0.9956	-0.0779	0.8576	-0.0689	0.8588	-0.1955	0.9088	-0.0481	0.9620	0.1474	0.8498

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG3631	CG3631	1630342_s_at	-0.0117	0.9815	0.7275	0.0774	0.3727	0.1267	-0.3603	0.4434	-0.2661	0.3035	0.0942	0.7317	-0.0229	0.9942	0.6035	0.3953	0.6264	0.3921
CG3679	CG3679	1630343_at	0.5523	0.4675	-1.3376	0.1196	-1.7208	0.0272	-0.6980	0.0899	2.1784	0.0001	2.8764	0.0000	-0.6317	0.8714	-0.3262	0.8845	0.3055	0.8862
CG6724 /// DyakCG6724	CG6724	1630344_at	0.6749	0.0682	0.3715	0.4909	0.6032	0.0250	0.3630	0.5848	0.9595	0.0120	0.5964	0.0533	0.1876	0.9032	0.7811	0.1517	0.5935	0.3027
CG32170 /// DereCG32170	CG32170	1630345_at	-0.9348	0.0158	-0.5754	0.0409	-1.1556	0.0077	-0.0363	0.9744	-0.0252	0.9515	0.0111	0.9738	0.3238	0.7215	0.1750	0.6768	-0.1488	0.7317
CG18477 /// CG31780	CG31780 /// CG1	1630346_at	-0.0729	0.6528	0.2805	0.0942	-0.0367	0.8414	-0.0461	0.9418	-0.3257	0.0735	-0.2796	0.0839	0.4028	0.4494	0.2925	0.2733	-0.1102	0.7264
CG5589 /// DmirCG5589	CG5589	1630347_at	0.4892	0.2379	0.3346	0.3450	0.4233	0.1629	0.1599	0.8671	0.4915	0.1437	0.3317	0.2770	0.0935	0.9679	0.4802	0.4359	0.3866	0.5535
CG5059	CG5059	1630348_a_at	0.6284	0.0574	1.0882	0.0220	0.2533	0.2706	-0.3264	0.4532	-0.3852	0.1025	-0.0588	0.8271	0.3435	0.7726	-0.0459	0.9558	-0.3893	0.4597
---	---	1630349_at	-0.1020	0.4802	-0.0377	0.7282	-0.0267	0.8849	-0.0181	0.9803	-0.0334	0.8892	-0.0153	0.9458	0.0213	0.9781	0.0035	0.9935	-0.0177	0.9453
---	---	1630350_at	0.2380	0.3470	0.0210	0.8366	0.0305	0.9106	0.0902	0.8432	0.0965	0.6015	0.0064	0.9761	-0.0577	0.9587	-0.0135	0.9814	0.0443	0.9173
CG31229	CG31229	1630351_at	-0.0936	0.6870	-0.3312	0.3636	-0.2310	0.1284	0.2767	0.4128	0.6198	0.0060	0.3431	0.0461	0.1180	0.9087	0.3361	0.3424	0.2181	0.5722
CG13445	CG13445	1630352_at	0.2908	0.2663	0.1424	0.4374	0.0582	0.7789	-0.0557	0.9221	0.1926	0.2644	0.2482	0.1055	0.0020	0.9994	-0.0376	0.9462	-0.0397	0.9344
CG31178 /// DsimCG31178	CG31178	1630353_at	-0.0877	0.5740	-0.0713	0.7915	0.2694	0.1425	0.1066	0.8716	0.0711	0.8010	-0.0355	0.8977	-0.1347	0.8141	-0.0977	0.7196	0.0370	0.9096
CG13353	CG13353	1630354_at	0.2805	0.1297	0.9239	0.1455	0.7520	0.0043	0.0299	0.9649	-0.1890	0.3021	-0.2189	0.1762	0.3712	0.7475	0.4059	0.4021	0.0346	0.9614
CG16890	CG16890	1630355_a_at	-0.1676	0.5249	0.1453	0.5364	0.5012	0.0303	0.2145	0.5072	-0.0692	0.7303	-0.2837	0.0674	0.0158	0.9914	0.2713	0.3856	0.2555	0.4329
CG13881 /// DmirCG13881	CG13881	1630356_at	0.1424	0.3132	0.2561	0.2706	-0.1098	0.5086	-0.1698	0.6489	-0.0819	0.6924	0.0879	0.6331	-0.0006	0.9998	0.0350	0.9046	0.0355	0.8918
---	---	1630357_at	0.1554	0.5241	0.0999	0.7204	0.3271	0.0751	-0.0706	0.8864	-0.1926	0.2504	-0.1220	0.4373	-0.3082	0.7872	-0.2577	0.6230	0.0504	0.9412
CG2162	CG2162	1630358_at	-0.5821	0.0460	0.4230	0.0934	-0.1220	0.6717	-0.2629	0.5756	-0.8615	0.0043	-0.5987	0.0132	0.4103	0.6749	0.1075	0.8432	-0.3028	0.4657
CG31810	CG31810	1630359_at	0.9990	0.6164	-3.1344	0.0158	-0.6149	0.0173	3.2855	0.0031	3.4129	0.0002	0.1274	0.8039	0.2990	0.9742	-1.2109	0.6232	-1.5099	0.5314
CG8202	CG8202	1630360_a_at	0.2491	0.1971	0.1386	0.2397	0.1903	0.3188	0.1465	0.6506	0.0149	0.9492	-0.1315	0.3599	0.1135	0.8903	-0.0212	0.9665	-0.1347	0.6956
nkd	Naked	1630361_at	-0.7303	0.1116	-0.5980	0.1230	-1.1407	0.0058	-0.1752	0.8508	-0.1802	0.6347	-0.0050	0.9906	0.3594	0.8049	-0.0346	0.9734	-0.3940	0.5399
CG18139	CG18139	1630362_at	0.3709	0.0494	0.7550	0.0372	0.5494	0.0341	0.0776	0.9319	-0.2079	0.4604	-0.2855	0.2386	0.2050	0.7387	0.1299	0.6472	-0.0751	0.8178
CG10469	CG10469	1630363_at	0.0348	0.9008	1.0187	0.1181	0.8125	0.0940	-0.5148	0.5156	-1.2264	0.0104	-0.7115	0.0610	-0.2876	0.8395	-0.3051	0.6282	-0.0176	0.9849
---	---	1630364_x_at	0.2019	0.2201	0.0865	0.6117	-0.0469	0.8662	0.0433	0.9603	0.0478	0.8770	0.0045	0.9875	0.0516	0.9447	0.0514	0.8724	-0.0001	0.9997
CG33330 /// CG3563	CG3563 /// CG33	1630365_at	0.2486	0.6267	-0.0271	0.9665	-0.6199	0.0447	-0.1706	0.7857	0.4828	0.0660	0.6534	0.0125	0.5313	0.7423	0.1314	0.8981	-0.3999	0.5814
CG31149	CG31149	1630366_at	-0.1035	0.6939	-0.3017	0.3634	-0.6052	0.0392	0.1687	0.8439	0.4318	0.1701	0.2631	0.3676	0.0224	0.9891	-0.0656	0.8988	-0.0879	0.8402
---	---	1630367_at	0.2970	0.2482	0.0018	0.9899	-0.2164	0.1692	-0.1087	0.7556	0.0470	0.7990	0.1557	0.2493	0.0917	0.9030	-0.1743	0.5414	-0.2661	0.3457
CG7856	CG7856	1630368_at	0.1112	0.4492	0.0979	0.5283	-0.0697	0.7718	-0.0583	0.9472	-0.0467	0.8899	0.0116	0.9704	0.1185	0.8217	0.0331	0.9225	-0.0853	0.7338
CG3847	CG3847	1630369_a_at	-0.3382	0.2845	1.0264	0.0200	0.9097	0.0010	-0.0705	0.9507	-0.9748	0.0084	-0.9042	0.0071	-0.0625	0.9589	0.4197	0.2158	0.4823	0.1912
CG9796 /// DyakCG9796	CG9796	1630370_at	0.0104	0.9620	0.5133	0.0386	0.6630	0.0041	0.3174	0.1732	0.2183	0.1205	-0.0991	0.4588	0.1656	0.8444	0.4347	0.1948	0.2691	0.4518
---	---	1630371_at	0.1259	0.6217	0.0067	0.9713	-0.3280	0.0417	-0.0101	0.9909	0.1189	0.5648	0.1290	0.4792	0.2210	0.7220	0.0956	0.7626	-0.1254	0.6604
---	---	1630372_s_at	-0.3814	0.6187	0.0247	0.9484	-0.5834	0.1172	-0.0610	0.9627	-0.1098	0.7966	-0.0487	0.9078	0.4197	0.8692	0.0804	0.9620	-0.3394	0.7767
CG4935	CG4935	1630373_at	-0.0278	0.9071	0.0872	0.4996	0.3948	0.0833	-0.3093	0.5511	-0.2118	0.4472	0.0975	0.7355	-0.7011	0.3926	-0.1453	0.7985	0.5558	0.2319
Cyp308a1	Cyp308a1	1630374_at	0.1936	0.4207	-0.0162	0.9228	0.0312	0.8793	0.0064	0.9956	-0.1198	0.6789	-0.1262	0.6248	0.0168	0.9914	-0.0445	0.9328	-0.0613	0.8923
Mec2	Mec2	1630375_at	-1.4213	0.1188	-3.3522	0.0632	-2.8230	0.0111	0.9900	0.3732	1.6901	0.0150	0.7001	0.2120	0.2804	0.9717	-0.1631	0.9614	-0.4434	0.8689
onecut	D-onecut	1630376_at	-0.2203	0.2429	-0.1506	0.1502	-0.0723	0.7207	0.1341	0.7604	-0.0739	0.7399	-0.2080	0.2185	-0.0654	0.9294	-0.0765	0.8005	-0.0111	0.9767
CG31928	CG31928	1630377_at	0.3079	0.2847	-0.0254	0.8012	-0.2537	0.3231	-0.1253	0.8773	0.1616	0.5942	0.2869	0.2514	0.0940	0.8973	-0.1505	0.6035	-0.2445	0.3807
frtz	fritz	1630378_at	0.0392	0.8874	-0.0802	0.7876	-0.2762	0.2541	-0.0473	0.9538	0.1915	0.4042	0.2387	0.2327	0.0617	0.9742	0.0471	0.9506	-0.0146	0.9849
CG12885	CG12885	1630379_s_at	0.3082	0.1323	0.3589	0.3232	0.3884	0.1281	0.0423	0.9445	-0.1291	0.4767	-0.1714	0.2741	0.0031	0.9984	-0.0592	0.8923	-0.0623	0.8772
CG3837	CG3837	1630380_at	-0.8475	0.0730	-1.6142	0.0088	-1.2892	0.0015	0.2095	0.7794	0.1228	0.7372	-0.0867	0.8040	0.0038	0.9990	-0.5483	0.2505	-0.5521	0.2798
Octbeta3R	Octbeta3R	1630381_at	0.1187	0.5188	0.0479	0.8397	0.4235	0.0273	0.1307	0.7327	-0.0415	0.8513	-0.1722	0.2629	-0.1436	0.7772	-0.1109	0.6425	0.0326	0.9152
---	---	1630382_at	0.0271	0.8890	0.0604	0.6734	-0.2190	0.1404	-0.1832	0.6936	0.0503	0.8581	0.2335	0.2252	0.0539	0.9457	0.0691	0.8258	0.0153	0.9659
---	---	1630383_at	-0.1129	0.6842	0.2475	0.0840	0.0692	0.7905	-0.0304	0.9763	-0.2698	0.2904	-0.2395	0.2964	0.1623	0.8122	0.1240	0.6971	-0.0383	0.9215
CG3430	CG3430	1630384_at	0.0977	0.5879	0.0128	0.9254	0.2954	0.1151	-0.0237	0.9711	0.5346	0.0068	0.5583	0.0033	-0.4374	0.5869	0.3260	0.3526	0.7634	0.0710
---	---	1630385_at	0.3041	0.1616	0.3110	0.2049	0.2696	0.1600	-0.0665	0.8915	-0.0503	0.8027	0.0163	0.9363	0.0586	0.9405	0.1305	0.6152	0.0719	0.8101
Upf3	Upf3	1630386_at	0.1023	0.5413	0.1394	0.2541	0.2425	0.1079	-0.0525	0.9376	0.0286	0.9173	0.0811	0.6944	-0.1399	0.7324	0.1107	0.5460	0.2506	0.1784
CG34031	CG34031	1630387_at	0.1693	0.6647	0.1225	0.2870	-0.1057	0.5625	-0.2085	0.5735	-0.0688	0.7627	0.1397	0.4339	0.0305	0.9835	0.0146	0.9821	-0.0159	0.9769
CG31050	CG31050	1630388_at	0.2826	0.0982	0.0305	0.8931	0.3383	0.1043	0.0489	0.9375	0.0438	0.8528	-0.0051	0.9824	-0.1651	0.7485	-0.1723	0.4301	-0.0072	0.9846
CG13178	CG13178	1630389_at	-0.0907	0.6407	0.0680	0.5799	-0.1200	0.5293	-0.0630	0.9108	-0.0852	0.6746	-0.0222	0.9193	0.0005	0.9998	0.0310	0.9405	0.0305	0.9331
CG5602	CG5602	1630390_at	-0.6725	0.2052	-0.8922	0.0874	-0.5268	0.1156	-0.2829	0.3812	0.2598	0.1527	0.5427	0.0058	-0.6294	0.7046	-0.0832	0.9404	0.5461	0.4434
---	---	1630391_at	0.0003	0.9991	0.0953	0.4843	0.0115	0.9511	-0.0423	0.9398	-0.1124	0.5129	-0.0700	0.6800	0.0597	0.9309	0.0456	0.8879	-0.0140	0.9664
CG10481	CG10481	1630392_at	0.1262	0.6602	0.0519	0.7680	0.0292	0.9136	-0.0547	0.9532	-0.0216	0.9548	0.0331	0.9157	0.0889	0.8874	0.0625	0.8444	-0.0263	0.9361
Hrs	Hepatocyte growtl	1630393_a_at	-0.4501	0.3219	-0.0240	0.9762	0.4599	0.1096	0.2949	0.4859	-0.2498	0.2708	-0.5447	0.0150	-0.1705	0.9545	0.2491	0.8307	0.4196	0.6596
---	---	1630394_at	-0.0522	0.8129	0.0414	0.7981	0.0721	0.6965	-0.0218	0.9715	-0.0170	0.9394	0.0048	0.9799	0.0281	0.9689	0.0025	0.9951	-0.0256	0.9217
eIF2B-beta	eIF2B-beta	1630395_at	0.4066	0.0349	1.0286	0.0109	0.9425	0.0003	0.0222	0.9675	-0.0414	0.8163	-0.0636	0.6704	0.2196	0.7215	0.5581	0.0512	0.3386	0.2073
CG4360	CG4360	1630396_at	0.2481	0.5769	0.9623	0.1074	0.6876	0.0169	-0.1719	0.7753	0.0399	0.9093	0.2118	0.3610	0.2456	0.8993	0.7941	0.2503	0.5485	0.4587
CG13625	CG13625	1630397_at	0.1938	0.2623	0.0145	0.9596	-0.0564	0.8703	-0.1978	0.6796	0.5013	0.0323	0.6991	0.0047	-0.0732	0.9705	0.4198	0.3994	0.4930	0.3406
CG6746 /// DyakCG6746	CG6746	1630398_at	0.3547	0.2033	-0.1294	0.6860	0.4142	0.0442	0.2259	0.5054	-0.0882	0.6662	-0.3141	0.0550	-0.3352	0.7493	-0.5794	0.1821	-0.2442	0.6132
Cad99C	Cad99C	1630399_at	-1.2759	0.0182	-1.5142	0.0079	-1.1894	0.0058	0.3619	0.7093	0.3707	0.4170	0.0088	0.9874	0.1859	0.8743	-0.0541	0.9420	-0.2401	0.6296
pxb	baikal	1630400_at	0.6554	0.0597	0.4432	0.1565	0.8918	0.0131	0.2055	0.7630	0.3218	0.2729	0.1163	0.7065	0.0314	0.9884	0.3025	0.5352	0.2712	0.5872
---	---	1630401_s_at	0.2278	0.2432	0.0684	0.6893	0.0169	0.9556	-0.0745	0.9067	0.0380	0.8906	0.1125	0.5811	-0.0717	0.9374	-0.1945	0.5119	-0.1228	0.7022
l(2)01289	Fusion2	1630402_at	-2.2964	0.0067	-2.7839	0.0296	-3.5664	0.0001	-0.2720	0.3793	0.2176	0.2124	0.4895	0.0078	0.3638	0.9142	-0.3924	0.7925	-0.7562	0.5476
CG12746	CG12746	1630403_s_at	0.0189	0.9417	-0.4947	0.2470	0.2446	0.3686	0.4348	0.0965	0.5713	0.0035	0.1365	0.3297	-0.1496	0.9298	0.0108	0.9935	0.1603	0.8151
CG34112	CG34112	1630404_at	1.5145	0.0091	1.5554	0.0395	3.3787	0.0000	0.7617	0.0660	-0.1512	0.5586	-0.9129	0.0019	-0.7551	0.5754	0.0118	0.9935	0.7670	0.2189
CycC	cyclin C	1630405_at	-0.0339	0.9107	0.1299	0.4185	0.4916	0.0185	0.0376	0.9491	-0.1051	0.5527	-0.1426	0.3438	-0.2478	0.7901	0.0300	0.9634	0.2778	0.4997
kappaB-Ras	kappaB-Ras	1630406_at	-0.0156	0.9628	0.0120	0.9642	0.1840	0.1914	0.1292	0.6506	0.4070	0.0099	0.2778	0.0314	0.0144	0.9934	0.4559	0.2380	0.4414	0.2837
CG1724	CG1724	1630407_at	0.2131	0.2024	0.0672	0.6709	0.0579	0.7239	0.0544	0.9199	0.0786	0.6768	0.0242	0.9034	0.0737	0.9441	-0.0229	0.9650	-0.0966	0.8040
CG7033 /// DyakCG7033	CG7033	1630408_s_at	0.0078	0.9868	-0.0621	0.8546	-0.4032	0.0696	0.1411	0.8609	0.5927	0.0458	0.4515	0.0817	0.4363	0.6955	0.5163	0.2505	0.0801	0.9016
CG9938	CG9938	1630409_at	0.0618	0.8506	-0.0293	0.9591	-0.7315	0.0530	-0.6307	0.1925	0.2845	0.3393	0.9152	0.0047	-0.1299	0.9581	0.0558	0.9615	0.1858	0.8307
CG32163	CG32163	1630410_at	0.0039	0.9871	0.5425	0.0541	0.6866	0.0049	-0.1659	0.7409	-0.7283	0.0062	-0.5624	0.0120	-0.1392	0.7810	-0.0706	0.7963	0.0686	0.7921

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9945	CG9945	1630411_at	-0.0572	0.8989	0.5191	0.0199	0.3290	0.1495	-0.0928	0.9030	-0.5054	0.0466	-0.4125	0.0655	0.0767	0.9626	0.0258	0.9734	-0.0509	0.9360
CG16873	CG16873	1630412_at	-1.6035	0.0178	-2.0283	0.0057	-1.8463	0.0001	-0.5575	0.2692	-0.7121	0.0249	-0.1546	0.6073	-0.6412	0.7305	-1.0562	0.1718	-0.4150	0.6294
---	---	1630413_s_at	0.0107	0.9528	0.0025	0.9836	0.1889	0.2536	0.0905	0.8097	-0.0506	0.7799	-0.1410	0.2978	0.0672	0.8940	0.1122	0.5711	0.0450	0.8533
CG13603 /// DyakCG13603	CG13603	1630414_at	-0.5981	0.0517	-0.2729	0.1526	0.0201	0.9036	-0.0411	0.9387	-0.7678	0.0011	-0.7267	0.0008	-0.2666	0.7726	-0.3309	0.3992	-0.0643	0.9065
Alh	Alhambra	1630415_at	-0.4343	0.3594	-1.3741	0.0202	-1.2433	0.0003	0.2251	0.7775	0.9528	0.0114	0.7277	0.0226	0.0587	0.9816	-0.1000	0.9053	-0.1587	0.8161
kin17	kin17	1630416_at	-0.3030	0.1259	-0.2229	0.4262	-0.0863	0.6256	-0.0053	0.9956	-0.1008	0.7279	-0.0955	0.7173	-0.1220	0.8692	-0.0517	0.9085	0.0704	0.8529
Spn	d-spinophilin	1630417_at	1.2765	0.0628	-0.1785	0.6723	0.0341	0.9533	0.1112	0.8856	1.0819	0.0023	0.9707	0.0022	-0.0205	0.9964	-0.1033	0.9507	-0.0827	0.9555
CG31374	CG31374	1630418_s_at	-2.6361	0.0010	-2.4591	0.0070	-2.5767	0.0000	-0.3284	0.4317	-0.8676	0.0033	-0.5392	0.0166	-0.2527	0.8972	-0.5501	0.4509	-0.2974	0.7177
CG7300	CG7300	1630419_a_at	0.4094	0.6863	0.0623	0.8051	-1.4047	0.0179	-1.0257	0.2721	-0.6530	0.2362	0.3727	0.4776	0.2793	0.9318	-0.8228	0.4352	-1.1021	0.3079
---	---	1630420_x_at	0.0046	0.9877	-0.3445	0.1685	-0.2059	0.3627	-0.0243	0.9860	0.3663	0.2628	0.3906	0.1787	-0.0731	0.8909	-0.0342	0.9120	0.0390	0.8880
CG33170	CG33170	1630421_at	-0.4965	0.1140	0.2784	0.2526	-0.2632	0.3151	-0.4875	0.3050	-0.9339	0.0058	-0.4464	0.0755	-0.0737	0.9514	-0.1728	0.6593	-0.0991	0.8247
CG18231	CG18231	1630422_at	0.0209	0.9056	-0.0174	0.8768	-0.1194	0.5933	-0.1777	0.7728	-0.0112	0.9780	0.1665	0.5035	-0.1978	0.7215	-0.0680	0.8212	0.1299	0.5995
CG4996	CG4996	1630423_at	0.0243	0.9535	0.0156	0.9684	-0.4306	0.0879	-0.1773	0.8196	0.2845	0.3604	0.4618	0.0901	0.2326	0.8363	0.3076	0.5234	0.0750	0.9067
---	---	1630424_at	0.2335	0.2825	0.2331	0.2992	0.2827	0.1032	0.1152	0.7929	-0.0333	0.8906	-0.1485	0.3689	0.1078	0.8875	0.1358	0.6658	0.0281	0.9435
CG34382	CG14165	1630425_at	0.0245	0.9071	0.1217	0.4312	0.0323	0.8601	0.1253	0.7947	0.0997	0.6515	-0.0256	0.9148	0.0835	0.8692	0.0904	0.6876	0.0069	0.9840
---	---	1630426_at	0.0069	0.9758	0.1043	0.4927	0.2608	0.1497	0.2248	0.5639	-0.1029	0.6474	-0.3278	0.0696	-0.0440	0.9498	-0.0745	0.7667	-0.0306	0.9155
---	---	1630427_at	0.0642	0.7043	0.2729	0.3102	0.2451	0.1039	-0.1132	0.7409	-0.0958	0.5557	0.0174	0.9252	0.0952	0.8802	0.2688	0.2499	0.1736	0.4875
---	---	1630428_at	0.0719	0.7642	0.0019	0.9885	-0.1141	0.5225	-0.0269	0.9711	0.1168	0.5649	0.1437	0.4139	0.0893	0.8940	0.0083	0.9870	-0.0810	0.7865
---	---	1630429_s_at	0.1386	0.8527	-0.1127	0.4324	0.1789	0.7380	-0.4396	0.4741	-0.3043	0.3671	0.1354	0.6964	-0.5909	0.8122	-0.5780	0.5975	0.0130	0.9945
CG30437	CG30437	1630430_a_at	-1.4551	0.0090	-2.8776	0.0108	-2.9524	0.0001	0.2439	0.8078	0.8805	0.0347	0.6366	0.0765	0.2624	0.8655	-0.3941	0.5414	-0.6565	0.3007
CG10417	CG10417	1630431_s_at	0.1622	0.3791	0.1794	0.5636	-0.0667	0.7705	-0.2057	0.5290	0.1948	0.2552	0.4005	0.0168	0.1983	0.8609	0.2495	0.6111	0.0512	0.9364
---	---	1630432_at	0.2138	0.3368	0.0610	0.8614	-0.0386	0.8846	-0.0765	0.9412	0.1899	0.5527	0.2664	0.3255	0.0563	0.9467	0.0007	0.9996	-0.0555	0.8729
CG11963	CG11963	1630433_at	0.1337	0.4726	0.7660	0.0354	0.8213	0.0049	-0.1115	0.8457	-1.0580	0.0010	-0.9466	0.0010	-0.1732	0.8427	-0.4726	0.1742	-0.2994	0.4114
CG31064	CG31064	1630434_a_at	0.1096	0.5292	0.2188	0.1982	0.0023	0.9953	0.0640	0.9436	0.2271	0.3930	0.1631	0.5161	0.1389	0.8331	0.2956	0.2657	0.1567	0.5965
---	---	1630435_at	-0.0314	0.9069	0.1428	0.3843	0.2182	0.1638	-0.1844	0.5498	-0.0837	0.6395	0.1007	0.5137	0.0101	0.9935	0.1905	0.5146	0.1804	0.5457
CG11269	CG11269	1630436_at	0.2459	0.5040	0.2571	0.2259	0.1844	0.3867	0.1634	0.7749	0.1261	0.6377	-0.0373	0.8968	0.0855	0.9523	0.1303	0.8153	0.0449	0.9408
CG15118	anon-fast-evolving	1630437_s_at	0.2276	0.1405	0.2511	0.0962	0.2071	0.2786	0.0803	0.8671	0.2112	0.2101	0.1309	0.4066	0.1309	0.8611	0.2839	0.3381	0.1530	0.6398
CG32409	CG32409	1630438_at	0.3231	0.4409	0.6640	0.1715	1.6034	0.0008	0.3885	0.6041	0.2129	0.6075	-0.1756	0.6522	-0.2495	0.8648	0.5649	0.3209	0.8145	0.1883
---	---	1630439_at	-0.8759	0.0129	-1.3151	0.0137	-1.0832	0.0012	0.0529	0.9343	0.6307	0.0057	0.5778	0.0051	-0.0289	0.9710	0.0323	0.9170	0.0612	0.8015
Mtr3	Mtr3	1630440_at	0.0910	0.7436	0.1611	0.7695	0.4706	0.0819	0.1964	0.7121	0.3024	0.2107	0.1059	0.6761	-0.0948	0.9679	0.3135	0.6438	0.4083	0.5355
CG12716	CG12716	1630441_at	0.0744	0.7475	-0.3711	0.0753	-0.3543	0.0758	0.2896	0.5129	0.3135	0.1778	0.0239	0.9344	-0.0556	0.9405	-0.1420	0.5531	-0.0864	0.7440
CG3598	CG3598	1630442_at	0.2247	0.2461	0.3945	0.0212	0.1907	0.3221	-0.1663	0.6013	-0.0317	0.8831	0.1346	0.3628	0.0610	0.9428	-0.0176	0.9672	-0.0786	0.8057
---	---	1630443_at	0.0554	0.7376	0.0419	0.6862	-0.0538	0.8142	-0.0083	0.9903	0.1000	0.5323	0.1083	0.4431	0.3118	0.5228	0.0292	0.9390	-0.2826	0.2449
---	---	1630444_at	0.1786	0.3637	0.0000	1.0000	0.0781	0.7205	0.0143	0.9865	0.1114	0.6254	0.0971	0.6461	0.0268	0.9821	0.0078	0.9900	-0.0190	0.9629
l(3)07882	lethal (3) 07882	1630445_at	0.0218	0.9602	-0.3390	0.4755	0.1121	0.5403	0.3064	0.6283	0.5436	0.0825	0.2372	0.4159	0.0349	0.9894	0.3071	0.5995	0.2722	0.6445
---	---	1630446_at	0.0142	0.9435	0.0375	0.8833	0.0958	0.5466	0.0922	0.8940	0.1843	0.4394	0.0921	0.7041	-0.1038	0.8480	0.1205	0.6191	0.2242	0.3322
CG3078 /// DmirCG3078	CG3078	1630447_at	-0.0045	0.9858	-0.1104	0.6121	-0.0447	0.7981	0.2310	0.5096	0.1896	0.3113	-0.0414	0.8471	-0.0782	0.9238	-0.1418	0.6312	-0.0636	0.8592
CG1024	CG1024	1630448_at	0.5187	0.0665	0.2885	0.1707	0.5706	0.0046	-0.1312	0.8515	0.1651	0.5477	0.2963	0.1932	-0.2208	0.7686	0.2187	0.5068	0.4395	0.1911
---	---	1630449_s_at	0.0999	0.6597	0.2779	0.0901	0.0330	0.8706	-0.1257	0.7952	-0.2666	0.1748	-0.1409	0.4494	0.2316	0.7324	0.0524	0.9064	-0.1792	0.5612
Thd1	Thd1	1630450_s_at	-0.2829	0.3985	0.9542	0.0047	0.1594	0.4136	-0.4945	0.1925	-1.2153	0.0008	-0.7208	0.0046	0.2999	0.7330	0.0331	0.9581	-0.2668	0.4944
CG30185	CG30185	1630451_at	0.3820	0.1896	0.0747	0.5667	0.3241	0.0946	-0.0214	0.9745	-0.5038	0.0083	-0.4824	0.0060	-0.2730	0.7220	-0.6430	0.0667	-0.3700	0.2743
---	---	1630452_at	0.1169	0.8606	-0.9403	0.2569	-1.8549	0.0021	0.0280	0.9909	2.1193	0.0028	2.0913	0.0018	0.7153	0.7070	0.9199	0.2346	0.2046	0.8411
CG32267	CG32267	1630453_at	0.2227	0.3467	0.6668	0.0673	0.5647	0.0294	-0.0571	0.9493	-0.6223	0.0212	-0.5652	0.0198	0.0068	0.9964	-0.1836	0.6333	-0.1905	0.6195
CG14086	CG14086	1630454_at	0.2032	0.2085	0.2456	0.1677	0.3314	0.0578	0.0532	0.9311	0.0024	0.9926	-0.0508	0.8039	0.0154	0.9914	0.1396	0.6950	0.1242	0.7323
RpL19	ribosomal protein	1630455_at	0.1271	0.3577	1.1952	0.0063	1.0972	0.0012	0.0229	0.9803	-0.9710	0.0018	-0.9939	0.0010	0.0841	0.8655	-0.0248	0.9387	-0.1089	0.6109
Mbs	MLC-phosphatase	1630456_at	-2.6235	0.0065	-2.5864	0.0321	-2.8187	0.0001	0.1677	0.7664	0.6165	0.0187	0.4488	0.0432	0.5431	0.8379	0.7804	0.4848	0.2373	0.8709
by	tensin	1630457_s_at	-0.9016	0.0157	0.2928	0.4486	0.1531	0.2903	-0.1757	0.6580	-1.3543	0.0003	-1.1786	0.0003	-0.1320	0.9056	-0.0930	0.8687	0.0389	0.9441
CheA7a	CheA7a	1630458_at	-0.0182	0.9419	-0.0033	0.9810	-0.0580	0.7137	-0.0195	0.9812	-0.0957	0.6723	-0.0762	0.7183	0.1279	0.7822	0.0153	0.9630	-0.1126	0.6005
CG33980	CG33980	1630459_at	0.4570	0.1682	0.5124	0.0301	0.3275	0.2376	0.0418	0.9567	-0.0928	0.6977	-0.1346	0.4999	0.1278	0.9059	0.1241	0.8022	-0.0036	0.9956
CG15873	CG15873	1630460_at	0.1285	0.4267	-0.0759	0.6067	-0.0325	0.8884	-0.1028	0.8189	0.0026	0.9918	0.1053	0.5393	-0.0019	0.9989	-0.0833	0.7481	-0.0813	0.7492
---	---	1630461_at	0.1245	0.4702	-0.0204	0.9096	-0.1311	0.5469	0.0563	0.9196	0.0723	0.7146	0.0160	0.9397	0.1871	0.7780	0.0004	0.9998	-0.1867	0.5403
Obp50e	Odorant-binding p	1630462_at	0.0620	0.7126	0.0028	0.9826	0.0336	0.8786	-0.0357	0.9608	-0.1079	0.6139	-0.0721	0.7276	0.0461	0.9640	0.0990	0.7678	0.0530	0.8891
---	---	1630463_at	-1.7973	0.0145	0.4941	0.1746	-0.0213	0.9677	-0.5878	0.6785	-2.3131	0.0048	-1.7253	0.0108	0.1990	0.8510	0.0138	0.9877	-0.1853	0.7067
CG33800	CG33800	1630464_at	0.1393	0.4968	0.0136	0.9174	0.1195	0.4110	0.0793	0.8822	0.0919	0.6556	0.0126	0.9572	-0.0651	0.9309	-0.0544	0.8760	0.0108	0.9777
---	---	1630465_at	0.1185	0.6635	-0.2782	0.2315	-0.2530	0.2804	-0.0346	0.9759	0.2318	0.4286	0.2664	0.2989	-0.0592	0.9462	-0.1856	0.4972	-0.1264	0.6607
l(2)k09022	CG10805	1630466_at	0.9591	0.0167	0.3308	0.5142	0.9068	0.0008	0.4698	0.2939	0.6416	0.0229	0.1718	0.4965	-0.1624	0.9251	0.0685	0.9411	0.2309	0.7266
Dh31	diuretic hormone 3	1630467_a_at	0.0806	0.6802	-0.0385	0.7892	0.0099	0.9585	0.2047	0.5363	0.0708	0.7260	-0.1339	0.4066	0.1460	0.8097	-0.0497	0.8940	-0.1957	0.4507
CG15247	CG15247	1630468_at	-2.1502	0.0011	-0.2848	0.3385	-1.3746	0.0062	-0.4522	0.6338	-1.7841	0.0032	-1.3318	0.0071	0.2932	0.7464	-0.0277	0.9649	-0.3209	0.4096
CG1091	CG1091	1630469_a_at	0.2003	0.3006	-0.1479	0.6639	-0.2116	0.2236	-0.0183	0.9782	0.5303	0.0080	0.5486	0.0041	0.0680	0.9577	0.2256	0.5501	0.1577	0.6944
CG9775	CG9775	1630470_s_at	-0.3267	0.3120	-0.6347	0.2224	-0.7146	0.0113	-0.1066	0.7777	0.3622	0.0291	0.4688	0.0058	0.0559	0.9851	0.0946	0.9306	0.0388	0.9704
CG4301	CG4301	1630471_at	-0.3081	0.7171	-0.6989	0.0901	-0.9054	0.0086	0.2308	0.8663	0.3168	0.5419	0.0860	0.8806	0.2711	0.9134	-0.2249	0.8500	-0.4960	0.5969
---	---	1630472_at	-0.0621	0.7922	0.0099	0.9440	0.0805	0.7084	0.0737	0.9199	0.0545	0.8513	-0.0193	0.9460	0.0344	0.9739	0.0506	0.8960	0.0161	0.9666
CG5468	CG5468	1630473_at	-0.0247	0.8828	0.0405	0.9142	0.1992	0.2583	0.1834	0.7281	-0.0079	0.9821	-0.1913	0.3819	-0.0182	0.9870	-0.0499	0.8967	-0.0316	0.9296
---	---	1630474_at	0.2628	0.2861	-0.0087	0.9633	-0.1829	0.4010	-0.0426	0.9627	0.2955	0.2227	0.3382	0.1182	0.1019	0.8692	0.0180	0.9647	-0.0839	0.7729
CG40269	CG40269	1630475_at	0.0114	0.9487	0.0335	0.7310	-0.0817	0.5792	-0.1158	0.7111	-0.1168	0.4261	-0.0011	0.9953	-0.0493	0.9317	-0.0992	0.6191	-0.0499	0.8310
nahoda	nahoda	1630476_s_at	0.2907	0.2861	0.1431	0.7061	0.0262	0.9467	0.3281	0.3013	0.3462	0.0665	0.0181	0.9379	0.3134	0.8270	0.1419	0.8649	-0.1715	0.8177
msl-3	male specifc letha	1630477_at	0.0356	0.9346	-0.4364	0.1194	-0.2582	0.5978	0.2485	0.8385	0.3149	0.5157	0.0664	0.9039	-0.1539	0.9462	-0.2770	0.7299	-0.1231	0.8972
---	---	1630478_at	0.0986	0.6187	-0.2457	0.2723	-0.0842	0.6840	0.1062	0.8734	0.1419	0.5702	0.0357	0.8978	-0.1554	0.8814	-0.1628	0.7205	-0.0074	0.9915
---	---	1630479_at	0.0972	0.6392	0.0283	0.9216	0.0931	0.5989	-0.1087	0.8299	-0.0732	0.7511	0.0356	0.8773	-0.0121	0.9913	-0.0166	0.9648	-0.0044	0.9913

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Obp59a	Odorant-binding p	1630480_at	-0.0922	0.5493	0.0559	0.5643	0.1298	0.5575	0.1073	0.7857	-0.0696	0.7124	-0.1770	0.2262	-0.0557	0.9474	0.0824	0.8021	0.1381	0.6186
CG33631	CG33631	1630481_at	0.2143	0.4188	-0.0082	0.9403	-0.0544	0.8211	-0.1319	0.7838	0.0348	0.8983	0.1667	0.3612	-0.1555	0.8680	-0.0983	0.8445	0.0572	0.9109
CG4161	CG4161	1630482_at	0.0339	0.8365	-0.0077	0.9563	0.1631	0.3372	-0.0224	0.9759	0.0099	0.9719	0.0323	0.8806	-0.0582	0.9174	-0.0188	0.9519	0.0394	0.8824
CG7546	CG7546	1630483_a_at	-0.0722	0.9000	0.0788	0.8963	0.1316	0.5458	0.2034	0.8140	0.4370	0.1976	0.2336	0.4706	0.1494	0.9476	0.5807	0.3949	0.4313	0.5557
---	---	1630484_at	0.1147	0.6009	-0.0029	0.9817	-0.1516	0.4062	0.0737	0.9243	0.1781	0.4620	0.1044	0.6645	0.1921	0.7464	-0.0362	0.9269	-0.2284	0.3716
CG32021	CG32021	1630485_at	0.1169	0.5784	-0.5743	0.1136	-0.5228	0.0285	-0.0860	0.9068	0.6084	0.0183	0.6944	0.0062	-0.0949	0.8461	-0.1152	0.5900	-0.0204	0.9431
Bteb2	Bteb2	1630486_at	0.1306	0.3917	0.1495	0.4737	0.1802	0.3625	-0.1822	0.5539	-0.0161	0.9455	0.1661	0.2452	-0.0642	0.9401	0.0350	0.9341	0.0992	0.7442
---	---	1630487_s_at	-0.1278	0.6209	-0.2682	0.1284	-0.3603	0.3468	-0.0488	0.9774	0.2643	0.5625	0.3131	0.4308	-0.0520	0.9545	-0.0135	0.9778	0.0385	0.9173
---	---	1630488_at	-0.0297	0.9242	0.2971	0.2482	0.1649	0.4418	0.0745	0.9254	-0.0549	0.8612	-0.1293	0.5888	0.1160	0.8968	0.1824	0.6093	0.0664	0.8846
CG4116	CG4116	1630489_at	0.2119	0.3208	0.1705	0.4096	0.3862	0.1519	-0.0791	0.9037	-0.1345	0.5570	-0.0554	0.8178	-0.1102	0.8875	0.0495	0.9122	0.1597	0.6129
---	---	1630490_at	0.0878	0.7251	0.1734	0.2795	-0.0942	0.5047	-0.0938	0.8707	-0.0807	0.7334	0.0132	0.9589	0.1468	0.7770	0.0914	0.7248	-0.0554	0.8481
---	---	1630491_at	0.0918	0.6838	-0.1575	0.2061	-0.0793	0.6869	-0.0006	0.9994	0.1425	0.4292	0.1431	0.3741	0.0092	0.9952	-0.1608	0.6873	-0.1700	0.6607
Tfb2	Tfb2	1630492_at	0.0110	0.9533	0.0356	0.7623	0.2358	0.1604	-0.0735	0.8550	0.0822	0.6066	0.1557	0.2315	-0.3266	0.6483	0.1099	0.7439	0.4365	0.1537
---	---	1630493_at	0.1285	0.3568	0.1950	0.4261	0.1051	0.6803	-0.1328	0.8479	-0.0920	0.7618	0.0408	0.8930	0.0364	0.9550	0.0612	0.7984	0.0247	0.9240
---	---	1630494_at	0.0348	0.8285	-0.2438	0.1894	-0.4464	0.1461	0.1422	0.8738	0.3980	0.2007	0.2558	0.3760	0.0572	0.9460	-0.0369	0.9275	-0.0941	0.7492
CG30362	CG30362	1630495_at	0.1339	0.4396	0.2085	0.2717	0.4076	0.0306	-0.0295	0.9603	-0.3421	0.0371	-0.3126	0.0341	-0.0697	0.9142	-0.0469	0.8866	0.0228	0.9433
spel1	spellchecker	1630496_a_at	-0.9041	0.0569	-0.4465	0.0226	-0.8097	0.0120	-0.4451	0.3892	-1.1529	0.0028	-0.7079	0.0147	-0.4186	0.7070	-1.1329	0.0391	-0.7143	0.1498
---	---	1630497_at	0.2262	0.2475	0.0327	0.8056	0.1999	0.4632	-0.1216	0.8449	0.0447	0.8846	0.1663	0.4400	-0.2621	0.7485	-0.1296	0.7539	0.1325	0.7416
---	---	1630498_at	0.1269	0.3543	0.0403	0.7218	0.2443	0.1128	-0.0083	0.9900	-0.0129	0.9531	-0.0046	0.9799	0.0032	0.9964	-0.0123	0.9668	-0.0155	0.9512
CG7551	CG7551	1630499_at	0.1351	0.6943	-0.0147	0.9273	0.2204	0.1439	0.0349	0.9749	0.0115	0.9786	-0.0234	0.9461	-0.0974	0.9340	0.0315	0.9590	0.1290	0.7683
---	---	1630500_at	-0.1764	0.4346	-0.2266	0.4227	-0.3130	0.0468	0.0634	0.8990	-0.0379	0.8595	-0.1014	0.5240	0.1313	0.8815	0.0294	0.9585	-0.1019	0.8045
Strn-Mlck	Stretchin	1630501_a_at	0.2592	0.1956	-0.2799	0.1188	-0.4144	0.0932	0.0905	0.8473	0.2971	0.0857	0.2066	0.1787	0.0669	0.9521	-0.2931	0.3726	-0.3600	0.2960
ana	medullaless	1630502_at	-2.3241	0.0020	-1.0189	0.2921	-2.6770	0.0048	-1.2510	0.1109	-1.7037	0.0039	-0.4527	0.2817	0.4336	0.9168	-0.2427	0.9124	-0.6763	0.6644
CG6045	CG6045	1630503_at	-0.0292	0.8750	-0.3269	0.3355	-0.4745	0.0725	0.1206	0.8342	0.3674	0.0930	0.2468	0.2059	0.2711	0.7768	0.0637	0.9216	-0.2074	0.6414
CG13830	CG13830	1630504_at	0.0711	0.7145	0.1061	0.6190	0.1378	0.4527	0.0964	0.8119	0.1236	0.4591	0.0273	0.8862	0.1638	0.8243	0.1745	0.5936	0.0107	0.9838
Cyp311a1	Cyp311a1	1630505_a_at	-0.4178	0.1010	-0.4404	0.0280	-0.5597	0.0063	-0.3367	0.5242	-0.4285	0.1202	-0.0919	0.7618	-0.0022	0.9984	-0.1037	0.6643	-0.1016	0.6673
CG3077	CG3077	1630506_at	0.0229	0.9535	0.1989	0.3187	0.0071	0.9891	0.2798	0.4201	0.2508	0.1900	-0.0290	0.9002	0.4812	0.7307	0.5054	0.3943	0.0242	0.9809
CG15204	CG15204	1630507_at	-0.1477	0.4144	0.1603	0.3432	0.2062	0.2625	0.0209	0.9755	-0.1447	0.3945	-0.1656	0.2675	0.1554	0.8400	0.1423	0.6895	-0.0131	0.9803
---	---	1630508_at	-0.0807	0.6957	-0.0579	0.8459	-0.2665	0.0769	-0.1448	0.7581	0.0811	0.7318	0.2259	0.2112	0.1187	0.9095	0.0574	0.9214	-0.0612	0.9056
Mp20	myophilin	1630509_at	-1.8544	0.0327	-2.0981	0.0249	-2.5592	0.0002	-0.2466	0.5735	-0.4092	0.0710	-0.1626	0.4409	0.3316	0.9301	-0.7161	0.5806	-1.0478	0.4004
---	---	1630510_at	0.2051	0.4294	0.1719	0.2008	0.1247	0.5774	0.0580	0.9584	0.0545	0.8906	-0.0034	0.9924	0.0844	0.9467	0.1911	0.6458	0.1067	0.8245
---	---	1630511_at	-0.0076	0.9806	-0.0037	0.9764	0.0597	0.8765	0.0763	0.9518	0.2824	0.4347	0.2061	0.5469	-0.1486	0.7707	0.0299	0.9312	0.1784	0.4189
CG8316	CG8316	1630512_at	0.3131	0.1290	-0.1403	0.2318	-0.1312	0.5466	-0.1542	0.7605	0.2881	0.1836	0.4423	0.0307	-0.2096	0.6955	-0.2054	0.3534	0.0043	0.9914
---	---	1630513_s_at	0.1875	0.4798	-0.1961	0.2793	-0.0549	0.7290	0.2463	0.4455	0.2123	0.2273	-0.0341	0.8688	0.1515	0.8532	-0.0751	0.8764	-0.2266	0.5206
Gprk1	G protein-coupled	1630514_at	-0.9943	0.0069	-0.6461	0.1388	-0.8529	0.0023	-0.3038	0.5125	-0.3595	0.1405	-0.0557	0.8435	-0.1536	0.8744	-0.1169	0.8132	0.0367	0.9452
Glt	glutactin	1630515_s_at	0.4159	0.2279	0.4617	0.6126	1.4068	0.0031	0.0343	0.9778	-0.6662	0.0426	-0.7005	0.0221	-0.9151	0.7070	-0.5374	0.6234	0.3777	0.7483
CG14615	CG14615	1630516_at	0.7160	0.0578	0.1165	0.5213	1.0712	0.0057	0.2558	0.5301	0.3930	0.0675	0.1373	0.5018	-0.4617	0.7220	-0.2443	0.6961	0.2174	0.7333
CG15072	CG15072	1630517_at	0.0155	0.9328	0.2811	0.1037	0.1689	0.3231	-0.2434	0.3301	-0.1510	0.3012	0.0924	0.5079	0.2113	0.7387	0.0622	0.8712	-0.1491	0.6062
---	---	1630518_at	0.0992	0.4845	0.2145	0.0949	0.3153	0.0809	-0.0694	0.8971	-0.0875	0.6624	-0.0181	0.9344	-0.0452	0.9543	0.0159	0.9664	0.0611	0.8348
Trip1	Trip1	1630519_at	0.3735	0.0485	0.9922	0.0088	1.2191	0.0001	0.1632	0.6465	-0.5476	0.0073	-0.7108	0.0014	-0.0358	0.9657	0.0438	0.8962	0.0796	0.7578
---	---	1630520_s_at	2.0172	0.0105	0.7065	0.3771	1.4813	0.0007	0.7206	0.1903	0.3196	0.3460	-0.4010	0.1776	-0.1933	0.9589	-1.1278	0.2766	-0.9344	0.3921
Rst(1)JH	Methoprene-tolera	1630521_at	-0.1931	0.6849	0.5138	0.3086	0.4514	0.0147	-0.0215	0.9792	-0.2352	0.2513	-0.2136	0.2446	0.1167	0.9635	0.2610	0.7505	0.1443	0.8779
ST6Gal	sialyltransferase-li	1630522_a_at	-0.0532	0.7570	0.1868	0.2018	0.3227	0.1220	-0.0157	0.9833	-0.2560	0.1311	-0.2403	0.1130	-0.0050	0.9976	0.1224	0.7972	0.1274	0.7748
CG12106	CG12106	1630523_at	0.4702	0.1649	-0.2642	0.2259	-0.3895	0.0855	0.0193	0.9873	0.8867	0.0085	0.8674	0.0057	0.1202	0.8454	-0.1087	0.7040	-0.2289	0.3787
---	---	1630524_at	-1.1374	0.0981	-1.2679	0.0974	-2.1174	0.0000	-0.0436	0.9819	0.6679	0.1391	0.7114	0.0806	0.6070	0.7506	0.3498	0.7050	-0.2572	0.7937
---	---	1630525_at	0.5223	0.0113	0.2950	0.1673	0.1179	0.6460	-0.0436	0.9518	0.0043	0.9889	0.0479	0.8357	0.3556	0.3800	0.0154	0.9664	-0.3401	0.1538
---	---	1630526_at	0.1707	0.4185	0.1153	0.4361	0.1486	0.4426	0.0762	0.8598	-0.0622	0.7322	-0.1384	0.3283	0.1385	0.8270	0.0401	0.9229	-0.0984	0.7484
CG8436	CG8436	1630527_at	0.6708	0.0069	0.3444	0.2409	0.0158	0.9316	-0.1622	0.7302	0.3615	0.0868	0.5237	0.0132	0.0865	0.9309	-0.0757	0.8688	-0.1622	0.6428
CG17751	CG17751	1630528_at	-0.1028	0.7659	-0.0026	0.9903	0.0648	0.8142	-0.1086	0.8908	-0.0813	0.8032	0.0273	0.9343	-0.0322	0.9816	0.0732	0.8751	0.1054	0.7848
CG11562	CG11562	1630529_at	0.1010	0.5890	-0.0879	0.8438	0.0254	0.8920	0.0759	0.9013	-0.1080	0.6247	-0.1839	0.3082	-0.0800	0.9589	-0.1701	0.7364	-0.0901	0.8779
CG9296	CG9296	1630530_at	-0.4212	0.1174	-0.2812	0.2350	-0.4293	0.1185	-0.0961	0.8987	0.3411	0.1651	0.4371	0.0523	0.0561	0.9689	0.4875	0.1797	0.4314	0.2665
RhoGAP93B	RhoGAP93B	1630531_at	-0.4941	0.2112	-0.7778	0.0149	-0.9936	0.0032	0.0335	0.9506	0.5687	0.0032	0.5353	0.0025	0.1530	0.9220	0.1766	0.7837	0.0236	0.9769
CG13091	CG13091	1630532_at	0.3422	0.9025	0.5667	0.7985	0.2242	0.3505	0.6044	0.8908	-1.4647	0.3296	-2.0691	0.1171	1.0604	0.8909	-1.0027	0.7802	-2.0632	0.4944
CG3221	CG3221	1630533_at	-1.0304	0.0054	-1.7283	0.0112	-2.0625	0.0001	0.0254	0.9865	0.8346	0.0296	0.8092	0.0213	0.0863	0.9095	0.0391	0.9275	-0.0472	0.8987
---	---	1630534_at	-0.0479	0.8500	0.1952	0.3466	0.0059	0.9815	-0.0961	0.8734	-0.1948	0.3621	-0.0987	0.6473	0.1508	0.8305	0.2232	0.4505	0.0724	0.8488
---	---	1630535_at	0.1344	0.4866	-0.0039	0.9767	-0.0457	0.8010	-0.0769	0.9130	0.0299	0.9230	0.1069	0.6393	-0.0555	0.9474	-0.0817	0.8034	-0.0261	0.9427
mod(mdg4)	Modifier67.2	1630536_at	-0.3905	0.0641	-0.0952	0.5437	-0.0902	0.7445	-0.0185	0.9857	-0.1878	0.4352	-0.1693	0.4377	-0.1555	0.9014	0.0302	0.9682	0.1857	0.7224
Dap160	Dynamin associat	1630537_a_at	0.3051	0.1744	0.5038	0.0920	0.6494	0.0037	0.1202	0.8140	0.3322	0.0991	0.2119	0.2407	0.0006	0.9998	0.4647	0.1382	0.4641	0.1651
Cpr60D	CG30163	1630538_at	0.1418	0.5225	0.0069	0.9610	0.2674	0.1330	-0.0271	0.9693	-0.0797	0.7033	-0.0526	0.7966	-0.2005	0.6955	-0.2195	0.2938	-0.0191	0.9494
Topors	Topoisomerase I-i	1630539_at	-0.2436	0.3296	-0.0295	0.9229	0.0453	0.8452	0.1700	0.6409	0.1398	0.4499	-0.0302	0.8867	0.0516	0.9734	0.2706	0.4875	0.2190	0.5907
yellow-e3	yellow-e3	1630540_at	0.1526	0.7058	0.1478	0.5322	-0.0240	0.9076	0.0610	0.9711	-0.2729	0.5527	-0.3338	0.4005	0.1031	0.9088	-0.1916	0.5664	-0.2947	0.3680
---	---	1630541_at	0.1902	0.1871	0.0019	0.9937	-0.0111	0.9646	0.0335	0.9550	0.0030	0.9899	-0.0305	0.8725	-0.0126	0.9943	-0.0089	0.9925	0.0037	0.9959
Hpr1	Hpr1	1630542_at	-0.0391	0.8527	0.2362	0.3599	0.5155	0.0043	0.0764	0.8671	-0.2974	0.0683	-0.3738	0.0177	-0.1802	0.8331	-0.0204	0.9734	0.1598	0.6861
CG5161	CG5161	1630543_at	-0.1223	0.4396	0.3064	0.3020	-0.0238	0.9091	0.0243	0.9705	-0.1284	0.4642	-0.1527	0.3189	0.3812	0.6483	0.4112	0.2077	0.0299	0.9493
---	---	1630544_at	-0.0178	0.9547	0.0000	1.0000	-0.0715	0.7083	0.0017	0.9986	0.0166	0.9554	0.0149	0.9534	-0.0132	0.9869	0.0774	0.7103	0.0905	0.6475
---	---	1630545_at	0.2289	0.2631	0.2101	0.0822	0.0700	0.7193	-0.0487	0.9377	0.0164	0.9518	0.0652	0.7403	0.0284	0.9717	-0.1116	0.5986	-0.1400	0.4970
CG15023	CG15023	1630546_at	0.0362	0.8515	0.1192	0.3913	0.0019	0.9944	-0.1572	0.6085	-0.2497	0.1041	-0.0925	0.5389	0.1778	0.7633	-0.0611	0.8578	-0.2389	0.3467
CG4723	CG4723	1630547_at	2.0779	0.0140	1.3695	0.0847	2.6026	0.0002	0.5343	0.5648	-0.0017	0.9980	-0.5361	0.2082	-0.8424	0.6955	-0.6710	0.4490	0.1714	0.8880
---	---	1630548_at	0.0062	0.9787	0.0278	0.8429	0.3431	0.0380	0.0455	0.9436	0.0471	0.8434	0.0015	0.9949	-0.1711	0.7893	-0.0180	0.9682	0.1531	0.5995

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1630549_at	0.1917	0.2269	0.0965	0.6223	0.1571	0.4648	-0.0802	0.9011	-0.0575	0.8288	0.0228	0.9293	-0.0584	0.9467	-0.0025	0.9973	0.0558	0.8774
CG5674	CG5674	1630550_a_at	-0.6107	0.0565	-0.7579	0.0772	-0.5588	0.0135	-0.1046	0.8350	-0.0346	0.8932	0.0700	0.7307	-0.3271	0.7726	-0.1683	0.7782	0.1588	0.7827
CG5290	CG5290	1630551_at	0.4449	0.5923	0.3541	0.4856	0.6002	0.0457	0.4333	0.8190	0.6560	0.3879	0.2227	0.7880	0.2876	0.8472	0.7286	0.2224	0.4410	0.4912
CG33980	CG33980	1630552_at	0.5185	0.0462	0.4007	0.1708	0.3955	0.0960	0.1556	0.7261	0.0707	0.7674	-0.0849	0.6845	0.2720	0.6955	0.0102	0.9856	-0.2618	0.3723
alphaPS5	alphaPS5	1630553_at	-0.2054	0.3106	-0.3212	0.1355	-0.2277	0.1693	0.1041	0.7929	0.1972	0.2214	0.0931	0.5599	-0.0909	0.9142	-0.0813	0.8382	0.0095	0.9847
l(3)neo38	lethal (3) neo38	1630554_at	0.1971	0.5459	0.0347	0.9002	0.6486	0.0164	0.1798	0.6584	0.0651	0.7839	-0.1147	0.5506	-0.4052	0.7506	-0.0836	0.9222	0.3216	0.5814
CG33346	CG33346	1630555_at	0.2036	0.5202	-0.0664	0.4843	-1.1341	0.0077	-0.9409	0.1956	0.2283	0.6439	1.1693	0.0098	-0.0949	0.8991	-0.1194	0.7020	-0.0245	0.9494
---	---	1630556_at	0.2209	0.1516	0.0793	0.5127	0.2051	0.1865	-0.0088	0.9895	-0.0292	0.8892	-0.0204	0.9137	-0.0824	0.8626	-0.0764	0.7317	0.0060	0.9849
CG11859	CG11859	1630557_at	-0.0981	0.7410	0.4731	0.2442	0.8025	0.0019	0.1108	0.8923	-0.6679	0.0230	-0.7787	0.0072	-0.2877	0.7856	-0.0393	0.9587	0.2484	0.6124
---	---	1630558_a_at	-0.0214	0.9424	0.0922	0.4787	-0.0439	0.8508	-0.0943	0.8067	-0.1195	0.4535	-0.0252	0.8907	0.0460	0.9514	-0.0236	0.9482	-0.0695	0.7954
MED8	Mediator complex	1630559_at	0.0476	0.8547	-0.2201	0.4156	-0.0931	0.5809	0.1042	0.7672	0.1937	0.1937	0.0895	0.5411	0.0620	0.9683	-0.0087	0.9925	-0.0707	0.9016
Gyc76C	receptor-type gua	1630560_s_at	-0.0792	0.7328	-0.0569	0.7625	-0.1305	0.4961	-0.2050	0.6202	-0.0226	0.9394	0.1824	0.3334	-0.2164	0.7070	-0.0631	0.8470	0.1533	0.5457
CG13641	CG13641	1630561_at	1.3717	0.0546	0.2806	0.7115	1.6609	0.0161	1.7854	0.0154	2.1142	0.0007	0.3288	0.3494	0.3931	0.9126	1.0643	0.3922	0.6712	0.6212
---	---	1630562_at	0.1994	0.4060	0.0055	0.9836	0.3617	0.0849	0.0875	0.9066	-0.0383	0.9085	-0.1258	0.6025	-0.1686	0.8193	-0.2181	0.4802	-0.0495	0.9057
CG6006	CG6006	1630563_at	-1.5135	0.0160	-0.0980	0.6426	-0.6039	0.0374	-0.5453	0.4633	-1.8074	0.0016	-1.2621	0.0046	0.0412	0.9816	-0.4707	0.2424	-0.5119	0.2338
---	---	1630564_at	0.2440	0.2303	0.1316	0.4148	0.0776	0.7845	0.0055	0.9956	0.0309	0.9335	0.0254	0.9381	0.1081	0.8909	-0.2107	0.4888	-0.3188	0.2964
CG17652	CG17652	1630565_at	-0.1048	0.5287	-0.3109	0.4105	-0.3935	0.0560	0.2941	0.3403	0.5085	0.0115	0.2144	0.1764	0.2928	0.7519	0.2206	0.6021	-0.0722	0.8947
CG15446	CG15446	1630566_at	0.2646	0.2200	0.0560	0.6157	0.2759	0.1233	0.1419	0.7604	0.1864	0.3616	0.0445	0.8482	-0.0755	0.9309	-0.0785	0.8349	-0.0030	0.9953
---	---	1630567_at	0.1067	0.5740	0.1535	0.3143	0.0603	0.7393	0.0960	0.8507	0.0816	0.7034	-0.0144	0.9511	0.0786	0.9029	0.0691	0.8216	-0.0095	0.9807
CG34040	CG34040	1630568_at	0.0743	0.7314	0.0130	0.9072	0.2208	0.2594	-0.0438	0.9592	0.0326	0.9177	0.0763	0.7562	-0.2572	0.6557	0.0089	0.9841	0.2661	0.2666
CG32668	CG32668	1630569_at	0.0032	0.9887	0.1691	0.4012	0.0947	0.6482	-0.1967	0.5735	-0.1962	0.2752	0.0005	0.9983	-0.0912	0.8875	0.0306	0.9376	0.1218	0.6398
CG13003	CG13003	1630570_at	-1.1452	0.0706	-0.6866	0.2925	-1.1538	0.0038	-0.3960	0.6506	-0.8290	0.0540	-0.4330	0.2532	0.0103	0.9964	-0.5047	0.3749	-0.5151	0.3848
CG14317	CG14317	1630571_at	0.1236	0.4662	-0.0225	0.8786	-0.0867	0.6715	-0.0834	0.9254	-0.1366	0.6500	-0.0532	0.8643	0.0244	0.9816	-0.0945	0.7184	-0.1189	0.6299
cpb	capping protein	1630572_at	-0.2035	0.2990	0.0030	0.9907	0.0556	0.7936	0.1288	0.7982	0.2030	0.3272	0.0742	0.7357	0.0153	0.9923	0.3675	0.3010	0.3522	0.3488
CG6498	CG6498	1630573_at	-0.5762	0.3207	0.6732	0.4268	0.7845	0.0035	0.0931	0.9311	-1.3393	0.0021	-1.4325	0.0009	-0.0417	0.9927	-0.0070	0.9988	0.0347	0.9842
CG3906	CG3906	1630574_at	0.0426	0.8944	0.0648	0.5450	0.0274	0.8950	0.0098	0.9879	-0.1555	0.3107	-0.1653	0.2242	0.0818	0.9538	-0.0122	0.9892	-0.0940	0.8646
CG1907	CG1907	1630575_at	0.3469	0.1412	0.4204	0.0448	0.1552	0.4263	-0.0607	0.9261	-0.2948	0.1335	-0.2340	0.1830	0.1992	0.8122	-0.2602	0.4565	-0.4594	0.2075
CG34362	CG12870	1630576_at	0.0167	0.9396	0.0534	0.8056	0.0120	0.9545	-0.0149	0.9838	-0.0541	0.7914	-0.0392	0.8388	0.0074	0.9943	-0.0320	0.9341	-0.0394	0.9061
CG17378	CG17378	1630577_at	-0.0658	0.7117	-0.0668	0.5963	0.1122	0.5620	0.1033	0.8064	0.0154	0.9514	-0.0880	0.6019	-0.1221	0.8235	-0.0607	0.8444	0.0614	0.8307
---	---	1630578_at	0.1033	0.5222	0.0662	0.6587	0.0539	0.7635	-0.0109	0.9935	-0.2092	0.4354	-0.1983	0.4121	0.1806	0.8331	-0.1045	0.8244	-0.2851	0.4386
Cdk8	Cyclin-dependent	1630579_at	0.3841	0.0780	-0.3008	0.4063	-0.2121	0.3115	0.0692	0.9436	0.4729	0.0889	0.4037	0.1027	0.0374	0.9816	-0.0880	0.8595	-0.1254	0.7635
---	---	1630580_at	0.0662	0.6587	0.0094	0.9507	0.3388	0.0780	0.1349	0.7941	0.1469	0.5097	0.0120	0.9639	-0.0356	0.9514	-0.0312	0.9075	0.0043	0.9874
CG32639	CG32639	1630581_at	-0.4686	0.0767	0.4801	0.5054	0.5955	0.0118	-0.2893	0.5336	-0.3114	0.1976	-0.0221	0.9416	-0.1763	0.9342	0.5820	0.3903	0.7583	0.2842
CG3967	CG3967	1630582_s_at	-1.1990	0.0014	-1.8773	0.0179	-1.8090	0.0001	-0.1611	0.8171	0.5624	0.0454	0.7235	0.0100	-0.0651	0.9677	-0.0117	0.9903	0.0534	0.9280
---	---	1630583_at	0.1793	0.3886	0.1420	0.5665	0.1450	0.3761	0.0046	0.9956	-0.0485	0.8718	-0.0530	0.8388	0.0398	0.9737	-0.0022	0.9984	-0.0420	0.9185
CG10657	retinaldehyde-bin	1630584_at	-2.9307	0.0010	-3.4102	0.0067	-3.7013	0.0000	0.2904	0.7003	0.5819	0.0936	0.2915	0.3607	0.6560	0.6749	0.1645	0.8539	-0.4915	0.4647
---	---	1630585_s_at	0.3008	0.0971	-0.5005	0.0682	-1.1729	0.0022	-0.0386	0.9777	0.4950	0.1440	0.5336	0.0808	-0.0335	0.9841	-0.3707	0.3173	-0.3372	0.3859
---	---	1630586_at	0.1979	0.3927	-0.0015	0.9907	0.3725	0.1187	0.1315	0.8590	-0.0008	0.9984	-0.1323	0.6260	-0.1624	0.7485	-0.0442	0.8867	0.1182	0.6099
Tehao	Tehao	1630587_at	0.2195	0.2057	0.1110	0.4457	-0.0402	0.8604	-0.0123	0.9922	-0.1637	0.5622	-0.1514	0.5579	0.0266	0.9777	-0.1874	0.4054	-0.2140	0.3601
---	---	1630588_at	0.0047	0.9844	-0.0427	0.8014	0.1691	0.2843	0.1085	0.8048	0.1685	0.3446	0.0600	0.7539	-0.1545	0.8012	0.0323	0.9376	0.1868	0.4805
CG17734	CG17734	1630589_at	-0.5702	0.0319	-0.3984	0.1647	-0.8308	0.0090	-0.3451	0.2397	-0.1153	0.5346	0.2298	0.1373	-0.0247	0.9898	-0.0542	0.9363	-0.0295	0.9622
CG18417	CG18417	1630590_at	1.8705	0.1708	0.4519	0.6639	1.9528	0.0004	0.3876	0.8836	-0.2338	0.8405	-0.6214	0.4670	-0.8752	0.8049	-0.9661	0.5283	-0.0909	0.9674
CG14460	CG14460	1630591_at	-0.0244	0.9141	-0.1926	0.5185	-0.0217	0.9061	0.2499	0.5902	0.2059	0.3938	-0.0440	0.8744	-0.0416	0.9636	-0.0839	0.7867	-0.0423	0.9023
---	---	1630592_at	0.0105	0.9574	-0.0419	0.6862	0.2539	0.1835	0.2898	0.4626	-0.0172	0.9566	-0.3070	0.1069	0.0002	0.9999	0.0021	0.9957	0.0019	0.9951
NLaz	Neural Lazarillo	1630593_at	0.2244	0.2534	1.3682	0.0075	1.5060	0.0003	0.0891	0.9117	-0.3029	0.2365	-0.3920	0.0868	0.0716	0.9588	0.7315	0.0721	0.6599	0.1192
---	---	1630594_at	-0.0244	0.8831	0.0303	0.8094	0.1188	0.5315	-0.0806	0.8422	-0.0558	0.7552	0.0247	0.8907	-0.1497	0.8141	-0.0332	0.9376	0.1165	0.6883
CG17601	CG17601	1630595_at	0.3644	0.1151	0.3206	0.2411	0.2329	0.2370	0.1355	0.8598	0.1137	0.7230	-0.0217	0.9503	0.1329	0.8276	-0.0027	0.9965	-0.1356	0.6225
CG14643	CG14643	1630596_at	0.9263	0.0210	0.6218	0.0837	0.1620	0.5516	-0.8108	0.1250	-0.5339	0.0937	0.2769	0.3423	-0.2328	0.8689	-0.6747	0.2113	-0.4420	0.4399
---	---	1630597_at	0.0701	0.7832	0.0278	0.8974	0.1744	0.3171	0.0711	0.9006	0.1150	0.5630	0.0439	0.8354	-0.0461	0.9309	-0.0148	0.9590	0.0314	0.8949
CG18179	CG18179	1630598_at	0.2187	0.4488	0.4258	0.3340	0.4517	0.0897	0.2547	0.6080	0.1984	0.4424	-0.0563	0.8447	0.6034	0.7220	0.6883	0.3259	0.0849	0.9331
CG31172	CG31172	1630599_at	0.0157	0.9232	-0.0112	0.9208	0.2047	0.3541	0.0812	0.8987	0.0981	0.6800	0.0169	0.9492	-0.0742	0.9405	0.0201	0.9682	0.0943	0.7994
Fst /// Scm	Frost /// Sex comb	1630600_at	0.0774	0.8307	0.2191	0.2433	0.1049	0.5529	0.0517	0.9649	-0.3312	0.2930	-0.3830	0.1690	0.0090	0.9939	-0.1369	0.6152	-0.1459	0.5910
CG15128	CG15128	1630601_at	0.0470	0.8239	0.0266	0.8824	0.1365	0.5389	0.0830	0.9080	-0.0386	0.9030	-0.1216	0.5989	0.0642	0.9400	0.0181	0.9664	-0.0460	0.8984
CG13842	CG13842	1630602_at	-0.0062	0.9748	-0.0466	0.8037	-0.0673	0.7064	-0.0761	0.8921	-0.0885	0.6797	-0.0124	0.9587	-0.0768	0.9137	-0.0450	0.9058	0.0318	0.9260
CG31352	CG31352	1630603_at	-1.6134	0.0177	0.0174	0.9883	-0.0618	0.7850	-0.0254	0.9857	-1.3767	0.0019	-1.3512	0.0012	0.0437	0.9928	0.5240	0.6673	0.4803	0.6968
---	---	1630604_s_at	0.1784	0.4402	0.0767	0.5221	0.3618	0.0387	-0.0274	0.9703	-0.1124	0.5803	-0.0849	0.6615	-0.0305	0.9775	0.1617	0.5440	0.1922	0.4678
CG16947	CG16947	1630605_at	-2.3601	0.0004	-2.9325	0.0027	-3.5925	0.0000	-0.1911	0.6888	-0.0091	0.9790	0.1820	0.3735	0.4145	0.7215	-0.4827	0.3077	-0.8972	0.1011
SH3PX1	SH3PX1	1630606_at	0.1401	0.6537	-0.1866	0.5052	-0.0131	0.9725	0.2898	0.5039	0.6915	0.0093	0.4017	0.0552	0.0757	0.9646	0.4318	0.3348	0.3561	0.4533
CG6938	CG6938	1630607_at	-0.0619	0.8063	0.0128	0.9476	0.2980	0.0696	0.1926	0.4779	-0.1713	0.2398	-0.3638	0.0130	0.0081	0.9943	0.0010	0.9994	-0.0070	0.9870
CG14608	CG14608	1630608_at	-0.0369	0.9079	0.1703	0.5539	0.1876	0.1863	0.1699	0.8164	0.0615	0.8722	-0.1084	0.7240	0.1033	0.8465	0.1672	0.4519	0.0639	0.8177
CG2225	CG2225	1630609_s_at	1.4241	0.0472	0.2683	0.4809	-0.4700	0.3407	0.4457	0.2840	2.0986	0.0001	1.6529	0.0002	0.9649	0.7092	0.4448	0.7200	-0.5201	0.6572
CG40177	CG40177	1630610_at	-0.0593	0.8116	-0.0360	0.7218	0.0349	0.8639	0.1461	0.6835	-0.0358	0.8713	-0.1819	0.2228	0.0265	0.9859	-0.1479	0.6949	-0.1744	0.6294
CG7181 /// DmauCG7181 /	CG7181	1630611_at	0.0834	0.8305	0.5693	0.1683	0.1253	0.5661	-0.2287	0.5197	-0.8087	0.0019	-0.5800	0.0048	0.2026	0.8997	-0.3610	0.5602	-0.5637	0.3545
---	---	1630612_at	-0.3914	0.7141	0.0982	0.4597	-0.1511	0.4814	-0.0919	0.9777	-0.7538	0.3631	-0.6619	0.3784	0.0469	0.9862	-0.2828	0.6844	-0.3297	0.6231
CG30279	CG30279	1630613_at	0.0302	0.8731	0.1293	0.4371	0.0961	0.5216	0.0196	0.9782	-0.0058	0.9827	-0.0254	0.9086	0.0587	0.9503	0.0268	0.9525	-0.0319	0.9348
---	---	1630614_s_at	0.5518	0.3718	-0.2473	0.6823	0.2070	0.7657	0.7948	0.1256	1.0278	0.0062	0.2329	0.4267	0.2130	0.9657	0.2354	0.9075	0.0224	0.9924
CG13602	CG13602	1630615_s_at	-0.0438	0.8380	-0.0712	0.6639	-0.1841	0.2660	0.0988	0.8589	0.0860	0.7101	-0.0128	0.9594	0.0412	0.9407	0.0243	0.9279	-0.0169	0.9435
Ugt37a1	UDP-glycosyltran	1630616_at	0.3191	0.2600	0.2636	0.1029	0.4245	0.0473	-0.0742	0.9234	-0.2844	0.2207	-0.2102	0.3224	-0.0125	0.9923	-0.0800	0.8411	-0.0676	0.8627
kek2	Kekkon-2	1630617_at	0.0257	0.8864	-0.0882	0.4187	0.0734	0.7242	0.0288	0.9621	0.0639	0.7364	0.0352	0.8507	-0.2231	0.8049	-0.1153	0.8131	0.1079	0.8178

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
dys	Dysfusion	1630618_at	0.0145	0.9792	0.5402	0.0721	0.9872	0.0067	-0.1362	0.8687	-0.5724	0.0535	-0.4362	0.0930	-0.3136	0.7811	0.0538	0.9460	0.3674	0.4683
Spn1	serpin 1	1630619_at	-1.4257	0.0032	-1.5506	0.0253	-0.9690	0.0009	0.1102	0.8567	0.1112	0.6521	0.0010	0.9973	-0.3935	0.7440	0.0980	0.8981	0.4915	0.3475
Pole2	Pole2	1630620_at	0.2519	0.6463	-0.0825	0.8790	-0.3541	0.4937	-0.3049	0.6890	0.4587	0.1920	0.7636	0.0239	-0.1969	0.9589	0.1667	0.9222	0.3636	0.7774
CG15212	CG15212	1630621_at	0.1453	0.3589	0.2629	0.1673	0.5020	0.0298	-0.1368	0.6823	-0.1048	0.5283	0.0321	0.8599	-0.0735	0.9137	0.1004	0.7086	0.1738	0.4791
Sirt6	Sirt6	1630622_at	0.6356	0.0401	0.2479	0.4129	0.6071	0.0241	-0.0138	0.9889	0.0706	0.8027	0.0844	0.7327	-0.2935	0.8049	-0.2580	0.6305	0.0355	0.9613
CG8058	CG8058	1630623_a_at	-1.0370	0.0057	0.1151	0.5362	-0.0047	0.9900	-0.4775	0.4191	-1.8179	0.0007	-1.3404	0.0014	-0.2763	0.7220	-0.3540	0.2714	-0.0777	0.8581
CG10151	CG10151	1630624_s_at	-0.0541	0.8744	0.1491	0.3983	-0.1750	0.2786	-0.2618	0.6368	-0.2575	0.3531	0.0043	0.9901	0.0885	0.9012	-0.0689	0.8468	-0.1574	0.5709
---	---	1630625_s_at	-0.0083	0.9732	-0.0071	0.9517	-0.0102	0.9585	-0.0137	0.9823	-0.0001	0.9996	0.0136	0.9397	0.0157	0.9869	-0.0853	0.7401	-0.1011	0.6731
pps	SNFs Protein Parl	1630626_at	0.1942	0.6335	0.2328	0.1551	0.0460	0.8067	0.2506	0.6908	0.0013	0.9977	-0.2493	0.3521	0.2155	0.8465	-0.0419	0.9557	-0.2574	0.6052
---	---	1630627_at	0.1818	0.3390	-0.0094	0.9517	-0.0900	0.6799	0.0846	0.8598	0.2697	0.1157	0.1851	0.2294	0.0713	0.9111	0.0178	0.9620	-0.0534	0.8562
CG18091	CG18091	1630628_at	0.1541	0.3790	0.5422	0.0775	0.4814	0.0067	-0.1496	0.7053	-0.1814	0.3250	-0.0317	0.8826	0.0827	0.8814	0.2124	0.3010	0.1297	0.5630
Atg8b	Autophagy-specifi	1630629_at	0.1325	0.5923	-0.0801	0.5498	0.1157	0.6147	0.2257	0.4815	0.1817	0.2943	-0.0440	0.8214	-0.0562	0.9514	0.0463	0.9121	0.1025	0.7445
CG6370	CG6370	1630630_at	0.8911	0.0016	0.5577	0.1850	0.6971	0.0045	0.1126	0.7970	0.5671	0.0072	0.4545	0.0117	-0.0172	0.9914	0.3151	0.3802	0.3323	0.3765
CG5882	CG5882	1630631_at	0.0850	0.6116	0.1658	0.3575	0.1252	0.5075	0.1416	0.6791	-0.0738	0.6872	-0.2154	0.1305	0.2511	0.7611	0.0637	0.9041	-0.1874	0.6150
CG2982	CG2982	1630632_s_at	0.4816	0.2212	-0.2260	0.4476	-0.2310	0.2387	-0.0020	0.9988	0.9286	0.0067	0.9306	0.0039	-0.1308	0.9036	0.1472	0.7515	0.2781	0.4938
CG13130	CG13130	1630633_at	0.1387	0.4504	0.0240	0.8263	0.2849	0.1249	0.1889	0.5755	0.2071	0.2300	0.0183	0.9314	-0.0056	0.9952	-0.0204	0.9567	-0.0148	0.9657
RpL9	anon-fast-evolving	1630634_s_at	0.3598	0.0304	0.9587	0.0055	0.9173	0.0160	0.3090	0.6506	-0.4791	0.1445	-0.7881	0.0156	0.1703	0.7783	-0.0404	0.9220	-0.2106	0.4357
CG7720	CG7720	1630635_s_at	-0.7630	0.0378	-1.0842	0.0338	-1.0957	0.0162	-0.4999	0.2977	-0.6336	0.0320	-0.1337	0.6449	-0.6736	0.6749	-0.8837	0.1747	-0.2102	0.7954
CG2446	CG2446	1630636_s_at	-0.6077	0.1827	-0.0162	0.9624	-0.3611	0.1211	-0.1568	0.8496	-0.0983	0.7897	0.0585	0.8682	0.2116	0.8814	0.5153	0.3285	0.3037	0.6027
Hsc70-4	heat-shock protein	1630637_s_at	-0.4456	0.0883	-0.0680	0.6723	-0.2590	0.0823	0.1214	0.7031	-0.0748	0.6507	-0.1961	0.1330	0.2656	0.7324	0.2758	0.4037	0.0102	0.9849
dre4	supressor of Ty el	1630638_at	-0.0133	0.9782	1.0288	0.0100	1.1959	0.0025	-0.0052	0.9961	-0.8435	0.0222	-0.8382	0.0142	-0.0699	0.9701	0.0726	0.9225	0.1425	0.8088
dsd	distracted	1630639_at	0.2899	0.6438	-0.4015	0.4409	-0.6528	0.0036	-0.0278	0.9712	0.6888	0.0045	0.7165	0.0022	0.3335	0.8814	0.0423	0.9782	-0.2912	0.7725
CG4594	CG4594	1630640_at	0.9821	0.0342	0.4226	0.0245	0.6358	0.0053	-0.2071	0.5779	0.1001	0.6395	0.3072	0.0735	-0.4586	0.7149	-0.4053	0.4388	0.0533	0.9428
CG11634	CG11634	1630641_at	-0.1001	0.5477	-0.0177	0.9198	-0.1942	0.2067	-0.1016	0.8393	-0.0237	0.9286	0.0780	0.6920	0.1256	0.8145	0.1405	0.5433	0.0149	0.9636
Pvf2	VEGF-related fact	1630642_at	-3.3520	0.0004	-2.0155	0.0214	-2.4426	0.0041	-0.0373	0.9665	-0.2254	0.3393	-0.1881	0.3814	0.4147	0.8825	1.1689	0.2575	0.7542	0.4978
ScpX	sterol carrier prote	1630643_at	0.8273	0.0375	0.1762	0.4901	0.5252	0.0087	0.0572	0.9259	0.0579	0.7988	0.0007	0.9978	-0.2264	0.8400	-0.3847	0.3999	-0.1583	0.7703
---	---	1630644_at	0.0178	0.9294	0.0237	0.8141	-0.2609	0.1096	-0.1229	0.7857	0.1611	0.4014	0.2840	0.0891	0.1124	0.8461	0.1602	0.5158	0.0478	0.8837
CG3528	CG3528	1630645_at	0.2346	0.2621	0.1475	0.2938	0.1717	0.2439	-0.1322	0.7492	-0.0920	0.6496	0.0402	0.8471	0.0541	0.9589	-0.1825	0.5634	-0.2367	0.4457
CG34345	CG11053	1630646_at	0.1884	0.4670	-0.0563	0.7261	-0.0948	0.6455	0.2131	0.6010	0.1466	0.4966	-0.0665	0.7689	0.1080	0.9305	-0.1408	0.7725	-0.2488	0.5568
---	---	1630647_at	-0.1697	0.5750	-0.0558	0.7285	0.1984	0.5475	-0.0429	0.9602	-0.0265	0.9338	0.0165	0.9544	-0.1655	0.8331	0.0617	0.9009	0.2272	0.5038
---	---	1630648_at	0.0984	0.7732	-0.1539	0.4044	-0.1054	0.4851	0.1127	0.9149	0.1525	0.6869	0.0398	0.9218	0.0736	0.9355	-0.0763	0.8460	-0.1499	0.6311
CG12520	CG12520	1630649_at	-0.0064	0.9758	0.1095	0.4758	-0.0009	0.9978	-0.0957	0.7857	-0.0725	0.6592	0.0232	0.8942	-0.1014	0.9142	-0.0481	0.9269	0.0534	0.9067
BicD	Bicaudal-D	1630650_at	-0.2141	0.4277	0.2507	0.5360	-0.1963	0.2600	-0.0769	0.8932	0.1501	0.4492	0.2269	0.1822	0.3443	0.7893	0.5768	0.2735	0.2325	0.7040
---	---	1630651_at	-0.2795	0.3306	0.0242	0.8107	-0.1026	0.5932	-0.0371	0.9608	-0.0979	0.6690	-0.0607	0.7871	-0.0262	0.9794	0.0602	0.8493	0.0864	0.7472
---	---	1630652_at	0.0165	0.9478	-0.0637	0.6067	0.0368	0.8429	0.0604	0.9191	0.0952	0.6422	0.0348	0.8714	-0.2275	0.7215	-0.0455	0.9125	0.1820	0.5173
Gs2	Glutamate-ammor	1630653_a_at	-0.6859	0.0372	-0.9108	0.0126	-0.1039	0.7509	0.4155	0.2830	0.1915	0.4148	-0.2239	0.2767	-0.5424	0.6272	-0.0883	0.9003	0.4541	0.3424
---	---	1630654_at	0.0154	0.9229	0.0134	0.9195	0.1073	0.4827	0.0886	0.8200	0.0185	0.9314	-0.0702	0.6571	-0.1811	0.7550	-0.0298	0.9404	0.1512	0.5629
CG33159	CG33159	1630655_at	3.5303	0.0004	2.3516	0.0014	2.9880	0.0000	0.1318	0.7149	0.0760	0.6872	-0.0559	0.7564	-0.1317	0.8864	-0.6400	0.0782	-0.5083	0.1651
---	---	1630656_at	-0.0458	0.8783	-0.0867	0.5032	0.0898	0.7404	-0.1540	0.8190	-0.0815	0.8012	0.0725	0.8064	-0.1171	0.9092	-0.0470	0.9353	0.0702	0.8884
CG11902	CG11902	1630657_at	0.0261	0.9242	0.1945	0.5472	-0.2274	0.2832	-0.0492	0.9154	0.1033	0.4989	0.1525	0.2415	0.2357	0.8400	0.2526	0.6270	0.0170	0.9837
CG32226 /// DmirCG32226	CG32226	1630658_at	0.2146	0.4641	0.4030	0.2201	0.5048	0.1125	0.0209	0.9863	0.1578	0.6244	0.1368	0.6467	-0.1876	0.9032	0.3465	0.5552	0.5341	0.3559
CG40081	CG40081	1630659_at	0.2003	0.2116	-0.0248	0.8079	0.0716	0.6890	-0.0287	0.9603	0.0554	0.7600	0.0841	0.5769	-0.2081	0.6903	-0.1159	0.6154	0.0921	0.7026
CG14033	CG14033	1630660_at	2.3331	0.0015	2.1742	0.0014	2.4466	0.0000	0.0766	0.9283	0.8142	0.0067	0.7376	0.0062	-0.2586	0.8202	0.6627	0.1489	0.9212	0.0852
---	---	1630661_at	0.8169	0.0791	-0.1638	0.5757	0.1376	0.4123	0.1126	0.9445	0.4662	0.3203	0.3536	0.4145	-0.1433	0.8940	-0.4999	0.1957	-0.3565	0.3828
---	---	1630662_at	0.0073	0.9723	-0.0372	0.7679	0.1370	0.4150	0.0132	0.9883	0.1366	0.5439	0.1234	0.5483	-0.0573	0.9514	-0.0307	0.9462	0.0266	0.9463
CG15219	CG15219	1630663_a_at	0.0879	0.6080	-0.0152	0.8824	-0.0014	0.9946	0.1028	0.7803	0.1134	0.4786	0.0106	0.9570	-0.1409	0.8455	-0.0721	0.8619	0.0688	0.8599
---	---	1630664_at	-0.1227	0.5923	0.0226	0.8741	0.2373	0.1664	-0.0169	0.9838	-0.0990	0.6525	-0.0821	0.6891	-0.1276	0.7677	-0.0883	0.6605	0.0393	0.8736
mthl10	Mth-like 10	1630665_at	0.1072	0.4685	-0.0520	0.7164	-0.1115	0.4820	0.0174	0.9777	0.1522	0.3282	0.1348	0.3372	0.1546	0.8379	0.0200	0.9689	-0.1346	0.6985
---	---	1630666_at	0.3762	0.0602	0.1847	0.2086	0.3337	0.1483	0.0376	0.9601	0.0959	0.6685	0.0583	0.7912	-0.0927	0.9012	-0.1175	0.7046	-0.0247	0.9487
CG34043	CG34043	1630667_at	-4.0123	0.0010	-4.6943	0.0014	-3.7833	0.0001	1.1070	0.3433	0.3220	0.6745	-0.7850	0.1911	-0.2147	0.8740	-0.4021	0.4505	-0.1874	0.7612
jagn	jagunal	1630668_a_at	-0.3336	0.2832	0.8567	0.0327	1.1304	0.0003	0.0475	0.9435	-0.8946	0.0013	-0.9421	0.0006	-0.2509	0.8202	0.3568	0.4280	0.6077	0.2016
CG7299	CG7299	1630669_at	-0.0841	0.6327	-0.1099	0.4409	0.2746	0.1665	0.0659	0.8961	-0.0417	0.8471	-0.1075	0.5026	-0.1307	0.8492	0.0671	0.8678	0.1978	0.5041
CG1368	CG1368	1630670_at	0.3736	0.3703	0.0718	0.6437	0.2821	0.1731	-0.0328	0.9627	-0.2396	0.1991	-0.2067	0.2177	-0.2393	0.7997	-0.3057	0.4355	-0.0663	0.9026
Patj	discs lost	1630671_s_at	-0.1755	0.6947	0.3942	0.3433	0.2205	0.2383	0.1658	0.7409	0.0352	0.9085	-0.1306	0.5465	0.2792	0.8534	0.5875	0.3327	0.3083	0.6457
bcn92	bcn92	1630672_at	-0.4578	0.0898	0.0960	0.6227	-0.0852	0.7193	0.2465	0.5841	0.0260	0.9367	-0.2205	0.2881	0.2937	0.7387	0.4701	0.1948	0.1764	0.6644
CG32240	CG32240	1630673_at	0.2019	0.3944	0.1297	0.6244	0.0805	0.7377	0.0708	0.9321	0.1171	0.6780	0.0463	0.8741	-0.0121	0.9939	0.0370	0.9505	0.0491	0.9231
Trax	Translin associate	1630674_at	0.0132	0.9737	0.2052	0.3313	0.3905	0.0667	0.1846	0.7596	-0.0637	0.8495	-0.2483	0.2903	-0.0314	0.9862	0.2680	0.5234	0.2994	0.4747
CG4956	CG4956	1630675_at	0.2061	0.4725	0.0171	0.8736	0.2380	0.1435	0.1670	0.7208	0.1170	0.6164	-0.0500	0.8363	-0.1256	0.8202	-0.1620	0.4893	-0.0364	0.9086
CG30116	CG30116	1630676_at	0.0706	0.9288	0.7751	0.2671	1.2151	0.0003	0.1330	0.8546	-0.2332	0.3858	-0.3662	0.1176	-0.3179	0.9093	0.3767	0.7467	0.6946	0.4992
CG15100	CG15100	1630677_at	0.8136	0.0103	0.5353	0.1330	0.8085	0.0030	0.5565	0.0978	1.0905	0.0007	0.5340	0.0091	0.3340	0.7423	0.7849	0.0795	0.4510	0.3048
---	---	1630678_at	-0.0165	0.9245	0.1655	0.4631	0.1202	0.6586	-0.0111	0.9883	-0.0834	0.6821	-0.0723	0.7006	-0.0228	0.9893	0.1262	0.7723	0.1490	0.7128
CG1961 /// DsimCG1961	CG1961	1630679_at	-2.0410	0.0626	-0.0810	0.9153	-2.0702	0.0069	-2.0628	0.0770	-2.3509	0.0048	-0.2881	0.6833	0.0954	0.9893	-0.5544	0.7554	-0.6499	0.6969
inx5	inx5	1630680_at	0.2306	0.3692	-0.0566	0.7434	0.2611	0.1715	0.1394	0.8164	0.2418	0.3096	0.1024	0.6776	-0.1290	0.8692	0.0014	0.9991	0.1304	0.7087
---	---	1630681_at	0.0692	0.6267	-0.0015	0.9923	0.3251	0.1601	0.2088	0.5461	0.0605	0.7805	-0.1483	0.3707	0.0933	0.8760	0.1888	0.4164	0.0955	0.7170
Z4	Z4	1630682_at	0.3264	0.2016	-0.0112	0.9515	-0.1145	0.6389	-0.0102	0.9917	0.2426	0.2398	0.2528	0.1692	-0.1243	0.9238	0.0961	0.8756	0.2204	0.6328
Patr-1	CG5208	1630683_at	0.3620	0.2662	1.5162	0.0162	0.8063	0.0097	-0.3828	0.3739	-0.2679	0.2726	0.1149	0.6461	0.3168	0.7980	1.0304	0.0661	0.7136	0.1911
CG13032	CG13032	1630684_at	0.0552	0.8025	-0.1127	0.6105	-0.0363	0.9036	0.0376	0.9695	0.3205	0.2038	0.2828	0.2114	0.0241	0.9853	0.0773	0.8501	0.0532	0.8976
CG1314	CG1314	1630685_at	-0.0160	0.9536	-0.1198	0.6319	0.1852	0.3312	0.0868	0.9013	0.0499	0.8680	-0.0368	0.8924	-0.1207	0.9007	-0.0566	0.9178	0.0642	0.8949
CG2186	CG2186	1630686_at	-0.0488	0.9280	0.5233	0.5031	-0.2019	0.5952	-0.4620	0.2561	-0.1689	0.5078	0.2931	0.1737	0.2383	0.9421	0.4015	0.7339	0.1632	0.9068

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1630687_at	-0.0109	0.9583	-0.1213	0.5894	0.1218	0.5649	-0.0497	0.9353	0.0639	0.7681	0.1135	0.5164	0.0305	0.9816	0.2327	0.4563	0.2023	0.5367
Hsp83	Enhancer of seve	1630688_at	-0.0683	0.9332	-0.0124	0.9721	-0.2381	0.3902	0.1228	0.9592	0.4561	0.4930	0.3333	0.6008	0.4291	0.7266	0.5547	0.2799	0.1256	0.8555
CG33061	CG33061	1630689_at	0.1655	0.4062	0.0938	0.6951	0.0739	0.6344	-0.1566	0.7232	-0.1181	0.5862	0.0385	0.8692	0.0370	0.9701	0.0369	0.9266	0.0000	0.9999
CG33477	CG33477	1630690_at	0.0640	0.7386	0.0235	0.8287	-0.0797	0.6763	-0.0362	0.9656	0.0048	0.9889	0.0410	0.8769	0.0301	0.9767	0.1136	0.6765	0.0835	0.7728
---	---	1630691_at	-0.3960	0.0901	-0.5187	0.0361	-0.4172	0.0198	0.0462	0.9386	0.0572	0.7899	0.0111	0.9597	-0.0185	0.9852	-0.1563	0.5084	-0.1378	0.5704
CG31789	CG31789	1630692_at	0.1852	0.3251	0.1699	0.1642	0.2782	0.0869	-0.0359	0.9603	-0.0931	0.6716	-0.0572	0.7910	-0.0266	0.9741	0.0552	0.8380	0.0818	0.7192
---	---	1630693_at	0.1249	0.5314	-0.0476	0.7623	0.4805	0.0091	0.3389	0.2438	0.2333	0.1733	-0.1056	0.5295	-0.0738	0.9101	-0.0457	0.8960	0.0282	0.9316
CG34394	CG15411	1630694_at	-0.0677	0.8272	0.4673	0.5536	0.8667	0.0804	0.0962	0.8816	0.0149	0.9647	-0.0813	0.7307	-0.1591	0.9538	0.7024	0.3768	0.8615	0.2997
CG1970	CG1970	1630695_at	-0.2211	0.5018	0.1829	0.6125	0.1091	0.5873	-0.3495	0.2511	-0.7103	0.0029	-0.3608	0.0335	-0.1794	0.9087	-0.1859	0.7882	-0.0065	0.9947
CG12972	CG12972	1630696_at	2.3504	0.0010	2.0515	0.0172	2.3031	0.0001	-0.1421	0.7658	0.2163	0.2917	0.3584	0.0531	-0.0401	0.9852	0.1091	0.8782	0.1492	0.8057
---	---	1630697_at	0.1784	0.2516	-0.0351	0.7823	-0.0198	0.9251	0.0618	0.8936	0.1385	0.3776	0.0768	0.6235	0.1222	0.8609	-0.1371	0.6556	-0.2592	0.3695
CG3192	CG3192	1630698_at	-0.0581	0.8846	-0.0359	0.8680	-0.0132	0.9554	-0.1049	0.8327	-0.4361	0.0283	-0.3312	0.0537	-0.2137	0.8425	-0.4532	0.2902	-0.2395	0.6148
betaInt-nu	integrin beta v	1630699_at	-2.4417	0.0004	-1.1342	0.1279	-2.3348	0.0000	-1.2374	0.0422	-1.2157	0.0044	0.0217	0.9597	-0.1149	0.9525	0.0768	0.9341	0.1917	0.7787
---	---	1630700_at	2.1125	0.0214	0.1807	0.8243	1.0947	0.0530	0.8611	0.1341	0.9568	0.0136	0.0957	0.8056	-0.4022	0.9291	-0.7965	0.6101	-0.3942	0.8306
CG10778	CG10778	1630701_at	-0.0863	0.7119	0.9181	0.0290	1.2265	0.0024	0.0365	0.9592	-0.4968	0.0158	-0.5334	0.0070	-0.2087	0.8814	0.4846	0.3534	0.6933	0.2130
CG14590	CG14590	1630702_at	-1.3494	0.0031	-0.6847	0.1477	-1.3184	0.0008	-0.5249	0.2977	-0.9074	0.0085	-0.3825	0.1476	0.0890	0.9683	-0.4230	0.4843	-0.5119	0.3953
CG8963	CG8963	1630703_at	0.3778	0.0743	0.9598	0.0101	0.7545	0.0015	-0.3386	0.2095	-0.2626	0.1044	0.0761	0.6488	-0.0838	0.9216	0.2909	0.2924	0.3747	0.2081
---	---	1630704_at	0.2824	0.4469	0.0236	0.9471	0.1441	0.4518	0.2293	0.6574	0.3429	0.1649	0.1136	0.6587	0.1612	0.8906	0.0526	0.9409	-0.1086	0.8484
CG4511	CG4511	1630705_at	-0.4740	0.0409	-0.9844	0.0248	-1.0517	0.0052	0.1962	0.7205	1.1272	0.0015	0.9309	0.0021	0.1969	0.8521	0.4618	0.2733	0.2649	0.5670
---	---	1630706_at	0.0580	0.7855	0.0176	0.8803	0.1708	0.3119	0.1600	0.7327	0.1223	0.5910	-0.0377	0.8791	0.0628	0.9152	0.1179	0.5897	0.0551	0.8318
edl	ETS-domain lacki	1630707_at	0.5966	0.0041	0.3706	0.1908	0.0779	0.8495	0.0798	0.8251	0.3538	0.0197	0.2740	0.0355	0.4067	0.7251	0.1311	0.8477	-0.2756	0.6095
---	---	1630708_at	0.1474	0.4210	0.1269	0.2763	-0.0758	0.6662	-0.0484	0.9463	-0.0303	0.9148	0.0181	0.9431	0.1269	0.7752	-0.0342	0.9075	-0.1611	0.4096
---	---	1630709_at	0.2225	0.2717	0.1945	0.3354	0.0227	0.9030	-0.0497	0.9507	0.0468	0.8738	0.0965	0.6784	0.1284	0.8740	0.1876	0.5776	0.0592	0.8921
GluRIIA	glutamate recepto	1630710_at	-2.2649	0.0012	-2.5584	0.0303	-3.3904	0.0001	-0.3577	0.2753	-0.1299	0.5283	0.2278	0.1861	0.5740	0.8013	-0.3025	0.8040	-0.8765	0.3695
CG11175	CG11175	1630711_at	-0.4189	0.2164	-0.3335	0.3232	-1.0160	0.0087	-0.1019	0.9042	0.0434	0.9096	0.1453	0.5999	0.2257	0.8655	-0.1095	0.8866	-0.3352	0.5499
---	---	1630712_at	0.2687	0.2882	0.1803	0.4253	-0.0923	0.6114	0.0457	0.9500	0.1730	0.4037	0.1274	0.5140	0.1874	0.7893	0.0538	0.9068	-0.1337	0.6867
SrpRbeta	Signal recognition	1630713_at	0.7902	0.0062	0.6734	0.2630	0.2518	0.1316	0.1259	0.7424	0.8434	0.0010	0.7175	0.0012	0.5626	0.6955	0.6558	0.2552	0.0933	0.9100
---	---	1630714_at	0.1678	0.3540	0.1438	0.3996	-0.0307	0.8765	-0.0465	0.9252	-0.0635	0.7117	-0.0170	0.9253	0.1356	0.8049	-0.0723	0.8055	-0.2079	0.3764
CG10588	CG10588	1630715_at	-0.0302	0.9011	-0.0929	0.5103	0.1222	0.4251	0.1020	0.8475	0.1863	0.3459	0.0843	0.6778	-0.1126	0.8202	-0.0895	0.6971	0.0232	0.9352
---	---	1630716_at	0.1437	0.4869	-0.1534	0.2534	0.0385	0.8381	0.1641	0.6506	0.3051	0.0822	0.1410	0.3842	0.0191	0.9848	0.0305	0.9335	0.0113	0.9751
cg	comb gap	1630717_s_at	0.9460	0.0057	-0.0790	0.7352	-0.8327	0.0551	-0.3322	0.4944	0.7643	0.0099	1.0965	0.0012	0.2313	0.8692	-0.1365	0.8616	-0.3678	0.5340
Ddr	Discoidin domain	1630718_at	0.0050	0.9812	-0.0773	0.6325	0.0908	0.6327	0.1475	0.7582	0.0406	0.8836	-0.1069	0.6073	0.0227	0.9824	0.0792	0.7912	0.0565	0.8555
CG32102	CG32102	1630719_at	0.1360	0.5707	-0.0042	0.9892	-0.0405	0.8506	0.1196	0.8546	0.0882	0.7542	-0.0314	0.9127	0.0696	0.9238	-0.0641	0.8461	-0.1336	0.6100
CG32249	CG32249	1630720_at	0.1878	0.3094	0.0670	0.5251	0.2126	0.3748	-0.1582	0.7970	-0.1502	0.5793	0.0080	0.9799	-0.1279	0.8122	-0.1103	0.6483	0.0176	0.9556
auxillin	auxilin	1630721_s_at	0.3475	0.0489	0.3360	0.1353	0.2210	0.2636	-0.1779	0.6059	0.0810	0.6821	0.2589	0.0946	-0.0822	0.9117	0.1076	0.7202	0.1897	0.4851
---	---	1630722_at	-0.0486	0.8397	0.0120	0.9680	-0.0848	0.7467	-0.0881	0.9218	-0.0704	0.8403	0.0178	0.9587	0.0116	0.9913	0.0506	0.8665	0.0391	0.8943
CG8708	CG8708	1630723_a_at	0.1031	0.6030	0.1094	0.4670	0.3505	0.0590	0.1009	0.8589	0.0781	0.7475	-0.0228	0.9274	-0.0961	0.8673	-0.0983	0.7029	-0.0022	0.9956
CG1890	CG1890	1630724_at	-0.1324	0.6840	-0.4197	0.2152	-0.1986	0.6131	0.3981	0.4291	0.7806	0.0123	0.3824	0.1211	0.0455	0.9894	0.4168	0.5825	0.3713	0.6311
CG14572	CG14572	1630725_at	0.2068	0.4938	-1.1280	0.0280	-0.7868	0.3368	0.3358	0.7803	1.3339	0.0157	0.9981	0.0329	-0.1691	0.9421	-0.1993	0.8353	-0.0302	0.9795
CG32652	CG32652	1630726_at	0.3464	0.0464	0.0398	0.7060	0.0591	0.7856	0.0134	0.9860	0.2245	0.2022	0.2111	0.1804	0.0166	0.9875	-0.0062	0.9911	-0.0228	0.9463
CG5070	CG5070	1630727_at	0.1168	0.5519	0.1676	0.3766	0.2283	0.3298	0.0551	0.9436	-0.0349	0.9103	-0.0900	0.7017	0.2044	0.8692	0.0902	0.9050	-0.1142	0.8599
phr	photolyase	1630728_a_at	-0.2110	0.3175	0.3963	0.1546	0.1182	0.5714	-0.4434	0.3619	-0.5951	0.0382	-0.1517	0.5833	-0.0511	0.9717	0.1000	0.8369	0.1511	0.7118
tefu	telomere fusion	1630729_at	-0.4326	0.0909	-0.6691	0.0324	-0.7083	0.0055	-0.0978	0.8959	0.5646	0.0299	0.6624	0.0094	-0.3100	0.7485	0.0422	0.9497	0.3523	0.4003
CG6808	CG6808	1630730_at	-0.4434	0.0658	-0.9088	0.0336	-0.6900	0.0261	-0.1016	0.8889	0.4160	0.0889	0.5176	0.0258	-0.2327	0.7291	0.0702	0.8613	0.3030	0.3047
CG14965	CG14965	1630731_at	-0.1582	0.5816	-0.0812	0.7133	-0.4332	0.1601	-0.3325	0.6228	0.0640	0.8879	0.3965	0.1856	-0.0391	0.9743	0.0752	0.8592	0.1143	0.7456
Trn-SR	Transportin-Serin	1630732_at	0.4887	0.0689	0.0451	0.7871	-0.0597	0.7656	-0.0171	0.9777	0.6488	0.0021	0.6659	0.0011	0.2555	0.7685	0.3038	0.4048	0.0483	0.9237
CG31203	CG31203	1630733_s_at	0.1489	0.3366	-0.0079	0.9429	0.1342	0.4760	0.1296	0.7187	0.1203	0.4814	-0.0094	0.9637	-0.2453	0.7307	-0.1301	0.7062	0.1152	0.7439
Cpr92A	CG6240	1630734_at	-0.0309	0.8831	-0.1623	0.5240	-0.1265	0.4852	0.0006	0.9994	0.1145	0.5300	0.1138	0.4854	-0.0654	0.9324	-0.1069	0.7029	-0.0415	0.9033
mRpL21	mitochondrial ribo	1630735_at	0.1163	0.5605	0.4698	0.0843	0.0707	0.8460	-0.0962	0.8508	-0.0534	0.8179	0.0428	0.8424	0.2637	0.8157	0.2457	0.6256	-0.0180	0.9821
nub	CG15488	1630736_at	0.0812	0.6987	-0.0178	0.8606	0.1213	0.5467	0.1371	0.6958	0.1100	0.5228	-0.0271	0.8883	0.0422	0.9717	-0.0885	0.8175	-0.1307	0.6883
CG4661	CG4661	1630737_at	0.0448	0.8377	-0.0200	0.8874	-0.1066	0.7669	-0.0208	0.9863	-0.1065	0.7606	-0.0857	0.7932	0.0469	0.9742	-0.0858	0.8665	-0.1327	0.7513
Pten	Pten	1630738_s_at	0.7136	0.0364	0.1789	0.6809	-0.0473	0.9136	-0.0508	0.9578	0.6475	0.0191	0.6983	0.0084	0.1945	0.9174	0.0631	0.9516	-0.1314	0.8818
---	---	1630739_at	0.1158	0.5322	0.1978	0.2486	-0.1443	0.4965	-0.0843	0.8905	-0.0660	0.7930	0.0183	0.9429	0.2077	0.7393	0.0697	0.8468	-0.1380	0.6296
CG4363 /// DyakCG4363	CG4363	1630740_at	0.2072	0.4601	-0.2513	0.3376	0.0196	0.9303	0.1735	0.7351	0.3012	0.1840	0.1277	0.5724	-0.1522	0.9013	-0.2288	0.6394	-0.0766	0.9023
PGRP-LB	Peptidoglycan rec	1630741_s_at	-1.8038	0.0008	-2.8133	0.0042	-1.8937	0.0017	0.7245	0.4034	1.1487	0.0256	0.4242	0.3385	-0.1235	0.8951	0.1420	0.7244	0.2655	0.4627
ppk28	pickpocket 28	1630742_at	-0.0251	0.9551	0.0671	0.6951	-0.0767	0.8024	-0.0032	0.9962	-0.0134	0.9698	-0.0103	0.9711	0.0480	0.9798	0.0471	0.9460	-0.0009	0.9992
CG11971	anon-fast-evolvin	1630743_at	0.3706	0.0482	0.0103	0.9666	-0.0833	0.6674	-0.1486	0.7389	0.2943	0.1336	0.4428	0.0201	-0.1220	0.9011	-0.0901	0.8551	0.0319	0.9494
---	---	1630744_at	0.1570	0.4163	0.0710	0.6528	0.1271	0.4485	0.0221	0.9803	-0.1119	0.6422	-0.1340	0.5198	-0.0786	0.9474	-0.0501	0.9312	0.0285	0.9564
CG9099	CG9099	1630745_at	0.0761	0.7026	0.2800	0.2034	0.4917	0.0256	-0.1861	0.6893	-0.2301	0.2906	-0.0441	0.8609	-0.1210	0.8846	0.1548	0.6524	0.2758	0.3921
CG31807	CG31807	1630746_at	0.1204	0.4664	-0.0282	0.7870	0.2439	0.3177	0.0842	0.8906	0.0800	0.7405	-0.0041	0.9873	-0.1108	0.8461	0.1159	0.6512	0.2267	0.3503
CG40445	CG40445	1630747_at	-0.0682	0.7573	0.0399	0.8240	0.1643	0.4315	0.0037	0.9956	-0.0922	0.7033	-0.0960	0.6590	0.0412	0.9689	0.1722	0.5509	0.1310	0.6609
CG32653	CG32653	1630748_at	0.3331	0.3680	-0.9402	0.0456	-0.7602	0.0956	0.3308	0.7803	1.2049	0.0227	0.8740	0.0516	0.0041	0.9984	-0.0346	0.9558	-0.0387	0.9431
CG11052	CG11052	1630749_at	-0.0478	0.8128	-0.1311	0.6128	0.0834	0.7124	0.1421	0.7795	-0.0650	0.8018	-0.2071	0.2773	-0.1540	0.8692	-0.1785	0.6656	-0.0245	0.9649
CG5323	CG5323	1630750_at	-0.0905	0.6326	-0.0878	0.6681	-0.1588	0.4608	-0.0110	0.9909	0.0875	0.7230	0.0986	0.6528	0.0031	0.9982	-0.0516	0.9054	-0.0547	0.8884
---	---	1630751_at	-0.1209	0.4204	-0.2319	0.2419	-0.0694	0.7869	0.3478	0.2747	0.3071	0.1014	-0.0408	0.8516	0.0095	0.9939	-0.0124	0.9821	-0.0219	0.9582
CG6982	CG6982	1630752_at	-0.3988	0.4144	0.5767	0.2240	0.3438	0.0460	-0.3027	0.7344	-0.7474	0.0631	-0.4448	0.2097	-0.0143	0.9952	0.4058	0.4600	0.4201	0.4567
ric8a	ric8a	1630753_at	0.9318	0.0016	1.1373	0.0234	0.7421	0.0014	-0.1171	0.7855	-0.0183	0.9439	0.0988	0.5760	-0.0490	0.9467	0.0112	0.9779	0.0602	0.8299
ORMDL	ORMDL	1630754_at	0.9314	0.0190	0.8781	0.0230	0.8005	0.0106	0.0733	0.9011	-0.1555	0.4319	-0.2288	0.1791	0.0646	0.9689	-0.2709	0.5501	-0.3355	0.4534
CG9226	CG9226	1630755_at	-0.3466	0.0371	0.0627	0.8393	-0.1752	0.2459	0.0159	0.9834	-0.1671	0.3470	-0.1830	0.2429	0.0837	0.8988	0.0868	0.7692	0.0031	0.9945

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Mipp2 /// raptor	dRaptor /// Multiph	1630756_at	-0.4036	0.4006	0.2591	0.4367	0.1556	0.3838	-0.0882	0.9061	-0.1256	0.6425	-0.0374	0.8974	0.0123	0.9964	0.5198	0.4258	0.5075	0.4533
---	---	1630757_at	0.1980	0.2420	0.0757	0.5246	0.0720	0.7866	-0.2872	0.4397	0.0504	0.8480	0.3376	0.0661	-0.3467	0.5754	-0.0482	0.9093	0.2985	0.2960
CG13033	CG13033	1630758_at	0.1753	0.4042	0.2428	0.2061	0.2689	0.2207	0.0111	0.9872	-0.0183	0.9381	-0.0294	0.8802	-0.1511	0.8692	-0.0861	0.8678	0.0650	0.8974
---	---	1630759_at	-0.0007	0.9973	0.0462	0.6899	-0.1859	0.2057	-0.0224	0.9791	0.1019	0.6730	0.1242	0.5506	0.1586	0.7644	0.0912	0.7269	-0.0674	0.8088
---	---	1630760_at	0.0743	0.6840	0.2710	0.2534	0.2529	0.2834	-0.2184	0.6820	-0.3262	0.1856	-0.1079	0.6776	0.0431	0.9742	0.0060	0.9935	-0.0371	0.9371
CG11854	CG11854	1630761_at	0.2884	0.0876	0.3036	0.4433	-0.1934	0.3106	0.0238	0.9838	0.0989	0.7601	0.0751	0.8047	0.3177	0.7324	-0.0056	0.9949	-0.3233	0.4301
CG9918	CG9918	1630762_at	-1.9047	0.0037	0.1997	0.3361	0.0537	0.8576	-0.1388	0.8908	-2.3142	0.0003	-2.1754	0.0002	-0.1299	0.8608	-0.4322	0.1400	-0.3023	0.3205
CG6845	CG6845	1630763_at	0.1564	0.4973	-0.0685	0.8768	-0.5957	0.0325	-0.4418	0.2248	-0.0371	0.9013	0.4047	0.0433	0.0097	0.9964	-0.2812	0.6259	-0.2910	0.6140
CG8683	CG8683	1630764_at	0.2603	0.4432	0.2083	0.3840	0.0386	0.8784	-0.0088	0.9937	0.6306	0.0113	0.6393	0.0064	0.1638	0.8956	0.6890	0.1377	0.5252	0.2757
CG15547	CG15547	1630765_at	-0.0342	0.8897	0.5088	0.2937	0.5001	0.0297	-0.0549	0.9465	-0.6222	0.0150	-0.5672	0.0138	-0.0019	0.9996	0.0638	0.9301	0.0657	0.9173
CG13800	CG13800	1630766_at	0.8312	0.0878	1.7753	0.0045	1.4406	0.0006	-0.2027	0.6558	-0.4545	0.0437	-0.2517	0.1971	0.1536	0.9400	0.4247	0.5141	0.2711	0.6993
CG31262	CG31262	1630767_at	0.3511	0.4846	-0.8159	0.0388	-0.2259	0.7057	0.2189	0.8090	1.2239	0.0054	1.0051	0.0079	-0.4790	0.8000	-0.0576	0.9647	0.4214	0.6195
achi /// vis	achintya /// visma	1630768_s_at	-0.6982	0.0110	0.0167	0.9612	0.0900	0.6307	0.0654	0.9345	-0.2155	0.3694	-0.2809	0.1800	-0.0644	0.9460	0.5118	0.0871	0.5761	0.0828
---	---	1630769_at	-0.1699	0.4207	0.0874	0.6615	0.2780	0.0711	0.0867	0.9013	-0.1058	0.6827	-0.1925	0.3559	-0.1478	0.7677	0.1128	0.6191	0.2606	0.2338
---	---	1630770_at	-0.2491	0.1625	-0.5363	0.0694	-0.7503	0.1196	0.0142	0.9904	0.3415	0.1908	0.3273	0.1620	-0.0120	0.9914	-0.0759	0.8185	-0.0640	0.8444
l(2)37Cc	transcription unit I	1630771_s_at	0.0502	0.7754	0.0182	0.9437	0.1714	0.3669	0.0032	0.9956	0.0603	0.7640	0.0571	0.7552	-0.2599	0.7187	0.0000	1.0000	0.2599	0.3921
sqz	squeeze	1630772_at	-0.5829	0.4437	-0.7259	0.0486	-1.1487	0.0045	-0.1090	0.9447	-0.2224	0.6621	-0.1134	0.8243	0.1713	0.9445	-0.1900	0.8518	-0.3612	0.6520
CG10899	CG10899	1630773_a_at	0.0413	0.8337	0.0778	0.4261	-0.0307	0.8589	0.0408	0.9329	0.0893	0.5634	0.0485	0.7566	0.1262	0.8320	0.1557	0.5471	0.0295	0.9325
sgg	Glycogen Synthas	1630774_s_at	-0.0750	0.8856	0.2500	0.5205	0.0761	0.7398	-0.0349	0.9705	-0.1927	0.4401	-0.1578	0.4947	0.2294	0.9092	0.4689	0.5259	0.2395	0.7761
---	---	1630775_at	0.0983	0.5580	0.1833	0.1419	0.1239	0.5279	0.1005	0.7937	0.0351	0.8652	-0.0654	0.6914	0.0973	0.8717	0.0870	0.7618	-0.0103	0.9785
CG1421	CG1421	1630776_at	0.0267	0.8923	0.0649	0.6542	-0.0111	0.9484	-0.0245	0.9649	-0.0241	0.9074	0.0004	0.9982	-0.0172	0.9869	0.0714	0.8185	0.0886	0.7487
---	---	1630777_at	-0.0304	0.8844	-0.2848	0.2789	-0.2676	0.1385	0.1110	0.8987	0.6114	0.0387	0.5004	0.0542	-0.0715	0.9342	0.1008	0.7626	0.1723	0.5585
CG10116	CG10116	1630778_at	-0.5688	0.8179	-0.1409	0.4209	0.0298	0.9162	0.0794	0.9924	-1.7044	0.3359	-1.7838	0.2555	0.0092	0.9994	-1.2032	0.5543	-1.2124	0.5579
CG2061	CG2061	1630779_s_at	-0.2561	0.1808	0.1713	0.6255	0.4924	0.0665	0.0827	0.8738	-0.1272	0.5057	-0.2099	0.1937	-0.1269	0.9339	0.2782	0.5897	0.4051	0.4114
hydra	hydra	1630780_a_at	0.0793	0.6973	-0.2547	0.2186	-0.3303	0.1271	-0.0755	0.9139	0.2198	0.3254	0.2954	0.1318	-0.2504	0.7202	-0.2182	0.4531	0.0322	0.9369
fs(2)ltoPP43	female sterile (2) l	1630781_at	0.4184	0.2305	-0.0825	0.7153	-0.1646	0.3826	-0.2834	0.6204	0.0927	0.7903	0.3761	0.1387	-0.3114	0.7362	-0.3740	0.3287	-0.0625	0.9086
CG34420	CG5917	1630782_at	0.1548	0.3313	-0.1398	0.3942	-0.0264	0.9076	0.1843	0.6337	-0.0020	0.9943	-0.1863	0.2765	0.0447	0.9503	-0.1244	0.5825	-0.1691	0.4411
---	---	1630783_at	0.1709	0.3788	0.0394	0.7288	-0.1736	0.2612	-0.0163	0.9777	0.0697	0.6761	0.0860	0.5497	0.0743	0.9011	-0.0745	0.7828	-0.1488	0.5141
CG9990	CG9990	1630784_a_at	-0.2543	0.5797	-0.9789	0.1424	-1.3307	0.0015	0.3091	0.6954	1.3851	0.0033	1.0760	0.0060	0.6407	0.7116	0.7924	0.2533	0.1518	0.8771
---	---	1630785_at	-0.0830	0.6515	-0.0280	0.8386	-0.1000	0.6146	-0.1154	0.8008	-0.1603	0.3964	-0.0448	0.8303	0.0743	0.9238	0.0299	0.9430	-0.0444	0.9020
Chd3	Chd3	1630786_at	-0.0773	0.8100	-0.4397	0.3811	-1.1170	0.0356	-0.5701	0.2876	0.6734	0.0392	1.2436	0.0017	-0.2188	0.9357	0.2875	0.7900	0.5063	0.5814
CG9879	CG9879	1630787_at	0.3676	0.0343	0.3623	0.1131	0.3074	0.1179	-0.0337	0.9540	-0.0547	0.7780	-0.0209	0.9148	0.0510	0.9340	0.0044	0.9925	-0.0465	0.8561
Pros26.4	26S proteasome s	1630788_at	-0.1130	0.5038	0.5315	0.1359	1.1430	0.0002	0.2205	0.5735	-0.5140	0.0188	-0.7345	0.0023	-0.3510	0.5875	0.2445	0.3900	0.5956	0.0755
CG5897	CG5897	1630789_at	0.2758	0.4230	0.1050	0.4914	0.1184	0.7361	0.0082	0.9943	-0.0338	0.9202	-0.0420	0.8826	0.0911	0.9562	-0.0870	0.9060	-0.1781	0.7484
---	---	1630790_s_at	0.0665	0.6792	0.0671	0.6463	0.0088	0.9801	-0.0035	0.9961	-0.2638	0.2575	-0.2603	0.2099	0.1695	0.8298	-0.0591	0.9075	-0.2285	0.5050
pyd	tamou	1630791_a_at	0.4318	0.4755	0.7770	0.2789	0.8533	0.0115	0.0698	0.9496	0.0446	0.9175	-0.0251	0.9486	-0.0704	0.9893	0.3209	0.8220	0.3913	0.7558
rept	reptin	1630792_at	-0.1990	0.4068	1.0227	0.0212	1.2799	0.0017	-0.0225	0.9835	-0.8727	0.0047	-0.8502	0.0032	-0.1373	0.9284	0.5700	0.2361	0.7073	0.1792
---	---	1630793_s_at	0.0373	0.8326	0.0309	0.8470	-0.2146	0.1739	-0.1775	0.6186	0.0104	0.9701	0.1879	0.2379	-0.0457	0.9611	-0.0220	0.9590	0.0238	0.9486
CG2097	CG2097	1630794_at	0.1589	0.3047	0.0308	0.9498	0.2949	0.0576	0.0240	0.9699	0.2241	0.1667	0.2002	0.1685	-0.2355	0.7953	-0.0130	0.9867	0.2225	0.5853
CG12125 /// DsimCG12125	CG12125	1630795_at	-0.2260	0.3648	-0.4220	0.1896	-0.4101	0.1000	0.0838	0.8562	0.4305	0.0179	0.3467	0.0277	0.0145	0.9943	0.1204	0.8546	0.1059	0.8672
CG31484	CG31484	1630796_at	0.2123	0.2815	-0.0775	0.8363	0.0432	0.7799	0.1097	0.9036	0.2137	0.4873	0.1039	0.7424	-0.0668	0.9342	-0.0924	0.7697	-0.0256	0.9434
CG32582	CG32582	1630797_at	1.2372	0.0250	1.6242	0.0081	1.6462	0.0002	0.2876	0.7067	0.1924	0.6194	-0.0952	0.8086	0.2536	0.8465	0.5122	0.3340	0.2586	0.6607
CG30077	CG30077	1630798_at	-0.2770	0.2650	-0.2746	0.1476	-0.4891	0.0815	0.2232	0.7135	0.3094	0.2663	0.0862	0.7796	0.4769	0.5754	0.3843	0.2973	-0.0926	0.8492
---	---	1630799_at	-0.0983	0.6926	-0.0023	0.9923	-0.0010	0.9972	0.0189	0.9838	-0.1232	0.5962	-0.1422	0.4848	-0.1176	0.8521	-0.1634	0.5501	-0.0458	0.8985
CG34040	CG34040	1630800_s_at	0.1445	0.5256	0.1187	0.4328	0.2348	0.2363	-0.1802	0.5498	-0.0964	0.5667	0.0838	0.5907	-0.1311	0.8806	0.0534	0.9175	0.1845	0.6074
Gsc	Muenster 72	1630801_at	-0.0186	0.9495	-0.0290	0.8527	0.1098	0.6029	0.0670	0.9387	0.0010	0.9980	-0.0660	0.8196	-0.0791	0.8943	-0.0370	0.9151	0.0421	0.8915
Cyp6d4	Cyp6d4	1630802_at	0.3021	0.2203	-0.3424	0.2833	0.0062	0.9757	0.0194	0.9852	-0.0095	0.9790	-0.0289	0.9193	-0.3494	0.6749	-0.6560	0.0683	-0.3066	0.3723
Taf10b	TBP-associated fa	1630803_at	-0.2844	0.3895	0.2985	0.4458	0.1269	0.5577	-0.2600	0.5688	-0.3240	0.1642	-0.0640	0.8077	-0.0347	0.9875	0.3505	0.4954	0.3852	0.4568
Oatp33Eb	Organic anion trar	1630804_at	2.7813	0.0150	2.0782	0.0445	3.1819	0.0006	0.8321	0.5626	0.3521	0.6749	-0.4800	0.4980	-0.5974	0.8461	-0.5612	0.6932	0.0362	0.9856
noi	noisette	1630805_at	-0.1141	0.5399	-0.2686	0.2626	0.1068	0.6263	0.1036	0.8174	0.3279	0.0657	0.2244	0.1527	-0.2571	0.7726	0.1414	0.7558	0.3985	0.3079
fs(1)K10	female sterile (1) l	1630806_at	-0.4158	0.1198	-0.6810	0.0303	-0.9253	0.0132	-0.0108	0.9922	0.3793	0.1033	0.3901	0.0647	0.1892	0.9012	0.1371	0.8602	-0.0522	0.9471
CG33111	CG33111	1630807_at	-0.1662	0.4829	-0.0954	0.7950	0.0362	0.8203	0.2041	0.5350	0.2608	0.1271	0.0567	0.7630	-0.0886	0.9416	0.2832	0.4495	0.3718	0.3307
rasp	sightless	1630808_at	0.6368	0.0238	0.5795	0.0713	0.6573	0.0023	-0.1756	0.6120	0.0933	0.6290	0.2689	0.0839	-0.1924	0.8510	0.0302	0.9652	0.2225	0.6257
CG40249	CG40249	1630809_at	0.0056	0.9806	-0.0485	0.7590	-0.0579	0.7786	0.1566	0.7140	0.1169	0.5806	-0.0396	0.8615	0.1461	0.7810	0.0671	0.8208	-0.0790	0.7647
---	---	1630810_at	0.0909	0.5689	0.2439	0.0750	0.0208	0.9299	-0.2028	0.5766	-0.0753	0.7299	0.1275	0.4719	-0.0319	0.9742	0.0216	0.9585	0.0535	0.8721
CG31624 /// DereCG31624	CG31624	1630811_at	0.0629	0.7553	-0.6709	0.0355	-0.2619	0.1021	0.3931	0.3016	0.6662	0.0096	0.2731	0.1708	-0.2335	0.6749	-0.1722	0.4530	0.0613	0.8310
---	---	1630812_at	0.3084	0.1707	-0.0007	0.9990	0.1188	0.7604	0.0583	0.9322	0.0610	0.8084	0.0027	0.9915	0.0945	0.9421	-0.2177	0.6120	-0.3122	0.4507
RpL11	Ribosomal protein	1630813_at	0.4677	0.0151	0.8316	0.0223	0.9180	0.0020	0.0389	0.9532	-0.3625	0.0520	-0.4014	0.0223	0.1682	0.7651	-0.0130	0.9748	-0.1813	0.4655
CG10353	CG10353	1630814_s_at	-0.8168	0.0440	-0.1424	0.4402	-0.6555	0.0239	-0.2428	0.6908	-0.1785	0.5577	0.0644	0.8434	0.2175	0.8541	0.4921	0.3002	0.2745	0.6005
CG14277	CG14277	1630815_at	0.1592	0.4076	-0.1739	0.4209	0.0852	0.7733	0.1103	0.8623	0.2259	0.3264	0.1156	0.6151	-0.0803	0.9220	-0.1274	0.6732	-0.0471	0.9006
CG7375 /// DyakCG7375	CG7375	1630816_at	0.2799	0.1782	0.7568	0.0134	0.7122	0.0012	0.1198	0.7419	-0.1321	0.4251	-0.2519	0.0800	0.2286	0.7215	0.4687	0.0912	0.2400	0.3828
CG17528	CG17528	1630817_s_at	-0.2067	0.3163	-0.0345	0.9406	0.1782	0.3425	0.1407	0.7874	0.3447	0.1073	0.2040	0.2945	-0.1171	0.9238	0.3478	0.3815	0.4649	0.2674
CG30178	CG30178	1630818_at	-0.1738	0.2991	-0.1700	0.4536	-0.0334	0.8659	0.1590	0.6257	-0.0589	0.7607	-0.2178	0.1291	-0.0326	0.9717	0.0378	0.9167	0.0704	0.8044
CG7046	CG7046	1630819_at	0.2140	0.2534	-0.0021	0.9864	0.0982	0.4929	0.0316	0.9599	0.0760	0.6876	0.0444	0.8113	0.0015	0.9992	-0.1638	0.4280	-0.1653	0.4370
CG17454	CG17454	1630820_at	0.3885	0.2521	0.7815	0.0385	0.1777	0.3679	-0.0711	0.8987	0.0729	0.7345	0.1440	0.3976	0.4331	0.6874	0.4713	0.2664	0.0382	0.9510
CG2680	CG2680	1630821_at	0.1547	0.8750	-0.3529	0.2110	-0.0036	0.9881	0.8180	0.2067	0.2476	0.5550	-0.5705	0.1011	0.4077	0.8513	-0.2327	0.8500	-0.6403	0.4913
CG13078	CG13078	1630822_at	0.1249	0.6779	-0.3763	0.2687	-0.2592	0.2927	0.1440	0.8198	0.2886	0.2452	0.1446	0.5532	-0.0137	0.9939	-0.2907	0.4997	-0.2771	0.5299
CG32210 /// DmirCG32210	CG32210	1630823_at	-0.3567	0.1477	-0.8707	0.0688	-1.0515	0.0024	-0.0868	0.9158	0.5299	0.0449	0.6167	0.0153	0.1524	0.8978	0.1297	0.8221	-0.0228	0.9738
CG33145	CG33145	1630824_a_at	-0.2223	0.2763	-0.7683	0.0342	-1.1622	0.0009	-0.2695	0.6202	0.6373	0.0265	0.9068	0.0034	-0.0528	0.9598	0.0591	0.8933	0.1119	0.7442

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG5798	CG5798	1630825_at	-0.2140	0.3304	0.1663	0.6285	0.1229	0.5164	-0.0278	0.9683	-0.1437	0.4495	-0.1159	0.5116	0.0623	0.9429	0.1616	0.5644	0.0992	0.7484
---	---	1630826_at	-0.3212	0.1410	-0.0787	0.6540	-0.0548	0.7916	0.0948	0.7895	0.0407	0.8229	-0.0541	0.7276	-0.1161	0.8409	0.0343	0.9275	0.1504	0.5499
---	---	1630827_s_at	-0.0497	0.9118	0.0850	0.7740	0.6136	0.0405	0.0020	0.9988	0.7284	0.0664	0.7264	0.0441	-0.4766	0.7363	0.8838	0.1394	1.3605	0.0607
---	---	1630828_at	0.0800	0.6386	0.0475	0.8550	0.1622	0.3325	0.1687	0.7020	0.0731	0.7635	-0.0956	0.6465	-0.0299	0.9816	0.0635	0.8721	0.0934	0.7749
PIP5K59B	PIP5K59B	1630829_at	-0.0125	0.9752	-0.0555	0.6025	-0.3482	0.1285	-0.1389	0.8649	-0.0247	0.9542	0.1142	0.7093	0.0103	0.9928	0.0611	0.8684	0.0508	0.8880
SpdS	Spermidine Synth	1630830_a_at	1.4680	0.0046	0.3864	0.1299	1.2556	0.0047	0.4275	0.4442	1.0660	0.0047	0.6384	0.0269	-0.2347	0.8680	0.0397	0.9652	0.2743	0.6476
CG7695	CG7695	1630831_at	1.6408	0.0015	2.8059	0.0036	2.7098	0.0005	-0.3029	0.8356	-1.4320	0.0176	-1.1292	0.0297	-0.0777	0.9514	0.0487	0.9353	0.1264	0.7754
---	---	1630832_at	0.0941	0.6657	0.0488	0.7336	0.3140	0.0943	0.1720	0.7023	0.0225	0.9399	-0.1495	0.4420	-0.0694	0.9460	-0.0015	0.9991	0.0679	0.8696
---	---	1630833_at	0.1160	0.6439	-0.1903	0.3859	-0.3737	0.0555	-0.0983	0.9029	0.1808	0.5129	0.2790	0.2327	0.1584	0.8386	-0.0652	0.8886	-0.2236	0.5003
SA-2	Stromalin-2	1630834_at	0.0827	0.6595	0.0764	0.5774	-0.1955	0.3470	-0.0718	0.8640	0.0409	0.8277	0.1127	0.4180	0.1279	0.7956	0.0215	0.9494	-0.1064	0.6342
CG8079	CG8079	1630835_a_at	0.7825	0.0745	1.7870	0.0575	0.4599	0.3368	-0.2064	0.7149	0.6014	0.0274	0.8079	0.0046	0.9932	0.6903	1.6333	0.1173	0.6401	0.5578
CG32973	CG32973	1630836_at	0.1274	0.4259	0.0714	0.7968	-0.0318	0.8445	0.0500	0.9462	0.2899	0.1630	0.2399	0.1982	0.1494	0.7768	0.1072	0.6710	-0.0422	0.8926
---	---	1630837_at	0.1655	0.2758	-0.2139	0.1940	-0.0383	0.8345	0.2188	0.5008	0.4317	0.0207	0.2129	0.1666	-0.0151	0.9862	-0.1515	0.4619	-0.1363	0.5245
CG33958	CG33958	1630838_at	-0.0374	0.9142	0.0179	0.8651	0.3403	0.0873	0.1758	0.7556	0.0748	0.8023	-0.1010	0.6930	-0.0855	0.9266	0.1047	0.7824	0.1902	0.5587
CG14804	CG14804	1630839_at	-0.3542	0.1470	-0.1289	0.3821	-0.2187	0.1336	0.0302	0.9507	0.1350	0.3272	0.1048	0.4086	0.1093	0.9246	0.4329	0.2474	0.3236	0.4114
Nfl	Nuclear factor l	1630840_at	-0.7317	0.0066	-0.4302	0.2296	-0.6155	0.0072	-0.1145	0.8921	-0.1478	0.6415	-0.0333	0.9235	0.0731	0.9101	0.0749	0.7971	0.0019	0.9965
CG8668	CG8668	1630841_at	0.8298	0.0232	1.2216	0.0818	1.2584	0.0044	-0.2096	0.5680	-0.5930	0.0076	-0.3834	0.0303	-0.2119	0.8909	-0.1339	0.8720	0.0779	0.9230
CG32640 /// CG32641	CG32641 /// CG3	1630842_s_at	-0.0601	0.7654	0.0071	0.9791	-0.2030	0.3048	-0.0574	0.9447	-0.1276	0.6282	-0.0702	0.7911	0.1292	0.8375	-0.0331	0.9363	-0.1623	0.5523
CG40159	CG40159	1630843_a_at	0.0970	0.6470	0.1560	0.2494	0.3830	0.0291	0.0463	0.9353	-0.1443	0.4056	-0.1907	0.2063	-0.1642	0.7464	-0.0428	0.8929	0.1214	0.6005
CG15293	CG15293	1630844_at	2.6141	0.0009	1.0773	0.1382	2.2907	0.0009	0.8288	0.4140	0.4636	0.4260	-0.3652	0.4999	-0.3642	0.8222	-1.0587	0.1211	-0.6944	0.3149
Sod	Cu-Zn superoxide	1630845_at	-0.0531	0.7968	0.4268	0.1630	0.5711	0.0076	0.0088	0.9922	-0.4264	0.0304	-0.4352	0.0176	-0.0933	0.9333	-0.1843	0.6322	-0.0910	0.8418
CG3020	CG3020	1630846_at	0.1773	0.4520	0.2338	0.3791	0.2271	0.1842	0.1728	0.7289	0.0551	0.8484	-0.1176	0.5991	0.1386	0.8424	0.1332	0.6747	-0.0054	0.9914
CG15930	CG15930	1630847_at	0.3555	0.2008	0.1356	0.4621	0.2545	0.2804	0.0838	0.9228	-0.0205	0.9576	-0.1043	0.7038	0.0640	0.9589	-0.1550	0.7022	-0.2190	0.5649
obst-l	CG32304	1630848_at	-0.0945	0.6356	0.0876	0.5902	0.2105	0.3127	0.1979	0.6822	0.0035	0.9913	-0.1944	0.3458	0.0930	0.9087	0.1751	0.5597	0.0821	0.8178
---	---	1630849_at	0.1909	0.3271	0.3377	0.1442	0.1168	0.4325	-0.2772	0.4596	-0.2558	0.2063	0.0213	0.9319	-0.0733	0.9092	-0.1715	0.4505	-0.0982	0.6978
CG30469	CG30469	1630850_at	-0.0642	0.7631	-0.1739	0.2442	0.0501	0.7669	0.2625	0.4861	0.1641	0.4323	-0.0984	0.6332	-0.0325	0.9725	0.0429	0.9062	0.0754	0.7915
CG12000	beta-type 4 subun	1630851_s_at	0.2754	0.1112	0.4376	0.0763	0.7733	0.0006	0.1510	0.7118	-0.0098	0.9733	-0.1607	0.3449	-0.1322	0.8122	0.0701	0.8151	0.2023	0.3898
FR	larval-opioid-recep	1630852_at	-0.0228	0.9329	-0.1907	0.3354	-0.1834	0.2541	0.0606	0.9380	0.2759	0.2274	0.2153	0.2995	0.0387	0.9710	-0.0508	0.9009	-0.0895	0.7774
Gr10a	Gustatory recepto	1630853_at	0.2359	0.2729	0.2328	0.3458	0.2844	0.0733	0.0760	0.8873	-0.0759	0.7189	-0.1519	0.3623	-0.0282	0.9816	-0.1826	0.5183	-0.1544	0.5988
CG32193	CG32193	1630854_at	0.0348	0.8738	-0.0734	0.7504	0.1292	0.5883	-0.0002	0.9999	-0.0827	0.7581	-0.0825	0.7317	-0.0718	0.9589	-0.0878	0.8764	-0.0160	0.9809
Ste12DOR /// Ste:CG3323	Stellate orphon ///	1630855_s_at	-0.2237	0.6980	-2.2938	0.0067	-1.7798	0.0188	0.4033	0.8074	2.1386	0.0066	1.7353	0.0102	-0.2595	0.8202	-0.3054	0.5296	-0.0459	0.9435
CG8677	CG8677	1630856_at	0.0159	0.9633	-0.1374	0.5311	-0.1623	0.6099	-0.0316	0.9858	0.0992	0.8522	0.1308	0.7728	-0.0730	0.9309	-0.0460	0.9129	0.0271	0.9434
NTPase	CD39-like NTPase	1630857_s_at	0.1952	0.2963	-0.1921	0.2532	0.0873	0.5984	-0.0625	0.9118	0.2096	0.2436	0.2720	0.0900	-0.2230	0.7464	-0.1820	0.5501	0.0410	0.9188
CG7816	CG7816	1630858_s_at	-0.6393	0.0041	-1.3488	0.0081	-0.7481	0.0237	0.3691	0.4356	0.3017	0.2464	-0.0675	0.8187	-0.1852	0.8313	-0.3207	0.3685	-0.1355	0.7461
Chro	Chromobox protei	1630859_s_at	1.2735	0.0101	1.5238	0.0218	1.4440	0.0017	-0.0987	0.9381	-0.8203	0.0353	-0.7216	0.0377	-0.0872	0.9589	-0.5873	0.2123	-0.5002	0.3141
scrt	scratch	1630860_at	-0.5130	0.2303	-0.2525	0.2495	-0.4511	0.0360	-0.4131	0.6908	-0.5944	0.2147	-0.1812	0.7235	-0.0719	0.9352	-0.2605	0.3434	-0.1886	0.5239
CG8314	CG8314	1630861_at	-0.9835	0.0057	-0.4250	0.0407	-0.3218	0.1378	0.1213	0.7982	-0.7632	0.0027	-0.8845	0.0009	-0.1127	0.8823	-0.0574	0.8942	0.0553	0.8900
CG18432	CG18432	1630862_at	0.0982	0.5843	0.1615	0.4019	0.2565	0.2962	0.0804	0.9040	0.0963	0.7003	0.0159	0.9542	-0.0116	0.9943	-0.0152	0.9836	-0.0036	0.9956
---	---	1630863_s_at	0.2946	0.0717	0.0953	0.5845	0.3232	0.1109	0.1594	0.6338	0.0666	0.7287	-0.0928	0.5686	-0.0897	0.8882	-0.1118	0.6750	-0.0221	0.9474
CG40209	CG40209	1630864_at	0.0751	0.7023	0.0204	0.8437	0.1391	0.3586	0.0374	0.9479	0.0551	0.7768	0.0177	0.9281	-0.0930	0.8589	-0.0558	0.8460	0.0372	0.8984
CG11337	CG11337	1630865_s_at	0.3339	0.0968	1.1407	0.0295	0.6509	0.1422	0.4062	0.4599	0.5775	0.0571	0.1712	0.5612	0.9065	0.3800	1.4276	0.0366	0.5211	0.3743
CG7502	CG7502	1630866_at	0.1708	0.6887	0.4306	0.1970	0.3419	0.0715	0.0552	0.9376	-0.4402	0.0387	-0.4954	0.0151	0.1459	0.9409	-0.1354	0.8764	-0.2814	0.6724
CG8119	CG8119	1630867_at	-0.0652	0.7805	0.2213	0.1567	0.2740	0.1204	0.0640	0.9165	-0.1839	0.3410	-0.2479	0.1429	0.1575	0.7953	0.1952	0.4457	0.0377	0.9158
---	---	1630868_s_at	0.0931	0.6778	0.0958	0.4733	0.0495	0.8128	0.0806	0.9149	0.0813	0.7753	0.0007	0.9982	0.0657	0.9357	0.1296	0.6449	0.0639	0.8474
---	---	1630869_at	0.2684	0.1746	0.2880	0.3079	0.0418	0.8328	-0.2334	0.5756	-0.0600	0.8225	0.1735	0.3810	-0.0407	0.9657	-0.0150	0.9719	0.0256	0.9431
sip2	septin interacting	1630870_s_at	0.1289	0.7178	-0.8434	0.2270	-1.3898	0.0109	-0.6192	0.2037	0.8701	0.0098	1.4893	0.0006	0.0675	0.9862	0.1005	0.9457	0.0330	0.9835
---	---	1630871_at	0.1110	0.4616	0.0095	0.9420	-0.1284	0.4814	0.1000	0.8074	0.1258	0.4593	0.0258	0.8955	0.1178	0.8461	0.0284	0.9438	-0.0894	0.7554
Gr36d	Gr36d	1630872_at	0.2696	0.2081	0.1397	0.4492	-0.0422	0.8067	0.0028	0.9962	0.1232	0.5528	0.1204	0.5198	-0.0231	0.9816	-0.0599	0.8360	-0.0368	0.9023
---	---	1630873_s_at	0.0791	0.6301	-0.1044	0.4347	0.2250	0.3038	0.3813	0.2500	0.3762	0.0596	-0.0051	0.9840	-0.1012	0.9064	-0.1126	0.7604	-0.0114	0.9832
Nckx30C	Nckx30C	1630874_s_at	0.4526	0.1089	0.2658	0.2833	0.6309	0.0265	0.1691	0.7293	-0.1922	0.3906	-0.3614	0.0673	-0.0291	0.9816	-0.1753	0.5465	-0.1462	0.6261
CG14417	CG14417	1630875_at	0.1664	0.4553	0.0172	0.8919	-0.0922	0.6420	0.0304	0.9610	0.1709	0.3086	0.1405	0.3580	0.0996	0.9112	0.0681	0.8810	-0.0315	0.9433
CG17025	CG17025	1630876_at	0.0192	0.9302	-0.1862	0.2011	-0.0323	0.8497	0.0770	0.8791	0.2079	0.2275	0.1309	0.4177	-0.1427	0.8122	-0.1385	0.6024	0.0042	0.9924
CG8455	CG8455	1630877_a_at	-0.6541	0.0048	-0.6963	0.0236	-0.5975	0.0157	-0.0808	0.8879	0.3207	0.0970	0.4015	0.0278	-0.3257	0.7116	0.0811	0.8779	0.4068	0.2841
CG1750	CG1750	1630878_a_at	0.3311	0.3266	0.4996	0.1133	0.0878	0.6316	-0.1627	0.7081	0.3061	0.1211	0.4688	0.0163	0.3326	0.7324	0.4574	0.2522	0.1248	0.8069
---	---	1630879_at	0.0060	0.9812	0.1910	0.2723	0.3574	0.0326	-0.0728	0.8955	-0.1422	0.4569	-0.0695	0.7228	0.0767	0.9055	0.1430	0.5543	0.0663	0.8179
CG12467 /// DmirCG12467	CG12467	1630880_at	0.3501	0.2237	-0.3204	0.3697	-0.1416	0.5028	-0.0131	0.9937	0.5795	0.0731	0.5925	0.0444	-0.1187	0.8461	-0.1407	0.5994	-0.0220	0.9505
CG1309	CG1309	1630881_at	-0.2666	0.2648	-0.1929	0.5776	0.0220	0.9118	-0.0538	0.9314	-0.1114	0.5815	-0.0577	0.7808	-0.3798	0.7324	-0.1422	0.8185	0.2376	0.6423
XNP	XNP	1630882_s_at	-0.2163	0.5532	-0.2554	0.0717	-0.4175	0.0238	0.1948	0.6558	0.5061	0.0242	0.3113	0.0972	0.2507	0.8235	0.3383	0.4785	0.0876	0.8918
CG33978	CG33978	1630883_at	-0.6544	0.0190	-0.8286	0.0894	-2.0559	0.0001	-0.3127	0.5863	0.3956	0.1748	0.7083	0.0149	0.1960	0.7893	-0.0983	0.8068	-0.2943	0.3610
Rad21	Rad21	1630884_at	-0.1417	0.3769	0.0067	0.9845	0.0739	0.6935	-0.1331	0.7351	0.0001	0.9997	0.1332	0.4158	-0.1635	0.8202	0.2329	0.4280	0.3963	0.2018
CG12534	CG12534	1630885_at	-0.2722	0.1574	-0.0348	0.8599	0.3365	0.0453	0.2394	0.6377	-0.1084	0.7061	-0.3479	0.1174	-0.1725	0.7633	-0.0193	0.9617	0.1532	0.5457
SelR	methionine-R-sulfo	1630886_at	-0.4542	0.0190	-0.3756	0.0563	-0.7897	0.0009	-0.1161	0.7507	0.0070	0.9775	0.1232	0.4051	0.1223	0.8331	0.1111	0.6803	-0.0112	0.9769
CG14998	CG14998	1630887_a_at	-0.8814	0.0421	-0.8475	0.2591	-0.9377	0.0012	-0.3363	0.2185	-0.1348	0.4275	0.2016	0.1685	-0.1461	0.9589	-0.1543	0.8971	-0.0082	0.9953
---	---	1630888_at	0.0582	0.7632	0.1756	0.4705	-0.0875	0.5776	-0.1900	0.6615	-0.1905	0.3701	-0.0005	0.9987	0.0349	0.9626	0.0076	0.9848	-0.0274	0.9231
Bro /// RhoGAP1A	RhoGAP1A /// Brc	1630889_at	0.0957	0.4976	0.0752	0.7087	0.2826	0.1292	0.0752	0.8921	0.0264	0.9197	-0.0488	0.8165	0.0108	0.9922	0.0690	0.8376	0.0582	0.8596
nullo	nullo	1630890_at	0.0030	0.9906	0.0198	0.8447	0.0581	0.7952	-0.1974	0.5251	-0.1088	0.5309	0.0886	0.5855	0.0027	0.9980	0.0716	0.8153	0.0689	0.8146
CG31010	CG31010	1630891_at	0.0435	0.8694	0.0902	0.6914	0.1135	0.5086	0.1101	0.8362	0.0603	0.8080	-0.0498	0.8282	0.0420	0.9616	0.0153	0.9698	-0.0267	0.9372
CG11208 /// DsecCG11208	CG11208	1630892_at	0.6020	0.0083	0.6455	0.0657	1.0951	0.0001	0.2383	0.3749	0.0580	0.7478	-0.1803	0.1838	-0.1149	0.8999	0.1791	0.6191	0.2940	0.3915
CG7639	CG7639	1630893_at	-1.0277	0.0104	-1.2690	0.0745	-1.0852	0.0005	-0.1953	0.4690	-0.0853	0.5899	0.1100	0.4187	-0.1528	0.9317	-0.1880	0.7963	-0.0352	0.9678

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1630894_s_at	0.0243	0.9275	0.0620	0.7801	0.1217	0.6023	-0.0204	0.9860	-0.0471	0.9011	-0.0267	0.9379	-0.0891	0.8541	-0.0045	0.9925	0.0846	0.7067
dpr11	dpr11	1630895_at	0.0071	0.9758	-0.2266	0.3626	-0.1383	0.3514	0.0047	0.9956	0.1871	0.3622	0.1824	0.3198	-0.0393	0.9710	0.0624	0.8721	0.1017	0.7454
CG2267	CG2267	1630896_at	0.0840	0.7117	-0.0160	0.8993	0.0647	0.7954	0.1311	0.8293	0.1773	0.4667	0.0462	0.8654	0.0226	0.9841	0.0022	0.9978	-0.0203	0.9562
GluClalpha	indefinite	1630897_at	0.0573	0.6911	0.0049	0.9796	0.0961	0.6338	0.0251	0.9730	0.0650	0.7688	0.0399	0.8521	-0.1348	0.7979	-0.1264	0.5880	0.0084	0.9821
---	---	1630898_at	-0.1465	0.3048	0.0735	0.5499	-0.0799	0.6772	-0.2313	0.4529	-0.1275	0.4656	0.1038	0.5234	-0.0314	0.9687	0.0859	0.7200	0.1173	0.6024
Fer1	48 related 1	1630899_at	0.0445	0.8066	0.0124	0.9187	0.0226	0.9084	0.1294	0.8155	0.1107	0.6513	-0.0187	0.9461	0.1337	0.8270	0.0755	0.8218	-0.0582	0.8635
Nhe1	Nhe1	1630900_at	-0.7542	0.0055	-0.7273	0.0758	-0.4633	0.0344	-0.1152	0.8244	0.0103	0.9735	0.1256	0.5199	-0.3233	0.7215	0.0771	0.8910	0.4003	0.3079
CG6422	CG6422	1630901_a_at	0.0822	0.6838	0.6248	0.1726	0.2390	0.1483	-0.1389	0.7302	-0.1074	0.5829	0.0315	0.8822	0.0509	0.9698	0.3693	0.2679	0.3184	0.3695
---	---	1630902_at	-0.1712	0.3537	-0.0581	0.6119	-0.0403	0.8551	-0.0120	0.9909	-0.1552	0.5241	-0.1432	0.5192	0.0820	0.8541	0.0656	0.7693	-0.0164	0.9492
---	---	1630903_s_at	0.1546	0.2826	0.0550	0.8075	-0.0119	0.9521	0.0737	0.8776	0.1693	0.3068	0.0956	0.5535	0.1565	0.7251	0.1029	0.6187	-0.0536	0.8247
---	---	1630904_x_at	0.1232	0.4528	-0.0391	0.7097	-0.1267	0.5295	0.0395	0.9580	0.2376	0.2306	0.1981	0.2685	0.0311	0.9701	-0.1724	0.4103	-0.2035	0.3503
---	---	1630905_at	0.0352	0.8474	-0.1369	0.3039	0.0127	0.9583	0.1173	0.7575	0.1412	0.3996	0.0239	0.9034	-0.1634	0.8076	-0.0506	0.9062	0.1127	0.7228
CG14441	CG14441	1630906_at	0.3408	0.3480	0.4801	0.1034	0.8552	0.0006	-0.0647	0.8707	-0.0694	0.6586	-0.0047	0.9785	-0.1348	0.8885	0.1464	0.7269	0.2812	0.4533
CG32847	CG32847	1630907_at	0.0843	0.6875	0.0216	0.8896	0.2985	0.0498	0.1111	0.8546	-0.1569	0.4955	-0.2680	0.1703	-0.1345	0.7979	-0.1679	0.4446	-0.0334	0.9121
CG13617 /// DbuzCG13617	CG13617	1630908_at	0.0349	0.8281	0.1180	0.2512	0.1412	0.4742	0.0616	0.9011	-0.0449	0.8239	-0.1066	0.4892	0.0204	0.9816	0.0786	0.7514	0.0582	0.8247
---	---	1630909_at	0.1437	0.5724	0.1964	0.4390	0.3196	0.1059	0.0569	0.9302	-0.1721	0.3767	-0.2290	0.1778	-0.1248	0.8889	-0.1282	0.7462	-0.0033	0.9954
---	---	1630910_at	-0.0967	0.6606	0.0409	0.6940	-0.0370	0.8566	-0.1353	0.6869	-0.0820	0.6389	0.0533	0.7544	-0.0978	0.8973	0.0189	0.9678	0.1167	0.7148
eIF5	anon-fast-evolving	1630911_s_at	-0.1841	0.2639	0.4518	0.1648	0.1347	0.4167	-0.0949	0.8028	-0.0087	0.9705	0.0862	0.5682	0.2331	0.7677	0.6478	0.0696	0.4147	0.2299
CG7789	CG7789	1630912_at	-0.2187	0.4069	0.2092	0.4441	0.3795	0.0246	-0.1079	0.8076	-0.4614	0.0171	-0.3536	0.0324	-0.2089	0.8114	-0.0314	0.9587	0.1775	0.6486
---	---	1630913_at	0.7042	0.0389	1.2794	0.1094	0.9875	0.0483	-0.1571	0.7233	-0.1797	0.3798	-0.0226	0.9262	0.0896	0.9816	0.4738	0.6106	0.3842	0.6892
neuroligin	neuroligin	1630914_s_at	-0.0659	0.7600	0.0450	0.8057	-0.2370	0.1717	-0.0685	0.9039	-0.0038	0.9892	0.0647	0.7461	0.2167	0.7136	0.1663	0.5141	-0.0504	0.8803
---	---	1630915_at	-0.0716	0.8138	-0.0493	0.6444	0.1051	0.4556	0.1476	0.7850	-0.0634	0.8196	-0.2110	0.2962	-0.1540	0.7893	-0.2497	0.2946	-0.0956	0.7338
Mkk4	JNK kinase 2	1630916_at	0.0353	0.9132	0.1260	0.4567	-0.2170	0.1813	0.0544	0.9314	0.3602	0.0603	0.3058	0.0721	0.4083	0.6749	0.3299	0.4021	-0.0784	0.8862
CG7889	CG7889	1630917_at	0.5982	0.0067	0.4650	0.0179	0.7059	0.0130	0.2044	0.5061	0.1770	0.2797	-0.0273	0.8880	-0.1946	0.7768	0.1042	0.7731	0.2987	0.3235
a6	a6	1630918_at	0.0115	0.9856	-0.1146	0.9103	-0.1695	0.4654	0.1073	0.8294	0.3247	0.0901	0.2174	0.2024	0.1474	0.9677	0.1539	0.9175	0.0065	0.9974
pip	pipe	1630919_at	-0.0793	0.9083	-0.1866	0.5879	-0.3759	0.1372	-0.0084	0.9956	0.0528	0.9223	0.0612	0.8953	0.1261	0.9589	-0.0735	0.9487	-0.1996	0.8184
Tsp42Ek	tetraspanin 42E	1630920_at	-0.1296	0.4464	-0.0456	0.6708	-0.2756	0.3030	-0.0934	0.8721	-0.0336	0.9044	0.0599	0.7911	-0.0564	0.9589	0.0123	0.9836	0.0687	0.8668
bbg	CG9587	1630921_at	0.7927	0.0261	0.6916	0.1290	0.8216	0.0227	0.0889	0.9089	-0.0340	0.9220	-0.1229	0.6260	-0.1280	0.9516	0.0241	0.9842	0.1521	0.8540
CG11538	CG11538	1630922_at	0.3158	0.6068	-0.7906	0.0695	-0.0991	0.7405	1.2630	0.0082	1.4261	0.0004	0.1630	0.4548	0.1723	0.9410	-0.0944	0.9350	-0.2667	0.7492
CG6945	CG6945	1630923_at	-0.2402	0.2319	-0.4113	0.1796	-0.9065	0.0021	-0.1549	0.6958	0.2097	0.2525	0.3646	0.0330	0.2530	0.7485	-0.0622	0.9036	-0.3153	0.3616
---	---	1630924_at	0.2277	0.5218	0.2060	0.2760	0.0956	0.6453	0.0544	0.9251	0.0377	0.8696	-0.0167	0.9385	0.2126	0.8049	-0.0641	0.9075	-0.2767	0.4569
Rpt1	AAA ATPase	1630925_at	0.0664	0.6299	0.2081	0.1274	0.3901	0.0381	-0.0123	0.9857	-0.0653	0.7175	-0.0531	0.7548	-0.1315	0.8215	0.1581	0.5285	0.2896	0.2497
CG13925	CG13925	1630926_at	0.1157	0.5491	0.0087	0.9359	0.2470	0.1472	0.0516	0.9433	-0.0302	0.9164	-0.0819	0.7050	-0.0392	0.9515	0.0390	0.8903	0.0782	0.7221
Pxd	Peroxidase	1630927_x_at	-0.5258	0.0453	-1.1215	0.0161	-0.9146	0.1024	0.2438	0.8817	0.7768	0.1628	0.5330	0.2931	-0.0640	0.9499	0.0110	0.9851	0.0750	0.8471
---	---	1630928_at	0.1278	0.5538	0.0156	0.8819	-0.0575	0.7748	0.0285	0.9814	0.0800	0.8291	0.0516	0.8815	0.0147	0.9898	-0.0358	0.9247	-0.0505	0.8772
CG5885 /// DyakCG5885	CG5885	1630929_at	1.0168	0.0051	1.3797	0.0145	1.4110	0.0001	-0.0428	0.9514	0.3118	0.1060	0.3546	0.0459	-0.1099	0.8991	0.6116	0.0714	0.7215	0.0607
fus	fusilli	1630930_a_at	0.5430	0.2738	0.4574	0.3741	0.3145	0.2309	-0.0850	0.9018	0.0083	0.9804	0.0933	0.6914	0.0986	0.9808	-0.1462	0.9156	-0.2449	0.8256
CG2843	CG2843	1630931_at	-0.2557	0.3150	0.2597	0.1132	0.5032	0.0753	0.1466	0.8380	0.1216	0.6939	-0.0249	0.9404	-0.0480	0.9794	0.5619	0.1720	0.6099	0.1752
CG7488	CG7488	1630932_at	-0.4464	0.0521	-0.2308	0.2995	-0.3114	0.0449	-0.2001	0.4770	-0.3951	0.0171	-0.1950	0.1476	-0.2241	0.7644	-0.1273	0.7272	0.0968	0.8006
CG6738	CG6738	1630933_at	-0.1953	0.8918	-0.1296	0.4281	-0.2130	0.3313	-0.2430	0.9380	-0.3388	0.7580	-0.0959	0.9328	0.0670	0.9816	-0.3588	0.6080	-0.4259	0.5369
---	---	1630934_at	0.8324	0.0376	-1.1668	0.1810	-1.3831	0.1375	0.1200	0.9300	1.3075	0.0065	1.1875	0.0060	0.3745	0.9420	-0.5620	0.7707	-0.9365	0.5788
CG2093	CG2093	1630935_at	-0.6008	0.0191	0.2066	0.4829	0.2494	0.1632	0.1184	0.8259	-0.2975	0.1507	-0.4158	0.0326	0.1684	0.8564	0.6384	0.1013	0.4700	0.2330
lola	longitudinals abse	1630936_at	-0.9471	0.0127	-0.9670	0.0871	-0.9606	0.0045	0.3212	0.6034	0.5083	0.1021	0.1871	0.5385	0.0429	0.9848	0.2639	0.6333	0.2210	0.6992
---	---	1630937_at	0.2185	0.3549	0.0661	0.5797	-0.3197	0.0956	-0.2425	0.5863	-0.1446	0.5543	0.0980	0.6803	0.0631	0.9400	-0.0359	0.9302	-0.0990	0.7399
CG14879	CG14879	1630938_a_at	-0.2659	0.3803	0.0081	0.9933	-0.7920	0.0872	0.0367	0.9761	1.2057	0.0028	1.1690	0.0019	0.9380	0.6955	1.4367	0.1460	0.4987	0.6463
---	---	1630939_s_at	0.2894	0.1027	0.0378	0.8120	-0.0721	0.7788	-0.0217	0.9758	0.0955	0.6243	0.1172	0.4838	0.0958	0.9320	-0.1805	0.6464	-0.2762	0.4613
Anp	andropin	1630940_at	0.1124	0.6197	0.0459	0.6641	-0.0852	0.6600	-0.1981	0.6086	-0.0418	0.8715	0.1564	0.3861	-0.0190	0.9898	-0.0681	0.8807	-0.0491	0.9107
sesB	lethal(1)9Ed	1630941_s_at	-0.4057	0.0412	0.0197	0.8835	-0.2420	0.1743	-0.1645	0.6189	-0.3934	0.0249	-0.2290	0.1188	0.0325	0.9762	-0.0776	0.8206	-0.1101	0.7087
CG5742	CG5742	1630942_at	0.0993	0.8102	-0.4140	0.0314	0.3146	0.3765	0.5319	0.5336	0.9384	0.0419	0.4065	0.3178	-0.1041	0.9467	0.4259	0.3620	0.5300	0.2804
sec3	sec3	1630943_at	-0.2486	0.1829	-0.1320	0.6431	0.0705	0.6742	0.0998	0.8189	0.0274	0.9077	-0.0724	0.6852	-0.1666	0.7644	0.1291	0.6096	0.2957	0.2263
CG7536	CG7536	1630944_s_at	0.7420	0.0057	0.4911	0.0844	0.0581	0.8718	-0.1228	0.8470	0.1682	0.4916	0.2911	0.1616	0.2119	0.8564	-0.0655	0.9321	-0.2774	0.5870
mnb	Minibrain	1630945_at	0.2669	0.2427	-0.1497	0.3862	-0.2961	0.1087	-0.4025	0.1921	0.2108	0.2606	0.6134	0.0037	-0.1627	0.7768	-0.0537	0.8794	0.1090	0.6914
---	---	1630946_at	-0.6719	0.0055	-0.5552	0.0586	-0.8573	0.0013	-0.0855	0.8162	-0.0803	0.6143	0.0051	0.9776	0.1554	0.8202	0.0212	0.9639	-0.1342	0.6607
---	---	1630947_s_at	0.0607	0.7131	0.0200	0.8671	0.1970	0.3098	0.1289	0.8103	0.1175	0.6189	-0.0114	0.9664	-0.1342	0.8049	0.0133	0.9723	0.1475	0.5391
---	---	1630948_s_at	0.4002	0.0806	-0.1148	0.7156	-0.0823	0.8135	0.1858	0.8791	1.2137	0.0106	1.0279	0.0135	0.1115	0.9457	0.2556	0.6322	0.1441	0.8153
Rheb	Rheb	1630949_s_at	0.0924	0.5441	0.4054	0.1365	0.3010	0.2239	0.0238	0.9745	-0.7513	0.0024	-0.7751	0.0013	0.0977	0.9514	-0.3524	0.4640	-0.4501	0.3578
nop5	nop5	1630950_at	1.3531	0.0233	0.6080	0.1071	0.2992	0.0608	0.3847	0.7191	1.5547	0.0070	1.1700	0.0150	0.5543	0.4415	0.8190	0.0512	0.2647	0.4962
CG18275	CG18275	1630951_at	0.1690	0.4997	0.1057	0.4524	0.3956	0.0494	-0.0266	0.9744	0.0502	0.8482	0.0768	0.7212	-0.2005	0.7506	0.1056	0.7400	0.3061	0.2757
CG11586	CG11586	1630952_at	-0.2883	0.1679	-0.1518	0.3180	-0.5320	0.0511	-0.0597	0.9346	0.1798	0.4175	0.2395	0.2124	-0.0083	0.9914	0.0603	0.7883	0.0685	0.7412
CG13298 /// DyakCG1329	CG13298	1630953_at	0.4533	0.0321	0.3471	0.1636	0.0703	0.7722	0.0117	0.9883	0.3207	0.0927	0.3090	0.0718	0.3771	0.6762	0.2394	0.5421	-0.1376	0.7513
---	---	1630954_at	0.0990	0.5387	0.1500	0.2734	0.2313	0.3136	-0.0992	0.8085	-0.0743	0.6913	0.0249	0.8991	-0.0726	0.9342	0.0355	0.9378	0.1081	0.7399
CG9870	CG9870	1630955_at	0.1900	0.5367	-0.2971	0.4941	-0.0106	0.9806	-0.1214	0.8340	0.2425	0.2719	0.3640	0.0671	-0.4634	0.7485	-0.3787	0.5544	0.0847	0.9198
CG34143	CG34143	1630956_at	0.0020	0.9965	-1.1899	0.0752	-0.9083	0.0470	-0.0595	0.9688	1.0698	0.0146	1.1293	0.0070	-0.2674	0.8425	-0.1574	0.8299	0.1100	0.8846
slif	slimfast	1630957_s_at	-0.1115	0.6939	0.5391	0.0409	0.6525	0.0060	0.2642	0.5774	-0.1070	0.7025	-0.3712	0.0878	0.1737	0.8471	0.5350	0.1432	0.3612	0.3475
Chmp1	Chmp1	1630958_at	-0.2448	0.3527	-0.3550	0.0650	-0.0124	0.9757	0.2222	0.5808	0.0487	0.8555	-0.1735	0.3584	-0.2032	0.8680	0.0507	0.9487	0.2539	0.6260
CG34118	CG34118	1630959_at	0.2708	0.0944	0.0668	0.6588	0.1915	0.3126	0.0049	0.9956	0.0704	0.7757	0.0655	0.7722	0.0086	0.9928	0.0058	0.9911	-0.0028	0.9946
CG6559	CG6559	1630960_at	-0.0199	0.9170	0.1000	0.5463	0.0990	0.5977	-0.1061	0.8049	-0.0672	0.7396	0.0388	0.8443	-0.0029	0.9990	0.1683	0.6474	0.1713	0.6389
CG17193	CG17193	1630961_at	0.2546	0.1133	0.0504	0.6535	0.3821	0.0265	0.1172	0.7604	0.2678	0.1040	0.1506	0.3165	-0.1276	0.8298	-0.0337	0.9326	0.0939	0.7439
---	---	1630962_at	0.1436	0.5291	0.0794	0.4087	0.0789	0.7330	0.0846	0.9108	-0.0084	0.9814	-0.0930	0.7138	0.1512	0.8270	0.0506	0.9089	-0.1006	0.7663

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Pbprp4	Pheromone-bindir	1630963_at	0.1551	0.5650	0.1140	0.4755	0.0792	0.6269	-0.1055	0.8671	-0.0844	0.7498	0.0211	0.9393	0.0377	0.9677	0.0287	0.9417	-0.0090	0.9834
CG15211	CG15211	1630964_at	-0.9817	0.0061	-0.4074	0.1835	-0.7568	0.0026	-0.0322	0.9558	0.0337	0.8701	0.0659	0.6864	0.4700	0.6557	0.6963	0.1124	0.2262	0.6270
CG32141	CG32141	1630965_at	-0.0624	0.7066	-0.0763	0.6025	-0.0089	0.9729	0.0855	0.8830	0.0566	0.8187	-0.0289	0.9034	-0.0879	0.8846	-0.0847	0.7567	0.0032	0.9937
---	---	1630966_at	0.0577	0.7611	0.0459	0.6575	0.1591	0.4526	0.2026	0.5978	-0.0100	0.9735	-0.2126	0.2207	-0.0768	0.9168	-0.0806	0.8045	-0.0038	0.9932
CG13385	CG13385	1630967_at	0.1882	0.2746	-0.0867	0.4909	0.0759	0.7894	-0.0609	0.9518	0.1015	0.7601	0.1625	0.5512	-0.0947	0.8997	-0.1280	0.6762	-0.0333	0.9316
CG13907	CG13907	1630968_at	-0.2964	0.1613	-0.2616	0.4560	-1.1654	0.0002	-0.6297	0.2330	-0.4543	0.1458	0.1754	0.5736	0.3487	0.3857	-0.2933	0.1742	-0.6420	0.0366
CG12206	CG12206	1630969_s_at	0.0825	0.6138	0.0635	0.6684	0.4933	0.0125	0.0623	0.8869	-0.0255	0.9004	-0.0878	0.5455	-0.1803	0.7100	-0.0648	0.8025	0.1154	0.5965
CG14973	CG14973	1630970_at	0.4041	0.0875	0.6146	0.0783	0.8526	0.0167	-0.0078	0.9956	-0.0534	0.8865	-0.0456	0.8907	-0.1766	0.9238	-0.0682	0.9451	0.1084	0.8969
Rbp9	female sterile(2)Bi	1630971_s_at	0.0361	0.8508	0.0603	0.8258	-0.1824	0.3285	-0.0578	0.9567	-0.0953	0.7841	-0.0375	0.9144	-0.0047	0.9964	-0.0259	0.9487	-0.0212	0.9514
CG4270	CG4270	1630972_at	0.0659	0.6762	-0.1934	0.3014	0.0806	0.7772	0.1628	0.7764	0.2780	0.2465	0.1153	0.6390	-0.0702	0.9246	0.0346	0.9306	0.1048	0.7087
IP3K2	Inositol 1,4,5-triph	1630973_at	0.2487	0.0894	0.2731	0.1298	0.5615	0.0055	0.1442	0.7451	-0.1295	0.5335	-0.2737	0.1145	0.1020	0.8795	0.1383	0.6166	0.0363	0.9189
---	---	1630974_at	0.0316	0.8711	-0.0617	0.5971	0.2677	0.1277	0.0895	0.8085	0.1637	0.2680	0.0741	0.6199	-0.2373	0.7633	0.1135	0.7787	0.3508	0.2984
CG2909 /// DsimCG2909	CG2909	1630975_at	0.0292	0.9278	-0.6252	0.0930	-0.1855	0.7467	0.2487	0.6558	1.2236	0.0015	0.9749	0.0024	-0.1509	0.9589	0.6175	0.4551	0.7684	0.3636
vir	virilizer	1630976_at	0.1465	0.7871	-0.3552	0.4793	-0.1116	0.6344	0.3736	0.3739	0.7151	0.0092	0.3414	0.1065	0.0996	0.9761	0.1804	0.8749	0.0808	0.9431
CG16771	CG16771	1630977_at	-0.3618	0.1074	0.1251	0.4567	0.6641	0.0554	-0.2957	0.5311	-0.7795	0.0074	-0.4838	0.0349	-0.7343	0.3362	-0.1488	0.7596	0.5855	0.1744
---	---	1630978_s_at	-0.0177	0.9547	-0.4874	0.0150	-0.6805	0.0126	0.0553	0.9578	0.5708	0.0450	0.5155	0.0434	0.0815	0.9340	-0.1373	0.6969	-0.2188	0.5001
CG13144	CG13144	1630979_at	-0.1170	0.5416	0.2623	0.2200	0.1566	0.5655	-0.0408	0.9558	-0.1160	0.5929	-0.0752	0.7234	-0.0890	0.9457	0.1126	0.8304	0.2015	0.6371
CG15464	CG15464	1630980_at	-0.1309	0.3853	0.1011	0.6748	0.1565	0.2743	-0.0083	0.9922	-0.1367	0.4448	-0.1284	0.4267	-0.0528	0.9589	0.1238	0.7090	0.1766	0.5695
CG2698	CG2698	1630981_at	-0.2899	0.1488	-0.2471	0.1312	-0.1565	0.2599	-0.0581	0.9029	-0.1426	0.3669	-0.0845	0.5846	-0.0722	0.9296	-0.1311	0.6458	-0.0590	0.8654
CG9890 /// DereCG9890	CG9890	1630982_at	0.2172	0.3755	-0.0043	0.9844	0.6585	0.0114	0.1924	0.5138	0.0478	0.8026	-0.1446	0.3064	-0.1655	0.8608	0.0387	0.9499	0.2041	0.6171
CG32736	CG32736	1630983_s_at	-0.2462	0.4230	-0.5366	0.1523	-0.8181	0.0013	-0.1626	0.7293	0.0769	0.7606	0.2395	0.2021	0.1042	0.9416	-0.1676	0.7455	-0.2718	0.5601
ATPsyn-beta	H+-ATPase beta s	1630984_at	-0.2562	0.2507	0.2248	0.3690	0.0553	0.8468	-0.2133	0.7015	-0.6855	0.0153	-0.4723	0.0444	-0.1654	0.8192	-0.2273	0.4421	-0.0619	0.8779
CG30441	CG30441	1630985_at	-1.2847	0.0430	-0.7316	0.1966	-1.1002	0.0216	0.0241	0.9904	-0.1249	0.8213	-0.1490	0.7581	0.0212	0.9912	-0.0352	0.9556	-0.0564	0.9162
Adk3	Adenylate kinase-	1630986_s_at	0.8012	0.0254	0.3835	0.2286	1.6371	0.0000	0.4227	0.2463	-0.3200	0.1369	-0.7426	0.0030	-0.8996	0.3517	-0.7379	0.1420	0.1617	0.7935
BG4	BG4	1630987_at	-0.2129	0.2654	-0.1758	0.4801	-0.3205	0.0821	0.0161	0.9838	-0.1338	0.4788	-0.1499	0.3686	0.0673	0.9589	-0.0612	0.9151	-0.1285	0.7652
CG31347	CG31347	1630988_at	0.0986	0.5179	-0.2646	0.3908	-0.1926	0.3250	-0.0566	0.9116	0.1666	0.3068	0.2232	0.1205	-0.0620	0.9701	-0.0288	0.9670	0.0332	0.9566
CG32442	CG32442	1630989_a_at	0.2488	0.4706	0.3622	0.0779	0.4709	0.0069	0.0829	0.8817	-0.3933	0.0449	-0.4762	0.0130	-0.0492	0.9764	-0.2671	0.5199	-0.2179	0.6144
bs	pruned	1630990_at	0.0208	0.9535	-0.1669	0.7366	-0.2329	0.4866	0.0105	0.9931	0.3583	0.1270	0.3478	0.0983	0.0939	0.9742	0.1269	0.9075	0.0330	0.9769
CG11007	CG11007	1630991_at	-1.4580	0.0044	0.6678	0.0314	0.3181	0.1422	-0.5737	0.1560	-2.3165	0.0001	-1.7429	0.0002	-0.0113	0.9952	0.0910	0.8890	0.1024	0.8619
CG7946 /// DsimCG7946	CG7946	1630992_at	-0.1579	0.5143	-0.0962	0.7899	0.3345	0.1241	-0.0329	0.9558	-0.1677	0.3014	-0.1349	0.3623	-0.4493	0.6994	-0.1074	0.8783	0.3419	0.5003
CG31414	CG31414	1630993_at	1.8622	0.0031	1.9814	0.0775	2.5167	0.0000	0.1226	0.8589	-0.4474	0.0741	-0.5700	0.0183	-0.1559	0.9689	-0.3509	0.7900	-0.1950	0.8925
---	---	1630994_x_at	0.1075	0.5345	-0.2531	0.1129	-0.0452	0.8234	0.1803	0.5633	0.4196	0.0170	0.2393	0.0946	-0.0561	0.9457	0.0030	0.9959	0.0591	0.8555
chif	chiffon	1630995_at	-0.1098	0.7861	0.0328	0.8079	-0.4625	0.0236	-0.2237	0.5336	0.2803	0.1341	0.5040	0.0093	0.3840	0.7644	0.3732	0.5057	-0.0108	0.9907
CG17723	CG17723	1630996_s_at	-1.1700	0.0096	-0.9552	0.0291	-0.7659	0.0078	0.1070	0.9297	0.1488	0.7185	0.0418	0.9235	-0.0610	0.9499	0.3407	0.2217	0.4017	0.1897
CG7429	CG7429	1630997_at	0.1027	0.6982	0.3012	0.2508	0.4393	0.0246	0.0647	0.9112	-0.0964	0.6399	-0.1611	0.3383	-0.0758	0.9411	0.2081	0.5266	0.2839	0.3820
---	---	1630998_at	0.2382	0.1899	0.0950	0.6732	0.1090	0.6956	-0.0681	0.8935	0.1400	0.4238	0.2081	0.1681	-0.1079	0.8823	-0.0476	0.9111	0.0603	0.8717
CG7724	CG7724	1630999_at	-2.1093	0.0015	-2.5988	0.0014	-2.1287	0.0001	0.3683	0.5633	0.0441	0.9228	-0.3242	0.2730	-0.1086	0.9342	-0.4748	0.2430	-0.3663	0.3921
CG9906	Calnexin	1631000_at	0.0058	0.9804	-0.0463	0.7193	-0.0254	0.9259	0.0627	0.8931	0.0941	0.5756	0.0315	0.8619	-0.0975	0.9291	-0.0362	0.9499	0.0613	0.9004
CG6393	CG6393	1631001_s_at	-0.0845	0.7513	-0.1590	0.4139	-0.1992	0.3609	-0.0727	0.9314	0.0188	0.9600	0.0915	0.7323	-0.1413	0.8455	-0.1195	0.7293	0.0218	0.9597
---	---	1631002_at	0.2242	0.3709	-0.0054	0.9902	0.1734	0.3190	0.1574	0.8265	0.1195	0.7083	-0.0379	0.9100	-0.0241	0.9848	-0.2275	0.4242	-0.2034	0.4903
CG31857	CG31857	1631003_at	0.2059	0.4471	-0.0275	0.8769	0.0889	0.6667	-0.0254	0.9722	0.0749	0.7240	0.1003	0.5806	-0.0396	0.9760	-0.0919	0.8256	-0.0523	0.9056
su(Hw)	suppressor of hair	1631004_s_at	0.2155	0.3107	0.7871	0.0611	0.2510	0.2377	-0.2654	0.6107	0.1360	0.6426	0.4014	0.0862	0.2636	0.7634	0.7042	0.0714	0.4405	0.2482
CG13038	CG13038	1631005_at	0.4290	0.0528	0.1184	0.2926	0.0822	0.7366	0.0315	0.9620	0.1290	0.4834	0.0974	0.5779	0.1371	0.8395	0.0594	0.8824	-0.0778	0.8240
CG10171	CG10171	1631006_a_at	-0.0934	0.5344	-0.0580	0.7291	0.1992	0.2355	-0.4407	0.1109	-0.5668	0.0050	-0.1261	0.4109	-0.4605	0.3166	-0.4333	0.0871	0.0272	0.9366
RnrL	ribonucleoside-dip	1631007_at	-0.1464	0.8346	-0.7005	0.5404	-0.8801	0.2651	-0.3484	0.5498	0.8038	0.0157	1.1522	0.0019	-0.3782	0.9467	0.1208	0.9658	0.4990	0.8166
---	---	1631008_at	0.0507	0.7680	0.1056	0.4037	0.2788	0.1905	0.1311	0.7865	0.1386	0.5116	0.0075	0.9761	0.0214	0.9862	0.0673	0.8662	0.0460	0.9067
Mes-4	Mes-4	1631009_at	0.6268	0.0417	0.2347	0.4407	0.2086	0.3166	-0.3441	0.2915	0.5139	0.0154	0.8580	0.0010	-0.2401	0.8331	0.2616	0.6068	0.5016	0.3027
---	---	1631010_at	-0.0382	0.9277	-0.2391	0.2126	-0.0524	0.7764	0.0956	0.8473	0.2181	0.2300	0.1225	0.4800	-0.0286	0.9816	-0.0944	0.7667	-0.0658	0.8444
CG11498	CG11498	1631011_at	0.0556	0.8378	0.0109	0.9194	0.1182	0.5469	0.0970	0.9228	-0.0991	0.7874	-0.1961	0.4987	-0.0175	0.9913	-0.0074	0.9925	0.0101	0.9849
cn	cinnabar	1631012_at	0.0567	0.7613	-0.0129	0.9007	0.2404	0.1193	0.0442	0.9436	-0.0364	0.8786	-0.0806	0.6630	-0.2154	0.7498	-0.1606	0.6012	0.0547	0.8904
mbc	Myoblast City	1631013_at	0.4195	0.0738	0.4807	0.1502	0.1933	0.6246	-0.1000	0.8908	-0.2207	0.3784	-0.1208	0.6293	0.1299	0.9457	-0.2285	0.7340	-0.3585	0.5581
---	---	1631014_at	0.2670	0.1994	0.2620	0.2407	0.3748	0.0491	-0.1673	0.6908	-0.2673	0.1672	-0.0999	0.6117	-0.1321	0.8571	-0.0856	0.8290	0.0465	0.9104
RpL5	ribosomal protein	1631015_s_at	0.0838	0.5506	0.0183	0.8546	0.0554	0.7252	-0.0203	0.9704	-0.0231	0.9045	-0.0028	0.9874	-0.0246	0.9717	-0.0200	0.9440	0.0046	0.9868
CG32822	CG32822	1631016_at	0.1563	0.3098	-0.1012	0.3408	0.0131	0.9507	0.0601	0.8915	0.4119	0.0131	0.3517	0.0160	-0.1827	0.7464	0.0492	0.8858	0.2319	0.3424
Cpr65Au	CG18778	1631017_at	0.8645	0.6326	0.1159	0.3851	-0.9943	0.0107	-1.2653	0.1857	-1.3526	0.0272	-0.0873	0.9002	-0.0769	0.9928	-1.7867	0.3706	-1.7099	0.4078
312	312	1631018_at	0.0024	0.9932	0.0079	0.9422	-0.0284	0.8999	-0.0156	0.9863	-0.0702	0.7945	-0.0546	0.8276	-0.1637	0.7839	-0.2156	0.3972	-0.0519	0.8831
CG5971	CG5971	1631019_at	0.1367	0.5248	0.0263	0.8292	0.1153	0.7738	-0.0671	0.9603	0.4884	0.1715	0.5556	0.0844	-0.0383	0.9859	0.2966	0.5591	0.3350	0.5045
CG34113	CG34113	1631020_at	-0.0425	0.8561	-0.1251	0.4924	-0.0728	0.7908	0.2728	0.5969	0.2033	0.4539	-0.0695	0.8132	0.0236	0.9816	0.0129	0.9748	-0.0107	0.9775
CG4771	CG4771	1631021_at	0.0633	0.8957	-0.6224	0.1776	-0.8815	0.0621	-0.4825	0.1028	0.8620	0.0011	1.3445	0.0001	-0.2458	0.9277	0.0151	0.9941	0.2609	0.8124
CG6209	CG6209	1631022_at	0.0776	0.6541	-0.0158	0.8964	0.0434	0.8119	0.0282	0.9648	0.1049	0.5656	0.0767	0.6619	0.0798	0.8541	0.0287	0.9175	-0.0512	0.8178
---	---	1631023_at	0.1560	0.4641	0.0062	0.9573	0.1203	0.5545	0.2041	0.7149	0.3367	0.1844	0.1326	0.6070	-0.1297	0.8016	-0.1089	0.6414	0.0208	0.9445
CG14617	CG14617	1631024_s_at	-0.2338	0.3788	-0.3328	0.0783	-0.5980	0.0161	-0.1298	0.8485	0.2904	0.2431	0.4203	0.0621	-0.0870	0.9088	0.0561	0.8850	0.1431	0.6202
Prosbeta5R1	Prosbeta5R1	1631025_at	0.1036	0.5802	0.1921	0.2662	0.1422	0.3448	0.0368	0.9412	0.0240	0.9026	-0.0128	0.9425	0.0154	0.9895	-0.0482	0.8952	-0.0635	0.8380
CG40346	CG40346	1631026_at	0.1767	0.3013	0.0196	0.8645	0.0922	0.6409	-0.0800	0.9068	0.1214	0.6191	0.2014	0.3159	-0.0199	0.9860	0.0281	0.9455	0.0480	0.8905
ara	iroquois	1631027_at	-0.1298	0.6849	-0.0527	0.6545	-0.0121	0.9533	0.1193	0.8605	-0.1439	0.5871	-0.2631	0.2276	-0.0117	0.9928	-0.2487	0.4103	-0.2370	0.4523
l(2)k09913	lethal (2) k09913	1631028_s_at	-0.0079	0.9784	-0.0134	0.9722	0.0333	0.8437	-0.0648	0.9148	-0.3188	0.0929	-0.2540	0.1317	-0.2280	0.8122	-0.3522	0.3666	-0.1243	0.7945
CG13776	CG13776	1631029_at	-0.3675	0.2540	-0.4966	0.1792	-0.3979	0.0466	-0.0305	0.9683	0.0598	0.8094	0.0904	0.6640	-0.1260	0.8680	-0.0775	0.8500	0.0485	0.9067
Dbp80	Helicase 80	1631030_at	-0.1326	0.5538	-0.1242	0.7165	0.0116	0.9595	-0.2769	0.3443	-0.1860	0.2724	0.0909	0.5900	-0.2438	0.7823	-0.0414	0.9467	0.2024	0.6228
---	---	1631031_at	0.1780	0.4338	0.0917	0.5525	0.1233	0.4150	-0.0175	0.9857	0.2388	0.2938	0.2563	0.2029	-0.0673	0.9142	-0.0173	0.9620	0.0500	0.8619

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14659	CG14659	1631032_at	0.0199	0.9236	0.0141	0.8940	0.0532	0.7521	0.1133	0.7753	0.0984	0.5862	-0.0149	0.9425	-0.0054	0.9952	-0.0589	0.8476	-0.0534	0.8548
---	---	1631033_s_at	0.2228	0.2398	0.1356	0.5196	-0.0256	0.9065	0.0111	0.9909	0.1363	0.5443	0.1252	0.5399	0.1994	0.7485	0.0326	0.9402	-0.1668	0.5511
Rca1	regulator of cyclin	1631034_at	0.2745	0.1425	0.0733	0.8741	-0.5746	0.0598	-0.1343	0.7225	0.3209	0.0634	0.4552	0.0096	0.4184	0.7233	0.0987	0.8983	-0.3196	0.5571
Gr36a	Gustatory recepto	1631035_at	-0.0223	0.9017	-0.1075	0.4895	0.1486	0.3456	0.1363	0.6755	0.0721	0.6797	-0.0642	0.6898	-0.3355	0.6724	0.0116	0.9852	0.3471	0.2923
CG11816	CG11816	1631036_at	0.0291	0.8800	-0.0975	0.6435	0.0276	0.8989	0.0592	0.9251	0.0256	0.9227	-0.0336	0.8806	-0.0435	0.9545	-0.0565	0.8546	-0.0131	0.9696
BthD	BthD selenoprotei	1631037_at	0.3753	0.2430	-0.2538	0.2873	0.1169	0.8407	0.0334	0.9759	0.1625	0.5825	0.1291	0.6448	-0.2803	0.8689	-0.3941	0.5800	-0.1138	0.9025
CG32773	CG32773	1631038_at	0.3996	0.1001	0.4202	0.0986	0.4993	0.0603	-0.0527	0.9346	-0.0594	0.7988	-0.0067	0.9776	0.0047	0.9976	0.0415	0.9402	0.0369	0.9380
CG33108	CG33108	1631039_at	-0.0804	0.6606	-0.0065	0.9669	0.0757	0.6940	0.0990	0.8540	0.2449	0.2054	0.1460	0.4219	-0.0034	0.9984	0.1625	0.6483	0.1659	0.6389
CG10632	CG10632	1631040_at	-0.3681	0.0338	-0.5862	0.1402	-0.1773	0.4814	0.2179	0.4776	0.1405	0.4062	-0.0773	0.6485	-0.2638	0.8016	-0.1875	0.7050	0.0764	0.8993
---	---	1631041_at	-0.0257	0.9092	0.0681	0.7918	-0.0525	0.7592	-0.1093	0.8578	-0.0110	0.9743	0.0984	0.6642	0.0293	0.9816	0.0939	0.8062	0.0646	0.8717
---	---	1631042_at	0.1470	0.4795	0.0322	0.7695	-0.0505	0.7578	-0.0086	0.9931	0.0604	0.7998	0.0689	0.7420	0.0335	0.9701	0.0560	0.8605	0.0225	0.9435
---	---	1631043_x_at	0.1303	0.4494	-0.1954	0.3303	-0.1793	0.2047	0.0505	0.9271	0.4944	0.0094	0.4439	0.0092	-0.0308	0.9701	0.0913	0.7017	0.1221	0.5870
CG34144	CG34144	1631044_at	0.1202	0.4611	0.0850	0.5982	0.0125	0.9615	-0.1269	0.6908	-0.0053	0.9809	0.1216	0.3734	-0.0576	0.9545	-0.0465	0.9194	0.0111	0.9832
CG3571	CG3571	1631045_a_at	0.0150	0.9658	0.6914	0.1088	0.7045	0.0038	-0.1909	0.6107	-0.5236	0.0130	-0.3328	0.0522	-0.1483	0.9239	0.2334	0.6876	0.3818	0.4776
CG6448	CG6448	1631046_s_at	-0.2565	0.2524	0.0147	0.9022	-0.2134	0.1865	-0.2500	0.5019	-0.3851	0.0560	-0.1350	0.4726	0.0760	0.9092	-0.1235	0.6270	-0.1994	0.4055
---	---	1631047_at	0.0283	0.9045	-0.2355	0.1151	0.0944	0.6011	0.2509	0.4752	0.2975	0.1142	0.0466	0.8291	-0.1366	0.8293	-0.0585	0.8779	0.0781	0.8128
CG14434	CG14434	1631048_at	1.0778	0.0121	0.8572	0.0040	0.7918	0.0090	0.1470	0.6650	0.3390	0.0426	0.1920	0.1838	0.2054	0.8800	0.3335	0.5398	0.1281	0.8482
Arp14D	Actin-related prote	1631049_at	0.2125	0.2906	0.3120	0.1849	0.4158	0.2112	-0.0168	0.9865	0.3997	0.0964	0.4165	0.0566	-0.1332	0.9342	0.6218	0.2168	0.7550	0.1701
CG5384	CG5384	1631050_at	0.0119	0.9610	1.2221	0.0073	1.4659	0.0002	0.1383	0.7915	-0.3995	0.0641	-0.5378	0.0124	-0.1725	0.8270	0.6059	0.0795	0.7784	0.0559
CG9323	CG9323	1631051_at	0.4065	0.2137	0.4557	0.0172	0.2603	0.1442	-0.1876	0.6202	0.1186	0.5602	0.3062	0.0719	0.0597	0.9643	0.2164	0.5660	0.1568	0.6961
CG17959	CG17959	1631052_at	-0.1965	0.3832	-0.0049	0.9767	0.0307	0.8582	-0.1617	0.7451	-0.1871	0.4046	-0.0253	0.9247	-0.0499	0.9657	0.1150	0.7584	0.1649	0.6248
Sgs7	group III	1631053_at	-0.0418	0.9533	-0.7468	0.4072	-0.7974	0.1369	-0.3680	0.8546	0.7173	0.3305	1.0853	0.0946	-0.1330	0.9737	-0.2028	0.8884	-0.0698	0.9621
CG7943	CG7943	1631054_s_at	-0.1223	0.5954	0.4508	0.0165	0.6580	0.0050	-0.1785	0.5633	-0.7205	0.0015	-0.5420	0.0031	-0.4167	0.5343	-0.1420	0.6889	0.2747	0.3941
Pka-C1	protein kinase A	1631055_at	-0.9607	0.1214	-1.9103	0.0177	-2.4080	0.0094	0.2482	0.6677	0.9779	0.0044	0.7297	0.0099	0.7382	0.8344	0.0413	0.9900	-0.6969	0.6619
CG17742	CG17742	1631056_at	0.1264	0.4879	0.1937	0.4032	0.1119	0.5336	0.0207	0.9819	-0.0022	0.9945	-0.0229	0.9315	0.0849	0.9092	0.0074	0.9900	-0.0775	0.8124
---	---	1631057_at	0.1446	0.3151	0.0234	0.8651	0.0886	0.6115	-0.0175	0.9777	-0.1274	0.4446	-0.1098	0.4713	-0.1176	0.8424	-0.1149	0.6673	0.0027	0.9951
Mgstl	Microsomal glutatl	1631058_s_at	0.4149	0.0907	0.4175	0.0527	0.7952	0.0021	0.0361	0.9528	-0.3707	0.0344	-0.4068	0.0148	-0.2502	0.7893	-0.3912	0.3106	-0.1410	0.7578
Pkc98E	protein kinase C	1631059_at	-0.9465	0.0301	-1.0183	0.0144	-1.2599	0.0011	-0.1679	0.7857	-0.0478	0.8902	0.1201	0.6480	0.0972	0.9555	-0.1105	0.8789	-0.2077	0.7155
---	---	1631060_at	0.0854	0.7417	0.0425	0.8005	-0.0004	0.9982	-0.0385	0.9500	-0.0009	0.9971	0.0376	0.8489	0.0464	0.9657	0.0524	0.9068	0.0061	0.9902
---	---	1631061_at	0.1875	0.3271	0.1281	0.4189	-0.0692	0.6803	-0.1464	0.7743	-0.0143	0.9653	0.1321	0.5254	0.1236	0.8374	0.0949	0.7439	-0.0287	0.9342
CG2023	CG2023	1631062_at	0.5923	0.0699	1.0355	0.0321	1.0866	0.0017	0.2683	0.6338	0.2399	0.3983	-0.0284	0.9341	0.3303	0.7271	0.7228	0.0881	0.3926	0.3503
Or24a	Olfactory receptor	1631063_at	0.1148	0.5477	-0.0470	0.6781	0.0864	0.6007	-0.1003	0.8164	-0.0104	0.9687	0.0899	0.5978	-0.1301	0.8202	-0.0012	0.9991	0.1289	0.6150
---	---	1631064_at	-0.0373	0.8675	0.1600	0.1852	-0.0026	0.9935	-0.0430	0.9639	-0.0170	0.9650	0.0261	0.9333	0.1303	0.8157	0.1905	0.4046	0.0602	0.8372
CG15480	CG15480	1631065_at	0.1407	0.3621	0.2343	0.2877	0.0358	0.8668	-0.0751	0.9020	0.0068	0.9816	0.0819	0.6940	0.0584	0.9409	0.0655	0.8431	0.0071	0.9852
rib	ribbon	1631066_at	-0.6439	0.7692	-0.9320	0.0126	-0.4159	0.1215	0.2550	0.9714	-0.6390	0.7669	-0.8940	0.6250	-0.3901	0.9061	-1.0398	0.3720	-0.6497	0.6117
CG5539	CG5539	1631067_at	0.2532	0.2512	0.1278	0.4107	0.0572	0.7744	-0.0138	0.9857	0.0684	0.7422	0.0822	0.6517	0.0500	0.9607	0.0223	0.9621	-0.0277	0.9440
CG2861	CG2861	1631068_a_at	0.2148	0.1936	0.1785	0.3102	0.1394	0.3525	0.0289	0.9626	0.0262	0.9083	-0.0027	0.9897	0.0689	0.9238	0.0267	0.9451	-0.0422	0.8982
---	---	1631069_a_at	-0.0197	0.9629	-0.2986	0.4227	-0.0774	0.6562	-0.1380	0.7409	-0.0670	0.7606	0.0709	0.7165	-0.1443	0.8967	-0.0710	0.9111	0.0733	0.8982
---	---	1631070_at	0.1037	0.6141	0.2878	0.2024	0.4361	0.1020	0.1913	0.6584	-0.0207	0.9466	-0.2120	0.2573	0.2215	0.7953	0.1292	0.7667	-0.0923	0.8407
---	---	1631071_s_at	0.1415	0.4236	-0.2051	0.3866	-0.0922	0.7350	0.1889	0.7749	0.5198	0.0637	0.3309	0.1799	-0.1326	0.8439	-0.0854	0.8117	0.0471	0.9020
CG9512	CG9512	1631072_at	-2.6160	0.0031	-1.7408	0.3715	-1.1333	0.1299	-0.0851	0.9658	-2.0742	0.0028	-1.9890	0.0020	-0.7958	0.8795	-1.1189	0.6026	-0.3231	0.9077
Clic	Chloride intracellu	1631073_at	-0.5473	0.0440	0.0028	0.9950	-0.5196	0.0047	-0.1836	0.5664	0.1213	0.4795	0.3049	0.0437	0.3775	0.6749	0.6604	0.0871	0.2830	0.4618
CG8435	CG8435	1631074_at	-0.4013	0.0471	0.2156	0.2433	0.3273	0.0767	-0.1048	0.8174	-0.4000	0.0320	-0.2952	0.0670	-0.1111	0.8956	0.2607	0.4041	0.3718	0.2557
Glu-RIB	glutamate recepto	1631075_at	0.0627	0.8097	0.0162	0.8920	0.0548	0.7710	0.0436	0.9610	-0.0455	0.8880	-0.0891	0.7268	0.1163	0.8202	-0.0769	0.7603	-0.1933	0.3712
CG13771	CG13771	1631076_at	0.5885	0.0277	0.2584	0.3190	0.2577	0.1783	-0.0150	0.9863	0.0097	0.9762	0.0248	0.9235	0.0434	0.9752	-0.1097	0.8037	-0.1531	0.6877
CG16957	CG16957	1631077_at	-0.0624	0.8138	-0.1240	0.5767	0.1249	0.5437	0.0664	0.9311	0.0033	0.9916	-0.0630	0.8043	-0.1589	0.8235	-0.0720	0.8614	0.0869	0.8129
zpg	zero population gr	1631078_at	0.8124	0.2366	-1.2296	0.2552	-0.8587	0.3102	0.0969	0.9046	1.8795	0.0003	1.7826	0.0002	-0.4287	0.9486	-0.0274	0.9949	0.4013	0.8830
---	---	1631079_at	0.1586	0.2786	0.1071	0.3626	0.1418	0.3976	-0.1643	0.5680	-0.1201	0.4301	0.0442	0.7878	-0.1475	0.7230	-0.0500	0.8356	0.0976	0.6152
---	---	1631080_at	0.0872	0.5311	0.1345	0.1913	0.1807	0.2925	-0.0089	0.9896	-0.0157	0.9463	-0.0068	0.9723	-0.0286	0.9735	0.0542	0.8527	0.0829	0.7346
CG17666	CG17666	1631081_a_at	0.0340	0.9056	0.1877	0.3181	0.3761	0.0575	-0.0268	0.9727	-0.0908	0.6831	-0.0639	0.7653	-0.1103	0.8215	0.0627	0.8117	0.1730	0.4016
B52	Serine/arginine ric	1631082_at	0.1283	0.3954	-0.0430	0.8259	0.1883	0.3332	-0.1042	0.8336	-0.0296	0.9096	0.0746	0.7076	-0.1635	0.8042	-0.0957	0.7760	0.0679	0.8471
l(1)G0144	lethal (1) G0144	1631083_at	-0.1111	0.5294	-0.0335	0.8696	0.2030	0.2965	0.0862	0.8370	0.1451	0.3677	0.0589	0.7266	-0.1321	0.8235	0.1678	0.5100	0.2999	0.2397
CG12976	CG12976	1631084_at	0.3633	0.1282	0.3610	0.1966	0.4581	0.0340	-0.1747	0.7469	-0.3697	0.1168	-0.1950	0.3727	-0.1811	0.7215	-0.1367	0.5360	0.0444	0.8774
Myo28B1	myosin VII 28B1-I	1631085_a_at	0.4151	0.0451	0.3573	0.2748	1.0257	0.0037	0.2093	0.7028	0.0851	0.7806	-0.1242	0.6297	-0.2347	0.7215	-0.1540	0.6028	0.0807	0.8157
---	---	1631086_at	0.2137	0.1791	-0.0040	0.9730	0.1738	0.2179	0.2290	0.4643	0.2175	0.1970	-0.0115	0.9573	0.1096	0.8235	0.0936	0.6805	-0.0160	0.9564
CG9809	CG9809	1631087_a_at	1.3697	0.0309	0.6075	0.4341	0.3636	0.2295	0.1661	0.9108	1.4468	0.0083	1.2807	0.0086	0.3967	0.8534	0.6629	0.4550	0.2662	0.8050
CG9578	CG9578	1631088_at	-0.2211	0.4747	0.2637	0.4125	0.4490	0.0435	-0.0091	0.9909	-0.1851	0.2789	-0.1761	0.2486	-0.2379	0.8480	0.2958	0.5897	0.5336	0.3134
---	---	1631089_at	0.2708	0.1565	0.0178	0.8665	-0.1083	0.5740	-0.0501	0.9163	0.1410	0.3514	0.1912	0.1489	0.0034	0.9978	-0.0835	0.8105	-0.0869	0.7853
CG9795 /// DyakCG9795	CG9795	1631090_a_at	-0.1162	0.6127	-0.1017	0.6732	-0.3284	0.0782	-0.0550	0.9376	0.0687	0.7830	0.1237	0.5378	-0.0356	0.9725	0.0110	0.9821	0.0466	0.8951
CG10928	CG10928	1631091_at	0.1137	0.5260	0.0214	0.8330	-0.1095	0.4713	-0.0637	0.9017	0.0434	0.8405	0.1072	0.5063	0.0637	0.9416	-0.1657	0.5552	-0.2295	0.4016
CG8032	CG8032	1631092_at	-0.4406	0.0407	0.5290	0.0366	0.1665	0.3846	-0.1948	0.6050	-0.5328	0.0125	-0.3380	0.0509	0.0929	0.9031	0.4201	0.1321	0.3272	0.2557
CG11837	CG11837	1631093_at	0.1744	0.5674	-0.3603	0.4223	-0.3340	0.2603	0.3006	0.6122	0.9292	0.0078	0.6286	0.0261	0.1072	0.9514	0.3414	0.5246	0.2342	0.6817
CG9339	CG9339	1631094_s_at	-0.4258	0.1789	-0.4289	0.1217	-1.9270	0.0000	-0.8064	0.0405	0.1819	0.4188	0.9883	0.0009	0.4574	0.3800	0.3407	0.2195	-0.1166	0.7205
Psc	Posterior sex com	1631095_at	0.3134	0.3070	0.1418	0.7623	-0.1259	0.6499	-0.0014	0.9991	0.2074	0.4987	0.2087	0.4472	0.1759	0.9296	0.2042	0.8048	0.0283	0.9778
CG6340	CG6340	1631096_a_at	-0.0563	0.7507	0.3652	0.1016	-0.3386	0.1415	-0.2985	0.5311	-0.0353	0.9197	0.2632	0.2374	0.1920	0.7485	0.1560	0.5574	-0.0359	0.9183
CG15153	CG15153	1631097_at	0.0589	0.7919	0.1588	0.4063	0.0341	0.8414	-0.0090	0.9937	-0.0857	0.7508	-0.0767	0.7570	0.0123	0.9898	-0.0176	0.9587	-0.0299	0.9159
CG7194	CG7194	1631098_at	-0.8567	0.0750	-0.5224	0.3381	-0.6993	0.0164	-0.9777	0.1033	-1.0974	0.0082	-0.1197	0.7575	-0.7367	0.6557	-0.8114	0.2116	-0.0747	0.9366
CG17807	CG17807	1631099_at	-0.2377	0.2083	-0.1537	0.5965	0.0464	0.7840	0.1230	0.7680	-0.1803	0.3139	-0.3033	0.0595	-0.0455	0.9746	-0.1140	0.8028	-0.0685	0.8889
Lkr	Drosokinin recepti	1631100_at	2.0183	0.0017	1.7045	0.0648	1.5705	0.0003	-0.1506	0.8196	0.0505	0.8850	0.2011	0.4050	0.2996	0.8751	0.0118	0.9941	-0.2878	0.7391

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG6480	CG6480	1631101_at	-0.0002	0.9994	0.2941	0.0457	0.6044	0.0131	0.1426	0.7944	0.2025	0.3742	0.0599	0.8118	-0.0820	0.9495	0.5443	0.1489	0.6263	0.1334
acj6	Abnormal chemos	1631102_a_at	0.1201	0.4894	-0.0019	0.9907	-0.2352	0.1482	-0.1440	0.6493	-0.0820	0.6300	0.0619	0.7006	0.1045	0.8692	-0.0376	0.9257	-0.1421	0.6030
---	---	1631103_at	0.3308	0.1190	0.0790	0.6761	0.2085	0.2003	-0.0292	0.9592	0.0319	0.8725	0.0611	0.6988	0.0221	0.9885	-0.0478	0.9277	-0.0699	0.8774
Fsh	Fsh-Tsh-like recep	1631104_s_at	1.7757	0.2772	-0.4710	0.0964	0.3094	0.2002	0.3574	0.4420	-0.4154	0.1011	-0.7728	0.0052	-0.5053	0.9467	-2.5336	0.2558	-2.0283	0.3915
Tsp96F	Tetraspanin 96F	1631105_at	-1.9871	0.0014	-2.0215	0.0077	-2.9830	0.0000	-0.6948	0.0325	-0.0117	0.9662	0.6831	0.0019	-0.0896	0.9589	-0.1565	0.8000	-0.0669	0.9199
Aats-ile	Isoleucyl-tRNA sy	1631106_s_at	0.4577	0.0111	0.5605	0.2112	0.3828	0.0587	0.0965	0.7845	0.4552	0.0078	0.3587	0.0136	0.2436	0.8270	0.5234	0.2460	0.2798	0.5733
---	---	1631107_at	0.0455	0.8420	-0.1323	0.4166	0.0738	0.7928	0.3113	0.4420	0.1551	0.5062	-0.1563	0.4543	0.1808	0.7721	0.1812	0.5102	0.0004	0.9992
---	---	1631108_at	-0.3272	0.1517	-0.1107	0.4804	-0.1967	0.2517	0.1316	0.7941	-0.0419	0.8789	-0.1735	0.3594	0.0205	0.9816	0.0000	1.0000	-0.0205	0.9435
CG13550	CG13550	1631109_at	0.6751	0.0186	0.0255	0.8458	0.0522	0.7868	0.0119	0.9904	0.7320	0.0058	0.7201	0.0038	0.0163	0.9912	0.1053	0.7710	0.0890	0.8073
CG10257	CG10257	1631110_at	0.2843	0.3115	0.0260	0.8189	-0.0354	0.8830	-0.1479	0.7493	0.1296	0.5497	0.2776	0.1218	-0.0270	0.9816	-0.1899	0.5042	-0.1629	0.5789
CG6879	CG6879	1631111_at	-1.6536	0.0151	-2.9021	0.0014	-2.7445	0.0001	0.5400	0.6325	1.5977	0.0110	1.0577	0.0388	0.1915	0.7726	0.2783	0.3158	0.0868	0.8025
---	---	1631112_at	2.7930	0.0113	2.6913	0.0341	4.5758	0.0000	1.4074	0.0154	0.3467	0.2525	-1.0607	0.0027	-0.5309	0.8671	0.2387	0.9004	0.7696	0.5642
---	---	1631113_s_at	-0.1146	0.5085	0.2077	0.1531	0.0761	0.6758	-0.0587	0.9130	-0.0341	0.8801	0.0245	0.9042	0.0373	0.9683	0.2137	0.3789	0.1765	0.4908
CG11489	CG11489	1631114_a_at	-1.3837	0.0035	-1.4778	0.0149	-1.8921	0.0001	-0.0083	0.9956	0.5581	0.1359	0.5664	0.0917	0.2294	0.8650	-0.0005	0.9999	-0.2299	0.7027
Obp8a	Odorant-binding p	1631115_at	-0.0820	0.8816	2.0279	0.0071	1.0554	0.0282	-0.9042	0.3147	-2.6792	0.0009	-1.7750	0.0033	0.2836	0.8192	-0.2418	0.6662	-0.5254	0.3112
Clp	Clipper	1631116_at	0.3098	0.2708	0.0608	0.8184	0.4580	0.0553	-0.0157	0.9863	0.0180	0.9554	0.0337	0.8974	-0.2468	0.7644	-0.1165	0.7916	0.1303	0.7484
---	---	1631117_at	0.1509	0.3469	0.0056	0.9816	-0.0655	0.6611	0.1124	0.8512	0.1021	0.6814	-0.0102	0.9702	0.1923	0.6898	0.0266	0.9326	-0.1657	0.4067
MAPk-Ak2	MAP kinase activa	1631118_s_at	-0.0455	0.7489	0.0735	0.4472	-0.1764	0.4919	-0.2946	0.4669	-0.1127	0.6464	0.1819	0.3671	-0.0144	0.9914	0.0184	0.9709	0.0327	0.9382
Cip4	Cip4	1631119_at	-0.6396	0.0285	0.5888	0.3284	0.7251	0.0777	0.1346	0.8863	-0.1248	0.7399	-0.2594	0.3793	-0.0466	0.9898	1.1042	0.1504	1.1508	0.1682
mbl	mindmelt	1631120_at	-1.1100	0.0599	-1.6748	0.2084	-1.7944	0.0004	-0.0567	0.9610	0.3732	0.2278	0.4299	0.1196	0.1019	0.9852	-0.2352	0.9011	-0.3371	0.8306
CG7267	CG7267	1631121_at	-1.2297	0.0126	0.1403	0.4870	-0.0130	0.9751	-0.0705	0.9619	-1.3330	0.0052	-1.2625	0.0040	0.0498	0.9775	0.0936	0.8765	0.0438	0.9411
CG4656	CG4656	1631122_at	-0.5031	0.0324	0.0028	0.9947	-0.4603	0.0870	-0.1503	0.7831	-0.2289	0.3153	-0.0786	0.7481	0.2998	0.7324	0.3165	0.3943	0.0166	0.9777
Corin	DCorin	1631123_at	0.0196	0.9283	-0.1019	0.5985	-0.3576	0.0601	-0.1409	0.8291	0.1184	0.6789	0.2593	0.2489	0.0466	0.9672	-0.0696	0.8684	-0.1162	0.7345
---	---	1631124_at	0.1645	0.6187	0.0127	0.9645	0.2639	0.1251	0.1379	0.8578	-0.1090	0.7372	-0.2470	0.3264	-0.1235	0.9246	-0.0833	0.8999	0.0401	0.9479
CG15737	CG15737	1631125_at	0.5922	0.4753	-1.3809	0.1787	-2.1657	0.0023	-0.8314	0.0437	2.3355	0.0001	3.1670	0.0000	-0.0554	0.9939	0.1946	0.9425	0.2500	0.9145
gcm	glide glide2	1631126_at	-0.1351	0.8053	-0.1469	0.3580	-0.1300	0.4494	0.0462	0.9857	-0.2692	0.6868	-0.3153	0.5887	0.0679	0.9555	-0.2128	0.5671	-0.2807	0.4407
CG10424 /// DyakCG10424	CG10424	1631127_at	0.4598	0.0162	0.7962	0.0165	0.6356	0.0025	-0.0384	0.9518	-0.1817	0.3111	-0.1433	0.3842	0.0585	0.9411	0.0955	0.7403	0.0370	0.9125
CG9016	CG9016	1631128_s_at	0.2171	0.3616	-0.0661	0.6946	-0.2157	0.2743	0.1470	0.7432	0.2981	0.1296	0.1511	0.4128	0.1956	0.8042	-0.0913	0.8353	-0.2869	0.3921
CG31612	CG31612	1631129_at	0.0832	0.8510	0.1287	0.7634	-0.3201	0.0863	-0.3696	0.5008	0.2351	0.4366	0.6047	0.0281	0.1551	0.9238	0.3111	0.5847	0.1560	0.8158
CG5745	CG5745	1631130_at	-0.1688	0.5580	0.3603	0.0949	0.5210	0.0154	0.1178	0.8897	-0.2332	0.4284	-0.3510	0.1665	0.1479	0.8515	0.4863	0.1374	0.3384	0.3108
CG2990	CG2990	1631131_a_at	-0.0294	0.8839	0.0443	0.9006	-0.1495	0.6411	-0.3592	0.2753	0.0785	0.7299	0.4377	0.0198	-0.1107	0.9523	0.1217	0.8786	0.2324	0.7113
---	---	1631132_s_at	0.1561	0.5245	0.1879	0.3014	0.3154	0.2072	0.3399	0.5149	0.1832	0.5341	-0.1567	0.5663	0.1226	0.9407	0.2091	0.7200	0.0865	0.9025
CG2712	Coding Region III	1631133_at	0.1072	0.6666	0.0501	0.7513	-0.0568	0.7684	0.1600	0.7271	0.3955	0.0582	0.2355	0.1986	0.2545	0.7464	0.3223	0.3198	0.0678	0.8812
Gef26	Gef26	1631134_at	-0.0241	0.9548	0.4500	0.1537	0.1977	0.2094	-0.1300	0.8251	-0.3018	0.1828	-0.1718	0.4204	0.1429	0.8744	0.1782	0.6437	0.0353	0.9433
CG30286	CG30286	1631135_at	-0.0546	0.7848	0.0033	0.9813	0.1104	0.5388	0.1643	0.7327	0.1701	0.4447	0.0058	0.9831	0.1125	0.8810	0.1338	0.6743	0.0214	0.9592
---	---	1631136_at	0.2577	0.1412	0.0650	0.6437	0.0810	0.7703	-0.0417	0.9639	0.0112	0.9763	0.0529	0.8507	0.0370	0.9535	-0.0407	0.8804	-0.0778	0.7149
viaf	viral IAP-associate	1631137_at	-0.7400	0.0088	-0.1314	0.4151	-0.4409	0.4392	0.2003	0.5515	0.3804	0.0355	0.1802	0.2482	0.5119	0.7947	1.1971	0.1406	0.6852	0.4132
CG8191	CG8191	1631138_at	0.0543	0.8275	0.2106	0.5334	0.3603	0.0287	-0.1167	0.7688	-0.1025	0.5731	0.0143	0.9468	-0.1222	0.8243	0.0105	0.9799	0.1327	0.5871
fs(2)ltoPP43	female sterile (2) l	1631139_a_at	0.3097	0.3861	1.0616	0.0096	1.5482	0.0001	0.1545	0.7929	-0.5549	0.0283	-0.7094	0.0060	-0.2794	0.7644	0.1523	0.7454	0.4317	0.2931
Rab9Fb	CG32670	1631140_at	0.0670	0.8621	-0.2067	0.2724	0.0648	0.7988	-0.0298	0.9803	0.2053	0.5086	0.2351	0.3891	-0.0131	0.9928	0.0198	0.9734	0.0328	0.9464
SF1	Splicing factor 1	1631141_at	-0.7238	0.2380	-0.0680	0.9536	-0.2799	0.1276	-0.1469	0.6197	-0.4346	0.0101	-0.2877	0.0360	0.1356	0.9775	0.2514	0.8782	0.1159	0.9431
Caps	Calcium activated	1631142_a_at	-0.0680	0.9206	0.0969	0.8731	-0.4838	0.0909	-0.3200	0.5172	0.1697	0.5451	0.4897	0.0418	0.4932	0.8395	0.4565	0.6833	-0.0367	0.9835
CG32169	CG32169	1631143_at	0.0229	0.8979	0.0586	0.7746	-0.0917	0.6684	0.0483	0.9532	-0.0436	0.8860	-0.0919	0.6995	-0.0195	0.9838	-0.1153	0.6264	-0.0958	0.6952
---	---	1631144_at	-0.0839	0.6945	0.0058	0.9675	0.1383	0.5076	0.2612	0.4200	0.1452	0.4354	-0.1160	0.5017	0.1261	0.8814	0.1975	0.5595	0.0714	0.8676
CG32073	CG32073	1631145_at	0.2181	0.4640	0.1517	0.3664	0.4300	0.0577	-0.0529	0.9375	-0.1705	0.3969	-0.1176	0.5393	-0.0125	0.9928	-0.0519	0.9164	-0.0393	0.9295
alph	alphabet	1631146_at	0.2537	0.2045	-0.1489	0.3704	0.0410	0.8813	0.0520	0.9463	0.2927	0.1770	0.2407	0.2174	-0.1636	0.8427	-0.1628	0.6612	0.0007	0.9992
CG14692	CG14692	1631147_at	0.0643	0.7150	0.1707	0.4622	0.2509	0.1898	0.0257	0.9641	0.0231	0.9134	-0.0026	0.9891	0.0162	0.9884	0.1352	0.5966	0.1190	0.6443
CG9673	CG9673	1631148_at	-0.3932	0.9143	-0.7797	0.0070	-0.0716	0.8405	0.4961	0.9425	-1.5495	0.4461	-2.0455	0.2445	-0.1903	0.9893	-1.9822	0.5179	-1.7920	0.5677
La	ribonucleoprotein	1631149_s_at	0.2808	0.4472	0.0250	0.9524	0.0709	0.7372	0.2637	0.7345	0.8020	0.0279	0.5384	0.0826	0.1909	0.8449	0.5583	0.1512	0.3674	0.3717
Or85c	Odorant receptor	1631150_at	0.1636	0.3759	0.0725	0.7118	-0.0351	0.8309	0.0630	0.9228	0.2228	0.2569	0.1598	0.3775	0.1140	0.9038	0.1770	0.6321	0.0630	0.8932
CG4951	CG4951	1631151_at	-0.7906	0.0110	-0.3735	0.1578	-0.3419	0.0641	-0.0787	0.9029	-0.4264	0.0466	-0.3477	0.0658	-0.0571	0.9445	0.0289	0.9425	0.0860	0.7661
Trxr-2	thioredoxin reduct	1631152_at	0.2092	0.1783	0.1877	0.2859	0.3695	0.0843	0.1188	0.7424	0.1151	0.4894	-0.0036	0.9856	-0.0486	0.9449	0.1555	0.4727	0.2041	0.3518
osk	oskar	1631153_at	0.9525	0.6766	-4.6408	0.0949	-1.7522	0.0825	1.6168	0.1551	5.1557	0.0002	3.5389	0.0006	-1.0502	0.9238	-0.3263	0.9557	0.7239	0.8837
CG13321	CG13321	1631154_at	-0.4282	0.7642	0.0683	0.8988	0.5133	0.3788	0.2830	0.9345	-0.6283	0.5649	-0.9113	0.3236	0.0591	0.9928	0.1118	0.9646	0.0527	0.9838
Ppr-Y	Ppr-Y	1631155_at	-0.0100	0.9535	0.0101	0.9244	0.0536	0.7471	0.0201	0.9757	0.0216	0.9230	0.0016	0.9938	-0.0664	0.8999	0.0044	0.9925	0.0708	0.7516
---	---	1631156_at	0.2509	0.2613	0.0642	0.5890	0.1699	0.2870	-0.0432	0.9435	0.0560	0.7908	0.0992	0.5606	0.0252	0.9816	0.0631	0.8546	0.0379	0.9121
CG7365	CG7365	1631157_at	1.7636	0.3791	-0.3542	0.1277	-0.2498	0.1148	0.5657	0.4979	0.4249	0.3442	-0.1407	0.7729	0.1879	0.9862	-1.8228	0.4505	-2.0107	0.4114
Acp53C14a	Acp53C14a	1631158_at	0.3357	0.0543	0.2401	0.3503	0.2986	0.0647	0.0026	0.9974	-0.0242	0.9409	-0.0268	0.9232	-0.1724	0.7230	0.0041	0.9935	0.1765	0.4058
---	---	1631159_at	0.2067	0.2358	-0.0809	0.5052	0.1106	0.4469	-0.0377	0.9518	-0.0634	0.7613	-0.0257	0.9027	0.0607	0.9421	-0.1354	0.6264	-0.1961	0.4620
---	---	1631160_at	0.0283	0.9056	-0.0917	0.5648	0.0452	0.7865	-0.0029	0.9961	0.0784	0.7130	0.0813	0.6717	-0.1274	0.8740	-0.1591	0.6414	-0.0317	0.9431
pelo	pelota	1631161_at	0.3030	0.2607	0.5240	0.0949	0.3047	0.2690	-0.1260	0.8234	0.0912	0.7211	0.2172	0.2726	0.0981	0.9409	0.3343	0.4097	0.2362	0.5886
---	---	1631162_at	-0.0002	0.9994	0.0491	0.6940	0.0115	0.9563	0.0438	0.9592	-0.0112	0.9746	-0.0550	0.8305	0.2064	0.7733	0.0325	0.9491	-0.1738	0.6038
---	---	1631163_at	0.1705	0.3197	-0.2598	0.1296	-0.1007	0.5211	0.1654	0.6416	0.2293	0.1836	0.0639	0.7307	0.0027	0.9984	-0.1624	0.5880	-0.1651	0.5834
CG12370	CG12370	1631164_a_at	0.0321	0.9761	0.0168	0.9392	-0.2275	0.4456	-0.2181	0.7015	-0.4946	0.0620	-0.2765	0.2372	0.0202	0.9964	-0.7408	0.4655	-0.7611	0.4631
CG4688	CG4688	1631165_at	-0.1139	0.6084	-0.1844	0.1430	-0.2399	0.1582	-0.0704	0.9319	0.1925	0.4499	0.2629	0.2309	0.0230	0.9800	-0.0072	0.9858	-0.0302	0.9157
CG17440	CG17440	1631166_at	0.0045	0.9868	-0.0628	0.6611	0.1160	0.5279	0.2768	0.3665	0.1209	0.5114	-0.1559	0.3270	0.0005	0.9999	0.1022	0.7891	0.1018	0.7774
CG5023	CG5023	1631167_at	-2.1035	0.0373	-1.2543	0.1522	-2.2780	0.0025	-0.9074	0.2140	-1.8286	0.0022	-0.9213	0.0270	0.1586	0.9778	-1.0606	0.4354	-1.2192	0.3813
CG5687	CG5687	1631168_at	-3.1052	0.0025	-1.5767	0.2451	-3.5157	0.0002	-1.5000	0.0416	-2.2265	0.0007	-0.7265	0.0524	0.4867	0.9032	-0.6730	0.6779	-1.1597	0.4392
CG34115	CG34115	1631169_at	-0.1248	0.4383	-0.1037	0.6403	0.1922	0.1934	0.1684	0.6825	-0.1398	0.4871	-0.3082	0.0742	-0.2900	0.7070	-0.3763	0.2254	-0.0863	0.8310

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1631170_at	0.1619	0.3463	0.0237	0.8180	0.1349	0.4577	-0.0456	0.9311	-0.0146	0.9500	0.0311	0.8626	-0.0962	0.8461	-0.0584	0.8297	0.0378	0.8926
CheB93a	CheB93a	1631171_at	-0.0829	0.7108	-0.0118	0.9101	0.0837	0.6061	0.1717	0.6130	0.1077	0.5577	-0.0640	0.7251	-0.0011	0.9996	-0.0022	0.9979	-0.0011	0.9982
CG7848	CG7848	1631172_at	0.0308	0.8743	0.0644	0.6108	0.1948	0.3195	-0.0632	0.9037	-0.1685	0.3272	-0.1053	0.5232	-0.1716	0.7697	-0.0397	0.9178	0.1320	0.6225
dob	CG5560	1631173_at	2.2347	0.0246	0.9584	0.4876	1.7302	0.0021	0.5700	0.5121	0.6676	0.1441	0.0976	0.8549	-0.2603	0.9665	-0.7126	0.7020	-0.4523	0.8264
CG9875	CG9875	1631174_at	0.1968	0.3193	0.2439	0.3584	-0.0724	0.6659	-0.1591	0.7422	-0.0743	0.7717	0.0848	0.7064	0.1169	0.8298	0.0536	0.8678	-0.0633	0.8247
CG10283	CG10283	1631175_a_at	-1.7532	0.0032	-1.1808	0.0291	-1.7424	0.0007	-0.2202	0.7658	-0.4588	0.1438	-0.2386	0.4184	0.3450	0.7686	0.0096	0.9935	-0.3354	0.5198
---	---	1631176_at	0.0810	0.7976	-0.0738	0.6651	0.0602	0.7372	0.1023	0.9071	0.1642	0.5941	0.0618	0.8505	-0.0238	0.9814	-0.0327	0.9231	-0.0090	0.9803
raptor	dRaptor	1631177_at	0.1831	0.5241	0.2706	0.1135	0.4188	0.0594	0.1054	0.8822	-0.1017	0.7194	-0.2072	0.3539	-0.2002	0.7956	-0.0109	0.9870	0.1893	0.5860
---	---	1631178_at	0.2288	0.2500	-0.0171	0.9086	0.1713	0.4182	0.0464	0.9436	0.1226	0.5443	0.0762	0.7003	0.1002	0.8689	-0.0005	0.9998	-0.1007	0.7096
CG6680	CG6680	1631179_s_at	-0.2203	0.3366	-1.2551	0.0132	-1.2424	0.0002	-0.0559	0.9154	0.5980	0.0040	0.6538	0.0016	-0.1385	0.8814	-0.4142	0.2249	-0.2757	0.4498
CG5728	CG5728	1631180_at	0.2976	0.4892	-0.3926	0.4092	-0.0500	0.8546	0.3765	0.5933	0.9351	0.0172	0.5587	0.0822	-0.0092	0.9964	0.2560	0.6394	0.2652	0.6257
CG3294	CG3294	1631181_a_at	-0.0713	0.8639	-0.2476	0.1475	-0.2751	0.3474	-0.1749	0.8271	-0.0732	0.8550	0.1017	0.7637	-0.2989	0.7644	-0.2846	0.5118	0.0143	0.9840
CG2983	CG2983	1631182_at	0.1414	0.4144	0.1391	0.2034	-0.0191	0.9331	0.0088	0.9922	-0.0312	0.9053	-0.0400	0.8566	0.1705	0.7464	-0.0614	0.8380	-0.2319	0.3089
RhoL	Rho-like	1631183_at	-1.3786	0.0151	-0.2433	0.5089	-0.4919	0.2473	-0.0589	0.9290	-0.8323	0.0019	-0.7734	0.0016	0.1270	0.9672	0.4505	0.6045	0.3235	0.7284
---	---	1631184_at	0.2118	0.2389	0.1069	0.3945	0.3125	0.1153	0.1619	0.5931	0.1541	0.3221	-0.0078	0.9679	-0.0276	0.9783	-0.0664	0.8372	-0.0388	0.9068
---	---	1631185_at	0.0688	0.6884	0.1633	0.3828	0.0905	0.6270	-0.0504	0.9314	-0.0811	0.6821	-0.0307	0.8814	-0.0197	0.9871	-0.0928	0.7903	-0.0731	0.8362
---	---	1631186_at	0.0519	0.8192	0.1097	0.4172	0.1994	0.2302	-0.0547	0.9314	0.0107	0.9709	0.0654	0.7514	-0.0644	0.9199	0.0925	0.7059	0.1569	0.4851
---	---	1631187_at	-0.1207	0.4651	0.0841	0.5781	0.2989	0.0500	0.2092	0.5499	-0.1170	0.5479	-0.3262	0.0488	-0.0815	0.8655	0.0125	0.9686	0.0940	0.6488
CG9098	CG9098	1631188_a_at	-0.9762	0.0067	0.2206	0.3236	-0.3632	0.1125	-0.6865	0.2463	-1.7524	0.0010	-1.0658	0.0053	0.0958	0.9449	-0.4603	0.2522	-0.5562	0.2056
---	---	1631189_s_at	0.0081	0.9699	0.1802	0.1314	0.0654	0.6959	-0.0873	0.8265	0.0298	0.8855	0.1171	0.4151	0.0761	0.8903	0.2552	0.2042	0.1791	0.3967
CG8089	CG8089	1631190_at	0.0296	0.9478	0.2494	0.4639	-0.5015	0.0975	-0.2668	0.4553	0.4164	0.0382	0.6832	0.0027	0.6687	0.7070	0.8038	0.2605	0.1352	0.8945
CG5003	CG5003	1631191_at	0.0372	0.8464	0.1233	0.2788	-0.0656	0.7050	-0.0660	0.9132	0.0207	0.9399	0.0867	0.6614	0.1159	0.8494	0.1950	0.4421	0.0791	0.7964
CG10887 /// DereCG10887	CG10887	1631192_at	0.2698	0.2983	0.0351	0.8638	0.0898	0.6658	0.0197	0.9852	0.0402	0.9030	0.0205	0.9461	0.1118	0.9101	-0.0326	0.9541	-0.1443	0.7160
CG3014 /// rig	CG3014 /// rigor m	1631193_at	-0.3101	0.1720	-0.2785	0.4397	-0.1150	0.7178	0.1869	0.8485	-0.1192	0.7835	-0.3061	0.3509	0.2949	0.7220	0.1785	0.6414	-0.1163	0.7810
CG10265	CG10265	1631194_at	-0.7169	0.0286	-0.7700	0.0509	-0.9204	0.0011	-0.1626	0.8280	0.2955	0.3086	0.4581	0.0763	-0.1459	0.8331	0.1210	0.7117	0.2669	0.3680
CG30156	CG30156	1631195_at	-0.1175	0.6404	-0.1701	0.5598	-0.3917	0.0692	-0.4607	0.2714	-0.3011	0.2224	0.1596	0.5023	-0.1821	0.7338	-0.0785	0.7818	0.1036	0.6830
MED10	Mediator complex	1631196_at	0.0425	0.7941	-0.2424	0.4605	-0.5026	0.0074	0.0627	0.8837	0.3043	0.0435	0.2416	0.0675	0.3246	0.6749	0.1294	0.7221	-0.1952	0.5615
CG8534	CG8534	1631197_at	0.1871	0.3705	0.3225	0.1870	0.5872	0.0044	0.0716	0.9116	0.0559	0.8294	-0.0157	0.9519	-0.1252	0.8737	0.2266	0.4700	0.3518	0.2721
Rs1	Rs1	1631198_at	0.2587	0.3927	0.0519	0.9046	0.4356	0.0976	0.1933	0.7469	0.5187	0.0524	0.3254	0.1636	-0.1309	0.9405	0.2788	0.6344	0.4097	0.4654
CG9941	CG9941	1631199_at	-0.7158	0.0337	-0.0652	0.7143	0.0189	0.9687	0.1132	0.9064	-0.2324	0.4733	-0.3457	0.2124	-0.1049	0.9589	0.5021	0.3813	0.6070	0.3089
Acp32CD	Accessory gland p	1631200_at	0.1841	0.3255	0.2585	0.1026	0.0865	0.6362	-0.1887	0.6010	-0.0643	0.7678	0.1243	0.4719	-0.0034	0.9976	0.1152	0.6762	0.1185	0.6602
---	---	1631201_at	0.2320	0.2568	0.0041	0.9791	0.0241	0.9008	0.1291	0.7777	0.2484	0.1902	0.1194	0.5189	0.0466	0.9657	-0.0100	0.9865	-0.0565	0.8877
---	---	1631202_s_at	-0.1088	0.5924	0.1841	0.4741	0.1784	0.3561	-0.1376	0.7857	-0.0818	0.7396	0.0558	0.8127	-0.2247	0.8202	0.1530	0.7544	0.3777	0.3708
CG6362	CG6362	1631203_at	0.0477	0.8385	0.1488	0.3218	-0.0272	0.8790	-0.0145	0.9857	-0.0484	0.8341	-0.0339	0.8738	0.0846	0.9257	-0.0530	0.9085	-0.1376	0.6787
Peritrophin-15b	Dm-peritrophin-15	1631204_at	0.6648	0.3377	0.1219	0.4257	0.3602	0.0849	0.0489	0.9451	0.0491	0.8521	0.0002	0.9995	-0.0231	0.9950	-0.1629	0.8990	-0.1398	0.9056
DyakCG6084	CG6084	1631205_a_at	-0.1576	0.4892	1.2872	0.0101	0.5720	0.0237	0.1551	0.7647	-0.2795	0.2046	-0.4346	0.0349	0.9700	0.2884	1.2415	0.0298	0.2714	0.5866
CG17717	CG17717	1631206_a_at	-0.2375	0.2425	-0.0852	0.6844	0.0071	0.9712	0.0704	0.9116	0.1241	0.5666	0.0537	0.8124	-0.0176	0.9844	0.1298	0.5377	0.1474	0.4815
elB	elbow B	1631207_at	-0.1453	0.8335	0.5999	0.4195	0.7549	0.0228	0.3023	0.6010	-0.3645	0.2083	-0.6668	0.0185	0.1888	0.9611	0.5545	0.6345	0.3657	0.7744
CG15525	CG15525	1631208_at	0.1935	0.2937	-0.1826	0.5594	-0.0506	0.7820	0.1903	0.6962	0.5424	0.0239	0.3521	0.0810	0.2270	0.7230	0.2434	0.3716	0.0164	0.9694
CG30293	CG30293	1631209_at	0.2605	0.3770	0.0270	0.9054	0.3760	0.0370	0.1847	0.6120	0.0649	0.7666	-0.1198	0.4936	0.0687	0.9509	0.1034	0.8076	0.0348	0.9401
CG14540	CG14540	1631210_at	0.1198	0.6258	0.0750	0.6169	-0.1998	0.2366	-0.0827	0.8791	0.0044	0.9879	0.0872	0.6485	0.2197	0.7893	0.0517	0.9261	-0.1680	0.6580
CG13245	CG13245	1631211_at	0.0887	0.7329	-0.1479	0.4135	0.0960	0.5438	0.2810	0.4799	0.1804	0.4114	-0.1006	0.6469	0.0295	0.9816	-0.0473	0.9144	-0.0769	0.8299
MED22	Mediator complex	1631212_at	-0.2750	0.0707	0.3165	0.1391	0.2445	0.2343	-0.1527	0.7039	-0.4164	0.0316	-0.2637	0.1096	-0.0837	0.9147	0.0469	0.9111	0.1306	0.6604
CG9581	CG9581	1631213_at	-0.2436	0.4964	-0.1663	0.7891	-0.0420	0.8602	0.1111	0.7857	0.1854	0.2745	0.0743	0.6728	0.1671	0.9460	0.3791	0.6408	0.2120	0.8208
CG1675	CG1675	1631214_at	-0.2669	0.3113	-0.2449	0.5728	-0.0843	0.6631	-0.1180	0.7091	-0.3277	0.0304	-0.2097	0.1029	-0.3031	0.8235	-0.2100	0.7554	0.0932	0.9056
CG32132	CG32132	1631215_at	0.0042	0.9841	0.0299	0.8019	-0.0079	0.9680	-0.1384	0.7604	-0.2135	0.2772	-0.0751	0.7180	-0.0006	0.9998	-0.0284	0.9341	-0.0279	0.9246
CG18375	CG18375	1631216_a_at	-0.2287	0.8034	0.5533	0.3849	-0.2594	0.3933	-0.2513	0.8094	-0.5486	0.1832	-0.2973	0.4467	0.4837	0.8655	0.2435	0.8824	-0.2402	0.8770
TpnC73F	troponin C	1631217_a_at	-1.8209	0.0361	-1.4299	0.1110	-2.0818	0.0021	-0.1733	0.7915	-1.1918	0.0017	-1.0185	0.0019	0.3903	0.9168	-0.8879	0.5071	-1.2781	0.3360
Doc1	Dorsocross	1631218_at	0.2637	0.5633	-0.0267	0.8099	0.1106	0.5348	0.1032	0.8879	-0.0090	0.9809	-0.1122	0.6613	0.2219	0.8870	-0.1263	0.8806	-0.3482	0.5766
---	---	1631219_at	0.0001	0.9996	0.0975	0.4962	-0.1544	0.5491	-0.0601	0.9470	0.0350	0.9220	0.0951	0.7276	0.0654	0.9112	-0.0273	0.9348	-0.0927	0.6875
CG9850	Dlg-interacting me	1631220_at	-0.0475	0.7915	0.1081	0.5536	-0.1042	0.4887	-0.0602	0.9380	-0.1737	0.4636	-0.1135	0.6236	0.0977	0.9246	-0.0936	0.8399	-0.1914	0.6060
---	---	1631221_at	0.1360	0.5086	0.1463	0.5296	-0.0585	0.8647	-0.0222	0.9852	-0.0170	0.9676	0.0052	0.9875	0.0199	0.9848	-0.0055	0.9925	-0.0254	0.9397
pdm3	pou domain motif	1631222_at	1.2008	0.0883	2.4370	0.0021	0.2321	0.1622	0.1798	0.7531	0.2812	0.2570	0.1014	0.6964	2.3243	0.1379	1.4500	0.0461	-0.8743	0.1992
CG2100	CG2100	1631223_at	-0.2961	0.2171	0.2805	0.2787	0.6091	0.0210	0.0112	0.9874	-0.4516	0.0159	-0.4627	0.0086	-0.3231	0.6999	0.1448	0.7200	0.4679	0.2057
Est-6	Esterase-6	1631224_at	-1.8181	0.0023	0.0375	0.8781	-0.3277	0.5341	-1.1957	0.1615	-1.9338	0.0035	-0.7380	0.1102	-0.9875	0.4443	-0.1556	0.8716	0.8319	0.2336
CG8317	CG8317	1631225_at	2.6342	0.0011	2.2435	0.0459	3.5959	0.0002	0.3558	0.8028	-0.4600	0.4366	-0.8159	0.1088	-1.0529	0.5754	-0.9457	0.2474	0.1071	0.9273
sds22	sds22	1631226_at	0.1103	0.4643	-0.5167	0.1402	-0.4041	0.0185	0.0421	0.9436	0.7504	0.0017	0.7083	0.0013	0.0227	0.9884	0.2027	0.5684	0.1800	0.6216
CG5793	CG5793	1631227_at	0.5979	0.1283	0.4334	0.2070	0.4159	0.0421	-0.1934	0.6580	-0.8031	0.0032	-0.6097	0.0065	-0.1495	0.9309	-0.9044	0.1108	-0.7549	0.1993
sPLA2	secretory Phosph	1631228_a_at	2.1567	0.0012	1.2342	0.1564	2.2777	0.0001	0.4284	0.5832	-0.2400	0.5773	-0.6684	0.0635	-0.4487	0.8194	-1.1305	0.1498	-0.6818	0.4058
---	---	1631229_at	-0.0023	0.9910	-0.0177	0.9113	0.3098	0.0825	0.0666	0.9311	-0.0544	0.8529	-0.1209	0.5928	-0.2732	0.5754	-0.2331	0.2734	0.0401	0.8947
CG40053	CG40053	1631230_at	0.0688	0.6908	-0.0720	0.5837	0.0306	0.8599	0.0867	0.8578	0.1241	0.4954	0.0374	0.8526	-0.0159	0.9853	0.0162	0.9605	0.0321	0.9061
fz	Frizzled	1631231_at	-1.3959	0.0092	-2.4576	0.0097	-2.3311	0.0000	-0.2344	0.7492	0.3667	0.2518	0.6011	0.0413	-0.4152	0.6496	-0.7042	0.0716	-0.2890	0.4472
CG14471	CG14471	1631232_at	0.3387	0.0675	0.0268	0.7843	0.2803	0.1006	0.0846	0.8469	-0.0333	0.8759	-0.1179	0.4343	-0.0526	0.9365	-0.0489	0.8678	0.0037	0.9921
Hr83	CG10296	1631233_at	0.0688	0.7659	-0.0126	0.9048	0.0923	0.6103	-0.1167	0.8337	-0.2090	0.3284	-0.0923	0.6753	-0.2650	0.5959	-0.2324	0.2736	0.0326	0.9152
CG18173	CG18173	1631234_at	0.0767	0.9298	-0.0324	0.7469	0.2519	0.1866	0.0898	0.9727	-0.4148	0.5550	-0.5046	0.4074	-0.0727	0.9701	-0.3404	0.5075	-0.2677	0.6171
CG9899	CG9899	1631235_at	-0.0058	0.9757	0.1136	0.5532	0.1012	0.6279	0.0378	0.9627	-0.1405	0.5436	-0.1783	0.3692	0.1439	0.8215	0.0658	0.8569	-0.0780	0.8123
lin-28	lin-28	1631236_at	-0.7120	0.0245	-0.0322	0.7512	-0.3456	0.0717	-0.0478	0.9376	-0.6784	0.0033	-0.6306	0.0027	0.0384	0.9677	-0.1335	0.6191	-0.1719	0.5092
CG13384	CG13384	1631237_s_at	-1.0287	0.0106	-1.1967	0.0117	-1.0882	0.0010	-0.2062	0.7228	0.0825	0.7977	0.2887	0.2182	-0.1326	0.9238	-0.2630	0.5929	-0.1304	0.8221
Hsp60B	Heat shock protei	1631238_at	0.3032	0.0718	0.2523	0.2223	0.0624	0.7617	0.0530	0.9116	0.0937	0.5651	0.0407	0.8109	0.2010	0.7215	0.0286	0.9402	-0.1723	0.4786

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31131 /// CG31253	CG31253 /// CG3	1631239_s_at	0.0584	0.7162	0.1276	0.3692	-0.0491	0.7706	-0.0684	0.8794	-0.0671	0.7055	0.0013	0.9948	0.0386	0.9514	0.0650	0.7802	0.0264	0.9194
OstStt3	Oligosaccharyl tra	1631240_at	0.8102	0.0017	0.8380	0.0122	0.9968	0.0004	0.0385	0.9507	0.2794	0.1034	0.2409	0.1158	-0.0814	0.8973	0.2698	0.2371	0.3512	0.1603
---	---	1631241_at	0.3658	0.4298	0.1110	0.7111	0.0960	0.6423	0.0797	0.9580	0.2632	0.5395	0.1836	0.6579	0.0624	0.9515	-0.0403	0.9353	-0.1026	0.7785
jumu	Domina	1631242_at	0.8088	0.0506	-0.1312	0.5952	0.3069	0.3592	-0.3426	0.7599	0.0946	0.8837	0.4372	0.3195	-0.8609	0.5228	-0.9261	0.1423	-0.0652	0.9431
l(3)05822	lethal (3) 05822	1631243_s_at	-0.7533	0.0301	-0.7637	0.0534	-0.9679	0.0010	-0.0574	0.9319	0.2531	0.2032	0.3105	0.0832	0.1054	0.9521	0.1369	0.8481	0.0316	0.9687
Pabp2	polyA-binding prot	1631244_a_at	0.2820	0.2392	1.5756	0.0362	0.0776	0.6769	0.0890	0.9071	0.7975	0.0066	0.7084	0.0068	1.4669	0.1628	1.9462	0.0134	0.4793	0.3683
CG9706	CG9706	1631245_s_at	0.7278	0.0087	0.5964	0.1670	0.7995	0.0018	0.1298	0.7845	1.1515	0.0006	1.0216	0.0005	-0.0605	0.9742	0.9897	0.0535	1.0503	0.0607
Fmrf	FMRFamide	1631246_at	0.1245	0.5223	0.3279	0.1499	0.2164	0.1846	-0.0377	0.9603	-0.0247	0.9298	0.0130	0.9591	-0.0374	0.9751	0.1346	0.6827	0.1720	0.5851
CG33341	CG33341	1631247_at	-0.1038	0.5694	-0.0713	0.5879	0.0451	0.8614	0.1606	0.6954	0.0751	0.7364	-0.0855	0.6633	-0.0297	0.9826	0.0263	0.9590	0.0560	0.8957
l(1)G0020	lethal (1) G0020	1631248_at	0.1244	0.8377	0.1695	0.7252	0.6881	0.1146	0.6273	0.4626	1.1039	0.0250	0.4765	0.2509	0.1476	0.9589	1.1114	0.1617	0.9638	0.2555
CG12713	CG12713	1631249_at	0.3160	0.1483	0.2124	0.3643	-0.0704	0.7005	-0.3030	0.4633	0.4936	0.0344	0.7967	0.0026	-0.0864	0.9416	0.3041	0.3994	0.3905	0.2964
ind	intermediate nerv	1631250_at	0.0948	0.5411	0.1356	0.4658	0.2961	0.1800	0.0571	0.9068	-0.0027	0.9902	-0.0598	0.7204	0.0579	0.9444	0.0575	0.8721	-0.0004	0.9992
CG32708	CG32708	1631251_at	0.4910	0.0630	-0.0658	0.8542	-0.1661	0.4435	0.0953	0.9040	0.8910	0.0049	0.7956	0.0049	0.1815	0.8461	0.2775	0.4810	0.0960	0.8463
GV1	GV1	1631252_a_at	0.0084	0.9861	0.4627	0.3013	0.6701	0.0061	0.1190	0.8111	-0.5765	0.0106	-0.6955	0.0027	-0.1049	0.9589	-0.1771	0.8153	-0.0722	0.9293
---	---	1631253_at	0.1170	0.6062	0.0384	0.7080	-0.1260	0.5625	0.0744	0.9295	0.2169	0.3906	0.1425	0.5581	-0.0579	0.9400	0.0607	0.8509	0.1186	0.6410
Gr9a	Gr9a	1631254_at	-0.0611	0.8231	-0.0236	0.8886	-0.0314	0.8999	0.0198	0.9857	-0.0309	0.9294	-0.0507	0.8605	0.1138	0.9095	0.1780	0.6458	0.0642	0.8951
---	---	1631255_at	-0.0441	0.8463	0.0313	0.7827	0.0245	0.9263	-0.0343	0.9683	-0.0984	0.7043	-0.0641	0.7993	0.0272	0.9849	0.1071	0.7978	0.0800	0.8514
Upf2	Upf2	1631256_at	0.0245	0.9609	0.6100	0.2141	1.1194	0.0011	0.1011	0.9284	-0.4215	0.2030	-0.5226	0.0800	-0.0715	0.9742	0.4592	0.3972	0.5308	0.3468
---	---	1631257_at	0.1619	0.4860	0.0537	0.6301	0.0601	0.8055	0.0939	0.8539	0.1328	0.4916	0.0389	0.8562	0.1525	0.7726	0.1042	0.6766	-0.0483	0.8736
CG30005	CG30005	1631258_at	0.1545	0.6710	0.4346	0.0544	0.6191	0.0638	-0.1936	0.5046	-0.5823	0.0033	-0.3888	0.0124	-0.3586	0.7506	-0.3636	0.4513	-0.0050	0.9953
---	---	1631259_s_at	0.1826	0.2863	0.3192	0.2458	0.0699	0.7068	-0.1566	0.7443	-0.2760	0.1883	-0.1193	0.5685	0.1863	0.7712	0.1531	0.6006	-0.0331	0.9316
Or56a	Odorant receptor	1631260_at	0.1095	0.6817	0.1015	0.5177	0.0621	0.7577	-0.0299	0.9659	0.0305	0.9045	0.0604	0.7660	-0.0634	0.9409	-0.0203	0.9628	0.0430	0.9063
CG1806	CG1806	1631261_at	-0.4059	0.1159	-0.5076	0.1232	-0.8871	0.0039	0.1042	0.7608	0.6001	0.0024	0.4959	0.0033	0.3692	0.7464	0.4018	0.4022	0.0327	0.9636
CG14866	CG14866	1631262_at	0.1192	0.5452	0.0097	0.9536	0.2227	0.1784	0.1237	0.7863	-0.0060	0.9827	-0.1297	0.4684	-0.0022	0.9984	-0.0926	0.7058	-0.0904	0.7128
CG4279	CG4279	1631263_at	0.3873	0.0530	0.3410	0.1388	0.3643	0.1744	-0.0586	0.9518	0.1206	0.6957	0.1792	0.4839	-0.1670	0.8609	0.0266	0.9664	0.1935	0.6392
CG2211	CG2211	1631264_at	0.9324	0.0063	0.5555	0.0226	0.5908	0.0157	0.0659	0.9387	0.1075	0.7119	0.0416	0.8903	-0.1577	0.7953	-0.1181	0.6772	0.0395	0.9108
CG6280	CG6280	1631265_at	0.2126	0.3005	0.1023	0.4980	0.1505	0.6634	-0.1365	0.8837	-0.1502	0.6792	-0.0137	0.9724	0.0019	0.9994	-0.1259	0.6966	-0.1279	0.6875
CG5840	CG5840	1631266_a_at	2.4441	0.0039	1.1348	0.2005	1.8543	0.0007	0.4212	0.7293	-0.3620	0.5344	-0.7831	0.1073	-0.3725	0.8681	-1.6659	0.0757	-1.2935	0.1691
CG32718	CG32718	1631267_at	0.0826	0.5982	-0.0357	0.8760	0.1181	0.5811	0.0686	0.9314	0.1470	0.5669	0.0784	0.7637	-0.0225	0.9884	0.1466	0.6966	0.1691	0.6389
ast	Star-recessive	1631268_at	0.1449	0.5158	0.8637	0.0435	1.1595	0.0028	0.1170	0.8738	-0.3004	0.2420	-0.4174	0.0711	-0.5092	0.6935	0.2462	0.6762	0.7555	0.1796
---	---	1631269_at	0.1080	0.6444	0.0241	0.8101	0.2532	0.2920	0.0253	0.9777	0.1129	0.6505	0.0876	0.7086	-0.1495	0.7215	-0.0735	0.7088	0.0760	0.6941
nht	nohitter	1631270_at	0.3327	0.0769	0.0830	0.5842	-0.0017	0.9943	-0.0573	0.9451	0.0799	0.7811	0.1372	0.5585	0.1446	0.7893	-0.0730	0.8054	-0.2176	0.3601
hang	hangover	1631271_a_at	-0.1998	0.6383	-0.5896	0.0459	-0.8109	0.0037	-0.0845	0.9011	0.4271	0.0571	0.5117	0.0175	0.0780	0.9657	0.0897	0.9052	0.0117	0.9878
---	---	1631272_at	0.0092	0.9778	-0.3715	0.1242	-0.2970	0.2834	0.0479	0.9619	0.3189	0.2282	0.2710	0.2558	-0.0572	0.9775	-0.0088	0.9935	0.0484	0.9431
CG5262	CG5262	1631273_at	-0.0023	0.9941	0.1607	0.6829	0.3393	0.0471	-0.2264	0.5863	-0.1946	0.3693	0.0319	0.9003	-0.3585	0.7113	-0.0001	1.0000	0.3584	0.3849
CG18095	gp150-like	1631274_at	0.2109	0.2663	-0.1629	0.2695	-0.1945	0.3994	0.2253	0.6325	0.4308	0.0651	0.2055	0.3292	0.1093	0.8270	-0.0539	0.8511	-0.1632	0.4523
---	---	1631275_at	0.1378	0.3463	0.0485	0.6492	0.0884	0.5817	-0.1134	0.8103	-0.0493	0.8383	0.0641	0.7548	-0.0404	0.9530	-0.1401	0.5079	-0.0996	0.6528
CG31389	CG31389	1631276_at	0.0123	0.9847	-0.0419	0.7923	0.0135	0.9578	-0.0967	0.9295	-0.4619	0.1475	-0.3652	0.2015	-0.0911	0.9535	-0.3300	0.4868	-0.2389	0.6328
alpha-Est4	fragment C	1631277_at	0.3529	0.1987	-0.0227	0.9254	-0.4725	0.0630	-0.1468	0.7958	-0.0973	0.7166	0.0496	0.8534	-0.0915	0.9677	-0.6474	0.2634	-0.5559	0.3676
---	---	1631278_a_at	0.2134	0.1788	0.1984	0.5032	0.1126	0.5557	-0.0800	0.8948	0.2966	0.1389	0.3767	0.0425	-0.0662	0.9467	0.2804	0.3397	0.3466	0.2668
CG4497	CG4497	1631279_at	-0.0976	0.6219	0.2492	0.1460	0.2933	0.2613	-0.0238	0.9865	-0.3059	0.3799	-0.2821	0.3698	0.1758	0.7644	0.1441	0.5855	-0.0317	0.9270
CG9095	CG9095	1631280_at	-0.4710	0.1210	-1.2029	0.0260	-1.1362	0.0003	0.2371	0.5587	0.4787	0.0293	0.2416	0.1943	-0.2303	0.8062	-0.4471	0.2385	-0.2168	0.6074
amd	dopa decarboxyla	1631281_a_at	-0.5636	0.0222	-1.6704	0.0048	-1.7678	0.0045	0.1500	0.9339	0.9104	0.0878	0.7605	0.1089	-0.1288	0.9095	-0.2442	0.5635	-0.1154	0.8182
CG31088	CG31088	1631282_at	0.0038	0.9889	0.0052	0.9731	-0.2179	0.4103	-0.0202	0.9857	0.0800	0.7986	0.1002	0.7120	0.0878	0.8909	0.0927	0.7439	0.0048	0.9906
---	---	1631283_at	0.0662	0.7145	-0.0367	0.7141	-0.0397	0.8380	-0.1083	0.8202	-0.1432	0.4617	-0.0349	0.8741	-0.0946	0.8608	-0.1288	0.5732	-0.0343	0.9089
---	---	1631284_at	0.0587	0.7841	0.1439	0.3889	0.0252	0.9041	0.0390	0.9610	-0.1062	0.6607	-0.1451	0.4755	0.2299	0.7131	0.1675	0.5378	-0.0623	0.8540
Tsp5D	Tetraspanin 5D	1631285_at	-0.5708	0.4294	0.1873	0.2712	0.0699	0.8071	-0.3097	0.8578	-1.9089	0.0085	-1.5992	0.0115	-0.0013	0.9998	-0.7863	0.1820	-0.7850	0.2136
CG7857	CG7857	1631286_at	0.4245	0.0291	0.0032	0.9885	0.1336	0.4314	0.1276	0.8000	0.6888	0.0054	0.5611	0.0082	-0.0438	0.9555	0.3136	0.1579	0.3573	0.1443
---	---	1631287_at	0.1244	0.4402	0.2520	0.0946	0.1519	0.3859	-0.0119	0.9912	-0.1594	0.5200	-0.1474	0.5131	-0.0361	0.9611	-0.0576	0.8356	-0.0215	0.9412
CG11696	CG11696	1631288_at	-0.0925	0.6898	0.3418	0.4973	0.3600	0.0435	0.1221	0.7695	-0.1511	0.4049	-0.2732	0.0844	0.0674	0.9774	0.2369	0.7082	0.1695	0.8040
CG12540	CG12540	1631289_at	0.1175	0.5223	0.1142	0.4558	0.0643	0.6887	-0.0052	0.9943	-0.0292	0.8858	-0.0240	0.8948	-0.0020	0.9990	-0.0894	0.7900	-0.0874	0.7836
CG15641	CG15641	1631290_at	0.0770	0.7159	0.2086	0.4620	0.1870	0.4704	0.0494	0.9515	-0.1071	0.6748	-0.1565	0.4607	0.3228	0.7644	0.1574	0.7798	-0.1654	0.7536
---	---	1631291_at	-2.3696	0.0006	-2.4116	0.0070	-2.7512	0.0000	-0.4850	0.2037	-0.4037	0.0805	0.0813	0.7428	-0.1277	0.8486	-0.5128	0.0795	-0.3851	0.1911
---	---	1631292_at	-0.0265	0.9040	-0.0427	0.6822	-0.1251	0.5552	-0.1680	0.7204	0.1026	0.6711	0.2705	0.1526	-0.2632	0.7436	-0.0362	0.9477	0.2270	0.5239
CG31600	CG31600	1631293_at	0.0433	0.8762	-0.0848	0.4239	0.4386	0.1572	0.2061	0.6510	0.0811	0.7581	-0.1251	0.5663	-0.3453	0.7220	0.0504	0.9402	0.3957	0.3543
CG33263	CG33263	1631294_at	0.1622	0.3358	0.0279	0.8112	0.1024	0.5484	0.0020	0.9983	-0.1085	0.6377	-0.1105	0.5914	-0.0328	0.9677	-0.1193	0.6011	-0.0865	0.7221
---	---	1631295_s_at	0.3575	0.1942	0.2201	0.0779	0.3803	0.0225	0.0779	0.9247	0.0725	0.8123	-0.0054	0.9861	0.0467	0.9588	-0.0017	0.9988	-0.0485	0.8887
RpS27	Ribosomal protein	1631296_at	0.2895	0.0845	0.4883	0.0266	0.4367	0.0093	0.0090	0.9902	-0.1708	0.2966	-0.1798	0.2153	-0.0303	0.9672	0.0008	0.9994	0.0311	0.9080
Mat89Bb	Maternal transcrip	1631297_at	-0.2962	0.1999	0.0663	0.6440	0.4403	0.0165	-0.0101	0.9904	-0.4821	0.0182	-0.4720	0.0124	-0.2968	0.6955	0.1424	0.6847	0.4393	0.1880
Or22b	Olfactory receptor	1631298_at	0.0459	0.8339	0.0247	0.8087	-0.1574	0.3929	-0.1258	0.8251	-0.0807	0.7603	0.0451	0.8605	0.1191	0.8541	-0.1591	0.5715	-0.2782	0.3082
CG31759	CG31759	1631299_s_at	0.8506	0.0019	0.2577	0.5902	0.4994	0.0192	0.2843	0.3739	0.9211	0.0009	0.6368	0.0027	0.0775	0.9736	0.3574	0.5531	0.2799	0.6533
Cyp18a1	Cytochrome P450	1631300_at	1.9892	0.0075	-0.5028	0.7310	0.5464	0.2269	0.8334	0.2789	1.3023	0.0114	0.4689	0.2503	-0.3439	0.9457	-1.3419	0.3695	-0.9980	0.5335
CG30398	CG30398	1631301_at	0.0302	0.9060	0.0759	0.6461	0.1453	0.5724	-0.1824	0.7128	-0.2628	0.2448	-0.0804	0.7403	0.0972	0.8940	0.1054	0.7400	0.0082	0.9849
CG16885	CG16885	1631302_at	-0.1345	0.5339	-0.1259	0.4881	-0.3188	0.2103	0.0448	0.9426	-0.0839	0.6780	-0.1287	0.4397	-0.0384	0.9809	-0.0547	0.9191	-0.0163	0.9766
NK7.1	NK7.1	1631303_s_at	0.5328	0.3122	-0.1694	0.6255	-0.2569	0.4172	0.1469	0.8967	0.9660	0.0179	0.8191	0.0223	0.2984	0.8609	0.2453	0.7695	-0.0532	0.9575
CG1968	CG1968	1631304_a_at	0.5883	0.0127	0.4899	0.0754	0.5949	0.0043	0.1516	0.7492	0.7982	0.0030	0.6466	0.0046	0.0665	0.9246	0.6626	0.0276	0.5962	0.0457
---	---	1631305_at	-0.2098	0.1821	-0.2956	0.2322	-0.1099	0.5398	0.1441	0.7854	0.1784	0.4291	0.0344	0.8956	-0.0994	0.8740	-0.0146	0.9719	0.0847	0.7689
CG30464	CG30464	1631306_at	0.2194	0.2627	0.1832	0.3430	0.2728	0.1278	-0.0101	0.9894	0.0956	0.6192	0.1057	0.5335	-0.0311	0.9717	0.0988	0.6910	0.1299	0.5814
CG11438	CG11438	1631307_at	-0.5953	0.2296	-0.8555	0.1746	-0.4062	0.4355	0.6173	0.5038	0.5545	0.2591	-0.0628	0.9163	0.3233	0.8956	0.2830	0.8123	-0.0403	0.9778

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG10737	Gene 1	1631308_at	-1.0111	0.0035	-0.7560	0.1251	-0.7004	0.0046	0.2465	0.5018	0.0703	0.7637	-0.1762	0.3214	0.1740	0.8882	0.2308	0.6488	0.0567	0.9312
CG7280	CG7280	1631309_at	0.3540	0.0342	0.3386	0.5406	0.3604	0.2056	-0.3877	0.3933	-0.6275	0.0225	-0.2398	0.3018	-0.3271	0.8236	-0.5595	0.3494	-0.2324	0.7426
---	---	1631310_at	0.1145	0.4918	0.2322	0.1849	0.3411	0.1335	-0.1267	0.7749	-0.3579	0.0579	-0.2312	0.1620	-0.0701	0.9116	-0.0569	0.8535	0.0132	0.9694
CG14351	CG14351	1631311_at	0.0440	0.7911	0.1582	0.2645	-0.1293	0.5483	-0.0587	0.9042	-0.0187	0.9331	0.0400	0.8247	0.2886	0.7142	0.1913	0.5827	-0.0973	0.8111
CG4073	CG4073	1631312_at	0.1242	0.4802	0.0557	0.7410	0.0864	0.7473	-0.0765	0.9026	0.0453	0.8653	0.1218	0.5381	0.0653	0.9486	0.1769	0.5818	0.1116	0.7492
CG2846	transcript C	1631313_at	0.1950	0.3630	0.6937	0.0275	0.4936	0.0059	-0.0368	0.9633	-0.2118	0.3252	-0.1750	0.3722	0.1096	0.8202	0.2220	0.2507	0.1124	0.6020
---	---	1631314_at	0.0193	0.9363	-0.2058	0.3377	-0.0030	0.9923	0.0687	0.9228	-0.0964	0.6965	-0.1651	0.4108	-0.2230	0.7464	-0.2385	0.4097	-0.0155	0.9727
CG12730	CG12730	1631315_at	-0.2428	0.3158	0.3576	0.1382	0.1956	0.2875	-0.4158	0.3784	-0.8305	0.0080	-0.4147	0.0833	-0.0766	0.8903	-0.0501	0.8631	0.0265	0.9266
CG6262	CG6262	1631316_a_at	0.1165	0.4947	0.1566	0.2787	0.1269	0.5400	-0.1025	0.8508	-0.1320	0.5340	-0.0295	0.9015	0.0008	0.9998	-0.0829	0.8137	-0.0837	0.7993
Or82a	Odorant receptor	1631317_at	0.1258	0.5041	-0.1539	0.3186	-0.0062	0.9756	0.0686	0.9029	0.1218	0.5305	0.0532	0.7945	0.0218	0.9862	-0.0456	0.9178	-0.0674	0.8562
l(2)35Di	lethal (2) 35Di	1631318_at	-0.5573	0.0653	-0.4439	0.1233	-0.2652	0.2309	-0.0672	0.9221	-0.6480	0.0078	-0.5809	0.0077	-0.2210	0.8379	-0.4929	0.2507	-0.2719	0.5649
CG32462	CG32462	1631319_at	-0.2387	0.3950	-0.0150	0.8834	0.1663	0.2920	0.0477	0.9538	-0.2595	0.2513	-0.3072	0.1262	-0.1002	0.9092	-0.0579	0.9036	0.0424	0.9215
Pp2C1	Protein phosphata	1631320_at	-0.3822	0.2648	0.4692	0.1660	0.7444	0.0181	0.0415	0.9666	-0.6612	0.0193	-0.7027	0.0091	-0.2725	0.8222	0.1499	0.8185	0.4223	0.4090
His1:CG31617 /// His1:CG3	His1:CG31617 ///	1631321_s_at	0.3964	0.2931	-1.4848	0.0884	-1.5636	0.0025	0.2280	0.7121	2.5581	0.0001	2.3301	0.0001	0.0384	0.9928	0.2777	0.8397	0.2393	0.8577
---	---	1631322_at	3.5287	0.0011	2.5971	0.0168	4.6884	0.0000	1.1523	0.0909	0.9264	0.0281	-0.2259	0.5686	-0.8435	0.6092	-0.3442	0.6581	0.4993	0.4970
CG2663	alpha tocopherol t	1631323_a_at	-1.8164	0.0665	0.0673	0.5555	0.0397	0.8662	-0.0311	0.9915	-2.5206	0.0024	-2.4895	0.0015	0.1336	0.9499	-0.3315	0.6270	-0.4651	0.4776
CG14735 /// DsimCG14735	CG14735	1631324_at	0.1685	0.3738	0.2344	0.2504	0.0057	0.9852	-0.0507	0.9523	-0.0552	0.8575	-0.0045	0.9875	0.1872	0.7485	0.0279	0.9451	-0.1593	0.5448
Sry-beta	serendipity beta	1631325_at	-0.0399	0.8333	-0.0720	0.5637	0.2083	0.3223	-0.0122	0.9902	-0.0773	0.7711	-0.0651	0.7926	-0.1428	0.7689	0.0163	0.9620	0.1591	0.4568
CG6416	CG6416	1631326_at	-2.1903	0.0022	-1.5577	0.0259	-2.7128	0.0001	-0.6494	0.1140	-0.2795	0.2591	0.3699	0.0935	0.1933	0.9011	-0.0596	0.9492	-0.2529	0.6885
mspo	m-spondin	1631327_at	-2.8849	0.0004	-3.6988	0.0014	-3.8773	0.0000	0.4108	0.5681	0.8727	0.0261	0.4620	0.1611	0.2718	0.7324	-0.0521	0.9221	-0.3239	0.3552
DnaJ-H	DnaJ homolog	1631328_s_at	-0.9413	0.0033	-0.3917	0.0751	-0.2440	0.2209	0.0233	0.9796	-0.3660	0.1005	-0.3894	0.0550	-0.0147	0.9918	0.3631	0.2566	0.3777	0.2729
---	---	1631329_at	0.0667	0.8077	0.0682	0.5632	0.0583	0.7971	-0.0077	0.9951	-0.0978	0.7443	-0.0901	0.7415	0.1074	0.8379	0.1473	0.5084	0.0399	0.8926
CG3408	CG3408	1631330_at	0.7858	0.0096	0.9627	0.1284	0.8842	0.0042	-0.3594	0.2478	-0.2010	0.2783	0.1584	0.3494	-0.3226	0.8317	-0.0647	0.9499	0.2579	0.7185
pxt	pxt	1631331_a_at	0.8003	0.6268	-2.3644	0.2552	-1.9920	0.0677	-0.2389	0.7564	3.2388	0.0001	3.4777	0.0000	-0.7955	0.9340	-0.1381	0.9808	0.6574	0.8740
---	---	1631332_at	0.2089	0.3525	-0.0322	0.8201	0.0294	0.8589	0.0563	0.9376	0.1102	0.6407	0.0539	0.8214	0.0185	0.9816	-0.0856	0.6853	-0.1041	0.6100
Adh	alcohol dehydroge	1631333_s_at	0.2447	0.2040	0.0413	0.8606	0.3604	0.0511	0.0518	0.9226	-0.0024	0.9916	-0.0542	0.7571	-0.2024	0.8202	-0.1967	0.6189	0.0058	0.9927
CG10260	CG10260	1631334_at	-0.9855	0.1928	-0.9634	0.4825	-0.8531	0.0654	0.2560	0.6998	0.2550	0.4176	-0.0010	0.9981	0.1334	0.9860	0.3024	0.9075	0.1691	0.9433
CG31275	late transcript	1631335_a_at	0.0143	0.9547	0.1077	0.4920	0.0431	0.8496	-0.2803	0.5724	-0.0774	0.8036	0.2029	0.3892	0.0485	0.9514	0.0445	0.9027	-0.0041	0.9924
---	---	1631336_at	-0.1121	0.6042	-0.2598	0.2300	0.0133	0.9560	0.1904	0.7376	0.0772	0.8026	-0.1132	0.6640	-0.0589	0.9530	-0.1048	0.7737	-0.0459	0.9109
CG33082	CG33082	1631337_at	-0.0627	0.8441	0.0972	0.4299	0.1147	0.6418	0.0979	0.9303	0.0094	0.9845	-0.0885	0.8111	-0.0595	0.9618	0.0561	0.9170	0.1156	0.7774
---	---	1631338_at	0.0617	0.7157	-0.0132	0.9672	0.0356	0.8709	0.0610	0.9375	0.0726	0.7935	0.0116	0.9679	-0.1565	0.8331	-0.1070	0.7784	0.0496	0.9077
---	---	1631339_s_at	-0.1068	0.5706	0.0045	0.9733	-0.1128	0.6231	-0.1736	0.7005	-0.0186	0.9520	0.1550	0.4250	-0.1172	0.8480	-0.0442	0.9094	0.0730	0.8178
Ggamma1	G protein gamma	1631340_at	-0.9658	0.0061	-1.8485	0.0149	-2.0024	0.0000	0.0527	0.9351	0.3029	0.1115	0.2502	0.1411	0.1799	0.8611	-0.3640	0.3739	-0.5439	0.2110
---	---	1631341_at	-0.0126	0.9548	0.0647	0.6159	0.1347	0.5619	0.0823	0.9228	-0.0298	0.9345	-0.1121	0.6767	-0.0826	0.8740	-0.0287	0.9291	0.0539	0.8354
Elongin-B	d-elongin B	1631342_at	0.3671	0.2289	0.4415	0.1015	0.3902	0.1570	0.0449	0.9425	0.2016	0.2570	0.1567	0.3336	0.0797	0.9701	0.1323	0.8612	0.0526	0.9444
CG15571	CG15571	1631343_at	0.0332	0.8693	0.0553	0.6506	0.0710	0.6506	0.0206	0.9760	0.0280	0.9045	0.0074	0.9716	0.0907	0.8558	0.1288	0.5469	0.0381	0.8921
msb1l	msb1l	1631344_at	0.2977	0.7951	-0.0779	0.9519	-1.0522	0.0685	-0.8262	0.3016	0.8315	0.0791	1.6577	0.0027	-0.3126	0.9558	-0.0585	0.9855	0.2541	0.9107
CrebB-17A	cAMP-regulated e	1631345_at	0.0079	0.9854	0.3686	0.4551	0.7461	0.0073	0.3517	0.6144	0.1771	0.6532	-0.1746	0.6244	0.1426	0.8653	0.6769	0.0637	0.5343	0.1393
CG30412	CG30412	1631346_a_at	-0.0307	0.9017	0.1853	0.3352	0.0917	0.6827	-0.0026	0.9956	-0.0182	0.9326	-0.0156	0.9344	-0.0689	0.9330	0.1764	0.5168	0.2453	0.3636
CG30393	CG30393	1631347_at	0.0714	0.7241	0.0043	0.9844	0.3452	0.0530	0.0982	0.8701	0.0764	0.7618	-0.0218	0.9334	-0.1486	0.8236	0.0333	0.9405	0.1819	0.5344
---	---	1631348_at	0.0338	0.8884	-0.0916	0.4514	-0.0797	0.7050	-0.0607	0.9422	0.0766	0.7927	0.1373	0.5589	-0.1084	0.8298	-0.0214	0.9501	0.0870	0.7154
---	---	1631349_s_at	-1.3261	0.0438	-3.1882	0.0061	-2.4050	0.0035	0.3504	0.8156	1.3511	0.0297	1.0007	0.0617	-0.4253	0.8395	-0.5455	0.5451	-0.1202	0.9197
CG18649	CG18649	1631350_at	-0.0276	0.8738	0.0476	0.7929	0.0275	0.9036	-0.0322	0.9562	0.0336	0.8728	0.0658	0.6924	0.0977	0.8655	0.1431	0.5523	0.0455	0.8861
CG12118	CG12118	1631351_s_at	-0.4418	0.1420	-0.2577	0.1816	-0.3110	0.2131	0.0036	0.9956	-0.6749	0.0033	-0.6785	0.0019	-0.0386	0.9848	-0.5132	0.2346	-0.4746	0.2993
---	---	1631352_at	-0.1718	0.3265	0.1174	0.5831	0.1328	0.4223	-0.0742	0.9011	-0.3086	0.1100	-0.2344	0.1748	0.0419	0.9611	0.0865	0.7692	0.0446	0.8933
ng4	new glue 4	1631353_at	0.1007	0.6553	-0.0134	0.9239	0.0271	0.8909	0.0079	0.9937	0.0030	0.9918	-0.0049	0.9855	-0.0817	0.8972	-0.1778	0.4514	-0.0962	0.7184
CG13679	CG13679	1631354_x_at	0.1385	0.4892	0.0448	0.6506	0.0852	0.5785	0.0538	0.9186	0.0022	0.9922	-0.0516	0.7737	0.0219	0.9862	-0.0662	0.8678	-0.0881	0.7961
mRpS28	mitochondrial ribo	1631355_at	0.0860	0.8041	0.4296	0.1113	0.4639	0.0174	0.2115	0.4551	-0.4610	0.0093	-0.6725	0.0011	0.0651	0.9717	-0.1303	0.8299	-0.1954	0.7030
scaf6	scaf6	1631356_at	-0.0554	0.7560	0.4630	0.3098	0.3121	0.2515	-0.0702	0.9117	-0.0454	0.8640	0.0247	0.9202	0.0731	0.9742	0.2523	0.6908	0.1792	0.7919
---	---	1631357_at	0.2792	0.1602	0.1166	0.7338	0.0103	0.9636	0.0769	0.9136	0.2060	0.3671	0.1291	0.5588	0.1849	0.8379	0.0650	0.9085	-0.1199	0.7865
CG32219	CG32219	1631358_at	0.0872	0.5988	0.1453	0.4966	0.1757	0.2804	-0.0343	0.9542	-0.0684	0.7194	-0.0341	0.8582	-0.0468	0.9683	-0.0326	0.9490	0.0143	0.9769
Vha100-2	Vha100-2	1631359_s_at	-0.9527	0.0099	-0.8475	0.0654	-1.1553	0.0001	-0.2471	0.3855	-0.4700	0.0106	-0.2228	0.1206	0.0332	0.9877	-0.4293	0.3680	-0.4624	0.3545
CG4133	CG4133	1631360_at	-0.0449	0.8667	0.0323	0.7622	0.0037	0.9869	0.0538	0.9518	-0.0694	0.8223	-0.1232	0.6246	-0.0095	0.9914	-0.0451	0.8679	-0.0356	0.8926
---	---	1631361_at	0.0226	0.9116	-0.0012	0.9930	0.0042	0.9860	-0.0445	0.9518	0.0800	0.7414	0.1244	0.5345	-0.1325	0.7768	0.0292	0.9275	0.1617	0.4370
CG11279	CG11279	1631362_at	0.2551	0.2125	0.4654	0.2611	0.6475	0.0252	0.2919	0.5528	0.1647	0.5469	-0.1271	0.6234	0.1692	0.8609	0.2268	0.5854	0.0576	0.9159
CG8902	CG8902	1631363_at	-1.0784	0.0148	-2.0635	0.0207	-1.9721	0.0058	-0.4686	0.3743	0.8164	0.0140	1.2850	0.0011	-0.5431	0.8270	-0.2076	0.8918	0.3355	0.7849
---	---	1631364_at	0.2045	0.2479	0.1609	0.1424	0.2132	0.3528	-0.0705	0.9314	-0.1750	0.4939	-0.1045	0.6822	0.0509	0.9340	0.0074	0.9842	-0.0435	0.8674
CG9760	CG9760	1631365_at	-0.0088	0.9719	-0.0348	0.8100	0.2815	0.1117	0.1736	0.7003	-0.1807	0.3950	-0.3543	0.0595	-0.0900	0.8740	-0.0712	0.8029	0.0188	0.9523
CG14752	CG14752	1631366_at	0.2743	0.3770	0.7517	0.0398	0.3222	0.0958	0.0532	0.9603	-0.3870	0.1752	-0.4401	0.0866	0.1270	0.8967	-0.0492	0.9326	-0.1762	0.6553
CG7828	CG7828	1631367_at	0.0270	0.8790	0.3217	0.0639	0.9324	0.0032	-0.0184	0.9835	-0.3420	0.0937	-0.3236	0.0773	-0.4777	0.3712	0.1019	0.7741	0.5796	0.0833
CG8108	CG8108	1631368_s_at	-0.2182	0.5900	0.3185	0.3387	0.4605	0.0121	-0.1202	0.6948	-0.1235	0.3946	-0.0033	0.9852	-0.2979	0.8331	0.3420	0.5855	0.6398	0.2938
CG7567	CG7567	1631369_at	0.5993	0.0140	0.4268	0.2592	0.5782	0.0080	0.1920	0.5311	0.2646	0.0978	0.0726	0.6612	-0.0620	0.9665	0.0521	0.9341	0.1141	0.8161
CG30026	CG30026	1631370_at	0.4574	0.3296	0.7543	0.1243	1.2623	0.0034	0.5964	0.3863	-0.1572	0.7304	-0.7537	0.0373	0.1679	0.9405	0.0922	0.9341	-0.0757	0.9382
Pde8	Phosphodiesteras	1631371_s_at	-2.8233	0.0104	-2.0971	0.0299	-3.1652	0.0000	-0.9660	0.1903	-0.4864	0.2790	0.4796	0.2307	-0.0294	0.9948	0.2248	0.8812	0.2542	0.8509
AP-2	stummelbein	1631372_a_at	0.1690	0.3665	0.2114	0.3409	0.1540	0.3029	-0.0013	0.9988	-0.1299	0.5543	-0.1286	0.5155	0.0437	0.9589	-0.0523	0.8792	-0.0960	0.7262
---	---	1631373_s_at	-0.1306	0.4970	0.1123	0.4037	0.0936	0.6255	-0.1106	0.8423	-0.2986	0.1461	-0.1880	0.3174	0.0020	0.9984	0.0799	0.7383	0.0779	0.7416
chico	flipper	1631374_at	0.1092	0.5885	0.0136	0.9779	-0.5149	0.0647	-0.3777	0.2939	-0.0160	0.9599	0.3617	0.0596	0.0964	0.9487	-0.2048	0.6820	-0.3012	0.5200
ImpE1	Y1-like	1631375_a_at	-0.1723	0.4001	-0.0342	0.8537	0.0119	0.9602	0.0161	0.9866	-0.0882	0.7518	-0.1043	0.6681	-0.0612	0.9421	-0.0395	0.9225	0.0217	0.9525
Pros35	20s proteasome	1631376_at	0.3501	0.0583	0.5494	0.0250	0.5786	0.0256	0.2613	0.4551	-0.0523	0.8266	-0.3136	0.0672	0.2146	0.7712	0.1586	0.6430	-0.0560	0.8976

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Adk1	Adenylate kinase-	1631377_a_at	-1.7111	0.0015	-0.0995	0.5779	-1.2312	0.0029	-0.2566	0.7018	-1.8717	0.0005	-1.6151	0.0005	0.9771	0.3712	0.1418	0.8615	-0.8353	0.1764
beat-Ia	triclops	1631378_at	0.0199	0.9344	-0.0142	0.9432	0.0059	0.9779	0.1050	0.7838	0.1011	0.5521	-0.0039	0.9849	0.0139	0.9913	0.0789	0.8216	0.0650	0.8508
ERp60	anon-fast-evolving	1631379_a_at	-0.3804	0.0409	-0.3630	0.0993	-0.2863	0.0810	0.0696	0.8950	0.1985	0.2594	0.1289	0.4340	-0.0264	0.9816	0.2616	0.2680	0.2879	0.2557
LanB1	Laminin	1631380_s_at	-1.8516	0.0004	-2.2878	0.0298	-2.5492	0.0000	-0.1920	0.5522	-0.0940	0.6092	0.0980	0.5506	-0.0035	0.9994	-0.5727	0.3438	-0.5692	0.3717
Uba2	ubiquitin-activating	1631381_at	-0.6653	0.0077	0.2741	0.1109	0.7721	0.0026	-0.1819	0.6673	-0.9297	0.0014	-0.7478	0.0022	-0.5166	0.3800	0.2033	0.5488	0.7198	0.0625
CG13085	CG13085	1631382_at	0.7934	0.0164	0.4044	0.1309	0.7821	0.0146	-0.2535	0.6120	-0.0422	0.9026	0.2114	0.3582	-0.5802	0.5955	-0.5345	0.2499	0.0457	0.9463
CG30472	CG30472	1631383_at	-0.0633	0.7313	-0.0357	0.7263	-0.0496	0.8267	-0.0380	0.9603	0.0628	0.8030	0.1008	0.6303	-0.0111	0.9923	-0.0270	0.9504	-0.0159	0.9691
giac	CG6589	1631384_at	0.1243	0.4624	-0.1652	0.1579	0.2891	0.1123	0.2153	0.5515	0.2115	0.2594	-0.0038	0.9873	-0.1638	0.7936	-0.1656	0.5471	-0.0018	0.9973
CG2889	CG2889	1631385_at	-0.2566	0.3573	0.2718	0.2352	0.6795	0.0092	0.3762	0.4930	-0.3914	0.1777	-0.7676	0.0102	0.0865	0.9333	0.2742	0.4031	0.1877	0.5984
---	---	1631386_at	0.3021	0.2161	0.0210	0.8432	0.1934	0.3468	0.2843	0.3553	0.3064	0.0828	0.0221	0.9185	0.0517	0.9405	0.0096	0.9806	-0.0420	0.8861
CG30344	CG30344	1631387_at	-0.2587	0.4349	-0.3880	0.1565	0.1485	0.5785	-0.2167	0.6338	-0.6499	0.0107	-0.4332	0.0370	-0.5890	0.6660	-0.7275	0.1718	-0.1386	0.8435
Rep	rab escort protein	1631388_at	-0.0878	0.7749	-0.2342	0.5537	-0.3262	0.2219	-0.0192	0.9759	0.3331	0.0396	0.3523	0.0200	0.0692	0.9734	0.2071	0.7259	0.1379	0.8307
---	---	1631389_at	0.3321	0.2396	-0.5401	0.1683	-0.8587	0.0042	0.3042	0.7187	1.1191	0.0105	0.8149	0.0255	0.1300	0.8903	0.0910	0.8504	-0.0390	0.9371
CG5919	CG5919	1631390_at	0.0662	0.7545	-0.2154	0.2671	0.1758	0.3023	0.2858	0.3793	-0.1307	0.4997	-0.4165	0.0195	0.0460	0.9618	-0.3012	0.2430	-0.3473	0.2130
---	---	1631391_at	-0.0780	0.6467	-0.0446	0.7525	-0.0989	0.6038	-0.0843	0.8578	0.1761	0.3014	0.2604	0.0851	-0.0504	0.9588	0.0508	0.9009	0.1012	0.7443
CG11089	CG11089	1631392_at	2.0029	0.0013	1.1223	0.1746	2.2393	0.0000	0.6372	0.2008	0.3392	0.2606	-0.2980	0.2717	-0.6283	0.7237	-0.6714	0.3744	-0.0431	0.9711
CG32459	CG32459	1631393_at	-0.0320	0.8545	0.0525	0.6183	0.1443	0.6240	0.0620	0.9518	-0.0434	0.9127	-0.1054	0.7279	0.0170	0.9862	0.0535	0.8664	0.0365	0.9067
CG31324	CG31324	1631394_at	-1.8629	0.1906	-0.6089	0.5175	-0.9424	0.0224	0.3749	0.9254	1.6902	0.1581	1.3154	0.2220	0.5755	0.8202	2.8027	0.0290	2.2272	0.0710
---	---	1631395_at	0.0688	0.7642	0.0285	0.9211	0.0048	0.9843	-0.0255	0.9734	-0.0430	0.8626	-0.0175	0.9404	0.0338	0.9831	0.0305	0.9608	-0.0034	0.9960
tos	tosca	1631396_at	0.1666	0.6991	-1.3302	0.0737	-1.3671	0.0063	0.0973	0.8723	1.5096	0.0003	1.4122	0.0002	0.1149	0.9776	0.2921	0.8186	0.1772	0.8949
---	---	1631397_at	0.1824	0.4135	0.1300	0.5678	0.3451	0.1324	-0.0279	0.9759	-0.0508	0.8649	-0.0229	0.9349	-0.0123	0.9928	-0.0606	0.8988	-0.0483	0.9118
Toll-4	Toll-like	1631398_at	0.2212	0.3120	-0.4678	0.0223	-0.1822	0.4636	0.1803	0.7664	0.6314	0.0229	0.4511	0.0554	-0.0948	0.8494	-0.1646	0.4258	-0.0698	0.7751
CG15771	CG15771	1631399_a_at	0.0921	0.6978	0.7739	0.0655	1.2703	0.0017	0.0290	0.9824	-0.8223	0.0157	-0.8514	0.0082	-0.3619	0.7230	-0.1225	0.8356	0.2394	0.6150
Cpr66D	CG32029	1631400_at	-1.2230	0.0017	-0.9583	0.1828	-1.7160	0.0011	-0.5013	0.4191	-0.2093	0.5731	0.2920	0.3555	-0.0977	0.8692	-0.0615	0.8494	0.0363	0.9118
CG14580	CG14580	1631401_at	0.1175	0.5346	0.0967	0.6128	0.0380	0.8643	-0.0969	0.8913	0.0087	0.9809	0.1056	0.6697	0.2310	0.7893	0.1532	0.7142	-0.0778	0.8755
Hr39	Hormone receptor	1631402_s_at	2.9739	0.0015	1.8930	0.0145	1.7438	0.0032	-0.0858	0.9598	0.9994	0.0324	1.0852	0.0146	-0.1424	0.9589	-0.1705	0.8821	-0.0281	0.9837
CG14445	CG14445	1631403_at	-0.0892	0.5216	0.1769	0.3409	0.3134	0.0477	-0.0774	0.8794	-0.1820	0.3015	-0.1046	0.5397	-0.0683	0.9029	-0.0212	0.9492	0.0470	0.8602
eIF3-S10	eIF3-S10	1631404_at	0.0579	0.8503	0.0830	0.8637	0.4321	0.0381	0.1654	0.6006	-0.1468	0.3662	-0.3122	0.0351	-0.2148	0.8928	-0.0403	0.9670	0.1745	0.8129
CG9508	CG9508	1631405_at	2.9122	0.0039	1.6678	0.0859	2.9539	0.0004	0.8781	0.3963	0.4808	0.4243	-0.3973	0.4728	-0.4286	0.8589	-0.6883	0.4903	-0.2597	0.8326
CG11414	CG11414	1631406_at	-0.0429	0.8422	-0.2650	0.0616	-0.3020	0.0702	0.2674	0.4064	0.3445	0.0587	0.0771	0.6806	0.1738	0.7770	0.1440	0.6139	-0.0298	0.9361
APC7	Anaphase Promot	1631407_a_at	0.3769	0.4649	-1.3070	0.1149	-1.1681	0.0950	0.0080	0.9953	1.4953	0.0007	1.4872	0.0005	-0.3651	0.9467	-0.0250	0.9942	0.3401	0.8800
SoxN	soxneuro	1631408_at	0.4451	0.6896	1.3257	0.1200	0.7050	0.0606	-0.6006	0.4846	-1.2760	0.0131	-0.6753	0.0992	0.1144	0.9845	-0.2364	0.9064	-0.3508	0.8320
---	---	1631409_at	0.0055	0.9834	-0.1674	0.3582	-0.0121	0.9607	0.0973	0.9042	0.3159	0.2293	0.2186	0.3680	-0.0022	0.9993	0.0907	0.8302	0.0928	0.8151
---	---	1631410_at	0.1577	0.3305	0.2768	0.1389	0.1203	0.4214	-0.0872	0.8572	-0.1273	0.4845	-0.0400	0.8415	0.1520	0.7893	0.0388	0.9175	-0.1132	0.6679
CG14944	CG14944	1631411_at	0.1203	0.4055	0.0115	0.9122	0.0967	0.5755	-0.0201	0.9755	-0.2584	0.1035	-0.2384	0.0936	0.1833	0.7154	-0.0491	0.8684	-0.2323	0.2841
Pat1	Protein interacting	1631412_at	0.6290	0.0059	0.7052	0.0457	1.1962	0.0022	-0.1116	0.9011	-0.3239	0.2683	-0.2122	0.4384	-0.3471	0.6749	-0.0842	0.8582	0.2629	0.4552
CG17267	CG17267	1631413_at	0.0212	0.9226	0.0528	0.8051	-0.0388	0.9051	-0.0607	0.9386	-0.0217	0.9503	0.0390	0.8871	0.1938	0.8421	0.0485	0.9402	-0.1453	0.7516
CG11313	CG11313	1631414_at	0.3072	0.2098	0.2946	0.2640	0.5275	0.1251	-0.1176	0.9154	-0.3044	0.3914	-0.1868	0.5898	0.0718	0.9071	0.0086	0.9841	-0.0632	0.8177
CG30414	CG30414	1631415_at	0.0500	0.7510	-0.0963	0.4927	0.0507	0.7958	0.0620	0.9260	0.0451	0.8652	-0.0169	0.9473	-0.0898	0.8869	-0.1384	0.5804	-0.0486	0.8800
---	---	1631416_at	-0.0147	0.9634	0.0575	0.5851	-0.0871	0.7081	0.0374	0.9627	-0.2030	0.3504	-0.2404	0.2080	0.0820	0.9137	-0.1021	0.7424	-0.1841	0.5021
CG6282	CG6282	1631417_s_at	-0.2331	0.2201	-0.2031	0.3751	0.0555	0.7600	-0.0453	0.9649	-0.0173	0.9676	0.0280	0.9341	-0.2569	0.6898	0.0020	0.9982	0.2589	0.3360
CG31394	CG31394	1631418_at	0.3607	0.0921	0.0633	0.6186	-0.1678	0.3871	-0.0796	0.8554	0.2244	0.1529	0.3040	0.0378	0.0536	0.9467	0.0293	0.9404	-0.0242	0.9433
Yp3	yolk protein	1631419_at	1.1867	0.0230	0.6345	0.2098	1.4253	0.0036	0.3477	0.7401	0.3849	0.4196	0.0371	0.9494	-0.4285	0.7230	-0.2566	0.6483	0.1719	0.7782
CG9509	CG9509	1631420_at	-0.3001	0.8748	0.0048	0.9936	-0.4459	0.1799	-0.1825	0.9673	-1.7222	0.1397	-1.5397	0.1399	0.3204	0.9421	-1.2643	0.3433	-1.5847	0.2622
---	---	1631421_at	0.2810	0.1575	0.1690	0.3312	0.0209	0.9206	-0.0111	0.9903	-0.0072	0.9809	0.0039	0.9875	0.1543	0.8425	0.0044	0.9950	-0.1500	0.6653
---	---	1631422_at	0.3674	0.0578	0.0890	0.5183	0.2537	0.1971	0.0556	0.9186	0.2403	0.1505	0.1847	0.2190	-0.1985	0.7726	-0.1842	0.5543	0.0143	0.9761
---	---	1631423_at	0.2065	0.1388	0.2376	0.2703	0.0816	0.7281	0.0513	0.9384	0.0834	0.7113	0.0321	0.8907	0.1399	0.8882	0.1184	0.8053	-0.0214	0.9703
---	---	1631424_at	-0.0315	0.8595	-0.0015	0.9905	0.2148	0.2443	0.0301	0.9598	0.1008	0.5480	0.0708	0.6619	-0.2145	0.7644	0.0727	0.8632	0.2872	0.3559
CG9231	CG9231	1631425_at	-0.5899	0.0577	-0.0494	0.8459	-0.2154	0.2060	-0.1874	0.6338	-0.7308	0.0033	-0.5434	0.0076	0.0377	0.9816	-0.1912	0.6414	-0.2290	0.5686
CG32412	CG32412	1631426_at	-1.0278	0.0041	-1.0244	0.0071	-0.4012	0.1156	0.2874	0.4643	0.2398	0.2609	-0.0476	0.8468	-0.2253	0.7893	0.3451	0.3259	0.5705	0.1409
synj	synaptojanin	1631427_s_at	-0.3039	0.4873	-0.6047	0.2516	-0.6443	0.0048	-0.3021	0.6953	-0.1280	0.7640	0.1742	0.6318	-0.1700	0.8967	-0.3400	0.4976	-0.1699	0.7647
---	---	1631428_s_at	0.0717	0.6564	0.2104	0.2117	0.1941	0.2613	-0.0532	0.9133	-0.2733	0.0780	-0.2201	0.1091	0.1524	0.8157	0.0769	0.8303	-0.0756	0.8237
---	---	1631429_at	0.0262	0.9250	-0.0347	0.7793	0.1363	0.3787	0.0265	0.9774	0.0212	0.9505	-0.0053	0.9852	-0.0993	0.9112	0.0221	0.9658	0.1215	0.7391
Cdc27	makos	1631430_at	1.1314	0.0876	0.4332	0.1074	-0.0339	0.9393	0.5687	0.1332	1.4608	0.0004	0.8920	0.0017	0.9046	0.6749	0.7988	0.3496	-0.1058	0.9316
CG33232	villin-like	1631431_at	-0.0179	0.9204	-0.0326	0.8153	0.0449	0.8610	-0.1023	0.9017	-0.0698	0.8403	0.0325	0.9214	-0.0064	0.9943	-0.0464	0.8787	-0.0399	0.8915
CG15786	CG15786	1631432_at	0.2590	0.1663	0.0807	0.5532	0.0328	0.8845	-0.1884	0.6954	0.0031	0.9922	0.1915	0.3474	-0.0442	0.9530	0.0203	0.9557	0.0645	0.8177
CG3335	CG3335	1631433_at	0.4995	0.0419	-0.3098	0.6349	0.2495	0.2058	0.3417	0.3953	0.8413	0.0036	0.4996	0.0216	-0.2887	0.8650	0.0208	0.9884	0.3095	0.6772
mRpS21	mitochondrial ribo	1631434_at	-0.1693	0.3164	0.5550	0.2603	0.2517	0.3140	0.0488	0.9649	-0.4611	0.1169	-0.5098	0.0576	0.2487	0.8400	0.2727	0.6166	0.0240	0.9766
CG7988	CG7988	1631435_at	0.1391	0.6961	0.1250	0.5709	0.5238	0.0189	0.0425	0.9651	-0.5880	0.0309	-0.6305	0.0146	-0.2961	0.7644	-0.4163	0.3066	-0.1202	0.8178
Mst98Cb	Male-specific RNA	1631436_at	0.0720	0.7386	0.1017	0.5334	0.1280	0.3981	-0.0746	0.8903	-0.2107	0.2463	-0.1360	0.4232	0.0633	0.9144	0.0376	0.9058	-0.0257	0.9291
CG30222	CG30222	1631437_at	0.1743	0.2657	-0.0514	0.8601	0.0936	0.6685	0.0541	0.9171	0.0332	0.8768	-0.0209	0.9150	-0.0641	0.9589	-0.0897	0.8556	-0.0256	0.9614
Or45a	Odorant receptor	1631438_a_at	0.0116	0.9487	-0.1889	0.1391	0.3181	0.1792	0.3144	0.2511	0.2417	0.1373	-0.0727	0.6688	-0.1155	0.8999	0.0586	0.9096	0.1742	0.6328
Odc2	Ornithine decarbo	1631439_at	0.8202	0.0298	0.9255	0.0355	1.7285	0.0019	-0.2102	0.5680	-0.1753	0.3628	0.0349	0.8761	-0.9880	0.4603	-0.3585	0.6280	0.6295	0.3712
CG14111	CG14111	1631440_at	0.3851	0.0354	0.1623	0.5845	0.2765	0.3083	0.1767	0.6327	0.0452	0.8472	-0.1314	0.4407	0.1846	0.8427	0.0181	0.9792	-0.1665	0.6963
CG7447	N-like	1631441_s_at	0.3415	0.0634	0.0471	0.8102	0.8117	0.0019	0.2395	0.4860	0.6425	0.0047	0.4029	0.0223	-0.3545	0.6898	0.2622	0.4825	0.6167	0.1240
Hex-t2	Hexokinase	1631442_at	0.1969	0.2153	0.0214	0.8590	-0.0554	0.7621	-0.0348	0.9451	0.0827	0.6007	0.1175	0.3793	0.0757	0.9112	0.0494	0.8877	-0.0263	0.9379
CG16815	CG16815	1631443_at	0.0043	0.9861	0.2057	0.3704	0.2189	0.1371	-0.0262	0.9639	-0.3261	0.0399	-0.2999	0.0358	0.0520	0.9653	-0.0343	0.9476	-0.0863	0.8307
Or35a	Odorant receptor	1631444_a_at	0.1069	0.6320	0.0293	0.7996	0.0381	0.8187	0.1949	0.5832	0.1064	0.5887	-0.0885	0.6321	0.0424	0.9545	-0.0808	0.7562	-0.1232	0.5995
CG32106	CG32106	1631445_at	-0.0445	0.8004	0.2129	0.4625	0.0221	0.9109	-0.1081	0.8671	-0.1816	0.4431	-0.0734	0.7708	0.1914	0.8236	0.1942	0.6119	0.0028	0.9965

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG10531	CG10531	1631446_at	-0.2390	0.1803	-0.2508	0.3662	-0.3549	0.2527	0.1215	0.8738	0.0679	0.8437	-0.0536	0.8626	-0.0726	0.9239	-0.1059	0.7149	-0.0334	0.9237
CG10307	CG10307	1631447_at	-1.9813	0.0042	-2.0637	0.0093	-2.6400	0.0000	0.0572	0.9345	-0.2512	0.2170	-0.3084	0.0910	0.4529	0.7387	-0.4577	0.4246	-0.9106	0.1410
Wnt10	Wnt10	1631448_at	-0.0425	0.8102	-0.0358	0.7193	-0.0716	0.6736	0.0109	0.9883	-0.0045	0.9867	-0.0154	0.9435	-0.0907	0.8901	-0.0279	0.9445	0.0628	0.8413
---	---	1631449_at	0.1103	0.5650	0.0016	0.9901	-0.0560	0.8024	-0.1408	0.7341	-0.0148	0.9567	0.1260	0.4724	-0.1319	0.8379	-0.1737	0.5283	-0.0418	0.9095
CG14860	CG14860	1631450_at	-0.4481	0.0341	0.0034	0.9818	-0.2654	0.1098	-0.0991	0.8196	-0.2020	0.2353	-0.1029	0.5355	-0.0290	0.9751	0.1417	0.5542	0.1707	0.4710
CG32181	CG32181	1631451_at	0.0113	0.9542	-0.1074	0.5406	0.1200	0.5576	0.0696	0.9311	0.1692	0.4948	0.0996	0.6875	-0.0190	0.9862	0.0273	0.9473	0.0464	0.8941
CG8665	CG8665	1631452_at	1.4015	0.0006	0.2563	0.4606	1.3917	0.0009	0.2571	0.5756	0.2187	0.3624	-0.0384	0.8915	-0.8348	0.3660	-0.9950	0.0637	-0.1603	0.7838
---	---	1631453_at	0.0597	0.7713	-0.0111	0.9273	0.0412	0.8345	-0.1501	0.7539	-0.1434	0.5120	0.0067	0.9798	-0.0029	0.9984	-0.0488	0.9129	-0.0459	0.9089
pio	PioPio	1631454_a_at	-1.9641	0.0045	-0.5368	0.1489	-1.4710	0.0042	-0.0575	0.9744	-0.8562	0.0533	-0.7986	0.0452	0.3004	0.8134	-0.0103	0.9935	-0.3107	0.5792
CG30493	CG30493	1631455_at	0.2315	0.3254	-0.1779	0.2825	0.1213	0.4881	0.0634	0.9045	0.3144	0.0701	0.2510	0.1016	-0.1546	0.8395	-0.0478	0.9225	0.1068	0.7739
CG6812	CG6812	1631456_at	-0.7044	0.0667	-0.2872	0.5845	-0.4246	0.0742	-0.7250	0.0840	-0.6478	0.0159	0.0772	0.7750	-0.6202	0.6898	-0.3106	0.6520	0.3096	0.6498
CG13462 /// CG3020	CG13462 /// CG3(	1631457_at	-0.0680	0.6547	0.1518	0.4179	-0.0596	0.7125	-0.1614	0.7605	-0.3025	0.1819	-0.1411	0.5244	0.0774	0.9095	-0.0005	0.9998	-0.0778	0.7873
CG32319	CG32319	1631458_at	0.0660	0.7360	0.1127	0.5902	0.0583	0.8142	-0.0198	0.9727	0.0097	0.9647	0.0296	0.8599	0.0479	0.9721	0.0007	0.9998	-0.0472	0.9212
---	---	1631459_at	0.2703	0.0810	0.1202	0.4209	-0.1465	0.4820	-0.0061	0.9951	0.0474	0.8539	0.0536	0.8109	0.1658	0.7506	-0.0040	0.9935	-0.1698	0.4631
Pof	Painting of fourth	1631460_s_at	0.0593	0.7977	0.3925	0.1825	0.2801	0.2834	-0.1046	0.8796	0.1061	0.6965	0.2107	0.3296	0.0460	0.9742	0.3210	0.3620	0.2750	0.4578
CG32061	CG32061	1631461_at	0.0362	0.8892	-0.2541	0.1998	0.1079	0.5226	0.1630	0.6738	0.1655	0.3757	0.0025	0.9913	-0.0577	0.9514	-0.0526	0.9019	0.0051	0.9920
---	---	1631462_at	-0.0431	0.7849	0.0317	0.7620	-0.0301	0.8618	0.0968	0.7850	0.0859	0.5893	-0.0109	0.9534	0.1300	0.9011	0.2340	0.5613	0.1040	0.8299
bcd	Bicoid	1631463_a_at	0.4072	0.4582	-1.6037	0.0416	-1.9298	0.1032	-1.0686	0.1162	1.8992	0.0014	2.9678	0.0002	-0.8258	0.8689	-0.0132	0.9988	0.8126	0.7154
CG7168	CG7168	1631464_at	-0.8485	0.0078	-0.3754	0.1765	-0.1961	0.3637	0.1223	0.8189	-0.3047	0.1435	-0.4270	0.0299	0.0006	0.9999	0.2042	0.6143	0.2036	0.6168
CG3919	CG3919	1631465_at	-0.9076	0.0030	0.1411	0.7950	-0.3703	0.1679	0.0007	0.9994	0.0200	0.9531	0.0193	0.9473	0.5085	0.7317	0.9428	0.1374	0.4343	0.5120
CG1523	CG1523	1631466_at	-0.0179	0.9401	0.0305	0.7585	0.4154	0.0278	0.2464	0.4102	0.0847	0.6502	-0.1616	0.2809	-0.0147	0.9913	0.2184	0.4391	0.2331	0.4177
CG15208	CG15208	1631467_at	0.1608	0.5044	0.0847	0.5684	0.2045	0.2900	0.0095	0.9937	0.0903	0.7658	0.0807	0.7725	-0.1990	0.7506	-0.0185	0.9664	0.1805	0.5187
tim	TIMELESS	1631468_a_at	0.8434	0.3909	-0.1303	0.7647	0.4403	0.1508	-0.7856	0.2156	-0.0962	0.8452	0.6894	0.0482	-1.5016	0.6389	-1.1535	0.3685	0.3481	0.8323
CG15459	CG15459	1631469_at	0.1244	0.5202	0.1473	0.2425	0.1567	0.4711	-0.1059	0.8111	-0.2432	0.1641	-0.1373	0.4014	0.0933	0.8999	0.1321	0.6595	0.0389	0.9176
CG12313	CG12313	1631470_at	0.1735	0.3830	-0.0035	0.9771	0.0469	0.8538	0.1897	0.7526	0.1585	0.5731	-0.0312	0.9225	0.1058	0.8668	0.0100	0.9834	-0.0958	0.7411
CG14889	CG14889	1631471_at	-0.1798	0.3801	-0.1910	0.1073	-0.2622	0.1422	-0.0562	0.9255	-0.0165	0.9517	0.0397	0.8501	-0.0082	0.9924	-0.0747	0.7529	-0.0665	0.7774
Cht4	Chitinase 4	1631472_at	0.3866	0.1573	-0.0044	0.9810	-0.2669	0.2612	-0.2751	0.6457	0.0270	0.9508	0.3021	0.2461	0.1369	0.8825	-0.1637	0.6768	-0.3006	0.4042
CG33169	CG33169	1631473_at	-0.3373	0.2529	0.0406	0.6871	-0.3508	0.2306	-0.5317	0.1590	-0.6756	0.0100	-0.1439	0.5155	0.0735	0.9514	0.0344	0.9528	-0.0391	0.9382
l(2)k05713	mitochondrial GPI	1631474_s_at	-0.4956	0.3487	0.1464	0.8925	1.3014	0.0013	0.6468	0.0976	-0.4469	0.0564	-1.0937	0.0008	-0.5181	0.8541	0.2755	0.8658	0.7937	0.5112
AttD	attacin	1631475_at	1.3793	0.5424	-3.7784	0.1609	-3.0724	0.0496	0.8125	0.6886	5.8219	0.0004	5.0094	0.0005	0.0527	0.9976	0.7212	0.9047	0.6685	0.9023
---	---	1631476_at	-0.0623	0.8042	-0.1180	0.3013	-0.1629	0.3995	-0.0562	0.9351	-0.0631	0.8007	-0.0069	0.9785	-0.0369	0.9732	-0.1188	0.6969	-0.0819	0.8044
Sdic1	Sdic:CG9580	1631477_a_at	0.0397	0.8034	0.0603	0.6130	0.1058	0.5576	0.0325	0.9538	-0.0108	0.9632	-0.0432	0.8020	0.1048	0.8283	0.1272	0.5473	0.0223	0.9371
---	---	1631478_at	-0.0060	0.9883	-0.1123	0.3101	-0.0395	0.8973	-0.0737	0.9088	-0.0204	0.9465	0.0532	0.8188	-0.0585	0.9499	-0.0877	0.8053	-0.0292	0.9401
l(3)73Ah	lethal(3)73Ah	1631479_at	0.4993	0.0837	0.8829	0.1035	0.9540	0.0011	0.1739	0.6699	-0.0969	0.6577	-0.2708	0.1167	0.1210	0.9462	0.3037	0.6011	0.1827	0.7760
CG2652	CG2652	1631480_at	0.0609	0.8009	0.3699	0.0703	0.3468	0.1863	-0.0894	0.9307	-0.1585	0.6377	-0.0692	0.8419	-0.0364	0.9677	-0.0433	0.9058	-0.0069	0.9852
Kr-h1	Krueppel-homolo(	1631481_a_at	-0.8299	0.4888	-0.1355	0.2949	-0.4691	0.1682	-0.4248	0.7138	0.4665	0.3884	0.8913	0.0625	-0.2369	0.9604	1.0386	0.4391	1.2755	0.3564
---	---	1631482_s_at	0.0215	0.9067	0.0454	0.8254	0.2364	0.2210	-0.0224	0.9753	0.1568	0.3839	0.1792	0.2578	-0.1092	0.8653	-0.0288	0.9451	0.0804	0.7951
CG11257	CG11257	1631483_at	-0.3632	0.1253	-0.1288	0.5373	-0.2044	0.1641	-0.2088	0.6986	-0.0048	0.9898	0.2040	0.3725	-0.0012	0.9998	0.1929	0.6469	0.1941	0.6414
CG15475	CG15475	1631484_at	0.1897	0.2100	0.0206	0.8560	0.0480	0.7704	0.0093	0.9931	0.0884	0.7132	0.0791	0.7206	0.0038	0.9967	-0.0605	0.8500	-0.0643	0.8259
ec	echinus	1631485_a_at	0.0420	0.8275	0.0081	0.9654	0.1916	0.2714	-0.0074	0.9941	-0.0139	0.9650	-0.0065	0.9799	-0.0618	0.9460	0.0214	0.9627	0.0831	0.8054
p38c	p38c	1631486_at	1.6489	0.0507	0.5318	0.5512	2.2460	0.0037	1.3543	0.0898	1.4747	0.0074	0.1204	0.8194	-0.2936	0.9441	0.3436	0.8397	0.6372	0.6398
CG32626	CG32626	1631487_s_at	0.0361	0.9059	-0.0934	0.5478	0.0083	0.9818	0.2648	0.6480	-0.0144	0.9735	-0.2792	0.2699	0.1259	0.8999	-0.1782	0.6596	-0.3041	0.4167
CG15618 /// DmirCG15618	CG15618	1631488_at	0.2030	0.8254	-0.0934	0.9491	-0.2276	0.4505	0.3676	0.2769	0.6859	0.0051	0.3183	0.0756	0.7658	0.8270	0.4364	0.8180	-0.3294	0.8641
---	---	1631489_at	0.1209	0.5010	0.0443	0.7696	0.3134	0.1506	0.0822	0.8794	0.1474	0.4435	0.0652	0.7447	-0.1432	0.8454	0.0221	0.9647	0.1652	0.6117
---	---	1631490_at	0.0683	0.7618	0.0441	0.7886	-0.0358	0.8257	0.0813	0.8836	-0.0066	0.9816	-0.0879	0.6528	0.0693	0.8825	-0.0312	0.9088	-0.1004	0.6016
CG9335	CG9335	1631491_at	-0.3869	0.1973	-0.2451	0.4244	-0.5551	0.0177	-0.2696	0.4908	-0.1832	0.3915	0.0864	0.6934	-0.0397	0.9831	-0.1803	0.7142	-0.1406	0.7838
CG17173	CG17173	1631492_at	-0.0076	0.9703	-0.3113	0.3457	-0.0779	0.6402	0.1704	0.7444	0.2081	0.3750	0.0377	0.8909	0.0327	0.9672	-0.0440	0.8850	-0.0767	0.7513
cathD	cathepsin D	1631493_at	-0.4440	0.0256	-0.2277	0.2287	-0.1004	0.5610	0.0080	0.9910	-0.5277	0.0051	-0.5356	0.0028	-0.1889	0.7588	-0.4000	0.1248	-0.2111	0.4243
---	---	1631494_at	0.1036	0.6690	0.1034	0.6226	0.0749	0.7500	-0.0512	0.9285	0.0078	0.9762	0.0591	0.7457	0.0461	0.9589	0.0562	0.8764	0.0101	0.9810
---	---	1631495_at	0.0920	0.6701	0.0747	0.5885	0.0396	0.8909	-0.0645	0.9311	-0.1133	0.6474	-0.0488	0.8483	-0.0661	0.9444	-0.0424	0.9257	0.0238	0.9526
CG4563	CG4563	1631496_at	0.0627	0.7938	0.0169	0.8797	0.2709	0.2194	-0.0816	0.9023	-0.2613	0.2253	-0.1797	0.3669	0.0361	0.9734	0.0807	0.8157	0.0446	0.9033
Tsp42Eo	tetraspanin 42E	1631497_at	-1.7191	0.0561	-2.2349	0.0444	-2.6051	0.0001	-0.2849	0.5839	-0.6754	0.0190	-0.3905	0.0989	0.2285	0.9589	-1.0809	0.3739	-1.3094	0.3026
fru	fru-satori	1631498_a_at	-0.9923	0.0256	-1.6423	0.0126	-1.6409	0.0002	0.0107	0.9910	0.6591	0.0072	0.6484	0.0046	-0.0412	0.9787	-0.3294	0.3490	-0.2882	0.4368
sws	swiss cheese	1631499_a_at	0.1566	0.7529	0.2690	0.5874	0.5180	0.0565	0.0147	0.9857	-0.1652	0.3950	-0.1799	0.2942	-0.2181	0.9295	0.0014	0.9998	0.2195	0.8255
mod(mdg4)	Modifier67.2	1631500_at	-0.7687	0.0267	-0.4959	0.1506	-0.5187	0.0209	0.1926	0.6718	-0.2591	0.2283	-0.4517	0.0270	0.2343	0.8013	-0.0458	0.9411	-0.2801	0.4852
CG40157	CG40157	1631501_at	0.0787	0.6780	-0.0219	0.8607	-0.0027	0.9903	-0.1149	0.7850	-0.0018	0.9945	0.1132	0.4974	-0.0904	0.8802	-0.0212	0.9543	0.0693	0.8044
wor	warniu	1631502_at	-0.1778	0.5934	-0.0243	0.9230	-0.0877	0.6877	-0.1922	0.8085	-0.5917	0.0644	-0.3995	0.1558	-0.1325	0.8978	-0.2335	0.5620	-0.1010	0.8354
---	---	1631503_at	-0.0857	0.7214	0.0577	0.5316	0.0787	0.7684	0.0343	0.9641	-0.2340	0.2491	-0.2684	0.1367	-0.0425	0.9772	-0.0930	0.8445	-0.0505	0.9169
---	---	1631504_at	0.1579	0.3334	-0.0498	0.6431	-0.0035	0.9859	0.0046	0.9951	0.0440	0.8108	0.0395	0.8140	-0.0676	0.9309	-0.0931	0.7584	-0.0255	0.9431
Sr-CIII	Scavenger Recep	1631505_at	0.0084	0.9784	0.1884	0.2676	0.1078	0.6664	-0.0646	0.9308	-0.0430	0.8831	0.0216	0.9363	-0.0904	0.8686	-0.0061	0.9900	0.0843	0.7338
---	---	1631506_at	-0.0131	0.9551	0.1670	0.3215	0.1466	0.3473	-0.0120	0.9904	-0.1304	0.5877	-0.1184	0.5925	-0.0712	0.9101	-0.1308	0.5825	-0.0597	0.8334
---	---	1631507_at	-0.0009	0.9980	0.2835	0.2916	0.0092	0.9660	-0.1827	0.7104	-0.0977	0.7053	0.0850	0.7212	0.1557	0.8875	0.2101	0.6382	0.0544	0.9233
Gr39a	Gustatory recepto	1631508_a_at	0.0840	0.6201	-0.0066	0.9529	0.0484	0.7950	0.0918	0.8084	0.1299	0.4019	0.0380	0.8244	-0.0411	0.9526	0.0048	0.9924	0.0458	0.8674
lt	light	1631509_a_at	-0.5659	0.0124	-0.4758	0.1787	-0.5683	0.0059	-0.2462	0.4857	-0.2949	0.1161	-0.0487	0.8195	-0.1271	0.8851	-0.1383	0.7161	-0.0112	0.9842
---	---	1631510_at	0.3043	0.1619	0.2842	0.1939	0.2917	0.0775	0.0572	0.9424	-0.2537	0.2625	-0.3108	0.1214	0.3154	0.4128	-0.0453	0.8785	-0.3608	0.1166
vas	vasa	1631511_at	0.0625	0.8450	0.0696	0.5406	-0.0307	0.9255	0.0434	0.9603	0.0581	0.8450	0.0147	0.9594	0.0083	0.9952	-0.0376	0.9432	-0.0459	0.9194
CG9377	CG9377	1631512_at	0.1851	0.4008	0.3156	0.0588	0.1922	0.4000	-0.0586	0.8908	-0.1807	0.2066	-0.1221	0.3545	0.1670	0.8122	0.0130	0.9810	-0.1541	0.6238
CG30000	CG30000	1631513_at	1.0946	0.0144	0.3612	0.1675	0.8481	0.0014	0.1016	0.9120	0.2601	0.3847	0.1585	0.5877	-0.3085	0.7436	-0.4017	0.3000	-0.0933	0.8577
CG6244	CG6244	1631514_at	0.0948	0.5727	0.2001	0.1898	0.1569	0.4631	-0.0785	0.8933	-0.1740	0.3832	-0.0955	0.6318	-0.0155	0.9869	0.0061	0.9900	0.0215	0.9433

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11877	CG11877	1631515_at	0.2080	0.5350	-0.0711	0.8837	-0.0462	0.8014	0.0561	0.9059	0.1536	0.3225	0.0974	0.5099	-0.0251	0.9928	-0.0974	0.9214	-0.0723	0.9347
skd	blind spot	1631516_s_at	-0.2725	0.6838	-0.1968	0.4675	-0.2235	0.6122	0.1428	0.8578	0.3112	0.2774	0.1683	0.5471	0.0030	0.9998	0.2308	0.8698	0.2278	0.8618
Mcm7	Minichromosome	1631517_at	0.5151	0.6300	-0.9817	0.4650	-0.7575	0.2609	-0.0982	0.9765	1.1982	0.1437	1.2963	0.0796	-0.3346	0.9611	-0.2976	0.9208	0.0371	0.9913
btsz	bitesize	1631518_at	0.0547	0.8102	0.3356	0.1896	0.0166	0.9532	-0.0368	0.9677	-0.2289	0.3469	-0.1921	0.3865	0.1971	0.8097	-0.0247	0.9648	-0.2218	0.5361
CG9143 /// DsecCG9143 ///	CG9143	1631519_at	0.5903	0.0768	0.4417	0.2606	1.0289	0.0195	0.4276	0.4386	0.7183	0.0254	0.2907	0.2874	-0.1982	0.9305	0.7603	0.2840	0.9585	0.2101
CG1845	CG1845	1631520_at	-0.7820	0.2453	-0.2714	0.8525	-0.2840	0.3651	0.0428	0.9675	-0.3916	0.1544	-0.4344	0.0800	-0.1999	0.9774	-0.0130	0.9981	0.1868	0.9360
CG33469	CG33469	1631521_at	0.1097	0.8041	-0.3265	0.5112	-0.2465	0.5755	0.1711	0.7669	0.8504	0.0050	0.6793	0.0082	-0.0826	0.9829	0.3253	0.7584	0.4079	0.6736
CG15213	CG15213	1631522_x_at	0.4320	0.5148	0.1670	0.3576	0.2142	0.4453	0.3189	0.5255	-0.0345	0.9252	-0.3535	0.1308	0.1823	0.9411	-0.2493	0.7961	-0.4316	0.5991
CG10824	CG10824	1631523_at	1.2224	0.0915	0.9112	0.3537	1.1550	0.0097	0.5406	0.5516	-0.6527	0.1629	-1.1934	0.0121	0.2124	0.9611	-0.9606	0.4285	-1.1730	0.3485
hth	dorsotonals	1631524_a_at	-0.7077	0.1062	-0.6167	0.1811	-1.5778	0.0018	0.0037	0.9973	0.5615	0.0703	0.5578	0.0476	0.9581	0.5142	0.6864	0.3185	-0.2717	0.7391
l(2)k10201	lethal (2) k10201	1631525_at	-0.5189	0.0274	-0.5584	0.0486	-0.4483	0.0507	0.3991	0.2753	0.2887	0.1778	-0.1103	0.6130	0.2246	0.7475	0.3624	0.2061	0.1378	0.6686
CG31284	CG31284	1631526_s_at	-3.7305	0.0005	-3.1050	0.0301	-3.9228	0.0003	-0.6333	0.3828	-1.3462	0.0062	-0.7129	0.0558	0.0451	0.9928	-0.6303	0.6118	-0.6754	0.5856
CG5642	CG5642	1631527_at	0.0263	0.8818	-0.0731	0.7886	0.0162	0.9303	0.1140	0.7293	-0.0336	0.8626	-0.1476	0.2671	0.0036	0.9978	-0.1485	0.6471	-0.1521	0.6366
---	---	1631528_at	-0.1711	0.5149	-0.1908	0.3956	0.0542	0.7939	0.2493	0.4904	0.2686	0.1612	0.0193	0.9360	0.0594	0.9640	0.1711	0.6607	0.1118	0.7940
Ser12	Serine protease 1	1631529_at	0.3741	0.3220	0.0580	0.5653	0.4956	0.0087	0.3135	0.2289	-0.0139	0.9518	-0.3274	0.0263	0.0507	0.9772	0.3178	0.4475	0.2671	0.5450
---	---	1631530_at	0.1730	0.4109	0.0146	0.9095	0.1480	0.4270	-0.0226	0.9803	-0.1789	0.4286	-0.1563	0.4479	-0.1749	0.7953	-0.1922	0.5123	-0.0173	0.9678
Srp72	Srp72	1631531_at	0.8031	0.0148	0.5363	0.1792	0.3743	0.1181	0.2645	0.5008	0.8434	0.0025	0.5789	0.0079	0.5321	0.6749	0.7035	0.1718	0.1714	0.7853
CG5282	CG5282	1631532_at	-0.1829	0.4498	-0.0127	0.9476	0.0729	0.7239	-0.0939	0.9136	-0.4321	0.1123	-0.3382	0.1645	-0.0676	0.9342	-0.0573	0.8787	0.0103	0.9816
Cyp6a22	Cytochrome P450	1631533_at	-0.3457	0.0891	-0.8962	0.2343	0.0559	0.7942	-0.3462	0.3863	-1.5207	0.0004	-1.1745	0.0005	-1.0898	0.5619	-1.9842	0.0415	-0.8943	0.3057
sfl	diet-wingless	1631534_at	-0.7484	0.3369	1.1680	0.3100	-0.0776	0.8668	-0.8464	0.4757	-0.7479	0.2414	0.0985	0.8977	0.2551	0.9514	1.0405	0.3909	0.7854	0.5456
Vinc	vinculin	1631535_at	-1.1507	0.1377	-1.2368	0.2773	-1.9383	0.0016	-0.0571	0.9298	0.5803	0.0086	0.6374	0.0034	0.7466	0.8363	0.5081	0.7829	-0.2385	0.9080
Dip3	Dorsal interacting	1631536_at	0.0253	0.9040	-0.3423	0.1463	-0.2418	0.3089	0.1356	0.8066	0.5078	0.0287	0.3722	0.0621	-0.2220	0.8395	0.0002	1.0000	0.2223	0.6476
CG31198	CG31198	1631537_at	-0.5665	0.8550	0.7132	0.1475	-2.6743	0.0014	-3.0183	0.2840	-2.7815	0.0937	0.2369	0.9058	0.2117	0.9862	-1.6174	0.5963	-1.8291	0.5457
---	---	1631538_at	-0.1474	0.3683	-0.0135	0.9588	-0.1403	0.3716	0.0169	0.9777	-0.1009	0.5279	-0.1178	0.3985	-0.0349	0.9643	0.0337	0.9175	0.0686	0.7836
CG14491	CG14491	1631539_at	-0.1322	0.5152	-0.1216	0.3755	0.1785	0.3426	0.1370	0.7202	0.0335	0.8857	-0.1035	0.5402	-0.1957	0.7644	0.0944	0.7787	0.2901	0.3011
CG15344	CG15344	1631540_at	0.0824	0.7600	0.1518	0.5004	0.1426	0.5211	0.0360	0.9774	0.0668	0.8705	0.0308	0.9362	-0.1341	0.8878	0.0946	0.8445	0.2288	0.5437
smt3	smt3	1631541_at	0.2123	0.2975	0.0931	0.5799	-0.1259	0.4278	0.1756	0.6499	0.4470	0.0247	0.2714	0.1032	0.3628	0.5228	0.3382	0.1995	-0.0247	0.9486
dve	defective proventr	1631542_a_at	-2.7105	0.0042	-3.4608	0.0075	-3.1266	0.0008	0.1306	0.9565	0.5331	0.4246	0.4024	0.5207	-0.2926	0.9168	-0.2709	0.8343	0.0217	0.9889
CG17286 /// DmirCG17286	CG17286	1631543_at	0.1493	0.8558	-1.3411	0.2397	-1.5566	0.0112	-0.4423	0.3553	1.2870	0.0013	1.7293	0.0002	-0.2047	0.9717	0.0364	0.9916	0.2410	0.9057
---	---	1631544_at	0.0617	0.7793	-0.0044	0.9883	-0.0241	0.8836	0.0895	0.8315	0.2358	0.1395	0.1464	0.3159	0.0732	0.9168	0.1200	0.6499	0.0467	0.8889
---	---	1631545_at	0.0776	0.6050	0.0026	0.9835	-0.1738	0.2846	-0.1949	0.5461	-0.0377	0.8645	0.1572	0.3003	-0.0437	0.9590	-0.0866	0.7729	-0.0429	0.8993
CG3501	CG3501	1631546_at	-0.0739	0.7663	0.4215	0.0568	0.7121	0.0106	0.2013	0.5680	-0.2025	0.2638	-0.4037	0.0205	0.0186	0.9898	0.3551	0.2518	0.3364	0.3079
---	---	1631547_s_at	0.0905	0.5710	-0.0696	0.5372	-0.2782	0.1601	-0.0770	0.9017	0.1821	0.3783	0.2592	0.1506	0.0547	0.9365	-0.0372	0.9111	-0.0919	0.7060
CG30196	CG30196	1631548_at	-1.8905	0.0062	-0.6974	0.2139	-0.5210	0.0261	-0.1340	0.8796	-1.3955	0.0015	-1.2615	0.0013	-0.0530	0.9816	-0.5518	0.2722	-0.4989	0.3519
CG18278 /// CG30059	CG30059 /// CG1	1631549_s_at	-0.8562	0.0230	-1.8419	0.0081	-1.3294	0.0002	-0.1085	0.8233	-0.1007	0.6323	0.0078	0.9736	-0.5862	0.4017	-1.0641	0.0276	-0.4778	0.2235
CG5874	CG5874	1631550_at	0.1387	0.8038	-0.1914	0.7731	-0.3662	0.2261	-0.0109	0.9935	0.8231	0.0072	0.8341	0.0040	0.1299	0.9657	0.3006	0.7567	0.1706	0.8774
CG40192	CG40192	1631551_at	0.2085	0.5275	-0.0585	0.8207	0.0097	0.9692	0.1062	0.8526	0.0957	0.6865	-0.0105	0.9682	0.0602	0.9734	-0.1915	0.7032	-0.2517	0.5988
pyr	pyramus	1631552_at	-1.0527	0.0104	-2.5130	0.0362	-2.7622	0.0006	0.4298	0.6644	1.5243	0.0068	1.0945	0.0178	0.8216	0.6935	0.2083	0.8615	-0.6133	0.4886
CG30463	CG30463	1631553_at	-0.1110	0.5244	0.0659	0.5990	0.0493	0.8101	-0.1652	0.5744	-0.3141	0.0436	-0.1489	0.2749	-0.0267	0.9816	-0.0169	0.9696	0.0098	0.9835
CG11473 /// DsimCG11473	CG11473	1631554_at	-0.2142	0.6070	-0.1543	0.3885	-0.2883	0.2402	0.0342	0.9777	-0.0245	0.9562	-0.0588	0.8685	0.0422	0.9816	-0.1750	0.7027	-0.2172	0.6192
CG10062	CG10062	1631555_at	-0.0158	0.9955	-0.6321	0.0220	-0.3775	0.2632	-0.2402	0.9497	-0.8053	0.4668	-0.5652	0.5944	-0.7333	0.9132	-1.5705	0.5199	-0.8371	0.7578
Cp18	Shell-18	1631556_at	-0.2215	0.5077	0.3513	0.0543	0.3783	0.0864	0.0360	0.9639	-0.5823	0.0136	-0.6184	0.0062	0.0280	0.9853	-0.0236	0.9670	-0.0516	0.9152
---	---	1631557_at	-0.0026	0.9922	-0.1098	0.4665	0.1446	0.5026	0.0778	0.9186	-0.0214	0.9517	-0.0992	0.6868	-0.0472	0.9467	-0.0891	0.7198	-0.0419	0.8880
Cpr65Ax2	CG18777	1631558_at	0.0364	0.8529	0.2455	0.2938	0.2443	0.1282	0.2839	0.5902	-0.0066	0.9875	-0.2905	0.2252	-0.0035	0.9976	0.0449	0.9110	0.0484	0.8925
---	---	1631559_s_at	0.1535	0.4699	0.2280	0.3634	0.0167	0.9490	-0.1475	0.7845	-0.0182	0.9557	0.1293	0.5545	-0.0198	0.9860	-0.0928	0.7565	-0.0731	0.8154
ferrochelatase	ferrochelatase	1631560_at	0.1284	0.5471	-0.1754	0.4457	-0.3186	0.5342	0.0223	0.9857	0.1522	0.6377	0.1299	0.6640	0.1368	0.9514	-0.1855	0.8340	-0.3223	0.6508
CG4607 /// DsimCG4607	CG4607	1631561_s_at	-0.3588	0.4592	0.5338	0.4732	0.6003	0.0855	-1.1633	0.1641	-2.1681	0.0019	-1.0048	0.0338	-1.1005	0.5765	-1.1062	0.1995	-0.0057	0.9976
Or88a	Odorant receptor	1631562_at	-0.0293	0.8800	-0.1915	0.1648	0.0725	0.8067	0.2950	0.4376	0.2652	0.2006	-0.0298	0.9055	-0.0467	0.9653	0.0443	0.9221	0.0911	0.7938
Gr21a	Gustatory recepto	1631563_at	-0.1083	0.6347	-0.0435	0.8672	0.1714	0.2506	0.1517	0.7790	-0.0782	0.7724	-0.2298	0.2554	-0.0750	0.9246	-0.0664	0.8551	0.0085	0.9847
Usp7	ubiquitin-specific p	1631564_at	-0.4048	0.2390	-0.3091	0.3907	-0.9374	0.0065	-0.1984	0.7764	0.2406	0.4259	0.4390	0.0927	0.4729	0.7241	0.3795	0.5283	-0.0935	0.9072
crc	cryptocephal	1631565_a_at	-0.1663	0.2604	-0.0454	0.7375	0.0540	0.7448	0.0110	0.9865	-0.0795	0.6543	-0.0905	0.5612	-0.0079	0.9928	-0.0011	0.9991	0.0068	0.9842
CG11069 /// DyakCG11069	CG11069	1631566_at	-0.0517	0.8676	-0.6770	0.1855	-0.8485	0.0076	-0.1089	0.8304	0.4866	0.0207	0.5955	0.0052	0.0017	0.9998	-0.1773	0.7779	-0.1790	0.7626
---	---	1631567_at	0.1556	0.3578	-0.0876	0.5393	0.1870	0.4016	0.1591	0.7857	0.0739	0.8031	-0.0852	0.7443	-0.0186	0.9862	0.0004	0.9998	0.0190	0.9558
sdt	anon-fast-evolving	1631568_a_at	-0.7695	0.3164	0.5925	0.4225	-0.4345	0.0589	-0.1818	0.5696	-0.6193	0.0033	-0.4375	0.0096	0.7975	0.7686	0.6717	0.5823	-0.1258	0.9375
Gp150	glycoprotein 150	1631569_s_at	-0.0053	0.9812	-0.0240	0.8618	-0.1736	0.2784	-0.1352	0.6734	-0.2186	0.1474	-0.0834	0.5811	-0.0271	0.9816	-0.3319	0.1868	-0.3049	0.2557
CG13438	CG13438	1631570_at	0.1121	0.6510	0.1096	0.3102	0.0974	0.6033	0.0409	0.9436	0.0250	0.9142	-0.0159	0.9385	0.0368	0.9816	0.1223	0.7741	0.0855	0.8488
spir	spire	1631571_a_at	-0.3713	0.3923	-0.7524	0.2240	0.0051	0.9916	0.3688	0.6673	0.4058	0.3278	0.0369	0.9423	-0.4185	0.8202	-0.0097	0.9954	0.4088	0.6166
CG17380	CG17380	1631572_at	0.0195	0.9055	-0.0983	0.3710	0.0172	0.9353	0.0880	0.8469	0.1739	0.3020	0.0859	0.6107	-0.0026	0.9976	-0.0111	0.9726	-0.0084	0.9777
wun	wunen	1631573_a_at	0.2525	0.3040	1.0797	0.0381	0.8266	0.0257	-0.0409	0.9631	-0.9591	0.0024	-0.9182	0.0017	0.2440	0.8553	-0.0587	0.9477	-0.3027	0.6093
---	---	1631574_at	-0.1271	0.7044	0.0287	0.7699	0.2626	0.2542	-0.0670	0.9247	-0.5437	0.0186	-0.4766	0.0202	0.0219	0.9899	-0.0833	0.8782	-0.1052	0.8247
U2A	lethal(2)43Ee	1631575_at	0.3461	0.1678	0.5242	0.2114	0.6810	0.0335	-0.0346	0.9610	-0.0295	0.9103	0.0050	0.9835	-0.0579	0.9737	0.1700	0.7350	0.2279	0.6231
CG2120	CG2120	1631576_at	0.0810	0.9405	0.0207	0.8523	0.0714	0.7027	-0.1439	0.9491	-0.6631	0.2967	-0.5192	0.3722	-0.1197	0.9653	-0.5933	0.4167	-0.4736	0.5413
CG3760 /// DyakCG3760	anon-fast-evolving	1631577_a_at	0.4866	0.0288	1.2313	0.0067	0.9952	0.0005	0.0557	0.9339	-0.2772	0.1581	-0.3329	0.0627	0.1861	0.7823	0.6562	0.0461	0.4701	0.1362
CG14213	CG14213	1631578_s_at	0.0858	0.5974	0.5718	0.1184	0.5760	0.0207	-0.0943	0.8685	-0.2863	0.1522	-0.1920	0.2915	-0.1109	0.8982	0.1648	0.6368	0.2757	0.4004
fry	Suppressor of GM	1631579_a_at	-0.0352	0.9215	0.0686	0.9173	-0.4307	0.0109	-0.0737	0.8856	0.2719	0.1165	0.3456	0.0335	0.2536	0.8870	0.3160	0.6686	0.0624	0.9471
---	---	1631580_at	0.2988	0.1138	0.1948	0.4574	0.1100	0.6203	-0.0956	0.8671	-0.0375	0.8907	0.0582	0.7976	0.1079	0.8635	-0.0043	0.9941	-0.1121	0.6875
CG18268	CG18268	1631581_at	-1.2807	0.0238	-2.7194	0.0163	-2.5063	0.0034	0.1920	0.9380	1.2339	0.0880	1.0419	0.1056	-0.0395	0.9914	-0.2559	0.8165	-0.2164	0.8416
CG15462	CG15462	1631582_at	-0.0837	0.6838	0.1269	0.5373	0.1596	0.3795	-0.0859	0.8189	-0.0610	0.7214	0.0249	0.8867	-0.0781	0.9421	0.2231	0.5157	0.3012	0.3764
Tsp66A	Tetraspanin 66A	1631583_at	-0.1001	0.6053	-0.1152	0.4606	-0.0944	0.6555	0.0451	0.9517	0.0943	0.6881	0.0492	0.8338	-0.0352	0.9752	0.1032	0.7573	0.1384	0.6461

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31189	CG31189	1631584_at	-1.2055	0.0153	-0.1719	0.4986	-0.9777	0.0190	-0.7123	0.4094	-1.2897	0.0151	-0.5774	0.1767	-0.0342	0.9816	-0.3727	0.2571	-0.3385	0.3332
CG12134 /// DereCG12134	CG12134	1631585_a_at	-0.0367	0.8908	0.0666	0.8964	0.1429	0.4911	0.1663	0.7742	0.2972	0.2194	0.1310	0.5895	0.1866	0.8655	0.4786	0.2593	0.2920	0.5284
CG15250	CG15250	1631586_at	-0.2138	0.3372	-0.3280	0.2185	-0.1424	0.3935	0.0130	0.9880	-0.1727	0.4092	-0.1858	0.3157	-0.1192	0.8882	-0.1858	0.5825	-0.0666	0.8779
---	---	1631587_at	0.0848	0.6606	0.1369	0.2990	0.1001	0.5044	0.0564	0.9098	0.1178	0.4757	0.0614	0.7142	0.1089	0.8461	0.1522	0.5241	0.0433	0.8915
Pgd	6-phosphoglucona	1631588_at	0.7553	0.0188	0.5419	0.3305	1.4634	0.0002	0.6214	0.0785	-0.2374	0.2464	-0.8588	0.0013	-0.3419	0.8331	-0.4072	0.5647	-0.0653	0.9444
---	---	1631589_at	0.0605	0.6705	0.0029	0.9818	0.2161	0.2479	0.0810	0.8485	0.0864	0.6131	0.0055	0.9776	-0.0903	0.8395	-0.0180	0.9508	0.0723	0.7305
---	---	1631590_at	0.4039	0.1715	0.2899	0.2481	0.0861	0.7046	-0.2136	0.7020	-0.1576	0.5693	0.0560	0.8506	-0.0528	0.9619	-0.2232	0.4794	-0.1703	0.6089
CG31935	CG31935	1631591_at	-0.6180	0.1755	0.5675	0.1046	1.0030	0.0004	0.0095	0.9937	-0.9954	0.0021	-1.0050	0.0012	-0.3337	0.8215	0.2593	0.7093	0.5930	0.3485
---	---	1631592_at	0.0877	0.6797	0.1468	0.5499	0.1518	0.3337	-0.1205	0.8164	-0.2766	0.1736	-0.1561	0.4138	0.1451	0.8650	0.0213	0.9701	-0.1238	0.7524
Fer3HCH	Ferritin 3 heavy ch	1631593_at	0.1537	0.3289	0.2617	0.1741	0.3057	0.0973	-0.0705	0.8640	-0.0129	0.9530	0.0576	0.7094	0.0116	0.9914	-0.0453	0.9064	-0.0569	0.8636
Pi3K92E	PI3 kinase	1631594_s_at	0.3889	0.1282	0.1042	0.4632	0.0471	0.8583	0.1948	0.7010	0.5036	0.0372	0.3087	0.1367	0.3021	0.7230	0.1077	0.8218	-0.1945	0.6216
CG13393	CG13393	1631595_at	0.7482	0.0125	1.3220	0.0108	1.6805	0.0001	0.0865	0.8189	-0.0109	0.9604	-0.0974	0.4909	-0.1352	0.8999	0.4986	0.1948	0.6338	0.1390
CG1332	CG1332	1631596_at	0.3481	0.1152	0.7385	0.0408	0.4263	0.0207	-0.1007	0.8299	-0.0127	0.9632	0.0881	0.6315	0.2030	0.8192	0.3541	0.3162	0.1511	0.7130
foxo	forkhead box, sub	1631597_at	-0.0846	0.8918	1.1764	0.0233	0.3778	0.3865	-0.7903	0.1033	-1.9762	0.0003	-1.1859	0.0014	-0.0473	0.9914	-0.7160	0.4466	-0.6686	0.4913
CG3184	CG3184	1631598_at	0.0961	0.6532	-0.1781	0.4022	0.0526	0.8614	0.0879	0.8891	0.4364	0.0447	0.3485	0.0678	-0.1307	0.8875	0.0831	0.8637	0.2138	0.5612
CG31976	CG31976	1631599_at	0.2179	0.4041	0.0160	0.9256	0.1936	0.3425	0.0565	0.9375	0.1303	0.5656	0.0738	0.7452	0.0725	0.9449	0.0913	0.8262	0.0188	0.9692
---	---	1631600_at	-0.1813	0.4140	0.1140	0.5135	0.0195	0.9259	-0.1540	0.7368	-0.1947	0.3462	-0.0407	0.8632	0.0177	0.9829	0.0227	0.9404	0.0050	0.9870
---	---	1631601_at	0.1736	0.3190	0.1112	0.5322	-0.1296	0.6673	0.0744	0.9218	0.2202	0.3447	0.1458	0.5098	0.0774	0.9238	0.1418	0.6209	0.0644	0.8519
CG14676	CG14676	1631602_at	-0.0406	0.8638	0.1762	0.1619	0.0260	0.8832	-0.2875	0.3328	-0.3197	0.0656	-0.0323	0.8736	-0.0395	0.9742	-0.0109	0.9857	0.0286	0.9471
CG34117	CG34117	1631603_at	-0.4776	0.0504	-0.0995	0.6484	-0.2904	0.1095	-0.0397	0.9421	-0.3450	0.0335	-0.3053	0.0349	0.2057	0.8222	-0.0297	0.9623	-0.2353	0.5610
CG9511	CG9511	1631604_at	0.9771	0.0187	0.6482	0.3675	1.2523	0.0040	0.2593	0.6808	-0.3400	0.2483	-0.5993	0.0301	-0.3913	0.8446	-0.7126	0.3760	-0.3213	0.7333
---	---	1631605_at	-0.0186	0.9308	0.1024	0.5247	-0.1498	0.3998	-0.1884	0.6284	-0.2138	0.2669	-0.0254	0.9132	0.0872	0.9273	0.0380	0.9402	-0.0491	0.9095
CG14285	CG14285	1631606_at	-1.4350	0.0095	-0.2597	0.3453	-0.6141	0.0038	-0.1229	0.7753	-0.6676	0.0037	-0.5447	0.0055	-0.0816	0.9555	0.0750	0.9084	0.1566	0.7492
---	---	1631607_at	0.1556	0.4169	-0.0120	0.9515	0.0817	0.6611	0.0452	0.9482	0.0899	0.6879	0.0447	0.8428	-0.0774	0.9095	-0.1259	0.6314	-0.0484	0.8846
CG6954	CG6954	1631608_at	0.1147	0.7142	0.1509	0.4376	-0.3028	0.1471	-0.2460	0.5947	-0.3118	0.1811	-0.0658	0.8033	0.2300	0.8298	-0.3040	0.5121	-0.5340	0.2523
---	---	1631609_at	-0.1156	0.6682	0.0103	0.9362	0.0297	0.9078	-0.0924	0.9213	-0.1255	0.6998	-0.0331	0.9240	0.1292	0.7910	0.0773	0.7541	-0.0519	0.8447
CG31886	CG31886	1631610_at	2.1398	0.0075	0.9970	0.0356	1.0530	0.0002	-0.2003	0.5379	0.2878	0.0911	0.4880	0.0069	-0.4061	0.8099	-0.9171	0.1779	-0.5110	0.4824
obst-H	CG33983	1631611_at	0.1335	0.5437	0.1394	0.2106	0.1349	0.6619	-0.1058	0.8102	-0.0882	0.6556	0.0177	0.9351	-0.0226	0.9898	0.1658	0.6977	0.1884	0.6441
CG17490	CG17490	1631612_at	0.6335	0.0593	0.8687	0.0846	0.1048	0.6717	-0.5665	0.2438	-0.1270	0.7018	0.4395	0.0879	0.0716	0.9589	0.0343	0.9577	-0.0373	0.9453
---	---	1631613_at	-0.0415	0.8643	-0.1999	0.3828	-0.2977	0.2366	-0.1746	0.7752	0.1573	0.5691	0.3319	0.1454	0.0124	0.9923	-0.0134	0.9809	-0.0258	0.9514
CG5190	CG5190	1631614_at	-0.4509	0.0459	0.4381	0.1694	0.5713	0.0058	-0.2289	0.6086	-0.9627	0.0019	-0.7337	0.0038	-0.2962	0.7215	0.1756	0.6414	0.4718	0.1990
Grip128	Grip128	1631615_at	-0.2157	0.3891	0.4189	0.0894	0.3844	0.2184	-0.3130	0.4259	-0.4652	0.0386	-0.1522	0.4592	-0.3327	0.7187	0.1121	0.8228	0.4448	0.2614
CG15308	CG15308	1631616_at	0.0944	0.7577	0.2193	0.3815	0.2228	0.1770	-0.0215	0.9838	-0.1366	0.6075	-0.1151	0.6437	-0.0265	0.9816	-0.0208	0.9628	0.0058	0.9899
CG13742	CG13742	1631617_at	-0.0566	0.8144	-1.0887	0.0088	-1.2800	0.0009	-0.1184	0.8834	1.1183	0.0024	1.2368	0.0010	0.0325	0.9892	0.0917	0.9046	0.0592	0.9330
PPP4R2r	Protein phosphata	1631618_s_at	0.3922	0.0819	0.5381	0.0671	0.6925	0.0044	-0.0376	0.9633	-0.0650	0.8080	-0.0274	0.9178	-0.1331	0.8533	0.0029	0.9972	0.1360	0.6756
MTF-1	MTF-1	1631619_a_at	0.3971	0.1713	-0.0291	0.8696	-0.2715	0.1341	-0.1714	0.7142	0.3381	0.1124	0.5096	0.0157	0.0666	0.9619	-0.0639	0.9157	-0.1305	0.7764
GlyP	glycogen phosphc	1631620_at	0.3496	0.2758	0.5325	0.5103	0.3657	0.2251	-0.0450	0.9753	-1.5271	0.0019	-1.4820	0.0013	0.1859	0.9460	-1.2320	0.1374	-1.4179	0.1192
egh	zeste-white 4	1631621_s_at	1.0859	0.0117	-0.7511	0.0088	-0.7782	0.0169	0.3610	0.5633	1.9923	0.0004	1.6313	0.0006	0.3455	0.6960	0.1637	0.6993	-0.1819	0.6534
CG7747	CG7747	1631622_at	-0.3391	0.1734	0.3557	0.2882	0.6582	0.0071	0.1399	0.7270	-0.4562	0.0192	-0.5962	0.0035	-0.2165	0.8076	0.2141	0.5855	0.4306	0.2614
---	---	1631623_at	0.1797	0.3175	0.2340	0.1648	0.0791	0.7288	-0.0564	0.9048	0.0269	0.8979	0.0834	0.5864	0.0361	0.9760	-0.0010	0.9995	-0.0371	0.9261
CG14626	CG14626	1631624_at	0.1893	0.3330	-0.9688	0.0162	-0.3769	0.0393	0.4581	0.2165	0.9643	0.0019	0.5062	0.0191	-0.3020	0.6988	-0.2401	0.4727	0.0618	0.8915
CG15563	CG15563	1631625_at	0.2088	0.4027	-0.0506	0.6562	0.0974	0.6023	0.1684	0.6666	0.1194	0.5518	-0.0490	0.8165	0.0931	0.8999	0.0185	0.9670	-0.0746	0.8268
aub	aubergine	1631626_at	0.5598	0.2387	-0.7891	0.0563	-0.5250	0.1310	0.1887	0.7630	1.3137	0.0011	1.1250	0.0013	-0.1110	0.9701	-0.2293	0.8153	-0.1182	0.9096
CG5335 /// Fbp1	CG5335 /// Protein	1631627_at	0.3146	0.0975	0.3393	0.0977	0.2313	0.1164	0.0748	0.9102	-0.2410	0.2534	-0.3158	0.0932	0.2099	0.7266	-0.2425	0.3358	-0.4524	0.1156
CG12896 /// Prx2540-1 /// F	AOP2-related /// p	1631628_s_at	1.9651	0.0010	0.8339	0.2550	1.6680	0.0000	0.4020	0.3991	-0.1193	0.6984	-0.5213	0.0360	-0.2185	0.9142	-1.2858	0.0826	-1.0673	0.1577
CG14654	CG14654	1631629_at	0.1479	0.2997	0.0788	0.6940	0.1736	0.3137	-0.1095	0.7604	-0.1601	0.3045	-0.0507	0.7653	-0.1324	0.8235	-0.0636	0.8500	0.0687	0.8240
Klp67A	kinesin like protein	1631630_at	-0.1500	0.7632	-0.3512	0.3973	-0.8351	0.0644	-0.3254	0.5909	0.5308	0.0848	0.8562	0.0079	-0.4556	0.8097	-0.0793	0.9504	0.3763	0.6592
Cpr47Ee	CG13222	1631631_at	-0.0077	0.9951	1.0218	0.0725	0.5300	0.6357	-0.2034	0.9185	-1.3026	0.0395	-1.0992	0.0494	0.0121	0.9994	-0.1094	0.9789	-0.1215	0.9727
BicC	Bicaudal-C	1631632_s_at	1.0588	0.4870	-2.4341	0.2088	-1.6855	0.0387	-0.0230	0.9937	3.0885	0.0007	3.1114	0.0004	-0.9791	0.8960	-0.8425	0.8166	0.1366	0.9756
GNBP2	Gram-negative ba	1631633_a_at	-0.6857	0.0971	-0.4789	0.1592	-0.4225	0.0992	-0.2078	0.6965	-0.5358	0.0359	-0.3280	0.1333	-0.3820	0.7220	-0.5883	0.1874	-0.2063	0.6853
---	---	1631634_at	0.0414	0.8099	-0.0329	0.7597	0.1246	0.4914	0.0455	0.9375	-0.0376	0.8658	-0.0831	0.6297	-0.0620	0.9142	-0.0698	0.7787	-0.0078	0.9821
CG7294	CG7294	1631635_at	-0.0362	0.9112	0.0044	0.9721	-0.0897	0.5398	-0.0468	0.9578	0.0481	0.8777	0.0948	0.7008	-0.0208	0.9862	0.0020	0.9987	0.0228	0.9548
CG13096	CG13096	1631636_at	0.3154	0.2169	0.8622	0.1194	1.3857	0.0008	0.2525	0.5465	-0.0872	0.7316	-0.3396	0.0813	-0.2724	0.8465	0.6090	0.2810	0.8814	0.1562
Cpr	P450 reductase	1631637_a_at	0.2284	0.2882	-0.0506	0.8786	0.3934	0.0330	0.2447	0.4679	0.1670	0.3691	-0.0777	0.6831	-0.2209	0.7893	-0.1654	0.6709	0.0554	0.9094
Pvf1	PDGF- and VEGF	1631638_at	0.2330	0.4264	-0.6519	0.0996	-0.0032	0.9905	0.0459	0.9380	0.1075	0.5648	0.0616	0.7406	-0.6408	0.6034	-0.6665	0.1982	-0.0258	0.9761
TpnC47D	troponin C	1631639_at	-2.7737	0.0625	-3.1691	0.0118	-3.4783	0.0000	-0.3219	0.5735	-1.3034	0.0017	-0.9815	0.0034	0.2251	0.9742	-1.5305	0.3674	-1.7556	0.3205
CG18004	CG18004	1631640_a_at	-0.2327	0.1373	0.2047	0.1947	0.3732	0.0484	-0.0584	0.8915	-0.4363	0.0089	-0.3779	0.0102	-0.0135	0.9914	0.2409	0.3805	0.2544	0.3764
---	---	1631641_at	0.1878	0.2663	0.1741	0.3849	-0.0626	0.7501	0.0817	0.8325	0.0978	0.5319	0.0162	0.9280	0.3079	0.6695	0.1210	0.7161	-0.1869	0.5437
CG4424	CG4424	1631642_at	0.2684	0.1758	0.0327	0.9171	-0.2964	0.1762	-0.0006	0.9994	0.6084	0.0040	0.6090	0.0024	0.1124	0.9226	0.1611	0.7111	0.0486	0.9271
---	---	1631643_at	0.1635	0.4869	0.3301	0.0986	0.4595	0.1385	-0.0777	0.9436	-0.3115	0.3284	-0.2338	0.4280	-0.1311	0.8246	-0.2030	0.4056	-0.0719	0.8148
CG31803	CG31803	1631644_at	0.0062	0.9845	-0.1392	0.4867	-0.0511	0.8467	0.1724	0.7885	0.1956	0.4782	0.0231	0.9445	-0.1090	0.8971	-0.0764	0.8611	0.0326	0.9411
CG3363	CG3363	1631645_at	0.2815	0.3217	-0.4013	0.1353	-0.4446	0.0376	0.0985	0.9060	0.5669	0.0415	0.4683	0.0556	0.1177	0.9235	0.0297	0.9647	-0.0880	0.8699
---	---	1631646_at	0.2257	0.2515	-0.0091	0.9361	-0.0473	0.8092	-0.0842	0.8611	0.0779	0.6949	0.1621	0.3004	0.0508	0.9421	-0.0172	0.9620	-0.0680	0.7933
---	---	1631647_at	0.1483	0.5131	0.1535	0.2552	0.2692	0.0740	-0.0257	0.9662	-0.0464	0.8144	-0.0206	0.9156	-0.1718	0.7485	0.0492	0.8807	0.2209	0.3475
Srp9	Srp9	1631648_at	0.5410	0.0413	0.5957	0.0294	0.8581	0.0065	0.0227	0.9704	0.1712	0.2726	0.1485	0.2915	-0.2246	0.8235	0.3434	0.4073	0.5680	0.1990
CG8271	CG8271	1631649_at	-0.0192	0.9889	-0.2600	0.2574	-0.4845	0.1032	0.0638	0.9873	-0.3175	0.7844	-0.3814	0.7057	0.2474	0.6389	-0.3278	0.1299	-0.5753	0.0422
DsimCG2056 /// spirit	CG2056	1631650_a_at	0.4305	0.1620	0.5995	0.1502	0.2330	0.3077	-0.1095	0.7923	-0.0060	0.9809	0.1034	0.5303	0.1807	0.9238	0.1429	0.8725	-0.0378	0.9684
CG14905	CG14905	1631651_at	-0.0846	0.6005	-0.0371	0.7546	0.1398	0.3609	0.1536	0.6861	0.0142	0.9567	-0.1394	0.3961	-0.0068	0.9932	0.0023	0.9957	0.0091	0.9756
CG7713	CG7713	1631652_at	-0.6623	0.0492	-0.2997	0.2275	-0.6518	0.0127	-0.0566	0.9247	-0.1977	0.2716	-0.1411	0.3970	0.3104	0.7899	0.2844	0.5897	-0.0260	0.9732

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15395	CG15395	1631653_at	0.1508	0.3224	0.0182	0.9278	0.1009	0.6496	0.0109	0.9878	-0.0379	0.8643	-0.0488	0.7962	-0.0216	0.9862	-0.0208	0.9644	0.0008	0.9990
CG31693	CG31693	1631654_at	0.2239	0.3278	0.0493	0.6997	-0.0769	0.6687	-0.2223	0.4116	-0.0691	0.6875	0.1532	0.2563	-0.0455	0.9653	-0.0680	0.8625	-0.0225	0.9550
CG9132	CG9132	1631655_a_at	0.2941	0.2325	0.3525	0.0908	0.3638	0.0285	-0.1548	0.6471	0.2695	0.0997	0.4243	0.0110	-0.1806	0.8386	0.2617	0.4833	0.4423	0.2420
---	---	1631656_at	-0.0346	0.8326	0.1306	0.3761	0.3872	0.1009	0.3578	0.2095	0.0925	0.6272	-0.2654	0.0840	0.0077	0.9950	0.1498	0.6270	0.1421	0.6436
---	---	1631657_at	-0.0231	0.9116	-0.1596	0.1973	-0.0795	0.5932	0.0100	0.9876	0.1549	0.3206	0.1448	0.2997	-0.1579	0.7768	0.0150	0.9701	0.1729	0.4869
---	---	1631658_at	0.1424	0.4183	-0.0455	0.6652	0.0197	0.9094	0.1514	0.6122	0.3657	0.0228	0.2143	0.1084	-0.1055	0.8235	-0.0556	0.8319	0.0499	0.8434
CG32704 /// DmirCG32704	CG32704	1631659_at	0.2063	0.3094	0.1382	0.3859	0.1563	0.2840	-0.0422	0.9438	-0.0487	0.8195	-0.0065	0.9761	-0.0256	0.9776	0.0445	0.8884	0.0701	0.7877
CG15065	CG15065	1631660_at	1.4117	0.1619	0.8773	0.3659	1.5870	0.0129	0.0724	0.9633	0.0585	0.9206	-0.0139	0.9788	-0.6772	0.8842	-0.4058	0.8718	0.2714	0.9109
---	---	1631661_at	0.1655	0.4496	0.0205	0.8392	0.2263	0.3613	0.0324	0.9598	-0.0399	0.8555	-0.0723	0.6819	-0.1587	0.8114	-0.1913	0.4983	-0.0326	0.9330
---	---	1631662_a_at	0.2296	0.2777	0.0487	0.7291	0.1240	0.6586	-0.1192	0.7937	-0.0226	0.9312	0.0966	0.6072	-0.0943	0.9503	-0.1813	0.7266	-0.0870	0.8880
mRpS24	mitochondrial ribo	1631663_at	-0.1929	0.3175	0.2072	0.4824	0.1071	0.6203	0.0333	0.9705	-0.0333	0.9180	-0.0666	0.7974	0.1021	0.9092	0.3493	0.2520	0.2472	0.4503
cngl	CNG channel-like	1631664_a_at	0.1621	0.4652	0.2971	0.0488	0.2879	0.1036	0.0939	0.8196	0.0841	0.6424	-0.0098	0.9613	0.0858	0.9199	0.2522	0.3738	0.1664	0.5899
Acp76A	Accessory gland p	1631665_at	-0.0539	0.7600	0.1061	0.3923	0.0672	0.8055	0.1142	0.8386	0.0430	0.8782	-0.0712	0.7565	0.0965	0.8888	0.1393	0.6166	0.0428	0.9056
CG9766	CG9766	1631666_s_at	-0.1493	0.5035	0.0832	0.5098	-0.0656	0.7331	-0.3100	0.3194	-0.4487	0.0217	-0.1387	0.4141	-0.0899	0.9174	-0.0866	0.8268	0.0032	0.9953
---	---	1631667_at	0.1239	0.4226	0.2271	0.3133	0.1241	0.4603	0.0319	0.9553	-0.0630	0.7248	-0.0948	0.5250	-0.0227	0.9776	-0.0776	0.7293	-0.0549	0.8204
CG17147	CG17147	1631668_at	0.0535	0.8333	0.0487	0.6494	0.0859	0.6248	0.0065	0.9945	0.0216	0.9407	0.0151	0.9533	-0.0074	0.9948	-0.0048	0.9935	0.0026	0.9954
CG32801	CG32801	1631669_at	0.0895	0.6094	0.4522	0.1287	0.2697	0.1703	-0.1647	0.6652	-0.2469	0.1721	-0.0822	0.6628	0.0290	0.9717	0.0114	0.9745	-0.0177	0.9517
Spn2	serpin 2	1631670_at	0.1179	0.4829	-0.0454	0.6391	0.0428	0.7865	0.0239	0.9745	0.0755	0.7270	0.0516	0.8043	-0.1345	0.7707	0.0157	0.9617	0.1502	0.4597
CG13827	CG13827	1631671_at	0.4540	0.1485	0.7708	0.0478	1.2461	0.0004	-0.0561	0.9375	-0.3361	0.1054	-0.2800	0.1306	-0.3385	0.7506	0.0860	0.9013	0.4244	0.3670
CG9867	CG9867	1631672_at	-1.2159	0.0017	-0.8558	0.0144	-0.8593	0.0003	0.1138	0.8155	-0.3372	0.0797	-0.4510	0.0165	0.0806	0.9152	-0.0463	0.9085	-0.1269	0.6579
CG14454 /// CG32453	CG32453 /// CG1	1631673_s_at	-0.0264	0.8979	0.4625	0.1051	0.1359	0.5226	-0.0143	0.9874	-0.3161	0.1437	-0.3018	0.1188	0.0701	0.9340	-0.0541	0.8924	-0.1242	0.6761
O-fut1	neurotic	1631674_at	-0.2765	0.1658	-0.0406	0.9219	-0.2453	0.1066	-0.2470	0.4259	-0.0600	0.7711	0.1870	0.2230	-0.0103	0.9950	0.1548	0.7202	0.1651	0.6950
Phm	Peptidyl gycine al	1631675_a_at	0.0330	0.8556	-0.1432	0.5566	0.1398	0.3714	0.1847	0.5628	0.1371	0.4170	-0.0475	0.7962	-0.0487	0.9589	0.0260	0.9521	0.0747	0.8256
CG4467	CG4467	1631676_at	-0.2609	0.3613	-0.3505	0.0346	-0.4247	0.0147	-0.1003	0.8883	-0.0170	0.9628	0.0833	0.7501	-0.0956	0.8940	-0.0169	0.9692	0.0787	0.8117
dalao	Brahma associate	1631677_at	0.0371	0.9544	-0.1190	0.9399	-0.1466	0.7785	0.0209	0.9803	-0.1424	0.5064	-0.1633	0.3858	0.2631	0.9634	-0.1933	0.9395	-0.4564	0.8138
---	---	1631678_at	0.0836	0.7150	0.0619	0.6233	0.1212	0.5301	0.0600	0.9108	-0.0958	0.6072	-0.1558	0.3123	0.0956	0.8965	0.0069	0.9914	-0.0888	0.7849
---	---	1631679_x_at	-0.1397	0.6108	0.0182	0.9479	-0.0208	0.9147	0.1418	0.7243	0.0537	0.8104	-0.0881	0.6347	0.1471	0.8395	0.1299	0.7008	-0.0172	0.9702
CG31878	CG31878	1631680_a_at	0.0270	0.9168	-0.0104	0.9755	-0.0904	0.5947	-0.0065	0.9956	0.0687	0.8446	0.0752	0.8047	-0.0119	0.9927	-0.0437	0.9259	-0.0317	0.9387
825-Oak /// CG32213 /// CC	CG32214 /// CG3	1631681_s_at	-0.1611	0.3539	-0.1084	0.4907	-0.4141	0.0606	-0.2115	0.4850	-0.1244	0.4604	0.0871	0.5900	0.0807	0.8963	-0.0601	0.8484	-0.1408	0.5649
twi	Twist	1631682_at	-0.9091	0.0955	-1.0349	0.0264	-1.2569	0.0079	0.1483	0.8678	-0.0532	0.9026	-0.2015	0.5078	0.2518	0.8494	-0.0747	0.9326	-0.3265	0.5788
---	---	1631683_at	-0.0377	0.8901	0.1671	0.2650	0.2616	0.1159	-0.0332	0.9491	-0.0902	0.5647	-0.0570	0.7111	-0.0026	0.9989	0.1714	0.5877	0.1740	0.5834
CG13014	CG13014	1631684_at	0.3281	0.0776	0.6197	0.0409	0.4503	0.0169	-0.2567	0.4259	-0.2929	0.0999	-0.0362	0.8616	-0.0982	0.9088	-0.0372	0.9402	0.0610	0.8834
---	---	1631685_at	0.0277	0.8784	0.4969	0.0447	0.4504	0.0674	-0.2409	0.5836	-0.4377	0.0551	-0.1969	0.3363	-0.1492	0.8740	0.0674	0.9054	0.2166	0.5819
CG34400	CG17368	1631686_at	0.0513	0.8099	0.0476	0.6935	-0.0284	0.8832	-0.0344	0.9727	0.0054	0.9889	0.0398	0.8946	0.0084	0.9946	-0.1779	0.5455	-0.1863	0.5299
RN-tre	tre oncogene-relat	1631687_at	-0.6363	0.1580	-0.6249	0.0887	-0.7948	0.0113	0.1452	0.8837	0.1032	0.8035	-0.0419	0.9193	0.2206	0.8744	0.0479	0.9585	-0.1727	0.7940
mal	bronzy	1631688_at	1.2978	0.0124	1.2606	0.0596	1.2839	0.0001	0.2686	0.3921	0.0950	0.6299	-0.1736	0.2775	0.1781	0.9305	0.0671	0.9499	-0.1110	0.9028
CG18482	CG18482	1631689_at	0.0912	0.7026	-0.0245	0.8522	0.1274	0.6057	0.1976	0.4751	0.2434	0.1013	0.0459	0.7820	0.0743	0.9503	0.0868	0.8595	0.0126	0.9836
CG9945	CG9945	1631690_a_at	0.2404	0.1741	0.6031	0.0898	0.5178	0.0723	0.1085	0.9011	-0.3071	0.2805	-0.4156	0.1001	0.0188	0.9913	-0.0272	0.9621	-0.0459	0.9233
CG16772	CG16772	1631691_at	1.0451	0.2771	0.7107	0.4103	1.1510	0.1272	0.0449	0.9915	1.4100	0.1112	1.3651	0.0860	-0.2544	0.9585	1.2653	0.3389	1.5197	0.2784
CG16798	CG16798	1631692_at	0.3777	0.0615	0.1529	0.3102	0.1338	0.4594	-0.0653	0.9186	0.0902	0.6895	0.1554	0.3956	0.0575	0.9400	0.0067	0.9898	-0.0508	0.8715
CG9616	CG9616	1631693_at	1.8481	0.0466	0.4975	0.3303	1.9728	0.0004	0.2513	0.8217	0.4483	0.3071	0.1970	0.6619	-0.9528	0.6389	-0.7390	0.3648	0.2139	0.8394
---	---	1631694_at	0.2235	0.2391	0.3836	0.1233	0.0790	0.6621	-0.1265	0.7690	-0.0740	0.7296	0.0525	0.7981	0.2663	0.7305	0.1629	0.6512	-0.1034	0.7963
CG13066	CG13066	1631695_at	-0.0327	0.8981	0.0226	0.9603	-0.1406	0.6231	0.1552	0.8676	-0.0488	0.9156	-0.2040	0.5232	0.0445	0.9589	-0.0415	0.9137	-0.0859	0.7654
MAN1	MAN1	1631696_s_at	0.3534	0.0629	0.2380	0.4782	0.2324	0.2106	-0.0741	0.9149	0.3094	0.1517	0.3835	0.0524	-0.0031	0.9984	0.1932	0.5544	0.1963	0.5523
Dro	Drosocin	1631697_at	0.4578	0.3617	0.4228	0.7014	-0.7966	0.2781	-0.6006	0.1462	1.4826	0.0005	2.0832	0.0001	0.3427	0.9564	0.9893	0.6082	0.6467	0.7560
CG16833	CG16833	1631698_s_at	-0.0283	0.9528	-1.1110	0.1420	-1.2673	0.0046	-0.2119	0.8342	1.4032	0.0039	1.6151	0.0013	-0.1234	0.9717	0.4324	0.6483	0.5558	0.5461
CG15436	CG15436	1631699_at	-0.0936	0.7425	-0.1024	0.8462	-0.1901	0.4426	0.1997	0.6699	0.7830	0.0046	0.5833	0.0103	0.0845	0.9734	0.5738	0.3446	0.4893	0.4457
CG18331	CG18331	1631700_at	0.4583	0.2187	0.4693	0.3423	0.1459	0.7593	-0.1676	0.9194	0.8352	0.0997	1.0028	0.0342	0.0436	0.9898	0.8034	0.2394	0.7598	0.2960
Cpr49Ac	CG8502	1631701_a_at	0.1500	0.6866	-0.0664	0.7167	0.0649	0.8643	0.2471	0.7717	0.1325	0.7576	-0.1147	0.7725	0.0650	0.9589	0.0044	0.9955	-0.0606	0.9016
CG15888	CG15888	1631702_at	0.1081	0.5633	0.1431	0.3180	0.0145	0.9445	0.0222	0.9753	0.0538	0.8023	0.0316	0.8791	0.2468	0.7324	0.0488	0.9193	-0.1981	0.5456
---	---	1631703_at	0.0695	0.7700	0.0570	0.5902	0.0230	0.9197	-0.0484	0.9436	-0.0425	0.8701	0.0059	0.9808	0.0141	0.9913	-0.0613	0.8749	-0.0754	0.8247
CG5805	CG5805	1631704_at	-0.3653	0.0527	-0.2049	0.0964	-0.4558	0.0181	-0.3527	0.2596	-0.3821	0.0452	-0.0294	0.8953	-0.1036	0.8650	-0.1963	0.4164	-0.0928	0.7411
---	---	1631705_at	0.1911	0.3611	0.1437	0.3828	-0.0197	0.9403	-0.1704	0.6327	-0.0433	0.8484	0.1272	0.4389	0.0200	0.9912	0.0077	0.9925	-0.0123	0.9842
CG13955	CG13955	1631706_at	-0.0996	0.5493	0.0391	0.8628	0.4833	0.0168	0.1159	0.7699	0.0008	0.9976	-0.1151	0.4702	-0.1255	0.8692	0.0477	0.9192	0.1732	0.5929
CG7370	CG7370	1631707_at	0.2533	0.1699	0.0036	0.9767	-0.1224	0.4543	-0.0707	0.8923	0.1382	0.4467	0.2088	0.1801	0.0252	0.9831	-0.0687	0.8575	-0.0938	0.7744
---	---	1631708_at	0.0180	0.9444	0.0924	0.6408	-0.0388	0.8458	0.0583	0.9449	-0.0699	0.8147	-0.1282	0.5953	0.1930	0.7633	0.1248	0.6747	-0.0682	0.8418
CG13980	CG13980	1631709_at	0.1466	0.4192	0.0073	0.9666	-0.0253	0.8867	0.0864	0.8288	0.0438	0.8188	-0.0426	0.8047	0.0633	0.9400	-0.1710	0.5226	-0.2342	0.3764
KaiRIA	bad reception	1631710_at	-2.3120	0.0006	-2.3245	0.0405	-2.8373	0.0000	-0.2162	0.4356	-0.6321	0.0022	-0.4159	0.0086	0.1319	0.9646	-0.6974	0.3749	-0.8293	0.3106
CG14823	CG14823	1631711_at	0.2819	0.1116	0.1068	0.3716	0.3323	0.2373	0.0641	0.8950	-0.0103	0.9676	-0.0744	0.6563	-0.1719	0.8395	-0.1051	0.8168	0.0668	0.8893
CG13398	CG13398	1631712_at	0.0760	0.8114	0.4199	0.1463	0.5182	0.0150	-0.0268	0.9745	-0.1311	0.5492	-0.1043	0.6122	-0.1526	0.8905	0.1521	0.7612	0.3047	0.4815
---	---	1631713_x_at	0.1163	0.4952	0.0113	0.9280	-0.0457	0.8234	-0.0530	0.9410	0.0941	0.6881	0.1471	0.4459	-0.1589	0.7726	-0.0697	0.8199	0.0892	0.7425
SP71	SP71	1631714_a_at	0.5544	0.0502	0.2512	0.3526	0.2773	0.2164	-0.2807	0.4728	0.0453	0.8712	0.3260	0.0840	-0.1338	0.9095	-0.2485	0.5742	-0.1147	0.8268
CG14872	CG14872	1631715_at	1.5114	0.3305	0.8870	0.4829	-2.9297	0.0007	-3.1386	0.0203	-0.2744	0.7479	2.8642	0.0016	1.0425	0.8439	-0.5611	0.8500	-1.6036	0.4753
CG2709	CG2709	1631716_at	0.0922	0.6803	0.1547	0.3848	-0.0608	0.7439	-0.1178	0.8671	-0.1281	0.6460	-0.0103	0.9733	-0.0734	0.9301	-0.0832	0.8123	-0.0098	0.9835
---	---	1631717_at	-0.0127	0.9676	0.0462	0.7086	-0.0276	0.8714	0.0368	0.9599	0.0307	0.9084	-0.0061	0.9799	0.0605	0.9472	0.0518	0.9033	-0.0087	0.9849
mRpL12	mitochondrial ribo	1631718_at	-0.1279	0.6908	0.6529	0.0303	0.5825	0.0157	0.0614	0.9136	-0.5422	0.0089	-0.6036	0.0033	0.0724	0.9746	0.4137	0.4753	0.3413	0.5724
CG31109	CG31109	1631719_at	-0.0072	0.9745	-0.3448	0.0734	-0.6201	0.0022	-0.0473	0.9308	0.4284	0.0151	0.4757	0.0057	0.2326	0.7215	0.1813	0.5200	-0.0513	0.8915
---	---	1631720_at	-0.0221	0.9535	-0.0195	0.8546	0.0029	0.9918	-0.0899	0.8540	0.0405	0.8611	0.1304	0.4308	-0.0662	0.9489	0.0607	0.8952	0.1270	0.7154
CkIIalpha-i1	CKII-alpha subuni	1631721_at	0.1497	0.6314	-0.1598	0.1556	-0.5768	0.0284	-0.0741	0.9260	0.4800	0.0507	0.5541	0.0180	0.2005	0.7893	0.0413	0.9353	-0.1592	0.6414

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1631722_at	0.5274	0.1424	-0.5903	0.3942	-0.9693	0.0071	-0.3883	0.1703	1.3898	0.0002	1.7781	0.0000	-0.1664	0.9491	0.1163	0.9240	0.2827	0.7526
CG5498	transcript 2	1631723_at	-0.1839	0.3521	0.1253	0.5031	0.1301	0.5829	-0.3343	0.2092	-0.8945	0.0007	-0.5601	0.0028	-0.2382	0.7588	-0.4297	0.1744	-0.1915	0.5814
CG17746	CG17746	1631724_s_at	-0.7066	0.0192	0.1719	0.3234	0.4609	0.1385	0.0524	0.9295	-0.6420	0.0037	-0.6944	0.0016	-0.1328	0.9309	0.1894	0.7466	0.3222	0.5403
CG4691	CG4691	1631725_at	0.2649	0.1724	0.0441	0.7991	0.0387	0.9006	-0.0127	0.9903	-0.0801	0.7755	-0.0674	0.7968	-0.0018	0.9994	-0.1033	0.7623	-0.1015	0.7568
CG31190	CG31190	1631726_at	0.1810	0.3909	0.0464	0.8668	-0.0356	0.8348	-0.0991	0.8930	-0.0743	0.8058	0.0248	0.9351	0.0343	0.9779	-0.0971	0.7949	-0.1314	0.6884
CG33121	CG33121	1631727_at	0.1230	0.4997	-0.0468	0.6836	0.0684	0.7344	-0.0111	0.9895	0.1323	0.5236	0.1433	0.4337	-0.0247	0.9742	0.0121	0.9689	0.0368	0.8871
PRL-1	PRL-1	1631728_s_at	-0.4484	0.0716	0.3716	0.3556	-0.7135	0.0040	-0.0660	0.9042	0.1357	0.4667	0.2017	0.2059	1.0248	0.2740	0.9371	0.0664	-0.0876	0.8925
CG13671	CG13671	1631729_at	0.3106	0.0835	0.1136	0.6428	0.2308	0.1877	-0.1170	0.7701	0.2428	0.1499	0.3598	0.0257	-0.1571	0.8326	0.0840	0.8395	0.2411	0.4498
CG12911	CG12911	1631730_at	-0.0399	0.7844	-0.0867	0.6342	-0.1118	0.5866	-0.1057	0.8107	-0.1713	0.3359	-0.0656	0.7256	-0.1848	0.7215	-0.1229	0.5902	0.0618	0.8178
CG1494	CG1494	1631731_at	0.0977	0.7064	-0.1946	0.3408	-0.0849	0.6648	0.1277	0.8028	0.2212	0.2836	0.0935	0.6595	-0.1147	0.8331	-0.1978	0.3735	-0.0831	0.7495
sdt	anon-fast-evolving	1631732_s_at	-0.0631	0.7848	0.3625	0.0564	0.2559	0.2427	-0.0432	0.9556	-0.1291	0.5658	-0.0859	0.6945	0.2188	0.7330	0.3152	0.2367	0.0965	0.7625
CG14817	CG14817	1631733_at	-0.2165	0.3651	-0.0538	0.8992	-0.1628	0.3410	-0.0423	0.9641	-0.1763	0.5024	-0.1340	0.5910	0.1220	0.9117	0.0229	0.9723	-0.0991	0.8424
---	---	1631734_at	-0.0759	0.6282	0.0902	0.5664	-0.0121	0.9521	0.0054	0.9956	-0.0554	0.8426	-0.0608	0.8015	0.0187	0.9816	0.0108	0.9719	-0.0078	0.9791
CG10445	CG10445	1631735_at	-0.1773	0.4399	-0.1668	0.4188	-0.1839	0.5208	0.0225	0.9857	0.3594	0.1971	0.3370	0.1773	-0.2410	0.7848	0.1065	0.8343	0.3475	0.3723
CG15720	CG15720	1631736_at	-0.0381	0.9181	0.0825	0.7086	0.0797	0.6753	0.0871	0.9220	0.1210	0.6952	0.0339	0.9178	0.1350	0.8668	0.3081	0.3239	0.1731	0.6155
---	---	1631737_at	0.0923	0.7378	0.2319	0.5524	-0.1112	0.7594	-0.3615	0.2266	0.1112	0.5628	0.4726	0.0098	0.1252	0.9503	0.2370	0.7339	0.1117	0.8918
CG6116	CG6116	1631738_at	-0.6557	0.0037	-0.1947	0.2215	-0.1812	0.4866	-0.0586	0.9346	-0.1485	0.5063	-0.0899	0.6853	0.0345	0.9775	0.2380	0.4103	0.2035	0.5008
RpL13A	Ribosomal protein	1631739_at	0.3378	0.0598	0.1585	0.2751	0.2592	0.1247	0.0993	0.8337	0.0867	0.6715	-0.0126	0.9562	0.1143	0.8122	-0.0348	0.9085	-0.1490	0.4683
CG30487	CG30487	1631740_at	0.0361	0.8546	0.1089	0.4427	0.0054	0.9843	0.0314	0.9726	0.0012	0.9971	-0.0302	0.9129	-0.0597	0.9421	-0.0890	0.7749	-0.0293	0.9344
Ptp4E	Protein tyrosine pl	1631741_a_at	-2.2658	0.0028	-0.2332	0.7054	-0.2433	0.2318	-0.2327	0.7924	-1.9948	0.0007	-1.7620	0.0007	-0.1473	0.9390	0.0705	0.9409	0.2177	0.7500
mod(mdg4)	Modifier67.2	1631742_s_at	-0.0574	0.8558	-0.2017	0.6094	-0.5662	0.0192	-0.2312	0.6324	-0.1993	0.4133	0.0319	0.9111	0.1559	0.9101	-0.2122	0.7027	-0.3681	0.4689
CG40172	CG40172	1631743_at	0.0877	0.5887	0.1165	0.4418	0.2196	0.3128	-0.0787	0.8578	-0.1958	0.2131	-0.1171	0.4285	-0.0155	0.9898	0.0516	0.8858	0.0670	0.8307
tj	female sterile(2)ec	1631744_at	0.1105	0.6258	0.1511	0.1915	0.3464	0.0876	0.1603	0.8244	-0.1234	0.7018	-0.2837	0.2597	-0.0390	0.9544	-0.0051	0.9911	0.0339	0.9025
CG6766	CG6766	1631745_at	0.6238	0.0047	0.8302	0.0818	1.0917	0.0004	-0.0066	0.9937	0.1726	0.3264	0.1791	0.2498	-0.2764	0.7588	0.4161	0.2520	0.6925	0.1013
CG8774	CG8774	1631746_a_at	-0.1327	0.9732	-0.0155	0.9173	-0.4876	0.0721	-0.2904	0.9705	-2.0637	0.3086	-1.7733	0.3340	0.0571	0.9964	-1.8706	0.5488	-1.9277	0.5403
---	---	1631747_at	0.0694	0.6980	0.1757	0.3632	0.1720	0.4035	0.0035	0.9956	0.0732	0.6924	0.0697	0.6794	0.0842	0.9330	0.1417	0.6949	0.0575	0.8957
---	---	1631748_at	0.1228	0.5184	0.0871	0.4867	-0.0112	0.9681	0.0127	0.9916	0.0562	0.8668	0.0435	0.8853	0.0120	0.9901	-0.0956	0.6876	-0.1076	0.6389
CG8928	CG8928	1631749_at	-0.3048	0.0626	-0.2466	0.4488	-0.4421	0.0453	-0.0899	0.8719	-0.1164	0.5812	-0.0265	0.9101	0.1224	0.8972	-0.1362	0.7424	-0.2586	0.4792
Oscp	Oligomycin sensiti	1631750_a_at	-0.1504	0.6770	0.1559	0.5120	0.0088	0.9744	-0.1699	0.7327	-0.5801	0.0171	-0.4102	0.0445	-0.1074	0.9457	-0.4011	0.3941	-0.2937	0.5612
CG10834	CG10834	1631751_at	-0.0988	0.6591	-0.0359	0.7241	0.2280	0.2273	0.0619	0.9436	0.0009	0.9979	-0.0609	0.8296	-0.2757	0.5989	-0.1164	0.6409	0.1593	0.5041
---	---	1631752_at	0.1538	0.3883	-0.1377	0.3686	-0.0308	0.8732	0.0062	0.9941	0.0570	0.7851	0.0508	0.7922	-0.0603	0.9199	-0.1618	0.4246	-0.1016	0.6419
CG15550	CG15550	1631753_at	0.1694	0.4168	0.0296	0.7703	-0.0793	0.6376	0.0211	0.9777	0.0072	0.9806	-0.0140	0.9549	-0.0837	0.8999	-0.1248	0.6382	-0.0411	0.9031
---	---	1631754_at	0.0007	0.9973	0.0183	0.8556	-0.1448	0.5827	-0.0849	0.9011	-0.0644	0.8141	0.0205	0.9404	-0.0179	0.9869	0.0441	0.9084	0.0620	0.8470
CG1236	CG1236	1631755_at	0.2449	0.3491	0.2958	0.0689	0.3775	0.0708	-0.1240	0.8190	0.0026	0.9934	0.1265	0.5426	-0.2128	0.8062	0.1391	0.7439	0.3519	0.3458
CG10859	CG10859	1631756_at	-0.0054	0.9856	-0.0269	0.8461	0.0697	0.6774	0.0437	0.9507	0.1027	0.6395	0.0589	0.7874	0.0676	0.8940	0.1074	0.5907	0.0397	0.8761
CG31882	CG31882	1631757_at	0.2304	0.3358	0.3846	0.0959	0.3703	0.1692	-0.0546	0.9426	-0.2149	0.3264	-0.1604	0.4285	-0.0123	0.9927	-0.0680	0.8764	-0.0557	0.8943
---	---	1631758_at	-0.0496	0.7350	0.0445	0.7886	-0.0197	0.9373	-0.0141	0.9858	0.0760	0.7225	0.0901	0.6310	-0.0004	0.9998	0.0322	0.9269	0.0327	0.9155
---	---	1631759_at	0.1620	0.5145	0.0067	0.9569	0.0462	0.7954	0.0634	0.9218	-0.0095	0.9749	-0.0729	0.7258	0.0485	0.9589	-0.0943	0.7710	-0.1428	0.6186
CG7768 /// DyakCG7768	CG7768	1631760_at	-1.0027	0.0487	-1.8594	0.0419	-1.8673	0.0007	-0.0457	0.9671	1.0606	0.0038	1.1062	0.0019	-0.1252	0.9598	0.0508	0.9650	0.1760	0.8461
CG31313	CG31313	1631761_at	0.2657	0.1513	-0.5509	0.4029	-0.0107	0.9686	0.3625	0.4139	0.2426	0.3310	-0.1199	0.6339	-0.2450	0.9044	-0.6202	0.3941	-0.3752	0.6366
---	---	1631762_at	0.0086	0.9692	-0.0068	0.9860	0.3047	0.0801	0.1639	0.7982	0.1480	0.6031	-0.0159	0.9611	-0.0941	0.8790	0.0614	0.8500	0.1555	0.5345
CG31793	CG31793	1631763_at	-0.4592	0.0627	-0.1902	0.2815	0.1234	0.5468	-0.1012	0.9036	-0.4463	0.0992	-0.3451	0.1527	-0.5701	0.3362	-0.2106	0.5166	0.3595	0.2674
CG30115	CG30115	1631764_a_at	-1.2401	0.0202	-1.0537	0.2762	-0.8346	0.0373	0.1678	0.8513	-0.0301	0.9509	-0.1980	0.5343	0.0315	0.9950	0.2581	0.8756	0.2267	0.8864
puc	puckered	1631765_at	-0.1930	0.6849	-0.3670	0.1904	-0.5705	0.0418	0.2653	0.5217	0.7700	0.0044	0.5047	0.0175	0.4780	0.7628	0.7086	0.2668	0.2305	0.7646
CG8818	CG8818	1631766_at	-0.8688	0.0014	-0.0937	0.6619	0.1639	0.5199	0.2559	0.5046	-0.6256	0.0084	-0.8816	0.0011	-0.0269	0.9894	0.1610	0.7466	0.1879	0.6881
CG13538 /// DereCG13538	CG13538	1631767_at	0.1683	0.3748	-0.0506	0.6895	0.0835	0.5697	0.0307	0.9649	0.0915	0.6662	0.0608	0.7675	-0.0199	0.9779	-0.0701	0.7204	-0.0502	0.8135
qkr58E-1	KH-domain	1631768_at	0.4899	0.0728	-0.0463	0.8514	0.4628	0.0237	-0.1232	0.8103	-0.0576	0.8210	0.0656	0.7711	-0.4533	0.7062	-0.4294	0.3823	0.0239	0.9762
put	activin receptor	1631769_at	-0.0296	0.9326	0.6901	0.0674	0.3184	0.1238	-0.3140	0.4861	-0.7484	0.0080	-0.4344	0.0488	0.0796	0.9550	-0.0811	0.8951	-0.1607	0.7345
Tango13	Transport and Gol	1631770_at	1.6430	0.0382	2.0702	0.0095	2.3172	0.0000	-0.0524	0.9544	-0.1396	0.6131	-0.0872	0.7481	-0.2150	0.9445	0.4947	0.6216	0.7097	0.4620
---	---	1631771_at	-0.0748	0.6971	-0.0535	0.6842	-0.0382	0.8285	0.0416	0.9493	0.0926	0.6523	0.0510	0.8040	0.0957	0.9004	0.0548	0.8963	-0.0409	0.9164
MTA1-like	MTA-like	1631772_a_at	0.8250	0.0575	0.9552	0.0303	0.3260	0.2373	-0.3395	0.3939	0.1383	0.5679	0.4779	0.0253	0.3594	0.7956	0.3410	0.5824	-0.0184	0.9848
Cpr30B	CG3818	1631773_at	0.1830	0.2838	0.1267	0.4834	0.0299	0.8811	0.0456	0.9528	-0.0307	0.9177	-0.0763	0.7384	0.1029	0.8953	-0.1072	0.7585	-0.2101	0.4865
Cpr66Cb	CG7076	1631774_at	1.2841	0.0400	0.5527	0.1112	0.5028	0.1019	-0.2219	0.7613	-0.4100	0.1899	-0.1881	0.5408	0.0788	0.9829	-0.8391	0.3022	-0.9178	0.2869
CG11137	CG11137	1631775_at	0.0247	0.9046	0.2107	0.4018	0.0010	0.9963	0.0429	0.9379	0.0354	0.8681	-0.0075	0.9704	0.2304	0.7338	0.2113	0.4743	-0.0191	0.9646
CG11968	CG11968	1631776_at	-0.1013	0.6608	-0.2475	0.2968	0.2039	0.2711	-0.0068	0.9931	-0.1526	0.3318	-0.1458	0.2992	-0.3703	0.6993	-0.2697	0.5183	0.1006	0.8459
fz4	frizzled 4	1631777_a_at	0.4312	0.0217	0.2145	0.3042	-0.0212	0.9314	-0.2262	0.6533	0.2448	0.3188	0.4710	0.0368	0.0827	0.9092	0.1136	0.6952	0.0309	0.9320
Drs-I	drosomycin-H	1631778_at	0.0686	0.7828	-0.0735	0.7304	0.2391	0.1621	0.0999	0.8903	0.0658	0.8314	-0.0341	0.9078	-0.0994	0.8706	-0.1664	0.5075	-0.0670	0.8256
ptr	proximal to raf	1631779_s_at	0.6114	0.2754	0.8011	0.1110	0.4562	0.0655	0.4173	0.5398	0.4937	0.1619	0.0763	0.8530	0.6653	0.6761	0.5939	0.3627	-0.0714	0.9387
C15	clawless	1631780_at	0.1122	0.4764	0.3256	0.3624	0.2377	0.2976	0.0613	0.9247	-0.0215	0.9385	-0.0828	0.6841	0.1561	0.8297	0.0348	0.9435	-0.1213	0.7259
CG18530	CG18530	1631781_at	1.0634	0.0277	0.6678	0.1554	1.4564	0.0541	-0.0014	0.9988	0.0896	0.7193	0.0910	0.6852	-0.3824	0.9238	-0.0122	0.9975	0.3702	0.8238
CG14312	CG14312	1631782_at	0.0651	0.7949	0.1190	0.4856	0.1070	0.6476	0.0056	0.9956	0.0003	0.9993	-0.0053	0.9859	0.0642	0.9412	0.1420	0.6259	0.0778	0.8177
CG4267	CG4267	1631783_at	-0.5962	0.1223	1.5835	0.0085	0.4654	0.1790	-0.0296	0.9675	-0.4615	0.0238	-0.4319	0.0196	1.1347	0.3800	1.9328	0.0290	0.7981	0.2782
CG13663	CG13663	1631784_at	0.1379	0.5696	0.6165	0.0286	0.8576	0.0028	0.0615	0.9441	-0.2232	0.3863	-0.2846	0.2052	-0.1215	0.8885	0.3687	0.2430	0.4901	0.1577
gwl	greatwall	1631785_at	0.6575	0.3361	-0.8976	0.1676	-1.1187	0.1096	0.3299	0.4188	1.7630	0.0002	1.4331	0.0002	0.3083	0.9514	0.1635	0.9467	-0.1448	0.9452
CG10064	CG10064	1631786_at	0.1240	0.5222	0.0437	0.7089	-0.1758	0.2459	0.0156	0.9863	-0.0163	0.9604	-0.0319	0.9039	0.0024	0.9990	-0.1168	0.7309	-0.1192	0.7189
---	---	1631787_at	0.2425	0.1739	0.0040	0.9864	0.0019	0.9944	0.0557	0.9300	0.1226	0.5367	0.0669	0.7374	-0.1002	0.8903	-0.1741	0.5406	-0.0739	0.8297
CG13121	CG13121	1631788_at	0.2051	0.3396	0.0952	0.4669	0.3071	0.0721	-0.2144	0.6615	-0.0794	0.7803	0.1350	0.5612	-0.0098	0.9913	-0.0600	0.8048	-0.0501	0.8341
---	---	1631789_at	0.1112	0.6033	0.1578	0.3141	0.3579	0.0497	-0.2002	0.7511	-0.1731	0.5578	0.0271	0.9363	0.0310	0.9721	0.0993	0.6952	0.0682	0.8041
CG8578	CG8578	1631790_at	-0.2678	0.1854	0.2455	0.2259	0.1729	0.3248	-0.0320	0.9627	-0.1819	0.3252	-0.1499	0.3734	0.0874	0.9238	0.3825	0.1948	0.2951	0.3458

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1631791_at	0.2607	0.1236	-0.0433	0.7430	0.6421	0.0265	0.0540	0.9136	0.0301	0.8873	-0.0240	0.8991	-0.3577	0.6584	-0.0869	0.8468	0.2708	0.4220
CG33274	CG33274	1631792_at	0.2861	0.1142	0.0810	0.5610	0.1626	0.3385	-0.0505	0.9314	-0.0468	0.8318	0.0037	0.9863	-0.0490	0.9467	-0.1687	0.4514	-0.1197	0.6190
---	---	1631793_at	0.2803	0.1199	0.0697	0.6473	0.0583	0.7948	0.0035	0.9962	0.1526	0.5694	0.1490	0.5389	0.0302	0.9689	-0.0618	0.8153	-0.0920	0.6817
CG6201	CG6201	1631794_at	-0.3564	0.3672	-1.6656	0.0230	-1.2200	0.0003	0.0570	0.9427	0.8179	0.0040	0.7610	0.0033	-0.4054	0.7611	-0.4924	0.3686	-0.0871	0.9096
CG2657	CG2657	1631795_at	0.0175	0.9410	-0.0400	0.6725	-0.1890	0.1858	-0.0275	0.9734	0.0766	0.7522	0.1041	0.6138	0.1334	0.7780	-0.0686	0.7901	-0.2020	0.3397
CG32543	CG32543	1631796_at	-0.1840	0.4964	0.3340	0.0683	-0.0089	0.9666	0.0303	0.9760	-0.2571	0.3050	-0.2875	0.1945	0.2140	0.7506	0.1757	0.5622	-0.0383	0.9232
Or1a	Odorant receptor	1631797_at	0.1723	0.5870	1.2445	0.0318	-0.2640	0.6569	-0.7949	0.1001	0.2383	0.4190	1.0332	0.0024	0.7665	0.7142	1.2802	0.1392	0.5137	0.5808
CG4774	CG4774	1631798_s_at	0.0581	0.8118	-0.0074	0.9711	0.2595	0.1058	0.0762	0.9110	-0.2061	0.3484	-0.2823	0.1423	-0.1263	0.8472	-0.1342	0.6512	-0.0079	0.9853
---	---	1631799_at	0.1613	0.3487	0.0088	0.9352	-0.0356	0.8716	0.1672	0.6113	0.2160	0.1875	0.0487	0.7915	0.2205	0.6898	0.1557	0.5075	-0.0648	0.8192
CG6690	CG6690	1631800_at	0.0194	0.9220	0.0538	0.6567	0.1372	0.4609	0.1474	0.7230	0.1002	0.6323	-0.0472	0.8251	0.0595	0.9521	0.0847	0.8303	0.0252	0.9523
CG15044	CG15044	1631801_at	-0.1017	0.9235	-0.0314	0.8450	0.0832	0.7332	0.0899	0.9745	-0.4726	0.5229	-0.5625	0.3826	0.0202	0.9940	-0.2341	0.7538	-0.2542	0.7208
---	---	1631802_at	-0.0395	0.8507	0.0142	0.9132	-0.1815	0.5589	-0.0834	0.9295	0.0609	0.8675	0.1443	0.6034	0.0945	0.9306	0.1633	0.6750	0.0688	0.8869
Lsp1alpha	Larval serum prot	1631803_at	0.1504	0.7477	0.4414	0.1698	-0.1379	0.3702	-0.1010	0.7857	-0.1576	0.3082	-0.0566	0.7304	0.5017	0.7215	0.2495	0.7065	-0.2521	0.7007
PQBP-1	Poly-glutamine tra	1631804_s_at	0.0589	0.7368	0.0568	0.5997	0.1497	0.3669	0.0698	0.8844	-0.1620	0.3290	-0.2318	0.1119	0.0044	0.9964	-0.1310	0.6068	-0.1354	0.5972
CG3397	CG3397	1631805_at	2.6333	0.0041	1.8495	0.0883	2.3354	0.0001	0.3318	0.6724	0.0828	0.8656	-0.2490	0.4825	-0.4125	0.8439	-0.5877	0.5071	-0.1752	0.8810
CG8693	CG8693	1631806_at	0.1189	0.6710	0.0192	0.9181	0.5382	0.0041	0.2997	0.5136	-0.2599	0.2872	-0.5596	0.0179	0.2148	0.8235	-0.0667	0.9144	-0.2815	0.4978
CG9386	CG9386	1631807_at	0.1444	0.5890	-0.1645	0.7512	0.0028	0.9926	-0.0107	0.9895	0.5652	0.0093	0.5759	0.0052	-0.1385	0.9421	0.1314	0.8760	0.2699	0.6748
---	---	1631808_at	0.2039	0.4477	0.2233	0.5412	0.2037	0.5153	-0.0866	0.8920	0.0453	0.8727	0.1319	0.5255	0.0015	0.9997	0.2523	0.5455	0.2508	0.5523
---	---	1631809_at	0.3006	0.0619	0.0767	0.7306	0.4868	0.0103	0.0807	0.9154	-0.0685	0.8163	-0.1492	0.5177	-0.1258	0.8461	-0.0399	0.9248	0.0859	0.7886
---	---	1631810_at	-0.0200	0.9492	-0.0314	0.9135	0.0745	0.6876	-0.0537	0.9558	0.0085	0.9823	0.0621	0.8340	-0.0633	0.9467	-0.0599	0.8867	0.0034	0.9947
---	---	1631811_at	0.0865	0.6803	0.0753	0.4988	-0.1025	0.6482	-0.1234	0.8244	-0.1213	0.6087	0.0021	0.9938	0.0690	0.9400	-0.0240	0.9590	-0.0930	0.7808
CG7295	CG7295	1631812_at	0.1526	0.2832	0.1655	0.3848	-0.0270	0.8865	-0.1504	0.7221	-0.1471	0.4589	0.0033	0.9889	0.0645	0.9309	-0.0647	0.8425	-0.1292	0.6186
CG3224	CG3224	1631813_at	0.2360	0.3271	0.1515	0.5849	0.4378	0.0330	0.1227	0.8251	-0.6196	0.0112	-0.7423	0.0030	-0.1824	0.8472	-0.5803	0.1382	-0.3979	0.3227
CG10738	CG10738	1631814_a_at	0.3091	0.2360	0.2953	0.1926	0.4207	0.0252	0.1498	0.8046	-0.1153	0.6755	-0.2651	0.2204	-0.0701	0.9016	-0.1946	0.3368	-0.1245	0.5733
CG14689	CG14689	1631815_at	0.3326	0.0875	-0.0291	0.7958	0.0853	0.6802	0.1497	0.7115	0.2466	0.1769	0.0969	0.5994	0.0746	0.9309	-0.0193	0.9664	-0.0939	0.7749
Akh	adipokinetic horm	1631816_at	0.0315	0.9020	0.1010	0.4849	0.1448	0.3854	0.1091	0.8155	0.0551	0.8080	-0.0540	0.7932	0.0369	0.9778	0.0229	0.9650	-0.0140	0.9777
Got1	Glutamate oxaloa	1631817_a_at	0.1154	0.4387	0.4043	0.0755	0.5213	0.0625	0.1437	0.8034	-0.7887	0.0053	-0.9324	0.0015	-0.0041	0.9978	-0.4055	0.2156	-0.4014	0.2527
---	---	1631818_at	-0.0004	0.9985	-0.0492	0.7943	0.0516	0.8071	0.1086	0.8485	0.0790	0.7486	-0.0296	0.9062	0.0769	0.8940	0.1408	0.5288	0.0639	0.8111
RpL29	60S ribosomal prc	1631819_s_at	0.3844	0.0535	1.6755	0.0120	1.5512	0.0004	0.1697	0.6820	-0.9368	0.0012	-1.1065	0.0004	0.1297	0.8875	0.0978	0.8301	-0.0319	0.9471
CG13747	CG13747	1631820_at	0.2141	0.2992	0.0721	0.7052	0.3721	0.1087	0.0651	0.9306	-0.0259	0.9330	-0.0909	0.6920	-0.1016	0.9238	-0.1614	0.6798	-0.0598	0.9023
Sr-CI	Scavenger Recep	1631821_at	-2.6329	0.0080	-3.9245	0.0108	-4.3988	0.0000	-0.4565	0.1545	-0.1131	0.5922	0.3434	0.0521	0.0187	0.9964	-1.3924	0.1886	-1.4110	0.2136
mus205	mutagen-sensitive	1631822_at	-0.0325	0.9144	-0.6807	0.0900	-0.5257	0.0057	0.0127	0.9872	0.5903	0.0076	0.5776	0.0049	-0.2087	0.8541	-0.0554	0.9409	0.1533	0.7815
CG34350 /// DmirCG8181	CG11824	1631823_at	0.0057	0.9830	0.2027	0.1916	0.0504	0.7936	-0.1225	0.8190	-0.1407	0.5296	-0.0181	0.9458	0.1582	0.7322	0.0628	0.8045	-0.0953	0.6567
CG6599	CG6599	1631824_at	0.1373	0.4494	0.1229	0.5426	-0.0607	0.7258	0.1765	0.6005	0.1326	0.4523	-0.0439	0.8194	0.2407	0.7707	0.1363	0.7424	-0.1044	0.8111
CG32167	CG32167	1631825_at	0.0170	0.9413	0.0676	0.7655	-0.0856	0.5738	0.0149	0.9796	0.0448	0.7980	0.0298	0.8574	0.0145	0.9913	-0.0691	0.8566	-0.0836	0.8056
---	---	1631826_at	0.1834	0.2018	0.0353	0.7334	0.1666	0.3243	0.0237	0.9704	0.0995	0.5695	0.0758	0.6501	0.0395	0.9467	0.0740	0.7200	0.0344	0.8896
CG15863 /// DereCG15863	CG15863	1631827_at	0.0994	0.6916	0.0457	0.7263	-0.2608	0.1092	-0.0748	0.9319	0.2124	0.4308	0.2872	0.2179	0.2183	0.7588	0.1413	0.6681	-0.0770	0.8395
CG8830 /// DyakCG8830	CG8830	1631828_s_at	0.2420	0.6235	0.2828	0.4376	0.0337	0.9162	-0.0345	0.9777	0.6180	0.0523	0.6526	0.0272	0.2955	0.8744	0.5911	0.4191	0.2956	0.7235
CG15719	CG15719	1631829_at	0.1628	0.2882	0.0647	0.6072	0.0726	0.6625	-0.0438	0.9384	0.0941	0.6064	0.1379	0.3688	-0.0690	0.9004	-0.0030	0.9943	0.0661	0.7836
---	---	1631830_at	-0.1263	0.5826	-0.0613	0.5749	0.2063	0.2151	0.0731	0.8738	-0.1940	0.2239	-0.2671	0.0649	-0.1482	0.7644	-0.0124	0.9716	0.1357	0.5345
Cog7	Cog7	1631831_at	0.6909	0.0271	0.4140	0.3961	0.2578	0.1767	0.2279	0.5633	0.8827	0.0018	0.6548	0.0040	0.3866	0.7686	0.6719	0.2145	0.2853	0.6354
Mlc-c	alkali light chain o	1631832_at	-0.1080	0.5885	0.6142	0.0100	0.5608	0.0189	0.0218	0.9774	-0.4520	0.0234	-0.4737	0.0120	0.1289	0.8510	0.2587	0.3500	0.1298	0.6770
mod(mdg4)	Modifier67.2	1631833_at	0.1271	0.5046	-0.1931	0.2640	-0.0986	0.5549	-0.0711	0.9042	-0.1409	0.4842	-0.0698	0.7355	-0.1435	0.8270	-0.3264	0.2181	-0.1828	0.5257
CG31098	CG31098	1631834_at	-1.3930	0.0071	-0.1416	0.8380	-1.5753	0.0004	-0.4062	0.3925	-0.8820	0.0062	-0.4758	0.0514	1.0473	0.4913	0.4790	0.5338	-0.5683	0.4582
CG6300	CG6300	1631835_at	0.0988	0.5803	-0.0412	0.7905	0.0067	0.9728	0.0466	0.9254	0.2624	0.0804	0.2158	0.1055	-0.0681	0.9142	-0.0487	0.8783	0.0194	0.9505
CG17568	CG17568	1631836_at	-0.0813	0.7712	0.0521	0.8201	0.0565	0.8147	-0.3222	0.4383	0.1163	0.6496	0.4385	0.0377	-0.1535	0.8882	0.2769	0.5140	0.4304	0.3048
Top2	topoisomerase II	1631837_at	0.1489	0.4686	0.2526	0.3885	0.6297	0.0594	0.0012	0.9988	0.1794	0.3683	0.1782	0.3157	-0.3592	0.7464	0.2888	0.5591	0.6480	0.1931
---	---	1631838_at	0.0373	0.9036	0.2266	0.1985	0.3277	0.1125	-0.0263	0.9728	-0.1369	0.4971	-0.1106	0.5593	-0.1292	0.8940	0.0718	0.8940	0.2010	0.6065
CG40169	CG40169	1631839_at	0.1928	0.2737	0.0186	0.8553	0.3253	0.0490	0.0921	0.7979	0.0642	0.7023	-0.0279	0.8713	-0.1089	0.8400	0.0212	0.9528	0.1302	0.5834
---	---	1631840_at	-0.0028	0.9886	-0.0149	0.9432	-0.1381	0.3725	-0.0422	0.9370	0.0573	0.7574	0.0995	0.5058	0.1775	0.6749	-0.0024	0.9949	-0.1799	0.3175
---	---	1631841_at	0.0212	0.9251	0.1951	0.5184	0.2164	0.1830	-0.0006	0.9994	-0.0526	0.7625	-0.0520	0.7393	0.0103	0.9948	0.0878	0.8625	0.0775	0.8745
---	---	1631842_at	0.0809	0.6186	0.1430	0.4234	-0.0018	0.9948	-0.1021	0.8389	0.0173	0.9513	0.1194	0.5111	-0.0083	0.9939	-0.0923	0.7400	-0.0840	0.7583
CG6749	CG6749	1631843_at	0.9234	0.1043	0.3393	0.2358	1.0705	0.0012	0.3160	0.5039	-0.1784	0.4994	-0.4944	0.0346	-0.0497	0.9852	-0.5341	0.3749	-0.4844	0.4434
Myo95E	Myosin 95E	1631844_at	0.0393	0.9405	-0.1720	0.5508	-0.4579	0.0708	-0.1751	0.7979	0.2813	0.3152	0.4564	0.0684	0.0482	0.9831	-0.0770	0.9271	-0.1252	0.8559
---	---	1631845_at	0.0913	0.5511	0.0009	0.9989	0.0612	0.7577	-0.0120	0.9852	0.0330	0.8617	0.0450	0.7789	-0.0453	0.9708	-0.0413	0.9341	0.0040	0.9945
CG14857	CG14857	1631846_at	-0.0098	0.9732	0.0109	0.9607	-0.0824	0.7042	0.0342	0.9603	-0.0628	0.7780	-0.0970	0.5999	0.0445	0.9710	-0.0889	0.8258	-0.1334	0.6978
---	---	1631847_at	0.0959	0.5564	0.1176	0.4798	0.0797	0.7352	0.0253	0.9761	-0.1252	0.5822	-0.1505	0.4463	0.1551	0.8755	0.0931	0.8633	-0.0620	0.9080
CG9235	CG9235	1631848_at	0.2804	0.2042	-0.0597	0.5971	-0.2554	0.1985	-0.0135	0.9889	0.0052	0.9880	0.0187	0.9476	0.2036	0.6592	-0.2985	0.1178	-0.5022	0.0422
heix	heixuedian	1631849_at	0.0895	0.7518	-0.1073	0.8340	-0.8266	0.0090	-0.0688	0.9319	1.0613	0.0017	1.1302	0.0008	0.5039	0.7116	0.7459	0.1767	0.2420	0.7049
CG40120	CG40120	1631850_at	0.1008	0.7583	-0.3736	0.1326	-0.2603	0.1178	0.2788	0.3744	0.4631	0.0173	0.1842	0.2481	-0.0994	0.9324	-0.1301	0.7787	-0.0307	0.9541
CG34354	CG12425	1631851_at	0.1849	0.3997	0.1208	0.4510	0.1248	0.7042	-0.1285	0.9116	-0.1326	0.7639	-0.0041	0.9928	0.0244	0.9831	-0.0993	0.7549	-0.1237	0.6719
---	---	1631852_at	0.1607	0.4041	-0.0237	0.8223	0.0755	0.7971	-0.0477	0.9704	0.1939	0.5837	0.2416	0.4301	0.0321	0.9717	0.0462	0.8864	0.0141	0.9661
CG6610	CG6610	1631853_at	0.4074	0.1357	0.7185	0.0755	0.6144	0.0040	-0.0447	0.9558	0.0689	0.7955	0.1135	0.6023	0.1932	0.8362	0.3785	0.3134	0.1853	0.6577
---	---	1631854_s_at	-0.0363	0.8294	0.0596	0.6776	0.0156	0.9617	-0.1633	0.7817	-0.1919	0.4499	-0.0286	0.9240	-0.0786	0.8751	-0.0762	0.7413	0.0024	0.9945
CG10864	CG10864	1631855_at	0.2428	0.4494	0.3353	0.1347	0.2674	0.1543	-0.2804	0.4858	-0.1360	0.5565	0.1444	0.4804	0.1133	0.9309	0.0993	0.8684	-0.0140	0.9846
RFeSP	Rieske iron-sulfur	1631856_a_at	-0.5062	0.0265	-0.0197	0.9791	-0.0555	0.8618	-0.1219	0.8678	-0.9731	0.0034	-0.8513	0.0037	-0.0256	0.9923	-0.4184	0.5100	-0.3928	0.5455
fd96Ca	forkhead domain (	1631857_at	-1.6190	0.0062	-1.3104	0.0093	-1.6387	0.0030	0.2029	0.8822	-0.0139	0.9839	-0.2168	0.6540	0.0337	0.9868	-0.1943	0.7111	-0.2280	0.6475
---	---	1631858_at	-0.0481	0.7739	0.0242	0.8160	0.4129	0.0419	0.2111	0.4975	0.0894	0.6252	-0.1217	0.4314	-0.0055	0.9964	0.1871	0.5471	0.1926	0.5399
CG33295	CG33295	1631859_at	0.2051	0.3776	0.1905	0.2290	0.6444	0.0042	0.0698	0.8844	-0.1780	0.2797	-0.2478	0.0905	-0.2611	0.7485	-0.0656	0.9009	0.1955	0.6005

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9447	CG9447	1631860_at	0.7103	0.0094	0.1534	0.3977	0.1157	0.6093	-0.1389	0.8732	-0.0256	0.9546	0.1132	0.7255	-0.0241	0.9848	-0.4041	0.1512	-0.3800	0.2094
cdc2	CDC2 kinase	1631861_at	0.3692	0.6135	-0.9854	0.3445	-1.3108	0.1696	-0.6421	0.4068	1.2785	0.0097	1.9207	0.0010	-0.5207	0.9260	-0.1934	0.9494	0.3274	0.8984
robl22E	robl22E	1631862_at	0.1110	0.4463	0.1675	0.3697	0.1831	0.3925	0.1364	0.7167	0.0107	0.9678	-0.1256	0.4345	0.0659	0.9503	0.0885	0.8335	0.0226	0.9612
chm	chameau	1631863_at	0.4325	0.1687	0.0448	0.8190	-0.0845	0.6953	-0.2101	0.6566	0.6315	0.0132	0.8417	0.0022	0.0066	0.9964	0.3355	0.3465	0.3289	0.3807
CG13713	CG13713	1631864_at	-0.0488	0.8388	0.0732	0.5918	0.0414	0.8409	0.0947	0.9060	-0.1075	0.7214	-0.2022	0.4010	0.1109	0.8086	0.0470	0.8586	-0.0639	0.7761
CG13064	CG13064	1631865_at	0.2187	0.2810	0.1894	0.4350	0.3570	0.0539	0.0933	0.9160	0.0954	0.7743	0.0021	0.9953	-0.1196	0.8882	-0.1740	0.6118	-0.0544	0.9025
CG14309	CG14309	1631866_at	0.1260	0.4485	-0.0639	0.7512	-0.0429	0.8466	-0.0248	0.9783	0.0248	0.9411	0.0496	0.8544	-0.0255	0.9848	-0.0671	0.8764	-0.0416	0.9198
ldgf5	Imaginal disc grov	1631867_at	1.8140	0.0115	1.1925	0.2045	2.1151	0.0000	-0.0863	0.9558	-0.2014	0.6748	-0.1151	0.8077	-0.7813	0.6927	-0.8956	0.2518	-0.1144	0.9189
Vap-33-1	Vap-33-1	1631868_at	-0.0469	0.9079	0.4144	0.3294	-0.2709	0.4453	-0.0483	0.9757	-0.0193	0.9746	0.0291	0.9533	0.4718	0.7322	0.3622	0.5585	-0.1096	0.8923
CG14053	CG14053	1631869_at	0.2821	0.0842	0.0884	0.5788	-0.0179	0.9408	0.0199	0.9753	0.1267	0.4344	0.1068	0.4724	0.0607	0.9467	-0.0723	0.8476	-0.1331	0.6567
CG5937 /// DmirCG5937	CG5937	1631870_at	-0.0907	0.6439	0.0936	0.6307	0.1152	0.4429	0.1257	0.7167	0.0130	0.9557	-0.1128	0.4483	0.0182	0.9913	0.0618	0.9037	0.0437	0.9243
ACXC	ACXC	1631871_at	0.1143	0.6553	-0.0185	0.8703	-0.1035	0.6638	0.0324	0.9711	0.1820	0.4378	0.1497	0.4895	-0.0395	0.9746	-0.0375	0.9404	0.0020	0.9977
CG7755	CG7755	1631872_at	0.0303	0.9088	0.0529	0.6148	0.1975	0.3902	0.1329	0.8590	0.0500	0.8910	-0.0829	0.7826	-0.0097	0.9939	0.1915	0.5018	0.2012	0.4825
CG3706	CG3706	1631873_at	-0.0382	0.9008	-0.0980	0.6500	0.0283	0.8832	0.0652	0.9011	0.0527	0.7998	-0.0125	0.9539	-0.1505	0.8882	-0.2017	0.6449	-0.0513	0.9271
---	---	1631874_at	0.0888	0.7453	-0.7504	0.0444	-0.5854	0.0349	0.2814	0.6169	0.9984	0.0044	0.7170	0.0118	0.0184	0.9913	-0.1270	0.7509	-0.1454	0.7007
CG1172 /// DbuzCG1172	CG1172	1631875_at	0.3435	0.4680	0.3370	0.2843	-0.0701	0.6412	-0.4208	0.5311	0.3097	0.3891	0.7304	0.0277	0.0381	0.9872	0.2888	0.6209	0.2507	0.6731
CG15266	CG15266	1631876_at	0.1251	0.7252	0.0041	0.9949	-0.1387	0.4098	0.1296	0.8817	0.5903	0.0519	0.4607	0.0840	0.2719	0.8425	0.5326	0.3300	0.2607	0.6711
Aats-his	Histidyl-tRNA synt	1631877_a_at	-0.2163	0.3965	0.6342	0.2410	0.9113	0.0042	0.1675	0.6856	-0.1916	0.3266	-0.3591	0.0431	-0.0077	0.9978	0.6260	0.3259	0.6337	0.3467
CG40451	CG40451	1631878_at	0.0711	0.7169	0.0221	0.8936	-0.1184	0.5585	-0.1055	0.8028	0.0656	0.7425	0.1711	0.2612	0.0335	0.9826	0.0504	0.9277	0.0169	0.9762
---	---	1631879_at	-0.0258	0.8973	-0.1559	0.4879	0.0398	0.8575	0.1663	0.6214	0.2089	0.2107	0.0427	0.8225	-0.1238	0.8425	-0.0216	0.9590	0.1022	0.7253
CG1492	CG1492	1631880_at	-0.5148	0.1416	-0.4024	0.2381	-0.9191	0.0203	-0.3249	0.5357	0.0286	0.9432	0.3534	0.1473	0.1417	0.9095	-0.1283	0.8255	-0.2700	0.5653
CG33464 /// sli	Slit /// CG33464	1631881_a_at	0.1037	0.5634	0.0761	0.6037	0.0428	0.8708	-0.0151	0.9834	-0.0516	0.8014	-0.0365	0.8507	0.0966	0.8940	-0.0078	0.9894	-0.1044	0.7375
CG10264	CG10264	1631882_at	0.0107	0.9715	0.0193	0.8724	-0.0076	0.9717	-0.1881	0.6790	-0.0793	0.7542	0.1088	0.6130	-0.0441	0.9646	0.0838	0.8080	0.1280	0.6597
---	---	1631883_at	0.0582	0.6886	0.1152	0.4356	0.1761	0.3218	0.0502	0.9459	-0.0099	0.9758	-0.0601	0.7981	-0.0183	0.9898	0.0453	0.9258	0.0636	0.8779
---	---	1631884_at	0.0812	0.7259	0.0618	0.5408	-0.3240	0.0509	-0.1934	0.6455	-0.1073	0.6381	0.0860	0.6873	0.1385	0.7953	-0.0035	0.9941	-0.1420	0.5488
CG4290	CG4290	1631885_at	-0.5087	0.2499	0.1067	0.8212	-0.1474	0.4687	-0.0579	0.9401	-0.2422	0.2793	-0.1843	0.3698	0.2692	0.8608	0.4147	0.5158	0.1454	0.8562
CG14135	CG14135	1631886_at	-0.1105	0.6049	-0.0484	0.6273	-0.1427	0.4733	-0.0810	0.8947	-0.2215	0.2794	-0.1405	0.4670	0.0301	0.9742	-0.1009	0.6962	-0.1311	0.5958
CG14718 /// DmauCG1471	CG14718	1631888_at	-0.0751	0.7274	-0.0922	0.6842	0.0268	0.9061	0.0356	0.9683	0.0615	0.8377	0.0259	0.9278	-0.1109	0.8461	-0.1034	0.6958	0.0075	0.9847
CG31752	CG31752	1631889_at	-0.1406	0.6437	-0.0765	0.4949	0.0753	0.7365	0.2285	0.5842	0.1313	0.5678	-0.0972	0.6590	0.0608	0.9495	0.0767	0.8432	0.0159	0.9714
CG6488 /// DyakCG6488	CG6488	1631891_at	0.5063	0.0305	0.9764	0.0895	0.8066	0.0027	0.1038	0.8518	0.1591	0.4491	0.0552	0.8078	0.2723	0.8202	0.6724	0.1621	0.4002	0.4303
CG32387 /// DmirCG32387	CG32387	1631893_at	0.1016	0.6855	-0.2804	0.1272	0.0508	0.8139	-0.0246	0.9803	0.0609	0.8439	0.0855	0.7397	-0.2048	0.6955	-0.1677	0.4426	0.0372	0.9025
---	---	1631895_s_at	-0.0211	0.9079	0.0542	0.6494	-0.0449	0.8155	-0.0905	0.8587	-0.0296	0.9074	0.0610	0.7610	-0.0136	0.9862	-0.0036	0.9935	0.0100	0.9716
Nup44A	nucleoporin-44A	1631897_s_at	0.8614	0.0049	0.3287	0.0900	0.5304	0.0126	0.0110	0.9909	0.2860	0.1691	0.2750	0.1400	-0.2440	0.6749	-0.2346	0.3146	0.0094	0.9816
CG6000	CG6000	1631899_a_at	0.1122	0.7169	0.4035	0.2410	0.3929	0.1361	-0.2052	0.7080	-0.4931	0.0533	-0.2879	0.1969	-0.2857	0.8344	-0.2325	0.7213	0.0531	0.9470
---	---	1631901_at	0.1166	0.4383	-0.1157	0.3142	-0.2978	0.0982	-0.0234	0.9698	0.2021	0.2014	0.2255	0.1111	-0.0764	0.8740	-0.0981	0.6312	-0.0217	0.9346
CG14062	CG14062	1631903_at	0.0014	0.9942	-0.0301	0.7672	0.1406	0.4020	-0.0187	0.9814	0.1782	0.3632	0.1969	0.2541	-0.1011	0.8480	-0.0071	0.9875	0.0941	0.7022
pyd	tamou	1631905_a_at	0.3014	0.4386	0.5897	0.1196	0.3241	0.2519	-0.5090	0.0840	-0.3447	0.0484	0.1643	0.2868	0.3102	0.8815	0.4289	0.6154	0.1187	0.9145
CG6038	CG6038	1631907_at	-0.3672	0.2181	0.0455	0.7496	-0.2708	0.2471	0.1738	0.7271	-0.1181	0.6357	-0.2919	0.1416	0.4606	0.6749	0.3979	0.3706	-0.0626	0.9197
bai	baiser	1631909_at	0.6344	0.0116	0.5954	0.0171	0.9325	0.0004	0.1451	0.6891	0.2409	0.1504	0.0958	0.5652	-0.1187	0.8745	0.3339	0.2430	0.4526	0.1522
---	---	1631911_at	0.0916	0.6791	0.1052	0.3742	0.2438	0.1086	-0.0013	0.9985	0.0254	0.8965	0.0267	0.8744	-0.1442	0.8339	-0.0887	0.8117	0.0555	0.8880
CG10301	CG10301	1631913_at	0.2681	0.2859	0.0044	0.9836	0.3523	0.0640	0.1122	0.7511	0.0570	0.7556	-0.0552	0.7385	-0.1889	0.7644	-0.1899	0.4810	-0.0010	0.9986
CG17776	CG17776	1631915_at	-0.2272	0.4592	0.2624	0.3690	0.3937	0.1032	0.0522	0.9313	-0.6135	0.0051	-0.6657	0.0021	0.0151	0.9950	-0.0267	0.9783	-0.0418	0.9575
Dys	Dystrophin-like pr	1631917_at	-0.1000	0.7261	-0.1737	0.4379	0.0391	0.8468	0.1651	0.8076	0.0842	0.7998	-0.0809	0.7889	-0.0700	0.9238	0.0915	0.7525	0.1615	0.5299
CG17078	CG17078	1631919_at	0.1120	0.7157	0.0033	0.9829	-0.0365	0.8551	-0.0553	0.9518	0.2924	0.2465	0.3477	0.1211	-0.0959	0.9585	0.2012	0.7362	0.2971	0.5862
CG14244	CG14244	1631921_at	0.2181	0.3629	0.2118	0.2940	-0.3258	0.2300	-0.2385	0.6506	0.0138	0.9720	0.2524	0.2712	0.1246	0.8651	-0.0734	0.8550	-0.1980	0.5167
---	---	1631923_at	0.0424	0.8391	0.0004	1.0000	-0.0335	0.8905	-0.0638	0.9345	0.1138	0.6610	0.1776	0.4061	-0.0286	0.9653	0.0483	0.8380	0.0768	0.6947
Mdr65	P-glycoprotein65-l	1631925_at	-0.4689	0.0787	0.4077	0.3996	-0.3541	0.1583	-0.3415	0.5631	-1.2636	0.0022	-0.9221	0.0053	0.5151	0.6749	-0.2455	0.6632	-0.7606	0.1618
---	---	1631927_at	0.2776	0.1248	0.4013	0.0179	-0.1485	0.5480	-0.2217	0.4861	-0.1723	0.3162	0.0494	0.7964	0.3558	0.6749	0.0285	0.9609	-0.3273	0.3622
---	---	1631929_at	0.3175	0.1982	0.2177	0.3902	-0.0528	0.8170	-0.0170	0.9860	0.0097	0.9776	0.0268	0.9218	0.0597	0.9506	-0.0739	0.8500	-0.1336	0.6683
Sdc	Dsyndecan	1631931_s_at	-0.7862	0.1021	-0.8228	0.0177	-1.0228	0.0032	-0.0529	0.9558	0.2714	0.2972	0.3243	0.1586	-0.2085	0.9092	0.0188	0.9895	0.2273	0.7638
CG8526	CG8526	1631933_at	-0.2876	0.6700	-1.1948	0.0574	-0.8671	0.0570	0.1605	0.9364	0.8729	0.1361	0.7124	0.1745	-0.3573	0.8270	-0.1320	0.8963	0.2253	0.7774
CG32487	CG32487	1631935_at	-0.0019	0.9930	-0.0468	0.8767	-0.1468	0.4179	-0.0543	0.9351	-0.0230	0.9338	0.0313	0.8949	0.0446	0.9717	-0.0231	0.9650	-0.0677	0.8764
CG31755	CG31755	1631937_at	0.0018	0.9961	-0.5850	0.0596	-0.5568	0.0295	-0.2292	0.5854	0.3949	0.0681	0.6241	0.0064	-0.2373	0.8657	-0.2535	0.6858	-0.0161	0.9855
---	---	1631938_at	0.1617	0.4675	-0.0028	0.9877	0.0456	0.7723	0.2149	0.6238	0.3088	0.1531	0.0939	0.6794	0.1517	0.7707	0.1086	0.6536	-0.0432	0.8879
Cp190	Centrosomal prote	1631940_s_at	-0.0071	0.9806	0.3543	0.1000	0.2160	0.2356	-0.1106	0.8578	-0.0633	0.8190	0.0473	0.8549	0.0440	0.9679	0.3308	0.2380	0.2867	0.3335
CG9425 /// DmirCG9425	CG9425	1631942_at	-0.5115	0.4038	-0.3441	0.7554	-0.2238	0.5279	0.1798	0.7138	0.2617	0.2419	0.0819	0.7326	0.3357	0.9409	0.6069	0.7020	0.2711	0.8884
---	---	1631944_at	0.1376	0.6627	0.0385	0.7014	0.1069	0.5002	0.0047	0.9943	0.0916	0.5503	0.0869	0.5318	-0.0565	0.9503	0.0309	0.9438	0.0874	0.7855
CG7173	CG7173	1631946_at	-0.0371	0.8664	0.1744	0.2288	0.2844	0.1043	-0.0621	0.9592	-0.1431	0.6962	-0.0811	0.8228	-0.0038	0.9978	0.0925	0.8218	0.0963	0.8007
Clk	dclock	1631948_s_at	0.2947	0.1908	-0.2778	0.7877	-0.7650	0.1426	-0.3105	0.2592	0.8652	0.0008	1.1757	0.0002	0.1070	0.9816	0.1883	0.9105	0.0814	0.9592
---	---	1631950_at	-0.0294	0.8804	0.1263	0.4262	-0.1632	0.3185	-0.2498	0.4734	-0.2605	0.1623	-0.0108	0.9637	-0.1150	0.8014	-0.0978	0.6350	0.0172	0.9487
---	---	1631952_at	0.3225	0.1485	0.2235	0.1149	0.1428	0.3285	0.0673	0.9247	0.1620	0.4675	0.0947	0.6697	0.1424	0.7485	0.0971	0.6344	-0.0453	0.8527
CG30375	CG30375	1631953_at	4.0003	0.0002	4.0749	0.0029	4.7729	0.0000	1.5173	0.0018	1.3155	0.0003	-0.2018	0.2288	0.8223	0.6557	1.4369	0.0694	0.6145	0.4142
Ranbp9	Ranbp9	1631954_at	0.7362	0.0087	0.1669	0.7402	0.4603	0.0570	0.2854	0.3056	0.8800	0.0008	0.5946	0.0024	-0.0508	0.9816	0.3547	0.5074	0.4055	0.4471
CG15145	CG15145	1631956_at	0.1197	0.5470	0.0784	0.5991	0.2882	0.1360	0.0519	0.9528	0.0023	0.9947	-0.0497	0.8605	-0.1451	0.8157	-0.1109	0.7027	0.0342	0.9231
CG40204	CG40204	1631958_at	0.2818	0.1944	0.3015	0.0811	-0.0276	0.9079	-0.1716	0.6869	0.0355	0.8953	0.2071	0.2440	0.2130	0.7485	0.1620	0.5889	-0.0510	0.8949
CG31526	CG31526	1631960_at	0.3497	0.0997	0.0393	0.8779	0.2125	0.3647	0.0793	0.9182	0.0214	0.9517	-0.0579	0.8300	-0.1159	0.8802	-0.2185	0.4609	-0.1026	0.7640
MtnD	Metallothionein D	1631962_at	-0.8280	0.0922	-0.6272	0.0521	-2.0158	0.0006	-0.7376	0.2876	0.2661	0.5452	1.0037	0.0139	-0.2320	0.8222	-0.3785	0.3706	-0.1464	0.7729
CG18262	CG18262	1631964_at	-0.0322	0.9220	0.0663	0.5743	0.6073	0.0356	0.2070	0.7477	-0.0692	0.8490	-0.2762	0.2765	-0.1854	0.8344	0.2081	0.5969	0.3936	0.2988
CG8005	CG8005	1631965_at	0.6164	0.0616	0.8503	0.0755	1.2868	0.0002	0.3507	0.3147	0.2743	0.1759	-0.0764	0.7255	0.0716	0.9653	0.6156	0.1512	0.5440	0.2338

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1631967_at	0.1782	0.2543	0.1397	0.3697	0.3156	0.3130	-0.1390	0.8725	-0.1754	0.5932	-0.0363	0.9218	0.1103	0.9340	0.0247	0.9724	-0.0855	0.8831
CG15221	CG15221	1631969_at	0.1216	0.7052	0.1176	0.5772	-0.0124	0.9533	-0.0729	0.9475	-0.1022	0.7887	-0.0293	0.9397	0.1153	0.9092	-0.1631	0.6844	-0.2784	0.4540
CG10907	CG10907	1631971_at	-0.0519	0.8856	-0.0025	0.9919	-0.1711	0.2664	-0.1173	0.8196	0.2046	0.3152	0.3219	0.0757	-0.1013	0.9238	0.1221	0.7786	0.2234	0.5490
Ote	otefin	1631973_at	-0.3142	0.1912	-0.4360	0.2816	-0.5837	0.0847	-0.3362	0.2438	0.1787	0.3011	0.5148	0.0056	-0.4119	0.8400	-0.0316	0.9843	0.3803	0.6827
CG15628	CG15628	1631975_at	0.3478	0.0613	-0.0512	0.6998	0.3473	0.0960	0.2191	0.6591	0.2084	0.3948	-0.0107	0.9713	-0.0428	0.9682	0.0864	0.8137	0.1292	0.6776
PH4alphaSG1	prolyl-4-hydroxyla	1631977_at	0.0769	0.6353	0.1164	0.6127	0.2894	0.0974	0.0535	0.9413	-0.0578	0.8251	-0.1112	0.5934	-0.0487	0.9640	0.0324	0.9454	0.0812	0.8247
MICAL-like	MICAL-like	1631979_at	-0.0817	0.6908	-0.1947	0.4857	-0.6988	0.0116	-0.0625	0.9252	0.5659	0.0120	0.6284	0.0045	0.4990	0.6749	0.5368	0.2461	0.0378	0.9563
CG14457	CG14457	1631981_at	-0.1252	0.5692	-0.0230	0.8413	0.2180	0.1816	0.1060	0.8827	-0.0395	0.9077	-0.1455	0.5446	-0.2853	0.5074	-0.1007	0.6495	0.1846	0.3779
CG30320	CG30320	1631983_at	0.1110	0.6479	-0.0559	0.6009	0.0992	0.5011	0.0834	0.9009	0.0457	0.8732	-0.0377	0.8831	-0.0552	0.9400	-0.1352	0.5647	-0.0800	0.7567
CG3639	CG3639	1631984_at	0.4432	0.2020	-0.0226	0.9096	0.0944	0.5135	-0.1296	0.8164	0.1377	0.5611	0.2673	0.1728	-0.1831	0.8270	-0.2227	0.5414	-0.0396	0.9353
CG16998	CG16998	1631986_at	0.1757	0.2835	0.0920	0.6221	0.3576	0.0667	0.1639	0.7271	0.1515	0.4929	-0.0125	0.9625	0.0479	0.9458	0.1525	0.4847	0.1046	0.6486
CG15555	CG15555	1631988_at	0.0400	0.8690	-0.0135	0.9130	-0.0520	0.8113	0.0941	0.8856	0.0425	0.8873	-0.0517	0.8403	0.0135	0.9914	0.0667	0.8564	0.0532	0.8850
Or33b	Olfactory receptor	1631990_at	-0.0297	0.9262	0.0833	0.6696	-0.0086	0.9729	0.0264	0.9797	0.1316	0.6423	0.1052	0.6914	0.0020	0.9990	0.0751	0.8151	0.0730	0.8112
CG5089	CG5089	1631991_at	0.2315	0.3251	0.0574	0.6062	0.3343	0.0900	0.1237	0.7556	0.0176	0.9445	-0.1061	0.5244	-0.0843	0.9457	-0.1429	0.7454	-0.0586	0.9095
Ald	aldolase	1631993_s_at	-0.2282	0.5399	0.0795	0.7487	-0.2260	0.3695	-0.0747	0.9300	-0.2566	0.3071	-0.1819	0.4365	0.2019	0.8736	0.0044	0.9977	-0.1975	0.7287
Mocs1	Mocs1	1631994_a_at	-2.2010	0.0005	-2.1771	0.0049	-2.2782	0.0000	-0.1765	0.7589	-0.2696	0.2776	-0.0931	0.7237	-0.0316	0.9816	-0.1948	0.5538	-0.1632	0.6311
mRpS34	mitochondrial ribo	1631996_at	-0.1595	0.3872	0.2436	0.4017	-0.1457	0.3531	-0.0822	0.8921	-0.2617	0.1951	-0.1795	0.3334	0.3291	0.6749	0.0666	0.8857	-0.2625	0.4226
CG3473	CG3473	1631998_at	0.2017	0.3938	0.0886	0.4685	0.2256	0.1569	0.0017	0.9987	-0.1243	0.5916	-0.1259	0.5460	-0.0746	0.8943	-0.1246	0.5754	-0.0501	0.8547
CG10710	CG10710	1631999_at	-2.0052	0.0030	-2.1386	0.0095	-2.6093	0.0001	-0.1337	0.7929	0.4037	0.0560	0.5374	0.0110	0.1281	0.9589	0.0598	0.9590	-0.0684	0.9445
CG7102	CG7102	1632001_at	-0.2319	0.1786	-0.1873	0.1609	0.1084	0.5985	-0.0320	0.9649	-0.0573	0.8122	-0.0253	0.9153	-0.2223	0.7491	-0.0663	0.8760	0.1560	0.6257
Ptp99A	Protein tyrosine pl	1632003_a_at	-1.2466	0.0409	-0.9899	0.0945	-1.2275	0.0013	0.1682	0.9011	0.2533	0.5954	0.0851	0.8681	0.4823	0.7215	0.4422	0.4303	-0.0401	0.9613
---	---	1632005_at	-0.0963	0.5473	0.0097	0.9702	0.1677	0.3139	0.0505	0.9353	0.0236	0.9271	-0.0269	0.9034	-0.0560	0.9220	0.0745	0.7381	0.1305	0.5095
CG7492	anon-fast-evolving	1632007_at	-0.2827	0.0892	0.1452	0.3996	-0.0037	0.9883	-0.0336	0.9532	-0.1984	0.2057	-0.1648	0.2430	-0.0026	0.9984	0.2903	0.2217	0.2929	0.2523
CG8044	CG8044	1632009_at	-0.2559	0.2362	0.0107	0.9204	-0.1183	0.4587	-0.2034	0.4998	-0.2927	0.0691	-0.0893	0.5724	0.0040	0.9964	-0.0294	0.9353	-0.0333	0.9159
CG31274 /// MESK4	CG31274 /// Mise	1632011_at	-0.4858	0.3543	-0.4903	0.0942	-1.8927	0.0020	-0.9038	0.1787	-0.3296	0.4344	0.5742	0.1130	0.3206	0.8943	-0.3382	0.7529	-0.6589	0.4824
CG40388 /// CG40390	CG40388 /// CG4(	1632013_s_at	-0.1794	0.4069	-0.0794	0.6094	0.0850	0.6152	-0.0692	0.9071	-0.0533	0.8240	0.0159	0.9478	-0.1146	0.8465	-0.0322	0.9350	0.0825	0.7761
beat-Vc	beat-Vc	1632015_at	0.1507	0.3621	0.0549	0.6366	0.2650	0.1651	0.1885	0.7254	0.0481	0.8812	-0.1404	0.5540	0.0865	0.8680	0.1696	0.4067	0.0831	0.7228
CG13990	CG13990	1632017_at	0.1197	0.6601	0.0694	0.5511	0.0600	0.8832	0.0049	0.9956	-0.0067	0.9822	-0.0117	0.9637	0.1010	0.9449	0.1119	0.8546	0.0109	0.9874
CG9008	CG9008	1632019_s_at	1.3495	0.0028	-1.0103	0.0719	-0.0046	0.9946	0.4544	0.4239	1.1470	0.0039	0.6927	0.0220	-0.4747	0.8049	-1.0903	0.1622	-0.6156	0.4584
Cyp6a20	Cyp6a20	1632021_at	-1.4441	0.0047	-1.0779	0.1332	-1.8766	0.0003	-0.9217	0.0465	-1.4784	0.0007	-0.5567	0.0263	-0.0282	0.9933	-0.9852	0.1906	-0.9570	0.2336
CG1909	CG1909	1632023_s_at	0.3388	0.5780	-0.1985	0.6997	1.1159	0.0217	0.2194	0.8794	0.1106	0.8656	-0.1088	0.8495	-1.0875	0.6077	-0.5126	0.5978	0.5749	0.5508
CG5180	CG5180	1632024_at	0.0088	0.9795	-0.3946	0.0832	-0.5758	0.0453	0.2040	0.6718	0.6338	0.0132	0.4298	0.0413	0.2643	0.7812	0.1019	0.8583	-0.1624	0.7307
CG4930	CG4930	1632026_at	-0.5126	0.0470	-0.4030	0.2938	-0.5108	0.0152	0.0149	0.9852	0.4437	0.0216	0.4288	0.0156	0.0107	0.9952	0.4583	0.2861	0.4476	0.3230
Gs1l	GS1-like protein	1632028_a_at	-0.1007	0.5738	0.6263	0.1449	0.8636	0.0102	-0.0844	0.9255	-1.2110	0.0017	-1.1266	0.0014	-0.2585	0.7932	-0.2979	0.4829	-0.0394	0.9463
CG31454	CG31454	1632030_at	0.5043	0.0649	-0.0114	0.9365	1.0482	0.0347	0.4712	0.1818	0.5178	0.0236	0.0466	0.8510	-0.5770	0.7046	-0.1344	0.8831	0.4426	0.5003
CG13188	CG13188	1632032_at	-0.4271	0.2249	-2.3175	0.0048	-1.7607	0.0003	0.4690	0.5902	1.4749	0.0058	1.0058	0.0191	-0.0349	0.9831	-0.2721	0.4875	-0.2372	0.5601
CG11737	CG11737	1632034_at	-0.2328	0.1927	0.2787	0.1047	0.4439	0.0497	-0.2867	0.4850	-0.7229	0.0061	-0.4362	0.0333	-0.3321	0.6301	-0.2395	0.3993	0.0926	0.7865
---	---	1632035_at	0.2684	0.1905	0.2395	0.3690	0.3100	0.1062	-0.1038	0.8500	-0.1690	0.4120	-0.0652	0.7670	-0.1102	0.8465	-0.0319	0.9328	0.0783	0.7790
---	---	1632037_at	0.0589	0.7766	0.0055	0.9645	0.1063	0.5509	0.0151	0.9819	0.0622	0.7364	0.0471	0.7880	-0.0371	0.9653	-0.0103	0.9808	0.0268	0.9341
CG4546	CG4546	1632039_at	-0.0463	0.8043	0.1368	0.3741	0.1157	0.5488	0.0369	0.9528	-0.1709	0.3275	-0.2077	0.1760	0.2152	0.7324	0.0334	0.9395	-0.1819	0.5181
DyakCG10731	CG10731	1632041_s_at	0.1713	0.3967	0.2153	0.5127	-0.0190	0.9315	-0.2639	0.4456	-0.0494	0.8391	0.2145	0.2034	0.0816	0.9411	0.1592	0.6762	0.0775	0.8637
shtd	shattered	1632043_a_at	0.7408	0.1410	0.4829	0.4356	0.6315	0.0141	0.0824	0.8932	0.5651	0.0136	0.4827	0.0165	-0.0377	0.9914	0.3021	0.7424	0.3399	0.6973
CG5697	CG5697	1632045_at	-0.2285	0.7423	0.0886	0.4475	0.0027	0.9916	-0.1134	0.9346	-0.4342	0.2921	-0.3208	0.3987	0.1486	0.9309	-0.3595	0.5324	-0.5081	0.3716
CG18271	CG18271	1632047_at	-0.3036	0.2050	-0.1093	0.4733	0.0642	0.7964	-0.0466	0.9475	-0.0703	0.7693	-0.0237	0.9230	-0.3686	0.5869	-0.1360	0.6876	0.2326	0.4569
CG13126	CG13126	1632049_at	-0.1677	0.4066	0.1675	0.6277	0.9476	0.0035	-0.1227	0.8189	-0.5561	0.0160	-0.4334	0.0283	-0.6033	0.5765	0.1162	0.8658	0.7195	0.1651
CG4119	CG4119	1632051_at	-0.0964	0.7849	0.3525	0.6029	0.7581	0.0255	-0.0232	0.9854	-0.0282	0.9469	-0.0050	0.9887	-0.3863	0.8521	0.4040	0.6695	0.7903	0.3695
CG15541	CG15541	1632053_at	0.2270	0.2726	0.0144	0.9022	0.0178	0.9593	0.1420	0.6718	0.1149	0.4901	-0.0271	0.8856	0.0557	0.9717	-0.0963	0.8574	-0.1520	0.7333
kappaTry	kappaTry	1632055_at	-0.0164	0.9842	0.0093	0.9817	0.0044	0.9879	0.2238	0.8102	-0.1538	0.7215	-0.3776	0.2570	0.1610	0.9375	-0.1641	0.8527	-0.3251	0.6389
bchs	beached	1632057_at	0.4965	0.3043	0.1348	0.7324	0.2562	0.2439	0.1022	0.9148	0.5265	0.0796	0.4244	0.1109	0.0925	0.9718	0.2158	0.7989	0.1233	0.8929
---	---	1632059_at	-0.1641	0.3309	0.1234	0.3751	-0.1546	0.2980	-0.2776	0.3353	-0.2777	0.0954	-0.0001	0.9995	0.0840	0.8767	0.0754	0.7687	-0.0086	0.9807
nAcRalpha-96Aa	Dalpha1	1632060_at	0.0723	0.6227	0.1115	0.4534	0.2862	0.2049	-0.0615	0.9223	0.0172	0.9518	0.0787	0.6940	-0.0054	0.9970	0.0805	0.8764	0.0859	0.8549
CG15454	CG15454	1632061_at	0.0423	0.8367	0.0272	0.8896	0.3923	0.0187	0.2449	0.4979	0.1302	0.5222	-0.1147	0.5397	-0.1668	0.7464	0.0783	0.7682	0.2451	0.2784
CG12075	CG12075	1632062_s_at	-0.2428	0.1547	-0.0499	0.8479	-0.1476	0.4165	0.1330	0.8342	-0.0790	0.7883	-0.2120	0.3362	0.0256	0.9848	-0.1194	0.7339	-0.1451	0.6567
CG11357 /// DyakCG11357	CG11357	1632063_s_at	-0.4918	0.1250	-1.1025	0.0117	-1.1601	0.0026	-0.2128	0.7115	-0.0026	0.9945	0.2102	0.3852	-0.1247	0.9454	-0.2273	0.7184	-0.1026	0.8919
PH4alphaNE1	prolyl-4-hydroxyla	1632064_at	0.1713	0.3565	0.3398	0.1551	0.4879	0.0324	0.0631	0.9507	-0.1854	0.5391	-0.2486	0.3344	0.0575	0.9266	0.0626	0.8153	0.0051	0.9874
CG2921	CG2921	1632065_at	-0.0542	0.7273	0.2362	0.3389	0.3377	0.0768	0.0869	0.8385	0.1759	0.2734	0.0890	0.5762	0.0206	0.9884	0.4169	0.1623	0.3963	0.2136
Hr4	Hr4	1632066_at	-0.0773	0.7797	0.0206	0.8401	0.1592	0.4728	0.1680	0.8302	-0.0981	0.7889	-0.2661	0.3317	0.0020	0.9994	0.0002	1.0000	-0.0018	0.9979
---	---	1632067_at	-1.7930	0.0046	-1.5847	0.0314	-2.7172	0.0000	-0.0510	0.9639	0.2139	0.4939	0.2649	0.3303	1.1742	0.3646	0.3497	0.6397	-0.8245	0.2495
---	---	1632068_at	0.0029	0.9896	0.2649	0.1312	0.2030	0.2013	0.0236	0.9676	-0.0008	0.9973	-0.0243	0.8945	0.1721	0.7644	0.1099	0.6844	-0.0622	0.8409
---	---	1632069_at	0.4465	0.0691	0.0117	0.9125	0.2278	0.1551	0.0111	0.9874	0.2119	0.2037	0.2009	0.1784	0.0971	0.8923	0.0774	0.8262	-0.0197	0.9597
Ugt58Fa	lectin-58Fg	1632070_at	-1.5415	0.0022	-1.1403	0.0162	-0.8633	0.0024	-0.4787	0.2129	-1.0240	0.0017	-0.5453	0.0165	-0.7353	0.3166	-0.5013	0.1802	0.2340	0.5676
CG8870	CG8870	1632071_at	-1.1896	0.0022	-0.1545	0.4736	-0.5022	0.0936	-0.5786	0.1607	-0.8097	0.0067	-0.2310	0.3096	-0.0306	0.9848	0.0269	0.9647	0.0575	0.9095
CG34134	CG34134	1632072_at	0.0898	0.7157	-0.4584	0.1767	0.0034	0.9897	-0.0435	0.9436	-0.6157	0.0045	-0.5722	0.0038	-0.6188	0.6389	-1.1947	0.0483	-0.5759	0.2982
CG8334	CG8334	1632073_at	-0.5645	0.3178	0.2460	0.6405	0.0411	0.8951	-0.0401	0.9790	-0.3386	0.3820	-0.2985	0.3970	0.0857	0.9760	0.3550	0.6314	0.2693	0.7323
luna	luna	1632074_at	-0.2334	0.2465	0.5850	0.2927	0.1809	0.4937	0.0831	0.9269	0.1457	0.6297	0.0626	0.8414	0.4370	0.7337	0.7932	0.1418	0.3563	0.5398
---	---	1632075_at	0.0075	0.9758	0.0766	0.5411	0.2501	0.2856	0.1227	0.8974	-0.0379	0.9313	-0.1606	0.6117	0.0712	0.9199	0.0498	0.8857	-0.0213	0.9493
CG6933	CG6933	1632076_a_at	-0.2492	0.4117	0.0961	0.5364	-0.3067	0.0919	-0.3341	0.4873	-0.3393	0.1845	-0.0052	0.9873	0.0249	0.9823	0.0288	0.9459	0.0038	0.9933
CG30268	CG30268	1632077_at	0.1026	0.6890	0.1039	0.3253	0.0258	0.9163	0.0439	0.9639	0.0265	0.9438	-0.0174	0.9573	0.1271	0.8283	0.2030	0.4032	0.0759	0.7981
CG10326	CG10326	1632078_at	-0.0194	0.9405	-0.1517	0.5253	-0.3199	0.1665	-0.1468	0.6763	-0.2133	0.1945	-0.0664	0.7029	0.1222	0.9342	-0.2821	0.5754	-0.4044	0.4044

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632079_s_at	0.0409	0.8547	0.0233	0.8520	0.1912	0.1846	0.1580	0.5914	0.0729	0.6676	-0.0851	0.5646	0.0225	0.9862	0.0221	0.9642	-0.0004	0.9992
Cpr67Fa1 /// Cpr67Fa2	CG18349 /// CG7	1632080_s_at	0.0027	0.9952	0.2280	0.1097	0.3134	0.1471	0.0454	0.9688	-0.1426	0.6792	-0.1881	0.5222	-0.0341	0.9774	0.0233	0.9603	0.0574	0.8813
CG32626	CG32626	1632081_s_at	0.1030	0.8478	0.7852	0.1527	0.9423	0.0247	0.0514	0.9744	-0.5188	0.1783	-0.5702	0.0996	-0.0869	0.9689	0.1950	0.7883	0.2819	0.6533
Ndg	Nidogen-like	1632082_at	-0.5111	0.0160	-1.0973	0.0284	0.0159	0.9792	0.2016	0.8770	0.6322	0.1560	0.4306	0.2877	-0.8218	0.5259	0.0397	0.9690	0.8615	0.1911
CG30432	CG30432	1632083_at	0.0788	0.6967	0.1553	0.2008	-0.0608	0.8147	-0.1684	0.7753	-0.1601	0.5469	0.0084	0.9785	0.0693	0.9036	0.0173	0.9600	-0.0520	0.8453
CG12378 /// DereCG12378	CG12378	1632084_at	-0.1130	0.7159	-0.1991	0.3627	-0.2330	0.3414	-0.0806	0.9387	0.3442	0.2644	0.4248	0.1205	-0.1814	0.8945	0.1165	0.8737	0.2979	0.5862
CG30095	CG30095	1632085_at	0.0488	0.7995	0.7011	0.2018	1.1345	0.0305	-0.0011	0.9988	-0.0988	0.6399	-0.0977	0.6077	-0.1945	0.9447	0.9015	0.2710	1.0959	0.2136
CG15740	CG15740	1632086_at	-0.0149	0.9358	0.0687	0.5523	0.1268	0.3952	0.0151	0.9834	0.0081	0.9750	-0.0070	0.9730	0.0972	0.8655	0.0264	0.9429	-0.0707	0.7983
CG9928	CG9928	1632087_at	2.4881	0.0022	1.7237	0.0503	2.5370	0.0001	0.1319	0.8915	0.2570	0.4459	0.1251	0.7171	-0.6610	0.7595	-0.2853	0.8061	0.3757	0.7135
---	---	1632088_at	0.2208	0.5617	-0.0162	0.9372	-0.2594	0.5396	-0.1131	0.8104	0.2960	0.1131	0.4091	0.0228	-0.0266	0.9916	-0.0073	0.9959	0.0194	0.9847
CG4045	CG4045	1632089_at	0.2103	0.3138	0.0644	0.7863	0.3079	0.1043	0.1862	0.7400	0.0969	0.7396	-0.0893	0.7370	-0.0190	0.9898	0.0411	0.9355	0.0600	0.8915
hale	hale-bopp	1632090_at	0.3173	0.3224	0.0695	0.4561	0.3831	0.0548	0.1857	0.6673	0.1334	0.5474	-0.0523	0.8233	-0.2432	0.7644	-0.2802	0.4107	-0.0370	0.9387
CG8192	CG8192	1632091_at	0.1327	0.4424	-0.2034	0.1744	-0.5372	0.0222	-0.0023	0.9967	0.5377	0.0086	0.5400	0.0051	0.2094	0.7726	0.0845	0.8393	-0.1250	0.7230
---	---	1632092_at	-0.0666	0.7108	-0.0611	0.6110	0.2046	0.4561	0.1602	0.7803	0.0409	0.9009	-0.1193	0.6230	-0.1047	0.9272	0.0842	0.8760	0.1889	0.6402
---	---	1632093_at	0.1277	0.4345	0.0373	0.7090	-0.0075	0.9690	-0.0749	0.8836	0.0137	0.9576	0.0886	0.6153	0.0723	0.8999	-0.0580	0.8380	-0.1303	0.5612
CG8862	CG8862	1632094_at	-0.0347	0.8964	0.0661	0.7942	-0.3392	0.1172	-0.0341	0.9666	0.0000	1.0000	0.0341	0.8945	0.2976	0.7220	0.1460	0.7211	-0.1516	0.7053
CG17198	CG17198	1632095_at	0.1610	0.3196	-0.0146	0.9399	0.0428	0.7975	0.0829	0.9057	0.0033	0.9916	-0.0795	0.7444	0.1456	0.7822	0.0020	0.9977	-0.1436	0.5508
CG10032	CG10032	1632096_at	0.0379	0.8517	0.3676	0.1650	0.3981	0.1144	0.1039	0.8350	-0.0457	0.8521	-0.1496	0.3936	0.2235	0.7726	0.2623	0.4280	0.0388	0.9338
CG15646	CG15646	1632097_at	-0.0674	0.7802	0.2097	0.3655	0.2542	0.1768	0.1834	0.7408	-0.3503	0.1492	-0.5337	0.0227	0.1267	0.8461	0.1178	0.6977	-0.0089	0.9846
CG4364	CG4364	1632098_at	0.1520	0.3850	0.9457	0.0103	1.4835	0.0001	0.4031	0.2006	-0.8306	0.0018	-1.2338	0.0002	-0.0666	0.9390	-0.0114	0.9821	0.0552	0.8800
Liprin-alpha	liprin-alpha	1632099_s_at	0.1538	0.7134	0.1154	0.7569	-0.3709	0.1078	-0.1487	0.8379	0.5343	0.0548	0.6831	0.0128	0.3553	0.7953	0.7333	0.1897	0.3779	0.5335
sina	anon-fast-evolving	1632100_s_at	0.2301	0.4588	0.1483	0.5394	-0.1640	0.4486	-0.1890	0.5631	0.2047	0.2205	0.3937	0.0167	0.3098	0.7810	0.3890	0.4167	0.0792	0.9056
---	---	1632101_at	-0.1216	0.5577	-0.1548	0.4767	-0.1705	0.3965	0.0427	0.9435	0.1611	0.3520	0.1184	0.4619	-0.0061	0.9964	0.0413	0.9309	0.0474	0.9086
CG15580	CG15580	1632102_a_at	0.0618	0.6748	0.0316	0.7631	-0.0542	0.7958	-0.0929	0.8000	-0.0366	0.8484	0.0563	0.7205	-0.1227	0.8202	-0.0499	0.8754	0.0728	0.7820
CG9769	CG9769	1632103_at	-0.0741	0.8831	1.2052	0.0042	1.6348	0.0003	0.2262	0.6869	-1.2304	0.0013	-1.4566	0.0004	-0.1699	0.9095	-0.0098	0.9935	0.1601	0.8056
---	---	1632104_at	0.2051	0.4313	0.0822	0.4462	0.1522	0.4695	0.0239	0.9807	-0.0259	0.9407	-0.0498	0.8598	0.1266	0.8801	0.0177	0.9741	-0.1089	0.7736
cib	ciboulot	1632105_a_at	-1.5715	0.0098	0.2630	0.2542	0.1085	0.6301	-0.1322	0.8891	-1.3158	0.0023	-1.1836	0.0021	-0.0577	0.9751	0.4830	0.2734	0.5406	0.2536
CG9747	CG9747	1632106_at	-2.2488	0.0070	-2.8431	0.0067	-2.0161	0.0017	0.6950	0.5735	0.1546	0.8506	-0.5404	0.3519	-0.1899	0.9208	-0.4482	0.4969	-0.2584	0.7235
---	---	1632107_at	0.0000	0.9997	0.0232	0.8778	0.0242	0.8999	-0.0739	0.9011	-0.0421	0.8689	0.0318	0.8906	-0.0844	0.8960	-0.0343	0.9277	0.0501	0.8779
CG30384	CG30384	1632108_at	0.1965	0.2133	0.0257	0.7969	-0.0567	0.7445	0.0552	0.8987	0.2589	0.0708	0.2038	0.1073	-0.0300	0.9760	-0.0053	0.9925	0.0246	0.9421
alphaCop	alpha-coatomer p	1632109_s_at	0.4714	0.0799	0.5642	0.0949	1.1317	0.0006	0.2060	0.6371	0.1622	0.4689	-0.0438	0.8608	-0.2294	0.8000	0.3485	0.3437	0.5779	0.1517
CG8228	CG8228	1632110_at	-0.3976	0.0416	-0.1379	0.7660	0.0685	0.7743	0.0108	0.9922	-0.0087	0.9804	-0.0194	0.9473	-0.1645	0.8882	0.2296	0.6280	0.3941	0.3810
Arr2	arrestin	1632111_at	0.1762	0.4201	0.0497	0.8380	-0.0955	0.6538	-0.0162	0.9872	-0.0744	0.8026	-0.0581	0.8338	0.0598	0.9616	-0.1011	0.8244	-0.1608	0.6711
---	---	1632112_at	0.1840	0.4467	-0.0025	0.9914	-0.0416	0.8190	0.1724	0.7152	0.1254	0.5938	-0.0470	0.8507	0.0839	0.8999	-0.1527	0.5523	-0.2366	0.3497
---	---	1632113_at	0.1371	0.5677	0.0065	0.9540	0.1002	0.5456	0.1332	0.7335	0.1834	0.2970	0.0503	0.7976	0.0766	0.8960	-0.0264	0.9404	-0.1030	0.6639
Cyp12a4	Cyp12a4	1632114_at	2.2342	0.0014	1.8827	0.0083	2.9191	0.0000	0.1679	0.8794	-0.9242	0.0228	-1.0921	0.0067	-0.7631	0.2485	-1.1654	0.0158	-0.4023	0.2338
---	---	1632115_at	-0.0815	0.7183	-0.0067	0.9750	0.1407	0.3619	0.3069	0.3544	0.2434	0.1972	-0.0635	0.7597	-0.1041	0.8940	0.0495	0.9111	0.1536	0.6222
CG33234	CG33234	1632116_at	0.0682	0.7508	0.2187	0.3575	0.2691	0.2089	-0.1835	0.7117	-0.3665	0.1032	-0.1830	0.3803	0.0142	0.9912	-0.0324	0.9387	-0.0466	0.8951
Vha16	ductin	1632117_s_at	-0.4773	0.0543	-0.6805	0.1146	-0.5713	0.0060	0.2445	0.4140	0.2917	0.0806	0.0472	0.8007	0.1826	0.8680	0.1261	0.8252	-0.0565	0.9248
CG31121	CG31121	1632118_s_at	-0.8501	0.0412	-1.1018	0.0677	-1.1242	0.0058	0.2933	0.6557	0.3224	0.3124	0.0291	0.9408	0.0543	0.9849	-0.2957	0.6883	-0.3500	0.6224
ltd	lightoid	1632119_s_at	-0.0591	0.8489	-0.2439	0.3582	-0.1195	0.7065	-0.3365	0.4669	-0.2251	0.3781	0.1115	0.6671	-0.2964	0.7686	-0.3652	0.3856	-0.0688	0.9074
CG15533	CG15533	1632120_at	0.2041	0.5251	0.2896	0.2821	0.3287	0.0937	0.0620	0.9412	-0.1157	0.6731	-0.1777	0.4318	0.0874	0.9309	0.0588	0.9068	-0.0286	0.9505
CG6416	CG6416	1632121_a_at	-1.6947	0.0105	-1.8666	0.0450	-2.7042	0.0009	-0.3228	0.3353	-0.4823	0.0209	-0.1595	0.3725	0.5307	0.8454	-0.6275	0.5988	-1.1581	0.3097
---	---	1632122_at	-0.0546	0.7632	0.0519	0.7712	-0.0303	0.8810	-0.0847	0.8325	-0.0492	0.7913	0.0354	0.8399	-0.0196	0.9875	0.0297	0.9485	0.0492	0.8985
CG12050	CG12050	1632123_at	0.8105	0.0048	0.7615	0.1357	0.9501	0.0040	0.2440	0.4610	0.5219	0.0107	0.2779	0.0839	0.0824	0.9726	0.4120	0.5076	0.3296	0.6124
CG32345	CG32345	1632124_at	-0.8691	0.0421	-0.7561	0.0110	-1.0070	0.0046	-0.3618	0.7042	-0.0644	0.9167	0.2973	0.4758	0.0008	0.9998	-0.0176	0.9682	-0.0184	0.9622
---	---	1632125_at	0.0724	0.7809	-0.1638	0.2140	0.1897	0.4147	0.2634	0.6280	0.3585	0.1779	0.0951	0.7416	-0.0930	0.9142	0.0004	0.9999	0.0933	0.7980
---	---	1632126_at	0.1317	0.4279	0.0406	0.6978	-0.0811	0.6887	0.0119	0.9875	0.0239	0.9252	0.0119	0.9589	0.0383	0.9624	-0.0784	0.7779	-0.1168	0.6317
CG40189	CG40189	1632127_at	-0.0046	0.9792	0.0362	0.8004	0.1542	0.4082	0.0642	0.8987	-0.0397	0.8529	-0.1039	0.5159	-0.0377	0.9499	-0.0032	0.9935	0.0344	0.8880
---	---	1632128_x_at	0.4371	0.0785	0.2091	0.2977	0.5617	0.0058	0.1496	0.6988	0.0875	0.6642	-0.0621	0.7466	-0.1199	0.8455	-0.1270	0.6429	-0.0071	0.9859
syd	sunday driver	1632129_a_at	-0.4007	0.4268	1.1817	0.1149	1.4851	0.0001	-0.1437	0.9186	-1.0715	0.0215	-0.9279	0.0246	-0.2959	0.8504	0.5332	0.4077	0.8291	0.2189
SRPK	SRPK	1632130_s_at	-0.5172	0.0833	-0.5118	0.0596	-0.7067	0.0022	-0.0236	0.9777	0.1540	0.4776	0.1776	0.3509	0.1595	0.8395	0.2967	0.3465	0.1372	0.7056
5-HT7	Serotonin recepto	1632131_at	0.0383	0.8695	-0.1199	0.4197	-0.1413	0.4829	-0.0275	0.9755	0.0892	0.7260	0.1167	0.5928	0.0392	0.9689	-0.1100	0.7139	-0.1492	0.5988
CG5151	CG5151	1632132_at	0.4932	0.0132	0.6047	0.2974	0.3177	0.2666	-0.2143	0.5461	0.1247	0.5243	0.3390	0.0450	0.4049	0.7953	0.5522	0.3922	0.1473	0.8646
---	---	1632133_at	-0.0180	0.9248	0.0672	0.5254	-0.0571	0.8104	-0.0706	0.8841	-0.0371	0.8646	0.0335	0.8618	0.0151	0.9893	0.0021	0.9977	-0.0130	0.9716
Axn	D-axin	1632134_s_at	0.1233	0.7850	-0.3780	0.3774	-0.5661	0.0885	-0.1972	0.8403	0.5923	0.1042	0.7895	0.0239	-0.1375	0.9514	0.0754	0.9449	0.2130	0.7909
fl(2)d	female lethal d	1632135_a_at	-0.0262	0.9251	-0.0072	0.9751	-0.1347	0.4193	-0.1154	0.8393	0.1172	0.6189	0.2326	0.2222	0.0980	0.9174	0.0780	0.8658	-0.0200	0.9678
CG7837	CG7837	1632136_at	0.0415	0.9036	0.0435	0.7074	-0.2934	0.1075	-0.1171	0.7967	0.1085	0.5890	0.2256	0.1707	0.2603	0.7707	0.1627	0.7052	-0.0977	0.8407
---	---	1632137_at	0.1722	0.3801	0.0134	0.9680	0.1624	0.3394	0.2708	0.6038	0.2603	0.3284	-0.0105	0.9742	0.2778	0.7116	0.1796	0.5897	-0.0982	0.7954
sa	spermatocytes arr	1632138_at	-0.0667	0.6842	-0.0377	0.7122	0.0573	0.8166	0.1304	0.7409	0.0127	0.9612	-0.1178	0.4776	-0.0323	0.9774	0.0449	0.9135	0.0771	0.8173
CG14174	CG14174	1632139_at	0.3012	0.2028	-0.3564	0.3902	-0.0261	0.9259	0.1070	0.9220	0.6375	0.0574	0.5305	0.0736	-0.2850	0.7464	0.0450	0.9402	0.3300	0.3842
---	---	1632140_at	0.1875	0.3035	0.0026	0.9836	-0.0021	0.9942	-0.0391	0.9518	0.0367	0.8791	0.0758	0.6896	-0.1006	0.8919	-0.0462	0.9148	0.0544	0.8870
CG18265 /// DereCG18265	CG18265	1632141_at	0.2259	0.3630	-0.1375	0.3609	-0.0796	0.7017	0.1535	0.6802	0.2005	0.2497	0.0470	0.8102	-0.1028	0.8917	-0.1766	0.5523	-0.0738	0.8374
Rpb5	Rpb5	1632142_at	0.3895	0.2068	0.7747	0.0903	0.8363	0.0043	0.1783	0.7293	-0.0094	0.9792	-0.1877	0.3798	0.0673	0.9666	0.2724	0.5411	0.2051	0.6577
CG10102	CG10102	1632143_at	0.1583	0.4336	0.0427	0.8634	0.3102	0.4133	0.1877	0.7595	0.3385	0.1964	0.1508	0.5624	-0.0759	0.9545	0.2432	0.5523	0.3191	0.4241
---	---	1632144_at	0.1042	0.4614	0.0255	0.8923	0.0165	0.9441	0.1506	0.6936	0.2875	0.1029	0.1369	0.4058	0.1513	0.7997	0.1197	0.6607	-0.0317	0.9266
CG31720	CG31720	1632145_a_at	0.2594	0.5563	-0.0123	0.9058	0.0689	0.7120	0.0632	0.9311	-0.0189	0.9529	-0.0821	0.7183	0.0576	0.9806	-0.4096	0.4270	-0.4672	0.3791
CG5508	CG5508	1632146_a_at	1.7329	0.0091	0.7524	0.2592	1.1031	0.0091	-0.2851	0.7947	-0.1691	0.7497	0.1161	0.8185	-0.7163	0.7506	-1.1832	0.2053	-0.4669	0.6545
And	Calmodulin-relate	1632147_at	0.0979	0.5223	0.0028	0.9828	0.2474	0.0970	0.0102	0.9873	0.0177	0.9352	0.0075	0.9689	-0.0581	0.9291	0.0298	0.9286	0.0879	0.7123

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632148_at	0.0634	0.7858	0.0165	0.8868	0.1360	0.4937	0.0470	0.9559	-0.0025	0.9941	-0.0495	0.8531	-0.0357	0.9816	0.0301	0.9590	0.0658	0.8921
CG17930	CG17930	1632149_at	-0.0142	0.9535	-0.2606	0.1301	0.1714	0.4759	0.1465	0.7647	0.1636	0.4521	0.0172	0.9484	-0.1500	0.8097	0.0582	0.8752	0.2082	0.4322
---	---	1632150_at	0.0405	0.8338	-0.0382	0.6849	0.0769	0.6935	0.2048	0.5008	0.1419	0.3904	-0.0630	0.7129	0.1849	0.7476	0.0649	0.8444	-0.1200	0.6473
Art7	Arginine methyltra	1632151_at	-0.3361	0.1259	-0.1892	0.6151	0.2876	0.1201	0.0835	0.8395	-0.4628	0.0093	-0.5463	0.0027	-0.3442	0.7280	-0.3420	0.4200	0.0022	0.9979
CG10992 /// DyakCG10992	cathepsin B /// CG	1632152_at	0.2854	0.1279	0.4284	0.1460	0.4353	0.0109	0.0832	0.8676	-0.1506	0.4064	-0.2338	0.1367	-0.0351	0.9666	-0.0791	0.7703	-0.0440	0.8857
---	---	1632153_at	0.1220	0.6533	0.0021	0.9905	0.0219	0.9345	0.0158	0.9852	-0.0639	0.7796	-0.0796	0.6873	0.1394	0.8378	-0.0507	0.9058	-0.1902	0.5145
CG14060 /// DmirCG14060	CG14060	1632154_at	0.4866	0.0743	0.1441	0.4206	-0.0497	0.8438	0.0298	0.9705	0.1615	0.4532	0.1317	0.5098	-0.0134	0.9940	-0.1489	0.7672	-0.1355	0.7830
CG13676	CG13676	1632155_at	-0.1934	0.2017	-0.2760	0.0921	-0.3104	0.0591	-0.0498	0.9303	0.1728	0.3056	0.2226	0.1345	-0.0867	0.8889	-0.1327	0.5930	-0.0461	0.8864
---	---	1632156_at	0.2967	0.2756	0.3128	0.0478	0.2457	0.2643	0.0552	0.9509	-0.1571	0.5582	-0.2123	0.3528	0.1461	0.8400	-0.0595	0.8929	-0.2056	0.5089
---	---	1632157_at	0.0748	0.6886	0.1441	0.3058	0.2949	0.0822	-0.0779	0.8866	-0.2409	0.1899	-0.1630	0.3344	-0.0539	0.9330	-0.0167	0.9613	0.0372	0.8949
EloA	Elongin A	1632158_a_at	0.2198	0.2617	0.4169	0.2120	0.2218	0.3421	-0.1122	0.8394	0.2064	0.3259	0.3186	0.0859	0.0973	0.9400	0.4004	0.3000	0.3031	0.4597
---	---	1632159_at	-0.0040	0.9849	0.1041	0.3181	0.1211	0.3941	-0.0499	0.9228	-0.0020	0.9928	0.0479	0.7830	0.1415	0.8141	0.1491	0.5679	0.0076	0.9849
CG15279	CG15279	1632160_s_at	0.3453	0.7178	0.5670	0.1181	-0.6963	0.1705	-1.2432	0.1787	-1.2164	0.0363	0.0268	0.9694	-0.0127	0.9984	-0.8769	0.4829	-0.8642	0.4948
---	---	1632161_at	-0.4504	0.0504	-0.8449	0.0289	-0.8101	0.0030	-0.3110	0.3793	0.0747	0.7541	0.3857	0.0377	-0.2715	0.7485	-0.2272	0.5469	0.0444	0.9299
CG15028	CG15028	1632162_a_at	-0.0016	0.9962	0.0918	0.5253	0.0955	0.6582	-0.0074	0.9956	-0.0782	0.8495	-0.0708	0.8478	-0.0425	0.9441	-0.0255	0.9303	0.0170	0.9471
CG8481	CG8481	1632163_at	-0.1387	0.5506	0.5211	0.0611	0.2279	0.3391	-0.3831	0.2501	-1.0135	0.0009	-0.6304	0.0042	0.0488	0.9814	-0.3508	0.4280	-0.3996	0.3803
scrib	scribble	1632164_at	-0.0819	0.7167	-0.6960	0.0311	-0.4295	0.0162	0.2694	0.4122	0.6073	0.0058	0.3379	0.0438	0.0021	0.9990	-0.1484	0.5854	-0.1505	0.5814
CG32058	CG32058	1632165_at	0.0189	0.9273	0.0300	0.7914	0.0548	0.7417	0.0050	0.9956	-0.0179	0.9469	-0.0229	0.9175	0.0417	0.9643	0.0547	0.8792	0.0131	0.9732
Plap	Phospholipase A2	1632166_at	0.1587	0.5673	0.2223	0.1391	0.7966	0.0031	0.3294	0.3909	0.0565	0.8395	-0.2729	0.1574	-0.1281	0.8798	0.2107	0.5322	0.3389	0.3079
---	---	1632167_at	0.0939	0.6088	-0.1087	0.3424	0.0733	0.7425	0.1042	0.7931	0.2037	0.2077	0.0994	0.5296	-0.1024	0.8378	-0.0444	0.8801	0.0580	0.8210
msopa	Drosophila male s	1632168_at	0.0487	0.8736	0.0345	0.8056	0.0586	0.8055	-0.0002	0.9997	-0.0169	0.9543	-0.0167	0.9473	-0.0800	0.9272	-0.0939	0.7954	-0.0139	0.9759
---	---	1632169_at	0.1508	0.3131	-0.2871	0.2691	0.0899	0.6208	0.1691	0.6499	0.3649	0.0478	0.1958	0.2240	-0.1391	0.8270	0.0475	0.9077	0.1866	0.5007
Cap-H2	Chromosome ass	1632170_a_at	0.2002	0.5188	0.7749	0.1587	0.5992	0.0092	-0.3028	0.5726	-0.5067	0.0676	-0.2039	0.4280	-0.0524	0.9822	0.1901	0.7769	0.2425	0.6877
CG3436	CG3436	1632171_a_at	-0.1046	0.6853	0.4032	0.0998	0.4096	0.0893	-0.0274	0.9761	-0.2457	0.2832	-0.2183	0.2892	0.0782	0.9309	0.2853	0.3146	0.2071	0.4916
---	---	1632172_at	0.0195	0.9346	0.2313	0.2228	0.0822	0.7694	0.0448	0.9489	-0.1588	0.4325	-0.2036	0.2454	0.0731	0.9309	0.0340	0.9390	-0.0391	0.9176
---	---	1632173_at	0.0196	0.9393	-0.0366	0.7193	-0.1315	0.5189	0.0436	0.9603	0.1050	0.6956	0.0614	0.8163	0.0418	0.9653	0.0009	0.9994	-0.0409	0.9081
---	---	1632174_at	-0.0169	0.9625	-0.0032	0.9796	0.0162	0.9473	0.1180	0.8107	-0.0638	0.7899	-0.1818	0.3063	-0.0521	0.9679	-0.0659	0.8988	-0.0139	0.9809
---	---	1632175_at	0.3211	0.0473	0.0419	0.8224	0.1285	0.6015	0.0762	0.8938	0.0797	0.7178	0.0035	0.9881	-0.0020	0.9994	-0.0590	0.9084	-0.0570	0.9023
CG5623	CG5623	1632176_at	0.0796	0.7638	0.0046	0.9818	0.3259	0.3509	-0.0026	0.9978	-0.0185	0.9598	-0.0159	0.9589	-0.3174	0.6984	-0.2275	0.5251	0.0899	0.8374
hth	dorsotonals	1632177_at	0.4965	0.3778	-0.5409	0.3866	-1.5367	0.0243	0.2838	0.6062	2.1005	0.0002	1.8167	0.0002	1.3718	0.6557	1.0339	0.4002	-0.3379	0.8287
enc	encore	1632178_at	-0.2867	0.6751	-0.5153	0.4924	-0.9886	0.0028	-0.4733	0.3462	0.1541	0.6333	0.6274	0.0235	0.0214	0.9963	0.0605	0.9710	0.0392	0.9816
Sptr	sepiapterin reduct	1632179_at	-0.7839	0.0087	-0.0015	0.9950	-0.1445	0.3418	-0.2397	0.3837	-1.0762	0.0003	-0.8364	0.0005	-0.0594	0.9449	-0.2100	0.4183	-0.1506	0.5880
CG7791	CG7791	1632180_at	-0.4036	0.0534	-0.0980	0.7492	-0.1746	0.2628	-0.0595	0.8891	-0.0275	0.8865	0.0320	0.8470	0.0790	0.9588	0.2964	0.4969	0.2174	0.6340
Or65c	Odorant receptor	1632181_at	0.2917	0.2288	0.0802	0.5245	0.0071	0.9714	-0.1227	0.8234	0.0070	0.9822	0.1297	0.5340	-0.0443	0.9487	0.0452	0.8785	0.0894	0.6973
CG12096	CG12096	1632182_at	0.0379	0.7939	0.2397	0.4622	0.4417	0.0096	0.0030	0.9960	0.2266	0.2234	0.2235	0.1787	-0.1543	0.8692	0.3649	0.3185	0.5192	0.1911
ru	Rhomboid 3	1632183_at	0.0329	0.8769	0.3015	0.2783	-0.1640	0.3256	0.0852	0.8815	0.0907	0.6827	0.0054	0.9825	0.4337	0.6660	0.2744	0.5120	-0.1592	0.7309
CG4953	CG4953	1632184_at	0.4122	0.2185	0.0787	0.7863	0.1351	0.3791	-0.1614	0.7217	0.1724	0.4133	0.3339	0.0703	-0.1594	0.8815	-0.1048	0.8516	0.0546	0.9231
scpr-A /// scpr-B /// scpr-C	SCP-containing p	1632185_s_at	0.4989	0.0145	0.8942	0.0205	0.8608	0.0283	0.0276	0.9704	0.1509	0.4451	0.1232	0.5002	0.2017	0.8882	0.7163	0.1722	0.5146	0.3552
---	---	1632186_at	-0.0077	0.9742	0.1350	0.4220	-0.0429	0.8218	-0.0942	0.8768	-0.2685	0.1969	-0.1743	0.3660	0.1606	0.7726	0.0634	0.8460	-0.0972	0.7217
CG9646	CG9646	1632187_at	-0.8467	0.0148	-0.1143	0.8240	0.6679	0.0324	0.1902	0.8604	-0.3577	0.3662	-0.5479	0.1127	-0.4593	0.6955	0.4909	0.3034	0.9502	0.0879
Hexo1	Hexosaminidase	1632188_at	-0.6341	0.0059	-1.2455	0.0445	-0.9017	0.0060	0.1319	0.8512	0.3176	0.2138	0.1857	0.4407	-0.2182	0.8427	-0.2675	0.5763	-0.0493	0.9379
CG3587	CG3587	1632189_at	-0.4494	0.1260	0.0651	0.8237	0.4273	0.0401	0.0118	0.9937	-0.1861	0.5803	-0.1978	0.5081	-0.0932	0.9238	0.4035	0.1995	0.4967	0.1528
Atx-1	Ataxin 1	1632190_at	0.3050	0.3345	0.0665	0.5894	-0.0509	0.8437	-0.1821	0.7507	-0.0323	0.9258	0.1497	0.5385	0.0579	0.9457	-0.1620	0.5446	-0.2199	0.3994
hts	hu-li tai shao	1632191_s_at	-0.4325	0.5059	-0.0318	0.8898	-1.3470	0.0274	-0.3590	0.3863	0.2672	0.2566	0.6262	0.0096	1.0728	0.6898	0.7192	0.5254	-0.3536	0.7853
CG14564	CG14564	1632192_at	0.2427	0.2544	0.2671	0.2539	0.2509	0.3049	-0.0805	0.8732	-0.2357	0.1759	-0.1552	0.3322	0.2692	0.6483	0.1226	0.6336	-0.1466	0.5614
---	---	1632193_at	-0.0615	0.7580	-0.1315	0.2939	0.2025	0.2358	0.1703	0.5917	0.0895	0.6145	-0.0808	0.6220	-0.1693	0.7726	-0.0046	0.9935	0.1647	0.5358
Nup98	Nup98	1632194_at	0.3227	0.3522	-0.1241	0.6611	0.1342	0.5708	0.1705	0.7582	0.6066	0.0191	0.4361	0.0461	0.0808	0.9657	0.3959	0.4355	0.3151	0.5587
CG4645 /// DsimCG4645 ///	CG4645	1632195_at	0.5355	0.0897	0.7461	0.0575	0.7777	0.0009	-0.0746	0.9191	0.2499	0.2701	0.3245	0.1067	-0.2061	0.8461	0.4404	0.2990	0.6465	0.1612
Abi	Abelson Interactin	1632196_at	-1.3107	0.0038	-0.1309	0.4654	-0.4964	0.0372	-0.3428	0.4865	-1.1224	0.0020	-0.7796	0.0060	-0.0767	0.9520	-0.0128	0.9870	0.0640	0.9033
eIF-2alpha	Eukaryotic initiatio	1632197_at	0.3186	0.2430	1.4377	0.0091	1.6511	0.0000	0.2326	0.6041	-0.8560	0.0033	-1.0885	0.0007	-0.0488	0.9742	0.2537	0.5183	0.3025	0.4371
CG13902	CG13902	1632198_at	-0.1634	0.4005	-0.4507	0.0755	-0.2540	0.0905	0.1976	0.5515	0.5445	0.0073	0.3469	0.0313	0.1053	0.9152	0.2757	0.4242	0.1704	0.6473
CG32390	CG32390	1632199_at	0.3621	0.1562	0.3238	0.0924	0.2679	0.2479	0.0890	0.8427	0.1380	0.4231	0.0489	0.7932	0.1375	0.8141	0.0326	0.9328	-0.1049	0.6973
Edem1	CG3810	1632200_s_at	-0.0873	0.7829	0.9962	0.0101	1.1649	0.0002	0.1946	0.6533	-0.1448	0.5120	-0.3395	0.0714	0.0107	0.9952	0.8944	0.0519	0.8837	0.0708
---	---	1632201_at	0.0728	0.7632	0.0389	0.7485	0.2182	0.3262	0.0122	0.9866	0.0094	0.9733	-0.0028	0.9901	-0.0650	0.9055	0.0525	0.8433	0.1175	0.5724
CG10616 /// DsimCG10616	CG10616	1632202_at	0.5917	0.0165	0.4368	0.3171	0.3272	0.0722	0.1140	0.7857	0.5685	0.0062	0.4546	0.0103	0.1218	0.9340	0.4055	0.3802	0.2837	0.5700
CG31706	CG31706	1632203_at	-0.4212	0.2825	0.4160	0.2200	-0.3548	0.2129	-0.5065	0.3625	-0.9450	0.0096	-0.4385	0.1202	0.0738	0.9545	-0.2846	0.4565	-0.3585	0.3597
Mmp1	Matrix metalloprot	1632204_at	0.1770	0.8453	-0.4365	0.6597	0.5274	0.4130	1.3835	0.2132	2.4866	0.0036	1.1031	0.0680	0.3374	0.9467	1.5483	0.3000	1.2109	0.4431
Dgk	Diacyl glycerol kin	1632205_at	-0.1395	0.4578	0.0011	0.9940	-0.0270	0.9118	-0.0597	0.9407	0.0220	0.9493	0.0817	0.7420	0.0470	0.9407	0.1119	0.5889	0.0649	0.7777
CG5491	CG5491	1632206_at	0.0250	0.9253	0.2277	0.3081	0.2483	0.1594	0.0842	0.9186	-0.0364	0.9185	-0.1207	0.6440	0.0173	0.9908	0.3079	0.3090	0.2907	0.3636
p130CAS	p130CAS	1632207_a_at	0.3837	0.0889	0.6809	0.3627	0.4551	0.2187	0.1236	0.8572	0.2955	0.2293	0.1719	0.4607	0.4381	0.8049	0.6245	0.3943	0.1864	0.8444
---	---	1632208_at	0.1798	0.4954	0.1150	0.4301	0.2748	0.2286	0.1558	0.7205	0.2302	0.2431	0.0744	0.7231	0.1114	0.8814	0.0065	0.9925	-0.1049	0.7495
CG7829	CG7829	1632209_at	0.3419	0.0913	0.5399	0.0149	0.2364	0.1204	-0.0854	0.8500	-0.0280	0.9018	0.0574	0.7478	0.2079	0.7464	0.1717	0.5439	-0.0362	0.9231
CG30091	CG30091	1632210_at	0.1014	0.5064	0.0396	0.8329	0.0079	0.9751	-0.0566	0.9500	-0.0604	0.8528	-0.0038	0.9905	0.0418	0.9530	0.0010	0.9991	-0.0408	0.8884
CG31829	CG31829	1632211_at	0.0645	0.6870	0.1380	0.5172	-0.1045	0.6890	0.0475	0.9603	0.0496	0.8836	0.0021	0.9948	0.3017	0.7485	0.0776	0.8980	-0.2241	0.6040
CG14401	CG14401	1632212_at	-1.3452	0.0008	-1.0829	0.0426	-1.5125	0.0028	-0.4099	0.3056	-0.8095	0.0051	-0.3996	0.0615	0.0098	0.9969	-0.5403	0.3739	-0.5502	0.3848
CG31292 /// CG3303 /// Dm	CG31292 /// CG3	1632213_s_at	-1.2862	0.0581	0.4972	0.3398	0.7829	0.0566	-0.2746	0.8589	-1.6560	0.0096	-1.3814	0.0132	-0.5429	0.7673	0.0022	0.9996	0.5451	0.4978
CG6138	CG6138	1632214_at	-0.0290	0.9061	-0.0738	0.6634	-0.0648	0.6955	0.1177	0.8738	0.1673	0.5477	0.0496	0.8705	0.0854	0.8760	0.0122	0.9734	-0.0732	0.7706
CG6296	CG6296	1632215_at	0.1059	0.6772	0.0127	0.9643	0.0290	0.8977	0.2282	0.6354	0.0073	0.9833	-0.2210	0.3034	0.0635	0.9246	-0.0565	0.8551	-0.1200	0.6216
CanA1	calcineurin	1632216_at	-2.8976	0.0041	-4.0974	0.0030	-3.8745	0.0000	-0.0394	0.9641	-0.1170	0.6558	-0.0776	0.7611	-0.4001	0.8609	-1.3480	0.1395	-0.9478	0.3149

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
gskt	gasket	1632217_at	0.0244	0.9091	0.2936	0.3354	0.3771	0.0445	-0.0767	0.9116	-0.1767	0.4363	-0.1000	0.6591	-0.0721	0.9324	0.0475	0.9093	0.1197	0.6941
---	---	1632218_s_at	0.3646	0.0833	0.0712	0.7400	0.2754	0.1149	0.2881	0.4833	0.1887	0.4032	-0.0994	0.6623	-0.1083	0.8850	-0.0002	1.0000	0.1081	0.7416
---	---	1632219_at	-0.0024	0.9934	0.0761	0.5265	-0.0082	0.9723	-0.1260	0.8074	-0.0137	0.9656	0.1123	0.5851	0.0757	0.9345	0.2618	0.3685	0.1861	0.5520
RpS26	ribosomal protein	1632220_s_at	0.3172	0.0621	0.7051	0.0085	0.7523	0.0107	0.0622	0.9376	-0.4104	0.0790	-0.4727	0.0304	0.1030	0.8379	-0.0209	0.9504	-0.1239	0.5733
CG12203 /// DyakCG12203	CG12203	1632221_at	-0.5448	0.0360	0.1095	0.6996	0.2189	0.5222	0.1468	0.8822	-1.1117	0.0063	-1.2585	0.0022	0.0314	0.9869	-0.3338	0.4479	-0.3652	0.4120
CG9238	CG9238	1632222_a_at	2.1591	0.0052	1.2227	0.1181	0.5871	0.2306	-0.3094	0.7409	0.1833	0.6995	0.4927	0.1800	0.3077	0.9257	-0.8433	0.4423	-1.1510	0.3064
dbe	dribble	1632223_at	0.0148	0.9547	0.3570	0.3622	0.6751	0.0238	0.1270	0.8162	-0.0454	0.8731	-0.1724	0.3865	-0.2437	0.8292	0.2446	0.6314	0.4883	0.3118
CG40137	---	1632224_at	0.2000	0.3688	-0.7953	0.0776	-0.7784	0.0049	-0.1170	0.8903	0.9587	0.0056	1.0757	0.0020	-0.1874	0.8283	-0.1840	0.6394	0.0034	0.9956
Las	Lipoic acid syntha	1632225_a_at	-0.2605	0.1923	0.1495	0.3387	0.5149	0.0096	0.1550	0.8222	-0.7309	0.0145	-0.8859	0.0037	-0.1249	0.8629	-0.0923	0.8052	0.0326	0.9361
CG1079	CG1079	1632226_at	-1.1333	0.0110	-1.3477	0.1301	-1.8691	0.0000	-0.3806	0.5910	-0.5744	0.1085	-0.1939	0.5890	0.2532	0.8849	-0.6431	0.3191	-0.8963	0.1993
CG16890	CG16890	1632227_at	-0.2015	0.5698	0.2559	0.6079	1.1653	0.0008	0.0096	0.9932	-0.8392	0.0034	-0.8488	0.0019	-0.5824	0.6898	-0.0556	0.9541	0.5267	0.3914
CG6733	CG6733	1632228_at	-0.4932	0.7653	-2.7869	0.0103	-2.8193	0.0000	0.3325	0.9228	0.8304	0.4378	0.4979	0.6382	0.3772	0.9056	-1.3602	0.2168	-1.7373	0.1532
---	---	1632229_at	0.0933	0.5922	0.1476	0.3546	-0.0015	0.9950	-0.0369	0.9558	-0.1125	0.5598	-0.0756	0.6870	0.0398	0.9467	-0.1557	0.3768	-0.1955	0.2917
---	---	1632230_at	0.0902	0.6928	0.1019	0.6110	0.1769	0.3586	0.1985	0.6669	0.1624	0.4801	-0.0362	0.8907	0.0511	0.9717	0.2158	0.5650	0.1647	0.6742
Mhcl	myosin XVII (PDZ	1632231_a_at	-1.8698	0.0041	-0.8105	0.0670	-0.9490	0.0008	0.2086	0.7349	-0.2531	0.3658	-0.4618	0.0628	-0.0036	0.9992	0.2455	0.6605	0.2490	0.6495
---	---	1632232_at	-1.6487	0.0137	-1.8045	0.0070	-1.0956	0.0215	0.6792	0.5552	0.4601	0.4574	-0.2191	0.7310	-0.0953	0.9238	-0.1638	0.6497	-0.0685	0.8792
CG14749	CG14749	1632233_at	0.2543	0.1273	-0.1095	0.4856	0.0069	0.9734	0.0192	0.9766	0.5601	0.0048	0.5410	0.0034	-0.0716	0.9352	0.2420	0.3802	0.3135	0.2798
Aats-tyr	Tyrosyl-tRNA synt	1632234_at	-0.1741	0.2937	0.9705	0.0375	1.3543	0.0001	0.3324	0.2596	-0.7415	0.0020	-1.0738	0.0003	-0.0493	0.9705	0.4602	0.1622	0.5095	0.1573
Aats-met	Methionyl-tRNA s	1632235_at	-0.1890	0.3983	0.3378	0.1865	0.3898	0.2720	-0.1963	0.7321	-0.2614	0.3111	-0.0652	0.8223	-0.1333	0.9056	0.3345	0.4032	0.4678	0.2622
CG10221	CG10221	1632236_at	0.7324	0.0072	0.5323	0.0554	0.9203	0.0012	0.0592	0.9218	0.7523	0.0023	0.6932	0.0019	-0.2450	0.7496	0.5580	0.0966	0.8030	0.0523
Rx	Retinal Homeobo	1632237_at	0.1032	0.6141	0.0037	0.9864	-0.1296	0.5091	0.0829	0.8422	0.0362	0.8578	-0.0466	0.7871	-0.0109	0.9914	-0.2010	0.3340	-0.1901	0.3831
CG13248	CG13248	1632238_at	-2.0840	0.0011	-1.7286	0.0470	-2.3698	0.0000	-0.5973	0.4509	-0.3442	0.4403	0.2531	0.5490	-0.1991	0.7893	-0.2392	0.4512	-0.0401	0.9271
Obp85a	Odorant-binding p	1632239_at	-0.2117	0.3787	0.0000	1.0000	-0.0919	0.6424	-0.0069	0.9956	-0.0179	0.9643	-0.0110	0.9730	0.0733	0.9128	0.1014	0.7027	0.0282	0.9321
mus81	mus81	1632240_at	0.4758	0.1991	0.0649	0.7618	-0.5524	0.0471	-0.9766	0.1378	0.2852	0.4951	1.2618	0.0043	-0.2774	0.7644	-0.0050	0.9955	0.2724	0.5030
CG14376	CG14376	1632241_at	0.2097	0.2259	-0.1183	0.5978	-0.0741	0.7702	-0.0168	0.9854	0.1243	0.5812	0.1411	0.4748	0.0346	0.9760	0.0505	0.9055	0.0159	0.9703
Syn1	Syntrophin-like 1	1632242_a_at	-0.2778	0.2550	0.3664	0.0681	-0.1083	0.6303	-0.0321	0.9745	-0.3025	0.2147	-0.2704	0.2164	0.3404	0.6749	0.0966	0.8228	-0.2438	0.4746
hd	humpty dumpty	1632243_at	0.2253	0.7150	-0.3273	0.7239	-0.1611	0.7119	-0.3587	0.3501	0.4796	0.0363	0.8384	0.0019	-0.7722	0.8298	-0.0919	0.9708	0.6803	0.6826
CG32244	CG32244	1632244_s_at	0.3544	0.1094	0.0621	0.6190	0.1147	0.6095	0.0328	0.9647	0.0611	0.7997	0.0283	0.9045	-0.1619	0.8192	-0.3119	0.2643	-0.1500	0.6311
Spn4	neuroserpin	1632245_a_at	0.2844	0.4592	0.5334	0.0794	1.2396	0.0004	0.4589	0.3812	0.2479	0.4170	-0.2110	0.4499	-0.2345	0.7485	0.5146	0.1061	0.7490	0.0557
CG14383	CG14383	1632246_at	0.1304	0.5471	0.0236	0.8285	0.0156	0.9552	-0.0868	0.9036	0.1178	0.6532	0.2046	0.3367	0.0721	0.8940	0.0598	0.8217	-0.0122	0.9686
fzr2	fizzy-related 2	1632247_at	-0.0462	0.8052	0.1520	0.4032	-0.0438	0.7910	-0.0903	0.8315	-0.0952	0.5887	-0.0049	0.9808	0.1902	0.7324	0.2031	0.3871	0.0129	0.9721
CycG	cyclin G1	1632248_s_at	-0.0373	0.9148	0.3223	0.1035	0.0455	0.7743	-0.0493	0.9247	-0.2208	0.1554	-0.1714	0.2197	0.1313	0.9061	0.2300	0.5860	0.0987	0.8472
---	---	1632249_at	0.0498	0.8295	0.1488	0.4389	-0.0364	0.8785	-0.0711	0.8723	-0.1408	0.3701	-0.0697	0.6621	0.1194	0.8999	0.0037	0.9965	-0.1156	0.7791
CG1409	CG1409	1632250_at	-0.0339	0.9060	-0.0066	0.9821	0.1063	0.4714	-0.0331	0.9633	-0.2145	0.2643	-0.1814	0.2955	-0.1157	0.8692	-0.0778	0.8374	0.0380	0.9224
CG6181	CG6181	1632251_s_at	0.0024	0.9962	0.1851	0.6625	0.1252	0.6221	-0.0150	0.9931	0.0550	0.9070	0.0701	0.8598	0.1169	0.9460	0.2468	0.6695	0.1299	0.8469
CG41435	CG41435	1632252_s_at	0.1938	0.2613	-0.0421	0.7066	0.1597	0.2907	0.1371	0.6718	0.0149	0.9481	-0.1222	0.3909	0.0821	0.9259	0.0436	0.9253	-0.0385	0.9237
---	---	1632253_at	0.1098	0.6425	-0.0899	0.4744	0.1026	0.5090	0.0770	0.9080	0.1488	0.5097	0.0718	0.7583	-0.0389	0.9589	-0.0806	0.7481	-0.0417	0.8870
CG30270	CG30270	1632254_a_at	0.0035	0.9883	0.0135	0.9095	-0.1089	0.5866	0.0357	0.9672	0.1056	0.6844	0.0700	0.7827	-0.0182	0.9895	0.0234	0.9617	0.0417	0.9175
---	---	1632255_s_at	-0.2872	0.4067	0.0362	0.7620	-0.0258	0.9084	-0.1421	0.7908	0.0466	0.8723	0.1887	0.3468	-0.2067	0.8069	0.1896	0.6156	0.3962	0.2778
Taf1	TFIID 230 kda sut	1632256_at	0.3861	0.3598	-0.0131	0.9664	-0.0662	0.7201	0.1064	0.8578	0.5301	0.0216	0.4237	0.0341	0.0535	0.9816	0.1626	0.8153	0.1091	0.8813
CG31183	CG31183	1632257_at	0.9858	0.0764	0.7098	0.2510	0.4332	0.2254	0.1532	0.7388	0.3361	0.0978	0.1829	0.3231	0.3517	0.9029	0.1322	0.9377	-0.2195	0.8775
CG33201	CG33201	1632258_s_at	-1.5931	0.0035	-1.7480	0.0077	-1.7989	0.0000	0.0687	0.9518	0.6641	0.0426	0.5953	0.0421	0.2193	0.7726	0.4758	0.1395	0.2565	0.4448
Tango9	Transport and Gol	1632259_at	0.0507	0.8054	-0.0074	0.9529	0.1016	0.5418	-0.2533	0.3744	-0.5859	0.0040	-0.3325	0.0289	-0.3330	0.5665	-0.6054	0.0419	-0.2724	0.3089
bru-2	bruno-2	1632260_s_at	0.0665	0.7642	0.2033	0.4842	-0.5096	0.1540	-0.2350	0.6202	0.1774	0.4731	0.4124	0.0553	0.2493	0.8206	0.1969	0.7028	-0.0524	0.9347
CG32732 /// DsimCG32732	CG32732	1632261_at	-0.0416	0.8367	-0.0789	0.8280	0.2166	0.2883	-0.0400	0.9518	-0.1858	0.3136	-0.1459	0.3900	-0.1427	0.8692	-0.1867	0.6097	-0.0440	0.9257
CG6833	CG6833	1632262_at	0.3465	0.0774	-0.4034	0.0986	-0.0529	0.7935	0.2600	0.3231	0.7897	0.0009	0.5297	0.0029	-0.1531	0.8203	0.0018	0.9988	0.1550	0.6067
CG1381	CG1381	1632263_at	0.5232	0.1629	0.3853	0.3516	0.7630	0.0266	0.3518	0.6354	0.3394	0.3611	-0.0124	0.9785	0.0724	0.9751	0.4198	0.4702	0.3474	0.5672
CG1764	CG1764	1632264_at	-0.7864	0.0171	-0.4316	0.1389	-0.3624	0.2471	0.1587	0.8137	-0.4131	0.1177	-0.5718	0.0240	-0.0160	0.9935	-0.1167	0.8472	-0.1007	0.8637
---	---	1632265_at	0.1727	0.6097	-0.3284	0.1587	-0.3355	0.1037	0.0935	0.9311	0.4244	0.1811	0.3309	0.2480	-0.1845	0.8076	-0.1621	0.6361	0.0224	0.9614
CG13367	CG13367	1632266_at	-0.2421	0.3126	-0.0697	0.8120	-0.0253	0.9366	0.0863	0.9018	-0.2106	0.3644	-0.2970	0.1427	0.0740	0.9657	-0.0797	0.9111	-0.1537	0.7799
eEF1delta	eEF1delta	1632267_a_at	0.6605	0.0045	0.6874	0.0280	0.7113	0.0078	0.0309	0.9655	0.4675	0.0219	0.4366	0.0181	-0.0402	0.9787	0.5038	0.1432	0.5441	0.1490
---	---	1632268_at	0.2784	0.1557	0.3626	0.1332	0.1470	0.4405	-0.0856	0.8897	-0.0173	0.9552	0.0683	0.7630	0.2011	0.7644	0.0895	0.8071	-0.1116	0.7333
RpS3	ribosomal protein	1632269_at	0.2733	0.0691	0.1310	0.2283	0.2812	0.1343	0.1467	0.6936	0.0674	0.7397	-0.0793	0.6563	0.1530	0.7628	-0.0069	0.9872	-0.1599	0.4647
---	---	1632270_s_at	0.0876	0.6528	-0.0121	0.9663	0.1900	0.2437	0.2306	0.4262	0.2959	0.0676	0.0653	0.6983	0.1169	0.8395	0.0842	0.7718	-0.0327	0.9204
CG7971	anon-fast-evolving	1632271_a_at	0.3731	0.4129	-0.0834	0.9293	0.0427	0.8127	0.1294	0.7777	0.4906	0.0186	0.3611	0.0411	0.0327	0.9939	0.1415	0.9256	0.1088	0.9350
Moca-cyp	Moca-cyp	1632272_a_at	-0.3576	0.4908	-0.4491	0.4467	-0.6086	0.0030	0.1502	0.7412	0.2800	0.1607	0.1298	0.5009	0.0850	0.9816	0.2057	0.8473	0.1207	0.9109
CG5789	CG5789	1632273_at	-0.6434	0.0584	-0.6590	0.0686	-0.4004	0.0821	-0.1720	0.7531	-0.4657	0.0537	-0.2937	0.1638	-0.3870	0.7230	-0.4733	0.3006	-0.0863	0.8934
CG8134	CG8134	1632274_at	-0.1672	0.5644	0.4339	0.2482	0.1996	0.2221	-0.1128	0.8350	-0.3925	0.0598	-0.2796	0.1258	0.1753	0.8999	0.2429	0.6673	0.0676	0.9234
CG8671	CG8671	1632275_a_at	-0.2898	0.2918	1.2036	0.0781	0.7993	0.0125	-0.3627	0.6679	-0.6693	0.0969	-0.3066	0.4159	0.0971	0.9589	0.7797	0.1431	0.6826	0.2249
---	---	1632276_at	0.4082	0.0900	0.0772	0.7722	0.1391	0.4704	0.0305	0.9639	0.2456	0.1648	0.2151	0.1751	0.0156	0.9928	-0.1684	0.7200	-0.1840	0.6853
HDAC6	histone deacetyla	1632277_a_at	0.8787	0.3869	0.9831	0.6405	1.4740	0.0026	0.4049	0.3034	-0.4820	0.0433	-0.8869	0.0019	-0.1250	0.9913	-0.3510	0.9231	-0.2260	0.9444
CG30193	CG30193	1632278_s_at	-0.6808	0.0691	0.3745	0.0626	-0.0878	0.7667	-0.3988	0.3248	-0.6919	0.0102	-0.2931	0.1624	0.1585	0.8882	0.3489	0.4083	0.1904	0.6861
Rac1	Rac1	1632279_at	-0.0532	0.8558	0.1667	0.4102	-0.4700	0.0309	-0.2396	0.6122	0.2065	0.3967	0.4461	0.0418	0.3217	0.6749	0.3765	0.2086	0.0549	0.8972
mRpS35	mitochondrial ribo	1632280_at	-0.1546	0.3305	-0.0801	0.8185	-0.0196	0.9459	0.2660	0.5507	0.0585	0.8446	-0.2076	0.3237	0.2646	0.7576	0.3465	0.3255	0.0820	0.8635
CG4266	CG4266	1632281_at	-0.1735	0.8219	-0.1911	0.8459	-0.3445	0.0919	-0.1917	0.7604	0.1828	0.5249	0.3745	0.1196	-0.0301	0.9952	0.2460	0.8830	0.2761	0.8547
---	---	1632282_s_at	0.0651	0.7909	0.3003	0.0538	0.0258	0.9187	-0.1451	0.7556	-0.1183	0.5890	0.0268	0.9127	0.1233	0.8903	0.0685	0.8884	-0.0548	0.9057
CG8950	CG8950	1632283_at	0.4738	0.0920	0.0728	0.4504	0.2829	0.1328	-0.0338	0.9704	0.8694	0.0035	0.9032	0.0018	-0.2434	0.7245	0.2693	0.3591	0.5127	0.1192
---	---	1632284_s_at	0.0150	0.9368	0.0417	0.7554	0.2338	0.1018	-0.0091	0.9887	-0.1846	0.2170	-0.1755	0.1896	-0.0006	0.9998	-0.1157	0.6408	-0.1151	0.6394
CG2165 /// CG34036	CG2165 /// CG34(	1632285_at	-1.3089	0.0047	-1.3852	0.0946	-1.7088	0.0000	-0.4691	0.0898	-0.1458	0.3764	0.3233	0.0324	-0.1565	0.9506	-0.1908	0.8531	-0.0343	0.9769

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15161	CG15161	1632286_at	0.1810	0.2085	0.0490	0.7020	0.0477	0.8583	0.0832	0.8244	0.0213	0.9167	-0.0620	0.6834	0.0422	0.9635	-0.0431	0.9111	-0.0854	0.7734
CG32113 /// DmirCG32113	CG32113	1632287_at	0.0200	0.9344	-0.2568	0.6025	-0.3384	0.1246	-0.1962	0.6375	0.2936	0.1487	0.4898	0.0154	-0.0176	0.9939	0.1789	0.7906	0.1965	0.7514
CDC45L	Transcription unit	1632288_at	-0.0403	0.8606	0.7885	0.0642	1.0097	0.0055	0.0299	0.9693	-0.3581	0.0760	-0.3880	0.0370	-0.1047	0.9441	0.5447	0.2135	0.6494	0.1752
CG33333	CG33333	1632289_at	-0.1233	0.4598	0.1127	0.5263	-0.0531	0.7412	-0.0960	0.8527	-0.1868	0.3259	-0.0908	0.6360	0.1372	0.8202	0.0142	0.9734	-0.1230	0.6457
---	---	1632290_at	0.1490	0.5710	-0.0460	0.7344	-0.0156	0.9422	0.0155	0.9857	0.1946	0.3247	0.1791	0.3122	-0.1746	0.8202	-0.0829	0.8487	0.0917	0.8178
Snap	soluble NSF attac	1632291_at	-0.8681	0.0110	-0.5167	0.1030	-0.9819	0.0003	-0.2941	0.4631	-0.5956	0.0139	-0.3015	0.1189	0.1236	0.9011	-0.2151	0.5776	-0.3387	0.3671
Myt1	Myt1	1632292_at	-0.2836	0.3686	-0.4801	0.0349	-0.5401	0.0059	-0.1050	0.8096	0.5433	0.0078	0.6484	0.0021	0.0093	0.9952	0.4201	0.2593	0.4108	0.3002
CG13576	CG13576	1632293_at	0.0932	0.5738	0.1542	0.3716	0.0035	0.9885	0.0732	0.8968	0.0715	0.7460	-0.0017	0.9942	0.1452	0.8395	0.0991	0.7886	-0.0461	0.9097
sens	senseless	1632294_at	-0.0091	0.9785	-0.1361	0.4126	0.1766	0.4020	0.1363	0.8513	0.1388	0.6407	0.0025	0.9940	-0.1642	0.8611	0.0759	0.8928	0.2401	0.5512
---	---	1632295_s_at	-0.5397	0.0398	-1.6496	0.0775	-1.8350	0.0002	-0.2118	0.6615	0.9322	0.0026	1.1440	0.0006	0.1278	0.9588	-0.1056	0.9221	-0.2334	0.7748
---	---	1632296_at	0.0607	0.7846	0.1354	0.4790	0.1439	0.4848	0.1658	0.7143	0.0527	0.8438	-0.1131	0.5837	-0.0281	0.9781	-0.0990	0.7263	-0.0708	0.8168
Atg2	Autophagy-specifi	1632297_at	0.1381	0.7843	0.4374	0.2887	0.6317	0.0058	0.0111	0.9860	-0.3009	0.0430	-0.3120	0.0236	-0.2461	0.8815	0.0032	0.9991	0.2493	0.7306
mfas	midline fasciclin	1632298_s_at	-3.0534	0.0005	-4.5194	0.0062	-3.9426	0.0000	0.4831	0.4590	0.6952	0.0540	0.2122	0.5393	0.0284	0.9922	-0.7591	0.2346	-0.7875	0.2501
alphaPS4	alphaPS4	1632299_at	-0.1233	0.5720	-0.0039	0.9810	0.4501	0.0472	0.1561	0.6760	0.1780	0.3158	0.0218	0.9193	-0.2406	0.7812	0.1568	0.7128	0.3973	0.3011
---	---	1632300_at	0.0415	0.8989	-0.1256	0.2490	0.0714	0.6780	0.1963	0.6580	0.1254	0.5887	-0.0709	0.7614	-0.0818	0.8802	-0.0681	0.7943	0.0137	0.9646
CG9536	CG9536	1632301_at	0.1600	0.4509	0.6462	0.0346	0.7744	0.0022	0.0992	0.7670	-0.0514	0.7639	-0.1506	0.2385	-0.0472	0.9737	0.3793	0.2639	0.4265	0.2409
CG40496	CG40496	1632302_at	0.1987	0.3834	0.0532	0.6367	0.0591	0.7334	0.0149	0.9860	0.0841	0.7070	0.0692	0.7403	0.2197	0.6927	0.1056	0.6768	-0.1142	0.6419
Trc8	Trc8	1632303_a_at	-0.9315	0.2529	0.2516	0.7861	0.3987	0.1058	0.0248	0.9729	-0.8757	0.0012	-0.9005	0.0006	-0.0777	0.9898	0.4538	0.7743	0.5315	0.7155
CG8026	CG8026	1632304_at	-0.4617	0.0763	-0.0287	0.8677	-0.0085	0.9688	-0.1097	0.8392	-0.1614	0.4414	-0.0518	0.8216	-0.3264	0.6660	0.0459	0.9231	0.3723	0.2365
---	---	1632305_at	0.2023	0.3507	0.1510	0.2734	0.3661	0.0713	0.0332	0.9610	-0.0579	0.7951	-0.0911	0.6222	-0.0267	0.9831	0.0372	0.9353	0.0639	0.8675
CG9467	CG9467	1632306_at	0.3244	0.2139	0.2923	0.0747	0.2988	0.1146	0.0668	0.9339	0.7657	0.0068	0.6989	0.0061	0.2026	0.7644	0.8288	0.0274	0.6262	0.0654
---	---	1632307_at	0.0421	0.8389	0.0748	0.7409	0.1699	0.3218	0.0113	0.9922	-0.1200	0.6654	-0.1313	0.5912	-0.0486	0.9449	0.0650	0.8148	0.1135	0.6216
CG10414	CG10414	1632308_at	0.6203	0.0331	-0.1556	0.6122	0.0753	0.6518	0.0606	0.9154	0.8204	0.0015	0.7598	0.0012	-0.0922	0.9270	0.0813	0.8594	0.1736	0.6268
As	Drosophila Angelr	1632309_at	0.3755	0.0802	0.3759	0.0447	0.4766	0.0173	0.1167	0.8327	0.4238	0.0487	0.3071	0.1013	0.1462	0.8354	0.5752	0.0665	0.4289	0.1651
---	---	1632310_at	-0.0009	0.9967	-0.0097	0.9273	0.0783	0.6458	0.2507	0.4728	0.1329	0.5026	-0.1178	0.5180	0.0359	0.9640	0.0942	0.6977	0.0582	0.8299
---	---	1632311_at	0.1139	0.5943	0.1275	0.2049	0.0534	0.8272	-0.0962	0.8738	0.0111	0.9733	0.1074	0.6166	-0.0338	0.9611	0.0689	0.7707	0.1028	0.6228
CG30353	CG30353	1632312_at	-0.0594	0.8025	0.1201	0.4248	0.2569	0.0924	0.0451	0.9338	0.0142	0.9523	-0.0309	0.8683	-0.0933	0.8823	0.0562	0.8684	0.1495	0.5581
CG1998	CG1998	1632313_at	0.4384	0.4869	-1.0615	0.1970	-0.1733	0.5354	0.1120	0.8578	0.3668	0.1018	0.2549	0.2032	-0.7063	0.7779	-1.1066	0.2972	-0.4003	0.7495
Osi8	Osiris	1632314_at	0.1234	0.5109	0.0089	0.9341	-0.1324	0.6390	-0.0483	0.9624	-0.0208	0.9592	0.0275	0.9351	0.0577	0.9640	-0.0699	0.8903	-0.1276	0.7492
CG13035	CG13035	1632315_at	0.0575	0.8102	-0.2699	0.2730	-0.1944	0.2497	0.0047	0.9956	0.2663	0.2316	0.2616	0.1886	-0.2315	0.7644	-0.2031	0.5529	0.0284	0.9510
CG16935 /// DmirCG16935	CG16935	1632316_at	0.0802	0.6134	0.0132	0.9524	0.2948	0.1199	-0.0424	0.9380	-0.5394	0.0054	-0.4971	0.0047	-0.1702	0.7424	-0.5212	0.0391	-0.3511	0.1389
CG3036	CG3036	1632317_at	0.4770	0.5244	1.6086	0.0077	1.3640	0.0011	0.0253	0.9931	0.6442	0.2581	0.6189	0.2235	0.2286	0.9294	1.6831	0.0637	1.4544	0.1156
CG12773	CG12773	1632318_at	0.2055	0.6404	0.4019	0.3381	0.4943	0.0075	0.1596	0.7678	0.3997	0.0831	0.2401	0.2445	0.1253	0.9619	0.7884	0.2643	0.6631	0.3782
CG18598	CG18598	1632319_at	-0.6848	0.0039	-0.0761	0.7247	-0.2374	0.2822	-0.3277	0.4324	-0.4576	0.0507	-0.1299	0.5669	-0.0394	0.9816	-0.0367	0.9531	0.0027	0.9974
CG32080	CG32080	1632320_at	0.1208	0.5246	-0.0977	0.5573	-0.0686	0.6879	-0.0488	0.9345	0.1244	0.5003	0.1731	0.2735	-0.0372	0.9588	-0.0766	0.7439	-0.0394	0.8861
mthl7	Mth-like 7	1632321_a_at	0.0228	0.8972	-0.0238	0.8691	-0.0017	0.9943	0.0637	0.9068	0.0048	0.9855	-0.0589	0.7589	0.0616	0.9441	0.0073	0.9903	-0.0543	0.8801
CG16953	CG16953	1632322_at	-0.5119	0.0199	-0.3975	0.0818	-0.6793	0.0033	-0.3726	0.1950	-0.3547	0.0466	0.0178	0.9343	-0.1972	0.7533	-0.3325	0.1985	-0.1354	0.6389
CG5451	CG5451	1632323_at	0.0195	0.9232	-0.2742	0.4857	-0.0752	0.6631	0.2333	0.4333	0.4075	0.0209	0.1741	0.2342	0.1581	0.8903	0.2991	0.4999	0.1410	0.7830
CG11201	CG11201	1632324_at	0.0517	0.7742	0.1858	0.3267	0.1592	0.5563	0.1269	0.8281	-0.0216	0.9491	-0.1485	0.4902	0.1946	0.7780	0.2336	0.4355	0.0391	0.9246
tio	tiptop	1632325_at	0.0477	0.7659	-0.0229	0.8333	0.0733	0.7811	0.2115	0.5008	0.2242	0.1764	0.0126	0.9523	-0.0245	0.9877	0.0456	0.9366	0.0701	0.8871
CG30105	CG30105	1632326_at	-0.3355	0.2119	-0.2614	0.4737	-0.3491	0.2962	-0.1463	0.8650	0.1801	0.5873	0.3264	0.2324	-0.0357	0.9894	0.2689	0.6619	0.3045	0.6141
CG4854	CG4854	1632327_at	0.1479	0.4339	0.0040	0.9781	-0.1340	0.5639	-0.0569	0.9451	0.3328	0.1564	0.3897	0.0676	0.2186	0.7230	0.1793	0.5181	-0.0394	0.9158
serp	Serpentine	1632328_at	0.8492	0.0462	0.8069	0.1673	0.6309	0.0139	-0.0812	0.9449	-0.4365	0.1905	-0.3553	0.2369	0.1000	0.9499	-0.4629	0.3158	-0.5629	0.2527
---	---	1632329_at	0.1592	0.5923	0.0786	0.4551	0.0691	0.6991	-0.0331	0.9759	0.0712	0.8356	0.1043	0.7136	-0.2262	0.7363	-0.3276	0.2374	-0.1014	0.7587
CG14323	CG14323	1632330_at	-0.7565	0.0691	-0.0328	0.8026	0.3352	0.0530	0.0660	0.9647	-0.6449	0.1016	-0.7109	0.0488	-0.2459	0.7070	-0.0127	0.9797	0.2332	0.4045
Toll-9	Toll-9	1632331_at	0.2671	0.1420	-0.0233	0.8977	-0.2678	0.0764	-0.1724	0.5621	0.0327	0.8718	0.2051	0.1307	0.0663	0.9314	-0.1987	0.4255	-0.2650	0.3009
CG5447	CG5447	1632332_at	0.1555	0.4772	0.9298	0.0324	0.7758	0.0043	-0.1420	0.7803	-0.5647	0.0157	-0.4227	0.0328	0.1728	0.8725	0.2907	0.5075	0.1180	0.8247
CG6108	CG6108	1632333_at	0.2646	0.3925	0.4989	0.1388	0.0984	0.7172	-0.2306	0.3744	0.2652	0.0735	0.4958	0.0033	0.0209	0.9928	0.2633	0.6634	0.2423	0.6909
---	---	1632334_at	0.0250	0.9011	-0.0590	0.6453	-0.1013	0.5765	0.0587	0.9434	0.1746	0.4738	0.1159	0.6252	0.0605	0.9142	0.1251	0.5426	0.0646	0.7820
RpL27	Ribosomal protein	1632335_at	0.5722	0.0467	1.0676	0.0255	0.8925	0.0066	0.0782	0.8721	-0.0855	0.6533	-0.1637	0.2840	0.0117	0.9939	0.0952	0.8359	0.0835	0.8509
CG15296	CG15296	1632336_at	-0.0106	0.9540	-0.1361	0.4120	0.0745	0.6698	0.1976	0.6387	0.2555	0.2136	0.0579	0.8030	-0.0793	0.8909	0.0797	0.7604	0.1590	0.4822
Sur	sulfonyl urea rece	1632337_at	-2.2272	0.0005	-2.3358	0.0077	-2.5345	0.0000	-0.2044	0.5067	-0.1286	0.4480	0.0758	0.6528	0.2792	0.7686	-0.2357	0.5810	-0.5149	0.2184
CG33523	MSP protein	1632338_a_at	-0.5942	0.0196	0.5576	0.3381	0.9965	0.0056	-0.0549	0.9567	-1.1191	0.0024	-1.0642	0.0017	-0.2977	0.8097	0.1356	0.8460	0.4333	0.4076
cher	filamin	1632339_s_at	-1.8214	0.0023	-0.4477	0.4609	-0.0996	0.7672	0.0292	0.9649	-0.7965	0.0016	-0.8257	0.0008	-0.2926	0.8814	0.6045	0.4164	0.8970	0.2479
---	---	1632340_at	0.4337	0.2024	0.1412	0.2579	0.3535	0.0386	0.0856	0.8517	0.0643	0.7440	-0.0213	0.9166	0.0096	0.9939	-0.0738	0.8393	-0.0834	0.8001
CG13559	CG13559	1632341_at	-0.2165	0.4404	-1.3369	0.0401	-0.7082	0.0242	0.6083	0.2638	0.6269	0.0559	0.0186	0.9635	-0.1055	0.9495	-0.3801	0.4417	-0.2746	0.6052
CG33080	CG33080	1632342_a_at	-0.7740	0.0240	0.8589	0.3741	0.4200	0.3310	-1.0245	0.1545	-2.8101	0.0004	-1.7856	0.0014	-0.4557	0.8465	-0.9165	0.3358	-0.4608	0.6644
Alas	Aminolevulinate s	1632343_at	-1.0583	0.0112	-0.5067	0.0442	0.0131	0.9572	0.2071	0.5142	-1.0878	0.0004	-1.2949	0.0001	-0.3289	0.7464	-0.3761	0.3706	-0.0472	0.9371
---	---	1632344_at	0.0553	0.8406	-0.1750	0.2734	-0.1326	0.6118	0.1686	0.8190	0.3965	0.1670	0.2279	0.3959	-0.1377	0.8320	-0.0928	0.7804	0.0449	0.9046
CG8353	CG8353	1632345_at	1.5996	0.0045	-0.4711	0.6545	0.3381	0.4536	0.6925	0.3359	1.2975	0.0079	0.6050	0.1028	-0.0778	0.9869	-0.6674	0.5578	-0.5896	0.6149
Kif19A	Kif19A	1632346_at	-0.1157	0.6207	-0.2073	0.4857	-0.3656	0.1212	0.0264	0.9803	0.0831	0.7925	0.0567	0.8493	0.1444	0.8718	-0.1029	0.8252	-0.2473	0.5008
CG9525	CG9525	1632347_at	0.0340	0.8989	-0.0326	0.7379	-0.0607	0.7226	0.0182	0.9777	0.0833	0.6577	0.0650	0.7129	0.0010	0.9998	-0.0064	0.9935	-0.0074	0.9889
CG9452	CG9452	1632348_at	-1.3388	0.0325	0.2340	0.5751	-0.3705	0.5695	-0.8472	0.4113	-0.8385	0.1407	0.0087	0.9904	-0.3887	0.8870	0.5045	0.6512	0.8932	0.3921
---	---	1632349_at	0.1769	0.3115	0.1296	0.3512	-0.0198	0.9303	-0.0251	0.9715	0.1396	0.4503	0.1647	0.3081	-0.0673	0.9238	0.0185	0.9620	0.0858	0.7513
CG17575 /// DyakCG17575	CG17575	1632350_at	0.1816	0.3634	0.0116	0.9132	0.0510	0.8142	0.0505	0.9380	0.0588	0.8008	0.0083	0.9718	0.0244	0.9895	-0.0209	0.9754	-0.0453	0.9341
Femcoat	FEMCOAT	1632351_at	0.1615	0.3519	0.0373	0.8215	0.1954	0.3034	0.0337	0.9641	0.1350	0.5248	0.1013	0.6165	-0.0649	0.9128	0.1545	0.4622	0.2194	0.3034
CG30007	CG30007	1632352_a_at	0.0317	0.9344	-0.6039	0.0949	-0.5094	0.0060	-0.0527	0.9482	0.9577	0.0021	1.0104	0.0010	-0.1965	0.8680	0.2431	0.6310	0.4396	0.3616
CG7884	CG7884	1632353_at	-3.1509	0.0010	-2.1462	0.0040	-1.6903	0.0080	0.4845	0.7773	-0.6263	0.3931	-1.1109	0.0833	-0.1244	0.8461	-0.0581	0.8778	0.0663	0.8444
---	---	1632354_at	-0.0195	0.9284	0.0649	0.5575	-0.1299	0.4870	0.0177	0.9755	0.0597	0.7139	0.0420	0.7894	-0.0123	0.9914	0.0400	0.9170	0.0523	0.8761

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
lack	Smurf	1632355_at	0.4252	0.2705	0.3862	0.0626	0.0616	0.7138	-0.1758	0.6081	0.3325	0.0575	0.5083	0.0060	0.2723	0.8305	0.3211	0.5653	0.0488	0.9479
Gr65a	Gustatory recepto	1632356_at	-0.0078	0.9758	-0.1968	0.1722	-0.1718	0.2524	-0.0771	0.8836	0.0456	0.8439	0.1228	0.4710	-0.0817	0.8924	-0.1508	0.5191	-0.0691	0.8009
CG40164	CG40164	1632357_x_at	0.1751	0.3467	-0.0039	0.9741	0.2043	0.3146	0.0348	0.9558	0.0492	0.8144	0.0144	0.9463	-0.0588	0.9400	-0.0728	0.8150	-0.0140	0.9696
CG32297	CG32297	1632358_at	0.0015	0.9952	-0.1166	0.5976	-0.0959	0.5993	-0.1628	0.6791	0.0057	0.9830	0.1686	0.3144	-0.1916	0.7848	-0.1410	0.6738	0.0506	0.9031
Takl2	Tak1-like 2	1632359_at	-0.0010	0.9983	-0.0012	0.9966	-0.0772	0.6423	-0.0009	0.9994	-0.1672	0.6662	-0.1664	0.6334	0.0189	0.9914	-0.2756	0.4656	-0.2944	0.4436
CG31038	CG31038	1632360_s_at	0.5380	0.1329	0.4393	0.3353	-0.6588	0.1119	-0.2926	0.6412	0.7179	0.0270	1.0105	0.0037	0.8849	0.6496	0.6326	0.4184	-0.2523	0.7878
retm	real-time	1632361_at	0.2190	0.7160	1.0969	0.0644	0.7304	0.0233	-0.0346	0.9777	-0.6056	0.0620	-0.5710	0.0505	0.2376	0.9298	0.2774	0.8045	0.0398	0.9769
CG13616	CG13616	1632362_at	0.1338	0.7030	-0.1467	0.5264	0.0389	0.8281	0.1355	0.8422	0.0785	0.7998	-0.0569	0.8455	-0.1437	0.8379	-0.1398	0.6604	0.0039	0.9941
---	---	1632363_at	0.1521	0.5528	0.1056	0.4282	-0.1556	0.4944	-0.1217	0.8197	0.0636	0.8050	0.1853	0.3334	0.0051	0.9964	-0.0577	0.8999	-0.0629	0.8786
Nup170	Nup170	1632364_at	0.0157	0.9736	0.1913	0.6799	0.3981	0.0293	0.0669	0.9022	0.1514	0.4042	0.0845	0.6414	-0.0892	0.9672	0.3392	0.5715	0.4284	0.4677
---	---	1632365_at	0.0975	0.4850	-0.0954	0.4867	0.3400	0.0488	0.0463	0.9196	0.1072	0.4647	0.0609	0.6783	-0.2124	0.6955	-0.0934	0.7240	0.1191	0.6311
---	---	1632366_at	0.3400	0.1456	0.3859	0.2458	0.1173	0.6129	-0.0261	0.9705	0.0383	0.8731	0.0644	0.7407	0.2029	0.7768	0.0894	0.8254	-0.1135	0.7495
Mst35Ba	protamine	1632367_at	0.0317	0.8816	-0.1984	0.2783	-0.3330	0.2248	-0.3161	0.5455	0.1040	0.7464	0.4201	0.0840	-0.1904	0.7848	-0.1150	0.7442	0.0754	0.8444
djl	don juan like	1632368_at	-0.1570	0.5548	-0.3001	0.1377	-0.0426	0.8438	0.1182	0.7975	0.1804	0.3427	0.0622	0.7613	-0.1046	0.9298	0.0707	0.9046	0.1753	0.6785
CG5142	CG5142	1632369_at	-0.3535	0.0504	-0.2458	0.2607	-0.6065	0.0305	-0.2911	0.3053	-0.2516	0.1305	0.0395	0.8366	-0.0554	0.9679	0.0814	0.8760	0.1368	0.7442
TfIIA-S	Transcription-fact	1632370_at	0.2546	0.3801	0.2570	0.1265	0.3214	0.1524	0.1395	0.7404	-0.3264	0.0799	-0.4659	0.0126	0.1230	0.8655	-0.2655	0.3465	-0.3885	0.2031
---	---	1632371_s_at	0.1655	0.5014	-0.0025	0.9836	0.0781	0.6562	0.0341	0.9515	-0.0038	0.9876	-0.0378	0.8296	-0.0305	0.9721	-0.0942	0.7065	-0.0636	0.8178
CG31076	CG31076	1632372_at	-1.5674	0.0024	-0.3345	0.1894	-1.2202	0.0116	-0.3693	0.5756	-1.3705	0.0025	-1.0012	0.0060	0.1611	0.9168	-0.4838	0.3534	-0.6450	0.2429
CG34015	CG34015	1632373_s_at	0.6019	0.0182	0.4862	0.1565	0.0625	0.7861	0.0164	0.9852	0.4167	0.0430	0.4002	0.0323	0.3899	0.6749	0.2183	0.5825	-0.1716	0.6761
CG6769	CG6769	1632374_at	0.0458	0.8979	0.0889	0.7556	0.6673	0.0122	0.4893	0.2492	0.2923	0.2458	-0.1970	0.3998	-0.1316	0.9226	0.2550	0.5929	0.3866	0.3981
---	---	1632375_at	0.0724	0.7658	-0.0469	0.6940	-0.0019	0.9944	0.0977	0.8507	0.2494	0.1852	0.1516	0.3883	0.0617	0.9421	-0.0704	0.8428	-0.1321	0.6392
CG5644	CG5644	1632376_at	0.1397	0.5698	-0.0915	0.6192	-0.0899	0.6505	0.0714	0.8908	0.0321	0.8920	-0.0393	0.8456	-0.1604	0.8510	-0.3133	0.3669	-0.1529	0.6980
CG8399	CG8399	1632377_at	-0.7706	0.0970	-0.8596	0.0604	-0.4450	0.1618	0.3401	0.3848	-0.4761	0.0381	-0.8161	0.0022	-0.1070	0.9651	-0.5737	0.3724	-0.4667	0.4903
---	---	1632378_at	0.4708	0.0679	-0.2885	0.2074	0.0092	0.9736	0.3758	0.2805	0.3840	0.0648	0.0082	0.9744	-0.2169	0.7808	-0.4447	0.1722	-0.2278	0.5158
---	---	1632379_at	0.0229	0.9011	-0.2313	0.2576	-0.1941	0.2909	-0.0779	0.8964	0.1556	0.4474	0.2334	0.1844	-0.0273	0.9816	-0.0454	0.9144	-0.0181	0.9640
mnb	Minibrain	1632380_at	-1.5205	0.0681	-0.8434	0.3741	-1.2102	0.0117	0.0628	0.9523	-0.5024	0.0860	-0.5652	0.0370	0.2983	0.9514	0.1316	0.9559	-0.1666	0.9350
dome	verstopft	1632381_at	-0.6131	0.2809	-0.9705	0.3357	-0.8279	0.0800	0.2923	0.6144	0.8118	0.0129	0.5195	0.0505	0.2294	0.9589	0.4711	0.7506	0.2417	0.8884
---	---	1632382_at	0.1170	0.4080	0.1340	0.5596	0.1631	0.2629	0.0972	0.8280	-0.0530	0.8014	-0.1502	0.3399	0.0700	0.8825	0.0622	0.7808	-0.0079	0.9783
Art1	Arginine methyltra	1632383_at	0.3573	0.1316	0.5296	0.1476	0.9254	0.0031	0.1611	0.7891	0.0975	0.7372	-0.0636	0.8206	-0.1895	0.8439	0.3714	0.3350	0.5609	0.1809
NC2beta	NC2beta	1632384_at	-0.1985	0.4238	0.0846	0.7699	0.2348	0.4453	0.2219	0.7380	0.1273	0.7069	-0.0946	0.7702	0.1928	0.8681	0.3663	0.4246	0.1735	0.7439
tai	Taiman	1632385_at	0.5163	0.6863	1.2930	0.2623	1.3489	0.0005	-0.2511	0.8156	-0.5878	0.1626	-0.3367	0.3918	0.0222	0.9977	0.4368	0.8528	0.4146	0.8515
---	---	1632386_at	0.2173	0.2245	0.0655	0.7927	0.2478	0.2902	-0.0394	0.9744	0.1534	0.6451	0.1928	0.4987	0.0705	0.9589	0.1447	0.7554	0.0742	0.8902
CG11983	CG11983	1632387_at	0.0816	0.7365	-0.2146	0.3622	0.0465	0.8035	0.0697	0.9011	0.2589	0.1527	0.1893	0.2466	-0.0357	0.9773	-0.0267	0.9557	0.0090	0.9849
CG34354	CG12425	1632388_at	-0.0491	0.8223	0.0979	0.5437	-0.1878	0.3082	-0.1942	0.5947	-0.2236	0.2265	-0.0293	0.8951	0.0796	0.9365	0.0148	0.9796	-0.0649	0.8800
CG14812	CG14812	1632389_at	0.0748	0.6735	-0.0385	0.8497	-0.0065	0.9809	0.0153	0.9838	-0.0397	0.8656	-0.0550	0.7809	0.0798	0.9441	-0.1581	0.6789	-0.2379	0.5041
cer	crammer	1632390_at	0.4981	0.0442	0.4487	0.3381	0.3769	0.0555	0.0144	0.9909	-0.0444	0.9063	-0.0588	0.8526	0.0864	0.9374	-0.0353	0.9499	-0.1217	0.7611
CG14234	CG14234	1632391_at	-0.0528	0.7444	0.0155	0.8807	-0.0235	0.9245	-0.0315	0.9632	0.0029	0.9910	0.0344	0.8722	-0.1142	0.8541	-0.0810	0.8048	0.0332	0.9246
Slob	giant slob	1632392_s_at	5.3933	0.0011	2.9399	0.0025	5.8901	0.0000	3.0776	0.0233	2.7055	0.0037	-0.3721	0.6161	-0.0875	0.9462	0.1201	0.8132	0.2076	0.6233
CG18301	CG18301	1632393_at	0.4498	0.1623	0.0903	0.4819	0.6064	0.0046	0.0018	0.9980	-0.0137	0.9569	-0.0154	0.9425	-0.3583	0.5754	-0.2917	0.2941	0.0667	0.8578
---	---	1632394_s_at	-0.4398	0.0289	-0.3138	0.1226	-0.6353	0.0018	-0.2620	0.3933	-0.0959	0.6156	0.1661	0.2889	0.0913	0.9064	0.1374	0.6497	0.0461	0.9049
Vha100-1	Vha100-1	1632395_s_at	-0.9782	0.0080	0.0975	0.8327	0.3231	0.0680	-0.0691	0.9329	-0.9121	0.0034	-0.8430	0.0029	-0.2665	0.8292	0.0228	0.9812	0.2893	0.6030
trsn	translin	1632396_at	0.2851	0.1343	0.0727	0.8520	0.1176	0.6030	-0.0747	0.9066	0.1181	0.5970	0.1928	0.2989	-0.0438	0.9816	-0.0205	0.9784	0.0233	0.9716
CG16868 /// DsecCG16868	CG16868	1632397_at	0.0780	0.7642	-0.3681	0.0613	-0.4186	0.0810	0.1372	0.8550	0.6505	0.0247	0.5133	0.0408	0.1830	0.7588	0.2008	0.4203	0.0179	0.9613
CG31735	CG31735	1632398_a_at	0.0150	0.9596	-0.0889	0.4712	0.0686	0.7524	-0.0553	0.9412	0.0382	0.8962	0.0935	0.6767	-0.0891	0.9174	-0.1477	0.6483	-0.0586	0.8864
CG34007	CG34007	1632399_at	-2.8983	0.0023	-3.3678	0.0027	-3.4541	0.0000	-0.0599	0.9627	0.1404	0.7239	0.2003	0.5501	-0.0381	0.9806	-0.2751	0.4185	-0.2370	0.5041
CG3841 /// nmo	CG3841 /// nemo	1632400_at	0.1452	0.3301	0.1062	0.4399	0.0395	0.8380	-0.0131	0.9864	0.1304	0.5064	0.1435	0.4083	0.0228	0.9872	-0.0317	0.9514	-0.0546	0.9031
CG15747	CG15747	1632401_at	-0.6701	0.0502	-0.4520	0.2657	-0.4149	0.1109	-0.0059	0.9952	-0.2394	0.2258	-0.2335	0.1861	-0.1762	0.8930	-0.0527	0.9470	0.1235	0.8435
CG6744	CG6744	1632402_at	0.3553	0.1988	0.4125	0.0880	0.5787	0.0032	0.0855	0.8350	0.0282	0.8940	-0.0573	0.7307	-0.1786	0.8541	-0.0316	0.9620	0.1470	0.7503
CG12263	CG12263	1632403_at	-0.0421	0.8053	0.0703	0.9013	-0.3332	0.1548	-0.2054	0.6360	0.5718	0.0153	0.7771	0.0023	0.0002	0.9999	0.5548	0.2824	0.5547	0.3083
rtet	tetracycline resist	1632404_at	-0.4062	0.0233	-0.4961	0.0256	-0.6158	0.0042	-0.1799	0.5407	0.0768	0.6567	0.2567	0.0641	0.0026	0.9978	0.0021	0.9973	-0.0005	0.9992
---	---	1632405_at	0.0152	0.9552	0.1049	0.3192	0.1601	0.4010	-0.0840	0.8419	-0.0285	0.8934	0.0555	0.7420	-0.0504	0.9514	0.0422	0.9095	0.0925	0.7417
CG9117	CG9117	1632406_at	-0.1435	0.5557	-0.0037	0.9924	-0.0446	0.7797	-0.1573	0.7204	-0.2247	0.2597	-0.0674	0.7564	-0.1815	0.8160	-0.0977	0.8153	0.0837	0.8378
Tre	Gustatory recepto	1632407_at	-0.0190	0.9495	0.1282	0.3430	-0.0075	0.9830	-0.0929	0.9039	-0.0784	0.7984	0.0145	0.9635	0.0468	0.9467	-0.0525	0.8578	-0.0993	0.6670
Dhc64C	dynein	1632408_at	0.1021	0.6448	0.0090	0.9659	0.0590	0.7026	-0.0855	0.8470	0.0141	0.9532	0.0996	0.5288	-0.1440	0.8228	-0.0616	0.8718	0.0823	0.8000
Pka-C1	protein kinase A	1632409_a_at	-0.6510	0.1342	-1.2760	0.0972	-1.5721	0.0089	0.1879	0.8384	0.8499	0.0220	0.6620	0.0385	0.4474	0.8558	0.1968	0.8963	-0.2506	0.8467
CG12093	CG12093	1632410_at	-0.2595	0.3972	-0.4425	0.3354	-0.3311	0.0470	0.0975	0.8076	0.2617	0.1009	0.1643	0.2519	0.0563	0.9816	0.1318	0.8624	0.0755	0.9197
---	---	1632411_at	-0.1060	0.5253	-0.0171	0.8803	0.1723	0.4453	0.0895	0.8336	-0.0340	0.8735	-0.1235	0.4127	-0.1292	0.7893	-0.0139	0.9672	0.1153	0.6021
---	---	1632412_s_at	-0.1307	0.5478	0.0403	0.8832	0.0160	0.9558	0.1584	0.8073	0.1163	0.6951	-0.0421	0.8916	0.1153	0.8940	-0.0942	0.8231	-0.2095	0.5369
CG5343	CG5343	1632413_at	-0.0325	0.8748	-0.2966	0.2926	-0.2757	0.1950	0.1029	0.8883	0.2533	0.3085	0.1504	0.5316	0.0259	0.9853	0.0173	0.9758	-0.0086	0.9868
CG8546	CG8546	1632414_at	0.1664	0.6061	-0.0267	0.8180	-0.0376	0.8399	-0.0639	0.9110	0.0834	0.6872	0.1473	0.3788	-0.0142	0.9913	-0.1011	0.7562	-0.0869	0.7909
Dcp2	CG6169	1632415_s_at	0.3440	0.4840	0.8173	0.0498	0.7404	0.0050	0.1297	0.9255	0.0791	0.8891	-0.0506	0.9215	0.1696	0.9012	0.4782	0.3269	0.3086	0.5615
---	---	1632416_at	0.0361	0.8693	-0.3174	0.1194	-0.5711	0.0187	-0.0496	0.9172	0.2720	0.0688	0.3216	0.0232	0.0179	0.9864	-0.0462	0.9009	-0.0641	0.8368
CG5021	CG5021	1632417_a_at	0.4345	0.0762	0.4737	0.0718	0.4256	0.0465	-0.0296	0.9744	0.3054	0.1662	0.3350	0.0910	0.1010	0.8846	0.3909	0.1374	0.2899	0.2855
pum	ovarette	1632418_s_at	-0.1247	0.8179	-0.8734	0.2787	-2.2903	0.0007	-0.3986	0.5128	1.4500	0.0016	1.8486	0.0004	0.9297	0.6955	0.7633	0.4410	-0.1664	0.9031
CG30061	CG30061	1632419_at	-0.5439	0.1587	0.0378	0.9298	0.8672	0.0440	-0.3086	0.7558	-0.7725	0.0739	-0.4640	0.2262	-0.9735	0.3712	0.0475	0.9617	1.0210	0.1190
CG1135	CG1135	1632420_at	-0.1847	0.3627	0.1485	0.3481	0.0519	0.8438	0.0831	0.8908	-0.4971	0.0226	-0.5802	0.0070	0.1530	0.8692	-0.2223	0.5711	-0.3753	0.3248
CG9531	CG9531	1632421_at	-0.8211	0.0310	-0.7137	0.0435	-0.9363	0.0002	-0.2840	0.5311	-0.2033	0.4026	0.0807	0.7549	0.0778	0.9412	-0.2035	0.5523	-0.2813	0.3986
CG12901 /// DsimCG12901	CG12901	1632422_s_at	0.0448	0.8158	0.2454	0.0808	0.1290	0.5202	-0.0084	0.9909	-0.0978	0.5691	-0.0895	0.5707	0.0220	0.9871	0.1066	0.7808	0.0846	0.8287
her	lethal(2)maternal	1632423_at	-0.0786	0.8043	-0.0063	0.9860	-0.3093	0.3491	-0.3408	0.3822	0.0470	0.8723	0.3878	0.0534	-0.1567	0.9238	0.2151	0.7305	0.3718	0.5054

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13995	CG13995	1632424_at	-0.6337	0.0364	0.0430	0.7483	-0.1727	0.3469	-0.1947	0.7160	-0.6972	0.0115	-0.5024	0.0288	-0.0445	0.9677	-0.0177	0.9704	0.0268	0.9471
---	---	1632425_at	0.3593	0.2062	0.1176	0.4389	0.3539	0.1494	0.1008	0.8337	0.0436	0.8531	-0.0572	0.7772	-0.0517	0.9751	-0.1051	0.8501	-0.0534	0.9242
---	---	1632426_at	0.1990	0.2861	-0.0121	0.9659	0.0175	0.9511	0.1186	0.8073	0.1308	0.5251	0.0122	0.9597	0.1117	0.8878	-0.0470	0.9188	-0.1588	0.6195
---	---	1632427_at	0.1705	0.4454	-0.1370	0.3762	0.0098	0.9654	0.0565	0.9422	0.2294	0.3052	0.1728	0.4023	0.0045	0.9974	0.0637	0.8919	0.0592	0.8923
CG11345	CG11345	1632428_at	0.1565	0.4846	0.0704	0.5408	-0.2683	0.3241	0.0027	0.9962	0.0033	0.9901	0.0006	0.9981	0.0529	0.9514	-0.1563	0.5668	-0.2092	0.4323
Rab7	Rab-protein 7	1632429_at	0.2047	0.2296	0.0448	0.6930	-0.0516	0.7922	-0.0803	0.8402	-0.1354	0.3728	-0.0550	0.7295	0.0564	0.9611	-0.2671	0.4041	-0.3235	0.3276
Tsf1	Transferrin	1632430_at	0.1502	0.8305	0.9758	0.6816	1.3179	0.0076	-0.1124	0.8967	-0.8438	0.0099	-0.7314	0.0116	-0.5382	0.9441	-0.2884	0.9391	0.2498	0.9387
Ance	anon-fast-evolving	1632431_s_at	-1.0083	0.0532	-0.5727	0.0296	-1.5486	0.0020	-0.8286	0.2501	-1.2494	0.0106	-0.4207	0.2773	0.2174	0.8854	-0.7765	0.1602	-0.9939	0.1162
CG9850	Dlg-interacting me	1632432_at	-1.4712	0.0085	-2.0465	0.0129	-2.3837	0.0002	-0.4804	0.2477	-0.3902	0.1121	0.0902	0.7330	-0.1859	0.9340	-0.9426	0.1731	-0.7567	0.3007
CG17598	CG17598	1632433_at	0.2151	0.6083	-0.2150	0.1230	-0.1848	0.3924	0.0375	0.9677	0.8839	0.0040	0.8464	0.0029	-0.0091	0.9963	0.4031	0.3311	0.4122	0.3470
l(3)70Da	lethal (3) 70Da	1632434_at	-0.1332	0.5872	0.4715	0.1574	0.6178	0.0034	-0.0790	0.9168	-0.3254	0.1635	-0.2463	0.2424	-0.1250	0.8882	0.3206	0.3236	0.4456	0.2037
CG7804	CG7804	1632435_at	0.1140	0.5036	0.0589	0.6477	0.1215	0.5835	0.0454	0.9435	-0.0068	0.9804	-0.0521	0.7975	-0.0525	0.9499	-0.0274	0.9460	0.0251	0.9431
---	---	1632436_at	0.1859	0.3436	0.3525	0.2535	0.2175	0.4313	-0.2001	0.7857	0.1103	0.7618	0.3104	0.2558	0.2665	0.8369	0.1929	0.7642	-0.0736	0.9198
CG31846	CG31846	1632437_at	0.1612	0.5576	-0.5900	0.0664	-0.4250	0.0400	0.2380	0.5008	0.6115	0.0065	0.3735	0.0333	0.0538	0.9777	-0.2157	0.6713	-0.2695	0.5816
CG2091	CG2091	1632438_at	0.3029	0.1581	0.1824	0.4523	0.0414	0.9011	-0.1823	0.7929	0.2103	0.4781	0.3926	0.1201	0.0051	0.9964	0.1533	0.5907	0.1482	0.6093
TpnC73F	troponin C	1632439_at	-0.6464	0.0407	-1.1935	0.0460	-1.6086	0.0019	-0.1799	0.6840	0.7750	0.0034	0.9549	0.0008	-0.0859	0.9530	0.0937	0.8807	0.1796	0.7144
CG7630	CG7630	1632440_at	-0.2401	0.3735	0.3918	0.2645	0.3538	0.0501	-0.1307	0.7931	-1.1049	0.0007	-0.9742	0.0007	-0.0989	0.9342	-0.3872	0.3000	-0.2883	0.4677
CG14669	CG14669	1632441_at	0.0691	0.8705	0.1221	0.4479	0.4337	0.0197	0.0861	0.8676	0.0499	0.8294	-0.0362	0.8645	-0.0079	0.9958	0.1704	0.6501	0.1783	0.6320
---	---	1632442_at	0.2778	0.2312	-0.1704	0.4435	-0.2821	0.2248	0.1895	0.7672	0.2638	0.3413	0.0742	0.8085	0.0198	0.9862	-0.1915	0.4588	-0.2113	0.4184
CG10274	CG10274	1632443_at	-0.3099	0.3940	0.5338	0.0901	0.3895	0.0207	-0.0130	0.9882	-0.3036	0.1362	-0.2906	0.1110	0.0342	0.9884	0.4654	0.3448	0.4312	0.4017
CG15170	CG15170	1632444_at	0.2329	0.1686	0.0719	0.5598	0.3845	0.0334	0.0709	0.9039	0.0877	0.6887	0.0169	0.9439	0.0175	0.9872	-0.1121	0.6861	-0.1296	0.6290
---	---	1632445_at	0.0852	0.5575	0.1932	0.3392	0.1202	0.4929	-0.0057	0.9940	-0.0273	0.8968	-0.0216	0.9078	0.0861	0.8768	0.0409	0.9009	-0.0452	0.8779
CG15742	CG15742	1632446_at	0.1666	0.3602	0.1517	0.6272	0.1945	0.2144	0.1198	0.7327	0.0700	0.6970	-0.0498	0.7728	-0.0110	0.9939	0.0855	0.8468	0.0965	0.8103
B52	Serine/arginine ric	1632447_at	0.2954	0.2432	-0.0396	0.6935	0.0787	0.6988	-0.0236	0.9803	0.0512	0.8692	0.0749	0.7728	-0.0698	0.9515	-0.2456	0.4865	-0.1758	0.6366
kel	Kelch	1632448_s_at	-0.4302	0.3118	-0.2784	0.1110	-0.7181	0.0066	-0.5302	0.4141	-0.6071	0.0928	-0.0769	0.8545	-0.2243	0.6557	-0.2580	0.1925	-0.0337	0.9055
---	---	1632449_at	-0.0968	0.5982	0.0679	0.6750	0.1280	0.4983	0.0139	0.9852	-0.1653	0.3319	-0.1792	0.2333	-0.0811	0.9238	-0.0014	0.9991	0.0797	0.8209
CG13494	CG13494	1632450_at	-0.0277	0.8720	-0.2102	0.5096	-0.0697	0.7496	-0.0040	0.9956	0.1572	0.4734	0.1612	0.4105	-0.1889	0.7822	-0.1496	0.6404	0.0392	0.9231
Cyp304a1	Cyp304a1	1632451_at	1.8281	0.2882	0.5186	0.8437	2.4514	0.0152	1.9883	0.0104	1.9672	0.0009	-0.0211	0.9658	0.1064	0.9943	0.9363	0.8372	0.8300	0.8494
CG18258	CG18258	1632452_at	0.0125	0.9827	1.3369	0.1601	1.2726	0.0250	-0.1863	0.9187	-0.4368	0.4555	-0.2506	0.6681	-0.1495	0.9672	0.9218	0.3236	1.0713	0.2784
CG16905	CG16905	1632453_at	0.9165	0.5835	-2.4263	0.4199	0.8819	0.3428	2.3755	0.2842	1.2402	0.3505	-1.1353	0.3425	-1.0876	0.9013	-2.0752	0.5360	-0.9876	0.8006
---	---	1632454_at	0.0462	0.8389	0.3172	0.1565	0.1914	0.2613	-0.0222	0.9714	-0.0337	0.8715	-0.0114	0.9545	0.0609	0.9168	0.0459	0.8740	-0.0150	0.9595
CG8680	CG8680	1632455_at	-0.0878	0.7150	0.1527	0.4522	0.1890	0.2633	-0.1255	0.7225	-0.7479	0.0013	-0.6224	0.0017	-0.2133	0.8192	-0.4053	0.2722	-0.1920	0.6399
---	---	1632456_s_at	0.3525	0.1652	-0.3283	0.1958	-0.0080	0.9830	0.0997	0.8902	0.2954	0.2209	0.1957	0.3827	-0.1053	0.9059	-0.1837	0.5880	-0.0784	0.8488
mam	mastermind	1632457_s_at	-0.8610	0.2368	-1.1980	0.1015	-2.3621	0.0000	-0.3334	0.4085	0.8805	0.0028	1.2139	0.0005	0.6223	0.8122	0.3042	0.8343	-0.3181	0.8138
Incenp	Inner centromere	1632458_at	0.0356	0.9548	-0.2474	0.7483	-1.0767	0.0713	-0.6123	0.2989	0.5629	0.1035	1.1751	0.0032	-0.3001	0.9405	-0.0355	0.9897	0.2646	0.8745
CG15166	CG15166	1632459_at	0.1192	0.4264	0.0603	0.7345	0.3304	0.1009	0.0190	0.9777	0.0554	0.7962	0.0364	0.8585	-0.0016	0.9994	0.0442	0.9347	0.0458	0.9211
---	---	1632460_at	0.0492	0.7911	0.0553	0.6302	0.3051	0.1556	0.1850	0.5271	0.1408	0.3680	-0.0442	0.7975	-0.0366	0.9721	0.1571	0.5750	0.1938	0.4805
CG31233	CG31233	1632461_at	-0.0946	0.9283	-0.0022	0.9913	0.1207	0.4491	0.1235	0.9507	-0.4191	0.4693	-0.5426	0.2785	-0.0564	0.9884	-0.4728	0.5923	-0.4164	0.6414
CG2185	CG2185	1632462_at	-0.7318	0.0412	0.2445	0.0688	0.4441	0.0147	-0.0962	0.8539	-0.5388	0.0115	-0.4426	0.0167	-0.2531	0.7708	0.3882	0.2814	0.6413	0.1190
---	---	1632463_at	0.2138	0.3178	-0.0228	0.8732	-0.1582	0.3491	-0.0968	0.8791	0.1058	0.6679	0.2027	0.3032	0.0512	0.9421	0.0002	0.9999	-0.0510	0.8577
Smg6	Smg6	1632464_at	0.7600	0.0053	0.2000	0.3705	0.3218	0.1519	0.0277	0.9696	0.3189	0.0888	0.2912	0.0826	-0.0488	0.9737	-0.1335	0.7598	-0.0847	0.8596
CG6424	CG6424	1632465_s_at	-1.2525	0.0037	-1.4437	0.0146	-1.9417	0.0001	-0.1424	0.8383	0.0553	0.8732	0.1978	0.4190	0.3138	0.7523	-0.1124	0.8457	-0.4262	0.3271
CG9243	CG9243	1632466_at	-0.1856	0.2956	0.2312	0.5104	0.0964	0.6935	-0.1418	0.8281	-0.1874	0.4756	-0.0456	0.8783	-0.0163	0.9933	0.0887	0.8929	0.1051	0.8570
---	---	1632467_at	-0.0096	0.9558	-0.0169	0.9093	0.0814	0.6122	0.0680	0.9048	0.0306	0.9045	-0.0374	0.8621	0.1086	0.7997	0.1335	0.4565	0.0248	0.9196
---	---	1632468_at	0.1713	0.4012	0.2402	0.2583	0.1039	0.6563	-0.0918	0.8830	-0.0909	0.7117	0.0009	0.9974	0.0678	0.9457	0.0071	0.9924	-0.0607	0.8834
---	---	1632469_at	-0.0013	0.9958	-0.2308	0.0668	-0.0093	0.9598	0.1046	0.8251	0.1855	0.3167	0.0809	0.6716	-0.0706	0.9238	-0.1774	0.4794	-0.1067	0.6973
SRm160	SRm160	1632470_at	-0.3967	0.5813	0.2729	0.8118	0.4038	0.2036	0.2661	0.7293	0.0764	0.8663	-0.1897	0.5801	0.2293	0.9588	0.6654	0.6132	0.4361	0.7580
CG12155	CG12155	1632471_at	0.1464	0.6328	0.3266	0.1499	0.1782	0.3286	0.1101	0.8449	0.0431	0.8758	-0.0670	0.7698	0.0895	0.9309	0.0792	0.8678	-0.0102	0.9849
CG4699 /// DyakCG4699	CG4699	1632472_s_at	-0.0967	0.8601	-0.3406	0.3529	-0.2965	0.1058	0.0901	0.8578	0.3484	0.0577	0.2583	0.1082	-0.0365	0.9898	-0.0154	0.9917	0.0211	0.9843
---	---	1632473_at	0.0321	0.8574	0.1433	0.3537	0.1253	0.5904	0.0548	0.9042	0.0375	0.8449	-0.0174	0.9240	-0.0263	0.9816	0.1451	0.6124	0.1714	0.5444
CG32755	CG32755	1632474_at	-0.8243	0.0217	-0.3814	0.2345	-0.7690	0.0219	0.1948	0.8776	0.2436	0.6105	0.0487	0.9268	-0.0914	0.9238	-0.0568	0.9075	0.0346	0.9382
msn	Mishappen	1632475_at	-1.3508	0.0650	-0.3355	0.7051	-0.6346	0.0397	-0.0527	0.9538	-0.3000	0.2269	-0.2473	0.2701	0.1552	0.9774	0.5183	0.7282	0.3631	0.8221
---	---	1632476_at	0.1251	0.4978	0.1029	0.4455	0.2343	0.2516	-0.0388	0.9538	0.0621	0.7795	0.1009	0.5808	-0.0096	0.9948	0.1143	0.7842	0.1238	0.7493
CG31708	CG31708	1632477_s_at	0.1390	0.5145	-0.1655	0.1830	0.0124	0.9658	0.0223	0.9722	0.2162	0.1743	0.1939	0.1743	-0.2734	0.7305	-0.0914	0.8427	0.1820	0.6189
Pkc53E	protein kinase C	1632478_a_at	-1.0757	0.0648	-0.3022	0.4851	-2.5013	0.0008	-1.2138	0.1403	-1.8405	0.0037	-0.6268	0.1584	0.6040	0.7506	-1.1492	0.1498	-1.7533	0.0702
Spase12	Spase 12-subunit	1632479_at	0.7634	0.0094	0.9777	0.0108	1.1522	0.0003	0.0679	0.8716	0.2419	0.0985	0.1740	0.1824	-0.1494	0.8494	0.3683	0.2430	0.5177	0.1409
---	---	1632480_at	0.0533	0.8054	0.0492	0.6561	0.0889	0.5868	-0.1390	0.7929	-0.1736	0.4366	-0.0347	0.8930	-0.1078	0.8480	-0.0161	0.9664	0.0916	0.7341
CG8602	CG8602	1632481_at	-0.0283	0.8973	0.1974	0.2068	0.1558	0.4145	-0.1615	0.6701	-0.3433	0.0605	-0.1818	0.2623	-0.2552	0.7220	-0.2101	0.5074	0.0452	0.9158
CG5554	CG5554	1632482_at	-0.2197	0.3562	0.1888	0.4833	0.6434	0.0043	0.3569	0.2299	-0.3514	0.0516	-0.7083	0.0016	-0.0004	0.9999	0.1023	0.8450	0.1027	0.8334
sop	Ribosome RS2	1632483_at	0.3479	0.0310	1.1266	0.0039	1.2175	0.0006	0.0617	0.9186	-0.5404	0.0108	-0.6020	0.0040	-0.0251	0.9742	0.0481	0.8571	0.0732	0.7437
CG2218	CG2218	1632484_at	-0.0561	0.8693	0.0047	0.9937	-0.0219	0.8932	-0.0506	0.9380	0.0604	0.7957	0.1110	0.5528	0.0322	0.9896	0.1868	0.7642	0.1546	0.8073
Nak	Numb-associated	1632485_a_at	-1.1212	0.0034	-0.0461	0.8210	-0.0198	0.9134	-0.1685	0.6010	-1.1444	0.0003	-0.9759	0.0003	-0.2885	0.7152	0.0608	0.9046	0.3494	0.3023
---	---	1632486_at	-0.0223	0.9125	-0.1582	0.3918	-0.1552	0.3961	-0.0849	0.8676	0.0891	0.6622	0.1740	0.2855	-0.0235	0.9816	-0.0072	0.9883	0.0163	0.9621
CG14721	CG14721	1632487_at	0.2468	0.1261	0.5982	0.0758	0.7904	0.0007	0.0312	0.9599	-0.1413	0.3972	-0.1725	0.2363	-0.1117	0.8609	0.2224	0.3768	0.3341	0.2117
---	---	1632488_a_at	-0.1715	0.2236	0.1689	0.4008	0.5743	0.0042	-0.0885	0.8462	-0.2340	0.1604	-0.1454	0.3435	-0.5562	0.2884	0.0017	0.9989	0.5579	0.0710
CG14881	CG14881	1632489_a_at	-0.0540	0.8714	-0.0480	0.7848	-0.0443	0.8011	-0.0766	0.8816	0.0568	0.7912	0.1334	0.4161	-0.2141	0.8362	-0.0306	0.9650	0.1834	0.7013
CG8008	CG8008	1632490_at	-1.9078	0.0073	-3.0177	0.0047	-3.1482	0.0000	-0.2608	0.5680	-0.7665	0.0062	-0.5057	0.0236	-0.0351	0.9914	-1.6364	0.0405	-1.6012	0.0585
CG32985	CG32985	1632491_at	0.2862	0.4199	-0.1030	0.4621	0.0220	0.9053	0.0194	0.9745	0.0923	0.5624	0.0729	0.6283	-0.0762	0.9555	-0.2171	0.6093	-0.1409	0.7582
CG5062	CG5062	1632492_at	0.0434	0.8309	0.2082	0.3823	0.2818	0.0778	-0.1610	0.6650	-0.2144	0.2265	-0.0534	0.7878	-0.1024	0.8513	-0.0187	0.9601	0.0836	0.7492

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632493_at	0.1295	0.5644	0.0510	0.6426	0.1201	0.7300	-0.1648	0.7081	0.0592	0.8114	0.2240	0.2125	0.0389	0.9781	0.1591	0.6711	0.1202	0.7582
Grip163	Grip163	1632494_at	-0.5841	0.0769	-0.2050	0.3758	-0.3175	0.1673	0.0528	0.9755	0.1349	0.7940	0.0821	0.8680	0.0259	0.9862	0.1701	0.6454	0.1442	0.7060
CG31867	CG31867	1632495_at	0.2407	0.2467	0.0229	0.8807	0.2180	0.3755	-0.1426	0.8233	-0.2583	0.3024	-0.1157	0.6519	-0.0847	0.9365	-0.2382	0.4881	-0.1535	0.6776
rad201	radiation-resistant	1632496_at	-0.3902	0.1082	-0.1193	0.4567	0.2546	0.2658	0.0311	0.9777	-0.2080	0.4757	-0.2391	0.3507	-0.2535	0.6898	0.0859	0.7922	0.3394	0.2107
Gug	Atrophin	1632497_a_at	-0.7713	0.4122	-0.9209	0.1336	-1.8335	0.0002	-0.4482	0.2471	0.3754	0.1017	0.8236	0.0025	0.5094	0.8769	0.3440	0.8431	-0.1653	0.9255
CG15629	CG15629	1632498_at	-0.0397	0.9388	-0.4808	0.0288	-0.2079	0.5065	0.2168	0.7432	-0.0671	0.8607	-0.2840	0.2804	-0.0909	0.9515	-0.4153	0.3416	-0.3244	0.4848
CG14796	CG14796	1632499_at	0.0421	0.9525	0.2448	0.5052	0.3823	0.3373	-0.4832	0.2114	-0.6466	0.0131	-0.1635	0.4577	-0.5195	0.8270	-0.1703	0.9105	0.3492	0.7592
CG10226	CG10226	1632500_at	-0.9442	0.0211	0.0868	0.4987	-0.5267	0.1175	-0.8790	0.1564	-1.3288	0.0045	-0.4498	0.1794	-0.1265	0.9405	-0.5781	0.2507	-0.4516	0.3960
---	---	1632501_at	-0.0219	0.9181	-0.0526	0.7324	0.3026	0.0826	-0.0332	0.9689	-0.2169	0.3347	-0.1837	0.3686	-0.0459	0.9598	-0.1810	0.4967	-0.1350	0.6294
---	---	1632502_s_at	0.0091	0.9606	0.1084	0.4435	0.2311	0.3814	0.0075	0.9921	-0.0302	0.8840	-0.0377	0.8299	-0.1400	0.8594	0.0117	0.9848	0.1517	0.6607
CG3792	CG3792	1632503_at	0.8367	0.0184	0.0496	0.9362	0.2695	0.1624	-0.0197	0.9854	1.0336	0.0019	1.0532	0.0011	-0.1992	0.8971	0.2417	0.7093	0.4408	0.4569
CG17637	CG17637	1632504_at	0.1859	0.2084	-0.1355	0.1849	0.0293	0.8773	-0.0491	0.9300	0.1113	0.5222	0.1604	0.2779	-0.0745	0.8692	-0.0913	0.6406	-0.0167	0.9471
---	---	1632505_at	0.0638	0.7243	0.1089	0.3938	0.2879	0.0785	0.1611	0.7208	0.0640	0.7980	-0.0971	0.6429	-0.0019	0.9991	-0.0306	0.9430	-0.0286	0.9387
---	---	1632506_at	-0.1285	0.5936	0.2306	0.2774	0.3889	0.2199	-0.0278	0.9715	-0.1765	0.3823	-0.1488	0.4214	-0.0780	0.9632	0.1592	0.7788	0.2372	0.6328
sick	sickie	1632507_at	-0.0970	0.7408	0.1355	0.3170	0.0773	0.8497	0.0304	0.9833	-0.1573	0.6775	-0.1877	0.5684	0.0684	0.9611	-0.0019	0.9991	-0.0704	0.8962
caps	capricious	1632508_s_at	-1.1424	0.1021	0.6185	0.3735	0.4554	0.0560	-0.1962	0.8606	-1.4456	0.0041	-1.2494	0.0047	0.1578	0.9545	0.5169	0.5425	0.3591	0.6896
CG4434	CG4434	1632509_at	0.0434	0.7869	0.0607	0.7009	-0.0711	0.6991	-0.0683	0.9238	0.0204	0.9495	0.0887	0.6940	0.0490	0.9547	0.0245	0.9519	-0.0245	0.9444
CG7332	CG7332	1632510_at	-0.6398	0.0256	-0.3053	0.3955	-0.2605	0.2643	0.1189	0.8967	-0.2231	0.4801	-0.3420	0.2051	0.0364	0.9831	0.1699	0.7063	0.1335	0.7764
CG12862	CG12862	1632511_at	0.1129	0.6266	0.0619	0.5718	0.1106	0.7225	-0.0013	0.9994	-0.1683	0.6300	-0.1670	0.5950	0.1100	0.9095	0.0554	0.9175	-0.0546	0.9092
CG11820	CG11820	1632512_at	-0.2992	0.0990	0.5275	0.1040	0.5794	0.0277	0.1026	0.8773	-0.6604	0.0110	-0.7630	0.0034	0.1184	0.8882	0.2718	0.3810	0.1535	0.6533
Neos	Neosin	1632513_at	0.0937	0.5380	-0.0248	0.9069	-0.0291	0.8850	0.0451	0.9413	0.3419	0.0556	0.2969	0.0617	-0.0064	0.9952	0.1344	0.6427	0.1408	0.6254
CG11905	CG11905	1632514_s_at	-0.0502	0.8112	0.1473	0.4467	0.0537	0.7717	-0.1708	0.5311	-0.2787	0.0548	-0.1078	0.4179	0.0167	0.9816	-0.0381	0.8767	-0.0548	0.7909
CG18619	CG18619	1632515_a_at	0.0624	0.8892	0.5920	0.0962	0.6824	0.0018	-0.3168	0.6560	-1.3260	0.0030	-1.0092	0.0061	-0.3694	0.6557	-0.8050	0.0391	-0.4356	0.2136
Rbsn	dRabenosyn	1632516_at	-0.4420	0.0349	0.1937	0.3078	0.3234	0.2589	0.1674	0.6221	-0.0452	0.8342	-0.2126	0.1581	0.0712	0.9611	0.6343	0.1238	0.5631	0.1911
CG7966	CG7966	1632517_at	2.2691	0.0019	1.6461	0.0021	2.7794	0.0000	0.5743	0.2977	0.2594	0.4403	-0.3149	0.2825	-0.3619	0.7423	-0.4163	0.3606	-0.0544	0.9338
CG5470	CG5470	1632518_at	0.2424	0.1080	0.0272	0.8688	0.0999	0.5300	0.0477	0.9375	0.0421	0.8541	-0.0056	0.9799	-0.1244	0.8145	-0.1229	0.5994	0.0015	0.9974
CG8366	CG8366	1632519_at	0.2066	0.1980	-0.1234	0.6400	-0.3043	0.0681	0.1523	0.6327	0.6483	0.0022	0.4960	0.0043	0.1895	0.8326	0.2202	0.5776	0.0307	0.9533
---	---	1632520_at	0.1799	0.3310	0.0150	0.9161	0.2514	0.1266	0.1937	0.6035	0.1834	0.3360	-0.0102	0.9658	-0.0355	0.9589	-0.0304	0.9212	0.0050	0.9874
CG4542	CG4542	1632521_at	0.2090	0.3195	0.5986	0.0168	1.1629	0.0009	0.0639	0.9436	-0.4732	0.0671	-0.5370	0.0267	-0.4660	0.3362	0.0365	0.9281	0.5025	0.0859
---	---	1632522_at	0.3629	0.1823	0.2681	0.3495	0.2203	0.3261	0.1281	0.8844	0.2490	0.4228	0.1208	0.7028	-0.0383	0.9829	-0.0595	0.9265	-0.0212	0.9732
---	---	1632523_at	0.1228	0.4771	0.1275	0.4298	0.1766	0.4599	0.0954	0.8544	0.0590	0.7977	-0.0364	0.8682	0.0463	0.9657	0.0109	0.9842	-0.0354	0.9310
Wnt4	Wnt oncogene an	1632524_at	-2.8320	0.0004	-3.4327	0.0011	-3.3367	0.0009	0.3846	0.7803	0.7389	0.1948	0.3544	0.5249	0.0657	0.9340	-0.1685	0.5191	-0.2342	0.3671
CG9925	CG9925	1632525_at	0.9952	0.3425	-1.6577	0.1357	-1.7267	0.0641	-0.3541	0.3176	2.5380	0.0000	2.8921	0.0000	-0.5094	0.9416	-0.2682	0.9388	0.2413	0.9360
Nc73EF	Neural conserved	1632526_s_at	-1.0011	0.0466	-0.6219	0.1375	-0.8497	0.0073	-0.1609	0.6869	-0.5305	0.0115	-0.3696	0.0332	-0.2021	0.9148	-0.3851	0.5827	-0.1829	0.8255
CG13705	CG13705	1632527_at	0.0921	0.6034	0.2121	0.3730	0.1604	0.2761	0.0709	0.9314	-0.1838	0.4738	-0.2547	0.2476	-0.0680	0.9324	-0.0707	0.8382	-0.0026	0.9953
---	---	1632528_at	0.1513	0.4979	-0.2769	0.2331	-0.4503	0.0836	-0.3577	0.2760	0.3465	0.0760	0.7042	0.0024	-0.0874	0.9212	-0.2274	0.4418	-0.1400	0.6630
tinc	tincar	1632529_at	0.2816	0.5856	-1.0871	0.1670	-0.6658	0.2078	-0.0233	0.9937	1.0593	0.0803	1.0826	0.0495	-0.2810	0.9135	-0.2409	0.8444	0.0401	0.9777
CG8776	CG8776	1632530_s_at	-2.0148	0.1163	-2.8622	0.0025	-2.8622	0.0004	-0.2179	0.8248	0.1081	0.8202	0.3260	0.3517	-0.1058	0.9894	-0.7423	0.6937	-0.6365	0.7422
CG5770 /// DyakCG5770	CG5770	1632531_at	0.1636	0.2737	0.0389	0.7445	-0.2724	0.2523	0.0127	0.9915	0.1899	0.4716	0.1771	0.4580	0.1979	0.7485	0.0407	0.9214	-0.1572	0.5756
wal	Electron transfer f	1632532_s_at	0.3185	0.1898	0.1295	0.4864	0.0960	0.6711	-0.1028	0.8640	-0.1350	0.5567	-0.0322	0.9001	-0.0868	0.9365	-0.3758	0.2582	-0.2890	0.4097
Timp	tissue inhibitor of	1632533_at	-1.1718	0.0012	-0.8684	0.1252	-1.5274	0.0001	-0.1965	0.5680	0.2475	0.1581	0.4440	0.0125	0.4966	0.6955	0.5749	0.2591	0.0783	0.9152
CG7448	CG7448	1632534_at	-0.0187	0.9432	0.0134	0.8979	0.1436	0.5250	-0.0742	0.8578	-0.0516	0.7724	0.0226	0.8991	-0.0548	0.9653	0.0542	0.9177	0.1090	0.7873
CG13074	CG13074	1632535_at	0.0974	0.7319	0.0729	0.6223	0.0160	0.9595	-0.1194	0.8940	0.1926	0.5479	0.3120	0.2435	-0.0530	0.9677	-0.0062	0.9935	0.0468	0.9220
CG4793	CG4793	1632536_at	0.2767	0.1519	0.0000	1.0000	-0.1424	0.4960	0.0430	0.9464	0.2836	0.1149	0.2406	0.1344	0.0853	0.9051	-0.0814	0.8053	-0.1667	0.5391
CdGAPr	CdGAPr	1632537_at	-0.1746	0.4981	0.0241	0.9187	-0.3671	0.1040	-0.2182	0.6218	-0.0080	0.9809	0.2102	0.2915	0.2184	0.8270	-0.0395	0.9528	-0.2579	0.5581
---	---	1632538_at	-0.0276	0.8836	0.1111	0.5277	0.1608	0.3273	0.0066	0.9937	-0.1301	0.4285	-0.1367	0.3484	0.0148	0.9914	0.1287	0.6957	0.1139	0.7346
GstE4	Glutathione S tran	1632539_at	-0.0861	0.5916	0.1702	0.4870	0.0450	0.8583	0.0016	0.9988	-0.1741	0.5527	-0.1757	0.5023	0.0413	0.9724	0.0509	0.9125	0.0095	0.9847
iotaTry	trypsin iota	1632540_at	-0.2448	0.7995	0.0119	0.9575	-0.1798	0.3461	0.2291	0.8942	-0.4042	0.5054	-0.6334	0.2179	0.2566	0.9142	-0.1435	0.9104	-0.4001	0.6567
Blimp-1	Blimp-1	1632541_at	0.2795	0.4976	-0.7871	0.1672	-0.6911	0.0902	0.1105	0.8074	0.6843	0.0033	0.5738	0.0043	0.1569	0.9672	-0.2876	0.8284	-0.4445	0.6883
CG13932	CG13932	1632542_at	0.1023	0.6770	0.1172	0.6227	0.2180	0.2560	0.0536	0.9599	0.0034	0.9934	-0.0502	0.8797	-0.0582	0.9309	0.0084	0.9841	0.0666	0.8003
CG9003	CG9003	1632543_at	-0.8653	0.0049	0.5757	0.1592	0.6816	0.0053	-0.2491	0.7028	-1.5359	0.0009	-1.2867	0.0012	-0.2657	0.7595	-0.0142	0.9841	0.2515	0.5050
---	---	1632544_at	0.1078	0.5034	0.0089	0.9639	0.0816	0.6216	0.0222	0.9734	0.0411	0.8471	0.0189	0.9252	0.0409	0.9460	0.0293	0.9175	-0.0116	0.9659
CG5522	CG5522	1632545_s_at	-0.6766	0.0157	-0.3627	0.5731	-0.8362	0.0236	-0.5267	0.1467	-0.1622	0.4799	0.3645	0.0665	-0.0625	0.9848	0.1883	0.8551	0.2508	0.7774
---	---	1632546_at	0.2908	0.3721	0.1394	0.3887	0.0147	0.9525	-0.2266	0.6634	-0.1181	0.6776	0.1085	0.6775	-0.0162	0.9914	-0.0027	0.9982	0.0135	0.9819
CG40045	CG40045	1632547_at	-0.0484	0.8513	0.0337	0.7748	-0.0790	0.6458	-0.1182	0.7753	-0.0506	0.8135	0.0676	0.7134	-0.0392	0.9759	0.0239	0.9633	0.0631	0.8834
Mes2	Mes2	1632548_a_at	1.2716	0.0506	1.5052	0.0179	1.1796	0.0008	-0.0638	0.9651	0.4682	0.2293	0.5320	0.1254	0.2910	0.8541	0.5960	0.3471	0.3050	0.6668
---	---	1632549_at	0.0790	0.6784	-0.0116	0.9560	0.2313	0.2062	0.1244	0.8512	-0.0384	0.9089	-0.1628	0.4797	-0.1224	0.8202	-0.0146	0.9684	0.1078	0.6557
unc-5	unc-5-like	1632550_at	-2.5453	0.0009	-2.1758	0.0225	-2.6743	0.0004	-0.6553	0.2870	-0.8294	0.0294	-0.1741	0.6386	-0.2833	0.8745	-0.5043	0.4852	-0.2210	0.7954
CG33969	CG33969	1632551_at	0.3116	0.2283	0.0699	0.8280	0.2944	0.0722	0.1236	0.8584	0.0967	0.7425	-0.0269	0.9297	-0.1277	0.8942	-0.2102	0.5823	-0.0825	0.8618
Oseg3	Oseg3	1632552_at	-0.0245	0.8855	0.0593	0.6067	0.1624	0.3742	0.0260	0.9675	-0.1966	0.2430	-0.2226	0.1370	0.0662	0.9201	0.0083	0.9844	-0.0578	0.8385
---	---	1632553_at	0.0782	0.7594	0.0297	0.8816	0.0516	0.7704	-0.1451	0.7023	-0.1236	0.5014	0.0215	0.9201	-0.0671	0.9514	-0.0137	0.9828	0.0535	0.9061
CG9839	CG9839	1632554_at	-0.6173	0.0231	-0.5754	0.0859	-1.0083	0.0007	-0.0573	0.9300	-0.2819	0.1394	-0.2246	0.1889	0.3207	0.7215	0.0543	0.9277	-0.2664	0.4978
---	---	1632555_s_at	-0.1094	0.7445	-0.0233	0.9165	0.0523	0.7709	0.1230	0.8299	-0.0851	0.7423	-0.2081	0.2951	0.0120	0.9940	0.1075	0.8255	0.0955	0.8402
CG2310	CG2310	1632556_s_at	-0.3664	0.1049	0.2872	0.3775	0.7428	0.0357	0.4364	0.1913	-0.0772	0.7478	-0.5136	0.0121	0.0410	0.9893	0.5953	0.3393	0.5544	0.3939
Rab30	Rab30	1632557_s_at	0.1906	0.5401	0.3722	0.0389	0.3605	0.1261	0.0860	0.8885	0.1407	0.5214	0.0546	0.8144	0.1008	0.9467	0.3337	0.4710	0.2329	0.6351
---	---	1632558_at	0.7405	0.0251	1.1959	0.0443	1.6261	0.0003	0.3294	0.5515	-0.1765	0.5682	-0.5060	0.0526	-0.0842	0.9657	0.2257	0.7036	0.3100	0.5792
rod	rough deal	1632559_at	-0.4060	0.0400	0.0364	0.8777	-0.3642	0.1025	-0.3410	0.4861	-0.0923	0.7689	0.2487	0.2940	-0.1367	0.9077	0.1014	0.8625	0.2381	0.5980
eys	eyes shut	1632560_at	0.2728	0.1095	0.3220	0.1477	0.3417	0.1462	0.0750	0.9032	-0.2004	0.3221	-0.2754	0.1215	-0.0390	0.9675	-0.0184	0.9647	0.0206	0.9536
CG30437	CG30437	1632561_at	0.0445	0.8154	-0.2977	0.0710	-0.4360	0.0250	0.1193	0.8028	0.5307	0.0132	0.4114	0.0242	0.1895	0.8236	0.0273	0.9628	-0.1623	0.6762

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632562_at	-0.1857	0.2912	0.1027	0.4737	-0.1263	0.4230	-0.1515	0.7368	-0.1916	0.3459	-0.0401	0.8630	0.0902	0.9092	0.1151	0.7200	0.0250	0.9492
---	---	1632563_at	0.3369	0.1208	0.1886	0.3555	-0.1385	0.4015	-0.3287	0.2871	-0.2543	0.1596	0.0745	0.6964	0.0365	0.9816	-0.3056	0.4021	-0.3421	0.3668
---	---	1632564_at	-0.0161	0.9454	-0.2293	0.4710	0.0434	0.8782	0.0943	0.9314	0.2158	0.5336	0.1215	0.7258	-0.0626	0.9348	-0.0022	0.9977	0.0604	0.8480
Bin1	Sin3A-associated	1632565_at	0.5036	0.0207	0.4886	0.1718	0.2390	0.1501	0.0546	0.9260	0.4125	0.0277	0.3579	0.0314	0.2604	0.7230	0.3668	0.2288	0.1064	0.7749
CG11318	CG11318	1632566_at	-0.1930	0.6687	0.2057	0.4551	0.4082	0.1733	0.1596	0.9048	-0.1760	0.7280	-0.3356	0.4061	-0.0101	0.9964	0.0771	0.9318	0.0872	0.9105
CG31342	CG31342	1632567_at	-0.2244	0.3652	0.0364	0.8808	-0.0207	0.9091	-0.1189	0.7803	-0.3562	0.0491	-0.2373	0.1315	0.0070	0.9964	-0.0059	0.9941	-0.0129	0.9837
CG3223	CG3223	1632568_at	-0.0374	0.8743	0.1940	0.0920	0.4746	0.0368	-0.0435	0.9412	-0.1804	0.2888	-0.1369	0.3812	-0.1767	0.8202	0.0700	0.8797	0.2467	0.4597
---	---	1632569_at	-0.1697	0.4622	0.0115	0.9479	-0.0881	0.6478	-0.0104	0.9932	-0.0231	0.9489	-0.0127	0.9669	0.0883	0.8692	0.1311	0.5613	0.0428	0.8848
CG2052	CG2052	1632570_at	0.1588	0.5441	0.0681	0.5152	0.2507	0.2252	-0.0412	0.9436	0.0930	0.6075	0.1343	0.3788	0.0513	0.9682	0.0086	0.9911	-0.0427	0.9278
---	---	1632571_at	-0.1849	0.4139	-0.0617	0.6659	0.1184	0.4280	-0.0308	0.9558	0.0391	0.8395	0.0700	0.6518	-0.2962	0.6955	0.0675	0.8813	0.3637	0.2638
---	---	1632572_at	0.1234	0.5525	0.2864	0.2968	0.1453	0.5993	-0.1015	0.8500	-0.1385	0.5018	-0.0370	0.8725	0.1955	0.8157	0.2226	0.5374	0.0271	0.9565
Mitf	Mitf	1632573_at	0.3371	0.1412	-0.3691	0.5074	-0.3916	0.1020	-0.1161	0.8578	0.3557	0.1264	0.4718	0.0314	0.1802	0.9155	-0.1925	0.7972	-0.3727	0.5515
CG6574	CG6574	1632574_at	-1.3861	0.0031	-0.8200	0.1078	-0.3411	0.1117	0.0590	0.9314	-0.6401	0.0076	-0.6991	0.0030	-0.4929	0.6496	0.0186	0.9822	0.5116	0.2606
---	---	1632575_at	0.3397	0.5080	0.2709	0.2552	0.4756	0.0106	0.2012	0.8006	-0.0729	0.8628	-0.2742	0.3518	-0.1661	0.8963	-0.2421	0.6387	-0.0760	0.9080
CG8142	CG8142	1632576_at	0.0549	0.8411	-0.5075	0.1774	0.0104	0.9807	-0.2334	0.6663	0.2674	0.3048	0.5008	0.0368	-0.7202	0.4415	-0.2665	0.6093	0.4537	0.3660
Su(var)3-9	Enhancer of varie	1632577_a_at	-0.0979	0.6171	1.3100	0.0179	1.3816	0.0000	0.0080	0.9912	-0.7667	0.0011	-0.7747	0.0006	-0.1180	0.9056	0.5902	0.1054	0.7082	0.0836
CG33251	CG33251	1632578_at	-0.0036	0.9897	-0.1367	0.5617	-0.1427	0.4787	-0.0164	0.9834	0.1427	0.4457	0.1591	0.3340	-0.0458	0.9467	-0.0049	0.9924	0.0409	0.8874
---	---	1632579_at	0.0687	0.6892	0.0917	0.5382	0.0816	0.6810	0.0897	0.8589	0.0833	0.6896	-0.0064	0.9776	-0.0795	0.9514	-0.0612	0.9178	0.0183	0.9762
CG12765	CG12765	1632580_at	-0.0728	0.7122	-0.3838	0.1388	-0.2470	0.1696	0.0861	0.8817	0.3433	0.0838	0.2572	0.1448	-0.1520	0.8692	0.0171	0.9796	0.1692	0.6742
CG1663	CG1663	1632581_at	-0.0901	0.6808	0.0221	0.9575	0.1076	0.5471	-0.0685	0.9196	0.0023	0.9941	0.0708	0.7554	-0.1347	0.9151	0.0941	0.8824	0.2288	0.6311
CG9270	CG9270	1632582_at	-0.2861	0.1561	0.0383	0.7714	-0.2139	0.4107	-0.2383	0.5451	-0.1575	0.4586	0.0808	0.7084	0.1022	0.8680	0.0313	0.9358	-0.0709	0.8153
CG13539	CG13539	1632583_at	-0.0680	0.7583	0.0201	0.8558	-0.0059	0.9788	-0.0251	0.9777	-0.1491	0.5308	-0.1239	0.5767	0.0354	0.9751	-0.0402	0.9275	-0.0756	0.8299
CG3344 /// DyakCG3344	CG3344	1632584_at	0.2393	0.4226	-0.0257	0.8012	-0.1552	0.4499	0.3026	0.4762	0.1919	0.4148	-0.1107	0.6356	0.1539	0.8472	-0.1199	0.7663	-0.2738	0.4192
---	---	1632585_at	0.3586	0.0675	0.0228	0.8218	0.0038	0.9902	-0.0179	0.9799	0.0959	0.6131	0.1138	0.4894	0.0105	0.9914	-0.0925	0.7036	-0.1030	0.6585
CG9951 /// DereCG9951 ///	CG9951 /// GA22	1632586_at	-0.5606	0.0211	-0.3588	0.1136	-0.0168	0.9474	0.2078	0.6013	0.0088	0.9777	-0.1990	0.2718	0.0423	0.9689	0.2573	0.3575	0.2150	0.4676
CG9133	CG9133	1632587_s_at	0.0617	0.7667	-0.0091	0.9632	-0.1696	0.2809	0.1172	0.7884	0.1665	0.3623	0.0493	0.8061	0.1634	0.8235	0.0083	0.9903	-0.1550	0.6366
---	---	1632588_at	-0.1077	0.5865	0.2249	0.1953	0.0998	0.6450	-0.1695	0.6854	-0.1695	0.3961	0.0000	0.9999	-0.0295	0.9787	0.1015	0.7369	0.1311	0.6388
CG7154	CG7154	1632589_at	0.0796	0.6626	0.1882	0.5693	0.2314	0.1494	0.0626	0.9300	0.1123	0.6299	0.0498	0.8358	-0.1017	0.9095	0.1357	0.7067	0.2374	0.4721
CG31320	CG31320	1632590_at	1.1034	0.0059	0.5036	0.2140	0.2490	0.2756	0.0853	0.9228	0.8098	0.0086	0.7245	0.0085	0.1682	0.8882	0.4247	0.3342	0.2565	0.5972
Phk-3	Pherokine 3	1632591_at	-0.3811	0.5969	-2.4199	0.0025	-2.6312	0.0000	-0.6715	0.3685	1.6707	0.0028	2.3423	0.0004	-0.5025	0.7743	-0.4240	0.6011	0.0786	0.9413
CG14632	CG14632	1632592_a_at	-0.1932	0.4481	0.1090	0.5557	0.1371	0.3710	-0.0011	0.9988	-0.0941	0.6896	-0.0930	0.6621	-0.0577	0.9168	0.0270	0.9289	0.0847	0.6892
CG32147	CG32147	1632593_at	-1.3641	0.0038	-1.0335	0.0270	-1.3329	0.0003	-0.0513	0.9223	-0.4121	0.0176	-0.3608	0.0191	0.1618	0.8909	-0.0329	0.9646	-0.1947	0.6939
CG1603	CG1603	1632594_at	0.1686	0.4144	0.3832	0.2915	0.3964	0.1272	0.0859	0.9289	0.1757	0.5648	0.0899	0.7737	-0.0613	0.9665	0.2805	0.4794	0.3418	0.3893
CG32237	CG32237	1632595_at	0.0112	0.9751	0.1535	0.5524	0.1220	0.5910	0.0809	0.9436	-0.1461	0.6963	-0.2270	0.4615	0.1119	0.8270	0.1174	0.6093	0.0054	0.9874
mRpL44	mitochondrial ribo	1632596_at	-0.1170	0.6775	0.1110	0.5829	-0.1278	0.4494	0.0636	0.8979	0.1812	0.2679	0.1176	0.4427	0.1726	0.8608	0.2753	0.4972	0.1028	0.8374
---	---	1632597_at	-0.0197	0.9220	-0.0115	0.9508	-0.0076	0.9773	-0.0272	0.9726	0.0842	0.7101	0.1114	0.5663	-0.1687	0.8533	-0.0981	0.8478	0.0705	0.8915
---	---	1632598_at	0.0606	0.7424	0.0030	0.9810	0.1023	0.6112	0.0252	0.9764	-0.0752	0.7657	-0.1003	0.6423	0.1201	0.8122	0.0718	0.7799	-0.0483	0.8602
---	---	1632599_at	0.0649	0.7856	0.0473	0.6579	0.1433	0.3647	-0.0874	0.8096	0.0422	0.8131	0.1295	0.3251	-0.2034	0.7780	-0.0515	0.9157	0.1519	0.6550
---	---	1632600_at	-0.0341	0.8608	0.2235	0.4262	0.2852	0.2257	0.2847	0.4356	0.0439	0.8684	-0.2407	0.1780	0.1595	0.8814	0.2122	0.6296	0.0527	0.9252
---	---	1632601_at	0.1944	0.4062	0.1103	0.3861	0.1888	0.2524	0.0285	0.9641	-0.0963	0.5997	-0.1247	0.4280	-0.0226	0.9816	-0.0394	0.9111	-0.0168	0.9597
AGO1	Argonaute1	1632602_s_at	0.3016	0.4157	0.8422	0.0890	0.7122	0.0155	-0.1076	0.8899	-0.0140	0.9725	0.0936	0.7384	-0.0664	0.9816	0.5363	0.4094	0.6027	0.3708
---	---	1632603_at	0.0602	0.7844	-0.0511	0.6286	0.0202	0.9149	0.0205	0.9773	0.0931	0.6343	0.0726	0.6937	-0.1812	0.7215	-0.1788	0.3935	0.0024	0.9948
---	---	1632604_at	0.0497	0.7642	0.0000	1.0000	0.1422	0.3688	0.1345	0.6908	0.1600	0.3114	0.0256	0.8915	0.0487	0.9296	0.0484	0.8396	-0.0003	0.9992
gammaSnap	gamma-soluble N	1632605_at	-0.7012	0.0075	-0.8032	0.0617	-0.3004	0.1307	0.2680	0.3053	0.3609	0.0267	0.0930	0.5353	-0.3366	0.7215	0.1774	0.6868	0.5140	0.2107
CG33169	CG33169	1632606_a_at	1.2577	0.0010	2.1034	0.0048	1.6624	0.0003	-0.0500	0.9345	-0.1836	0.3118	-0.1336	0.4285	0.3623	0.7241	0.6276	0.1495	0.2653	0.5755
CG33189	CG33189	1632607_at	0.2249	0.4040	0.0827	0.5873	0.1861	0.2676	0.0333	0.9671	0.0292	0.9215	-0.0042	0.9874	-0.0444	0.9689	0.1534	0.6319	0.1978	0.5257
---	---	1632608_at	0.1557	0.3844	-0.0039	0.9828	0.1723	0.3828	0.0160	0.9830	0.0174	0.9456	0.0014	0.9949	-0.1491	0.7464	-0.0524	0.8428	0.0966	0.6458
---	---	1632609_s_at	0.1697	0.3874	0.1300	0.3736	0.3494	0.0434	0.0338	0.9596	-0.0758	0.7075	-0.1095	0.5189	-0.0738	0.9011	0.0363	0.9144	0.1101	0.6389
Gr98c	Gustatory recepto	1632610_at	0.0529	0.8207	0.0926	0.6298	0.2275	0.2761	0.0927	0.8794	0.0122	0.9702	-0.0806	0.7174	-0.0740	0.9467	0.0651	0.8967	0.1391	0.7134
CG15136	CG15136	1632611_at	0.0135	0.9728	0.1665	0.3796	-0.1305	0.4736	-0.0361	0.9780	-0.1035	0.8012	-0.0674	0.8626	0.1637	0.8609	-0.0061	0.9941	-0.1699	0.6848
Fmr1	Fragile X	1632612_s_at	-0.0570	0.8754	0.3487	0.1657	0.2202	0.1946	-0.0104	0.9945	0.0596	0.8918	0.0700	0.8519	0.2014	0.8472	0.6256	0.1420	0.4242	0.3439
Spn6	serpin 6	1632613_at	-0.8252	0.0418	-1.0027	0.0303	-0.1277	0.8229	0.8735	0.1462	1.4419	0.0027	0.5684	0.0816	0.0715	0.9848	1.2545	0.1382	1.1830	0.1897
---	---	1632614_at	0.1209	0.4442	0.2151	0.3539	-0.0157	0.9341	0.0083	0.9937	-0.0435	0.8727	-0.0518	0.8241	0.1645	0.7715	0.0770	0.8043	-0.0876	0.7534
CG6893	CG6893	1632615_at	-0.0062	0.9758	0.0809	0.4261	0.2618	0.1624	0.0224	0.9761	-0.1633	0.3898	-0.1857	0.2661	-0.0017	0.9992	0.0512	0.8810	0.0529	0.8672
CG18094	CG18094	1632616_at	0.0944	0.8613	0.2194	0.2941	0.1000	0.7292	0.0629	0.9620	-0.2148	0.5698	-0.2777	0.3932	-0.1620	0.7673	-0.1330	0.5880	0.0290	0.9282
---	---	1632617_at	0.1347	0.6646	0.1098	0.4964	0.1445	0.6814	0.0535	0.9422	0.0328	0.9106	-0.0207	0.9366	0.0521	0.9653	0.0271	0.9590	-0.0250	0.9568
cno	lips	1632618_a_at	-2.6178	0.0006	-0.6578	0.0287	-1.6135	0.0002	-0.9126	0.0976	-1.9790	0.0005	-1.0664	0.0037	0.0552	0.9657	-0.0906	0.8478	-0.1458	0.7087
---	---	1632619_at	0.2449	0.2007	-0.2294	0.1933	-0.0027	0.9919	0.4133	0.1390	0.4537	0.0152	0.0404	0.8346	0.0054	0.9958	-0.0320	0.9325	-0.0374	0.9086
VGlut	Vesicular glutama	1632620_at	0.1578	0.4939	0.3063	0.0356	0.1368	0.5838	-0.1279	0.8074	-0.3534	0.0916	-0.2255	0.2281	-0.0205	0.9862	-0.1214	0.6952	-0.1009	0.7506
CG12946	CG12946	1632621_at	0.5895	0.2490	0.2509	0.6431	0.8806	0.0546	0.2771	0.7121	0.0450	0.9249	-0.2322	0.4750	-0.1040	0.9741	-0.1744	0.8765	-0.0704	0.9491
nAcRalpha-18C	---	1632622_at	0.8812	0.0334	0.3586	0.4111	1.9409	0.0007	0.0725	0.9125	-0.1309	0.5613	-0.2034	0.2811	-1.3906	0.3362	-0.5329	0.4983	0.8576	0.2784
Cyt-b5-r	Cytochrome b5-re	1632623_at	2.6660	0.0035	1.5383	0.0258	2.8179	0.0001	0.9244	0.3628	0.3921	0.5247	-0.5323	0.3121	-0.3194	0.7953	-0.7308	0.1495	-0.4114	0.4404
cort	cortex	1632624_at	-0.1243	0.7764	-1.7750	0.0288	-1.5379	0.0047	0.1820	0.8836	1.8330	0.0019	1.6510	0.0018	-0.0507	0.9877	0.0885	0.9407	0.1391	0.8915
CG8960	CG8960	1632625_a_at	-0.0037	0.9905	-0.4328	0.1361	-0.2065	0.3801	0.2545	0.6075	0.2999	0.2271	0.0453	0.8776	-0.0057	0.9970	-0.2403	0.5246	-0.2346	0.5415
CG17712	CG17712	1632626_at	-0.1186	0.4699	-0.0732	0.7492	0.0574	0.8677	0.0344	0.9745	0.1961	0.4759	0.1617	0.5279	-0.0013	0.9998	0.2740	0.4996	0.2753	0.5007
CG31897	CG31897	1632627_at	0.0978	0.5130	0.0668	0.6962	0.1072	0.5421	-0.0063	0.9937	-0.0132	0.9557	-0.0069	0.9723	-0.1969	0.7475	0.0413	0.9178	0.2382	0.3708
CG15911	CG15911	1632628_at	0.2425	0.3568	0.0338	0.8634	-0.0581	0.8218	0.0099	0.9912	0.0690	0.7678	0.0591	0.7865	-0.0107	0.9946	-0.0977	0.8407	-0.0870	0.8524
CG10804	CG10804	1632629_a_at	-0.0963	0.7367	-0.1861	0.3799	0.0066	0.9884	0.1298	0.7659	0.1461	0.4502	0.0163	0.9447	-0.2090	0.8644	-0.0721	0.9259	0.1368	0.8235
CG7039	CG7039	1632630_at	0.4201	0.0176	0.7400	0.1212	0.6281	0.0087	-0.0206	0.9774	0.1509	0.4101	0.1715	0.2868	0.0271	0.9875	0.4937	0.1925	0.4666	0.2501

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632631_at	0.1637	0.3816	0.0566	0.6556	0.0838	0.7010	-0.0906	0.8732	-0.1620	0.4262	-0.0714	0.7360	0.0605	0.9467	-0.0434	0.9198	-0.1039	0.7455
CG4140	CG4140	1632632_at	-0.0343	0.8892	0.0289	0.8757	0.1302	0.5374	-0.1092	0.8526	0.2410	0.2595	0.3502	0.0687	-0.2200	0.8049	0.2140	0.5855	0.4340	0.2576
TpnC25D	TpnC25D	1632633_at	-2.6584	0.0360	-2.4716	0.0345	-3.1517	0.0001	-0.2009	0.6908	-1.4946	0.0004	-1.2937	0.0004	0.3654	0.9407	-1.3148	0.3773	-1.6803	0.2837
---	---	1632634_at	0.0966	0.6440	0.0908	0.6973	0.1188	0.4164	-0.0144	0.9852	0.0448	0.8423	0.0592	0.7552	-0.0282	0.9852	0.0753	0.8807	0.1035	0.8110
---	---	1632635_at	0.0441	0.7986	0.0321	0.7562	0.1833	0.3984	0.0816	0.8908	0.1160	0.5940	0.0344	0.8846	-0.1489	0.7990	0.0168	0.9664	0.1657	0.5145
CG2145	CG2145	1632636_at	-0.3151	0.1404	-1.7632	0.0419	-0.9716	0.0003	0.1328	0.7228	0.2641	0.1162	0.1313	0.4037	-0.5622	0.6922	-1.0712	0.0795	-0.5090	0.3921
hiw	highwire	1632637_at	-0.0409	0.8654	0.0226	0.8230	-0.0927	0.5459	-0.0883	0.8402	-0.1631	0.3256	-0.0748	0.6587	0.1480	0.8222	0.0345	0.9377	-0.1135	0.7135
Arp8	CG7846	1632638_at	-0.2773	0.4899	0.0802	0.9323	0.0817	0.8583	-0.1787	0.5445	-0.2108	0.1657	-0.0321	0.8571	-0.1149	0.9816	0.0374	0.9865	0.1523	0.9189
CG13941	CG13941	1632639_at	1.6370	0.0026	0.5292	0.3458	0.8167	0.3616	0.5502	0.4873	2.1896	0.0009	1.6394	0.0018	0.3087	0.9495	1.1495	0.4258	0.8408	0.5862
CG9590	CG9590	1632640_at	0.4400	0.0238	0.6738	0.0278	0.6387	0.0034	0.0632	0.9082	0.3782	0.0376	0.3150	0.0495	-0.0176	0.9898	0.5566	0.0711	0.5742	0.0836
CG2104	CG2104	1632641_at	0.1652	0.4492	0.1550	0.3755	0.1550	0.4706	0.1176	0.8385	-0.0602	0.8247	-0.1777	0.3739	0.1770	0.7305	0.0224	0.9499	-0.1546	0.4978
CG2862	CG2862	1632642_a_at	-0.4608	0.0204	0.6027	0.1772	0.2342	0.4258	-0.3850	0.5242	-1.2552	0.0027	-0.8702	0.0084	-0.2163	0.7677	-0.0673	0.8794	0.1490	0.6577
CG10949	CG10949	1632643_at	-0.2081	0.2802	0.0000	1.0000	-0.0736	0.7430	-0.0571	0.9234	0.1198	0.5308	0.1769	0.2743	-0.0114	0.9922	0.3560	0.1694	0.3674	0.1911
cpo	couch potato	1632644_s_at	-0.5318	0.3891	-0.6830	0.1198	-1.2527	0.0089	0.0616	0.9688	0.7245	0.0795	0.6629	0.0731	0.6102	0.7506	0.5363	0.5312	-0.0739	0.9491
CG13330	CG13330	1632645_at	0.1474	0.4908	-0.0492	0.7861	0.0865	0.6316	0.1451	0.7405	0.1484	0.4604	0.0033	0.9890	0.0450	0.9588	0.0005	0.9998	-0.0445	0.8933
Ptth	prothoracicotropic	1632646_at	-0.6021	0.0054	-0.2590	0.0775	-0.3186	0.0630	0.1842	0.6354	0.0649	0.7793	-0.1194	0.5195	0.0645	0.8909	0.0131	0.9647	-0.0515	0.8166
---	---	1632647_s_at	0.0057	0.9804	0.0789	0.6871	0.1794	0.3028	0.0004	0.9995	-0.0609	0.7595	-0.0612	0.7306	0.0873	0.9118	0.0365	0.9350	-0.0508	0.8947
---	---	1632648_at	-0.0545	0.8515	-0.0568	0.7853	-0.2534	0.1602	-0.0533	0.9413	-0.0114	0.9716	0.0419	0.8621	0.1498	0.8686	0.0966	0.8408	-0.0532	0.9147
RluA-2	RluA-2	1632649_at	-0.5545	0.0497	-0.8483	0.1952	-0.8538	0.0132	-0.1598	0.7929	0.7131	0.0112	0.8729	0.0027	-0.2396	0.8815	0.3132	0.6391	0.5528	0.3807
CG5867	CG5867	1632650_at	0.0956	0.7075	-0.3317	0.5727	-0.9738	0.0040	-0.2704	0.6580	0.0642	0.8689	0.3347	0.2019	0.4472	0.7644	-0.3454	0.6126	-0.7927	0.2287
sut4	sugar transporter	1632651_s_at	0.0242	0.8973	-0.0528	0.6325	0.0927	0.6488	0.1154	0.8417	0.0551	0.8431	-0.0603	0.8033	-0.0905	0.8608	-0.0063	0.9885	0.0842	0.7202
CG30022 /// CG40100	CG30022 /// CG4(	1632652_s_at	0.3297	0.3378	-0.0997	0.7280	0.0628	0.7174	0.0087	0.9943	0.4505	0.0848	0.4418	0.0615	0.0004	0.9999	0.0997	0.8407	0.0993	0.8307
CG31555	CG31555	1632653_at	-0.5715	0.0647	0.1101	0.4432	-0.3568	0.0329	-0.2755	0.5832	-0.4738	0.0671	-0.1983	0.4057	0.1224	0.8270	0.2004	0.3859	0.0780	0.7782
l(2)37Cg	lethal (2) 37Cg	1632654_at	0.2198	0.5755	0.3235	0.1902	0.3978	0.1640	0.2610	0.6506	0.3033	0.2777	0.0423	0.8992	0.2165	0.8692	0.5208	0.3087	0.3044	0.5871
blue	bluestreak	1632655_at	-0.0514	0.8885	0.4428	0.5385	1.0405	0.0020	0.4125	0.4586	-0.3272	0.2805	-0.7396	0.0140	-0.1960	0.9333	0.2834	0.7485	0.4795	0.5448
CG12112	CG12112	1632656_at	-0.0349	0.8892	-0.2997	0.1522	-0.1788	0.6362	0.2964	0.3493	0.7648	0.0021	0.4683	0.0114	0.1272	0.9467	0.4215	0.4729	0.2943	0.6366
CG7166	CG7166	1632657_at	0.1977	0.3198	-0.0429	0.7060	-0.0608	0.7660	0.0302	0.9680	0.2446	0.2142	0.2144	0.2254	-0.0957	0.8940	-0.0347	0.9353	0.0610	0.8636
l(1)G0196	lethal (1) G0196	1632658_a_at	-0.0273	0.9302	0.1256	0.7886	-0.1047	0.6784	-0.0840	0.9067	-0.3087	0.1779	-0.2247	0.2805	0.0848	0.9611	-0.0005	0.9999	-0.0853	0.8984
NPC1b	NPC1b	1632659_at	0.0230	0.9181	-0.0498	0.6625	-0.1128	0.5557	0.0160	0.9819	0.1521	0.3725	0.1360	0.3786	-0.0156	0.9901	-0.0601	0.8786	-0.0445	0.9065
CkIIbeta2	Casein kinase II b	1632660_at	0.2749	0.1488	0.0000	1.0000	0.0750	0.7684	0.0280	0.9790	0.1708	0.5480	0.1428	0.5892	-0.0680	0.9467	-0.1142	0.7569	-0.0461	0.9142
---	---	1632661_at	0.2007	0.1876	0.1124	0.3924	0.1140	0.6558	-0.0011	0.9988	0.0321	0.8999	0.0331	0.8804	-0.0214	0.9841	-0.0261	0.9476	-0.0047	0.9914
CG31007	CG31007	1632662_at	-0.0077	0.9706	0.0637	0.6399	0.0636	0.7600	0.1160	0.8571	0.0719	0.7997	-0.0441	0.8714	0.0880	0.8909	0.1405	0.5847	0.0525	0.8717
CG7407	CG7407	1632663_at	-0.2090	0.3090	-0.5208	0.0288	-0.2922	0.0685	-0.0463	0.9351	0.0146	0.9531	0.0609	0.7355	-0.3522	0.4920	-0.2930	0.2233	0.0592	0.8549
Osi12	Osiris	1632664_at	-0.5207	0.1687	-1.8569	0.0021	-1.3274	0.0280	0.5804	0.6844	1.3578	0.0462	0.7774	0.1890	-0.2188	0.7152	-0.2568	0.2915	-0.0379	0.9121
CG6340	CG6340	1632665_a_at	-0.5943	0.0509	-0.5277	0.4645	-0.2356	0.2277	0.0359	0.9540	-0.1555	0.3727	-0.1914	0.2102	-0.3643	0.8461	-0.1598	0.8870	0.2045	0.8354
LanB2	Laminin	1632666_at	-2.0083	0.0009	-1.6657	0.0243	-1.7496	0.0003	-0.1155	0.8000	-0.5464	0.0097	-0.4309	0.0169	0.1420	0.9441	-0.0884	0.9275	-0.2304	0.7440
Gapdh2	G3-P dehydrogen	1632667_s_at	-0.1422	0.4351	-0.5702	0.0269	-0.2322	0.1120	-0.1261	0.6886	-0.2410	0.0985	-0.1150	0.3969	-0.5030	0.3712	-0.7672	0.0365	-0.2642	0.3921
---	---	1632668_at	0.4164	0.0549	-0.0790	0.6652	-0.3013	0.1696	0.2770	0.2693	0.6043	0.0024	0.3273	0.0216	0.0274	0.9898	-0.0226	0.9772	-0.0500	0.9360
Mcm2	Minichromosome	1632669_at	0.1518	0.8447	-0.9508	0.4608	-1.1833	0.1428	-0.6109	0.3901	1.1548	0.0112	1.7657	0.0010	-0.3222	0.9589	0.1536	0.9587	0.4758	0.8310
CG9028	CG9028	1632670_at	-0.3328	0.2201	0.1417	0.5702	-0.3752	0.0857	-0.1121	0.8628	0.1828	0.4459	0.2949	0.1518	0.1970	0.8344	0.4299	0.2538	0.2329	0.5768
---	---	1632671_at	0.1828	0.3606	-0.0242	0.8106	-0.0729	0.8061	-0.0855	0.9098	0.0395	0.9052	0.1250	0.6056	-0.0100	0.9939	-0.2335	0.4073	-0.2235	0.4457
lbk	kekkon-like	1632672_at	-0.7928	0.0179	-0.8612	0.0751	-1.3053	0.0001	0.0101	0.9909	0.5814	0.0090	0.5713	0.0058	0.5667	0.5754	0.6560	0.1395	0.0893	0.8854
CG13738	CG13738	1632673_at	0.1040	0.6675	0.0180	0.8649	0.0645	0.7556	-0.0064	0.9943	0.1404	0.4727	0.1467	0.3988	-0.1266	0.8761	0.0637	0.8911	0.1903	0.5722
CG31047	CG31047	1632674_at	0.0705	0.7370	0.0096	0.9579	0.1332	0.5035	0.0532	0.9345	-0.0281	0.9162	-0.0814	0.6832	0.0181	0.9848	0.0082	0.9842	-0.0099	0.9769
CG5455	CG5455	1632675_s_at	-0.1000	0.7131	0.4432	0.2802	0.5140	0.0163	-0.2368	0.4751	-0.5307	0.0096	-0.2940	0.0678	-0.1255	0.9260	0.0421	0.9533	0.1676	0.7464
CG11897	CG11897	1632676_s_at	0.9795	0.0139	-0.6722	0.2847	-0.2852	0.1474	-0.0467	0.9528	1.2652	0.0006	1.3119	0.0003	-0.3539	0.8270	-0.4498	0.5158	-0.0959	0.9173
GM130	GM130	1632677_a_at	0.8961	0.0138	0.9109	0.0188	0.7535	0.0031	0.1143	0.8545	0.6805	0.0091	0.5662	0.0126	0.3281	0.7644	0.7118	0.1268	0.3837	0.4148
S2P	S2P	1632678_at	-0.3628	0.0387	0.0032	0.9821	0.3339	0.0530	0.0197	0.9772	-0.1266	0.4699	-0.1463	0.3407	-0.2718	0.5978	0.1572	0.4997	0.4290	0.0931
esn	espinas	1632679_s_at	-0.1969	0.4321	-0.3711	0.1196	-0.0621	0.6958	-0.0441	0.9518	-0.4928	0.0223	-0.4488	0.0206	-0.1800	0.8564	-0.6609	0.1103	-0.4808	0.2544
Pgant35A	Polypeptide N-ace	1632680_at	-0.5271	0.1614	0.4642	0.4267	0.6881	0.0104	-0.1924	0.5681	-1.0508	0.0005	-0.8584	0.0006	-0.2720	0.8744	0.0911	0.9329	0.3631	0.6188
grn	lethal(3)84Fa	1632681_at	0.0395	0.9155	-0.2504	0.0981	0.1201	0.5027	0.1194	0.8897	0.1029	0.7661	-0.0165	0.9639	-0.1625	0.7726	-0.1801	0.4547	-0.0177	0.9596
---	---	1632682_at	0.2472	0.1377	0.0964	0.4807	0.5142	0.0319	0.1860	0.7732	0.1840	0.5233	-0.0020	0.9953	-0.0117	0.9913	0.1774	0.4285	0.1891	0.4078
copiaGIP	gag-int-pol	1632683_s_at	0.4546	0.0808	0.3577	0.0356	0.4280	0.0211	0.1376	0.7790	0.1205	0.5866	-0.0171	0.9473	0.1165	0.7893	0.0301	0.9170	-0.0864	0.6715
CG4575 /// gt	CG4575 /// giant	1632684_s_at	0.2930	0.1998	0.2792	0.1707	0.0784	0.7645	-0.0193	0.9803	-0.0629	0.7883	-0.0437	0.8435	0.1931	0.8009	0.0437	0.9316	-0.1493	0.6668
CG32603	CG32603	1632685_at	0.2269	0.4069	0.3376	0.2634	0.4176	0.0519	-0.0602	0.9480	-0.2700	0.3018	-0.2098	0.3818	0.0277	0.9816	-0.0575	0.8746	-0.0852	0.7764
---	---	1632686_at	-0.1202	0.6258	-0.1203	0.5191	-0.0156	0.9379	0.0066	0.9956	-0.0432	0.9292	-0.0497	0.9046	0.0097	0.9922	0.0915	0.7248	0.0818	0.7527
mib1	Mind Bomb	1632687_at	0.1036	0.6964	-0.1684	0.5536	-0.0699	0.7902	0.2964	0.5914	-0.0072	0.9868	-0.3036	0.2258	0.0420	0.9829	-0.2759	0.5585	-0.3179	0.4944
CG11594	CG11594	1632688_s_at	2.0108	0.0007	1.2667	0.1889	1.8371	0.0005	-0.0894	0.8977	-0.8203	0.0042	-0.7309	0.0041	-0.7025	0.7464	-1.5476	0.1005	-0.8451	0.3705
CG1307	hypothetical 23.1k	1632689_at	-0.0198	0.9504	0.6756	0.0394	1.0890	0.0025	-0.0255	0.9715	-0.5638	0.0078	-0.5383	0.0057	-0.4030	0.7220	0.1667	0.7770	0.5698	0.2525
CG13049	CG13049	1632690_a_at	0.1547	0.3651	0.0871	0.6171	0.1408	0.4337	-0.0786	0.8791	-0.0622	0.7688	0.0165	0.9397	-0.0933	0.8767	-0.0625	0.8440	0.0308	0.9237
R	Enhancer of faf	1632691_at	-0.1321	0.7153	-0.3310	0.0536	-0.3323	0.0677	0.3330	0.2435	0.5590	0.0060	0.2261	0.1322	0.2167	0.8612	0.3092	0.5579	0.0926	0.8933
---	---	1632692_at	-0.0812	0.6652	0.1705	0.1254	0.2529	0.2766	-0.0134	0.9860	-0.1770	0.3425	-0.1637	0.3282	-0.0080	0.9952	0.1096	0.7692	0.1176	0.7409
Lcp65Ac	Lcp65Ac	1632693_at	-0.0784	0.5762	-0.2796	0.4778	0.1546	0.3783	0.0592	0.9441	-0.0561	0.8586	-0.1152	0.6413	-0.2260	0.7576	-0.1051	0.7833	0.1209	0.7328
CG31918	CG31918	1632694_at	-0.1231	0.6340	-0.3213	0.2566	-0.2882	0.1644	-0.0416	0.9603	0.2151	0.3370	0.2566	0.1921	-0.1659	0.8814	-0.1246	0.8220	0.0414	0.9440
CG18585	CG18585	1632695_at	0.1006	0.7828	-0.0747	0.5230	0.4650	0.0847	0.2625	0.6506	0.1452	0.6439	-0.1172	0.6899	-0.3746	0.6584	-0.2211	0.5399	0.1535	0.6877
CG15323 /// DsimCG15323	CG15323	1632696_at	0.3154	0.1688	0.1034	0.4815	0.0136	0.9623	-0.0786	0.9311	0.1128	0.7153	0.1913	0.4461	0.0399	0.9521	-0.0402	0.8903	-0.0801	0.7239
---	---	1632697_at	0.2732	0.1789	0.0156	0.9058	-0.1671	0.6019	-0.1987	0.7840	0.1359	0.6925	0.3346	0.2109	0.0274	0.9845	0.0812	0.8500	0.0538	0.9020
ush	u-shaped	1632698_at	0.0475	0.9466	1.5888	0.0832	0.2651	0.3358	-0.0356	0.9753	-0.2837	0.3165	-0.2480	0.3331	1.1886	0.6389	1.3284	0.1861	0.1398	0.9218
---	---	1632699_s_at	-0.1223	0.8529	-0.0209	0.9523	1.5296	0.0038	0.1357	0.8578	-0.8063	0.0101	-0.9420	0.0030	-1.1036	0.6749	-0.5366	0.6417	0.5671	0.6210

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18324	CG18324	1632700_a_at	-0.4086	0.3269	-0.3484	0.1144	-0.5321	0.0070	-0.3388	0.1555	-0.5440	0.0034	-0.2052	0.1109	-0.3092	0.8012	-0.6434	0.1948	-0.3341	0.5355
Pal	Peptidyl-alpha-hy	1632701_at	-0.3031	0.3885	0.0237	0.8730	-0.0881	0.6182	-0.1486	0.8791	-0.6453	0.0573	-0.4967	0.0953	0.1513	0.8395	-0.2528	0.4074	-0.4041	0.2094
CG14387	CG14387	1632702_at	0.1505	0.3565	0.0233	0.8763	-0.0828	0.6955	0.0394	0.9610	0.0245	0.9367	-0.0149	0.9572	-0.0221	0.9831	-0.0781	0.7950	-0.0560	0.8575
CG17003	CG17003	1632703_at	0.1842	0.4892	0.1028	0.4819	0.1565	0.5301	-0.0139	0.9874	0.0581	0.8267	0.0719	0.7540	0.0576	0.9683	0.1052	0.8377	0.0476	0.9287
CG4663	CG4663	1632704_at	0.3000	0.1781	0.0306	0.8885	0.4410	0.0399	0.1394	0.7390	-0.0398	0.8707	-0.1792	0.2830	-0.1566	0.8425	-0.1480	0.6820	0.0086	0.9868
ninaD	neither inactivatio	1632705_at	-1.5693	0.0833	-0.4583	0.1732	-1.5318	0.0017	-0.6754	0.4012	-1.4621	0.0066	-0.7866	0.0548	-0.0385	0.9919	-1.0043	0.2367	-0.9658	0.2848
CG13054	CG13054	1632706_at	0.0755	0.7807	0.0174	0.9323	0.1430	0.5184	0.0264	0.9790	0.0017	0.9962	-0.0247	0.9362	0.0359	0.9717	-0.0369	0.9275	-0.0728	0.8194
CG12379	CG12379	1632707_at	-0.0841	0.6443	0.0892	0.8007	0.1927	0.5290	0.3482	0.5304	0.2617	0.3763	-0.0865	0.7893	0.2129	0.8395	0.2646	0.5591	0.0517	0.9321
CG10510	CG10510	1632708_at	0.2568	0.2031	0.1390	0.4016	0.2692	0.2312	0.1028	0.8532	-0.0013	0.9962	-0.1041	0.6079	0.0559	0.9487	-0.1351	0.6310	-0.1910	0.4779
---	---	1632709_at	-0.0077	0.9678	0.2266	0.1961	0.0032	0.9896	0.0662	0.8817	0.0061	0.9793	-0.0601	0.7086	0.2978	0.7070	0.1679	0.6408	-0.1298	0.7333
CG4572 /// DpseGA18267	CG4572 /// GA18	1632710_s_at	0.1590	0.3323	0.3146	0.1029	0.3952	0.0146	-0.0311	0.9603	-0.6131	0.0036	-0.5820	0.0027	-0.1715	0.7464	-0.4714	0.0564	-0.2999	0.2018
CG7864	CG7864	1632711_at	0.4736	0.0089	0.3495	0.0752	0.4419	0.0136	-0.0066	0.9937	0.0102	0.9687	0.0169	0.9352	-0.1505	0.7733	-0.2241	0.3097	-0.0736	0.7847
CG17836 /// DbuzCG1783	CG17836	1632712_s_at	0.1548	0.5045	-0.1437	0.6745	-0.0911	0.6830	0.1402	0.7556	0.5382	0.0133	0.3980	0.0295	0.0960	0.9588	0.2950	0.5910	0.1991	0.7391
nos	nanos	1632713_at	1.1917	0.4023	-2.1771	0.3934	-1.6484	0.0962	-0.7734	0.4174	3.3854	0.0004	4.1588	0.0001	-1.3644	0.8603	-0.0459	0.9942	1.3185	0.7060
CG15711	CG15711	1632714_at	0.1278	0.5534	-0.0187	0.8603	0.1397	0.5914	0.0534	0.9639	0.0533	0.9018	-0.0001	0.9996	0.0520	0.9324	0.0420	0.8824	-0.0100	0.9743
---	---	1632715_at	1.7126	0.0051	-0.1232	0.6748	-0.3846	0.1950	0.4843	0.5854	1.7791	0.0028	1.2948	0.0070	0.5452	0.7307	-0.2513	0.7567	-0.7965	0.2574
CG5509	CG5509	1632716_at	-0.0189	0.9377	0.0044	0.9721	-0.1893	0.1825	-0.0055	0.9952	-0.1339	0.4877	-0.1284	0.4615	0.1860	0.6955	0.1531	0.4363	-0.0329	0.9033
CG14341	CG14341	1632717_a_at	-0.0042	0.9863	-0.6275	0.0406	-0.6642	0.0121	0.1415	0.6869	0.9119	0.0007	0.7704	0.0008	0.1599	0.8692	0.2465	0.5414	0.0866	0.8656
---	---	1632718_at	0.0464	0.8682	-0.0829	0.6105	0.0428	0.7813	0.0883	0.8640	0.0589	0.7930	-0.0294	0.8936	0.1322	0.8465	0.0605	0.8821	-0.0717	0.8432
CecC	Cecropin	1632719_at	0.0654	0.9627	-4.1066	0.1406	-2.1514	0.2960	2.2260	0.5376	5.1419	0.0143	2.9160	0.0848	0.4894	0.9689	1.2071	0.7578	0.7177	0.8699
LysX	Lysozyme X	1632720_at	0.3823	0.1376	-0.9056	0.0210	-0.2128	0.5056	0.6474	0.1832	1.0998	0.0035	0.4525	0.0862	-0.1321	0.9342	-0.2233	0.7018	-0.0912	0.8982
Nle	notchless	1632721_at	0.2599	0.2824	0.0214	0.9625	0.6964	0.0223	0.3115	0.3084	0.4598	0.0187	0.1484	0.3710	-0.2807	0.8305	0.2040	0.7551	0.4847	0.3878
CG13679	CG13679	1632722_x_at	0.6196	0.0345	0.3037	0.0949	0.3918	0.0920	0.0592	0.9110	0.1263	0.4716	0.0672	0.7045	0.1389	0.9298	0.0231	0.9806	-0.1158	0.8636
scaf6	scaf6	1632723_at	0.3682	0.0244	0.0122	0.9220	0.1980	0.2214	0.0973	0.7956	0.1680	0.2742	0.0707	0.6550	-0.0722	0.9013	-0.0659	0.8123	0.0063	0.9852
alpha-Cat	alpha-catenin	1632724_at	-0.3802	0.0535	-0.1923	0.5744	-0.6461	0.0022	-0.2023	0.5357	0.1690	0.3284	0.3713	0.0231	0.2330	0.8202	0.3527	0.3951	0.1198	0.8188
CG6691	CG6691	1632725_at	0.0189	0.9220	-0.0118	0.9233	0.1188	0.3948	-0.0069	0.9943	0.0170	0.9546	0.0239	0.9229	-0.0634	0.9276	0.1025	0.6876	0.1658	0.4824
Or67a	Odorant receptor	1632726_at	0.0704	0.6462	-0.1311	0.3510	0.0557	0.7597	0.1728	0.7138	0.1524	0.5004	-0.0204	0.9392	-0.0348	0.9643	-0.1098	0.6272	-0.0750	0.7566
CG14006	CG14006	1632727_at	-0.0554	0.7063	-0.1560	0.2882	0.0630	0.7668	-0.1169	0.7906	-0.0259	0.9180	0.0910	0.6176	-0.2995	0.4377	-0.1051	0.6280	0.1944	0.3475
Gcn2	Gcn2	1632728_at	-0.1955	0.5036	-0.1085	0.5906	-0.2691	0.1454	-0.4624	0.2177	-0.4843	0.0382	-0.0220	0.9375	-0.2501	0.8009	-0.3199	0.4380	-0.0698	0.9026
dmrt93B	doublesex-Mab re	1632729_at	0.4486	0.0407	0.0362	0.7417	0.1466	0.6329	0.1098	0.8908	0.0986	0.7595	-0.0113	0.9731	0.2509	0.7464	-0.1113	0.7819	-0.3621	0.2816
CG14013	CG14013	1632730_at	-0.0059	0.9806	0.0638	0.5347	-0.0349	0.8625	-0.0914	0.8350	-0.1208	0.4877	-0.0294	0.8814	0.0591	0.9342	0.0571	0.8567	-0.0020	0.9963
CG8613	CG8613	1632731_at	0.4443	0.0227	-0.3098	0.2962	0.0184	0.9358	-0.1389	0.7577	-0.1838	0.3509	-0.0449	0.8407	-0.4535	0.6272	-0.9196	0.0401	-0.4662	0.2485
sofe	sister of feo	1632732_at	-0.1569	0.7922	-0.6662	0.4402	-1.0567	0.1367	-0.2641	0.5636	0.7045	0.0094	0.9686	0.0014	0.0685	0.9901	0.0684	0.9734	-0.0002	0.9999
Hr4	Hr4	1632733_at	0.2303	0.2826	0.0394	0.7097	-0.3585	0.1508	-0.0802	0.8636	0.0405	0.8508	0.1207	0.4389	0.2326	0.7061	-0.0229	0.9566	-0.2554	0.3299
bnb	bangles and bead	1632734_s_at	0.4732	0.3367	1.4951	0.0651	1.0044	0.0150	-0.4050	0.7388	-2.0144	0.0034	-1.6094	0.0056	0.2154	0.8889	-0.8747	0.1332	-1.0901	0.0962
---	---	1632735_at	-0.0842	0.6090	-0.0340	0.7767	-0.0343	0.8614	-0.0192	0.9803	-0.0733	0.7399	-0.0542	0.7973	0.0242	0.9742	0.0161	0.9587	-0.0081	0.9784
CG6617	CG6617	1632736_at	-0.2574	0.1208	0.6168	0.0117	0.6223	0.0061	-0.0717	0.8907	-0.8181	0.0014	-0.7463	0.0012	-0.0536	0.9405	0.0447	0.8903	0.0983	0.6883
---	---	1632737_at	0.0062	0.9812	0.1580	0.2555	0.1993	0.4710	0.0208	0.9763	-0.3032	0.0823	-0.3241	0.0432	0.1574	0.8275	0.0171	0.9737	-0.1403	0.6719
---	---	1632738_at	0.2961	0.0857	-0.0651	0.6330	-0.2460	0.2505	0.0082	0.9924	0.0558	0.7993	0.0476	0.8134	-0.0262	0.9816	-0.1449	0.5972	-0.1187	0.6731
CG32536	CG32536	1632739_at	-0.4783	0.3310	-0.3060	0.4675	-0.0305	0.9008	0.4311	0.2876	0.1681	0.5051	-0.2630	0.2183	0.0423	0.9900	0.3197	0.7007	0.2774	0.7442
CG31818	CG31818	1632740_at	-0.0081	0.9768	0.0579	0.7127	0.0890	0.6303	0.1757	0.6533	0.0949	0.6556	-0.0807	0.6828	0.1849	0.7726	0.0675	0.8611	-0.1174	0.7060
---	---	1632741_at	0.0746	0.7642	0.1413	0.4488	-0.1045	0.5199	0.0511	0.9436	0.0854	0.7208	0.0343	0.8883	0.1761	0.8222	0.0761	0.8684	-0.1000	0.8002
eyg	eye gone	1632742_at	-0.0473	0.9043	0.0957	0.5001	-0.0656	0.8128	-0.1348	0.8217	-0.1407	0.5781	-0.0059	0.9844	-0.0575	0.9611	-0.0188	0.9733	0.0386	0.9337
dre4	supressor of Ty el	1632743_a_at	0.1120	0.8257	0.3643	0.0677	0.7124	0.0192	0.2932	0.4940	-0.1424	0.5624	-0.4356	0.0391	0.0014	0.9998	0.2056	0.7707	0.2042	0.7610
if	alphaPS2 integrin	1632744_a_at	-1.4771	0.0047	-2.2984	0.0056	-2.7056	0.0000	-0.2254	0.7511	0.4586	0.1359	0.6841	0.0214	0.1593	0.9046	-0.2565	0.6177	-0.4158	0.3941
CG3610	CG3610	1632745_at	0.1534	0.2756	0.1662	0.3094	0.0606	0.7878	-0.1104	0.8822	-0.1176	0.6868	-0.0071	0.9825	-0.0471	0.9478	-0.2452	0.2430	-0.1981	0.3764
CG13466 /// CG13467	CG13466 /// CG1	1632746_at	0.1278	0.4265	0.0597	0.5884	0.0699	0.7865	0.0334	0.9603	0.0624	0.7693	0.0290	0.8912	0.1180	0.8235	0.0474	0.8807	-0.0707	0.7879
CG11164	CG11164	1632747_at	0.1607	0.6868	-0.1549	0.6639	0.0489	0.8964	-0.0774	0.9223	0.6368	0.0159	0.7142	0.0058	-0.2179	0.9101	0.1017	0.9260	0.3196	0.6713
CG15084	CG15084	1632748_at	-0.0322	0.8892	-0.3437	0.1913	-0.6945	0.0021	-0.2018	0.6908	0.2705	0.2505	0.4723	0.0319	0.0628	0.9522	-0.0521	0.9152	-0.1149	0.7516
Eig71Ek	Eig71Ek	1632749_at	-0.0916	0.5947	0.0335	0.9049	-0.0251	0.8889	0.0038	0.9956	0.0472	0.8490	0.0434	0.8444	-0.0105	0.9914	0.0826	0.7453	0.0931	0.6993
---	---	1632750_at	0.1993	0.2524	0.2273	0.2060	0.3928	0.1412	0.1092	0.7447	-0.0129	0.9531	-0.1221	0.3688	-0.0690	0.9677	0.0683	0.9225	0.1373	0.8017
CG6301	CG6301	1632751_at	0.7368	0.0210	0.2868	0.3321	0.3940	0.1122	-0.0813	0.8942	-0.0051	0.9871	0.0762	0.7253	-0.1369	0.9029	-0.2105	0.6322	-0.0737	0.8949
Ast	drostatin-A4	1632752_at	-0.2051	0.8512	-0.8950	0.0411	-0.7688	0.0186	0.2149	0.9293	0.2621	0.7573	0.0472	0.9584	-0.0949	0.9698	-0.5172	0.4203	-0.4223	0.5341
CG3980	CG3980	1632753_a_at	-0.8391	0.0221	-0.9046	0.0288	-0.9314	0.0003	-0.3214	0.2871	0.2785	0.1155	0.5999	0.0033	-0.1703	0.8494	0.2644	0.4878	0.4347	0.2606
CG33283	CG33283	1632754_at	-0.0373	0.8621	0.0732	0.5540	0.1181	0.4517	-0.0019	0.9978	-0.0560	0.7803	-0.0541	0.7669	-0.0729	0.9168	0.0790	0.7951	0.1520	0.5518
CG31266	CG31266	1632755_at	-0.2370	0.9564	0.0370	0.7165	-0.1917	0.3605	-0.0984	0.9932	-2.2753	0.3234	-2.1769	0.2903	0.1599	0.9913	-1.9723	0.5373	-2.1321	0.5027
CG33249	CG33249	1632756_at	-0.7030	0.4492	-0.2905	0.8278	-0.3769	0.2763	0.2705	0.5539	-0.0181	0.9592	-0.2886	0.1707	0.5399	0.9011	0.7927	0.6472	0.2528	0.9086
CG11760	CG11760	1632757_a_at	0.3437	0.1386	0.0770	0.7723	-0.3834	0.1764	-0.0630	0.8819	0.4890	0.0055	0.5520	0.0019	0.3418	0.7779	0.1743	0.7921	-0.1674	0.7915
CG13872	CG13872	1632758_at	-0.0717	0.6284	-0.1151	0.4523	0.0413	0.8702	0.2316	0.5756	0.1838	0.3986	-0.0479	0.8454	-0.0541	0.9148	0.0585	0.7925	0.1126	0.5482
CG15864	CG15864	1632759_at	0.0001	0.9994	-0.0332	0.8156	0.2340	0.1646	0.2354	0.5149	0.0964	0.6548	-0.1390	0.4389	-0.0683	0.9142	0.0310	0.9291	0.0993	0.6853
---	---	1632760_at	0.1100	0.4288	-0.0115	0.9517	0.2639	0.0838	0.1358	0.6457	0.0875	0.5713	-0.0483	0.7572	0.0812	0.9330	0.0724	0.8680	-0.0087	0.9857
CG34106	CG34106	1632761_at	-0.1306	0.4578	-0.0336	0.7400	0.0556	0.7704	0.0141	0.9840	-0.0797	0.6711	-0.0938	0.5652	0.0206	0.9776	-0.0652	0.7551	-0.0858	0.6498
CG3176 /// CG32817	CG3176 /// CG32	1632762_s_at	-0.0216	0.9069	-0.0886	0.7047	0.0948	0.6892	-0.0342	0.9608	-0.1544	0.4216	-0.1201	0.5011	-0.2816	0.7677	-0.2824	0.4954	-0.0008	0.9992
CG33506	CG33506	1632763_at	-0.0441	0.8314	0.0895	0.8135	-0.0549	0.8438	-0.3537	0.2217	-0.2321	0.1778	0.1216	0.4586	-0.3272	0.7464	-0.1916	0.6883	0.1356	0.7918
Tdc2	Tyrosine decarbo	1632764_at	0.2735	0.2589	0.2767	0.1354	0.3236	0.0651	-0.1403	0.8028	-0.0262	0.9340	0.1141	0.6166	-0.0799	0.9168	0.0151	0.9734	0.0950	0.7553
CG6623	CG6623	1632765_at	-0.1848	0.2683	0.0473	0.8993	-0.3497	0.0636	-0.1951	0.5311	0.0897	0.6186	0.2848	0.0546	0.1808	0.8384	0.2805	0.4421	0.0996	0.8287
---	---	1632766_at	-0.4480	0.0738	0.0424	0.6805	-0.0153	0.9473	-0.0741	0.9116	-0.6169	0.0106	-0.5428	0.0115	0.0642	0.9238	-0.0085	0.9841	-0.0727	0.7814
Gr68a	Gustatory recepto	1632767_at	0.2042	0.3257	-0.0686	0.6841	0.0884	0.6147	0.0650	0.9314	0.1627	0.4894	0.0978	0.6775	0.0007	0.9998	-0.0761	0.8448	-0.0768	0.8310
CG13002	CG13002	1632768_at	-0.0750	0.8259	0.1635	0.3801	0.0972	0.7434	-0.0924	0.8578	-0.1180	0.5543	-0.0256	0.9089	-0.0111	0.9946	0.1242	0.7998	0.1353	0.7621

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632769_at	0.4727	0.0838	0.2987	0.1277	0.2488	0.2693	-0.0624	0.9218	0.0096	0.9743	0.0720	0.7253	0.0994	0.8940	0.1067	0.7424	0.0074	0.9868
---	---	1632770_at	0.3606	0.5250	0.6200	0.4741	0.1955	0.4291	-0.4827	0.5886	-0.1163	0.8431	0.3664	0.3832	-0.0727	0.9848	0.1278	0.9233	0.2006	0.8562
mRpS29	mitochondrial ribo	1632771_at	-0.2249	0.2139	-0.3519	0.1684	-0.2113	0.2153	0.2005	0.5311	-0.0493	0.8112	-0.2497	0.0941	0.0553	0.9555	-0.1931	0.5184	-0.2484	0.3981
CG15203	CG15203	1632772_at	1.1862	0.0024	0.5865	0.3170	1.2381	0.0012	0.0942	0.9311	-0.6420	0.0497	-0.7362	0.0181	-0.4782	0.7187	-1.0586	0.0711	-0.5805	0.3027
---	---	1632773_a_at	0.1730	0.3005	0.2561	0.2338	0.2504	0.1519	-0.0188	0.9803	-0.0023	0.9933	0.0165	0.9435	0.0426	0.9611	0.0828	0.7874	0.0402	0.9057
CG12065 /// DsimCG12065	CG12065	1632774_s_at	-0.7833	0.0565	0.0457	0.7837	0.2346	0.3097	0.3849	0.3553	-0.4524	0.0613	-0.8372	0.0027	0.0661	0.9775	0.2935	0.6209	0.2275	0.7154
CG12854	CG12854	1632775_at	0.2078	0.5168	-0.1045	0.5624	-0.6372	0.0606	-0.0364	0.9651	0.3313	0.1356	0.3677	0.0677	0.3504	0.7823	0.0493	0.9561	-0.3011	0.6117
CSN5	JUN activation do	1632776_at	0.5339	0.0749	0.6971	0.0893	0.7019	0.0014	-0.0709	0.8817	-0.3433	0.0415	-0.2724	0.0649	-0.0544	0.9742	0.0085	0.9931	0.0629	0.9131
CG8204	CG8204	1632777_at	-0.6989	0.0367	-0.3290	0.2388	-0.1007	0.7386	0.1189	0.8161	-0.1797	0.3814	-0.2986	0.0957	-0.0674	0.9701	0.0636	0.9309	0.1310	0.8190
---	---	1632778_at	0.1737	0.3936	0.0064	0.9745	-0.0542	0.7495	0.0721	0.9247	0.2194	0.3462	0.1473	0.5040	-0.0256	0.9778	-0.0026	0.9959	0.0230	0.9408
---	---	1632779_at	0.1416	0.4795	-0.1904	0.1919	0.1172	0.5262	0.2822	0.5680	0.3499	0.1648	0.0677	0.8129	-0.0536	0.9246	0.0772	0.7260	0.1308	0.5100
---	---	1632780_at	0.1421	0.3609	-0.0498	0.7494	0.0123	0.9522	0.1530	0.7929	0.2112	0.3829	0.0582	0.8293	-0.0107	0.9928	-0.2407	0.3651	-0.2300	0.4031
Abd-B	Abdominal-B	1632781_s_at	0.3790	0.2475	-0.0067	0.9913	-0.4584	0.0676	-0.1997	0.6015	0.5127	0.0159	0.7124	0.0022	-0.0336	0.9915	0.1306	0.9075	0.1642	0.8655
CG7675	CG7675	1632782_a_at	-0.2859	0.6676	-1.2345	0.0870	-1.0662	0.0105	0.8747	0.1956	0.9639	0.0263	0.0893	0.8507	0.2798	0.9095	-0.1395	0.9178	-0.4193	0.6578
Su(var)2-10	zinc finger-contair	1632783_a_at	0.0746	0.8093	0.6488	0.0144	0.2136	0.2952	-0.4262	0.3744	-0.0909	0.7824	0.3352	0.1660	0.1027	0.8940	0.4767	0.1042	0.3740	0.2110
cnk	connector enhanc	1632784_at	-0.4611	0.2655	-0.1120	0.6748	-0.4565	0.0148	-0.1131	0.8022	-0.0661	0.7606	0.0470	0.8188	0.1721	0.8999	0.2483	0.6512	0.0762	0.9122
---	---	1632785_at	0.0662	0.7769	0.1309	0.4960	-0.1782	0.3225	0.0199	0.9803	0.0607	0.8023	0.0408	0.8595	0.0823	0.9421	-0.0330	0.9531	-0.1153	0.7800
---	---	1632786_at	0.0188	0.9522	-0.0378	0.7304	-0.0224	0.9404	0.0481	0.9548	0.0610	0.8368	0.0129	0.9649	0.0120	0.9913	-0.1123	0.6536	-0.1243	0.6152
Pka-C2	cAMP-dependent	1632787_a_at	0.2849	0.1263	0.1467	0.3923	0.0658	0.7065	-0.0011	0.9988	0.2453	0.2306	0.2464	0.1777	0.0319	0.9657	-0.0557	0.8353	-0.0876	0.6941
---	---	1632788_at	-0.0249	0.9226	0.0891	0.5530	0.1310	0.4010	-0.0497	0.9380	-0.0221	0.9330	0.0276	0.9022	0.0340	0.9589	0.1555	0.4065	0.1214	0.5422
CG3701	CG3701	1632789_at	0.0622	0.6994	0.0542	0.7030	0.0495	0.8500	-0.0848	0.8903	0.0186	0.9517	0.1035	0.6220	0.0076	0.9929	0.0708	0.7779	0.0632	0.7980
dream	Strica	1632790_at	2.1367	0.0018	1.7979	0.0014	1.7789	0.0005	0.4478	0.1787	0.8622	0.0020	0.4144	0.0294	0.5636	0.6592	0.5987	0.2361	0.0350	0.9637
CG32479	CG32479	1632791_at	0.4733	0.2289	0.1474	0.3997	0.0993	0.5746	-0.0038	0.9956	0.8229	0.0025	0.8267	0.0015	0.1234	0.9390	0.5422	0.2584	0.4187	0.4085
---	---	1632792_at	0.3466	0.3790	0.3134	0.3674	-0.1654	0.6622	-0.3218	0.5038	0.1298	0.6511	0.4517	0.0521	0.0858	0.9717	0.2272	0.7578	0.1414	0.8618
CG12723	CG12723	1632793_at	0.2255	0.3309	0.4097	0.1341	-0.0986	0.5315	-0.1655	0.6808	0.0720	0.7459	0.2375	0.1552	0.2884	0.7215	0.0656	0.8964	-0.2228	0.5299
CG11526	CG11526	1632794_a_at	-0.1046	0.5983	-0.2469	0.0586	-0.4259	0.1333	-0.0074	0.9956	0.1147	0.7395	0.1221	0.6907	0.1405	0.7811	-0.0359	0.9157	-0.1764	0.4322
---	---	1632795_at	0.0216	0.9106	0.0136	0.8979	0.1424	0.3315	0.0149	0.9848	0.0069	0.9800	-0.0080	0.9711	-0.0428	0.9589	0.0509	0.8820	0.0936	0.7307
EcR	ecdysone recepto	1632796_s_at	-0.0009	0.9988	0.3608	0.2483	-0.1185	0.6853	-0.4746	0.1649	-0.5089	0.0224	-0.0343	0.8900	0.0793	0.9775	-0.1147	0.9111	-0.1940	0.8162
---	---	1632797_a_at	0.1085	0.5146	0.0060	0.9796	-0.1267	0.4759	-0.1674	0.6718	-0.1046	0.6121	0.0628	0.7589	0.0228	0.9761	-0.0565	0.8123	-0.0793	0.6985
Sry-alpha	serendipity-alpha	1632798_at	0.1640	0.5598	-0.0629	0.6071	-0.0859	0.5932	-0.1207	0.7007	0.0058	0.9793	0.1265	0.3378	0.0089	0.9917	-0.1431	0.4875	-0.1520	0.4669
---	---	1632799_at	0.6541	0.0137	0.1840	0.3756	0.0351	0.8281	0.0548	0.9311	0.5027	0.0145	0.4479	0.0148	0.1403	0.8705	-0.0155	0.9806	-0.1558	0.6800
CG15278	CG15278	1632800_at	-0.1595	0.4063	0.0988	0.4964	-0.1309	0.3547	-0.0606	0.9130	-0.2173	0.2139	-0.1567	0.3279	0.0092	0.9952	-0.0935	0.8478	-0.1027	0.8178
---	---	1632801_at	0.3678	0.1006	0.0510	0.6213	0.1851	0.3654	0.0137	0.9860	0.0909	0.6642	0.0772	0.6902	0.0854	0.9199	-0.0317	0.9460	-0.1171	0.7192
Jhl-26	Juvenile hormone	1632802_at	-1.5829	0.0039	-1.4195	0.0936	-0.6920	0.0174	-0.4704	0.3625	-2.0117	0.0003	-1.5413	0.0005	-1.2323	0.3712	-1.7725	0.0391	-0.5402	0.4841
CG3719	thioredoxin	1632803_at	-0.8387	0.0152	-1.2696	0.0262	-1.0826	0.0023	0.4376	0.3775	0.6077	0.0370	0.1701	0.5345	0.1816	0.8379	0.2059	0.5972	0.0242	0.9640
CG7175	CG7175	1632804_at	0.0847	0.6595	1.1215	0.0093	0.9446	0.0003	-0.3819	0.1140	-0.9045	0.0004	-0.5226	0.0025	-0.1388	0.7810	0.1087	0.6425	0.2475	0.2681
CG9573	CG9573	1632805_at	0.1881	0.4166	-0.1276	0.4478	0.0106	0.9751	0.2290	0.6010	0.4146	0.0635	0.1856	0.3600	0.0187	0.9892	-0.0586	0.8885	-0.0773	0.8307
---	---	1632806_s_at	-0.0129	0.9737	-0.0947	0.6151	-0.1437	0.4821	0.1737	0.7138	0.2022	0.3587	0.0285	0.9137	0.0035	0.9976	-0.0244	0.9507	-0.0279	0.9360
Dhap-at	Glycerol 3-phosph	1632807_at	0.4873	0.1221	0.2520	0.0564	-0.2703	0.1053	-0.1708	0.6580	0.6764	0.0039	0.8472	0.0009	0.2875	0.7726	0.2294	0.6150	-0.0582	0.9211
CG6012	CG6012	1632808_at	0.2474	0.5820	0.1085	0.4100	-0.4787	0.1553	-0.3422	0.5858	-0.2209	0.5143	0.1214	0.7218	0.0670	0.9474	-0.4741	0.1280	-0.5412	0.1156
CG14762	gp150-like	1632809_at	2.3243	0.0021	0.7920	0.5309	2.1258	0.0007	0.5754	0.5311	1.1471	0.0249	0.5717	0.1814	-0.8196	0.7810	-0.3677	0.8258	0.4519	0.7578
CG7386	CG7386	1632810_at	-0.7654	0.0091	-0.0550	0.7681	0.4132	0.2081	0.2834	0.5617	-0.2570	0.3092	-0.5404	0.0239	-0.2073	0.7220	0.1872	0.4487	0.3944	0.1390
Andorra	Andorra	1632811_at	-0.1320	0.5811	0.0746	0.5153	-0.1494	0.3291	0.0511	0.9441	-0.1026	0.6651	-0.1536	0.4332	0.2662	0.7423	0.2954	0.3778	0.0293	0.9514
Eo	Ecdysone oxidase	1632812_at	0.0392	0.9045	0.3863	0.0931	0.0123	0.9646	-0.2263	0.7497	-0.3522	0.2549	-0.1259	0.6980	0.2376	0.7215	0.0203	0.9647	-0.2173	0.4457
CG11975	CG11975	1632813_at	0.4213	0.0608	0.9543	0.0340	1.7183	0.0001	0.3489	0.3353	-0.7613	0.0039	-1.1102	0.0005	-0.4227	0.6557	-0.2639	0.5075	0.1588	0.7154
plu	plutonium	1632814_at	0.4576	0.4256	-0.8238	0.1302	-0.4072	0.3846	0.2145	0.7753	1.4403	0.0015	1.2258	0.0018	-0.3828	0.9159	-0.0904	0.9650	0.2924	0.8599
CG16952	CG16952	1632815_at	-0.5307	0.1179	-0.0595	0.8576	-0.1596	0.3030	0.0084	0.9937	-0.1361	0.5392	-0.1445	0.4607	0.0910	0.9495	0.2620	0.5591	0.1710	0.7253
Es2	Es2	1632816_at	-0.1957	0.4287	-0.2971	0.0430	-0.4156	0.0668	0.1346	0.8164	0.2896	0.2024	0.1550	0.4754	0.2157	0.7644	0.2300	0.4566	0.0143	0.9769
---	---	1632817_at	0.2113	0.3001	0.0529	0.7853	0.1073	0.6143	0.3130	0.5255	0.2959	0.2531	-0.0171	0.9586	0.0945	0.8688	0.0672	0.8189	-0.0273	0.9316
CG32281	CG32281	1632818_at	0.7972	0.0032	0.0092	0.9905	0.4756	0.0243	0.1594	0.6836	0.5637	0.0081	0.4043	0.0211	-0.1102	0.9487	-0.0296	0.9727	0.0806	0.9095
CG31224	lethal (3) c00083	1632819_at	0.1775	0.5865	-0.0762	0.4906	-0.6719	0.0299	-0.1570	0.7379	0.3110	0.1324	0.4680	0.0198	0.3352	0.7826	0.0626	0.9409	-0.2726	0.6311
---	---	1632820_at	-0.0274	0.9019	-0.0982	0.5667	0.0214	0.9347	0.0424	0.9559	0.1878	0.3769	0.1454	0.4603	0.1323	0.8202	0.1368	0.5889	0.0045	0.9915
CG3287	CG3287	1632821_a_at	-0.5794	0.0644	-0.1860	0.6895	0.2044	0.4666	0.1576	0.7604	-0.2583	0.2464	-0.4158	0.0427	-0.1640	0.9365	0.2025	0.8114	0.3665	0.6016
gig	gigas	1632822_at	-0.2105	0.5248	-0.3740	0.3747	-0.4212	0.0421	-0.0325	0.9695	0.2249	0.3078	0.2574	0.1867	-0.0942	0.9621	0.0788	0.9277	0.1730	0.7954
CG17788	CG17788	1632823_at	0.1447	0.5377	0.1031	0.2979	0.3305	0.1175	0.1345	0.7803	0.0993	0.6616	-0.0352	0.8818	-0.1111	0.8042	0.1169	0.5479	0.2280	0.2358
Fs	dFollistatin	1632824_at	0.3704	0.2212	1.2386	0.0902	0.5468	0.0058	0.3243	0.4968	0.0847	0.7824	-0.2396	0.2956	1.1722	0.3338	1.2243	0.0714	0.0521	0.9539
---	---	1632825_at	0.1455	0.4679	0.0836	0.5651	0.3412	0.0432	0.1061	0.8379	0.0095	0.9750	-0.0966	0.6253	-0.0043	0.9964	0.1155	0.6483	0.1198	0.6328
CG17237	CG17237	1632826_at	0.0016	0.9958	0.0720	0.5175	0.0621	0.8152	-0.1430	0.7368	-0.0392	0.8758	0.1038	0.5810	0.0063	0.9964	0.0090	0.9903	0.0027	0.9965
mus210	xeroderma pigmer	1632827_a_at	-0.3126	0.1445	0.1370	0.2103	0.0517	0.8642	-0.2222	0.5516	-0.8118	0.0021	-0.5896	0.0052	-0.1981	0.8202	-0.2266	0.5434	-0.0285	0.9552
CG3033	CG3033	1632828_at	-0.2155	0.3944	0.1302	0.3939	0.7645	0.0013	0.1769	0.6663	-0.3065	0.1158	-0.4834	0.0135	-0.3454	0.6660	0.0762	0.8671	0.4216	0.2107
CG10730	CG10730	1632829_at	0.0881	0.6563	0.1584	0.3942	0.0556	0.7558	-0.0870	0.8349	-0.0741	0.6806	0.0129	0.9484	0.1801	0.7215	0.1567	0.4598	-0.0234	0.9369
G-oalpha47A	brokenhearted	1632830_s_at	2.3132	0.0019	1.7078	0.0217	0.6758	0.1477	-0.8887	0.2105	0.2531	0.5851	1.1419	0.0095	0.1826	0.9400	-0.2383	0.8017	-0.4209	0.5984
---	---	1632831_at	0.1365	0.5135	-0.0504	0.6325	0.0880	0.6595	-0.1041	0.8605	0.0022	0.9942	0.1064	0.6232	-0.0893	0.8317	-0.0278	0.9185	0.0615	0.7620
---	---	1632832_at	-0.4505	0.0206	-1.7136	0.0089	-1.4743	0.0019	0.2524	0.7969	1.4775	0.0036	1.2252	0.0050	0.0625	0.9772	0.0188	0.9848	-0.0436	0.9520
CG7431	CG7431	1632833_at	0.2387	0.3281	-0.2067	0.1461	-0.1353	0.3723	0.3196	0.4270	0.3496	0.1142	0.0300	0.9111	0.0076	0.9952	-0.2847	0.3203	-0.2924	0.3307
zetaCOP	zetaCOP	1632834_a_at	0.6294	0.0674	1.1117	0.0150	1.0282	0.0004	0.0231	0.9744	0.0630	0.7640	0.0399	0.8435	0.0454	0.9806	0.4753	0.2265	0.4299	0.3027
CG9376	CG9376	1632835_at	0.0009	0.9973	0.0271	0.8680	0.0719	0.6739	-0.0119	0.9883	0.1337	0.4966	0.1456	0.4036	-0.0599	0.9462	0.0562	0.8850	0.1161	0.6990
---	---	1632836_at	1.9004	0.0004	2.2671	0.0022	2.9578	0.0000	0.3434	0.1601	-0.3160	0.0394	-0.6594	0.0010	-0.3289	0.6749	0.0007	0.9998	0.3296	0.3171
CG2129	CG2129	1632837_at	0.2902	0.3077	0.5819	0.1195	0.6806	0.0028	0.1323	0.8064	0.2379	0.2700	0.1056	0.6300	-0.0283	0.9901	0.3177	0.5411	0.3460	0.5032

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
ATPsyn-gamma	F1ATP synthase	1632838_at	-0.1611	0.3723	0.1062	0.6589	-0.2409	0.3245	-0.1729	0.7579	-0.4895	0.0477	-0.3166	0.1406	0.1790	0.8292	-0.2156	0.5531	-0.3946	0.2738
CG8312	CG8312	1632839_a_at	-0.6419	0.2373	-0.3682	0.7111	-0.3959	0.1292	-0.1355	0.7929	-0.5331	0.0193	-0.3976	0.0405	0.0148	0.9977	-0.0856	0.9646	-0.1004	0.9510
fus	fusilli	1632840_s_at	-0.3257	0.5230	0.4961	0.5315	0.3057	0.2013	0.0200	0.9876	-0.5563	0.0730	-0.5763	0.0422	0.2174	0.9467	0.2639	0.8444	0.0466	0.9766
Hsp70Bc	heat shock 70	1632841_x_at	-0.9834	0.6454	-0.9451	0.1202	-0.8830	0.5802	1.2442	0.8330	1.6979	0.4681	0.4537	0.8619	0.5596	0.8917	0.8443	0.6093	0.2847	0.8932
---	---	1632842_at	0.0119	0.9548	0.1406	0.2861	-0.2036	0.2505	-0.3907	0.1462	-0.1858	0.2549	0.2049	0.1580	-0.0473	0.9709	0.0083	0.9911	0.0556	0.9023
Ucp4C	Ucp4C	1632843_at	-0.1105	0.8727	0.2213	0.2114	0.1637	0.5545	-0.3613	0.6566	-0.3245	0.4218	0.0368	0.9395	-0.0184	0.9939	0.1334	0.8664	0.1518	0.8307
CG12004	anon-fast-evolving	1632844_s_at	0.6267	0.0208	0.5037	0.0444	0.5855	0.0149	0.0286	0.9683	0.1706	0.3741	0.1421	0.4192	0.0134	0.9939	0.1426	0.7544	0.1292	0.7749
Hr39	Hormone receptor	1632845_at	0.4045	0.0206	0.2608	0.1671	1.2921	0.0001	0.0739	0.8899	0.0263	0.9167	-0.0476	0.8135	-0.5856	0.1628	-0.0818	0.7456	0.5038	0.0559
alpha4GT1	alpha4GT1	1632846_at	0.1665	0.4539	0.0338	0.8699	0.1187	0.4913	-0.1471	0.6177	0.0651	0.7011	0.2122	0.1059	-0.2982	0.7220	-0.1069	0.8187	0.1913	0.6212
CG13484	CG13484	1632847_at	-0.1886	0.4544	0.0914	0.6325	-0.1616	0.3592	-0.0228	0.9741	0.0738	0.7066	0.0966	0.5661	0.1234	0.8870	0.2827	0.3758	0.1594	0.6491
nmdyn-D6	nmdyn-D6	1632848_at	0.3968	0.0648	0.6507	0.0348	0.8015	0.0013	-0.1271	0.8053	-0.3035	0.1381	-0.1764	0.3499	-0.3493	0.4866	-0.1759	0.4848	0.1735	0.4970
nol	no optic lobe	1632849_at	0.2483	0.1808	-0.0669	0.6123	-0.0372	0.8625	-0.0576	0.9311	0.1457	0.4778	0.2033	0.2485	-0.0373	0.9657	-0.0529	0.8746	-0.0156	0.9640
CG7338 /// DmirCG7338	CG7338	1632850_at	0.0890	0.7509	0.1156	0.8510	0.5039	0.0061	0.3919	0.4909	0.2999	0.3296	-0.0920	0.7865	-0.0479	0.9831	0.3119	0.5753	0.3598	0.5120
---	---	1632851_at	0.2454	0.2193	0.2557	0.1547	-0.0661	0.7439	-0.1478	0.6125	-0.0512	0.7717	0.0966	0.4917	0.2188	0.7746	0.0888	0.8431	-0.1300	0.7320
Gs1	glutamine synthet	1632852_s_at	0.9600	0.0463	0.4043	0.4240	0.8181	0.0087	-0.1887	0.8948	-0.0673	0.9197	0.1215	0.8216	-0.7350	0.5126	-0.6757	0.1900	0.0593	0.9365
---	---	1632853_at	-0.0876	0.7567	-0.0624	0.7078	0.0725	0.6758	0.0129	0.9922	0.0653	0.8522	0.0524	0.8682	-0.0330	0.9691	-0.0042	0.9935	0.0288	0.9254
Prm	paramyosin	1632854_s_at	-2.7038	0.0053	-2.0607	0.0074	-2.6808	0.0006	-0.2813	0.6854	-1.3839	0.0019	-1.1027	0.0031	0.4044	0.8882	-0.6067	0.5994	-1.0111	0.3636
CG7352	CG7352	1632855_at	-0.1029	0.7125	0.0227	0.8880	-0.3751	0.0374	-0.2928	0.5008	-0.0604	0.8380	0.2323	0.2614	-0.0140	0.9940	-0.0569	0.9341	-0.0429	0.9433
CG33528	CG33528	1632856_s_at	0.3607	0.0393	-0.0252	0.8120	0.0114	0.9518	-0.0407	0.9451	0.0230	0.9223	0.0637	0.7231	-0.0756	0.9092	-0.1451	0.5541	-0.0695	0.8111
CG30088	CG30088	1632857_at	-0.1196	0.4840	-0.1201	0.4524	0.1280	0.4378	0.0941	0.8257	0.1839	0.2681	0.0898	0.5870	-0.1658	0.7697	0.0810	0.7883	0.2467	0.3171
Xe7	Xe7	1632858_a_at	-0.2340	0.3287	-0.0288	0.9409	0.0222	0.9133	0.0594	0.9300	0.0343	0.8997	-0.0252	0.9174	-0.0299	0.9831	0.1602	0.6497	0.1901	0.5820
fru	fru-satori	1632859_a_at	-0.0030	0.9936	0.2059	0.1442	0.2696	0.2010	0.1129	0.9225	-0.1174	0.7818	-0.2303	0.4920	-0.0812	0.9030	0.0292	0.9404	0.1104	0.6787
Cpr64Aa	CG15006	1632860_at	0.0919	0.5439	0.0122	0.9181	-0.1883	0.2694	0.0118	0.9893	0.0301	0.9144	0.0183	0.9412	-0.0055	0.9952	-0.0383	0.9120	-0.0329	0.9164
msta	msta	1632861_at	-1.6314	0.0035	-0.0302	0.8986	-0.2184	0.2353	-0.3116	0.5496	-1.5937	0.0005	-1.2821	0.0008	0.0702	0.9460	-0.0530	0.9129	-0.1233	0.7333
---	---	1632862_at	0.1434	0.4152	0.0527	0.5901	0.2388	0.1815	0.0584	0.8987	0.0950	0.5567	0.0366	0.8301	-0.0744	0.9306	0.1558	0.5967	0.2302	0.4114
---	---	1632863_at	-0.1578	0.4148	0.0595	0.5765	0.2316	0.1066	-0.0098	0.9924	-0.1940	0.3749	-0.1842	0.3484	0.0970	0.8480	0.1043	0.6496	0.0073	0.9838
CG13258	CG13258	1632864_at	0.1752	0.5545	0.2525	0.3636	-0.1210	0.5193	-0.0567	0.9412	-0.0889	0.7299	-0.0322	0.9031	0.1477	0.8472	-0.0978	0.8132	-0.2455	0.4572
CG12974	CG12974	1632865_s_at	3.3075	0.0161	3.6422	0.0042	4.2426	0.0000	1.5654	0.3840	0.6182	0.5741	-0.9472	0.3053	1.0199	0.4415	1.1831	0.0950	0.1632	0.8551
---	---	1632866_at	0.0822	0.7255	0.1755	0.4010	-0.0474	0.8582	0.0930	0.8140	0.0873	0.6110	-0.0057	0.9766	0.1852	0.8270	0.1305	0.7558	-0.0546	0.9104
---	---	1632867_at	-0.0220	0.9036	0.3702	0.1988	-0.2020	0.2541	-0.2836	0.3909	-0.3390	0.0719	-0.0554	0.7908	0.5064	0.3500	0.2336	0.4107	-0.2728	0.3553
wg	Wingless	1632868_a_at	-0.8962	0.0020	-0.1961	0.4477	-0.5998	0.0148	-0.3980	0.3359	-0.6249	0.0171	-0.2269	0.2947	-0.1734	0.7644	-0.1699	0.4997	0.0035	0.9935
CG32532	CG32532	1632869_at	0.4466	0.0478	-0.0025	0.9940	0.1055	0.6120	0.0094	0.9937	0.1568	0.5479	0.1474	0.5336	-0.2087	0.7997	-0.2143	0.5491	-0.0056	0.9924
CG32364	CG32364	1632870_at	-1.4373	0.0030	-2.3011	0.0052	-2.0075	0.0003	0.0103	0.9949	0.6776	0.0450	0.6672	0.0307	-0.1351	0.9238	-0.0743	0.9192	0.0608	0.9255
---	---	1632871_at	0.0037	0.9929	0.0039	0.9739	0.0610	0.7606	-0.0155	0.9857	-0.1558	0.4356	-0.1403	0.4389	0.1180	0.8972	-0.1807	0.6215	-0.2987	0.3898
---	---	1632872_at	0.7427	0.0948	-0.0951	0.4951	0.3751	0.0965	0.0824	0.8671	0.1288	0.4795	0.0464	0.8132	-0.3330	0.7997	-0.5372	0.3128	-0.2042	0.7456
MtnA	metallothionein N	1632873_at	-0.4883	0.0109	0.2460	0.4187	-0.0754	0.8152	-0.3592	0.4500	-1.1052	0.0019	-0.7460	0.0067	-0.1083	0.9342	-0.3789	0.3575	-0.2705	0.5420
CG6602	CG6602	1632874_at	-0.7207	0.4331	-0.0991	0.6020	0.0196	0.9246	0.0527	0.9860	-0.8868	0.2138	-0.9394	0.1404	0.0183	0.9939	-0.3909	0.4915	-0.4092	0.4755
Grip71	Grip71	1632875_at	-0.6883	0.0650	-0.9452	0.0421	-1.0038	0.0341	-0.2839	0.6041	0.4899	0.0777	0.7739	0.0077	-0.5330	0.7689	-0.0184	0.9923	0.5146	0.5251
CG4896	CG4896	1632876_a_at	-0.0965	0.7466	-0.6796	0.0711	-0.7493	0.0042	-0.1583	0.7724	0.5890	0.0187	0.7473	0.0039	-0.0497	0.9792	-0.0091	0.9928	0.0406	0.9471
CG2906	CG2906	1632877_a_at	0.5945	0.0183	0.4573	0.3079	0.3184	0.2121	0.2593	0.5735	0.3212	0.1721	0.0619	0.8172	0.3907	0.7464	0.1939	0.7466	-0.1969	0.7378
CG14252	CG14252	1632878_at	-0.0019	0.9944	0.0214	0.9633	-0.1057	0.5935	-0.3885	0.5863	0.0265	0.9612	0.4150	0.2013	-0.1114	0.8870	0.0672	0.8725	0.1786	0.5659
---	---	1632879_at	0.1769	0.2879	-0.0891	0.5843	-0.0355	0.8561	-0.1322	0.6763	0.0240	0.9077	0.1563	0.2424	-0.0492	0.9567	0.0259	0.9505	0.0751	0.8168
CG13073	CG13073	1632880_a_at	0.0389	0.8078	0.0730	0.6025	0.0415	0.8425	-0.1862	0.5067	-0.1064	0.4954	0.0798	0.5912	-0.1183	0.8901	0.0536	0.9141	0.1718	0.6188
---	---	1632881_at	-0.0967	0.5779	-0.7609	0.0622	-0.9457	0.1065	-0.2719	0.8628	0.5746	0.3096	0.8465	0.0909	-0.0058	0.9950	0.0385	0.9036	0.0443	0.8745
CG11671	CG11671	1632882_at	0.5368	0.2295	0.1874	0.4088	0.4595	0.1804	0.1957	0.7255	-0.0656	0.8391	-0.2613	0.2454	0.1019	0.9721	-0.2798	0.7485	-0.3817	0.6318
CG31467	CG31467	1632883_at	-0.1512	0.5519	-0.0166	0.9610	0.3482	0.1433	-0.0542	0.9610	-0.3213	0.2819	-0.2672	0.3240	-0.3916	0.6935	-0.1935	0.6673	0.1981	0.6533
CG33967	CG33967	1632884_at	0.0289	0.9120	0.6600	0.0910	0.5487	0.0517	-0.0443	0.9558	-0.5216	0.0231	-0.4773	0.0210	0.1240	0.9238	0.0762	0.9085	-0.0477	0.9371
Msi	male-specific IDG	1632885_at	0.3611	0.0866	0.1794	0.5201	0.1401	0.3647	-0.0227	0.9774	0.0104	0.9729	0.0330	0.8862	-0.0144	0.9914	-0.0980	0.7971	-0.0836	0.8247
CG13247	CG13247	1632886_at	-1.4706	0.0029	-2.0274	0.0084	-2.6188	0.0005	-0.2754	0.7929	1.2169	0.0117	1.4923	0.0028	0.1418	0.9016	0.1864	0.6939	0.0447	0.9395
---	---	1632887_at	0.0659	0.7933	-0.1136	0.3868	-0.1756	0.2708	0.0336	0.9759	0.2580	0.3555	0.2244	0.3760	0.0737	0.9095	0.1586	0.5068	0.0849	0.7499
CG31687 /// CG31688	CG31688 /// CG3	1632888_s_at	0.0170	0.9478	0.0075	0.9755	0.2275	0.2450	-0.0391	0.9603	-0.1160	0.6111	-0.0769	0.7279	-0.0708	0.9527	0.0305	0.9590	0.1014	0.8204
dpr9	dpr9	1632889_at	0.0242	0.8989	-0.2109	0.2634	-0.0039	0.9891	0.0035	0.9956	0.1707	0.2958	0.1672	0.2498	-0.1230	0.8609	-0.0391	0.9306	0.0838	0.8098
Sobp	Sine oculis-bindin	1632890_at	-1.7534	0.0098	-0.4396	0.3618	-1.0845	0.0013	-0.1978	0.8444	-0.4890	0.1856	-0.2912	0.3999	0.1746	0.9101	0.1658	0.8172	-0.0089	0.9924
Hrb27C	linha	1632891_s_at	-0.1120	0.8043	1.0527	0.0901	0.6375	0.0084	-0.1856	0.8064	-0.4200	0.1610	-0.2345	0.4033	0.3696	0.8298	0.8169	0.2408	0.4473	0.5587
---	---	1632892_at	-0.0011	0.9955	0.0674	0.5514	0.1929	0.2111	0.1693	0.5633	0.0078	0.9734	-0.1615	0.2308	-0.0154	0.9831	-0.0324	0.8995	-0.0169	0.9433
CG14116	CG14116	1632893_at	0.0491	0.8888	0.1594	0.4355	-0.1127	0.5444	-0.1268	0.7579	-0.1583	0.3811	-0.0315	0.8803	0.1936	0.7981	0.1465	0.6765	-0.0471	0.9147
CG12121	CG12121	1632894_at	0.2639	0.2750	-0.0903	0.7810	-0.1455	0.3416	-0.0619	0.9376	0.7352	0.0073	0.7971	0.0030	0.1486	0.8930	0.4410	0.2671	0.2924	0.4925
CG31146	CG31146	1632895_at	-0.5195	0.0230	-0.4048	0.0235	-0.1590	0.3402	0.4662	0.2463	0.2607	0.2761	-0.2055	0.3474	-0.2372	0.7187	0.0500	0.9048	0.2872	0.3032
CG18166	CG18166	1632896_at	0.1802	0.5611	-0.3685	0.3087	-0.6027	0.0251	-0.2524	0.7795	0.2288	0.5720	0.4812	0.1497	0.1070	0.8564	-0.0151	0.9701	-0.1221	0.6392
Golgin84	Golgin84	1632897_at	0.1599	0.3986	0.7713	0.0288	0.7335	0.0028	-0.0304	0.9677	-0.4467	0.0312	-0.4163	0.0266	0.0731	0.9449	0.4045	0.1920	0.3313	0.3089
CG33978	CG33978	1632898_at	-0.8990	0.0591	-0.5615	0.2684	-1.4956	0.0016	-0.6345	0.1403	-0.0430	0.9045	0.5915	0.0192	0.2678	0.8740	0.0308	0.9796	-0.2370	0.7562
CG6986	CG6986	1632899_s_at	0.0732	0.7829	0.0610	0.6541	0.0479	0.7862	-0.0276	0.9745	-0.0295	0.9214	-0.0019	0.9945	0.0537	0.9308	-0.0906	0.6844	-0.1443	0.4851
CG34346	CG11072	1632900_at	-0.2210	0.3496	-0.1806	0.2315	-0.1741	0.3545	-0.0465	0.9466	0.0835	0.7135	0.1301	0.4891	-0.0004	0.9999	0.1541	0.6794	0.1545	0.6734
c(3)G	crossover suppre	1632901_s_at	0.4453	0.5467	0.0512	0.8396	-0.4820	0.0774	-0.5808	0.5597	0.5705	0.2669	1.1513	0.0200	-0.0258	0.9946	0.1480	0.9156	0.1739	0.8881
---	---	1632902_s_at	0.0633	0.8639	-1.3524	0.0247	-1.0171	0.0418	0.0363	0.9860	1.2115	0.0231	1.1752	0.0164	-0.2885	0.8012	-0.1815	0.7459	0.1071	0.8650
---	---	1632903_at	0.1155	0.4298	-0.2866	0.0843	-0.3538	0.1092	0.2935	0.2501	0.6574	0.0019	0.3639	0.0155	0.2354	0.6960	0.1923	0.4520	-0.0431	0.9025
CG10354	CG10354	1632904_at	0.3402	0.2496	-0.0482	0.8936	0.1133	0.4895	0.2309	0.5455	0.5544	0.0127	0.3235	0.0702	0.0550	0.9761	0.2790	0.5459	0.2240	0.6389
---	---	1632905_at	0.2085	0.5152	0.2963	0.1050	0.1410	0.3482	-0.1056	0.8725	-0.2034	0.3854	-0.0978	0.6832	-0.0015	0.9994	-0.0833	0.8262	-0.0817	0.8209
---	---	1632906_at	0.1194	0.6225	0.1299	0.5489	-0.0210	0.9087	0.0289	0.9761	0.0071	0.9839	-0.0217	0.9416	-0.0008	0.9998	-0.0033	0.9953	-0.0025	0.9956

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13204	CG13204	1632907_a_at	-1.2189	0.0027	-0.5052	0.0836	-0.9032	0.0089	0.1275	0.8342	-0.1204	0.6422	-0.2479	0.2306	0.1181	0.8846	0.0404	0.9350	-0.0777	0.8444
Lerp	lysosomal enzyme	1632908_s_at	1.0458	0.0195	1.1679	0.0286	1.3738	0.0001	-0.1362	0.7857	-0.7286	0.0046	-0.5925	0.0070	-0.3322	0.8049	-0.5216	0.3416	-0.1894	0.7754
---	---	1632909_at	0.0360	0.8389	-0.0141	0.9562	0.2666	0.0667	0.1190	0.8385	-0.0667	0.8041	-0.1857	0.3559	-0.0324	0.9621	-0.0520	0.8386	-0.0196	0.9414
CG7512	CG7512	1632910_at	0.1411	0.7169	0.2425	0.1513	0.2471	0.1400	-0.1634	0.8649	-0.3767	0.2708	-0.2133	0.5186	0.0186	0.9898	-0.1651	0.6265	-0.1837	0.5834
CG33493	CG33493	1632911_at	2.2048	0.0012	0.9840	0.2788	2.2343	0.0002	0.6136	0.2969	0.0525	0.9134	-0.5611	0.0717	-0.3164	0.8873	-0.8774	0.2775	-0.5610	0.5211
CG13667	CG13667	1632912_s_at	-0.0992	0.5300	0.0155	0.9713	0.1778	0.3686	0.0233	0.9693	0.1373	0.3926	0.1140	0.4389	-0.1409	0.8603	0.1364	0.7072	0.2774	0.3967
CG10959	CG10959	1632913_at	-0.1079	0.6186	-0.1130	0.5825	0.1979	0.3479	0.2381	0.5766	0.6984	0.0071	0.4603	0.0267	-0.3317	0.6824	0.2544	0.4471	0.5862	0.1156
l(2)dtl	lethal-(2)-denticlel	1632914_at	-0.2190	0.3209	-0.1985	0.7140	-0.7111	0.0543	-0.4817	0.1551	-0.1532	0.4754	0.3285	0.0751	0.1379	0.9635	-0.0305	0.9844	-0.1684	0.8800
---	---	1632915_s_at	0.5352	0.2248	0.0153	0.9840	-1.4299	0.0071	-0.7467	0.2506	1.1655	0.0093	1.9123	0.0006	0.6636	0.7423	0.5003	0.5800	-0.1634	0.8891
CG31357	CG31357	1632916_at	-0.5474	0.0318	-0.4491	0.0505	-0.4835	0.0551	-0.2682	0.5353	-0.7511	0.0059	-0.4829	0.0248	-0.2300	0.8049	-0.6778	0.0897	-0.4478	0.2622
beta4GalT7	glycosaminoglyca	1632917_at	-0.0609	0.8731	-0.1315	0.6026	-0.1054	0.7287	-0.1193	0.8211	0.4575	0.0334	0.5768	0.0077	-0.1000	0.9555	0.3021	0.5852	0.4021	0.4564
CG18746	CG18746	1632918_at	0.1609	0.3580	0.0531	0.6286	-0.2475	0.2724	-0.1489	0.6820	0.0747	0.7015	0.2236	0.1375	0.0323	0.9771	-0.0210	0.9620	-0.0533	0.8836
mats	mob as tumor sup	1632919_at	0.4950	0.0846	0.6716	0.1144	0.7412	0.0070	0.0532	0.9311	-0.2054	0.2595	-0.2586	0.1101	-0.0857	0.9589	0.0958	0.8903	0.1815	0.7410
CG10700	CG10700	1632920_at	0.0289	0.8926	-0.0111	0.9220	-0.0250	0.8992	-0.1081	0.8020	-0.0881	0.6507	0.0200	0.9252	0.0221	0.9857	-0.1211	0.7002	-0.1433	0.6328
---	---	1632921_at	-0.5267	0.1456	-0.5101	0.0847	-0.0986	0.6566	-0.1394	0.8297	-0.4361	0.0805	-0.2968	0.1802	-0.1489	0.7953	-0.2553	0.2714	-0.1064	0.6886
NFAT	Misexpression Su	1632922_at	-0.0764	0.9127	0.5190	0.1247	0.5048	0.0520	-0.1948	0.5335	-0.4461	0.0142	-0.2513	0.0860	-0.1683	0.9467	0.1769	0.8721	0.3452	0.6861
CG14661	CG14661	1632923_at	-0.9968	0.1474	-1.8307	0.0595	-1.9800	0.0000	0.2897	0.6835	0.1510	0.6905	-0.1387	0.6893	0.4317	0.8627	-0.5681	0.5959	-0.9998	0.3307
rho-4	rhomboid-4	1632924_at	-0.1373	0.4258	-0.5777	0.0275	-0.3017	0.4334	0.4145	0.5539	0.6294	0.0832	0.2149	0.5446	-0.1416	0.8049	0.0404	0.9125	0.1820	0.4620
Gr57a	Gustatory recepto	1632925_at	-0.0059	0.9832	-0.2178	0.1435	-0.0794	0.6882	0.1442	0.7835	0.2520	0.2454	0.1079	0.6248	-0.0042	0.9967	-0.0181	0.9663	-0.0138	0.9727
CG33639	CG33639	1632926_at	0.5105	0.0376	-0.0847	0.5262	0.0376	0.8823	0.0680	0.9133	0.1507	0.4635	0.0828	0.6893	-0.0242	0.9862	-0.1620	0.6430	-0.1378	0.7027
CG8359	CG8359	1632927_at	0.4927	0.0316	0.2714	0.3283	-0.1626	0.4149	-0.1382	0.8111	0.2512	0.2760	0.3894	0.0614	0.2925	0.6824	0.0255	0.9587	-0.2670	0.3782
CG9523	CG9523	1632928_at	0.9822	0.0128	1.0069	0.0402	0.9599	0.0013	0.4031	0.4117	1.1215	0.0023	0.7184	0.0101	0.4247	0.6749	1.1364	0.0290	0.7116	0.1238
---	---	1632929_at	0.2579	0.1559	0.0693	0.7745	0.2053	0.2366	0.0259	0.9777	0.0327	0.9189	0.0068	0.9808	-0.0507	0.9552	-0.1087	0.7198	-0.0579	0.8694
CG14506	CG14506	1632930_at	0.0234	0.9173	0.0455	0.7679	-0.0481	0.7652	-0.0592	0.9039	-0.0821	0.6475	-0.0229	0.9054	0.1214	0.8479	0.1245	0.6684	0.0031	0.9947
cta	concertina	1632931_at	0.0117	0.9641	-0.0455	0.8047	-0.6095	0.0137	-0.1657	0.7358	0.4513	0.0445	0.6170	0.0073	0.2793	0.7062	0.2964	0.3255	0.0171	0.9714
CG33171	CG33171	1632932_a_at	-2.7780	0.0022	-2.8170	0.0198	-3.2939	0.0002	0.0692	0.9351	0.3385	0.1748	0.2692	0.2306	0.6076	0.8016	0.5158	0.6380	-0.0919	0.9486
CG6865	CG6865	1632933_at	0.0742	0.8012	0.3247	0.2534	0.1589	0.4925	-0.0142	0.9917	0.0159	0.9711	0.0301	0.9313	0.1273	0.8943	0.1185	0.7930	-0.0088	0.9874
---	---	1632934_at	-0.0533	0.7443	0.0495	0.7201	0.2386	0.1714	0.0772	0.8676	-0.0927	0.6054	-0.1700	0.2470	-0.1234	0.8940	0.1044	0.8172	0.2278	0.5335
CG13300	CG13300	1632935_at	0.2089	0.4578	0.2629	0.3747	-0.0273	0.8853	-0.1275	0.7349	0.1040	0.5649	0.2314	0.1205	0.2647	0.5236	0.2010	0.3000	-0.0637	0.7891
alpha-Man-IIb	alpha-mannosidas	1632936_at	0.2847	0.4386	-0.5141	0.1330	-0.9616	0.0028	-0.5913	0.2187	0.6491	0.0318	1.2404	0.0012	-0.0742	0.9538	-0.1264	0.7897	-0.0522	0.9198
CadN2	DN-cadh-like	1632937_at	0.2163	0.3224	0.3052	0.0651	0.2033	0.3264	-0.0937	0.8939	-0.0611	0.8391	0.0326	0.9084	0.0771	0.9467	-0.0550	0.9205	-0.1321	0.7464
Obp47a	Odorant-binding p	1632938_at	0.1916	0.1653	-0.0014	0.9914	0.1431	0.4443	0.0376	0.9436	0.2032	0.1767	0.1655	0.2213	-0.1157	0.8049	-0.0273	0.9291	0.0884	0.6800
shep	alan shepard	1632939_s_at	0.1642	0.4764	0.3248	0.0550	0.3024	0.1422	0.0503	0.9345	-0.1270	0.5057	-0.1773	0.2774	0.1630	0.7644	0.0222	0.9526	-0.1408	0.5672
pip	pipe	1632940_a_at	0.1652	0.3758	0.0234	0.8202	0.2709	0.0972	0.0259	0.9782	-0.1682	0.5103	-0.1941	0.3871	-0.1448	0.8062	-0.1541	0.5500	-0.0093	0.9829
---	---	1632941_at	-0.0477	0.7578	0.1584	0.1677	0.0901	0.6824	-0.1932	0.6473	-0.0693	0.7795	0.1239	0.5349	-0.1440	0.7220	-0.0040	0.9924	0.1400	0.4243
CG13060	CG13060	1632942_at	0.0183	0.9211	-0.0659	0.5267	0.0466	0.7766	-0.0165	0.9777	0.0248	0.9045	0.0413	0.8061	-0.1272	0.8236	-0.0186	0.9622	0.1086	0.6785
bru-3	bruno-3	1632943_a_at	-2.1046	0.0004	-1.5123	0.1907	-2.5671	0.0001	-0.7027	0.0174	-0.9611	0.0004	-0.2584	0.0640	0.2559	0.9296	-0.2212	0.8677	-0.4771	0.6375
CG31488	CG31488	1632944_at	-0.0349	0.8816	-0.0517	0.7042	0.0896	0.6204	0.0357	0.9504	0.0769	0.6702	0.0412	0.8185	-0.1518	0.7932	-0.1057	0.7012	0.0461	0.8913
Msp-300	nesprin	1632945_at	-0.4903	0.2494	-0.6533	0.1608	-1.9722	0.0092	-0.2928	0.5932	0.8723	0.0078	1.1651	0.0013	1.0880	0.6749	0.8573	0.4164	-0.2307	0.8707
Stlk	Ste20-like kinase	1632946_at	-0.6755	0.0353	-0.5509	0.0811	-0.8656	0.0010	-0.2419	0.7278	-0.1301	0.7194	0.1118	0.7384	0.0246	0.9856	-0.1279	0.7183	-0.1526	0.6473
axo	axotactin	1632947_at	-0.7368	0.0071	-0.1937	0.5145	-0.9089	0.0022	-0.2040	0.7631	-0.3393	0.2430	-0.1353	0.6519	0.2117	0.7726	0.1247	0.7346	-0.0871	0.8256
CG32241	CG32241	1632948_x_at	0.2046	0.4049	0.0917	0.5012	0.3477	0.0793	-0.0116	0.9942	-0.0954	0.8160	-0.0838	0.8238	-0.0723	0.9223	-0.0522	0.8830	0.0201	0.9538
Rad1	Rad1	1632949_at	0.1686	0.6294	0.0742	0.6053	-0.3227	0.0554	-0.6641	0.2500	0.0970	0.8202	0.7611	0.0211	-0.2140	0.7953	-0.0043	0.9956	0.2097	0.5682
Or10a	Odorant receptor	1632950_at	0.2011	0.2720	-0.1147	0.3823	0.1413	0.3934	0.0546	0.9300	0.1197	0.5410	0.0651	0.7403	-0.4106	0.3517	-0.3538	0.1345	0.0568	0.8524
---	---	1632951_at	-0.0553	0.7563	0.1594	0.4501	-0.1075	0.5148	-0.0591	0.9295	-0.1817	0.3625	-0.1226	0.5177	0.0568	0.9467	-0.0010	0.9994	-0.0577	0.8648
CG4598	CG4598	1632952_at	0.6880	0.0340	0.3201	0.2672	0.6346	0.0054	-0.1851	0.6366	-0.2923	0.1273	-0.1072	0.5758	-0.4153	0.6557	-0.5599	0.1382	-0.1446	0.7416
CG13250	CG13250	1632953_at	-0.9780	0.0035	-1.2395	0.0032	-1.2956	0.0032	0.3777	0.5647	0.7611	0.0312	0.3834	0.2023	0.2285	0.8215	0.4350	0.2871	0.2065	0.6485
JTBR	anon-fast-evolving	1632954_at	-0.2065	0.4157	-0.1913	0.3514	-0.3373	0.0432	0.0309	0.9619	-0.0861	0.6556	-0.1170	0.4719	0.1436	0.8882	-0.0840	0.8787	-0.2277	0.5785
PhKgamma	phosphorylase kin	1632955_at	1.7457	0.0066	-0.0619	0.9137	0.2184	0.2787	-0.0100	0.9950	0.4184	0.1901	0.4284	0.1340	-0.3646	0.8331	-1.4370	0.0661	-1.0724	0.1618
CG3021	Transcription unit	1632956_at	0.2005	0.2356	0.3629	0.1896	0.2266	0.1810	0.0374	0.9567	0.2097	0.2591	0.1723	0.3053	0.2187	0.7363	0.4103	0.1374	0.1916	0.5045
---	---	1632957_s_at	-0.0387	0.8401	-0.0572	0.6065	0.0561	0.8516	0.1844	0.5931	0.1616	0.3654	-0.0228	0.9148	-0.0869	0.9441	0.0580	0.9209	0.1449	0.7373
CG15675	CG15675	1632958_a_at	0.0196	0.9576	0.0724	0.9086	0.2387	0.5231	0.4978	0.0711	0.6877	0.0019	0.1900	0.1819	0.4578	0.8222	0.9202	0.2582	0.4624	0.6123
CG8854	CG8854	1632959_at	-0.6574	0.0448	0.2156	0.3217	-0.0323	0.9030	-0.1829	0.8280	-1.0667	0.0061	-0.8838	0.0086	0.0206	0.9893	-0.1654	0.6310	-0.1859	0.5834
Eh	Eclosion hormone	1632960_at	0.0579	0.7318	0.0283	0.8041	0.0487	0.8114	-0.0294	0.9619	-0.0015	0.9947	0.0279	0.8856	0.0560	0.9398	-0.0668	0.8218	-0.1229	0.6154
Shaw	Shaker cognate w	1632961_at	-1.2987	0.0107	-2.2672	0.0075	-2.4463	0.0000	0.0597	0.9570	1.1791	0.0027	1.1194	0.0020	0.0717	0.9357	-0.1218	0.7027	-0.1935	0.5100
RpL36A	Ribosomal protein	1632962_at	0.4482	0.0270	0.7124	0.0200	1.0149	0.0019	0.1521	0.7625	-0.2614	0.2257	-0.4136	0.0388	0.0840	0.8889	-0.0233	0.9496	-0.1074	0.6621
CG4963	anon-fast-evolving	1632963_at	-0.7848	0.0250	-1.0135	0.0445	-1.4934	0.0001	-0.2031	0.5854	0.3776	0.0515	0.5807	0.0052	0.1797	0.8767	0.0139	0.9884	-0.1658	0.7495
CG15678	CG15678	1632964_at	-0.5987	0.6084	-2.3516	0.0107	-2.1697	0.0003	0.7496	0.7149	3.6170	0.0031	2.8674	0.0052	0.5664	0.7810	1.7407	0.0665	1.1743	0.2054
CG13251	CG13251	1632965_at	0.1838	0.4259	-0.1374	0.3679	0.0149	0.9645	0.2039	0.7305	0.3809	0.1485	0.1770	0.4839	-0.1003	0.9365	-0.0767	0.8981	0.0235	0.9695
CG8213	CG8213	1632966_at	0.0397	0.8030	0.0495	0.6715	0.0394	0.8596	-0.1009	0.8507	-0.0416	0.8723	0.0592	0.7865	-0.0624	0.9142	-0.0666	0.7932	-0.0042	0.9902
M6	M6	1632967_s_at	-0.1789	0.6733	1.0201	0.1546	1.3319	0.0002	-0.1709	0.8589	-1.3814	0.0027	-1.2105	0.0028	-0.2830	0.8378	-0.1095	0.8981	0.1735	0.8015
CG6053	CG6053	1632968_at	0.1707	0.2265	-0.0596	0.6895	-0.0011	0.9959	0.1587	0.6015	0.1771	0.2499	0.0184	0.9223	0.0441	0.9577	-0.0475	0.8893	-0.0917	0.7333
ACXA	adenylyl cyclase	1632969_at	0.1908	0.3176	0.0134	0.9000	0.3428	0.0897	0.1156	0.7679	0.1337	0.4384	0.0181	0.9293	-0.0814	0.9238	-0.0802	0.8294	0.0012	0.9982
CG15336	CG15336	1632970_at	0.1156	0.6116	0.2601	0.1576	0.2641	0.1058	-0.0302	0.9624	-0.0759	0.6986	-0.0457	0.8116	0.0997	0.8951	0.0238	0.9590	-0.0759	0.8307
CG12895	CG12895	1632971_at	-0.4606	0.1495	0.4740	0.0574	0.2699	0.1363	-0.1626	0.6457	-1.0185	0.0005	-0.8559	0.0006	0.1307	0.8930	-0.0173	0.9789	-0.1480	0.7175
Nep3	Neprilysin 3	1632972_at	-1.7765	0.0069	-0.3304	0.4134	-1.2361	0.0083	-0.8977	0.1751	-1.4992	0.0035	-0.6015	0.0935	-0.1714	0.9309	-0.2574	0.7305	-0.0861	0.9231
mod(mdg4)	Modifier67.2	1632973_at	-0.1234	0.6252	0.0665	0.8680	-0.4083	0.0265	-0.1629	0.7415	-0.1146	0.6381	0.0482	0.8483	0.1368	0.9066	0.1364	0.7962	-0.0004	0.9995
CG30015	CG30015	1632974_s_at	0.2202	0.6775	-0.3774	0.5197	-0.5193	0.1058	0.2157	0.7915	0.0003	0.9994	-0.2153	0.4999	0.2753	0.9206	-0.5235	0.5973	-0.7988	0.3994
CG4398	CG4398	1632975_at	-2.3965	0.0400	-3.2982	0.0052	-3.8857	0.0000	-0.3183	0.5398	-0.2840	0.2982	0.0343	0.9175	-0.0683	0.9913	-1.4800	0.2430	-1.4117	0.2964

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1632976_s_at	0.1876	0.2614	0.1214	0.4227	0.1182	0.6256	-0.0729	0.9223	0.0231	0.9447	0.0959	0.6850	-0.1739	0.7464	-0.0036	0.9941	0.1703	0.4655
Poxn	pox-neuro	1632977_at	-0.0290	0.8631	0.0273	0.7968	0.2049	0.1564	0.1379	0.6454	0.0248	0.9013	-0.1131	0.4033	-0.0312	0.9611	-0.0482	0.8407	-0.0170	0.9460
CG32207	CG32207	1632978_at	-1.6259	0.0276	-1.2070	0.1820	-2.0491	0.0007	-0.3869	0.5068	-0.6415	0.0440	-0.2546	0.3730	0.4199	0.8541	-0.4268	0.6849	-0.8467	0.3807
CG11784	CG11784	1632979_at	-0.3448	0.3193	-0.0001	1.0000	0.2180	0.1886	0.2152	0.7290	-0.4439	0.1087	-0.6591	0.0160	-0.0330	0.9892	-0.0315	0.9702	0.0015	0.9990
CG3902	CG3902	1632980_at	0.2553	0.5681	0.0986	0.8951	0.4374	0.0222	0.1351	0.8028	-0.3507	0.1050	-0.4858	0.0205	-0.2072	0.9449	-0.5325	0.5800	-0.3253	0.7560
kl-2	male fertility factor	1632981_s_at	0.0148	0.9589	0.0692	0.6675	0.2770	0.1033	0.1244	0.7939	0.0598	0.8023	-0.0647	0.7612	0.0065	0.9946	0.0926	0.7213	0.0861	0.7422
CG31021	CG31021	1632982_at	0.0968	0.5499	0.0163	0.9549	0.2875	0.1014	-0.0261	0.9705	-0.0160	0.9528	0.0101	0.9650	-0.0787	0.9246	0.0516	0.9037	0.1304	0.6677
l(3)82Fd	late puff gene at 8	1632983_s_at	-1.0892	0.0032	0.2822	0.4595	-0.4337	0.0608	-0.5038	0.2789	-0.8619	0.0077	-0.3581	0.1424	0.3439	0.7644	0.4264	0.3685	0.0826	0.9020
CG6503 /// DyakCG6503	putative noncodin	1632984_s_at	2.9661	0.0019	1.5340	0.1380	2.8132	0.0002	1.2431	0.2471	0.5582	0.3925	-0.6849	0.2294	-0.1407	0.9653	-0.9251	0.2669	-0.7844	0.3764
CG1893	CG1893	1632985_at	-0.3322	0.0505	-0.8496	0.0314	-1.0998	0.0003	-0.0193	0.9777	0.3913	0.0287	0.4106	0.0148	0.2389	0.7423	0.0183	0.9718	-0.2205	0.4869
CG15088	CG15088	1632986_a_at	0.9000	0.3892	0.2926	0.1310	0.2596	0.2397	0.0786	0.9311	0.0242	0.9517	-0.0544	0.8616	-0.0931	0.9837	-0.6586	0.5469	-0.5655	0.6166
---	---	1632987_at	-0.1118	0.4741	-0.0126	0.9050	-0.1271	0.4560	0.0227	0.9677	-0.0134	0.9517	-0.0361	0.8296	0.1396	0.7726	0.0475	0.8721	-0.0921	0.6870
CG13308 /// DereCG13308	CG13308	1632988_at	-0.0352	0.8380	-0.0329	0.7519	-0.2090	0.3119	0.0918	0.8671	0.0814	0.7181	-0.0104	0.9664	-0.0406	0.9677	-0.1583	0.5698	-0.1176	0.6877
mth	methuselah	1632989_a_at	-0.4054	0.0406	-0.1616	0.6894	-0.5903	0.0042	-0.5240	0.0820	-0.5920	0.0053	-0.0680	0.7129	-0.0953	0.9445	-0.2720	0.5259	-0.1766	0.7039
---	---	1632990_at	0.0483	0.8111	0.1182	0.5728	0.1278	0.4633	-0.1477	0.6811	-0.1244	0.4773	0.0233	0.9086	0.0478	0.9677	-0.0653	0.8846	-0.1131	0.7520
CG1418 /// DereCG1418 ///	CG1418 /// GA12	1632991_at	0.4136	0.1560	1.5039	0.0155	1.5685	0.0001	0.1792	0.6338	-0.0283	0.9124	-0.2075	0.2082	0.0969	0.9474	1.0354	0.0418	0.9385	0.0729
CG11381	CG11381	1632992_at	0.1718	0.3905	-0.0110	0.9515	-0.1599	0.2990	-0.2026	0.5093	0.0507	0.7995	0.2533	0.0825	0.0138	0.9898	-0.1349	0.5634	-0.1488	0.5241
CG6372	CG6372	1632993_at	0.0513	0.7263	-0.1842	0.3996	0.0648	0.7865	0.1528	0.8342	0.2260	0.4301	0.0732	0.8145	-0.0266	0.9852	-0.1472	0.6883	-0.1206	0.7495
CG11000	CG11000	1632994_at	-0.0413	0.8506	0.0827	0.5661	-0.0358	0.8517	0.1478	0.6473	0.0063	0.9792	-0.1415	0.3216	0.1415	0.7597	0.0991	0.6382	-0.0424	0.8699
---	---	1632995_at	0.2555	0.1464	0.1802	0.2507	0.0189	0.9531	-0.0577	0.9302	-0.0091	0.9758	0.0486	0.8251	0.1341	0.8461	-0.0219	0.9629	-0.1561	0.6133
CG34353	CG12274	1632996_at	0.0983	0.7054	0.0022	0.9948	0.0753	0.6890	-0.0114	0.9894	0.1552	0.4511	0.1665	0.3604	-0.0085	0.9929	0.1545	0.5120	0.1630	0.4888
CG34127	CG34127	1632997_at	0.2555	0.1684	0.0785	0.5103	-0.1384	0.4138	-0.1289	0.7400	-0.0175	0.9446	0.1114	0.4984	-0.0570	0.9380	-0.1427	0.5501	-0.0857	0.7442
D12	D12	1632998_at	0.5495	0.0163	-0.0240	0.8430	-0.1881	0.4312	-0.1242	0.7526	0.7214	0.0021	0.8456	0.0006	0.0128	0.9928	0.1183	0.7578	0.1055	0.7815
SIP3	Syntaxin Interactir	1632999_at	1.1570	0.0134	1.1855	0.0852	0.9074	0.0266	-0.2292	0.6854	-0.2042	0.4560	0.0250	0.9393	0.1755	0.9450	0.1261	0.9158	-0.0494	0.9659
CadN	N-cadherin	1633000_a_at	-0.1407	0.4379	0.0104	0.9496	-0.1562	0.4477	0.0101	0.9909	-0.0107	0.9716	-0.0208	0.9305	0.0400	0.9665	-0.0295	0.9425	-0.0694	0.8264
CG2200	CG2200	1633001_at	0.3733	0.1115	0.6964	0.0083	0.9475	0.0003	0.1858	0.6010	-0.1492	0.4201	-0.3350	0.0426	-0.0104	0.9935	0.3055	0.2722	0.3159	0.2869
CG2196	CG2196	1633002_at	-1.0262	0.6170	0.2108	0.7192	0.2893	0.2097	0.3422	0.9598	-2.0980	0.2375	-2.4402	0.1225	0.5016	0.8882	-0.2928	0.8785	-0.7944	0.5767
CG30349	CG30349	1633003_at	0.4941	0.1556	-0.0958	0.8143	0.3202	0.1054	0.3356	0.5735	0.8088	0.0160	0.4731	0.0835	-0.0369	0.9841	0.2381	0.5959	0.2750	0.5359
CG3875	CG3875	1633004_at	0.1309	0.3805	-0.1488	0.3331	0.0790	0.6869	0.1463	0.6654	0.1654	0.3106	0.0191	0.9235	-0.0639	0.9199	-0.1010	0.6713	-0.0371	0.9010
CG6903	CG6903	1633005_at	-0.1059	0.6713	0.3882	0.0369	0.7490	0.0050	-0.0384	0.9451	-1.1569	0.0003	-1.1185	0.0002	-0.3048	0.7533	-0.5297	0.1871	-0.2250	0.6140
nab	nab	1633006_at	0.2433	0.3543	-0.0211	0.9523	0.0362	0.8708	0.0526	0.9441	0.0763	0.7673	0.0237	0.9280	0.0180	0.9898	-0.1319	0.6842	-0.1499	0.6318
---	---	1633007_at	-0.1099	0.5769	-0.2475	0.1998	-0.0039	0.9903	0.0795	0.8835	0.0885	0.6741	0.0090	0.9689	-0.1998	0.8141	-0.1077	0.8132	0.0921	0.8364
Cyp6u1	Cyp6u1	1633008_at	-0.7633	0.0035	-0.8386	0.1303	-0.4287	0.0155	0.0801	0.8908	0.1020	0.6422	0.0220	0.9268	-0.3476	0.7893	-0.1168	0.8831	0.2308	0.7105
CG6640	CG6640	1633009_a_at	0.4497	0.2691	-0.8831	0.0370	0.1716	0.6742	1.1725	0.0233	1.2104	0.0019	0.0379	0.9105	0.1871	0.9305	-0.2404	0.7779	-0.4274	0.5612
CG30382 /// Prosalpha6	CG30382 /// 20s p	1633010_s_at	0.2413	0.3366	0.6208	0.0391	0.8866	0.0016	0.0246	0.9715	-0.1506	0.4013	-0.1752	0.2656	-0.2092	0.8270	0.1465	0.7584	0.3557	0.3828
crp	cropped	1633011_at	-0.4045	0.5509	-0.2338	0.4846	-0.5370	0.1148	-0.1084	0.8776	-0.0077	0.9826	0.1007	0.6913	0.2675	0.9330	0.0878	0.9587	-0.1797	0.8982
CG3662	CG3662	1633012_at	-0.2160	0.2800	-0.1854	0.3191	-0.2664	0.1399	0.0198	0.9774	-0.2095	0.2175	-0.2293	0.1306	0.0658	0.9562	-0.2091	0.5647	-0.2749	0.4400
---	---	1633013_at	0.0854	0.6553	0.0507	0.7065	0.2010	0.2537	-0.0465	0.9436	-0.0310	0.9040	0.0155	0.9475	-0.2122	0.7485	-0.2031	0.4825	0.0092	0.9847
---	---	1633014_at	0.0474	0.8908	0.0764	0.4622	0.4884	0.0237	0.1334	0.8436	0.0107	0.9783	-0.1227	0.6310	-0.3808	0.5126	0.0017	0.9989	0.3825	0.1901
CG32158	CG32158	1633015_at	0.1236	0.4060	0.1390	0.3967	0.1375	0.4511	0.1662	0.6171	0.0115	0.9638	-0.1547	0.3047	0.1058	0.8439	-0.1211	0.6071	-0.2269	0.3141
stau	Staufen	1633016_a_at	-1.3330	0.0017	-1.4767	0.0083	-0.9768	0.0012	0.3680	0.3056	0.0679	0.7932	-0.3000	0.1114	-0.0398	0.9816	0.0097	0.9914	0.0496	0.9264
Tsf3	Transferrin 3	1633017_at	-0.1958	0.5165	-0.4623	0.2605	-0.1373	0.8106	0.8294	0.1747	1.2643	0.0052	0.4349	0.1880	0.3460	0.8815	0.9750	0.2536	0.6291	0.4944
CG32681	CG32681	1633018_at	0.1089	0.4604	0.0206	0.8399	-0.0285	0.8875	-0.0858	0.8180	-0.0447	0.8027	0.0410	0.8030	0.0988	0.8882	0.0125	0.9794	-0.0863	0.7830
Spf45	CG17540	1633019_a_at	0.1279	0.4763	-0.0108	0.9791	0.0529	0.7905	0.0964	0.8686	0.1590	0.4573	0.0626	0.7855	-0.0252	0.9862	0.0112	0.9870	0.0364	0.9387
unk	unkempt	1633020_at	0.1908	0.6356	-0.3461	0.0777	-0.3960	0.3091	0.1532	0.7845	0.2149	0.3619	0.0617	0.8130	0.1980	0.9246	-0.2552	0.7593	-0.4532	0.5367
Spn	d-spinophilin	1633021_s_at	1.3071	0.0546	-0.3644	0.6151	-0.0056	0.9922	0.2178	0.9234	0.9807	0.1497	0.7629	0.2122	-0.1378	0.9653	-0.6614	0.4355	-0.5237	0.5612
CG1316 /// DyakCG1316	CG1316	1633022_at	-0.5117	0.3251	0.8881	0.1392	1.3751	0.0001	-0.3495	0.7221	-1.9368	0.0018	-1.5873	0.0025	-0.7426	0.5765	-0.3206	0.6207	0.4220	0.4991
Acp70A	sex peptide	1633023_at	0.1082	0.7023	-0.2470	0.1667	0.1237	0.4177	0.1101	0.9108	0.0943	0.8048	-0.0158	0.9679	-0.1834	0.7036	-0.0847	0.7093	0.0987	0.6476
CG14520	CG14520	1633024_at	0.0329	0.8810	-0.0513	0.6226	0.0422	0.8558	0.0125	0.9889	0.0221	0.9418	0.0096	0.9710	-0.0747	0.9117	-0.0210	0.9578	0.0537	0.8646
CG4627	CG4627	1633025_at	-0.1761	0.3840	-0.2733	0.1642	-0.0760	0.8076	-0.1703	0.7235	-0.0965	0.6999	0.0738	0.7561	-0.1984	0.8009	-0.0859	0.8472	0.1125	0.7691
CG31158	CG31158	1633026_a_at	-0.8609	0.3077	-0.5175	0.4129	-1.1135	0.0008	0.0019	0.9988	0.2985	0.2735	0.2966	0.2220	0.5966	0.8298	0.6471	0.6017	0.0504	0.9784
---	---	1633027_at	0.0144	0.9555	-0.2100	0.2381	-0.1388	0.4609	0.0844	0.8234	0.1235	0.4155	0.0391	0.8135	-0.1180	0.8608	-0.1259	0.6747	-0.0080	0.9852
---	---	1633028_at	0.2636	0.2421	0.0063	0.9575	0.1829	0.2620	0.0029	0.9961	0.0483	0.8380	0.0454	0.8284	-0.1150	0.7953	-0.0613	0.7932	0.0536	0.8178
CG10177	MLCK-like	1633029_at	-0.0291	0.9528	-0.0997	0.6015	-0.1286	0.5347	0.1711	0.8358	0.0480	0.9122	-0.1232	0.7095	-0.0432	0.9835	-0.2212	0.6823	-0.1779	0.7495
---	---	1633030_at	0.0830	0.7458	0.1292	0.4185	0.2192	0.3343	0.1898	0.6699	0.0727	0.7757	-0.1171	0.5785	0.0245	0.9858	0.1598	0.6314	0.1353	0.6929
CG8299	CG8299	1633031_at	-0.2315	0.6142	0.0865	0.6840	0.0214	0.9356	0.0459	0.9745	-0.4553	0.1973	-0.5012	0.1129	-0.0170	0.9901	-0.1833	0.5630	-0.1663	0.6095
CG8177 /// DmirCG8177	CG8177	1633032_s_at	-1.5434	0.0022	-2.0046	0.0014	-2.6521	0.0000	-0.0508	0.9461	0.3656	0.0858	0.4164	0.0351	0.4106	0.6272	-0.1678	0.6673	-0.5785	0.1313
CG1732	CG1732	1633033_s_at	-0.6990	0.0369	-0.2257	0.1018	-0.4188	0.1202	-0.2519	0.8274	-0.5459	0.2182	-0.2940	0.4897	-0.1980	0.8032	-0.1011	0.8132	0.0969	0.8138
Fer1HCH	Ferritin	1633034_s_at	-0.0102	0.9555	-0.2536	0.2098	-0.0495	0.7647	0.0376	0.9376	0.0759	0.6267	0.0383	0.8079	-0.1711	0.7677	-0.3188	0.1798	-0.1477	0.5702
---	---	1633035_at	0.1105	0.5260	0.1156	0.3873	-0.0334	0.8585	-0.2001	0.6325	-0.0520	0.8445	0.1481	0.4423	0.0983	0.8637	-0.0443	0.8980	-0.1426	0.5581
CG32495 /// GS	CG32495 /// Gluta	1633036_s_at	-0.4166	0.0352	0.6621	0.4237	1.1803	0.0004	0.1250	0.7857	-0.8883	0.0014	-1.0133	0.0005	-0.2824	0.8909	0.1940	0.8551	0.4764	0.5626
CG10669	CG10669	1633037_at	-0.1001	0.7771	0.1999	0.2032	0.3517	0.0737	-0.0279	0.9819	-0.5423	0.0676	-0.5145	0.0542	0.0521	0.9515	0.1593	0.5552	0.1072	0.7134
CG5863	CG5863	1633038_at	-0.0711	0.6314	-0.0862	0.5101	-0.1550	0.4775	-0.2532	0.5855	-0.0641	0.8314	0.1891	0.3899	-0.0146	0.9911	0.1218	0.6783	0.1363	0.6318
CG5646	CG5646	1633039_at	-1.8231	0.0010	-1.8913	0.0048	-1.8055	0.0001	-0.1465	0.6237	-0.5653	0.0032	-0.4188	0.0074	-0.2271	0.8243	-0.6624	0.1238	-0.4353	0.3175
---	---	1633040_at	0.1059	0.6056	-0.0069	0.9642	0.0916	0.6417	-0.0382	0.9592	0.0116	0.9698	0.0498	0.8239	0.0266	0.9816	-0.0056	0.9924	-0.0322	0.9237
---	---	1633041_at	0.0560	0.8042	0.0173	0.9103	-0.0401	0.8538	-0.0932	0.8508	-0.0451	0.8471	0.0481	0.8135	0.1165	0.8628	0.0776	0.8287	-0.0388	0.9175
CG7265	CG7265	1633042_at	0.5031	0.0458	0.3384	0.3845	0.4262	0.0398	0.0179	0.9774	0.0652	0.7158	0.0473	0.7837	-0.0844	0.9588	-0.0971	0.8823	-0.0127	0.9859
CG31159	CG31159	1633043_at	-0.6432	0.0502	0.2025	0.6403	0.4836	0.1525	-0.0268	0.9825	-0.1546	0.6279	-0.1278	0.6678	-0.2647	0.8479	0.5469	0.3269	0.8116	0.1823
CG17280 /// DyakCG17280	CG17280	1633044_at	-0.3842	0.1856	0.1799	0.4532	0.1000	0.7023	-0.2541	0.4433	-0.8400	0.0014	-0.5859	0.0039	-0.2090	0.8653	-0.3124	0.5422	-0.1033	0.8771

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1633045_at	0.0712	0.6563	0.1518	0.3999	-0.0022	0.9921	-0.2289	0.5735	-0.0239	0.9356	0.2050	0.2770	-0.0447	0.9589	0.0183	0.9647	0.0630	0.8443
CG7239	CG7239	1633046_at	-0.6205	0.0250	-0.3825	0.0884	-0.2346	0.2549	0.1818	0.5766	-0.0202	0.9317	-0.2020	0.1754	0.0064	0.9964	0.1118	0.7787	0.1054	0.7839
CG1809	CG1809	1633047_at	0.2801	0.4212	0.4198	0.3320	0.4020	0.0425	0.3118	0.4566	-0.0032	0.9920	-0.3150	0.1190	-0.0392	0.9848	-0.0821	0.9064	-0.0429	0.9466
CG8193	CG8193	1633048_at	-1.5510	0.0926	-3.3894	0.0097	-3.3010	0.0000	0.1818	0.8776	0.5253	0.1924	0.3434	0.3565	0.2211	0.9598	-1.2988	0.2829	-1.5199	0.2397
CG11159 /// DmirCG11159	CG11159 /// GA1(	1633049_at	0.2017	0.5789	-0.1283	0.5088	0.1264	0.4502	0.0833	0.8990	0.2540	0.2370	0.1707	0.3923	-0.3387	0.7307	-0.0910	0.8807	0.2477	0.5814
CG12813	CG12813	1633050_at	-0.0822	0.7644	0.0106	0.9312	0.0013	0.9959	0.2789	0.6082	0.0498	0.8932	-0.2291	0.3614	0.2299	0.7118	0.1163	0.6905	-0.1136	0.6956
CG13741 /// DyakCG13741	CG13741	1633051_at	1.0637	0.0311	0.2501	0.5664	0.0086	0.9876	-0.6706	0.0899	0.6732	0.0107	1.3438	0.0004	-0.5652	0.7720	-0.0986	0.9411	0.4665	0.6024
---	---	1633052_at	-0.1518	0.3625	-0.1102	0.4718	-0.1084	0.5480	0.0077	0.9922	-0.0942	0.5954	-0.1018	0.5186	-0.1654	0.7726	-0.1066	0.7005	0.0587	0.8534
IM1	Immune induced r	1633053_at	1.5849	0.0868	0.8457	0.4907	1.5516	0.0329	0.3074	0.8671	0.5743	0.3858	0.2669	0.6943	-0.3401	0.9555	-0.1481	0.9597	0.1920	0.9384
CG14057	CG14057	1633054_at	-1.5641	0.0014	-3.0842	0.0028	-2.1101	0.0001	0.9023	0.0644	1.8169	0.0004	0.9146	0.0038	0.1007	0.9291	0.4425	0.2116	0.3418	0.3629
CG10874	CG10874	1633055_at	-0.2159	0.4006	0.2080	0.2314	0.3825	0.0581	-0.0706	0.9345	-0.4124	0.1032	-0.3418	0.1296	-0.1833	0.7726	-0.0585	0.8804	0.1248	0.6740
CG7387	CG7387	1633056_at	0.0162	0.9226	0.1324	0.3676	0.1529	0.3267	0.0582	0.9339	-0.0619	0.8072	-0.1201	0.5532	0.2422	0.7070	0.1817	0.5141	-0.0605	0.8647
CG2871	CG2871	1633057_at	0.1732	0.4119	-0.3045	0.1602	-0.7054	0.0672	-0.1914	0.9037	0.6176	0.2293	0.8090	0.0800	-0.0917	0.9147	-0.1989	0.5232	-0.1072	0.7568
halo	halo	1633058_at	0.0581	0.7869	0.0615	0.6485	-0.2309	0.1357	0.0159	0.9819	0.1793	0.2819	0.1634	0.2750	0.1406	0.7997	0.0602	0.8481	-0.0804	0.7649
CG6357	CG6357	1633059_at	0.0088	0.9818	0.2581	0.3342	0.1874	0.3390	-0.0282	0.9761	0.0304	0.9254	0.0586	0.8253	0.1795	0.8494	0.4128	0.2775	0.2334	0.5771
Cyp6g2	Cyp6g2	1633060_at	0.2575	0.4195	0.0532	0.5766	0.1683	0.4580	-0.0419	0.9649	0.0230	0.9517	0.0650	0.8212	-0.0365	0.9775	0.0086	0.9901	0.0451	0.9162
Mgat2	Mgat2	1633061_at	-1.0637	0.0028	-0.7467	0.0956	-1.0483	0.0061	-0.2545	0.7579	-0.8593	0.0236	-0.6048	0.0605	0.0744	0.9640	-0.4550	0.2942	-0.5294	0.2527
CG32791	CG32791	1633062_at	0.2577	0.2576	0.0689	0.6654	-0.1684	0.5168	-0.2659	0.4597	0.1367	0.5064	0.4027	0.0286	0.0915	0.9246	0.0604	0.9036	-0.0311	0.9460
---	---	1633063_at	0.0537	0.7178	0.2710	0.1354	0.4092	0.0312	-0.0760	0.8897	-0.2559	0.1607	-0.1800	0.2773	-0.0569	0.9462	0.0701	0.8382	0.1271	0.6443
CG12985	CG12985	1633064_at	-0.1947	0.3314	0.1820	0.3945	0.2456	0.1188	-0.0777	0.9135	-0.4262	0.0622	-0.3485	0.0844	-0.1414	0.8814	-0.0320	0.9576	0.1094	0.8046
C901	C901	1633065_at	0.0384	0.8856	0.1864	0.3041	0.2323	0.1336	-0.0269	0.9603	-0.1427	0.3211	-0.1157	0.3785	-0.0375	0.9769	0.0402	0.9352	0.0777	0.8435
CG13215	CG13215	1633066_at	0.9116	0.4619	-1.3249	0.0090	-0.7000	0.0040	0.9268	0.0231	0.9969	0.0015	0.0701	0.7721	-0.0142	0.9989	-1.1404	0.4828	-1.1262	0.4938
CG34376	CG13845	1633067_at	2.6682	0.0055	1.0646	0.2360	2.5219	0.0002	0.7867	0.4346	0.2975	0.6288	-0.4891	0.3332	-0.7315	0.7726	-1.0449	0.3268	-0.3133	0.8182
CG4750	CG4750	1633068_at	-0.0403	0.8118	0.2671	0.2170	0.1660	0.5405	0.2390	0.4543	0.0157	0.9517	-0.2232	0.1480	0.3012	0.7142	0.2658	0.4380	-0.0355	0.9413
CG10277	CG10277	1633069_s_at	-0.4019	0.3224	0.0600	0.6100	-0.1703	0.3409	-0.2744	0.4163	-0.7128	0.0032	-0.4384	0.0169	-0.1355	0.9226	-0.2751	0.5719	-0.1397	0.8044
CG30222	CG30222	1633070_at	0.3655	0.2627	0.0031	0.9940	0.1518	0.4976	0.0173	0.9889	0.1185	0.7245	0.1012	0.7453	-0.1533	0.8599	-0.1667	0.6655	-0.0134	0.9821
---	---	1633071_at	-0.1121	0.6079	-0.1465	0.2532	-0.1271	0.4240	0.0847	0.8871	-0.0189	0.9505	-0.1036	0.6111	-0.0080	0.9922	-0.0043	0.9924	0.0037	0.9909
---	---	1633072_at	0.2634	0.1751	-0.3788	0.0691	-0.3632	0.0791	0.0507	0.9098	0.4429	0.0079	0.3923	0.0082	-0.1480	0.8609	-0.2973	0.3756	-0.1492	0.6955
---	---	1633073_at	0.2502	0.2255	0.0967	0.4958	0.0051	0.9843	-0.0044	0.9956	0.0033	0.9907	0.0077	0.9756	0.2523	0.7631	-0.0498	0.9278	-0.3020	0.3953
Moe	DMoesin	1633074_a_at	-1.1840	0.0023	-0.6797	0.0158	-1.2239	0.0003	-0.1379	0.6869	0.0398	0.8467	0.1777	0.2110	0.4019	0.6927	0.5818	0.1512	0.1800	0.7001
---	---	1633075_at	-0.0205	0.9248	0.0105	0.9233	0.0540	0.7577	-0.0153	0.9857	-0.0248	0.9255	-0.0095	0.9685	0.0387	0.9718	0.1512	0.6139	0.1125	0.7236
CG15092 /// CG34020	CG15092 /// CG3	1633076_at	1.5254	0.0019	1.4005	0.1110	1.6581	0.0001	-0.0202	0.9860	-0.3019	0.2765	-0.2817	0.2563	-0.1587	0.9499	-0.5069	0.5140	-0.3482	0.6708
Osi20	Osiris	1633077_at	0.0660	0.6647	0.2397	0.2605	0.2291	0.1641	-0.0735	0.8550	-0.1004	0.5138	-0.0269	0.8759	0.0583	0.9296	0.0689	0.7990	0.0105	0.9754
CG10268	CG10268	1633078_at	-0.5105	0.0608	-0.3334	0.0298	-0.1500	0.7201	-0.1318	0.8837	-0.0293	0.9493	0.1025	0.7606	-0.2070	0.8806	0.1264	0.8634	0.3334	0.5490
CG17472 /// DyakCG17472	CG17472	1633079_at	0.0954	0.8243	0.0144	0.9394	0.0458	0.8014	0.1252	0.7982	0.2164	0.2784	0.0912	0.6573	0.0413	0.9832	-0.0339	0.9647	-0.0752	0.9057
CG14835	CG14835	1633080_at	0.1050	0.7066	-0.0201	0.8425	0.0563	0.8166	-0.0813	0.9276	0.1465	0.6191	0.2278	0.3512	-0.0699	0.9503	-0.0247	0.9644	0.0452	0.9211
l(1)G0196	lethal (1) G0196	1633081_at	-0.0864	0.7545	0.0680	0.8630	-0.3845	0.1338	-0.3024	0.5136	-0.3789	0.1196	-0.0766	0.7781	0.1338	0.9246	-0.3059	0.5333	-0.4396	0.3648
CG8449	CG8449	1633082_at	0.7926	0.0275	-0.2908	0.5981	0.4438	0.1449	0.3882	0.5392	0.8404	0.0190	0.4522	0.1235	-0.3954	0.8009	-0.2130	0.7962	0.1824	0.8237
mRpS33	mitochondrial ribo	1633083_at	-0.3365	0.1588	0.0076	0.9855	-0.1626	0.3446	0.0070	0.9943	-0.0509	0.8511	-0.0579	0.8049	0.2164	0.7707	0.3858	0.2084	0.1694	0.6188
---	---	1633084_at	0.0582	0.6839	0.2292	0.1284	0.3082	0.0537	0.1125	0.7389	0.1124	0.4708	-0.0001	0.9995	-0.0195	0.9811	0.0687	0.7391	0.0881	0.6410
---	---	1633085_at	0.1775	0.4980	0.0313	0.7853	0.2521	0.1459	0.0255	0.9777	-0.1235	0.6377	-0.1489	0.5098	-0.1246	0.8207	-0.1466	0.5385	-0.0220	0.9449
CG1213	CG1213	1633086_s_at	2.7332	0.0030	0.9789	0.3437	2.3562	0.0001	0.0756	0.9754	-0.0822	0.9218	-0.1578	0.8144	-1.2548	0.6145	-1.9480	0.0795	-0.6932	0.5358
Octbeta2R	Octbeta2R	1633087_at	-1.0705	0.0041	-0.3533	0.0651	-0.1875	0.2236	-0.0557	0.9559	-0.9252	0.0054	-0.8695	0.0042	-0.0676	0.9092	-0.1160	0.6049	-0.0485	0.8604
CG4733 /// DereCG4733 ///	CG4733	1633088_at	-2.0517	0.0006	-1.8675	0.0507	-1.9495	0.0002	0.0591	0.9506	-0.2340	0.3848	-0.2931	0.2128	0.1843	0.9108	0.0324	0.9739	-0.1519	0.8379
lola	longitudinals abse	1633089_a_at	-0.7635	0.1727	1.1531	0.0863	0.8522	0.0064	0.1121	0.9491	-1.1076	0.0310	-1.2197	0.0131	0.3149	0.8814	0.8744	0.2610	0.5595	0.5041
CG14330	CG14330	1633090_at	-0.0081	0.9804	0.0099	0.9299	-0.1630	0.3070	-0.1452	0.7777	0.0373	0.8990	0.1824	0.3545	0.0752	0.9405	0.0740	0.8660	-0.0013	0.9982
CG1789	CG1789	1633091_at	0.0845	0.6990	0.0801	0.8217	0.1245	0.5485	-0.0368	0.9584	0.2664	0.1494	0.3032	0.0702	-0.0525	0.9761	0.2175	0.6325	0.2700	0.5444
cup	cup	1633092_at	0.2260	0.5452	0.1410	0.3885	0.2397	0.4205	-0.1262	0.7924	-0.0699	0.7646	0.0562	0.7975	-0.1647	0.9352	0.0495	0.9629	0.2142	0.7788
CG12983 /// DmirCG12983	CG12983	1633093_at	-0.3412	0.0617	0.0663	0.6702	0.0267	0.8865	-0.1350	0.7512	-0.3727	0.0491	-0.2377	0.1493	-0.0732	0.9189	0.0442	0.9068	0.1174	0.6601
toy	twin of eyeless	1633094_a_at	0.0170	0.9155	-0.0609	0.5842	0.0699	0.7032	0.0035	0.9956	0.0441	0.8218	0.0406	0.8191	-0.2490	0.6551	-0.1127	0.6414	0.1362	0.5649
ec	echinus	1633095_at	0.1405	0.4772	-0.1481	0.4514	0.1067	0.6758	0.0649	0.9154	0.2024	0.2944	0.1375	0.4459	-0.1343	0.8650	-0.1111	0.7723	0.0232	0.9596
---	---	1633096_at	0.1012	0.5298	-0.0929	0.4986	0.0061	0.9795	-0.0911	0.8281	0.0777	0.6721	0.1688	0.2436	-0.1386	0.7893	-0.0410	0.9035	0.0975	0.6894
CG14770	CG14770	1633097_at	0.1156	0.5891	0.1241	0.4215	0.3182	0.2417	0.2159	0.4586	0.1107	0.5064	-0.1052	0.4848	-0.0122	0.9923	0.0859	0.8255	0.0981	0.7792
---	---	1633098_at	-0.0295	0.9109	-0.0189	0.9126	0.0575	0.7727	-0.0305	0.9603	-0.0196	0.9309	0.0109	0.9573	-0.1043	0.9095	-0.0276	0.9590	0.0767	0.8570
CG3876	CG3876	1633099_at	0.0857	0.7573	0.8618	0.0205	0.6661	0.0018	-0.1938	0.6113	-0.5309	0.0131	-0.3372	0.0528	-0.0515	0.9717	0.2934	0.4242	0.3449	0.3623
Trp1	Translocation prot	1633100_a_at	-0.1038	0.6532	0.7283	0.0478	0.3800	0.0688	-0.1037	0.8422	0.0080	0.9790	0.1117	0.5588	0.1702	0.8611	0.8301	0.0568	0.6599	0.1271
---	---	1633101_at	0.0978	0.7967	0.0253	0.8767	-0.1287	0.4596	0.2468	0.4585	0.2337	0.1923	-0.0131	0.9544	0.1560	0.8692	0.0858	0.8734	-0.0702	0.8921
Ubc84D	Ubiquitin conjugat	1633102_at	0.0552	0.9009	-0.0260	0.9287	-0.0775	0.7797	0.1143	0.9223	-0.0053	0.9916	-0.1196	0.7567	0.0214	0.9846	-0.1055	0.7054	-0.1269	0.6328
CG14357	CG14357	1633103_at	0.1587	0.2503	0.0727	0.5902	0.1139	0.5306	0.0170	0.9803	-0.0178	0.9414	-0.0348	0.8588	-0.0217	0.9816	-0.1191	0.5711	-0.0974	0.6506
---	---	1633104_at	0.0489	0.7748	-0.0164	0.8874	-0.0262	0.9036	-0.1009	0.8794	-0.0831	0.7602	0.0178	0.9505	-0.0972	0.8270	-0.0485	0.8481	0.0488	0.8362
---	---	1633105_at	-0.0512	0.8327	-0.0232	0.8741	0.1218	0.4367	-0.0678	0.8815	-0.1825	0.2393	-0.1148	0.4303	-0.0010	0.9997	-0.0246	0.9531	-0.0236	0.9489
CG6662	CG6662	1633106_at	-0.7068	0.3230	-1.5916	0.0755	-1.5528	0.0000	0.0451	0.9518	0.5270	0.0189	0.4818	0.0171	-0.0201	0.9964	-0.2922	0.8377	-0.2721	0.8411
CG33204	CG33204	1633107_at	0.1458	0.3301	0.1236	0.4743	0.1997	0.3014	0.0374	0.9515	-0.0534	0.7977	-0.0908	0.5925	0.2001	0.7464	-0.0044	0.9941	-0.2045	0.4472
CG10083	Debrin-like	1633108_at	-0.3535	0.0366	-0.7162	0.0220	-0.3314	0.0374	-0.0134	0.9857	-0.1493	0.3964	-0.1359	0.3941	-0.3860	0.3162	-0.6103	0.0236	-0.2243	0.2822
CG16971 /// CG33229	CG16971 /// CG3	1633109_at	-0.2713	0.5144	0.0418	0.9596	-0.3100	0.0640	-0.3806	0.5842	-0.2470	0.5097	0.1336	0.7237	-0.3557	0.8424	-0.3348	0.6823	0.0209	0.9856
CG3353	CG3353	1633110_at	0.5486	0.0915	-0.4587	0.3755	-0.5529	0.0029	0.2076	0.7539	1.1001	0.0031	0.8925	0.0047	0.3481	0.7707	0.2020	0.7346	-0.1461	0.8184
Porin2	Porin2	1633111_at	0.1087	0.5597	0.3375	0.2141	0.1744	0.4318	-0.1008	0.8485	-0.0452	0.8579	0.0556	0.7976	0.0148	0.9922	0.2413	0.4969	0.2265	0.5340
moody	CG4322	1633112_at	0.3138	0.2607	-0.5867	0.4204	-0.0339	0.9178	0.2883	0.5068	0.5402	0.0267	0.2519	0.2214	-0.2374	0.9142	-0.3497	0.6844	-0.1122	0.9164
CG6876	CG6876	1633113_at	-0.0898	0.5567	0.2652	0.1472	0.1793	0.3049	-0.0956	0.8078	-0.2176	0.1628	-0.1220	0.4035	0.0003	0.9999	0.0911	0.7672	0.0907	0.7564

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1633114_at	0.1167	0.5431	0.1202	0.6597	-0.2897	0.1738	-0.1009	0.8732	0.0135	0.9686	0.1144	0.6051	0.2605	0.7116	0.0172	0.9723	-0.2434	0.4148
C3G	C3G	1633115_a_at	-1.4751	0.0096	-2.1892	0.0142	-2.0591	0.0001	0.0701	0.9549	0.4952	0.1390	0.4251	0.1558	-0.0831	0.9725	-0.2710	0.6908	-0.1879	0.7981
---	---	1633116_at	0.0841	0.5532	0.0156	0.9013	0.2735	0.2741	0.1273	0.8200	0.0718	0.7883	-0.0554	0.8239	-0.1010	0.8846	0.0080	0.9886	0.1090	0.7144
CG12395	CG12395	1633117_at	0.1258	0.5289	0.0608	0.7098	0.0791	0.6664	-0.0893	0.8498	-0.1393	0.4344	-0.0500	0.7956	-0.0068	0.9950	-0.1033	0.7200	-0.0965	0.7409
CG5768	CG5768	1633118_at	0.2642	0.1406	0.1807	0.2743	0.1688	0.3248	-0.1661	0.6738	-0.0062	0.9822	0.1599	0.3498	-0.0731	0.9324	-0.0388	0.9291	0.0342	0.9295
---	---	1633119_at	0.0167	0.9441	0.0105	0.9215	0.4252	0.1118	-0.0382	0.9639	-0.2067	0.3616	-0.1686	0.4174	-0.4377	0.5134	-0.2040	0.5431	0.2337	0.4825
---	---	1633120_at	-0.0137	0.9697	-0.1426	0.2818	-0.0264	0.8757	0.0398	0.9626	0.0105	0.9762	-0.0294	0.9148	-0.0369	0.9741	-0.1212	0.7003	-0.0843	0.8044
---	---	1633121_at	0.0580	0.7688	0.0315	0.7708	0.1243	0.4026	0.0673	0.8974	-0.0149	0.9534	-0.0822	0.6411	0.0517	0.9367	0.1652	0.4203	0.1135	0.6092
---	---	1633122_at	-0.0761	0.6565	0.0010	0.9949	-0.0233	0.9187	-0.0623	0.8841	-0.1786	0.2189	-0.1163	0.3900	-0.1038	0.8553	-0.0719	0.8123	0.0318	0.9211
CG7201	CG7201	1633123_at	-6.1965	0.0005	-7.0322	0.0007	-6.3803	0.0001	0.1334	0.9081	0.6020	0.1034	0.4686	0.1558	-0.6401	0.8049	-0.5502	0.6380	0.0899	0.9525
CG6026	CG6026	1633124_at	-0.0669	0.7816	0.1871	0.4129	0.1257	0.5576	0.0031	0.9956	-0.1417	0.4376	-0.1448	0.3735	0.0935	0.9101	0.0548	0.9037	-0.0387	0.9242
CG40236	CG40236	1633125_at	-0.0300	0.8793	0.1080	0.4645	0.0399	0.8058	-0.2123	0.5362	-0.1378	0.4607	0.0745	0.6919	0.0822	0.9088	0.0191	0.9634	-0.0630	0.8467
CG4596	CG4596	1633126_at	-1.0194	0.1053	-1.8933	0.0028	-1.3080	0.0165	0.3184	0.8067	0.8299	0.1074	0.5116	0.2731	-0.3302	0.8465	-0.1967	0.8356	0.1335	0.8915
---	---	1633127_at	0.1604	0.5467	0.1070	0.4955	0.0677	0.7261	0.1073	0.7845	-0.0118	0.9611	-0.1191	0.4330	-0.0458	0.9511	-0.1320	0.5689	-0.0862	0.7334
CG15471	CG15471	1633128_at	0.4306	0.0719	0.0202	0.8969	0.2620	0.2603	0.2011	0.6257	0.2739	0.1770	0.0727	0.7403	-0.1779	0.7726	-0.0807	0.8123	0.0972	0.7484
CG7183	CG7183	1633129_at	-0.4745	0.0781	0.1634	0.4579	0.1326	0.5984	-0.0988	0.8384	-0.4455	0.0224	-0.3467	0.0393	-0.1539	0.8940	0.1770	0.7200	0.3309	0.4573
CG34035	CG34035	1633130_at	-1.4669	0.1752	0.2464	0.8618	-0.0430	0.9560	-0.7861	0.3353	-1.6833	0.0043	-0.8971	0.0396	-0.6285	0.9030	-0.1486	0.9620	0.4800	0.8414
CG32164 /// CG32165	CG32165 /// CG3	1633131_s_at	0.2798	0.2181	0.6867	0.1226	0.4081	0.0217	0.0014	0.9988	-0.0711	0.7743	-0.0725	0.7443	0.2597	0.7981	0.3023	0.4861	0.0427	0.9431
fry	Suppressor of GM	1633132_a_at	0.0039	0.9902	0.0914	0.5251	-0.0245	0.9046	-0.0078	0.9937	-0.0656	0.7803	-0.0578	0.7903	0.1576	0.8037	0.1280	0.6582	-0.0295	0.9361
---	---	1633133_at	0.2552	0.1574	0.2333	0.1341	0.1814	0.2765	-0.1452	0.7511	-0.0970	0.6683	0.0483	0.8313	0.0566	0.9142	0.0209	0.9445	-0.0357	0.8880
CG30099	CG30099	1633134_at	0.0354	0.8693	0.1839	0.4055	0.2996	0.0520	-0.0487	0.9346	0.0045	0.9868	0.0532	0.7861	-0.0601	0.9544	0.0099	0.9873	0.0700	0.8633
CG18854	CG18854	1633135_s_at	-0.5960	0.0969	-0.8085	0.3629	-0.9914	0.0272	0.0283	0.9639	0.1811	0.2772	0.1528	0.3106	0.3194	0.9225	-0.0304	0.9900	-0.3498	0.7872
CG13035	CG13035	1633136_a_at	0.2647	0.1451	-0.0219	0.8274	0.3830	0.0605	0.2668	0.3237	0.1363	0.3980	-0.1304	0.3686	-0.0620	0.9238	0.0500	0.8709	0.1120	0.6311
eIF4E-6	eIF4E-6	1633137_at	0.1194	0.6434	0.2417	0.1700	0.1178	0.4684	-0.1735	0.6591	-0.2555	0.1743	-0.0820	0.6779	0.1183	0.8791	0.0828	0.8362	-0.0356	0.9329
CG32251	CG32251	1633138_at	0.1510	0.4351	-0.1951	0.4927	-0.4858	0.0547	-0.4107	0.5199	0.5567	0.0975	0.9673	0.0068	-0.1489	0.9151	0.0522	0.9477	0.2011	0.7154
e(y)1	enhancer of yellow	1633139_at	0.0366	0.8285	0.5894	0.0406	0.8869	0.0004	0.0867	0.8449	-0.5785	0.0045	-0.6652	0.0015	-0.1118	0.8510	0.2148	0.3738	0.3266	0.2056
---	---	1633140_at	-0.0323	0.8402	-0.0321	0.7899	-0.0103	0.9645	-0.0500	0.9251	0.0176	0.9387	0.0676	0.6862	0.0634	0.9405	0.0748	0.8303	0.0114	0.9785
---	---	1633141_at	-0.2693	0.2245	0.0074	0.9669	-0.2483	0.1379	0.0415	0.9435	-0.0901	0.6244	-0.1316	0.3924	0.2057	0.7956	0.0322	0.9521	-0.1735	0.6310
CG17681	CG17681	1633142_at	0.4256	0.0331	0.3700	0.1230	0.0944	0.5513	-0.1497	0.6338	0.0633	0.7250	0.2131	0.1211	-0.0762	0.9341	-0.0322	0.9462	0.0440	0.9140
Dyrk3	unknown-telomeri	1633143_s_at	0.0910	0.8333	-0.3433	0.4599	-0.5506	0.0127	-0.2820	0.6046	0.0616	0.8650	0.3436	0.1601	-0.0853	0.9590	-0.3471	0.4763	-0.2618	0.6124
jbug	filamin	1633144_at	-1.4758	0.0360	-1.3354	0.1121	-1.3349	0.0004	0.3704	0.6041	0.5057	0.1581	0.1353	0.7247	0.2087	0.9459	0.6677	0.4820	0.4591	0.6468
PGRP-LF	PRGP-like	1633145_at	0.2543	0.6034	-2.1380	0.0019	-1.9342	0.0019	0.3303	0.6582	2.5796	0.0003	2.2493	0.0003	-0.0964	0.9737	-0.0034	0.9991	0.0931	0.9252
---	---	1633146_at	0.0578	0.8305	-0.2614	0.1374	-0.3815	0.0495	0.2431	0.6198	0.4596	0.0633	0.2165	0.3316	0.0662	0.9243	-0.0704	0.8153	-0.1366	0.5811
CG10659	CG10659	1633147_at	0.0907	0.6708	-0.0780	0.5103	0.0307	0.8854	0.1237	0.8028	-0.0514	0.8407	-0.1751	0.3340	-0.0396	0.9779	-0.0738	0.8807	-0.0342	0.9433
---	---	1633148_at	0.0205	0.9152	0.0067	0.9710	0.2126	0.2331	0.0481	0.9380	0.0038	0.9892	-0.0443	0.8300	-0.0387	0.9514	-0.0078	0.9829	0.0310	0.9056
Gs1l	GS1-like protein	1633149_at	0.3332	0.1707	0.7882	0.1027	0.7316	0.0077	-0.0093	0.9943	-0.7354	0.0134	-0.7261	0.0086	-0.0039	0.9990	-0.3105	0.5024	-0.3066	0.5131
CG12677	CG12677	1633150_a_at	0.2167	0.1862	-0.0884	0.5895	0.2147	0.1670	0.0713	0.8967	0.1145	0.5572	0.0432	0.8350	-0.1990	0.7779	-0.2252	0.4694	-0.0262	0.9515
---	---	1633151_at	0.2064	0.3005	0.1559	0.5112	0.1980	0.4273	0.0941	0.8836	0.0734	0.7818	-0.0207	0.9394	0.0694	0.9499	0.1584	0.6605	0.0890	0.8299
mod(mdg4)	Modifier67.2	1633152_at	-0.1743	0.2908	0.0071	0.9540	0.2994	0.0814	0.1098	0.7851	-0.1342	0.4358	-0.2441	0.1000	-0.1764	0.7230	-0.0038	0.9935	0.1726	0.4301
cdc14	CG7134	1633153_at	-0.0063	0.9860	0.2135	0.5608	0.0666	0.7920	-0.2505	0.6080	-0.2341	0.3474	0.0163	0.9582	0.1005	0.9416	0.1957	0.6779	0.0951	0.8653
CG11281	CG11281	1633154_at	0.0963	0.6803	-0.0909	0.6869	-0.0627	0.7742	0.0354	0.9656	0.0502	0.8605	0.0148	0.9573	-0.0636	0.9441	-0.0809	0.8218	-0.0174	0.9661
---	---	1633155_at	0.1871	0.5014	0.0670	0.6680	0.1956	0.2562	-0.0249	0.9772	0.1066	0.6559	0.1315	0.5239	-0.0745	0.9390	0.0094	0.9877	0.0839	0.8237
CG31537	CG31537	1633156_at	0.1073	0.6049	0.0318	0.8424	-0.1025	0.6136	-0.0159	0.9860	-0.0539	0.8449	-0.0381	0.8803	0.1182	0.8650	-0.0372	0.9335	-0.1553	0.6024
igl	igloo	1633157_a_at	-1.8507	0.0426	-0.1896	0.3014	-1.2372	0.0097	-0.9788	0.4597	-1.8094	0.0200	-0.8306	0.1947	0.1665	0.8608	-0.1201	0.8060	-0.2866	0.4677
CG10761	CG10761	1633158_at	0.1204	0.4206	0.0030	0.9832	0.1500	0.4814	-0.0651	0.9039	-0.0595	0.7795	0.0056	0.9799	-0.2252	0.7707	-0.1941	0.5800	0.0311	0.9466
CG11900	CG11900	1633159_at	-0.0474	0.8539	0.1037	0.5667	-0.2738	0.2057	-0.2989	0.5079	-0.0295	0.9298	0.2694	0.2082	0.1578	0.8425	0.1608	0.6497	0.0030	0.9958
CG14736	CG14736	1633160_a_at	0.1944	0.2947	0.0160	0.8768	0.0340	0.8644	-0.0430	0.9441	0.0512	0.8139	0.0942	0.5924	-0.0678	0.8967	-0.0041	0.9925	0.0637	0.7827
CG14074	CG14074	1633161_at	0.2463	0.5870	0.2742	0.2717	0.0172	0.9482	-0.1429	0.8708	0.1117	0.7619	0.2546	0.3699	0.1106	0.9623	0.2013	0.8118	0.0907	0.9195
CG13532	CG13532	1633162_at	0.0708	0.7511	-0.0968	0.7322	-0.0856	0.5994	-0.0594	0.9528	0.0961	0.7733	0.1555	0.5719	0.0525	0.9455	0.0067	0.9896	-0.0458	0.8859
l(1)G0431	lethal (1) G0431	1633163_at	0.0764	0.8478	0.1355	0.5084	0.7858	0.0025	0.4946	0.3487	0.1920	0.5567	-0.3026	0.2687	-0.1616	0.8454	0.2420	0.4868	0.4036	0.2528
CG10625	CG10625	1633164_s_at	-2.9877	0.0036	-1.1626	0.1562	-2.0130	0.0004	-0.4774	0.4162	-1.4251	0.0018	-0.9477	0.0064	0.0193	0.9952	-0.4135	0.6209	-0.4328	0.6052
CG17275	CG17275	1633165_at	0.0477	0.7524	-0.2109	0.3813	0.1888	0.2670	0.1305	0.7293	0.0140	0.9556	-0.1166	0.4666	-0.0720	0.9365	-0.1190	0.7134	-0.0470	0.9055
CG12942	CG12942	1633166_at	0.3397	0.2762	-0.4509	0.1151	-0.4500	0.0232	-0.0169	0.9803	0.7970	0.0012	0.8139	0.0006	-0.0527	0.9741	0.0001	1.0000	0.0528	0.9231
CG32496 /// CG6788	CG32496 /// CG6	1633167_s_at	0.1839	0.2137	0.0913	0.6084	-0.0126	0.9682	-0.1771	0.7298	-0.2144	0.3603	-0.0373	0.8924	0.1074	0.8608	-0.1532	0.5522	-0.2607	0.3008
CG2574	CG2574	1633168_at	0.1126	0.5920	-0.1379	0.4262	0.0176	0.9191	-0.0289	0.9744	0.0933	0.7114	0.1222	0.5723	-0.1018	0.9101	-0.1039	0.7986	-0.0021	0.9974
dlt	Vanaso	1633169_at	-0.1332	0.7196	0.0711	0.7346	0.1315	0.6316	0.4032	0.2753	0.8055	0.0037	0.4023	0.0449	0.3974	0.7387	0.9307	0.0795	0.5333	0.3031
lectin-30A	Lectin30A	1633170_at	0.0567	0.7497	0.1515	0.3681	0.2415	0.2417	-0.1409	0.7749	-0.1482	0.4937	-0.0073	0.9776	-0.0635	0.9447	-0.0420	0.9225	0.0214	0.9573
CG13690	CG13690	1633171_at	0.3126	0.3511	-0.0820	0.6544	-0.0273	0.9507	-0.1668	0.7647	0.4458	0.0644	0.6126	0.0114	-0.1100	0.9558	0.2530	0.6971	0.3630	0.5518
---	---	1633172_at	0.0116	0.9528	0.0695	0.5200	0.1840	0.2533	-0.0667	0.9171	-0.2046	0.3071	-0.1379	0.4631	-0.1067	0.8202	-0.0407	0.8850	0.0660	0.7734
CG34401	CG17757	1633173_s_at	0.0608	0.7964	0.7278	0.0120	0.7551	0.0016	-0.1612	0.7275	-0.4431	0.0388	-0.2819	0.1262	0.0000	0.9999	0.1968	0.4271	0.1968	0.4408
stg	string	1633174_at	-1.9743	0.0611	-2.2133	0.1249	-2.7179	0.0032	-0.7850	0.5140	-0.0764	0.9319	0.7086	0.2124	-0.3537	0.9514	-0.4314	0.8556	-0.0777	0.9776
---	---	1633175_at	-0.0046	0.9847	0.1808	0.2788	-0.0027	0.9905	0.0451	0.9507	0.0789	0.7401	0.0338	0.8883	0.1183	0.8940	0.1981	0.5715	0.0799	0.8526
Ilp1	Drosophila insulin	1633176_at	0.0566	0.8084	0.1151	0.4709	0.1000	0.5028	-0.1461	0.6823	-0.1410	0.4106	0.0051	0.9802	0.0657	0.9374	-0.0611	0.8686	-0.1268	0.6553
---	---	1633177_at	0.1578	0.4294	-0.1334	0.3895	-0.0950	0.7282	0.2285	0.5229	0.2436	0.1914	0.0151	0.9494	0.0805	0.9365	-0.0835	0.8478	-0.1640	0.6328
sub	double-or-nothing	1633178_at	-0.1514	0.4352	-0.0503	0.8796	-0.2416	0.2007	-0.2055	0.5933	-0.0242	0.9298	0.1814	0.3065	0.0223	0.9916	0.0246	0.9782	0.0023	0.9979
CG11391	CG11391	1633179_at	-0.1198	0.6228	0.1296	0.4578	-0.0050	0.9885	-0.0769	0.9316	-0.2201	0.4260	-0.1431	0.5924	-0.0520	0.9619	0.0792	0.8481	0.1313	0.6985
exu	exuperantia	1633180_at	0.1455	0.3248	-0.0067	0.9777	0.1178	0.4411	0.2403	0.4861	0.3833	0.0439	0.1430	0.4066	-0.0047	0.9963	0.0356	0.9162	0.0402	0.8932
osm-1	osm-1	1633181_at	0.0958	0.5636	0.1189	0.5460	0.2194	0.4774	-0.0991	0.8794	-0.2756	0.2160	-0.1766	0.3954	-0.0670	0.9467	-0.0506	0.9152	0.0164	0.9725
CG31661	CG31661	1633182_at	0.2031	0.2695	0.0465	0.8385	0.2005	0.1727	0.0778	0.9222	0.0499	0.8758	-0.0279	0.9247	-0.0934	0.8882	0.0052	0.9928	0.0986	0.7305

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13134	CG13134	1633183_at	-0.0255	0.8855	-0.0011	0.9966	-0.0325	0.8847	-0.2235	0.5735	-0.0844	0.7215	0.1391	0.4715	-0.0540	0.9653	-0.0618	0.9023	-0.0078	0.9880
CG9389	CG9389	1633184_at	0.1697	0.3885	0.0322	0.9062	0.1592	0.3010	0.0449	0.9375	0.1442	0.3978	0.0994	0.5409	-0.0778	0.9246	-0.1114	0.7267	-0.0336	0.9312
dpr14	dpr14	1633185_at	0.4022	0.1661	-0.1774	0.4019	0.3455	0.0859	0.1694	0.6465	0.3668	0.0460	0.1974	0.2177	-0.0796	0.9092	0.0014	0.9989	0.0810	0.7815
CG9153	CG9153	1633186_s_at	-0.8463	0.0442	0.6103	0.0843	0.4106	0.0459	-0.3122	0.5068	-0.8876	0.0044	-0.5754	0.0180	-0.0488	0.9816	0.7102	0.1395	0.7590	0.1459
CG6619	CG6619	1633187_at	-0.1975	0.4449	-0.2613	0.1910	-0.0140	0.9533	0.4039	0.3016	0.1361	0.5862	-0.2678	0.1921	0.0279	0.9816	-0.1413	0.6314	-0.1691	0.5587
Grip84	gamma-tubulin rin	1633188_a_at	-1.4646	0.0039	-0.4498	0.2444	-0.0375	0.8782	0.1203	0.8546	-0.3889	0.1013	-0.5092	0.0246	-0.4070	0.7741	0.3706	0.5650	0.7776	0.2221
CG13646	CG13646	1633189_at	0.0062	0.9827	0.2657	0.2611	0.2094	0.1906	0.0750	0.8863	-0.0197	0.9383	-0.0948	0.5944	0.0951	0.9445	-0.0451	0.9462	-0.1402	0.7734
CG31278	CG31278	1633190_at	0.0995	0.6369	0.4326	0.1584	0.2943	0.1157	-0.2326	0.5357	-0.3873	0.0529	-0.1547	0.3961	-0.0234	0.9816	0.1943	0.3972	0.2177	0.3616
---	---	1633191_at	-0.1527	0.6856	0.2327	0.3472	0.6190	0.0178	0.1770	0.7188	-0.2705	0.2239	-0.4475	0.0321	-0.4041	0.6955	-0.0196	0.9797	0.3844	0.3791
CG34406	NAT1	1633192_at	0.2637	0.3768	0.1519	0.2733	0.0884	0.6034	-0.0196	0.9863	0.0859	0.7984	0.1055	0.7174	0.0000	0.9999	0.0557	0.8850	0.0557	0.8778
CheA29a	CheA29a	1633193_at	0.1052	0.5811	-0.1451	0.5582	-0.0717	0.7279	0.2102	0.6480	0.1893	0.4090	-0.0209	0.9397	0.0189	0.9882	0.0489	0.9085	0.0300	0.9387
CG16848	CG16848	1633194_at	0.2257	0.1250	-0.0504	0.7218	-0.1516	0.4051	-0.0423	0.9368	0.0255	0.9052	0.0679	0.6761	-0.0638	0.9030	-0.1337	0.4948	-0.0699	0.7502
CG8065	CG8065	1633195_at	0.1702	0.3115	-0.0407	0.7571	0.1360	0.4167	0.0625	0.9068	0.0111	0.9672	-0.0514	0.7888	-0.0304	0.9653	-0.1105	0.5778	-0.0801	0.7030
alpha-Est2	fragment B	1633196_at	3.3726	0.0035	1.3333	0.1748	2.4029	0.0001	0.4755	0.7104	0.5426	0.3612	0.0671	0.9252	-0.7619	0.7230	-1.4415	0.1248	-0.6795	0.4818
CG32024	CG32024	1633197_at	-0.1033	0.8512	-0.0517	0.7333	0.1049	0.6673	-0.0402	0.9778	-0.4171	0.2641	-0.3770	0.2600	-0.3282	0.7136	-0.3267	0.3726	0.0014	0.9985
5-HT2	Serotonin recepto	1633198_a_at	0.0606	0.7979	0.1463	0.2290	0.1071	0.4804	-0.1420	0.7982	-0.0498	0.8671	0.0922	0.6970	-0.1059	0.8465	-0.0549	0.8624	0.0510	0.8647
Mst98Ca	Male-specific RNA	1633199_at	0.2344	0.4266	-0.0790	0.7301	-0.2113	0.2946	0.1107	0.8636	0.3219	0.1575	0.2112	0.3097	-0.0136	0.9916	0.0185	0.9712	0.0321	0.9402
CG9449	CG9449	1633200_at	2.3014	0.0044	1.1232	0.0798	2.5597	0.0001	0.5451	0.5502	-0.2773	0.5890	-0.8225	0.0554	-0.7367	0.6749	-1.3830	0.0689	-0.6464	0.3756
CG8245	CG8245	1633201_at	-0.9664	0.0057	-0.1200	0.3998	-0.2812	0.0749	0.0287	0.9744	-0.7165	0.0059	-0.7452	0.0029	-0.0320	0.9738	-0.0946	0.7351	-0.0626	0.8372
---	---	1633202_at	0.0431	0.8869	-0.0266	0.8287	0.1487	0.5011	-0.0512	0.9586	-0.0448	0.9003	0.0064	0.9849	0.0113	0.9928	0.0077	0.9903	-0.0036	0.9945
---	---	1633203_s_at	-0.0072	0.9801	-0.0753	0.5419	-0.1427	0.3592	0.0248	0.9759	0.0911	0.6939	0.0662	0.7644	0.0153	0.9914	0.0559	0.9075	0.0406	0.9254
CG10971	CG10971	1633204_a_at	-0.7399	0.0452	-0.8674	0.0321	-1.1487	0.0020	-0.1397	0.8000	0.6730	0.0091	0.8126	0.0023	-0.0343	0.9898	0.5404	0.3322	0.5747	0.3262
DAT	Dopamine transpo	1633205_at	-0.3460	0.2990	0.0508	0.7298	0.0880	0.6690	-0.1184	0.9154	-0.6155	0.0798	-0.4971	0.1107	-0.0855	0.9174	-0.1673	0.5818	-0.0818	0.8184
CG9780	CG9780	1633206_at	0.3520	0.3716	1.2467	0.2652	1.1709	0.0630	-0.2970	0.8678	-1.6958	0.0152	-1.3988	0.0215	-0.1048	0.9841	-0.7811	0.5289	-0.6763	0.5994
CG31906	CG31906	1633207_at	-0.5658	0.0251	-0.7056	0.0442	-0.9503	0.0036	0.2240	0.7589	0.4465	0.1522	0.2225	0.4520	0.1955	0.8692	0.0378	0.9617	-0.1577	0.7755
---	---	1633208_at	0.5275	0.2377	1.7895	0.0616	2.2239	0.0020	0.1963	0.8340	-0.0348	0.9478	-0.2310	0.4985	-0.3067	0.9152	1.5397	0.1287	1.8464	0.1013
eIF3-S9	eIF3-S9	1633209_s_at	0.2661	0.1401	0.4718	0.0590	0.5667	0.0048	0.1250	0.7143	-0.1580	0.3155	-0.2829	0.0475	0.1532	0.8122	0.0960	0.7656	-0.0573	0.8729
CG10674 /// DyakCG10674	CG10674	1633210_at	-0.0445	0.8692	0.5698	0.0478	0.7564	0.0057	0.0195	0.9796	-0.2660	0.1508	-0.2854	0.0864	-0.1840	0.8438	0.3315	0.3768	0.5156	0.2012
CG7795	CG7795	1633211_a_at	-0.1227	0.4716	-0.2315	0.3449	-0.1839	0.3367	0.1033	0.8074	0.1719	0.3127	0.0686	0.6996	0.0180	0.9862	-0.0351	0.9264	-0.0531	0.8669
CG31226	CG31226	1633212_s_at	-0.0131	0.9633	-0.0566	0.7676	0.1405	0.6313	0.2888	0.4110	0.2675	0.1671	-0.0213	0.9286	0.0738	0.9441	0.0960	0.8169	0.0222	0.9622
U2af50	U2 small nuclear r	1633213_at	-0.0020	0.9936	-0.0350	0.8535	-0.0876	0.5809	-0.0387	0.9406	-0.1431	0.3456	-0.1044	0.4593	0.0314	0.9831	-0.0633	0.9018	-0.0947	0.8234
CG4835	CG4835	1633214_at	0.1331	0.4120	-0.0580	0.8273	0.1460	0.5383	0.0223	0.9722	0.1042	0.5464	0.0818	0.6146	-0.1405	0.8692	-0.0926	0.8405	0.0479	0.9194
CG12006	CG12006	1633215_at	0.3817	0.0723	0.3098	0.1481	0.4675	0.0093	-0.1736	0.6360	0.0945	0.6401	0.2681	0.0956	-0.2740	0.6557	0.0387	0.9197	0.3128	0.2226
Gr93c	Gustatory recepto	1633216_at	0.0834	0.6757	0.2458	0.1485	0.0893	0.6997	-0.1568	0.6188	0.0083	0.9732	0.1651	0.2408	0.0963	0.8953	0.1743	0.5411	0.0779	0.8187
CG30101	CG30101	1633217_at	0.3355	0.2124	0.6115	0.0381	0.6804	0.0039	-0.2832	0.6086	-0.6060	0.0358	-0.3228	0.1927	-0.0114	0.9914	0.1423	0.5501	0.1537	0.5176
UGP	UTP-G-1-P uridyly	1633218_a_at	2.6169	0.0039	-0.1515	0.6223	1.3303	0.0052	0.2131	0.7701	0.8259	0.0155	0.6128	0.0336	-1.1542	0.4207	-1.8034	0.0404	-0.6492	0.4044
---	---	1633219_at	0.1600	0.6123	-0.1053	0.4342	0.1512	0.4799	0.3341	0.5470	0.3786	0.1852	0.0445	0.8975	0.0439	0.9816	0.0232	0.9739	-0.0207	0.9756
CG16739	CG16739	1633220_at	0.2511	0.2157	0.0086	0.9372	0.1511	0.4233	0.1027	0.8405	0.1272	0.5319	0.0245	0.9161	0.0021	0.9994	-0.1415	0.7339	-0.1436	0.7234
---	---	1633221_at	0.0068	0.9795	0.0526	0.8672	0.1335	0.5054	-0.0764	0.9353	-0.4013	0.1466	-0.3249	0.1898	0.0366	0.9764	-0.1513	0.6394	-0.1879	0.5499
CG6073	CG6073	1633222_at	-0.0247	0.9588	-0.5173	0.3486	0.1431	0.6778	0.1858	0.8399	0.7842	0.0299	0.5984	0.0555	-0.3688	0.8444	0.1865	0.8624	0.5553	0.4863
---	---	1633223_at	0.1166	0.5147	-0.0514	0.6748	0.0300	0.9024	0.1064	0.8074	0.1948	0.2647	0.0884	0.6161	0.1271	0.8634	-0.0208	0.9658	-0.1479	0.6425
CG9631	CG9631	1633224_at	1.5541	0.0075	0.1089	0.8941	1.2477	0.0004	0.4107	0.5633	0.9230	0.0195	0.5123	0.1145	-0.7023	0.7116	-0.5296	0.5183	0.1727	0.8699
CG32278	CG32278	1633225_at	-0.2924	0.1958	-0.2404	0.2715	-0.3268	0.0811	-0.0533	0.9110	-0.2084	0.1655	-0.1551	0.2533	-0.0221	0.9898	-0.2015	0.6020	-0.1794	0.6457
hts	hu-li tai shao	1633226_at	-0.1322	0.6371	-0.3799	0.0823	-0.0812	0.7597	0.0333	0.9759	0.1589	0.5912	0.1256	0.6551	-0.1930	0.7215	-0.0328	0.9275	0.1602	0.4978
CG14945	CG14945	1633227_s_at	1.9737	0.0057	0.7922	0.1304	1.2596	0.0007	0.2523	0.7433	0.3122	0.3677	0.0598	0.8815	-0.2135	0.8956	-0.6640	0.2558	-0.4505	0.4738
---	---	1633228_at	0.1792	0.3777	0.1250	0.5339	0.2625	0.1877	0.1848	0.6338	0.0339	0.8963	-0.1508	0.3942	-0.0300	0.9862	0.0626	0.9188	0.0927	0.8577
Hr38	Hormone receptor	1633229_a_at	0.2738	0.1673	-0.0508	0.7995	0.0413	0.8780	0.0691	0.9311	0.1896	0.4338	0.1205	0.6100	0.0750	0.9296	0.0542	0.8933	-0.0208	0.9582
CG13457	CG13457	1633230_a_at	0.1141	0.5573	0.1687	0.1836	-0.0353	0.8584	-0.0440	0.9622	-0.2111	0.4068	-0.1671	0.4788	0.0716	0.9405	-0.0477	0.9170	-0.1193	0.7223
Trf2	TBP-like factor	1633231_a_at	0.1787	0.4119	0.3025	0.4949	-0.3472	0.1098	-0.4476	0.1483	0.3891	0.0437	0.8368	0.0010	0.2969	0.7893	0.4655	0.3140	0.1686	0.7578
CG18358	CG18358	1633232_at	-0.9781	0.0012	-0.8351	0.0478	-1.1602	0.0043	0.1839	0.7828	-0.3191	0.2485	-0.5030	0.0469	0.4253	0.6749	-0.2766	0.5296	-0.7019	0.1300
CG15817	CG15817	1633233_a_at	0.2125	0.3801	-0.2488	0.0798	-0.6097	0.0305	-0.0515	0.9598	0.6107	0.0302	0.6622	0.0136	0.1515	0.8510	0.0687	0.8867	-0.0828	0.8463
CG2807	CG2807	1633234_at	0.4144	0.0169	-0.0594	0.7261	0.3167	0.0836	0.1026	0.8660	0.4035	0.0637	0.3009	0.1157	-0.2956	0.6557	-0.0607	0.8745	0.2350	0.3898
---	---	1633235_at	0.0499	0.8347	0.1331	0.4661	0.2195	0.1326	0.1513	0.6341	0.0028	0.9901	-0.1485	0.2945	0.0550	0.9545	-0.0193	0.9670	-0.0743	0.8366
---	---	1633236_at	-0.0257	0.9057	0.0504	0.6288	0.1197	0.4569	0.0052	0.9943	-0.1117	0.4868	-0.1169	0.4145	-0.0762	0.9126	-0.0472	0.8980	0.0290	0.9329
Idgf1	Imaginal disc grov	1633237_at	-0.5026	0.4076	-1.1967	0.0052	-0.1762	0.4037	0.8509	0.2654	1.0447	0.0286	0.1938	0.6822	-0.1749	0.8202	0.3819	0.2156	0.5568	0.1156
GstE8 /// GstE9	Glutathione S tran	1633238_at	0.8363	0.1827	0.1182	0.7127	0.2016	0.2664	0.2452	0.4961	0.3484	0.0708	0.1033	0.5885	-0.0005	0.9999	-0.6162	0.4426	-0.6156	0.4592
r-cup	ryder cup	1633239_at	0.0072	0.9704	-0.0151	0.8998	-0.0243	0.9008	0.0882	0.8174	-0.1083	0.4908	-0.1965	0.1411	0.0532	0.9421	0.0238	0.9485	-0.0294	0.9246
---	---	1633240_at	0.2690	0.0932	0.4614	0.1121	-0.0516	0.8257	-0.3086	0.4592	-0.3586	0.1115	-0.0500	0.8484	0.2788	0.6955	0.1257	0.7058	-0.1531	0.6298
CG31340	CG31340	1633241_at	-1.2223	0.0036	-1.3580	0.0087	-1.6365	0.0002	-0.6029	0.1194	-0.0333	0.9190	0.5695	0.0144	-0.2180	0.7116	-0.1271	0.6325	0.0909	0.7492
CG3964	CG3964	1633242_a_at	0.3415	0.3206	0.0472	0.6488	0.1490	0.5672	-0.0421	0.9733	-0.0798	0.8426	-0.0377	0.9210	-0.0992	0.9144	-0.0361	0.9454	0.0631	0.8879
sced	scrambled	1633243_at	0.0096	0.9889	-1.1372	0.0221	-1.6369	0.0112	-0.2320	0.6854	0.9589	0.0041	1.1909	0.0010	0.1134	0.9767	-0.1437	0.9206	-0.2570	0.8247
CG31551	CG31551	1633244_at	0.2211	0.2165	0.0384	0.8209	0.2426	0.1321	0.0124	0.9857	-0.0188	0.9307	-0.0313	0.8619	-0.0476	0.9421	0.0094	0.9796	0.0570	0.8221
CG15529	CG15529	1633245_at	-0.8853	0.0054	-0.1152	0.7864	-0.7636	0.0026	-0.4483	0.5644	-0.8347	0.0430	-0.3864	0.2850	0.1347	0.8988	-0.1683	0.7040	-0.3030	0.4545
Dif	Dorsal-related imr	1633246_s_at	0.0014	0.9954	-0.1195	0.4924	-0.4814	0.0242	-0.2617	0.3056	0.1385	0.3673	0.4002	0.0097	0.0873	0.9011	0.0842	0.7943	-0.0030	0.9946
CG7006 /// DyakCG7006	CG7006	1633247_at	0.7405	0.0669	-0.2041	0.7116	0.1448	0.4853	0.7162	0.2871	1.5751	0.0027	0.8589	0.0233	0.3204	0.8049	0.3398	0.5471	0.0194	0.9835
CG11229	CG11229	1633248_at	-0.0517	0.7658	-0.2269	0.3278	-0.3272	0.0418	-0.1512	0.7803	0.0632	0.8223	0.2143	0.2931	-0.0544	0.9243	-0.1505	0.4300	-0.0961	0.6399
DNApol-alpha180	DNA polymerase	1633249_at	0.2660	0.2549	-0.6264	0.1760	-0.7849	0.0528	-0.2421	0.7604	0.9383	0.0139	1.1803	0.0029	-0.5033	0.6955	-0.1140	0.8830	0.3893	0.4836
CG30332	CG30332	1633250_at	0.2223	0.2904	-0.0111	0.9304	-0.0880	0.6231	-0.2347	0.5242	0.0074	0.9801	0.2421	0.1597	-0.0554	0.9238	-0.0011	0.9989	0.0543	0.8223
CG15829	CG15829	1633251_at	0.1744	0.5525	-0.3423	0.0456	0.3894	0.0728	0.4286	0.1659	0.1634	0.3970	-0.2652	0.1133	-0.0600	0.9515	-0.2051	0.4999	-0.1451	0.6484

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31714	CG31714	1633252_at	-1.6052	0.0066	-0.9982	0.0894	-1.0950	0.0008	-0.0012	0.9994	0.1748	0.6687	0.1760	0.6318	0.0119	0.9912	-0.1255	0.5855	-0.1374	0.5498
CG2991	CG2991	1633253_s_at	-0.3513	0.1541	0.4687	0.2252	-0.2663	0.4203	-0.2725	0.7010	-0.5346	0.1009	-0.2621	0.3861	0.4198	0.6780	0.1612	0.7439	-0.2586	0.5615
CG7787 /// DyakCG7787	CG7787	1633254_at	0.4509	0.0647	0.0629	0.7793	-0.1923	0.2265	-0.1379	0.6972	0.4190	0.0184	0.5568	0.0031	0.1710	0.8293	0.1396	0.7111	-0.0313	0.9466
CG33325	CG33325	1633255_at	0.0456	0.7841	0.1366	0.3189	0.1840	0.2454	-0.0571	0.9319	-0.1355	0.5223	-0.0784	0.7105	0.0910	0.8344	0.0322	0.9077	-0.0588	0.7841
---	---	1633256_at	0.1416	0.4727	0.0590	0.6403	0.1722	0.2334	-0.0080	0.9924	-0.0633	0.7640	-0.0553	0.7772	-0.0737	0.9246	-0.0609	0.8684	0.0128	0.9756
CG9973	CG9973	1633257_at	-0.0879	0.7484	-0.0386	0.7587	0.0188	0.9157	0.1001	0.8841	-0.0502	0.8718	-0.1503	0.5056	0.0571	0.9401	-0.1526	0.5299	-0.2097	0.3817
---	---	1633258_at	0.2582	0.2256	0.0337	0.8257	0.1500	0.3139	0.1445	0.7929	0.1200	0.6322	-0.0245	0.9290	-0.0820	0.8802	-0.1307	0.5477	-0.0487	0.8577
CG31406	CG31406	1633259_a_at	0.0390	0.8751	-0.0187	0.8742	0.1922	0.2458	-0.0024	0.9962	-0.0642	0.7367	-0.0619	0.7202	-0.1204	0.8424	-0.1210	0.6553	-0.0006	0.9992
CG31715	CG31715	1633260_at	-0.0423	0.8741	1.1286	0.0057	0.8597	0.0062	-0.1234	0.8800	-0.2236	0.4413	-0.1002	0.7397	0.0202	0.9921	0.7339	0.1164	0.7137	0.1483
CG9107	CG9107	1633261_a_at	0.7668	0.0210	0.4507	0.3591	0.4282	0.0788	0.2014	0.7409	0.7068	0.0167	0.5054	0.0416	-0.0127	0.9952	0.2334	0.6713	0.2461	0.6458
Lim1	---	1633262_s_at	1.6894	0.0101	0.6213	0.0745	0.0289	0.9615	0.2555	0.8281	1.3304	0.0102	1.0749	0.0160	0.7009	0.6897	0.3406	0.6647	-0.3603	0.6389
CG3121	CG3121	1633263_at	-0.0850	0.7318	0.1078	0.2981	-0.0421	0.8058	-0.3812	0.3501	-0.4080	0.0828	-0.0268	0.9253	-0.0585	0.9514	-0.0370	0.9350	0.0214	0.9582
CG13024	CG13024	1633264_at	-2.4959	0.0101	-3.9666	0.0016	-3.5461	0.0000	0.3739	0.7579	0.8543	0.1013	0.4804	0.3110	0.1241	0.9467	-0.7328	0.1821	-0.8570	0.1558
---	---	1633265_s_at	0.1348	0.4211	0.4573	0.0250	0.3868	0.0391	-0.0612	0.9162	-0.2851	0.1148	-0.2238	0.1663	-0.0137	0.9916	-0.0339	0.9462	-0.0201	0.9650
BG642378	BG642378	1633266_at	0.1115	0.5411	-0.0164	0.8742	0.0270	0.8765	-0.0918	0.8330	0.1082	0.5424	0.2000	0.1751	-0.0119	0.9898	0.0428	0.8787	0.0547	0.8237
CG12970	CG12970	1633267_a_at	0.2272	0.2434	-0.1759	0.2995	0.1944	0.3112	0.4080	0.1573	0.6856	0.0029	0.2776	0.0770	0.0256	0.9893	0.2100	0.6253	0.1844	0.6719
CG10806	serine protease	1633268_s_at	2.6445	0.3516	-0.4107	0.0962	-0.1741	0.5027	0.2099	0.9130	-0.3740	0.5680	-0.5839	0.2874	-0.0835	0.9952	-3.5719	0.3002	-3.4884	0.3369
CG9140	CG9140	1633269_at	-0.6766	0.0084	0.3353	0.1926	0.4909	0.0379	-0.1980	0.6455	-1.2515	0.0005	-1.0535	0.0006	-0.3624	0.6557	-0.2031	0.5544	0.1594	0.6541
---	---	1633270_at	0.3748	0.1236	0.0782	0.7811	0.3558	0.3570	0.0416	0.9558	0.2330	0.2488	0.1914	0.2947	0.1368	0.9396	0.2025	0.7598	0.0657	0.9330
Sox15	Sox box protein 1!	1633271_at	0.2320	0.3642	0.0000	1.0000	0.0905	0.6134	-0.0209	0.9826	0.0736	0.7883	0.0945	0.6886	-0.1573	0.7506	-0.1454	0.5047	0.0119	0.9709
CG9090 /// DsecCG9090 ///	CG9090	1633272_at	-2.0648	0.0299	-1.1591	0.0048	-1.8831	0.0001	-0.3410	0.4881	-1.5871	0.0005	-1.2461	0.0008	0.3220	0.9142	-0.8576	0.4094	-1.1797	0.2777
---	---	1633273_at	-0.0863	0.6598	-0.0428	0.6935	0.1504	0.3709	0.1499	0.7285	-0.0951	0.6645	-0.2451	0.1549	-0.0468	0.9525	-0.0288	0.9402	0.0180	0.9578
CG10158	CG10158	1633274_at	-0.3034	0.1086	0.0230	0.9028	-0.3398	0.0310	-0.1405	0.6954	-0.1536	0.3635	-0.0130	0.9503	0.1517	0.7859	0.1950	0.4107	0.0433	0.8949
CG31562	CG31562	1633275_at	0.2161	0.4152	0.1810	0.4506	-0.0683	0.7923	0.0415	0.9695	0.0005	0.9991	-0.0410	0.9030	0.1796	0.7485	-0.0647	0.8397	-0.2443	0.3163
---	---	1633276_at	0.1859	0.3135	-0.0364	0.7596	-0.0859	0.6878	-0.0058	0.9956	0.0568	0.8449	0.0626	0.8042	-0.0213	0.9893	-0.2669	0.4202	-0.2456	0.4755
RpL40	ubiquitin 52-AA ex	1633277_at	0.1956	0.2824	1.4068	0.0199	1.1557	0.0045	0.1977	0.7424	-0.8887	0.0056	-1.0864	0.0014	0.1238	0.8444	0.1095	0.7088	-0.0143	0.9714
CG34404	CG18496	1633278_at	-0.1108	0.5689	-0.0603	0.6695	-0.1565	0.4134	0.0717	0.9145	0.2012	0.3449	0.1295	0.5249	0.2994	0.5126	0.1975	0.3571	-0.1019	0.6689
CG33090	CG33090	1633279_s_at	0.0118	0.9535	0.2269	0.2816	0.3999	0.1168	-0.1900	0.8246	-0.3220	0.3401	-0.1320	0.7069	-0.1609	0.8114	0.0611	0.8785	0.2220	0.4387
CG12581	CG12581	1633280_s_at	0.4120	0.1914	0.6713	0.0785	0.8684	0.0031	-0.0212	0.9794	-0.2143	0.2928	-0.1931	0.2908	-0.0618	0.9742	0.2443	0.6408	0.3062	0.5486
CG32776	CG32776	1633281_at	0.2340	0.1474	-0.1783	0.3133	-0.2491	0.1729	0.0424	0.9627	0.5778	0.0245	0.5353	0.0211	0.0041	0.9964	0.0662	0.7891	0.0621	0.7944
CG15708	CG15708	1633282_at	0.1500	0.5876	0.0885	0.4230	0.1850	0.3597	0.1000	0.8594	-0.0696	0.7771	-0.1696	0.3657	-0.0698	0.9277	-0.2375	0.3327	-0.1677	0.5257
CG6764	CG6764	1633283_at	0.2502	0.1047	1.0315	0.0126	0.9918	0.0012	0.3740	0.1526	-0.6296	0.0027	-1.0036	0.0002	0.2247	0.7220	0.0487	0.9064	-0.1761	0.5355
eIF6	eIF6	1633284_at	0.3392	0.4214	0.1434	0.7383	0.1448	0.3368	0.3776	0.5617	0.7649	0.0295	0.3873	0.1937	0.3437	0.7644	0.5801	0.2126	0.2365	0.6476
y	yellow	1633285_at	0.0899	0.6679	-0.0376	0.7127	-0.0658	0.7328	-0.0635	0.9209	-0.0413	0.8723	0.0222	0.9253	-0.0071	0.9952	-0.0643	0.8678	-0.0572	0.8779
CG32372	CG32372	1633286_at	1.0011	0.2520	1.2713	0.1937	0.8580	0.0311	-0.5910	0.6041	-0.6495	0.2594	-0.0585	0.9344	-0.3067	0.9449	-0.4045	0.8166	-0.0978	0.9599
Adam	Complementation	1633287_at	0.2061	0.1914	0.5791	0.0461	0.4407	0.0158	-0.0338	0.9515	0.1144	0.4752	0.1482	0.2846	0.0731	0.9174	0.4517	0.0782	0.3786	0.1479
CG31763	CG31763	1633288_at	0.4793	0.2883	0.3643	0.4858	0.2102	0.4216	0.1623	0.6493	0.2939	0.0889	0.1316	0.4140	0.5066	0.8202	0.3691	0.7285	-0.1375	0.9135
Or85f	Odorant receptor	1633289_at	-0.0348	0.8796	0.1098	0.5873	0.1423	0.4616	-0.0712	0.8987	-0.1928	0.2960	-0.1216	0.4863	-0.0503	0.9734	-0.0824	0.8769	-0.0321	0.9512
---	---	1633290_a_at	-2.0910	0.0102	-1.4613	0.0035	-1.8140	0.0024	0.2402	0.7310	0.1573	0.6555	-0.0830	0.8135	0.2360	0.8206	0.4583	0.2722	0.2223	0.6328
dy	dusky	1633291_at	-1.6072	0.0296	-0.2950	0.3258	-1.1783	0.0011	-0.5724	0.5311	-1.3692	0.0113	-0.7968	0.0648	0.2076	0.8008	-0.1909	0.6011	-0.3985	0.2611
CG4300	CG4300	1633292_a_at	-0.0515	0.7570	0.6745	0.0666	1.0635	0.0014	0.0390	0.9540	-0.4636	0.0207	-0.5026	0.0089	-0.3397	0.7196	0.2853	0.4743	0.6250	0.1393
CG10157	CG10157	1633293_at	0.3186	0.2775	0.9759	0.1411	1.0871	0.0044	-0.3348	0.6429	-1.7316	0.0011	-1.3968	0.0016	-0.4970	0.7464	-1.0673	0.1078	-0.5702	0.3915
CG2812	CG2812	1633294_at	-1.2185	0.0011	-0.7275	0.0707	-0.7049	0.0098	0.2992	0.5037	-0.2819	0.2341	-0.5810	0.0140	0.0711	0.9427	0.0902	0.8208	0.0191	0.9666
mthl13	methuselah-like 1	1633295_a_at	0.0254	0.9013	-0.0301	0.7783	0.0494	0.8050	0.1666	0.7325	0.2363	0.2790	0.0698	0.7712	-0.2318	0.6935	-0.1888	0.4336	0.0430	0.8972
CG13293	CG13293	1633296_at	0.2262	0.2409	0.1504	0.5842	0.4429	0.0087	0.1736	0.5317	0.0050	0.9816	-0.1686	0.1925	-0.0391	0.9778	0.0055	0.9941	0.0446	0.9231
---	---	1633297_at	0.2215	0.8025	-0.2535	0.6276	-0.4578	0.1997	-0.0830	0.9602	0.4557	0.3013	0.5387	0.1665	-0.1297	0.9742	-0.2976	0.8220	-0.1678	0.9048
mei-P22	meiotic P22	1633298_at	0.2196	0.1559	-0.2600	0.3776	-0.0094	0.9635	0.1793	0.5756	0.3458	0.0416	0.1665	0.2602	0.0037	0.9964	-0.0783	0.7347	-0.0820	0.7141
lz	lozenge	1633299_at	0.6279	0.5202	0.9261	0.2921	0.5395	0.0140	0.0439	0.9705	-0.3171	0.3032	-0.3610	0.1848	0.4780	0.8940	-0.0198	0.9942	-0.4978	0.7484
CG13282	CG13282	1633300_at	-3.3876	0.0016	-3.8892	0.0077	-4.0439	0.0000	0.0443	0.9733	0.1478	0.6943	0.1035	0.7751	0.0476	0.9898	-0.5696	0.4999	-0.6172	0.4683
CG3436	CG3436	1633301_at	-0.2641	0.1795	-0.5000	0.1648	-0.3903	0.2150	-0.2407	0.6698	0.3240	0.2260	0.5647	0.0265	-0.4088	0.7464	0.0206	0.9841	0.4294	0.4301
CG40211	CG40211	1633302_a_at	-0.0138	0.9467	0.0305	0.7602	0.0968	0.5619	0.0633	0.8908	-0.0098	0.9676	-0.0732	0.6462	0.0976	0.8692	0.0046	0.9935	-0.0930	0.7287
CG2989	CG2989	1633303_at	-0.2334	0.3709	-0.1258	0.5052	-0.2132	0.2327	0.4073	0.2413	0.3501	0.0895	-0.0572	0.8046	0.2145	0.7485	0.1765	0.5530	-0.0379	0.9231
p24-1	p24-related-1	1633304_at	0.8857	0.0205	2.0798	0.0048	2.2218	0.0000	0.2599	0.5046	-0.2999	0.1450	-0.5598	0.0089	-0.0024	0.9994	0.9526	0.0677	0.9550	0.0852
Obp99a	Odorant-binding p	1633305_at	2.6553	0.0132	1.6870	0.2198	3.9544	0.0003	1.1812	0.3256	0.5743	0.4260	-0.6069	0.3423	-1.0915	0.7565	-0.4551	0.8132	0.6364	0.7026
ms(2)35Ci	male sterile(2)46A	1633306_at	-0.2859	0.2849	0.0864	0.5348	0.0374	0.8339	0.0265	0.9761	-0.2089	0.3510	-0.2354	0.2323	0.0277	0.9764	0.1464	0.5287	0.1186	0.6228
CG13720	CG13720	1633307_at	0.1535	0.4669	0.0000	1.0000	-0.0660	0.6853	-0.0280	0.9695	-0.0068	0.9809	0.0211	0.9268	0.1235	0.8262	0.0716	0.8123	-0.0519	0.8666
CG6660	CG6660	1633308_at	0.5726	0.6903	-1.3281	0.4982	1.0990	0.1708	1.8378	0.2334	0.7656	0.4187	-1.0723	0.1896	-0.5589	0.9467	-1.0050	0.7319	-0.4461	0.8982
CG40374	CG40374	1633309_at	0.1799	0.3418	0.0555	0.6100	0.3953	0.1189	0.0871	0.8189	0.0190	0.9277	-0.0681	0.6590	-0.0754	0.8940	-0.0925	0.6969	-0.0171	0.9545
CG15483	CG15483	1633310_at	0.1065	0.7075	0.3558	0.1501	0.2094	0.2920	-0.0906	0.8933	-0.1746	0.4560	-0.0840	0.7267	0.0649	0.9514	0.0802	0.8550	0.0153	0.9762
Scgdelta	Sarcoglycan delta	1633311_at	-0.2047	0.3543	-0.0791	0.5548	0.1156	0.6812	-0.0741	0.9380	0.0192	0.9642	0.0933	0.7612	-0.1818	0.7220	0.0284	0.9349	0.2102	0.3457
CG6962	CG6962	1633312_at	-0.0503	0.8162	-1.0402	0.0866	-0.4318	0.2459	0.0530	0.9728	0.8595	0.0334	0.8065	0.0275	-0.6082	0.6557	-0.1228	0.8754	0.4854	0.3844
---	---	1633313_at	-2.3373	0.0019	-0.4802	0.2390	-1.2403	0.0050	-0.6892	0.2915	-1.5395	0.0026	-0.8504	0.0215	-0.1851	0.7779	-0.2677	0.3346	-0.0826	0.8161
CG15605	CG15605	1633314_at	0.0665	0.6804	0.0052	0.9709	0.1043	0.6555	-0.0857	0.9371	-0.0222	0.9632	0.0635	0.8625	-0.0633	0.9460	-0.1120	0.7354	-0.0486	0.9020
CG2006	CG2006	1633315_at	0.4136	0.0980	0.1782	0.5401	0.0088	0.9849	-0.0378	0.9705	0.6134	0.0269	0.6512	0.0132	0.1142	0.9352	0.4103	0.3472	0.2960	0.5299
CG30490	CG30490	1633316_at	0.3055	0.0797	0.0124	0.9388	0.2258	0.1782	0.0040	0.9956	0.0461	0.8128	0.0421	0.8130	-0.0551	0.9249	-0.0718	0.7599	-0.0168	0.9517
CG40216	CG40216	1633317_at	0.1453	0.5856	-0.1740	0.2496	-0.0945	0.6228	-0.0318	0.9674	0.0427	0.8754	0.0745	0.7355	-0.0025	0.9984	-0.0975	0.7462	-0.0950	0.7484
CG6854	CG6854	1633318_at	0.1239	0.4927	-0.0480	0.6363	0.0609	0.7657	0.2224	0.5376	0.2849	0.1295	0.0625	0.7623	0.0937	0.9168	0.0130	0.9825	-0.0807	0.8373
CG14956	CG14956	1633319_at	-0.3565	0.4060	-0.6694	0.3228	-0.9061	0.0080	-0.4573	0.3909	-0.0794	0.8372	0.3779	0.1588	-0.1450	0.9589	-0.2336	0.8262	-0.0886	0.9374
Srp54k	Signal recognition	1633320_at	0.5701	0.0221	1.1921	0.0232	1.4738	0.0001	-0.1390	0.7304	-0.3636	0.0490	-0.2246	0.1627	-0.2406	0.7893	0.3996	0.2830	0.6402	0.1277

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11769	CG11769	1633321_at	0.1348	0.4735	0.0495	0.7450	0.2761	0.1753	0.1300	0.7617	0.0452	0.8500	-0.0848	0.6568	-0.0126	0.9939	-0.0932	0.8516	-0.0806	0.8674
CG2972	CG2972	1633322_at	0.5824	0.0230	0.4601	0.3389	0.5995	0.0187	0.1905	0.6541	0.5106	0.0210	0.3201	0.0821	-0.0644	0.9742	0.2972	0.5755	0.3616	0.4863
CG3916	CG3916	1633323_at	0.2740	0.4828	0.5564	0.0098	0.4389	0.0460	0.0416	0.9649	-0.1444	0.6036	-0.1860	0.4352	0.0164	0.9928	0.2729	0.5259	0.2565	0.5612
CG12590	CG12590	1633324_at	0.1843	0.2669	-0.0205	0.8418	0.3097	0.0924	0.3018	0.4221	0.3573	0.0869	0.0555	0.8132	-0.1149	0.8283	0.0224	0.9504	0.1373	0.5605
---	---	1633325_at	0.0221	0.9017	-0.0276	0.8886	0.1213	0.7144	0.1233	0.8837	0.1139	0.7364	-0.0094	0.9793	-0.0417	0.9816	0.0794	0.8857	0.1212	0.7919
CG15356	CG15356	1633326_at	0.0881	0.8247	0.3020	0.1538	0.4890	0.0129	0.1459	0.7673	-0.1872	0.3832	-0.3331	0.0773	0.0470	0.9530	0.1046	0.6965	0.0575	0.8517
CG11023	CG11023	1633327_at	-0.0245	0.9467	-0.0019	0.9879	0.1022	0.5210	-0.2008	0.7380	-0.0862	0.7908	0.1146	0.6800	-0.0553	0.9405	-0.0183	0.9617	0.0370	0.9065
Rpl12	Rpl12	1633328_at	-0.0015	0.9973	-0.3548	0.0857	-0.3416	0.0960	0.0746	0.9422	0.4739	0.1045	0.3993	0.1254	-0.1287	0.9012	0.0825	0.8794	0.2112	0.6052
yellow-h	yellow-h	1633329_at	-1.6528	0.0021	-1.5141	0.1187	-2.1009	0.0000	-0.3454	0.3744	-0.9228	0.0021	-0.5774	0.0103	0.0557	0.9884	-0.6840	0.4073	-0.7397	0.3847
---	---	1633330_at	0.1288	0.4785	0.0014	0.9913	0.1532	0.3964	-0.1142	0.7529	-0.0019	0.9934	0.1123	0.4479	-0.1819	0.7726	-0.0975	0.7677	0.0843	0.7978
dom	domino	1633331_at	0.6182	0.1608	0.4593	0.0664	-0.0331	0.8976	-0.2055	0.6086	0.5716	0.0120	0.7771	0.0018	0.0609	0.9824	0.2810	0.6955	0.2202	0.7677
---	---	1633332_at	0.1678	0.4060	0.0372	0.7017	0.0261	0.9067	0.0490	0.9314	0.0275	0.9069	-0.0215	0.9175	0.0854	0.8846	-0.0805	0.7653	-0.1659	0.4710
CG7231	CG7231	1633333_a_at	-0.0617	0.8794	-1.2792	0.0230	-1.3448	0.0001	-0.0337	0.9658	0.3813	0.0699	0.4151	0.0330	-0.0743	0.9589	-0.7532	0.0795	-0.6789	0.1303
CG15822	CG15822	1633334_at	0.3987	0.5258	-0.2832	0.7007	0.0141	0.9792	-0.0261	0.9934	0.2215	0.7625	0.2476	0.7007	-0.3179	0.8940	-0.4368	0.6545	-0.1189	0.9231
Sos	lethal 25 in the bla	1633335_at	-0.2184	0.7168	-0.1987	0.8172	-0.4107	0.1803	0.0050	0.9956	0.3883	0.1481	0.3833	0.1108	0.3551	0.8940	0.3582	0.7653	0.0030	0.9989
CG32560 /// DmirCG32560	CG32560	1633336_at	-1.0062	0.0737	0.2752	0.7853	-0.1843	0.4814	-0.3779	0.2876	-1.0163	0.0012	-0.6384	0.0052	0.0828	0.9869	0.2926	0.8518	0.2098	0.8934
CG17386	CG17386	1633337_at	0.0459	0.8023	0.1161	0.6807	0.2614	0.1334	0.2201	0.6615	0.0696	0.8157	-0.1505	0.5201	0.0280	0.9814	0.0644	0.8546	0.0364	0.9176
mRpS23	mRpS23	1633338_at	-0.1483	0.4444	-0.3731	0.2109	-0.5137	0.0174	-0.0619	0.9371	0.1074	0.6796	0.1692	0.4287	-0.0967	0.9063	-0.1334	0.6847	-0.0367	0.9283
---	---	1633339_at	0.0165	0.9222	-0.1906	0.3496	0.0344	0.8709	0.1142	0.7795	0.3268	0.0583	0.2126	0.1599	-0.0553	0.9342	-0.0359	0.9138	0.0194	0.9486
CG1887	CG1887	1633340_at	0.0057	0.9818	0.2533	0.1944	-0.5683	0.0685	-0.3052	0.2792	-0.3194	0.0591	-0.0142	0.9461	0.2783	0.8023	0.0916	0.8940	-0.1867	0.7228
dac	dachshund	1633341_s_at	0.0754	0.9287	1.1795	0.2536	0.0809	0.8451	-0.5037	0.6166	-1.1969	0.0253	-0.6932	0.1212	0.5157	0.8740	0.0150	0.9959	-0.5007	0.7325
---	---	1633342_at	0.0223	0.9218	0.0263	0.8754	0.1533	0.5211	0.3945	0.2506	0.1291	0.5579	-0.2654	0.1453	-0.0783	0.8999	-0.0786	0.7802	-0.0003	0.9992
CG5250	CG5250	1633343_at	0.1531	0.3435	-0.0307	0.8212	0.0136	0.9477	0.1179	0.8424	0.1653	0.4728	0.0474	0.8530	0.0137	0.9892	0.0346	0.9157	0.0208	0.9434
Cpr92F	CG5494	1633344_at	0.1985	0.3112	0.2149	0.3575	0.2534	0.1907	0.0050	0.9943	-0.0671	0.6900	-0.0720	0.6303	-0.0091	0.9940	-0.0194	0.9663	-0.0104	0.9833
Scr	sex-combs reduce	1633345_at	0.0912	0.7026	-0.0875	0.4097	0.1051	0.4556	0.1813	0.6354	0.1885	0.3212	0.0072	0.9756	-0.0484	0.9441	-0.0743	0.7707	-0.0258	0.9295
CG7351	CG7351	1633346_at	0.3512	0.1444	-0.0121	0.9721	0.1875	0.2829	-0.0461	0.9345	0.2576	0.1166	0.3037	0.0450	-0.2556	0.7726	-0.1178	0.8123	0.1378	0.7544
CG34001	CG34001	1633347_at	0.5908	0.0239	1.0504	0.0165	0.6999	0.0124	-0.0694	0.9154	0.2606	0.2013	0.3300	0.0734	0.2396	0.8049	0.4249	0.2790	0.1852	0.6773
---	---	1633348_s_at	0.2117	0.3903	0.4103	0.0935	0.3972	0.0491	0.0831	0.8794	-0.1571	0.4160	-0.2402	0.1499	0.1263	0.8870	0.1195	0.7672	-0.0068	0.9903
TfIIFbeta	TFIIF small subun	1633349_at	0.4701	0.0281	0.2193	0.3715	0.2753	0.0903	0.0204	0.9745	0.3509	0.0320	0.3305	0.0261	-0.0597	0.9545	0.0851	0.8359	0.1448	0.6644
---	---	1633350_at	0.0054	0.9783	0.0974	0.3764	-0.1774	0.4196	-0.0176	0.9777	0.0613	0.7510	0.0789	0.6352	0.2346	0.7220	0.1141	0.7260	-0.1205	0.7030
CG10384	CG10384	1633351_at	0.1597	0.4663	-0.0960	0.4078	-0.0344	0.8665	0.0339	0.9639	0.0528	0.8340	0.0189	0.9385	-0.1668	0.7241	-0.2804	0.1584	-0.1136	0.6052
Imp	KH-domain protein	1633352_at	0.0334	0.8344	-0.0676	0.5662	-0.0172	0.9403	0.0377	0.9466	0.0392	0.8495	0.0015	0.9940	0.0196	0.9816	-0.0348	0.9068	-0.0544	0.8228
CAP	CAP	1633353_s_at	-0.3728	0.1674	-0.0153	0.8884	-0.2218	0.1278	-0.2876	0.4511	-0.4746	0.0296	-0.1870	0.3229	-0.1723	0.8424	-0.1450	0.7209	0.0273	0.9575
Cyp6v1	Cyp6v1	1633354_at	0.1273	0.6595	1.0625	0.0523	0.0686	0.7618	0.3077	0.2977	0.3987	0.0294	0.0910	0.6027	1.1509	0.2553	1.1942	0.0405	0.0433	0.9517
CG10802	CG10802	1633355_at	-0.4138	0.2716	0.6830	0.0536	1.1794	0.0041	-0.0603	0.9477	-0.7299	0.0123	-0.6696	0.0111	-0.5506	0.6557	0.4107	0.4073	0.9612	0.0892
nAcRalpha-96Ab	Dalpha2	1633356_s_at	0.1701	0.4222	0.2387	0.2191	0.2529	0.2641	0.0334	0.9619	-0.1007	0.6244	-0.1341	0.4408	0.0062	0.9952	-0.0608	0.8556	-0.0670	0.8259
CG7910	CG7910	1633357_at	0.1994	0.9061	-1.6423	0.4966	1.5679	0.1692	2.1285	0.2810	0.4239	0.7613	-1.7045	0.1009	-1.0323	0.8953	-1.4732	0.6425	-0.4410	0.9125
---	---	1633358_at	0.0243	0.9261	-0.0497	0.6426	0.1067	0.6465	0.1417	0.8578	0.2513	0.3906	0.1095	0.7183	0.0185	0.9852	0.1260	0.6061	0.1075	0.6659
Traf3	Traf3	1633359_a_at	-0.4395	0.2364	-0.3642	0.2911	-0.8366	0.0009	-0.2393	0.7614	0.1586	0.6818	0.3979	0.1886	0.1240	0.8740	0.1174	0.7454	-0.0067	0.9894
CG13999	CG13999	1633360_at	-0.0111	0.9629	0.2197	0.1740	0.2504	0.2132	-0.1422	0.7760	-0.1777	0.4090	-0.0356	0.8862	0.0417	0.9734	0.0599	0.8962	0.0182	0.9691
CG15025	CG15025	1633361_at	-0.0877	0.6868	-0.0721	0.5836	0.1343	0.3854	0.0299	0.9637	0.0110	0.9676	-0.0189	0.9288	-0.1580	0.7970	0.0496	0.8983	0.2075	0.4322
CG17565	CG17565	1633362_at	-0.3720	0.2789	0.4048	0.0917	0.9466	0.0013	0.1966	0.7205	-0.4352	0.0812	-0.6318	0.0120	-0.2735	0.7633	0.4661	0.2043	0.7396	0.0860
CG6428	CG6428	1633363_at	0.8108	0.0146	0.0411	0.8546	0.4463	0.0265	0.4127	0.1913	0.7853	0.0024	0.3726	0.0361	0.0272	0.9841	0.0292	0.9531	0.0020	0.9975
CG14184	CG14184	1633364_at	0.0937	0.5506	-0.1030	0.6314	-0.2251	0.4165	-0.1396	0.7627	-0.2353	0.2358	-0.0956	0.6384	0.0365	0.9852	-0.2998	0.5191	-0.3363	0.4684
CG7385	CG7385	1633365_at	-0.0252	0.9150	0.1644	0.3937	0.2278	0.2782	0.0096	0.9893	-0.0974	0.5811	-0.1070	0.4919	0.0503	0.9751	0.1490	0.7518	0.0987	0.8467
CG12455	CG12455	1633366_at	-1.9496	0.0012	-1.1215	0.0200	-1.9863	0.0001	-0.9942	0.0154	-0.4734	0.0333	0.5208	0.0141	-0.0092	0.9964	0.0472	0.9505	0.0564	0.9320
CG6340	CG6340	1633367_at	0.1259	0.5561	0.4205	0.0310	0.2292	0.1861	-0.2919	0.5096	-0.2597	0.2700	0.0322	0.9098	0.1184	0.8331	0.0726	0.8117	-0.0457	0.8874
tlk	Tousled-like kinas	1633368_at	-0.0787	0.6454	-0.0482	0.6453	-0.1335	0.4087	-0.0163	0.9803	-0.0195	0.9313	-0.0033	0.9873	0.0318	0.9644	-0.0164	0.9590	-0.0482	0.8447
CG6184	CG6184	1633369_s_at	-0.1784	0.2645	0.1614	0.4032	0.4526	0.0446	0.0639	0.9186	-0.3745	0.0566	-0.4385	0.0192	0.0020	0.9986	0.0147	0.9671	0.0127	0.9697
---	---	1633370_s_at	-0.3700	0.4544	-0.8230	0.0364	-0.3375	0.1928	0.4070	0.4815	1.0668	0.0050	0.6598	0.0253	-0.1796	0.9242	0.5496	0.3738	0.7292	0.2622
CG17684	CG40072	1633371_at	-0.0443	0.8306	0.0101	0.9273	-0.0269	0.8834	0.0827	0.8968	0.1149	0.6235	0.0321	0.8992	0.0270	0.9742	0.1825	0.3757	0.1556	0.4742
---	---	1633372_at	-0.0750	0.6881	0.0540	0.6296	-0.0947	0.7660	-0.3210	0.5255	-0.2142	0.4356	0.1068	0.7022	0.0027	0.9984	-0.0063	0.9925	-0.0090	0.9847
CG2750	CG2750	1633373_at	0.0543	0.7468	0.0433	0.8062	-0.0406	0.8583	-0.0597	0.8959	0.0007	0.9976	0.0604	0.7045	-0.0179	0.9898	0.1029	0.7836	0.1208	0.7259
---	---	1633374_at	0.0610	0.6961	0.0832	0.4099	0.0182	0.9268	-0.0156	0.9801	0.0198	0.9247	0.0354	0.8374	0.0718	0.9445	0.0695	0.8760	-0.0022	0.9972
CG5026	CG5026	1633375_at	-0.2124	0.5038	-0.0232	0.9220	-0.3792	0.0636	-0.2714	0.6391	-0.1696	0.5816	0.1018	0.7384	0.3047	0.7628	0.1187	0.8319	-0.1861	0.6886
---	---	1633376_at	0.0145	0.9528	0.1296	0.2260	0.1707	0.2834	-0.0192	0.9808	-0.0468	0.8530	-0.0276	0.9071	-0.0023	0.9990	0.0723	0.8501	0.0746	0.8336
---	---	1633377_at	0.1817	0.2525	-0.0180	0.9517	0.1045	0.6590	0.0896	0.9371	0.2057	0.5627	0.1161	0.7441	0.0827	0.9342	0.0517	0.9170	-0.0310	0.9444
Lectin-galC1	galactin	1633378_at	0.5306	0.7587	-1.7948	0.4658	1.4995	0.0758	2.7687	0.1749	0.9685	0.4518	-1.8002	0.1029	-0.6540	0.9465	-1.4361	0.6552	-0.7821	0.8334
cnc	cap and collar	1633379_s_at	-0.0972	0.8274	-0.3501	0.1791	-0.5958	0.0082	0.0075	0.9922	0.2715	0.0826	0.2640	0.0617	0.1540	0.9309	-0.0237	0.9829	-0.1778	0.7983
CG12128	CG12128	1633380_at	-0.1645	0.4900	0.6828	0.0795	1.1535	0.0005	0.1361	0.7931	-0.5559	0.0169	-0.6921	0.0038	-0.1417	0.8480	0.3167	0.2901	0.4585	0.1612
Zpr1	Zpr1	1633381_at	0.3163	0.1095	0.0741	0.8687	0.2856	0.1343	0.2219	0.5633	0.5680	0.0112	0.3461	0.0541	0.0434	0.9816	0.3893	0.3695	0.3459	0.4463
CG8920	CG8920	1633382_s_at	-0.2979	0.2865	0.2681	0.0755	-0.2539	0.1125	-0.0431	0.9427	-0.0193	0.9366	0.0238	0.9085	0.2764	0.7644	0.4906	0.1979	0.2142	0.6144
toe	twin of eyegone	1633383_at	0.0292	0.9031	0.2815	0.4262	0.1336	0.6200	0.0352	0.9695	-0.2953	0.2127	-0.3305	0.1187	0.0482	0.9812	-0.0252	0.9727	-0.0733	0.9030
Mpp6	M-phase phospho	1633384_at	0.0996	0.6849	-0.1417	0.4786	-0.0135	0.9580	-0.0863	0.8738	0.2694	0.1522	0.3557	0.0416	-0.1083	0.8973	0.1154	0.7578	0.2237	0.4898
Karl	Karl	1633385_at	-2.7309	0.0073	-3.2083	0.0259	-4.0917	0.0000	-0.4962	0.4599	-0.8712	0.0247	-0.3751	0.2513	0.2827	0.9498	-1.3363	0.3022	-1.6190	0.2403
mthl8	Mth-like 8	1633386_s_at	0.1672	0.3630	-0.0451	0.7400	0.2971	0.2077	0.0639	0.9293	-0.0716	0.7795	-0.1355	0.5054	-0.1367	0.8510	-0.2244	0.4591	-0.0877	0.8152
bves	bves	1633387_at	-0.4466	0.2382	0.0490	0.9399	-0.7560	0.0027	0.0612	0.9375	0.4190	0.0676	0.3578	0.0788	0.8200	0.5126	0.7666	0.1849	-0.0534	0.9491
---	---	1633388_at	0.0787	0.6973	0.2048	0.3036	0.4922	0.0268	-0.0660	0.8899	-0.1999	0.2092	-0.1338	0.3631	-0.2718	0.7681	0.0366	0.9534	0.3083	0.4443
CG12404	CG12404	1633389_at	0.7341	0.0150	0.9294	0.0305	1.1201	0.0002	0.0583	0.9071	0.0129	0.9569	-0.0455	0.8017	-0.1921	0.8284	0.1682	0.6855	0.3604	0.3468

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9056	CG9056	1633390_at	-0.3591	0.5548	-0.0192	0.9767	-0.1032	0.5851	-0.0952	0.9108	-0.0801	0.8096	0.0151	0.9648	0.0557	0.9898	0.4360	0.6605	0.3803	0.7083
Irc	Immune-regulatec	1633391_at	0.2024	0.4082	-1.2106	0.0365	-0.7929	0.0069	-0.0067	0.9943	0.7751	0.0033	0.7818	0.0019	-0.3669	0.7485	-0.5224	0.2645	-0.1554	0.7887
---	---	1633392_at	0.0581	0.8271	-0.1574	0.4475	-0.2410	0.1219	-0.0928	0.8623	0.1704	0.3826	0.2633	0.1212	-0.1230	0.8298	0.0336	0.9291	0.1567	0.5358
CG6499	CG6499	1633393_at	-0.1404	0.5698	-0.0986	0.4737	0.1882	0.1794	0.3268	0.2661	0.1478	0.4090	-0.1790	0.2516	-0.0361	0.9751	-0.0094	0.9870	0.0267	0.9470
CG13900 /// DmirCG13900	CG13900	1633394_a_at	0.8897	0.0636	0.8120	0.0478	0.6637	0.0114	0.1692	0.8202	0.7480	0.0186	0.5788	0.0336	0.4323	0.7324	0.6483	0.2168	0.2161	0.7271
---	---	1633395_at	0.0210	0.9458	0.0009	0.9980	-0.2361	0.2626	-0.1000	0.8967	0.0313	0.9298	0.1313	0.6083	0.1391	0.8682	-0.0024	0.9985	-0.1415	0.7035
btd	buttonhead	1633396_at	-2.8158	0.0021	-0.3798	0.1564	-2.6649	0.0002	-2.0631	0.0358	-2.9143	0.0008	-0.8512	0.0804	-0.0042	0.9990	-0.4606	0.3926	-0.4564	0.4113
---	---	1633397_at	0.0912	0.6138	0.1481	0.3420	0.1749	0.3708	0.0471	0.9466	0.0419	0.8736	-0.0052	0.9837	-0.0081	0.9939	0.0721	0.8229	0.0802	0.7836
CG2616	CG2616	1633398_at	0.7748	0.0056	0.6254	0.1160	1.1490	0.0002	-0.0150	0.9889	-0.0207	0.9554	-0.0057	0.9856	-0.6724	0.1628	0.1642	0.4958	0.8366	0.0238
DnaJ-60	DnaJ-like-60	1633399_at	-0.6461	0.0451	-0.4787	0.0747	-0.8492	0.0037	-0.0961	0.8852	-0.5800	0.0185	-0.4839	0.0248	0.1731	0.8609	-0.4002	0.3040	-0.5734	0.1767
cue	cueball	1633400_at	1.0698	0.0114	0.5837	0.0201	0.3901	0.2650	-0.0101	0.9917	0.6524	0.0072	0.6625	0.0040	0.0501	0.9862	0.0613	0.9550	0.0112	0.9924
Cyp12d1-d /// Cyp12d1-p	Cyp12d1-p /// Cyp	1633401_s_at	2.0217	0.0020	1.8418	0.0048	3.4620	0.0000	0.7686	0.1659	-1.7438	0.0009	-2.5124	0.0001	-0.4880	0.6955	-1.7237	0.0204	-1.2357	0.0559
CG31684	CG31684	1633402_at	0.2094	0.3969	0.0525	0.6581	0.1261	0.6356	-0.0179	0.9854	-0.0775	0.7696	-0.0596	0.8102	-0.0478	0.9619	0.0278	0.9506	0.0756	0.8299
Socs44A	Suppressor of Cyt	1633403_at	0.0072	0.9709	0.1895	0.4045	-0.0955	0.7647	-0.0057	0.9943	0.4094	0.0238	0.4150	0.0140	0.2868	0.7953	0.6034	0.1795	0.3166	0.5123
CG9965	CG9965	1633404_at	0.0908	0.5934	0.1813	0.2378	-0.0210	0.9084	0.0440	0.9518	0.0705	0.7712	0.0264	0.9143	0.0654	0.9412	0.0717	0.8481	0.0063	0.9886
CG4288	CG4288	1633405_s_at	0.8637	0.1128	0.2467	0.3474	0.3432	0.1556	-0.0476	0.9592	-0.1195	0.6675	-0.0720	0.7932	-0.0157	0.9946	-0.2158	0.7221	-0.2000	0.7434
---	---	1633406_at	-0.1295	0.4582	0.0084	0.9700	-0.0629	0.7915	-0.1403	0.7278	-0.2194	0.2234	-0.0791	0.6749	0.0152	0.9913	0.0116	0.9829	-0.0035	0.9945
CG6959	connectin-like	1633407_at	-1.4726	0.0012	-0.7504	0.0472	-1.6485	0.0006	-0.1727	0.7355	-0.3676	0.1048	-0.1949	0.3501	0.7918	0.3712	0.2686	0.5972	-0.5231	0.2869
CG14424	CG14424	1633408_at	0.2432	0.2266	0.0767	0.5230	0.0287	0.9130	0.0051	0.9956	0.0289	0.9220	0.0238	0.9263	0.0379	0.9611	-0.1243	0.5888	-0.1622	0.4687
CG10598	CG10598	1633409_a_at	0.1517	0.4473	0.0728	0.5348	0.2078	0.3219	-0.0655	0.8987	-0.2002	0.2345	-0.1347	0.3892	0.1095	0.8609	-0.1178	0.6743	-0.2273	0.3807
CG2975	CG2975	1633410_at	-1.8108	0.0018	-0.9019	0.0372	-1.4076	0.0014	-0.3711	0.6247	-0.8497	0.0312	-0.4786	0.1531	0.4111	0.7215	0.0505	0.9492	-0.3606	0.4726
CG12546 /// CG14452	CG14452 /// CG1	1633411_s_at	-0.1082	0.6849	0.0122	0.9612	0.1796	0.2505	0.0318	0.9672	-0.1680	0.4231	-0.1998	0.2755	-0.0940	0.8692	-0.1638	0.4753	-0.0699	0.7980
CG3249	PKA anchor prote	1633412_s_at	0.5486	0.1329	0.6049	0.1149	0.6067	0.0558	-0.0046	0.9956	-0.0827	0.7315	-0.0781	0.7214	0.0763	0.9742	0.0742	0.9373	-0.0021	0.9987
---	---	1633413_s_at	-0.0410	0.7960	0.0500	0.7234	0.2009	0.2721	0.0431	0.9502	0.0626	0.7892	0.0194	0.9348	0.0302	0.9677	0.0605	0.8123	0.0303	0.9109
CG4328	CG4328	1633414_at	0.2899	0.1142	0.1043	0.5810	0.0644	0.7546	-0.1014	0.8385	-0.0337	0.8945	0.0678	0.7355	0.0739	0.9421	-0.0317	0.9504	-0.1056	0.7756
dpr18	dpr18	1633415_at	-0.1586	0.6753	-0.5081	0.1081	-0.6305	0.0171	-0.0423	0.9677	0.2561	0.3621	0.2983	0.2262	-0.2576	0.8270	-0.3787	0.4355	-0.1211	0.8461
---	---	1633416_s_at	0.2451	0.3442	0.0560	0.6303	0.2098	0.2457	0.2448	0.5735	0.2192	0.3331	-0.0256	0.9253	0.1093	0.8846	-0.0635	0.8764	-0.1728	0.5684
gnu	giant nuclei	1633417_at	0.7118	0.5879	-2.5238	0.1204	-1.4173	0.1018	0.3535	0.4455	3.3576	0.0000	3.0041	0.0000	-0.8646	0.8882	-0.1025	0.9811	0.7620	0.7810
CG9666	CG9666	1633418_at	-0.1137	0.6224	0.5655	0.0999	0.7754	0.0029	0.0167	0.9863	-0.6478	0.0138	-0.6645	0.0075	-0.1736	0.8342	0.1177	0.7829	0.2913	0.4075
CG7341	CG7341	1633419_s_at	0.6922	0.0289	0.3091	0.1474	0.5170	0.0140	-0.2370	0.6954	-0.1471	0.6381	0.0899	0.7709	-0.1955	0.7070	-0.0620	0.8287	0.1335	0.5615
ldlCp	ldlCp-related prote	1633420_at	0.3043	0.2861	-0.1242	0.8423	-0.7327	0.0091	-0.4672	0.6015	0.3144	0.5101	0.7816	0.0570	0.2835	0.7644	-0.0116	0.9895	-0.2951	0.4791
CG8584	CG8584	1633421_at	0.0708	0.7722	-0.0179	0.8899	0.0248	0.9082	-0.0088	0.9915	0.0544	0.7991	0.0631	0.7361	0.0192	0.9848	0.0186	0.9617	-0.0006	0.9992
lola	longitudinals abse	1633422_a_at	-0.6777	0.3988	0.5146	0.3754	0.4497	0.0960	0.1123	0.9540	-0.2376	0.6999	-0.3499	0.4959	0.2169	0.9305	0.7846	0.3152	0.5676	0.4944
---	---	1633423_at	0.0784	0.7888	0.1468	0.5119	0.3884	0.0545	0.0335	0.9745	-0.0733	0.8183	-0.1068	0.6918	-0.3266	0.7100	-0.0925	0.8546	0.2340	0.5463
CG33298	CG33298	1633424_at	0.1474	0.3389	0.2687	0.2255	0.1181	0.6008	-0.1398	0.7479	-0.1736	0.3701	-0.0338	0.8804	-0.0662	0.9499	-0.0784	0.8550	-0.0122	0.9819
CG2191	CG2191	1633425_at	-0.9987	0.5833	-0.0189	0.9771	-0.1634	0.4666	-0.1268	0.9833	-1.7356	0.2057	-1.6087	0.1899	0.0629	0.9922	-0.8004	0.6053	-0.8634	0.5763
Cpr78Cb	CG7663	1633426_at	0.1036	0.6787	-0.0917	0.6036	0.0940	0.7399	-0.0574	0.9610	0.0534	0.9018	0.1108	0.7439	-0.1119	0.8692	-0.2353	0.3774	-0.1234	0.6804
CG7582	CG7582	1633427_at	-1.5917	0.0049	-3.3234	0.0014	-3.2363	0.0000	-0.2411	0.7398	0.7808	0.0228	1.0219	0.0042	-0.1779	0.9064	-0.8683	0.1151	-0.6904	0.2198
prc	Pericardin	1633428_at	-0.2397	0.4053	0.0000	1.0000	0.1881	0.3555	0.0591	0.9538	-0.3950	0.1535	-0.4541	0.0703	-0.0467	0.9589	-0.1553	0.5717	-0.1086	0.7128
mof	males-absent on t	1633429_at	-0.5617	0.0082	0.3852	0.2579	0.6472	0.0317	0.1206	0.8822	-0.8368	0.0093	-0.9574	0.0030	-0.4375	0.6955	-0.2063	0.6883	0.2312	0.6399
TfIIS	lethal(2)35Cf	1633430_at	0.0087	0.9633	0.0754	0.7509	-0.1639	0.3123	-0.0386	0.9602	0.1299	0.5527	0.1684	0.3688	0.2567	0.6226	0.3162	0.1410	0.0596	0.8287
---	---	1633431_at	-0.0413	0.8223	-0.0222	0.8458	-0.0156	0.9359	0.1863	0.6354	0.0910	0.6790	-0.0953	0.6273	0.0736	0.9056	0.0316	0.9279	-0.0420	0.8912
CG9657	CG9657	1633432_at	0.1307	0.4601	0.1144	0.6099	0.2813	0.2250	0.0040	0.9956	-0.0780	0.7138	-0.0819	0.6671	-0.0769	0.9503	-0.1835	0.6458	-0.1066	0.8162
CG12866	CG12866	1633433_at	0.0065	0.9802	0.1495	0.3757	-0.0196	0.9362	-0.0236	0.9803	-0.0947	0.7274	-0.0711	0.7841	0.0389	0.9677	-0.0752	0.8216	-0.1141	0.6861
CG18327	CG18327	1633434_at	1.8850	0.0015	1.2474	0.0959	2.9662	0.0001	0.5114	0.3766	0.0420	0.9270	-0.4694	0.1075	-0.9964	0.5259	-0.4976	0.5246	0.4988	0.5296
CG12061	CG12061	1633435_a_at	0.2310	0.3454	-0.0002	1.0000	0.0094	0.9756	0.1377	0.8738	0.3334	0.2716	0.1957	0.5001	-0.0423	0.9697	0.0156	0.9748	0.0578	0.8831
CG14149	CG14149	1633436_at	-0.1226	0.8153	-0.3803	0.1055	-0.6189	0.0175	-0.2302	0.8251	-0.0318	0.9576	0.1984	0.6302	-0.1358	0.9460	-0.3948	0.5321	-0.2590	0.7040
---	---	1633437_at	-0.0717	0.7833	-0.1184	0.4262	-0.1480	0.4232	0.0769	0.9061	-0.0360	0.9009	-0.1129	0.5913	0.0381	0.9701	-0.0594	0.8734	-0.0975	0.7456
---	---	1633438_at	0.1766	0.4284	0.0614	0.6522	-0.0771	0.7329	0.0535	0.9474	0.0916	0.7334	0.0380	0.8895	-0.0256	0.9742	-0.0215	0.9462	0.0041	0.9902
Galpha73B	GTP-binding prote	1633439_at	3.6042	0.0002	2.2316	0.0044	3.4747	0.0000	1.3583	0.0033	1.7210	0.0001	3.6627	0.0482	0.0827	0.9238	0.2702	0.3352	0.1875	0.5367
CG10263	CG10263	1633440_a_at	-0.0072	0.9897	0.3412	0.5120	0.4283	0.2139	0.2022	0.4643	-0.0869	0.5940	-0.2891	0.0374	0.1800	0.9589	0.3059	0.8103	0.1259	0.9262
---	---	1633441_at	0.0764	0.7221	0.0038	0.9800	0.0965	0.6231	0.3080	0.3166	0.1244	0.5124	-0.1836	0.2555	0.1307	0.8415	0.0797	0.8197	-0.0510	0.8903
PebII	Protein ejaculaton	1633442_at	-0.0031	0.9906	-0.0898	0.5362	0.1787	0.2537	-0.0135	0.9863	-0.0432	0.8594	-0.0297	0.8942	-0.3341	0.3517	-0.0586	0.8132	0.2755	0.1744
CG2082	CG2082	1633443_s_at	-2.9598	0.0008	-3.9647	0.0055	-4.1931	0.0000	0.4567	0.1583	0.3368	0.0867	-0.1198	0.5306	0.5465	0.7559	-0.4754	0.5404	-1.0219	0.1929
CG32767 /// CG40450	CG32767 /// CG4(	1633444_s_at	0.1426	0.5737	0.0175	0.8642	-0.0260	0.8891	0.0236	0.9641	0.1342	0.3451	0.1106	0.3944	0.0934	0.9098	0.1409	0.6628	0.0475	0.9067
CG8541	CG8541	1633445_at	0.2422	0.2580	-0.0563	0.8546	0.0124	0.9756	0.1572	0.8128	0.1860	0.5009	0.0288	0.9283	0.1880	0.8940	-0.0522	0.9507	-0.2401	0.6781
CG6981	CG6981	1633446_s_at	1.0310	0.5491	-0.8194	0.5498	-0.5105	0.4777	0.5573	0.8794	0.6189	0.6631	0.0615	0.9689	0.1879	0.9826	-1.2446	0.5488	-1.4325	0.4854
Mctp	Multiple C2 domai	1633447_at	0.9496	0.0385	-0.1503	0.8108	0.4342	0.2290	0.2329	0.6086	0.5979	0.0170	0.3650	0.0756	-0.3488	0.8937	-0.4844	0.6449	-0.1356	0.9194
CG11926	CG11926	1633448_at	-0.6108	0.0157	0.0488	0.6682	0.2877	0.0802	0.0287	0.9681	-0.6201	0.0060	-0.6487	0.0029	-0.0736	0.9309	0.1329	0.6604	0.2065	0.4683
---	---	1633449_at	0.0212	0.9588	0.0557	0.8292	0.4278	0.0209	0.2869	0.6988	0.1109	0.7899	-0.1760	0.6091	-0.0170	0.9912	0.0437	0.9285	0.0607	0.8866
CG16781	CG16781	1633450_at	0.0122	0.9547	-0.0151	0.9432	0.0302	0.9041	-0.0064	0.9956	0.0184	0.9568	0.0249	0.9296	-0.0825	0.9246	0.0188	0.9689	0.1013	0.7674
CG32634	CG32634	1633451_at	0.1628	0.6038	0.1183	0.7623	-0.2168	0.5715	-0.3848	0.4786	0.2696	0.3647	0.6544	0.0202	-0.0138	0.9964	0.1772	0.8649	0.1910	0.8407
Rya-r44F	ryanodine recepto	1633452_a_at	-1.9798	0.0059	-2.2860	0.0324	-2.9283	0.0000	-0.0446	0.9641	0.6307	0.0247	0.6754	0.0116	0.6693	0.7644	0.4224	0.6918	-0.2469	0.8375
---	---	1633453_s_at	0.2090	0.3609	0.0636	0.5821	0.0135	0.9484	-0.1909	0.5932	-0.1158	0.5530	0.0751	0.6933	-0.0538	0.9467	-0.0968	0.7339	-0.0430	0.8982
Pfrx	phosphofructokina	1633454_s_at	1.0479	0.1333	1.0752	0.3636	1.6920	0.0032	0.2110	0.5948	-0.5483	0.0143	-0.7593	0.0020	-0.2953	0.9514	-0.5031	0.7741	-0.2078	0.9168
Vps16B	Vps16B	1633455_at	-0.3949	0.0829	-0.3340	0.0993	-0.1506	0.5755	0.1275	0.7440	-0.0031	0.9901	-0.1306	0.4159	-0.1467	0.7644	-0.0259	0.9384	0.1208	0.5819
---	---	1633456_at	0.2360	0.1351	0.0442	0.8222	0.2271	0.2806	0.0883	0.9252	0.0988	0.7712	0.0104	0.9766	-0.0539	0.9409	-0.0924	0.7200	-0.0386	0.9020
CG15144	CG15144	1633457_at	0.1005	0.5789	-0.0539	0.6202	0.0902	0.6642	0.1663	0.6515	0.1301	0.4854	-0.0362	0.8614	0.0305	0.9774	-0.1308	0.6324	-0.1613	0.5471
swa	swallow	1633458_at	0.5574	0.6149	-1.6313	0.3370	-1.1682	0.1884	0.2158	0.8220	2.6530	0.0003	2.4373	0.0002	-0.3803	0.9659	0.2345	0.9512	0.6148	0.8409

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
fau	fau	1633459_a_at	-1.7650	0.0277	-0.7704	0.2322	-1.8211	0.0049	-0.4503	0.3206	-2.0181	0.0002	-1.5677	0.0003	0.8088	0.7877	-1.0067	0.4302	-1.8155	0.1808
CG6034	CG6034	1633460_at	-0.4031	0.6140	-3.0245	0.0047	-2.8521	0.0000	0.2156	0.7589	1.1830	0.0029	0.9674	0.0041	-0.0472	0.9914	-1.2775	0.1957	-1.2304	0.2447
slik	polo kinase kinas	1633461_at	1.1684	0.0281	0.4906	0.3523	0.2829	0.1494	0.0552	0.9336	0.4390	0.0315	0.3838	0.0345	0.3338	0.7770	-0.0137	0.9905	-0.3476	0.5154
CG31365	CG31365	1633462_at	0.1907	0.5256	0.2629	0.1086	0.3519	0.0723	0.0481	0.9603	0.2927	0.2503	0.2447	0.2871	-0.0128	0.9938	0.2691	0.4563	0.2819	0.4457
CG30463	CG30463	1633463_s_at	1.4316	0.0060	2.7969	0.0020	1.2467	0.0104	0.2256	0.7443	-0.0364	0.9321	-0.2620	0.3474	1.7485	0.1953	1.3096	0.0795	-0.4389	0.5612
---	---	1633464_at	0.0667	0.7955	0.0282	0.8453	0.3853	0.0630	0.0480	0.9558	-0.0694	0.8096	-0.1175	0.6204	-0.0844	0.9441	0.0148	0.9829	0.0993	0.8287
Cp7Fa	Chorion protein a	1633465_at	0.0711	0.6532	0.2737	0.0939	0.5381	0.0183	0.0687	0.9311	-0.2847	0.2224	-0.3535	0.0909	0.0039	0.9970	-0.0083	0.9872	-0.0122	0.9760
CG7728	CG7728	1633466_at	0.4618	0.0993	0.2512	0.4986	0.5477	0.0266	0.3369	0.5140	0.6092	0.0328	0.2723	0.2684	-0.0917	0.9487	0.3364	0.4300	0.4281	0.3271
CG30459	CG30459	1633467_a_at	0.2561	0.3396	0.1450	0.3186	0.2047	0.2424	-0.0374	0.9558	-0.0478	0.8390	-0.0104	0.9642	-0.0776	0.9317	-0.1398	0.6655	-0.0621	0.8753
CG40373	CG40373	1633468_x_at	0.3221	0.2631	0.0586	0.7871	0.3504	0.0658	0.3185	0.5813	0.3917	0.1816	0.0732	0.8278	0.0394	0.9589	0.1796	0.4096	0.1402	0.5461
---	---	1633469_at	0.2664	0.0880	0.0663	0.5902	0.2290	0.3574	-0.0634	0.9023	0.0418	0.8472	0.1051	0.5169	-0.2670	0.6955	-0.1323	0.6813	0.1348	0.6677
RN-tre	tre oncogene-relat	1633470_at	-0.5471	0.1485	0.6237	0.0283	1.0014	0.0021	-0.0482	0.9653	-1.0768	0.0039	-1.0285	0.0028	-0.1176	0.9420	0.3718	0.4565	0.4894	0.3362
Prx2540-2	Peroxiredoxin 254	1633471_at	0.7943	0.0111	-0.0676	0.6776	-0.4640	0.1020	-0.4394	0.4141	0.3732	0.2097	0.8126	0.0088	-0.2749	0.7092	-0.6123	0.0661	-0.3374	0.2869
---	---	1633472_at	-0.0221	0.9284	0.2227	0.3853	0.0592	0.7747	-0.1431	0.7562	-0.2806	0.1563	-0.1374	0.4660	-0.0156	0.9914	-0.0175	0.9738	-0.0019	0.9977
Ald	aldolase	1633473_s_at	0.7300	0.0042	0.6944	0.1742	0.5623	0.1708	-0.0144	0.9924	-0.4234	0.1768	-0.4089	0.1454	0.0447	0.9875	-0.6283	0.3167	-0.6730	0.3079
CG32603	CG32603	1633474_x_at	0.1689	0.5328	0.2954	0.3312	0.2694	0.0863	-0.1696	0.8064	-0.3637	0.1841	-0.1940	0.4561	-0.0851	0.9330	-0.1553	0.6614	-0.0703	0.8709
---	---	1633475_at	0.1470	0.3856	0.0552	0.6587	0.3045	0.0470	0.0826	0.8721	0.0360	0.8800	-0.0466	0.8180	-0.0957	0.8270	-0.0745	0.7247	0.0212	0.9344
Acox57D-d	acyl co-enzyme A	1633476_at	1.8965	0.0559	1.1693	0.1233	1.9204	0.0002	0.0931	0.9772	-0.1242	0.9102	-0.2174	0.8073	-0.6742	0.6824	-0.7332	0.2630	-0.0590	0.9510
CG10496	CG10496	1633477_at	0.3629	0.0613	0.5377	0.0784	1.0729	0.0003	0.1201	0.8546	-0.0277	0.9338	-0.1477	0.5203	-0.1347	0.8342	0.0105	0.9839	0.1452	0.6150
---	---	1633478_at	0.1498	0.3854	0.0858	0.4675	-0.0370	0.8209	-0.0928	0.8327	0.0647	0.7425	0.1575	0.2978	0.1747	0.6955	0.0344	0.9047	-0.1403	0.4684
CG4270	CG4270	1633479_a_at	0.1626	0.3271	0.0433	0.7741	-0.0491	0.8006	-0.0140	0.9852	0.0756	0.6996	0.0896	0.5999	0.0278	0.9821	-0.0808	0.8343	-0.1086	0.7442
CG5392	CG5392	1633480_at	0.0079	0.9895	-0.1544	0.4524	-0.6782	0.0032	-0.3883	0.6202	-0.2384	0.5720	0.1499	0.7183	-0.1408	0.9024	-0.5322	0.1900	-0.3914	0.3631
CG14394	CG14394	1633481_at	-1.1833	0.0073	-0.9208	0.0295	-1.3107	0.0012	-0.3020	0.4317	-0.4195	0.0515	-0.1175	0.5759	0.0169	0.9939	-0.0530	0.9496	-0.0699	0.9214
---	---	1633482_at	0.1304	0.5437	0.0720	0.5096	0.1200	0.6722	-0.0776	0.9487	-0.1502	0.6990	-0.0725	0.8530	0.0498	0.9550	0.1515	0.5800	0.1017	0.7307
CG14207	small heat shock p	1633483_a_at	-1.6218	0.0045	-1.3192	0.0442	-2.0338	0.0001	-0.1697	0.6718	0.0763	0.7314	0.2460	0.1456	0.5295	0.7387	0.4813	0.4840	-0.0483	0.9612
CG34462 /// Cpr66Ca	CG7072	1633484_at	0.7424	0.7104	-1.5048	0.0170	0.2817	0.2397	1.7896	0.0189	0.9367	0.0276	-0.8529	0.0258	-0.1094	0.9914	-1.3110	0.5994	-1.2016	0.6328
CG32828	CG32828	1633485_at	-0.0886	0.7306	0.1010	0.4882	0.1527	0.4679	-0.0697	0.9295	-0.2180	0.3539	-0.1483	0.5053	-0.0046	0.9964	0.0121	0.9808	0.0167	0.9673
mRpL3	mitochondrial ribo	1633486_at	-0.0655	0.8519	0.7095	0.0720	0.7334	0.0059	-0.0825	0.8935	-0.5595	0.0143	-0.4769	0.0174	-0.1771	0.8875	0.0757	0.9170	0.2528	0.6168
CG30109	CG30109	1633487_at	-0.0469	0.8113	0.1085	0.5561	0.6690	0.1096	0.3988	0.4643	0.3379	0.2531	-0.0609	0.8599	-0.0539	0.9853	0.3934	0.5825	0.4473	0.5282
aay	astray	1633488_at	-0.5225	0.2412	-0.4401	0.0451	-0.0416	0.9231	0.3498	0.3425	0.5187	0.0227	0.1689	0.3916	0.0108	0.9978	0.7347	0.4167	0.7239	0.4387
beat-Ic	beaten path Ic	1633489_at	-0.0290	0.8526	-0.3298	0.0463	0.0057	0.9806	0.1864	0.5037	0.2394	0.1040	0.0530	0.7385	-0.1862	0.8298	-0.1171	0.7994	0.0691	0.8900
CG15638	CG15638	1633490_at	0.1436	0.3632	-0.0790	0.5059	-0.1210	0.6680	0.0612	0.9370	0.2756	0.2187	0.2144	0.2921	0.1075	0.8480	-0.0201	0.9588	-0.1276	0.6133
CG31865 /// CG31866	CG31866 /// CG3	1633491_s_at	0.0257	0.9284	0.4081	0.2085	0.4721	0.0992	0.0422	0.9259	-0.2955	0.0371	-0.3376	0.0135	-0.0683	0.9701	0.0788	0.9125	0.1471	0.7933
hgo	homogentisate 1,2	1633492_at	0.5278	0.6279	-0.0009	0.9960	0.1303	0.5336	0.0177	0.9833	0.0148	0.9589	-0.0029	0.9905	-0.2700	0.9467	-0.6088	0.6450	-0.3388	0.8247
---	---	1633493_at	0.4353	0.0457	0.1786	0.3563	0.1586	0.3335	0.0021	0.9981	0.0352	0.9045	0.0331	0.8974	0.0124	0.9895	-0.0416	0.8845	-0.0540	0.8278
---	---	1633494_at	0.1711	0.4354	-0.0200	0.9399	-0.3464	0.2194	0.0000	0.9999	0.0157	0.9717	0.0156	0.9658	0.1453	0.8802	-0.1865	0.6414	-0.3318	0.3807
---	---	1633495_at	-0.0900	0.5642	0.0292	0.8074	0.2512	0.2085	-0.1030	0.8074	-0.2509	0.1347	-0.1479	0.3371	-0.1671	0.7644	-0.1075	0.6820	0.0596	0.8432
---	---	1633496_at	0.3037	0.1185	0.0656	0.6288	0.2374	0.1654	-0.0304	0.9626	0.0877	0.6500	0.1181	0.4687	0.0202	0.9895	-0.0686	0.8821	-0.0888	0.8255
CG33092	CG33092	1633497_s_at	0.2034	0.3936	-0.0310	0.7943	0.1137	0.4672	0.0580	0.9255	0.0829	0.7002	0.0249	0.9125	-0.0663	0.9374	-0.0945	0.7693	-0.0282	0.9394
CG7188	CG7188	1633498_at	-1.4665	0.0148	-1.9524	0.0156	-2.2134	0.0003	-0.0451	0.9777	0.9835	0.0230	1.0287	0.0119	0.1799	0.8884	0.3801	0.4303	0.2002	0.7138
retinin	retinin	1633499_at	0.5408	0.3402	-0.0560	0.7350	0.1885	0.3362	0.0451	0.9592	0.2305	0.3260	0.1853	0.3881	-0.1134	0.9589	-0.1713	0.8352	-0.0579	0.9465
---	---	1633500_at	-0.1718	0.2487	0.0244	0.8661	0.1053	0.4806	0.0877	0.8636	-0.0231	0.9278	-0.1108	0.5306	-0.0270	0.9793	0.0868	0.7618	0.1138	0.6601
tws	phosphoprotein pl	1633501_s_at	0.3767	0.3301	0.6024	0.1564	-0.1761	0.3261	-0.5176	0.0840	-0.3475	0.0507	0.1701	0.2791	0.2283	0.8943	-0.1768	0.8380	-0.4051	0.5518
CG34375	CG13834	1633502_at	0.1411	0.5320	0.1852	0.3096	0.1648	0.3954	-0.0958	0.8837	-0.0541	0.8521	0.0417	0.8745	0.0941	0.8930	0.0310	0.9409	-0.0631	0.8515
CG12171	CG12171	1633503_at	0.2169	0.5838	1.2926	0.0056	1.4429	0.0005	-0.1248	0.7803	-1.1303	0.0005	-1.0054	0.0005	-0.2891	0.8386	-0.0916	0.9196	0.1975	0.7749
---	---	1633504_at	0.2771	0.3204	-0.0541	0.7026	0.0778	0.6416	-0.0091	0.9937	0.1937	0.4358	0.2029	0.3575	-0.0119	0.9928	0.0779	0.8477	0.0898	0.8046
ALiX	ALG-2 interacting	1633505_at	-0.5186	0.1091	-0.4553	0.0839	-0.1986	0.3462	0.2322	0.5912	0.1525	0.5099	-0.0796	0.7355	-0.0847	0.9588	0.1675	0.7584	0.2521	0.6068
CG6091	CG6091	1633506_s_at	-0.0882	0.9315	0.7164	0.2808	0.4152	0.1563	0.2578	0.7929	-0.0554	0.9215	-0.3132	0.4004	0.6417	0.8049	0.7809	0.4794	0.1392	0.9257
---	---	1633507_at	0.1303	0.4025	0.0376	0.7583	0.0589	0.7571	-0.0010	0.9988	0.0186	0.9337	0.0196	0.9195	-0.0193	0.9848	-0.0598	0.8504	-0.0406	0.8993
CG31700	CG31700	1633508_at	0.1331	0.3441	0.0390	0.7696	0.0880	0.5764	-0.0141	0.9825	0.1077	0.4937	0.1217	0.3804	0.1052	0.8825	0.0347	0.9363	-0.0705	0.8385
CG32720	CG32720	1633509_at	0.1235	0.5081	0.0426	0.7848	0.1190	0.5393	0.0609	0.9375	0.3182	0.1561	0.2573	0.2008	0.0041	0.9976	0.0439	0.9259	0.0398	0.9230
CG17141	CG17141	1633510_at	0.1554	0.3854	-0.1341	0.6553	-0.3970	0.0380	-0.0906	0.9043	0.1906	0.4553	0.2811	0.1990	0.0487	0.9677	-0.0009	0.9996	-0.0496	0.9095
---	---	1633511_at	0.1094	0.4213	-0.0102	0.9555	0.2015	0.3139	0.1090	0.7949	0.1114	0.5441	0.0024	0.9912	-0.1177	0.8270	-0.0012	0.9989	0.1165	0.6315
toy	twin of eyeless	1633512_at	-0.0366	0.8545	-0.1162	0.5197	0.0401	0.8748	0.2446	0.6338	0.1269	0.6556	-0.1177	0.6513	-0.1922	0.8480	-0.1500	0.7671	0.0422	0.9431
CG11474	CG11474	1633513_a_at	-0.2619	0.2472	0.6488	0.0582	0.9866	0.0103	-0.0929	0.8990	-1.2361	0.0009	-1.1432	0.0008	-0.4146	0.6749	-0.3570	0.3648	0.0576	0.9171
---	---	1633514_at	-0.0486	0.7748	0.0344	0.8786	-0.0054	0.9848	0.2437	0.4728	0.1104	0.5767	-0.1333	0.4372	0.1420	0.8472	0.2088	0.5085	0.0667	0.8699
ago	archipelago	1633515_s_at	0.5357	0.0148	-0.0457	0.7801	-0.0715	0.7737	-0.1297	0.8249	0.4231	0.0656	0.5528	0.0144	0.0219	0.9831	-0.0220	0.9557	-0.0439	0.8949
CG34409	CG31411	1633516_at	-0.3694	0.1623	-0.2425	0.2730	-0.5830	0.0098	0.2252	0.6854	0.2439	0.3534	0.0187	0.9547	0.2959	0.6824	0.1273	0.7050	-0.1686	0.5984
CG5726	CG5726	1633517_at	0.0868	0.7874	-0.3130	0.0935	-0.1405	0.5766	0.3639	0.4800	0.6440	0.0274	0.2801	0.2598	0.1959	0.8270	0.2273	0.5630	0.0313	0.9525
---	---	1633518_at	0.1731	0.4883	0.4115	0.0772	0.0111	0.9629	-0.1604	0.7433	-0.1781	0.4246	-0.0176	0.9486	0.3572	0.6272	0.2304	0.4565	-0.1268	0.7154
---	---	1633519_at	0.2776	0.2421	0.0341	0.7613	0.2898	0.1674	0.0069	0.9943	0.0922	0.7057	0.0853	0.7020	-0.2476	0.7287	-0.0566	0.9040	0.1910	0.5576
CG32027	CG32027	1633520_at	0.2176	0.5383	0.7661	0.0198	0.3620	0.0759	-0.8088	0.0840	-0.8070	0.0097	0.0018	0.9958	-0.1392	0.9242	-0.2602	0.6191	-0.1210	0.8470
CG40000	CG40000	1633521_at	0.1217	0.5575	0.0171	0.8862	0.1035	0.5359	-0.0388	0.9592	-0.0404	0.8789	-0.0015	0.9953	-0.0458	0.9467	0.0012	0.9989	0.0470	0.8636
CG8968	CG8968	1633522_at	0.2297	0.1893	0.2052	0.2431	0.0996	0.7158	-0.1339	0.7967	-0.1302	0.5680	0.0037	0.9887	0.2083	0.7095	0.0823	0.7731	-0.1260	0.6168
Pbgs	Porphobilinogen s	1633523_at	0.0366	0.9432	0.5609	0.0750	0.8644	0.0007	0.0786	0.8640	-0.5391	0.0060	-0.6177	0.0019	-0.1717	0.9238	-0.0131	0.9925	0.1586	0.8310
CalpA	Calpain A	1633524_a_at	-1.3626	0.0035	-0.3218	0.0775	-0.6052	0.0028	-0.3989	0.4046	-1.1894	0.0017	-0.7906	0.0059	-0.0134	0.9914	0.0559	0.8885	0.0693	0.8435
---	---	1633525_at	-0.0603	0.7318	0.1104	0.5902	0.0371	0.8437	-0.1393	0.6413	-0.1295	0.3868	0.0099	0.9573	0.0787	0.9142	0.0771	0.8169	-0.0015	0.9979
Sox15	Sox box protein 15	1633526_at	-0.0548	0.7319	0.1683	0.1485	0.0157	0.9403	-0.0398	0.9436	-0.0092	0.9703	0.0306	0.8704	0.1403	0.7997	0.1384	0.5679	-0.0019	0.9965
Cap	Chromosome-ass	1633527_a_at	-0.4564	0.1963	0.0353	0.8580	-0.0488	0.8706	0.1271	0.8274	0.0651	0.8144	-0.0619	0.8056	0.1261	0.9467	0.6019	0.2775	0.4759	0.4135

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
exd	extradenticle	1633528_s_at	-0.2256	0.1928	0.1072	0.7125	-0.3283	0.1018	-0.2924	0.4122	-0.1757	0.3848	0.1166	0.5473	0.0481	0.9677	0.2035	0.5342	0.1553	0.6471
---	---	1633529_s_at	0.0777	0.6392	0.1779	0.2775	0.0987	0.6021	0.1207	0.8164	0.2293	0.2642	0.1086	0.5961	-0.0233	0.9831	-0.0037	0.9943	0.0196	0.9575
HGTX	HGTX	1633530_at	0.0614	0.7770	-0.0169	0.8819	-0.0577	0.7984	0.0485	0.9251	-0.0226	0.9167	-0.0711	0.6568	-0.0471	0.9611	-0.0463	0.9120	0.0008	0.9990
CG34381	CG14004	1633531_at	-1.7118	0.0036	-0.0681	0.5665	-0.6855	0.1272	-0.3519	0.6338	-1.6974	0.0013	-1.3455	0.0022	0.0700	0.9679	-0.1434	0.8114	-0.2134	0.6728
CG5767	CG5767	1633532_at	0.2441	0.2872	0.3561	0.0354	-0.0123	0.9533	-0.1359	0.7229	-0.2350	0.1719	-0.0991	0.5624	0.2126	0.8065	-0.1843	0.6394	-0.3969	0.2869
CG6329	CG6329	1633533_at	-2.1874	0.0039	-2.0321	0.0658	-2.6859	0.0000	-0.3818	0.6081	-0.5001	0.1778	-0.1183	0.7749	-0.1835	0.9012	-0.6562	0.2117	-0.4728	0.3921
mura	murashka	1633534_s_at	0.3757	0.5581	0.1793	0.7631	-1.1742	0.0133	-0.7696	0.2823	0.3782	0.3801	1.1478	0.0087	0.2439	0.9309	-0.1200	0.9341	-0.3640	0.7247
VhaAC39	VhaAC39	1633535_at	-0.9764	0.0107	-0.5907	0.0752	-0.8849	0.0016	-0.2778	0.2868	-0.6347	0.0023	-0.3569	0.0173	-0.0264	0.9895	-0.2421	0.5880	-0.2158	0.6340
CG4630	CG4630	1633536_at	-0.7052	0.0052	-1.3255	0.0019	-1.1308	0.0071	0.1004	0.9285	0.6538	0.0525	0.5534	0.0643	-0.1275	0.8814	-0.1608	0.6512	-0.0332	0.9429
---	---	1633537_at	-0.2955	0.1973	0.1077	0.4746	0.0594	0.7349	-0.0518	0.9247	-0.1765	0.2853	-0.1248	0.4163	0.1852	0.7697	0.1352	0.6421	-0.0500	0.8921
---	---	1633538_at	0.2416	0.1869	0.1476	0.3308	0.0963	0.5890	0.0767	0.8809	0.1397	0.4400	0.0630	0.7370	0.0483	0.9380	0.0738	0.7456	0.0255	0.9234
Hsp60D	Hsp60D	1633539_s_at	0.3044	0.1940	0.1136	0.4871	0.2873	0.1534	0.0009	0.9988	-0.0645	0.7371	-0.0655	0.7034	-0.0283	0.9742	-0.0525	0.8619	-0.0243	0.9361
CG8147	CG8147	1633540_at	3.7360	0.0043	0.9852	0.5805	1.9604	0.0012	0.1605	0.9528	-0.4403	0.5882	-0.6008	0.3863	-0.8901	0.8461	-3.2014	0.0966	-2.3113	0.2338
---	---	1633541_at	0.3236	0.1066	0.1296	0.5762	0.0866	0.5730	0.0168	0.9857	0.1708	0.4375	0.1540	0.4401	0.1169	0.8270	-0.0455	0.8874	-0.1624	0.4795
CG13717	CG13717	1633542_at	-1.7238	0.0035	-0.2909	0.5781	-0.6314	0.0365	0.0343	0.9834	-1.4165	0.0035	-1.4508	0.0019	0.2796	0.8609	-0.0297	0.9797	-0.3093	0.6577
Cdc42	Cdc42-like	1633543_s_at	0.4014	0.0335	0.7410	0.0207	0.7625	0.0033	0.0520	0.9247	0.1018	0.5670	0.0498	0.7868	0.1219	0.8909	0.3576	0.2693	0.2357	0.4978
CG7349	CG7349	1633544_at	-0.3336	0.2051	-0.1291	0.3337	-0.2668	0.1243	0.0276	0.9627	-0.0791	0.6556	-0.1067	0.4762	0.0085	0.9939	0.0081	0.9876	-0.0004	0.9992
PGRP-SD	PGRP-SD	1633545_at	1.4593	0.0211	-0.2967	0.7406	-0.9195	0.0487	-0.8773	0.2642	1.4374	0.0082	2.3147	0.0006	-0.4180	0.8815	-0.0622	0.9726	0.3558	0.7785
CG5197	CG5197	1633546_at	-0.0187	0.9206	0.0770	0.5352	-0.1494	0.4684	-0.0993	0.8676	-0.0513	0.8507	0.0479	0.8434	0.1495	0.7485	0.1169	0.5755	-0.0326	0.9055
CG13211	CG13211	1633547_a_at	0.6796	0.0636	0.9690	0.0717	1.0103	0.0021	0.1667	0.8816	-0.0589	0.9114	-0.2256	0.5464	0.2121	0.8276	0.3294	0.4177	0.1173	0.8178
---	---	1633548_at	0.8647	0.0368	0.4417	0.0990	0.3030	0.1458	0.0419	0.9491	0.0753	0.7237	0.0334	0.8776	0.2829	0.8192	-0.0251	0.9782	-0.3079	0.5667
e(r)	enhancer of rudim	1633549_s_at	-0.1463	0.3967	0.2253	0.1100	0.0618	0.8077	0.0766	0.9262	-0.2616	0.2974	-0.3382	0.1267	0.2370	0.6898	0.0823	0.7874	-0.1546	0.5493
Ubx	bithorax	1633550_a_at	1.1546	0.0981	0.2366	0.4935	0.9825	0.0036	0.0595	0.9255	0.2461	0.1990	0.1865	0.2834	-0.5249	0.8236	-0.4101	0.7142	0.1148	0.9341
Osi11	Osiris	1633551_at	0.1673	0.3729	0.1663	0.2384	0.1084	0.6389	0.0672	0.9242	0.1225	0.5955	0.0553	0.8174	0.0424	0.9667	0.0209	0.9628	-0.0215	0.9551
---	---	1633552_at	-0.0072	0.9747	-0.1580	0.6460	-0.1144	0.6698	0.0784	0.9506	0.1179	0.7806	0.0395	0.9263	0.0760	0.8869	0.0978	0.6536	0.0219	0.9374
SP1173	SP1173	1633553_s_at	-1.3328	0.0085	0.0072	0.9785	-1.2673	0.0002	-0.1800	0.7850	-0.8305	0.0086	-0.6505	0.0156	0.9109	0.3166	0.3056	0.5439	-0.6053	0.2249
cos	costal2	1633554_at	0.4744	0.1051	-0.0819	0.7009	-0.8164	0.0126	-0.0859	0.8571	0.6934	0.0025	0.7793	0.0009	0.5067	0.6749	0.0865	0.9085	-0.4201	0.4016
---	---	1633555_at	-0.0396	0.8223	0.0556	0.8201	0.2540	0.4138	0.2366	0.6533	0.1046	0.7240	-0.1320	0.6071	-0.0989	0.9101	0.2164	0.5020	0.3153	0.3248
Dad	Daughters agains	1633556_s_at	-1.7341	0.0033	-1.2360	0.0404	-1.1917	0.0038	0.0355	0.9680	-0.6576	0.0120	-0.6931	0.0057	0.0397	0.9901	-0.1000	0.9282	-0.1397	0.8857
CG5414 /// DmirCG5414	CG5414	1633557_at	-0.1474	0.5428	-0.3316	0.4050	-0.1951	0.4648	-0.2742	0.6041	0.2501	0.3574	0.5244	0.0349	-0.4622	0.6749	-0.1106	0.8602	0.3516	0.4504
CG14997	CG14997	1633558_s_at	-0.3887	0.4360	-1.7339	0.0151	-1.2141	0.0116	0.1799	0.8747	0.6597	0.0967	0.4798	0.1754	-0.3379	0.8540	-0.7402	0.3133	-0.4023	0.6202
CG8331	CG8331	1633559_at	0.4438	0.0611	1.0017	0.0081	0.9990	0.0005	0.0252	0.9649	-0.2848	0.0644	-0.3100	0.0302	-0.0864	0.9168	0.1919	0.5157	0.2783	0.3427
CG16903	CG16903	1633560_at	-0.3756	0.4455	0.1266	0.6912	0.6207	0.0062	0.1793	0.7409	-0.5576	0.0272	-0.7369	0.0049	-0.2667	0.8875	0.0892	0.9377	0.3559	0.6398
klu	klumphuss	1633561_at	0.3681	0.1327	-0.0378	0.8450	0.1426	0.4207	0.0224	0.9838	0.1290	0.6587	0.1066	0.6949	-0.0116	0.9916	-0.1343	0.6237	-0.1227	0.6562
CG14569	CG14569	1633562_at	-0.0063	0.9783	0.0976	0.5813	0.3496	0.0416	0.0457	0.9445	-0.0054	0.9845	-0.0511	0.8090	-0.0769	0.9400	0.0866	0.8376	0.1635	0.6298
CG1637	CG1637	1633563_at	-0.7884	0.1408	-1.2468	0.0157	-0.3805	0.2115	0.1124	0.9247	-0.3920	0.2716	-0.5044	0.1108	-0.8076	0.5754	-0.8446	0.1795	-0.0369	0.9704
---	---	1633564_at	-0.0155	0.9427	-0.2049	0.3705	-0.1807	0.4382	0.1185	0.7814	0.1840	0.3032	0.0655	0.7298	-0.0906	0.8775	-0.1214	0.6209	-0.0308	0.9220
CG7741	CG7741	1633565_at	0.2224	0.2028	0.0668	0.7992	0.0025	0.9916	-0.2291	0.4631	0.3701	0.0355	0.5991	0.0027	-0.0238	0.9874	0.1763	0.6314	0.2001	0.5811
CG7694	CG7694	1633566_at	0.3252	0.1093	0.5382	0.0218	0.5975	0.0051	-0.0132	0.9838	-0.1053	0.5162	-0.0921	0.5383	-0.0624	0.9366	0.0725	0.8244	0.1349	0.6150
---	---	1633567_at	-0.0665	0.7700	-0.0285	0.7793	0.1271	0.5773	0.1017	0.8217	0.1258	0.4959	0.0241	0.9099	-0.0044	0.9980	0.1361	0.7673	0.1404	0.7484
CG17683	CG17683	1633568_s_at	0.5756	0.0707	0.3301	0.2583	0.2370	0.3096	-0.0546	0.9343	0.2007	0.3072	0.2553	0.1409	0.0934	0.9466	-0.0299	0.9652	-0.1234	0.8138
---	---	1633569_s_at	0.2645	0.3644	-0.0121	0.9204	0.2381	0.1378	-0.0207	0.9857	0.1210	0.6938	0.1417	0.5973	-0.1345	0.8609	-0.0810	0.8478	0.0534	0.9013
CG1688	CG1688	1633570_at	-0.0448	0.8636	-0.1092	0.2799	0.0024	0.9950	-0.0192	0.9777	0.0755	0.7015	0.0947	0.5785	0.2077	0.8472	0.2670	0.5711	0.0593	0.9231
---	---	1633571_at	0.0452	0.8078	-0.0874	0.7215	-0.0451	0.8522	-0.0416	0.9436	-0.1452	0.4012	-0.1036	0.5270	0.0802	0.9452	-0.0433	0.9402	-0.1235	0.7640
CG13210	CG13210	1633572_at	0.4383	0.0318	0.7500	0.0301	1.2643	0.0083	0.3131	0.4861	0.1864	0.4562	-0.1267	0.6000	-0.3342	0.7633	0.3541	0.4452	0.6883	0.1618
att-ORFA /// att-ORFB	alternative testis t	1633573_a_at	-0.8172	0.0036	0.0285	0.9515	0.1221	0.7408	-0.2466	0.7248	-1.9874	0.0004	-1.7408	0.0004	-0.3526	0.7733	-1.1180	0.0568	-0.7653	0.1744
CG31820	CG31820	1633574_at	0.0730	0.6328	0.0048	0.9713	0.0939	0.6134	-0.1375	0.8196	-0.0527	0.8656	0.0847	0.7391	-0.1744	0.7506	-0.0653	0.8353	0.1091	0.6711
---	---	1633575_at	0.1819	0.3970	0.3251	0.0688	0.3730	0.1436	-0.3676	0.3855	-0.3087	0.1958	0.0590	0.8293	-0.2646	0.7500	0.0864	0.8613	0.3510	0.3343
CG1910	anon-fast-evolving	1633576_s_at	1.0461	0.0115	1.6672	0.0108	0.2823	0.4165	-0.7710	0.1897	-0.4223	0.2349	0.3487	0.2779	0.5177	0.6287	0.1923	0.7036	-0.3254	0.4824
---	---	1633577_at	-0.1797	0.2384	-0.0950	0.4310	-0.0238	0.9025	0.0580	0.9216	-0.0260	0.9167	-0.0840	0.6453	-0.1297	0.7644	-0.0327	0.9063	0.0970	0.6189
PpN58A	Protein phosphata	1633578_a_at	0.3747	0.0353	0.0955	0.6410	0.1901	0.2560	0.0740	0.8721	0.1338	0.4190	0.0597	0.7273	-0.0671	0.9514	0.0183	0.9739	0.0854	0.8374
CG5045	CG5045	1633579_at	-0.1713	0.4473	0.6524	0.0291	0.8567	0.0005	0.1427	0.6908	-0.8229	0.0011	-0.9655	0.0004	-0.0285	0.9831	0.0811	0.8469	0.1096	0.7578
CG18208	CG18208	1633580_at	-0.1480	0.5278	0.1196	0.4629	-0.0577	0.7279	-0.1295	0.8315	-0.3092	0.1806	-0.1797	0.4066	0.2692	0.7118	0.1796	0.5778	-0.0896	0.8128
CG32571	CG32571	1633581_at	0.0597	0.7315	0.5214	0.0221	0.2358	0.1620	-0.1168	0.7908	-0.1746	0.3390	-0.0578	0.7708	0.1865	0.7220	0.2797	0.1948	0.0932	0.7096
Ih	putative Ih-channe	1633582_at	0.7156	0.0832	0.4485	0.1387	0.7646	0.0910	-0.3944	0.4482	-0.4356	0.1221	-0.0412	0.9039	-0.6203	0.7241	-0.6648	0.3739	-0.0444	0.9696
Glu-Rl	glutamate recepto	1633583_at	0.1148	0.5798	0.1476	0.2603	0.0346	0.9042	-0.1858	0.6533	-0.0684	0.7762	0.1173	0.5489	-0.0357	0.9823	-0.0706	0.9004	-0.0349	0.9468
CG17843	CG17843	1633584_at	-0.4819	0.1061	0.8210	0.0876	0.3010	0.2308	-0.4707	0.5735	-1.5780	0.0038	-1.1073	0.0112	0.1340	0.9503	-0.1921	0.8176	-0.3261	0.6367
CG12550	CG12550	1633585_s_at	0.1196	0.5283	0.0979	0.6114	0.1580	0.3792	-0.0092	0.9917	0.0201	0.9394	0.0293	0.8945	0.0203	0.9871	0.0325	0.9445	0.0122	0.9791
Atg12	Autophagy-specifi	1633586_at	-0.2116	0.4294	-0.7695	0.0601	-0.9555	0.0004	0.1957	0.6457	0.3905	0.0623	0.1948	0.2979	0.3931	0.6955	-0.1046	0.8546	-0.4977	0.2454
---	---	1633587_at	0.2928	0.1319	0.1350	0.4157	0.1333	0.4619	0.0772	0.9108	0.3304	0.1283	0.2532	0.1932	0.0231	0.9853	-0.0330	0.9432	-0.0562	0.8874
---	---	1633588_at	0.4068	0.0689	0.2694	0.1663	0.1373	0.4559	-0.0474	0.9479	-0.0141	0.9639	0.0333	0.8915	-0.0340	0.9666	0.0417	0.8980	0.0757	0.7629
CG1552	CG1552	1633589_a_at	0.2134	0.1613	0.0917	0.5213	0.0788	0.6814	-0.0534	0.9529	-0.1883	0.4698	-0.1349	0.5882	0.0301	0.9816	-0.0683	0.8684	-0.0984	0.7754
---	---	1633590_at	0.2163	0.1926	-0.1051	0.5416	-0.0993	0.5645	-0.0082	0.9922	0.2729	0.1139	0.2811	0.0716	0.0418	0.9611	0.0126	0.9763	-0.0293	0.9312
gcm2	glide glide2	1633591_at	-0.1513	0.3255	0.0808	0.4658	0.1625	0.4678	0.2273	0.3817	0.0182	0.9298	-0.2091	0.1101	0.0999	0.9101	0.1955	0.5531	0.0956	0.8039
CREG	cellular repressor	1633592_a_at	0.8962	0.0049	0.2507	0.4149	0.7958	0.0092	0.3599	0.4174	-0.3707	0.1310	-0.7305	0.0059	-0.2313	0.8202	-1.0081	0.0391	-0.7769	0.1002
CG32713 /// CG33223 /// D	CG32713 /// CG3	1633593_s_at	0.1913	0.4393	0.0467	0.6587	0.1849	0.5613	0.1093	0.8979	0.2148	0.4610	0.1055	0.7234	-0.0082	0.9965	0.1346	0.8356	0.1427	0.8123
Rpn7	Rpn7	1633594_at	0.4716	0.0234	0.2380	0.4203	0.4003	0.0886	0.2429	0.4898	0.3639	0.0565	0.1210	0.5019	0.0291	0.9862	0.3027	0.4270	0.2737	0.4882
---	---	1633595_at	0.0349	0.8251	0.2468	0.1419	0.4423	0.0236	-0.0048	0.9956	-0.1650	0.3781	-0.1601	0.3397	-0.1135	0.7779	0.0421	0.8631	0.1556	0.3838
CG12681	CG12681	1633596_at	0.2211	0.2918	0.0991	0.5145	-0.0653	0.7203	-0.2029	0.6084	-0.0070	0.9814	0.1959	0.2753	0.0559	0.9238	-0.1246	0.5411	-0.1805	0.3683

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	**Gene Title**	**Affy ID**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**
CG14290 /// DbuzCG1429(	CG14290	1633597_at	0.0302	0.8961	-0.0296	0.9108	0.2393	0.3328	-0.1176	0.8189	-0.7178	0.0043	-0.6002	0.0057	-0.3841	0.6955	-0.6787	0.1061	-0.2946	0.4889
CG32221	CG32221	1633598_at	0.1896	0.2482	0.0189	0.9297	0.0802	0.6029	0.0428	0.9314	0.3529	0.0246	0.3101	0.0266	0.0274	0.9816	0.2252	0.4093	0.1978	0.4865
Pepck	Phosphoenolpyru	1633599_a_at	1.8998	0.0369	1.1243	0.3096	2.2153	0.0003	0.5147	0.5917	0.6225	0.1990	0.1078	0.8483	-0.4207	0.9095	-0.2775	0.8845	0.1432	0.9382
---	---	1633600_at	0.1213	0.5422	0.1047	0.4547	0.0701	0.7492	0.0683	0.9225	0.0532	0.8463	-0.0151	0.9553	0.0584	0.9365	-0.0836	0.7667	-0.1419	0.5642
Dscam	Down syndrome c	1633601_at	-0.0357	0.8295	0.2087	0.4760	0.3315	0.0877	0.0542	0.9426	-0.0828	0.7438	-0.1371	0.5076	0.0848	0.9152	0.1959	0.4935	0.1111	0.7264
Eaf6	Eaf6	1633602_at	-0.0345	0.8831	-0.6485	0.0435	-0.3071	0.1178	0.2151	0.4573	0.7515	0.0012	0.5364	0.0031	0.0172	0.9898	0.1930	0.5214	0.1758	0.5682
---	---	1633603_at	0.2647	0.3832	0.2488	0.0725	0.4949	0.0264	-0.0072	0.9937	-0.1452	0.4394	-0.1380	0.4155	-0.0325	0.9589	-0.1253	0.4917	-0.0929	0.6284
CG11828	CG11828	1633604_at	0.0550	0.8147	-0.0509	0.8062	-0.0080	0.9776	0.0189	0.9857	0.2493	0.3109	0.2304	0.2968	-0.0036	0.9982	0.0853	0.8500	0.0890	0.8306
CG14100	CG14100	1633605_at	-0.2687	0.2400	-0.1881	0.1792	0.3325	0.1333	0.0849	0.9069	-0.3429	0.1418	-0.4278	0.0467	-0.4930	0.4912	-0.1394	0.7304	0.3536	0.3205
CG33199 /// CG8229 /// Dy	CG8229 /// CG33	1633606_s_at	-0.8832	0.0227	0.0529	0.6544	-0.0153	0.9558	-0.1839	0.7018	-1.2789	0.0006	-1.0951	0.0006	-0.0602	0.9727	-0.2874	0.5300	-0.2272	0.6328
CG2444	CG2444	1633607_at	0.3761	0.0944	0.2511	0.0599	-0.0495	0.8002	-0.0115	0.9893	-0.0265	0.9232	-0.0150	0.9525	0.2108	0.7589	-0.2559	0.3669	-0.4667	0.1358
---	---	1633608_at	0.0984	0.6767	0.2296	0.1604	0.3510	0.1567	-0.0611	0.9310	-0.1579	0.4627	-0.0968	0.6486	-0.0621	0.9514	-0.0449	0.9234	0.0172	0.9702
Gas41	Gas41	1633609_at	0.2406	0.1686	0.4140	0.1961	-0.0172	0.9240	-0.3510	0.2359	-0.0131	0.9604	0.3379	0.0381	0.1035	0.8999	0.1243	0.7200	0.0209	0.9625
CG12419	CG12419	1633610_at	0.1528	0.3791	0.0313	0.7559	-0.0095	0.9711	-0.0434	0.9570	-0.0723	0.7831	-0.0289	0.9125	0.1152	0.8378	-0.0139	0.9714	-0.1291	0.6038
CG32043	CG32043	1633611_s_at	-1.7857	0.0015	-1.2460	0.0031	-1.3884	0.0001	-0.0574	0.9423	-0.4688	0.0440	-0.4115	0.0472	0.0156	0.9898	0.1101	0.7166	0.0945	0.7577
ps	pasilla	1633612_at	0.3941	0.5095	1.8303	0.0213	0.2970	0.6146	-0.9083	0.1209	-0.7081	0.0500	0.2002	0.5674	0.8111	0.7644	0.9842	0.3866	0.1731	0.9132
CG2765 /// DsimCG2765	CG2765	1633613_at	0.5241	0.2068	-0.1218	0.6858	0.1817	0.3937	0.3940	0.5407	0.7703	0.0293	0.3763	0.2095	0.0483	0.9769	0.0657	0.9149	0.0174	0.9778
---	---	1633614_at	0.1046	0.4795	0.0463	0.7142	0.3279	0.0883	-0.0479	0.9461	-0.0954	0.6757	-0.0475	0.8358	-0.1199	0.8270	-0.0988	0.6985	0.0211	0.9471
CG15330	CG15330	1633615_at	0.0082	0.9757	0.0713	0.5191	0.5207	0.1025	0.1697	0.7402	0.0041	0.9899	-0.1656	0.4345	-0.0265	0.9893	0.0394	0.9526	0.0658	0.9080
CG7650	CG7650	1633616_at	0.2043	0.3905	0.5585	0.0466	0.5840	0.0061	-0.0909	0.8967	-0.3402	0.1389	-0.2493	0.2275	0.1058	0.8494	0.0254	0.9462	-0.0804	0.7659
CG15116	CG15116	1633617_at	0.2568	0.1370	-0.0815	0.7421	0.3045	0.0801	0.2879	0.3294	0.3148	0.0684	0.0269	0.8956	-0.2548	0.6749	-0.1609	0.5428	0.0939	0.7486
CG14297 /// DereCG14297	CG14297 /// GA12	1633618_at	-0.0073	0.9890	0.1616	0.4558	0.0079	0.9711	0.0387	0.9763	0.0851	0.8368	0.0465	0.9046	0.1219	0.8141	0.1276	0.5732	0.0057	0.9868
---	---	1633619_at	0.2750	0.2514	-0.1215	0.2266	-0.0707	0.7327	0.1948	0.5821	0.2683	0.1357	0.0734	0.6992	0.0874	0.9076	-0.0446	0.9144	-0.1320	0.6471
---	---	1633620_at	-0.0008	0.9972	-0.1140	0.4299	0.0523	0.7738	-0.0779	0.9293	0.2204	0.4036	0.2983	0.1934	-0.0988	0.8513	0.0214	0.9508	0.1202	0.6094
CG7876	CG7876	1633621_at	0.0799	0.6976	0.2538	0.1852	-0.0169	0.9552	-0.1310	0.8196	-0.1809	0.4386	-0.0499	0.8483	0.2039	0.7779	0.0750	0.8642	-0.1289	0.7155
CG3008	CG3008	1633622_at	0.1913	0.3536	0.6356	0.0455	0.8301	0.0045	0.1312	0.7604	-0.0071	0.9801	-0.1383	0.4249	-0.0609	0.9701	0.4797	0.2329	0.5406	0.2120
sano	serrano	1633623_s_at	0.1966	0.2234	0.1786	0.4735	-0.0704	0.6630	-0.0021	0.9983	-0.0595	0.8235	-0.0575	0.8109	0.3102	0.5754	-0.0186	0.9634	-0.3288	0.2051
alphaTub84D	alpha-Tubulin	1633624_s_at	0.3649	0.4310	-0.0765	0.7379	-0.7240	0.0012	-0.5919	0.0654	0.5709	0.0081	1.1628	0.0002	0.1994	0.8192	0.1286	0.7668	-0.0708	0.8857
PitsIre	PITSLRE	1633625_s_at	-0.0234	0.9226	-0.1479	0.4692	-0.2851	0.0807	0.0034	0.9956	0.3052	0.0422	0.3018	0.0281	0.0579	0.9588	0.1647	0.6216	0.1068	0.7689
CG14104	CG14104	1633626_at	-0.0188	0.9609	0.0701	0.8553	-0.0621	0.7380	-0.0351	0.9733	-0.1985	0.4620	-0.1633	0.5143	0.0694	0.9589	-0.1099	0.8317	-0.1793	0.6731
CG13023	CG13023	1633627_at	-0.0498	0.8266	0.1049	0.4461	-0.0120	0.9674	-0.0654	0.9518	-0.0233	0.9580	0.0421	0.9075	-0.0072	0.9952	0.0497	0.9041	0.0569	0.8761
---	---	1633628_at	0.1857	0.4107	-0.1750	0.4120	-0.0477	0.8286	0.1731	0.7556	0.1613	0.5283	-0.0119	0.9688	0.0575	0.9523	-0.1997	0.5018	-0.2572	0.3833
CG11030	CG11030	1633629_at	-0.1032	0.7145	1.0776	0.0695	1.5830	0.0006	0.1140	0.8932	-0.8287	0.0105	-0.9427	0.0035	-0.3980	0.7822	0.4459	0.4828	0.8438	0.1990
CG4324	CG4324	1633630_at	-0.3613	0.0703	-0.3530	0.1922	-0.6068	0.0164	-0.2955	0.5163	-0.0560	0.8586	0.2394	0.2693	0.2592	0.7230	0.1273	0.7293	-0.1320	0.7130
CG9682	CG9682	1633631_at	0.0500	0.8090	0.0050	0.9659	0.0154	0.9348	0.0902	0.8512	0.1124	0.5508	0.0223	0.9173	0.0035	0.9964	-0.0382	0.9027	-0.0417	0.8813
CG32944	CG32944	1633632_at	-0.5223	0.0585	-0.1342	0.2819	-0.2997	0.1446	0.0688	0.9436	0.1411	0.6496	0.0723	0.8171	0.0175	0.9914	0.1268	0.7903	0.1093	0.8178
CG40274	CG40274	1633633_s_at	-0.0519	0.8633	-0.8405	0.0787	-0.9394	0.0267	-0.1815	0.8875	0.8570	0.0525	1.0385	0.0154	-0.0263	0.9895	0.0965	0.8708	0.1228	0.8127
grau	grauzone	1633634_at	-0.4278	0.1065	-0.1145	0.3729	-0.1376	0.4920	-0.1258	0.7960	-0.2019	0.3124	-0.0761	0.7179	-0.1045	0.8692	0.0797	0.8053	0.1842	0.4799
CG7879 /// DmirCG7879	CG7879	1633635_s_at	-0.3556	0.3459	-0.0605	0.8263	0.0003	0.9989	0.1882	0.8794	-0.1249	0.8112	-0.3131	0.4310	0.0080	0.9950	0.0325	0.9451	0.0244	0.9525
CG10633 /// DmirCG10633	CG10633	1633636_at	0.0361	0.8438	-0.2009	0.1279	-0.3577	0.0399	0.0088	0.9922	0.2382	0.1900	0.2294	0.1593	0.1098	0.8611	-0.0581	0.8734	-0.1679	0.5322
CG17026	CG17026	1633637_at	-1.8536	0.0091	-2.9564	0.0126	-2.8064	0.0000	-0.4811	0.2667	0.1316	0.6477	0.6126	0.0156	-0.4847	0.8128	-0.8515	0.3013	-0.3668	0.6984
---	---	1633638_s_at	0.1220	0.5528	-0.0108	0.9585	0.1152	0.4870	0.0534	0.9377	0.1394	0.5097	0.0860	0.6811	-0.0339	0.9717	-0.0323	0.9341	0.0016	0.9974
Cyp28d1	Cyp28d1	1633639_at	2.1946	0.0076	0.3211	0.3799	0.8258	0.0290	0.1040	0.9251	0.0334	0.9456	-0.0706	0.8549	-0.3943	0.8472	-1.7501	0.0618	-1.3557	0.1393
---	---	1633640_at	-0.0050	0.9804	0.1957	0.3130	0.0882	0.7136	-0.0123	0.9872	-0.1040	0.6067	-0.0917	0.6246	0.1091	0.9096	-0.0909	0.8450	-0.2000	0.5845
CG15611	CG15611	1633641_a_at	-1.2405	0.0057	-0.6089	0.0737	-0.6597	0.0144	0.1871	0.7556	-0.3141	0.2239	-0.5012	0.0371	0.3488	0.7633	0.2014	0.7158	-0.1474	0.8033
CG14189	CG14189	1633642_at	0.2042	0.3389	0.2030	0.4388	-0.0737	0.7516	-0.0375	0.9705	-0.0141	0.9720	0.0234	0.9425	0.0543	0.9525	-0.1026	0.7506	-0.1568	0.5907
CG31033	CG31033	1633643_a_at	0.2676	0.1991	0.2424	0.0917	0.1027	0.7354	-0.1934	0.5461	0.0205	0.9298	0.2139	0.1484	0.0977	0.9449	-0.0432	0.9504	-0.1410	0.7813
CG13151	CG13151	1633644_at	-0.0849	0.7488	0.1132	0.6152	0.2902	0.1533	-0.0725	0.8942	0.0564	0.8006	0.1289	0.4529	-0.0363	0.9841	0.3995	0.3236	0.4358	0.3042
CG13512 /// DyakCG13512	CG13512	1633645_at	0.0435	0.8603	-0.1590	0.6492	-0.1974	0.6223	-0.0220	0.9863	0.1601	0.6409	0.1821	0.5431	-0.0820	0.9734	0.0733	0.9404	0.1553	0.8413
inaF	inaF	1633646_at	0.3345	0.0920	0.0474	0.6528	0.0154	0.9299	-0.0796	0.8718	0.0225	0.9252	0.1021	0.5446	0.0149	0.9913	-0.1857	0.5191	-0.2006	0.4856
Cpr49Ah	CG8515	1633647_at	-0.1629	0.2532	-0.0296	0.8796	0.0918	0.6362	0.0952	0.8905	-0.2237	0.3426	-0.3190	0.1219	-0.0345	0.9653	-0.0060	0.9898	0.0285	0.9224
sns	sticks and stones	1633648_at	-0.0215	0.9083	0.1692	0.2683	0.2354	0.1367	-0.0732	0.8872	-0.2512	0.1475	-0.1780	0.2560	-0.1212	0.8141	0.0040	0.9935	0.1251	0.5811
trpl	trp-like	1633649_s_at	-0.0440	0.8478	-0.0070	0.9506	0.0016	0.9943	0.0360	0.9678	-0.0096	0.9790	-0.0457	0.8682	-0.0201	0.9816	-0.0142	0.9631	0.0060	0.9847
CG5380	CG5380	1633650_at	0.2121	0.2824	0.0631	0.8614	0.0483	0.8437	-0.1543	0.6885	0.2512	0.1573	0.4054	0.0192	-0.1323	0.9108	0.0643	0.9221	0.1967	0.6669
Tk	tachykinin-3	1633651_at	-0.6406	0.6973	0.1743	0.2941	0.2069	0.4706	-0.2300	0.9603	-1.3809	0.2675	-1.1509	0.3066	-0.0710	0.9848	-0.3606	0.6969	-0.2895	0.7614
CG12784	CG12784	1633652_at	-0.1274	0.4552	0.0516	0.7683	0.1252	0.5055	0.0415	0.9518	-0.1186	0.5665	-0.1601	0.3646	0.0329	0.9683	0.1216	0.6019	0.0886	0.7205
CG8647	CG8647	1633653_at	-0.0152	0.9684	0.1842	0.2755	0.0321	0.8992	-0.2425	0.4821	-0.2707	0.1410	-0.0282	0.8991	-0.1240	0.8861	-0.1177	0.7649	0.0063	0.9913
---	---	1633654_s_at	0.3795	0.2934	0.2213	0.6827	0.6306	0.0312	0.0524	0.9601	0.1111	0.7299	0.0587	0.8543	-0.3184	0.8692	-0.0490	0.9696	0.2694	0.7655
---	---	1633655_at	0.2151	0.3098	0.0527	0.7074	-0.0752	0.7475	-0.1737	0.7974	-0.0107	0.9800	0.1630	0.5446	0.1028	0.9246	-0.1941	0.6181	-0.2969	0.4243
CG13894	CG13894	1633656_at	-0.2434	0.2092	0.2217	0.2942	0.1043	0.5769	0.0616	0.9559	-0.2911	0.3435	-0.3527	0.1909	0.1274	0.8215	0.1750	0.4596	0.0475	0.8821
CG31533	CG31533	1633657_at	0.1190	0.6275	0.1340	0.3181	0.1597	0.3522	0.0946	0.8325	0.0029	0.9903	-0.0917	0.5895	0.0210	0.9862	0.1430	0.6303	0.1220	0.6881
CG33233	CG33233	1633658_a_at	-0.0968	0.6015	0.0443	0.6976	0.3067	0.1530	0.2139	0.5229	0.0058	0.9822	-0.2081	0.1838	-0.0293	0.9677	0.0349	0.9046	0.0641	0.7749
CG11349	CG11349	1633659_at	-0.1818	0.4049	0.0601	0.6128	-0.1455	0.4188	-0.1284	0.8217	-0.2707	0.2218	-0.1424	0.5063	-0.0445	0.9589	-0.0974	0.7277	-0.0530	0.8699
Orc5	lethal(2)34Df	1633660_at	0.2397	0.2833	0.0410	0.8835	0.1203	0.5757	0.1117	0.8470	0.4959	0.0273	0.3842	0.0482	-0.2703	0.7506	0.0493	0.9326	0.3196	0.3895
CG11377	CG11377	1633661_at	0.3300	0.1350	0.9721	0.0328	1.3549	0.0013	0.0510	0.9608	-0.1890	0.5201	-0.2400	0.3438	-0.3780	0.6960	0.6455	0.1199	1.0235	0.0499
CG8290	CG8290	1633662_a_at	-0.1260	0.6376	0.0804	0.6244	0.0905	0.8024	-0.0972	0.9254	-0.1418	0.6930	-0.0446	0.9061	0.0165	0.9913	0.0078	0.9911	-0.0087	0.9856
---	---	1633663_at	-0.1007	0.5768	0.0843	0.6940	0.2052	0.2366	0.0167	0.9852	-0.0718	0.7642	-0.0885	0.6705	-0.0864	0.8692	-0.0665	0.8052	0.0199	0.9463
bc10	bc10	1633664_at	0.4626	0.3253	0.9616	0.0177	0.2990	0.5123	0.3231	0.5311	0.3036	0.2609	-0.0195	0.9548	0.8061	0.7202	0.7210	0.4391	-0.0851	0.9485
lic	licorne	1633665_at	-0.0433	0.8530	-0.0033	0.9901	-0.2914	0.1368	-0.2454	0.5138	0.1698	0.4054	0.4153	0.0273	-0.0632	0.9495	0.0659	0.8764	0.1291	0.6963

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG30037	CG30037	1633666_at	0.3321	0.1394	-0.0201	0.9059	0.3584	0.1249	0.2202	0.4910	-0.1299	0.4654	-0.3501	0.0293	-0.0670	0.9359	-0.1388	0.6264	-0.0718	0.8287
Lip2	Lipase 2	1633667_at	0.3174	0.1853	-0.0055	0.9857	0.4744	0.0313	0.0810	0.8289	0.1161	0.4361	0.0352	0.8298	-0.3117	0.7768	-0.2977	0.5495	0.0140	0.9852
---	---	1633668_at	0.1583	0.4515	0.1128	0.4739	0.1466	0.5201	0.0204	0.9819	-0.0255	0.9320	-0.0459	0.8526	0.0216	0.9845	-0.0639	0.8500	-0.0855	0.7678
CG31798	CG31798	1633669_at	0.0355	0.9003	-0.0279	0.8233	0.2685	0.1475	0.0967	0.8869	-0.1972	0.4036	-0.2939	0.1515	-0.0887	0.8903	-0.1452	0.5670	-0.0565	0.8577
---	---	1633670_at	0.1041	0.5385	-0.1383	0.7972	-0.1053	0.6586	0.1236	0.8738	0.2200	0.4343	0.0964	0.7416	0.2801	0.8270	0.1838	0.7787	-0.0963	0.8949
---	---	1633671_at	0.1760	0.2393	-0.0883	0.6600	0.0682	0.7704	-0.0655	0.9361	0.0884	0.7578	0.1539	0.5054	-0.1184	0.8192	-0.0981	0.6762	0.0203	0.9453
CG12992	CG12992	1633672_at	1.3153	0.0108	1.7541	0.0849	0.9810	0.0195	0.3350	0.4551	0.6490	0.0157	0.3140	0.1473	1.1040	0.6496	1.2634	0.1849	0.1594	0.9065
ATPsyn-beta	H+-ATPase beta s	1633673_a_at	-0.2523	0.6710	2.4573	0.0077	2.4299	0.0003	-0.4698	0.5702	-2.9891	0.0003	-2.5193	0.0003	-0.4507	0.8099	-0.2350	0.8166	0.2156	0.8255
CG31150	CG31150	1633674_at	0.3885	0.4168	-0.3855	0.6115	0.3201	0.1685	-0.0423	0.9558	0.4152	0.0481	0.4575	0.0208	-0.7845	0.7215	-0.2987	0.7971	0.4858	0.6231
CG30008	CG30008	1633675_at	0.6807	0.7213	-2.1461	0.4895	1.6366	0.0949	2.4269	0.3127	1.0913	0.4560	-1.3356	0.2947	-1.3895	0.8728	-1.6733	0.6536	-0.2838	0.9527
CG40169	CG40169	1633676_a_at	-0.0485	0.7989	-0.1546	0.3761	-0.0416	0.8067	-0.0746	0.8732	-0.0089	0.9719	0.0658	0.7008	-0.0754	0.8961	-0.2011	0.3389	-0.1256	0.5846
eap	Ebony Activating I	1633677_at	0.4696	0.1317	0.7482	0.0062	1.3510	0.0002	-0.3481	0.2949	-1.1019	0.0007	-0.7538	0.0018	-0.9304	0.1902	-0.7235	0.0665	0.2069	0.5995
CG40154	CG40154	1633678_at	0.3318	0.0898	-0.0033	0.9812	0.1218	0.4728	0.1124	0.7795	0.4444	0.0158	0.3320	0.0334	-0.2019	0.7307	-0.2077	0.4050	-0.0058	0.9885
---	---	1633679_at	0.0499	0.7319	0.0982	0.5146	0.0041	0.9876	-0.1457	0.6908	-0.1345	0.4455	0.0112	0.9587	0.0360	0.9589	-0.0014	0.9986	-0.0374	0.8896
CG12818	CG12818	1633680_at	0.3889	0.2488	-0.0119	0.9291	-0.3315	0.1932	0.0116	0.9864	0.5634	0.0052	0.5518	0.0034	0.2405	0.8472	0.1261	0.8617	-0.1144	0.8684
CG2519	CG2519	1633681_at	-0.6443	0.0343	-0.8570	0.0120	-1.1465	0.0008	-0.2438	0.5639	0.0100	0.9763	0.2539	0.1899	0.0125	0.9943	-0.3011	0.4531	-0.3137	0.4464
l(1)G0232	lethal (1) G0232	1633682_a_at	-0.3141	0.5301	0.3778	0.3486	0.1539	0.5358	-0.2509	0.6964	-0.2718	0.3694	-0.0209	0.9559	0.0091	0.9976	0.5522	0.4255	0.5431	0.4475
CG9602	CG9602	1633683_at	0.2167	0.2845	-0.1026	0.6123	0.2030	0.2352	0.0902	0.8951	0.0720	0.7965	-0.0182	0.9496	-0.2140	0.7681	-0.1282	0.7151	0.0858	0.8243
CG32444 /// DsimCG32444	aldose 1-epimera	1633684_at	0.6904	0.0217	2.3935	0.0751	2.5766	0.0015	-0.2114	0.9005	-2.6472	0.0012	-2.4358	0.0010	-0.4361	0.8746	-1.2423	0.2380	-0.8062	0.4741
---	---	1633685_at	0.0632	0.7061	0.0895	0.4522	-0.2439	0.1428	-0.1970	0.6226	-0.1375	0.5137	0.0595	0.7889	0.0700	0.9101	0.0109	0.9783	-0.0591	0.8310
CG3225	CG3225	1633686_at	-0.1533	0.6214	0.1167	0.4228	0.0761	0.7833	0.1632	0.8140	0.3189	0.2448	0.1556	0.5659	0.0959	0.9449	0.4964	0.2222	0.4005	0.3559
CG13611	CG13611	1633687_at	-0.1055	0.5778	0.0451	0.6594	0.0081	0.9681	0.0861	0.8408	-0.0773	0.6715	-0.1634	0.2562	0.0094	0.9934	0.1961	0.4507	0.1867	0.4865
---	---	1633688_at	0.0329	0.8666	-0.0467	0.7026	0.4023	0.0300	0.1940	0.6755	0.0411	0.8894	-0.1530	0.4586	-0.0948	0.8930	0.0226	0.9587	0.1174	0.6863
CG40368	CG40368	1633689_at	0.0261	0.8741	-0.0881	0.5096	0.0625	0.7096	0.0303	0.9562	-0.0421	0.8210	-0.0724	0.6354	-0.1770	0.7266	-0.1207	0.6053	0.0563	0.8401
CG8323	CG8323	1633690_at	0.3220	0.0648	0.3940	0.1697	0.7353	0.0291	-0.1618	0.5683	-0.1523	0.3008	0.0096	0.9587	-0.5503	0.6483	-0.2410	0.6473	0.3093	0.5460
CG32262	CG32262	1633691_at	-0.0245	0.9091	0.4062	0.1475	0.6220	0.0038	0.1681	0.5988	-0.2248	0.1612	-0.3929	0.0144	0.0214	0.9898	0.1225	0.7836	0.1011	0.8227
CG15240	CG15240	1633692_at	0.2079	0.4201	-0.0657	0.6212	0.0336	0.9180	0.1453	0.7708	0.1829	0.3974	0.0376	0.8802	-0.0269	0.9901	0.0671	0.9287	0.0939	0.8858
CG13814	CG13814	1633693_at	0.0708	0.7693	0.0195	0.8454	0.0250	0.9235	0.0067	0.9937	0.0896	0.6505	0.0829	0.6467	0.1482	0.8655	-0.0351	0.9505	-0.1833	0.6266
---	---	1633694_at	0.2726	0.2234	-0.0667	0.7360	0.5298	0.0290	0.1590	0.7680	0.0474	0.8765	-0.1116	0.6338	-0.3384	0.5200	-0.1756	0.4959	0.1629	0.5391
---	---	1633695_at	1.1601	0.0117	2.3257	0.0054	2.6370	0.0000	0.1076	0.8190	0.1039	0.6081	-0.0037	0.9873	-0.1081	0.9412	0.5169	0.2389	0.6250	0.1914
TM4SF	Transmembrane 4	1633696_at	-0.3428	0.2508	-0.7367	0.0083	-0.4839	0.0061	0.1485	0.7266	0.1770	0.3621	0.0285	0.9007	-0.1992	0.7751	-0.3878	0.1849	-0.1887	0.5540
---	---	1633697_at	0.0291	0.9232	-0.2811	0.2965	0.0525	0.7869	0.2406	0.5913	0.2472	0.2794	0.0065	0.9817	-0.0743	0.9467	0.0457	0.9325	0.1199	0.7571
l(3)neo18	lethal (3) neo18	1633698_at	-0.2068	0.4494	0.3917	0.1217	0.4538	0.0863	0.0962	0.8844	-0.9721	0.0019	-1.0683	0.0008	0.0766	0.9558	-0.3311	0.4067	-0.4077	0.3212
CG9962	CKIa-like	1633699_at	0.1709	0.4580	0.0094	0.9864	0.1469	0.5333	0.0162	0.9853	0.0435	0.8671	0.0274	0.9086	-0.0486	0.9792	-0.1492	0.7763	-0.1007	0.8570
Rpd3	Suppressor of var	1633700_at	0.2014	0.5289	0.5357	0.1648	0.5527	0.0282	0.0922	0.8422	0.5315	0.0082	0.4393	0.0117	-0.0528	0.9816	0.7450	0.1495	0.7978	0.1562
CG16985	CG16985	1633701_at	-0.0762	0.8030	-0.1862	0.6942	0.0572	0.7978	0.0230	0.9774	-0.3620	0.0723	-0.3850	0.0375	-0.3571	0.8016	-0.5043	0.3873	-0.1472	0.8467
CG12849	CG12849	1633702_at	0.0984	0.6169	0.1497	0.2851	0.3241	0.0397	0.0553	0.9248	-0.0232	0.9247	-0.0785	0.6671	-0.0206	0.9862	0.0458	0.9111	0.0664	0.8463
Pbprp2	Pheromone-bindir	1633703_s_at	0.1487	0.7563	0.3879	0.1368	-0.0448	0.9065	-0.4853	0.5434	-0.7217	0.0830	-0.2364	0.5646	0.1756	0.9296	-0.2908	0.6839	-0.4664	0.4822
CG7045 /// DsimCG7045	CG7045	1633704_at	-0.0770	0.6423	-0.0372	0.7151	-0.0692	0.6599	0.0893	0.8189	-0.0597	0.7401	-0.1490	0.2797	0.0729	0.8795	0.0310	0.9137	-0.0420	0.8635
---	---	1633705_at	0.0200	0.9173	0.0609	0.6110	0.0410	0.7931	0.1111	0.7690	-0.1057	0.5386	-0.2168	0.1290	0.0049	0.9964	0.0056	0.9935	0.0007	0.9992
CG7714	CG7714	1633706_at	0.5902	0.2908	0.1776	0.3264	0.2032	0.4394	-0.3457	0.4908	-0.1657	0.5656	0.1800	0.4801	-0.0865	0.9775	-0.5106	0.4881	-0.4240	0.5808
CG12824	CG12824	1633707_at	-0.1707	0.8652	0.2066	0.1365	-0.0122	0.9674	-0.3910	0.8094	-0.8129	0.2058	-0.4219	0.4955	-0.1344	0.9467	-0.4532	0.4694	-0.3188	0.6317
Gr59b	Gustatory recepto	1633708_at	0.2302	0.2644	-0.0218	0.9002	0.0123	0.9508	0.0064	0.9932	0.0221	0.9158	0.0157	0.9319	-0.0497	0.9775	-0.0714	0.9111	-0.0217	0.9732
CG2772	CG2772	1633709_at	-0.1847	0.4115	-0.2045	0.1702	-0.3394	0.0576	0.0919	0.8897	0.0001	0.9997	-0.0918	0.6936	0.1548	0.7758	-0.1057	0.6875	-0.2605	0.2832
ChLD3	CG17905	1633710_at	-0.2648	0.6120	-0.0198	0.8519	-0.4345	0.0228	0.0214	0.9883	-0.3602	0.3052	-0.3817	0.2204	0.2847	0.8222	-0.2564	0.6555	-0.5411	0.3141
CG31012 /// DmadCG3101	CG31012	1633711_a_at	-1.4265	0.0778	-0.5974	0.2176	-0.0607	0.8453	0.4910	0.6122	-0.4696	0.3425	-0.9606	0.0342	0.0347	0.9926	0.3898	0.6820	0.3551	0.7128
CG31997	CG31997	1633712_at	-0.5374	0.0203	-0.2089	0.2005	-0.2632	0.3065	0.1683	0.7188	-0.2656	0.2083	-0.4339	0.0296	0.2815	0.7633	0.0334	0.9590	-0.2481	0.5488
CG33260 /// Trl	GAGA factor /// C	1633713_at	0.3454	0.0535	0.8073	0.1445	0.8668	0.0037	-0.1320	0.7845	-0.2798	0.1578	-0.1478	0.4285	-0.0973	0.9441	0.2864	0.5084	0.3837	0.3723
eys	eyes shut	1633714_at	0.2920	0.1827	0.0991	0.5659	0.1807	0.3599	-0.1789	0.6705	-0.0262	0.9252	0.1527	0.4134	-0.0481	0.9633	-0.0910	0.8031	-0.0429	0.9127
CG18769	CG18769	1633715_s_at	0.9912	0.2328	0.6862	0.0917	0.5709	0.0122	0.2429	0.5140	0.2583	0.1866	0.0154	0.9506	0.4281	0.8760	-0.0509	0.9791	-0.4790	0.6867
CG10747	CG10747	1633716_at	-1.4500	0.0009	-0.8766	0.0229	-0.8546	0.0009	-0.0920	0.8362	-0.7721	0.0015	-0.6801	0.0015	-0.0093	0.9943	-0.2262	0.4487	-0.2169	0.4824
CG2199	CG2199	1633717_a_at	0.7620	0.0270	0.3113	0.4759	0.4863	0.0058	-0.1026	0.8875	0.5732	0.0270	0.6758	0.0082	-0.2544	0.8236	0.0463	0.9517	0.3008	0.5501
CG5355	CG5355	1633718_at	0.3400	0.1085	0.6451	0.2193	0.7532	0.0012	0.0316	0.9586	0.2464	0.1220	0.2148	0.1315	-0.0929	0.9587	0.5166	0.2825	0.6096	0.2341
Ugt86Dg	Ugt86Dg	1633719_at	0.1404	0.4041	0.1924	0.1241	-0.0084	0.9641	0.0055	0.9946	0.0950	0.6137	0.0896	0.6018	0.1227	0.7810	0.0733	0.7439	-0.0494	0.8389
CG8795	CG8795	1633720_s_at	-1.9781	0.0025	-0.5390	0.1388	-1.2575	0.0005	-0.6229	0.1526	-1.1819	0.0016	-0.5590	0.0258	-0.0331	0.9589	-0.0224	0.9401	0.0107	0.9695
---	---	1633721_at	0.0399	0.8295	0.1848	0.3486	0.0688	0.6753	-0.1137	0.7949	-0.2292	0.1965	-0.1155	0.5002	0.0587	0.9457	-0.0045	0.9941	-0.0632	0.8517
---	---	1633722_at	0.2148	0.4379	0.0385	0.7401	-0.1164	0.4385	0.0380	0.9569	0.1359	0.4932	0.0979	0.6081	0.2135	0.7691	0.0764	0.8583	-0.1371	0.6881
mav	maverick	1633723_at	-2.0848	0.0023	-2.6094	0.0321	-3.0639	0.0000	-0.3829	0.3129	-0.1856	0.4185	0.1973	0.3310	0.1330	0.9676	-0.6678	0.4354	-0.8008	0.3631
grim	grim	1633724_at	0.1717	0.4655	0.0019	0.9882	0.1627	0.4728	-0.0248	0.9744	0.1658	0.3898	0.1906	0.2599	-0.0567	0.9689	0.0266	0.9657	0.0833	0.8692
---	---	1633725_at	0.1844	0.2377	0.2076	0.1601	0.0461	0.8277	-0.0291	0.9677	0.2336	0.2147	0.2627	0.1187	-0.0051	0.9964	0.0953	0.7837	0.1003	0.7560
CG14082	CG14082	1633726_s_at	0.0696	0.7751	0.1044	0.4266	0.2246	0.1393	0.0887	0.8968	0.0303	0.9231	-0.0584	0.8194	-0.0814	0.9092	0.0780	0.8127	0.1594	0.5523
wnd	CG8789	1633727_s_at	0.7081	0.0315	0.0268	0.9515	0.0358	0.8663	-0.0592	0.9370	0.5128	0.0261	0.5719	0.0102	0.0252	0.9898	-0.1598	0.7562	-0.1850	0.7020
CG12728	CG12728	1633728_at	-0.0685	0.8348	0.8786	0.0394	0.9928	0.0380	-0.0767	0.9165	-0.7481	0.0062	-0.6714	0.0061	-0.1097	0.9599	0.3738	0.5652	0.4835	0.4498
Eps-15	Eps-15	1633729_a_at	-0.6909	0.1018	-0.3981	0.6477	-0.1798	0.3756	0.0534	0.9098	-0.1262	0.4137	-0.1796	0.1787	-0.1759	0.9547	0.0484	0.9758	0.2243	0.8474
dik	diskette	1633730_at	-0.3197	0.1765	-0.3415	0.5432	-0.4570	0.1055	-0.1111	0.8498	-0.2353	0.2734	-0.1241	0.5557	0.0018	0.9998	-0.1925	0.8180	-0.1943	0.8044
wee	wee	1633731_at	0.3183	0.3230	-0.0856	0.7957	-0.2336	0.3155	-0.0005	0.9997	0.8165	0.0257	0.8170	0.0161	0.2787	0.7997	0.3635	0.4268	0.0848	0.8926
CG14713	CG14713	1633732_at	-0.7237	0.0599	1.0905	0.2158	0.3413	0.1793	-0.1376	0.9130	-0.9666	0.0233	-0.8290	0.0277	0.5621	0.7485	0.7859	0.2738	0.2238	0.8056
svr	carboxypeptidase	1633733_a_at	-0.1008	0.8553	0.3535	0.5334	0.2288	0.1440	-0.1158	0.8633	-0.2940	0.2173	-0.1782	0.4267	0.0699	0.9840	0.3124	0.7340	0.2426	0.8001
CG7261	CG7261	1633734_at	0.2293	0.2377	-0.2439	0.2173	-0.1158	0.5489	0.3420	0.2455	1.0034	0.0006	0.6613	0.0020	0.0481	0.9677	0.4610	0.1382	0.4129	0.2079

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG1407	CG1407	1633735_a_at	-0.3152	0.4211	0.5241	0.0809	0.6635	0.0162	-0.2522	0.7525	-0.8195	0.0259	-0.5672	0.0696	-0.1032	0.9514	0.2739	0.6152	0.3770	0.4740
CG10333	CG10333	1633736_at	-0.0010	0.9964	-0.0745	0.8459	0.0076	0.9757	-0.0456	0.9376	0.2966	0.0835	0.3422	0.0323	-0.1062	0.9400	0.2196	0.6392	0.3258	0.4655
---	---	1633737_at	0.1024	0.7642	0.0928	0.5031	-0.0672	0.7936	-0.1153	0.8777	0.0986	0.7464	0.2139	0.3675	0.1200	0.8731	0.1394	0.6695	0.0194	0.9649
Gr64f	Gustatory recepto	1633738_at	0.2384	0.1485	0.0573	0.7159	0.1224	0.5992	0.0833	0.8796	0.0953	0.6540	0.0120	0.9597	0.0347	0.9816	0.0582	0.9085	0.0235	0.9616
EfSec	EfSec	1633739_at	0.4491	0.1441	0.6084	0.0884	0.9602	0.0006	-0.1029	0.9068	0.0413	0.9169	0.1442	0.6146	-0.4396	0.5754	0.1106	0.8094	0.5502	0.1479
CG1885	CG1885	1633740_at	-0.4746	0.0270	0.3008	0.1419	0.3599	0.2402	-0.2874	0.3580	-0.5149	0.0111	-0.2276	0.1526	-0.2839	0.7597	0.2882	0.4640	0.5721	0.1651
CR31032	CR31032	1633741_at	0.5103	0.0418	-0.1919	0.5185	0.0938	0.5585	0.0384	0.9599	0.3438	0.0870	0.3054	0.0881	-0.3311	0.7062	-0.5053	0.1542	-0.1742	0.6605
CG40164	CG40164	1633742_at	0.1328	0.5590	-0.0163	0.9176	0.0678	0.7007	0.0854	0.9117	0.2067	0.4126	0.1213	0.6262	-0.0177	0.9862	-0.0490	0.8824	-0.0313	0.9211
RpII15	RNA polymerase	1633743_at	-0.0103	0.9740	0.4556	0.1937	0.5685	0.0201	0.0377	0.9704	-0.5460	0.0411	-0.5837	0.0198	-0.2037	0.8510	-0.1143	0.8528	0.0894	0.8846
---	---	1633744_at	0.1598	0.4049	0.1810	0.3361	0.4554	0.0674	0.0699	0.9017	-0.1357	0.4806	-0.2055	0.2112	-0.0131	0.9923	0.0143	0.9797	0.0274	0.9495
shot	kakapo	1633745_a_at	-0.1620	0.5874	-0.9579	0.0380	-1.2429	0.0010	0.2633	0.7278	1.2059	0.0044	0.9426	0.0080	0.5162	0.6092	0.4556	0.2775	-0.0606	0.9194
a	arc	1633746_s_at	2.5023	0.0022	2.0492	0.0155	2.9068	0.0000	0.6223	0.3171	0.2769	0.4652	-0.3454	0.2940	-0.1324	0.9499	0.1023	0.9161	0.2347	0.7456
CG15048	CG15048	1633747_at	0.1895	0.2193	0.0114	0.9569	0.1934	0.2337	0.0822	0.8776	0.0353	0.8865	-0.0470	0.8214	-0.1195	0.8386	-0.1836	0.4479	-0.0642	0.8334
CG2014	CG2014	1633748_at	-0.0500	0.9060	0.2595	0.0634	0.4236	0.0145	-0.0033	0.9956	-0.3060	0.0749	-0.3027	0.0514	-0.1382	0.9112	0.0167	0.9842	0.1549	0.7615
CG6337	CG6337	1633749_at	1.2663	0.4957	0.8475	0.4728	-4.2108	0.0002	-4.3124	0.0033	-0.4264	0.5386	3.8860	0.0003	0.7158	0.9340	-0.7458	0.8407	-1.4616	0.6228
PNUTS	PNUTS	1633750_s_at	-0.2015	0.4173	-0.0471	0.9189	-0.2051	0.4538	-0.3690	0.3886	-0.3034	0.2077	0.0656	0.8091	-0.2539	0.8461	-0.2626	0.6586	-0.0087	0.9924
CG10903	CG10903	1633751_at	0.0265	0.9588	-0.0826	0.8817	0.3112	0.2684	0.2634	0.5633	0.4428	0.0624	0.1793	0.4126	-0.1770	0.9405	0.2419	0.7924	0.4189	0.5942
CG34413	CG32836	1633752_at	-0.6822	0.0588	-0.2738	0.3420	-0.3027	0.1389	0.0137	0.9866	-0.3651	0.0711	-0.3789	0.0406	-0.2107	0.8395	-0.2505	0.5823	-0.0398	0.9471
otk	Off-track	1633753_at	-0.3696	0.1477	-0.6338	0.0676	-0.3224	0.1709	0.4694	0.4464	0.2807	0.4190	-0.1887	0.5719	0.0162	0.9928	-0.0909	0.8807	-0.1071	0.8426
CG5626	CG5626	1633754_at	0.1294	0.6519	-0.1258	0.4399	-0.1256	0.6765	-0.1021	0.9080	0.3952	0.1607	0.4973	0.0537	0.0742	0.9445	0.2732	0.3909	0.1989	0.5612
CG7706 /// DyakCG7706	CG7706	1633755_at	-0.0588	0.8016	-0.0984	0.4304	-0.0023	0.9931	0.1086	0.7947	0.4341	0.0189	0.3255	0.0388	-0.0179	0.9913	0.4320	0.1925	0.4499	0.2090
---	---	1633756_at	0.2037	0.2409	0.0677	0.5258	0.1292	0.3998	-0.0124	0.9847	-0.0895	0.5731	-0.0771	0.6023	-0.0661	0.9306	-0.0988	0.7271	-0.0327	0.9231
CG10188	CG10188	1633757_s_at	1.7859	0.0013	0.8435	0.0122	0.6223	0.1226	0.0753	0.8803	0.7813	0.0015	0.7060	0.0014	0.1196	0.9555	-0.2906	0.6754	-0.4102	0.5332
Tsp86D	Tetraspanin 86D	1633758_at	-0.5108	0.0401	0.2832	0.2442	0.9929	0.0084	0.3374	0.2438	-0.4218	0.0222	-0.7592	0.0010	-0.2395	0.8481	0.4029	0.4392	0.6424	0.2338
larp	meteor	1633759_s_at	0.0674	0.9442	0.9258	0.1747	0.8209	0.0014	-0.0297	0.9847	-0.4644	0.1843	-0.4347	0.1659	0.2239	0.9503	0.3743	0.7743	0.1504	0.9189
CG10399	CG10399	1633760_at	0.2358	0.3390	0.1287	0.2837	0.4940	0.0568	-0.0141	0.9860	-0.0681	0.7601	-0.0540	0.7966	-0.1955	0.8270	-0.1284	0.7784	0.0670	0.8949
CG14930	CG14930	1633761_at	-0.0484	0.8654	0.0356	0.7733	0.1119	0.5945	0.1680	0.6901	0.0725	0.7553	-0.0955	0.6321	0.0120	0.9926	-0.0376	0.9366	-0.0496	0.9031
---	---	1633762_at	0.1073	0.5981	-0.0045	0.9787	0.3045	0.0798	0.0523	0.9517	0.0906	0.7482	0.0383	0.8930	-0.1679	0.7464	0.0682	0.8117	0.2361	0.2964
aph-1	presenilin enhano	1633763_at	0.2386	0.2860	0.3597	0.1086	0.4831	0.0179	-0.0543	0.9311	-0.2831	0.1220	-0.2288	0.1630	-0.0694	0.9405	-0.0910	0.8053	-0.0215	0.9589
CG4949 /// DyakCG4949	CG4949	1633764_a_at	0.4997	0.1219	0.3749	0.1118	0.1463	0.5250	-0.2637	0.6120	0.0924	0.7675	0.3561	0.1247	0.0082	0.9950	0.0886	0.8244	0.0804	0.8344
asparagine-synthetase	---	1633765_at	1.1054	0.0845	1.1543	0.0555	2.7064	0.0012	1.2701	0.2429	1.3782	0.0386	0.1081	0.8879	-0.2053	0.9421	1.4876	0.0966	1.6929	0.0878
CG5241	CG5241	1633766_at	-1.5898	0.0022	0.0093	0.9907	0.1496	0.7355	0.2947	0.5381	-0.9471	0.0033	-1.2418	0.0006	-0.1021	0.9611	0.3470	0.5747	0.4491	0.4573
Hrp59	heterogeneous nu	1633767_at	0.1072	0.5151	0.4231	0.1367	0.2032	0.2890	0.0272	0.9688	0.1800	0.3257	0.1529	0.3568	0.2632	0.7307	0.5650	0.0950	0.3019	0.3708
CG31814	CG31814	1633768_at	-0.3353	0.1133	0.0420	0.6770	0.0135	0.9443	-0.0024	0.9973	-0.2181	0.2578	-0.2158	0.2091	-0.0802	0.8599	0.1051	0.5875	0.1853	0.3208
CG4613	CG4613	1633769_at	1.3667	0.0233	0.1361	0.4979	0.7038	0.0267	0.6104	0.2955	1.1590	0.0055	0.5486	0.0754	0.2298	0.8874	0.1426	0.8678	-0.0872	0.9173
CG11967	CG11967	1633770_at	1.7943	0.0082	0.7522	0.0336	1.7047	0.0002	0.4739	0.3102	0.4500	0.1002	-0.0239	0.9448	-0.4620	0.7768	-0.6093	0.3787	-0.1474	0.8779
CG2017	GP-1 related	1633771_s_at	-0.4079	0.2645	0.9198	0.0818	1.4329	0.0001	0.1382	0.7293	-1.0417	0.0006	-1.1799	0.0002	-0.3309	0.8202	0.2584	0.7019	0.5893	0.3360
CG32564	CG32564	1633772_at	0.6394	0.1984	-0.1811	0.3582	-0.3221	0.6772	-0.3773	0.6837	0.2895	0.5283	0.6668	0.0842	-0.3464	0.8991	-0.5153	0.6382	-0.1688	0.9032
Keap1	Keap1	1633773_s_at	0.5532	0.2373	-0.2094	0.6714	-0.1188	0.6597	-0.0209	0.9857	0.4920	0.0665	0.5129	0.0371	-0.0417	0.9895	-0.1219	0.9050	-0.0802	0.9318
CG32152	CG32152	1633774_at	0.0280	0.8993	0.1022	0.5384	0.1507	0.5365	-0.1139	0.8143	-0.1068	0.6123	0.0071	0.9763	-0.1821	0.8282	-0.0242	0.9664	0.1579	0.6843
CG31665	fibropellin	1633775_at	0.0628	0.7125	-0.0393	0.6994	-0.1324	0.4123	0.0455	0.9361	0.0537	0.7923	0.0082	0.9688	-0.0084	0.9939	-0.1663	0.5074	-0.1579	0.5377
Mst35Ba /// Mst35Bb	protamine	1633776_s_at	0.0541	0.8382	-0.0217	0.9501	-0.5070	0.1066	-0.1200	0.8800	0.3226	0.2366	0.4426	0.0716	0.0614	0.9589	0.0828	0.8584	0.0213	0.9659
CG11221	CG11221	1633777_at	-2.3361	0.0037	-1.9541	0.0217	-2.5398	0.0000	0.0970	0.8578	0.1200	0.5664	0.0230	0.9230	0.2757	0.8940	0.0575	0.9642	-0.2183	0.8209
gb	CG6070	1633778_at	1.5924	0.0012	0.6841	0.2894	1.1174	0.0026	0.0402	0.9666	0.7365	0.0105	0.6963	0.0082	-0.4832	0.7893	-0.1882	0.8611	0.2950	0.7410
loj	logjam	1633779_s_at	0.5973	0.0244	0.6762	0.0518	0.7845	0.0038	0.1987	0.5842	0.2518	0.1715	0.0531	0.7981	0.0737	0.9588	0.3866	0.3185	0.3129	0.4457
CG13793	CG13793	1633780_at	0.1726	0.8091	0.0502	0.8920	-0.2223	0.6900	-0.7055	0.5492	-1.3340	0.0354	-0.6285	0.2503	-0.3486	0.8611	-1.1767	0.1399	-0.8282	0.3168
---	---	1633781_at	0.1155	0.6050	0.0896	0.6072	0.1490	0.3428	0.1239	0.8327	0.0221	0.9469	-0.1018	0.6591	0.2131	0.7707	0.0534	0.9115	-0.1597	0.6354
---	---	1633782_at	0.2345	0.1724	0.0849	0.6286	-0.0681	0.7762	-0.0901	0.8800	0.0376	0.8920	0.1277	0.5198	0.2057	0.8689	0.0766	0.9204	-0.1291	0.8362
CG5921	CG5921	1633783_a_at	0.4138	0.2707	0.4220	0.0960	0.2968	0.0609	-0.2039	0.6791	-0.2386	0.3051	-0.0347	0.9003	-0.1087	0.9377	-0.3506	0.4182	-0.2419	0.6052
CG8379	CG8379	1633784_at	0.5599	0.0108	0.5262	0.2483	0.3343	0.0511	-0.1004	0.8029	0.4276	0.0161	0.5280	0.0038	0.1737	0.8700	0.4200	0.3087	0.2463	0.5862
epsilonCOP	epsilonCOP	1633785_at	0.7590	0.0099	0.4276	0.2010	0.6932	0.0012	0.1173	0.7135	0.9307	0.0004	0.8134	0.0004	0.0069	0.9964	0.5862	0.1321	0.5794	0.1591
---	---	1633786_at	-0.0183	0.9173	-0.0142	0.9022	-0.1103	0.5151	-0.0104	0.9863	-0.0618	0.7124	-0.0514	0.7424	0.0284	0.9812	0.0263	0.9499	-0.0021	0.9965
---	---	1633787_at	0.1543	0.3454	0.1549	0.3567	0.4348	0.0308	0.1929	0.7749	0.2684	0.3505	0.0755	0.8130	-0.1506	0.7893	-0.0580	0.8628	0.0927	0.7381
CG15186	CG15186	1633788_s_at	-2.6876	0.0015	-3.0577	0.0057	-3.9235	0.0000	-0.8945	0.1162	-0.8129	0.0257	0.0815	0.8357	-0.0249	0.9952	-1.0196	0.2982	-0.9947	0.3343
Hand	Hand	1633789_at	0.3180	0.6562	0.1041	0.8544	0.1805	0.4266	0.0744	0.9482	-0.1273	0.7366	-0.2017	0.5170	-0.1836	0.9506	-0.4948	0.5980	-0.3112	0.7598
CG30043	CG30043	1633790_at	-0.0252	0.9936	-0.0183	0.8885	-0.2977	0.1767	-0.1701	0.9777	-1.4867	0.3435	-1.3166	0.3527	0.0609	0.9950	-1.2904	0.5825	-1.3513	0.5649
KH1	KH1	1633791_at	0.0118	0.9480	0.0552	0.7060	0.4170	0.0516	0.2313	0.5320	-0.1592	0.4260	-0.3906	0.0314	-0.3899	0.3800	-0.2099	0.3795	0.1799	0.4752
osa	eyelid	1633792_a_at	-0.3032	0.6326	-0.6046	0.3383	-0.4137	0.0881	0.3393	0.3298	0.7350	0.0039	0.3957	0.0351	0.1016	0.9816	0.3311	0.7860	0.2296	0.8577
unc-13-4A	unc-13-4A	1633793_at	-0.1892	0.3902	-0.3518	0.1309	-0.1359	0.4549	-0.0241	0.9838	0.0590	0.8704	0.0831	0.7859	-0.4111	0.6145	-0.2642	0.4482	0.1468	0.7071
smi21F	smell impaired 21	1633794_a_at	-0.5051	0.0559	-0.6462	0.0226	-0.4861	0.1357	0.2704	0.6615	0.1859	0.5602	-0.0844	0.7993	0.1348	0.9330	0.0946	0.9040	-0.0402	0.9565
CG17124	CG17124	1633795_a_at	1.1126	0.0309	1.0319	0.1816	1.5307	0.0005	0.0375	0.9860	-1.1565	0.0265	-1.1940	0.0146	-0.4502	0.8145	-1.1658	0.1392	-0.7156	0.3803
CG11425	CG11425	1633796_at	1.4172	0.0489	0.4023	0.4549	1.6786	0.0855	0.3303	0.4324	0.4075	0.0793	0.0772	0.7599	-0.4519	0.9226	-0.3174	0.8867	0.1345	0.9507
Rh2	rhodopsin	1633797_at	0.3249	0.1793	0.0046	0.9770	-0.0151	0.9473	0.0789	0.9011	0.4240	0.0446	0.3451	0.0634	0.0764	0.9401	0.0429	0.9320	-0.0335	0.9392
yps	Ypsilon Schachtel	1633798_at	-0.8107	0.2282	-0.2693	0.6492	-0.9723	0.0196	0.0128	0.9894	-0.1706	0.4598	-0.1834	0.3698	0.6881	0.7822	0.5077	0.6686	-0.1804	0.9031
CG12010	CG12010	1633799_a_at	0.1870	0.4649	0.1136	0.4861	0.0142	0.9573	-0.3426	0.4786	-0.1072	0.7227	0.2355	0.3189	-0.2506	0.7779	-0.1542	0.7307	0.0964	0.8459
CG10089	MKP-like	1633800_s_at	-0.0583	0.7551	-0.0068	0.9781	-0.0722	0.7001	-0.0766	0.9029	-0.0106	0.9733	0.0659	0.7694	0.1816	0.7230	0.1739	0.4317	-0.0077	0.9838
CG9171	CG9171	1633801_s_at	1.0116	0.1075	1.7434	0.0237	0.2045	0.2782	-0.0483	0.9705	0.4216	0.2081	0.4699	0.1167	1.5055	0.3162	1.2087	0.1251	-0.2968	0.7426
CG12650	CG12650	1633802_at	-0.4393	0.0218	-0.7838	0.0464	-0.7897	0.0039	0.0080	0.9951	0.7966	0.0082	0.7886	0.0052	-0.0976	0.9400	0.2636	0.5246	0.3612	0.3791
Pgm	Phosphogluconat	1633803_at	1.3288	0.0011	1.5063	0.0411	2.5884	0.0002	0.4964	0.4643	-1.1082	0.0092	-1.6045	0.0011	-0.4048	0.7726	-0.7161	0.2168	-0.3113	0.6296

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
RhoGDI	RhoGDI	1633804_at	0.0596	0.7528	-0.0326	0.9233	-0.4431	0.0096	0.0541	0.9220	0.3936	0.0262	0.3395	0.0305	0.3607	0.6749	0.3749	0.2764	0.0142	0.9811
sax	saxophone	1633805_a_at	-0.1269	0.5344	-0.0421	0.8314	-0.1048	0.6478	-0.1132	0.7969	-0.1532	0.4035	-0.0400	0.8468	-0.0193	0.9898	-0.0649	0.8942	-0.0456	0.9197
---	---	1633806_at	0.1328	0.5660	0.4023	0.0683	0.5537	0.0049	0.2798	0.6122	0.2008	0.4888	-0.0790	0.7993	0.0631	0.9589	0.3686	0.2644	0.3055	0.3828
l(3)neo38	lethal (3) neo38	1633807_at	0.7596	0.1139	0.2679	0.3994	-0.0278	0.9378	-0.1108	0.9228	0.2325	0.5276	0.3433	0.2704	0.3228	0.8374	-0.2362	0.7599	-0.5590	0.3953
---	---	1633808_at	0.3398	0.1047	0.0494	0.5977	0.1621	0.3762	0.0844	0.8707	0.0437	0.8527	-0.0407	0.8466	0.1205	0.8767	-0.0397	0.9350	-0.1602	0.6228
CG13046	CG13046	1633809_at	0.0733	0.7423	-0.0942	0.5256	0.1959	0.5021	0.1974	0.7589	0.0578	0.8747	-0.1396	0.6183	0.0291	0.9852	-0.0965	0.8397	-0.1256	0.7604
CG8793	CG8793	1633810_at	-0.0598	0.7460	0.2781	0.4364	0.5626	0.0039	0.1679	0.7556	-0.0276	0.9328	-0.1955	0.3604	-0.1122	0.9056	0.2969	0.3733	0.4091	0.2449
CG13962	CG13962	1633811_at	0.0380	0.8381	0.0397	0.8782	-0.0865	0.6743	0.1262	0.8578	0.1895	0.4747	0.0633	0.8265	0.0013	0.9994	0.0464	0.8781	0.0451	0.8739
CG9080	CG9080	1633812_at	-1.3703	0.0069	-2.1988	0.0063	-1.7343	0.0006	-0.2523	0.6763	2.1436	0.0002	2.3959	0.0001	-0.9427	0.4196	1.1643	0.0757	2.1070	0.0311
---	---	1633813_at	0.1372	0.3734	0.0023	0.9918	0.1012	0.6415	0.0142	0.9838	0.0448	0.8282	0.0306	0.8738	0.0532	0.9616	-0.0167	0.9751	-0.0699	0.8637
CG3842	CG3842	1633814_at	0.8613	0.0418	0.3178	0.3860	0.4268	0.0677	-0.0138	0.9865	-0.0398	0.8811	-0.0259	0.9150	-0.1066	0.9549	-0.3921	0.4868	-0.2856	0.6311
CG31191	CG31191	1633815_at	-0.1159	0.5107	-0.3567	0.0768	-0.0506	0.7577	0.1211	0.7278	0.2631	0.0922	0.1420	0.3173	-0.1226	0.8235	-0.0788	0.7795	0.0439	0.8891
CG17181	CG17181	1633816_at	-0.5845	0.0335	0.0415	0.6806	-0.0579	0.8223	-0.0623	0.9538	-0.5573	0.0621	-0.4950	0.0635	0.0779	0.9521	0.0070	0.9935	-0.0709	0.8947
DJ-1alpha	DJ-1alpha	1633817_at	0.4189	0.0745	0.2007	0.4199	0.3803	0.0638	-0.0327	0.9712	0.0423	0.8920	0.0749	0.7668	-0.0408	0.9778	-0.1461	0.7124	-0.1053	0.8046
CG7368	CG7368	1633818_at	-0.1022	0.6649	-0.1438	0.4335	-0.1260	0.4166	0.3291	0.4166	0.1690	0.4722	-0.1601	0.4501	0.0955	0.9092	-0.0122	0.9829	-0.1077	0.7527
---	---	1633819_at	-0.0649	0.6713	-0.0029	0.9816	0.0381	0.8147	0.0478	0.9370	0.1316	0.4739	0.0837	0.6428	0.0503	0.9309	0.0595	0.8053	0.0092	0.9756
Fad2	Fad2	1633820_at	1.2877	0.4702	-1.4539	0.6650	2.2141	0.0445	1.9019	0.4034	0.4034	0.7961	-1.4985	0.1863	-1.9612	0.8270	-2.4706	0.5259	-0.5094	0.9215
B52	Serine/arginine ric	1633821_at	0.5177	0.0155	0.2159	0.7619	0.6738	0.0461	0.2055	0.8058	0.8358	0.0204	0.6303	0.0406	-0.1112	0.9611	0.6804	0.2723	0.7917	0.2336
CG7519	CG7519	1633822_at	0.5097	0.0461	0.4042	0.1681	0.4491	0.0485	0.1307	0.8115	0.0824	0.7495	-0.0483	0.8478	0.1722	0.8461	0.0282	0.9628	-0.1440	0.7309
CG5781	CG5781	1633823_at	-0.0025	0.9896	0.0175	0.8802	0.1083	0.6701	0.0485	0.9538	-0.0207	0.9518	-0.0691	0.7874	-0.1796	0.7440	-0.0699	0.8168	0.1098	0.6628
CG32710	CG32710	1633824_at	0.3279	0.0944	0.3575	0.1917	0.0530	0.8022	-0.1362	0.7161	-0.0437	0.8432	0.0925	0.5880	0.3137	0.6749	0.0766	0.8551	-0.2371	0.4544
CG14221	CG14221	1633825_at	0.0320	0.8981	-0.0591	0.7278	0.0730	0.6892	0.0587	0.9491	0.0457	0.8964	-0.0131	0.9679	-0.0662	0.9142	-0.0368	0.9111	0.0293	0.9211
Iris	Iris	1633826_at	2.8781	0.0016	1.4526	0.0652	3.0677	0.0002	0.8961	0.3628	0.8341	0.1357	-0.0620	0.9279	-0.7241	0.6927	-0.6320	0.3963	0.0921	0.9306
CG14327	CG14327	1633827_at	-0.0905	0.6113	-0.1510	0.4898	-0.0316	0.8566	0.1575	0.7451	0.1716	0.4366	0.0141	0.9586	-0.0862	0.8650	0.0185	0.9550	0.1047	0.6311
---	---	1633828_at	-0.1284	0.4315	-0.1807	0.1998	0.1296	0.4444	0.0839	0.8636	0.0146	0.9549	-0.0693	0.7048	-0.0994	0.8814	-0.1032	0.7259	-0.0037	0.9933
---	---	1633829_at	-0.1125	0.6176	-0.0509	0.6350	-0.0857	0.7185	0.1549	0.7409	-0.0412	0.8811	-0.1961	0.2947	0.0214	0.9875	0.0793	0.8519	0.0579	0.8918
CG10195	CG10195	1633830_at	0.0291	0.9122	-0.1141	0.6021	0.0884	0.5974	-0.3479	0.4162	-0.3455	0.1437	0.0024	0.9935	-0.4167	0.6034	-0.3546	0.2960	0.0621	0.8969
CG33658	CG33658	1633831_at	-0.0003	0.9988	0.0325	0.7438	-0.1141	0.5098	-0.0634	0.9116	-0.0074	0.9790	0.0560	0.7809	0.0498	0.9503	0.0052	0.9925	-0.0446	0.8918
CG32143 /// ome	CG32143 /// omeg	1633832_a_at	-0.1531	0.5592	-0.4178	0.2352	-0.1904	0.5637	0.2382	0.7982	0.4928	0.1896	0.2546	0.4794	-0.2349	0.8062	0.0304	0.9631	0.2653	0.5270
CG18213	CG18213	1633833_at	-0.6933	0.0990	-1.2137	0.0165	-1.4415	0.0086	-0.0697	0.9642	0.9289	0.0316	0.9986	0.0148	-0.0667	0.9848	0.1780	0.8725	0.2447	0.7940
CG9701	CG9701	1633834_at	2.7757	0.0030	3.2640	0.0080	4.6421	0.0000	1.7554	0.0290	0.1840	0.7168	-1.5715	0.0025	0.0123	0.9967	0.3417	0.6885	0.3295	0.6986
CG2694	CG2694	1633835_s_at	-0.0205	0.9581	-0.1115	0.6550	-0.3909	0.0251	-0.2709	0.3793	0.3363	0.0581	0.6072	0.0028	0.0697	0.9515	0.2989	0.3769	0.2292	0.5271
Fkbp13	Fkbp13	1633836_a_at	2.2505	0.0006	1.1905	0.0049	1.8664	0.0000	0.6250	0.1162	1.1187	0.0014	0.4937	0.0289	0.0686	0.8956	0.0627	0.8000	-0.0059	0.9849
Obp50c	Odorant-binding p	1633837_at	0.1084	0.5756	0.0873	0.6740	0.1644	0.3636	0.0604	0.9218	0.0465	0.8470	-0.0139	0.9532	0.2192	0.7644	-0.0166	0.9755	-0.2358	0.4617
---	---	1633838_at	0.0442	0.8053	0.0719	0.5364	-0.0947	0.5455	-0.1395	0.6988	-0.0494	0.8094	0.0901	0.5863	0.0036	0.9970	0.0239	0.9501	0.0203	0.9512
CG14830	CG14830	1633839_at	-0.5029	0.0421	0.1657	0.7038	-0.4006	0.1058	-0.1553	0.7141	-0.4682	0.0237	-0.3129	0.0726	0.3908	0.7472	0.2395	0.6748	-0.1513	0.8124
CG8026	CG8026	1633840_a_at	-0.7029	0.2059	-0.1000	0.7954	0.1473	0.6882	0.0653	0.9745	0.0799	0.9083	0.0146	0.9812	-0.2856	0.8513	0.6911	0.2533	0.9767	0.1479
yellow-e2	yellow-e2	1633841_at	0.1683	0.4578	0.1363	0.2806	0.3835	0.0440	0.0209	0.9745	-0.0359	0.8656	-0.0568	0.7441	-0.1130	0.8513	0.0014	0.9989	0.1144	0.6761
CG3546	CG3546	1633842_at	0.3375	0.0722	0.1287	0.6524	0.2023	0.3223	0.1317	0.8546	0.0976	0.7529	-0.0342	0.9142	-0.0316	0.9742	-0.0314	0.9353	0.0002	0.9997
CG31672	CG31672	1633843_at	-0.0019	0.9977	-0.5310	0.4924	-0.8498	0.0127	-0.3462	0.5639	0.6307	0.0463	0.9769	0.0043	0.1608	0.9651	0.3495	0.7703	0.1887	0.8889
CG31469	CG31469	1633844_at	-0.5704	0.3164	-0.0065	0.9927	0.1875	0.4787	-0.2818	0.6822	-0.4775	0.1347	-0.1956	0.5315	-0.3945	0.8331	-0.0097	0.9956	0.3847	0.6476
CG15896	CG15896	1633845_at	-0.1572	0.5454	0.1349	0.8453	0.0497	0.9131	-0.0344	0.9500	-0.1536	0.3193	-0.1192	0.4008	0.0384	0.9922	0.0302	0.9868	-0.0082	0.9959
CG15523	CG15523	1633846_at	0.9855	0.0045	0.5451	0.1233	0.6025	0.0141	0.0251	0.9756	1.0764	0.0008	1.0513	0.0005	0.0170	0.9935	0.5892	0.1984	0.5722	0.2437
CG11284	CG11284	1633847_at	-0.0562	0.7518	-0.0288	0.9099	0.0653	0.6889	0.0345	0.9562	-0.0023	0.9926	-0.0368	0.8521	-0.0883	0.8714	0.1472	0.5106	0.2355	0.2918
CG40133	CG40133	1633848_at	-0.0205	0.9116	0.0959	0.5173	-0.0166	0.9239	0.0404	0.9530	0.0816	0.7079	0.0412	0.8507	0.0879	0.8446	0.1636	0.3669	0.0758	0.7154
CG31559	CG31559	1633849_at	-0.0601	0.8097	-0.1651	0.3280	-0.0177	0.9375	0.0747	0.9380	-0.1202	0.7132	-0.1948	0.4674	-0.1423	0.8284	-0.2011	0.4729	-0.0588	0.8745
CG18156	CG18156	1633850_at	0.2125	0.5466	-0.2288	0.1005	-0.1638	0.3507	0.0347	0.9805	0.3736	0.2846	0.3389	0.2791	-0.1272	0.8395	-0.1184	0.6820	0.0088	0.9841
CG31231	CG31231	1633851_at	0.1003	0.5759	0.0284	0.7796	0.1777	0.4952	-0.0749	0.9247	-0.0891	0.7530	-0.0142	0.9621	0.0001	0.9999	-0.1048	0.7868	-0.1048	0.7749
---	---	1633852_at	-0.7940	0.3194	-0.1596	0.4103	-0.3757	0.1221	-0.2382	0.9138	-0.9800	0.1558	-0.7419	0.2323	-0.1508	0.8480	-0.3079	0.3346	-0.1571	0.6578
---	---	1633853_at	0.2677	0.1043	-0.0110	0.9176	-0.0567	0.7412	0.1901	0.5735	0.2756	0.1099	0.0855	0.6273	0.1310	0.8305	-0.0127	0.9782	-0.1436	0.6021
DNApol-alpha50	DNA polymerase	1633854_at	0.2262	0.5487	-0.2454	0.4954	0.0265	0.9539	-0.3527	0.6869	-0.2640	0.5460	0.0887	0.8507	-0.6707	0.6483	-0.6600	0.2491	0.0107	0.9921
lid	little imaginal disc	1633855_s_at	0.1251	0.7434	-0.4956	0.1559	-0.4017	0.0420	0.5174	0.2330	1.0184	0.0028	0.5011	0.0365	0.4338	0.7230	0.5522	0.2787	0.1184	0.8648
CG11858	CG11858	1633856_at	0.2802	0.0845	0.4085	0.1014	0.6468	0.0159	-0.0099	0.9883	-0.1703	0.2793	-0.1605	0.2544	-0.0996	0.9077	0.1662	0.6111	0.2657	0.3921
CG13659	CG13659	1633857_at	2.4452	0.0007	1.1530	0.0409	2.2172	0.0001	1.3600	0.0265	2.1221	0.0004	0.7621	0.0188	0.2507	0.7720	0.7520	0.0637	0.5013	0.1990
CG4286	CG4286	1633858_at	0.0694	0.7899	0.3602	0.1280	0.0932	0.6133	-0.1070	0.8793	-0.1625	0.5254	-0.0555	0.8414	0.0422	0.9717	0.0315	0.9477	-0.0107	0.9835
---	---	1633859_at	0.2446	0.1509	0.0936	0.5728	0.2846	0.1276	0.1955	0.5255	0.1264	0.4509	-0.0691	0.6822	0.1680	0.7196	0.1054	0.6114	-0.0626	0.7864
CG12078	CG12078	1633860_at	0.1201	0.5423	-0.1807	0.5698	0.0755	0.6442	0.1741	0.6656	0.1424	0.4789	-0.0317	0.8906	-0.1582	0.8858	-0.2004	0.6601	-0.0422	0.9428
CG8145	CG8145	1633861_at	-0.8288	0.0089	-0.8804	0.0235	-0.7455	0.0078	-0.0895	0.8244	0.2404	0.1224	0.3299	0.0263	-0.2725	0.7726	-0.0226	0.9739	0.2499	0.5612
CG12934	CG12934	1633862_at	0.3562	0.0279	0.0348	0.7966	0.1729	0.4311	0.2209	0.5633	0.3070	0.1165	0.0861	0.6746	-0.0903	0.9246	-0.1983	0.5531	-0.1080	0.7749
---	---	1633863_at	-0.0196	0.9288	-0.0632	0.5391	0.0498	0.8121	0.1080	0.8550	-0.0153	0.9623	-0.1233	0.5593	-0.0644	0.9296	-0.0242	0.9499	0.0403	0.9023
CG14608	CG14608	1633864_at	-0.1929	0.1747	-0.0223	0.8266	-0.0047	0.9826	-0.0369	0.9471	-0.1413	0.3792	-0.1044	0.4870	-0.1432	0.7686	-0.0100	0.9784	0.1332	0.5416
CG16734	CG16734	1633865_at	0.0058	0.9895	-0.0791	0.6164	-0.1129	0.6795	-0.4637	0.3597	0.0130	0.9771	0.4767	0.0677	-0.4574	0.7305	-0.1230	0.8796	0.3344	0.5807
Su(z)2	Aristopedoid	1633866_at	-0.3681	0.2061	-0.5481	0.0409	-0.6730	0.0214	-0.0337	0.9755	0.2502	0.3604	0.2839	0.2371	0.0320	0.9848	0.0371	0.9526	0.0050	0.9944
CG31923	CG31923	1633867_at	0.1688	0.2869	0.0249	0.8212	0.0031	0.9903	-0.0589	0.9154	-0.0361	0.8758	0.0227	0.9146	0.1051	0.8576	-0.0857	0.7671	-0.1908	0.4361
---	---	1633868_at	0.2632	0.1966	0.0000	1.0000	0.2048	0.3162	0.1061	0.9030	0.1301	0.6880	0.0239	0.9468	0.0048	0.9963	-0.0778	0.7730	-0.0825	0.7456
Dfd	deformed	1633869_at	0.2043	0.3850	0.1398	0.2415	0.3745	0.0604	0.1332	0.8366	-0.0464	0.8878	-0.1796	0.4321	0.1275	0.7753	0.0119	0.9703	-0.1156	0.5736
CG9436	CG9436	1633870_at	0.2938	0.3101	-0.5228	0.3636	0.6427	0.0062	0.5380	0.0978	0.7369	0.0031	0.1989	0.2475	-0.4175	0.7485	0.0409	0.9646	0.4584	0.4149
CG34377	CG13859	1633871_at	-0.1170	0.5246	0.1106	0.4272	0.0369	0.8553	-0.0676	0.9300	-0.4032	0.0758	-0.3357	0.0952	0.0687	0.9296	-0.0159	0.9696	-0.0846	0.7756
CG15673	CG15673	1633872_at	0.7141	0.2260	1.5044	0.0329	1.0738	0.0756	0.1684	0.9218	0.4017	0.4562	0.2333	0.6642	0.6090	0.7768	1.0789	0.2286	0.4699	0.6389

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1633873_at	-0.0363	0.8622	-0.0569	0.5719	0.1900	0.4513	0.1500	0.6218	0.0204	0.9228	-0.1296	0.3506	-0.1770	0.7896	-0.0544	0.8986	0.1225	0.6982
CG12460	CG12460	1633874_at	0.0469	0.8534	-0.0189	0.8585	-0.1455	0.3161	0.0268	0.9704	0.0767	0.7209	0.0498	0.8102	0.0701	0.9342	-0.0205	0.9639	-0.0905	0.7782
---	---	1633875_at	0.0736	0.6149	0.1453	0.3843	0.0767	0.6486	-0.1065	0.8578	0.0234	0.9393	0.1298	0.5324	0.0134	0.9862	-0.0152	0.9585	-0.0287	0.9057
CG7110	CG7110	1633876_at	-1.7995	0.0012	-0.4463	0.0673	-1.7562	0.0011	-1.2730	0.1163	-2.5774	0.0008	-1.3044	0.0096	0.1972	0.7677	-1.3227	0.0052	-1.5200	0.0039
CG30358	CG30358	1633877_at	-0.0052	0.9802	0.0131	0.9095	0.0491	0.7782	-0.0428	0.9314	-0.0493	0.7824	-0.0065	0.9716	0.0070	0.9952	0.0476	0.9171	0.0406	0.9206
MESR4	Misexpression suj	1633878_at	-0.1480	0.6845	0.5569	0.2164	0.4326	0.0146	-0.0938	0.9314	-0.2227	0.5152	-0.1289	0.7057	0.0805	0.9623	0.5151	0.2584	0.4346	0.3711
prd	paired	1633879_a_at	0.1394	0.4840	-0.0543	0.6697	-0.0775	0.7452	-0.0387	0.9705	0.1061	0.7372	0.1447	0.5895	0.0048	0.9959	-0.0339	0.9178	-0.0387	0.8946
CG8533	CG8533	1633880_s_at	-0.3515	0.1201	-0.4029	0.0703	-0.6313	0.0044	-0.0999	0.7929	0.0539	0.7724	0.1538	0.2710	-0.0760	0.9420	-0.0392	0.9402	0.0368	0.9347
CG1776	MLCK-like	1633881_at	-0.3952	0.1687	-0.1178	0.8461	-1.2607	0.0006	-0.2835	0.3863	0.1355	0.4848	0.4191	0.0196	0.7752	0.5790	0.4195	0.5216	-0.3557	0.6005
Rgk1	Rgk1	1633882_at	-0.0966	0.6232	0.5840	0.1180	-0.0054	0.9881	-0.2440	0.6333	-0.2520	0.3205	-0.0080	0.9799	0.3083	0.7215	0.4141	0.2373	0.1058	0.8140
CG9609	CG9609	1633883_at	-0.2473	0.2084	0.6044	0.1534	0.7576	0.0019	-0.1821	0.8364	-0.4701	0.1556	-0.2881	0.3449	-0.2281	0.6898	0.3196	0.1584	0.5477	0.0559
Rtf1	CG10955	1633884_at	-0.3665	0.2457	0.7609	0.1503	0.8810	0.0004	0.0158	0.9934	-0.6683	0.0752	-0.6842	0.0456	-0.0149	0.9927	0.5205	0.1498	0.5354	0.1719
CG6950	CG6950	1633885_s_at	0.1200	0.5290	0.5662	0.0351	0.9328	0.0027	-0.0968	0.8155	-0.8682	0.0008	-0.7714	0.0008	-0.4291	0.6272	-0.2754	0.4589	0.1537	0.7128
CG7949	CG7949	1633886_at	0.3775	0.0725	0.1706	0.1893	0.2938	0.0969	0.0478	0.9361	0.0768	0.7030	0.0291	0.8893	-0.1071	0.8940	-0.0994	0.7912	0.0076	0.9874
CG13857	CG13857	1633887_at	0.3588	0.2462	0.1356	0.4953	0.1426	0.5677	0.0775	0.9491	0.1011	0.8094	0.0235	0.9568	0.0989	0.8425	-0.1141	0.6009	-0.2130	0.3081
kirre	dumbfounded	1633888_s_at	-0.1721	0.5523	-0.1891	0.6020	-0.7418	0.0190	-0.0618	0.9603	0.3313	0.3218	0.3931	0.1817	0.1540	0.9112	-0.0047	0.9973	-0.1587	0.7849
---	---	1633889_at	0.1757	0.3868	-0.0164	0.8802	0.1788	0.3227	-0.1496	0.7975	0.0088	0.9804	0.1584	0.4826	-0.2292	0.7633	-0.0646	0.8895	0.1646	0.6318
CG34384	CG14462	1633890_at	-0.0112	0.9493	0.0010	0.9945	0.0346	0.8599	-0.0139	0.9826	0.0330	0.8675	0.0470	0.7776	-0.0547	0.9238	0.0234	0.9395	0.0781	0.7149
CG15431	CG15431	1633891_at	0.1252	0.5740	-0.1541	0.4252	0.3177	0.0646	0.1871	0.7558	-0.0539	0.8759	-0.2410	0.3066	-0.0699	0.9168	-0.0918	0.7339	-0.0218	0.9464
---	---	1633892_at	-0.0877	0.5974	0.1102	0.4461	0.1727	0.3219	-0.2191	0.4568	-0.1713	0.2872	0.0478	0.7888	-0.1475	0.8193	0.0114	0.9819	0.1589	0.5720
CG31217 /// DvirCG31217	predicted gene W	1633893_at	-0.4684	0.0246	-1.3017	0.0251	-1.2994	0.0003	-0.1599	0.5735	-0.2971	0.0470	-0.1372	0.2991	-0.1477	0.9149	-1.0773	0.0476	-0.9296	0.0893
CG10543	CG10543	1633894_a_at	-0.3597	0.4694	0.2970	0.3944	0.5462	0.0126	0.0784	0.9393	-0.5433	0.0728	-0.6216	0.0282	0.0902	0.9691	0.5615	0.3454	0.4712	0.4557
H2.0	Homeodomain pr	1633895_at	0.3692	0.1082	0.3976	0.1228	0.3664	0.0294	-0.2286	0.5217	-0.2023	0.2831	0.0263	0.9078	0.0631	0.9441	-0.0148	0.9747	-0.0779	0.8178
---	---	1633896_at	0.0827	0.5460	0.0167	0.8703	0.0640	0.7482	-0.1065	0.8115	-0.0210	0.9330	0.0855	0.6375	0.0623	0.8960	0.0236	0.9341	-0.0387	0.8707
Fer3	ntwist	1633897_at	0.3120	0.0915	0.2899	0.3973	0.2222	0.3426	0.0807	0.9182	-0.0710	0.8123	-0.1517	0.5176	0.0865	0.9342	-0.0840	0.8569	-0.1705	0.6392
---	---	1633898_at	0.0683	0.6957	0.0790	0.6396	0.1437	0.4759	-0.0715	0.9228	-0.0530	0.8548	0.0186	0.9476	-0.0621	0.9309	0.0749	0.7988	0.1370	0.5789
---	---	1633899_at	0.0409	0.8941	-0.1063	0.6015	-0.1205	0.5107	0.1144	0.8424	0.2757	0.1950	0.1613	0.4202	0.0140	0.9914	0.0033	0.9961	-0.0106	0.9835
CG14044	CG14044	1633900_at	-0.0599	0.7230	-0.0985	0.4887	-0.0395	0.8912	0.1640	0.7678	0.0677	0.8174	-0.0963	0.6981	0.0042	0.9956	-0.0022	0.9959	-0.0064	0.9837
---	---	1633901_at	0.0402	0.8803	-0.1303	0.3985	-0.3508	0.0344	-0.1380	0.8221	0.1360	0.6044	0.2740	0.1997	0.1162	0.9101	-0.0364	0.9507	-0.1525	0.7134
---	---	1633902_at	0.0785	0.7063	0.0027	0.9828	0.1304	0.4070	0.1695	0.5419	-0.0137	0.9500	-0.1831	0.1552	0.2007	0.7770	0.0486	0.9188	-0.1521	0.6457
---	---	1633903_s_at	0.0793	0.7546	0.1688	0.3779	0.1472	0.5035	-0.0698	0.9488	-0.1488	0.6663	-0.0790	0.8185	0.0199	0.9841	0.0325	0.9275	0.0126	0.9709
CG5467	CG5467	1633904_at	-0.4663	0.4520	-0.3283	0.6235	-0.4676	0.0181	-0.0945	0.9228	0.0033	0.9939	0.0979	0.7618	0.0581	0.9898	0.0507	0.9758	-0.0074	0.9965
RpL28	Ribosomal protein	1633905_at	1.5845	0.0015	1.0952	0.0996	0.9458	0.0104	-0.0709	0.9488	0.5449	0.0793	0.6159	0.0329	0.0483	0.9848	0.2654	0.6768	0.2171	0.7434
CG6485	CG6485	1633906_at	0.0649	0.8171	0.0355	0.7371	-0.0561	0.8537	-0.2065	0.6869	-0.1356	0.6038	0.0710	0.7902	0.0213	0.9848	-0.1430	0.5990	-0.1644	0.5415
---	---	1633907_at	0.0699	0.6772	0.0516	0.6391	-0.0105	0.9569	-0.1172	0.7981	0.0587	0.7965	0.1759	0.2955	0.0848	0.8814	0.1275	0.5800	0.0426	0.8869
Pdf	Pigment dispersin	1633908_at	-0.1310	0.5706	-0.0529	0.6522	0.1659	0.3276	0.1584	0.6338	-0.2110	0.1948	-0.3694	0.0197	-0.2279	0.7644	-0.0670	0.8850	0.1609	0.6403
---	---	1633909_at	0.0764	0.7050	-0.0045	0.9913	-0.0153	0.9514	-0.0892	0.8810	0.1316	0.5474	0.2209	0.2250	0.0266	0.9862	0.0611	0.9091	0.0345	0.9435
---	---	1633910_at	0.2409	0.3470	0.1884	0.3130	0.1592	0.4336	-0.1068	0.8117	-0.1969	0.2697	-0.0901	0.6167	0.1018	0.8940	0.0033	0.9960	-0.0984	0.7691
CG31550	CG31550	1633911_a_at	-0.7087	0.0214	-0.3323	0.6111	-0.3270	0.1009	-0.1290	0.8544	-0.3177	0.2081	-0.1887	0.4269	-0.2564	0.8775	0.0549	0.9590	0.3113	0.6567
CG2113	CG2113	1633912_at	0.2069	0.2073	0.0000	1.0000	-0.0523	0.8034	0.1017	0.8532	0.1212	0.5726	0.0195	0.9369	0.1158	0.8128	0.0640	0.8055	-0.0518	0.8412
a10	Pherokine 1	1633913_at	-0.0833	0.7484	0.1224	0.4946	0.4089	0.1115	-0.0294	0.9803	-0.3314	0.2463	-0.3019	0.2376	-0.1617	0.8692	-0.1873	0.6626	-0.0257	0.9646
Gal	beta-galactosidas	1633914_at	-2.0762	0.0010	-0.9720	0.0351	-1.8067	0.0005	-0.8948	0.1661	-1.7284	0.0017	-0.8336	0.0257	-0.0504	0.9689	-0.8143	0.0391	-0.7639	0.0607
RnpS1	RRM-domain prot	1633915_at	-0.0068	0.9802	0.4189	0.0237	0.4526	0.0266	0.2206	0.5455	0.2396	0.2036	0.0190	0.9354	0.0448	0.9689	0.5737	0.0721	0.5289	0.1156
CG18605	CG18605	1633916_at	0.1190	0.4954	0.2358	0.1843	0.1896	0.3226	0.1486	0.7313	-0.1227	0.5550	-0.2713	0.1150	0.0611	0.9520	-0.0034	0.9961	-0.0645	0.8755
---	---	1633917_at	0.2144	0.3403	-0.1950	0.1426	-0.0671	0.7132	0.0127	0.9883	0.3075	0.1286	0.2948	0.1036	-0.0022	0.9978	-0.1346	0.4699	-0.1324	0.4857
CG7115	CG7115	1633918_s_at	-0.8293	0.0041	-0.9513	0.0979	-1.3825	0.0002	-0.1612	0.7656	0.2759	0.2309	0.4371	0.0404	0.2584	0.8270	0.1873	0.7443	-0.0710	0.9165
l(3)mbn	lethal(3)malignant	1633919_s_at	0.1275	0.5620	-0.0080	0.9507	-0.0815	0.6792	0.1084	0.8053	0.0892	0.6513	-0.0192	0.9286	0.0721	0.9193	-0.0606	0.8551	-0.1327	0.6121
CG5715	CG5715	1633920_at	-0.0621	0.7723	-0.0769	0.4784	-0.1978	0.2348	-0.0498	0.9353	0.0234	0.9267	0.0731	0.7017	0.0392	0.9499	-0.0509	0.8382	-0.0901	0.6536
CG30050	CG30050	1633921_at	0.1357	0.4256	-0.0675	0.5968	0.1604	0.4139	-0.0140	0.9852	0.0834	0.6660	0.0974	0.5624	-0.2955	0.5754	-0.1559	0.5266	0.1396	0.5814
---	---	1633922_at	-0.0395	0.7895	0.0756	0.5841	0.1458	0.4997	0.0840	0.8816	0.0250	0.9258	-0.0590	0.7830	0.0031	0.9976	0.0097	0.9816	0.0066	0.9852
CG1749	CG1749	1633923_at	1.6441	0.0018	1.3664	0.0138	1.3997	0.0001	0.3281	0.3045	0.8835	0.0013	0.5554	0.0060	0.2551	0.7770	0.6231	0.1162	0.3681	0.3601
---	---	1633924_s_at	0.0861	0.6892	0.0961	0.4520	0.2661	0.1470	0.1289	0.7711	0.0104	0.9713	-0.1185	0.5119	0.0794	0.8700	0.0284	0.9265	-0.0510	0.8354
---	---	1633925_at	0.0191	0.9297	0.0242	0.8981	0.1899	0.3686	0.1421	0.6825	0.0672	0.7214	-0.0748	0.6550	0.0182	0.9898	0.0014	0.9991	-0.0168	0.9704
CG16837	CG16837	1633926_at	0.2502	0.2684	0.0305	0.8571	0.2237	0.2378	0.0043	0.9956	-0.0039	0.9897	-0.0082	0.9734	-0.1510	0.7953	-0.1357	0.6019	0.0153	0.9665
beat-IIIc	beat-IIIc	1633927_a_at	-2.5995	0.0005	-2.8496	0.0045	-3.1560	0.0002	0.0273	0.9921	0.4403	0.4489	0.4131	0.4318	-0.1601	0.8202	-0.0998	0.7802	0.0603	0.8786
CG15721	CG15721	1633928_at	0.1037	0.6169	0.1232	0.4639	0.0308	0.9098	0.0411	0.9580	0.5487	0.0159	0.5076	0.0138	0.1756	0.8694	0.2998	0.4954	0.1243	0.8161
---	---	1633929_at	0.2213	0.2679	0.0848	0.6033	-0.0685	0.7901	-0.0683	0.9491	-0.0375	0.9289	0.0308	0.9338	0.0472	0.9487	-0.0874	0.7317	-0.1346	0.5630
CG1394	CG1394	1633930_at	0.1035	0.4869	0.0732	0.5039	0.1147	0.4826	0.0984	0.8543	-0.0033	0.9903	-0.1017	0.5999	0.0671	0.9117	-0.0176	0.9605	-0.0847	0.7299
VhaSFD	Vacuolar H[+]-ATI	1633931_a_at	-0.6223	0.0107	-0.5666	0.0639	-1.0654	0.0002	-0.2728	0.3193	-0.3694	0.0283	-0.0966	0.5354	0.1245	0.8692	-0.2580	0.3884	-0.3826	0.2231
---	---	1633932_x_at	0.0804	0.6982	-0.0035	0.9810	0.1418	0.3730	0.0073	0.9934	0.1334	0.4535	0.1261	0.4324	-0.0999	0.8815	0.0605	0.8678	0.1604	0.5556
CG31423	CG31423	1633933_at	-0.1559	0.4156	-0.1411	0.3605	-0.0402	0.8113	-0.0095	0.9904	0.0630	0.7654	0.0726	0.6950	-0.1329	0.7922	-0.1000	0.6717	0.0329	0.9109
CG8417 /// DyakCG8417	CG8417	1633934_at	-0.2597	0.3805	0.6006	0.0270	0.9662	0.0057	-0.0835	0.9136	-1.2933	0.0008	-1.2098	0.0006	-0.4059	0.6749	-0.2785	0.4932	0.1274	0.7872
skf	skiff	1633935_at	0.7887	0.1416	1.5030	0.0349	0.8834	0.0028	0.2808	0.4365	0.0025	0.9930	-0.2783	0.1144	0.8424	0.6557	0.7953	0.2847	-0.0471	0.9675
sca	scabrous	1633936_a_at	0.0885	0.5411	0.1258	0.4884	0.0926	0.5883	-0.1414	0.6673	-0.1824	0.2458	-0.0410	0.8174	-0.0100	0.9943	-0.1218	0.7501	-0.1117	0.7674
---	---	1633937_at	0.0908	0.5463	0.0003	1.0000	0.0498	0.7480	0.1069	0.7492	-0.0185	0.9282	-0.1254	0.3481	-0.0209	0.9821	-0.1394	0.5367	-0.1186	0.6121
CG14914	CG14914	1633938_at	0.0870	0.6227	0.0543	0.6058	-0.0603	0.8585	-0.0096	0.9940	-0.0413	0.9092	-0.0317	0.9210	0.2185	0.7628	0.1223	0.7248	-0.0961	0.7907
CG13563	CG13563	1633939_at	0.1216	0.5470	-0.0571	0.6183	0.0358	0.8671	0.1937	0.7349	0.3248	0.1990	0.1311	0.6096	-0.0526	0.9340	-0.0295	0.9269	0.0231	0.9347
CG9393	CG9393	1633940_at	-0.1869	0.2848	0.1982	0.3828	0.5452	0.0234	-0.0102	0.9883	-0.2608	0.1044	-0.2506	0.0830	-0.2068	0.8049	0.1717	0.6524	0.3785	0.2949
CG13569	CG13569	1633941_a_at	0.0445	0.7536	-0.1864	0.3914	0.0043	0.9860	0.2074	0.6122	0.3159	0.1224	0.1085	0.5992	0.0699	0.9333	0.1241	0.6762	0.0543	0.8814

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1633942_at	-0.0431	0.8805	-0.0809	0.5001	0.2127	0.1629	0.2442	0.6506	0.2663	0.3114	0.0221	0.9466	-0.1048	0.8870	0.0835	0.8165	0.1884	0.5141
---	---	1633943_at	0.1322	0.5281	-0.1943	0.1698	-0.2638	0.2358	0.1130	0.8738	0.2992	0.2265	0.1862	0.4222	0.1011	0.9174	-0.0686	0.8895	-0.1697	0.6411
CG17362	CG17362	1633944_at	0.0190	0.9148	-0.0198	0.8495	0.1595	0.4549	-0.1227	0.8076	-0.2810	0.1600	-0.1583	0.3972	-0.0749	0.9174	-0.1197	0.6634	-0.0448	0.8959
CG40382	CG40382	1633945_at	-0.0147	0.9311	-0.1857	0.2773	-0.0088	0.9670	0.2109	0.5008	0.3915	0.0259	0.1806	0.2233	-0.0157	0.9896	0.0776	0.8123	0.0933	0.7484
CG31955	CG31955	1633946_at	0.4668	0.2843	0.2258	0.1022	0.4351	0.0461	0.3866	0.5149	1.1454	0.0039	0.7588	0.0148	0.1070	0.9489	0.7970	0.1155	0.6900	0.1911
Cyp6a18	Cyp6a18	1633947_at	-0.5495	0.5789	-0.0257	0.8158	-0.1089	0.5440	0.2874	0.9266	-0.4543	0.6701	-0.7416	0.3944	0.1004	0.9238	-0.1687	0.6537	-0.2691	0.4498
Egfr	EGF receptor	1633948_a_at	0.8579	0.0637	0.9896	0.0990	1.9257	0.0004	0.3305	0.4694	-0.4676	0.0615	-0.7981	0.0040	-0.5181	0.7810	-0.3157	0.7371	0.2024	0.8444
CG15152	CG15152	1633949_at	0.1675	0.8958	0.0076	0.9523	-0.0798	0.7348	-0.2993	0.8640	-0.6784	0.2771	-0.3790	0.5316	-0.0453	0.9922	-0.5523	0.6213	-0.5071	0.6511
---	---	1633950_at	0.0754	0.6337	0.0500	0.6416	0.0067	0.9756	0.0739	0.9011	0.1842	0.3478	0.1103	0.5631	0.1623	0.7677	0.1497	0.5374	-0.0126	0.9721
DDB1	Drosophila damag	1633951_at	0.4048	0.0243	0.0833	0.8885	0.1169	0.4843	0.0298	0.9637	0.4833	0.0127	0.4536	0.0103	-0.0316	0.9898	0.1310	0.8569	0.1627	0.7991
CG7080	CG7080	1633952_at	0.5046	0.0378	0.0256	0.7999	-0.0769	0.7185	0.2119	0.6202	0.3649	0.0875	0.1530	0.4459	0.0672	0.9238	-0.0320	0.9315	-0.0992	0.7039
CG14744	CG14744	1633953_at	-0.0278	0.9481	0.0812	0.6629	0.1271	0.4406	0.1233	0.8479	-0.0595	0.8449	-0.1828	0.4046	-0.1158	0.9305	-0.0271	0.9696	0.0887	0.8788
CG14881	CG14881	1633954_at	-0.4542	0.2725	-0.0443	0.7831	0.1314	0.5849	-0.2508	0.6893	-0.3539	0.2233	-0.1031	0.7424	-0.3726	0.5519	0.0739	0.8527	0.4465	0.1498
pcl	scute beta	1633955_at	-6.1987	0.0008	-8.1453	0.0011	-6.2961	0.0001	1.0610	0.4317	0.7134	0.3435	-0.3476	0.6488	-0.5954	0.8479	-0.8982	0.4983	-0.3028	0.8577
CG7995	CG7995	1633956_s_at	0.6447	0.0754	-1.2019	0.0834	-0.4264	0.2134	-0.2639	0.3739	0.5195	0.0081	0.7834	0.0008	-0.9780	0.6496	-1.2192	0.1504	-0.2413	0.8251
CG5316	CG5316	1633957_s_at	-0.2196	0.2781	-0.2541	0.1889	-0.5166	0.0054	-0.0420	0.9339	0.2089	0.1593	0.2509	0.0633	0.2151	0.8326	0.1661	0.7371	-0.0490	0.9343
SamDC	S-adenosylmethio	1633958_at	0.5136	0.0167	0.4120	0.0508	0.5764	0.0083	-0.1653	0.6576	-0.1337	0.4753	0.0316	0.8815	-0.1990	0.6927	-0.0985	0.6660	0.1005	0.6533
CG9766	CG9766	1633959_s_at	0.1002	0.8803	-1.5225	0.0283	-0.4932	0.2495	0.1415	0.9249	1.2081	0.0153	1.0666	0.0162	-0.8847	0.6955	-0.4370	0.6820	0.4476	0.6669
CG5823	CG5823	1633960_at	-0.2361	0.2285	0.2144	0.3212	0.3293	0.0574	-0.2107	0.5859	-0.4529	0.0292	-0.2421	0.1684	-0.3254	0.6732	-0.1398	0.6876	0.1856	0.5746
cni	spindle-G	1633961_s_at	-0.0830	0.7849	0.4698	0.0478	0.3237	0.0485	-0.0407	0.9436	-0.5188	0.0075	-0.4781	0.0064	0.0293	0.9848	-0.0701	0.8924	-0.0994	0.8209
CG14815	CG14815	1633962_a_at	0.4337	0.1014	0.3680	0.5157	0.8179	0.0015	0.2301	0.6493	-0.2133	0.3950	-0.4434	0.0477	-0.2394	0.8717	-0.3299	0.5988	-0.0905	0.9109
nAcRalpha-30D	Dalpha6	1633963_a_at	0.0665	0.7924	-0.4361	0.1426	-0.1405	0.4536	0.2592	0.4786	0.3562	0.0723	0.0969	0.6276	-0.0733	0.9653	-0.1912	0.7072	-0.1178	0.8366
CG3626 /// DmirCG3626	CG3626	1633964_at	-0.0384	0.9577	0.4771	0.2117	0.3248	0.0680	-0.1998	0.6409	-0.5227	0.0207	-0.3230	0.0850	-0.1263	0.9677	-0.1797	0.8792	-0.0534	0.9657
CG6020	CG6020	1633965_at	-0.3664	0.1526	-0.1555	0.5992	0.1188	0.6618	-0.0105	0.9937	-0.6197	0.0318	-0.6092	0.0216	-0.2167	0.8303	-0.4279	0.3000	-0.2112	0.6423
CG30366	CG30366	1633966_at	0.1177	0.6187	0.0045	0.9699	0.2118	0.4550	0.0675	0.9470	0.1073	0.7533	0.0398	0.9086	-0.0172	0.9913	-0.0182	0.9741	-0.0011	0.9988
CG3756	CG3756	1633967_at	0.2012	0.6554	0.1337	0.6850	0.8143	0.0045	0.2826	0.6513	0.2267	0.4732	-0.0559	0.8762	-0.3836	0.7644	0.2489	0.6827	0.6325	0.2622
os	unpaired	1633968_at	0.1273	0.4691	0.0789	0.6044	0.0115	0.9711	-0.0583	0.9387	-0.1688	0.4661	-0.1105	0.6246	-0.2509	0.7979	-0.1372	0.7888	0.1137	0.8247
Osi23	Osiris	1633969_at	-0.0108	0.9630	-0.0551	0.6024	0.1308	0.4772	0.2523	0.3374	0.1869	0.2167	-0.0654	0.6802	-0.0275	0.9742	0.0954	0.6834	0.1228	0.5819
---	---	1633970_at	0.1781	0.2026	0.0590	0.6811	0.1801	0.1971	0.0275	0.9600	0.1280	0.3839	0.1005	0.4598	-0.0618	0.9174	-0.0216	0.9488	0.0402	0.8874
CG31036	CG31036	1633971_at	0.1122	0.6619	0.0463	0.8699	0.1487	0.4477	0.1060	0.8578	0.0677	0.7945	-0.0383	0.8787	0.1900	0.8215	0.1436	0.7200	-0.0464	0.9231
CG13727	CG13727	1633972_at	0.1624	0.4290	0.0419	0.8462	-0.2047	0.3543	-0.2428	0.5470	-0.0863	0.7229	0.1566	0.4234	-0.0023	0.9986	-0.1639	0.5199	-0.1616	0.5332
ox	Ubiquinol-cytochr	1633973_at	-0.2708	0.3896	0.5140	0.2158	0.4720	0.0818	-0.2914	0.3691	-1.4222	0.0002	-1.1308	0.0003	-0.1604	0.9296	-0.3828	0.5289	-0.2223	0.7426
---	---	1633974_at	0.0824	0.6849	-0.0143	0.8922	0.1257	0.4092	0.1300	0.7760	-0.0012	0.9965	-0.1312	0.4699	-0.0717	0.9243	0.0162	0.9689	0.0879	0.7640
CG12214	CG12214	1633975_s_at	-0.3598	0.1282	0.5984	0.2688	0.6881	0.0312	-0.0922	0.8891	-1.4355	0.0004	-1.3433	0.0003	-0.1341	0.9449	-0.4642	0.4301	-0.3300	0.6031
---	---	1633976_at	0.0540	0.7490	0.0726	0.6625	0.1537	0.3024	0.0110	0.9860	-0.0910	0.5695	-0.1019	0.4683	0.0387	0.9737	0.0396	0.9318	0.0009	0.9988
---	---	1633977_at	0.2207	0.4978	-0.1002	0.4811	-0.0943	0.6876	0.1035	0.8197	0.3880	0.0362	0.2845	0.0757	0.0160	0.9913	-0.0598	0.8963	-0.0758	0.8474
CG40085	CG40085	1633978_at	0.1662	0.4279	0.1057	0.5267	0.3641	0.0501	0.1864	0.6084	0.0652	0.7660	-0.1212	0.4874	-0.0266	0.9829	-0.0045	0.9941	0.0221	0.9571
---	---	1633979_at	0.3169	0.1373	-0.0038	0.9864	-0.1904	0.5135	-0.0055	0.9956	0.1569	0.5467	0.1624	0.4817	0.0462	0.9754	-0.1630	0.6937	-0.2092	0.5969
CG14969	CG14969	1633980_at	-0.5368	0.0505	-0.4590	0.2963	-0.5012	0.0100	-0.2390	0.4751	-0.3430	0.0605	-0.1039	0.5584	-0.1820	0.8814	-0.3220	0.5020	-0.1401	0.8056
---	---	1633981_at	0.0260	0.9019	0.1001	0.4622	0.3877	0.0428	-0.0161	0.9873	-0.1083	0.6984	-0.0922	0.7212	-0.1285	0.7893	-0.0078	0.9841	0.1207	0.5797
CG32088	CG32088	1633982_at	0.3663	0.0739	-0.0193	0.9511	0.3544	0.0292	-0.1279	0.8102	-0.1000	0.6789	0.0279	0.9137	-0.1574	0.7644	-0.1075	0.6536	0.0499	0.8635
CG32046	CG32046	1633983_a_at	1.1342	0.0075	1.2631	0.0110	1.4294	0.0008	-0.2226	0.7232	-0.5190	0.0701	-0.2964	0.2445	-0.4809	0.6955	-0.3722	0.4794	0.1087	0.8761
Gr85a	Gustatory recepto	1633984_at	0.2320	0.1865	-0.1551	0.4297	0.0209	0.9200	-0.0198	0.9759	0.0638	0.7377	0.0836	0.6083	-0.2141	0.7893	-0.1031	0.8157	0.1111	0.7836
---	---	1633985_at	-0.1501	0.3358	-0.9559	0.0161	-1.1132	0.0003	-0.0769	0.9004	0.5568	0.0128	0.6337	0.0043	-0.0550	0.9558	-0.1893	0.5266	-0.1343	0.6693
CG5466	CG5466	1633986_at	-0.3805	0.3115	-0.3334	0.2141	-1.2429	0.0022	-0.5121	0.0899	0.2471	0.1599	0.7592	0.0012	-0.0406	0.9913	-0.1307	0.9108	-0.0900	0.9330
---	---	1633987_at	0.2284	0.2078	0.1124	0.4044	0.1314	0.5125	0.0416	0.9542	-0.0248	0.9294	-0.0664	0.7604	0.1117	0.8400	-0.0661	0.8262	-0.1778	0.4523
CG40211	CG40211	1633988_at	-0.1610	0.3627	-0.0307	0.8433	0.0088	0.9635	0.1569	0.6869	0.1408	0.4546	-0.0161	0.9431	-0.0420	0.9441	0.0330	0.9058	0.0750	0.7144
CG1124 /// DyakCG1124	CG1124	1633989_at	-1.4245	0.0012	-2.7311	0.0101	-2.6724	0.0000	0.0314	0.9677	0.2165	0.2943	0.1851	0.3214	0.2250	0.8870	-0.8947	0.1321	-1.1198	0.0940
adat	adat	1633990_at	0.0148	0.9405	-0.8444	0.0689	-1.0008	0.0124	0.3194	0.6447	1.3771	0.0024	1.0577	0.0045	-0.0685	0.9309	-0.0765	0.8197	-0.0079	0.9849
CG31445	CG31445	1633991_at	-0.9484	0.0131	-0.0063	0.9960	-0.6929	0.0203	-0.5107	0.1956	-1.3665	0.0006	-0.8558	0.0025	0.1457	0.9522	-0.4186	0.5889	-0.5642	0.4554
CG2137	CG2137	1633992_at	0.3039	0.0730	0.0520	0.8489	0.0072	0.9721	0.0325	0.9610	0.0003	0.9992	-0.0322	0.8779	0.0311	0.9742	-0.3186	0.1727	-0.3497	0.1701
CG30126	CG30126	1633993_at	-0.0310	0.8717	0.0707	0.6492	0.0879	0.6061	0.0732	0.8791	-0.1395	0.4090	-0.2127	0.1457	0.0354	0.9771	-0.0061	0.9925	-0.0415	0.9176
CG34402	CG17793	1633994_at	0.2717	0.3980	-0.1095	0.4567	-0.1529	0.3181	-0.0240	0.9647	0.1077	0.4722	0.1317	0.3118	0.0279	0.9832	-0.2202	0.4893	-0.2481	0.4374
CG13087	CG13087	1633995_at	0.0572	0.8152	0.1034	0.4034	0.2845	0.1804	-0.0243	0.9838	-0.0204	0.9610	0.0040	0.9907	-0.0828	0.9499	0.0531	0.9325	0.1358	0.7655
---	---	1633996_at	-0.0938	0.5154	-0.0149	0.8924	-0.0565	0.7185	-0.0779	0.8851	-0.1677	0.3699	-0.0898	0.6314	0.0738	0.8724	0.0211	0.9425	-0.0527	0.8161
CG5938	CG5938	1633997_s_at	-0.6300	0.0117	-0.2970	0.2730	-0.3272	0.0889	0.0128	0.9863	-0.3044	0.0930	-0.3172	0.0542	-0.0583	0.9651	-0.0211	0.9719	0.0372	0.9409
---	---	1633998_s_at	0.7905	0.0400	0.6641	0.4282	0.2112	0.2907	-0.3038	0.5863	-0.1801	0.5566	0.1236	0.6776	0.1821	0.9460	-0.3602	0.6949	-0.5423	0.5270
---	---	1633999_at	0.0542	0.7497	0.0068	0.9810	0.1601	0.2911	0.1223	0.8686	0.1836	0.5045	0.0613	0.8376	-0.1253	0.8344	-0.0354	0.9282	0.0899	0.7541
CG1847	CG1847	1634000_a_at	-0.0380	0.8197	-0.0022	0.9923	-0.1475	0.4539	-0.0883	0.8628	-0.0298	0.9058	0.0585	0.7710	0.0940	0.9142	0.0691	0.8739	-0.0249	0.9538
CG9062	CG9062	1634001_at	0.2613	0.4206	0.3896	0.1673	0.8186	0.0008	0.1258	0.8485	0.1965	0.4305	0.0707	0.7932	-0.1469	0.9087	0.5960	0.1742	0.7428	0.1313
CG13315 /// DyakCG13315	CG13315	1634002_at	1.4331	0.0234	1.6381	0.1057	2.4187	0.0001	0.7465	0.2939	-0.8079	0.0576	-1.5543	0.0021	-0.0629	0.9875	-0.4681	0.6310	-0.4052	0.6848
CG32685	CG32685	1634003_at	0.1489	0.7020	0.3790	0.4986	-0.0433	0.8670	-0.2117	0.6854	0.1433	0.5879	0.3550	0.1017	0.2369	0.9061	0.4270	0.5747	0.1901	0.8348
CG13028	CG13028	1634004_at	-0.4180	0.0442	0.0946	0.6112	-0.1716	0.3849	0.0408	0.9649	-0.1674	0.5229	-0.2082	0.3579	0.0117	0.9906	0.0253	0.9404	0.0136	0.9655
CG7906	CG7906	1634005_at	0.2211	0.1964	0.1368	0.3407	0.2103	0.3141	0.1249	0.7929	0.0267	0.9220	-0.0982	0.6176	0.2021	0.7677	0.2152	0.4633	0.0131	0.9775
ATPCL	ATP citrate lyase	1634006_s_at	0.1252	0.4021	-0.0895	0.8161	-0.1862	0.2306	-0.0763	0.8837	0.4245	0.0249	0.5008	0.0075	-0.0822	0.9589	0.3340	0.4655	0.4162	0.3703
---	---	1634007_at	0.1658	0.3575	0.0615	0.5624	0.0511	0.8476	-0.0464	0.9422	0.0233	0.9272	0.0697	0.7187	-0.0568	0.9333	-0.1076	0.6483	-0.0509	0.8577
CG34041	CG34041	1634008_at	0.0145	0.9405	-0.2174	0.4267	0.1093	0.4919	0.0509	0.9436	0.1777	0.4019	0.1268	0.5276	-0.2585	0.6728	-0.0833	0.7870	0.1752	0.4925
---	---	1634009_at	-0.2650	0.2342	-0.5777	0.0145	-0.9416	0.0035	-0.2614	0.3739	-0.1035	0.5631	0.1579	0.2933	0.0544	0.9737	-0.4209	0.2842	-0.4753	0.2574
---	---	1634010_at	0.0936	0.7850	-0.0240	0.8736	0.0500	0.8574	-0.1311	0.8742	-0.0447	0.9114	0.0864	0.7892	-0.1154	0.8786	-0.2071	0.4871	-0.0916	0.7940

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
HLHmdelta	split locus enhanc	1634011_at	-0.3076	0.3686	-0.1854	0.3885	-0.0420	0.8736	0.0479	0.9757	-0.1428	0.7553	-0.1907	0.6257	0.0264	0.9751	0.0034	0.9941	-0.0230	0.9371
CG5002	CG5002	1634012_at	-1.9918	0.0459	-1.9308	0.0200	-1.4101	0.0003	-0.1119	0.9228	-0.0108	0.9833	0.1011	0.7965	-0.6000	0.8461	-0.1173	0.9550	0.4827	0.7439
mRpL17	mitochondrial ribo	1634013_at	-0.1907	0.5696	0.3513	0.3803	0.4350	0.0269	0.0682	0.9283	-0.6305	0.0123	-0.6987	0.0047	0.0291	0.9913	0.1273	0.8660	0.0982	0.8939
CG31773	CG31773	1634014_at	0.1183	0.4539	0.0379	0.7529	0.1985	0.2204	-0.0383	0.9683	-0.0925	0.7580	-0.0542	0.8526	-0.0880	0.8686	0.0007	0.9994	0.0887	0.7071
CG5077	CG5077	1634015_s_at	-0.2325	0.6348	0.0673	0.8667	-0.1768	0.4930	0.1918	0.6122	0.3377	0.0776	0.1459	0.4110	0.5262	0.7633	0.7193	0.3135	0.1930	0.8354
CG2781	CG2781	1634016_at	1.0837	0.4207	-1.5948	0.0647	0.4686	0.3819	1.6364	0.1422	1.1887	0.0797	-0.4477	0.4879	-0.2263	0.9775	-1.3647	0.4852	-1.1384	0.5758
CG12017	CG12017	1634017_at	-0.1490	0.3538	-0.1195	0.4658	-0.0426	0.8281	-0.0431	0.9596	0.0524	0.8574	0.0955	0.6827	-0.1472	0.7726	0.0347	0.9194	0.1819	0.4203
CG34028	CG34028	1634018_at	0.6972	0.2405	0.9844	0.6072	1.8605	0.0002	0.1846	0.6321	0.2276	0.2267	0.0430	0.8434	-0.3416	0.9589	0.6234	0.7963	0.9649	0.6391
CG2064	CG2064	1634019_at	0.2643	0.2248	0.1862	0.4742	0.8063	0.0115	0.1129	0.8327	-0.1785	0.3909	-0.2914	0.1075	-0.5331	0.6287	-0.2440	0.6218	0.2891	0.5520
mei-P26	mei-P26	1634020_at	1.0609	0.0105	0.1644	0.7132	-0.8956	0.1067	-0.2001	0.7140	1.3093	0.0008	1.5094	0.0003	0.8920	0.6955	0.5218	0.6093	-0.3702	0.7355
Psi	P-element somatic	1634021_a_at	0.4311	0.7292	0.3871	0.4714	-0.3376	0.1184	-0.2674	0.6936	0.6874	0.0353	0.9548	0.0053	0.4902	0.9147	0.8190	0.6382	0.3288	0.8801
CG17745	CG17745	1634022_at	0.2619	0.1174	-0.0321	0.8827	-0.0597	0.6941	0.2492	0.3924	0.4439	0.0148	0.1947	0.1838	0.0889	0.9242	0.0415	0.9341	-0.0474	0.9121
CG14544	CG14544	1634023_at	0.2627	0.3832	0.5836	0.2598	1.1245	0.0044	0.2729	0.7511	0.0437	0.9338	-0.2291	0.5295	-0.1566	0.9514	0.3569	0.6713	0.5135	0.5176
---	---	1634024_at	0.1827	0.2347	0.1441	0.4267	0.1755	0.2497	0.2148	0.6086	0.0798	0.7499	-0.1350	0.5066	0.0199	0.9816	0.0260	0.9279	0.0061	0.9843
CG13010	CG13010	1634025_at	-0.0056	0.9842	0.0598	0.7078	0.2891	0.1188	0.1449	0.7121	0.1195	0.5295	-0.0254	0.9063	-0.1317	0.8609	0.1279	0.7106	0.2595	0.4017
---	---	1634026_at	-0.0425	0.8128	-0.0438	0.6944	-0.1935	0.4460	0.1144	0.7059	0.1119	0.4375	-0.0026	0.9880	0.0774	0.9467	-0.0412	0.9421	-0.1186	0.7751
CG12715	CG12715	1634027_at	0.2810	0.1740	0.0159	0.8859	0.0542	0.7370	-0.0324	0.9602	0.0766	0.6959	0.1090	0.5076	-0.0622	0.9400	-0.2304	0.3552	-0.1682	0.5299
Fmr1	Fragile X	1634028_s_at	-0.9348	0.0884	-0.0936	0.7689	-0.4647	0.1619	-0.0999	0.9196	-0.2989	0.3317	-0.1989	0.4947	0.3213	0.8903	0.5922	0.5140	0.2709	0.7983
Ugt86Dj	Ugt86Dj	1634029_at	-0.1542	0.8889	-0.0513	0.7739	-0.1290	0.4683	-0.1425	0.9254	-0.1786	0.7399	-0.0361	0.9502	0.0601	0.9893	-0.3599	0.7439	-0.4200	0.6857
CG40390	CG40390	1634030_at	0.2215	0.3047	0.0411	0.8638	0.2461	0.1230	0.2266	0.4337	0.0658	0.7224	-0.1608	0.2613	-0.0494	0.9677	-0.0971	0.8169	-0.0477	0.9155
Best2	Bestrophin 2	1634031_at	-0.5387	0.2123	0.4342	0.1726	-0.5249	0.0124	-0.3319	0.4442	-0.5148	0.0362	-0.1829	0.4068	0.6577	0.5765	0.5319	0.3043	-0.1258	0.8562
---	---	1634032_at	0.1231	0.5934	0.1866	0.1239	0.0622	0.8197	-0.1060	0.8967	-0.0470	0.8984	0.0591	0.8495	0.1460	0.8756	0.0682	0.9022	-0.0777	0.8745
Glut1	glucose transporte	1634033_s_at	-1.0208	0.0078	-0.5839	0.0436	-0.2275	0.1968	-0.2195	0.5928	-0.8251	0.0027	-0.6055	0.0064	-0.6051	0.4120	-0.2647	0.5172	0.3404	0.3967
RhoGAP68F	RhoGAP68F	1634034_at	0.3912	0.2097	0.5198	0.0494	0.6393	0.0127	-0.2502	0.5515	-0.4430	0.0471	-0.1928	0.3322	-0.2955	0.8235	-0.2952	0.6173	0.0003	0.9997
Ephrin	ephrin	1634035_s_at	-0.3158	0.3255	0.2468	0.5929	-0.3751	0.0861	-0.6857	0.0999	-0.0614	0.8471	0.6243	0.0121	-0.0032	0.9994	0.4590	0.4333	0.4622	0.4448
CG8788	CG8788	1634036_at	1.9422	0.0020	2.0511	0.0077	2.3940	0.0001	0.1999	0.8432	-0.5955	0.1108	-0.7954	0.0257	-0.1248	0.9243	-0.4103	0.3358	-0.2855	0.5361
Ssdp	Sequence-specific	1634037_s_at	-0.3713	0.3314	-0.3092	0.3821	-0.1655	0.6997	-0.1438	0.8009	-0.0409	0.8962	0.1029	0.6660	-0.2490	0.9149	0.1184	0.9273	0.3673	0.6842
CG14856	CG14856	1634038_a_at	0.0778	0.6379	-0.0828	0.4063	0.1537	0.5630	0.0714	0.8908	0.0310	0.8965	-0.0404	0.8407	-0.0443	0.9717	-0.0314	0.9507	0.0129	0.9807
nerfin-1	nerfin 1	1634039_at	-0.4097	0.6291	-0.0294	0.7689	0.1665	0.4082	0.1604	0.9441	-0.4101	0.5621	-0.5706	0.3425	-0.1479	0.9306	-0.3098	0.5968	-0.1618	0.8122
---	---	1634040_at	0.1461	0.3980	0.0066	0.9604	0.3554	0.0371	0.0008	0.9994	0.1187	0.6025	0.1179	0.5676	-0.2798	0.5421	-0.0747	0.7769	0.2051	0.3538
CG31548	CG31548	1634041_at	-0.1024	0.7485	0.4668	0.2441	0.6245	0.0180	-0.1114	0.8610	-1.1862	0.0009	-1.0748	0.0008	-0.4386	0.7215	-0.8189	0.1181	-0.3803	0.4776
---	---	1634042_at	-0.1412	0.4211	0.2178	0.1953	0.0494	0.8547	-0.0212	0.9803	-0.1668	0.4352	-0.1456	0.4551	-0.1220	0.8940	0.0031	0.9975	0.1251	0.7492
CG5721 /// DmirCG5721	CG5721	1634043_at	0.1437	0.3572	0.7840	0.0256	0.4797	0.0092	0.0507	0.9254	-0.2169	0.1811	-0.2676	0.0690	0.2866	0.6749	0.3931	0.1562	0.1065	0.7464
bib	big brain	1634044_at	0.0136	0.9629	-0.0424	0.6687	0.1263	0.5105	0.0046	0.9956	-0.2026	0.3854	-0.2072	0.3188	-0.1170	0.8465	-0.1378	0.6073	-0.0208	0.9527
CG10748	CG10748	1634045_at	0.3621	0.1340	-0.0897	0.3755	0.0613	0.7242	0.3056	0.4142	0.3979	0.0598	0.0922	0.6697	-0.1105	0.8940	-0.2707	0.3706	-0.1602	0.6294
---	---	1634046_at	0.3671	0.2376	0.0600	0.5845	0.3608	0.0503	-0.0275	0.9733	-0.0425	0.8744	-0.0150	0.9537	-0.1350	0.8657	0.0036	0.9961	0.1387	0.6973
CG9028	CG9028	1634047_a_at	-0.0971	0.5769	-0.0022	0.9940	0.0511	0.7397	0.2050	0.6006	0.0305	0.9114	-0.1745	0.3329	0.0778	0.8999	0.1432	0.5469	0.0654	0.8178
bgm	bubblegum	1634048_a_at	2.2192	0.0008	1.4146	0.0286	2.3227	0.0001	0.8942	0.1551	0.7249	0.0610	-0.1693	0.6690	-0.1194	0.9239	-0.2170	0.6312	-0.0976	0.8578
CG17626	CG17626	1634049_at	0.1249	0.6188	0.0935	0.4690	0.1451	0.4248	0.0916	0.9171	0.1394	0.6492	0.0479	0.8818	0.0606	0.9491	0.1952	0.5041	0.1346	0.6607
CG17122	CG17122	1634050_at	0.0555	0.8444	-0.0058	0.9710	0.2262	0.3396	0.0119	0.9895	-0.0143	0.9643	-0.0261	0.9187	-0.1101	0.8521	-0.0212	0.9587	0.0889	0.7522
---	---	1634051_at	0.3035	0.1727	0.0784	0.5334	-0.0035	0.9910	0.2351	0.4356	0.3171	0.0594	0.0820	0.6262	0.0572	0.9514	-0.0182	0.9682	-0.0755	0.8287
mod(mdg4)	Modifier67.2	1634052_s_at	-0.3219	0.1336	-0.0721	0.5916	-0.1192	0.4546	-0.0309	0.9661	-0.1835	0.3459	-0.1526	0.3906	0.1661	0.7953	0.0688	0.8519	-0.0973	0.7534
CG8408	CG8408	1634053_at	-0.2500	0.1229	0.8331	0.0638	0.5833	0.0094	0.1455	0.6856	-0.3407	0.0471	-0.4863	0.0065	0.5823	0.4128	0.8196	0.0532	0.2373	0.5585
ninaC	Droninac	1634054_at	0.2086	0.2645	0.0517	0.7349	0.3063	0.0956	0.0459	0.9343	0.1585	0.3407	0.1126	0.4684	-0.1488	0.7644	-0.0956	0.6837	0.0532	0.8432
Mat1	Mat1	1634055_at	-0.3228	0.1216	-0.1856	0.1195	-0.0986	0.5488	-0.0088	0.9903	-0.0571	0.7664	-0.0484	0.7871	-0.1323	0.8270	0.0525	0.8867	0.1848	0.4835
lbe	ladybird	1634056_at	0.1443	0.5686	0.1001	0.3219	0.3363	0.0529	0.1662	0.6010	0.0577	0.7640	-0.1085	0.4771	-0.0276	0.9657	0.0513	0.8196	0.0789	0.6771
CG7707	CG7707	1634057_at	0.0342	0.8693	-0.1245	0.4131	-0.0784	0.6959	0.0203	0.9825	0.1674	0.4569	0.1471	0.4748	-0.0673	0.9117	-0.0981	0.6826	-0.0308	0.9174
---	---	1634058_at	0.0733	0.6768	0.2402	0.2857	0.0365	0.8810	-0.0009	0.9988	0.0015	0.9951	0.0024	0.9906	0.1881	0.8122	0.1652	0.6425	-0.0229	0.9622
CG12543	CG12543	1634059_at	0.3522	0.0952	0.0866	0.5015	0.2993	0.2199	-0.0557	0.9185	0.0212	0.9281	0.0770	0.6567	0.0302	0.9737	-0.0043	0.9935	-0.0345	0.9118
---	---	1634060_at	0.0365	0.8992	0.0158	0.9136	0.1260	0.5269	0.0381	0.9671	0.0072	0.9839	-0.0310	0.9162	-0.0224	0.9752	0.0702	0.7349	0.0926	0.6294
rdgC	retinal degeneratio	1634061_a_at	-1.3264	0.0183	-0.2349	0.3716	-0.6834	0.0235	-0.4781	0.4167	-1.0501	0.0069	-0.5720	0.0544	0.0454	0.9816	0.0898	0.8845	0.0444	0.9411
CG14218	CG14218	1634062_at	0.0490	0.8735	-0.0127	0.9364	0.0033	0.9880	0.0949	0.8174	0.1176	0.4868	0.0227	0.9075	0.0534	0.9618	-0.0127	0.9829	-0.0662	0.8745
stumps	heartbroken	1634063_a_at	-1.2773	0.0048	-1.6783	0.0096	-1.8932	0.0002	-0.2721	0.4907	-0.4961	0.0261	-0.2240	0.2342	-0.0375	0.9901	-1.0347	0.1212	-0.9972	0.1568
CG13311	CG13311	1634064_at	0.0364	0.9105	-0.0085	0.9382	0.1066	0.5827	-0.0912	0.8897	-0.1327	0.5783	-0.0415	0.8730	-0.0395	0.9679	-0.2128	0.4094	-0.1733	0.5257
Eno	phosphopyruvate	1634065_s_at	-0.4486	0.0490	-0.3310	0.4958	-0.3150	0.1327	-0.0216	0.9819	-0.5018	0.0336	-0.4802	0.0257	-0.1685	0.9142	-0.4048	0.4656	-0.2363	0.6993
---	---	1634066_at	0.0569	0.7769	0.3022	0.1252	0.2640	0.1736	-0.0332	0.9630	-0.1149	0.5773	-0.0818	0.6814	-0.1270	0.8395	-0.0207	0.9617	0.1063	0.7154
---	---	1634067_at	-0.0937	0.5256	0.2041	0.2178	0.1145	0.5096	-0.0387	0.9376	-0.1024	0.5014	-0.0638	0.6717	0.1892	0.7770	0.1841	0.5425	-0.0051	0.9924
CG1631	CG1631	1634068_at	0.0377	0.8142	0.0779	0.6013	-0.2101	0.2434	-0.2414	0.5681	-0.0517	0.8545	0.1897	0.3405	0.0478	0.9599	0.0889	0.7933	0.0411	0.9121
CG40109	CG40109	1634069_at	-1.4534	0.0089	-1.1553	0.2024	-1.3897	0.0193	0.0191	0.9940	-0.4537	0.3982	-0.4727	0.3218	0.1445	0.9521	-0.1630	0.8749	-0.3075	0.7040
CG4101	CG4101	1634070_at	-0.9199	0.0023	-0.6254	0.0234	-0.5764	0.0472	0.0221	0.9789	-0.1002	0.6729	-0.1224	0.5500	-0.0081	0.9964	0.1872	0.7024	0.1954	0.6839
kek1	Kekkon	1634071_at	-0.0996	0.8160	0.7879	0.3638	0.3347	0.2277	0.0057	0.9952	-0.1067	0.6207	-0.1124	0.5584	0.6845	0.7644	0.8428	0.3693	0.1583	0.9050
Hmgs	HMG-CoA syntha	1634072_s_at	0.3665	0.1037	0.3308	0.2861	0.6350	0.0047	0.2428	0.4122	0.3683	0.0321	0.1255	0.4133	-0.0049	0.9982	0.3360	0.4598	0.3409	0.4653
CG13466	CG13466	1634073_at	-0.8805	0.0019	-0.2169	0.1838	-0.7320	0.0080	-0.7259	0.0796	-1.2387	0.0009	-0.5128	0.0248	-0.1723	0.7633	-0.2949	0.2018	-0.1226	0.6328
yin	oligopeptide trans	1634074_a_at	0.1938	0.7511	-0.5077	0.0260	-0.8032	0.1196	0.0615	0.9733	1.8655	0.0023	1.8040	0.0016	0.1944	0.9487	0.9583	0.2722	0.7640	0.4061
dp	Dumpy	1634075_at	0.7532	0.4674	-1.1980	0.0302	-3.0639	0.0067	-0.3789	0.5744	2.7542	0.0002	3.1332	0.0001	1.4871	0.7307	0.7861	0.7072	-0.7010	0.7433
CG12057	CG12057	1634076_at	0.1530	0.6632	0.1427	0.6473	0.2812	0.1291	0.0347	0.9749	-0.0902	0.7844	-0.1249	0.6563	-0.0603	0.9735	-0.0658	0.9264	-0.0055	0.9945
CG4789	CG4789	1634077_at	0.0664	0.7832	0.2918	0.1797	0.2102	0.3798	0.1400	0.8386	0.0641	0.8480	-0.0760	0.7926	0.1859	0.8331	0.1518	0.7183	-0.0341	0.9471
Sirt7	Sirt7	1634078_at	0.3959	0.0373	0.5447	0.2285	0.4669	0.0850	-0.2469	0.5633	-0.2346	0.2886	0.0124	0.9645	-0.1385	0.8882	-0.0427	0.9428	0.0958	0.8379
CG32568	CG32568	1634079_at	0.2022	0.4321	-0.2172	0.4345	-0.2608	0.1572	-0.0036	0.9956	0.3295	0.1065	0.3331	0.0712	-0.0361	0.9831	-0.1540	0.7393	-0.1179	0.8080

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14537	CG14537	1634080_at	-0.0249	0.9466	-0.1728	0.5370	0.0538	0.7439	0.1553	0.8544	0.1244	0.7273	-0.0309	0.9351	0.1520	0.8013	0.1527	0.5630	0.0007	0.9990
RfC38	DNA replication fa	1634081_at	0.2585	0.1089	0.1019	0.7334	0.1203	0.5919	-0.1005	0.7604	0.6229	0.0018	0.7234	0.0006	-0.0047	0.9978	0.4798	0.1894	0.4845	0.2148
eIF4E-4	eIF4E-4	1634082_at	0.1578	0.5275	0.0028	0.9933	-0.0494	0.7732	-0.0455	0.9345	0.1520	0.3604	0.1975	0.1741	0.1137	0.8875	0.0786	0.8481	-0.0351	0.9342
Nf1	neurofibromatosis	1634083_at	-0.4764	0.1283	-0.4854	0.2732	-0.9649	0.0445	-0.1599	0.8834	0.3399	0.3695	0.4998	0.1299	0.2614	0.8906	0.0514	0.9650	-0.2100	0.8138
CG4213	CG4213	1634084_at	-2.1760	0.0010	-1.9556	0.0055	-2.0897	0.0002	0.1294	0.6988	0.0288	0.8902	-0.1006	0.4948	0.3191	0.7726	0.1220	0.8535	-0.1971	0.7154
CG31204	CG31204	1634085_at	0.2627	0.3381	0.5735	0.0494	0.2617	0.3128	0.0664	0.9473	0.0395	0.9213	-0.0269	0.9393	0.4393	0.4562	0.2869	0.3322	-0.1524	0.6403
faf	fat facets	1634086_a_at	-0.2265	0.6663	0.7254	0.1698	0.9507	0.0046	0.0724	0.9536	-0.3562	0.3002	-0.4287	0.1582	0.1195	0.9515	0.8190	0.1523	0.6994	0.2532
CG1311	CG1311	1634087_at	-0.3771	0.2354	-1.4142	0.0395	-0.9190	0.0095	0.2955	0.6114	1.0148	0.0048	0.7193	0.0136	-0.1014	0.9611	0.0790	0.9326	0.1803	0.7983
---	---	1634088_at	0.0926	0.4955	0.0069	0.9534	0.0173	0.9253	0.0373	0.9592	0.0830	0.7088	0.0457	0.8358	0.0689	0.9076	0.0585	0.8359	-0.0104	0.9756
---	---	1634089_at	0.4258	0.0483	0.0743	0.6770	0.2156	0.2497	-0.0812	0.9247	0.0898	0.7711	0.1710	0.4848	-0.0350	0.9717	-0.1607	0.5259	-0.1256	0.6328
CG14633	CG14633	1634090_at	-0.0787	0.7529	-0.0237	0.9061	0.2057	0.2495	0.1246	0.7825	0.0404	0.8704	-0.0842	0.6636	-0.0905	0.9016	0.0135	0.9778	0.1040	0.7366
Cp19	Shell-19	1634091_at	-0.0593	0.7871	0.1989	0.3324	0.1117	0.6124	-0.2285	0.6854	-0.4669	0.0769	-0.2384	0.3182	-0.1105	0.8141	-0.1387	0.4861	-0.0282	0.9168
CG3654	CG3654	1634092_at	0.3765	0.3955	-0.5012	0.0440	-0.3689	0.2171	0.3322	0.4140	1.0866	0.0012	0.7544	0.0034	0.1220	0.9555	0.1936	0.8153	0.0716	0.9360
CG32017	CG32017	1634093_at	-0.8235	0.0434	-2.0955	0.0106	-1.9175	0.0002	0.2391	0.7799	1.1077	0.0079	0.8686	0.0142	0.0447	0.9722	0.0463	0.9278	0.0016	0.9979
pita	spotted dick	1634094_a_at	-0.5146	0.1230	0.3372	0.2687	0.8923	0.0016	0.1970	0.6955	-0.8147	0.0047	-1.0117	0.0011	-0.1466	0.8736	0.0552	0.9222	0.2018	0.6043
Ef1beta	Elongation factor	1634095_at	0.3625	0.0511	0.6568	0.0225	0.7525	0.0011	0.0642	0.9011	-0.1914	0.2538	-0.2556	0.0876	0.0311	0.9717	0.1435	0.5283	0.1125	0.6334
Aldh	Acetaldehyde deh	1634096_at	0.3565	0.0374	0.4424	0.0172	0.6661	0.0024	-0.1199	0.7409	-0.3539	0.0342	-0.2339	0.1015	-0.3527	0.3324	-0.3293	0.0945	0.0234	0.9300
CG11659 /// CG6300	CG11659 /// CG6	1634097_s_at	-0.0345	0.9040	-0.0431	0.6724	-0.0445	0.7859	-0.0513	0.9470	-0.1870	0.4047	-0.1356	0.5230	-0.0473	0.9589	-0.3317	0.1948	-0.2845	0.2960
---	---	1634098_at	0.0238	0.9311	0.1077	0.3714	0.0380	0.8275	-0.1208	0.7161	0.0018	0.9937	0.1226	0.3819	-0.0055	0.9963	0.0721	0.8313	0.0776	0.8028
rdgB	receptor-degener	1634099_at	-1.1856	0.0177	-2.2616	0.0112	-1.8232	0.0001	-0.1962	0.7409	0.2502	0.3470	0.4464	0.0599	-0.7186	0.6749	-0.8649	0.1998	-0.1463	0.8774
CG13865	CG13865	1634100_at	-0.1855	0.3969	0.4573	0.0786	0.8775	0.0096	-0.1732	0.6533	-0.7581	0.0024	-0.5849	0.0045	-0.4150	0.6927	-0.0174	0.9833	0.3977	0.3701
CG3075	NF-YC-like	1634101_at	-0.1359	0.6896	0.0270	0.9700	0.2912	0.2089	0.1733	0.6247	-0.0238	0.9227	-0.1971	0.2092	-0.0788	0.9816	0.0407	0.9766	0.1196	0.9121
sls	D-Titin	1634102_at	-2.1771	0.0043	-1.1004	0.3495	-2.2755	0.0017	-0.7372	0.3068	-1.1220	0.0163	-0.3848	0.3198	0.3632	0.9324	-0.1554	0.9449	-0.5186	0.7441
CG15122	CG15122	1634103_at	0.0902	0.6261	0.0774	0.6228	0.1731	0.4362	0.2557	0.5863	0.1624	0.5243	-0.0933	0.7136	0.0230	0.9862	0.0367	0.9402	0.0137	0.9769
Top1	topoisomerase I	1634104_a_at	0.8501	0.0098	1.0824	0.0088	1.1127	0.0003	-0.1734	0.6937	0.3008	0.1381	0.4742	0.0173	-0.1132	0.8819	0.6809	0.0413	0.7942	0.0417
---	---	1634105_s_at	0.0326	0.8758	-0.0004	0.9998	0.1189	0.4955	0.1540	0.6998	-0.0162	0.9529	-0.1702	0.3064	-0.0193	0.9869	-0.0287	0.9477	-0.0094	0.9837
CG11686	CG11686	1634106_at	0.8982	0.0113	2.4908	0.0014	1.2689	0.0126	-0.0183	0.9774	-0.4765	0.0089	-0.4583	0.0063	1.1810	0.3800	1.1118	0.1343	-0.0692	0.9470
Prx6005	Peroxiredoxin 600	1634107_at	-0.0093	0.9696	0.1615	0.4701	0.1774	0.3126	-0.0118	0.9867	-0.4181	0.0239	-0.4063	0.0168	0.0789	0.9426	-0.1822	0.6152	-0.2611	0.4554
CG17991	CG17991	1634108_at	0.0358	0.9117	0.0669	0.6899	-0.0201	0.9045	-0.0171	0.9853	-0.0388	0.8906	-0.0217	0.9328	0.1175	0.9076	-0.0062	0.9940	-0.1237	0.7691
bnl	fibroblast growth f	1634109_a_at	-0.3080	0.1042	0.0868	0.6581	0.0489	0.8218	-0.0107	0.9935	-0.2249	0.3882	-0.2142	0.3600	0.1326	0.8564	0.0586	0.8962	-0.0740	0.8474
CG17304 /// DereCG17304	CG17304	1634110_at	-0.1596	0.4979	-0.0689	0.6286	-0.0126	0.9554	-0.0034	0.9960	-0.1371	0.5289	-0.1337	0.4952	0.1810	0.7506	0.0935	0.7442	-0.0874	0.7560
---	---	1634111_at	0.0630	0.8639	-0.5941	0.1792	0.0064	0.9891	0.2264	0.7170	0.4952	0.0829	0.2688	0.2971	-0.3235	0.7644	-0.2171	0.6673	0.1064	0.8591
CG5174	CG5174	1634112_a_at	0.2545	0.2617	0.5198	0.0447	0.6125	0.0090	0.1664	0.6654	0.0894	0.6685	-0.0770	0.6896	0.1073	0.9142	0.3923	0.2430	0.2849	0.4243
gk	geko	1634113_at	1.1692	0.0325	0.4229	0.4878	1.5731	0.0005	1.0784	0.1109	1.5495	0.0030	0.4711	0.1878	-0.0588	0.9848	0.7401	0.2808	0.7989	0.2757
CG31517	CG31517	1634114_at	-0.0553	0.8696	-0.3962	0.4496	-0.2428	0.3267	-0.0623	0.9034	0.1724	0.3041	0.2346	0.1129	-0.1060	0.9665	0.0561	0.9596	0.1621	0.8519
Cpr31A	CG33302	1634115_a_at	-0.0518	0.7987	-0.1590	0.3798	0.2097	0.4362	-0.0262	0.9782	-0.0651	0.8360	-0.0389	0.8952	-0.1206	0.8326	-0.0761	0.8018	0.0445	0.8918
CG7158	CG7158	1634116_at	-0.2257	0.4302	-0.3375	0.3813	-0.1776	0.4215	0.0547	0.9518	0.0119	0.9757	-0.0428	0.8856	-0.0862	0.9589	-0.0459	0.9508	0.0403	0.9512
CG6204	CG6204	1634117_at	0.3270	0.1470	0.0255	0.9522	0.0115	0.9647	0.1618	0.6854	0.5758	0.0081	0.4140	0.0209	0.0858	0.9589	0.1474	0.8117	0.0616	0.9252
CG15149	CG15149	1634118_at	-0.1654	0.3596	-0.0263	0.8105	-0.1020	0.5709	-0.1813	0.6886	-0.0217	0.9438	0.1596	0.4161	-0.0430	0.9598	0.0390	0.9175	0.0820	0.7749
CG14892	CG14892	1634119_at	0.2469	0.1799	0.2451	0.0962	0.1785	0.3455	0.1901	0.6015	0.0500	0.8308	-0.1400	0.4155	0.1469	0.8202	-0.0638	0.8642	-0.2108	0.4436
Crtp	Caldesmon-relate	1634120_at	0.1260	0.5379	-0.1317	0.3975	0.2817	0.1966	-0.1604	0.7979	-0.0323	0.9278	0.1281	0.6192	-0.1468	0.8611	-0.1891	0.6026	-0.0423	0.9288
Obp57c	Odorant-binding p	1634121_at	-3.3377	0.0055	-0.1465	0.8535	-2.0743	0.0081	-1.7922	0.1462	-3.5093	0.0013	-1.7170	0.0179	0.2033	0.9391	-0.4304	0.6270	-0.6337	0.4557
CG31223	CG31223	1634122_at	0.1227	0.5677	0.0782	0.8518	0.2587	0.3015	-0.0592	0.9037	0.1774	0.2661	0.2366	0.0952	-0.3664	0.7848	0.1532	0.8446	0.5197	0.3807
CG14304	CG14304	1634123_at	0.0065	0.9732	0.0772	0.5926	-0.0559	0.7499	-0.1689	0.6424	-0.0349	0.8834	0.1339	0.4192	0.0674	0.9095	-0.0669	0.8057	-0.1343	0.5488
CG30428	CG30428	1634124_at	0.0036	0.9961	1.1636	0.0446	0.7578	0.0161	0.0906	0.9266	0.0065	0.9888	-0.0842	0.7996	0.4652	0.7633	1.1076	0.0952	0.6424	0.3299
CG30440	CG30440	1634125_at	-0.8789	0.0055	-2.5036	0.0092	-2.4750	0.0002	0.1210	0.9300	1.2646	0.0078	1.1435	0.0073	0.0962	0.9421	-0.2385	0.5818	-0.3346	0.4220
CG31788	CG31788	1634126_at	-0.0210	0.9116	-0.0273	0.8273	-0.0005	0.9980	0.0714	0.8761	0.1048	0.5346	0.0335	0.8556	0.0465	0.9514	0.1439	0.5431	0.0974	0.7011
CG13029 /// DereCG13029	CG13029	1634127_a_at	0.0530	0.7632	-0.2855	0.1267	-0.2018	0.3007	0.1305	0.8584	0.2538	0.3435	0.1233	0.6501	-0.0229	0.9852	-0.0798	0.8319	-0.0569	0.8833
CG15876	CG15876	1634128_at	0.1163	0.4561	0.0203	0.8419	0.0742	0.7058	-0.1498	0.7400	-0.0500	0.8461	0.0998	0.6218	0.0831	0.9030	0.1189	0.6633	0.0358	0.9171
CG9663	CG9663	1634129_at	0.8374	0.0371	0.6045	0.2012	0.2133	0.3085	-0.3031	0.5128	-0.2666	0.2786	0.0364	0.9012	0.0991	0.9515	-0.2476	0.6425	-0.3468	0.4970
CG6767	CG6767	1634130_a_at	1.3247	0.0051	0.4409	0.2180	0.4757	0.0877	0.1402	0.8189	0.3797	0.1121	0.2395	0.2676	0.1439	0.9467	-0.4942	0.4512	-0.6382	0.3437
Mnt	Mnt	1634131_s_at	0.7963	0.1866	0.6949	0.0958	0.0366	0.9447	0.1872	0.7582	1.3008	0.0011	1.1136	0.0012	0.8676	0.6955	1.0669	0.2351	0.1993	0.8732
CG33697	CG33697	1634132_at	0.1933	0.2681	-0.0045	0.9865	0.1965	0.5261	0.1524	0.8550	0.2267	0.4722	0.0743	0.8292	-0.0967	0.9238	-0.1124	0.7907	-0.0157	0.9769
CG8004	CG8004	1634133_at	-0.3919	0.0590	0.1872	0.4567	0.0099	0.9729	-0.2519	0.6615	-0.7491	0.0144	-0.4972	0.0485	-0.0474	0.9775	-0.0535	0.9341	-0.0061	0.9929
CG6114	SNF1-related	1634134_at	-0.1095	0.6581	-0.3141	0.1128	-0.4301	0.0584	-0.1840	0.8045	-0.0405	0.9223	0.1435	0.6369	-0.1814	0.8270	-0.3326	0.3178	-0.1512	0.6890
CG12012	CG12012	1634135_at	0.4668	0.0208	0.7925	0.0164	0.8852	0.0057	0.3091	0.4270	-0.2620	0.2212	-0.5710	0.0097	0.1738	0.8331	0.0502	0.9266	-0.1236	0.7548
CG2157	CG2157	1634136_at	-1.8577	0.0182	0.4127	0.4602	-0.6185	0.0725	-0.9739	0.1919	-1.9833	0.0018	-1.0093	0.0207	0.3092	0.8666	0.2107	0.8270	-0.0985	0.9224
---	---	1634137_at	0.0253	0.9144	0.0692	0.5514	0.3010	0.1472	0.0786	0.8852	-0.0143	0.9584	-0.0930	0.6214	-0.1227	0.8331	-0.0987	0.7228	0.0240	0.9434
CG7333	CG7333	1634138_at	-5.7845	0.0004	-7.3130	0.0011	-6.2344	0.0000	0.3645	0.7611	0.8116	0.1148	0.4471	0.3452	-0.6210	0.8062	-0.5428	0.6350	0.0782	0.9596
Cyp301a1	Cyp301a1	1634139_at	0.6140	0.7022	-2.6161	0.0110	-1.8401	0.0041	1.0567	0.3276	2.2913	0.0038	1.2346	0.0343	-0.0489	0.9959	-1.4517	0.5228	-1.4028	0.5444
CG11155	CG11155	1634140_a_at	-2.1641	0.0025	-1.6921	0.0321	-1.7811	0.0002	-0.4110	0.3328	-1.0202	0.0022	-0.6092	0.0133	-0.3003	0.8270	-0.6350	0.2489	-0.3347	0.5816
Parp	poly(ADP-ribose)	1634141_a_at	-0.3431	0.3598	0.0201	0.9337	-0.1662	0.5135	-0.3236	0.6749	-0.2811	0.4570	0.0424	0.9240	-0.1772	0.8270	0.0150	0.9811	0.1922	0.5988
CG6403	CG6403	1634142_at	0.2465	0.3196	0.1306	0.3712	0.3574	0.0380	0.1074	0.8764	-0.2222	0.3603	-0.3296	0.1200	0.0113	0.9914	-0.0622	0.8395	-0.0735	0.7876
Cyp6w1	Cyp6w1	1634143_at	2.4506	0.0384	2.0785	0.1677	2.8200	0.0004	-0.0296	0.9922	0.0795	0.9271	0.1090	0.8815	-0.7953	0.8505	-0.4037	0.8709	0.3916	0.8654
CG34365	CG12958	1634144_at	-0.1393	0.5870	0.0341	0.7634	-0.4879	0.0359	-0.1130	0.8449	0.0282	0.9243	0.1412	0.4876	0.2188	0.7726	0.0429	0.9350	-0.1760	0.6180
CG14968	CG14968	1634145_s_at	-0.8835	0.0069	1.2785	0.0607	0.0530	0.9243	-1.3000	0.1755	-2.0339	0.0047	-0.7340	0.1571	-0.0137	0.9939	0.1778	0.7050	0.1915	0.6734
CG15926	CG15926	1634146_at	-0.4616	0.1416	-0.1242	0.7321	-0.9384	0.0007	-0.2662	0.5293	-0.1357	0.5696	0.1305	0.5473	0.5755	0.5352	0.2710	0.5543	-0.3045	0.5038
homer	d-Homer	1634147_a_at	-0.6321	0.1976	0.1648	0.5180	0.0751	0.7081	-0.2271	0.7349	-0.2310	0.4561	-0.0038	0.9917	-0.0781	0.9689	0.5180	0.3127	0.5961	0.2749
CG4842	CG4842	1634148_at	-0.2303	0.6664	-0.2448	0.6419	-0.4177	0.1500	-0.1986	0.5854	-0.1617	0.3956	0.0369	0.8641	-0.2387	0.9308	-0.0865	0.9515	0.1522	0.9019

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
SMC2	SMC2	1634149_at	-0.0784	0.7828	-0.2925	0.0678	-0.1059	0.6494	-0.2977	0.5921	0.0066	0.9879	0.3044	0.2276	-0.5796	0.3056	-0.1245	0.7098	0.4551	0.1498
Hr78	X receptor at 78E/	1634150_s_at	-0.1176	0.5506	0.4194	0.1564	0.5011	0.0097	-0.0059	0.9956	-0.2749	0.2310	-0.2690	0.1896	-0.0868	0.9416	0.3953	0.2625	0.4822	0.2084
CG40251	CG40251	1634151_at	-0.0610	0.6936	-0.0029	0.9816	-0.0205	0.9275	-0.0219	0.9672	0.1095	0.4511	0.1314	0.2995	0.0009	0.9994	0.0247	0.9350	0.0237	0.9277
GstD5	Glutathione S tran	1634152_at	0.3895	0.3832	1.6594	0.0951	0.8252	0.0057	-0.2711	0.5376	0.4756	0.0443	0.7468	0.0039	0.3831	0.8882	1.5632	0.1297	1.1801	0.2668
CG14052	CG14052	1634153_at	0.0795	0.6972	0.1475	0.3155	0.2443	0.1377	-0.2672	0.6041	-0.2962	0.2557	-0.0290	0.9272	-0.1113	0.8494	-0.0450	0.9037	0.0662	0.8287
CG15570	CG15570	1634154_at	0.4902	0.0397	0.5158	0.0771	0.4912	0.0207	-0.0741	0.8908	-0.0701	0.7445	0.0041	0.9858	-0.0278	0.9860	0.0547	0.9216	0.0825	0.8597
CG3565	CG3565	1634155_at	0.0314	0.8831	0.1339	0.4399	-0.0035	0.9883	0.0327	0.9620	0.0614	0.7810	0.0286	0.8959	0.2097	0.7707	0.2004	0.5283	-0.0093	0.9849
CG5604	CG5604	1634156_at	0.0359	0.8758	-0.5714	0.0419	-0.6132	0.0355	0.0994	0.9011	0.8086	0.0079	0.7092	0.0086	0.1157	0.9032	0.1783	0.6344	0.0626	0.8949
Bap60	brahma associate	1634157_at	-0.5744	0.0584	0.4480	0.3344	0.8002	0.0057	0.0517	0.9339	-1.0687	0.0006	-1.1205	0.0003	-0.2929	0.8331	0.0758	0.9350	0.3687	0.5461
CG6164	CG6164	1634158_at	-4.6290	0.0003	-4.2566	0.0048	-3.7166	0.0002	-0.1074	0.9413	0.6711	0.1073	0.7785	0.0433	-0.7458	0.6749	0.9212	0.1985	1.6670	0.0607
CG31917	CG31917	1634159_s_at	-0.0449	0.9132	1.1399	0.0328	1.0192	0.0153	-0.0217	0.9884	-0.7507	0.0408	-0.7289	0.0293	0.0191	0.9933	0.3829	0.4692	0.3638	0.5003
---	---	1634160_at	0.0379	0.9095	0.0609	0.7714	0.1578	0.4511	0.1158	0.8967	-0.0570	0.8853	-0.1728	0.5486	-0.0956	0.8906	-0.1808	0.5041	-0.0852	0.7841
CG12314	CG12314	1634161_at	0.0885	0.8037	-0.0311	0.9058	0.3230	0.4889	0.0805	0.9339	0.1645	0.5983	0.0839	0.7932	-0.2537	0.9142	0.1946	0.8678	0.4483	0.6152
CG40198	CG40198	1634162_at	-0.2827	0.1960	0.1305	0.4734	-0.1465	0.6364	-0.0621	0.9094	-0.9472	0.0007	-0.8851	0.0006	0.1131	0.9132	-0.4906	0.1647	-0.6037	0.1289
---	---	1634163_at	0.0352	0.8430	0.0241	0.9099	0.1589	0.4684	-0.1069	0.8633	-0.1175	0.6333	-0.0106	0.9694	-0.1195	0.8331	0.0891	0.7500	0.2086	0.3878
SIP1	Syntaxin Interactir	1634164_at	0.5914	0.0081	0.5688	0.1648	0.2843	0.1379	0.0708	0.9090	0.2189	0.2694	0.1481	0.4225	0.1616	0.8049	0.0082	0.9887	-0.1534	0.6027
CG11378	CG11378	1634165_at	-0.2013	0.5137	0.4602	0.3150	0.5592	0.0719	-0.2287	0.7018	-0.8643	0.0076	-0.6356	0.0177	-0.2483	0.8903	-0.0941	0.9296	0.1542	0.8618
---	---	1634166_at	0.1501	0.6508	0.2927	0.3769	0.1073	0.5489	-0.1683	0.8084	-0.2398	0.3973	-0.0715	0.8191	0.0791	0.9330	-0.0357	0.9411	-0.1148	0.7418
sage	salivary gland-exp	1634167_a_at	-0.0432	0.9196	0.2513	0.6189	0.0143	0.9729	-0.1444	0.8707	0.0637	0.8775	0.2080	0.4818	0.2751	0.7697	0.4247	0.2707	0.1496	0.7448
CG5515	CG5515	1634168_s_at	-0.0763	0.6299	0.5075	0.0216	0.8041	0.0069	-0.0111	0.9872	-0.3936	0.0252	-0.3825	0.0177	-0.1641	0.8903	0.3649	0.4094	0.5290	0.2528
---	---	1634169_at	0.1287	0.5467	0.0712	0.5548	-0.0741	0.6891	-0.1312	0.8038	-0.0830	0.7390	0.0481	0.8434	0.1232	0.8553	-0.1246	0.6876	-0.2478	0.3820
jbug	filamin	1634170_a_at	-2.4917	0.0661	-2.0510	0.1302	-2.0131	0.0012	0.5617	0.4815	0.4659	0.2782	-0.0959	0.8463	0.4755	0.9331	0.9999	0.6059	0.5245	0.8171
CG12231	CG12231	1634171_at	0.1571	0.4588	0.0972	0.4115	0.1812	0.3531	-0.1664	0.6578	0.1257	0.5097	0.2921	0.0722	-0.1972	0.7611	-0.1643	0.5636	0.0329	0.9316
TfIIB	Transcription factc	1634172_at	0.1960	0.2848	0.3168	0.1904	-0.1098	0.4392	-0.1884	0.6368	-0.2018	0.3073	-0.0134	0.9573	0.2149	0.7095	-0.0604	0.8551	-0.2752	0.2707
Trap1	Hsp90-related pro	1634173_at	-0.1011	0.5740	-0.4492	0.2730	-0.2332	0.1559	-0.0482	0.9313	0.0015	0.9951	0.0497	0.7874	-0.2403	0.7947	-0.3199	0.4041	-0.0797	0.8800
CG9702	CG9702	1634174_at	-0.2045	0.5017	-0.0003	1.0000	-0.0863	0.6344	-0.1538	0.8281	-0.3839	0.1573	-0.2301	0.3587	-0.0417	0.9589	-0.1119	0.6536	-0.0702	0.8004
Mctp	Multiple C2 domai	1634175_at	0.1359	0.4136	0.1166	0.4214	-0.0481	0.7819	0.0629	0.9422	0.0351	0.9193	-0.0278	0.9267	0.0971	0.9095	-0.0269	0.9577	-0.1240	0.7192
MED9	Mediator complex	1634176_a_at	-0.3319	0.0836	-0.2719	0.1089	0.0469	0.8429	0.0842	0.8605	-0.1547	0.3790	-0.2390	0.1191	-0.1076	0.8609	-0.1763	0.4861	-0.0687	0.8247
CG31055	CG31055	1634177_a_at	-0.8124	0.0279	-0.4987	0.1141	-1.4724	0.0042	-0.8597	0.3952	-0.0392	0.9640	0.8206	0.1058	0.2005	0.7220	0.1477	0.5591	-0.0529	0.8695
CG4351	CG4351	1634178_at	0.2922	0.2794	-0.0513	0.7555	-0.4006	0.1215	-0.5696	0.2020	0.1972	0.4844	0.7668	0.0071	-0.3336	0.7266	-0.1793	0.7000	0.1543	0.7447
CG34139	CG34139	1634179_at	0.2211	0.3224	0.0105	0.9203	-0.3089	0.1571	-0.1393	0.7949	0.0378	0.8994	0.1771	0.3815	0.0493	0.9564	-0.1672	0.5374	-0.2165	0.4114
CG9664	CG9664	1634180_s_at	0.2896	0.4578	-0.5375	0.0374	-0.2688	0.2110	-0.1210	0.9120	0.3987	0.2534	0.5197	0.0947	-0.4922	0.6695	-0.6536	0.1498	-0.1614	0.7663
CG11203	CG11203	1634181_at	-0.0793	0.6713	-0.0546	0.8447	-0.1395	0.4995	-0.0148	0.9857	-0.0485	0.8425	-0.0338	0.8800	0.1246	0.8745	-0.1279	0.7199	-0.2525	0.4275
CG11593	CG11593	1634182_at	0.0472	0.8034	-0.3926	0.0602	-0.3076	0.2952	0.1112	0.8545	0.6050	0.0135	0.4938	0.0198	0.0211	0.9914	0.2397	0.6232	0.2186	0.6567
CG4374	CG4374	1634183_at	0.1301	0.6029	0.0057	0.9745	0.2307	0.3357	0.0958	0.9252	0.0133	0.9780	-0.0825	0.8109	-0.2065	0.6955	-0.0434	0.8922	0.1631	0.4683
jp	junctophilin	1634184_a_at	-2.2800	0.0010	-2.0623	0.0155	-3.0900	0.0000	-0.1491	0.8143	0.2911	0.2455	0.4402	0.0535	0.3216	0.7230	0.0454	0.9429	-0.2762	0.4978
CG32053	CG32053	1634185_at	-0.0317	0.9280	-0.1364	0.4032	-0.1213	0.4954	-0.1485	0.8194	-0.1924	0.4703	-0.0439	0.8845	0.0127	0.9928	-0.1909	0.5922	-0.2037	0.5672
CG17119	CG17119	1634186_a_at	0.4586	0.7508	-0.2241	0.6688	-0.1988	0.6680	0.2902	0.9256	0.5793	0.5672	0.2891	0.7821	0.0807	0.9914	-0.4247	0.8445	-0.5054	0.7931
---	---	1634187_x_at	-1.1991	0.1854	-1.1729	0.0302	-0.9515	0.4406	0.1329	0.9627	0.2131	0.8218	0.0802	0.9313	-0.2284	0.9767	0.2230	0.9407	0.4515	0.8515
tim	TIMELESS	1634188_a_at	1.1685	0.0144	0.1793	0.1782	0.0084	0.9755	-0.8971	0.0206	-0.1818	0.3904	0.7153	0.0029	-0.6769	0.5126	-1.1341	0.0418	-0.4572	0.3640
CG10338	CG10338	1634189_at	-0.0497	0.8307	0.4014	0.1746	0.4971	0.0644	-0.1033	0.8650	-0.0643	0.8092	0.0390	0.8786	-0.0862	0.9380	0.4661	0.1647	0.5523	0.1390
CG6138	CG6138	1634190_a_at	0.0863	0.6697	0.0049	0.9785	0.3182	0.1343	0.0548	0.9374	0.0218	0.9404	-0.0330	0.8915	-0.1994	0.6749	-0.0641	0.7942	0.1353	0.5041
CG6398 /// DyakCG6398	CG6398	1634191_at	0.0840	0.7529	0.9917	0.0544	1.1912	0.0008	-0.1559	0.7135	-0.8411	0.0018	-0.6851	0.0027	-0.3366	0.7689	0.2431	0.6458	0.5796	0.2523
CG30281	CG30281	1634192_at	2.7231	0.0070	1.1406	0.1820	2.9445	0.0007	1.5794	0.2438	0.6114	0.4661	-0.9680	0.1769	-0.3836	0.8692	-1.0328	0.2501	-0.6492	0.5007
CG13142	CG13142	1634193_at	0.3586	0.2699	0.3103	0.4102	0.1210	0.7656	-0.5031	0.2943	-0.3736	0.1831	0.1295	0.6578	-0.1206	0.9571	-0.3188	0.6449	-0.1983	0.7967
snf	splicing necessary	1634194_at	-0.1051	0.6357	0.1467	0.6112	0.2871	0.1292	0.0636	0.9110	-0.2050	0.2572	-0.2686	0.0953	-0.2293	0.8141	-0.0285	0.9655	0.2008	0.6428
CG11093 /// DsimCG11093	CG11093	1634195_at	-0.2221	0.4564	-0.0757	0.7793	-0.3789	0.1498	-0.0455	0.9647	0.0780	0.8174	0.1234	0.6615	0.1838	0.8521	0.0727	0.9075	-0.1111	0.8286
CG9777	CG9777	1634196_at	-0.2851	0.4119	-0.2961	0.0574	-0.0492	0.8162	0.0913	0.8841	-0.0582	0.8306	-0.1495	0.4599	-0.0815	0.9657	0.0044	0.9979	0.0859	0.9032
CG11576	CG11576	1634197_at	-1.2857	0.0034	-0.3110	0.4262	-0.1840	0.3353	-0.1355	0.7621	-0.8165	0.0020	-0.6810	0.0027	-0.3612	0.7387	0.0037	0.9980	0.3649	0.4393
lwf	lots wife	1634198_at	0.1200	0.4374	0.0271	0.8661	0.0213	0.9113	-0.0060	0.9937	0.0155	0.9459	0.0215	0.9086	-0.0580	0.9390	-0.1119	0.6616	-0.0540	0.8600
CG15220	CG15220	1634199_at	-0.0620	0.8382	-0.6901	0.0681	-0.2893	0.3880	-0.0172	0.9895	0.5766	0.0609	0.5938	0.0354	-0.4199	0.7220	-0.0298	0.9719	0.3901	0.4524
CG9413	CG9413	1634200_a_at	-2.5244	0.0010	-1.9704	0.0303	-2.6223	0.0005	-0.1507	0.7850	-0.1689	0.4778	-0.0182	0.9496	0.4712	0.8283	0.3054	0.7886	-0.1658	0.8947
CG13841	CG13841	1634201_at	-0.7322	0.0636	0.0496	0.6750	-0.4275	0.1139	-0.3011	0.6257	-0.6925	0.0308	-0.3914	0.1506	-0.1014	0.8541	-0.1011	0.6915	0.0003	0.9992
---	---	1634202_at	0.1911	0.3249	0.0336	0.7689	0.0067	0.9791	0.1252	0.7278	0.1266	0.4491	0.0014	0.9943	0.1486	0.8236	0.0667	0.8636	-0.0819	0.8123
CG17377	CG17377	1634203_at	0.1296	0.5109	0.0056	0.9810	0.0444	0.8242	0.0137	0.9860	0.0182	0.9470	0.0045	0.9850	0.0163	0.9862	-0.1249	0.5897	-0.1412	0.5403
Dh /// p16-ARC	p16-ARC /// CRF-	1634204_at	-0.2054	0.2649	-0.0464	0.6425	0.0969	0.6423	0.1852	0.5311	0.2930	0.0615	0.1079	0.4607	0.0554	0.9651	0.4750	0.1498	0.4196	0.2298
CG15254	CG15254	1634205_at	-0.3469	0.9210	-0.0810	0.6505	-0.2154	0.4367	-0.0361	0.9956	-2.0162	0.2888	-1.9801	0.2419	0.0479	0.9964	-1.7223	0.5329	-1.7702	0.5257
CG2124 /// DereCG2124 ///	CG2124	1634206_at	0.1012	0.8167	-0.0173	0.9533	-0.2923	0.2235	-0.4577	0.3294	0.2178	0.4393	0.6754	0.0132	-0.0295	0.9898	0.2306	0.6764	0.2601	0.6282
CG16799 /// DmirCG16799	CG16799 /// GA1/	1634207_at	-1.3062	0.0017	-1.9860	0.0164	-1.6370	0.0003	0.3283	0.2596	0.0010	0.9966	-0.3273	0.0404	0.0336	0.9898	-0.8077	0.1345	-0.8413	0.1459
plx	pollux	1634208_a_at	-1.7288	0.0005	-2.0555	0.0128	-2.7822	0.0000	-0.4945	0.1332	-0.9394	0.0013	-0.4449	0.0204	0.0512	0.9764	-1.2835	0.0222	-1.3347	0.0320
pim	pimples	1634209_at	-0.6725	0.3526	-1.9493	0.1121	-2.0829	0.0210	-0.2383	0.8532	1.2598	0.0148	1.4981	0.0041	0.0250	0.9976	-0.0122	0.9988	-0.0372	0.9902
CG15649	CG15649	1634210_at	0.1444	0.4219	0.0953	0.4947	0.1272	0.4931	0.0697	0.8732	0.0235	0.9114	-0.0462	0.7854	-0.0531	0.9309	-0.0298	0.9233	0.0233	0.9338
arr	LDLR-like	1634211_at	0.3671	0.3888	0.4313	0.1485	0.0318	0.9325	-0.0973	0.9149	0.0964	0.7799	0.1937	0.4754	0.1517	0.9450	0.1138	0.9115	-0.0379	0.9704
CG6905	CG6905	1634212_at	-0.1060	0.7933	-0.1955	0.5540	0.1031	0.5007	0.1865	0.6968	0.2111	0.3467	0.0246	0.9274	-0.0365	0.9893	0.1400	0.8583	0.1765	0.7964
CG13139	CG13139	1634213_at	-0.9421	0.0196	-0.5840	0.1353	-0.5132	0.0887	0.3424	0.5255	-0.2546	0.3781	-0.5970	0.0263	0.1965	0.8446	0.1407	0.7824	-0.0558	0.9216
Ctr1C	Copper transporte	1634214_a_at	-0.1313	0.5579	-0.0275	0.8659	0.0661	0.7284	0.2225	0.4962	0.1241	0.4954	-0.0984	0.5666	-0.0220	0.9885	0.0777	0.8685	0.0997	0.8078
---	---	1634215_at	-0.0611	0.6908	-0.0015	0.9913	-0.0543	0.7479	-0.0188	0.9803	0.0253	0.9223	0.0441	0.8353	-0.0017	0.9992	-0.0065	0.9901	-0.0048	0.9902
Adgf-C	Adenosine deamir	1634216_at	-0.4757	0.3523	-0.2144	0.4022	-0.7679	0.0634	-0.5793	0.4998	-1.0948	0.0239	-0.5155	0.2032	-0.0056	0.9989	-0.8602	0.1522	-0.8547	0.1911
Crk	Crk	1634217_s_at	-0.3175	0.1593	0.1017	0.3258	-0.1427	0.3187	-0.2089	0.6504	-0.2801	0.2060	-0.0712	0.7720	0.0525	0.9489	0.1130	0.6773	0.0605	0.8472

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG17270	CG17270	1634218_at	-0.2319	0.2929	-1.0224	0.0280	-1.2292	0.0010	-0.2911	0.4568	0.4034	0.0611	0.6945	0.0038	-0.1318	0.8882	-0.5701	0.1114	-0.4383	0.2287
Jafrac1	thioredoxin peroxi	1634219_a_at	-0.2006	0.4619	-0.7388	0.0287	-0.6327	0.0070	0.2396	0.6580	0.6695	0.0183	0.4299	0.0676	0.0496	0.9683	0.1211	0.7579	0.0715	0.8709
CG13362	CG13362	1634220_at	-0.0015	0.9973	-0.3857	0.1654	-0.4031	0.3066	0.1386	0.9302	0.6496	0.1600	0.5110	0.2187	-0.0595	0.9626	-0.0367	0.9492	0.0228	0.9655
CG13632	CG13632	1634221_at	0.0223	0.9105	0.2766	0.2109	0.1254	0.4885	0.0875	0.8584	-0.1577	0.3826	-0.2452	0.1191	0.2515	0.6898	0.1031	0.7323	-0.1484	0.5929
CG18596	CG18596	1634222_at	0.2710	0.2529	-0.1010	0.7101	-0.5331	0.0897	-0.2971	0.5754	0.7348	0.0148	1.0319	0.0019	0.0812	0.9653	0.4391	0.3718	0.3579	0.4888
---	---	1634223_at	-0.0198	0.9309	0.1592	0.5093	-0.0220	0.9065	-0.0836	0.8897	-0.1777	0.3909	-0.0940	0.6524	-0.1161	0.8608	-0.1081	0.7245	0.0080	0.9849
CG9997	CG9997	1634224_at	0.1013	0.5506	0.0410	0.7921	0.2874	0.1562	0.1636	0.7303	-0.0040	0.9899	-0.1676	0.3959	-0.1589	0.8042	-0.0889	0.7897	0.0700	0.8355
CG13193	CG13193	1634225_at	0.1074	0.7049	-0.3704	0.1706	-0.2500	0.1343	-0.1159	0.8315	0.0643	0.8006	0.1802	0.3438	-0.1472	0.8298	-0.3163	0.2518	-0.1691	0.5803
CG17249	CG17249	1634226_at	0.0749	0.7785	0.1705	0.6169	0.2725	0.2252	0.1290	0.8155	0.1577	0.4886	0.0287	0.9137	0.0452	0.9774	0.3501	0.3366	0.3048	0.4266
CG9517	CG9517	1634227_a_at	-1.3995	0.0302	-0.9056	0.0395	-1.7279	0.0042	0.0132	0.9956	-0.3316	0.6507	-0.3448	0.5973	0.0930	0.9390	-0.2903	0.4372	-0.3833	0.3163
---	---	1634228_at	0.1946	0.2620	-0.0358	0.8553	0.2158	0.2714	-0.0611	0.8979	0.2002	0.1990	0.2613	0.0651	-0.1238	0.8978	-0.0418	0.9425	0.0821	0.8618
CG8713	CG8713	1634229_at	-2.4112	0.0034	-2.4321	0.0074	-2.6194	0.0000	0.0618	0.9635	0.5431	0.1270	0.4813	0.1299	0.1593	0.8973	0.4022	0.3738	0.2429	0.6254
grp	grapes	1634230_s_at	0.1818	0.5340	-0.2508	0.4170	-0.3981	0.1071	-0.1435	0.7781	0.5755	0.0146	0.7190	0.0032	-0.1695	0.9030	0.0764	0.9231	0.2459	0.6533
---	---	1634231_at	0.0320	0.8807	0.0579	0.7072	-0.1609	0.6099	-0.0988	0.8408	-0.1016	0.6163	-0.0029	0.9901	-0.0281	0.9816	0.0325	0.9425	0.0606	0.8717
mRpL13	mitochondrial ribo	1634232_at	-0.2000	0.5777	-0.0484	0.8760	-0.7047	0.0037	-0.2221	0.6130	-0.0643	0.8136	0.1578	0.4479	0.4485	0.7187	0.0043	0.9980	-0.4442	0.3977
---	---	1634233_at	0.1451	0.3594	0.0130	0.9375	0.0772	0.7356	0.0258	0.9758	0.0283	0.9226	0.0025	0.9921	-0.0887	0.8972	-0.0828	0.7985	0.0058	0.9886
---	---	1634234_s_at	0.0303	0.8927	0.2397	0.3760	0.1905	0.2319	-0.0129	0.9860	-0.1561	0.3779	-0.1432	0.3703	0.0230	0.9871	-0.0097	0.9887	-0.0327	0.9431
CG34388	CG14842	1634235_at	0.1382	0.6618	0.0201	0.8507	-0.0047	0.9816	0.2510	0.6086	0.2134	0.3983	-0.0375	0.8992	0.0216	0.9901	-0.0317	0.9601	-0.0532	0.9197
CG32815	CG32815	1634236_at	0.4016	0.2050	0.4046	0.2228	0.2136	0.3670	0.0137	0.9937	-0.0877	0.8500	-0.1015	0.8007	0.2509	0.7726	0.0536	0.9277	-0.1973	0.6257
l(2)34Fa	lethal (2) 34Fa	1634237_at	1.9618	0.0051	0.3683	0.2444	1.1863	0.0005	0.6913	0.3171	1.1561	0.0114	0.4648	0.1967	0.0074	0.9970	-0.2302	0.6519	-0.2376	0.6389
CG9628	CG9628	1634238_a_at	0.5561	0.1333	0.0812	0.8378	1.1628	0.0005	-0.0709	0.9060	-0.7284	0.0034	-0.6575	0.0032	-1.1305	0.3162	-1.2762	0.0519	-0.1457	0.8477
CG14205	CG14205	1634239_at	0.0056	0.9847	-0.0071	0.9497	0.3482	0.1837	0.1221	0.8057	-0.0193	0.9479	-0.1414	0.4481	-0.1502	0.8298	-0.0193	0.9682	0.1308	0.6863
CG5107	CG5107	1634240_at	1.0282	0.1355	-0.3996	0.3926	-0.4089	0.1691	0.0779	0.9182	0.8203	0.0046	0.7423	0.0043	0.0527	0.9913	-0.7833	0.4351	-0.8360	0.4116
CG4860	CG4860	1634241_at	0.7019	0.0433	0.2929	0.4292	0.9467	0.0846	0.6077	0.5128	0.6739	0.1671	0.0662	0.9111	-0.3143	0.8870	0.1388	0.9139	0.4531	0.6128
---	---	1634242_at	-0.0874	0.6837	-0.0263	0.8306	0.1131	0.5746	-0.0852	0.9011	-0.1315	0.5869	-0.0463	0.8585	-0.1851	0.8049	-0.0274	0.9582	0.1576	0.6395
CG4673	CG4673	1634243_a_at	-0.3015	0.0695	0.1755	0.3458	0.2041	0.1663	0.0484	0.9339	-0.4388	0.0187	-0.4873	0.0072	0.0050	0.9952	0.1593	0.4192	0.1543	0.4504
CG31639	CG31639	1634244_s_at	0.1425	0.5334	0.1307	0.5811	0.1310	0.4435	-0.0605	0.9362	0.0435	0.8836	0.1040	0.6485	-0.1578	0.8076	-0.0767	0.8304	0.0811	0.8065
---	---	1634245_at	-0.3892	0.0560	0.0098	0.9669	0.0463	0.8324	-0.0272	0.9774	-0.0453	0.8869	-0.0181	0.9517	0.1754	0.8680	0.4749	0.2389	0.2996	0.4869
CG11319	CG11319	1634246_at	1.8920	0.0413	2.4808	0.0165	1.4297	0.0077	-0.0414	0.9873	-1.3641	0.0355	-1.3226	0.0257	0.8929	0.7142	-0.7608	0.4570	-1.6537	0.1335
---	---	1634247_at	0.1523	0.3799	-0.0690	0.5296	-0.1727	0.3328	0.0457	0.9339	0.3218	0.0516	0.2762	0.0601	0.1111	0.8655	0.0612	0.8684	-0.0499	0.8897
---	---	1634248_at	-0.1886	0.4869	-0.3400	0.6564	-0.1114	0.6274	-0.1884	0.8000	0.3128	0.3015	0.5012	0.0649	-0.2522	0.9063	0.3501	0.6824	0.6023	0.4457
Pdsw	Pdsw	1634249_s_at	-0.1445	0.5693	0.1209	0.7783	-0.1246	0.6886	-0.0604	0.9562	-0.6410	0.0378	-0.5806	0.0360	0.1856	0.8999	-0.3973	0.4699	-0.5829	0.2960
Act57B	Actin 57B	1634250_at	-0.6129	0.0668	-0.9828	0.0559	-1.0540	0.0009	-0.0973	0.7981	-0.1245	0.4361	-0.0271	0.8815	-0.0202	0.9943	-0.4651	0.4868	-0.4449	0.5145
CG12853	CG12853	1634251_at	0.1129	0.4256	0.1277	0.3882	0.0649	0.7071	-0.1167	0.7322	-0.1129	0.4779	0.0039	0.9844	0.0480	0.9498	0.1136	0.6429	0.0657	0.8161
---	---	1634252_at	0.1076	0.6009	-0.0214	0.8817	0.0457	0.8009	0.0302	0.9705	0.1145	0.6236	0.0843	0.7035	0.0984	0.8599	0.1071	0.6645	0.0088	0.9818
CG9917	CG9917	1634253_at	-0.6640	0.0216	-0.4748	0.1391	-0.5606	0.0187	-0.2189	0.5539	-0.0537	0.8223	0.1652	0.3466	-0.1202	0.9296	0.0307	0.9663	0.1509	0.7713
CG15055	CG15055	1634254_at	0.2101	0.4147	0.1728	0.5339	0.1467	0.5199	-0.0985	0.9060	-0.1399	0.6430	-0.0414	0.8982	-0.0061	0.9964	-0.0330	0.9541	-0.0269	0.9586
CG12721	CG12721	1634255_at	0.2296	0.1153	-0.2982	0.4718	-0.4830	0.0220	-0.2993	0.4551	0.4799	0.0341	0.7792	0.0025	-0.1963	0.8160	-0.0779	0.8760	0.1184	0.7734
stg1	CG33670	1634256_a_at	-0.0672	0.6980	0.1197	0.7820	0.1107	0.7449	0.1439	0.7121	0.2067	0.2439	0.0627	0.7423	0.0916	0.9689	0.2337	0.7459	0.1422	0.8599
CG5162	CG5162	1634257_at	1.9686	0.0011	1.1741	0.0925	1.7258	0.0002	0.3100	0.5285	0.1382	0.6292	-0.1718	0.4838	-0.3375	0.8283	-0.6455	0.3087	-0.3080	0.6630
mod(mdg4)	Modifier67.2	1634258_at	-0.6404	0.0197	-0.1115	0.8292	0.1197	0.4915	-0.2622	0.6666	-0.2393	0.4253	0.0228	0.9503	-0.4961	0.6749	0.2390	0.6479	0.7351	0.1528
CG9083	CG9083	1634259_at	0.3858	0.2070	0.0356	0.7740	0.2485	0.3484	0.0693	0.9317	0.0057	0.9879	-0.0636	0.8151	0.1109	0.8815	-0.0232	0.9613	-0.1341	0.6652
---	---	1634260_at	0.1513	0.3080	0.0685	0.6710	0.1545	0.4341	-0.0235	0.9777	0.0962	0.6983	0.1197	0.5769	0.1388	0.8608	-0.0271	0.9590	-0.1659	0.6289
Cyp312a1	Cyp312a1	1634261_at	1.8567	0.0387	3.4557	0.0091	2.1823	0.0007	-0.9805	0.4455	-2.3120	0.0061	-1.3314	0.0398	0.2001	0.9445	-0.7999	0.3437	-0.9999	0.2651
CycK	cyclin K	1634262_s_at	-0.1455	0.4432	0.6165	0.0096	0.8120	0.0005	0.0897	0.8532	-0.6152	0.0047	-0.7049	0.0015	0.0551	0.9390	0.3260	0.1392	0.2709	0.2388
lectin-21Cb	Lectin21Cb	1634263_at	0.0642	0.7169	0.0436	0.7324	0.2670	0.1841	0.0689	0.9200	-0.1266	0.5768	-0.1955	0.3034	-0.0511	0.9515	0.0526	0.8853	0.1038	0.7154
CG13202	CG13202	1634264_at	0.0756	0.5962	-0.0420	0.8423	0.0023	0.9916	0.0194	0.9774	-0.0588	0.7722	-0.0782	0.6540	0.0885	0.8960	0.0098	0.9841	-0.0787	0.7964
Acp95EF	Accessory gland-s	1634265_at	0.5242	0.0404	0.2594	0.4973	-0.1365	0.7282	-0.0063	0.9962	0.8407	0.0594	0.8470	0.0375	0.2342	0.8270	0.3555	0.4268	0.1213	0.8299
---	---	1634266_at	-0.0007	0.9973	0.0834	0.6379	0.1978	0.2254	0.0398	0.9649	-0.0336	0.9215	-0.0735	0.7868	-0.1359	0.8738	-0.0684	0.8886	0.0675	0.8834
CG30495	CG30495	1634267_at	-0.2744	0.2738	-0.2177	0.4112	-0.3234	0.0725	-0.0579	0.9017	-0.1126	0.4789	-0.0547	0.7379	-0.0365	0.9845	-0.1580	0.7499	-0.1215	0.8162
---	---	1634268_at	0.0448	0.8803	0.0101	0.9731	-0.0507	0.8531	0.0720	0.9314	0.2324	0.3600	0.1604	0.5027	0.0776	0.9521	0.1689	0.7002	0.0913	0.8572
CG13784	CG13784	1634269_at	-0.0861	0.8330	0.5376	0.0558	0.3272	0.2766	-0.3352	0.3443	-0.5166	0.0197	-0.1814	0.3312	-0.1182	0.9495	0.0870	0.9194	0.2052	0.7484
CG5037	CG5037	1634270_at	-1.1606	0.0080	-0.2708	0.2762	-0.1416	0.5396	-0.2547	0.4455	-1.1627	0.0004	-0.9080	0.0006	-0.1517	0.8953	-0.0979	0.8735	0.0538	0.9282
Def	defensin	1634271_at	-3.0689	0.0008	-2.9358	0.0179	-2.1285	0.0012	0.0042	0.9956	-0.0458	0.8906	-0.0500	0.8612	-0.7124	0.7506	0.1637	0.9120	0.8761	0.3757
CG18516	CG18516	1634272_at	0.0685	0.7322	0.0407	0.6935	0.0069	0.9841	-0.0934	0.8578	-0.0368	0.8842	0.0566	0.7880	0.0955	0.9340	-0.0572	0.9204	-0.1527	0.7134
---	---	1634273_s_at	0.0974	0.5558	0.2048	0.3398	0.0588	0.8599	-0.1114	0.7981	-0.0857	0.6693	0.0257	0.9036	0.1809	0.8235	0.0731	0.8792	-0.1078	0.7884
CG14305	CG14305	1634274_s_at	-0.1012	0.7032	-0.0051	0.9794	0.2714	0.1571	-0.0364	0.9482	0.0344	0.8685	0.0708	0.6637	-0.1600	0.8395	0.0935	0.8280	0.2534	0.4507
---	---	1634275_s_at	0.2965	0.2544	-0.0180	0.8883	-0.2415	0.1498	-0.1351	0.6652	-0.1550	0.3030	-0.0199	0.9124	0.1250	0.8192	-0.1782	0.4246	-0.3033	0.1990
Adar	Adenosine deamir	1634276_a_at	-0.3857	0.1831	0.3570	0.4163	0.3233	0.1555	-0.1408	0.8903	-0.3745	0.2760	-0.2337	0.4724	0.1546	0.8882	0.5764	0.1512	0.4218	0.3181
CG1602	CG1602	1634277_at	0.2270	0.2426	-0.2218	0.4856	-0.2086	0.2874	-0.0550	0.9487	0.5900	0.0216	0.6449	0.0090	0.0060	0.9967	0.0448	0.9402	0.0388	0.9394
CG6409	CG6409	1634278_at	0.8505	0.6351	-0.8361	0.4058	-1.1663	0.0262	0.7065	0.4500	-0.1075	0.8731	-0.8140	0.0747	1.1338	0.8477	-1.8082	0.4676	-2.9420	0.2485
CG11529	CG11529	1634279_at	-0.3078	0.3668	-0.5735	0.0195	-1.1663	0.0002	-0.1351	0.8507	0.7793	0.0095	0.9144	0.0028	0.0735	0.9405	0.1731	0.5907	0.0996	0.7815
---	---	1634280_at	0.1681	0.4671	0.0762	0.5190	0.1994	0.1590	0.1372	0.7550	0.1672	0.3896	0.0299	0.8955	0.0631	0.9309	0.1112	0.6701	0.0481	0.8813
CG12343	CG12343	1634281_at	-0.3009	0.1373	-0.1750	0.3223	-0.2686	0.1253	-0.2119	0.6360	-0.3726	0.0911	-0.1607	0.4369	-0.0659	0.9330	-0.2297	0.3474	-0.1639	0.5358
CR32745 /// Top3beta	Topoisomerase 3	1634282_s_at	0.0025	0.9969	0.7451	0.1841	0.6299	0.0039	0.0514	0.9649	-0.4908	0.1142	-0.5422	0.0558	0.0154	0.9952	0.2059	0.8045	0.1905	0.8129
CG3921	CG3921	1634283_at	0.1959	0.5250	0.4945	0.1648	0.1999	0.3113	0.1578	0.8073	0.1451	0.6078	-0.0128	0.9688	0.6344	0.5754	0.4679	0.3470	-0.1665	0.7836
CG14639 /// DyakCG1463	CG14639	1634284_at	0.6577	0.2290	0.6629	0.0351	0.7510	0.0015	-0.1362	0.7030	-0.2982	0.0725	-0.1620	0.2755	-0.3088	0.8069	-0.3930	0.4592	-0.0843	0.9080
CheB93b	CheB93b	1634285_at	0.0253	0.8904	-0.0731	0.5074	0.0731	0.7289	0.1370	0.8140	0.1745	0.4661	0.0375	0.8913	-0.0757	0.9168	0.0885	0.7730	0.1643	0.5355
CG15554	CG15554	1634286_at	-0.0548	0.8530	-0.0399	0.7022	-0.1660	0.4191	-0.0055	0.9956	-0.0162	0.9674	-0.0108	0.9727	-0.0223	0.9895	-0.1421	0.7300	-0.1198	0.7749

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13737	CG13737	1634287_at	0.6231	0.0149	0.1799	0.2894	0.7392	0.0012	0.0835	0.9182	-0.1745	0.5078	-0.2580	0.2493	-0.1699	0.7697	-0.1323	0.6165	0.0376	0.9121
---	---	1634288_at	-0.1939	0.5163	-1.4167	0.1012	-1.1123	0.0081	-0.1087	0.9422	1.0859	0.0192	1.1946	0.0077	-0.3106	0.8795	-0.2763	0.7734	0.0343	0.9783
rb	ruby	1634289_at	-0.0788	0.8272	0.0405	0.7825	-0.3396	0.0425	-0.1061	0.8073	0.1917	0.2698	0.2978	0.0581	0.2865	0.7707	0.3453	0.4051	0.0588	0.9192
mew	PS1 integrin	1634290_a_at	0.2049	0.6868	2.2295	0.0269	1.0839	0.0152	-0.6093	0.3744	-0.8080	0.0433	-0.1986	0.6125	0.4763	0.8192	1.2258	0.1423	0.7495	0.3898
CG11814 /// DmirCG11814	CG11814	1634291_at	-0.2729	0.5911	0.0437	0.9599	0.4464	0.1175	0.2149	0.8000	-0.2916	0.4073	-0.5064	0.0973	-0.1617	0.9598	0.0266	0.9893	0.1883	0.8797
CG15234	CG15234	1634292_at	-0.1145	0.5181	-0.0057	0.9739	-0.1057	0.4835	-0.0639	0.8875	-0.0527	0.7741	0.0112	0.9542	0.2682	0.5954	0.1772	0.4242	-0.0910	0.7167
---	---	1634293_at	0.1487	0.3655	0.2071	0.1822	0.4038	0.0254	-0.0528	0.9452	0.1016	0.6850	0.1544	0.4565	-0.2442	0.7220	0.1340	0.6834	0.3781	0.2149
AP-1gamma	AP-1gamma	1634294_s_at	-0.1418	0.7769	0.8898	0.1091	1.1914	0.0014	-0.0944	0.9457	-0.1669	0.7141	-0.0725	0.8766	-0.4325	0.8331	0.8523	0.3043	1.2848	0.1573
hoe2	hoepel2	1634295_at	-2.3644	0.0024	-0.9159	0.1163	-1.4355	0.0001	-0.2796	0.8251	-1.6186	0.0061	-1.3390	0.0086	0.3610	0.7478	-0.0678	0.9279	-0.4289	0.3807
CG17646	CG17646	1634296_s_at	1.6919	0.0030	0.5829	0.3542	1.1772	0.0009	-0.0074	0.9943	0.2315	0.2801	0.2389	0.2099	-0.5825	0.7541	-0.7610	0.3245	-0.1785	0.8643
Cyp318a1	Cyp318a1	1634297_at	-0.2292	0.7475	0.1002	0.4947	0.5255	0.0426	-0.0890	0.9639	-1.1413	0.0358	-1.0523	0.0317	-0.4216	0.7953	-0.8707	0.1875	-0.4491	0.5304
CG12375	CG12375	1634298_at	0.0858	0.7067	0.0649	0.7087	0.1938	0.1966	0.0066	0.9937	-0.2954	0.0937	-0.3020	0.0589	-0.1396	0.8541	-0.2937	0.3358	-0.1542	0.6476
CG7810	CG7810	1634299_at	0.0332	0.9002	0.5524	0.0401	1.1393	0.0086	-0.0587	0.9530	-0.3837	0.1607	-0.3250	0.1849	-0.5466	0.6496	0.2383	0.6531	0.7849	0.1362
CG4025	CG4025	1634300_at	-0.2157	0.4134	0.0391	0.7938	0.8148	0.0046	0.3575	0.3301	-0.3444	0.1042	-0.7019	0.0036	-0.3242	0.7464	-0.0289	0.9664	0.2953	0.4978
CG14017	CG14017	1634301_at	-0.3913	0.0217	0.1306	0.5018	-0.2792	0.2414	-0.0987	0.8816	-0.3274	0.1459	-0.2287	0.2614	0.2456	0.7644	0.2256	0.5300	-0.0200	0.9701
CG14516	CG14516	1634302_s_at	-1.1353	0.0129	-1.4335	0.0730	-1.4424	0.0008	0.3074	0.5407	0.9572	0.0039	0.6498	0.0133	0.3070	0.8692	0.7001	0.3342	0.3932	0.6233
hpo	hippo	1634303_at	0.1686	0.3828	0.0039	0.9903	-0.1680	0.2623	-0.2437	0.4861	-0.0612	0.7884	0.1825	0.2792	0.0374	0.9777	-0.0260	0.9613	-0.0634	0.8853
CG17233	CG17233	1634304_a_at	-0.2051	0.5228	0.0074	0.9884	0.0205	0.9231	-0.0056	0.9956	0.0109	0.9748	0.0165	0.9534	0.0870	0.9657	0.1942	0.7691	0.1072	0.8861
CheB42b	CheB42b	1634305_at	0.2286	0.3140	0.1823	0.1581	0.2286	0.2373	0.0891	0.9042	0.0156	0.9669	-0.0735	0.7854	0.0607	0.9407	0.0956	0.7476	0.0349	0.9197
Klp98A	Kinesin-like protei	1634306_at	-0.3403	0.2188	-0.4074	0.2209	-0.8531	0.0022	-0.1790	0.8090	0.1254	0.7132	0.3044	0.2502	0.0629	0.9604	-0.0763	0.8821	-0.1392	0.7346
Graf	Graf	1634307_at	0.3468	0.0807	0.0276	0.9115	-0.0418	0.8239	-0.1931	0.5794	-0.1636	0.3691	0.0294	0.8903	-0.1708	0.8049	-0.4600	0.1173	-0.2892	0.3293
---	---	1634308_at	0.1167	0.5576	-0.1178	0.6000	-0.0025	0.9916	0.1518	0.7596	0.1651	0.4562	0.0133	0.9599	0.0845	0.8973	0.0464	0.9002	-0.0381	0.9097
CG14322	CG14322	1634309_at	-0.1654	0.5477	-0.6559	0.0834	-0.8339	0.0077	0.0592	0.9517	0.9763	0.0040	0.9171	0.0032	0.0894	0.9514	0.2825	0.5324	0.1931	0.6881
TpnC4	TpnC4	1634310_at	-1.7563	0.0265	-3.5866	0.0011	-2.9538	0.0007	0.2455	0.8968	1.3596	0.0362	1.1140	0.0507	-0.1608	0.9029	-0.1757	0.7593	-0.0149	0.9849
CG14408 /// DsimCG14408	CG14408	1634311_at	0.1930	0.7420	0.2376	0.6406	0.3272	0.0919	0.2459	0.6755	-0.1759	0.5543	-0.4219	0.0879	0.0654	0.9862	-0.0199	0.9924	-0.0853	0.9433
---	---	1634312_at	0.2488	0.2155	-0.0601	0.6531	0.0836	0.7209	0.1839	0.6605	0.2867	0.1515	0.1028	0.6130	0.0983	0.9048	0.0164	0.9739	-0.0819	0.8255
---	---	1634313_s_at	0.2934	0.1112	1.0942	0.0074	1.1687	0.0004	-0.0996	0.8642	-0.8257	0.0027	-0.7261	0.0027	-0.0102	0.9928	-0.0066	0.9917	0.0036	0.9941
Idh	NADP-IDH	1634314_s_at	0.1941	0.4045	0.7130	0.0101	1.1664	0.0004	0.0334	0.9727	-0.5926	0.0227	-0.6261	0.0111	-0.3167	0.6749	-0.1181	0.7449	0.1985	0.5421
CG32982	CG32982	1634315_a_at	-1.9103	0.0026	-1.6788	0.0181	-1.8314	0.0003	0.2659	0.8194	0.3545	0.4553	0.0887	0.8682	0.1587	0.8869	0.3840	0.3456	0.2254	0.6166
CG1582 /// DmirCG1582	CG1582	1634316_at	-0.1397	0.7507	0.0503	0.9153	0.1736	0.4599	0.0932	0.9247	0.3249	0.2746	0.2317	0.4004	0.3599	0.7889	0.6479	0.2395	0.2879	0.6389
---	---	1634317_at	-0.0967	0.5598	-0.0484	0.7851	-0.2406	0.2798	0.0184	0.9874	0.0575	0.8778	0.0390	0.9086	-0.0495	0.9407	0.0622	0.8161	0.1117	0.6149
CG33110	CG33110	1634318_at	-0.2370	0.7754	-0.2723	0.5570	-0.3554	0.0379	-0.0028	0.9988	-0.5533	0.1531	-0.5505	0.1125	-0.0675	0.9898	-0.8557	0.4667	-0.7882	0.5153
CG14316	CG14316	1634319_at	0.0673	0.6777	0.0688	0.4689	0.1038	0.6237	0.1168	0.7293	-0.0273	0.8943	-0.1441	0.2915	0.1289	0.7726	0.0296	0.9222	-0.0993	0.6328
CG9548	CG9548	1634320_at	0.3902	0.2663	0.4017	0.1866	0.4682	0.0834	0.0780	0.8844	0.3022	0.0987	0.2241	0.1697	0.0593	0.9816	0.3683	0.5257	0.3091	0.6085
CG2051	CG2051	1634321_s_at	-0.2906	0.1789	0.4900	0.0529	1.0370	0.0007	0.1410	0.7850	-0.4868	0.0297	-0.6278	0.0060	-0.3474	0.7230	0.3675	0.3738	0.7148	0.1210
---	---	1634322_at	0.1388	0.4062	-0.0466	0.6732	0.1064	0.5714	0.1803	0.7028	0.2071	0.3484	0.0269	0.9195	0.0119	0.9877	-0.0518	0.8185	-0.0637	0.7495
CG17191	CG17191	1634323_at	0.1972	0.3602	-0.0118	0.9591	-0.0060	0.9797	0.1405	0.7850	0.1254	0.5871	-0.0150	0.9557	0.2740	0.7697	0.2297	0.5855	-0.0443	0.9367
CG34029	CG34029	1634324_at	0.0243	0.8987	0.0860	0.5515	0.0998	0.5278	-0.0637	0.9130	-0.0834	0.6934	-0.0197	0.9306	0.0312	0.9742	0.1054	0.6941	0.0742	0.7964
MESK2	Misexpression sup	1634325_a_at	-2.6904	0.0009	-1.3290	0.0350	-2.0756	0.0001	-0.6524	0.2501	-1.1500	0.0052	-0.4976	0.0979	-0.1584	0.8981	-0.1965	0.7053	-0.0381	0.9528
---	---	1634326_at	-0.0904	0.6381	-0.1059	0.5311	-0.0078	0.9715	0.2086	0.5777	0.0754	0.7396	-0.1333	0.4639	-0.0305	0.9775	-0.1211	0.6686	-0.0906	0.7603
---	---	1634327_at	-0.0033	0.9895	0.3533	0.0444	0.2662	0.1797	-0.1387	0.7857	-0.2060	0.3360	-0.0674	0.7725	-0.0732	0.9412	0.1097	0.7695	0.1829	0.5767
---	---	1634328_at	0.2723	0.1626	-0.0747	0.5248	0.1214	0.5106	0.1501	0.7647	0.2423	0.2592	0.0922	0.6805	-0.1078	0.8901	-0.1224	0.7119	-0.0146	0.9749
CG32579	CG32579	1634329_at	0.1582	0.5546	-1.1960	0.0101	-0.7410	0.0335	0.2482	0.6338	1.1442	0.0017	0.8960	0.0028	-0.2175	0.8409	-0.2411	0.6144	-0.0236	0.9727
ndl	nudel	1634330_at	0.2821	0.1362	0.0182	0.9168	-0.1212	0.4752	0.0384	0.9620	0.1842	0.4019	0.1457	0.4734	0.1789	0.7686	0.0376	0.9269	-0.1413	0.6122
l(2)k14710	lethal (2) k14710	1634331_at	-0.3636	0.0747	-0.3137	0.1048	-0.5676	0.0231	-0.1530	0.7121	-0.0167	0.9528	0.1363	0.4420	-0.0465	0.9589	-0.0068	0.9906	0.0397	0.9109
---	---	1634332_s_at	-0.1723	0.6775	0.0423	0.8864	-0.2241	0.1824	-0.0333	0.9755	-0.1712	0.5492	-0.1379	0.6075	0.0179	0.9916	-0.0859	0.8821	-0.1038	0.8393
---	---	1634333_at	0.1807	0.3751	-0.0815	0.5158	0.3193	0.0372	0.3164	0.3793	0.2217	0.2791	-0.0948	0.6525	-0.1159	0.8875	-0.1990	0.5336	-0.0831	0.8303
CG6015	CG6015	1634334_at	-0.0399	0.8804	0.3629	0.1110	0.6043	0.0063	0.2162	0.6236	-0.0487	0.8656	-0.2649	0.1743	-0.0141	0.9913	0.3460	0.1718	0.3600	0.1911
---	---	1634335_at	-0.0998	0.5460	0.1459	0.3659	0.1786	0.3748	-0.1245	0.8164	-0.1925	0.3701	-0.0681	0.7696	0.0636	0.9095	0.1533	0.4417	0.0897	0.6844
bam	bag-of-marbles	1634336_at	0.2347	0.3090	0.0132	0.9845	-0.5456	0.2709	-1.1329	0.0233	0.5047	0.0637	1.6376	0.0003	-0.4820	0.8331	0.1142	0.9404	0.5963	0.5523
---	---	1634337_s_at	0.2726	0.5326	0.1259	0.8583	-0.8521	0.1053	-0.3583	0.3016	0.5999	0.0102	0.9583	0.0008	0.7379	0.7726	0.5154	0.6762	-0.2225	0.8837
CG12550	CG12550	1634338_x_at	0.2700	0.2276	0.0159	0.8777	-0.0521	0.7916	-0.0217	0.9781	0.0221	0.9398	0.0438	0.8521	-0.0219	0.9816	-0.1858	0.4032	-0.1639	0.4814
CG32694	CG32694	1634339_at	-2.2990	0.0069	-1.8437	0.0321	-1.6450	0.0002	0.2753	0.8483	-0.4351	0.4237	-0.7103	0.1292	0.0674	0.9467	0.1866	0.5634	0.1192	0.7369
CG11970	CG11970	1634340_at	-0.0577	0.9150	-0.1679	0.3380	0.0308	0.8865	0.2153	0.6135	0.2483	0.2464	0.0330	0.8978	0.0917	0.9589	0.1735	0.7781	0.0818	0.9057
CG6129	CG6129	1634341_a_at	1.5638	0.0304	1.7471	0.0361	2.1930	0.0014	0.1044	0.8950	0.3614	0.1670	0.2571	0.2791	-0.1765	0.9589	0.4198	0.7020	0.5963	0.5629
CG15744	CG15744	1634342_s_at	-0.2686	0.2650	0.0222	0.9460	0.1849	0.3957	0.0512	0.9255	-0.1692	0.3109	-0.2204	0.1344	-0.0347	0.9851	0.2573	0.5618	0.2920	0.5050
CG2241	CG2241	1634343_at	0.0761	0.6390	-0.1165	0.5937	0.2812	0.1056	0.2972	0.2581	0.2111	0.1727	-0.0861	0.5788	-0.0111	0.9922	0.2046	0.4280	0.2157	0.4114
l(2)35Bg	transcription unit E	1634344_at	-0.4400	0.0670	-0.3777	0.1878	-0.0761	0.8127	0.2418	0.5948	0.3562	0.1210	0.1144	0.6262	-0.1566	0.9246	0.2920	0.6218	0.4486	0.4270
Nox	NADPH oxidase	1634345_at	0.3393	0.2931	-0.0153	0.9229	0.0555	0.8008	0.0415	0.9603	0.2918	0.1870	0.2503	0.2081	0.1108	0.8875	-0.0834	0.8304	-0.1942	0.5299
CG13239	CG13239	1634346_at	0.1083	0.6015	0.2241	0.1486	0.3489	0.1456	-0.1267	0.7278	-0.0212	0.9254	0.1054	0.5010	0.0528	0.9717	0.1137	0.8177	0.0609	0.9077
---	---	1634347_at	0.1985	0.3617	0.1169	0.5315	0.1536	0.2777	0.0129	0.9880	0.1155	0.5964	0.1026	0.6117	0.0147	0.9898	0.0794	0.7922	0.0647	0.8307
---	---	1634348_s_at	0.0914	0.5377	-0.1068	0.5451	0.0745	0.7328	-0.0387	0.9558	0.0979	0.6418	0.1367	0.4383	-0.1915	0.7230	-0.1536	0.5248	0.0379	0.9061
fs(1)M3	female sterile (1) I	1634349_at	0.4195	0.4800	-0.9613	0.0390	-0.1866	0.6871	0.5396	0.0584	1.2825	0.0002	0.7428	0.0009	-0.3297	0.8963	-0.2580	0.8386	0.0718	0.9582
sob	odd skipped-relate	1634350_at	-2.8872	0.0004	-0.5945	0.3398	-2.4115	0.0009	-1.7904	0.1034	-2.7209	0.0022	-0.9306	0.1092	0.1642	0.8963	-0.2958	0.5465	-0.4600	0.3406
CG7860	CG7860	1634351_at	-0.5789	0.0592	-0.1788	0.5311	-0.3415	0.1637	-0.2442	0.7752	-0.8462	0.0259	-0.6020	0.0633	0.0994	0.9125	-0.0886	0.8353	-0.1880	0.5790
---	---	1634352_at	-0.0045	0.9861	-0.1093	0.4944	0.0355	0.8822	0.1000	0.9004	0.1395	0.6283	0.0395	0.8992	0.1054	0.8903	0.1743	0.5636	0.0689	0.8535
Lcp3	cuticle p. III	1634353_at	0.4456	0.1745	0.4012	0.1238	0.5809	0.0048	0.0280	0.9757	-0.0158	0.9633	-0.0437	0.8681	-0.0509	0.9742	-0.0781	0.8918	-0.0272	0.9622
---	---	1634354_at	-0.0192	0.9511	0.0269	0.7990	0.2783	0.1104	0.0516	0.9160	-0.0502	0.7872	-0.1019	0.4849	-0.1376	0.8960	-0.0714	0.9058	0.0662	0.9028
CG17404	CG17404	1634355_at	-0.0147	0.9613	-0.1678	0.4854	0.1331	0.5975	0.0064	0.9956	0.1175	0.6748	0.1111	0.6630	-0.0954	0.8903	-0.0074	0.9897	0.0879	0.7745

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1634356_at	0.2384	0.1377	0.1002	0.4388	0.1126	0.4848	-0.1344	0.7121	-0.0927	0.6120	0.0417	0.8252	-0.0526	0.9514	-0.1964	0.4404	-0.1438	0.5995
---	---	1634357_at	0.0028	0.9925	-0.1275	0.4305	0.0938	0.6661	0.1951	0.5357	0.2265	0.1662	0.0314	0.8713	-0.2022	0.7387	-0.0731	0.8307	0.1292	0.6410
---	---	1634358_s_at	0.2671	0.1223	0.1688	0.2162	-0.0453	0.8618	-0.0856	0.9116	0.0068	0.9851	0.0924	0.7219	0.1585	0.7628	0.0358	0.9157	-0.1227	0.6029
Borr	Borealin	1634359_a_at	-0.0732	0.9561	-1.7490	0.2712	-2.2039	0.0307	-0.6151	0.7010	1.8419	0.0206	2.4569	0.0034	-0.2522	0.9816	-0.0938	0.9832	0.1584	0.9630
---	---	1634360_at	0.0683	0.7108	0.1178	0.6723	0.0391	0.8055	-0.0644	0.9133	-0.0744	0.7348	-0.0099	0.9669	0.0941	0.9330	0.0435	0.9402	-0.0506	0.9186
CG31315	CG31315	1634361_at	0.1314	0.4402	0.0152	0.9302	0.1036	0.5904	-0.0264	0.9649	0.0171	0.9411	0.0434	0.8092	-0.1574	0.8217	-0.0262	0.9558	0.1313	0.6817
caz	cabeza	1634362_a_at	0.0014	0.9966	0.0974	0.5564	0.1567	0.4181	-0.1061	0.8636	0.5154	0.0265	0.6215	0.0073	-0.1865	0.8400	0.5359	0.1497	0.7224	0.0922
---	---	1634363_at	0.0009	0.9980	0.0401	0.7900	-0.1702	0.4278	-0.0048	0.9961	-0.0117	0.9801	-0.0068	0.9858	0.1856	0.7677	0.0522	0.8960	-0.1334	0.6419
---	---	1634364_s_at	-0.8728	0.0644	-0.8820	0.0854	-1.4229	0.0005	0.2541	0.6046	0.7156	0.0110	0.4614	0.0431	0.7278	0.5765	0.7014	0.2194	-0.0264	0.9777
CG2278	CG2278	1634365_at	0.1668	0.4961	0.4201	0.0527	-0.0722	0.6940	-0.0719	0.9029	-0.1555	0.4305	-0.0837	0.6742	0.1504	0.8062	0.0524	0.8879	-0.0980	0.7409
CG32185	CG32185	1634366_at	0.5846	0.2773	0.0484	0.9278	-0.4801	0.1696	-0.3113	0.5217	0.7428	0.0106	1.0541	0.0014	0.3800	0.8461	-0.0217	0.9905	-0.4017	0.6428
---	---	1634367_at	0.0029	0.9915	0.0439	0.6960	0.0625	0.7012	-0.0701	0.9353	-0.0285	0.9380	0.0417	0.8915	0.0246	0.9773	0.0049	0.9918	-0.0196	0.9444
CG40268	CG40268	1634368_at	0.0371	0.8435	-0.1785	0.5598	-0.0375	0.8190	-0.0567	0.9380	0.1400	0.5440	0.1967	0.3138	-0.2763	0.6955	-0.0125	0.9821	0.2638	0.3803
---	---	1634369_at	0.3546	0.1132	0.1840	0.3416	0.0675	0.7936	-0.0681	0.9011	0.0316	0.8962	0.0997	0.5709	0.1086	0.9001	0.0528	0.9151	-0.0558	0.8985
pnr	pannier	1634370_a_at	2.3474	0.0010	2.2027	0.0014	2.2503	0.0000	0.0872	0.9467	0.2830	0.4620	0.1959	0.5964	-0.0839	0.9156	0.0150	0.9748	0.0989	0.7560
CG14020	CG14020	1634371_at	0.1102	0.5653	0.1087	0.5894	0.1147	0.4617	-0.1209	0.7514	-0.0328	0.8829	0.0881	0.5956	0.0440	0.9421	0.0541	0.8247	0.0101	0.9717
Ddr	Discoidin domain	1634372_at	-0.0176	0.9467	0.0103	0.9515	-0.1484	0.5576	0.2034	0.7400	0.0280	0.9437	-0.1754	0.4991	0.1920	0.7707	-0.0321	0.9441	-0.2242	0.4399
CG13492	CG13492	1634373_a_at	-0.2514	0.9422	0.2331	0.4442	0.3533	0.5701	-0.4637	0.9412	-1.5421	0.4029	-1.0784	0.5387	-0.7649	0.9296	-1.4289	0.6377	-0.6640	0.8548
CG33138	CG33138	1634374_at	0.6179	0.0632	0.4412	0.4568	0.6289	0.0379	0.0584	0.9540	-1.2155	0.0018	-1.2738	0.0009	-0.1098	0.9679	-1.3217	0.0795	-1.2119	0.1269
---	---	1634375_at	0.2416	0.1132	0.0413	0.7194	-0.0742	0.6395	-0.0007	0.9994	0.1132	0.5506	0.1140	0.5017	0.0163	0.9860	-0.0898	0.7000	-0.1061	0.6328
CG15307	CG15307	1634376_at	-1.0961	0.0133	-0.2871	0.0563	-1.3192	0.0046	-0.3300	0.6663	-0.6027	0.0986	-0.2728	0.4250	0.3036	0.6955	0.0700	0.8817	-0.2335	0.4892
CG14646	CG14646	1634377_at	-0.1454	0.4426	0.3299	0.4358	0.2854	0.1488	0.0528	0.9247	0.0899	0.6294	0.0371	0.8483	0.0110	0.9958	0.5528	0.2385	0.5418	0.2798
---	---	1634378_at	0.1256	0.4191	0.0720	0.5835	-0.0688	0.7377	0.1639	0.6584	0.4020	0.0315	0.2381	0.1336	0.0954	0.8472	0.1253	0.5622	0.0300	0.9165
fru	fru-satori	1634379_a_at	-0.0235	0.9146	0.0065	0.9655	0.1148	0.7015	0.0230	0.9826	0.1380	0.6121	0.1150	0.6521	0.1587	0.8734	0.1145	0.8231	-0.0441	0.9360
CG32065	CG32065	1634380_at	0.0880	0.7445	0.0097	0.9343	0.0843	0.6882	-0.0260	0.9749	-0.0638	0.7980	-0.0378	0.8753	0.0502	0.9548	-0.0808	0.8099	-0.1310	0.6389
CG2818	CG2818	1634381_a_at	-0.0153	0.9430	0.2759	0.4259	0.1783	0.2732	-0.1926	0.6936	0.0268	0.9327	0.2195	0.2821	-0.0121	0.9940	0.2631	0.4939	0.2753	0.4776
CG31675	CG31675	1634382_at	-1.7173	0.0034	-1.7554	0.0389	-2.5084	0.0000	-0.2093	0.5836	-0.1975	0.3147	0.0117	0.9616	0.3716	0.8395	-0.3451	0.6827	-0.7168	0.3572
Jhl-21	Jhl-21	1634383_a_at	1.1164	0.0010	1.1363	0.0354	0.4890	0.0758	0.0890	0.8589	0.8107	0.0016	0.7217	0.0016	0.6650	0.5126	0.8651	0.0843	0.2001	0.7134
CG13426	CG13426	1634384_at	-0.2273	0.3269	0.2078	0.6681	0.4567	0.0997	0.1368	0.8028	-0.0280	0.9277	-0.1648	0.4223	0.0397	0.9862	0.5534	0.2727	0.5137	0.3375
CG15524	CG15524	1634385_at	-0.2578	0.3805	-0.0855	0.5552	-0.0038	0.9932	-0.2232	0.7225	-0.1952	0.5153	0.0280	0.9364	-0.2877	0.7485	0.1053	0.8367	0.3930	0.3128
Rab4	Rab-protein 4	1634386_s_at	-0.2869	0.1977	-0.4440	0.1597	-0.2312	0.1429	0.0363	0.9436	-0.1438	0.3331	-0.1801	0.1676	-0.2257	0.7780	-0.2350	0.5141	-0.0093	0.9868
CG14691	CG14691	1634387_at	0.0933	0.6141	0.0402	0.8215	-0.1242	0.4507	0.0169	0.9893	0.1022	0.7651	0.0853	0.7892	0.1442	0.7677	0.0280	0.9326	-0.1162	0.6021
CG30197	CG30197	1634388_at	-0.1775	0.4583	-0.0918	0.5407	0.0520	0.8413	0.3287	0.4509	-0.0638	0.8326	-0.3926	0.0669	0.2052	0.8202	0.0663	0.9075	-0.1389	0.7457
---	---	1634389_at	0.1431	0.4271	0.2755	0.1145	0.2121	0.3002	0.0832	0.8736	0.0068	0.9804	-0.0764	0.6888	0.0910	0.8885	0.1688	0.5042	0.0777	0.7918
Nacalpha	Alpha NAC	1634390_s_at	0.1799	0.2025	0.1042	0.4271	0.1352	0.3358	-0.0395	0.9416	-0.0026	0.9903	0.0369	0.8346	-0.0179	0.9816	-0.0847	0.6642	-0.0667	0.7441
CG18109	CG18109	1634391_at	0.0904	0.5648	-0.0240	0.8160	0.0308	0.8710	-0.0770	0.8809	0.0854	0.6680	0.1624	0.3072	-0.0090	0.9914	-0.0082	0.9824	0.0008	0.9982
CG14257	CG14257	1634392_at	0.2853	0.2002	0.2145	0.4639	0.0471	0.8233	0.0984	0.8872	-0.0164	0.9633	-0.1149	0.6314	0.1423	0.8400	-0.0166	0.9734	-0.1589	0.6129
Klp10A	Klp10A	1634393_s_at	0.3563	0.4546	1.0169	0.1304	0.9687	0.0089	0.0913	0.9351	0.2773	0.4169	0.1861	0.5709	0.2265	0.9309	1.1866	0.1512	0.9602	0.2765
Or67d	Or67d	1634394_at	0.1025	0.5511	-0.0965	0.4257	-0.0741	0.6488	-0.0199	0.9777	0.0636	0.7722	0.0836	0.6586	-0.0428	0.9588	-0.0597	0.8487	-0.0169	0.9596
yellow-g2	yellow-g2	1634395_at	0.4185	0.0167	0.1267	0.4974	-0.0903	0.5936	0.1147	0.8107	0.2699	0.1538	0.1552	0.3775	-0.0254	0.9848	-0.0223	0.9647	0.0031	0.9953
CG3303 /// DmirCG3303	CG3303	1634396_at	-0.7997	0.0080	-0.4903	0.1721	-0.4426	0.0866	-0.0603	0.9311	-0.4330	0.0411	-0.3727	0.0475	-0.1705	0.8815	-0.2005	0.6820	-0.0300	0.9628
---	---	1634397_at	0.0981	0.5698	-0.0659	0.5720	-0.0112	0.9699	-0.0695	0.9436	-0.0875	0.7998	-0.0180	0.9594	-0.0521	0.9266	-0.0176	0.9531	0.0345	0.8929
Dl	Overflow	1634398_a_at	-2.0921	0.0007	-1.4954	0.0226	-2.1781	0.0000	-0.0372	0.9708	-0.6562	0.0196	-0.6190	0.0156	0.5278	0.5754	-0.0244	0.9726	-0.5523	0.2094
CG18367	CG18367	1634399_at	-3.8221	0.0169	-4.0436	0.0179	-4.2827	0.0000	-0.4882	0.0821	-1.4103	0.0001	-0.9221	0.0004	-0.1007	0.9901	-1.5269	0.3780	-1.4262	0.4318
---	---	1634400_at	-0.0244	0.9044	-0.0477	0.7313	-0.0434	0.7999	0.0771	0.8868	0.0927	0.6532	0.0156	0.9464	0.0046	0.9952	-0.0147	0.9613	-0.0193	0.9394
fig	fos intronic gene	1634401_at	0.0806	0.6562	0.0822	0.5625	-0.1682	0.2902	-0.1578	0.6936	0.0269	0.9175	0.1847	0.2690	0.0816	0.9467	-0.0118	0.9877	-0.0934	0.8461
Scgbeta	Sarcoglycan beta	1634402_at	-1.6099	0.0094	-0.6482	0.1097	-0.6688	0.0021	0.3615	0.4337	-0.7397	0.0106	-1.1012	0.0011	0.3274	0.7387	-0.0673	0.9170	-0.3947	0.3576
alpha-Adaptin	alpha-adaptin	1634403_at	-0.8229	0.0739	-0.7876	0.0456	-1.1580	0.0006	-0.0261	0.9704	0.3409	0.0614	0.3670	0.0298	0.2864	0.8461	0.3142	0.6345	0.0278	0.9771
CG4573	CG4573	1634404_at	-0.3354	0.2259	0.0202	0.9236	0.3217	0.0657	-0.2198	0.5758	-0.5233	0.0177	-0.3035	0.0922	-0.5288	0.3712	-0.1849	0.5897	0.3439	0.2993
Pak	Pak1 kinase	1634405_s_at	0.3364	0.3832	0.5618	0.0330	0.2896	0.1716	-0.1162	0.7753	0.0759	0.6991	0.1921	0.2072	0.2185	0.8940	0.3228	0.6250	0.1043	0.9023
CG32982	CG32982	1634406_at	-1.3999	0.0016	-1.3899	0.0161	-1.3620	0.0003	0.5189	0.3645	0.3826	0.2407	-0.1363	0.6902	0.3498	0.6557	0.2528	0.4280	-0.0970	0.8039
CG1950	CG1950	1634407_at	0.1348	0.4122	0.0579	0.6147	-0.1282	0.4947	-0.2947	0.4221	-0.1993	0.3337	0.0954	0.6485	0.0307	0.9741	-0.0864	0.7544	-0.1171	0.6389
CG10209	CG10209	1634408_at	0.1192	0.4496	-0.2056	0.1849	-0.4074	0.0588	0.2393	0.4410	0.5907	0.0048	0.3514	0.0281	0.2984	0.6695	0.2541	0.3565	-0.0443	0.9093
CG31775	CG31775	1634409_at	0.0120	0.9699	0.2622	0.4581	1.1413	0.1335	0.1135	0.8794	0.1035	0.7299	-0.0100	0.9753	-0.9442	0.7215	0.1740	0.9193	1.1182	0.3235
CG11560	CG11560	1634410_at	-0.0717	0.7423	-0.1318	0.4956	-0.2671	0.2603	0.0030	0.9960	0.2856	0.1289	0.2826	0.0939	0.2082	0.8541	0.2922	0.5495	0.0839	0.8951
CG6506	CG6506	1634411_at	0.3760	0.1162	0.0497	0.7660	0.1943	0.2842	0.1344	0.6823	0.3566	0.0276	0.2222	0.1038	-0.0294	0.9816	0.0979	0.7817	0.1273	0.6877
CG2135	CG2135	1634412_at	0.1953	0.3005	0.4683	0.2611	0.8259	0.0048	0.0281	0.9749	-0.4764	0.0358	-0.5044	0.0178	-0.3691	0.6955	-0.2275	0.5802	0.1416	0.7516
run	runt	1634413_at	-0.0760	0.7941	0.0803	0.5408	0.1478	0.3878	0.1498	0.6493	0.0246	0.9116	-0.1253	0.3975	0.0753	0.9156	0.1359	0.6080	0.0606	0.8496
mRpL23	mitochondrial ribo	1634414_at	-0.4685	0.2441	-0.5870	0.0698	-0.6360	0.0569	-0.2030	0.7028	-0.3128	0.1991	-0.1098	0.6655	-0.2038	0.8680	-0.6152	0.1922	-0.4114	0.4055
CG12685	CG12685	1634415_at	-0.1947	0.3231	0.0344	0.8886	-0.1973	0.2032	-0.1918	0.5038	-0.1451	0.3485	0.0467	0.7828	0.0186	0.9848	0.0198	0.9548	0.0012	0.9979
---	---	1634416_x_at	-0.1947	0.3875	-0.0026	0.9855	0.0208	0.9108	-0.0215	0.9755	-0.0009	0.9973	0.0206	0.9221	-0.1372	0.8215	0.2349	0.3397	0.3721	0.1651
CG3683 /// DyakCG3683	CG3683	1634417_s_at	-0.0506	0.8945	0.2210	0.4119	0.3120	0.1851	-0.0342	0.9727	-0.5948	0.0246	-0.5605	0.0197	-0.0078	0.9971	-0.2445	0.6730	-0.2367	0.6817
CG33281	CG33281	1634418_at	-1.5336	0.0113	-1.9090	0.0052	-2.1129	0.0000	0.0542	0.9834	0.2035	0.7776	0.1493	0.8257	0.1914	0.7266	-0.0043	0.9935	-0.1956	0.4148
Tango14	Transport and Gol	1634419_at	-0.0379	0.9024	0.7623	0.0974	0.6140	0.0121	0.0353	0.9664	-0.3709	0.0916	-0.4062	0.0442	0.1598	0.9030	0.4531	0.3299	0.2932	0.5629
---	---	1634420_at	0.0790	0.6243	0.0000	1.0000	0.0029	0.9894	0.1635	0.5735	0.2162	0.1433	0.0527	0.7415	0.0609	0.9095	0.0054	0.9900	-0.0555	0.8158
CG8414	CG8414	1634421_at	0.3153	0.3825	-0.2167	0.5124	0.0709	0.7301	0.2507	0.5270	0.9517	0.0015	0.7010	0.0033	-0.1371	0.9309	0.2670	0.6270	0.4041	0.4405
CG31213	CG31213	1634422_at	-0.0832	0.6788	-0.0608	0.7818	-0.1042	0.5138	0.0069	0.9937	0.1505	0.4113	0.1435	0.3834	-0.1256	0.8270	-0.0057	0.9921	0.1199	0.6398
CG32408	CG32408	1634423_at	0.3273	0.0580	0.2417	0.2158	-0.0369	0.8710	0.0380	0.9506	0.0668	0.7380	0.0288	0.8862	0.0428	0.9713	-0.0864	0.8248	-0.1293	0.6969
---	---	1634424_at	0.0083	0.9761	0.0744	0.5240	0.2605	0.1859	0.0548	0.9253	-0.1278	0.4871	-0.1826	0.2461	-0.0347	0.9816	0.1122	0.7863	0.1468	0.6888

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1634425_at	0.0672	0.7304	0.0418	0.6546	0.1589	0.2719	0.0392	0.9445	0.0226	0.9201	-0.0165	0.9341	-0.0197	0.9848	0.0535	0.8684	0.0733	0.7889
---	---	1634426_at	0.0672	0.6859	-0.1137	0.5461	-0.2393	0.1703	-0.1329	0.7590	-0.0417	0.8652	0.0912	0.6314	0.0864	0.8999	0.0078	0.9884	-0.0787	0.7964
CG15461	CG15461	1634427_at	-1.3012	0.0059	-1.9117	0.0085	-1.9419	0.0002	0.1308	0.8085	0.5807	0.0148	0.4499	0.0271	0.0089	0.9964	-0.2672	0.5523	-0.2761	0.5430
CG5924	CG5924	1634428_at	0.1475	0.5021	0.2820	0.4209	0.4785	0.0083	-0.2473	0.4183	-0.4232	0.0208	-0.1759	0.2478	-0.4056	0.7033	-0.2349	0.6222	0.1706	0.7390
CG8360	CG8360	1634429_at	0.5980	0.0048	0.3782	0.5004	0.8469	0.0018	0.2978	0.3420	-0.1017	0.6083	-0.3995	0.0215	-0.2200	0.8740	-0.3288	0.5649	-0.1088	0.8844
CG14614	CG14614	1634430_at	0.1651	0.7166	0.1052	0.5877	-0.6169	0.0609	-0.6640	0.0657	0.2297	0.2712	0.8937	0.0012	-0.0214	0.9943	0.0925	0.9339	0.1139	0.9056
CG15071	CG15071	1634431_at	0.2958	0.1969	0.4756	0.2133	0.6817	0.0076	-0.2172	0.5068	-0.1177	0.5242	0.0995	0.5620	-0.0936	0.9589	0.1026	0.8924	0.1962	0.7416
---	---	1634432_at	0.0889	0.6381	0.0504	0.7060	0.0394	0.8617	0.0772	0.8897	0.0130	0.9635	-0.0642	0.7514	0.1957	0.7726	0.0268	0.9552	-0.1690	0.5882
scu	scully	1634433_at	0.6217	0.0550	0.3897	0.6358	0.3969	0.1646	0.0230	0.9778	-0.2495	0.2421	-0.2725	0.1513	-0.0674	0.9862	-0.5526	0.5629	-0.4851	0.6211
CG5359	CG5359	1634434_at	-0.7184	0.0857	-1.0330	0.0432	-0.6390	0.0319	-0.0073	0.9956	0.6398	0.1213	0.6471	0.0822	-0.5104	0.6749	0.1092	0.8804	0.6196	0.2351
---	---	1634435_at	-0.0493	0.8285	0.1173	0.3730	-0.1368	0.4683	0.0232	0.9777	0.0687	0.7906	0.0455	0.8530	0.0366	0.9535	-0.0381	0.8877	-0.0747	0.7239
Ast-C	drostatin-C	1634436_at	-0.5310	0.5576	-0.4934	0.0457	-0.1689	0.2923	0.4597	0.8073	-0.0034	0.9977	-0.4631	0.5251	-0.1271	0.8558	-0.1654	0.5855	-0.0383	0.9231
CG10973	CG10973	1634437_at	0.3905	0.0544	0.1448	0.5387	-0.0023	0.9921	0.1835	0.6321	0.5613	0.0094	0.3778	0.0316	0.3276	0.5765	0.3311	0.1991	0.0035	0.9944
Pph13	Munster	1634438_at	0.1293	0.5244	0.3964	0.0435	0.1939	0.3578	0.0569	0.9435	-0.0333	0.9167	-0.0902	0.7075	0.1314	0.8049	0.1584	0.4848	0.0270	0.9312
CG33256	CG33256	1634439_at	-0.5366	0.1233	0.0991	0.5566	0.0258	0.8813	-0.0735	0.8671	-0.4026	0.0174	-0.3291	0.0255	0.0222	0.9898	-0.0481	0.9343	-0.0704	0.8884
Eip74EF	ecdysone inducibl	1634440_s_at	0.8898	0.0370	-0.1964	0.6311	-1.0097	0.0268	-0.4357	0.5461	0.1653	0.7015	0.6010	0.0742	0.4851	0.7726	-0.8717	0.2110	-1.3568	0.0940
---	---	1634441_at	-0.0057	0.9755	0.0082	0.9418	0.0516	0.7840	0.0612	0.8990	-0.0502	0.7951	-0.1114	0.4548	-0.0775	0.8611	-0.0108	0.9714	0.0667	0.7484
smo	smoothened	1634442_at	0.2760	0.1665	0.0052	0.9857	-0.1846	0.2259	-0.0518	0.9269	0.2630	0.1181	0.3148	0.0433	-0.0016	0.9994	-0.1761	0.6044	-0.1746	0.6117
Rab1	Rab-protein 1	1634443_a_at	-0.4862	0.0437	-0.3463	0.2424	-0.5271	0.0158	-0.0006	0.9994	-0.2178	0.2313	-0.2172	0.1814	0.3167	0.6957	0.0225	0.9682	-0.2942	0.3994
CG12736	CG12736	1634444_at	-0.8562	0.0020	-0.3287	0.0865	-0.0913	0.6887	0.0520	0.9314	-0.7122	0.0028	-0.7642	0.0012	-0.2497	0.5659	-0.0739	0.7504	0.1758	0.3817
CG32750	CG32750	1634445_at	-1.2201	0.0015	1.8997	0.0321	0.8244	0.0353	-1.4016	0.0417	-3.1947	0.0002	-1.7931	0.0008	-0.3670	0.7757	-0.1356	0.8612	0.2314	0.7134
CG30065	CG30065	1634446_at	0.0474	0.8347	0.0601	0.7042	0.1248	0.5430	0.0243	0.9777	-0.0119	0.9732	-0.0362	0.8926	0.0282	0.9848	0.0399	0.9404	0.0117	0.9837
CG31140	CG31140	1634447_at	-2.3851	0.0005	-2.4763	0.0100	-2.9294	0.0001	-0.0003	0.9997	0.1485	0.5413	0.1488	0.4932	0.3283	0.8400	-0.0013	0.9998	-0.3296	0.6476
---	---	1634448_at	0.1471	0.4062	0.0936	0.3801	0.0479	0.7994	-0.0284	0.9726	0.0074	0.9809	0.0358	0.8826	0.2141	0.7381	0.1371	0.6414	-0.0770	0.8204
CG9459	CG9459	1634449_at	0.5802	0.6392	-0.9554	0.6064	1.2390	0.1450	1.3717	0.3420	0.5657	0.5221	-0.8059	0.2832	-0.6070	0.9374	-0.8242	0.7842	-0.2172	0.9491
---	---	1634450_a_at	0.1981	0.2997	-0.0116	0.9698	0.1768	0.6250	-0.2601	0.4131	-0.1146	0.5455	0.1454	0.3716	-0.3689	0.7464	-0.2016	0.7120	0.1673	0.7652
CG12236	CG12236	1634451_at	-0.4841	0.2696	-0.1751	0.6823	-0.4221	0.1185	0.0597	0.9603	-0.4383	0.1703	-0.4981	0.0840	0.1351	0.9330	-0.0211	0.9829	-0.1562	0.8039
endoA	endophilin	1634452_s_at	-0.9399	0.0045	-1.4118	0.0047	-1.6347	0.0000	-0.0591	0.9228	0.4313	0.0264	0.4903	0.0095	0.2176	0.7893	-0.0028	0.9980	-0.2204	0.5476
g	garnet	1634453_at	0.3197	0.5269	0.1120	0.5661	-0.6444	0.1679	-0.0401	0.9699	0.5886	0.0372	0.6286	0.0179	0.5982	0.7611	0.3891	0.6683	-0.2091	0.8424
Bruce	Bruce	1634454_at	-0.0154	0.9748	0.8202	0.1811	0.8658	0.0015	-0.0078	0.9956	-0.4794	0.2738	-0.4715	0.2271	-0.0696	0.9677	0.2758	0.5597	0.3454	0.4596
CG14406	CG14406	1634455_at	0.0464	0.8376	0.1850	0.1688	0.2525	0.1058	-0.0244	0.9772	-0.1609	0.4560	-0.1366	0.4922	-0.0804	0.9142	0.0794	0.8151	0.1598	0.5637
CG34371	CG13548	1634456_at	-0.0940	0.5445	0.4051	0.0233	0.1153	0.5680	-0.3176	0.4420	-0.3665	0.1033	-0.0489	0.8516	0.3006	0.6557	0.4058	0.1356	0.1052	0.7371
CG12702	CG12702	1634457_at	-0.0739	0.6993	-0.0409	0.7968	0.1706	0.5994	-0.0406	0.9387	0.1122	0.4886	0.1528	0.2739	-0.1519	0.9142	0.0835	0.9130	0.2354	0.6617
CG31635	CG31635	1634458_at	0.1278	0.6727	0.1683	0.3039	0.1216	0.5609	-0.0190	0.9819	-0.0698	0.7688	-0.0509	0.8209	0.0106	0.9939	0.0416	0.9310	0.0311	0.9414
---	---	1634459_at	0.2478	0.2062	0.2319	0.0641	0.2577	0.2558	0.0518	0.9257	0.1533	0.3699	0.1016	0.5344	-0.0008	0.9998	0.2120	0.5457	0.2128	0.5488
---	---	1634460_at	0.1434	0.4661	-0.1476	0.4010	-0.1075	0.5857	0.0894	0.8676	0.2941	0.1196	0.2047	0.2283	-0.0304	0.9816	0.0072	0.9923	0.0376	0.9293
DIP2	DISCO Interacting	1634461_at	0.0636	0.7473	0.3640	0.2497	0.3550	0.1684	-0.0477	0.9598	0.0724	0.8128	0.1201	0.6356	-0.1845	0.8404	0.3194	0.3922	0.5039	0.2054
CG3822	CG3822	1634462_at	0.1101	0.4860	0.0373	0.8307	0.0795	0.6209	0.0684	0.8883	-0.0078	0.9757	-0.0762	0.6528	0.0716	0.9309	-0.0275	0.9499	-0.0991	0.7492
CG7992	CG7992	1634463_at	-0.2480	0.8827	-0.0749	0.6791	-0.1831	0.5040	0.0611	0.9882	-0.8591	0.3812	-0.9203	0.2896	-0.0388	0.9928	-0.5230	0.6156	-0.4842	0.6434
RpS19b	Ribosomal protein	1634464_at	0.0018	0.9952	0.1157	0.3781	0.0962	0.5772	-0.0846	0.8971	-0.1736	0.4368	-0.0890	0.6940	0.0691	0.9226	0.0098	0.9828	-0.0593	0.8444
GluClalpha	indefinite	1634465_a_at	-0.0297	0.8918	-0.1200	0.4737	-0.0835	0.7202	0.0535	0.9302	-0.0141	0.9576	-0.0676	0.7236	-0.0366	0.9645	-0.1381	0.5523	-0.1014	0.6766
CG9507	CG9507	1634466_at	0.1736	0.4380	0.0583	0.8240	0.1095	0.5010	0.1600	0.7149	0.0028	0.9922	-0.1572	0.3983	0.0188	0.9898	-0.1469	0.6768	-0.1657	0.6291
---	---	1634467_at	0.1008	0.6379	-0.1298	0.6431	-0.2651	0.1578	-0.0381	0.9744	0.0338	0.9326	0.0719	0.8241	0.1402	0.8236	-0.0177	0.9672	-0.1579	0.5704
CG13397	CG13397	1634468_at	-0.2942	0.3101	-2.1208	0.0135	-1.5427	0.0003	0.1440	0.8512	0.4847	0.0845	0.3406	0.1727	-0.4492	0.7220	-1.2915	0.0391	-0.8423	0.1426
CG40244	CG40244	1634469_at	-0.1063	0.8638	-0.3488	0.6755	-0.2629	0.4478	0.1190	0.9542	0.6009	0.2944	0.4820	0.3559	-0.1537	0.9522	0.1242	0.9170	0.2779	0.7541
---	---	1634470_at	0.1224	0.4449	0.0298	0.8886	0.0519	0.8027	0.1056	0.7539	0.1209	0.4217	0.0153	0.9321	0.0900	0.8886	0.0305	0.9379	-0.0595	0.8484
CG33214	CG33214	1634471_at	0.0102	0.9735	0.1158	0.5492	0.2627	0.1768	-0.0137	0.9852	-0.0013	0.9955	0.0124	0.9534	-0.1592	0.8882	0.1432	0.7877	0.3024	0.4917
CG9114	CG9114	1634472_at	0.0007	0.9984	-0.0943	0.4321	0.2780	0.1485	0.0802	0.8550	-0.0493	0.7998	-0.1296	0.3774	-0.3041	0.7036	-0.2107	0.5446	0.0934	0.8234
---	---	1634473_at	0.0217	0.9226	0.0345	0.7943	0.2568	0.1197	0.0266	0.9666	-0.0654	0.7396	-0.0920	0.5801	-0.2407	0.7230	-0.0740	0.8550	0.1667	0.5986
Pop2	Pop2	1634474_s_at	-0.8092	0.0129	0.1940	0.4835	0.0755	0.6458	-0.1378	0.7604	-0.6713	0.0050	-0.5335	0.0084	-0.0208	0.9913	0.2275	0.5860	0.2483	0.5514
CG30477	CG30477	1634475_at	-0.0563	0.7178	-0.1055	0.4882	-0.1201	0.5740	0.1158	0.8358	0.0900	0.7123	-0.0258	0.9209	-0.0880	0.8769	-0.0948	0.7050	-0.0069	0.9849
---	---	1634476_at	0.0953	0.6096	0.0818	0.5755	0.3593	0.1068	0.1588	0.6615	0.0295	0.9019	-0.1292	0.4285	-0.2463	0.7464	-0.0941	0.8234	0.1522	0.6630
CG31177	CG31177	1634477_at	0.5573	0.1091	0.1967	0.2298	1.1146	0.0008	-0.0506	0.9604	-0.2245	0.4277	-0.1739	0.5103	-0.5625	0.6557	-0.3174	0.5543	0.2451	0.6602
---	---	1634478_at	0.0515	0.7620	-0.0832	0.5004	0.1165	0.4724	-0.0329	0.9603	-0.0144	0.9562	0.0185	0.9328	-0.0299	0.9644	-0.0116	0.9689	0.0183	0.9431
---	---	1634479_at	0.0406	0.8464	-0.1462	0.3828	-0.3258	0.0595	0.1982	0.5902	0.2935	0.1148	0.0953	0.6144	0.0863	0.8692	0.0390	0.9021	-0.0473	0.8635
---	---	1634480_at	0.1337	0.3608	0.0454	0.6937	-0.0688	0.7032	0.1698	0.6936	0.1343	0.5272	-0.0354	0.8806	0.2637	0.4511	-0.0206	0.9445	-0.2843	0.1443
RhoGAP18B	RhoGAP18B	1634481_at	0.4481	0.4136	0.8085	0.0281	1.2417	0.0032	0.2410	0.6791	-0.4883	0.0754	-0.7293	0.0095	-0.2987	0.8540	-0.0960	0.9275	0.2027	0.8033
CG8918	CG8918	1634482_at	0.1072	0.5773	0.1767	0.3514	0.1817	0.2386	-0.0217	0.9838	-0.1336	0.6329	-0.1119	0.6671	-0.0870	0.9246	-0.0950	0.8117	-0.0080	0.9872
CG12584 /// CG34357	CG14652	1634483_at	0.2146	0.4626	0.0558	0.6302	0.2771	0.2714	-0.0436	0.9659	0.0818	0.8047	0.1253	0.6523	-0.1397	0.9016	0.0311	0.9639	0.1708	0.7154
---	---	1634484_at	0.0592	0.7764	0.2428	0.2386	-0.0234	0.9121	-0.1299	0.8218	-0.0791	0.7696	0.0507	0.8454	0.0980	0.8610	0.0261	0.9433	-0.0720	0.7919
---	---	1634485_at	0.0051	0.9784	-0.0354	0.7275	0.0974	0.6220	0.0627	0.9017	0.1989	0.2265	0.1362	0.3703	-0.1355	0.8192	0.1139	0.6716	0.2495	0.3154
CG30047	CG30047	1634486_at	-0.3204	0.9251	0.2172	0.4259	-0.8300	0.1181	-0.9109	0.8498	-2.0719	0.2371	-1.1610	0.4884	0.0876	0.9942	-1.2642	0.6772	-1.3519	0.6464
tau	tau	1634487_at	-1.0740	0.0044	-1.5207	0.0104	-2.4596	0.0000	-0.2050	0.6005	-0.0794	0.7306	0.1257	0.5085	0.6968	0.5790	-0.4194	0.4656	-1.1162	0.0852
CG33111 /// ofs	eIF-4G-like protein	1634488_s_at	-0.1607	0.7260	0.3425	0.3574	0.1788	0.4710	-0.2768	0.7231	-0.3485	0.3297	-0.0716	0.8616	-0.0350	0.9912	0.3041	0.6666	0.3392	0.6225
CG14816	CG14816	1634489_at	0.2502	0.3107	0.7250	0.0188	0.7623	0.0022	0.2254	0.6247	-0.1233	0.6268	-0.3488	0.0877	0.1144	0.8837	0.4276	0.1420	0.3132	0.3032
CG31835	CG31835	1634490_at	0.1404	0.5634	0.2467	0.3516	0.2571	0.2097	0.0300	0.9732	0.0890	0.7306	0.0590	0.8130	0.0764	0.9365	0.2071	0.5071	0.1307	0.6986
---	---	1634491_at	0.1716	0.3740	0.0971	0.4810	0.3208	0.1519	0.2226	0.6202	-0.0875	0.7399	-0.3101	0.1196	0.1127	0.8571	-0.1833	0.4828	-0.2960	0.2622
CG7597 /// DmirCG7597	CG7597	1634492_s_at	0.3688	0.4331	0.3709	0.0425	-0.1401	0.4195	-0.1403	0.7293	0.4831	0.0157	0.6234	0.0030	0.3742	0.7731	0.4995	0.3669	0.1254	0.8669
LysS	Lysozyme S	1634493_at	0.0388	0.8872	0.2834	0.3757	0.0668	0.7610	-0.0522	0.9523	-0.1931	0.4413	-0.1410	0.5535	0.1182	0.8882	0.0030	0.9973	-0.1152	0.7526

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG17648	CG17648	1634494_at	-0.2992	0.2758	0.1029	0.4889	-0.0692	0.6630	-0.1451	0.6338	-0.2499	0.0942	-0.1048	0.4577	0.0920	0.9246	0.1070	0.7951	0.0150	0.9769
lola	longitudinals abse	1634495_s_at	-0.6482	0.1854	0.2501	0.7346	-0.1414	0.6606	0.1928	0.8578	-0.1709	0.7001	-0.3637	0.2974	0.6781	0.6958	0.7897	0.2645	0.1116	0.9121
CG32232	CG32232	1634496_at	0.2897	0.0565	0.0692	0.5594	0.0646	0.6832	0.0111	0.9863	0.1285	0.4263	0.1174	0.4222	-0.0331	0.9646	-0.0474	0.8667	-0.0143	0.9613
---	---	1634497_at	0.1290	0.3500	0.0275	0.7982	0.0999	0.5014	-0.0045	0.9956	0.0739	0.7097	0.0784	0.6596	-0.0883	0.9235	-0.0677	0.8760	0.0206	0.9630
CG17152	CG17152	1634498_at	0.1426	0.4296	0.3789	0.0613	0.2052	0.4486	-0.1211	0.8153	-0.2342	0.2508	-0.1131	0.5767	0.1152	0.9168	0.0258	0.9671	-0.0894	0.8577
CG8565	CG8565	1634499_at	0.2608	0.1963	-0.0402	0.9108	-0.3024	0.3700	-0.2678	0.7118	0.3076	0.3598	0.5755	0.0548	-0.0656	0.9679	-0.1132	0.8471	-0.0476	0.9371
---	---	1634500_at	0.0087	0.9684	0.0188	0.9049	0.1935	0.3163	-0.0128	0.9860	0.0138	0.9578	0.0266	0.8999	-0.1224	0.8825	0.0201	0.9691	0.1425	0.6835
CG16979	CG16979	1634501_at	0.0284	0.9149	0.3783	0.4543	0.2687	0.1025	-0.0247	0.9658	0.2565	0.0924	0.2812	0.0444	0.0626	0.9752	0.5913	0.2181	0.5287	0.2994
CG13404	CG13404	1634502_at	-0.1074	0.5820	0.1749	0.3522	0.2563	0.1028	0.0344	0.9567	-0.2413	0.1550	-0.2758	0.0725	0.1035	0.8461	0.1687	0.4392	0.0652	0.8084
CG15695	CG15695	1634503_at	-1.4652	0.0009	-1.0745	0.0075	-0.2777	0.2342	0.3110	0.4238	0.5624	0.0171	0.2514	0.1900	-0.5008	0.5228	1.0396	0.0280	1.5404	0.0114
sol	small optic lobe	1634504_a_at	-0.5667	0.3332	-0.3566	0.8080	-0.1203	0.7939	-0.1314	0.7277	-0.5057	0.0094	-0.3743	0.0214	-0.1818	0.9778	-0.1814	0.9425	0.0003	0.9999
---	---	1634505_at	0.1621	0.2952	0.0649	0.6190	0.2178	0.2288	0.2630	0.4140	0.2162	0.2257	-0.0468	0.8171	0.1496	0.7633	0.1395	0.5176	-0.0101	0.9760
---	---	1634506_s_at	0.0104	0.9683	-0.0082	0.9498	0.0089	0.9614	-0.0395	0.9462	-0.1098	0.5335	-0.0703	0.6845	0.0906	0.9224	-0.0398	0.9353	-0.1304	0.7067
CG17549	CG17549	1634507_s_at	-0.7635	0.0037	-1.1867	0.0409	-0.7753	0.0024	0.1900	0.6010	-0.1046	0.6015	-0.2946	0.0735	-0.2284	0.8192	-0.5241	0.1865	-0.2957	0.4848
---	---	1634508_at	-0.1489	0.4062	-0.0938	0.4163	-0.0478	0.7995	0.0573	0.8905	0.1014	0.4876	0.0441	0.7750	-0.0432	0.9515	0.1203	0.5983	0.1635	0.4597
fwd	four wheel drive	1634509_s_at	0.5665	0.1267	1.2083	0.0266	0.8381	0.0038	0.1537	0.8196	0.2257	0.4059	0.0720	0.8083	0.4223	0.7366	0.8218	0.1268	0.3994	0.4697
CG17324	CG17324	1634510_at	0.1263	0.5225	-0.0034	0.9828	0.1401	0.3736	0.0612	0.9066	0.0078	0.9762	-0.0534	0.7733	-0.0266	0.9841	-0.0701	0.8690	-0.0436	0.9169
CG33125	CG33125	1634511_at	0.4386	0.0452	-0.0328	0.7465	0.0983	0.6146	0.1656	0.6197	0.3812	0.0294	0.2156	0.1454	-0.0502	0.9398	-0.0712	0.7741	-0.0210	0.9412
CG5226	CG5226	1634512_at	-1.1608	0.0099	-1.3089	0.0762	-1.5539	0.0048	0.2888	0.6615	0.4721	0.1343	0.1834	0.5569	0.2255	0.8692	-0.0579	0.9485	-0.2835	0.6318
---	---	1634513_at	0.1218	0.6941	-0.2948	0.1522	-0.4729	0.0702	-0.0097	0.9943	0.3554	0.2173	0.3651	0.1565	0.0184	0.9901	-0.0235	0.9650	-0.0418	0.9263
CG33258	CG33258	1634514_at	0.3078	0.9305	0.4413	0.4670	-4.1868	0.0002	-3.8485	0.2217	-1.3752	0.4864	2.4733	0.1408	0.6336	0.9514	-1.2762	0.7199	-1.9098	0.5612
CG8093 /// DyakCG8093	CG8093	1634515_at	0.0715	0.9535	0.0244	0.8433	0.0357	0.8749	0.0213	0.9931	-0.4588	0.3555	-0.4801	0.2755	0.0419	0.9928	-0.5740	0.6106	-0.6159	0.5834
---	---	1634516_at	0.1906	0.3113	0.2007	0.4612	0.0775	0.7155	-0.0424	0.9506	-0.0295	0.9105	0.0129	0.9573	0.1529	0.8222	0.1210	0.7059	-0.0319	0.9360
---	---	1634517_at	-0.3054	0.0646	-0.0791	0.5325	-0.0906	0.5793	-0.0249	0.9705	-0.1175	0.5212	-0.0926	0.5912	-0.0334	0.9589	-0.0302	0.9170	0.0032	0.9924
CG13886	CG13886	1634518_at	-0.1160	0.6670	0.0266	0.9221	0.3254	0.2284	-0.1745	0.8398	-1.1905	0.0037	-1.0161	0.0045	-0.2812	0.7848	-0.8438	0.0717	-0.5626	0.2202
---	---	1634519_at	0.4659	0.0541	0.1187	0.3736	0.7392	0.0064	-0.0079	0.9937	-0.1289	0.5543	-0.1209	0.5417	-0.4287	0.5651	-0.5491	0.1089	-0.1204	0.7567
vn	defective dorsal di	1634520_at	-0.8295	0.0424	-0.8209	0.2018	-1.7901	0.0001	-0.2849	0.7149	0.2391	0.5287	0.5240	0.0993	0.4051	0.8202	0.1504	0.8866	-0.2547	0.7640
zetaCOP	zetaCOP	1634521_at	-0.4243	0.1473	-0.1223	0.7968	-0.5165	0.0642	-0.0692	0.9380	0.2218	0.4079	0.2910	0.2114	0.2586	0.8423	0.3726	0.4953	0.1141	0.8736
Paps	paps synthetase	1634522_s_at	0.1781	0.7180	1.3977	0.0039	1.4348	0.0003	0.3449	0.2620	0.6801	0.0035	0.3352	0.0452	0.3234	0.8387	1.8569	0.0276	1.5335	0.0559
---	---	1634523_at	0.1562	0.4452	0.0498	0.8313	0.1960	0.2011	0.0866	0.8794	0.1786	0.3692	0.0921	0.6457	-0.0521	0.9653	0.1859	0.5860	0.2380	0.4756
DmirCG17149 /// Hdm	Flavoprotein /// C(	1634524_s_at	0.2324	0.3115	-0.0384	0.8799	-0.2378	0.2626	-0.0278	0.9774	0.8787	0.0039	0.9064	0.0021	0.0716	0.9515	0.3732	0.2740	0.3016	0.4016
CG7738	CG7738	1634525_at	-3.0007	0.0218	-3.0129	0.0129	-3.2764	0.0001	0.1386	0.9247	-0.3455	0.4480	-0.4840	0.2172	0.3242	0.9412	-0.4182	0.8127	-0.7424	0.6133
qua	quail	1634526_a_at	1.9742	0.0018	1.6361	0.0177	1.9022	0.0010	0.0390	0.9629	1.0664	0.0012	1.0273	0.0009	-0.2225	0.9142	0.6302	0.3758	0.8527	0.2557
CG40005 /// cta	concertina /// CG4	1634527_s_at	0.1271	0.6168	0.2681	0.3104	0.1671	0.4557	-0.2471	0.6754	0.1817	0.5423	0.4287	0.0848	-0.0260	0.9816	0.4344	0.1043	0.4604	0.1146
CG8412	CG8412	1634528_at	0.2002	0.2719	0.4546	0.4882	0.1855	0.3071	-0.0900	0.8307	0.1772	0.2729	0.2672	0.0664	0.1954	0.9149	0.4841	0.4505	0.2888	0.6835
CG3290	CG3290	1634529_at	-1.4547	0.5068	0.0819	0.8009	-0.6906	0.0766	-0.2433	0.9761	-1.6836	0.4193	-1.4403	0.4493	0.3077	0.8919	-0.5510	0.5329	-0.8587	0.3230
CG13040	CG13040	1634530_at	0.3261	0.3098	0.2315	0.3307	0.7370	0.0007	0.1650	0.7941	-0.1298	0.6556	-0.2948	0.2003	-0.3821	0.6749	-0.1414	0.7425	0.2407	0.5332
CG8750	CG8750	1634531_at	-0.0923	0.7260	0.0454	0.7344	-0.0355	0.8259	0.0378	0.9532	-0.0053	0.9838	-0.0432	0.8305	0.1332	0.8465	0.2102	0.4694	0.0770	0.8307
---	---	1634532_at	0.1194	0.5430	-0.0724	0.5344	-0.1221	0.4814	0.0713	0.8885	0.1553	0.3738	0.0840	0.6303	0.1755	0.7485	0.1869	0.4215	0.0114	0.9757
UbcD10	Ubiquitin conjugat	1634533_at	-0.4216	0.1543	-0.2122	0.4922	0.2454	0.2297	0.1920	0.6405	-0.2589	0.1951	-0.4509	0.0203	-0.2611	0.7739	0.1218	0.8119	0.3829	0.3457
mei-W68	meiotic W68	1634534_at	0.1425	0.3834	-0.1160	0.5905	0.0689	0.6952	0.2220	0.5978	0.3638	0.0878	0.1418	0.4848	-0.0287	0.9677	-0.0011	0.9989	0.0276	0.9158
---	---	1634535_at	0.1477	0.4781	0.0627	0.7224	-0.1094	0.4619	0.0602	0.9086	0.2488	0.1356	0.1887	0.2072	0.0724	0.9199	0.0381	0.9198	-0.0343	0.9189
CG6885	CG6885	1634536_at	0.3764	0.0532	0.0455	0.7261	0.0576	0.7859	0.0320	0.9593	0.2079	0.2065	0.1760	0.2341	-0.0750	0.9246	-0.0848	0.8016	-0.0097	0.9835
CG12502	CG12502	1634537_at	0.1747	0.3730	0.2237	0.4402	0.2579	0.1607	0.1257	0.8089	0.0322	0.9108	-0.0935	0.6630	0.1075	0.9277	0.0684	0.9084	-0.0391	0.9431
---	---	1634538_at	0.0530	0.8138	0.0175	0.9188	0.0024	0.9924	-0.0757	0.8636	0.0377	0.8528	0.1134	0.4407	0.1329	0.8650	0.2224	0.4856	0.0895	0.8178
---	---	1634539_at	-0.0368	0.8099	0.0214	0.8329	0.1510	0.3399	-0.0573	0.9138	-0.0317	0.8883	0.0255	0.8991	-0.0729	0.8973	-0.0560	0.8445	0.0169	0.9548
Cyp316a1	Cyp316a1	1634540_at	0.0174	0.9306	0.0009	0.9950	0.2460	0.1634	0.0175	0.9819	-0.0019	0.9943	-0.0194	0.9308	-0.1215	0.8395	0.0665	0.8420	0.1880	0.4617
CG3271	CG3271	1634541_a_at	-0.1566	0.3120	-0.3777	0.2851	-0.2434	0.2378	0.0341	0.9592	0.4835	0.0136	0.4494	0.0116	0.0161	0.9914	0.3116	0.3259	0.2955	0.3764
Sc2	Ubiquitin-63E	1634542_at	0.7046	0.0230	-0.0341	0.9078	0.1007	0.6021	0.2166	0.5301	0.4823	0.0154	0.2657	0.0981	0.2099	0.8318	-0.2458	0.5711	-0.4557	0.2835
CG9203 /// DmirCG9203	CG9203	1634543_at	-0.3498	0.0397	-0.5675	0.0223	-0.8020	0.0013	-0.2984	0.3016	0.3506	0.0450	0.6490	0.0019	-0.1125	0.8461	-0.0152	0.9692	0.0974	0.7179
---	---	1634544_at	0.0646	0.7018	-0.0158	0.9375	0.0524	0.8082	-0.0729	0.9339	-0.0700	0.8309	0.0029	0.9928	-0.1729	0.7707	-0.0106	0.9828	0.1623	0.5439
CG41465	CG41465	1634545_at	0.0193	0.9309	0.0810	0.4809	-0.2498	0.1606	-0.2789	0.2785	-0.1568	0.3054	0.1220	0.3853	0.1521	0.8141	0.0067	0.9915	-0.1454	0.6141
Tig	tiggrin	1634546_at	-0.1147	0.8922	-0.1703	0.8452	-0.0142	0.9595	-0.0197	0.9932	0.9261	0.0475	0.9458	0.0280	-0.1170	0.9816	0.8210	0.4475	0.9379	0.3921
CG32506	CG32506	1634547_at	-0.1801	0.4504	-0.0661	0.8396	-0.1983	0.2575	-0.0913	0.9093	-0.1231	0.6714	-0.0318	0.9191	0.0592	0.9653	-0.0372	0.9501	-0.0964	0.8354
CG10029	CG10029	1634548_at	0.2209	0.1932	0.1187	0.5103	0.0897	0.6560	0.0955	0.8909	0.0306	0.9253	-0.0649	0.8052	-0.0469	0.9330	-0.0241	0.9326	0.0228	0.9264
CG11750	CG11750	1634549_at	-0.5334	0.3094	-0.0582	0.8564	-0.4103	0.0297	0.1240	0.9319	0.2034	0.6800	0.0794	0.8773	0.4841	0.5754	0.6045	0.1212	0.1204	0.7964
hyx	hyrax	1634550_at	-0.1801	0.4747	-0.0613	0.7445	0.2399	0.1495	0.1914	0.6506	-0.2377	0.2455	-0.4292	0.0268	-0.1535	0.8461	-0.0432	0.9347	0.1103	0.7755
CG6188	CG6188	1634551_at	3.4704	0.0439	1.2621	0.5637	2.8456	0.0024	0.1428	0.9722	1.4824	0.1451	1.3396	0.1411	-1.3319	0.8446	-0.6716	0.8631	0.6603	0.8555
TepIV	Thiolester contain	1634552_at	-0.7030	0.0482	-0.5203	0.1960	-0.7080	0.0043	-0.2610	0.3420	-0.0691	0.7046	0.1918	0.1737	-0.0849	0.9659	0.1404	0.8474	0.2253	0.7085
CG13602	CG13602	1634553_at	-0.2867	0.2818	-0.7601	0.1584	-0.7045	0.0322	0.2558	0.8304	0.7392	0.1048	0.4834	0.2378	0.0943	0.8608	0.0705	0.7947	-0.0237	0.9371
GstD8	Glutathione S tran	1634554_at	0.1852	0.2861	0.1149	0.5978	0.1492	0.4226	-0.0805	0.8990	-0.0428	0.8765	0.0377	0.8782	0.0022	0.9987	0.1353	0.6063	0.1330	0.6152
CG11033	CG11033	1634555_at	0.0579	0.8626	-0.3227	0.0543	-0.5886	0.0090	0.0807	0.9158	0.5112	0.0387	0.4305	0.0488	0.3090	0.6749	0.1697	0.6060	-0.1393	0.6817
---	---	1634556_at	-0.1245	0.6503	0.0842	0.5189	0.1456	0.5811	-0.0886	0.9023	-0.0731	0.7990	0.0155	0.9586	-0.0075	0.9964	0.2277	0.5855	0.2352	0.5746
Rhp	Rhophilin	1634557_at	-0.3704	0.0489	0.1338	0.7512	0.6155	0.0203	0.1453	0.7067	-0.3131	0.0781	-0.4584	0.0109	-0.3034	0.7746	0.2423	0.6241	0.5457	0.2525
egh	zeste-white 4	1634558_s_at	0.5514	0.0769	0.3392	0.3703	0.6241	0.0275	0.1178	0.8650	0.5385	0.0352	0.4207	0.0592	-0.0113	0.9952	0.4212	0.3259	0.4325	0.3386
CG7213	CG7213	1634559_at	0.0236	0.8939	0.0789	0.6064	0.1115	0.5677	0.0675	0.9036	-0.0952	0.6381	-0.1627	0.3241	-0.0007	0.9998	-0.1322	0.6248	-0.1316	0.6263
140up	upstream of Rpll1	1634560_at	-0.5099	0.0527	0.1232	0.7594	-0.0034	0.9916	-0.4889	0.1883	-0.6602	0.0103	-0.1714	0.4155	-0.3941	0.7464	-0.1957	0.7453	0.1984	0.7362
Tango5	Transport and Gol	1634561_s_at	-0.4107	0.1832	1.1207	0.0310	1.6106	0.0003	0.1462	0.8742	-1.0081	0.0078	-1.1543	0.0025	-0.3511	0.7092	0.6121	0.1233	0.9632	0.0523
Vrp1	solitary	1634562_s_at	0.2144	0.6274	-0.0132	0.9853	-0.2826	0.1579	-0.0048	0.9949	0.2165	0.1521	0.2213	0.1024	0.1599	0.9421	-0.1414	0.8852	-0.3013	0.6877

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG3764	CG3764	1634563_at	0.4981	0.0188	-0.0530	0.8307	-0.4383	0.0288	-0.0630	0.9218	0.3134	0.1042	0.3764	0.0364	0.1304	0.8541	-0.2475	0.3899	-0.3778	0.2132
---	---	1634564_at	0.0447	0.8056	-0.0016	0.9909	-0.0731	0.7038	0.0168	0.9807	0.0756	0.6990	0.0588	0.7501	0.0436	0.9672	-0.0707	0.8543	-0.1144	0.7163
CG4957	CG4957	1634565_at	0.1569	0.3602	-0.0072	0.9854	-0.0573	0.7859	-0.2429	0.6010	-0.1027	0.7034	0.1401	0.5402	-0.1061	0.9063	-0.1819	0.5972	-0.0758	0.8577
---	---	1634566_at	-0.0536	0.7903	0.1610	0.2947	0.2759	0.1277	-0.0629	0.8950	-0.2365	0.1343	-0.1736	0.2213	-0.0107	0.9913	0.1126	0.5972	0.1234	0.5612
gce	germ cell-express	1634567_at	-0.6631	0.1785	0.7953	0.1820	0.9546	0.0004	0.0388	0.9777	-1.3311	0.0030	-1.3698	0.0016	0.0124	0.9959	0.2166	0.7406	0.2042	0.7505
Cpr67Fa2	CG18349	1634568_at	0.0383	0.8289	0.0326	0.8423	0.1681	0.2784	0.0362	0.9580	0.0589	0.7908	0.0228	0.9187	-0.1082	0.8875	0.0231	0.9617	0.1313	0.6761
CG34051	CG34051	1634569_at	0.1125	0.5015	-0.0880	0.4893	0.1403	0.4149	0.2644	0.4136	0.2822	0.1142	0.0179	0.9362	0.0059	0.9952	0.0868	0.7506	0.0810	0.7626
CG5375 /// DyakCG5375	CG5375	1634570_at	0.0585	0.7708	-0.0501	0.8525	0.2127	0.2116	0.0262	0.9666	0.0686	0.7193	0.0424	0.8195	-0.1965	0.7726	-0.0509	0.9085	0.1456	0.6414
---	---	1634571_at	-0.2252	0.2573	0.0211	0.8400	0.0008	0.9972	0.0090	0.9937	-0.0845	0.7446	-0.0936	0.6835	0.0331	0.9616	0.0332	0.9098	0.0002	0.9997
CG6459	CG6459	1634572_at	0.0284	0.9071	-0.2591	0.5170	0.1656	0.5202	0.1242	0.7718	0.3342	0.0671	0.2100	0.1923	-0.2799	0.8235	0.0945	0.9058	0.3744	0.4869
grh	Grainyhead	1634573_a_at	-0.8540	0.0449	-1.9693	0.0191	-2.5837	0.0000	-0.0405	0.9774	0.9920	0.0121	1.0325	0.0061	0.4056	0.7464	-0.3591	0.5123	-0.7647	0.1734
Strn-Mlck	Stretchin	1634574_a_at	-1.3688	0.0283	-1.2334	0.0223	-1.2473	0.0003	-0.1564	0.7950	0.0491	0.8810	0.2055	0.3634	-0.0691	0.9816	0.2916	0.6930	0.3607	0.6106
CG11037	CG11037	1634575_at	0.3809	0.0400	0.0993	0.5664	0.1225	0.4318	0.1169	0.7409	0.1211	0.4540	0.0042	0.9831	0.1363	0.8093	0.1040	0.6937	-0.0323	0.9199
N	Notch	1634576_at	-2.6055	0.0009	-0.7998	0.2515	-2.3764	0.0005	-1.3842	0.0492	-1.8295	0.0015	-0.4452	0.2176	-0.1732	0.9101	-0.4153	0.4483	-0.2421	0.6886
CG8974	CG8974	1634577_s_at	-0.0199	0.9250	0.1073	0.5978	-0.1194	0.5609	-0.0391	0.9422	-0.3058	0.0516	-0.2667	0.0564	0.1588	0.8714	-0.2039	0.6270	-0.3628	0.3642
CG30419	CG30419	1634578_at	0.3342	0.1624	0.2464	0.1035	0.3062	0.0619	-0.0163	0.9857	-0.1433	0.5045	-0.1270	0.5199	-0.0230	0.9816	-0.1004	0.6868	-0.0774	0.7654
CG4747	CG4747	1634579_at	0.5048	0.1165	1.2426	0.0073	0.8961	0.0025	-0.3296	0.5507	-0.0729	0.8437	0.2567	0.3246	0.0206	0.9893	0.6442	0.0637	0.6236	0.0859
bsk	Jun kinase	1634580_at	0.3206	0.1772	0.0720	0.7927	0.0341	0.8938	-0.0579	0.9247	0.4143	0.0316	0.4722	0.0115	0.0756	0.9653	0.1904	0.7200	0.1148	0.8472
---	---	1634581_at	-0.0215	0.9125	0.0968	0.4505	0.0179	0.9467	-0.0107	0.9926	0.0865	0.7593	0.0972	0.6948	0.0233	0.9826	0.1260	0.6380	0.1027	0.7128
CG32736	CG32736	1634582_at	-0.1962	0.3855	0.1020	0.5694	-0.0881	0.7281	-0.1356	0.7576	-0.3435	0.0718	-0.2079	0.2186	0.2184	0.7752	0.0010	0.9996	-0.2173	0.5332
CG4822	CG4822	1634583_s_at	1.8286	0.0091	-0.0172	0.9868	0.5025	0.2043	-0.6185	0.4022	0.1538	0.7546	0.7723	0.0425	-1.1352	0.5967	-1.6593	0.0890	-0.5242	0.6052
CG31875	CG31875	1634584_at	0.1004	0.6676	-0.2247	0.2320	-0.3229	0.2719	-0.0847	0.8899	0.4759	0.0278	0.5606	0.0085	-0.1130	0.9457	-0.0358	0.9652	0.0773	0.9109
CG13575 /// DmirCG13575	CG13575	1634585_at	0.2559	0.2746	0.0520	0.6026	-0.1409	0.3815	-0.1137	0.7278	-0.0342	0.8581	0.0795	0.5864	0.0072	0.9933	-0.0697	0.7734	-0.0769	0.7355
pip	pipe	1634586_at	0.2584	0.2143	-0.0617	0.8741	0.1400	0.5655	-0.1340	0.8770	-0.1164	0.7399	0.0176	0.9626	-0.1396	0.9112	-0.3752	0.3922	-0.2356	0.6225
Mnf	Mnf	1634587_s_at	-0.3279	0.2941	0.1486	0.5765	-0.5576	0.1124	-0.1934	0.6886	-0.5208	0.0277	-0.3274	0.1013	0.5466	0.7154	-0.0566	0.9557	-0.6032	0.3497
Sema-1a	semiphorin	1634588_at	0.0585	0.6818	-0.1474	0.2592	-0.1207	0.4399	0.0743	0.8578	0.3138	0.0425	0.2395	0.0755	0.0777	0.9156	0.0600	0.8679	-0.0177	0.9622
CG11152 /// DsimCG11152	CG11152	1634589_at	0.1667	0.3127	0.1346	0.5132	0.1757	0.3648	0.2054	0.7138	0.1425	0.6124	-0.0629	0.8291	0.1037	0.8655	0.1033	0.7090	-0.0004	0.9992
CG13885	CG13885	1634590_at	-0.0230	0.9354	0.2568	0.3503	0.3985	0.0722	-0.0276	0.9759	-0.0684	0.8044	-0.0408	0.8782	0.0002	0.9999	0.1449	0.6915	0.1447	0.6877
CG11878	CG11878	1634591_at	-0.1755	0.9551	-1.4648	0.0290	-1.6044	0.0422	0.4824	0.8215	-0.3192	0.7469	-0.8017	0.2880	0.3888	0.9756	-1.8546	0.5706	-2.2434	0.4851
ECSIT	ECSIT	1634592_at	-0.1777	0.4588	0.1202	0.7161	-0.2242	0.2083	-0.0650	0.9116	0.0470	0.8443	0.1121	0.5353	0.2400	0.7707	0.3462	0.3134	0.1062	0.8052
CG40228	CG40228	1634593_at	-0.1236	0.5552	0.2239	0.3791	-0.2076	0.1818	-0.3092	0.5735	-0.3073	0.2783	0.0019	0.9958	0.1697	0.7392	0.0578	0.8439	-0.1119	0.6317
CG17738	CG17738	1634594_at	0.3815	0.2382	-0.0447	0.7819	-0.0130	0.9640	0.2382	0.6063	0.1862	0.4395	-0.0520	0.8469	0.0388	0.9816	-0.2377	0.5329	-0.2764	0.4669
CG10934	CG10934	1634595_at	-0.1128	0.4288	-0.0856	0.4945	0.1249	0.4138	-0.0704	0.8707	-0.2132	0.1573	-0.1427	0.2982	-0.3332	0.5074	-0.1730	0.4753	0.1603	0.5196
---	---	1634596_at	-0.0742	0.7378	0.0617	0.6081	0.1105	0.5231	0.0208	0.9777	-0.1408	0.4649	-0.1616	0.3390	-0.0774	0.9296	0.0943	0.7903	0.1717	0.5676
CG33970	CG33970	1634597_a_at	-0.4474	0.5350	-1.8109	0.0425	-1.6991	0.0003	0.0040	0.9956	0.2719	0.2343	0.2679	0.1893	-0.0213	0.9964	-1.1336	0.2775	-1.1123	0.3139
CG1599	CG1599	1634598_a_at	0.0211	0.9100	-0.3605	0.0442	-0.4376	0.0331	0.1691	0.6197	0.4916	0.0109	0.3225	0.0399	0.1296	0.8472	0.0565	0.8903	-0.0732	0.8372
ctp	dynein	1634599_s_at	-1.3203	0.0075	-0.3304	0.2938	-0.3791	0.0910	0.2808	0.6219	-0.1516	0.6308	-0.4324	0.0878	0.1975	0.8705	0.7898	0.1084	0.5923	0.2350
CG5347	CG5347	1634600_at	-0.0871	0.6193	-0.0107	0.9229	0.1677	0.4223	-0.1297	0.7803	0.0739	0.7469	0.2036	0.2394	-0.2189	0.7215	0.0257	0.9505	0.2447	0.3569
CG9684	CG9684	1634601_at	-0.1396	0.7507	-0.9504	0.0082	-0.9993	0.0002	-0.1275	0.7225	0.7974	0.0011	0.9250	0.0004	-0.1493	0.9220	-0.0112	0.9925	0.1380	0.8287
---	---	1634602_at	0.1416	0.5661	-0.0109	0.9187	-0.0093	0.9751	-0.0621	0.9098	-0.0351	0.8800	0.0270	0.8975	0.0545	0.9530	-0.1845	0.5191	-0.2390	0.3953
---	---	1634603_at	0.3228	0.2420	0.2698	0.0628	0.2143	0.2806	0.0327	0.9744	0.0842	0.7793	0.0515	0.8588	0.0614	0.9333	-0.0005	0.9998	-0.0619	0.8354
mit(1)15	Zeste-White 10	1634604_at	0.0436	0.8969	0.6309	0.0138	0.7173	0.0021	0.1403	0.7230	0.1343	0.4722	-0.0060	0.9787	0.1617	0.8802	0.8147	0.0664	0.6529	0.1399
---	---	1634605_at	0.2885	0.1590	-0.0536	0.8012	-0.0054	0.9831	0.2455	0.5038	0.3624	0.0657	0.1169	0.5387	0.0213	0.9872	-0.0880	0.8250	-0.1093	0.7541
mmy	cystic	1634606_a_at	0.2730	0.1618	0.3335	0.0270	0.5328	0.0214	0.0050	0.9956	-0.3669	0.1063	-0.3719	0.0704	-0.2228	0.7436	-0.3055	0.2714	-0.0827	0.8178
CG14160	CG14160	1634607_at	0.1867	0.4501	0.2400	0.3225	0.3622	0.0785	-0.0631	0.9538	-0.0196	0.9663	0.0435	0.9034	-0.1187	0.8202	-0.0837	0.7381	0.0350	0.9057
CG30383	CG30383	1634608_at	0.0741	0.7573	-0.1680	0.4109	0.1901	0.3802	0.0530	0.9567	0.3008	0.2534	0.2478	0.2981	-0.1955	0.7230	-0.0541	0.8734	0.1413	0.5767
CG31111	CG31111	1634609_at	0.1878	0.2189	0.2319	0.2610	0.3159	0.1293	0.0172	0.9819	0.2095	0.2501	0.1922	0.2377	-0.1134	0.8446	0.1411	0.5729	0.2545	0.2964
---	---	1634610_at	0.7729	0.0384	0.1934	0.4591	0.0518	0.9356	-0.3815	0.3772	0.4447	0.0714	0.8262	0.0033	0.0299	0.9929	-0.1090	0.9325	-0.1389	0.9004
CG34370	CG13499	1634611_at	0.4682	0.0849	0.0803	0.5281	0.2997	0.1463	0.2969	0.5593	0.2663	0.3126	-0.0305	0.9240	0.1132	0.8643	-0.0420	0.9184	-0.1552	0.5814
CG14310	CG14310	1634612_a_at	0.2474	0.2521	0.1174	0.4366	0.1702	0.3621	0.0352	0.9620	0.0349	0.8966	-0.0003	0.9991	-0.2457	0.6749	-0.1041	0.7007	0.1416	0.5794
CG40002	CG40002	1634613_s_at	-0.1565	0.6097	0.0062	0.9823	-0.0205	0.9474	-0.1992	0.7959	-0.3195	0.3125	-0.1203	0.7183	-0.1940	0.8521	-0.1699	0.7345	0.0241	0.9709
hfw	singed wings	1634614_at	-0.1124	0.5260	0.2187	0.1573	0.4037	0.0331	0.0593	0.9371	-0.4803	0.0347	-0.5396	0.0135	-0.2095	0.7215	-0.0022	0.9976	0.2073	0.4114
CG8509	CG8509	1634615_at	0.1660	0.3271	0.0917	0.4829	0.2789	0.2903	0.0176	0.9803	-0.0749	0.7083	-0.0926	0.5944	0.0512	0.9689	0.0811	0.8678	0.0300	0.9510
TfIIEbeta	Transcription factc	1634616_at	0.0150	0.9588	-0.0464	0.8767	-0.1013	0.5510	-0.0637	0.9314	0.1056	0.6717	0.1693	0.4068	0.1386	0.8940	0.0382	0.9507	-0.1004	0.8380
ImpE1	Y1-like	1634617_at	0.1837	0.3796	0.0000	1.0000	0.3860	0.0653	0.0952	0.8478	0.0275	0.9137	-0.0677	0.7269	-0.2512	0.7220	-0.1196	0.7311	0.1316	0.6925
---	---	1634618_at	-0.0268	0.8601	0.0220	0.8278	-0.0216	0.9162	-0.0350	0.9527	-0.2348	0.1481	-0.1997	0.1697	-0.1219	0.8650	-0.1655	0.5855	-0.0436	0.9121
IP3K1	Inositol 1,4,5-triph	1634619_at	-0.0105	0.9673	-0.1648	0.6971	-0.5109	0.2985	-0.1921	0.8132	-0.0380	0.9330	0.1541	0.6390	0.1403	0.9530	-0.1613	0.8739	-0.3016	0.7053
stl	stall	1634620_a_at	2.6398	0.0073	1.7427	0.0287	2.7770	0.0001	0.5159	0.3433	0.4513	0.1485	-0.0646	0.8599	-0.4714	0.8400	-0.3909	0.7256	0.0806	0.9525
CG13599	CG13599	1634621_at	0.0760	0.7730	-0.0749	0.5909	-0.1164	0.6533	-0.0767	0.9017	0.0869	0.7094	0.1636	0.3815	0.1464	0.8973	0.0951	0.8754	-0.0513	0.9313
---	---	1634622_at	0.2157	0.1449	0.1542	0.4248	0.0282	0.9557	-0.1966	0.7753	-0.1390	0.6724	0.0576	0.8639	0.0552	0.9717	-0.0913	0.8687	-0.1465	0.7457
Cyp6a14	Cyp6a14	1634623_a_at	2.6024	0.0005	1.2723	0.0393	1.7116	0.0004	0.3179	0.4455	0.2403	0.2936	-0.0776	0.7542	-0.0538	0.9835	-0.6960	0.2336	-0.6422	0.2993
CG16719	CG16719	1634624_at	-0.0736	0.6249	0.0252	0.9227	0.0121	0.9570	-0.0170	0.9852	-0.1158	0.6042	-0.0988	0.6360	0.0615	0.9357	-0.0295	0.9402	-0.0909	0.7442
Dnz1	DNZDHHC/NEW1	1634625_at	-0.7958	0.0233	-0.8440	0.0405	-1.1856	0.0001	-0.3551	0.5304	0.0790	0.8341	0.4341	0.0996	-0.0536	0.9653	-0.0271	0.9608	0.0266	0.9548
CG14766	CG14766	1634626_at	-0.1038	0.5107	0.0025	0.9840	0.0957	0.5137	0.0862	0.8327	0.0007	0.9977	-0.0855	0.5796	-0.0805	0.8692	0.0142	0.9648	0.0947	0.6531
CG40311	CG40311	1634627_at	-0.2599	0.1344	0.2624	0.1510	-0.0112	0.9559	-0.1693	0.7278	-0.2491	0.2540	-0.0798	0.7327	0.0951	0.8837	0.1867	0.4505	0.0916	0.7484
CG30392 /// DwilCG30392	CG30392	1634628_at	-0.0972	0.6991	0.2865	0.2126	0.2483	0.2505	-0.0868	0.8578	-0.4040	0.0282	-0.3172	0.0474	0.1070	0.9391	0.1359	0.8062	0.0289	0.9639
CG13359	CG13359	1634629_at	0.1193	0.4556	0.1410	0.3538	-0.1122	0.5089	-0.1585	0.6698	-0.0332	0.8897	0.1253	0.4519	0.0743	0.8815	0.0566	0.8197	-0.0177	0.9471
CG5707	CG5707	1634630_at	1.0679	0.0035	1.3446	0.0421	2.0141	0.0000	0.1374	0.7235	-0.4614	0.0166	-0.5988	0.0031	-0.3472	0.7697	-0.0635	0.9379	0.2837	0.6016
CG3105	CG3105	1634631_at	-0.0375	0.9416	-0.7282	0.0096	-0.6429	0.0267	0.2500	0.6101	0.6390	0.0176	0.3890	0.0784	-0.0688	0.9816	-0.2268	0.7682	-0.1580	0.8460

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14416	CG14416	1634632_at	-0.1849	0.5532	0.1357	0.5762	0.3815	0.0416	0.1022	0.8856	-0.1411	0.5913	-0.2433	0.2629	0.1198	0.8930	0.3117	0.3349	0.1919	0.5874
---	---	1634633_s_at	1.5259	0.0372	0.3172	0.8037	0.7174	0.2095	-0.4648	0.6457	0.2999	0.5709	0.7647	0.0822	-0.8239	0.8122	-0.8646	0.5672	-0.0407	0.9856
---	---	1634634_at	0.0573	0.7131	0.2616	0.0695	0.1278	0.4713	-0.0628	0.9149	-0.0344	0.8906	0.0285	0.8977	0.0219	0.9778	0.0621	0.7927	0.0402	0.8744
---	---	1634635_at	0.4046	0.0483	0.0000	1.0000	0.1787	0.2762	0.0855	0.8578	0.1527	0.3829	0.0672	0.7116	-0.0058	0.9946	-0.0959	0.6519	-0.0901	0.6734
CG6426	CG6426	1634636_at	2.7684	0.0004	1.7151	0.0045	3.7170	0.0000	1.6293	0.0290	1.0058	0.0217	-0.6235	0.0870	-0.1179	0.8012	-0.0276	0.9282	0.0903	0.6719
dUTPase	Deoxyuridine triph	1634637_a_at	0.4314	0.4264	-0.1140	0.6518	0.1740	0.7427	-0.1020	0.9375	0.4447	0.2408	0.5467	0.1058	-0.4006	0.8655	-0.1369	0.9274	0.2637	0.8240
CG9025 /// DmirCG9025 ///	CG9025 /// GA21	1634638_at	0.0290	0.9273	0.0684	0.7814	0.0522	0.7353	-0.0831	0.8822	-0.1653	0.4016	-0.0822	0.6822	0.0003	0.9999	0.0666	0.8918	0.0663	0.8844
---	---	1634639_at	-0.0341	0.8810	0.1313	0.2730	0.0012	0.9959	0.0329	0.9774	0.0261	0.9513	-0.0068	0.9850	0.1284	0.8655	0.2250	0.4603	0.0966	0.7898
Cyp4ac1	Cyp4ac1	1634640_at	2.1578	0.0474	0.5895	0.6012	1.5134	0.0098	0.6188	0.4126	0.5946	0.1532	-0.0242	0.9635	-0.1662	0.9814	-0.9615	0.5414	-0.7953	0.6254
Cyt-b5	Cyt-b5	1634641_a_at	0.3637	0.0434	0.7641	0.0187	1.2599	0.0002	0.1777	0.6702	-0.5271	0.0159	-0.7048	0.0026	-0.1450	0.8160	0.0969	0.7505	0.2419	0.3616
---	---	1634642_at	0.1415	0.5045	0.0512	0.7923	0.1352	0.3623	0.0049	0.9956	0.0637	0.7722	0.0588	0.7710	-0.1303	0.8714	-0.0936	0.8226	0.0366	0.9346
CG12303	CG12303	1634643_at	0.2591	0.1232	0.0207	0.8363	0.1212	0.5529	-0.0058	0.9943	-0.0470	0.8363	-0.0411	0.8406	0.0169	0.9901	-0.1380	0.6811	-0.1549	0.6328
---	---	1634644_at	0.3072	0.1859	0.1897	0.2279	0.3285	0.0601	-0.1263	0.7649	-0.0054	0.9839	0.1209	0.4825	-0.0178	0.9869	0.0526	0.8821	0.0704	0.8183
CG31245	CG31245	1634645_at	-0.0545	0.7906	-0.0306	0.7948	0.1583	0.4596	-0.0510	0.9538	-0.1030	0.7121	-0.0519	0.8530	0.0219	0.9746	0.0701	0.7229	0.0481	0.8237
---	---	1634646_at	0.0243	0.9210	0.2361	0.4578	-0.0238	0.9030	-0.0943	0.8084	-0.0730	0.6792	0.0213	0.9097	0.2526	0.7726	0.2195	0.5852	-0.0331	0.9505
atms	antimeros	1634647_at	-0.1404	0.5399	-0.0015	0.9913	-0.1807	0.2289	-0.0608	0.9017	0.4757	0.0096	0.5365	0.0034	0.1235	0.8940	0.4719	0.1604	0.3484	0.3253
Cpr47Ec	CG9077	1634648_at	0.0656	0.8231	0.1812	0.2560	0.2411	0.1600	-0.1171	0.8498	-0.1722	0.4635	-0.0551	0.8296	-0.0585	0.9545	-0.0348	0.9429	0.0237	0.9563
mys	lethal myospheroi	1634649_at	-0.6553	0.0459	-0.1588	0.6020	-0.9406	0.0008	-0.3566	0.4128	-0.1237	0.6489	0.2329	0.2865	0.2574	0.7241	0.2940	0.3404	0.0366	0.9344
galectin	galectin	1634650_at	-0.1904	0.3231	-0.0880	0.6135	-0.1174	0.4031	-0.2566	0.4517	-0.2047	0.2712	0.0519	0.8035	-0.2268	0.6955	-0.1175	0.6626	0.1093	0.6875
Acp53C14c	Acp53C14c	1634651_at	0.2307	0.2535	0.1169	0.4984	0.1326	0.5119	0.1869	0.7984	0.2300	0.4561	0.0431	0.9039	0.1310	0.8461	0.1126	0.7241	-0.0184	0.9640
CG7656	CG7656	1634652_a_at	-0.6112	0.0216	-0.6199	0.0179	-0.6923	0.0156	-0.0208	0.9715	0.0165	0.9378	0.0373	0.8241	0.0143	0.9939	0.1508	0.7550	0.1365	0.7754
Ptp10D	Protein tyrosine pl	1634653_a_at	-0.6562	0.0944	0.1932	0.3638	-0.3346	0.2321	-0.2917	0.6919	-0.9805	0.0112	-0.6888	0.0313	0.1723	0.8882	0.0701	0.9222	-0.1023	0.8668
poe	pushover	1634654_at	0.0328	0.9220	-0.0827	0.6190	0.0831	0.6772	0.2738	0.5336	0.2611	0.2573	-0.0127	0.9653	0.1591	0.8874	0.1523	0.7642	-0.0068	0.9924
CG3894	dmNeural-like	1634655_at	-0.2225	0.2600	0.1156	0.5395	0.2971	0.1270	-0.0309	0.9687	-0.4927	0.0237	-0.4618	0.0194	-0.1708	0.7707	-0.0861	0.7787	0.0847	0.7741
---	---	1634656_s_at	-0.2064	0.2340	0.2953	0.1489	0.2020	0.2101	0.0708	0.8676	-0.3795	0.0199	-0.4503	0.0057	0.1902	0.7961	0.2334	0.4520	0.0432	0.9194
CG5011	CG5011	1634657_at	0.2783	0.7245	-0.1579	0.4457	-0.0974	0.6619	0.1437	0.9457	-0.0364	0.9689	-0.1800	0.7878	0.1311	0.9589	-0.3875	0.6218	-0.5186	0.4931
nemy	no extended mem	1634658_a_at	2.3732	0.0325	1.8456	0.0053	2.0853	0.0005	0.6488	0.5255	-0.3206	0.5812	-0.9693	0.0480	0.4790	0.8608	-0.8132	0.4639	-1.2922	0.2557
Dl	Overflow	1634659_at	-1.8284	0.0012	-2.0697	0.0223	-3.1211	0.0000	-0.3022	0.5680	0.0250	0.9513	0.3272	0.1764	0.7124	0.5978	-0.2094	0.7723	-0.9218	0.1477
mrt	martik	1634660_at	-2.1063	0.0011	-1.0452	0.1850	-0.6567	0.0017	-0.0698	0.9108	-1.2413	0.0004	-1.1715	0.0003	-0.2987	0.8589	-0.0114	0.9941	0.2873	0.7067
ImpE1	Y1-like	1634661_at	0.2007	0.2213	0.1940	0.3973	0.1209	0.7056	-0.0974	0.8605	-0.1214	0.5714	-0.0240	0.9215	0.0567	0.9522	0.0891	0.8070	0.0325	0.9348
Cyp313a1	Cyp313a1	1634662_at	0.0795	0.7713	0.0176	0.9290	-0.1846	0.3697	-0.2526	0.4139	0.0074	0.9780	0.2601	0.0898	-0.0087	0.9939	-0.2777	0.2507	-0.2690	0.2964
CG7206	CG7206	1634663_at	-0.1538	0.4475	-0.2480	0.2554	0.1371	0.5738	-0.0015	0.9985	-0.0751	0.6946	-0.0736	0.6709	-0.2350	0.8202	-0.2964	0.4997	-0.0614	0.9173
Cyp309a2	Cyp309a2	1634664_at	2.9329	0.0043	1.5609	0.1651	2.6093	0.0001	1.0157	0.1829	0.2843	0.5687	-0.7314	0.0779	-0.0195	0.9964	-1.0618	0.3417	-1.0423	0.3764
CG33476	CG33476	1634665_at	-0.0777	0.7468	0.0471	0.6615	-0.0571	0.7255	-0.0933	0.8571	0.1188	0.5527	0.2121	0.2009	-0.0366	0.9751	0.1910	0.5249	0.2276	0.4457
---	---	1634666_at	0.0507	0.8706	-0.2188	0.1669	-0.0869	0.6611	-0.0389	0.9569	0.0575	0.8101	0.0964	0.6248	-0.0349	0.9831	-0.2385	0.5479	-0.2036	0.6189
CG11099 /// DsecCG11099	CG11099	1634667_at	-1.9120	0.0013	-0.7432	0.0100	-1.3609	0.0003	-0.4750	0.2789	-1.2647	0.0011	-0.7897	0.0052	0.0750	0.9112	-0.2361	0.3090	-0.3111	0.2110
Ice	Ice	1634668_at	0.1772	0.5320	-0.1751	0.3710	-0.4582	0.0547	-0.0652	0.9435	0.1457	0.6117	0.2109	0.3818	0.1051	0.9330	-0.2027	0.6407	-0.3077	0.4568
CG32333 /// DmirCG32333	CG32333	1634669_at	-1.2793	0.0041	-0.8787	0.0830	-1.8727	0.0000	-0.5915	0.0660	-0.0896	0.6717	0.5018	0.0091	0.1341	0.8400	0.1251	0.6824	-0.0090	0.9846
CG32714	CG32714	1634670_at	-2.8331	0.0005	-1.4466	0.1792	-3.9627	0.0003	-1.5064	0.0342	-1.6475	0.0021	-0.1411	0.7377	1.0023	0.7215	-0.2990	0.8539	-1.3013	0.2842
CG41136	CG41136	1634671_a_at	-0.3255	0.1535	-1.0277	0.0182	-0.5227	0.0171	0.3447	0.5405	0.7037	0.0245	0.3590	0.1699	-0.0849	0.9280	-0.0560	0.9054	0.0289	0.9465
Srp68	Srp68	1634672_at	0.8123	0.0028	0.9877	0.0395	0.9195	0.0009	0.1591	0.6916	0.3757	0.0482	0.2166	0.1911	0.2011	0.7872	0.5180	0.1089	0.3169	0.3299
---	---	1634673_at	0.0174	0.9413	0.1677	0.3632	0.3045	0.1350	-0.0244	0.9819	-0.0083	0.9823	0.0160	0.9597	-0.0622	0.9401	0.1104	0.7012	0.1726	0.5163
CG9989	CG9989	1634674_at	2.4055	0.0070	-0.1311	0.8845	1.2445	0.1402	1.6846	0.2946	2.2245	0.0264	0.5399	0.5636	0.0181	0.9965	-0.4101	0.7445	-0.4281	0.7253
CG3774	CG3774	1634675_s_at	0.3120	0.2193	0.0272	0.9345	0.1654	0.3223	-0.0750	0.8897	-0.3851	0.0395	-0.3100	0.0590	-0.2213	0.8480	-0.6521	0.1634	-0.4308	0.3820
---	---	1634676_at	0.1294	0.4276	-0.0107	0.9437	0.0394	0.8517	0.0698	0.9255	0.1378	0.5704	0.0680	0.7868	0.0958	0.8956	0.0565	0.8858	-0.0393	0.9170
CG34145	CG34145	1634677_at	0.1056	0.5513	0.1083	0.5774	0.2148	0.1815	0.0423	0.9371	-0.0061	0.9800	-0.0484	0.7805	-0.1014	0.9092	0.0856	0.8389	0.1870	0.5756
CG13108	CG13108	1634678_at	-0.1083	0.5982	-0.1934	0.4193	-0.3853	0.0380	-0.1326	0.7658	0.0835	0.7025	0.2161	0.2019	0.0184	0.9875	-0.0472	0.9075	-0.0657	0.8472
CG33980	CG33980	1634679_at	0.2697	0.2237	0.1973	0.2329	0.0995	0.5399	-0.0925	0.8581	-0.0640	0.7768	0.0286	0.8991	0.0661	0.9400	0.0786	0.8248	0.0125	0.9769
---	---	1634680_at	0.0609	0.7300	0.1923	0.2271	0.4272	0.0305	0.0146	0.9819	-0.0921	0.5825	-0.1067	0.4660	-0.0561	0.9618	0.1387	0.7065	0.1948	0.5744
Ucp4B	Ucp4B	1634681_a_at	-0.0561	0.7141	0.1037	0.4099	0.2382	0.3646	0.1783	0.7949	-0.0089	0.9827	-0.1872	0.4834	-0.0434	0.9717	0.0554	0.9080	0.0989	0.7933
---	---	1634682_at	0.1180	0.5598	0.0541	0.6528	0.0135	0.9677	0.0855	0.8546	0.1039	0.5691	0.0184	0.9290	0.0743	0.9421	-0.0846	0.8425	-0.1589	0.6389
Mad	Mothers against D	1634683_at	0.4236	0.4243	1.4197	0.0138	0.9614	0.0010	-0.0372	0.9647	-0.3634	0.1023	-0.3262	0.1008	0.4154	0.7677	0.5924	0.3066	0.1770	0.8102
CG7272	CG7272	1634684_at	-1.3916	0.0026	-1.2790	0.0060	-1.2192	0.0007	-0.4180	0.1188	-0.6375	0.0028	-0.2195	0.1269	-0.5112	0.6660	-0.5954	0.1995	-0.0842	0.8984
RpL3	Ribosomal protein	1634685_at	0.3400	0.2622	0.2980	0.2178	0.6187	0.0299	0.1589	0.7851	0.1128	0.6807	-0.0461	0.8706	0.1066	0.9581	0.3899	0.5057	0.2833	0.6419
CG9109	CG9109	1634686_at	0.1495	0.5697	0.2934	0.1578	0.2496	0.2090	0.0613	0.9147	-0.0727	0.7288	-0.1339	0.4249	0.1241	0.8837	0.0340	0.9485	-0.0901	0.8237
CG15019	CG15019	1634687_at	0.4460	0.1074	0.5111	0.2496	0.7047	0.0079	0.0769	0.9249	-0.2400	0.3338	-0.3169	0.1460	-0.0642	0.9677	-0.1202	0.8256	-0.0561	0.9220
RabX5	RabX5	1634688_at	0.2630	0.1153	0.2616	0.1983	0.2135	0.1939	-0.0147	0.9883	-0.0498	0.8725	-0.0351	0.9005	0.0908	0.8795	0.0753	0.7943	-0.0155	0.9636
---	---	1634689_at	0.0294	0.9131	-0.0058	0.9745	0.0060	0.9843	-0.1821	0.6763	-0.1160	0.6066	0.0662	0.7701	-0.1654	0.8331	-0.1199	0.7598	0.0455	0.9197
---	---	1634690_at	0.1914	0.6916	-0.3215	0.5435	-0.4451	0.3538	-0.4250	0.3752	0.4861	0.0756	0.9111	0.0034	-0.2229	0.9352	-0.0560	0.9690	0.1669	0.8897
CG30020	CG30020	1634691_a_at	0.3645	0.0883	0.2146	0.3611	0.1025	0.5477	0.0326	0.9683	0.3251	0.1278	0.2925	0.1250	0.1289	0.8480	0.2081	0.4613	0.0791	0.8209
CG18336	CG18336	1634692_at	0.1186	0.8094	0.4155	0.1088	0.8427	0.0339	-0.1607	0.8350	-0.0845	0.8154	0.0762	0.8178	-0.2758	0.8270	0.2737	0.6293	0.5495	0.3071
CG5122	CG5122	1634693_at	-0.0811	0.6573	-0.1258	0.4154	0.0119	0.9625	0.4138	0.3778	0.2530	0.3491	-0.1608	0.5349	-0.0166	0.9839	0.0464	0.8569	0.0630	0.7749
CG12169	CG12169	1634694_at	0.0694	0.7595	-0.1282	0.4154	-0.0787	0.6569	-0.0706	0.9242	-0.1360	0.5706	-0.0655	0.7926	-0.1422	0.8395	-0.1254	0.7007	0.0168	0.9697
lds	factor 2	1634695_at	-0.7998	0.0791	-0.6218	0.0624	-0.8894	0.0006	-0.2244	0.5844	0.1285	0.5696	0.3529	0.0627	0.0313	0.9892	0.2548	0.6218	0.2235	0.6701
Nup133	CG6958	1634696_at	0.1500	0.3013	-0.3386	0.4986	-0.0902	0.6792	-0.1598	0.6338	0.3698	0.0321	0.5296	0.0041	-0.3007	0.8331	-0.0758	0.9366	0.2249	0.7437
CG32667	CG32667	1634697_at	2.2002	0.0005	1.7611	0.1065	3.3240	0.0001	0.8535	0.1109	-0.0597	0.8897	-0.9132	0.0069	-0.7980	0.7436	-0.6433	0.5501	0.1547	0.9127
---	---	1634698_at	0.0687	0.7700	0.1122	0.4416	0.0845	0.6552	0.0569	0.9062	0.0912	0.5901	0.0343	0.8489	0.0756	0.9148	0.1333	0.6154	0.0577	0.8582
CG11919 /// CG18003	CG11919 /// CG18	1634699_a_at	2.4713	0.0071	1.4772	0.0818	3.2126	0.0002	0.5591	0.6082	-0.9696	0.0781	-1.5287	0.0078	-0.9069	0.6898	-1.5084	0.1116	-0.6015	0.5412
CG8258	CG8258	1634700_at	0.1742	0.6692	-0.1335	0.3973	0.1356	0.7155	0.5465	0.3953	0.9316	0.0182	0.3851	0.2344	0.3563	0.7953	0.6401	0.2507	0.2838	0.6484

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG3713	CG3713	1634701_at	-0.1978	0.3984	-0.6300	0.0988	-0.6681	0.0231	0.0314	0.9803	0.6556	0.0408	0.6242	0.0317	0.0547	0.9408	-0.0996	0.6977	-0.1542	0.5163
CG7227	CG7227	1634702_at	1.7103	0.0633	0.9537	0.2500	1.6178	0.0023	0.1257	0.9168	0.3676	0.3321	0.2418	0.5005	-0.3151	0.9451	-0.1341	0.9519	0.1810	0.9245
---	---	1634703_at	0.1642	0.4311	0.0487	0.7122	0.1362	0.5422	-0.0923	0.9149	-0.0092	0.9816	0.0831	0.7826	-0.2057	0.7220	-0.0761	0.8123	0.1297	0.6263
ATP7	ATP7	1634704_at	-0.9163	0.0599	-0.1493	0.6841	-0.4237	0.2150	-0.2033	0.7121	-0.4715	0.0631	-0.2682	0.2313	-0.1403	0.9514	0.2243	0.7921	0.3645	0.6172
CG10341	CG10341	1634705_at	0.5538	0.1074	0.4469	0.2392	0.5307	0.0139	0.4785	0.2299	0.7297	0.0084	0.2512	0.2367	0.3492	0.7464	0.5385	0.2217	0.1893	0.7135
---	---	1634706_at	0.0683	0.6868	0.0627	0.6810	0.1476	0.3139	0.0092	0.9893	0.0718	0.6901	0.0626	0.7059	-0.0123	0.9913	0.0689	0.8218	0.0812	0.7659
Gfat1	Glutamine:fructos	1634707_s_at	1.1742	0.0043	0.4359	0.1746	0.9596	0.0156	0.1866	0.8424	0.3202	0.3691	0.1336	0.7188	-0.2684	0.8097	-0.3734	0.4184	-0.1050	0.8637
---	---	1634708_s_at	-1.6168	0.0018	-0.3892	0.2246	-0.3204	0.1998	-0.2615	0.7149	-1.0062	0.0083	-0.7446	0.0189	-0.2461	0.7485	-0.1858	0.5929	0.0603	0.8932
CG12402	CG12402	1634709_at	-0.0933	0.6249	-0.1561	0.4176	-0.0449	0.7914	-0.0231	0.9790	-0.2550	0.2417	-0.2319	0.2338	-0.1002	0.8680	-0.1594	0.5192	-0.0592	0.8470
---	---	1634710_at	0.1281	0.4504	-0.1773	0.3179	0.0696	0.8004	0.2598	0.4959	0.2824	0.1607	0.0226	0.9272	-0.0200	0.9898	0.0375	0.9451	0.0575	0.9020
---	---	1634711_at	0.1966	0.3305	-0.0240	0.8259	0.0906	0.5368	-0.1452	0.7461	-0.0700	0.7665	0.0752	0.7187	-0.1035	0.8744	-0.0569	0.8774	0.0466	0.8949
CLIP-190	CLIP-190	1634712_s_at	-0.4536	0.1525	-0.1137	0.8079	-0.2388	0.2549	-0.1625	0.7327	-0.0919	0.7070	0.0706	0.7612	-0.1344	0.9409	0.1631	0.8246	0.2974	0.6224
CG5532	CG5532	1634713_at	0.1164	0.6326	0.5715	0.0610	0.4321	0.0261	0.1342	0.7225	-0.5795	0.0053	-0.7137	0.0013	0.2010	0.8202	-0.1145	0.8061	-0.3155	0.3939
cactin	cactin	1634714_at	-0.4057	0.3310	-0.4644	0.3393	-0.0404	0.8522	0.3098	0.4902	0.3874	0.1044	0.0776	0.7687	-0.1080	0.9666	0.3600	0.6267	0.4679	0.5131
---	---	1634715_at	-0.3635	0.2517	-0.7235	0.0763	-0.6694	0.0129	0.2954	0.7507	0.7274	0.0743	0.4319	0.2325	-0.2014	0.8270	-0.1603	0.7111	0.0411	0.9387
CG32843	CG32843	1634716_s_at	-0.1690	0.4761	0.0118	0.9564	0.0209	0.9249	0.0723	0.9319	0.0403	0.9083	-0.0319	0.9175	0.0310	0.9717	0.0960	0.6994	0.0650	0.8114
CG17580	CG17580	1634717_at	0.0298	0.8958	0.0808	0.7275	0.2291	0.2721	-0.0535	0.9314	0.0133	0.9616	0.0668	0.7355	-0.1805	0.8194	0.0487	0.9233	0.2292	0.4944
CG7963	CG7963	1634718_at	-0.0693	0.6557	0.0085	0.9399	-0.0137	0.9579	-0.0544	0.9313	-0.2053	0.2716	-0.1509	0.3806	-0.0203	0.9869	-0.0135	0.9790	0.0068	0.9878
CG9920	CG9920	1634719_at	0.0282	0.8693	0.0355	0.8042	-0.0289	0.8905	-0.0401	0.9620	-0.0808	0.7629	-0.0408	0.8776	-0.0083	0.9916	-0.0437	0.8684	-0.0354	0.8906
CG11151 /// DyakCG11151	transcript 2 /// CG	1634720_at	-0.0677	0.8556	0.5385	0.0793	1.0351	0.0011	0.0238	0.9745	-1.3636	0.0002	-1.3875	0.0001	-0.5001	0.6955	-0.7538	0.1498	-0.2538	0.6634
kl-5	Y-linked flagellar	1634721_at	-0.0014	0.9957	-0.0902	0.5604	0.0210	0.9049	-0.0964	0.8535	-0.0895	0.6774	0.0069	0.9770	-0.1315	0.8122	-0.1507	0.5259	-0.0191	0.9525
mbf1	multiprotein bridgi	1634722_s_at	0.3153	0.3442	1.1347	0.0083	1.0760	0.0002	0.5470	0.1319	0.3483	0.1091	-0.1987	0.3172	0.5697	0.5126	1.1623	0.0280	0.5926	0.1808
CG14280	CG14280	1634723_at	-0.1753	0.4753	-0.0547	0.6913	-0.0199	0.9046	-0.0045	0.9956	-0.2184	0.3491	-0.2139	0.3037	-0.1829	0.7726	-0.2240	0.4067	-0.0412	0.9122
CG12499	CG12499	1634724_at	0.9213	0.0164	0.1781	0.7938	0.3483	0.1729	0.1140	0.9064	0.9277	0.0097	0.8138	0.0107	-0.0253	0.9923	0.3358	0.6086	0.3611	0.5808
beat-IV	beat-IV	1634725_at	0.0573	0.7787	-0.2599	0.0664	-0.6493	0.0024	-0.2013	0.6050	0.2816	0.1490	0.4829	0.0137	-0.1170	0.8270	-0.1133	0.6408	0.0037	0.9924
CG18478 /// CG31827	CG31827 /// CG1	1634726_s_at	0.2582	0.2023	0.0579	0.6033	0.1317	0.4365	-0.0106	0.9889	0.0604	0.7757	0.0709	0.7014	-0.0639	0.9576	-0.0556	0.9157	0.0083	0.9875
thr	anarchist	1634727_a_at	0.1107	0.8327	-0.8151	0.1896	-1.2023	0.0023	0.0251	0.9857	1.5168	0.0013	1.4916	0.0008	-0.0407	0.9893	0.0088	0.9949	0.0495	0.9571
CG17821	CG17821	1634728_at	0.1402	0.4879	0.0060	0.9672	-0.0099	0.9627	-0.0282	0.9666	-0.0871	0.6649	-0.0589	0.7614	0.0369	0.9742	-0.1925	0.5170	-0.2295	0.4357
l(2)37Cd	lethal (2) 37Cd	1634729_at	-0.1504	0.3538	-0.0398	0.8833	0.1383	0.5724	-0.2958	0.3171	-0.3963	0.0293	-0.1005	0.5540	-0.1902	0.7644	-0.0218	0.9608	0.1683	0.5499
---	---	1634730_s_at	0.1876	0.3334	-0.2231	0.1084	-0.0523	0.7925	0.0886	0.8180	0.1753	0.2458	0.0867	0.5609	-0.0006	0.9998	-0.1673	0.4483	-0.1667	0.4648
Cyp4p3	Cyp4p3	1634731_at	-1.1179	0.0207	-0.7654	0.1758	-0.9436	0.0023	-0.4469	0.4690	-0.1812	0.6244	0.2658	0.3904	-0.1794	0.9298	0.1935	0.8224	0.3729	0.5988
---	---	1634732_at	-0.0427	0.8961	0.0026	0.9831	-0.0001	0.9995	-0.2329	0.4553	-0.1435	0.4099	0.0893	0.5989	-0.0557	0.9599	0.0100	0.9877	0.0657	0.8779
CG3831 /// DmirCG3831	CG3831	1634733_at	-1.2267	0.0279	-2.9215	0.0123	-2.7839	0.0000	-0.1330	0.8512	0.7916	0.0085	0.9245	0.0025	-0.1943	0.9466	-0.8469	0.3209	-0.6526	0.4703
CG12636 /// CG33482	CG12636 /// DS07	1634734_at	0.1005	0.8003	0.1158	0.6117	0.2890	0.2212	-0.1011	0.8834	-0.2259	0.3436	-0.1248	0.5954	-0.3203	0.5134	-0.1118	0.6673	0.2085	0.3868
---	---	1634735_at	-0.0567	0.7573	0.0020	0.9905	0.0409	0.7944	0.0227	0.9801	-0.1491	0.5217	-0.1718	0.3988	0.0411	0.9467	0.0500	0.8422	0.0089	0.9762
hts	hu-li tai shao	1634736_at	0.4341	0.1794	-1.0766	0.1059	-1.0778	0.0402	0.0900	0.9252	1.6983	0.0005	1.6083	0.0004	-0.1835	0.9611	0.0119	0.9961	0.1954	0.8926
CG12753	CG12753	1634737_at	-0.5302	0.0113	-0.0457	0.7101	-0.2685	0.2103	-0.1227	0.7042	-0.2405	0.1048	-0.1177	0.3942	-0.0339	0.9742	0.3078	0.2168	0.3417	0.2056
CG31751	constitutive transc	1634738_s_at	0.9601	0.0110	0.8067	0.1001	0.8070	0.0117	-0.4054	0.3828	-0.1471	0.6109	0.2584	0.2758	-0.3414	0.7772	-0.3356	0.5401	0.0058	0.9951
Pfk	6-phosphofructoki	1634739_a_at	-0.4839	0.0419	-0.4523	0.2415	0.0294	0.8583	-0.1089	0.8623	-0.7601	0.0054	-0.6511	0.0065	-0.4615	0.6483	-0.7657	0.0741	-0.3041	0.4683
tth	toothreen	1634740_at	0.4992	0.3383	0.5998	0.1560	0.5823	0.0193	0.1472	0.8220	0.1952	0.4635	0.0480	0.8736	0.2340	0.9142	0.2925	0.7439	0.0585	0.9576
---	---	1634741_at	0.0880	0.6351	-0.0197	0.8447	0.0780	0.6146	-0.0284	0.9603	-0.0517	0.7793	-0.0233	0.8991	0.0245	0.9831	-0.0434	0.9175	-0.0679	0.8445
Sodh-2	Sorbitol dehydrog	1634742_at	-1.8311	0.0014	-2.0659	0.0286	-2.2378	0.0004	-0.0052	0.9961	-0.3051	0.3695	-0.2999	0.3236	0.1999	0.9161	-0.4743	0.4805	-0.6742	0.3170
CG15292	CG15292	1634743_at	0.2101	0.2461	0.0555	0.5572	0.2803	0.0699	-0.0430	0.9346	-0.0788	0.6517	-0.0359	0.8412	0.0060	0.9948	-0.0680	0.8017	-0.0741	0.7640
---	---	1634744_at	0.2646	0.4326	0.0274	0.8599	0.1038	0.5040	0.1559	0.7271	0.2265	0.2594	0.0706	0.7441	-0.2492	0.7597	-0.0395	0.9429	0.2098	0.5630
---	---	1634745_at	0.1486	0.3658	0.1139	0.5831	0.1852	0.3071	-0.0697	0.8921	-0.0595	0.7741	0.0101	0.9629	0.0097	0.9916	0.0070	0.9872	-0.0027	0.9945
CG3589	CG3589	1634746_at	0.1272	0.4866	0.5371	0.1642	0.8575	0.0009	-0.1233	0.7447	-0.7100	0.0020	-0.5868	0.0027	-0.2020	0.8207	-0.0116	0.9877	0.1904	0.6328
Ddr	Discoidin domain	1634747_at	0.0010	0.9964	-0.1285	0.4785	-0.0623	0.7404	0.1543	0.7482	-0.0274	0.9252	-0.1817	0.3444	-0.0924	0.8945	-0.1513	0.5847	-0.0590	0.8641
dwg	zeste-white 5	1634748_at	0.2421	0.2735	0.5525	0.2730	0.8378	0.0021	0.1974	0.7138	-0.5473	0.0328	-0.7447	0.0053	-0.0782	0.9679	-0.1306	0.8546	-0.0524	0.9431
CG32686	CG32686	1634749_a_at	0.1147	0.4501	0.0118	0.9304	0.1440	0.4430	0.1615	0.6015	0.0960	0.5695	-0.0655	0.6895	0.0723	0.8743	0.0425	0.8665	-0.0299	0.9045
CG34033	CG34033	1634750_at	0.1131	0.4829	0.0740	0.7531	0.1435	0.5148	-0.0379	0.9675	0.0416	0.9020	0.0795	0.7687	-0.2877	0.6483	-0.1107	0.7002	0.1770	0.5023
---	---	1634751_at	0.2574	0.2261	-0.0555	0.6988	0.1146	0.5105	0.1756	0.7039	0.2034	0.3470	0.0278	0.9148	-0.0153	0.9914	-0.0809	0.8539	-0.0657	0.8800
CG14085	CG14085	1634752_a_at	0.2092	0.4288	-0.0487	0.7846	0.0227	0.9338	0.0477	0.9676	0.1264	0.7260	0.0787	0.8226	0.1126	0.8796	-0.1540	0.6144	-0.2666	0.3631
CG5327	CG5327	1634753_at	-0.0856	0.6186	-0.2226	0.2713	-0.0684	0.7293	0.1262	0.8217	0.3045	0.1613	0.1783	0.3775	-0.0443	0.9635	0.0247	0.9533	0.0691	0.8368
---	---	1634754_at	0.4220	0.0510	0.5186	0.1026	0.3771	0.0517	-0.0908	0.8803	-0.0331	0.9080	0.0576	0.8039	0.0485	0.9521	0.0567	0.8684	0.0082	0.9841
CG31053	CG31053	1634755_at	0.1787	0.2824	0.3062	0.2656	0.0166	0.9686	-0.2647	0.3175	-0.1560	0.3185	0.1087	0.4575	-0.0094	0.9959	-0.0495	0.9416	-0.0401	0.9460
nAcRalpha-80B	Dalpha4	1634756_at	1.4489	0.0009	0.7595	0.0703	1.0109	0.0021	-0.0973	0.7772	-0.0060	0.9789	0.0914	0.5098	-0.0753	0.9619	-0.3242	0.4676	-0.2489	0.5988
Oda	gut feeling	1634757_a_at	-0.3316	0.1377	0.0613	0.7042	-0.5011	0.0122	-0.1134	0.7929	-0.2142	0.2234	-0.1008	0.5631	0.3157	0.5126	0.0900	0.7362	-0.2257	0.3360
CG34393	CG15405	1634758_at	0.0505	0.7708	0.0776	0.7862	-0.1504	0.4323	0.0705	0.9215	0.1948	0.3767	0.1243	0.5595	0.1355	0.8673	0.0638	0.8936	-0.0717	0.8667
---	---	1634759_at	0.3185	0.2180	-0.0213	0.8540	0.4071	0.0188	0.1637	0.7030	0.1973	0.3252	0.0335	0.8864	-0.0729	0.9101	-0.0827	0.7669	-0.0098	0.9793
---	---	1634760_at	0.1496	0.5493	0.0102	0.9521	0.0550	0.8033	0.1145	0.8645	0.1463	0.5696	0.0318	0.9124	-0.0708	0.9305	-0.1009	0.7435	-0.0301	0.9344
CG9472	CG9472	1634761_at	0.1689	0.3959	0.0663	0.5554	0.1775	0.3117	-0.0434	0.9397	-0.0499	0.8080	-0.0065	0.9751	-0.1539	0.8744	0.0176	0.9799	0.1714	0.6861
CG15096	CG15096	1634762_a_at	0.7069	0.0181	0.1910	0.5247	0.6100	0.1442	-0.2921	0.6905	-0.1552	0.6905	0.1370	0.7020	-0.8007	0.5754	-0.7945	0.2043	0.0062	0.9956
CG3769	CG3769	1634763_at	0.1442	0.4447	-0.0237	0.8672	0.2313	0.2263	0.0727	0.9011	0.2635	0.1613	0.1908	0.2623	-0.0293	0.9742	0.0483	0.8815	0.0775	0.7686
RpL23	Ribosomal protein	1634764_at	0.0457	0.7730	0.1418	0.2884	0.2630	0.0836	0.0092	0.9883	-0.1688	0.2556	-0.1779	0.1777	-0.0499	0.9342	-0.0328	0.9129	0.0171	0.9505
CG14739	CG14739	1634765_at	0.0187	0.9192	-0.2215	0.1525	-0.0715	0.7422	0.1754	0.6634	0.1561	0.4352	-0.0193	0.9361	0.1106	0.8717	0.0021	0.9981	-0.1084	0.7259
Ser	Beaded	1634766_at	-1.1418	0.1118	-1.6749	0.0073	-2.1875	0.0000	-0.3786	0.3381	-0.1320	0.5955	0.2465	0.2271	-0.1304	0.9542	-0.5448	0.4094	-0.4143	0.5579
CG6126	CG6126	1634767_at	0.8220	0.3686	-0.2691	0.3996	0.1904	0.5655	0.0405	0.9777	-0.4598	0.1995	-0.5002	0.1190	-0.4226	0.8979	-1.4898	0.2102	-1.0672	0.3921
CG12009	CG12009	1634768_at	0.2752	0.2386	0.0539	0.6457	0.3468	0.0446	0.0632	0.8905	0.0073	0.9757	-0.0559	0.7355	-0.1412	0.8202	-0.0295	0.9430	0.1117	0.6955
CG6888	CG6888	1634769_at	-0.0132	0.9612	0.0261	0.8767	-0.2580	0.0879	-0.2062	0.7018	0.0224	0.9517	0.2286	0.3092	0.0429	0.9672	-0.0384	0.9294	-0.0813	0.8109

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG32377	CG32377	1634770_at	-0.1081	0.6142	0.0449	0.7519	0.0973	0.5950	-0.0568	0.9436	-0.1344	0.5890	-0.0776	0.7564	-0.1388	0.7644	-0.0575	0.8218	0.0812	0.7130
CG8184	CG8184	1634771_a_at	1.2297	0.2560	0.5522	0.5113	-0.0311	0.9564	0.0073	0.9956	1.6377	0.0015	1.6304	0.0009	0.5978	0.8846	0.9152	0.5805	0.3174	0.8813
CG1288 /// DbuzCG1288	CG1288	1634772_at	0.2296	0.2328	-0.0430	0.8919	-0.0464	0.8577	-0.1130	0.9022	0.0770	0.8415	0.1900	0.5090	-0.1831	0.7464	-0.1851	0.4380	-0.0020	0.9965
SsRbeta	Signal sequence r	1634773_at	1.1894	0.0035	2.2680	0.0057	2.3569	0.0000	0.1224	0.7556	0.0590	0.7727	-0.0634	0.7262	-0.1545	0.8940	0.9610	0.0476	1.1155	0.0441
---	---	1634774_at	-0.0899	0.5619	-0.0775	0.5385	0.1209	0.4603	0.0662	0.8835	-0.0371	0.8525	-0.1033	0.4809	0.0636	0.9142	-0.0407	0.8942	-0.1043	0.6389
CG33235	CG33235	1634775_at	0.1246	0.5164	-0.2633	0.1175	-0.0987	0.5583	0.1339	0.7795	0.4291	0.0372	0.2953	0.0956	-0.0873	0.8940	-0.1244	0.6408	-0.0371	0.9121
CG7509	CG7509	1634776_at	0.2114	0.4107	0.0989	0.5842	0.0396	0.9036	-0.0346	0.9693	0.0870	0.7530	0.1216	0.6027	0.1453	0.9014	0.0271	0.9696	-0.1182	0.8261
CG13297	CG13297	1634777_at	0.0339	0.8527	0.0268	0.8976	-0.0284	0.8589	-0.0114	0.9880	-0.0991	0.6131	-0.0877	0.6285	0.0730	0.9238	0.0857	0.7846	0.0127	0.9746
CG3092 /// DsimCG3092	CG3092	1634778_at	0.2207	0.2824	0.2887	0.3823	0.2124	0.3045	0.0473	0.9380	0.1872	0.2995	0.1400	0.4005	0.0885	0.9309	0.1712	0.6305	0.0828	0.8444
Haspin	Haspin	1634779_at	-0.7392	0.1115	-0.9842	0.0328	-1.3858	0.0001	-0.2766	0.7556	-0.1177	0.8026	0.1589	0.6934	0.1822	0.8611	-0.1582	0.7525	-0.3404	0.4322
CG7579	CG7579	1634780_at	-0.0009	0.9966	0.0123	0.9215	0.1099	0.5579	0.0081	0.9931	0.0176	0.9500	0.0095	0.9679	-0.0596	0.9229	0.0024	0.9959	0.0620	0.8032
---	---	1634781_at	0.0497	0.8654	-0.0857	0.6698	0.1271	0.5444	0.0170	0.9883	-0.0169	0.9686	-0.0339	0.9195	-0.0848	0.9330	-0.1393	0.7029	-0.0545	0.9028
Cctgamma	predicted gene Y	1634782_s_at	-0.2307	0.5261	-0.5250	0.2282	-0.3633	0.1838	0.4106	0.5353	0.9961	0.0111	0.5855	0.0617	0.2520	0.8609	0.7549	0.1798	0.5029	0.3953
CG11455 /// CG3436 /// Dy	CG3436 /// CG11	1634783_s_at	-0.2175	0.4304	0.1546	0.5445	0.1913	0.3177	-0.0823	0.8939	-0.7071	0.0050	-0.6248	0.0052	0.0249	0.9885	-0.2120	0.5959	-0.2369	0.5499
yemalpha	yemanuclein-alph	1634784_at	0.1887	0.6067	-0.4176	0.6008	-0.4580	0.0129	-0.3055	0.3219	0.5022	0.0127	0.8077	0.0010	-0.1879	0.9467	-0.1097	0.9353	0.0782	0.9474
---	---	1634785_at	0.0618	0.6934	-0.0374	0.8928	0.0072	0.9734	0.0557	0.9333	0.2772	0.1561	0.2215	0.2069	-0.1689	0.8202	-0.0216	0.9655	0.1472	0.6577
lectin-28C	Lectin28C	1634786_at	0.0715	0.8745	-1.1225	0.1942	-1.1160	0.0042	-0.2068	0.7747	-0.4341	0.1492	-0.2273	0.4222	-0.1380	0.9657	-1.5169	0.0881	-1.3789	0.1385
CG32638	CG32638	1634787_at	-1.0147	0.0216	-0.2202	0.5185	-0.4094	0.2400	0.0020	0.9984	-0.6069	0.0115	-0.6089	0.0069	0.1938	0.9340	0.2294	0.8123	0.0356	0.9762
CG4872	CG4872	1634788_at	0.7370	0.0157	1.1531	0.0438	1.8277	0.0001	0.2894	0.3443	-0.0520	0.8113	-0.3414	0.0371	-0.3271	0.7726	0.4458	0.3526	0.7729	0.1432
CG30151	CG30151	1634789_at	-1.6719	0.0014	-0.0230	0.9242	-0.2167	0.5211	-0.4800	0.5355	-1.4942	0.0037	-1.0142	0.0127	-0.1393	0.8845	0.1662	0.6818	0.3055	0.4078
---	---	1634790_at	-0.0413	0.8979	-0.2687	0.0910	0.1898	0.3994	0.2628	0.5415	0.1588	0.5009	-0.1039	0.6531	-0.2725	0.7215	-0.2036	0.5411	0.0690	0.8723
---	---	1634791_at	-0.0235	0.8999	-0.6438	0.0200	-0.4796	0.0386	0.2545	0.5766	0.7172	0.0085	0.4627	0.0340	0.1295	0.8461	0.0519	0.9004	-0.0776	0.8205
CG17838	CG17838	1634792_s_at	0.7894	0.0443	1.3613	0.0576	0.2148	0.6463	-0.5063	0.3068	-0.0358	0.9299	0.4705	0.0730	0.5036	0.7822	0.4760	0.5644	-0.0276	0.9837
---	---	1634793_at	0.0564	0.8064	-0.0508	0.6286	-0.3266	0.1353	-0.3576	0.4962	0.0149	0.9729	0.3725	0.1330	0.0091	0.9939	-0.0717	0.8376	-0.0808	0.7978
CG10924	CG10924	1634794_at	0.3961	0.7832	-0.2695	0.7887	0.8271	0.1576	0.1150	0.9834	-0.1466	0.9340	-0.2616	0.8585	-1.1554	0.6960	-1.0043	0.4210	0.1511	0.9316
Prosalpha7	20s proteasome	1634795_a_at	0.3024	0.1091	0.6858	0.0171	0.6916	0.0089	0.0149	0.9863	-0.2476	0.2271	-0.2625	0.1515	0.0446	0.9555	0.3008	0.1845	0.2562	0.2869
CG17219	CG17219	1634796_at	0.9774	0.0022	0.5502	0.3716	0.6688	0.0455	-0.1203	0.8940	0.7128	0.0255	0.8331	0.0080	-0.2195	0.9093	0.1850	0.8408	0.4046	0.5792
---	---	1634797_s_at	-0.1644	0.4211	-0.0500	0.7324	0.1020	0.6631	0.1467	0.6869	-0.0209	0.9297	-0.1677	0.2715	0.0155	0.9928	0.1172	0.8256	0.1017	0.8444
Rpp30	RNaseP protein p	1634798_at	0.3157	0.2975	-0.2425	0.5836	0.2205	0.2596	0.0855	0.9116	0.3479	0.1510	0.2624	0.2285	-0.2677	0.8571	-0.1576	0.8487	0.1101	0.8949
ppk16	pickpocket 16	1634799_at	0.1016	0.5906	0.0496	0.6849	0.1014	0.6270	0.1490	0.8073	0.1196	0.6643	-0.0294	0.9215	-0.0788	0.9420	-0.0202	0.9710	0.0586	0.8984
Or43b	Olfactory receptor	1634800_at	-0.3295	0.2538	-0.2991	0.1721	-0.3571	0.1433	-0.1008	0.9154	-0.1977	0.5339	-0.0969	0.7675	-0.0802	0.9514	-0.2258	0.5847	-0.1455	0.7456
CG30184	CG30184	1634801_at	0.2546	0.1519	0.1982	0.4979	0.0507	0.8645	-0.2000	0.5311	-0.1658	0.3271	0.0342	0.8605	0.0713	0.9589	0.0264	0.9679	-0.0450	0.9350
CG40188	CG40188	1634802_at	0.0072	0.9804	-0.0052	0.9686	-0.0418	0.8366	0.0352	0.9649	-0.0432	0.8798	-0.0783	0.7326	-0.0728	0.9324	-0.1247	0.6853	-0.0519	0.8915
CG34006	CG34006	1634803_at	0.0314	0.8441	-0.0235	0.8292	0.0242	0.9275	-0.1300	0.7842	-0.1826	0.3604	-0.0527	0.8112	-0.2113	0.7215	-0.1299	0.6265	0.0814	0.7810
CG15819	CG15819	1634804_at	-1.9035	0.0012	-2.7843	0.0171	-3.4905	0.0000	0.1212	0.8800	0.2994	0.2787	0.1782	0.5000	0.5764	0.7363	-0.7351	0.3002	-1.3115	0.1080
CG14642	CG14642	1634805_at	4.8714	0.0024	3.1948	0.0027	5.5595	0.0000	1.3079	0.2896	0.7329	0.3184	-0.5751	0.3941	-0.5654	0.7092	-0.3189	0.6425	0.2465	0.7347
CG34003	CG34003	1634806_at	0.6878	0.0600	-0.0318	0.9771	0.9084	0.0035	0.0565	0.9640	0.3131	0.3545	0.2565	0.4074	-0.9132	0.6531	-0.4756	0.5826	0.4376	0.6189
---	---	1634807_at	0.0501	0.7441	-0.0137	0.8944	0.0932	0.5680	0.1164	0.7787	0.0346	0.8807	-0.0818	0.6441	-0.1000	0.9238	-0.0648	0.9036	0.0352	0.9431
lectin-21Ca	Lectin21Ca	1634808_at	0.1648	0.4023	0.0738	0.5054	-0.0276	0.8781	-0.2024	0.6327	-0.0733	0.7689	0.1291	0.5227	-0.0768	0.9309	-0.1083	0.7484	-0.0315	0.9371
CG30194 /// DereCG30194	CG30194	1634809_s_at	-1.1712	0.0513	-0.1893	0.1074	-0.8007	0.0110	0.0034	0.9988	-0.2562	0.6377	-0.2596	0.5935	0.1768	0.8191	0.1819	0.5860	0.0051	0.9924
---	---	1634810_at	0.3200	0.0859	0.0803	0.4630	0.3613	0.1057	0.0830	0.8337	0.0429	0.8183	-0.0401	0.8126	-0.0004	0.9999	0.1123	0.7906	0.1127	0.7772
CG33060	CG33060	1634811_at	0.2303	0.1257	-0.1817	0.5032	-0.2262	0.3358	0.0171	0.9860	0.2735	0.2398	0.2564	0.2177	0.0515	0.9657	-0.1826	0.5909	-0.2341	0.4814
CG7095	CG7095	1634812_at	0.3579	0.0774	0.0792	0.4581	0.1745	0.3729	0.0403	0.9649	0.0904	0.7574	0.0501	0.8605	-0.0577	0.9092	-0.1091	0.5622	-0.0513	0.8182
CG12641	CG12641	1634813_at	-3.7645	0.0006	-1.9055	0.0092	-2.6389	0.0000	-0.3523	0.6354	-1.1019	0.0092	-0.7496	0.0294	-0.1453	0.8380	-0.1083	0.7567	0.0370	0.9274
CG7083	CG7083	1634814_at	0.2671	0.4060	0.9094	0.0833	0.7656	0.1261	-0.2715	0.6441	-1.2065	0.0021	-0.9350	0.0038	-0.2753	0.8810	-0.6933	0.3097	-0.4181	0.5758
CG31104	CG31104	1634815_at	-0.6643	0.5513	-0.0862	0.6525	-0.2481	0.1636	-0.1148	0.9749	-1.0453	0.2423	-0.9305	0.2457	-0.0794	0.9814	-0.4680	0.5329	-0.3886	0.6171
CG8401	CG8401	1634816_at	0.3063	0.0761	0.0434	0.7558	0.1080	0.4787	-0.0403	0.9339	0.0525	0.7593	0.0928	0.4998	-0.0533	0.9357	-0.1356	0.5338	-0.0823	0.7323
or	orange	1634817_at	0.0570	0.7851	0.0526	0.7882	0.1730	0.2650	-0.0023	0.9978	0.2776	0.1778	0.2799	0.1292	-0.0915	0.8903	0.2957	0.2181	0.3872	0.1462
CG2165	CG2165	1634818_s_at	-1.0122	0.0180	-0.0217	0.9828	-0.1945	0.2495	-0.5730	0.1185	-0.7441	0.0056	-0.1711	0.4072	-0.3157	0.8655	0.3861	0.6312	0.7018	0.3597
Cpr76Ba	CG9283	1634819_at	0.0222	0.9425	0.0290	0.7731	-0.0676	0.8156	0.0535	0.9375	-0.0562	0.8210	-0.1096	0.5802	0.1618	0.8814	-0.0376	0.9557	-0.1995	0.6566
Pros29	20s proteasome	1634820_at	0.2389	0.1629	0.6221	0.0249	0.8233	0.0020	0.1676	0.7556	-0.3182	0.1673	-0.4857	0.0270	0.0480	0.9611	0.2293	0.4028	0.1813	0.5335
CG30468	CG30468	1634821_at	-0.0345	0.8962	-0.0560	0.6100	0.0429	0.7981	0.0429	0.9376	-0.0390	0.8513	-0.0819	0.6138	-0.0644	0.9296	-0.2505	0.2722	-0.1862	0.4448
CG10344	CG10344	1634822_a_at	-0.5231	0.0266	-0.2254	0.1130	-0.0604	0.7236	0.1180	0.7701	0.0565	0.7859	-0.0616	0.7379	-0.1896	0.8461	0.2606	0.5310	0.4502	0.2778
mRpS25	mitochondrial ribo	1634823_at	-0.3000	0.3854	0.0621	0.6518	0.1399	0.5977	0.2202	0.6013	-0.0061	0.9845	-0.2263	0.2343	0.0657	0.9814	0.4461	0.4600	0.3803	0.5488
CG1677	CG1677	1634824_a_at	0.2136	0.5132	0.2214	0.5152	0.3321	0.0299	0.1428	0.8551	0.0092	0.9823	-0.1336	0.6465	0.0840	0.9640	0.2535	0.6427	0.1696	0.7755
---	---	1634825_at	0.1921	0.3382	-0.2844	0.1635	-0.2866	0.2416	-0.0379	0.9677	0.1107	0.6962	0.1485	0.5375	-0.0241	0.9875	-0.3207	0.3424	-0.2966	0.4011
CG13337	CG13337	1634826_at	0.0735	0.6406	0.1075	0.2887	0.3098	0.1946	-0.0002	0.9999	-0.1126	0.5094	-0.1124	0.4622	-0.0899	0.9217	0.1012	0.7912	0.1911	0.5533
---	---	1634827_at	0.0704	0.7368	-0.1155	0.4517	-0.0795	0.5994	0.0432	0.9316	0.0977	0.5453	0.0545	0.7357	-0.1414	0.8049	-0.1682	0.4925	-0.0269	0.9360
pgant8	polypeptide GalN/	1634828_at	-0.3131	0.2457	-0.7010	0.1853	-0.3184	0.2045	0.1300	0.8770	0.3364	0.2447	0.2064	0.4481	-0.2386	0.8480	-0.3737	0.4802	-0.1352	0.8373
---	---	1634829_at	0.4482	0.1112	-0.0382	0.8423	0.2227	0.3822	0.0747	0.9314	0.2696	0.2995	0.1949	0.4184	-0.1206	0.8326	-0.2752	0.2289	-0.1546	0.5361
CG10164	CG10164	1634830_at	0.0116	0.9594	0.1653	0.2010	0.2122	0.2376	-0.1162	0.7281	-0.1015	0.5240	0.0147	0.9369	-0.0877	0.8903	0.1754	0.4696	0.2631	0.2855
antdh	antdh	1634831_at	0.3877	0.4346	-0.0823	0.6431	0.1561	0.5130	0.1007	0.8990	-0.0419	0.9077	-0.1425	0.5824	-0.0294	0.9928	-0.4085	0.6108	-0.3791	0.6389
---	---	1634832_at	-0.0240	0.9014	0.0350	0.7583	-0.1820	0.3625	-0.3149	0.4331	-0.2620	0.2341	0.0529	0.8338	-0.0793	0.9404	-0.0007	0.9998	0.0786	0.8519
PQBP-1	Poly-glutamine tra	1634833_at	0.3936	0.0230	0.0028	0.9836	-0.1319	0.4618	-0.1464	0.6327	0.2212	0.1387	0.3676	0.0140	0.1662	0.8013	-0.0281	0.9499	-0.1942	0.4958
CG4053	CG4053	1634834_at	-0.0725	0.9547	-0.0293	0.8762	-0.0305	0.8583	-0.0173	0.9956	-0.7310	0.2942	-0.7137	0.2498	0.1137	0.9742	-0.4721	0.6191	-0.5858	0.5299
CG15880	CG15880	1634835_at	-0.1580	0.5671	-0.0911	0.5375	0.0274	0.9046	0.1629	0.8346	-0.0895	0.8052	-0.2524	0.3517	0.1656	0.7822	0.1192	0.6789	-0.0464	0.8957
CG9317	CG9317	1634836_a_at	0.0618	0.8675	0.3356	0.3623	0.4922	0.0387	0.1236	0.9004	-0.1020	0.7951	-0.2256	0.4559	-0.0591	0.9742	0.0447	0.9519	0.1038	0.8645
CG40124	CG40124	1634837_at	-0.1272	0.4786	-0.1820	0.3848	-0.3136	0.1552	0.1922	0.6533	0.5104	0.0215	0.3182	0.0850	0.0499	0.9577	0.1820	0.5064	0.1321	0.6428
CG10638	CG10638	1634838_at	-0.6241	0.0183	-0.3322	0.0982	-0.1029	0.5213	0.1952	0.5896	0.3730	0.0476	0.1778	0.2841	-0.3488	0.7230	0.3025	0.4868	0.6514	0.1517

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG11103	CG11103	1634839_at	-0.2292	0.3580	0.0446	0.7923	0.2552	0.0889	0.0281	0.9732	-0.2194	0.2935	-0.2476	0.1799	-0.2731	0.7307	-0.0304	0.9567	0.2427	0.4913
CG32248	CG32248	1634840_at	0.1702	0.2845	0.2660	0.2047	0.6074	0.0205	0.0353	0.9622	-0.0590	0.8101	-0.0943	0.6440	-0.0410	0.9495	0.0293	0.9222	0.0703	0.7513
---	---	1634841_at	0.0406	0.8798	0.2420	0.0808	0.3255	0.1180	0.0555	0.9028	-0.0380	0.8418	-0.0935	0.5103	0.0133	0.9862	0.1281	0.4934	0.1147	0.5511
CG11160	CG11160	1634842_a_at	-3.2532	0.0005	-2.8550	0.0011	-3.6997	0.0000	-0.7341	0.0965	-1.1622	0.0016	-0.4281	0.0673	0.0718	0.9683	-0.7188	0.1331	-0.7905	0.1300
CG31737	CG31737	1634843_at	-0.0059	0.9816	0.1313	0.4979	0.3021	0.1620	-0.1271	0.8479	-0.2572	0.2934	-0.1301	0.5920	-0.0255	0.9816	0.0341	0.9310	0.0596	0.8519
CG9577	CG9577	1634844_at	0.1463	0.3945	-0.2514	0.6166	0.2020	0.3227	0.0577	0.9329	-0.1472	0.4881	-0.2049	0.2612	-0.3642	0.7485	-0.5959	0.2005	-0.2317	0.6571
Tsp42Ef	tetraspanin 42E	1634845_at	-0.9053	0.0269	-0.8570	0.0747	-0.8460	0.0033	0.0187	0.9838	-0.0694	0.7899	-0.0882	0.6937	0.1008	0.9589	0.0979	0.9079	-0.0029	0.9979
l(1)G0156	lethal (1) G0156	1634846_a_at	-0.8514	0.0466	-0.6937	0.0077	-1.0789	0.0065	-0.2674	0.5096	-0.5416	0.0199	-0.2742	0.1518	0.1733	0.9164	-0.5353	0.3358	-0.7086	0.2299
CG34382	CG14165	1634847_at	0.1631	0.4386	0.1545	0.3685	0.0356	0.9019	-0.0192	0.9840	-0.0240	0.9407	-0.0048	0.9863	0.1799	0.8235	0.0710	0.8831	-0.1089	0.7838
DsimCG16705 /// SPE	spz processing en	1634848_at	0.3257	0.1702	1.2148	0.0443	1.2132	0.0213	0.0476	0.9803	-1.5531	0.0056	-1.6008	0.0029	-0.0380	0.9885	-0.6921	0.2116	-0.6540	0.2691
CG9247	CG9247	1634849_at	0.1300	0.6926	0.4449	0.2783	-0.0382	0.8579	-0.4013	0.3576	-0.1952	0.4514	0.2062	0.3700	0.1833	0.8801	0.2127	0.6820	0.0293	0.9661
---	---	1634850_at	0.0721	0.7732	0.0576	0.5779	0.3315	0.1064	-0.0079	0.9943	-0.0732	0.7996	-0.0653	0.8047	0.0800	0.9400	0.0840	0.8501	0.0040	0.9945
---	---	1634851_at	-0.2620	0.1803	-0.1091	0.4140	-0.4868	0.0180	-0.0241	0.9759	-0.1765	0.3777	-0.1525	0.4033	-0.1493	0.8270	-0.1756	0.5523	-0.0263	0.9471
CG14851	CG14851	1634852_at	-0.1755	0.3759	0.3673	0.0415	0.3106	0.0703	-0.1471	0.7795	-0.3167	0.1446	-0.1697	0.4037	0.1453	0.7893	0.3547	0.1287	0.2094	0.3803
CG18011	CG18011	1634853_at	0.3790	0.1261	0.5739	0.1884	0.5094	0.0103	-0.3125	0.4861	-0.2300	0.3453	0.0824	0.7523	-0.2057	0.8528	0.1098	0.8634	0.3155	0.5061
CG16989	CG16989	1634854_at	-0.6825	0.1208	0.4412	0.6110	0.6695	0.0019	0.5080	0.1253	-1.0346	0.0009	-1.5426	0.0001	0.1724	0.9515	0.0874	0.9499	-0.0850	0.9433
CG9813	CG9813	1634855_s_at	-0.4173	0.0354	-0.4918	0.0674	-1.3562	0.0001	-0.6001	0.0762	-0.0481	0.8495	0.5520	0.0074	0.0523	0.9521	-0.0020	0.9985	-0.0543	0.8787
Hdc	histidine decarbox	1634856_at	-0.2802	0.1995	-0.1132	0.3556	-0.1104	0.6127	-0.0216	0.9744	0.0379	0.8580	0.0595	0.7314	-0.0016	0.9994	-0.0860	0.7825	-0.0844	0.7764
---	---	1634857_at	0.2576	0.3769	0.3408	0.1476	0.3727	0.0270	0.0001	0.9999	-0.0069	0.9790	-0.0070	0.9731	0.0442	0.9768	0.0996	0.8377	0.0553	0.9118
wkd	whacked	1634858_s_at	-0.6110	0.0287	-0.2194	0.4315	-0.6940	0.0036	-0.1683	0.7225	-0.5820	0.0143	-0.4138	0.0370	0.1538	0.8270	-0.3635	0.2021	-0.5174	0.1156
r2d2	r2d2	1634859_at	0.5837	0.1091	0.8733	0.0521	0.5247	0.0123	-0.1179	0.8968	0.1994	0.5335	0.3173	0.2374	0.2128	0.8222	0.4178	0.2722	0.2049	0.6296
CG34123	CG34123	1634860_at	0.1471	0.5404	0.4073	0.0488	0.6816	0.0032	-0.0207	0.9819	-0.3224	0.1361	-0.3017	0.1190	-0.4083	0.6557	-0.0772	0.8850	0.3312	0.3807
CG17177	CG17177	1634861_at	-0.0024	0.9904	-0.0084	0.9389	-0.1011	0.4956	-0.0364	0.9436	-0.0354	0.8522	0.0010	0.9958	0.0151	0.9868	-0.0735	0.7722	-0.0887	0.7040
CG10470	transcription unit 2	1634862_at	0.0263	0.9203	0.1202	0.6882	-0.0040	0.9852	0.0559	0.9108	0.4235	0.0153	0.3676	0.0173	0.2008	0.8202	0.4967	0.1622	0.2960	0.4303
CG33107	CG33107	1634863_at	-0.4115	0.1994	-0.4423	0.0804	-0.3289	0.0545	-0.2774	0.4643	0.1533	0.4747	0.4307	0.0269	-0.3376	0.6749	0.0947	0.8318	0.4324	0.2136
CRMP	dihydropyrimidine	1634864_at	0.0268	0.8769	-0.0386	0.7435	0.1013	0.5400	0.1752	0.5907	0.1841	0.2671	0.0089	0.9663	-0.0769	0.9092	0.0830	0.7779	0.1599	0.5239
---	---	1634865_at	0.3637	0.3125	-0.3722	0.2934	-0.0023	0.9932	0.1097	0.8472	0.3580	0.0870	0.2484	0.1817	-0.3062	0.7737	-0.4011	0.3759	-0.0950	0.8800
CG4880	CG4880	1634866_at	0.4032	0.0668	0.4227	0.1871	0.5418	0.0113	-0.0950	0.8791	0.1338	0.5602	0.2288	0.2306	-0.2649	0.7611	0.0525	0.9275	0.3174	0.3921
CG13169	CG13169	1634867_at	0.1393	0.4727	0.0800	0.7086	0.0713	0.7616	-0.0699	0.9311	0.0588	0.8489	0.1287	0.5888	-0.0304	0.9816	-0.0130	0.9815	0.0174	0.9697
CG13016	CG13016	1634868_at	0.6093	0.0322	0.5890	0.1247	0.5574	0.0082	-0.1164	0.7807	0.1317	0.4680	0.2481	0.1091	0.0113	0.9948	0.1874	0.6749	0.1760	0.6941
Thiolase	thiolase	1634869_at	0.6481	0.0351	0.5184	0.4165	0.6106	0.1350	0.0727	0.9276	-0.2180	0.3701	-0.2907	0.1714	-0.1108	0.9742	-0.4037	0.6633	-0.2929	0.7655
CG13031	CG13031	1634870_at	0.1509	0.5184	0.0271	0.8437	0.3135	0.0713	0.0540	0.9507	0.0433	0.8952	-0.0107	0.9715	-0.1569	0.7810	-0.1565	0.5393	0.0004	0.9992
---	---	1634871_s_at	-0.9670	0.0557	0.1178	0.8050	-0.5720	0.0997	-0.4757	0.4333	-0.7350	0.0344	-0.2594	0.4034	0.1386	0.9495	0.3301	0.6410	0.1915	0.8130
CG4975	CG4975	1634872_at	0.2576	0.2107	0.2572	0.5567	0.6064	0.0109	0.2207	0.5813	-0.1086	0.6339	-0.3294	0.0734	-0.1718	0.9138	-0.0668	0.9404	0.1049	0.8908
---	---	1634873_at	-0.3889	0.1256	-0.8482	0.0939	-0.4101	0.1261	-0.1379	0.8743	0.1424	0.6814	0.2803	0.3106	-0.4065	0.6955	-0.4072	0.3285	-0.0007	0.9992
AR-2	Allatostatin Recep	1634874_at	0.6088	0.2788	1.1150	0.1970	0.0857	0.8655	-0.3235	0.7208	-0.5548	0.1730	-0.2313	0.5686	0.6384	0.8069	0.0064	0.9989	-0.6320	0.5866
---	---	1634875_at	0.1593	0.3634	0.0563	0.7224	0.2024	0.1797	0.0083	0.9932	0.0465	0.8525	0.0381	0.8650	-0.0990	0.8609	0.0012	0.9991	0.1002	0.6941
CG4537	CG4537	1634876_at	0.1606	0.3089	0.2901	0.3956	0.1995	0.2235	-0.1131	0.8376	-0.2204	0.2916	-0.1073	0.6079	-0.0911	0.9053	-0.0736	0.8432	0.0175	0.9659
CG15546	CG15546	1634877_at	-1.7062	0.0106	-3.1836	0.0080	-1.6643	0.0013	0.9742	0.3836	1.3515	0.0387	0.3773	0.5411	-0.3815	0.5228	-0.2798	0.3158	0.1017	0.7580
CG15127	CG15127	1634878_at	0.4877	0.0485	0.5804	0.0157	0.4073	0.0643	-0.0242	0.9808	-0.2238	0.3705	-0.1996	0.3786	0.1691	0.8344	-0.0030	0.9977	-0.1722	0.6350
CG8519	CG8519	1634879_at	-2.6603	0.0080	-1.8445	0.0057	-3.0682	0.0000	-0.8197	0.0492	-0.9425	0.0028	-0.1228	0.6117	0.1054	0.9741	-0.5098	0.5378	-0.6152	0.4533
CG12902	CG12902	1634880_s_at	-0.1177	0.5547	0.2362	0.4248	0.0135	0.9441	-0.1770	0.7111	-0.2510	0.2483	-0.0740	0.7548	0.0341	0.9816	0.1407	0.6917	0.1066	0.7749
l(2)NC136	lethal (2) NC136	1634881_at	0.4886	0.0286	0.5713	0.2191	0.4005	0.0385	-0.0090	0.9889	0.3501	0.0269	0.3591	0.0152	0.2972	0.7628	0.3082	0.4531	0.0110	0.9866
Acp63F	Accessory gland p	1634882_at	0.6747	0.1047	0.0360	0.7322	0.5520	0.1001	-0.0185	0.9777	0.0982	0.5864	0.1167	0.4577	-0.4578	0.7644	-0.7552	0.2217	-0.2974	0.6711
CG6750	CG6750	1634883_at	-0.2228	0.3463	0.4985	0.0745	0.6374	0.0056	0.0991	0.8299	-0.2684	0.1278	-0.3676	0.0279	-0.0254	0.9853	0.3793	0.2168	0.4047	0.2189
CG31635	CG31635	1634884_s_at	-0.5725	0.0981	-0.0729	0.7617	-0.0997	0.6902	-0.2466	0.6869	-0.4188	0.1393	-0.1721	0.5353	-0.1949	0.8465	0.0245	0.9723	0.2195	0.6263
Or67b	Odorant receptor	1634885_at	0.0653	0.6767	0.2181	0.3466	-0.1948	0.2883	-0.1175	0.8771	-0.0164	0.9678	0.1011	0.7156	0.1696	0.7726	0.1617	0.5363	-0.0079	0.9848
---	---	1634886_at	0.0496	0.7923	0.0525	0.6190	0.0804	0.6661	-0.0895	0.8373	-0.0331	0.8796	0.0564	0.7523	0.0268	0.9816	0.0722	0.8471	0.0454	0.9055
Spase18-21	Spase 18/21-subu	1634887_at	0.8117	0.0041	0.7958	0.0236	1.1627	0.0002	0.2393	0.4669	0.3341	0.0635	0.0948	0.5941	-0.2139	0.7220	0.3260	0.1884	0.5399	0.0710
CG16752 /// DsimCG16752	CG16752	1634888_at	0.3208	0.2560	0.1052	0.4402	0.2060	0.3761	0.0022	0.9978	0.0022	0.9942	0.0000	0.9999	-0.0493	0.9821	-0.0034	0.9988	0.0459	0.9491
AlCR2	Drosophila allatos	1634889_at	0.3081	0.2007	0.1316	0.4843	0.3717	0.0365	-0.0202	0.9759	0.0012	0.9960	0.0214	0.9153	-0.0109	0.9943	0.0402	0.9431	0.0511	0.9161
UK114	UK114	1634890_at	2.8103	0.0011	1.2001	0.2947	2.1164	0.0001	0.6235	0.1613	0.6800	0.0200	0.0566	0.8573	-0.1972	0.9589	-0.8599	0.4287	-0.6627	0.5658
CG14200	CG14200	1634891_at	-0.2642	0.5205	0.5789	0.2024	0.7289	0.0297	-0.1357	0.9051	-0.6224	0.0923	-0.4867	0.1386	-0.2109	0.8736	0.2049	0.7361	0.4158	0.4370
CG15865	CG15865	1634892_at	0.4552	0.0354	0.1418	0.3745	-0.1352	0.4686	-0.3557	0.4174	-0.0334	0.9220	0.3223	0.1370	0.1475	0.8270	-0.0535	0.9002	-0.2010	0.4892
Gpdh	5' gene	1634893_at	-0.9512	0.0030	-1.7293	0.0110	-1.8077	0.0002	0.1886	0.6455	0.6217	0.0078	0.4331	0.0230	0.0592	0.9689	-0.2155	0.6109	-0.2747	0.5041
CG34110	CG34110	1634894_at	0.1618	0.4277	0.2026	0.1373	0.2131	0.3426	-0.1239	0.7929	-0.2036	0.2929	-0.0797	0.6921	0.0298	0.9775	0.0650	0.8500	0.0352	0.9194
Rab5	Rab-protein 5	1634895_s_at	-0.3293	0.1160	-0.5415	0.0108	-0.6649	0.0090	-0.0730	0.9234	0.1085	0.6781	0.1815	0.3941	-0.0053	0.9964	-0.1051	0.7400	-0.0998	0.7492
CG12798	CG12798	1634896_at	-0.0479	0.8510	0.0353	0.7599	0.1261	0.4881	-0.1123	0.8408	-0.0143	0.9643	0.0980	0.6485	-0.1071	0.8586	0.0080	0.9872	0.1151	0.6630
---	---	1634897_at	0.3275	0.1250	0.1082	0.4557	0.0828	0.5970	-0.1219	0.7067	0.0327	0.8675	0.1545	0.2454	-0.0085	0.9943	-0.0722	0.8481	-0.0636	0.8611
MBD-like	methyl-CpG-bindi	1634898_a_at	0.1776	0.2773	-0.2404	0.2921	-0.0746	0.7733	0.0822	0.8800	0.4053	0.0365	0.3231	0.0567	0.0229	0.9898	-0.0390	0.9507	-0.0618	0.9096
CG6512	CG6512	1634899_a_at	-0.0452	0.8406	-0.2581	0.4399	-0.0403	0.9051	0.0583	0.9518	0.3972	0.1389	0.3389	0.1585	-0.2816	0.8215	0.1362	0.8471	0.4178	0.4322
CG13244	CG13244	1634900_at	-0.1364	0.4712	0.0409	0.6648	0.3460	0.0402	-0.0832	0.8738	-0.3781	0.0447	-0.2950	0.0734	-0.4220	0.3162	-0.0391	0.9068	0.3829	0.1156
---	---	1634901_at	-0.0234	0.9125	-0.0744	0.6035	0.2511	0.1893	0.2442	0.4503	0.1293	0.4821	-0.1149	0.4957	-0.0354	0.9653	0.1192	0.6124	0.1546	0.4978
---	---	1634902_at	0.1026	0.6390	0.0778	0.7479	0.1161	0.4147	0.0987	0.8899	0.1159	0.6699	0.0171	0.9551	0.0188	0.9816	0.0921	0.6394	0.0733	0.7220
CG9981	CG9981	1634903_at	-0.0881	0.6713	0.1268	0.6405	-0.0560	0.8170	0.0073	0.9946	-0.0946	0.7224	-0.1020	0.6672	0.0648	0.9618	0.0276	0.9647	-0.0373	0.9434
RhoGEF2	Shar pei	1634904_s_at	-0.5838	0.0334	-0.4412	0.0487	-0.7119	0.0038	-0.0373	0.9672	-0.0299	0.9284	0.0074	0.9799	0.1206	0.8331	-0.0619	0.8499	-0.1825	0.4597
---	---	1634905_at	0.2558	0.1734	0.2211	0.2246	0.1462	0.5206	-0.3098	0.3765	-0.1548	0.4526	0.1550	0.4004	-0.0274	0.9816	-0.1683	0.5715	-0.1409	0.6422
alphaTub84D /// Tm1	alpha-Tubulin /// T	1634906_at	-0.5872	0.0656	-0.1496	0.6518	-1.0336	0.0027	-0.3847	0.2885	0.1196	0.6109	0.5043	0.0163	0.5434	0.6824	0.5813	0.2714	0.0379	0.9622
---	---	1634907_s_at	-0.0229	0.9060	-0.0300	0.7921	0.1928	0.3788	0.0879	0.8589	0.0400	0.8645	-0.0479	0.8112	-0.2310	0.7230	-0.0459	0.9157	0.1850	0.5314

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1634908_at	-0.0331	0.8723	0.1087	0.7166	0.1953	0.2501	0.1716	0.6535	0.0229	0.9296	-0.1486	0.3845	0.0553	0.9717	0.2336	0.5800	0.1783	0.6857
CG12975	CG12975	1634909_at	0.6496	0.0453	0.1178	0.7595	0.3909	0.1843	0.3283	0.4779	0.8788	0.0045	0.5505	0.0213	0.0410	0.9848	0.1995	0.7105	0.1585	0.7767
CG15803	CG15803	1634910_at	1.2796	0.0500	2.1970	0.0914	2.5344	0.0001	0.0783	0.8698	-0.1175	0.5064	-0.1958	0.1900	-0.0932	0.9869	0.8347	0.5401	0.9279	0.4913
---	---	1634911_at	0.2252	0.3230	0.0868	0.6228	0.2616	0.2809	0.1897	0.5419	0.1094	0.5256	-0.0803	0.6260	0.0314	0.9672	0.0099	0.9783	-0.0214	0.9394
mod	modulo	1634912_at	0.4365	0.1111	0.0332	0.9515	0.1579	0.5408	0.1655	0.7225	0.7422	0.0045	0.5766	0.0084	0.1116	0.9589	0.5038	0.4032	0.3922	0.5415
pes	peste	1634913_s_at	0.6885	0.0032	0.7509	0.0428	0.5632	0.0186	-0.1683	0.6010	0.0233	0.9183	0.1916	0.1856	0.1439	0.8736	0.0944	0.8445	-0.0494	0.9197
---	---	1634914_at	0.1649	0.3906	0.1581	0.5625	0.3166	0.1366	-0.0132	0.9937	-0.0023	0.9963	0.0110	0.9799	-0.0014	0.9994	0.0651	0.8372	0.0665	0.8221
---	---	1634915_at	0.0385	0.8962	0.1111	0.4345	0.1403	0.4973	-0.0059	0.9956	-0.1859	0.4356	-0.1801	0.4003	0.0342	0.9589	0.0001	1.0000	-0.0342	0.8957
CG8910	CG8910	1634916_at	0.0611	0.7283	-0.0322	0.8259	0.1734	0.3161	-0.0217	0.9761	-0.0085	0.9758	0.0132	0.9538	-0.2416	0.6660	-0.1250	0.6041	0.1166	0.6318
RpL23A	Ribosomal protein	1634917_at	0.0555	0.7065	-0.0364	0.7679	-0.0197	0.9058	0.0043	0.9955	0.0577	0.7400	0.0534	0.7363	0.0040	0.9964	-0.0501	0.8792	-0.0541	0.8575
CG32276	CG32276	1634918_a_at	0.6738	0.0033	0.2640	0.0615	0.4366	0.0229	0.0968	0.8217	0.5589	0.0058	0.4621	0.0083	0.0749	0.8722	0.3011	0.1085	0.2262	0.2341
CG11123 /// DmirCG11123	CG11123	1634919_at	0.5634	0.0990	-0.1314	0.7637	0.2153	0.2018	0.3086	0.5740	0.8909	0.0074	0.5823	0.0284	-0.0829	0.9589	0.1108	0.8610	0.1936	0.7030
---	---	1634920_at	-0.7977	0.0387	-0.8180	0.1162	-1.2308	0.0002	-0.1665	0.7028	0.1768	0.3898	0.3433	0.0594	0.2003	0.9076	0.0469	0.9631	-0.1535	0.8458
CG13198	CG13198	1634921_at	0.0290	0.9143	0.0860	0.4718	-0.0259	0.9156	-0.0923	0.8676	-0.1433	0.4852	-0.0511	0.8183	0.1819	0.8167	0.0263	0.9613	-0.1556	0.6574
CG32795	CG32795	1634922_s_at	0.5825	0.0803	1.0803	0.0239	1.1194	0.0010	-0.3447	0.4504	-1.3636	0.0007	-1.0189	0.0014	-0.3895	0.7220	-0.6852	0.1394	-0.2957	0.5498
CG12996	CG12996	1634923_at	0.1477	0.3378	0.3431	0.3051	0.3743	0.0583	0.0511	0.9472	-0.1190	0.6236	-0.1701	0.4026	0.0997	0.8655	0.0491	0.8850	-0.0506	0.8729
Spt6	Spt6	1634924_at	0.2480	0.5444	0.0094	0.9901	0.3725	0.0635	0.1315	0.7743	0.5119	0.0158	0.3804	0.0339	-0.0907	0.9796	0.3944	0.6519	0.4852	0.5695
Tango2	Transport and Gol	1634925_at	0.2976	0.3251	0.1125	0.4968	0.5072	0.0561	-0.0917	0.9273	-0.2276	0.4675	-0.1359	0.6620	-0.5262	0.3400	-0.4695	0.1160	0.0566	0.8888
---	---	1634926_at	0.0932	0.6853	0.0670	0.6702	-0.1045	0.7048	-0.0070	0.9943	0.2530	0.2101	0.2600	0.1500	0.1478	0.8940	0.1084	0.8481	-0.0394	0.9464
CG30177	CG30177	1634927_at	0.1052	0.4588	0.1047	0.4598	0.2008	0.3177	-0.0073	0.9921	-0.1420	0.3469	-0.1348	0.3189	-0.0586	0.9291	-0.0996	0.6727	-0.0410	0.8886
ImpE2	Ecdysone-inducib	1634928_at	0.1332	0.4152	-0.0116	0.9189	0.1724	0.2999	0.0374	0.9573	-0.0603	0.7923	-0.0977	0.6034	0.0146	0.9872	-0.1834	0.3669	-0.1980	0.3529
---	---	1634929_at	0.1882	0.2712	0.0796	0.4305	0.1112	0.4445	-0.2888	0.4762	-0.2836	0.1902	0.0052	0.9850	-0.0833	0.8705	-0.0565	0.8353	0.0267	0.9237
---	---	1634930_at	0.5057	0.0515	0.4081	0.2748	0.4500	0.0428	0.1457	0.7983	0.0339	0.9167	-0.1117	0.6372	-0.0026	0.9984	-0.0278	0.9476	-0.0252	0.9444
CG6340	CG6340	1634931_at	0.5682	0.0857	-0.3075	0.0444	-0.8809	0.1444	-0.2935	0.4998	1.3251	0.0006	1.6186	0.0002	0.2326	0.9457	0.1438	0.9303	-0.0888	0.9514
---	---	1634932_at	0.2534	0.3089	-0.0032	0.9871	-0.1102	0.5643	-0.0920	0.9074	0.0293	0.9351	0.1213	0.6428	0.0886	0.9221	-0.1273	0.7088	-0.2159	0.4876
CG17207	CG17207	1634933_s_at	0.1308	0.6443	0.3593	0.0568	0.3513	0.1366	-0.2367	0.6325	-0.1708	0.5045	0.0659	0.8087	0.0426	0.9588	0.1141	0.6458	0.0715	0.7956
CG8765	CG8765	1634934_at	-0.1442	0.6856	-0.1741	0.2734	-0.2903	0.1506	0.1277	0.7950	0.4123	0.0454	0.2846	0.1111	0.2568	0.7677	0.2661	0.4791	0.0093	0.9874
CG1648	CG1648	1634935_a_at	1.1927	0.0049	0.4584	0.6651	1.0228	0.0108	0.3837	0.6144	0.0860	0.8645	-0.2977	0.4022	-0.1598	0.9677	-0.6914	0.5251	-0.5316	0.6389
CG14972	CG14972	1634936_at	0.0536	0.8083	0.0699	0.7092	0.1019	0.5699	0.0898	0.8813	-0.0998	0.6683	-0.1895	0.3082	-0.0554	0.9398	-0.1129	0.6425	-0.0576	0.8409
CG17824	CG17824	1634937_at	-0.0043	0.9871	0.2151	0.2708	0.2374	0.2643	0.0005	0.9994	-0.1957	0.3742	-0.1962	0.3180	0.2142	0.7689	0.3245	0.2794	0.1103	0.7577
---	---	1634938_s_at	0.1378	0.3331	0.2249	0.1937	0.4116	0.0168	0.0001	0.9999	-0.0353	0.8745	-0.0354	0.8569	0.0173	0.9848	0.0058	0.9903	-0.0115	0.9716
CG13077	CG13077	1634939_at	1.7137	0.0024	2.2685	0.0126	2.0367	0.0022	-0.2636	0.8215	-0.4065	0.3830	-0.1429	0.7775	-0.1522	0.9246	0.0106	0.9935	0.1628	0.8046
kraken	kraken	1634940_at	0.1005	0.6808	-0.2350	0.7239	-0.1844	0.5467	0.4576	0.0976	0.6170	0.0033	0.1594	0.2774	0.4728	0.7872	0.2274	0.8150	-0.2454	0.7815
---	---	1634941_at	0.0781	0.6470	0.1610	0.3859	0.2891	0.1621	-0.0347	0.9677	0.0046	0.9896	0.0392	0.8834	0.0077	0.9939	0.0632	0.8380	0.0555	0.8524
CG5114	CG5114	1634942_at	0.8391	0.0207	0.0349	0.9517	0.3014	0.1855	0.1401	0.8605	1.0297	0.0041	0.8896	0.0047	0.0297	0.9913	0.2915	0.6209	0.2618	0.6602
RfC40	Replication-factor-	1634943_at	-0.0316	0.9066	-0.0894	0.4761	-0.2429	0.3868	-0.0248	0.9777	0.4848	0.0337	0.5097	0.0172	0.0659	0.9710	0.3190	0.4952	0.2532	0.6051
l(2)tid	lethal (2) tumorou	1634944_at	-0.8527	0.1222	-0.1784	0.8878	-0.1098	0.7382	0.0863	0.8747	-0.6126	0.0069	-0.6989	0.0023	0.0359	0.9946	0.1952	0.9204	0.1593	0.9271
CG7830	CG7830	1634945_at	0.7779	0.0044	0.6806	0.0483	0.9655	0.0026	0.0187	0.9814	0.2388	0.2117	0.2201	0.1987	-0.1153	0.8744	0.2734	0.3284	0.3887	0.1993
---	---	1634946_at	0.2192	0.4952	-0.3344	0.4033	-0.5774	0.1249	-0.1678	0.9247	0.8822	0.0971	1.0500	0.0342	0.1569	0.9092	0.2157	0.6945	0.0587	0.9318
---	---	1634947_s_at	0.0549	0.7906	0.1314	0.3752	0.1896	0.3028	-0.0689	0.8640	0.0262	0.8926	0.0951	0.4866	0.0546	0.9521	0.1610	0.5778	0.1064	0.7347
luna	luna	1634948_at	0.1881	0.7285	0.8282	0.2111	-0.3302	0.1442	-0.4700	0.1401	0.0571	0.8104	0.5270	0.0085	0.5731	0.7387	0.8646	0.2198	0.2915	0.7259
hep	hemipterous	1634949_at	-0.9246	0.2613	-0.5168	0.4129	-0.8086	0.0041	-0.0167	0.9894	0.0543	0.8904	0.0710	0.8297	0.1909	0.9589	0.3675	0.7755	0.1766	0.9031
CG12785	CG12785	1634950_at	0.0485	0.9405	-0.1068	0.8948	0.3143	0.3014	0.4153	0.7293	1.0434	0.0562	0.6281	0.1894	-0.0105	0.9976	0.7159	0.3571	0.7264	0.3723
---	---	1634951_at	0.2904	0.1867	0.0192	0.9010	0.2763	0.2497	-0.0957	0.8578	-0.0088	0.9767	0.0869	0.6636	-0.0575	0.9621	-0.0834	0.8584	-0.0258	0.9578
CG15094	CG15094	1634952_s_at	-0.1815	0.4846	0.4594	0.1415	0.8520	0.0004	0.1675	0.7891	-0.0590	0.8610	-0.2265	0.3340	-0.2905	0.7464	0.5668	0.1337	0.8573	0.0607
CG3795	CG3795	1634953_at	0.0307	0.8601	0.1484	0.3652	0.4293	0.0436	0.2474	0.3541	0.1313	0.4026	-0.1160	0.4166	0.0357	0.9788	0.1866	0.5711	0.1509	0.6564
CG4756	CG4756	1634954_at	-0.1529	0.6908	-0.5352	0.2753	-0.9621	0.0013	-0.2726	0.5832	0.1083	0.7114	0.3809	0.0921	-0.0906	0.9515	-0.2781	0.5542	-0.1875	0.7128
mtacp1	NADH-ubiquinone	1634955_at	-0.0165	0.9632	0.4363	0.1817	0.4845	0.0345	-0.1330	0.8084	-1.0508	0.0012	-0.9177	0.0013	-0.0895	0.9555	-0.4703	0.3009	-0.3808	0.4266
CG15022	CG15022	1634956_at	0.1203	0.6115	-0.0294	0.8397	-0.1616	0.5631	0.0134	0.9902	0.2668	0.2734	0.2534	0.2445	0.0279	0.9824	-0.0734	0.8551	-0.1013	0.7672
Dh	CRF-like peptide	1634957_at	0.1071	0.7050	-0.7325	0.0238	-0.1261	0.8149	0.9992	0.1595	1.4317	0.0059	0.4326	0.2641	0.1748	0.9056	0.3739	0.4903	0.1992	0.7442
CG9455	CG9455	1634958_at	2.6568	0.0008	2.7055	0.0039	3.0412	0.0000	0.2310	0.5628	0.2579	0.2062	0.0269	0.9142	-0.1886	0.9112	0.2212	0.7579	0.4098	0.5137
TfIIA-S-2	TfIIA-S-2	1634959_at	-0.0025	0.9898	-0.0424	0.8252	0.0934	0.6569	0.1428	0.7887	0.1475	0.5262	0.0047	0.9863	-0.0098	0.9939	0.0522	0.9047	0.0620	0.8699
CG14375	CG14375	1634960_at	2.0593	0.0307	1.4273	0.0522	1.7764	0.0009	0.8147	0.5680	-0.2560	0.7711	-1.0707	0.1010	0.3494	0.8445	-0.7749	0.2722	-1.1243	0.1498
Vha68-2	Vha68-2	1634961_s_at	-0.8170	0.0092	-0.9901	0.0245	-1.1750	0.0001	-0.1493	0.6327	-0.2112	0.1659	-0.0619	0.7017	0.0064	0.9964	-0.2398	0.5020	-0.2462	0.4930
hts	hu-li tai shao	1634962_s_at	-0.7502	0.0123	-0.4798	0.1584	-1.4557	0.0018	-0.2236	0.4529	-0.1404	0.3973	0.0832	0.6108	0.6886	0.6702	0.1397	0.8821	-0.5489	0.4090
lgs	legless	1634963_at	-0.0842	0.7145	0.0714	0.7375	-0.4001	0.0268	-0.2526	0.3381	0.0653	0.7105	0.3179	0.0273	0.2628	0.7215	0.2388	0.4437	-0.0240	0.9575
CG7794	CG7794	1634964_at	0.1013	0.6839	0.1696	0.4886	0.1676	0.3878	-0.0465	0.9518	0.0089	0.9789	0.0554	0.8191	0.0789	0.9558	0.1266	0.8135	0.0477	0.9344
---	---	1634965_at	0.0884	0.6401	0.0008	0.9962	0.1133	0.4416	-0.0108	0.9893	0.0726	0.7364	0.0833	0.6604	0.0139	0.9869	-0.0442	0.8708	-0.0581	0.8039
Aly	Aly	1634966_a_at	0.0354	0.9578	1.1686	0.0136	0.8762	0.0042	-0.0829	0.9311	-0.6841	0.0230	-0.6012	0.0247	0.1938	0.9238	0.5126	0.4475	0.3188	0.6627
---	---	1634967_at	0.2093	0.2593	0.0921	0.6192	-0.0744	0.6441	0.0135	0.9860	-0.0205	0.9342	-0.0340	0.8700	-0.0527	0.9495	-0.1970	0.4249	-0.1443	0.5846
---	---	1634968_at	0.2924	0.4060	-0.0305	0.8453	-0.2558	0.5705	-0.0606	0.9672	0.1407	0.7613	0.2013	0.6058	0.0798	0.9739	-0.0375	0.9698	-0.1173	0.8870
Pkd2	Polycystic kidney	1634969_at	0.0517	0.7276	-0.0601	0.6286	-0.0255	0.9118	-0.1084	0.8241	-0.0299	0.9091	0.0784	0.6925	-0.1326	0.8122	-0.1873	0.4164	-0.0547	0.8559
CG3634	CG3634	1634970_at	0.0190	0.9535	0.0019	0.9912	0.1473	0.5354	0.1272	0.8841	0.1095	0.7583	-0.0177	0.9621	-0.0143	0.9914	0.1989	0.5299	0.2132	0.5003
CG6465	CG6465	1634971_at	-0.0113	0.9838	-1.2841	0.0213	-0.4990	0.1904	0.0741	0.9726	1.1386	0.0396	1.0645	0.0332	-0.7567	0.5933	-0.1307	0.8850	0.6260	0.3276
CG4802	CG4802	1634972_at	1.0391	0.0035	0.6980	0.1074	1.3272	0.0004	0.2803	0.2989	0.1064	0.5292	-0.1739	0.2207	-0.2437	0.8192	-0.1952	0.6883	0.0485	0.9367
nuf	nuclear-fallout	1634973_s_at	0.2429	0.4007	0.1152	0.4435	-0.1075	0.5826	-0.2331	0.5174	-0.0596	0.7982	0.1735	0.3157	0.1724	0.8292	-0.0086	0.9911	-0.1810	0.6166
---	---	1634974_at	-0.0344	0.8576	0.1978	0.2552	0.1642	0.3984	0.1189	0.7507	-0.0347	0.8723	-0.1535	0.2975	0.0631	0.9495	0.1398	0.6694	0.0767	0.8387
SF2	SF2	1634975_at	-0.0039	0.9892	0.2757	0.0488	0.2973	0.0581	-0.0101	0.9922	0.2709	0.2147	0.2810	0.1499	0.0376	0.9793	0.6859	0.0609	0.6483	0.0859
Pif2	PFTAIRE-interacti	1634976_at	-0.1379	0.4021	0.1176	0.3510	-0.0826	0.7327	-0.2451	0.4141	-0.3293	0.0529	-0.0842	0.6184	-0.0635	0.9409	-0.0086	0.9873	0.0549	0.8788

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Tsp42Er	tetraspanin 42E	1634977_at	1.3845	0.1234	0.2716	0.3769	1.2620	0.0070	0.3819	0.7230	-0.1808	0.7553	-0.5627	0.1947	-0.5327	0.8446	-1.1530	0.2874	-0.6203	0.6052
CG13117	CG13117	1634978_at	-0.7859	0.0101	-0.5446	0.1020	-0.2601	0.3277	0.1655	0.7468	-0.4157	0.0665	-0.5812	0.0109	0.0982	0.9643	0.1518	0.8509	0.0536	0.9486
CG18600	CG18600	1634979_at	0.3879	0.3211	-0.0858	0.9122	0.1704	0.4454	0.0949	0.9319	0.5624	0.0894	0.4675	0.1122	-0.0893	0.9774	0.2175	0.8218	0.3068	0.7134
CG18490	CG18490	1634980_s_at	-0.1067	0.9104	0.0316	0.9855	-0.3726	0.2977	0.0002	0.9999	0.4277	0.0982	0.4275	0.0674	0.4511	0.9296	0.6505	0.7354	0.1993	0.9316
CG9010 /// DguaCG9010 //	CG9010	1634981_at	0.2515	0.1469	0.3051	0.1584	0.4209	0.1324	-0.1017	0.8792	-0.4124	0.0733	-0.3108	0.1259	0.0280	0.9852	-0.1116	0.8000	-0.1396	0.7228
CG7523	CG7523	1634982_at	0.1758	0.2441	-0.1128	0.3845	-0.2738	0.1308	0.2424	0.3328	0.2096	0.1453	-0.0328	0.8443	0.4556	0.3338	-0.1280	0.6325	-0.5836	0.0559
CG32521	CG32521	1634983_s_at	-0.1855	0.6876	-0.1888	0.8888	0.2815	0.0667	-0.3434	0.2661	-0.4295	0.0265	-0.0862	0.6469	-0.7805	0.7953	-0.4402	0.7755	0.3403	0.8303
---	---	1634984_at	-0.0118	0.9642	-0.2163	0.3050	0.0045	0.9873	-0.0091	0.9946	0.0008	0.9985	0.0099	0.9781	-0.1129	0.8270	-0.1183	0.6102	-0.0054	0.9874
pHCl	pHCl	1634985_at	-0.3423	0.3287	-0.1253	0.4266	-0.1743	0.3199	-0.1125	0.9068	-0.1611	0.6427	-0.0487	0.8959	0.0689	0.9277	-0.0008	0.9996	-0.0696	0.8235
CG32821	CG32821	1634986_at	-0.1769	0.4051	0.2924	0.3776	-0.3540	0.1006	-0.2582	0.6166	-0.1772	0.5167	0.0811	0.7767	0.4695	0.5228	0.4178	0.2181	-0.0517	0.9158
CG34119	CG34119	1634987_at	0.1555	0.3200	0.1715	0.3181	0.3217	0.0667	0.0817	0.8959	-0.0652	0.7967	-0.1469	0.4510	0.0315	0.9672	-0.0150	0.9643	-0.0465	0.8606
CG17352	CG17352	1634988_a_at	0.2507	0.3515	0.1713	0.5305	0.4804	0.2208	0.0307	0.9768	0.0437	0.9026	0.0130	0.9683	-0.1307	0.9511	0.0333	0.9760	0.1640	0.8372
Acon	m-Aconitase	1634989_at	-0.6928	0.0073	0.1855	0.3755	-0.1631	0.4539	-0.2513	0.5972	-0.9327	0.0030	-0.6814	0.0073	0.0745	0.9109	-0.0941	0.7332	-0.1686	0.4912
---	---	1634990_s_at	0.1208	0.6170	0.0543	0.6067	0.0976	0.6610	-0.0568	0.9465	-0.0795	0.7841	-0.0227	0.9393	-0.0375	0.9742	0.0251	0.9573	0.0626	0.8686
CG8083	CG8083	1634991_at	-0.8047	0.0139	-1.6112	0.0418	-1.5934	0.0004	0.2850	0.6247	1.4153	0.0011	1.1303	0.0017	0.1733	0.9092	0.4997	0.3397	0.3264	0.5669
CG9821 /// DsimCG9821	CG9821	1634992_s_at	-0.2372	0.5017	-0.7395	0.0818	-0.7129	0.0495	0.2256	0.6585	0.4050	0.0989	0.1794	0.4365	0.2243	0.9056	0.0508	0.9641	-0.1736	0.8414
CG40498	CG40498	1634993_at	-0.0815	0.7416	0.0000	1.0000	-0.0752	0.7772	-0.0449	0.9582	-0.0878	0.7422	-0.0429	0.8724	0.0401	0.9776	0.0730	0.8814	0.0330	0.9444
RacGAP84C	RnRacGAP	1634994_a_at	-0.0829	0.6987	0.3063	0.3662	0.0598	0.7473	-0.0328	0.9673	-0.1145	0.6243	-0.0817	0.7142	0.1195	0.9092	0.2609	0.4943	0.1414	0.7419
CG3788	CG3788	1634995_at	-0.1030	0.6633	-0.1519	0.5744	0.0174	0.9629	-0.0348	0.9610	-0.0729	0.7478	-0.0381	0.8639	-0.2195	0.8846	-0.1187	0.8858	0.1008	0.8984
---	---	1634996_at	0.2820	0.0749	0.0218	0.8343	0.5194	0.0092	0.1056	0.7950	0.2568	0.1197	0.1512	0.3166	-0.2313	0.6749	-0.0180	0.9620	0.2134	0.3601
Sgs5	Salivary gland sec	1634997_at	0.0864	0.6676	0.1213	0.3741	0.3003	0.1918	0.2112	0.5644	0.2195	0.2443	0.0084	0.9715	0.0043	0.9978	0.1846	0.6270	0.1803	0.6340
CG9269	CG9269	1634998_at	0.0293	0.8601	-0.4215	0.1036	-0.2263	0.3314	0.1599	0.7538	0.4457	0.0483	0.2858	0.1454	-0.1922	0.8049	-0.1741	0.6167	0.0181	0.9712
---	---	1634999_at	0.1339	0.4122	0.0982	0.6100	-0.0630	0.8053	-0.1307	0.7117	0.1613	0.3206	0.2921	0.0473	0.0498	0.9717	-0.0011	0.9996	-0.0510	0.9176
Glycogenin	Glycogenin	1635000_at	-1.9449	0.0021	-1.6203	0.0938	-2.2869	0.0001	-0.5934	0.0884	-0.9868	0.0012	-0.3934	0.0371	0.1772	0.9636	-0.5999	0.5934	-0.7771	0.4779
CG9727	CG9727	1635001_at	0.0828	0.5723	0.3685	0.3266	0.6954	0.0084	0.1796	0.6425	-0.1094	0.5940	-0.2890	0.0867	-0.0334	0.9816	0.1577	0.6593	0.1911	0.5814
shd	shade	1635002_at	-0.0434	0.8906	0.0089	0.9343	0.1191	0.5170	-0.1845	0.5873	-0.3287	0.0615	-0.1442	0.3667	-0.0072	0.9939	0.0658	0.8191	0.0730	0.7799
CG6697	CG6697	1635003_at	0.7640	0.0196	0.4834	0.1494	0.4045	0.0638	0.0934	0.7887	0.2533	0.0781	0.1599	0.2107	0.2642	0.7823	0.1455	0.7738	-0.1187	0.8178
---	---	1635004_at	0.1181	0.4315	-0.0864	0.4420	0.1312	0.5677	0.0851	0.8500	0.0135	0.9557	-0.0716	0.6765	0.0142	0.9916	-0.2519	0.4258	-0.2662	0.4093
Rdl	GABA receptor	1635005_a_at	0.2527	0.4578	-0.1660	0.4268	-0.2239	0.3770	-0.0624	0.9298	0.1660	0.4403	0.2285	0.2185	-0.0341	0.9816	-0.2494	0.4796	-0.2153	0.5578
CG7077	CG7077	1635006_at	-0.8696	0.0288	-0.3691	0.0456	-0.5754	0.0479	0.0164	0.9877	-0.1507	0.5825	-0.1672	0.4892	-0.2042	0.7848	-0.1667	0.6312	0.0375	0.9342
Sulf1	Sulfated	1635007_at	-1.3469	0.0034	-1.8178	0.0842	-2.2029	0.0003	-0.3209	0.5070	0.3919	0.1253	0.7128	0.0080	0.1588	0.9614	0.0358	0.9841	-0.1230	0.9231
Cyp9b2	fly plexin a	1635008_at	1.6449	0.0016	2.4203	0.0014	2.8487	0.0006	-0.1143	0.9496	-1.1991	0.0253	-1.0848	0.0241	-0.4106	0.7070	-0.4440	0.3158	-0.0334	0.9597
CG8500	CG8500	1635009_at	-0.2308	0.6034	-0.3031	0.4209	-0.4981	0.0510	-0.2158	0.8636	0.3112	0.5092	0.5270	0.1853	-0.1746	0.8981	-0.1520	0.8174	0.0225	0.9777
CG40057	CG40057	1635010_at	0.1146	0.5429	-0.0053	0.9721	-0.0591	0.7853	-0.0481	0.9488	-0.0006	0.9985	0.0475	0.8437	-0.0649	0.9357	-0.0622	0.8612	0.0027	0.9953
Tom20	Translocase of ou	1635011_at	0.0711	0.7766	0.4513	0.2084	0.4047	0.0748	0.2119	0.7023	-0.2187	0.4030	-0.4305	0.0621	0.1659	0.8825	0.2389	0.6011	0.0730	0.9023
---	---	1635012_x_at	0.1140	0.5285	-0.0060	0.9639	0.2124	0.2344	-0.0492	0.9322	-0.0565	0.7880	-0.0072	0.9730	-0.1167	0.8156	-0.1504	0.4729	-0.0337	0.9057
CG10722	CG10722	1635013_s_at	-0.5534	0.2682	-0.3210	0.3996	0.2363	0.5976	-0.1741	0.8979	-0.2385	0.6310	-0.0644	0.9045	-0.6145	0.7822	-0.0319	0.9870	0.5826	0.5670
---	---	1635014_at	0.0537	0.7692	0.0068	0.9519	-0.0561	0.7381	-0.1221	0.7205	0.0034	0.9885	0.1255	0.3798	-0.0638	0.9246	-0.0777	0.7815	-0.0139	0.9675
CG15157	CG15157	1635015_at	0.1310	0.5199	0.0433	0.7080	0.2581	0.1780	-0.0035	0.9956	-0.1722	0.4217	-0.1687	0.3793	-0.1834	0.7644	-0.0213	0.9605	0.1620	0.5556
CG13090	CG13090	1635016_at	0.2159	0.3223	0.1953	0.2055	0.3578	0.0345	-0.1632	0.7167	-0.1659	0.4347	-0.0027	0.9916	-0.3365	0.5954	-0.2293	0.4067	0.1072	0.7391
---	---	1635017_at	-0.2228	0.1661	-0.5872	0.1385	-0.4610	0.0253	0.0169	0.9826	0.3272	0.0742	0.3103	0.0599	-0.1709	0.7464	0.0503	0.8751	0.2212	0.3336
Sin1	CG10105	1635018_at	-0.3015	0.3023	0.2761	0.1968	0.5126	0.0714	0.0211	0.9819	-0.1684	0.4647	-0.1894	0.3497	-0.3194	0.7770	0.2790	0.5905	0.5984	0.2363
CG12971	CG12971	1635019_at	0.1731	0.5300	0.4540	0.0726	0.3811	0.0655	0.1338	0.7451	0.0619	0.7768	-0.0719	0.7065	0.1993	0.8305	0.3736	0.3259	0.1743	0.6861
phr6-4	photolyase	1635020_s_at	-0.0756	0.8212	-0.2989	0.3859	-0.3213	0.1919	-0.2229	0.5445	0.5397	0.0121	0.7626	0.0016	-0.4193	0.7990	0.0796	0.9425	0.4989	0.4824
disp	dispatched	1635021_at	-0.8394	0.0635	-0.4086	0.1501	-0.8770	0.0007	-0.3404	0.2753	0.0601	0.7897	0.4005	0.0232	0.0026	0.9994	0.3153	0.4297	0.3127	0.4475
---	---	1635022_at	0.3266	0.1480	0.1274	0.4966	0.0094	0.9691	-0.1861	0.7205	0.0043	0.9899	0.1904	0.3819	0.0644	0.9400	0.0846	0.7988	0.0202	0.9575
CG3560	CG3560	1635023_at	-0.2171	0.3126	-0.0790	0.8293	-0.2816	0.1923	-0.1806	0.7380	-0.4557	0.0604	-0.2751	0.1969	-0.0200	0.9914	-0.3613	0.3760	-0.3414	0.4218
---	---	1635024_at	-0.0262	0.9072	0.0736	0.5882	0.1091	0.5692	0.0796	0.9101	-0.1401	0.5613	-0.2197	0.2773	0.0189	0.9824	0.0075	0.9842	-0.0114	0.9701
l(1)G0255	lethal (1) G0255	1635025_a_at	-0.8420	0.0041	-0.2254	0.1896	-0.1089	0.6018	-0.1608	0.7409	-1.0304	0.0012	-0.8696	0.0015	-0.2006	0.7565	-0.2643	0.3216	-0.0637	0.8581
tap	biparous	1635026_at	-0.5211	0.6470	-0.0387	0.7216	0.0719	0.7408	0.1560	0.9683	-0.7207	0.5059	-0.8767	0.3517	-0.0540	0.9689	-0.3320	0.3448	-0.2780	0.4564
CG31800	CG31800	1635027_at	-0.0952	0.6635	-0.0052	0.9722	0.1095	0.4951	0.0032	0.9956	-0.1691	0.2534	-0.1723	0.1911	-0.0428	0.9724	0.0143	0.9799	0.0571	0.8936
---	---	1635028_s_at	-2.3682	0.0065	-2.4563	0.0270	-3.4575	0.0001	-0.5191	0.3829	-0.7215	0.0382	-0.2024	0.5376	0.2123	0.9653	-1.2148	0.3397	-1.4270	0.2884
---	---	1635029_at	0.1558	0.4193	0.0102	0.9364	0.0104	0.9718	0.0758	0.8817	0.2158	0.2140	0.1400	0.3862	0.0350	0.9823	0.0420	0.9434	0.0070	0.9914
CG12868	CG12868	1635030_at	-0.2984	0.2571	0.0443	0.8759	0.0162	0.9712	0.3199	0.4554	0.4833	0.0436	0.1635	0.4573	0.3959	0.7215	0.9232	0.0665	0.5273	0.2757
CG11147	CG11147	1635031_s_at	-2.9809	0.0116	-3.8952	0.0074	-4.0379	0.0002	0.0432	0.9599	-0.3790	0.0935	-0.4222	0.0426	0.3541	0.9405	-1.1859	0.4126	-1.5400	0.3027
CG3526	CG3526	1635032_a_at	0.1587	0.3231	-0.1036	0.3170	0.2488	0.3742	0.0565	0.9451	0.0999	0.7122	0.0434	0.8760	0.0185	0.9870	0.0322	0.9395	0.0137	0.9730
CG30083	CG30083	1635033_at	1.9266	0.0041	0.0171	0.9828	1.2698	0.0180	0.3662	0.7595	0.4137	0.4369	0.0475	0.9404	-0.9402	0.6483	-1.4505	0.0908	-0.5103	0.5624
---	---	1635034_at	-0.0121	0.9610	-0.1112	0.3496	-0.2443	0.1087	-0.0909	0.8810	0.0689	0.7841	0.1598	0.4082	-0.0641	0.8940	-0.0183	0.9499	0.0457	0.8426
CG9384	CG9384	1635035_at	-1.2213	0.0043	-1.6373	0.0082	-1.9118	0.0000	-0.0636	0.9346	0.1414	0.5675	0.2050	0.3268	0.0223	0.9869	-0.2569	0.4016	-0.2792	0.3800
---	---	1635036_at	0.0198	0.9080	-0.0563	0.6069	-0.2941	0.0585	-0.2250	0.5233	0.0110	0.9696	0.2359	0.1518	0.0688	0.8981	-0.0114	0.9734	-0.0802	0.7240
CG4022	CG4022	1635037_at	0.1546	0.6392	-0.9325	0.0084	-1.5729	0.0020	0.0807	0.8902	1.4981	0.0002	1.4174	0.0002	0.6817	0.5228	0.3189	0.5479	-0.3629	0.4892
---	---	1635038_s_at	-0.0682	0.6579	-0.0658	0.6083	0.1347	0.3555	-0.0017	0.9978	0.0058	0.9801	0.0075	0.9689	-0.1502	0.7992	0.0296	0.9404	0.1798	0.4805
MED16	Mediator complex	1635039_at	-0.7865	0.0107	-0.0693	0.6402	0.2124	0.4426	-0.1653	0.8180	-0.6042	0.0382	-0.4389	0.0823	-0.5118	0.3800	-0.0414	0.9353	0.4703	0.1660
didum	myosin V	1635040_a_at	0.5958	0.0213	0.4135	0.2141	0.0241	0.9040	-0.0609	0.9314	0.5701	0.0144	0.6311	0.0055	0.2133	0.8222	0.4633	0.2226	0.2501	0.5488
CG32225	CG32225	1635041_at	-0.2794	0.1798	0.0832	0.6926	-0.0874	0.5975	0.1581	0.8164	-0.2298	0.4034	-0.3880	0.1044	0.0244	0.9816	0.0090	0.9848	-0.0154	0.9666
ia2	ia2	1635042_s_at	-0.0318	0.8497	0.0255	0.8796	-0.0732	0.8583	-0.1585	0.8626	-0.0913	0.8229	0.0672	0.8599	0.0758	0.9238	0.1063	0.7257	0.0305	0.9342
CG14964 /// DmirCG14964	CG14964	1635043_at	-1.6140	0.0022	-0.1790	0.3459	-0.5997	0.1016	-0.0411	0.9627	-0.7925	0.0055	-0.7514	0.0041	-0.0267	0.9848	0.0100	0.9875	0.0367	0.9331
Hsp26	heat shock proteir	1635044_at	0.1971	0.9262	-0.8619	0.0447	-1.3108	0.2689	0.7872	0.8551	3.3275	0.0393	2.5403	0.0710	1.3006	0.7953	2.1103	0.3013	0.8097	0.7384
CG6271	CG6271	1635045_at	0.0193	0.9155	0.2596	0.2832	0.2915	0.0965	0.0314	0.9731	-0.0597	0.8411	-0.0911	0.7091	-0.0434	0.9717	0.1978	0.5368	0.2412	0.4457

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18869 /// DsimCG18869	CG18869	1635046_at	0.4103	0.2423	0.0079	0.9428	0.0044	0.9858	-0.1048	0.8979	0.1711	0.5533	0.2759	0.2517	-0.0055	0.9976	-0.2434	0.5544	-0.2379	0.5695
trio	trio	1635047_s_at	-0.1409	0.6803	-0.4773	0.2461	-0.3539	0.2051	0.1017	0.9486	0.4012	0.3726	0.2995	0.4752	-0.1920	0.8284	0.0052	0.9949	0.1972	0.6225
---	---	1635048_at	0.1779	0.3098	0.0643	0.5765	0.1329	0.5591	-0.0632	0.9419	0.0229	0.9509	0.0861	0.7481	0.0448	0.9635	0.0567	0.8824	0.0119	0.9777
CG13877	CG13877	1635049_at	-0.6698	0.0888	-1.2284	0.1088	-0.6874	0.0397	0.2357	0.6557	0.1436	0.6069	-0.0920	0.7363	-0.3035	0.8806	-0.4264	0.6052	-0.1228	0.9086
---	---	1635050_at	0.0838	0.6842	0.3132	0.0412	0.1503	0.4424	-0.0683	0.9116	-0.1601	0.4275	-0.0918	0.6476	0.1825	0.7485	0.0806	0.7903	-0.1018	0.7085
Herp	CG14536	1635051_a_at	0.2592	0.3257	0.5518	0.0804	0.7212	0.0021	-0.0028	0.9961	0.2129	0.2483	0.2157	0.1890	-0.0860	0.9481	0.5004	0.1948	0.5864	0.1651
CG15747	CG15747	1635052_a_at	0.1091	0.6303	0.1501	0.5769	0.1469	0.4848	0.0949	0.8852	0.3086	0.1643	0.2137	0.2903	0.1622	0.8806	0.5522	0.1689	0.3899	0.3586
---	---	1635053_at	0.0586	0.7916	0.0950	0.4768	0.3282	0.0615	-0.0946	0.8776	0.0178	0.9546	0.1124	0.5935	-0.1587	0.7707	0.0930	0.7311	0.2517	0.2956
CG12069	PKA-like	1635054_at	0.1629	0.4387	0.0509	0.6248	0.2523	0.1289	0.1500	0.7401	-0.0202	0.9452	-0.1702	0.3517	0.0225	0.9734	0.0270	0.9162	0.0045	0.9866
CG3306	CG3306	1635055_at	0.0858	0.7748	0.0456	0.6556	0.4701	0.0302	0.1151	0.8107	-0.0087	0.9768	-0.1237	0.5000	-0.1840	0.7822	0.0350	0.9402	0.2190	0.4620
CG18810	CG18810	1635056_at	0.1273	0.5167	0.0373	0.8190	0.2649	0.2795	0.1627	0.7081	0.1480	0.4754	-0.0147	0.9535	0.0748	0.9246	0.0054	0.9929	-0.0694	0.8352
CG15445	CG15445	1635057_s_at	0.2567	0.1776	0.7564	0.0751	0.9306	0.0004	0.2308	0.4846	-0.1211	0.5169	-0.3519	0.0328	0.0265	0.9875	0.3546	0.3397	0.3281	0.3977
CG6937	CG6937	1635058_at	0.6213	0.0775	0.4571	0.0949	0.7110	0.0014	0.2298	0.6947	0.4556	0.0926	0.2257	0.3654	0.0742	0.9552	0.3749	0.3178	0.3007	0.4497
---	---	1635059_at	-0.0857	0.5706	0.2978	0.0615	0.2687	0.1059	-0.2519	0.3147	-0.3062	0.0428	-0.0543	0.7291	0.0313	0.9705	0.0557	0.8499	0.0244	0.9351
CG15068	CG15068	1635060_at	2.1687	0.0020	2.2001	0.1059	3.7826	0.0002	1.2492	0.1462	-0.7335	0.1561	-1.9827	0.0019	-0.4115	0.9087	-0.7704	0.5622	-0.3588	0.8209
---	---	1635061_at	-0.6648	0.0213	0.5531	0.0136	-0.0177	0.9482	-0.3451	0.3166	-0.9500	0.0013	-0.6049	0.0055	0.2121	0.8145	0.2382	0.5414	0.0261	0.9622
Smg1	FRAP-related	1635062_at	0.1836	0.3367	-0.4689	0.4209	-0.2534	0.1851	0.0680	0.9247	0.5413	0.0205	0.4733	0.0225	-0.1216	0.9340	-0.2514	0.6166	-0.1298	0.8255
snRNP70K	U1 snRNP-specifi	1635063_at	-0.3782	0.4528	0.5442	0.4607	0.3112	0.3586	-0.0676	0.9098	-0.3145	0.0965	-0.2469	0.1427	0.2857	0.9280	0.7613	0.4667	0.4757	0.6734
---	---	1635064_at	-0.0424	0.8463	-0.1888	0.3910	-0.2167	0.4684	0.0421	0.9754	0.2313	0.5099	0.1892	0.5652	0.0281	0.9717	0.0344	0.9115	0.0063	0.9849
CG7025 /// DyakCG7025	CG7025	1635065_at	0.1493	0.4191	0.0382	0.8676	0.1858	0.4806	-0.1193	0.8350	-0.1616	0.4762	-0.0423	0.8679	-0.2709	0.7322	-0.3659	0.2629	-0.0950	0.8234
CG9063	CG9063	1635066_at	-0.2847	0.3825	-0.0061	0.9828	-0.0558	0.7434	0.1023	0.8350	-0.0455	0.8508	-0.1478	0.3931	0.1426	0.8814	0.3202	0.3720	0.1776	0.6524
CG14909	CG14909	1635067_at	0.3227	0.0505	0.2474	0.3015	0.2806	0.2377	-0.0048	0.9956	0.0258	0.9378	0.0306	0.9130	0.0698	0.9492	-0.0213	0.9686	-0.0911	0.8243
---	---	1635068_at	0.3617	0.0452	0.0308	0.8059	0.0557	0.7989	-0.0045	0.9956	0.1273	0.5460	0.1318	0.4804	0.0031	0.9970	-0.1776	0.3669	-0.1807	0.3803
---	---	1635069_at	0.0350	0.8926	0.2673	0.1147	0.1820	0.3230	0.1032	0.8449	-0.1640	0.4132	-0.2672	0.1225	0.2183	0.8049	0.1440	0.7401	-0.0743	0.8835
CG9953	CG9953	1635070_at	0.7116	0.0030	0.9857	0.0144	0.7325	0.0035	0.3276	0.2506	0.1101	0.5469	-0.2174	0.1510	0.5149	0.3712	0.3496	0.2430	-0.1652	0.6216
CG14950	CG14950	1635071_at	0.1846	0.3390	0.1116	0.4514	0.0997	0.5520	0.1911	0.7380	0.0813	0.7928	-0.1098	0.6776	-0.0452	0.9653	0.0433	0.9204	0.0885	0.7909
CG7214	CG7214	1635072_at	1.9250	0.1754	0.0563	0.7968	0.1923	0.2607	0.1857	0.7845	-0.1294	0.6865	-0.3151	0.2082	0.0684	0.9923	-1.5407	0.3338	-1.6091	0.3368
CG17108 /// DsimCG17108	CG17108	1635073_at	2.5777	0.0081	0.4678	0.4284	2.0266	0.0003	0.7689	0.5255	0.2007	0.7965	-0.5682	0.3267	-0.7545	0.6955	-1.9824	0.0391	-1.2279	0.1551
---	---	1635074_s_at	0.0991	0.7858	-0.7615	0.0470	-0.9234	0.0032	-0.0676	0.9592	0.5894	0.0904	0.6570	0.0406	0.1652	0.9095	-0.1350	0.8481	-0.3002	0.5869
CG6852	CG6852	1635075_at	-0.0155	0.9692	0.4819	0.0695	0.3473	0.0688	-0.0469	0.9441	-0.4551	0.0253	-0.4083	0.0250	0.0168	0.9928	-0.1788	0.7200	-0.1956	0.6853
CG12674	CG12674	1635076_at	0.0053	0.9819	0.0915	0.3528	0.1819	0.2344	-0.0719	0.9186	-0.1414	0.5451	-0.0695	0.7723	-0.0062	0.9944	0.0919	0.6912	0.0980	0.6607
---	---	1635077_at	0.2653	0.3683	0.1646	0.3079	0.3442	0.0644	0.0869	0.8180	-0.0483	0.7897	-0.1352	0.3157	-0.1448	0.8943	-0.0792	0.8976	0.0656	0.9077
CG10617 /// DmirCG10617	CG10617	1635078_at	-0.0695	0.7644	-0.0963	0.5715	-0.1853	0.1964	-0.0936	0.8424	-0.0823	0.6797	0.0113	0.9593	-0.0879	0.8999	-0.1776	0.5042	-0.0897	0.7647
CG1550	CG1550	1635079_at	-0.3100	0.1519	0.5106	0.0375	0.9899	0.0004	0.2997	0.2654	-0.8006	0.0010	-1.1003	0.0002	-0.1095	0.8465	0.1282	0.6097	0.2376	0.3221
CG6059	CG6059	1635080_at	0.6708	0.0160	0.0947	0.4827	0.4028	0.0844	0.0098	0.9952	-0.1419	0.7077	-0.1517	0.6549	-0.1874	0.7772	-0.3833	0.1699	-0.1959	0.5141
---	---	1635081_at	0.1386	0.3638	0.2268	0.2959	-0.0835	0.6398	0.0574	0.9071	-0.0696	0.7033	-0.1270	0.3880	0.2951	0.6898	0.0293	0.9521	-0.2658	0.3921
---	---	1635082_at	0.1958	0.2482	0.0496	0.6105	-0.0295	0.8614	-0.1389	0.6718	-0.1644	0.2943	-0.0255	0.8908	0.0829	0.8689	-0.1477	0.4598	-0.2306	0.2614
CG32458	CG32458	1635083_at	0.1418	0.5608	0.6079	0.0986	0.1971	0.4106	0.0670	0.9436	0.1688	0.5597	0.1018	0.7214	0.6633	0.2884	0.6522	0.0660	-0.0111	0.9841
CG31413	CG31413	1635084_at	0.0638	0.6997	-0.0190	0.9304	0.0434	0.8136	-0.0390	0.9558	0.0231	0.9305	0.0621	0.7653	-0.0226	0.9848	0.0012	0.9991	0.0239	0.9491
---	---	1635085_at	0.1640	0.3271	-0.0415	0.7695	0.1453	0.5461	0.2265	0.5785	0.2130	0.3124	-0.0136	0.9591	0.0849	0.8999	-0.0024	0.9972	-0.0873	0.7625
CG4666 /// DyakCG4666	CG4666	1635086_at	1.3552	0.1963	2.2433	0.0141	2.5772	0.0039	1.2932	0.3348	-0.2060	0.8352	-1.4992	0.0370	0.9989	0.7305	0.8482	0.5076	-0.1507	0.9313
CG30352	CG30352	1635087_at	0.0098	0.9753	0.0718	0.5054	-0.0987	0.6511	-0.0967	0.8653	-0.0990	0.6658	-0.0023	0.9925	-0.0435	0.9778	0.0880	0.8675	0.1315	0.7612
CG1444	CG1444	1635088_at	0.9718	0.0431	0.8356	0.0321	1.6765	0.0007	0.3493	0.6141	-0.0714	0.8772	-0.4207	0.1745	-0.5515	0.5200	-0.1190	0.8248	0.4325	0.3027
tun	tungus	1635089_at	0.6576	0.0508	0.8356	0.0872	0.6635	0.0153	-0.1431	0.7604	0.8081	0.0026	0.9511	0.0008	-0.0185	0.9928	0.9455	0.0661	0.9640	0.0786
---	---	1635090_at	0.1888	0.1877	0.0932	0.4236	0.0437	0.8064	-0.0008	0.9988	-0.0189	0.9267	-0.0181	0.9197	0.0590	0.9457	0.0223	0.9590	-0.0366	0.9194
---	---	1635091_at	0.3359	0.0929	-0.0886	0.6498	0.1904	0.2102	0.2182	0.6006	0.2837	0.1740	0.0655	0.7786	-0.0497	0.9587	-0.1652	0.5523	-0.1155	0.6969
CG33557	CG33557	1635092_at	-1.3699	0.0139	0.0693	0.7494	-0.1463	0.4845	-0.1092	0.9071	-1.5732	0.0009	-1.4640	0.0007	-0.0009	0.9998	-0.0466	0.9462	-0.0456	0.9387
CG32369	CG32369	1635093_a_at	0.5980	0.4363	1.0857	0.0625	0.2229	0.2656	0.5514	0.1394	0.5358	0.0255	-0.0155	0.9572	1.3591	0.5126	1.0813	0.2520	-0.2778	0.8209
---	---	1635094_at	-0.0082	0.9728	0.0175	0.9542	0.0454	0.8320	-0.0343	0.9668	-0.0268	0.9296	0.0075	0.9781	0.0043	0.9971	0.2058	0.4813	0.2015	0.4977
CG14586	CG14586	1635095_at	0.0837	0.6434	0.1984	0.2761	0.1307	0.4945	-0.0185	0.9834	0.0078	0.9801	0.0263	0.9151	-0.0308	0.9762	0.0978	0.7363	0.1287	0.6316
lola	longitudinals abse	1635096_at	0.8012	0.0071	-0.1914	0.7647	-0.7404	0.1340	-0.1952	0.7024	1.6997	0.0003	1.8949	0.0001	0.3249	0.9095	0.4450	0.6986	0.1201	0.9331
---	---	1635097_at	0.2732	0.1004	-0.1742	0.2494	0.0607	0.8100	-0.0583	0.9558	0.1022	0.7641	0.1605	0.5691	-0.1716	0.8062	-0.1041	0.7692	0.0674	0.8602
CG4789	CG4789	1635098_a_at	0.0948	0.7143	0.7217	0.0460	0.8785	0.0276	0.2376	0.7328	-0.5582	0.0748	-0.7958	0.0116	0.0616	0.9765	0.1262	0.8550	0.0645	0.9259
CG3590	CG3590	1635099_at	0.1081	0.6470	-0.0191	0.9660	0.0354	0.9176	0.0642	0.9262	0.1975	0.3520	0.1333	0.5075	0.0517	0.9816	0.2605	0.6458	0.2089	0.7253
---	---	1635100_at	0.1974	0.2499	0.0821	0.5040	0.2573	0.1701	-0.0008	0.9994	-0.0066	0.9826	-0.0058	0.9825	-0.1075	0.8202	-0.2597	0.1669	-0.1523	0.4450
pad	poils au dos	1635101_at	0.1793	0.5351	0.0879	0.5221	0.5547	0.0087	0.1374	0.7670	0.1454	0.4816	0.0080	0.9738	-0.4125	0.6749	0.0466	0.9430	0.4591	0.2743
syt	Synaptotagmin	1635102_at	-0.0235	0.9584	-0.0791	0.7852	0.1252	0.4222	0.2024	0.8251	0.0159	0.9772	-0.1865	0.6012	0.0068	0.9946	0.0930	0.7393	0.0862	0.7534
trr	trithorax-related	1635103_at	0.1943	0.6457	-0.3287	0.6714	-0.0563	0.8362	0.1684	0.8164	0.7063	0.0215	0.5379	0.0412	-0.2196	0.9340	0.1223	0.9272	0.3420	0.7205
CG33936	CG33936	1635104_s_at	-0.3559	0.3852	0.5053	0.0345	-0.0052	0.9814	-0.2546	0.5988	-0.4332	0.0777	-0.1787	0.4359	0.1328	0.9400	0.2432	0.6862	0.1104	0.8801
---	---	1635105_at	0.5338	0.0097	0.2085	0.3986	0.0316	0.8516	-0.1212	0.7688	0.0057	0.9823	0.1269	0.4400	0.0051	0.9963	-0.0396	0.9120	-0.0446	0.8893
CG30111	CG30111	1635106_at	-0.0115	0.9474	-0.3674	0.2750	-0.3814	0.0911	0.0681	0.9351	0.3275	0.1812	0.2594	0.2402	0.1651	0.8206	0.0077	0.9911	-0.1574	0.6297
---	---	1635107_at	0.3381	0.1589	0.1106	0.3352	-0.0486	0.8478	-0.2782	0.5505	-0.0907	0.7519	0.1876	0.4035	0.0358	0.9742	-0.0444	0.9158	-0.0802	0.8116
CG5964	CG5964	1635108_at	-0.0566	0.8552	0.0023	0.9865	0.1076	0.6779	0.1322	0.8189	0.0139	0.9685	-0.1182	0.6019	-0.0045	0.9971	-0.0671	0.8760	-0.0626	0.8779
CG5888	CG5888	1635109_at	-1.5276	0.0020	-2.0856	0.0159	-2.7401	0.0000	-0.4957	0.2535	-0.0988	0.7470	0.3969	0.0861	0.2598	0.8023	-0.4266	0.3105	-0.6864	0.1409
Cyp6a13	Cyp6a13	1635110_at	0.9899	0.0043	0.9321	0.0075	1.1116	0.0004	0.0180	0.9854	-0.1922	0.3961	-0.2101	0.2931	-0.2188	0.7241	-0.0751	0.8349	0.1438	0.6193
CG3358	CG3358	1635111_s_at	0.0702	0.8226	0.1442	0.6894	0.1173	0.5071	-0.0361	0.9558	0.1609	0.3692	0.1970	0.2089	-0.0258	0.9914	0.2561	0.6454	0.2818	0.6100
CG34143	CG34143	1635112_at	0.4470	0.0489	-0.0388	0.7995	0.1785	0.2637	0.0753	0.8526	0.2118	0.1478	0.1365	0.3065	-0.1155	0.8940	-0.1920	0.5762	-0.0765	0.8570
CG34104	CG34104	1635113_at	-0.2916	0.3697	-0.2208	0.2937	-0.6678	0.0182	-0.3249	0.6607	0.1402	0.7373	0.4651	0.1411	0.0859	0.9467	0.1331	0.7852	0.0472	0.9319
GABA-B-R3	metabotropic GAE	1635114_a_at	-0.2414	0.2211	0.0860	0.4476	0.0320	0.9003	-0.0903	0.8577	-0.3540	0.0541	-0.2637	0.1010	-0.0183	0.9903	-0.0739	0.8740	-0.0556	0.9023

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG30325	CG30325	1635115_at	0.1172	0.4117	0.0000	1.0000	0.0000	0.9998	0.0638	0.8948	-0.0004	0.9987	-0.0641	0.7046	-0.0627	0.9056	-0.0484	0.8518	0.0142	0.9588
CG7518	CG7518	1635116_a_at	-1.0259	0.2581	-0.5489	0.7097	-0.4774	0.1803	0.2330	0.5626	-0.0575	0.8240	-0.2905	0.1140	0.1265	0.9869	0.3840	0.8794	0.2575	0.9164
fu	fused	1635117_at	-0.0005	0.9994	-0.2969	0.1107	-0.3561	0.0630	0.0793	0.8794	0.4094	0.0298	0.3300	0.0450	0.2328	0.8609	0.2291	0.7065	-0.0038	0.9973
CG14658	CG14658	1635118_at	0.3305	0.1231	0.3004	0.1746	0.3945	0.1573	0.0194	0.9848	-0.0245	0.9414	-0.0439	0.8716	-0.1284	0.8692	-0.1246	0.7299	0.0038	0.9945
---	---	1635119_at	0.7926	0.0722	0.2103	0.7435	0.2411	0.1929	-0.1373	0.8817	0.7167	0.0305	0.8541	0.0090	-0.1609	0.9398	0.2355	0.7653	0.3964	0.5667
CG8446	CG8446	1635120_at	0.5538	0.0327	1.3210	0.0083	1.5829	0.0011	0.1544	0.8038	-0.8548	0.0051	-1.0091	0.0015	-0.0872	0.9530	-0.1953	0.6949	-0.1081	0.8495
Vps20	Vps20	1635121_a_at	-0.1381	0.6212	0.0613	0.8485	0.3849	0.0910	0.1941	0.5199	-0.1957	0.2161	-0.3899	0.0139	-0.1599	0.8940	0.0483	0.9467	0.2082	0.6677
---	---	1635122_at	0.1332	0.4136	-0.0263	0.7950	-0.1120	0.6196	0.3427	0.2966	0.4556	0.0260	0.1129	0.5516	0.0334	0.9635	0.0668	0.7883	0.0334	0.9031
glu	gluon	1635123_at	-0.1043	0.5447	-0.1763	0.4974	-0.1644	0.4112	-0.1826	0.6325	0.3087	0.0985	0.4913	0.0102	-0.3637	0.7215	0.1552	0.7621	0.5189	0.2398
ase	Asense	1635124_at	-0.1700	0.5693	-0.0385	0.7074	-0.1097	0.4711	-0.0849	0.8605	-0.1822	0.2969	-0.0973	0.5712	-0.0635	0.9467	-0.1040	0.7692	-0.0405	0.9197
CG6206	CG6206	1635125_a_at	1.3098	0.0021	0.5614	0.0311	0.8739	0.0009	0.1677	0.7161	0.3745	0.0768	0.2068	0.2761	-0.2413	0.7464	-0.4789	0.1321	-0.2376	0.4683
---	---	1635126_a_at	-1.0307	0.0843	-0.3606	0.5248	-1.4333	0.0004	0.1418	0.9068	-0.0070	0.9901	-0.1488	0.7203	1.2289	0.3985	0.6998	0.3685	-0.5291	0.5243
---	---	1635127_at	-1.0676	0.0145	-1.0128	0.0818	-1.9386	0.0015	-0.3379	0.6056	0.1885	0.6010	0.5264	0.0754	0.3280	0.7810	0.0192	0.9842	-0.3088	0.5680
ash2	many abnormal di	1635128_a_at	-0.6857	0.0118	-0.2628	0.0507	-0.3400	0.0576	-0.1055	0.8584	-1.0159	0.0012	-0.9104	0.0012	0.0164	0.9898	-0.4912	0.0761	-0.5076	0.0889
CG6332	CG6332	1635129_at	0.4375	0.0981	0.0290	0.8861	0.1512	0.4732	0.1457	0.6888	0.1227	0.4877	-0.0230	0.9101	-0.0254	0.9914	-0.1197	0.8725	-0.0943	0.8949
SamDC	S-adenosylmethio	1635130_a_at	0.1542	0.5748	0.9832	0.0163	1.3085	0.0110	-0.0032	0.9987	-1.0118	0.0175	-1.0087	0.0110	-0.2447	0.8609	0.0093	0.9941	0.2539	0.6817
CG4525	CG4525	1635131_at	0.1695	0.4173	-0.2005	0.3311	0.0202	0.9178	0.0214	0.9768	0.0852	0.6796	0.0638	0.7439	-0.1141	0.8878	-0.0881	0.8248	0.0260	0.9514
CG4580	CG4580	1635132_at	-0.1914	0.2699	-0.0118	0.9298	-0.4191	0.0606	-0.3688	0.4420	-0.1025	0.7399	0.2663	0.2586	0.0193	0.9848	0.0942	0.7198	0.0749	0.7822
---	---	1635133_at	0.0080	0.9713	-0.1135	0.5051	0.0012	0.9960	0.1209	0.8738	0.1409	0.6336	0.0200	0.9525	0.0012	0.9994	-0.0434	0.8749	-0.0446	0.8599
---	---	1635134_at	-0.0906	0.6056	-0.1436	0.1697	-0.0215	0.9231	0.1804	0.6171	0.1715	0.3498	-0.0089	0.9688	0.0491	0.9514	0.0119	0.9780	-0.0371	0.9096
---	---	1635135_at	-0.1479	0.8706	-0.8642	0.1594	-1.4198	0.0060	0.0469	0.9671	1.4677	0.0011	1.4208	0.0007	0.5324	0.8541	0.6683	0.6003	0.1359	0.9350
CG30471	CG30471	1635136_at	0.1469	0.5763	-0.0652	0.7385	0.1421	0.4147	0.1707	0.7271	0.2177	0.3272	0.0470	0.8534	-0.0121	0.9923	-0.0190	0.9682	-0.0069	0.9879
CG5946	CG5946	1635137_a_at	-0.1813	0.2487	-0.4612	0.0176	-0.1599	0.3608	-0.0422	0.9445	-0.0197	0.9363	0.0225	0.9154	-0.3569	0.3844	-0.3865	0.0968	-0.0296	0.9246
Fur1	Furin-like proteas	1635138_at	-0.8126	0.1827	-1.2125	0.0452	-2.1074	0.0003	-0.1476	0.8816	0.9669	0.0115	1.1146	0.0037	0.7286	0.7116	0.6407	0.4359	-0.0879	0.9394
CG13340	CG13340	1635139_at	0.0148	0.9625	0.0658	0.5540	0.2510	0.1787	0.0495	0.9339	-0.1580	0.3781	-0.2075	0.1848	0.0052	0.9964	-0.0621	0.8889	-0.0673	0.8684
CG15515	CG15515	1635140_at	0.0415	0.9188	0.0967	0.4364	-0.1749	0.2986	-0.0114	0.9880	-0.0445	0.8469	-0.0331	0.8753	0.2063	0.8521	0.0393	0.9587	-0.1670	0.7514
CG14675	CG14675	1635141_at	0.1370	0.4335	0.0606	0.7329	0.0418	0.8590	0.1239	0.8215	0.1330	0.5649	0.0091	0.9727	0.1242	0.8380	-0.0640	0.8527	-0.1882	0.4683
---	---	1635142_at	-0.0002	0.9994	0.1032	0.4517	-0.1872	0.2764	0.0658	0.8942	0.1061	0.5493	0.0403	0.8297	0.1423	0.8390	-0.0440	0.9222	-0.1862	0.5403
Aut1	Autophagy-specifi	1635143_at	0.1644	0.4060	0.3424	0.0683	0.7015	0.0026	0.1515	0.7053	-0.2877	0.1162	-0.4392	0.0156	-0.1719	0.7893	-0.0361	0.9341	0.1358	0.6434
klg	klingon	1635144_at	0.0582	0.7768	0.1047	0.4829	0.0731	0.7108	-0.1884	0.7478	-0.1116	0.7053	0.0768	0.7880	-0.1149	0.8097	-0.0098	0.9783	0.1051	0.6225
---	---	1635145_at	0.3832	0.1120	-0.0001	1.0000	0.2032	0.3132	0.1258	0.8794	0.3324	0.2425	0.2066	0.4391	-0.1622	0.8141	-0.2353	0.4037	-0.0731	0.8407
---	---	1635146_at	0.0915	0.5680	-0.0371	0.7026	0.0231	0.9061	-0.0810	0.8738	-0.0166	0.9505	0.0644	0.7370	-0.0313	0.9651	-0.0339	0.9080	-0.0026	0.9940
---	---	1635147_at	0.0813	0.6735	-0.0986	0.4624	0.0587	0.7865	-0.0289	0.9666	0.1243	0.5184	0.1532	0.3580	-0.0879	0.9142	-0.0632	0.8781	0.0247	0.9512
ninaG	ninaG	1635148_at	0.3199	0.1854	0.0136	0.9204	0.0029	0.9927	0.1142	0.7970	0.1652	0.3695	0.0511	0.8009	-0.0023	0.9990	-0.0151	0.9763	-0.0128	0.9777
CG32117	CG32117	1635149_at	0.0452	0.8803	0.0431	0.7111	0.0256	0.8931	0.0497	0.9475	-0.1052	0.6622	-0.1549	0.4386	0.0820	0.9309	0.0131	0.9821	-0.0689	0.8654
---	---	1635150_at	0.3855	0.1621	0.0932	0.5244	0.1658	0.5245	0.0032	0.9962	0.0997	0.7069	0.0965	0.6893	0.0196	0.9912	-0.0744	0.8852	-0.0940	0.8354
COQ7	COQ7	1635151_at	-1.5324	0.0062	-1.7861	0.0083	-1.0042	0.0095	0.5433	0.5168	-0.3748	0.4106	-0.9181	0.0280	-0.2493	0.8160	-0.4962	0.2456	-0.2469	0.6043
CG12376	CG12376	1635152_at	0.1411	0.4635	-0.0119	0.9113	0.0477	0.7738	0.1049	0.7795	0.1239	0.4430	0.0190	0.9209	-0.1170	0.9066	-0.0462	0.9353	0.0709	0.8854
Gfat2	Glutamine:fructos	1635153_at	0.1895	0.5223	-0.0878	0.8458	0.3054	0.1512	0.2975	0.3294	0.8119	0.0015	0.5144	0.0064	0.0537	0.9831	0.5635	0.3217	0.5098	0.3921
---	---	1635154_s_at	-0.1207	0.5111	0.2615	0.0780	-0.1242	0.4412	-0.3499	0.1580	-0.3821	0.0190	-0.0322	0.8532	0.2183	0.6955	0.2036	0.3720	-0.0147	0.9661
Tim17b2	Translocase inner	1635155_a_at	0.0442	0.8009	0.0034	0.9829	-0.1037	0.6038	0.0188	0.9777	0.0923	0.6323	0.0735	0.6851	0.1872	0.7485	0.0417	0.9125	-0.1455	0.5811
Pka-R1	cAMP-dependent	1635156_s_at	-1.1891	0.0115	-0.2629	0.5385	-1.0501	0.0104	-0.4131	0.3004	-0.8871	0.0033	-0.4739	0.0313	0.3473	0.8513	0.0180	0.9924	-0.3293	0.7015
CG33057	CG33057	1635157_at	-0.0573	0.8142	0.0933	0.7479	-0.4919	0.0313	-0.1033	0.8979	0.0626	0.8555	0.1660	0.5176	0.2407	0.7686	0.0951	0.8383	-0.1456	0.7099
CG13924	CG13924	1635158_at	0.1417	0.3849	0.2932	0.3099	0.3691	0.0546	-0.1183	0.8288	-0.2692	0.1981	-0.1509	0.4459	-0.0704	0.9221	-0.0459	0.8989	0.0245	0.9423
---	---	1635159_at	0.1028	0.5803	-0.0084	0.9476	0.1280	0.4342	-0.0001	0.9999	-0.0725	0.6902	-0.0724	0.6590	-0.0324	0.9635	-0.0698	0.7645	-0.0374	0.8880
CG17763	CG17763	1635160_at	0.0046	0.9877	0.1169	0.4305	0.2954	0.0858	0.0890	0.8796	-0.0076	0.9804	-0.0967	0.6402	-0.0217	0.9841	0.1917	0.4355	0.2134	0.3939
CG7532	CG7532	1635161_at	-0.4779	0.5038	-0.7473	0.0237	-0.6594	0.0682	0.0208	0.9781	-0.2069	0.2897	-0.2278	0.1880	-0.1714	0.9405	-0.5970	0.3922	-0.4255	0.5708
CG15314	CG15314	1635162_at	0.0524	0.8344	0.2172	0.1868	0.0074	0.9734	-0.2212	0.5280	-0.0105	0.9705	0.2108	0.1977	-0.0128	0.9916	0.0027	0.9972	0.0155	0.9714
CG5731	CG5731	1635163_at	-0.5520	0.2368	-1.6193	0.0452	-1.3256	0.0042	0.2852	0.8217	0.3120	0.5575	0.0268	0.9659	-0.0251	0.9923	-0.5699	0.3307	-0.5447	0.3782
---	---	1635164_s_at	0.2644	0.1924	-0.1570	0.4022	-0.2613	0.4166	-0.2813	0.6633	0.1114	0.7640	0.3927	0.1543	-0.1065	0.8956	-0.0638	0.8845	0.0427	0.9189
CG12090 /// CG33971	CG12090 /// CG3	1635165_a_at	1.3849	0.0781	0.8740	0.3392	1.0059	0.0040	0.1136	0.8908	0.3000	0.2851	0.1864	0.4835	-0.1152	0.9824	-0.1403	0.9425	-0.0252	0.9902
CG34400	CG17368	1635166_at	-0.0118	0.9535	0.1410	0.3535	-0.0254	0.9112	-0.0373	0.9621	-0.1761	0.4130	-0.1388	0.4862	0.0693	0.8870	0.0703	0.7439	0.0010	0.9979
---	---	1635167_at	0.1821	0.3355	0.0183	0.9148	0.1320	0.5578	0.0909	0.9218	0.0587	0.8738	-0.0322	0.9252	-0.0585	0.9495	-0.1049	0.7459	-0.0464	0.9030
CG5872	CG5872	1635168_at	0.1565	0.7145	-0.1207	0.7106	-0.1176	0.7315	-0.3731	0.5757	0.2067	0.5769	0.5798	0.0617	-0.2953	0.7464	-0.0106	0.9901	0.2846	0.4730
---	---	1635169_at	0.3369	0.3310	0.1136	0.6169	0.0712	0.6840	0.0435	0.9468	0.1437	0.4540	0.1002	0.5869	-0.0512	0.9807	-0.0942	0.8878	-0.0430	0.9469
CG2901 /// CG7755	CG7755 /// CG29	1635170_at	0.1048	0.5301	0.0112	0.9308	-0.0805	0.7006	-0.0880	0.9065	-0.0328	0.9226	0.0553	0.8426	-0.0268	0.9717	-0.0425	0.8746	-0.0156	0.9525
---	---	1635171_at	0.2038	0.4860	0.1373	0.4914	0.3612	0.1100	0.0301	0.9647	-0.1064	0.5869	-0.1364	0.4179	-0.2588	0.7464	-0.1604	0.6695	0.0984	0.8168
---	---	1635172_at	0.1744	0.4290	0.2164	0.3437	0.0277	0.9167	0.0282	0.9834	0.0407	0.9252	0.0125	0.9741	0.2712	0.7095	0.0058	0.9935	-0.2653	0.3921
pgant5	polypeptide GalN/	1635173_at	0.3045	0.2691	1.4271	0.0185	0.0395	0.8281	-0.1258	0.7719	-0.3965	0.0367	-0.2707	0.0970	1.1245	0.1902	0.7374	0.1054	-0.3870	0.3939
---	---	1635174_at	-0.0519	0.7632	0.0218	0.8794	0.0658	0.7416	0.0769	0.9057	-0.0305	0.9174	-0.1074	0.6128	0.0226	0.9781	0.1071	0.6093	0.0845	0.6990
CG17121	CG17121	1635175_at	-1.2070	0.0097	-0.7029	0.1410	-0.9064	0.0013	-0.0366	0.9753	-0.7070	0.0224	-0.6704	0.0173	0.1631	0.9092	-0.1710	0.7861	-0.3341	0.5299
---	---	1635176_at	0.2905	0.2290	0.5183	0.0108	0.4420	0.1830	0.0749	0.9035	-0.2599	0.1913	-0.3348	0.0645	0.2144	0.8076	0.0704	0.8988	-0.1440	0.7323
CG7770 /// DyakCG7770	CG7770	1635177_at	0.1209	0.4404	0.3912	0.1053	0.2524	0.1008	0.1501	0.7083	0.2417	0.1845	0.0915	0.6237	0.4304	0.4415	0.5925	0.0637	0.1621	0.6059
CG33299	CG33299	1635178_at	0.1295	0.6618	0.0425	0.6799	0.1289	0.6301	0.1193	0.8589	0.1312	0.6247	0.0119	0.9688	0.0326	0.9816	-0.0054	0.9941	-0.0381	0.9350
---	---	1635179_at	0.1843	0.2303	-0.0242	0.8503	0.1429	0.3910	0.0907	0.8471	0.2390	0.1634	0.1483	0.3487	-0.0808	0.8903	-0.0396	0.9068	0.0413	0.8918
CG32194	CG32194	1635180_at	-0.1451	0.4393	-0.1006	0.4149	-0.3240	0.1163	-0.1641	0.7432	-0.0914	0.7217	0.0727	0.7637	-0.1467	0.7826	-0.0520	0.8749	0.0947	0.7154
CG3337	CG3337	1635181_at	-0.1873	0.5126	0.3504	0.3908	0.2011	0.2833	-0.2587	0.6402	-0.5042	0.0656	-0.2455	0.3194	0.0549	0.9717	0.0504	0.9353	-0.0045	0.9948
wts	warts	1635182_at	-0.2668	0.6392	-0.6484	0.0791	-0.6363	0.0082	0.0302	0.9723	0.4778	0.0349	0.4476	0.0289	0.0828	0.9775	0.0025	0.9994	-0.0802	0.9342
Spn43Ab	Serine protease ir	1635183_at	0.3314	0.0648	0.2323	0.5825	0.3995	0.0738	-0.1976	0.6679	-0.3407	0.1190	-0.1431	0.4976	-0.3601	0.7436	-0.5001	0.2643	-0.1400	0.8050

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1635184_at	0.4374	0.0831	0.2470	0.4056	0.1502	0.6538	-0.1997	0.7888	0.1114	0.7612	0.3110	0.2580	-0.0228	0.9884	-0.0914	0.8438	-0.0686	0.8837
CG7560	CG7560	1635185_at	2.4650	0.0011	0.8073	0.2706	2.5578	0.0007	1.1039	0.0458	1.4490	0.0014	0.3451	0.2194	-0.6641	0.7739	-0.1790	0.9075	0.4851	0.6546
Syx13	syntaxin	1635186_at	-0.4937	0.0342	-0.3588	0.1680	-0.4140	0.0159	0.1140	0.8112	0.0764	0.7303	-0.0376	0.8649	0.2243	0.7628	0.2481	0.4213	0.0238	0.9575
CG1646	CG1646	1635187_s_at	0.2399	0.1947	0.4651	0.2176	0.3463	0.0707	0.0665	0.9066	-0.2715	0.1350	-0.3380	0.0440	0.0782	0.9514	-0.0321	0.9590	-0.1102	0.8138
CG6675	CG6675	1635188_at	-0.3648	0.1237	-0.0042	0.9771	-0.3630	0.0768	-0.2929	0.5336	-0.5737	0.0274	-0.2808	0.2008	-0.0059	0.9943	-0.0770	0.7275	-0.0711	0.7481
Drs	Drosomycin	1635189_at	-0.0191	0.9090	0.6142	0.0527	0.3955	0.1217	-0.3377	0.5917	-0.6940	0.0364	-0.3563	0.2118	-0.0164	0.9845	-0.0038	0.9931	0.0126	0.9640
---	---	1635190_at	0.2305	0.2453	0.1892	0.1175	0.2829	0.1247	-0.0846	0.8736	-0.0235	0.9282	0.0611	0.7628	0.0513	0.9467	0.0124	0.9755	-0.0388	0.9031
Tsp42Ed	tetraspanin 42E	1635191_at	0.0941	0.6633	-0.4537	0.0473	-0.2712	0.2302	0.0683	0.9108	0.4776	0.0215	0.4093	0.0257	-0.0303	0.9848	0.0099	0.9905	0.0401	0.9372
svp	seven-up	1635192_at	0.1012	0.5678	-0.1040	0.3593	0.1618	0.2990	0.0607	0.8987	0.0835	0.6306	0.0229	0.9030	-0.1035	0.8609	-0.0826	0.7779	0.0209	0.9500
---	---	1635193_at	-0.0587	0.7194	0.0301	0.8547	0.0065	0.9727	-0.1119	0.7604	0.0041	0.9868	0.1159	0.4332	-0.1383	0.8202	-0.1467	0.5798	-0.0084	0.9842
CG13869 /// DsecCG13869	CG13869	1635194_at	0.4422	0.0696	-0.2752	0.1896	0.2288	0.1664	0.4691	0.1028	0.4679	0.0136	-0.0012	0.9955	-0.0149	0.9906	-0.1348	0.6408	-0.1200	0.6839
---	---	1635195_at	-0.0597	0.8032	0.1759	0.3426	0.1613	0.4733	-0.0306	0.9696	-0.0526	0.8446	-0.0219	0.9314	0.0277	0.9869	0.2029	0.6256	0.1753	0.6795
CG6836	CG6836	1635196_at	-1.2624	0.1915	-1.6908	0.0875	-2.6853	0.0014	-0.9180	0.1270	-1.1101	0.0085	-0.1921	0.5960	0.2393	0.9677	-1.3960	0.3620	-1.6353	0.3056
---	---	1635197_at	0.2515	0.1132	-0.0406	0.7187	0.0246	0.8999	-0.1030	0.8164	0.1291	0.4790	0.2322	0.1348	-0.0679	0.9457	-0.1345	0.6894	-0.0666	0.8674
CG10483	CG10483	1635198_at	-1.9940	0.0108	-2.2630	0.0183	-2.7465	0.0001	-0.2303	0.5363	-0.4037	0.0439	-0.1734	0.3289	0.4706	0.8122	-0.4852	0.5776	-0.9558	0.2614
CG6014	CG6014	1635199_at	-2.3242	0.0016	-1.8478	0.0197	-2.3955	0.0000	-0.4522	0.3925	0.0837	0.8223	0.5359	0.0494	0.1510	0.8967	0.5694	0.1749	0.4184	0.3476
CG11261	CG11261	1635200_at	-0.1124	0.7799	-0.2265	0.4174	-0.1264	0.7453	0.0414	0.9755	0.2717	0.4244	0.2304	0.4593	-0.1577	0.9011	-0.0286	0.9704	0.1291	0.8247
CG15824	CG15824	1635201_at	0.3466	0.2434	-0.0437	0.8002	-0.0347	0.8345	-0.0104	0.9903	0.0451	0.8546	0.0554	0.7943	-0.0242	0.9848	-0.2507	0.3669	-0.2265	0.4353
CG5640	CG5640	1635202_s_at	0.1530	0.4257	0.2360	0.5326	0.0458	0.8133	-0.2649	0.4586	-0.2161	0.2666	0.0488	0.8244	-0.0729	0.9619	-0.0328	0.9628	0.0401	0.9466
Dip2	dorsal interacting	1635203_at	-1.4408	0.0025	-1.8245	0.0135	-2.2070	0.0000	-0.0995	0.9046	0.2679	0.3278	0.3674	0.1263	0.1887	0.8425	-0.0809	0.8857	-0.2696	0.5045
CG13568	CG13568	1635204_at	-0.3141	0.1731	-0.2455	0.3392	-0.7853	0.0080	-0.1185	0.8743	-0.1552	0.5843	-0.0367	0.9078	0.0412	0.9742	-0.0831	0.8487	-0.1243	0.7353
CG10267	CG10267	1635205_at	0.0013	0.9964	0.4112	0.0711	-0.1095	0.5449	-0.4735	0.1462	-0.2183	0.2696	0.2551	0.1457	-0.0216	0.9862	0.2304	0.4147	0.2521	0.3859
CG32541 /// DsimCG32541	CG32541	1635206_at	0.0030	0.9915	0.0448	0.7623	0.2104	0.2237	-0.0795	0.8725	-0.0793	0.6877	0.0002	0.9995	-0.2687	0.7644	-0.0547	0.9275	0.2141	0.5990
CG32557	CG32557	1635207_at	0.2345	0.3778	0.2560	0.2345	0.3755	0.0469	-0.1253	0.8485	-0.3463	0.1468	-0.2210	0.3106	-0.0218	0.9851	0.0512	0.8981	0.0729	0.8261
CG9934	CG9934	1635208_at	0.1087	0.6310	0.4248	0.0817	0.2811	0.1674	-0.0423	0.9445	-0.0685	0.7380	-0.0262	0.9003	0.1670	0.8113	0.2301	0.4251	0.0631	0.8699
---	---	1635209_at	-0.1944	0.3334	0.1176	0.4296	0.0589	0.7674	0.0382	0.9558	-0.2361	0.2022	-0.2743	0.0983	-0.0518	0.9342	-0.0380	0.9009	0.0138	0.9628
Ppn	papilin	1635210_a_at	-2.5423	0.0011	-4.4054	0.0039	-4.5414	0.0000	-0.2724	0.3281	-0.1118	0.5123	0.1606	0.2699	-0.0955	0.9689	-1.8031	0.0276	-1.7075	0.0422
sbb	scribble	1635211_at	-0.2306	0.8326	-1.1660	0.2210	-1.5347	0.0030	0.0888	0.9218	1.0473	0.0031	0.9585	0.0027	0.4197	0.9246	0.1824	0.9402	-0.2373	0.9084
CG40176	CG40176	1635212_at	-0.0204	0.9234	0.1236	0.4341	-0.0652	0.6708	0.1158	0.7408	0.1523	0.3292	0.0365	0.8358	0.1980	0.6955	0.1731	0.3999	-0.0249	0.9322
CG40470	CG40470	1635213_at	0.2756	0.2052	0.0331	0.8399	0.0981	0.6027	0.1298	0.8485	0.3399	0.1683	0.2101	0.3568	0.0749	0.8846	0.0450	0.8721	-0.0299	0.9118
mus308	Nuclease-3	1635214_at	0.3922	0.0631	-0.4176	0.0270	-0.2369	0.2571	0.1744	0.7023	0.8291	0.0028	0.6547	0.0047	-0.0688	0.9449	-0.0587	0.8963	0.0101	0.9843
CG2059	CG2059	1635215_at	-0.3492	0.0698	-0.4007	0.1791	-0.2740	0.0924	0.1873	0.5832	-0.0710	0.7263	-0.2583	0.0959	-0.0605	0.9589	-0.1798	0.6209	-0.1194	0.7608
CG1407	CG1407	1635216_at	-0.5363	0.0209	0.1392	0.6343	0.3055	0.1829	-0.1289	0.8396	-0.5935	0.0209	-0.4647	0.0360	-0.2334	0.8009	0.0491	0.9363	0.2825	0.4791
Rgk2	Rgk2	1635217_at	0.0466	0.7722	0.2322	0.2131	0.0749	0.6121	-0.0374	0.9602	-0.1047	0.6333	-0.0673	0.7540	0.1571	0.7644	0.0637	0.8254	-0.0934	0.7053
CG11009	CG11009	1635218_at	-0.5949	0.0050	-0.4582	0.0599	-0.3696	0.0339	-0.0589	0.8923	-0.0987	0.5241	-0.0399	0.8073	-0.1689	0.7707	-0.0183	0.9645	0.1506	0.5648
CG12883	CG12883	1635219_at	-0.0470	0.8993	0.5509	0.1012	1.2739	0.0020	0.1286	0.8337	-0.5398	0.0278	-0.6685	0.0068	-0.3017	0.7686	0.2764	0.5431	0.5781	0.2059
Or83a	Odorant receptor	1635220_at	0.0622	0.7213	-0.0267	0.9359	-0.1947	0.2554	0.1131	0.7803	0.2679	0.1108	0.1548	0.3123	0.1316	0.8400	0.0891	0.7922	-0.0425	0.9092
CG32811	CG32811	1635221_at	-0.0069	0.9758	0.0790	0.5425	0.2015	0.1963	0.0482	0.9436	-0.0475	0.8509	-0.0957	0.6303	0.0514	0.9342	0.1152	0.5863	0.0637	0.7907
CG1513	CG1513	1635222_at	-0.6592	0.2699	0.5079	0.3614	0.7240	0.0022	-0.1303	0.9125	-0.6438	0.0875	-0.5134	0.1247	-0.1642	0.9449	0.7618	0.2714	0.9260	0.2136
CG30285	CG30285	1635223_at	-1.6420	0.0006	-1.5827	0.0145	-2.2582	0.0001	0.1285	0.8337	-0.0845	0.7614	-0.2131	0.3135	0.7405	0.2892	-0.0172	0.9796	-0.7577	0.0759
Jra	Jun oncogene	1635224_s_at	-0.7547	0.0238	-0.3880	0.1937	0.0426	0.8993	0.6129	0.1054	0.2680	0.2320	-0.3449	0.0862	0.2206	0.8760	0.6768	0.2064	0.4561	0.4172
l(2)37Cb	lethal (2) 37Cb	1635225_at	0.0589	0.8342	-0.0135	0.9003	0.1236	0.6430	0.0278	0.9744	0.4825	0.0281	0.4547	0.0228	-0.1694	0.8480	0.3274	0.3667	0.4967	0.2012
lox2	lysyl oxidase-like 2	1635226_at	-0.0805	0.6356	-0.0140	0.8948	0.0123	0.9621	0.1708	0.6578	-0.0139	0.9600	-0.1847	0.2687	0.0553	0.9665	-0.3028	0.3779	-0.3581	0.3171
ImpL3	lactic DH	1635227_at	-1.6995	0.0037	-2.6177	0.0099	-2.1330	0.0018	0.4428	0.4223	1.1321	0.0036	0.6894	0.0198	0.1056	0.9741	0.3377	0.7078	0.2321	0.8152
---	---	1635228_s_at	0.2009	0.5302	0.0427	0.6809	0.0712	0.6790	0.0077	0.9922	0.2083	0.1983	0.2006	0.1666	0.1089	0.9092	0.0617	0.9054	-0.0473	0.9194
Elo68alpha	CG32072	1635229_at	0.0897	0.8116	-0.0361	0.7473	0.0970	0.7571	0.1888	0.8267	0.0761	0.8616	-0.1127	0.7565	0.1032	0.9514	-0.0700	0.9291	-0.1732	0.7663
---	---	1635230_at	-0.0865	0.5953	-0.0027	0.9828	-0.0019	0.9943	0.0196	0.9761	0.0174	0.9407	-0.0022	0.9915	-0.1044	0.8494	-0.0128	0.9739	0.0916	0.7247
CG14943	CG14943	1635231_at	0.1951	0.2431	0.1163	0.7218	0.2092	0.3195	-0.0910	0.8931	0.1306	0.5956	0.2216	0.2773	0.0513	0.9657	0.2127	0.5214	0.1614	0.6392
CG13983	CG13983	1635232_at	-0.0628	0.7422	0.0735	0.6895	0.1027	0.6316	0.0328	0.9603	-0.1590	0.3794	-0.1919	0.2260	-0.0777	0.9092	0.1139	0.6694	0.1917	0.4405
CG15258	CG15258	1635233_at	0.0569	0.7112	-0.3000	0.3179	-0.1595	0.4031	0.0116	0.9909	0.2551	0.2534	0.2436	0.2219	-0.1498	0.7720	-0.0734	0.7906	0.0764	0.7664
Mcm3	Minichromosome	1635234_at	0.3769	0.4843	0.0181	0.9829	-0.4258	0.1914	-0.1989	0.7079	0.9847	0.0025	1.1836	0.0007	-0.3750	0.9130	0.1084	0.9558	0.4834	0.7243
---	---	1635235_at	0.1529	0.4720	0.0819	0.7807	-0.1504	0.3199	-0.1669	0.6990	-0.1510	0.4661	0.0160	0.9494	0.1336	0.8814	-0.0352	0.9496	-0.1688	0.6463
CG15690	CG15690	1635236_at	0.1882	0.2817	0.1041	0.4034	0.2462	0.1283	-0.0240	0.9704	0.0558	0.7793	0.0797	0.6354	-0.1615	0.7307	0.0517	0.8511	0.2132	0.3019
Ing3	Ing3	1635237_at	-0.5217	0.3301	0.0031	0.9998	-0.3233	0.2903	0.2919	0.5756	-0.0312	0.9338	-0.3231	0.1756	0.4823	0.8461	0.3588	0.7737	-0.1235	0.9316
Mbs	MLC-phosphatase	1635238_s_at	-0.8776	0.0206	-0.7298	0.0553	-1.1243	0.0002	-0.3786	0.2654	0.2852	0.1539	0.6638	0.0036	0.0370	0.9848	0.4429	0.2916	0.4059	0.3614
CG13901	CG13901	1635239_at	-0.4234	0.1327	-0.2262	0.2071	-0.0873	0.6214	0.1399	0.7003	0.1976	0.2369	0.0577	0.7511	-0.0275	0.9816	0.3035	0.2518	0.3310	0.2456
Deaf1	deaf-1	1635240_at	-0.1083	0.8514	0.3506	0.2550	0.3940	0.1148	0.1128	0.8518	0.0781	0.7669	-0.0347	0.8954	0.1326	0.9619	0.5997	0.4388	0.4671	0.5702
---	---	1635241_at	0.0637	0.7833	0.1533	0.4072	0.1656	0.2774	-0.0582	0.9387	-0.1002	0.6946	-0.0420	0.8725	-0.1311	0.8097	-0.1450	0.5398	-0.0139	0.9674
---	---	1635242_at	0.1534	0.5571	0.1111	0.4783	0.1100	0.6173	0.0171	0.9873	0.0828	0.7930	0.0657	0.8221	-0.0631	0.9467	-0.0229	0.9620	0.0401	0.9197
CG31371	CG31371	1635243_at	0.2632	0.3200	0.0815	0.5991	0.1062	0.5394	-0.0599	0.9445	0.1158	0.6797	0.1756	0.4490	0.0777	0.8972	0.0684	0.8132	-0.0092	0.9803
---	---	1635244_at	-0.0191	0.9232	-0.0904	0.4411	0.0248	0.9046	-0.0671	0.9329	-0.1036	0.7033	-0.0365	0.8976	-0.1086	0.8609	-0.2160	0.3802	-0.1074	0.7025
CG9196 /// DlitCG9196 /// D	CG9196 /// GA21(	1635245_at	0.3728	0.2140	0.0946	0.5179	0.2835	0.0941	-0.0170	0.9854	-0.0981	0.6790	-0.0811	0.7131	-0.1367	0.8284	-0.0045	0.9941	0.1322	0.6419
Nrk	neurospecific rece	1635246_at	0.1755	0.4109	0.0446	0.6683	-0.0431	0.7832	0.0165	0.9781	-0.0641	0.7189	-0.0805	0.6002	-0.1303	0.7697	-0.2581	0.1606	-0.1278	0.5183
CG6013	CG6013	1635247_at	-0.2252	0.1777	0.1524	0.5233	0.0577	0.7375	0.1536	0.7229	0.0264	0.9228	-0.1272	0.4983	-0.0341	0.9821	0.3333	0.3340	0.3674	0.3089
CG17144	CG17144	1635248_at	-0.4286	0.1599	0.0115	0.9574	-0.4014	0.0651	-0.2897	0.4553	0.2345	0.2669	0.5242	0.0131	0.0741	0.9589	0.5917	0.1432	0.5176	0.2253
---	---	1635249_at	0.1963	0.4119	-0.0959	0.4522	0.0545	0.7535	0.0143	0.9834	0.1488	0.3574	0.1345	0.3561	-0.1233	0.8479	-0.1169	0.6983	0.0064	0.9878
CG31120	CG31120	1635250_at	0.2658	0.4150	0.3725	0.3649	0.5193	0.0141	0.2855	0.3053	0.3605	0.0347	0.0750	0.6585	0.2468	0.8298	0.4817	0.3033	0.2349	0.6507
CG32553	CG32553	1635251_at	0.1619	0.6302	-0.1095	0.5912	0.0183	0.9148	0.2247	0.4586	0.1348	0.4277	-0.0899	0.5837	-0.0064	0.9964	-0.1364	0.7618	-0.1300	0.7656
Actn	Alpha-actinin	1635252_a_at	-1.4543	0.0250	-0.0959	0.5617	-1.8459	0.0007	-0.2091	0.6114	-0.3620	0.0806	-0.1528	0.4299	1.5031	0.3324	1.0654	0.1769	-0.4377	0.6171

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
l(1)G0334	Pyruvate dehydro	1635253_a_at	0.0078	0.9727	0.5799	0.0941	0.6007	0.0061	-0.0575	0.9252	-0.8147	0.0017	-0.7572	0.0014	-0.0511	0.9689	-0.2995	0.3743	-0.2483	0.4851
CG17662	CG17662	1635254_at	0.0124	0.9612	0.0872	0.5068	0.0084	0.9822	-0.0208	0.9761	-0.0710	0.7247	-0.0502	0.7962	0.2014	0.7981	0.0864	0.8471	-0.1150	0.7639
CG9173	CG9173	1635255_at	-0.0056	0.9758	0.1625	0.4530	0.0568	0.7241	-0.1435	0.6321	-0.3009	0.0457	-0.1574	0.2306	0.0547	0.9604	-0.0002	1.0000	-0.0549	0.8982
CG1275	CG1275	1635256_s_at	-0.7313	0.0468	-0.5177	0.1121	0.3595	0.0655	0.2354	0.4643	-0.1924	0.2716	-0.4278	0.0139	-0.7087	0.3390	-0.0484	0.9402	0.6603	0.1271
CG3407	CG3407	1635257_at	-0.1477	0.5475	0.0747	0.5514	-0.3591	0.1696	-0.1482	0.7598	0.2052	0.3316	0.3534	0.0615	0.4293	0.5126	0.4011	0.1874	-0.0281	0.9491
---	---	1635258_s_at	-0.0957	0.8195	-2.2883	0.0386	-2.0673	0.0004	-0.1375	0.8915	1.2099	0.0043	1.3474	0.0016	-0.4180	0.8400	-0.8619	0.3006	-0.4439	0.6311
Mov34	Mov34	1635259_at	0.0207	0.9175	0.3685	0.0770	0.5601	0.0031	0.1438	0.7161	0.0663	0.7583	-0.0775	0.6810	-0.1571	0.8270	0.2988	0.3036	0.4559	0.1534
Stam	Signal transducin	1635260_at	0.3373	0.4985	0.6540	0.2311	0.8007	0.0068	0.2663	0.4511	-0.0498	0.8407	-0.3160	0.0684	0.0686	0.9848	0.1876	0.8680	0.1190	0.9150
CG33262	CG33262	1635261_at	-0.0014	0.9964	0.0802	0.5190	0.0389	0.8703	-0.2464	0.5148	-0.1952	0.3360	0.0511	0.8216	-0.0240	0.9779	-0.0250	0.9404	-0.0010	0.9979
---	---	1635262_at	0.0955	0.6318	0.0247	0.8416	0.3333	0.0584	0.0307	0.9524	-0.0813	0.6034	-0.1120	0.3990	-0.2840	0.6955	-0.1535	0.6314	0.1305	0.6913
CG11825	CG11825	1635263_at	0.0831	0.9005	-1.0271	0.0960	-0.7959	0.0341	0.0306	0.9884	0.7596	0.1282	0.7291	0.1029	-0.1619	0.9457	-0.4663	0.5330	-0.3045	0.7067
lilli	lilliputian	1635264_s_at	0.5414	0.3030	0.8152	0.0688	-0.0743	0.6955	-0.6046	0.0965	-0.2218	0.3021	0.3828	0.0494	0.1378	0.9365	0.1323	0.8624	-0.0055	0.9953
CG31782	CG31782	1635265_a_at	0.0577	0.6886	-0.3653	0.0562	-0.5328	0.1178	-0.2154	0.5710	0.5544	0.0117	0.7698	0.0016	-0.1242	0.9246	-0.0593	0.9327	0.0649	0.9155
GV1	GV1	1635266_at	0.3076	0.3891	-0.2405	0.6244	-0.5369	0.1029	-0.2251	0.6084	0.0968	0.7033	0.3220	0.1017	0.0316	0.9916	-0.4267	0.5607	-0.4583	0.5335
Tsp66E	Tetraspanin 66E	1635267_s_at	0.4249	0.3423	0.6564	0.5006	0.2032	0.6065	-0.6675	0.2602	-0.7219	0.0457	-0.0544	0.8976	-0.3341	0.9057	-0.5004	0.6497	-0.1662	0.9057
CG31806	CG31806	1635268_at	0.0099	0.9545	0.1440	0.4033	0.1836	0.2035	0.0310	0.9627	-0.0045	0.9869	-0.0355	0.8630	0.0171	0.9912	0.1269	0.7247	0.1098	0.7619
CG5435	CG5435	1635269_at	0.0600	0.8382	0.1778	0.2245	0.2723	0.0765	0.0058	0.9937	-0.1393	0.3609	-0.1451	0.2825	-0.0827	0.9277	-0.0504	0.9125	0.0323	0.9383
---	---	1635270_at	0.1152	0.7739	-0.9950	0.0759	-0.1529	0.5239	0.9073	0.1495	1.1689	0.0084	0.2616	0.4690	0.1581	0.8441	-0.1975	0.5699	-0.3556	0.2960
CG30431	CG30431	1635271_at	2.0699	0.0630	1.6220	0.3882	3.0252	0.0002	0.5322	0.4930	0.4604	0.2671	-0.0718	0.8833	-0.7248	0.8981	0.3329	0.9188	1.0576	0.6398
CG10205	CG10205	1635272_a_at	-0.3632	0.0909	0.2922	0.1961	0.3539	0.2165	-0.0806	0.9343	-0.3248	0.2590	-0.2443	0.3536	-0.0461	0.9775	0.2032	0.6269	0.2493	0.5419
kek5	kekkon-like	1635273_s_at	1.5196	0.1851	1.5148	0.0908	0.6264	0.3771	0.1413	0.8164	1.1556	0.0012	1.0142	0.0012	0.9859	0.8235	1.3431	0.4729	0.3571	0.8880
---	---	1635274_at	-0.0710	0.7730	0.0671	0.6702	0.2676	0.4122	0.0628	0.9314	0.0009	0.9977	-0.0618	0.7981	0.0679	0.9606	0.2080	0.6152	0.1401	0.7527
---	---	1635275_at	-0.1081	0.5623	0.1872	0.3365	0.2920	0.1483	0.0023	0.9981	-0.2119	0.3637	-0.2142	0.3014	-0.0606	0.9390	0.1640	0.5167	0.2246	0.3705
CG11437	CG11437	1635276_at	0.1020	0.6075	0.2082	0.1035	0.3158	0.1144	-0.0280	0.9777	-0.0432	0.8997	-0.0152	0.9611	-0.0512	0.9467	-0.0008	0.9995	0.0504	0.8707
---	---	1635277_at	0.5685	0.1642	0.1334	0.5849	0.2266	0.4769	0.4241	0.2129	0.4666	0.0302	0.0425	0.8588	0.3229	0.8270	0.0931	0.9225	-0.2298	0.7451
CG4972	CG4972	1635278_at	0.1066	0.5114	0.1844	0.6311	0.1666	0.2833	-0.1384	0.7647	0.2547	0.1945	0.3930	0.0331	-0.1426	0.9029	0.3447	0.4141	0.4873	0.2676
pont	dpontin	1635279_at	0.4656	0.0666	0.7345	0.1314	0.7896	0.0047	0.1274	0.7647	-0.0167	0.9515	-0.1441	0.3941	0.0492	0.9831	0.3388	0.5431	0.2897	0.6150
CG4685	CG4685	1635280_at	-0.3153	0.1890	-0.1103	0.4126	0.3452	0.0520	0.0376	0.9508	-0.3823	0.0314	-0.4199	0.0134	-0.2485	0.7644	-0.1062	0.8174	0.1423	0.7235
CG8237	CG8237	1635281_at	0.5582	0.0233	0.3110	0.3054	0.7522	0.0012	0.0627	0.9015	0.4757	0.0110	0.4129	0.0127	-0.2100	0.8202	0.3327	0.3760	0.5427	0.1823
CG7142	CG7142	1635282_at	1.4934	0.0267	1.3178	0.2258	1.4669	0.0194	0.7408	0.0500	0.8721	0.0026	0.1313	0.5356	0.8769	0.8141	1.0971	0.4861	0.2202	0.9176
CG31839	CG31839	1635283_at	-0.3377	0.1021	-1.1604	0.0520	-0.3586	0.0377	0.1877	0.5897	0.4370	0.0216	0.2493	0.1130	-0.6266	0.4674	-0.4270	0.3158	0.1997	0.6772
HDAC4	HDAC4	1635284_a_at	-0.2629	0.7684	-0.2512	0.6009	-1.1955	0.0008	-0.2316	0.7604	0.7089	0.0365	0.9405	0.0068	0.8659	0.7644	0.7245	0.5715	-0.1414	0.9342
CG32461	CG32461	1635285_at	0.3214	0.1167	0.1104	0.6581	-0.0469	0.8715	-0.1355	0.7478	0.2316	0.2062	0.3670	0.0328	0.0025	0.9994	-0.1166	0.8231	-0.1191	0.8079
CG31068	CG31068	1635286_at	0.3048	0.1485	-0.1199	0.4655	0.0051	0.9860	0.1375	0.7585	0.4238	0.0345	0.2863	0.0952	0.0237	0.9869	-0.2646	0.4182	-0.2883	0.3898
CG15366	CG15366	1635287_at	-0.0634	0.7739	-0.2467	0.1999	-0.5760	0.0042	-0.2326	0.5128	0.5088	0.0146	0.7414	0.0016	0.0288	0.9779	0.2277	0.3470	0.1989	0.4357
---	---	1635288_at	-0.0764	0.7197	0.0339	0.7445	-0.0670	0.7201	-0.1539	0.7028	-0.1661	0.3815	-0.0122	0.9587	-0.0134	0.9893	0.0970	0.6762	0.1103	0.6239
---	---	1635289_at	0.0748	0.7474	0.3415	0.0625	0.1683	0.4093	-0.2357	0.6013	-0.4553	0.0505	-0.2195	0.2865	0.0094	0.9950	-0.0002	1.0000	-0.0096	0.9861
CG11418	CG11418	1635290_at	0.1226	0.6156	-0.1287	0.8263	0.0473	0.8166	0.1021	0.8609	0.4589	0.0351	0.3567	0.0601	-0.1526	0.9296	0.2888	0.6310	0.4414	0.4399
Zn72D	Zn72D	1635291_a_at	-0.1197	0.7975	-0.0305	0.9629	-0.3428	0.0681	-0.1793	0.6082	0.2045	0.2448	0.3838	0.0227	0.0808	0.9816	0.3766	0.6614	0.2959	0.7439
CG34396	CG15654	1635292_at	-1.7181	0.0018	-1.1686	0.0635	-1.7927	0.0000	-0.2441	0.6327	-0.4607	0.0688	-0.2166	0.3457	0.1970	0.9088	0.2520	0.7175	0.0550	0.9487
CG8336	CG8336	1635293_s_at	-0.2293	0.2662	0.9811	0.0402	1.1451	0.0018	0.1117	0.8589	-0.6686	0.0100	-0.7803	0.0030	-0.0679	0.9717	0.6138	0.1822	0.6817	0.1752
CG9312	CG9312	1635294_at	0.2831	0.6365	0.1080	0.5607	-0.0399	0.9190	-0.2549	0.7599	-0.3077	0.4034	-0.0527	0.9027	0.0111	0.9966	-0.3293	0.6512	-0.3405	0.6377
Imd	myoblasts incomp	1635295_a_at	-0.2801	0.2738	-0.7049	0.0918	-0.3919	0.3432	0.6376	0.3193	0.8942	0.0247	0.2565	0.4694	0.3112	0.7142	0.3333	0.3340	0.0221	0.9666
---	---	1635296_at	-0.0571	0.7260	0.0094	0.9382	0.0094	0.9640	-0.0249	0.9705	-0.0819	0.6768	-0.0569	0.7630	-0.0421	0.9589	-0.0519	0.8764	-0.0099	0.9793
---	---	1635297_at	0.2767	0.1225	-0.1198	0.3937	-0.0552	0.8392	-0.1947	0.6954	0.0720	0.7980	0.2667	0.1927	-0.1819	0.8446	-0.3125	0.4093	-0.1306	0.7706
CG13458	CG13458	1635298_at	-0.0052	0.9768	-0.0562	0.7090	-0.2044	0.3274	-0.0331	0.9568	0.2218	0.1752	0.2549	0.0840	-0.0009	0.9997	-0.0374	0.9128	-0.0365	0.9056
Rev1	Rev1	1635299_at	-0.0001	0.9998	-0.0018	0.9962	-0.4571	0.0695	-0.0833	0.8894	0.4295	0.0386	0.5128	0.0117	0.3299	0.7152	0.4569	0.2116	0.1270	0.7754
CG1268	CG1268	1635300_at	-0.8519	0.0056	-0.0849	0.4934	-0.3461	0.2457	0.0193	0.9909	-0.5240	0.1547	-0.5433	0.1000	-0.0896	0.9298	0.0344	0.9492	0.1240	0.7442
CG32095	CG32095	1635301_at	0.0330	0.8889	-0.3718	0.0958	-0.3811	0.0570	0.0374	0.9491	0.4549	0.0132	0.4174	0.0119	0.0630	0.9495	0.0832	0.8334	0.0202	0.9630
mus101	mutagen-sensitive	1635302_at	-0.0309	0.8968	0.0595	0.9158	0.0882	0.7739	-0.1370	0.7803	0.0574	0.8219	0.1945	0.2931	-0.2068	0.8854	0.0837	0.9213	0.2905	0.6188
RpS13	Minute(2)32A	1635303_at	0.3097	0.1605	1.5372	0.0066	1.6228	0.0001	-0.0371	0.9753	-1.6292	0.0007	-1.5921	0.0004	0.0297	0.9816	-0.0275	0.9505	-0.0573	0.8779
CG13954	CG13954	1635304_at	0.1885	0.4795	0.0941	0.5059	0.1320	0.4104	0.0862	0.9012	-0.0370	0.9054	-0.1232	0.5843	0.1317	0.8270	-0.1082	0.7053	-0.2399	0.3569
Trl	GAGA factor	1635305_s_at	0.2236	0.3642	0.5760	0.2554	-0.0702	0.6648	-0.4937	0.1658	-0.0459	0.8726	0.4478	0.0282	0.2237	0.8534	0.3887	0.4351	0.1650	0.7790
CG4650	CG4650	1635306_at	-0.0441	0.8185	0.0320	0.7525	0.2464	0.1303	0.0313	0.9688	0.0225	0.9407	-0.0089	0.9727	-0.2012	0.6898	0.0580	0.8331	0.2592	0.2237
CG4662	CG4662	1635307_at	0.7790	0.1826	0.5571	0.2882	0.3385	0.0619	0.2324	0.8671	0.6535	0.1807	0.4211	0.3501	0.3546	0.7485	0.3399	0.4804	-0.0147	0.9849
CG5768	CG5768	1635308_a_at	-0.0962	0.4964	-0.1269	0.4523	-0.1191	0.4509	0.1080	0.8100	0.2016	0.2606	0.0935	0.6025	0.0653	0.9296	0.0384	0.9170	-0.0268	0.9360
---	---	1635309_at	0.1584	0.5605	-0.0532	0.7127	-0.3550	0.2297	-0.3238	0.6202	0.3552	0.2780	0.6790	0.0275	-0.1115	0.8395	0.0969	0.7056	0.2084	0.3711
CG31688	CG31688	1635310_s_at	-0.0266	0.9742	1.1887	0.4931	-0.2159	0.7688	-0.2817	0.6818	-0.6057	0.0619	-0.3240	0.2601	0.8698	0.8692	0.5889	0.8343	-0.2809	0.9231
---	---	1635311_at	-0.0702	0.6972	0.0408	0.7099	-0.0319	0.9057	-0.0152	0.9826	0.0400	0.8506	0.0552	0.7572	0.0704	0.9171	0.1568	0.5181	0.0864	0.7492
---	---	1635312_at	0.0183	0.9474	-0.0257	0.8437	0.0486	0.8146	0.1022	0.8776	0.1408	0.5666	0.0386	0.8864	-0.1422	0.8270	-0.0702	0.8500	0.0720	0.8345
CG10405	CG10405	1635313_at	0.1532	0.4601	0.3822	0.0982	0.6081	0.0126	0.1226	0.7749	-0.0432	0.8529	-0.1658	0.3094	0.0687	0.9098	0.0690	0.8028	0.0004	0.9992
maf-S	maf-S	1635314_at	-0.1790	0.3036	0.1611	0.5525	0.0534	0.8308	0.1736	0.7045	-0.4610	0.0354	-0.6345	0.0055	0.2879	0.7506	-0.1211	0.8097	-0.4091	0.3041
---	---	1635315_at	0.1476	0.4054	0.0776	0.4962	0.0983	0.6298	0.1260	0.8076	0.2300	0.2670	0.1040	0.6196	0.1416	0.8181	-0.0354	0.9291	-0.1770	0.4978
---	---	1635316_at	0.0380	0.8176	0.0572	0.6695	0.0839	0.6395	0.0591	0.9218	-0.0290	0.9083	-0.0880	0.6348	0.0747	0.9225	0.0952	0.7525	0.0205	0.9548
Art9	Arginine methyltra	1635317_at	0.3429	0.2249	0.0946	0.3811	0.2325	0.1077	0.1014	0.8734	0.0324	0.9167	-0.0690	0.7789	-0.1334	0.8395	-0.1890	0.4969	-0.0556	0.8800
Odc1	ornithine decarbo	1635318_at	2.4143	0.0008	2.2170	0.0151	3.8547	0.0000	1.0952	0.0741	-0.1572	0.7008	-1.2524	0.0027	-0.7259	0.6955	-0.4507	0.5880	0.2752	0.7625
CG18476	CG18476	1635319_at	-0.0057	0.9892	-0.1010	0.3740	-0.0043	0.9852	0.0510	0.9466	0.0772	0.7646	0.0262	0.9210	0.0377	0.9826	0.0509	0.9363	0.0131	0.9846
---	---	1635320_s_at	0.1981	0.2648	-0.0146	0.9581	0.2193	0.2039	0.0570	0.9345	0.0717	0.7704	0.0147	0.9550	-0.0060	0.9964	-0.0210	0.9655	-0.0151	0.9745
CG14688	CG14688	1635321_at	0.1991	0.6725	0.2335	0.1053	0.1639	0.4175	0.3310	0.5785	1.1864	0.0030	0.8554	0.0077	0.4137	0.7485	1.2202	0.0476	0.8065	0.1651

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13322	CG13322	1635322_a_at	0.4028	0.1667	0.4425	0.2320	0.0902	0.6878	-0.0273	0.9724	0.2122	0.2838	0.2396	0.1723	0.2088	0.8745	0.3391	0.5300	0.1303	0.8453
---	---	1635323_at	0.0261	0.8962	0.0242	0.8107	0.0277	0.8663	0.0144	0.9854	-0.0141	0.9572	-0.0285	0.8934	-0.0218	0.9862	-0.0192	0.9687	0.0026	0.9963
---	---	1635324_at	0.2295	0.3264	-0.0758	0.5873	0.2530	0.1652	0.2254	0.4317	0.3330	0.0407	0.1076	0.4719	0.1539	0.8191	0.1074	0.7381	-0.0465	0.9025
Gap1	sextra	1635325_a_at	-0.5902	0.0665	-0.2817	0.1684	-0.3123	0.2374	0.2033	0.6015	0.2156	0.2746	0.0123	0.9599	0.1243	0.9305	0.4488	0.3162	0.3245	0.4965
---	---	1635326_at	0.2694	0.1470	0.0969	0.4922	-0.0340	0.8952	-0.2232	0.5777	0.0233	0.9366	0.2464	0.1786	-0.0260	0.9829	-0.1066	0.7439	-0.0806	0.8155
CG31206 /// DereCG31206	CG31206	1635327_at	0.1926	0.2030	-0.2081	0.3377	0.0199	0.9344	0.1444	0.7205	0.3429	0.0638	0.1985	0.2248	-0.2252	0.7826	-0.2342	0.5234	-0.0089	0.9874
Oamb	octopamine recep	1635328_at	-2.2645	0.0022	-1.2940	0.0089	-1.6880	0.0026	-0.6637	0.1787	-1.3120	0.0018	-0.6483	0.0236	-0.1603	0.9396	-0.2858	0.6952	-0.1255	0.8880
CG7638	CG7638	1635329_at	-0.1749	0.5121	-0.0154	0.9254	0.1086	0.7027	-0.3237	0.2501	-0.1369	0.4285	0.1868	0.2112	-0.3063	0.7305	0.0373	0.9509	0.3436	0.3800
CG5428	CG5428	1635330_at	1.2921	0.0059	0.4220	0.3991	1.4104	0.0007	0.0870	0.9068	-0.3413	0.1522	-0.4283	0.0503	-0.8629	0.6483	-1.1450	0.1345	-0.2821	0.7492
Cpr49Af	CG8510	1635331_at	-0.2135	0.3955	0.0771	0.6528	0.1310	0.7245	-0.0029	0.9962	-0.4040	0.0515	-0.4010	0.0342	0.0649	0.9657	-0.0245	0.9714	-0.0894	0.8694
---	---	1635332_at	0.1158	0.5048	0.0548	0.6909	-0.3301	0.1436	-0.0781	0.8902	0.0410	0.8703	0.1191	0.5166	0.1208	0.8553	-0.0656	0.8625	-0.1864	0.5095
---	---	1635333_at	0.1765	0.3630	0.0722	0.4633	0.3032	0.0768	0.1627	0.6591	-0.0041	0.9881	-0.1668	0.2990	-0.1035	0.8424	-0.0675	0.8053	0.0359	0.9028
CG10732 /// DmirCG10732	CG10732	1635334_a_at	-2.5792	0.0005	-3.0449	0.0045	-3.1927	0.0000	0.1179	0.8967	0.2125	0.5005	0.0946	0.7752	0.3126	0.7576	-0.0515	0.9404	-0.3641	0.4016
CG12018	CG12018	1635335_at	0.0478	0.9043	0.0602	0.5723	0.1721	0.5192	-0.0845	0.8436	0.1661	0.2984	0.2506	0.0781	-0.1559	0.8882	0.2043	0.6534	0.3602	0.3967
comm2	comm2	1635336_at	-0.5651	0.2140	-0.3795	0.2691	-1.1968	0.0048	-0.3598	0.7168	-1.6799	0.0036	-1.3201	0.0063	0.3014	0.8202	-1.6694	0.0222	-1.9708	0.0236
---	---	1635337_at	-0.0544	0.7951	0.0851	0.6240	0.0370	0.8709	-0.0127	0.9883	0.0003	0.9992	0.0130	0.9597	-0.0144	0.9893	0.1425	0.5411	0.1569	0.4978
retinophilin /// uta	undertaker	1635338_a_at	-0.1491	0.4363	0.0817	0.4607	0.1356	0.5828	0.0181	0.9883	-0.0635	0.8699	-0.0815	0.8041	-0.0483	0.9467	0.1162	0.6270	0.1645	0.4740
CG15725	CG15725	1635339_at	-0.0148	0.9603	0.0627	0.5978	0.0795	0.6768	-0.0700	0.9201	-0.2111	0.3282	-0.1411	0.4883	0.0223	0.9860	0.0719	0.8546	0.0496	0.8993
tud	tudor	1635340_at	-0.0901	0.6770	-0.5409	0.2035	-1.2246	0.0011	-0.2891	0.3553	0.4384	0.0225	0.7275	0.0015	0.3829	0.7475	-0.0337	0.9678	-0.4166	0.4163
CG31220	CG31220	1635341_at	0.2201	0.2238	0.2025	0.4350	-0.0159	0.9324	-0.1333	0.8000	-0.0103	0.9750	0.1229	0.5560	0.1580	0.7697	0.1041	0.6844	-0.0539	0.8575
CG10306 /// DyakCG10306	CG10306	1635342_at	0.6183	0.0067	0.4783	0.0617	0.4228	0.0194	0.2244	0.4191	0.1203	0.4526	-0.1041	0.4777	0.3387	0.5765	0.0589	0.8821	-0.2798	0.3205
CkIIalpha-i3	CKII-alpha subuni	1635343_a_at	-0.4159	0.2916	-1.3505	0.0189	-1.0208	0.0008	0.1494	0.7735	1.0598	0.0011	0.9105	0.0013	-0.1530	0.9216	0.2463	0.6673	0.3993	0.4569
Ppox	protoporphyrinoge	1635344_at	-0.5182	0.0360	0.0047	0.9817	0.0942	0.5411	-0.0076	0.9931	-0.2825	0.1028	-0.2749	0.0777	-0.1910	0.7783	0.2013	0.5090	0.3922	0.2056
---	---	1635345_at	0.0895	0.6491	-0.1040	0.5362	0.2260	0.1274	0.1868	0.6565	0.1358	0.5283	-0.0510	0.8244	-0.1627	0.7783	-0.0479	0.8997	0.1148	0.6776
CG15160	CG15160	1635346_at	-0.1027	0.8262	-0.4056	0.1961	-0.0935	0.6326	0.2732	0.4998	0.4314	0.0490	0.1582	0.4312	-0.1061	0.9400	-0.0357	0.9605	0.0704	0.9065
CG8557 /// DmirCG8557 ///	CG8557	1635347_a_at	-0.5884	0.3646	-1.2826	0.1141	-1.5581	0.0001	-0.3433	0.3381	0.6691	0.0067	1.0125	0.0007	-0.0473	0.9916	-0.0390	0.9846	0.0083	0.9965
CG8086	CG8086	1635348_at	-0.1958	0.7824	0.5326	0.6423	0.4880	0.0255	0.0476	0.9774	-0.6761	0.1005	-0.7236	0.0537	0.1293	0.9761	0.3936	0.7481	0.2643	0.8422
nAcRbeta-96A	Dbeta2	1635349_a_at	0.2633	0.1132	0.1978	0.4148	-0.0757	0.6120	-0.0191	0.9790	0.2150	0.2330	0.2341	0.1454	-0.0282	0.9742	-0.0032	0.9943	0.0250	0.9349
---	---	1635350_at	0.0325	0.8636	0.1683	0.3445	0.2754	0.0612	0.0670	0.9098	-0.1952	0.3032	-0.2622	0.1172	0.0301	0.9665	0.0334	0.9085	0.0033	0.9924
---	---	1635351_at	0.1564	0.3308	0.1181	0.2464	0.0052	0.9791	-0.0377	0.9538	0.1564	0.3925	0.1942	0.2242	-0.0529	0.9514	0.0209	0.9609	0.0738	0.8171
Vha44	vacuolar ATPase	1635352_s_at	-1.1710	0.0186	-1.0006	0.0908	-1.7225	0.0001	-0.4276	0.1153	-0.6187	0.0033	-0.1912	0.1889	0.2591	0.8706	-0.3987	0.5469	-0.6578	0.3089
---	---	1635353_at	-0.0837	0.7445	0.1115	0.5977	0.1530	0.4273	0.1043	0.8551	0.1093	0.6363	0.0050	0.9850	-0.1703	0.7644	0.0405	0.9141	0.2108	0.3921
Mst36Fb	CG31791	1635354_at	0.0135	0.9608	0.0461	0.6866	0.0693	0.7517	-0.1380	0.7927	-0.1554	0.4873	-0.0174	0.9488	0.0261	0.9831	0.0294	0.9499	0.0033	0.9948
Xbp1	X box binding prot	1635355_a_at	-0.3274	0.4762	0.6193	0.1818	0.6160	0.0123	-0.0484	0.9470	-0.5881	0.0115	-0.5397	0.0103	0.0061	0.9990	0.3580	0.6746	0.3519	0.6761
CG8816	CG8816	1635356_at	0.1593	0.6712	-0.0752	0.8215	-0.3029	0.1015	0.1114	0.9057	0.5926	0.0557	0.4811	0.0780	0.3211	0.7506	0.3141	0.4753	-0.0070	0.9924
CG11146	CG11146	1635357_at	0.0663	0.8090	-0.0611	0.6214	0.0658	0.6698	0.1022	0.8842	0.0973	0.7279	-0.0049	0.9873	0.0223	0.9816	0.0000	1.0000	-0.0223	0.9431
CG2116	CG2116	1635358_at	-0.0044	0.9886	0.4557	0.0907	0.5831	0.0032	0.1633	0.7182	-0.1470	0.4954	-0.3103	0.0909	-0.0155	0.9898	0.3836	0.1418	0.3991	0.1577
---	---	1635359_at	-0.0320	0.8774	0.1666	0.3652	0.2408	0.2366	0.0290	0.9716	-0.0964	0.6827	-0.1255	0.5344	0.0737	0.9296	0.1468	0.6093	0.0731	0.8287
CG10830	CG10830	1635360_at	-2.3501	0.0018	-0.3036	0.5883	-1.8221	0.0074	-1.1445	0.1968	-2.0826	0.0030	-0.9381	0.0548	0.2017	0.8870	-0.1465	0.8376	-0.3483	0.5346
CG18806	CG18806	1635361_at	0.7516	0.0088	0.4985	0.0676	1.0587	0.0031	-0.0858	0.8578	-0.1178	0.5201	-0.0320	0.8757	-0.2628	0.8461	-0.2378	0.7080	0.0250	0.9777
CG13654	CG13654	1635362_at	0.1379	0.6853	-0.1251	0.7863	0.8591	0.0296	0.2964	0.6876	-0.2842	0.4222	-0.5806	0.0616	-0.5930	0.6955	-0.6060	0.3311	-0.0130	0.9904
Jupiter	Jupiter	1635363_a_at	-1.6348	0.0113	-3.1165	0.0047	-3.3479	0.0000	-0.1552	0.7104	0.8843	0.0014	1.0395	0.0004	0.1362	0.9571	-0.6181	0.3820	-0.7543	0.3070
CG30343	CG30343	1635364_at	-0.3158	0.5693	-0.9360	0.0417	-0.6334	0.1621	0.0072	0.9956	0.4047	0.2975	0.3975	0.2502	-0.3061	0.8768	-0.2014	0.8478	0.1047	0.9211
CG5198	CG5198	1635365_at	-0.1739	0.5983	0.0137	0.9010	0.1831	0.2125	-0.0225	0.9803	0.0748	0.7805	0.0973	0.6744	-0.2844	0.6824	0.3520	0.2043	0.6364	0.0625
CG5614	CG5614	1635366_at	0.1244	0.3865	0.1240	0.3910	0.0303	0.8931	-0.1181	0.7839	-0.0452	0.8442	0.0729	0.6962	0.1821	0.7739	0.0554	0.8903	-0.1267	0.6731
fs(1)h	rancor	1635367_at	0.0642	0.7114	-0.0450	0.8507	-0.3342	0.2053	-0.1192	0.8578	0.4365	0.0710	0.5557	0.0175	0.0482	0.9756	0.3059	0.4246	0.2578	0.5225
CG11398	CG11398	1635368_at	-0.1761	0.6748	0.0193	0.8936	-0.3145	0.2702	-0.1787	0.8994	0.3139	0.5224	0.4926	0.2331	0.1224	0.9405	0.4386	0.3743	0.3162	0.5520
---	---	1635369_s_at	0.1750	0.3224	-0.0002	1.0000	-0.0752	0.6557	-0.1217	0.7795	0.1186	0.5409	0.2403	0.1353	-0.0448	0.9549	-0.1113	0.6620	-0.0664	0.8178
Tsp42El	tetraspanin 42E	1635370_at	-0.4913	0.0659	0.7785	0.0435	0.5135	0.1345	-0.1876	0.7870	-0.9961	0.0048	-0.8085	0.0074	0.1381	0.9291	0.3665	0.4711	0.2285	0.6779
CG31323	CG31323	1635371_at	-3.1872	0.0028	-2.3921	0.0103	-2.6172	0.0000	-0.0784	0.9012	-0.5571	0.0145	-0.4787	0.0171	0.1506	0.9618	0.0277	0.9877	-0.1229	0.9194
Zeelin1	Zeelin1	1635372_a_at	-1.3010	0.0354	-1.8248	0.0236	-2.5689	0.0001	-0.1759	0.5357	-0.1088	0.4829	0.0671	0.6621	0.6265	0.7822	-0.6775	0.5009	-1.3039	0.2051
CG14045	CG14045	1635373_at	0.1675	0.3796	0.2171	0.0786	0.5448	0.0242	0.1908	0.7469	-0.0150	0.9702	-0.2058	0.3916	0.0315	0.9672	-0.0450	0.8765	-0.0765	0.7437
CG31728	CG31728	1635374_at	-0.4660	0.1578	-0.0703	0.4717	-0.0550	0.8513	-0.0467	0.9466	-0.5359	0.0144	-0.4892	0.0132	0.0399	0.9816	-0.0540	0.9291	-0.0939	0.8501
CG5630	CG5630	1635375_at	0.3187	0.2143	0.5123	0.1916	0.7397	0.0332	-0.1444	0.8424	0.1255	0.6834	0.2698	0.2671	-0.3232	0.8270	0.3363	0.6080	0.6594	0.2960
CG9086 /// DmirCG9086	CG9086	1635376_at	0.5129	0.2735	0.6643	0.0654	0.4841	0.1159	-0.2338	0.7613	-0.0233	0.9632	0.2105	0.5103	0.1273	0.9449	0.2574	0.6772	0.1301	0.8578
CG13344	CG13344	1635377_at	-0.1520	0.4716	0.2262	0.4953	0.2845	0.1302	-0.1565	0.8350	-0.1058	0.7550	0.0507	0.8803	-0.1423	0.9101	0.4455	0.3094	0.5878	0.2110
CG7177 /// DmirCG7177	CG7177	1635378_at	0.0308	0.9458	-0.1240	0.7461	-0.4099	0.0823	0.0173	0.9884	0.7170	0.0193	0.6998	0.0134	0.2223	0.8802	0.4352	0.4397	0.2128	0.7439
CG31300	CG31300	1635379_at	-1.2974	0.0053	0.0607	0.9034	-1.0027	0.0121	-0.3323	0.6749	-1.2984	0.0048	-0.9660	0.0110	0.8811	0.4128	0.0503	0.9587	-0.8308	0.1752
CG30413	CG30413	1635380_at	0.0170	0.9562	0.4642	0.0908	0.0972	0.5809	-0.1866	0.5628	-0.1865	0.2610	0.0001	0.9995	0.1184	0.8731	0.2454	0.3945	0.1270	0.6965
CRMP	dihydropyrimidine	1635381_a_at	0.1345	0.5467	0.2002	0.2881	0.3775	0.0493	0.0550	0.9533	-0.2390	0.3644	-0.2939	0.2019	-0.0357	0.9816	0.1292	0.7506	0.1650	0.6578
dlg1	Discs large	1635382_at	-0.6680	0.0413	-0.6858	0.3130	-1.0910	0.0041	0.0788	0.9387	0.4033	0.1768	0.3245	0.2271	0.3137	0.8215	0.3769	0.5283	0.0631	0.9373
vnd	ventral nerve cord	1635383_at	0.0692	0.7283	-0.1962	0.4044	0.0764	0.7379	0.1719	0.7321	0.1817	0.4336	0.0099	0.9716	-0.1079	0.8331	-0.1048	0.6533	0.0031	0.9935
ZIP1	Zinc/iron regulate	1635384_at	0.1238	0.6148	-0.0228	0.8819	-0.1346	0.5123	0.0580	0.9260	0.1857	0.3300	0.1277	0.4754	0.0625	0.9400	-0.0851	0.7897	-0.1476	0.5868
CG10225	CG10225	1635385_at	-0.0639	0.7531	-0.0617	0.6980	-0.0150	0.9522	0.0129	0.9838	0.0272	0.8918	0.0143	0.9378	-0.0874	0.9342	0.0219	0.9689	0.1092	0.7886
---	---	1635386_s_at	0.1003	0.5179	0.1408	0.5012	0.0664	0.7442	-0.0741	0.9117	-0.1572	0.4792	-0.0831	0.7129	0.0933	0.8609	0.0436	0.8893	-0.0497	0.8602
CG14913	CG14913	1635387_at	0.1221	0.3995	0.1202	0.4097	0.1553	0.3095	0.0779	0.8473	0.1487	0.3227	0.0707	0.6425	0.0046	0.9952	0.0044	0.9924	-0.0003	0.9992
RpS9	Ribosomal protein	1635388_s_at	0.2115	0.1648	0.2618	0.0857	0.3250	0.0331	0.0560	0.9008	-0.0101	0.9635	-0.0662	0.6605	0.0139	0.9867	0.0236	0.9400	0.0097	0.9745
Idgf4	Imaginal disc grov	1635389_s_at	-1.0685	0.0020	-0.8407	0.0595	-0.6464	0.0021	-0.0543	0.9036	-0.4269	0.0103	-0.3726	0.0116	-0.1565	0.8541	-0.2140	0.5601	-0.0575	0.9057
CG9331	CG9331	1635390_s_at	1.1621	0.0029	1.0946	0.0996	1.1158	0.0071	-0.1303	0.8897	-0.6001	0.0605	-0.4698	0.0947	-0.1342	0.9469	-0.6662	0.2634	-0.5320	0.3972

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1635391_at	-0.1401	0.3725	0.0637	0.5757	-0.0129	0.9474	0.0112	0.9866	-0.1313	0.4366	-0.1425	0.3392	0.1128	0.8521	0.0505	0.8898	-0.0623	0.8461
---	---	1635392_at	0.1188	0.5803	-0.0327	0.7946	0.0360	0.8361	0.1793	0.5008	0.2052	0.1429	0.0260	0.8762	-0.0037	0.9980	-0.0869	0.8450	-0.0831	0.8431
Eip75B	Ecdysone-induce	1635393_s_at	-0.7646	0.3122	0.1357	0.8776	0.6237	0.0291	-0.1771	0.7010	-0.7306	0.0050	-0.5535	0.0105	-0.3086	0.9407	0.2210	0.9110	0.5296	0.7149
CG31301	CG31301	1635394_at	-0.0783	0.7323	-0.3743	0.2555	-0.3292	0.0493	-0.1504	0.6908	0.0758	0.7079	0.2262	0.1478	-0.2381	0.7893	-0.2077	0.6093	0.0304	0.9549
---	---	1635395_at	-0.0343	0.8633	0.0941	0.5661	0.0200	0.9408	0.0671	0.8967	-0.0153	0.9518	-0.0824	0.6367	0.1078	0.8973	0.1160	0.7544	0.0082	0.9868
CG30203	CG30203	1635396_at	0.1384	0.5816	-0.0037	0.9772	0.0466	0.8024	0.0596	0.9311	0.2182	0.2846	0.1586	0.4001	-0.1074	0.9238	-0.0135	0.9853	0.0938	0.8453
mthl12	Mth-like 12	1635397_at	0.0641	0.7401	-0.0426	0.7722	-0.1070	0.6122	-0.1218	0.8686	0.1087	0.7186	0.2305	0.3310	-0.0022	0.9990	-0.0672	0.8467	-0.0650	0.8418
Scp2	Calexcitin	1635398_at	0.1316	0.8741	0.0000	1.0000	0.1556	0.4316	-0.0745	0.9749	-0.4766	0.4285	-0.4021	0.4660	-0.2536	0.8494	-0.5100	0.3458	-0.2564	0.6729
CG11006	CG11006	1635399_s_at	-0.3767	0.2867	0.5683	0.3996	0.5845	0.0221	-0.0215	0.9833	-0.5401	0.0287	-0.5186	0.0214	0.0689	0.9816	0.3150	0.6932	0.2460	0.7662
dtr	defective transmit	1635400_at	0.0009	0.9965	-0.0391	0.6937	0.1128	0.6237	0.2698	0.4592	0.2989	0.1289	0.0291	0.9027	-0.0196	0.9831	0.0376	0.9093	0.0572	0.8345
---	---	1635401_at	0.1679	0.3243	-0.0206	0.8694	-0.0061	0.9836	0.2348	0.5492	0.2210	0.2784	-0.0138	0.9573	0.0155	0.9875	-0.0550	0.8595	-0.0705	0.7936
---	---	1635402_at	0.1169	0.4520	0.4351	0.1855	0.1903	0.3899	-0.2083	0.6382	-0.1645	0.4682	0.0438	0.8624	0.0376	0.9717	0.0597	0.8749	0.0221	0.9525
shf	shifted	1635403_at	0.6569	0.0553	-2.2251	0.0324	-1.1978	0.0029	0.8671	0.1505	1.7775	0.0011	0.9104	0.0130	-0.1961	0.9357	-1.1145	0.1418	-0.9184	0.2527
---	---	1635404_at	0.1823	0.4781	-0.0909	0.6249	-0.2574	0.1379	0.0037	0.9956	0.0694	0.7027	0.0657	0.6914	0.1484	0.9112	-0.0057	0.9959	-0.1541	0.7831
---	---	1635405_at	0.1660	0.3132	-0.0727	0.7767	-0.2860	0.0651	0.0644	0.9116	0.3750	0.0463	0.3106	0.0617	0.1832	0.7726	0.0332	0.9404	-0.1500	0.6129
Pkg21D	cGMP-dependent	1635406_at	0.7383	0.4494	-0.5174	0.0346	0.0013	0.9958	0.5127	0.1526	0.4120	0.0612	-0.1007	0.6529	0.1748	0.9657	-0.7238	0.5209	-0.8986	0.4167
CG32772	CG32772	1635407_at	-0.4619	0.0442	0.7615	0.1813	0.7700	0.0106	-0.1818	0.7140	-1.0189	0.0016	-0.8371	0.0022	-0.1465	0.9101	0.2639	0.5905	0.4105	0.3851
CG13700	CG13700	1635408_at	0.0521	0.8446	-0.0723	0.7708	0.0281	0.8910	0.1087	0.8605	0.0062	0.9851	-0.1025	0.6556	0.0143	0.9898	-0.0221	0.9561	-0.0364	0.9155
RnrS	ribonucleoside-dip	1635409_at	1.1080	0.0139	0.1543	0.6226	0.3736	0.1423	-0.1635	0.7807	0.3678	0.1299	0.5313	0.0230	-0.2926	0.8235	-0.4594	0.3922	-0.1668	0.8007
CG12766	CG12766	1635410_at	-0.0091	0.9889	-0.4413	0.3801	-0.1440	0.4470	-0.2293	0.6120	-0.1766	0.4539	0.0527	0.8404	-0.5877	0.6955	-0.6850	0.2641	-0.0972	0.9118
CG32652	CG32652	1635411_at	0.1342	0.3964	0.2029	0.5132	-0.1940	0.5461	0.0527	0.9599	0.0519	0.8892	-0.0008	0.9981	0.1667	0.8719	-0.0040	0.9972	-0.1707	0.7128
CG7422	CG7422	1635412_at	0.6697	0.5246	-1.0071	0.0223	-0.1867	0.3127	0.8785	0.0904	0.7610	0.0206	-0.1175	0.7165	-0.0732	0.9898	-1.0340	0.4034	-0.9609	0.4569
CG32266	CG32266	1635413_at	0.3311	0.1299	0.2518	0.3634	-0.0245	0.8978	0.0500	0.9470	-0.1091	0.6476	-0.1592	0.4233	0.0749	0.8953	-0.1307	0.5552	-0.2056	0.3465
---	---	1635414_at	0.0524	0.8387	0.0934	0.6651	-0.1280	0.5749	-0.1759	0.7596	0.1451	0.5890	0.3211	0.1448	0.0602	0.9457	0.0790	0.8216	0.0188	0.9621
---	---	1635415_at	0.1912	0.3686	-0.0004	1.0000	0.1347	0.4872	0.0904	0.9247	0.0879	0.8026	-0.0025	0.9945	-0.0059	0.9959	-0.1635	0.5538	-0.1576	0.5751
CG31100	CG31100	1635416_at	0.6206	0.1357	0.6063	0.1853	0.2399	0.6019	0.4207	0.5859	1.1091	0.0123	0.6884	0.0547	0.8646	0.6483	1.1757	0.1269	0.3111	0.7199
CG9025 /// DmirCG9025 ///	CG9025 /// GA21	1635417_a_at	0.1078	0.6618	-0.0598	0.7090	0.0626	0.7445	0.2150	0.7025	0.0455	0.8979	-0.1695	0.4949	-0.0122	0.9893	-0.0254	0.9337	-0.0132	0.9622
CG18558	CG18558	1635418_at	-0.0295	0.9109	0.1746	0.3098	0.0424	0.8224	-0.3048	0.4012	-0.2907	0.1504	0.0141	0.9568	0.1028	0.8270	0.0991	0.6425	-0.0037	0.9915
Gycbeta100B	Guanylyl cyclase	1635419_at	0.1196	0.6080	0.0632	0.5372	0.2659	0.3299	0.1082	0.8698	0.1863	0.4374	0.0782	0.7571	0.1517	0.8465	0.1949	0.5673	0.0433	0.9231
CG14760	CG14760	1635420_at	0.2604	0.2845	0.1480	0.3538	0.3146	0.0743	0.0253	0.9777	0.0326	0.9200	0.0073	0.9799	0.0037	0.9963	-0.0131	0.9628	-0.0168	0.9435
Syx16	syntaxin	1635421_at	-0.8446	0.0538	-0.2521	0.6698	-0.0064	0.9805	0.2580	0.4501	-0.7548	0.0024	-1.0127	0.0004	-0.0096	0.9976	-0.0432	0.9698	-0.0337	0.9757
CG9147	CG9147	1635422_s_at	-0.4482	0.1867	-0.0891	0.7973	0.0408	0.8500	0.0176	0.9857	-0.5721	0.0188	-0.5897	0.0102	-0.0588	0.9816	0.0271	0.9782	0.0860	0.9095
Hsc70-3	immunoglobulin bi	1635423_s_at	0.5074	0.0370	0.4233	0.0361	0.5638	0.0039	0.1158	0.7608	0.4284	0.0171	0.3126	0.0393	-0.0161	0.9914	0.5312	0.0959	0.5473	0.1146
CG32063	CG32063	1635424_s_at	-0.0193	0.9326	-0.1121	0.4458	-0.1980	0.3438	-0.0519	0.9518	0.0902	0.7522	0.1421	0.5459	-0.0132	0.9901	0.0052	0.9925	0.0183	0.9553
---	---	1635425_s_at	0.1854	0.3343	0.0142	0.8936	-0.1124	0.5192	0.0644	0.9153	0.1737	0.3695	0.1093	0.5592	0.0732	0.9298	-0.0122	0.9806	-0.0854	0.7918
CG11291	CG11291	1635426_at	0.0783	0.6751	0.1594	0.2343	0.0836	0.6735	-0.1065	0.7479	-0.1030	0.4984	0.0035	0.9850	-0.0217	0.9816	0.1221	0.5824	0.1438	0.5092
gukh	GUKholder	1635427_a_at	0.9829	0.2663	1.4608	0.2283	1.2772	0.0019	-0.2205	0.4568	0.0711	0.7011	0.2916	0.0476	-0.0283	0.9964	0.4816	0.8119	0.5099	0.7827
CG12863	CG12863	1635428_at	-0.3225	0.1957	-0.3628	0.1493	-0.6731	0.0056	-0.3760	0.1949	-0.1787	0.3124	0.1973	0.2077	-0.0765	0.9503	-0.2926	0.4217	-0.2160	0.5789
CG33095	CG33095	1635429_at	0.8186	0.0026	0.2461	0.3745	0.0030	0.9916	-0.1541	0.6897	0.5361	0.0094	0.6902	0.0018	0.1004	0.9340	0.0554	0.9277	-0.0450	0.9342
CG6791	CG6791	1635430_a_at	-0.2921	0.3499	0.5787	0.1088	0.9890	0.0096	0.1059	0.9314	-0.8126	0.0336	-0.9185	0.0127	-0.1264	0.9467	0.2915	0.6382	0.4179	0.4799
CG40092	CG40092	1635431_at	0.2478	0.1047	0.0166	0.9561	0.0461	0.7815	0.0443	0.9599	0.1662	0.4927	0.1219	0.5989	-0.0942	0.8479	-0.1578	0.4391	-0.0636	0.7964
CG14607	CG14607	1635432_at	-0.0973	0.7255	-0.0443	0.8459	0.0428	0.8291	0.1620	0.7576	-0.0089	0.9798	-0.1709	0.4181	0.0768	0.9284	0.0405	0.9269	-0.0363	0.9241
---	---	1635433_at	0.3481	0.1362	-0.0095	0.9540	0.3055	0.0693	0.0318	0.9745	0.1699	0.5054	0.1381	0.5631	-0.1465	0.8012	-0.0990	0.7213	0.0475	0.8870
---	---	1635434_at	0.1539	0.4772	0.0030	0.9922	0.5217	0.0045	0.1841	0.6135	-0.0833	0.6913	-0.2674	0.1014	0.0856	0.9266	0.0171	0.9739	-0.0684	0.8667
CG30278	CG30278	1635435_at	0.1427	0.4468	-0.0880	0.4475	-0.0487	0.8452	-0.0824	0.8902	0.1151	0.5985	0.1975	0.2743	-0.0387	0.9619	-0.0805	0.7697	-0.0418	0.8926
---	---	1635436_at	-0.6433	0.0604	-0.3390	0.0854	-0.7513	0.0146	-0.2426	0.6504	0.0603	0.8583	0.3029	0.1886	0.0840	0.8940	0.0145	0.9706	-0.0695	0.8128
CG11920	CG11920	1635437_at	0.3365	0.3347	-0.1734	0.6431	-0.1725	0.3569	0.2147	0.8048	0.4934	0.1525	0.2788	0.3858	0.2118	0.8434	0.0191	0.9815	-0.1926	0.6939
---	---	1635438_at	0.0432	0.8228	-0.0142	0.8895	-0.2444	0.1333	-0.0674	0.9142	0.2305	0.2419	0.2979	0.0903	-0.1106	0.8427	0.0742	0.7985	0.1848	0.4322
CG10063	CG10063	1635439_at	0.0825	0.5640	0.0932	0.4856	0.0876	0.6935	0.2947	0.3742	0.1029	0.6196	-0.1918	0.2547	0.2644	0.6749	0.1528	0.5786	-0.1116	0.7007
---	---	1635440_at	0.1203	0.6363	0.0924	0.5271	0.1607	0.4240	0.1544	0.7539	0.0317	0.9153	-0.1227	0.5595	0.1194	0.8097	-0.0175	0.9590	-0.1369	0.5271
CG33337	CG33337	1635441_at	0.1158	0.5064	-0.0423	0.7891	0.0189	0.9244	-0.0072	0.9922	0.0182	0.9317	0.0253	0.8879	-0.0025	0.9984	0.0183	0.9639	0.0208	0.9516
GRHR	hormone receptor	1635442_a_at	2.9125	0.0089	1.7941	0.0419	2.5067	0.0014	0.3600	0.8856	0.1526	0.8952	-0.2073	0.8300	-0.3946	0.8725	-1.0658	0.2532	-0.6713	0.5050
CG18539	CG18539	1635443_at	0.2098	0.3770	0.1459	0.4514	0.1062	0.5699	0.0477	0.9436	-0.0034	0.9902	-0.0511	0.8135	0.0727	0.9117	0.0644	0.8356	-0.0083	0.9835
CG32663	CG32663	1635444_at	0.3853	0.3518	1.0582	0.2402	0.6758	0.0620	-0.6378	0.0821	-0.3140	0.1375	0.3238	0.0882	-0.1939	0.9557	0.4723	0.6762	0.6662	0.5340
sls	D-Titin	1635445_at	-1.7853	0.0215	-1.9043	0.0386	-3.2196	0.0007	-0.2710	0.4793	-0.0728	0.7675	0.1982	0.2863	0.8779	0.7707	-0.2481	0.8979	-1.1260	0.3898
CG15043 /// DyakCG15043	CG15043	1635446_at	0.2145	0.4574	-0.0717	0.6444	0.4393	0.0460	0.6934	0.0198	0.4835	0.0085	-0.2099	0.1351	-0.0206	0.9914	-0.0829	0.8963	-0.0623	0.9161
CG4827	5' nucleotidase pr	1635447_at	-1.1757	0.2166	-0.4577	0.0753	-0.4755	0.1539	0.3268	0.8132	-1.1284	0.0445	-1.4552	0.0096	0.2748	0.9238	-0.7039	0.4655	-0.9787	0.3149
nmd	Mitochondrial sort	1635448_at	0.0154	0.9478	-0.1617	0.3118	-0.1275	0.4260	-0.0697	0.8917	0.2448	0.1525	0.3145	0.0463	-0.1001	0.8870	0.0066	0.9915	0.1067	0.7223
Dip-B	Dipeptidase B	1635449_s_at	-1.0045	0.0023	-1.4331	0.0063	-1.5104	0.0001	0.0196	0.9803	0.2550	0.1778	0.2354	0.1655	0.0654	0.9445	-0.1249	0.6952	-0.1903	0.5209
smp-30	Drosophila cold a	1635450_a_at	2.9425	0.0026	1.8230	0.1702	1.5283	0.0081	0.2374	0.9280	-1.3985	0.0749	-1.6359	0.0268	0.4174	0.8917	-2.6473	0.0427	-3.0647	0.0422
---	---	1635451_s_at	0.1319	0.4315	-0.0095	0.9381	-0.0892	0.5489	-0.1190	0.7596	0.0490	0.8128	0.1680	0.2648	0.0628	0.9064	0.0147	0.9629	-0.0481	0.8444
CG14897	CG14897	1635452_a_at	0.2700	0.2117	-0.0077	0.9493	-0.1023	0.6700	0.0239	0.9756	0.1099	0.5994	0.0859	0.6649	0.0804	0.9174	-0.0954	0.7709	-0.1758	0.5355
CG13095 /// DyakCG13095	CG13095	1635453_at	1.2818	0.4044	-0.6216	0.0971	0.1187	0.7676	0.7621	0.2038	0.9786	0.0147	0.2165	0.5397	-0.3694	0.9555	-1.2268	0.5422	-0.8575	0.6877
CG12213	CG12213	1635454_a_at	-0.2378	0.3531	0.2155	0.1878	0.3081	0.1042	-0.1316	0.7794	-0.5605	0.0113	-0.4290	0.0222	-0.0345	0.9717	0.0190	0.9628	0.0535	0.8717
CG14529	CG14529	1635455_at	0.6609	0.0591	0.4228	0.1187	0.3518	0.0453	0.1369	0.7796	0.0387	0.8880	-0.0982	0.6380	0.1824	0.8744	-0.3147	0.4997	-0.4972	0.2869
CG31235	CG31235	1635456_at	-0.0029	0.9872	0.1345	0.3582	0.3237	0.1853	-0.0380	0.9436	-0.2844	0.0657	-0.2464	0.0731	0.0012	0.9996	0.0340	0.9407	0.0328	0.9344
CG31736	CG31736	1635457_at	-0.0752	0.7108	-0.0057	0.9767	-0.0065	0.9765	0.0208	0.9777	-0.0785	0.7239	-0.0993	0.6058	-0.0202	0.9877	0.0650	0.8779	0.0851	0.8173
---	---	1635458_at	0.0477	0.8041	0.0691	0.4979	0.1170	0.4718	-0.0429	0.9376	0.0608	0.7482	0.1036	0.4992	0.0437	0.9530	0.1379	0.5509	0.0943	0.7034
CG40156	CG40156	1635459_at	0.2481	0.2629	0.0688	0.4744	0.1503	0.4731	0.0016	0.9988	-0.0160	0.9609	-0.0176	0.9492	0.1046	0.8972	-0.0602	0.8924	-0.1648	0.6128

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Mhc	myosin	1635460_a_at	-1.4166	0.0079	-1.6791	0.0092	-2.3691	0.0001	-0.2514	0.2905	-0.3320	0.0256	-0.0807	0.5594	0.3026	0.8603	-0.5941	0.3805	-0.8968	0.2130
CG3362	CG3362	1635461_at	-0.2523	0.3255	0.2059	0.4906	0.3223	0.0595	-0.0047	0.9952	-0.4789	0.0095	-0.4741	0.0060	-0.0134	0.9943	0.1459	0.7813	0.1593	0.7456
rho	veinlet	1635462_at	1.0420	0.2809	2.3160	0.0185	-0.4745	0.0762	-1.0194	0.1532	0.1601	0.7562	1.1795	0.0086	1.9029	0.3712	1.3986	0.2018	-0.5043	0.6877
CG8407	CG8407	1635463_at	-0.0362	0.8880	0.0558	0.6076	0.2704	0.2184	-0.0260	0.9745	-0.0722	0.7635	-0.0462	0.8413	-0.0953	0.8885	0.0308	0.9404	0.1261	0.6476
---	---	1635464_at	0.1690	0.2482	-0.0185	0.9201	0.0252	0.9053	0.1968	0.7023	0.1957	0.4240	-0.0011	0.9973	0.0860	0.8940	0.0005	0.9998	-0.0855	0.7582
---	---	1635465_at	0.1882	0.4551	0.1106	0.4256	0.3368	0.0905	0.0842	0.8715	0.1222	0.5281	0.0379	0.8577	0.1057	0.8968	0.1344	0.6945	0.0287	0.9464
Uch-L3	ubiquitin C-termin	1635466_at	-0.1053	0.5241	0.6932	0.0575	0.9876	0.0011	0.1341	0.7929	-0.4288	0.0448	-0.5629	0.0089	-0.2677	0.7392	0.3589	0.2770	0.6265	0.1013
CG7381	CG7381	1635467_a_at	-0.6790	0.0486	-1.0819	0.0096	-0.8925	0.0091	0.0139	0.9922	0.2606	0.3884	0.2467	0.3637	-0.1118	0.9365	-0.2210	0.6458	-0.1092	0.8474
CG7730	CG7730	1635468_a_at	0.0281	0.9487	0.0180	0.9714	-0.1993	0.7970	-0.6396	0.0555	0.1350	0.5060	0.7745	0.0015	-0.3392	0.9400	0.1416	0.9497	0.4808	0.7640
CG18228 /// DsimCG18228	CG18228	1635469_at	-0.4227	0.0191	0.5950	0.0955	0.1555	0.3624	-0.2860	0.4815	-0.8887	0.0024	-0.6027	0.0081	0.0793	0.9296	-0.1289	0.6945	-0.2082	0.4889
GNBP3	Gram-negative ba	1635470_at	0.3339	0.0670	0.2758	0.6417	1.3511	0.0023	0.3285	0.3433	-0.7802	0.0029	-1.1087	0.0004	-0.7919	0.6695	-0.9001	0.2130	-0.1082	0.9168
CG7304	CG7304	1635471_at	-0.0727	0.7043	0.0667	0.5379	0.1280	0.3626	0.0207	0.9777	-0.0698	0.7444	-0.0905	0.6234	0.0363	0.9611	0.0740	0.7716	0.0376	0.8951
---	---	1635472_at	0.0688	0.7875	0.1406	0.5105	0.3537	0.0865	-0.0258	0.9774	-0.1447	0.5460	-0.1189	0.5942	-0.2351	0.7644	-0.0068	0.9931	0.2283	0.5139
Apf	diadenosine tetra	1635473_at	-0.1705	0.3723	0.3095	0.1440	0.2156	0.2057	-0.1315	0.7271	-0.3302	0.0553	-0.1986	0.1876	-0.1132	0.8650	0.0882	0.7906	0.2014	0.4634
jet	jetlag	1635474_at	0.2311	0.1211	-0.0358	0.8192	0.0896	0.5897	0.0915	0.8405	0.1805	0.2922	0.0889	0.6027	0.0023	0.9990	-0.0374	0.9354	-0.0397	0.9211
corn	cornetto	1635475_at	1.0217	0.0220	0.5243	0.1243	1.0712	0.0018	0.3877	0.5617	0.4337	0.2037	0.0459	0.9117	-0.4879	0.6749	-0.2027	0.7142	0.2853	0.5814
CG9752	CG9752	1635476_at	-0.7874	0.1863	-1.4450	0.0319	-1.8313	0.0093	-0.2869	0.4141	0.4755	0.0230	0.7623	0.0017	-0.1183	0.9764	0.0576	0.9714	0.1760	0.8933
---	---	1635477_at	0.2102	0.4731	-0.2692	0.0617	0.1103	0.5089	0.3628	0.3921	0.4126	0.0854	0.0497	0.8585	-0.1010	0.8689	-0.0332	0.9318	0.0678	0.8235
---	---	1635478_at	0.1190	0.5225	0.0641	0.6751	0.1624	0.2915	0.0771	0.8526	0.0801	0.6333	0.0030	0.9874	0.0906	0.8884	0.1330	0.6093	0.0424	0.8998
CG32056	CG32056	1635479_a_at	-0.2167	0.3669	0.4359	0.1190	0.3053	0.2151	-0.2484	0.5735	-0.3893	0.0851	-0.1410	0.5189	-0.0482	0.9742	0.3525	0.3272	0.4007	0.2930
---	---	1635480_at	0.2140	0.3047	0.0383	0.8974	0.0863	0.5915	0.1144	0.8035	0.1030	0.6117	-0.0114	0.9604	-0.0832	0.8767	-0.0457	0.8787	0.0375	0.8957
CG6659	CG6659	1635481_a_at	-0.1421	0.6344	0.2319	0.5557	0.6392	0.0425	-0.0388	0.9672	-0.0883	0.7665	-0.0495	0.8637	-0.2555	0.8732	0.4471	0.4875	0.7026	0.2800
CG2976	CG2976	1635482_at	0.2971	0.1816	0.2961	0.1015	0.3288	0.1832	-0.0922	0.8248	0.1528	0.3526	0.2450	0.0897	-0.0893	0.9466	0.2923	0.4754	0.3816	0.3569
CG16884 /// DsimCG16884	CG16884	1635483_at	-0.1468	0.7096	-0.1437	0.3450	0.1000	0.6471	0.1736	0.8578	-0.1668	0.6748	-0.3404	0.2801	0.0917	0.8611	0.0642	0.8164	-0.0275	0.9252
---	---	1635484_at	0.1030	0.6145	0.0765	0.5874	0.0130	0.9474	0.0037	0.9956	0.0730	0.6761	0.0694	0.6628	0.0092	0.9943	0.0831	0.8319	0.0739	0.8444
Eip93F	Eip93F	1635485_at	0.4352	0.2249	-0.0092	0.9314	0.1303	0.7109	0.1181	0.8589	0.4745	0.0522	0.3564	0.0955	0.0735	0.9742	0.1581	0.8319	0.0846	0.9121
CG33468	CG33468	1635486_at	-0.2924	0.5298	-1.7955	0.0759	-2.2575	0.0011	0.2232	0.6823	2.7095	0.0001	2.4863	0.0000	0.5222	0.8465	1.0634	0.3305	0.5412	0.6567
CG3534	CG3534	1635487_at	1.4050	0.0008	0.4233	0.0594	0.8232	0.0009	0.1320	0.7025	0.5208	0.0063	0.3888	0.0141	-0.2263	0.7644	-0.4134	0.1838	-0.1871	0.5838
CG31660	CG31660	1635488_at	0.1032	0.5344	0.0043	0.9745	-0.0046	0.9830	0.0948	0.8611	0.1414	0.4874	0.0466	0.8338	0.0108	0.9914	0.0292	0.9341	0.0184	0.9525
---	---	1635489_at	0.3311	0.0633	0.2292	0.3532	0.0703	0.7732	-0.2465	0.6597	0.0175	0.9666	0.2641	0.2773	0.2476	0.6955	0.0868	0.7891	-0.1608	0.5587
---	---	1635490_at	-0.0981	0.5660	-0.1775	0.5407	-0.0082	0.9677	0.1097	0.9136	0.0552	0.8994	-0.0545	0.8856	-0.0254	0.9831	-0.1291	0.6745	-0.1037	0.7451
TRAM	TRAM	1635491_s_at	1.7977	0.0011	1.8212	0.0025	2.0377	0.0000	0.4975	0.0754	0.5681	0.0046	0.0706	0.6766	0.3553	0.6832	0.6596	0.0795	0.3043	0.4051
CG33259	CG33259	1635492_at	-0.0308	0.9271	0.0416	0.7722	0.0089	0.9624	0.1376	0.7969	0.1047	0.6717	-0.0328	0.9000	0.0095	0.9914	0.0038	0.9935	-0.0057	0.9856
---	---	1635493_s_at	-0.1106	0.6232	-0.0298	0.7922	-0.0666	0.7339	0.1127	0.7556	0.0323	0.8766	-0.0805	0.6108	0.0291	0.9800	0.0287	0.9459	-0.0004	0.9992
Bc	Black Cell	1635494_at	-1.2480	0.2140	-3.5222	0.0370	-3.4808	0.0000	-0.2619	0.7031	0.6997	0.0342	0.9616	0.0053	-0.1769	0.9816	-1.3579	0.4258	-1.1810	0.5050
DopR2	dopamine recepto	1635495_a_at	-0.5095	0.1474	0.0036	0.9768	-0.2512	0.1362	-0.2527	0.6130	-0.5067	0.0486	-0.2541	0.2603	0.0062	0.9950	-0.0041	0.9935	-0.0103	0.9777
---	---	1635496_at	0.2008	0.4708	0.1972	0.2172	0.1584	0.4408	0.0370	0.9506	-0.0675	0.7255	-0.1045	0.5116	0.1228	0.8461	0.0186	0.9652	-0.1042	0.7262
---	---	1635497_at	0.1391	0.4548	-0.0007	1.0000	-0.2076	0.1642	-0.1972	0.5480	-0.0182	0.9407	0.1790	0.2400	0.1232	0.8235	-0.0366	0.9178	-0.1598	0.5022
CG9259 /// DyakCG9259	CG9259	1635498_at	-1.5592	0.3986	-0.0413	0.8954	0.0063	0.9821	0.5302	0.8930	-1.9725	0.1320	-2.5027	0.0397	0.3862	0.9457	-0.6882	0.7275	-1.0743	0.5518
---	---	1635499_at	-0.0891	0.6233	0.0208	0.8703	0.2071	0.1638	0.1127	0.8485	-0.0455	0.8732	-0.1582	0.4345	0.0215	0.9841	0.0637	0.8478	0.0422	0.9004
pros	a la voile et a la v	1635500_a_at	-0.4182	0.7441	-0.2245	0.1485	-0.2247	0.2604	0.1057	0.9727	-0.1717	0.8656	-0.2774	0.7374	-0.1755	0.8097	-0.2912	0.3253	-0.1157	0.7411
p120ctn	p120 catenin	1635501_at	-0.4698	0.0457	-0.6825	0.0108	-1.1807	0.0001	-0.2180	0.4902	0.2297	0.1700	0.4477	0.0097	0.3390	0.6145	0.1826	0.5414	-0.1565	0.6129
CG8311	CG8311	1635502_at	1.2951	0.0087	0.7141	0.2277	1.0939	0.0003	0.1944	0.5859	0.2929	0.1064	0.0985	0.5885	-0.2500	0.8940	-0.3093	0.6949	-0.0593	0.9523
Vm34Ca	Vitelline membran	1635503_at	0.5922	0.5430	-0.7119	0.0664	-0.2592	0.2650	0.8331	0.2172	1.0665	0.0167	0.2334	0.5601	-0.1194	0.9816	-0.6771	0.5825	-0.5577	0.6578
---	---	1635504_s_at	-0.1264	0.4069	0.0373	0.7192	0.1141	0.6121	0.0619	0.9380	-0.0182	0.9582	-0.0801	0.7539	-0.0280	0.9807	0.1094	0.7002	0.1373	0.6128
CG17274	CG17274	1635505_a_at	0.2513	0.1132	0.0390	0.8360	0.2803	0.1149	0.0035	0.9956	0.1066	0.6177	0.1031	0.5941	0.0080	0.9952	0.0852	0.8353	0.0771	0.8439
CG11777	CG11777	1635506_at	-0.3688	0.1446	0.1725	0.3233	-0.2613	0.0722	-0.3079	0.4073	-0.2726	0.1845	0.0353	0.8859	0.1591	0.8033	0.3324	0.1961	0.1733	0.5361
CG4269	CG4269	1635507_at	0.4726	0.4576	0.7768	0.0854	0.3459	0.4791	-0.2739	0.8402	0.3083	0.5734	0.5823	0.1983	0.2630	0.9142	0.7862	0.3479	0.5232	0.5658
Taf10	TBP-associated fa	1635508_at	-0.2266	0.4157	0.3416	0.2935	0.0810	0.6959	-0.0187	0.9860	-0.2182	0.4025	-0.1995	0.3978	0.2285	0.8141	0.3767	0.3353	0.1483	0.7492
---	---	1635509_at	0.5753	0.4886	-0.0715	0.9132	0.4129	0.0669	0.6217	0.0741	0.5756	0.0115	-0.0461	0.8427	0.1435	0.9775	-0.0461	0.9841	-0.1896	0.9094
CG34025	CG34025	1635510_at	0.0794	0.7068	-0.1363	0.3731	0.2449	0.1638	0.2137	0.6041	0.1663	0.4400	-0.0474	0.8434	-0.1891	0.7230	-0.0771	0.7891	0.1120	0.6533
---	---	1635511_at	0.1477	0.2849	-0.1338	0.1939	0.1354	0.4469	0.4155	0.2171	0.3534	0.0814	-0.0621	0.7827	0.1156	0.8478	-0.0377	0.9231	-0.1534	0.5634
CG11893	CG11893	1635512_at	-0.3791	0.3537	0.5397	0.2620	-0.9992	0.1430	-0.2843	0.5539	1.0544	0.0020	1.3386	0.0005	1.2297	0.6749	1.9162	0.1164	0.6865	0.5958
CG11982	CG11982	1635513_at	-0.2451	0.1379	0.2139	0.2852	0.4822	0.0187	0.1300	0.7104	-0.6076	0.0032	-0.7376	0.0008	-0.0332	0.9748	0.0212	0.9614	0.0544	0.8779
CG1273	CG1273	1635514_at	0.2444	0.4321	-0.1413	0.5651	-0.0014	0.9958	0.0295	0.9840	0.0970	0.8202	0.0675	0.8656	-0.0614	0.9408	-0.2327	0.3510	-0.1713	0.5227
---	---	1635515_at	-0.0746	0.8095	0.0255	0.8976	0.3542	0.0538	0.1655	0.7556	-0.0628	0.8288	-0.2283	0.2690	-0.3460	0.6483	-0.1617	0.6227	0.1844	0.5691
CG13972	CG13972	1635516_at	0.5361	0.0437	0.1860	0.1205	0.1643	0.6781	0.2793	0.5502	0.3279	0.1729	0.0485	0.8630	0.1268	0.9238	-0.0933	0.8828	-0.2201	0.6403
CG1667	CG1667	1635517_at	0.5594	0.2776	-1.1439	0.0048	-0.8561	0.0476	0.2717	0.8633	1.5583	0.0131	1.2866	0.0184	-0.0552	0.9741	-0.1273	0.8137	-0.0721	0.9006
nonA	no-on-transient A	1635518_at	-0.2045	0.6730	0.1528	0.7020	-0.5662	0.0416	-0.3095	0.5183	0.0121	0.9755	0.3216	0.1531	0.3349	0.8425	0.2459	0.7703	-0.0890	0.9257
CG13950	CG13950	1635519_at	2.9340	0.0004	3.0725	0.0100	4.3398	0.0000	0.5294	0.2100	0.0876	0.7744	-0.4418	0.0556	-0.3043	0.8680	0.6525	0.3589	0.9568	0.2081
---	---	1635520_at	0.2770	0.2263	0.0640	0.6686	0.0287	0.9459	0.0997	0.8839	0.3296	0.1548	0.2299	0.2738	0.2361	0.8444	0.0828	0.9137	-0.1533	0.7975
CG4618	CG4618	1635521_at	-0.3511	0.1242	-0.9925	0.0223	-0.5982	0.1175	0.4488	0.1106	0.9253	0.0007	0.4765	0.0070	0.1106	0.9589	0.4338	0.4943	0.3233	0.6279
santa-maria	CG12789	1635522_a_at	0.7077	0.0729	-0.0312	0.9507	-0.1669	0.5328	-0.7774	0.1876	-0.6646	0.0656	0.1128	0.7751	-0.5231	0.6724	-1.3388	0.0290	-0.8157	0.1313
CG6479	CG6479	1635523_a_at	0.5997	0.0118	0.0093	0.9606	0.2123	0.1495	-0.0688	0.8975	0.6452	0.0038	0.7140	0.0015	-0.3332	0.4912	-0.0881	0.7525	0.2451	0.3088
CG30427	CG30427	1635524_at	0.1528	0.7786	-0.0835	0.6994	-0.0938	0.6680	-0.0343	0.9695	0.0743	0.7930	0.1086	0.6484	-0.0439	0.9891	-0.3013	0.6876	-0.2574	0.7391
CG9672	CG9672	1635525_at	-0.1152	0.8387	-0.1242	0.4262	0.1065	0.6411	-0.0478	0.9602	-0.3111	0.2139	-0.2634	0.2424	-0.1739	0.9296	-0.2502	0.7368	-0.0763	0.9321
CG15390	CG15390	1635526_at	0.2664	0.4754	0.3522	0.4928	0.1161	0.4695	0.0176	0.9873	0.4787	0.0743	0.4611	0.0565	0.3865	0.7644	0.7468	0.1548	0.3603	0.5257
CG5039	CG5039	1635527_at	-0.1404	0.3915	0.4718	0.1957	0.4495	0.0365	-0.0401	0.9592	-0.2745	0.1792	-0.2344	0.2013	-0.0225	0.9816	0.2616	0.2107	0.2841	0.2084
RpS10b	Ribosomal protein	1635528_s_at	0.1968	0.2407	1.4875	0.0063	1.3228	0.0011	0.0689	0.9425	-1.1737	0.0019	-1.2425	0.0009	0.1032	0.8909	0.0436	0.9220	-0.0596	0.8773

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1635529_at	0.4544	0.0153	0.2692	0.1878	0.3794	0.0863	0.2207	0.5470	0.0786	0.7224	-0.1421	0.4242	0.0616	0.9298	-0.1277	0.5971	-0.1893	0.4114
CG1249	CG1249	1635530_at	0.3308	0.1267	0.1800	0.4811	0.0232	0.9098	0.0701	0.8902	0.0602	0.7678	-0.0098	0.9635	0.3404	0.6557	0.1121	0.7660	-0.2283	0.4785
rost	rolling stone	1635531_at	0.5777	0.1749	-0.0704	0.9317	0.4394	0.0778	0.4155	0.5926	1.0963	0.0130	0.6808	0.0570	0.0259	0.9928	0.5629	0.4073	0.5369	0.4462
---	---	1635532_at	-0.0112	0.9555	0.0525	0.8228	0.2791	0.1252	0.1026	0.8578	-0.1187	0.5953	-0.2213	0.2277	-0.0851	0.9095	-0.0425	0.9180	0.0426	0.9081
---	---	1635533_s_at	0.1035	0.5832	-0.0673	0.5664	0.0318	0.8478	0.0542	0.9254	0.0270	0.9114	-0.0272	0.8968	0.0232	0.9775	-0.0744	0.7439	-0.0977	0.6394
---	---	1635534_at	-0.0135	0.9382	0.0975	0.4211	0.0882	0.7267	0.1479	0.6533	0.0812	0.6500	-0.0667	0.6893	0.1118	0.9029	0.1832	0.6068	0.0714	0.8736
CG1287	CG1287	1635535_at	0.1196	0.4417	0.0698	0.4935	-0.0615	0.7704	-0.0704	0.8735	-0.0942	0.5702	-0.0237	0.8977	0.0375	0.9717	0.0354	0.9350	-0.0022	0.9965
CG14126	CG14126	1635536_at	-0.1165	0.6399	0.2172	0.3916	0.0249	0.9179	-0.0859	0.8791	-0.1276	0.5342	-0.0418	0.8516	0.0934	0.9238	0.0344	0.9476	-0.0590	0.8934
asf1	anti-silencing fact	1635537_at	-0.1487	0.4753	0.1214	0.4870	0.0710	0.8053	-0.0524	0.9255	0.0920	0.6244	0.1444	0.3518	-0.0768	0.9521	0.4352	0.2404	0.5120	0.2051
CG11210	CG11210	1635538_at	0.1735	0.6576	1.1857	0.0776	1.1274	0.0004	0.0315	0.9763	-0.5425	0.0449	-0.5740	0.0229	0.0608	0.9816	0.3880	0.5246	0.3272	0.6049
---	---	1635539_at	0.1295	0.5181	-0.0753	0.4612	0.1722	0.3374	0.1117	0.8138	0.1326	0.5009	0.0209	0.9272	0.0895	0.9011	0.1279	0.6586	0.0385	0.9162
---	---	1635540_at	0.0231	0.9226	0.1183	0.4620	0.0736	0.7442	0.1417	0.7235	0.1053	0.5926	-0.0364	0.8626	0.1653	0.8480	0.0370	0.9498	-0.1283	0.7574
MESK2	Misexpression su	1635541_s_at	-0.2059	0.4632	0.3293	0.3365	-0.2242	0.4342	-0.2131	0.7028	-0.3881	0.1289	-0.1751	0.4730	0.3251	0.7810	0.2696	0.6191	-0.0554	0.9372
l(3)02640	lethal (3) 02640	1635542_at	-0.4880	0.0768	-0.9523	0.0115	-0.5928	0.1066	0.0113	0.9924	0.4993	0.0493	0.4880	0.0349	-0.2838	0.8049	0.1445	0.8153	0.4284	0.3807
---	---	1635543_at	0.0752	0.8172	-0.0233	0.9050	-0.0735	0.7010	0.1580	0.8373	0.4210	0.1437	0.2631	0.3181	0.1974	0.8427	0.3053	0.4586	0.1079	0.8351
---	---	1635544_s_at	-2.6556	0.0157	-1.7121	0.1210	-3.0282	0.0003	-0.7767	0.0898	-2.2899	0.0001	-1.5132	0.0004	0.5382	0.8906	-1.2921	0.3726	-1.8303	0.2298
---	---	1635545_at	0.0704	0.6787	0.2405	0.1173	0.2223	0.1738	-0.0295	0.9603	-0.0731	0.6841	-0.0436	0.8052	-0.0046	0.9959	0.0604	0.8213	0.0650	0.7915
na	halothane-resistar	1635546_a_at	-0.0953	0.5498	0.1045	0.5237	0.3503	0.0654	0.1503	0.7599	-0.1222	0.5964	-0.2725	0.1481	0.0827	0.9064	0.1539	0.5613	0.0711	0.8218
Aats-trp	Tryptophanyl-tRN	1635547_a_at	-0.0241	0.9352	0.8611	0.0873	0.6401	0.0033	0.1037	0.8523	0.2010	0.3267	0.0974	0.6384	0.3477	0.7337	1.1828	0.0290	0.8352	0.0906
CG15281	CG15281	1635548_s_at	2.3037	0.0163	1.6345	0.0827	1.9920	0.0030	0.2007	0.9339	0.0954	0.9232	-0.1053	0.9021	-0.0884	0.9831	-0.4882	0.6437	-0.3998	0.7154
TotA	Turandot	1635549_at	2.4704	0.0161	0.0484	0.9764	3.0632	0.0070	2.6126	0.1663	1.4699	0.1983	-1.1427	0.2697	-0.2706	0.9545	-0.8684	0.5522	-0.5978	0.7013
CG10317	CG10317	1635550_at	0.2358	0.2918	-0.1525	0.5180	0.1371	0.4604	0.3055	0.2438	0.2979	0.0596	-0.0076	0.9689	-0.0215	0.9862	-0.1166	0.7151	-0.0951	0.7734
CG8386	CG8386	1635551_at	-0.1429	0.4970	0.4902	0.1194	0.5613	0.0115	0.1287	0.7880	-0.1581	0.4376	-0.2868	0.1019	0.1610	0.8215	0.5736	0.0697	0.4127	0.1893
CG31472	CG31472	1635552_s_at	-0.5699	0.0738	0.1361	0.5103	-0.3154	0.0538	-0.2354	0.6457	-0.6928	0.0129	-0.4574	0.0451	0.1735	0.8599	-0.0447	0.9449	-0.2182	0.6089
---	---	1635553_at	-0.0873	0.7159	0.0734	0.4567	0.0742	0.7412	-0.1622	0.8156	-0.2662	0.3375	-0.1041	0.7204	-0.0506	0.9717	-0.2280	0.5426	-0.1774	0.6463
---	---	1635554_at	-0.0869	0.6268	0.2174	0.2773	-0.1104	0.4662	-0.1612	0.6936	-0.0946	0.6577	0.0666	0.7448	0.1024	0.8680	0.1730	0.4875	0.0706	0.8167
CG10495	CG10495	1635555_at	0.0164	0.9623	-0.3382	0.3769	-0.2225	0.1621	-0.0465	0.9467	0.4771	0.0234	0.5236	0.0097	-0.1872	0.8761	0.0219	0.9796	0.2091	0.6875
---	---	1635556_at	-0.0397	0.8601	0.0856	0.6888	-0.0837	0.6425	-0.0888	0.8545	-0.1788	0.3113	-0.0900	0.6110	0.1333	0.8395	0.0367	0.9326	-0.0966	0.7576
CG3880	CG3880	1635557_at	0.2972	0.1698	-0.0006	0.9994	0.1460	0.3819	0.2260	0.4741	0.2946	0.0837	0.0687	0.7010	-0.0541	0.9589	0.0176	0.9724	0.0717	0.8555
CG17724 /// seq	CG17724 /// sequ	1635558_s_at	1.7800	0.1120	0.2044	0.7167	-1.2486	0.1787	-0.5694	0.4715	1.5358	0.0041	2.1052	0.0006	0.9634	0.8362	0.1749	0.9557	-0.7885	0.7159
CG34365	CG12958	1635559_at	0.1850	0.4911	0.0228	0.8158	0.1520	0.3779	0.1932	0.5680	0.1307	0.4722	-0.0626	0.7379	0.0507	0.9672	-0.0614	0.9009	-0.1121	0.7687
CG40263	CG40263	1635560_at	-0.0351	0.8862	-0.2003	0.4010	0.0040	0.9882	-0.2064	0.5515	-0.1246	0.5106	0.0818	0.6595	-0.4165	0.6821	-0.2884	0.5027	0.1281	0.8006
---	---	1635561_at	0.0871	0.5813	0.0860	0.5965	0.2808	0.1865	-0.0870	0.9115	-0.0868	0.7703	0.0002	0.9995	-0.3161	0.7131	-0.2747	0.4437	0.0415	0.9344
---	---	1635562_at	-0.7854	0.0568	-1.3297	0.0105	-1.4238	0.0169	0.3166	0.7857	1.0094	0.0421	0.6927	0.1063	0.1876	0.8097	0.0182	0.9738	-0.1694	0.6278
CG6694	CG6694	1635563_at	-0.2016	0.2692	-0.2262	0.5369	-0.3861	0.0417	-0.4672	0.1671	-0.1053	0.6490	0.3619	0.0534	-0.2331	0.8270	-0.1335	0.8165	0.0996	0.8646
CG7133	CG7133	1635564_at	0.2888	0.0983	0.5264	0.1787	0.4346	0.0260	-0.1816	0.6572	-0.4733	0.0242	-0.2917	0.0966	0.0368	0.9816	-0.0571	0.9124	-0.0939	0.8247
CG32060	CG32060	1635565_at	0.2019	0.4086	0.0519	0.6054	0.4286	0.0511	0.1185	0.7605	0.0860	0.6480	-0.0325	0.8682	0.0597	0.9534	0.1936	0.5404	0.1339	0.6892
Kap-alpha3	Importin alpha-3	1635566_s_at	0.5660	0.0250	-0.0109	0.9882	0.0417	0.8284	-0.3475	0.2581	0.3933	0.0372	0.7408	0.0015	-0.3147	0.7853	-0.1727	0.7774	0.1420	0.8182
swm	second mitotic wa	1635567_s_at	0.0968	0.8195	0.5539	0.2730	-0.2217	0.4008	-0.1570	0.7604	-0.0820	0.7556	0.0750	0.7564	0.4872	0.7726	0.3120	0.7053	-0.1753	0.8526
CG10080	CG10080	1635568_at	0.2673	0.4025	0.1831	0.4442	0.6403	0.0164	0.2185	0.7507	0.3731	0.2100	0.1545	0.6083	-0.1099	0.9142	0.3769	0.2764	0.4868	0.1990
---	---	1635569_at	0.0892	0.6217	-0.0006	0.9990	-0.1089	0.4621	0.1557	0.7187	0.1624	0.4233	0.0067	0.9785	-0.0326	0.9775	-0.1221	0.6932	-0.0895	0.7838
CrebB-17A	cAMP-regulated e	1635570_a_at	0.1615	0.6083	0.6163	0.1190	0.2536	0.1136	-0.0562	0.9507	0.0761	0.8064	0.1323	0.6023	0.4278	0.6749	0.7231	0.0934	0.2953	0.4948
---	---	1635571_at	-0.1268	0.6203	-0.0036	0.9884	0.2169	0.1601	0.1262	0.7368	0.0043	0.9868	-0.1219	0.4355	-0.0486	0.9717	0.1189	0.7779	0.1675	0.6495
CG30058	CG30058	1635572_at	-0.2226	0.4864	-0.0435	0.8333	-0.1052	0.6123	-0.0001	0.9999	0.0692	0.8107	0.0693	0.7911	-0.0025	0.9990	0.1951	0.5123	0.1976	0.5093
CG12488	CG12488	1635573_at	-1.3531	0.0417	-0.8126	0.0600	-1.3821	0.0017	-0.1535	0.9441	0.0737	0.9333	0.2272	0.7357	-0.1775	0.9225	0.0677	0.9451	0.2452	0.7205
CG33347	CG33347	1635574_at	0.0999	0.5139	0.0162	0.9153	-0.0268	0.9004	-0.0641	0.9186	0.0965	0.6577	0.1607	0.3671	-0.0956	0.9238	-0.0578	0.9086	0.0378	0.9350
CG40032	CG40032	1635575_at	0.1211	0.6398	-0.1386	0.4033	0.0698	0.6769	-0.0845	0.8803	0.0964	0.6566	0.1809	0.2988	-0.2530	0.7062	-0.0397	0.9279	0.2133	0.4605
Nplp2	neuropeptide-like	1635576_at	1.2140	0.0049	1.1248	0.0435	1.6366	0.0005	0.4205	0.4356	0.2869	0.3405	-0.1336	0.6634	-0.1703	0.8867	-0.1817	0.7248	-0.0113	0.9874
---	---	1635577_at	0.0145	0.9414	0.1343	0.5052	0.1979	0.3921	-0.0455	0.9538	-0.1864	0.3975	-0.1408	0.4939	0.0091	0.9929	0.0747	0.8126	0.0656	0.8306
CG31975	CG31975	1635578_at	0.2523	0.3157	0.1075	0.3327	0.3575	0.0732	0.2285	0.5891	0.0645	0.8079	-0.1640	0.4141	-0.0246	0.9809	-0.0005	0.9996	0.0240	0.9382
CG2209	CG2209	1635579_at	0.0944	0.6832	-0.0378	0.8585	-0.1814	0.3353	0.0057	0.9943	0.0951	0.6208	0.0895	0.6100	0.0681	0.9514	-0.0160	0.9784	-0.0841	0.8418
Cbl	Cbl	1635580_at	-0.8900	0.1863	-1.1092	0.0798	-1.6332	0.0003	-0.5837	0.2905	0.3478	0.2878	0.9315	0.0069	-0.2744	0.8978	-0.0657	0.9602	0.2087	0.8361
CG8300	CG8300	1635581_at	-1.1825	0.0141	-1.3125	0.0278	-1.6997	0.0046	0.0705	0.9248	0.0928	0.7224	0.0223	0.9362	0.3214	0.8882	-0.2235	0.8487	-0.5448	0.5488
---	---	1635582_at	0.0281	0.8888	0.1168	0.5127	0.0143	0.9473	-0.0120	0.9866	-0.0798	0.6889	-0.0678	0.7135	0.1682	0.8303	-0.0919	0.8346	-0.2602	0.4411
fz2	D-frizzled2	1635583_s_at	-0.4558	0.4601	-0.0195	0.9823	-1.5181	0.0482	-0.7942	0.2114	-0.4809	0.2069	0.3133	0.3754	0.6205	0.8637	0.0044	0.9996	-0.6161	0.7045
Tm1	Tropomyosin	1635584_s_at	-1.5549	0.0368	-1.1775	0.1204	-2.2421	0.0010	-0.5557	0.2008	-1.3217	0.0010	-0.7660	0.0063	0.5776	0.8379	-0.8828	0.4475	-1.4604	0.2235
CG9264	CG9264	1635585_at	-0.4729	0.1367	-0.1799	0.3446	-0.3204	0.1149	-0.0481	0.9592	-0.2532	0.3109	-0.2051	0.3691	-0.0226	0.9853	-0.1145	0.7250	-0.0918	0.7839
CG14148	CG14148	1635586_at	0.1510	0.4544	0.4932	0.0234	-0.1162	0.6423	-0.2598	0.5608	-0.2867	0.2091	-0.0270	0.9235	0.2849	0.7440	0.1752	0.6660	-0.1097	0.8088
CG4050	CG4050	1635587_s_at	0.4659	0.0777	0.6528	0.0945	0.0529	0.7393	-0.2612	0.4770	0.4659	0.0260	0.7272	0.0022	0.3789	0.7440	0.6515	0.1705	0.2726	0.6046
CG14977	CG14977	1635588_at	-0.1118	0.5530	-0.2660	0.4097	-0.3273	0.0495	0.0763	0.8692	0.0551	0.7801	-0.0212	0.9149	0.1004	0.9168	-0.1367	0.7200	-0.2371	0.4908
CG7920 /// DsimCG7920	CG7920	1635589_at	-0.2747	0.0834	0.1080	0.6341	0.2890	0.1483	-0.0292	0.9608	-0.7131	0.0016	-0.6839	0.0011	-0.2679	0.7506	-0.3550	0.3137	-0.0871	0.8524
CG34109	CG34109	1635590_a_at	1.1660	0.4129	-2.2711	0.0020	0.3109	0.2212	2.7669	0.0001	2.6188	0.0000	-0.1481	0.4748	0.0144	0.9990	-0.9150	0.6600	-0.9294	0.6479
NijA	ninjurin A	1635591_at	-1.2500	0.0319	-1.7051	0.0056	-1.5044	0.0007	-0.1263	0.8070	-0.4957	0.0240	-0.3693	0.0495	-0.2441	0.9108	-0.9202	0.2189	-0.6761	0.3921
vav	vav	1635592_at	-0.1776	0.5231	-0.5333	0.0441	-0.8274	0.0021	-0.1324	0.7531	0.6234	0.0052	0.7559	0.0013	0.0314	0.9831	0.1157	0.7910	0.0843	0.8515
Cyp6d2	Cyp6d2	1635593_at	1.2499	0.0014	1.0300	0.0105	2.3450	0.0000	0.5341	0.2461	0.2835	0.3032	-0.2506	0.3123	-0.6054	0.3771	0.1703	0.6857	0.7757	0.0761
CG5255	CG5255	1635594_at	0.1479	0.4547	0.1935	0.4651	0.1806	0.2094	-0.0148	0.9863	-0.0854	0.7196	-0.0706	0.7511	-0.0569	0.9116	-0.0809	0.6916	-0.0240	0.9234
CG10996	CG10996	1635595_at	0.1081	0.6682	0.1287	0.3388	0.0466	0.8151	0.0311	0.9672	0.0886	0.6994	0.0575	0.7973	0.0821	0.9095	0.1014	0.7364	0.0194	0.9588
CG6568	CG6568	1635596_at	-0.0570	0.7715	0.5500	0.0349	0.4923	0.0087	-0.2351	0.4455	-0.8391	0.0010	-0.6040	0.0025	0.0261	0.9816	0.0025	0.9972	-0.0235	0.9491
Snap24	Synapse protein 2	1635597_at	-0.9794	0.0068	0.1578	0.6856	0.4212	0.0859	0.0205	0.9863	-0.9502	0.0062	-0.9707	0.0034	-0.1392	0.9174	0.3198	0.5055	0.4589	0.3357

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18266	CG18266	1635598_at	0.1749	0.4840	0.0268	0.8640	0.1570	0.3018	-0.0459	0.9380	0.0450	0.8405	0.0909	0.6023	0.0291	0.9742	-0.0440	0.8929	-0.0731	0.7793
Or98a	Odorant receptor	1635599_at	0.0842	0.6655	-0.1101	0.5280	0.1345	0.4918	0.0561	0.9397	0.1381	0.5503	0.0821	0.7200	-0.2157	0.7810	-0.1088	0.7985	0.1069	0.7909
---	---	1635600_at	0.0555	0.7576	-0.0204	0.9337	0.0034	0.9922	0.0236	0.9777	0.0336	0.9114	0.0099	0.9710	0.0681	0.9457	-0.0002	1.0000	-0.0683	0.8641
CG13420	CG13420	1635601_at	0.5594	0.1727	0.2748	0.3747	0.9558	0.0131	-0.0196	0.9826	-0.0307	0.9169	-0.0111	0.9669	-0.5977	0.7196	-0.2463	0.7692	0.3514	0.6366
CG16956	CG16956	1635602_at	0.2355	0.3137	0.0492	0.8423	0.2747	0.1863	0.0285	0.9777	0.0397	0.9134	0.0112	0.9725	-0.1921	0.7707	-0.1189	0.7090	0.0732	0.8375
CG17202 /// DyakCG17202	CG17202	1635603_at	0.0337	0.8728	0.6127	0.0524	0.8002	0.0010	0.3166	0.3816	-0.0383	0.8902	-0.3549	0.0562	0.0330	0.9689	0.5459	0.0384	0.5129	0.0560
CG15923	CG15923	1635604_at	0.0874	0.6167	-0.0859	0.5275	-0.0876	0.5796	-0.1137	0.7539	-0.1271	0.4345	-0.0134	0.9463	-0.0222	0.9816	-0.1740	0.4475	-0.1519	0.5274
CG32694	CG32694	1635605_at	-0.0975	0.7016	-0.2941	0.0416	-0.3076	0.2236	0.0369	0.9702	0.1863	0.4783	0.1494	0.5436	0.2523	0.7606	0.1570	0.6848	-0.0953	0.8260
dsf	dissatisfaction	1635606_at	0.1285	0.6369	0.0309	0.7568	-0.0597	0.7936	0.0650	0.9351	0.1109	0.6800	0.0459	0.8681	0.0256	0.9816	0.0068	0.9900	-0.0188	0.9565
CG31742	AE003657 - prote	1635607_at	0.2099	0.2986	0.1019	0.3652	0.2878	0.1790	-0.0865	0.8841	0.0547	0.8319	0.1412	0.4614	-0.0041	0.9964	0.1628	0.4224	0.1668	0.4220
Nlp	nucleoplasmin-1	1635608_at	0.3859	0.0505	0.2284	0.2186	-0.1353	0.5503	-0.0439	0.9436	0.5143	0.0110	0.5582	0.0045	0.2644	0.7220	0.3225	0.2938	0.0581	0.8932
CG40294	CG40294	1635609_at	-0.2499	0.1781	-0.4773	0.1321	-0.5072	0.2000	-0.0365	0.9847	0.0186	0.9780	0.0551	0.9172	0.0402	0.9831	-0.2137	0.6512	-0.2539	0.5835
CG8310	CG8310	1635610_at	-1.1275	0.0140	-0.8130	0.1673	-0.8929	0.0008	-0.0867	0.8939	-0.3924	0.0723	-0.3057	0.1124	-0.1711	0.8331	-0.2250	0.5199	-0.0539	0.9081
---	---	1635611_a_at	0.3601	0.1519	0.3834	0.0648	0.2650	0.1370	0.0672	0.9116	-0.1433	0.4747	-0.2105	0.2198	0.0679	0.9400	-0.2109	0.4466	-0.2787	0.3248
---	---	1635612_at	0.0521	0.8521	-0.0486	0.7194	0.2361	0.1533	0.0803	0.8358	0.0340	0.8595	-0.0463	0.7750	-0.0628	0.9449	-0.1122	0.7234	-0.0494	0.8958
CG5079	CG5079	1635613_at	0.0251	0.8726	-0.0554	0.5835	-0.1552	0.4618	-0.1086	0.7247	-0.0109	0.9579	0.0977	0.4565	0.0660	0.9087	0.0007	0.9994	-0.0653	0.7879
---	---	1635614_at	0.0902	0.6683	0.0790	0.4229	0.0487	0.8126	-0.0352	0.9598	0.0840	0.6877	0.1192	0.4968	-0.0660	0.9499	0.0400	0.9363	0.1060	0.7745
---	---	1635615_at	0.1110	0.5968	0.1291	0.4155	0.3128	0.1346	0.1645	0.6338	0.0649	0.7468	-0.0996	0.5497	-0.1600	0.8609	-0.0105	0.9898	0.1495	0.7205
CG2118	CG2118	1635616_a_at	-0.0572	0.7830	0.3959	0.2076	1.2220	0.0066	-0.1489	0.7046	-0.7826	0.0019	-0.6337	0.0027	-0.7577	0.5126	-0.2043	0.7579	0.5535	0.3299
---	---	1635617_at	-0.1015	0.5496	0.0130	0.9575	-0.0239	0.9108	0.0459	0.9464	-0.0120	0.9679	-0.0579	0.7878	0.0064	0.9959	0.0781	0.8394	0.0717	0.8459
Dok	Downstream of kir	1635618_at	-0.0788	0.6565	0.1059	0.5553	-0.1234	0.7442	-0.5635	0.2397	-0.3239	0.2543	0.2396	0.3579	-0.2515	0.8009	-0.1232	0.8196	0.1283	0.7983
cnn	oblivious	1635619_a_at	-1.2235	0.0108	-2.2420	0.0014	-2.4764	0.0000	-0.4176	0.4568	0.1327	0.7056	0.5503	0.0482	-0.3089	0.7464	-0.9179	0.0446	-0.6090	0.1562
---	---	1635620_at	-0.2787	0.1572	-0.1835	0.4243	-0.3287	0.1946	-0.0119	0.9909	-0.0095	0.9786	0.0025	0.9928	0.0466	0.9672	-0.1224	0.7209	-0.1690	0.5995
CG13994	CG13994	1635621_at	0.2208	0.3676	-0.0665	0.7891	-0.1742	0.2212	0.1471	0.6942	0.4455	0.0182	0.2985	0.0575	0.2162	0.8049	0.1646	0.6868	-0.0516	0.9186
---	---	1635622_at	0.0907	0.5444	0.2756	0.0891	0.4131	0.0325	0.0638	0.8865	-0.0013	0.9951	-0.0651	0.6810	-0.0406	0.9588	0.2036	0.3380	0.2442	0.2784
Cpr73D	CG9665	1635623_at	0.7963	0.0492	-0.0234	0.9517	0.1966	0.2346	0.0017	0.9986	-0.2847	0.1604	-0.2863	0.1145	-0.2245	0.7464	-0.8832	0.0241	-0.6587	0.0607
---	---	1635624_at	0.2737	0.1526	0.2549	0.1166	0.6374	0.0110	0.0972	0.8265	-0.0743	0.7061	-0.1715	0.2659	-0.2474	0.7644	-0.0605	0.9095	0.1869	0.6189
---	---	1635625_at	0.1712	0.4086	-0.0070	0.9515	0.1787	0.3036	0.1388	0.8149	-0.0069	0.9839	-0.1458	0.5195	-0.0485	0.9486	-0.1139	0.6408	-0.0655	0.8153
---	---	1635626_s_at	0.0988	0.5367	0.1392	0.4053	0.1824	0.3010	-0.0621	0.8815	-0.0206	0.9175	0.0415	0.7944	-0.0366	0.9793	-0.0581	0.9068	-0.0214	0.9640
CSN3	COP9 complex hc	1635627_at	-0.0114	0.9612	0.4031	0.0911	0.5499	0.0042	0.0376	0.9405	-0.3311	0.0297	-0.3687	0.0119	-0.1861	0.7644	0.0365	0.9291	0.2226	0.4008
Nipsnap	Nipsnap	1635628_s_at	0.7616	0.0050	0.9315	0.0052	0.8665	0.0007	-0.1102	0.7774	-0.5869	0.0041	-0.4766	0.0062	-0.0731	0.9030	-0.5460	0.0365	-0.4730	0.0679
---	---	1635629_at	0.1308	0.6307	-0.1417	0.5244	0.1171	0.4592	0.1882	0.7556	0.2702	0.3044	0.0820	0.7767	0.0297	0.9816	0.0576	0.8993	0.0279	0.9471
spir	spire	1635630_a_at	-0.1745	0.7285	-1.2561	0.0307	-1.2476	0.0100	-0.0450	0.9825	0.9261	0.0581	0.9711	0.0313	-0.1664	0.9291	-0.3097	0.6345	-0.1433	0.8549
rab3-GAP	rab3-GAP	1635631_a_at	0.3492	0.2130	0.1220	0.7149	0.1853	0.3192	0.1867	0.6327	0.6296	0.0062	0.4429	0.0177	0.1388	0.9095	0.3360	0.4391	0.1972	0.6815
CG18363	CG18363	1635632_at	-0.0192	0.9405	-0.0044	0.9826	0.0455	0.7807	-0.1086	0.7689	-0.0636	0.7290	0.0450	0.7981	-0.1343	0.8427	-0.0964	0.7786	0.0379	0.9211
bin3	bicoid-interacting	1635633_s_at	-1.1329	0.0544	-0.4848	0.5958	-1.4665	0.0033	-0.0880	0.8604	0.2170	0.2257	0.3050	0.0610	0.8779	0.7464	0.7847	0.5082	-0.0932	0.9543
CG7172	CG7172	1635634_at	-0.0817	0.6576	0.6718	0.1051	0.3968	0.1223	-0.2404	0.6579	-0.7435	0.0117	-0.5031	0.0377	0.1780	0.7997	0.2034	0.4991	0.0253	0.9512
Msr-110	Engrailed nuclear	1635635_a_at	-0.5071	0.0967	-0.6162	0.0157	-1.0584	0.0001	-0.0222	0.9734	-0.0637	0.7436	-0.0415	0.8249	0.3369	0.7215	-0.1904	0.6614	-0.5273	0.2056
to	takeout	1635636_at	-1.0830	0.2578	-0.2533	0.9504	-3.1898	0.0009	-1.0569	0.3405	-1.2149	0.0611	-0.1580	0.8284	1.7876	0.8202	-0.4366	0.9326	-2.2242	0.5073
CG9773	CG9773	1635637_at	0.1178	0.7262	0.3030	0.3961	0.9031	0.0042	0.0563	0.9346	0.0002	0.9995	-0.0561	0.8053	-0.4588	0.7215	0.2257	0.7128	0.6845	0.2211
tomosyn	tomosyn	1635638_s_at	-0.0539	0.8856	-0.4756	0.2552	-0.4916	0.0064	0.1195	0.7589	0.7395	0.0017	0.6200	0.0022	0.0843	0.9677	0.2850	0.6310	0.2007	0.7513
Eip74EF	ecdysone inducibl	1635639_a_at	-0.0194	0.9841	0.0382	0.9752	-0.0482	0.9144	-0.6759	0.0976	-1.0420	0.0018	-0.3660	0.0881	-0.4231	0.9127	-0.7840	0.5855	-0.3609	0.8332
---	---	1635640_at	0.0312	0.8495	0.0384	0.8537	0.2892	0.1623	0.1538	0.7432	0.0915	0.7012	-0.0624	0.7871	-0.2914	0.7136	-0.0869	0.8471	0.2045	0.5612
---	---	1635641_at	0.0710	0.7503	0.6014	0.2434	0.4943	0.1691	0.1246	0.7762	-0.0879	0.6741	-0.2124	0.1941	0.1424	0.9421	0.3279	0.6118	0.1855	0.8003
CG31082	CG31082	1635642_at	0.0357	0.8977	0.2551	0.4417	-0.4134	0.0854	-0.3450	0.3616	0.0333	0.9114	0.3783	0.0536	0.2957	0.7946	0.2713	0.5911	-0.0244	0.9739
Rab23	Rab23	1635643_at	0.2034	0.7632	-0.6833	0.3389	-1.5766	0.0311	-0.9961	0.0492	0.9005	0.0082	1.8966	0.0002	-0.1859	0.9717	0.0661	0.9783	0.2520	0.8915
CG33203	CG33203	1635644_at	-1.9735	0.0020	-1.8594	0.0029	-3.0749	0.0000	-0.8909	0.0270	-0.3085	0.1578	0.5824	0.0098	0.0694	0.9653	-0.2258	0.6259	-0.2951	0.5083
CG9319	CG9319	1635645_at	-0.0077	0.9774	-0.2830	0.1565	-0.0689	0.6572	-0.0674	0.8875	-0.1357	0.4125	-0.0684	0.6850	-0.3230	0.6749	-0.6475	0.0609	-0.3245	0.3097
---	---	1635646_at	0.0461	0.8062	-0.0330	0.7489	0.0384	0.8425	-0.1005	0.8449	0.0046	0.9876	0.1051	0.5776	-0.1166	0.8608	-0.0069	0.9911	0.1097	0.7155
CG2617	CG2617	1635647_at	0.5892	0.0230	0.1156	0.4801	0.4613	0.0567	-0.1918	0.7422	0.1947	0.4674	0.3865	0.0919	-0.5128	0.3646	-0.3387	0.2430	0.1741	0.5868
CG14490	CG14490	1635648_at	-0.8613	0.0280	-0.1415	0.3716	-0.3342	0.2284	0.1994	0.8388	-0.1990	0.6259	-0.3983	0.2258	0.0476	0.9761	0.0550	0.9276	0.0074	0.9914
CG8038	CG8038	1635649_at	-0.2523	0.2691	0.2507	0.0859	0.6402	0.0304	0.1875	0.7121	-0.4783	0.0445	-0.6658	0.0068	-0.1804	0.8546	-0.0533	0.9347	0.1271	0.7947
---	---	1635650_at	-0.1509	0.6244	-0.0445	0.7257	-0.0480	0.9053	-0.1415	0.8875	0.1047	0.8003	0.2462	0.4382	-0.0258	0.9762	0.0573	0.8377	0.0831	0.7262
CG30195	CG30195	1635651_at	-1.5737	0.0022	-0.6831	0.0159	-1.3963	0.0003	-0.3638	0.3068	-0.9265	0.0017	-0.5627	0.0093	-0.1465	0.8953	-0.4370	0.2764	-0.2905	0.4998
CG3585 /// DrepCG3585	CG3585	1635652_at	-0.2779	0.7661	-0.2424	0.7783	-0.3114	0.1368	0.1091	0.9422	0.6057	0.1551	0.4966	0.1935	0.2131	0.9589	0.5580	0.6694	0.3448	0.8152
---	---	1635653_at	-0.0085	0.9697	0.0383	0.7334	0.1442	0.3670	0.0888	0.8645	-0.0522	0.8210	-0.1410	0.4144	-0.0069	0.9938	-0.0059	0.9884	0.0009	0.9979
CG11679	CG11679	1635654_at	0.0677	0.6855	0.4044	0.1191	0.6136	0.0058	-0.0830	0.8578	-0.3043	0.0694	-0.2213	0.1339	-0.2696	0.7230	-0.0617	0.9016	0.2079	0.5492
cv-2	crossveinless 2	1635655_at	-0.6668	0.0341	0.4151	0.2453	0.2058	0.4595	-0.0287	0.9803	-0.5112	0.0762	-0.4825	0.0628	0.2208	0.8386	0.5303	0.2193	0.3095	0.5041
---	---	1635656_at	0.2513	0.1166	0.0042	0.9791	0.0228	0.9011	-0.0572	0.9130	0.1881	0.2567	0.2453	0.0964	-0.0578	0.9416	0.0210	0.9587	0.0788	0.7853
SK	small conductanc	1635657_s_at	0.1425	0.3363	-0.0773	0.5372	0.3707	0.1026	0.3640	0.3521	0.2805	0.2091	-0.0835	0.7276	-0.1106	0.8870	-0.1090	0.7530	0.0016	0.9979
CG3592 /// DmirCG3592 ///	CG3592	1635658_at	-0.0174	0.9170	0.0067	0.9671	-0.1428	0.4401	0.0445	0.9381	0.0119	0.9632	-0.0326	0.8687	-0.1315	0.7726	-0.1283	0.5297	0.0032	0.9924
crn	yolky	1635659_at	-0.3405	0.2507	0.2013	0.3294	0.7117	0.0032	0.3214	0.2896	-0.0953	0.6319	-0.4167	0.0171	-0.2037	0.8439	0.3550	0.3981	0.5587	0.2090
---	---	1635660_at	0.0576	0.7416	-0.0433	0.7634	0.0310	0.8592	0.0301	0.9647	-0.1060	0.5873	-0.1361	0.4179	0.0924	0.8869	0.0240	0.9507	-0.0684	0.8221
CG14777	CG14777	1635661_at	-0.5962	0.0169	-1.1154	0.0162	-0.6449	0.0026	0.0470	0.9346	0.4913	0.0108	0.4443	0.0103	-0.3398	0.6531	0.1138	0.7521	0.4536	0.1591
CG32468	CG32468	1635662_at	0.0876	0.6507	0.0516	0.8071	-0.0403	0.8437	-0.0861	0.8732	-0.0403	0.8710	0.0458	0.8298	0.0533	0.9498	-0.0181	0.9650	-0.0714	0.8202
CG8320	CG8320	1635663_at	1.2837	0.0144	0.3727	0.0974	0.2723	0.3752	0.2254	0.7977	0.8580	0.0242	0.6327	0.0516	0.2903	0.7215	-0.1153	0.7837	-0.4055	0.2497
Bzd	Buzidau	1635664_at	-0.3245	0.1224	-0.6708	0.0186	-0.7554	0.0034	0.0815	0.8405	0.2952	0.0555	0.2138	0.1126	0.1929	0.7810	-0.1119	0.7544	-0.3047	0.3206
Tdc1	Tyrosine decarbo	1635665_at	-0.7574	0.6833	-3.1662	0.0014	-3.4340	0.0002	-0.4054	0.7021	0.5973	0.2182	1.0027	0.0289	0.1510	0.9862	-1.7786	0.3778	-1.9295	0.3621
His2B:CG17949 /// His2B:C	His2B:CG33910 //	1635666_at	0.0075	0.9952	-2.8803	0.0395	-1.8305	0.0146	0.5464	0.3775	2.9350	0.0002	2.3886	0.0002	-0.5066	0.9088	-0.0663	0.9823	0.4403	0.8222

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31108	CG31108	1635667_at	0.2904	0.1042	0.1860	0.6689	0.1365	0.4437	-0.1813	0.6990	0.1685	0.4522	0.3498	0.0716	-0.1280	0.9089	0.1922	0.6578	0.3202	0.4301
thoc5	thoc5	1635668_at	0.4198	0.0274	0.0876	0.8686	-0.1845	0.2710	-0.0039	0.9956	0.6557	0.0037	0.6596	0.0022	0.2000	0.8751	0.2692	0.6152	0.0693	0.9197
mirr	iroquois	1635669_a_at	0.4930	0.0408	0.0572	0.8281	0.0296	0.9118	0.1399	0.8029	0.2259	0.3205	0.0860	0.7189	0.1448	0.8956	-0.2977	0.4794	-0.4425	0.2964
CG11601	CG11601	1635670_at	-0.0890	0.7179	-0.2008	0.4099	-0.1382	0.3434	0.0198	0.9759	-0.2003	0.2149	-0.2201	0.1269	-0.0437	0.9742	-0.2605	0.4355	-0.2168	0.5391
CG33290	CG33290	1635671_at	0.1439	0.4970	-0.0420	0.6778	0.0383	0.8545	-0.2261	0.5683	-0.1277	0.5608	0.0984	0.6373	-0.0765	0.9309	-0.0918	0.8018	-0.0153	0.9726
CG34411	CG32637	1635672_at	-0.1761	0.6308	0.0598	0.6424	-0.0002	0.9991	-0.0975	0.9112	-0.3735	0.1750	-0.2760	0.2688	0.0126	0.9916	-0.0424	0.9238	-0.0550	0.8880
---	---	1635673_s_at	0.0019	0.9954	-0.0583	0.5884	0.1982	0.4190	0.1371	0.8381	0.0810	0.7928	-0.0561	0.8473	-0.2253	0.7485	-0.1275	0.7058	0.0978	0.7827
CG6901	CG6901	1635674_at	-0.0235	0.9170	0.0307	0.8893	0.0536	0.7313	-0.0215	0.9732	-0.0269	0.9019	-0.0054	0.9785	0.0125	0.9938	-0.0720	0.8868	-0.0845	0.8523
scro	scarecrow	1635675_at	-0.1477	0.4958	-0.0338	0.8422	0.1084	0.6053	0.0851	0.9149	0.0366	0.9167	-0.0484	0.8682	-0.1079	0.9238	-0.0232	0.9702	0.0847	0.8641
---	---	1635676_at	-0.1085	0.5880	0.1153	0.3576	-0.0019	0.9945	-0.0462	0.9311	-0.1647	0.2963	-0.1185	0.4177	0.1631	0.8049	0.2651	0.3259	0.1020	0.7492
sty	Sprouty	1635677_a_at	-1.5582	0.0175	-1.8556	0.0582	-2.1313	0.0001	-0.1168	0.9110	0.7523	0.0303	0.8691	0.0102	0.1940	0.9309	0.4537	0.5509	0.2596	0.7567
CG11299	CG11299	1635678_at	0.0144	0.9658	0.9203	0.0200	1.0853	0.0158	-0.3854	0.5183	-1.5564	0.0011	-1.1710	0.0021	-0.4922	0.6551	-0.4914	0.2489	0.0008	0.9992
lab	Labial	1635679_at	-0.0161	0.9277	0.0437	0.7352	0.2215	0.2530	0.1048	0.8791	-0.0400	0.9013	-0.1448	0.5256	-0.1259	0.8145	0.0362	0.9162	0.1621	0.4815
CG30486	CG30486	1635680_at	0.1140	0.6625	-0.0528	0.7054	0.3726	0.0677	0.2561	0.3228	0.1213	0.4373	-0.1348	0.3272	-0.0588	0.9460	0.0121	0.9799	0.0709	0.8307
CG4615	CG4615	1635681_at	-0.4725	0.0646	-0.3955	0.0456	-0.4551	0.0258	-0.4084	0.2435	-0.0524	0.8446	0.3560	0.0583	-0.3516	0.6695	0.1233	0.7544	0.4749	0.1760
Sucb	Sucb	1635682_at	0.0685	0.7632	-0.0887	0.5918	0.0355	0.8668	-0.0059	0.9950	-0.0947	0.6578	-0.0888	0.6494	-0.2037	0.8160	-0.2874	0.4258	-0.0837	0.8600
CG6136	CG6136	1635683_at	-1.8164	0.0072	-1.3103	0.0614	-1.4197	0.0005	0.0223	0.9922	-0.6636	0.1684	-0.6859	0.1122	0.0082	0.9964	-0.3581	0.3620	-0.3662	0.3720
unc-13	lethal (4) ry16	1635684_a_at	-1.3531	0.0010	-2.1735	0.0067	-2.3744	0.0001	-0.3431	0.5539	0.0091	0.9834	0.3522	0.1872	-0.2124	0.8692	-0.8130	0.1199	-0.6006	0.2614
---	---	1635685_at	0.2454	0.3378	0.1691	0.4154	0.1898	0.2495	0.0557	0.9603	-0.0649	0.8675	-0.1206	0.6966	0.1977	0.6749	0.1003	0.6394	-0.0974	0.6463
CG14110	CG14110	1635686_at	0.0349	0.8846	0.0139	0.8990	-0.0168	0.9250	0.0642	0.9297	-0.0348	0.9060	-0.0991	0.6528	0.0596	0.9057	-0.0681	0.7504	-0.1277	0.4957
CG8490	CG8490	1635687_at	0.6425	0.0417	0.7745	0.0128	0.7357	0.0015	0.0636	0.9116	-0.1286	0.5009	-0.1922	0.2366	0.1940	0.8042	0.4111	0.1925	0.2171	0.5252
CG15210	CG15210	1635688_at	0.5169	0.1591	1.5804	0.0179	2.4397	0.0024	0.5053	0.1879	-0.0690	0.8099	-0.5743	0.0135	-0.4452	0.8553	0.9854	0.3120	1.4306	0.1779
Lcp2	cuticle p. II-	1635689_at	0.1118	0.5706	0.0524	0.6840	0.1906	0.3539	0.0771	0.8903	0.1562	0.4196	0.0792	0.6873	-0.0039	0.9964	0.0070	0.9872	0.0110	0.9740
---	---	1635690_at	0.4260	0.0228	0.2070	0.3433	0.5295	0.0135	0.0233	0.9744	0.0560	0.7963	0.0327	0.8762	-0.0448	0.9409	0.0099	0.9758	0.0548	0.8138
br	broad complex	1635691_at	-0.3164	0.3318	-2.2164	0.0027	-1.3816	0.0373	0.4802	0.7225	1.7604	0.0110	1.2802	0.0268	-0.4253	0.6898	-0.2249	0.6312	0.2004	0.6728
Treh	trehalase	1635692_s_at	-0.9407	0.0013	-0.4329	0.3955	-1.2834	0.0034	-0.6539	0.0414	-0.9994	0.0007	-0.3455	0.0370	0.1661	0.9405	-0.4482	0.5312	-0.6143	0.3831
CG32946	CG32946	1635693_at	0.1888	0.4879	-0.1154	0.4725	-0.1685	0.3153	-0.1242	0.7507	-0.0395	0.8586	0.0847	0.6238	-0.0110	0.9946	-0.1328	0.7787	-0.1218	0.7923
CG12907	CG12907	1635694_at	0.0409	0.7921	-0.0564	0.5985	0.1882	0.3749	0.1885	0.5932	0.1390	0.4559	-0.0495	0.8053	-0.2432	0.7663	-0.1091	0.8070	0.1342	0.7378
Sdic3	Sdic:CG32823	1635695_at	0.2098	0.3638	0.3464	0.1551	0.1223	0.5196	-0.0977	0.8856	0.0075	0.9823	0.1052	0.6588	0.1407	0.8461	0.1455	0.6586	0.0048	0.9924
---	---	1635696_s_at	0.6441	0.4520	-0.2864	0.7377	-1.2663	0.0236	0.0604	0.9745	1.2683	0.0146	1.2079	0.0112	0.8527	0.7739	0.1080	0.9593	-0.7447	0.5871
---	---	1635697_at	0.0806	0.6520	-0.0643	0.7373	0.0865	0.6827	0.0835	0.8987	0.0298	0.9214	-0.0537	0.8259	-0.0157	0.9898	-0.0384	0.9277	-0.0227	0.9523
---	---	1635698_at	0.0003	0.9994	0.0407	0.7730	0.0115	0.9629	0.0571	0.9201	0.0549	0.7966	-0.0022	0.9919	-0.0154	0.9885	0.0152	0.9684	0.0306	0.9237
---	---	1635699_at	-0.0349	0.8182	0.2328	0.3394	0.3442	0.1266	-0.1038	0.8493	-0.0683	0.7767	0.0355	0.8804	-0.0274	0.9848	0.2023	0.5509	0.2297	0.4932
CG31792	CG31792	1635700_at	0.1452	0.6875	-0.0102	0.9569	-0.1978	0.3248	0.0936	0.9242	0.1388	0.6798	0.0452	0.8991	0.0016	0.9994	-0.2527	0.3236	-0.2543	0.3462
GstD3	Glutathione S tran	1635701_at	-0.8721	0.0632	-0.9600	0.0369	-1.3799	0.0002	-0.4849	0.2411	-0.1150	0.6800	0.3699	0.0909	0.0644	0.9764	-0.2725	0.6281	-0.3369	0.5403
Caf1-180	Caf1-180	1635702_a_at	0.1690	0.3959	-0.0756	0.7341	-0.0209	0.9474	-0.2127	0.5038	0.2748	0.1028	0.4875	0.0067	-0.2521	0.7707	0.1258	0.7824	0.3779	0.3162
l(3)82Fd	late puff gene at 8	1635703_s_at	-0.5828	0.1491	0.7537	0.0917	-0.5189	0.0392	-0.3041	0.5319	-0.4149	0.1023	-0.1108	0.6769	0.9431	0.3800	0.6978	0.2236	-0.2453	0.7149
CG1622	CG1622	1635704_at	-0.0745	0.6818	-0.0769	0.8709	0.1190	0.6951	-0.0128	0.9834	0.1028	0.4871	0.1155	0.3758	-0.3079	0.8378	0.0407	0.9682	0.3486	0.5995
CG12858	CG12858	1635705_at	-0.2834	0.2071	-0.0604	0.6265	-0.1254	0.5303	-0.1151	0.8340	-0.2139	0.3086	-0.0988	0.6441	-0.1577	0.8236	-0.1002	0.7842	0.0576	0.8874
mRpL20	mitochondrial ribo	1635706_at	-0.0127	0.9732	0.1541	0.6342	-0.1665	0.2530	-0.1111	0.8738	0.0337	0.9214	0.1449	0.5428	0.0110	0.9962	0.0564	0.9450	0.0454	0.9486
---	---	1635707_at	0.0614	0.6997	0.1221	0.2995	-0.0283	0.8773	-0.0368	0.9550	-0.0237	0.9231	0.0131	0.9536	0.1032	0.8349	0.0212	0.9496	-0.0820	0.7264
CG14803	CG14803	1635708_at	-0.0853	0.6234	-0.0986	0.8608	0.6976	0.0069	-0.0308	0.9803	-0.2059	0.5267	-0.1752	0.5617	-0.6620	0.6483	-0.1359	0.8695	0.5261	0.3808
CG7359	CG7359	1635709_at	1.3109	0.0021	0.7794	0.0222	0.6453	0.0120	-0.1251	0.7850	1.1358	0.0005	1.2608	0.0002	-0.0188	0.9914	0.7160	0.0689	0.7349	0.0828
CG15470	CG15470	1635710_at	0.0904	0.6902	0.0034	0.9864	0.2643	0.1479	0.0668	0.9068	-0.1399	0.4672	-0.2067	0.2095	0.0246	0.9829	0.0323	0.9402	0.0077	0.9852
Ank	ankyrin	1635711_s_at	-0.5218	0.0211	-0.4265	0.2421	-1.1112	0.0008	-0.3255	0.2856	-0.0337	0.8897	0.2918	0.0710	0.2368	0.7956	0.1315	0.7819	-0.1053	0.8274
CG2247	CG2247	1635712_at	0.7521	0.0088	0.1408	0.6888	-0.1208	0.6258	-0.2405	0.5008	0.6006	0.0074	0.8411	0.0010	0.0404	0.9816	0.0318	0.9628	-0.0087	0.9902
CG13185	CG13185	1635713_at	1.3509	0.0151	0.4743	0.4456	0.4903	0.1718	0.2616	0.6929	1.6029	0.0009	1.3413	0.0011	0.1264	0.9717	0.6969	0.4410	0.5705	0.5511
fdl	fused lobes	1635714_s_at	-0.2674	0.6641	0.2864	0.1821	0.2186	0.3359	-0.0805	0.8830	-0.0262	0.9204	0.0544	0.7952	0.0243	0.9939	0.5320	0.4539	0.5078	0.4876
His4:CG31611 /// His4:CG3	His4:CG31611 ///	1635715_at	1.2286	0.4379	-2.5889	0.1950	-0.8307	0.4736	0.6364	0.4728	3.4050	0.0003	2.7685	0.0004	-1.0319	0.9011	-0.4477	0.9263	0.5842	0.8889
beat-VII /// CG14248	beat-VII /// CG142	1635716_at	-2.3730	0.0016	-2.9063	0.0052	-2.2201	0.0000	0.4659	0.5832	0.8875	0.0456	0.4216	0.2798	-0.1235	0.8611	0.0303	0.9485	0.1538	0.6186
---	---	1635717_x_at	0.2396	0.5045	0.1361	0.6840	0.0480	0.8729	-0.1358	0.8817	0.2282	0.4845	0.3640	0.1903	0.0396	0.9871	0.0925	0.9151	0.0529	0.9463
bur	Enhancer of GMR	1635718_at	1.0829	0.0136	0.5432	0.0536	0.2382	0.2372	0.0886	0.8666	0.8711	0.0013	0.7824	0.0013	0.4401	0.7220	0.2973	0.5941	-0.1428	0.8287
Eig71Ef	Gene V	1635719_at	0.2156	0.2210	-0.2101	0.1193	0.0615	0.7963	0.0624	0.9305	0.2055	0.3341	0.1431	0.4726	0.0037	0.9970	-0.1114	0.6853	-0.1151	0.6677
---	---	1635720_at	-0.0818	0.7019	0.1074	0.3636	0.1168	0.5672	-0.1535	0.8196	-0.1679	0.5547	-0.0144	0.9658	-0.0374	0.9643	0.0011	0.9991	0.0384	0.9023
CG15383	CG15383	1635721_at	0.1775	0.2942	0.1665	0.3083	0.3770	0.0306	0.2424	0.4979	0.1578	0.4201	-0.0846	0.6670	0.0831	0.8967	0.0650	0.8391	-0.0181	0.9582
---	---	1635722_at	0.1533	0.2800	-0.0270	0.8391	0.0384	0.8302	-0.0516	0.9087	0.0792	0.6189	0.1308	0.3180	0.0053	0.9964	-0.0507	0.9051	-0.0559	0.8827
CG33129	CG33129	1635723_s_at	0.0704	0.7394	1.2420	0.0224	1.3239	0.0013	-0.2879	0.6822	-0.6952	0.0400	-0.4073	0.1628	-0.4609	0.5754	0.5180	0.1512	0.9789	0.0433
CG14292 /// DereCG14292	CG14292 /// GA1	1635724_at	-0.8396	0.6068	-0.2167	0.8452	0.0033	0.9885	0.5109	0.9017	-1.3484	0.3200	-1.8593	0.1193	0.4222	0.8968	-0.6893	0.5925	-1.1115	0.3719
unc-115	unc-115	1635725_a_at	-0.6308	0.2560	0.1804	0.7523	-0.2813	0.1036	-0.1364	0.7230	-0.3305	0.0609	-0.1941	0.2112	0.3411	0.8814	0.4400	0.6414	0.0989	0.9351
---	---	1635726_at	0.2619	0.0984	-0.0227	0.8319	-0.0015	0.9942	0.1211	0.7949	0.1037	0.6234	-0.0174	0.9412	0.1605	0.7142	-0.0583	0.8038	-0.2187	0.2497
Cyp4g15	Cyp4g15	1635727_a_at	-0.1547	0.7468	-0.1715	0.8230	-0.0614	0.8503	0.0820	0.9507	-0.8003	0.0366	-0.8823	0.0155	0.0827	0.9831	-0.8298	0.3571	-0.9126	0.3323
ps	pasilla	1635728_a_at	0.8475	0.0723	0.7099	0.1007	-0.4388	0.4086	-0.3185	0.5531	0.8639	0.0080	1.1824	0.0012	0.9936	0.6557	0.7921	0.3760	-0.2015	0.8680
Ptp61F	protein tyrosine pl	1635729_a_at	1.3441	0.0160	1.3168	0.0241	1.1268	0.0284	-0.5646	0.2753	-0.0574	0.8897	0.5073	0.0663	-0.2812	0.9101	0.0001	1.0000	0.2813	0.7931
CG30069	CG30069	1635730_at	-0.8986	0.0485	-1.2277	0.0831	-1.5458	0.0001	0.1748	0.8705	0.9270	0.0216	0.7522	0.0321	0.5595	0.6898	0.5651	0.3107	0.0056	0.9954
CG6785	CG6785	1635731_at	0.2392	0.4119	0.0574	0.6205	0.4580	0.0172	0.0016	0.9988	-0.2293	0.3185	-0.2309	0.2589	-0.1409	0.8541	-0.4107	0.1801	-0.2698	0.4017
CG6915 /// DmirCG6915	CG6915	1635732_at	-1.0258	0.0298	-1.3244	0.0148	-1.4445	0.0096	0.3379	0.8424	0.9250	0.1410	0.5870	0.3056	0.2300	0.7498	0.2568	0.3994	0.0268	0.9510
CG31439	CG31439	1635733_x_at	0.2804	0.3458	0.1723	0.4113	-0.1055	0.8131	-0.1475	0.8732	0.0353	0.9393	0.1828	0.5649	0.2449	0.8306	0.1894	0.7346	-0.0555	0.9343
---	---	1635734_at	0.0912	0.6142	-0.0079	0.9687	-0.0663	0.6814	-0.1269	0.7749	0.0148	0.9569	0.1418	0.4143	-0.0421	0.9701	-0.1106	0.7439	-0.0684	0.8548
---	---	1635735_at	-0.0141	0.9417	-0.0776	0.6940	-0.3952	0.0604	-0.1385	0.7135	0.1198	0.5094	0.2582	0.0925	0.1168	0.9186	0.0874	0.8764	-0.0294	0.9588

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG5157	CG5157	1635736_at	0.1906	0.2254	0.0727	0.7879	0.1358	0.3842	-0.2036	0.5363	-0.1137	0.5366	0.0899	0.6034	-0.1264	0.8472	-0.0852	0.8084	0.0412	0.9121
CG6701	CG6701	1635737_at	0.3468	0.0543	-0.8114	0.0174	-0.1603	0.3718	0.4059	0.1749	0.9453	0.0009	0.5394	0.0058	-0.2300	0.7606	-0.1167	0.7569	0.1134	0.7558
Cng	CNG-channel	1635738_at	-0.7587	0.0257	0.4297	0.1276	0.1523	0.4220	-0.6514	0.1683	-1.6313	0.0006	-0.9799	0.0032	-0.1225	0.8513	-0.1010	0.7543	0.0215	0.9553
CG33960	CG33960	1635739_at	-0.0203	0.9167	-0.0251	0.8073	0.0941	0.5523	0.1574	0.6615	0.1178	0.5196	-0.0396	0.8420	-0.1521	0.8016	-0.0574	0.8740	0.0948	0.7449
Hmu	hemomucin	1635740_at	0.2677	0.5639	0.6521	0.0257	0.9279	0.0015	0.2679	0.5956	0.0490	0.8873	-0.2189	0.3512	0.0529	0.9812	0.4799	0.3022	0.4269	0.3830
Patsas	Patsas	1635741_a_at	-0.0561	0.7739	-0.4103	0.1849	-0.0650	0.7701	0.1239	0.8449	0.1944	0.4201	0.0705	0.7869	-0.3242	0.7230	0.0615	0.9207	0.3856	0.3444
Cad74A	Cad74A	1635742_s_at	-0.2084	0.1932	-0.3260	0.0452	-0.1708	0.2752	-0.0499	0.9251	0.1342	0.4133	0.1841	0.1948	-0.2626	0.5700	-0.1883	0.3534	0.0744	0.7560
Vang	Strabismus	1635743_at	0.3497	0.3330	0.4840	0.2131	-0.1057	0.5544	-0.0640	0.9425	0.0954	0.7511	0.1594	0.5152	0.4455	0.6710	0.1689	0.7317	-0.2766	0.5358
CG9342 /// DyakCG9342	CG9342	1635744_at	2.1214	0.0183	0.8158	0.3819	2.0953	0.0001	0.3988	0.4821	1.1736	0.0030	0.7748	0.0118	-0.9437	0.7464	-0.1369	0.9451	0.8068	0.5334
CG4389	CG4389	1635745_a_at	0.5279	0.0230	0.1479	0.6028	0.5480	0.0058	0.0414	0.9380	0.0772	0.6643	0.0358	0.8431	-0.2649	0.7450	-0.2765	0.4172	-0.0116	0.9841
---	---	1635746_at	-0.1309	0.4772	0.0350	0.7758	0.1053	0.6520	-0.0371	0.9507	-0.0812	0.6666	-0.0441	0.8135	-0.0466	0.9683	0.0341	0.9460	0.0807	0.8372
mei-41	meiotic 41	1635747_at	0.2765	0.6656	0.2694	0.2915	0.8979	0.0017	-0.1040	0.9098	-0.1920	0.5391	-0.0880	0.7872	-0.1696	0.8692	0.0655	0.9178	0.2351	0.5919
CG15322	CG15322	1635748_at	0.1148	0.5188	0.2488	0.2634	0.2005	0.5869	-0.0379	0.9518	-0.0829	0.6776	-0.0450	0.8207	0.1275	0.9238	0.0493	0.9451	-0.0782	0.8982
---	---	1635749_at	0.3661	0.0723	0.0895	0.5365	0.0796	0.6864	-0.1079	0.8140	0.1320	0.4858	0.2399	0.1360	-0.1042	0.8472	-0.0722	0.8005	0.0320	0.9176
Gr93a	Gustatory recepto	1635750_at	0.1520	0.3459	-0.0150	0.8984	0.1042	0.5324	0.1710	0.5311	0.2290	0.1074	0.0580	0.6995	-0.1911	0.6868	-0.0738	0.7457	0.1172	0.5681
CG32069	CG32069	1635751_at	0.2741	0.3035	0.8003	0.1103	0.6233	0.0105	0.3493	0.4024	0.5045	0.0371	0.1553	0.4855	0.3358	0.7644	0.9604	0.0637	0.6246	0.2090
CG4520	CG4520	1635752_at	0.3252	0.2156	0.2506	0.4214	0.2068	0.3236	-0.0607	0.9436	0.0442	0.8943	0.1048	0.6821	0.1091	0.9368	0.1830	0.7105	0.0739	0.9025
CG12362	CG12362	1635753_s_at	0.2725	0.1077	0.1384	0.3233	0.1033	0.6627	0.0088	0.9931	0.0365	0.8920	0.0277	0.9084	0.1147	0.8427	-0.0120	0.9774	-0.1267	0.6209
tgo	PRD gene 7	1635754_at	-0.5662	0.0307	-0.2037	0.5483	-0.5970	0.0040	-0.2363	0.5199	-0.2507	0.1921	-0.0144	0.9534	0.1242	0.9061	0.0361	0.9525	-0.0881	0.8578
CG10741	CG10741	1635755_a_at	-1.2058	0.1746	0.2720	0.8226	0.2483	0.4751	0.0140	0.9922	-1.0698	0.0040	-1.0838	0.0023	0.1487	0.9816	0.4382	0.8220	0.2894	0.8880
CG32039	CG32039	1635756_at	-0.6423	0.0492	-0.2084	0.5089	-0.6867	0.0021	-0.1686	0.7982	-0.3935	0.1395	-0.2248	0.3608	0.2770	0.7215	0.0713	0.8782	-0.2056	0.5490
laza	CG11440	1635757_at	2.5221	0.0012	1.9716	0.0297	3.0414	0.0001	0.3572	0.6338	-0.3156	0.4047	-0.6728	0.0467	-0.6852	0.7464	-0.8921	0.3087	-0.2069	0.8613
TBPH	TBPH	1635758_s_at	-0.0886	0.6800	-0.0292	0.8803	-0.5621	0.0066	-0.1803	0.5068	0.0974	0.5254	0.2776	0.0377	0.3748	0.6660	0.2244	0.5404	-0.1505	0.7020
CG13775	CG13775	1635759_at	-0.1001	0.6807	-0.2430	0.3690	0.0282	0.9162	0.1692	0.8791	0.1996	0.6385	0.0304	0.9496	-0.0441	0.9591	-0.0578	0.8685	-0.0137	0.9715
Orc2	lethal(3)88Ab	1635760_at	-0.1543	0.4701	-0.2285	0.7118	-0.5757	0.1519	-0.6255	0.0998	0.4548	0.0477	1.0803	0.0007	-0.0729	0.9841	0.2705	0.7963	0.3435	0.7138
wdn	pourquoi-pas?	1635761_at	0.6826	0.0127	0.3705	0.4692	0.6491	0.0275	0.1744	0.6572	0.1684	0.3832	-0.0060	0.9799	-0.1328	0.9441	-0.0806	0.9291	0.0522	0.9485
CG6401	CG6401	1635762_at	-0.0144	0.9613	-0.0008	0.9969	0.2880	0.2199	0.2763	0.5978	0.4213	0.1101	0.1451	0.5827	-0.1451	0.8827	0.2709	0.4797	0.4160	0.2834
CG13226	CG13226	1635763_at	-0.1307	0.5642	-0.9426	0.0311	-0.7195	0.0780	0.2233	0.8640	0.9066	0.0550	0.6834	0.0985	-0.0725	0.9056	-0.1263	0.5864	-0.0538	0.8486
CG14696 /// DvarCG14696	CG14696	1635764_at	-0.6983	0.0585	-0.3071	0.2489	0.2222	0.4448	0.1123	0.8707	-0.4657	0.0579	-0.5780	0.0155	-0.2261	0.8236	-0.0760	0.9068	0.1501	0.7587
CG31161	CG31161	1635765_at	-0.0638	0.7663	0.0755	0.5995	-0.1957	0.3193	0.0020	0.9988	-0.1994	0.4921	-0.2013	0.4389	0.2168	0.6749	0.0987	0.6757	-0.1181	0.6051
Fs(2)Ket	importin beta	1635766_at	2.1109	0.0013	0.5808	0.2634	-0.5883	0.0151	-0.2854	0.5997	2.3803	0.0001	2.6657	0.0000	0.7297	0.5461	0.8053	0.1441	0.0755	0.9230
CG4313	CG4313	1635767_at	-1.6916	0.0004	-0.2251	0.2572	-1.3532	0.0009	-0.9242	0.0457	-1.6380	0.0004	-0.7137	0.0087	0.0884	0.9400	-0.3057	0.3903	-0.3941	0.2883
CG4670	CG4670	1635768_at	-1.5378	0.0075	-2.1411	0.0091	-1.6479	0.0001	0.2021	0.5793	0.2186	0.2415	0.0166	0.9433	-0.0986	0.9679	-0.2848	0.6974	-0.1862	0.8182
CG8773	CG8773	1635769_at	0.0924	0.7578	-0.0260	0.8136	-0.0254	0.8989	0.0286	0.9688	0.0348	0.8920	0.0062	0.9793	-0.0857	0.9342	-0.2065	0.5538	-0.1208	0.7527
CG31097	CG31097	1635770_at	0.1478	0.7329	0.0550	0.8525	-0.2580	0.1715	-0.1507	0.8794	-0.0330	0.9485	0.1177	0.7505	-0.0409	0.9774	-0.2436	0.4835	-0.2027	0.5757
CG32306	CG32306	1635771_a_at	-0.6883	0.0125	-1.4587	0.0407	-1.5706	0.0000	0.0920	0.8815	0.5932	0.0122	0.5012	0.0157	0.3189	0.7822	0.0184	0.9846	-0.3004	0.5691
CG9287	CG9287	1635772_at	0.0873	0.6287	-0.0278	0.7957	0.0848	0.6435	0.1697	0.6835	0.2697	0.1613	0.1001	0.6086	0.1904	0.7726	0.0307	0.9462	-0.1597	0.5972
CG10144	CG10144	1635773_at	-0.0666	0.7735	0.3116	0.1648	0.2119	0.4291	0.1301	0.8254	0.3467	0.1279	0.2166	0.2955	0.1393	0.9064	0.6683	0.1199	0.5290	0.2282
CG15835	CG15835	1635774_at	0.8056	0.0163	0.4943	0.1648	-0.1045	0.6188	-0.2959	0.5067	0.8118	0.0051	1.1077	0.0008	0.1489	0.8903	0.3147	0.4355	0.1658	0.7156
CG15207	CG15207	1635775_at	0.2632	0.2636	0.1046	0.4435	0.2018	0.1797	0.1237	0.7915	0.1191	0.5622	-0.0046	0.9850	-0.0798	0.9093	-0.0879	0.7723	-0.0081	0.9846
---	---	1635776_at	-0.0335	0.8706	0.0169	0.8925	-0.1573	0.3483	-0.0594	0.9099	0.0933	0.6117	0.1527	0.3145	-0.0833	0.8884	-0.0416	0.9027	0.0416	0.8921
CG11041	CG11041	1635777_at	0.0080	0.9791	-0.2568	0.1136	-0.3316	0.0810	-0.1025	0.7895	0.2276	0.1459	0.3301	0.0268	-0.0553	0.9611	-0.0124	0.9841	0.0429	0.9228
CG8771	CG8771	1635778_at	0.5994	0.0355	0.1460	0.7896	0.0612	0.7826	-0.0835	0.9247	0.6609	0.0205	0.7444	0.0074	-0.0240	0.9922	0.2844	0.6408	0.3084	0.6100
CG14888	CG14888	1635779_at	-0.0314	0.9017	0.0293	0.8786	0.0461	0.7815	0.0922	0.8507	-0.0678	0.7485	-0.1599	0.3255	0.1079	0.8650	0.1481	0.5805	0.0402	0.9086
---	---	1635780_at	0.3897	0.1261	0.0792	0.4151	0.3221	0.1679	0.0973	0.8773	0.0902	0.7201	-0.0071	0.9793	-0.1963	0.7506	-0.0495	0.9022	0.1469	0.6053
Ikb1	Ikb1	1635781_s_at	-0.5527	0.0893	-0.0184	0.8884	0.0014	0.9953	-0.1286	0.8143	-0.2544	0.2389	-0.1258	0.5535	-0.0169	0.9916	0.3657	0.3157	0.3826	0.3171
CG32692	CG32692	1635782_at	-0.4304	0.1009	-1.6999	0.0027	-1.5501	0.0022	0.2574	0.7958	1.2507	0.0082	0.9933	0.0139	0.2158	0.7215	0.0267	0.9487	-0.1891	0.4710
CG3119	CG3119	1635783_a_at	0.2533	0.4115	0.2505	0.2988	0.0701	0.8055	-0.1940	0.5079	-0.2672	0.0866	-0.0732	0.6474	0.3231	0.7387	0.0817	0.8929	-0.2414	0.5811
CG32230	CG32230	1635784_a_at	-0.0392	0.8707	0.3789	0.2554	0.1282	0.6366	-0.0359	0.9677	-0.7386	0.0074	-0.7027	0.0055	0.1371	0.8953	-0.2980	0.4421	-0.4351	0.2784
CG14368	CG14368	1635785_at	0.0826	0.6567	0.0841	0.5185	0.0888	0.5521	-0.0310	0.9538	-0.0861	0.5878	-0.0552	0.7231	-0.0005	0.9998	-0.0550	0.8605	-0.0545	0.8512
---	---	1635786_at	0.0313	0.8661	0.5187	0.0623	0.1808	0.4452	-0.0739	0.9036	-0.2874	0.1452	-0.2135	0.2288	0.2180	0.7215	0.1439	0.6011	-0.0741	0.8178
CG15630	CG15630	1635787_at	-0.3567	0.2793	0.0759	0.4203	0.3445	0.0836	-0.0116	0.9943	-0.8610	0.0230	-0.8494	0.0153	-0.0410	0.9589	-0.1207	0.6191	-0.0798	0.7612
SelR	methionine-R-sulfi	1635788_a_at	-0.9311	0.0023	-0.9419	0.0047	-1.2093	0.0002	-0.1571	0.6360	-0.0238	0.9171	0.1333	0.3770	0.0639	0.9457	-0.0199	0.9657	-0.0838	0.8112
CG6630	CG6630	1635789_at	-0.7666	0.0075	-0.4816	0.1155	-0.1656	0.5329	0.2018	0.8084	-0.0481	0.9174	-0.2499	0.4201	-0.1001	0.8510	-0.0041	0.9935	0.0961	0.6955
---	---	1635790_at	0.0335	0.8450	-0.0689	0.5619	0.0246	0.9053	0.1395	0.7235	0.1814	0.3124	0.0419	0.8376	-0.0011	0.9994	0.0437	0.8678	0.0448	0.8524
CG11470	CG11470	1635791_at	0.0978	0.5604	-0.0786	0.5660	-0.1456	0.3836	0.0799	0.8872	0.2022	0.2921	0.1223	0.5046	0.0796	0.9309	0.0596	0.8922	-0.0200	0.9638
Ela	elastin-like	1635792_a_at	2.3275	0.0032	1.9486	0.0411	3.6201	0.0000	0.9636	0.1395	0.6772	0.0869	-0.2864	0.4397	-0.4148	0.8202	0.3233	0.7058	0.7382	0.3439
CG11413	CG11413	1635793_at	0.1076	0.6356	0.0803	0.5118	0.1825	0.4407	0.0704	0.9271	-0.0911	0.7372	-0.1615	0.4588	0.0097	0.9947	-0.0457	0.9328	-0.0554	0.9057
CG12535	CG12535	1635794_a_at	0.1846	0.2025	0.2188	0.0952	0.1952	0.3873	-0.0182	0.9857	0.0971	0.7194	0.1153	0.6257	0.0904	0.8504	0.0985	0.6483	0.0080	0.9809
---	---	1635795_at	0.0320	0.9089	0.1075	0.7083	0.3096	0.1285	-0.0764	0.8794	-0.1334	0.4562	-0.0571	0.7622	-0.2433	0.7810	0.1198	0.8053	0.3631	0.3513
cad	caudal	1635796_a_at	-0.5165	0.8411	-2.2051	0.2218	-2.3629	0.0121	-0.5585	0.8462	1.2245	0.2480	1.7830	0.0628	-0.3192	0.9816	-0.5770	0.9085	-0.2578	0.9563
CG14982	CG14982	1635797_at	0.2832	0.0602	-0.0386	0.8343	-0.0023	0.9935	0.0999	0.8872	0.2076	0.3952	0.1076	0.6627	0.0346	0.9779	-0.2343	0.4302	-0.2689	0.3800
CG17364	CG17364	1635798_a_at	0.3255	0.1078	0.1239	0.4987	0.3425	0.0972	0.1677	0.7202	0.1245	0.5887	-0.0432	0.8609	0.0598	0.9465	0.0348	0.9353	-0.0250	0.9471
HP1b	HP1b	1635799_at	0.0132	0.9422	-0.1254	0.6559	0.2635	0.0895	0.0214	0.9683	0.0792	0.6095	0.0578	0.6964	-0.2564	0.7506	0.0347	0.9505	0.2912	0.4086
CG5431	CG5431	1635800_at	-0.6934	0.0830	-0.6688	0.1190	-1.0577	0.0069	-0.1880	0.7444	-0.0766	0.8058	0.1115	0.6728	-0.0123	0.9964	-0.1990	0.7992	-0.1867	0.8041
CG33673	CG33673	1635801_at	0.3461	0.1208	-0.0252	0.8389	0.3142	0.0602	0.2655	0.4356	0.3153	0.0920	0.0498	0.8132	-0.1372	0.8756	0.1487	0.7012	0.2858	0.4114
CG9279 /// DmirCG9279	CG9279	1635802_s_at	0.1963	0.4723	0.1874	0.2940	0.0545	0.7916	-0.0397	0.9639	-0.3138	0.1747	-0.2740	0.1861	0.1337	0.8541	-0.2827	0.3310	-0.4164	0.1883
bowl	bowel	1635803_s_at	-0.2494	0.6936	0.4393	0.4552	-0.2276	0.2169	0.2038	0.8162	0.0119	0.9816	-0.1919	0.5754	0.7660	0.7677	0.8338	0.4479	0.0678	0.9673
nito	One Twenty Two	1635804_at	-0.0600	0.8856	-0.1776	0.5811	-0.1517	0.5901	-0.2084	0.7482	0.0263	0.9508	0.2347	0.3688	-0.2922	0.8215	-0.3217	0.5698	-0.0295	0.9717

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG3045	CG3045	1635805_at	-0.5726	0.1319	-0.3779	0.4935	-0.6802	0.0383	0.2071	0.8362	0.9384	0.0213	0.7313	0.0373	0.5379	0.7423	0.9867	0.1445	0.4488	0.5380
Cpr78Ca	CG11310	1635806_at	0.1440	0.6371	0.2578	0.3525	0.1841	0.3501	0.0067	0.9956	0.0143	0.9698	0.0076	0.9799	-0.0268	0.9851	-0.0744	0.8734	-0.0477	0.9162
CG17739	CG17739	1635807_at	0.0685	0.8848	-0.6316	0.5822	0.2632	0.2290	0.1399	0.8402	0.5648	0.0361	0.4249	0.0692	-0.7140	0.7936	-0.1451	0.9376	0.5689	0.6457
CG8790	CG8790	1635808_s_at	0.4193	0.3584	-0.4390	0.2574	0.2491	0.2680	0.1542	0.5948	0.0673	0.6901	-0.0869	0.5469	-0.5198	0.7644	-0.7911	0.2627	-0.2712	0.7484
pip	pipe	1635809_at	0.1468	0.3675	-0.0296	0.8915	0.2145	0.2258	-0.0174	0.9838	0.0279	0.9200	0.0453	0.8427	-0.1147	0.8480	0.0260	0.9490	0.1407	0.5991
CG7806	CG7806	1635810_at	-0.0609	0.8218	-0.2798	0.1952	0.1476	0.4294	0.1912	0.7223	0.4245	0.0813	0.2333	0.2875	-0.2401	0.6832	0.1768	0.4729	0.4169	0.1190
rux	roughex	1635811_at	-0.2513	0.1266	-0.6163	0.0177	-0.4973	0.0691	0.3172	0.4567	0.5358	0.0279	0.2186	0.2960	0.0638	0.9467	0.1386	0.6614	0.0747	0.8380
CG16965	CG16965	1635812_at	1.0484	0.2573	-1.0844	0.3369	0.0620	0.8904	0.7495	0.3347	1.7989	0.0026	1.0494	0.0169	-0.4842	0.9238	-0.4910	0.8244	-0.0068	0.9983
mia	TAF6-like	1635813_at	-0.8304	0.0325	-0.9298	0.0151	-1.1575	0.0009	0.1718	0.8005	0.5127	0.0645	0.3409	0.1627	0.0954	0.9368	0.2551	0.5172	0.1597	0.7087
---	---	1635814_at	0.1050	0.6646	0.2184	0.2318	0.2716	0.1128	-0.0944	0.8830	-0.1441	0.5385	-0.0498	0.8435	-0.2373	0.7230	-0.1224	0.7054	0.1148	0.7246
zetaTry	zetaTrypsin	1635815_at	-0.3748	0.9250	-0.0652	0.5563	-0.3340	0.3176	-0.1609	0.9860	-2.4975	0.2497	-2.3366	0.2277	-0.0493	0.9970	-2.2766	0.4925	-2.2273	0.5067
---	---	1635816_at	0.2949	0.1938	-0.7976	0.0203	-0.2873	0.2529	0.2926	0.4979	1.0849	0.0013	0.7923	0.0029	-0.2610	0.7644	-0.1994	0.6109	0.0616	0.9031
CG11131	CG11131	1635817_at	0.2978	0.3112	0.3238	0.2221	0.5104	0.0151	-0.0890	0.9218	-0.0225	0.9556	0.0665	0.8301	0.1207	0.8680	0.0504	0.9085	-0.0704	0.8491
CG4713	CG4713	1635818_at	0.2234	0.1519	-0.1644	0.3798	-0.1158	0.6270	0.0727	0.8932	0.6014	0.0059	0.5287	0.0062	-0.0089	0.9952	0.2523	0.4843	0.2612	0.4742
sty	Sprouty	1635819_at	-1.6528	0.0162	-1.6473	0.0307	-2.5050	0.0000	-0.0006	0.9995	0.6710	0.0160	0.6716	0.0098	0.7062	0.7147	0.6660	0.4038	-0.0402	0.9751
---	---	1635820_at	-0.0363	0.8156	0.0878	0.4655	-0.0783	0.6811	-0.1144	0.7581	-0.1508	0.3539	-0.0364	0.8435	0.1083	0.8744	-0.0336	0.9390	-0.1419	0.6254
MED28	Mediator complex	1635821_at	0.4891	0.0659	-0.0324	0.9389	0.0864	0.6679	-0.1514	0.7735	0.3640	0.0991	0.5154	0.0171	-0.1001	0.9409	-0.0555	0.9341	0.0446	0.9394
eag	ether-a-go-go	1635822_at	0.3669	0.0343	0.1636	0.5158	0.4701	0.0105	0.1681	0.6224	0.1547	0.3721	-0.0134	0.9502	-0.1025	0.8802	-0.0519	0.8926	0.0507	0.8879
---	---	1635823_at	-0.0461	0.8844	0.0481	0.7265	0.1278	0.4027	-0.0837	0.9116	-0.0681	0.8193	0.0156	0.9589	-0.1192	0.8326	-0.0352	0.9233	0.0840	0.7560
Tip60	Tip60	1635824_at	0.2093	0.2716	0.4575	0.0432	0.3033	0.1066	-0.0521	0.9351	-0.0991	0.6399	-0.0470	0.8274	0.0477	0.9679	0.1865	0.5749	0.1388	0.6893
TFAM	mitochondrial tran	1635825_a_at	-0.0364	0.8503	0.8204	0.0100	0.8790	0.0563	0.1068	0.8875	-0.7341	0.0112	-0.8409	0.0036	0.0335	0.9898	0.2438	0.6910	0.2103	0.7390
CG9940 /// DmirCG9940	CG9940	1635826_s_at	0.2741	0.5038	0.3086	0.2871	0.4243	0.2347	-0.1148	0.9311	-0.2134	0.6197	-0.0986	0.8231	-0.2701	0.8814	-0.2813	0.7240	-0.0112	0.9924
Stat92E	marelle	1635827_s_at	0.8809	0.0103	0.6073	0.0233	0.6684	0.0018	-0.1980	0.5932	0.1386	0.4816	0.3365	0.0503	-0.1382	0.8510	0.0063	0.9928	0.1445	0.6607
Chd64	Chd64	1635828_at	-0.1905	0.2429	0.2078	0.2515	0.1471	0.4297	0.0472	0.9371	0.0833	0.6739	0.0362	0.8585	0.0956	0.8889	0.4759	0.0787	0.3804	0.1651
---	---	1635829_s_at	-0.1022	0.7040	-0.9060	0.0855	-1.1043	0.0147	0.0719	0.9669	0.9839	0.0368	0.9120	0.0319	0.1589	0.8967	0.3144	0.5042	0.1554	0.7727
CG8105	CG8105	1635830_at	0.1286	0.4955	-0.0354	0.7512	-0.0027	0.9909	-0.0282	0.9641	0.0750	0.6949	0.1033	0.5234	0.0537	0.9495	0.0376	0.9247	-0.0162	0.9659
CG14036	CG14036	1635831_at	0.7116	0.3743	-0.1203	0.8662	0.1754	0.7509	-0.0527	0.9603	0.7914	0.0118	0.8441	0.0053	-0.5342	0.8825	-0.3588	0.8492	0.1754	0.9271
CG12413	CG12413	1635832_at	0.0209	0.8977	0.0320	0.8068	0.2789	0.1000	-0.0511	0.9115	-0.0915	0.5589	-0.0404	0.8044	-0.1167	0.8692	0.1403	0.6497	0.2570	0.3791
CG4198	CG4198	1635833_at	-0.0827	0.8557	-0.9352	0.0582	-0.4388	0.3414	0.2591	0.8272	0.6387	0.1604	0.3796	0.3682	-0.2995	0.8202	-0.2562	0.6700	0.0433	0.9553
CG17169	CG17169	1635834_at	0.0982	0.5962	0.1006	0.4669	0.2275	0.2520	0.1262	0.7397	0.0708	0.7153	-0.0554	0.7624	0.0068	0.9950	-0.0351	0.9275	-0.0419	0.9016
CG30374	CG30374	1635835_at	0.1732	0.4054	0.0990	0.4570	0.3850	0.0653	0.0740	0.8877	0.0420	0.8552	-0.0321	0.8784	-0.1287	0.8222	0.0006	0.9996	0.1293	0.6149
CG8072	CG8072	1635836_at	0.2236	0.3573	-0.0481	0.6906	-0.0397	0.8492	0.2012	0.5724	0.3687	0.0480	0.1676	0.3111	-0.0033	0.9986	0.1345	0.7459	0.1378	0.7343
CG40203	CG40203	1635837_at	0.0056	0.9795	0.0623	0.5255	0.2758	0.0843	-0.0377	0.9470	-0.0927	0.5964	-0.0549	0.7539	-0.1527	0.8042	-0.0248	0.9519	0.1279	0.6421
CG13545	CG13545	1635838_at	0.4123	0.0873	0.5111	0.0179	0.4107	0.1025	0.0496	0.9586	-0.0393	0.9105	-0.0890	0.7438	-0.0351	0.9741	-0.1123	0.7090	-0.0772	0.8161
---	---	1635839_at	0.2828	0.2906	0.1278	0.5256	0.6130	0.0146	-0.1396	0.7854	-0.4971	0.0265	-0.3574	0.0622	-0.1569	0.8461	-0.1816	0.6093	-0.0246	0.9595
msl-2	male specifc letha	1635840_s_at	-0.4338	0.2272	-0.0776	0.8741	0.1710	0.4406	-0.1554	0.7982	-0.5268	0.0388	-0.3713	0.0916	-0.2010	0.8987	0.0847	0.9273	0.2857	0.6495
---	---	1635841_at	0.0166	0.9315	0.0892	0.6682	0.1882	0.3042	0.0666	0.9045	-0.1523	0.4129	-0.2188	0.1741	0.0505	0.9405	0.0918	0.6966	0.0413	0.8861
CG9311 /// DmirCG9311	CG9311	1635842_at	-0.2255	0.5121	-0.2124	0.3992	-0.0901	0.6563	0.2531	0.6175	0.4509	0.0769	0.1978	0.4010	0.0856	0.9445	0.4375	0.2224	0.3518	0.3572
CG9144	CG9144	1635843_at	-0.2266	0.1792	-0.1208	0.4882	-0.0657	0.6756	0.0284	0.9675	0.0409	0.8656	0.0125	0.9573	-0.0626	0.9246	0.1226	0.6071	0.1852	0.4132
CG15887	CG15887	1635844_at	0.0474	0.7964	0.0832	0.5865	0.2594	0.2116	0.0102	0.9893	-0.1557	0.3847	-0.1659	0.2954	-0.1643	0.8202	-0.0246	0.9605	0.1397	0.6677
CG30480	CG30480	1635845_at	-0.0071	0.9714	-0.0984	0.5952	0.0164	0.9404	0.2409	0.5917	0.2683	0.2393	0.0275	0.9218	0.0066	0.9952	0.2265	0.4138	0.2199	0.4448
orb	oo18 RNA-binding	1635846_a_at	1.3288	0.2645	-2.5613	0.0884	-2.0374	0.0355	0.6393	0.0942	3.5569	0.0000	2.9176	0.0000	-0.3966	0.9647	-0.5776	0.8642	-0.1810	0.9588
Vm26Aa	Vitelline membran	1635847_at	0.6237	0.5720	-1.1608	0.0108	-0.2989	0.3139	1.0610	0.1482	1.1678	0.0166	0.1068	0.8332	-0.2653	0.9502	-1.0389	0.4056	-0.7736	0.5630
Inos	myo-inositol-1-phc	1635848_at	-1.2499	0.0012	-1.7468	0.0215	-2.3090	0.0000	-0.0680	0.8686	0.1921	0.1827	0.2602	0.0497	0.1612	0.8589	-0.5687	0.1245	-0.7299	0.0834
CG9804	CG9804	1635849_at	-0.2508	0.2534	0.2170	0.3499	0.0222	0.8989	-0.0829	0.8495	-0.2148	0.1750	-0.1319	0.3696	0.1282	0.8832	0.3298	0.3071	0.2017	0.5670
---	---	1635850_at	0.1106	0.5387	-0.1512	0.3580	-0.0626	0.7009	0.1414	0.6311	0.2293	0.1109	0.0879	0.5325	0.0152	0.9899	-0.1077	0.7213	-0.1229	0.6683
CG15352	CG15352	1635851_at	-0.6276	0.1626	-0.6136	0.1163	-0.9131	0.0012	-0.0654	0.9435	0.5423	0.0436	0.6077	0.0172	-0.0879	0.9503	0.1954	0.6762	0.2833	0.5192
Wnt5	wnt-oncogene-ana	1635852_at	0.4431	0.0717	0.3803	0.1716	0.2942	0.2303	0.0629	0.9436	0.1295	0.6505	0.0667	0.8166	0.4460	0.3712	0.1936	0.4832	-0.2524	0.3636
---	---	1635853_at	0.0710	0.6976	0.0734	0.5262	0.1344	0.3935	-0.0222	0.9774	-0.0763	0.7334	-0.0541	0.8007	-0.0360	0.9611	-0.0002	0.9999	0.0358	0.8998
CG40351	CG40351	1635854_s_at	-0.0552	0.9019	0.3949	0.4333	-0.2362	0.3314	-0.3362	0.4568	-0.0549	0.8652	0.2813	0.1977	0.1874	0.9142	0.2904	0.6647	0.1030	0.9025
CG30076	CG30076	1635855_at	0.1636	0.3819	0.1126	0.5283	-0.0437	0.8233	-0.1209	0.8546	0.0689	0.8159	0.1898	0.3922	0.0111	0.9923	-0.0794	0.8225	-0.0906	0.7764
CG13163	CG13163	1635856_at	-0.0666	0.8091	0.6836	0.0847	0.4651	0.0738	-0.2135	0.7409	-0.5250	0.0683	-0.3115	0.2218	0.1752	0.8108	0.2488	0.4084	0.0736	0.8511
CG10919	CG10919	1635857_at	0.0070	0.9842	0.0294	0.9010	0.0069	0.9709	0.1408	0.7735	0.1884	0.3693	0.0476	0.8407	-0.0353	0.9816	0.0111	0.9871	0.0464	0.9202
Nhe2	Nhe2	1635858_s_at	-0.9009	0.0136	0.1765	0.3353	-0.3301	0.2443	-0.3826	0.4083	-1.0031	0.0029	-0.6205	0.0150	0.1269	0.9071	0.1417	0.7604	0.0148	0.9824
---	---	1635859_s_at	0.1344	0.5266	0.1531	0.3433	-0.0393	0.8277	-0.0606	0.9302	-0.0462	0.8627	0.0144	0.9556	0.1108	0.8890	0.0048	0.9941	-0.1060	0.7588
---	---	1635860_at	0.2454	0.2596	0.0830	0.6811	0.2793	0.3014	-0.1498	0.8074	-0.1447	0.5866	0.0052	0.9867	-0.1863	0.8012	-0.1213	0.7361	0.0651	0.8761
Tpr2	Tetratricopeptide r	1635861_a_at	-0.4294	0.1914	0.3882	0.1811	0.5341	0.0077	-0.0420	0.9705	-0.2603	0.3786	-0.2183	0.4201	-0.0281	0.9848	0.5175	0.1287	0.5456	0.1368
CG33120	CG33120	1635862_at	3.1739	0.0024	1.4945	0.0316	3.1043	0.0002	0.6130	0.6597	1.0070	0.1305	0.3940	0.5485	-0.9052	0.5126	-0.6942	0.2753	0.2110	0.7886
CG7185 /// DrepCG7185	CG7185	1635863_at	0.0208	0.9804	-0.5393	0.4791	-0.5884	0.0729	0.1243	0.7018	0.7172	0.0012	0.5929	0.0016	0.3021	0.9340	0.2576	0.8765	-0.0445	0.9821
---	---	1635864_at	0.0376	0.8424	-0.1706	0.1648	0.0596	0.8162	0.0748	0.8676	0.2812	0.0781	0.2064	0.1444	-0.3067	0.6557	-0.0993	0.7668	0.2075	0.4683
---	---	1635865_at	-0.2000	0.3683	-0.2023	0.3693	-0.4158	0.0158	-0.0635	0.9300	0.5238	0.0214	0.5873	0.0080	-0.0427	0.9507	0.1719	0.3959	0.2146	0.3061
---	---	1635866_at	0.1143	0.5663	-0.0404	0.7938	0.1083	0.4812	0.1210	0.7140	0.1453	0.3430	0.0244	0.8925	0.0386	0.9659	0.0896	0.7579	0.0510	0.8775
---	---	1635867_at	0.0959	0.5598	0.1083	0.5657	-0.1100	0.5631	-0.1194	0.8074	0.0388	0.8814	0.1582	0.3813	0.2416	0.6955	0.1328	0.6319	-0.1088	0.7060
CG6295	CG6295	1635868_at	1.0223	0.3969	0.0728	0.6518	0.2907	0.3291	0.1395	0.8621	0.2563	0.3815	0.1169	0.6983	-0.2661	0.9523	-0.7074	0.6229	-0.4412	0.7799
---	---	1635869_at	0.3320	0.1457	0.2269	0.1996	0.3259	0.2001	0.0758	0.9339	0.0503	0.8906	-0.0256	0.9393	0.1753	0.8421	0.0651	0.9064	-0.1102	0.8013
CG1628	CG1628	1635870_at	0.7429	0.0817	0.7360	0.2543	1.7764	0.0009	0.0265	0.9777	-0.1632	0.5078	-0.1898	0.3796	-0.7781	0.6960	-0.1399	0.9149	0.6382	0.4647
beta'Cop	beta'-coatomer pr	1635871_at	0.3028	0.2309	1.1478	0.0328	1.4674	0.0001	0.2792	0.4346	-0.2166	0.2716	-0.4959	0.0123	-0.0750	0.9489	0.8326	0.0391	0.9076	0.0441
---	---	1635872_at	0.1316	0.5389	0.3370	0.1131	0.7257	0.0364	0.3779	0.5779	-0.2605	0.4739	-0.6384	0.0458	0.1429	0.8363	0.1577	0.6044	0.0149	0.9732
CG34030	CG34030	1635873_at	-0.0005	0.9978	-0.2214	0.1835	-0.1221	0.5631	-0.0186	0.9757	0.0010	0.9963	0.0196	0.9143	-0.1567	0.8193	-0.2761	0.3134	-0.1194	0.7053

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG2794	CG2794	1635874_at	0.3874	0.0614	-0.0609	0.6127	0.0934	0.6760	0.0213	0.9753	0.3899	0.0278	0.3685	0.0223	-0.1314	0.8395	-0.0267	0.9505	0.1047	0.7318
mr	morula	1635875_at	-0.4932	0.0154	-0.1939	0.3098	-0.3095	0.1792	-0.2128	0.4858	-0.0695	0.7123	0.1433	0.3360	-0.2096	0.7485	0.1772	0.5422	0.3868	0.1902
---	---	1635876_at	0.0199	0.9351	0.1031	0.6031	-0.2537	0.1997	-0.0421	0.9496	0.0710	0.7469	0.1131	0.5318	0.1789	0.7606	0.1181	0.6616	-0.0609	0.8472
---	---	1635877_at	0.3393	0.0474	0.1326	0.5308	0.1911	0.2151	0.0435	0.9482	0.0368	0.8840	-0.0067	0.9776	0.0204	0.9841	-0.0737	0.8031	-0.0941	0.7192
CG17571 /// DyakCG17571	CG17571	1635878_s_at	0.5229	0.4915	-0.1925	0.2615	-0.1438	0.4944	0.0520	0.9649	0.0773	0.8508	0.0253	0.9496	-0.1857	0.9445	-0.6349	0.4300	-0.4492	0.6045
---	---	1635879_at	0.0273	0.9119	0.1507	0.2730	0.0666	0.6887	-0.1234	0.7845	-0.0366	0.8836	0.0868	0.6517	0.0289	0.9814	0.1422	0.6144	0.1133	0.7005
CG15710	CG15710	1635880_at	0.1985	0.1695	-0.0092	0.9331	0.2395	0.4306	0.2680	0.5557	0.1095	0.6829	-0.1585	0.4790	-0.0798	0.9273	0.0244	0.9590	0.1042	0.7526
Nmnat	Nicotinamide mon	1635881_at	-0.2569	0.2523	-0.2536	0.2611	-0.1414	0.5576	0.0078	0.9937	0.2284	0.2578	0.2206	0.2204	-0.1564	0.8461	0.0809	0.8611	0.2372	0.4889
---	---	1635882_at	0.0348	0.9116	0.1584	0.3936	0.1273	0.3663	-0.0823	0.8791	-0.1149	0.5626	-0.0326	0.8809	0.2291	0.7893	0.3090	0.3903	0.0799	0.8694
---	---	1635883_at	-0.3696	0.1493	0.0356	0.7261	-0.0587	0.7246	-0.0229	0.9860	-0.3426	0.2566	-0.3197	0.2355	0.0924	0.8395	0.0933	0.6483	0.0009	0.9980
---	---	1635884_at	0.2026	0.1647	-0.1071	0.3007	-0.1004	0.5244	0.2144	0.4612	0.2791	0.0787	0.0647	0.6964	-0.1967	0.7215	-0.1575	0.5102	0.0392	0.9025
---	---	1635885_s_at	0.0959	0.5813	-0.0008	0.9962	0.1516	0.3856	0.0082	0.9909	-0.0092	0.9696	-0.0175	0.9270	0.0398	0.9626	-0.0306	0.9353	-0.0704	0.8071
---	---	1635886_s_at	1.2091	0.0186	1.6145	0.0918	1.2571	0.0038	0.2056	0.8376	0.6724	0.0763	0.4668	0.1644	0.7977	0.7043	1.3062	0.1353	0.5085	0.5862
CG30094	CG30094	1635887_at	-0.1169	0.5470	0.1702	0.7025	0.0078	0.9658	0.0849	0.8507	0.1584	0.3436	0.0735	0.6669	0.3623	0.7215	0.4919	0.2361	0.1297	0.8044
---	---	1635888_at	0.1708	0.3206	-0.0602	0.6971	-0.1407	0.4312	-0.1163	0.8070	0.0468	0.8485	0.1631	0.3476	0.0388	0.9672	-0.0962	0.7413	-0.1351	0.6140
---	---	1635889_at	-0.0084	0.9699	-0.1395	0.4504	0.0329	0.8495	0.1746	0.6311	0.1603	0.3784	-0.0143	0.9495	0.0209	0.9816	0.0405	0.8879	0.0196	0.9434
capaR	capa receptor	1635890_at	0.1650	0.5634	-0.0368	0.7968	0.1948	0.3650	0.0457	0.9620	0.1155	0.6942	0.0697	0.8079	-0.1901	0.8395	0.0563	0.9270	0.2464	0.5475
Eig71Eh	Eig71Eh	1635891_at	0.0660	0.7663	-0.0388	0.6899	-0.1163	0.4149	0.1890	0.5642	0.1773	0.2968	-0.0117	0.9567	0.0135	0.9922	-0.0251	0.9617	-0.0387	0.9296
CG12321	CG12321	1635892_at	0.1337	0.5235	-0.0013	0.9993	0.3525	0.1238	0.2361	0.5902	0.0771	0.7737	-0.1591	0.4482	-0.0499	0.9717	0.0258	0.9647	0.0757	0.8736
alpha-Est8	fragment I	1635893_at	-1.7509	0.0008	-0.0420	0.9536	-1.7391	0.0167	-1.5351	0.1194	-2.6072	0.0018	-1.0721	0.0490	0.0233	0.9943	-0.8611	0.2328	-0.8844	0.2528
CG30148	CG30148	1635894_at	0.4199	0.2748	-0.6299	0.5185	-0.0336	0.9139	-0.1173	0.9072	-0.4628	0.1515	-0.3455	0.2339	-0.7233	0.7726	-1.4679	0.1584	-0.7447	0.5032
CG31133	CG31133	1635895_at	-0.1985	0.2275	0.0359	0.8540	0.5162	0.0195	-0.0948	0.8667	-0.2896	0.1440	-0.1948	0.2789	-0.5952	0.1628	-0.0860	0.7349	0.5092	0.0559
---	---	1635896_at	0.0504	0.8517	-0.0775	0.5811	-0.1277	0.5356	0.1361	0.8058	0.0147	0.9656	-0.1215	0.5809	0.0802	0.9291	-0.0195	0.9678	-0.0997	0.7703
CG15399	CG15399	1635897_at	-0.5389	0.0120	-0.0563	0.6736	-0.4513	0.0985	-0.3694	0.3352	-0.9203	0.0022	-0.5508	0.0131	-0.0201	0.9913	-0.5030	0.1622	-0.4829	0.2110
CG14856	CG14856	1635898_at	-0.6875	0.0106	-0.2084	0.0859	-0.4463	0.0264	-0.0571	0.9540	-0.3218	0.2372	-0.2647	0.2817	-0.0368	0.9666	-0.0557	0.8649	-0.0189	0.9546
---	---	1635899_s_at	-0.0969	0.6996	0.0340	0.7709	-0.0813	0.6878	0.0788	0.9149	-0.0035	0.9914	-0.0823	0.7403	0.0792	0.9333	0.1159	0.7445	0.0367	0.9313
Thor	insulin-stimulated	1635900_at	0.9127	0.0013	1.9913	0.0031	1.9427	0.0002	0.0781	0.9116	-1.1755	0.0008	-1.2536	0.0004	0.2425	0.8222	-0.0386	0.9588	-0.2811	0.5547
CG17197	CG17197	1635901_at	0.2541	0.2187	0.1320	0.6020	0.2273	0.2150	-0.0189	0.9854	0.0454	0.8855	0.0643	0.8066	0.0000	0.9999	-0.0572	0.8890	-0.0572	0.8813
---	---	1635902_at	-0.0072	0.9820	0.0452	0.6452	0.0203	0.9521	-0.0910	0.9228	-0.0148	0.9733	0.0762	0.8124	0.0055	0.9963	-0.0156	0.9704	-0.0210	0.9525
Pk92B	Apoptotic signal-r	1635903_at	-0.5032	0.0287	-0.6643	0.0435	-0.7482	0.0015	-0.0537	0.9339	0.1835	0.3425	0.2372	0.1613	0.0665	0.9246	-0.0185	0.9620	-0.0849	0.7526
BtbVII	BTB protein VII	1635904_s_at	-0.0319	0.9458	-0.1697	0.8287	-0.4912	0.0250	-0.1534	0.6641	0.2698	0.1091	0.4232	0.0128	0.0373	0.9918	0.1551	0.9050	0.1178	0.9197
CG31532	CG31532	1635905_at	0.1576	0.6372	-0.1179	0.5600	0.0170	0.9522	0.1955	0.8087	0.1968	0.5709	0.0013	0.9976	0.0753	0.9499	-0.0521	0.9265	-0.1274	0.7561
---	---	1635906_at	-0.0476	0.8693	0.0747	0.5720	0.1989	0.2102	0.0875	0.8967	-0.0802	0.7634	-0.1677	0.4192	-0.1270	0.8270	-0.0872	0.7692	0.0398	0.9057
CG7414	CG7414	1635907_at	-1.0509	0.0630	0.0767	0.9451	0.5652	0.0171	0.3144	0.5172	-1.0713	0.0022	-1.3858	0.0005	-0.1075	0.9816	0.2216	0.8796	0.3291	0.7871
CG11071 /// DmirCG11071	CG11071	1635908_at	0.2490	0.2699	0.3777	0.0859	0.1572	0.3716	-0.0781	0.8297	-0.0496	0.7665	0.0285	0.8605	0.0872	0.9061	0.1087	0.7200	0.0215	0.9545
cic	fettucine	1635909_at	1.1500	0.2879	1.4076	0.1202	-0.2860	0.6743	-0.7244	0.1447	0.6652	0.0337	1.3896	0.0008	0.9929	0.8086	0.9869	0.5854	-0.0060	0.9987
CG7331	CG7331	1635910_at	0.2838	0.1543	-0.0964	0.7362	0.0955	0.7010	-0.0967	0.8405	0.0023	0.9932	0.0990	0.5803	-0.2319	0.8202	-0.2285	0.6139	0.0034	0.9965
CG10347	CG10347	1635911_s_at	0.5176	0.0125	0.8285	0.0562	0.7067	0.0017	-0.0130	0.9838	0.1028	0.5078	0.1158	0.3970	0.1558	0.8521	0.3529	0.2942	0.1971	0.5963
Rh7	rhodopsin	1635912_at	0.2091	0.2218	0.1733	0.4558	0.2717	0.1313	0.0433	0.9331	-0.0139	0.9517	-0.0572	0.7249	0.0923	0.9451	-0.1366	0.7883	-0.2288	0.6005
CG10814	CG10814	1635913_at	0.3988	0.4273	0.0253	0.9771	0.2922	0.4148	0.6551	0.1787	0.9361	0.0074	0.2810	0.2931	0.1825	0.9421	0.2798	0.7668	0.0973	0.9282
CG10413	CG10413	1635914_at	-1.4068	0.0020	-2.3033	0.0019	-2.2166	0.0001	0.1301	0.8727	0.7787	0.0136	0.6486	0.0182	0.1273	0.8222	0.0386	0.9157	-0.0887	0.7442
SmD3	gut feeling	1635915_at	0.5146	0.0208	0.6519	0.1502	0.7389	0.0498	0.2702	0.6564	0.0802	0.8281	-0.1900	0.4987	0.1352	0.9330	0.4439	0.3802	0.3087	0.5731
---	---	1635916_at	-0.0749	0.6535	0.0294	0.7873	-0.0235	0.9073	0.0145	0.9860	-0.0331	0.8985	-0.0476	0.8254	0.0244	0.9775	0.1049	0.6345	0.0805	0.7307
RpL35	Ribosomal protein	1635917_at	0.7293	0.0588	0.5228	0.2754	0.5672	0.0347	0.2264	0.7278	0.1444	0.6612	-0.0820	0.8025	0.1103	0.9499	-0.0179	0.9865	-0.1282	0.8486
CG34401	CG17757	1635918_at	-0.3152	0.6347	-0.5643	0.5825	-1.0732	0.0080	0.0493	0.9705	0.4552	0.1811	0.4059	0.1838	0.5738	0.8378	0.3011	0.8478	-0.2728	0.8555
CG13255	CG13255	1635919_at	0.5026	0.5470	-0.2282	0.5472	-0.0139	0.9679	0.0217	0.9906	-0.6918	0.0868	-0.7134	0.0520	-0.1731	0.9619	-1.3637	0.1657	-1.1906	0.2558
CG15226	CG15226	1635920_at	-0.0561	0.8180	-0.0021	0.9916	-0.0002	0.9996	0.1128	0.8714	0.1176	0.6710	0.0048	0.9874	0.0156	0.9914	0.0493	0.9270	0.0336	0.9434
CG3502	CG3502	1635921_at	0.2033	0.3048	-0.0004	0.9999	0.1679	0.2503	0.0387	0.9422	0.0748	0.6648	0.0360	0.8358	-0.0934	0.8689	-0.0583	0.8481	0.0351	0.9095
---	---	1635922_at	-0.1049	0.7010	0.1922	0.4185	0.2025	0.2894	0.1358	0.8288	0.0146	0.9689	-0.1211	0.6217	-0.0173	0.9914	0.1365	0.7331	0.1538	0.6859
sunz	sungrazer	1635923_at	0.0506	0.8158	-0.0614	0.7074	0.1042	0.5994	0.2632	0.5502	0.1582	0.5112	-0.1049	0.6563	-0.0492	0.9611	-0.0320	0.9443	0.0172	0.9678
Teh3	tipE homolog 3	1635924_at	-0.0432	0.8025	-0.0158	0.9266	-0.0028	0.9922	0.0434	0.9466	-0.0793	0.7078	-0.1227	0.4835	0.0624	0.9296	-0.0528	0.8718	-0.1152	0.6399
Pk34A	Pk34A	1635925_at	0.2322	0.1633	-0.0667	0.7378	0.1401	0.3762	0.1195	0.7233	0.1390	0.3708	0.0196	0.9161	-0.1748	0.7775	-0.1560	0.5841	0.0189	0.9621
Cpr76Bb	CG9290	1635926_at	-0.0609	0.7545	0.0579	0.5663	-0.0441	0.8054	-0.1052	0.8117	0.0642	0.7595	0.1693	0.2842	-0.1204	0.8692	0.0464	0.9180	0.1667	0.5930
CG9746	CG9746	1635927_at	-0.3038	0.1948	-0.0611	0.8512	0.2182	0.2732	0.0442	0.9445	-0.1723	0.3520	-0.2165	0.1817	-0.1755	0.8779	0.1393	0.8061	0.3148	0.4913
Pka-C3	Protein-kinase-like	1635928_a_at	-1.1350	0.0285	0.1674	0.8136	-1.3057	0.0029	-1.0389	0.1028	-1.8533	0.0011	-0.8144	0.0236	0.2519	0.9112	-0.5238	0.5288	-0.7757	0.3475
---	---	1635929_at	-0.0712	0.8314	0.1544	0.3268	0.1350	0.5477	-0.1080	0.8852	-0.1906	0.4727	-0.0827	0.7676	-0.0261	0.9845	0.0808	0.8411	0.1069	0.7568
btn	buttonless	1635930_at	-0.0060	0.9799	-0.1272	0.4174	-0.1185	0.4474	0.0176	0.9755	0.1563	0.2672	0.1387	0.2736	0.0567	0.9400	0.0261	0.9445	-0.0305	0.9231
CG11563	CG11563	1635931_at	0.8230	0.0271	0.1735	0.8152	0.4209	0.1869	0.3335	0.5612	0.9767	0.0060	0.6432	0.0228	0.1056	0.9717	0.2457	0.7930	0.1400	0.8918
CG8202	CG8202	1635932_at	-0.0832	0.6605	-0.0011	0.9940	0.0713	0.7355	-0.0404	0.9436	-0.0896	0.6172	-0.0493	0.7860	-0.1452	0.8692	0.0138	0.9841	0.1590	0.6833
CG4496	CG4496	1635933_at	-0.2690	0.3320	-0.2458	0.4808	-0.9052	0.0371	-0.5480	0.2990	-0.0004	0.9994	0.5476	0.0528	-0.1005	0.9535	-0.1484	0.8250	-0.0479	0.9469
spag	spaghetti	1635934_at	0.1021	0.5200	-0.2190	0.4857	-0.2330	0.1895	-0.0231	0.9705	0.4537	0.0118	0.4768	0.0057	-0.0598	0.9647	0.0097	0.9906	0.0694	0.8897
CG34039	CG34039	1635935_at	0.3756	0.0279	0.1777	0.3507	0.1147	0.4490	0.0793	0.8650	0.0892	0.6277	0.0099	0.9620	0.1171	0.8000	-0.0021	0.9961	-0.1192	0.5605
CG13822	CG13822	1635936_at	-0.0021	0.9973	-0.7767	0.0779	-1.0565	0.0041	-0.0381	0.9715	-0.1836	0.5222	-0.1455	0.5898	0.1040	0.9725	-1.0857	0.1432	-1.1897	0.1441
CG4500	CG4500	1635937_at	1.3624	0.2482	0.2450	0.5153	0.2714	0.1042	0.0897	0.9436	-0.2336	0.5495	-0.3233	0.3296	-0.0197	0.9970	-1.3549	0.2840	-1.3352	0.3171
Ork1	Open rectifier K[+]	1635938_s_at	-1.7943	0.0197	-1.0705	0.2725	-1.6043	0.0017	-0.8669	0.1329	-1.4332	0.0023	-0.5663	0.0704	-0.3546	0.9149	-0.7480	0.5374	-0.3934	0.7745
CG9641	CG9641	1635939_a_at	-0.3202	0.2964	-0.5126	0.3295	-0.5069	0.1358	-0.0827	0.9247	0.7046	0.0151	0.7873	0.0056	-0.0663	0.9831	0.5441	0.4605	0.6103	0.4114
CG5367	CG5367	1635940_at	0.0192	0.9288	-0.0483	0.7950	0.2540	0.1343	0.1699	0.7121	0.1381	0.5377	-0.0317	0.8997	-0.0478	0.9710	0.0416	0.9382	0.0894	0.8305
---	---	1635941_at	-0.2089	0.2747	-0.2554	0.1121	-0.2792	0.0882	-0.0880	0.8732	0.0624	0.7899	0.1504	0.3988	-0.0021	0.9990	-0.1036	0.7058	-0.1016	0.7113
---	---	1635942_at	-0.0965	0.4897	0.0727	0.5103	0.1037	0.5961	-0.0752	0.8931	-0.0674	0.7634	0.0078	0.9732	0.0034	0.9975	0.0963	0.7345	0.0929	0.7416

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG10947	CG10947	1635943_s_at	-0.8215	0.0297	0.0510	0.8013	-0.1420	0.3533	-0.0811	0.9260	-0.5118	0.0561	-0.4306	0.0696	0.0506	0.9717	0.1678	0.6765	0.1172	0.7853
Syb	synaptobrevin	1635944_a_at	0.0281	0.8680	0.9030	0.0174	0.8476	0.0036	0.2100	0.6338	-0.4030	0.0660	-0.6130	0.0073	0.2344	0.7230	0.4338	0.1299	0.1994	0.5003
CG15394	CG15394	1635945_at	-2.2224	0.0030	-4.2789	0.0019	-3.3144	0.0000	0.3290	0.6457	2.2395	0.0004	1.9105	0.0004	-0.4542	0.7644	0.3493	0.6065	0.8035	0.2219
CG17923	CG17923	1635946_at	-0.0254	0.8736	-0.1764	0.3081	-0.1390	0.5540	0.0205	0.9833	0.2038	0.3705	0.1833	0.3737	-0.1462	0.8122	-0.0621	0.8612	0.0841	0.7798
CG9603	CG9603	1635947_at	-0.1729	0.3090	0.2584	0.2331	0.0982	0.7214	-0.0646	0.9254	-0.5684	0.0139	-0.5039	0.0144	0.1712	0.8461	-0.1501	0.7198	-0.3213	0.3878
---	---	1635948_at	-0.0321	0.8558	0.1135	0.2884	0.2512	0.1333	0.0180	0.9808	-0.0325	0.8959	-0.0506	0.8047	-0.3074	0.5128	-0.1038	0.6792	0.2036	0.3801
---	---	1635949_at	0.1390	0.3657	0.0176	0.9227	0.0503	0.7660	0.0807	0.8384	0.0870	0.5899	0.0063	0.9730	0.0650	0.9411	0.0003	0.9999	-0.0647	0.8548
Su(P)	Suppressor of ref(	1635950_at	-0.2045	0.2387	0.3713	0.1292	0.6616	0.0171	0.1567	0.8103	-0.5575	0.0382	-0.7141	0.0083	-0.0103	0.9923	0.2196	0.3697	0.2300	0.3689
CG7069	CG7069	1635951_at	0.1579	0.6049	-0.0045	0.9849	-0.0276	0.9037	0.0530	0.9436	0.1815	0.4108	0.1286	0.5393	0.0736	0.9516	0.0224	0.9708	-0.0512	0.9194
l(2)efl	lethal (2) essentia	1635952_at	-0.0692	0.6993	-0.0634	0.6452	-0.0636	0.7595	0.1594	0.7008	0.1195	0.5603	-0.0398	0.8577	-0.0412	0.9742	-0.0387	0.9402	0.0025	0.9965
CG31524	CG31524	1635953_at	0.1127	0.5986	-0.0682	0.5561	0.0438	0.8050	0.0752	0.8943	0.0545	0.8157	-0.0207	0.9286	0.0924	0.8846	-0.1196	0.6487	-0.2121	0.3898
lva	Lava Lamp	1635954_at	0.5435	0.3808	0.7078	0.2024	1.0893	0.0010	0.2765	0.8017	0.9991	0.0312	0.7226	0.0701	0.0171	0.9952	1.0540	0.1349	1.0369	0.1651
CG3883	CG3883	1635955_at	0.1580	0.3175	-0.0805	0.5431	-0.0729	0.6569	0.0120	0.9864	0.4929	0.0112	0.4809	0.0076	-0.0235	0.9816	0.1998	0.3720	0.2233	0.3404
CG10089	MKP-like	1635956_at	0.1598	0.5640	0.1879	0.3376	0.0897	0.6339	0.0724	0.9311	0.0737	0.8080	0.0013	0.9966	0.1373	0.8270	0.1241	0.6669	-0.0131	0.9748
---	---	1635957_at	0.0513	0.7348	0.1078	0.5385	0.0508	0.7807	-0.1324	0.7592	-0.0924	0.6594	0.0400	0.8526	-0.0248	0.9816	-0.0248	0.9508	-0.0001	0.9999
---	---	1635958_at	-0.4206	0.0829	-0.9304	0.1170	-0.7897	0.0135	0.2327	0.6936	0.8676	0.0073	0.6349	0.0173	0.3193	0.7726	0.4735	0.3036	0.1542	0.7838
CG13461	CG13461	1635959_at	0.2426	0.3228	0.0589	0.6339	0.2927	0.1643	-0.0061	0.9937	0.0102	0.9666	0.0164	0.9328	0.0255	0.9831	-0.0727	0.8502	-0.0982	0.7654
slmb	pingiel	1635960_at	-0.6040	0.0774	0.0805	0.6587	-0.0243	0.8938	-0.0555	0.9266	-0.1392	0.4604	-0.0836	0.6549	0.1459	0.8958	0.5151	0.1995	0.3692	0.3844
CG33472	CG33472	1635961_at	-0.8140	0.0116	-0.3685	0.0459	-1.1245	0.0008	-0.1096	0.8385	-0.0635	0.7974	0.0461	0.8432	-0.0197	0.9862	-0.0789	0.8215	-0.0592	0.8674
CG12609	CG12609	1635962_a_at	0.1632	0.6495	-0.0500	0.7881	0.1851	0.3686	0.0208	0.9894	-0.1413	0.7417	-0.1620	0.6671	-0.1703	0.8167	-0.1283	0.7080	0.0419	0.9197
CG7990	CG7990	1635963_a_at	-1.4653	0.0044	-0.4893	0.3862	-2.1554	0.0020	-1.2877	0.0466	-1.2385	0.0056	0.0492	0.9125	0.1291	0.9626	-0.4282	0.5983	-0.5573	0.4805
niki	nimA-like kinase	1635964_at	0.5052	0.0340	0.1335	0.4071	0.3404	0.0744	0.0864	0.8794	0.0839	0.7096	-0.0025	0.9917	0.0912	0.8608	-0.0594	0.8304	-0.1506	0.4863
CG17917	CG17917	1635965_at	0.4600	0.0414	0.3080	0.0953	0.2152	0.1578	-0.0244	0.9777	0.0533	0.8540	0.0777	0.7485	0.1515	0.7872	0.0032	0.9950	-0.1483	0.5597
---	---	1635966_at	0.1009	0.7323	0.1389	0.3918	0.2689	0.1681	0.0111	0.9883	-0.0442	0.8463	-0.0553	0.7771	-0.1171	0.9084	0.0232	0.9689	0.1403	0.7346
qm	geranylgeranyl py	1635967_at	0.0426	0.9092	0.5560	0.0807	0.0057	0.9858	-0.2865	0.5199	-0.0886	0.7499	0.1978	0.3581	0.2982	0.7633	0.3879	0.3349	0.0897	0.8709
CG3604	CG3604	1635968_at	-2.6572	0.0938	-4.1945	0.0014	-3.5646	0.0001	0.1590	0.9017	0.3194	0.4635	0.1604	0.7180	-0.1290	0.9871	-1.0042	0.6075	-0.8752	0.6588
CG11590	CG11590	1635969_at	0.0736	0.6971	-0.3046	0.3317	-0.0522	0.7789	-0.0045	0.9956	0.3071	0.0868	0.3116	0.0553	-0.1595	0.8494	-0.1052	0.8170	0.0544	0.9101
Sin	Sex-lethal interact	1635970_at	0.2653	0.1821	0.3137	0.3657	0.1827	0.3697	0.2344	0.5522	0.6198	0.0089	0.3855	0.0408	0.3645	0.6635	0.6452	0.0716	0.2807	0.4139
CG31525	CG31525	1635971_at	0.2416	0.1595	0.0801	0.5744	0.1644	0.4439	-0.0221	0.9777	0.0274	0.9220	0.0496	0.8282	0.0614	0.9405	0.0425	0.9135	-0.0189	0.9591
CG15014	CG15014	1635972_at	0.2957	0.0837	0.0302	0.9582	0.0519	0.7334	-0.1481	0.6301	0.4993	0.0060	0.6473	0.0012	-0.2599	0.8192	0.0075	0.9941	0.2673	0.5906
CG11196	CG11196	1635973_at	0.1069	0.5726	0.0178	0.8606	0.1254	0.5496	-0.0869	0.8462	-0.0532	0.7913	0.0336	0.8605	-0.1045	0.8521	-0.0439	0.9009	0.0606	0.8370
Pros54	54 kda mu particle	1635974_at	-0.2090	0.2441	0.6419	0.0168	0.8876	0.0006	0.1819	0.6004	-0.4132	0.0259	-0.5951	0.0032	0.0256	0.9816	0.4623	0.0722	0.4367	0.1096
---	---	1635975_s_at	0.4771	0.2991	0.0271	0.9502	-0.2330	0.2658	-0.0015	0.9987	0.0707	0.7566	0.0722	0.7227	0.3027	0.8298	-0.3042	0.6325	-0.6069	0.3141
CG9977	CG9977	1635976_at	-2.0771	0.0074	-0.3645	0.5406	-0.2245	0.3873	-0.1187	0.9491	-1.2712	0.0227	-1.1524	0.0213	-0.1960	0.9238	0.4307	0.5470	0.6266	0.3723
---	---	1635977_at	0.0221	0.9233	-0.1655	0.3389	-0.1789	0.3722	0.0097	0.9917	0.1372	0.5015	0.1275	0.4921	-0.0254	0.9779	-0.0278	0.9369	-0.0025	0.9950
CG8229 /// DyakCG8229	CG8229	1635978_at	-0.7707	0.0142	0.0266	0.8672	-0.3886	0.0674	-0.4242	0.2596	-1.1042	0.0011	-0.6801	0.0052	0.0903	0.9301	-0.3024	0.3526	-0.3926	0.2555
CG7600	CG7600	1635979_at	0.2320	0.2367	0.7602	0.0269	0.6741	0.0016	-0.0236	0.9745	-0.0615	0.7803	-0.0378	0.8585	0.1732	0.7893	0.4575	0.1047	0.2843	0.3139
CG12991	CG12991	1635980_s_at	0.4293	0.3578	0.8553	0.0626	1.0764	0.0021	0.3439	0.2492	0.0285	0.9077	-0.3155	0.0506	0.2023	0.9235	0.5863	0.3914	0.3840	0.6061
growl	growl	1635981_a_at	-0.1077	0.5666	0.0589	0.6064	-0.3017	0.2179	-0.0703	0.8889	0.0010	0.9969	0.0712	0.6873	0.3817	0.5754	0.1712	0.6007	-0.2105	0.5088
---	---	1635982_at	0.1375	0.3175	0.2169	0.1726	0.2714	0.1368	0.1072	0.8231	-0.0459	0.8489	-0.1531	0.3736	-0.0482	0.9411	-0.0093	0.9798	0.0389	0.8889
nompB	no mechanorecep	1635983_a_at	0.1148	0.5511	-0.0832	0.6178	0.1627	0.3533	0.0536	0.9263	0.0900	0.6459	0.0364	0.8577	-0.0434	0.9701	0.0449	0.9225	0.0882	0.8088
Neu3	Meltrin-like	1635984_at	-0.5454	0.1146	0.8733	0.0301	1.2217	0.0001	-0.1213	0.8642	-1.1844	0.0013	-1.0631	0.0013	-0.1084	0.9302	0.4068	0.2971	0.5152	0.2140
CG40154	CG40154	1635985_at	0.1232	0.4510	0.0350	0.7247	-0.0265	0.9114	-0.0823	0.8932	-0.0151	0.9607	0.0673	0.7645	-0.0341	0.9589	-0.0624	0.7886	-0.0282	0.9121
---	---	1635986_at	0.1637	0.5986	-0.0280	0.7941	0.2002	0.2903	0.0965	0.8578	0.0067	0.9816	-0.0898	0.6553	-0.0324	0.9831	-0.1167	0.7901	-0.0843	0.8524
CG12116	CG12116	1635987_at	1.5608	0.0040	0.8744	0.0656	2.0544	0.0007	0.6554	0.3953	0.1958	0.6946	-0.4597	0.2367	-0.3228	0.7810	-0.4189	0.3992	-0.0961	0.8893
---	---	1635988_at	0.0654	0.7735	0.4359	0.0504	-0.0594	0.7712	-0.2320	0.5633	-0.1524	0.4798	0.0796	0.7165	0.1694	0.7945	0.3262	0.2168	0.1568	0.5907
Os9	Olfactory-specific	1635989_at	2.7531	0.0012	1.5630	0.0745	3.2909	0.0001	0.8142	0.3847	0.3429	0.5476	-0.4713	0.3303	-0.9386	0.5754	-0.8284	0.2538	0.1102	0.9162
Or65b	Odorant receptor (	1635990_at	0.0334	0.9046	0.0441	0.8479	0.0761	0.7587	0.0681	0.9441	0.0438	0.9095	-0.0243	0.9444	0.0118	0.9923	0.0367	0.9353	0.0249	0.9503
Osi5	Osiris	1635991_at	0.1569	0.3951	-0.1319	0.5902	-0.0218	0.9148	0.0483	0.9313	0.2040	0.2161	0.1557	0.2991	-0.1082	0.8465	-0.1532	0.5242	-0.0451	0.8879
sno	Strawberry Notch	1635992_at	0.1260	0.6838	0.1433	0.4703	0.3420	0.1743	0.1359	0.7352	0.0358	0.8808	-0.1001	0.5730	-0.0269	0.9898	-0.0302	0.9664	-0.0034	0.9965
tll	TAILLESS	1635993_at	-0.1081	0.6035	0.1821	0.2025	0.0023	0.9946	-0.0386	0.9647	-0.1954	0.4125	-0.1568	0.4766	0.1555	0.8611	-0.0430	0.9409	-0.1985	0.6095
CG32547	CG32547	1635994_at	0.2371	0.1751	0.1317	0.5124	-0.0011	0.9964	0.0576	0.9247	0.1820	0.3265	0.1244	0.4748	0.0738	0.9514	-0.0141	0.9841	-0.0879	0.8511
Mms19	Mms19	1635995_a_at	-0.0307	0.8478	0.4072	0.3905	0.0914	0.5961	-0.0315	0.9610	0.0951	0.6169	0.1266	0.4331	0.2158	0.8494	0.5602	0.2191	0.3444	0.4815
CG6638	CG6638	1635996_at	0.5021	0.0213	0.2693	0.0560	0.6562	0.0031	0.1469	0.7167	0.3155	0.0877	0.1686	0.3145	-0.1256	0.8731	0.0208	0.9676	0.1463	0.6630
Tango1	Transport and Gol	1635997_a_at	0.4827	0.2411	0.2912	0.3322	0.4737	0.0550	0.2143	0.6451	0.9672	0.0019	0.7529	0.0034	0.0439	0.9862	0.7245	0.2066	0.6806	0.2668
Cpr67B	CG3672	1635998_at	-2.5608	0.0117	-3.4084	0.0220	-3.3407	0.0000	0.0529	0.9329	-0.6796	0.0039	-0.7325	0.0017	0.1567	0.9776	-1.2876	0.3225	-1.4443	0.2933
Obp56b	Odorant-binding p	1635999_at	0.4445	0.0535	0.1016	0.4118	0.0947	0.6509	0.0401	0.9518	0.1715	0.3608	0.1314	0.4488	-0.0955	0.9011	-0.1472	0.6281	-0.0517	0.8936
CG13640	CG13640	1636000_at	0.1711	0.4136	0.2441	0.2772	-0.0676	0.7594	-0.0980	0.8340	-0.0629	0.7676	0.0352	0.8650	0.2051	0.7478	0.1458	0.6150	-0.0593	0.8694
CG18186	CG18186	1636001_at	0.1097	0.6044	-0.1034	0.4746	-0.0718	0.7621	0.0381	0.9610	0.1398	0.5283	0.1016	0.6323	0.0774	0.9409	-0.2192	0.5118	-0.2966	0.3711
CG4281 /// DmirCG4281	CG4281	1636002_at	0.1831	0.3375	-0.0075	0.9645	-0.0741	0.7336	0.0951	0.8156	0.3577	0.0330	0.2625	0.0697	0.3471	0.6749	0.2700	0.4317	-0.0771	0.8655
CG6388	CG6388	1636003_at	0.1486	0.4404	0.1292	0.7124	0.3048	0.2243	0.1010	0.8732	0.1988	0.3762	0.0979	0.6684	0.0479	0.9754	0.2996	0.4303	0.2517	0.5299
---	---	1636004_at	0.1552	0.2793	-0.1705	0.3923	0.1369	0.3733	0.3157	0.4596	0.2688	0.2460	-0.0469	0.8619	-0.1331	0.8157	-0.1648	0.4969	-0.0317	0.9224
CG14495	CG14495	1636005_at	0.0116	0.9891	0.0709	0.6213	-0.2641	0.2356	-0.2698	0.7576	-0.4191	0.2669	-0.1493	0.7074	0.1920	0.9333	-0.0639	0.9586	-0.2558	0.7643
CG17766 /// DereCG17766	CG17766	1636006_at	-0.5874	0.1299	0.1525	0.7759	-0.0776	0.7058	-0.4861	0.1883	-0.2316	0.3052	0.2545	0.2026	-0.1196	0.9636	0.5065	0.4875	0.6262	0.3898
Ance-2	Ance-2	1636007_at	-0.0409	0.8152	-0.1685	0.4129	-0.0428	0.8111	0.1470	0.7135	0.2485	0.1707	0.1015	0.5763	0.0018	0.9994	0.0929	0.7971	0.0911	0.7907
CG31828	CG31828	1636008_at	-0.0571	0.7759	0.0688	0.5542	0.0047	0.9839	-0.0335	0.9569	-0.0875	0.6391	-0.0540	0.7689	0.0475	0.9621	0.1116	0.7299	0.0642	0.8608
CG12910	CG12910	1636009_at	-0.0507	0.9308	-0.1031	0.6940	0.0239	0.9369	0.0209	0.9917	-0.3922	0.3616	-0.4131	0.2778	-0.1630	0.8971	-0.4658	0.3066	-0.3028	0.5403
---	---	1636010_at	0.0303	0.8976	-0.1783	0.5241	0.0342	0.8713	0.1230	0.8362	0.2506	0.2665	0.1276	0.5660	0.0314	0.9775	-0.0345	0.9353	-0.0659	0.8467
CG1308	CG1308	1636011_at	0.0969	0.5164	0.0153	0.8976	0.1674	0.3927	-0.1147	0.8735	0.1162	0.6845	0.2308	0.3106	0.1048	0.8380	0.1891	0.3623	0.0843	0.7262

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG13437 /// DafsCG13437	CG13437 /// GA12	1636012_at	0.1859	0.3219	0.0182	0.8542	0.3472	0.0550	0.2268	0.6406	0.1641	0.5134	-0.0627	0.8150	-0.1481	0.7726	-0.1467	0.5209	0.0014	0.9977
Actn3	open reading fram	1636013_at	0.2953	0.1025	0.1030	0.5431	0.0455	0.8822	-0.0146	0.9857	0.2670	0.1436	0.2816	0.0862	0.0437	0.9657	0.0810	0.8199	0.0373	0.9211
CG31419	CG31419	1636014_at	0.1922	0.2482	0.3829	0.0448	-0.0658	0.6769	-0.0636	0.9249	0.0504	0.8479	0.1140	0.5704	0.1958	0.6749	0.1348	0.4843	-0.0610	0.7851
CG32850	CG32850	1636015_s_at	0.8270	0.5491	-3.0152	0.0398	-1.6032	0.0017	1.6543	0.0144	2.6437	0.0002	0.9894	0.0071	0.2347	0.9767	-1.2234	0.5411	-1.4581	0.4627
CG15510	CG15510	1636016_at	-0.0903	0.5947	0.1946	0.2993	0.1480	0.4700	-0.1519	0.7949	-0.2863	0.2289	-0.1343	0.5695	-0.0539	0.9305	0.0381	0.8991	0.0920	0.6728
---	---	1636017_at	0.0489	0.7833	0.1951	0.1961	-0.1709	0.3647	-0.0826	0.8645	-0.0156	0.9517	0.0670	0.7135	0.1830	0.7953	0.2272	0.4424	0.0442	0.9142
---	---	1636018_at	0.0683	0.6520	0.0782	0.5983	0.1422	0.3410	-0.1181	0.7596	-0.0512	0.8012	0.0669	0.7020	-0.0705	0.8903	0.0484	0.8546	0.1189	0.5587
CG14061 /// DyakCG14061	CG14061	1636019_at	0.1098	0.5168	0.0932	0.6177	0.1114	0.6349	-0.0140	0.9884	0.0467	0.8789	0.0607	0.8149	-0.2162	0.7147	-0.0270	0.9462	0.1892	0.4582
Adf1	nalyot	1636020_s_at	0.3186	0.0648	0.2634	0.2343	-0.1061	0.7032	-0.0540	0.9353	0.0118	0.9694	0.0657	0.7602	0.3082	0.7910	-0.1236	0.8551	-0.4318	0.3921
CG11666	CG11666	1636021_at	0.0205	0.9450	0.1765	0.2671	0.0587	0.7045	-0.0209	0.9744	-0.1152	0.4869	-0.0943	0.5420	0.0787	0.9291	0.0656	0.8721	-0.0131	0.9769
CG30267	CG30267	1636022_at	0.1180	0.4294	0.0615	0.7493	0.1052	0.6552	0.1233	0.8794	0.1384	0.6595	0.0151	0.9658	-0.0454	0.9689	-0.0343	0.9451	0.0111	0.9835
Cdep	Cdep	1636023_at	0.6802	0.1155	0.7848	0.1541	0.5619	0.0122	-0.0048	0.9956	0.2505	0.3887	0.2553	0.3238	0.1199	0.9588	0.1523	0.8678	0.0323	0.9751
CG40441	CG40441	1636024_at	-0.0312	0.8749	0.2513	0.3878	0.2810	0.1150	-0.0179	0.9749	-0.0513	0.7589	-0.0334	0.8346	0.0450	0.9776	0.1935	0.6414	0.1486	0.7362
Ras64B	ras oncogene	1636025_at	-0.5995	0.0269	-0.3334	0.2224	-0.7604	0.0011	-0.1726	0.5735	-0.2390	0.1262	-0.0664	0.6864	0.2203	0.7953	0.0736	0.8865	-0.1467	0.7154
beta-Spec	beta-spectrin	1636026_at	-0.5315	0.4148	-0.2663	0.6487	-0.5581	0.0145	0.1183	0.9035	0.4120	0.1897	0.2937	0.3059	0.4560	0.8421	0.7552	0.4182	0.2992	0.7899
DsimCG9245 /// Pis	CG9245	1636027_at	-0.0611	0.7245	0.0829	0.4895	0.0914	0.5739	0.1562	0.6409	0.1342	0.4277	-0.0220	0.9112	0.1923	0.6749	0.2058	0.2642	0.0135	0.9614
CG6723	CG6723	1636028_at	0.2215	0.2746	-0.0857	0.5437	0.0663	0.7274	0.1990	0.7453	0.2870	0.2897	0.0880	0.7660	0.1412	0.8222	0.0854	0.7968	-0.0559	0.8745
CG9172	CG9172	1636029_s_at	-0.2458	0.4206	0.6735	0.0235	0.5171	0.0433	-0.0144	0.9866	-0.9129	0.0018	-0.8985	0.0011	-0.0509	0.9775	-0.3988	0.3397	-0.3479	0.4285
CG32626	CG32626	1636030_s_at	0.4471	0.0301	-0.0054	0.9770	0.1651	0.3941	0.1532	0.6926	0.3712	0.0443	0.2179	0.1728	-0.0609	0.9168	-0.1221	0.5636	-0.0612	0.8039
CG1294	CG1294	1636031_at	-0.3005	0.3618	-0.9201	0.0071	-0.9222	0.0447	0.1399	0.9182	0.4835	0.2493	0.3437	0.3744	-0.1174	0.8909	-0.1172	0.7635	0.0002	0.9997
CG15327 /// DsimCG15327	CG15327	1636032_at	0.0639	0.7444	-0.0233	0.8718	0.0229	0.8970	0.1026	0.8738	0.0942	0.7185	-0.0084	0.9761	-0.0862	0.8956	-0.1344	0.6080	-0.0482	0.8866
CG31368	CG31368	1636033_at	0.2997	0.2453	0.0190	0.9713	-0.0971	0.6664	-0.1632	0.8046	0.4912	0.0655	0.6544	0.0132	-0.0837	0.9589	0.1884	0.7200	0.2721	0.5771
dom	domino	1636034_at	0.6129	0.0366	0.1328	0.6597	-0.1811	0.3677	-0.0835	0.8908	0.5481	0.0152	0.6316	0.0049	0.0315	0.9893	-0.0664	0.9325	-0.0979	0.8837
beat-IIa	beaten path IIa	1636035_at	-0.4209	0.0818	-0.1037	0.4614	-0.1014	0.6217	-0.0027	0.9985	-0.1190	0.7299	-0.1163	0.7091	-0.1360	0.8283	-0.0462	0.9085	0.0897	0.7715
CG1240 /// DyakCG1240	CG1240	1636036_at	-0.2539	0.1344	-0.2674	0.1329	-0.2597	0.0943	0.1164	0.7293	0.3020	0.0498	0.1856	0.1665	0.1090	0.8541	0.2445	0.3012	0.1356	0.6045
CG31195	CG31195	1636037_at	-0.3696	0.4291	0.9431	0.0575	0.8871	0.0031	0.0870	0.9466	-0.1339	0.7613	-0.2209	0.5387	0.0947	0.9653	1.2241	0.0591	1.1294	0.0893
CG1115	CG1115	1636038_at	-0.3360	0.0640	-0.1925	0.2277	-0.1296	0.5226	0.1235	0.7229	-0.2286	0.1437	-0.3521	0.0204	0.0926	0.9088	-0.1663	0.5825	-0.2588	0.3775
Arp53D	Actin-related prote	1636039_at	0.2180	0.1903	0.1524	0.1559	0.1920	0.3659	-0.0191	0.9803	0.0810	0.7123	0.1001	0.5999	-0.0359	0.9705	0.0007	0.9996	0.0366	0.9151
CG10527 /// DyakCG10527	CG10527	1636040_at	-1.0712	0.0107	-0.4261	0.1992	-0.8440	0.0085	0.0263	0.9744	-0.1153	0.5926	-0.1416	0.4477	0.4116	0.7485	0.4050	0.4655	-0.0066	0.9946
---	---	1636041_at	0.0425	0.8595	-0.0150	0.8895	0.1646	0.3971	0.2206	0.5680	0.1864	0.3543	-0.0342	0.8838	-0.0831	0.9267	-0.0247	0.9600	0.0584	0.8870
Gr23a	Gustatory recepto	1636042_at	0.2035	0.2773	0.0426	0.7722	0.0809	0.7554	0.0157	0.9872	0.1887	0.4356	0.1730	0.4299	0.1054	0.9046	0.1577	0.6473	0.0523	0.9055
---	---	1636043_at	0.1292	0.4076	-0.0704	0.4906	0.2297	0.3123	0.1015	0.8281	0.2672	0.1360	0.1657	0.3109	-0.0368	0.9705	0.0632	0.8552	0.1000	0.7296
Slh	SLY-1 homologou	1636044_at	0.3599	0.1733	1.0336	0.0190	1.3086	0.0001	0.2945	0.4633	0.1801	0.4233	-0.1144	0.6003	0.0665	0.9535	0.8764	0.0332	0.8100	0.0557
---	---	1636045_at	0.1207	0.4787	0.0396	0.7023	0.1508	0.3153	-0.1163	0.7425	-0.0978	0.5602	0.0185	0.9230	0.0111	0.9913	-0.0231	0.9462	-0.0342	0.9064
CG32499	CG32499	1636046_at	0.0930	0.7055	-0.0013	0.9948	-0.1000	0.6160	-0.0505	0.9436	0.1170	0.5996	0.1674	0.3737	0.1308	0.8680	-0.0118	0.9842	-0.1426	0.6763
tinc	tincar	1636047_s_at	-0.0957	0.6938	-0.0295	0.8813	-0.2430	0.2718	-0.0659	0.9117	-0.0831	0.7030	-0.0172	0.9417	-0.0463	0.9742	-0.2038	0.5882	-0.1575	0.6881
Sld5	Sld5	1636048_at	0.4168	0.0334	0.5356	0.0545	0.4374	0.0129	-0.0574	0.9098	0.0807	0.6570	0.1381	0.3503	-0.1367	0.8439	0.2573	0.3571	0.3940	0.1911
---	---	1636049_at	0.1046	0.6838	0.3212	0.3095	0.1284	0.5485	0.0090	0.9932	0.0613	0.8122	0.0523	0.8257	0.0457	0.9646	0.0443	0.9178	-0.0014	0.9979
comr	cookie monster	1636050_at	0.0076	0.9795	0.0703	0.6273	0.1642	0.3609	-0.0226	0.9777	-0.0579	0.8218	-0.0353	0.8850	-0.0106	0.9939	-0.0161	0.9773	-0.0055	0.9924
CG14456	CG14456	1636051_at	-0.0710	0.7472	0.1113	0.5610	0.6166	0.0080	0.0037	0.9956	-0.3973	0.0440	-0.4010	0.0271	-0.2851	0.7230	-0.0739	0.8831	0.2112	0.5676
l(1)dd4	gamma-tubulin rin	1636052_at	-0.5193	0.0345	-0.3824	0.3704	-0.4217	0.0291	-0.0738	0.8950	0.1771	0.3489	0.2509	0.1292	0.0023	0.9994	0.2705	0.5835	0.2682	0.5892
CG30044	CG30044	1636053_s_at	0.2031	0.3650	-0.2082	0.4493	0.0967	0.5926	0.1093	0.8050	0.2220	0.2070	0.1127	0.5081	-0.3475	0.4415	-0.2236	0.3295	0.1239	0.6259
CG33678	CG33678	1636054_at	0.1042	0.6853	-0.1377	0.3083	-0.1068	0.5826	0.2223	0.6652	0.1164	0.6772	-0.1059	0.6798	-0.0932	0.9088	-0.1841	0.5403	-0.0909	0.7931
Hsc70-2	Heat shock protei	1636055_at	-0.6654	0.0207	-0.0137	0.8975	0.0151	0.9700	-0.3667	0.4428	-0.7487	0.0113	-0.3821	0.1006	-0.1673	0.8292	-0.0476	0.9265	0.1197	0.7494
RpL18A	ribosomal protein	1636056_at	0.4288	0.0158	1.4154	0.0113	1.4758	0.0003	0.2145	0.5680	-0.7019	0.0039	-0.9164	0.0008	0.0652	0.9092	-0.0527	0.8493	-0.1180	0.5866
CG9572	CG9572	1636057_at	-2.1595	0.0133	-2.9945	0.0440	-2.8402	0.0008	0.8487	0.3405	1.7861	0.0048	0.9374	0.0465	0.6280	0.8386	0.8647	0.5100	0.2366	0.8921
CG31439	CG31439	1636058_at	0.3709	0.1097	0.0673	0.7492	-0.0143	0.9575	0.1660	0.6785	0.1957	0.3010	0.0297	0.8951	0.1857	0.8283	-0.1467	0.7206	-0.3325	0.3671
CG9689	CG9689	1636059_at	-1.9517	0.0008	-2.1029	0.0108	-2.3795	0.0001	0.0947	0.8856	0.3861	0.0852	0.2914	0.1432	0.0282	0.9913	0.0645	0.9404	0.0362	0.9636
---	---	1636060_at	0.0944	0.5477	-0.1906	0.5349	0.3137	0.1967	0.3539	0.3553	0.2209	0.3188	-0.1330	0.5340	0.0091	0.9933	-0.0375	0.9231	-0.0466	0.8918
---	---	1636061_at	-0.1863	0.6553	-0.1327	0.7275	-0.9478	0.0043	-0.2827	0.4586	0.0171	0.9554	0.2998	0.1036	0.4137	0.7812	-0.0614	0.9528	-0.4751	0.4764
amos	Rough eye	1636062_at	0.3198	0.1135	0.0575	0.7093	0.1124	0.4796	0.0492	0.9380	0.1378	0.4778	0.0886	0.6422	0.1419	0.8122	0.0594	0.8634	-0.0825	0.7774
CG34355 /// DbuzCG12492	CG12492 /// GA11	1636063_at	-2.4798	0.0053	-1.8108	0.0674	-1.9233	0.0005	0.0706	0.9803	-0.1645	0.8532	-0.2350	0.7540	0.0293	0.9913	0.3400	0.5673	0.3107	0.6110
sec8	sec8	1636064_at	-0.7397	0.0747	0.1792	0.3939	0.0937	0.6479	0.0177	0.9854	-0.3466	0.1129	-0.3643	0.0663	0.0895	0.9589	0.5865	0.2130	0.4971	0.3175
---	---	1636065_at	-0.0171	0.9297	0.0680	0.6244	0.1829	0.3937	-0.1955	0.7774	-0.1129	0.7399	0.0826	0.7989	0.0493	0.9545	0.0806	0.8048	0.0312	0.9299
---	---	1636066_s_at	-0.1233	0.6091	0.3113	0.2579	0.2144	0.4736	-0.2325	0.5859	-0.3729	0.0875	-0.1404	0.5026	0.0840	0.9467	0.5575	0.1408	0.4734	0.2338
CG5934	CG5934	1636067_at	-0.0651	0.7574	0.2845	0.2838	-0.0563	0.8008	0.2492	0.4142	0.4814	0.0119	0.2322	0.1240	0.4954	0.5754	0.6581	0.1062	0.1627	0.7191
AGO2	Argonaute2	1636068_a_at	0.0757	0.6675	-0.3035	0.1811	-0.2468	0.2499	-0.1110	0.8231	0.4245	0.0347	0.5355	0.0080	-0.1159	0.8760	0.1559	0.6156	0.2718	0.3626
---	---	1636069_at	0.0881	0.6838	0.0037	0.9781	0.0453	0.7868	0.0710	0.8872	0.0635	0.7528	-0.0076	0.9713	-0.0247	0.9824	-0.0823	0.8045	-0.0576	0.8672
CG33911	CG33911	1636070_at	0.1679	0.4753	0.0399	0.6915	0.1214	0.5871	0.2199	0.5719	0.2732	0.1669	0.0533	0.8123	0.1571	0.7726	0.2869	0.2023	0.1298	0.6040
Cbp80	cap binding protei	1636071_a_at	0.4697	0.1730	0.6310	0.1315	0.4987	0.0176	-0.0795	0.9314	0.3129	0.2524	0.3925	0.1066	0.0285	0.9898	0.4962	0.2698	0.4678	0.3262
---	---	1636072_at	0.2957	0.2349	0.0927	0.5998	-0.0192	0.9303	-0.0378	0.9477	0.0056	0.9816	0.0435	0.8127	-0.0165	0.9898	-0.1210	0.6891	-0.1045	0.7371
CG32344	CG32344	1636073_at	0.2901	0.3023	-0.2674	0.6488	-0.0832	0.6751	0.1727	0.7380	0.6638	0.0107	0.4911	0.0242	-0.0238	0.9916	0.1423	0.8445	0.1661	0.7980
beat-Vb	beat-Vb	1636074_at	-0.0393	0.8800	0.0670	0.6257	0.1614	0.5829	-0.1447	0.8294	-0.2009	0.4539	-0.0561	0.8510	-0.2327	0.7644	-0.1370	0.7184	0.0957	0.8173
CG31453	CG31453	1636075_at	-0.5042	0.4648	-0.8213	0.3584	-0.9727	0.0335	-0.2710	0.5726	0.0685	0.8238	0.3395	0.1204	-0.1773	0.9741	-0.3138	0.8678	-0.1365	0.9426
tutl	turtle	1636076_a_at	0.0185	0.9220	0.3239	0.1121	0.0999	0.6236	-0.0643	0.9154	0.0781	0.7234	0.1424	0.4222	0.0017	0.9990	0.1477	0.4757	0.1460	0.4876
---	---	1636077_at	0.0258	0.9199	-0.0299	0.7958	0.0502	0.8161	-0.0252	0.9744	0.0556	0.8113	0.0808	0.6825	0.0049	0.9978	0.0583	0.9266	0.0534	0.9230
CG13670	CG13670	1636078_at	0.1717	0.3695	-0.0541	0.5926	0.1245	0.4539	0.1057	0.8650	0.0516	0.8578	-0.0540	0.8296	-0.0606	0.9199	-0.1500	0.4694	-0.0894	0.6941
---	---	1636079_at	0.0755	0.7367	0.1255	0.3855	0.0641	0.6767	0.0630	0.9030	0.1305	0.4548	0.0675	0.7028	-0.0874	0.8815	0.1145	0.6384	0.2019	0.3807
jim	jim	1636080_a_at	-0.2676	0.6500	0.0760	0.8841	-0.2483	0.3184	-0.1530	0.8070	-0.2091	0.4155	-0.0562	0.8454	0.0868	0.9823	0.0237	0.9910	-0.0631	0.9622

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Ino80	Ino80	1636081_at	0.3014	0.0984	0.1771	0.3983	0.1989	0.1908	-0.0103	0.9909	0.1501	0.4601	0.1604	0.3734	0.0932	0.8349	0.0979	0.6254	0.0047	0.9874
CG14212	CG14212	1636082_at	-0.1687	0.4359	-0.0385	0.8371	-0.0228	0.9118	0.2227	0.6371	0.1941	0.4170	-0.0286	0.9198	0.0567	0.9487	-0.0357	0.9326	-0.0923	0.7634
---	---	1636083_at	0.0558	0.7490	0.0218	0.8569	0.1054	0.6562	0.0235	0.9682	0.1485	0.3435	0.1250	0.3814	0.0056	0.9956	0.1864	0.4380	0.1809	0.4683
CG6497	CG6497	1636084_at	0.4910	0.0384	0.0955	0.5884	0.2949	0.1534	0.1666	0.6010	-0.0006	0.9979	-0.1672	0.2459	0.0342	0.9826	-0.0710	0.8949	-0.1052	0.8150
CG18143	CG18143	1636085_at	0.6071	0.0058	0.5644	0.0318	0.8970	0.0013	0.0964	0.8423	-0.2739	0.1265	-0.3703	0.0290	-0.2088	0.6955	-0.1918	0.3842	0.0170	0.9582
CG14584	CG14584	1636086_at	0.0126	0.9429	0.1291	0.4126	0.2061	0.2212	-0.0452	0.9470	-0.1569	0.4304	-0.1116	0.5554	0.0654	0.9092	0.0888	0.7000	0.0234	0.9345
CG31872 /// DsimCG31872	CG31872	1636087_at	0.1175	0.4472	-0.0126	0.9055	0.0399	0.8452	0.0455	0.9255	0.0293	0.8808	-0.0161	0.9281	-0.1171	0.8330	-0.0759	0.7932	0.0412	0.8964
Dll	Distalless	1636088_at	-1.0916	0.0051	0.0465	0.8270	-0.6504	0.0136	-0.3931	0.5735	-1.2476	0.0048	-0.8545	0.0156	-0.0560	0.9585	-0.0749	0.8550	-0.0189	0.9661
comm	commissureless	1636089_at	-0.5979	0.1668	0.0054	0.9903	-0.7260	0.0013	-0.2650	0.6122	-0.8495	0.0067	-0.5844	0.0210	0.3170	0.7953	-0.4819	0.3340	-0.7989	0.1447
sv	sparkling	1636090_a_at	0.0203	0.9539	-0.0238	0.8988	0.0960	0.5935	-0.0685	0.8914	-0.1021	0.5746	-0.0336	0.8637	-0.0083	0.9935	-0.0923	0.7293	-0.0840	0.7514
fj	four-jointed	1636091_at	0.3628	0.5469	0.7178	0.0349	1.0507	0.0004	-0.1039	0.9496	-0.6285	0.1676	-0.5245	0.1996	-0.3218	0.7492	-0.3269	0.4475	-0.0051	0.9946
wapl	parallel sister chrc	1636092_a_at	-0.5490	0.2649	0.2182	0.8091	0.3110	0.1605	-0.0990	0.9048	-0.4283	0.1100	-0.3294	0.1695	-0.2625	0.9246	0.2546	0.8385	0.5170	0.6052
ERR	estrogen-related r	1636093_a_at	0.4659	0.0412	0.1126	0.7093	0.1714	0.3659	-0.1911	0.7949	0.1845	0.5685	0.3756	0.1596	-0.3738	0.6749	-0.3141	0.3761	0.0597	0.9055
CG14838 /// DmirCG14838	CG14838	1636094_at	-0.0993	0.5604	0.1102	0.3623	0.0191	0.9414	-0.0808	0.8534	-0.1605	0.3146	-0.0798	0.6189	0.0303	0.9816	-0.0073	0.9918	-0.0376	0.9278
CG4619	CG4619	1636095_at	0.1618	0.7632	-0.2937	0.2916	-0.3509	0.1006	0.1118	0.9308	0.2682	0.4985	0.1564	0.6922	-0.0952	0.9571	-0.5667	0.2430	-0.4716	0.3626
CG8924	CG8924	1636096_at	-0.1912	0.5820	-0.9679	0.0899	-1.3513	0.0261	-0.4993	0.2729	1.0880	0.0025	1.5873	0.0003	-0.2596	0.9223	-0.0825	0.9535	0.1772	0.8837
CG14243	CG14243	1636097_at	0.2133	0.2773	0.0349	0.7297	-0.0124	0.9552	-0.0814	0.9011	-0.0340	0.9079	0.0473	0.8471	-0.0195	0.9816	-0.2119	0.2195	-0.1924	0.2960
CG5144	CG5144	1636098_at	-0.0720	0.8252	0.3330	0.1080	0.5449	0.0080	-0.2599	0.3454	-0.6170	0.0030	-0.3571	0.0201	-0.4301	0.3836	0.0382	0.9296	0.4683	0.1192
mbl	mindmelt	1636099_s_at	0.2343	0.1818	0.1292	0.5189	-0.8655	0.0245	-0.5352	0.3172	0.5738	0.0692	1.1090	0.0026	0.3472	0.7464	0.2269	0.6446	-0.1203	0.8334
CG32416	CG32416	1636100_at	0.1867	0.2429	0.1328	0.4654	0.0265	0.8914	0.0331	0.9603	0.1389	0.4539	0.1058	0.5447	-0.1226	0.8202	-0.0369	0.9156	0.0857	0.7409
CG31729	CG31729	1636101_a_at	-0.0271	0.9338	0.4533	0.0653	0.3393	0.0941	0.1062	0.8507	0.0277	0.9231	-0.0786	0.7200	0.1269	0.9031	0.4176	0.2522	0.2906	0.4597
CG8671	CG8671	1636102_at	0.2017	0.4360	0.0951	0.4371	0.0365	0.8403	0.1250	0.7229	0.1808	0.2570	0.0558	0.7452	0.0328	0.9816	-0.0963	0.8222	-0.1291	0.7307
nocturnin	Dnocturnin	1636103_a_at	0.6245	0.0535	0.0666	0.8805	0.5521	0.0062	0.1373	0.8508	-0.3276	0.2168	-0.4649	0.0549	-0.4104	0.7152	-0.9556	0.0637	-0.5452	0.2619
---	---	1636104_at	-1.3113	0.0923	-1.1390	0.1974	-2.0820	0.0001	-0.5481	0.2654	-0.2894	0.3269	0.2587	0.3307	0.1154	0.9816	-0.3047	0.8258	-0.4201	0.7269
CG13675	CG13675	1636105_at	0.1400	0.3736	-0.1081	0.4984	-0.1029	0.6507	0.2022	0.6854	0.2972	0.1977	0.0951	0.6964	0.0710	0.9306	-0.0256	0.9517	-0.0966	0.7499
alt	CG18212	1636106_s_at	0.5602	0.1189	0.6280	0.0535	-0.2880	0.1391	-0.2663	0.3127	0.5624	0.0039	0.8287	0.0005	0.6556	0.5754	0.8046	0.1260	0.1490	0.8189
CG14440	CG14440	1636107_at	-0.2457	0.5472	-0.4184	0.1833	-0.4575	0.0398	0.1056	0.8155	0.4124	0.0281	0.3067	0.0578	0.0200	0.9939	0.2326	0.7200	0.2126	0.7442
CG17033	CG17033	1636108_at	-0.0659	0.6812	-0.0257	0.8439	0.0042	0.9852	-0.0258	0.9649	0.0524	0.7839	0.0782	0.6250	-0.0212	0.9860	0.0825	0.8132	0.1037	0.7381
CG15458	CG15458	1636109_at	0.1111	0.6232	0.2201	0.0688	0.4095	0.0715	0.0695	0.9375	0.0120	0.9761	-0.0575	0.8470	0.0932	0.8425	0.0432	0.8754	-0.0500	0.8379
---	---	1636110_at	-0.0705	0.7097	-0.2509	0.1267	-0.1237	0.7338	0.2173	0.7838	0.6034	0.0677	0.3862	0.1851	-0.0824	0.9445	0.0997	0.8352	0.1822	0.6366
---	---	1636111_at	0.1532	0.3355	0.0013	0.9924	-0.2128	0.2247	-0.1592	0.7380	0.1550	0.4822	0.3142	0.0949	-0.0553	0.9405	-0.0734	0.7999	-0.0181	0.9552
CG33521	anon-fast-evolving	1636112_s_at	0.2743	0.6643	0.6548	0.0588	-0.1814	0.4092	-0.1858	0.8512	-0.3532	0.3348	-0.1674	0.6531	0.6808	0.6749	0.2794	0.7158	-0.4014	0.5744
---	---	1636113_at	0.2822	0.3281	0.2358	0.3094	0.0900	0.5691	-0.0371	0.9610	0.0179	0.9531	0.0550	0.8104	0.0368	0.9816	-0.0381	0.9469	-0.0748	0.8746
---	---	1636114_at	0.4045	0.1245	0.4270	0.1815	0.5109	0.0140	0.0289	0.9753	-0.2985	0.1843	-0.3273	0.1047	-0.0097	0.9939	-0.1032	0.7671	-0.0934	0.7838
---	---	1636115_at	-0.0121	0.9407	0.1454	0.4047	-0.2391	0.1441	-0.1354	0.7121	-0.1576	0.3536	-0.0222	0.9127	0.1939	0.7230	0.0771	0.7945	-0.1168	0.6443
---	---	1636116_s_at	0.2056	0.3100	-0.1798	0.2013	0.0322	0.9332	0.1318	0.8573	0.2469	0.3564	0.1151	0.6749	0.0005	0.9998	-0.1072	0.6966	-0.1077	0.6901
---	---	1636117_at	0.2096	0.2788	0.0551	0.6160	-0.0015	0.9959	0.1300	0.7949	0.2646	0.1916	0.1346	0.4903	0.0687	0.9445	-0.0094	0.9877	-0.0781	0.8362
CG13065	CG13065	1636118_at	0.5428	0.0508	0.1154	0.4761	0.3419	0.1442	0.0289	0.9759	0.1496	0.5527	0.1208	0.6096	-0.0936	0.8909	-0.2284	0.3669	-0.1349	0.6270
CG1468	CG1468	1636119_at	2.7001	0.0055	0.8778	0.3829	2.6974	0.0003	1.3921	0.1119	1.1523	0.0348	-0.2398	0.6586	-0.4491	0.8885	-0.5485	0.6842	-0.0994	0.9533
---	---	1636120_at	0.0140	0.9425	0.2140	0.1884	0.0091	0.9629	-0.1251	0.8074	-0.0797	0.7411	0.0454	0.8482	0.1045	0.8480	0.0790	0.7778	-0.0256	0.9360
Lsm11	CG12924	1636121_at	0.1971	0.5698	0.0142	0.9345	0.1775	0.4035	-0.1207	0.8621	-0.0316	0.9278	0.0891	0.7384	-0.1707	0.8628	-0.2745	0.5032	-0.1038	0.8372
Socs16D	Suppressor of Cyt	1636122_at	-1.0942	0.0609	0.1903	0.7894	0.2546	0.1350	-0.2752	0.6506	-1.1135	0.0033	-0.8383	0.0070	-0.3228	0.9007	0.1580	0.9144	0.4809	0.6388
CG7870	CG7870	1636123_at	0.8905	0.0077	1.2249	0.0140	1.8226	0.0000	0.2369	0.3863	0.0253	0.9058	-0.2116	0.1243	-0.3572	0.6984	0.3167	0.4103	0.6739	0.1179
---	---	1636124_at	0.1577	0.5986	0.0142	0.8896	0.1253	0.4715	0.0337	0.9766	-0.0017	0.9965	-0.0354	0.9187	-0.1853	0.7677	-0.0876	0.7932	0.0977	0.7495
granny-smith	granny smith	1636125_a_at	0.2262	0.2513	0.7736	0.0195	0.9005	0.0009	-0.0203	0.9860	-0.4394	0.1086	-0.4191	0.0880	-0.2929	0.6741	-0.0255	0.9556	0.2673	0.3543
CG8290	CG8290	1636126_at	-0.1058	0.8327	0.0667	0.8582	-0.2835	0.0940	-0.3493	0.4653	0.1852	0.4958	0.5344	0.0286	-0.0071	0.9978	0.2672	0.6686	0.2744	0.6536
---	---	1636127_at	0.3057	0.3856	-0.0200	0.9196	0.2096	0.3466	-0.0335	0.9620	0.0245	0.9252	0.0580	0.7806	0.0708	0.9589	-0.0153	0.9838	-0.0861	0.8673
CG13838	CG13838	1636128_at	0.2452	0.2650	0.1347	0.3828	0.0248	0.9049	0.0121	0.9866	0.0988	0.6109	0.0867	0.6297	0.1855	0.7644	0.0521	0.8940	-0.1334	0.6389
CG3645	CG3645	1636129_s_at	0.0089	0.9653	-0.3839	0.1594	-0.1541	0.3549	0.1376	0.6633	0.3858	0.0190	0.2481	0.0692	-0.0768	0.9011	0.1153	0.6394	0.1921	0.4054
Arf72A	Arflike at 72A	1636130_at	0.3390	0.1045	0.4996	0.1224	0.3159	0.0488	0.1496	0.7028	0.2672	0.1368	0.1176	0.4974	0.3506	0.6749	0.4898	0.1418	0.1392	0.7157
lig3	DNA ligase III	1636131_at	0.3846	0.0230	0.5152	0.2440	0.5175	0.0205	0.0598	0.8899	0.3032	0.0424	0.2434	0.0634	0.0646	0.9640	0.4438	0.2371	0.3792	0.3411
---	---	1636132_at	1.4460	0.0473	0.7392	0.3089	-0.0043	0.9944	-0.5327	0.5311	1.3869	0.0079	1.9196	0.0011	0.4155	0.8610	0.5700	0.5755	0.1545	0.9077
---	---	1636133_at	0.1547	0.5309	-0.0795	0.7555	0.2566	0.1119	0.1878	0.5893	0.2774	0.1149	0.0896	0.6167	-0.2796	0.6955	-0.0808	0.8431	0.1988	0.5304
Or83b	Odorant receptor	1636134_at	-0.0376	0.8378	0.0303	0.8195	0.0069	0.9792	-0.0233	0.9838	0.1460	0.6206	0.1693	0.5098	0.2265	0.7741	0.1976	0.5855	-0.0289	0.9523
CG13931	CG13931	1636135_at	-0.1422	0.6180	0.4023	0.1968	0.1987	0.2743	-0.2261	0.7028	-0.3151	0.2483	-0.0891	0.7663	0.0728	0.9390	0.1724	0.5778	0.0996	0.7729
---	---	1636136_at	0.0863	0.5521	-0.0216	0.9187	-0.0298	0.8598	0.0986	0.8265	0.2218	0.1988	0.1232	0.4516	0.0450	0.9449	-0.0183	0.9542	-0.0633	0.7886
dare	ADXRDTASE	1636137_at	0.2543	0.2187	0.5639	0.2332	0.7655	0.0165	-0.0019	0.9985	0.1325	0.5656	0.1344	0.5165	-0.2698	0.8305	0.5115	0.3206	0.7814	0.1641
CG12509	CG12509	1636138_at	0.0398	0.8693	-0.0211	0.8374	0.0821	0.6458	-0.0497	0.9311	-0.1986	0.2386	-0.1489	0.3334	0.1206	0.8215	-0.1016	0.6805	-0.2222	0.3271
---	---	1636139_at	0.0728	0.6866	0.0340	0.7158	-0.0809	0.5755	-0.0776	0.8526	-0.1125	0.4778	-0.0349	0.8419	0.0009	0.9994	-0.0375	0.8789	-0.0385	0.8655
CG3338	CG3338	1636140_at	0.0089	0.9796	0.1592	0.5698	0.2296	0.2833	0.0992	0.8604	0.1313	0.5460	0.0321	0.8950	0.1212	0.9237	0.4604	0.2507	0.3392	0.4258
gro	Groucho	1636141_s_at	-0.7333	0.1359	-0.3090	0.2256	-0.5549	0.0097	-0.3273	0.3722	-0.2346	0.2591	0.0927	0.6662	-0.0299	0.9914	0.0821	0.9296	0.1120	0.8897
CG7872	CG7872	1636142_at	0.0935	0.5969	0.4496	0.0307	0.7769	0.0160	0.2153	0.6988	-0.0197	0.9588	-0.2350	0.3116	-0.2291	0.8069	0.4096	0.2787	0.6387	0.1313
CG14720	CG14720	1636143_at	-0.1251	0.4183	-0.0969	0.6438	0.4093	0.0145	0.3858	0.3048	-0.0037	0.9903	-0.3895	0.0531	-0.0159	0.9848	0.0118	0.9701	0.0277	0.9161
CG10217 /// DyakCG10217	CG10217	1636144_s_at	-0.4395	0.0546	0.0212	0.9547	-0.1399	0.5755	-0.2143	0.5249	-0.1630	0.3658	0.0513	0.7962	-0.0347	0.9862	0.3245	0.4976	0.3592	0.4560
CG7219	CG7219	1636145_at	-1.3379	0.0049	-1.2383	0.0679	-0.2847	0.4310	0.5500	0.4221	0.1113	0.8123	-0.4387	0.1947	-0.1022	0.9589	0.5537	0.3152	0.6560	0.2622
Nrg	neuroglian	1636146_at	-0.7282	0.0292	-0.0110	0.9767	-0.9324	0.0079	-0.1703	0.6578	-0.1513	0.4286	0.0189	0.9343	0.7327	0.5665	0.5508	0.3268	-0.1819	0.7935
lectin-29Ca	Lectin29Ca	1636147_at	0.0359	0.8337	0.0213	0.9184	0.0157	0.9337	-0.0298	0.9586	0.0013	0.9951	0.0311	0.8599	-0.0011	0.9996	0.0283	0.9451	0.0295	0.9342
CG6553	CG6553	1636148_at	-1.8880	0.0397	-3.1959	0.0014	-3.2151	0.0001	0.1545	0.9558	0.9917	0.1845	0.8372	0.2129	-0.0866	0.9764	-0.6054	0.3780	-0.5187	0.4745
CG31705	CG31705	1636149_at	0.4024	0.3131	-0.4921	0.5478	-0.1089	0.5383	0.1681	0.5376	0.3572	0.0205	0.1891	0.1362	-0.2807	0.9168	-0.5123	0.6053	-0.2317	0.8460

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1636150_at	0.0477	0.8892	0.2647	0.3744	0.2279	0.1793	-0.0708	0.8909	-0.1601	0.3671	-0.0893	0.6111	-0.0853	0.9215	-0.0586	0.8893	0.0267	0.9472
CG32196	CG32196	1636151_at	-0.5287	0.0111	-0.6996	0.0162	-0.5356	0.0066	-0.1086	0.7795	0.2726	0.0916	0.3812	0.0163	-0.2678	0.5913	0.2012	0.3501	0.4691	0.0710
CG11138	CG11138	1636152_a_at	-0.4358	0.4968	-0.1038	0.8943	-1.0767	0.0167	-0.2849	0.5008	0.2111	0.3544	0.4961	0.0214	0.8495	0.7485	0.8654	0.4406	0.0158	0.9936
CG6984	CG6984	1636153_at	1.0804	0.0159	0.3686	0.0677	0.3750	0.0423	-0.2435	0.5869	0.1670	0.4852	0.4105	0.0494	-0.1621	0.8617	-0.4787	0.1900	-0.3166	0.4090
CG30090	CG30090	1636154_at	1.0617	0.0360	-0.0699	0.8840	0.6337	0.1391	0.5500	0.4409	0.4498	0.2490	-0.1002	0.8207	-0.2631	0.8991	-0.6093	0.4246	-0.3463	0.6817
CG14669	CG14669	1636155_at	0.1625	0.3126	0.0072	0.9659	0.1443	0.5205	-0.0234	0.9814	0.0281	0.9337	0.0515	0.8531	0.0318	0.9657	0.0211	0.9479	-0.0107	0.9723
CG7372	CG7372	1636156_at	0.9737	0.0591	0.3238	0.3839	0.2173	0.1632	-0.4884	0.4712	0.0270	0.9604	0.5155	0.1130	-0.3800	0.6728	-0.6017	0.1047	-0.2218	0.5642
CG16912	anon-fast-evolving	1636157_at	-0.2383	0.2442	-0.2101	0.4705	0.1341	0.6134	-0.0309	0.9777	0.1034	0.7482	0.1343	0.6271	-0.3448	0.7070	0.0456	0.9411	0.3904	0.3211
CG8492	CG8492	1636158_at	0.0614	0.6911	-0.4709	0.0659	-0.2176	0.3391	0.2886	0.3433	0.5291	0.0092	0.2405	0.1238	-0.1038	0.8873	-0.1113	0.7248	-0.0074	0.9865
---	---	1636159_at	0.3080	0.1746	0.1087	0.4974	0.1677	0.3285	0.0839	0.8678	-0.0005	0.9984	-0.0844	0.6453	0.0811	0.9201	0.0094	0.9865	-0.0716	0.8366
cdc23	cdc23	1636160_at	0.1856	0.2861	0.3537	0.2316	0.7328	0.0047	-0.1773	0.6522	-0.3381	0.0770	-0.1608	0.3570	-0.5860	0.4337	-0.0993	0.8539	0.4867	0.2292
CG3371	CG3371	1636161_at	-0.6605	0.0490	-0.1671	0.4830	-0.2329	0.1308	-0.0158	0.9883	-0.2707	0.2930	-0.2549	0.2678	0.0559	0.9635	0.1502	0.6864	0.0942	0.8209
CG10349	domain II	1636162_a_at	0.1066	0.6243	-0.0594	0.8437	0.0434	0.8128	0.2309	0.6086	0.2216	0.3359	-0.0093	0.9736	0.0801	0.9421	-0.0327	0.9525	-0.1128	0.7799
Arr1	arrestin	1636163_at	0.2579	0.5333	0.0406	0.7613	-0.0914	0.6222	-0.2085	0.7121	-0.0706	0.8277	0.1378	0.5917	0.0120	0.9952	-0.2330	0.6458	-0.2449	0.6268
CG3107	CG3107	1636164_s_at	0.1171	0.6187	0.5728	0.1934	0.1977	0.2535	-0.5359	0.1532	-0.4372	0.0585	0.0987	0.6776	-0.0887	0.9457	0.0010	0.9996	0.0896	0.8621
CG31002	CG31002	1636165_at	0.1806	0.4481	-0.1561	0.5338	0.0277	0.8915	0.1368	0.7431	0.1357	0.4795	-0.0011	0.9963	-0.0574	0.9611	-0.1545	0.6699	-0.0971	0.8112
---	---	1636166_at	-0.0108	0.9717	0.1021	0.5647	-0.1018	0.5226	-0.0723	0.9376	0.0033	0.9933	0.0756	0.8030	0.2110	0.7215	0.1881	0.4503	-0.0229	0.9474
---	---	1636167_at	0.0059	0.9748	0.1117	0.4621	0.1553	0.4636	0.1175	0.7409	-0.0746	0.6758	-0.1921	0.1688	0.0681	0.9409	-0.1373	0.6601	-0.2055	0.4851
CG13101	CG13101	1636168_s_at	0.6093	0.1110	1.3741	0.0179	0.8013	0.0057	-0.1384	0.7630	-0.0121	0.9686	0.1263	0.5056	0.3274	0.8215	0.7709	0.1874	0.4435	0.4764
---	---	1636169_at	0.0988	0.5637	0.2590	0.1301	0.1504	0.5113	-0.0933	0.8738	-0.2408	0.2398	-0.1475	0.4443	-0.0050	0.9963	0.0551	0.8685	0.0601	0.8418
CG13607	CG13607	1636170_at	2.4071	0.0070	2.5968	0.0263	3.3193	0.0001	0.2550	0.8550	-0.1093	0.8703	-0.3642	0.4430	-0.7285	0.7633	-0.1076	0.9477	0.6209	0.5615
Cpr47Ed	CG9076	1636171_at	0.0373	0.8979	-0.1404	0.5528	0.0165	0.9263	0.0784	0.8942	0.1332	0.5242	0.0548	0.8042	-0.0199	0.9898	-0.0325	0.9519	-0.0127	0.9830
CG12699 /// DyakCG12699	CG12699	1636172_at	0.1387	0.6227	0.0227	0.9173	-0.0188	0.9261	-0.0716	0.9068	0.0214	0.9398	0.0931	0.6461	-0.0340	0.9792	-0.1446	0.6599	-0.1106	0.7492
Mpk2	p38a MAP kinase	1636173_s_at	-0.5664	0.0086	-0.3551	0.3705	-0.7674	0.0225	-0.1801	0.5478	-0.0490	0.7968	0.1311	0.3579	0.0604	0.9793	0.0199	0.9844	-0.0405	0.9582
GstD9	Glutathione S tran	1636174_at	0.2798	0.2139	0.2129	0.4078	0.1106	0.7757	-0.2584	0.4094	0.2594	0.1341	0.5178	0.0058	-0.2039	0.8961	0.3129	0.6157	0.5168	0.3855
CG9084	CG9084	1636175_a_at	1.1069	0.2248	0.8628	0.2088	1.3633	0.0236	0.3408	0.8584	0.2697	0.7396	-0.0711	0.9338	-0.0853	0.9862	-0.1609	0.9265	-0.0756	0.9628
eIF-2beta	Eukaryotic initiatio	1636176_at	0.3371	0.1066	0.5206	0.0632	0.6947	0.0054	0.1743	0.6515	0.2683	0.1485	0.0940	0.6201	0.0775	0.9291	0.5383	0.0714	0.4608	0.1313
CG32394	CG32394	1636177_at	0.0824	0.6053	0.0240	0.8982	-0.0041	0.9869	0.0823	0.8209	0.1854	0.1890	0.1031	0.4397	0.1648	0.8222	0.0552	0.9064	-0.1096	0.7558
Scamp	anon-fast-evolving	1636178_at	0.3421	0.4036	0.5072	0.0500	0.4514	0.1043	-0.0985	0.8545	-0.0259	0.9235	0.0725	0.7259	0.2598	0.8692	0.3826	0.5613	0.1228	0.8872
CG31459	CG31459	1636179_at	0.5378	0.0087	0.1912	0.4032	0.3519	0.0437	0.0896	0.8690	0.1507	0.4484	0.0611	0.7725	-0.2321	0.7220	-0.1852	0.5214	0.0469	0.9031
yip2	yippee interacting	1636180_at	1.0736	0.0049	0.9054	0.0221	1.4025	0.0001	-0.0572	0.9474	-0.1440	0.5887	-0.0867	0.7424	-0.5514	0.3362	-0.3301	0.2627	0.2213	0.4846
CG18669	CG18669	1636181_at	0.0184	0.9483	-0.0381	0.8120	0.1054	0.4767	-0.0133	0.9860	0.0723	0.7247	0.0856	0.6347	-0.0217	0.9862	0.0562	0.8963	0.0779	0.8301
CG10948	CG10948	1636182_a_at	-0.0878	0.7263	0.0724	0.8101	-0.6232	0.0134	-0.4557	0.1717	0.0072	0.9809	0.4629	0.0186	0.2324	0.8140	0.2871	0.4917	0.0547	0.9227
CG32198	CG32198	1636183_at	0.0554	0.8155	-0.0091	0.9331	-0.2590	0.2233	-0.0764	0.9018	0.1629	0.4352	0.2393	0.1822	-0.0570	0.9552	-0.0083	0.9900	0.0486	0.9056
CG31813	CG31813	1636184_at	0.2368	0.7150	0.2521	0.1050	-0.2515	0.3275	-0.0917	0.9626	-0.5158	0.3275	-0.4241	0.3775	0.2109	0.8940	-0.3033	0.6380	-0.5142	0.3941
kay	shroud	1636185_at	0.1287	0.4753	0.3434	0.1494	0.1435	0.5827	0.0097	0.9894	-0.0444	0.8391	-0.0541	0.7728	0.4498	0.6695	0.2704	0.5426	-0.1794	0.7077
Fer2LCH	Ferritin	1636186_s_at	-0.0220	0.9030	-0.3044	0.0870	-0.2932	0.0566	-0.0525	0.9116	-0.0472	0.7977	0.0052	0.9778	0.0207	0.9898	-0.2805	0.4246	-0.3012	0.4016
CG33798	CG33798	1636187_at	0.2380	0.2073	0.0051	0.9826	0.1627	0.4653	-0.0061	0.9952	0.0219	0.9407	0.0280	0.9100	-0.1533	0.8093	-0.1806	0.5071	-0.0274	0.9412
---	---	1636188_at	0.1043	0.6359	0.0173	0.8773	0.2914	0.0919	-0.0628	0.8915	0.0361	0.8585	0.0988	0.5076	-0.1524	0.8215	-0.0944	0.7875	0.0580	0.8788
polo	Polo protein kinas	1636189_at	0.6192	0.6351	-1.9941	0.3011	-2.1691	0.0623	-0.5908	0.7225	2.4007	0.0071	2.9916	0.0016	-0.4295	0.9679	-0.2401	0.9590	0.1894	0.9639
CG13392	CG13392	1636190_at	-0.0179	0.9598	0.8409	0.0524	0.8070	0.0198	-0.4217	0.2109	-0.9992	0.0011	-0.5775	0.0071	-0.4411	0.7322	-0.1656	0.8174	0.2755	0.6422
CG16742	CG16742	1636191_a_at	-0.3559	0.2780	-0.2768	0.4466	-0.6843	0.0116	-0.1005	0.8794	0.4599	0.0478	0.5605	0.0135	0.0810	0.9545	0.3322	0.4242	0.2512	0.5699
CG18178 /// DyakCG18178	CG18178	1636192_at	-0.0280	0.9405	-0.5021	0.2186	-0.1385	0.6002	0.4541	0.4815	0.9705	0.0125	0.5163	0.0944	0.0927	0.9521	0.3582	0.4391	0.2655	0.5930
CG13893	CG13893	1636193_at	0.0836	0.7833	0.8042	0.0798	1.0764	0.0040	0.1735	0.7303	-0.8484	0.0034	-1.0218	0.0009	-0.2545	0.8270	-0.3490	0.4802	-0.0945	0.8874
---	---	1636194_s_at	-0.0793	0.6272	0.0059	0.9721	-0.0789	0.8320	0.0051	0.9956	-0.0164	0.9686	-0.0216	0.9494	-0.0947	0.8494	0.1131	0.6101	0.2078	0.3271
---	---	1636195_at	0.4730	0.2005	-0.0672	0.7288	0.0370	0.9139	0.0167	0.9900	0.1699	0.6049	0.1532	0.6130	-0.0572	0.9743	-0.2624	0.5800	-0.2052	0.6755
CG6891 /// CG6900	CG6900 /// CG688	1636196_s_at	-1.1485	0.0014	-0.5415	0.3780	-0.7585	0.2903	-0.0176	0.9852	0.4943	0.0289	0.5119	0.0157	0.1813	0.9717	1.0541	0.4016	0.8728	0.5083
CG31013	CG31013	1636197_at	0.1119	0.4702	-0.1216	0.4857	0.0101	0.9647	0.1689	0.6566	0.3153	0.0854	0.1465	0.3877	0.0374	0.9643	0.0052	0.9924	-0.0322	0.9176
CG1265	CG1265	1636198_at	-0.6932	0.4119	-1.1683	0.0269	-1.4570	0.0005	-0.0313	0.9653	-0.8435	0.0018	-0.8122	0.0012	-0.0686	0.9898	-1.5342	0.1646	-1.4656	0.2136
dpr15	dpr15	1636199_at	0.0810	0.6272	0.0334	0.7522	-0.1952	0.3078	-0.1149	0.8016	0.0410	0.8656	0.1558	0.3556	0.1567	0.7707	0.0023	0.9969	-0.1543	0.5173
---	---	1636200_at	-0.0976	0.5147	-0.1461	0.5026	-0.1418	0.5173	0.0351	0.9573	0.1721	0.3333	0.1370	0.4010	0.1259	0.8823	0.2122	0.5283	0.0863	0.8332
CG11865	CG11865	1636201_at	-0.0486	0.8562	-0.1730	0.4206	0.0006	0.9974	0.2155	0.6169	0.1255	0.5926	-0.0901	0.6902	-0.0705	0.8953	0.0184	0.9548	0.0889	0.6877
ImpL2	neural and ectode	1636202_s_at	0.7212	0.1062	0.3326	0.6893	0.4912	0.4901	0.3555	0.8671	0.7949	0.2934	0.4394	0.5508	0.2375	0.9409	0.5031	0.6401	0.2655	0.8310
B-H1	BarH1	1636203_at	0.0125	0.9555	-0.1812	0.2508	0.1664	0.2537	-0.0389	0.9380	-0.0421	0.8183	-0.0032	0.9863	-0.3160	0.5074	-0.1841	0.4103	0.1319	0.5834
---	---	1636204_at	0.2454	0.6872	0.2138	0.2216	-0.3656	0.2150	-0.7075	0.1279	0.1640	0.5893	0.8715	0.0047	0.0715	0.9836	0.6181	0.4408	0.5466	0.5141
CG14896	CG14896	1636205_at	-0.2264	0.6438	-0.5273	0.1673	-0.3849	0.2173	-0.0104	0.9937	0.3747	0.1349	0.3850	0.0872	-0.0876	0.9574	0.1146	0.8584	0.2022	0.6963
CG17737 /// DyakCG17737	CG17737	1636206_at	-0.0579	0.7739	-0.0604	0.7928	-0.1726	0.4316	0.1572	0.6972	0.0690	0.7562	-0.0882	0.6445	0.2398	0.7810	-0.0436	0.9425	-0.2834	0.4634
CG15435	CG15435	1636207_at	-0.1184	0.4895	0.3958	0.0372	0.3749	0.1518	0.0207	0.9688	-0.0909	0.5366	-0.1115	0.3806	0.0773	0.9396	0.2775	0.3669	0.2002	0.5448
CG15097	CG15097	1636208_a_at	-0.8245	0.0069	-0.7053	0.0190	-0.5579	0.0325	0.1693	0.6663	0.0586	0.7986	-0.1107	0.5470	-0.0683	0.9544	0.0005	0.9998	0.0688	0.8864
---	---	1636209_at	-0.0355	0.8365	0.2731	0.0549	0.1787	0.2695	-0.1910	0.5311	-0.3051	0.0596	-0.1140	0.4479	0.0933	0.8599	0.1877	0.3695	0.0944	0.6881
CG10191	CG10191	1636210_at	0.0167	0.9804	-0.1120	0.7124	-0.6415	0.0876	-0.2746	0.7663	-0.2510	0.5541	0.0236	0.9621	0.2393	0.8941	-0.2524	0.7525	-0.4917	0.4822
CG12782	CG12782	1636211_at	0.3061	0.1427	0.0982	0.4955	0.1088	0.6642	-0.1303	0.7825	-0.0489	0.8465	0.0814	0.6914	0.0004	0.9999	-0.0961	0.7272	-0.0965	0.7220
CG31824	CG31824	1636212_at	0.0687	0.7667	-0.1150	0.4654	-0.1734	0.2751	0.1205	0.8342	0.1293	0.5855	0.0088	0.9738	0.1023	0.8446	-0.1013	0.6686	-0.2036	0.3565
---	---	1636213_at	0.6604	0.0399	0.0118	0.9120	0.9211	0.0405	0.0816	0.9011	0.1441	0.5247	0.0625	0.7932	-0.8142	0.6557	-0.4936	0.5232	0.3207	0.7004
---	---	1636214_at	0.0592	0.7992	-0.0781	0.4809	0.2281	0.1333	0.1894	0.5642	0.1564	0.3643	-0.0331	0.8669	-0.2058	0.7423	-0.0669	0.8546	0.1390	0.6262
---	---	1636215_at	1.3518	0.0294	0.5530	0.0291	2.7797	0.0001	2.0764	0.0423	0.9156	0.1105	-1.1608	0.0315	0.0067	0.9950	0.0535	0.8807	0.0468	0.8915
CG12932	CG12932	1636216_at	0.3787	0.1322	0.2459	0.0990	0.0581	0.8525	0.0926	0.9186	0.1538	0.6180	0.0612	0.8495	0.1528	0.8379	0.1060	0.7798	-0.0468	0.9121
CG30503	CG30503	1636217_at	1.7159	0.0148	0.6721	0.2581	1.1629	0.0020	0.1338	0.9068	0.2533	0.5170	0.1195	0.7689	-0.3370	0.8680	-0.7978	0.3066	-0.4607	0.5913
CG7813	CG7813	1636218_at	0.0811	0.5934	0.0025	0.9876	0.2040	0.2392	0.1492	0.6936	0.1104	0.5578	-0.0387	0.8484	0.0839	0.8461	0.0158	0.9565	-0.0681	0.7394

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Ac13E	adenylyl cyclase	1636219_at	-1.4036	0.0011	-0.6663	0.1883	-1.5769	0.0000	-0.6275	0.0311	-0.8437	0.0008	-0.2162	0.1267	0.2286	0.8444	0.0128	0.9905	-0.2158	0.6826
CG10587	CG10587	1636220_at	0.2507	0.1192	0.2386	0.2554	-0.1256	0.5677	-0.1058	0.8395	0.0528	0.8319	0.1586	0.3806	0.0849	0.8972	0.0166	0.9670	-0.0683	0.8240
tkv	thick veins	1636221_a_at	1.9269	0.0011	1.7598	0.0108	2.0232	0.0000	-0.2797	0.6471	-0.0433	0.9167	0.2364	0.3900	-0.4324	0.6496	-0.1540	0.7299	0.2784	0.4850
Mef2	Complementation	1636222_at	-0.2410	0.1766	-0.1529	0.5757	0.0121	0.9564	0.0618	0.9435	0.0899	0.7613	0.0281	0.9258	-0.0915	0.9291	0.0592	0.9075	0.1507	0.6800
Hex-t1	Hexokinase	1636223_at	0.1046	0.7004	-0.0324	0.8984	-0.1844	0.4531	0.1017	0.8791	0.1961	0.4025	0.0944	0.6936	0.0666	0.9352	-0.0027	0.9969	-0.0693	0.8341
---	---	1636224_at	0.1886	0.2052	-0.0944	0.4375	-0.0689	0.7762	0.0482	0.9619	0.1585	0.5850	0.1103	0.6943	0.0338	0.9683	-0.0618	0.8380	-0.0956	0.7046
tan	tantalus	1636225_at	0.0322	0.9019	0.0761	0.7054	0.0641	0.7379	-0.0748	0.8590	0.2557	0.0925	0.3306	0.0230	-0.0961	0.9340	0.2888	0.4341	0.3848	0.3088
---	---	1636226_at	0.2988	0.0604	0.3477	0.1252	0.3017	0.1697	-0.0186	0.9774	-0.1639	0.3110	-0.1453	0.3197	0.1884	0.8049	-0.0300	0.9531	-0.2183	0.5067
CG8197	CG8197	1636227_at	0.3296	0.1667	0.0489	0.6935	-0.1706	0.2493	-0.1381	0.6578	0.1830	0.2207	0.3211	0.0247	-0.0025	0.9971	-0.0311	0.9054	-0.0286	0.9031
Cp36	defective chorion	1636228_at	-0.0278	0.9706	0.0800	0.5564	-0.1566	0.3176	-0.1210	0.8578	-0.2870	0.2379	-0.1660	0.4726	0.0865	0.9726	-0.2544	0.7275	-0.3409	0.6166
---	---	1636229_at	0.2174	0.2627	0.0600	0.6408	0.0543	0.7788	0.1530	0.7492	0.2012	0.3426	0.0482	0.8411	-0.0112	0.9884	0.0494	0.8213	0.0606	0.7527
---	---	1636230_at	0.0080	0.9715	-0.0408	0.7707	0.0779	0.6767	0.1875	0.6251	0.1572	0.4236	-0.0303	0.8945	0.0093	0.9914	0.0621	0.8091	0.0528	0.8336
CG3781	CG3781	1636231_at	0.6540	0.0233	0.5890	0.0248	0.7200	0.0024	0.0600	0.9228	0.1576	0.4154	0.0976	0.6062	-0.0593	0.9523	0.0336	0.9450	0.0929	0.7964
CG34391	CG15214	1636232_at	0.1235	0.5664	0.0486	0.6069	0.1730	0.4429	-0.0376	0.9647	-0.0580	0.8413	-0.0204	0.9418	-0.0596	0.9467	-0.0871	0.8031	-0.0275	0.9431
CG16995	CG16995	1636233_at	0.3181	0.1377	-0.1416	0.3258	-0.4327	0.0871	-0.2041	0.5931	0.2065	0.2897	0.4105	0.0257	-0.1420	0.8903	-0.2251	0.5835	-0.0831	0.8736
CG7366	CG7366	1636234_at	0.0264	0.8808	0.0345	0.8218	0.1803	0.3928	0.0103	0.9921	-0.0248	0.9337	-0.0351	0.8893	-0.2411	0.6824	-0.0735	0.8153	0.1677	0.5041
CG15754	CG15754	1636235_at	0.1455	0.5329	0.2628	0.1400	0.0195	0.9149	-0.1502	0.7647	-0.1401	0.5437	0.0101	0.9700	0.1142	0.8875	0.0674	0.8764	-0.0468	0.9107
CG2111	CG2111	1636236_at	0.3308	0.0685	0.1263	0.4267	0.0091	0.9663	0.0672	0.9108	0.1265	0.5319	0.0593	0.7781	0.1634	0.7506	0.1722	0.4317	0.0089	0.9809
l(1)G0222	lethal (1) G0222	1636237_s_at	-0.1195	0.5943	-0.1745	0.6112	-0.1598	0.5567	-0.2150	0.6099	-0.2482	0.2410	-0.0332	0.8961	-0.1013	0.9400	-0.1584	0.7439	-0.0571	0.9197
CG4911	CG4911	1636238_at	-0.0207	0.9542	-0.4583	0.0870	-0.8686	0.0221	-0.7790	0.1840	0.5969	0.0930	1.3759	0.0018	-0.1116	0.9467	0.1317	0.8493	0.2434	0.6581
---	---	1636239_at	0.1279	0.6287	-0.1355	0.5093	0.1528	0.5044	0.2023	0.6354	0.1408	0.5283	-0.0615	0.7932	-0.0844	0.9291	0.0705	0.8718	0.1549	0.6389
CG7678	CG7678	1636240_at	0.2021	0.3688	0.1279	0.2778	0.1214	0.6240	-0.0584	0.9376	-0.0176	0.9567	0.0409	0.8745	0.1088	0.8331	-0.0241	0.9451	-0.1329	0.5612
CG4738	CG4738	1636241_at	0.2201	0.4349	-0.0756	0.6151	-0.0340	0.8583	-0.0352	0.9620	0.5344	0.0140	0.5696	0.0064	0.0567	0.9717	0.1648	0.7273	0.1080	0.8348
Ilp6	Drosophila insulin	1636242_at	2.3972	0.0183	0.8400	0.2040	1.6797	0.0007	0.8710	0.1054	1.2056	0.0033	0.3346	0.2423	-0.0380	0.9939	-0.2161	0.8939	-0.1781	0.9057
Dcp1	Decapping proteir	1636243_at	-0.4437	0.1173	0.1340	0.3771	0.4204	0.0227	0.0319	0.9603	-0.2387	0.1581	-0.2706	0.0765	-0.1874	0.8270	0.3712	0.2878	0.5585	0.1479
CG5191	CG5191	1636244_s_at	-0.1664	0.6635	-1.2137	0.1129	-1.0340	0.0010	0.1684	0.6238	0.4722	0.0133	0.3038	0.0507	-0.0925	0.9774	-0.5973	0.4355	-0.5048	0.5325
T48	48' transcript	1636245_at	-1.5477	0.0017	-1.2078	0.0394	-1.3602	0.0005	-0.2234	0.6533	-1.0971	0.0015	-0.8736	0.0023	-0.2400	0.8889	-0.6189	0.3259	-0.3789	0.5820
CG8009 /// DyakCG8009	CG8009	1636246_a_at	-0.1330	0.4360	-0.5102	0.1327	-0.4554	0.0360	0.1466	0.7143	0.2331	0.2007	0.0865	0.6461	-0.0013	0.9996	-0.1503	0.6711	-0.1490	0.6690
CG32354	CG32354	1636247_at	-0.8845	0.0140	-0.5251	0.0103	-1.0827	0.0021	-0.0818	0.9436	-0.5498	0.0943	-0.4680	0.1097	0.2650	0.6955	-0.2852	0.3043	-0.5502	0.0891
CG15628	CG15628	1636248_at	-0.4540	0.0605	-0.3255	0.0440	-0.6063	0.0075	-0.2124	0.6086	-0.1166	0.6117	0.0958	0.6568	-0.1228	0.8320	-0.0357	0.9258	0.0871	0.7536
7B2	7B2	1636249_at	-1.4891	0.0154	-3.0629	0.0014	-2.5220	0.0002	0.8778	0.0976	1.1102	0.0043	0.2323	0.4260	0.2139	0.8953	-0.2542	0.7120	-0.4681	0.4558
---	---	1636250_at	0.2761	0.3395	0.6255	0.0510	0.3491	0.2968	-0.3026	0.6988	-0.6099	0.0901	-0.3073	0.3510	0.3456	0.8016	-0.1211	0.8843	-0.4667	0.4285
Tsp	Thrombospondin	1636251_at	0.4836	0.0736	0.0659	0.8124	0.2416	0.2675	0.1030	0.8392	0.0619	0.7899	-0.0411	0.8521	-0.1578	0.8386	-0.2107	0.5246	-0.0529	0.9048
---	---	1636252_at	-0.2907	0.1456	0.0539	0.7348	0.2459	0.2311	0.0456	0.9338	-0.2215	0.1662	-0.2670	0.0665	-0.1147	0.8940	0.0908	0.8319	0.2055	0.5454
CG9330	CG9330	1636253_at	-0.0527	0.7133	0.0484	0.7344	0.4282	0.0317	0.0988	0.7929	-0.0796	0.6425	-0.1784	0.1909	-0.2182	0.7036	0.0027	0.9961	0.2209	0.3683
rdx	roadkill	1636254_s_at	0.3201	0.3972	1.1626	0.0529	0.4949	0.0623	-0.0939	0.8946	-0.0676	0.8178	0.0263	0.9272	0.6057	0.6955	0.7027	0.2569	0.0970	0.9125
CG7997	CG7997	1636255_s_at	2.9821	0.0012	2.2781	0.0038	3.6178	0.0000	0.9635	0.1503	0.2427	0.5811	-0.7207	0.0512	-0.3824	0.7070	-0.4248	0.3066	-0.0424	0.9434
---	---	1636256_s_at	-0.0909	0.5610	-0.0035	0.9828	-0.0028	0.9909	0.0391	0.9423	0.0944	0.5708	0.0553	0.7393	0.0269	0.9774	0.0990	0.6912	0.0721	0.7841
CG10688	CG10688	1636257_at	1.5067	0.0007	1.1271	0.0220	1.3581	0.0001	0.1605	0.5842	0.7218	0.0012	0.5613	0.0021	-0.0874	0.9314	0.4514	0.1584	0.5389	0.1313
CG8550	CG8550	1636258_at	2.1335	0.0107	0.5758	0.1450	2.7190	0.0005	0.7372	0.0863	0.8856	0.0045	0.1484	0.5459	-1.1206	0.5905	-0.2672	0.8255	0.8534	0.3683
CG34050	CG34050	1636259_s_at	0.5369	0.0648	0.0383	0.7025	-0.1120	0.4979	-0.0703	0.9243	0.2168	0.3356	0.2872	0.1459	0.0499	0.9611	-0.1345	0.6747	-0.1844	0.5454
---	---	1636260_at	0.0762	0.6001	0.1422	0.6286	0.2953	0.1182	0.0296	0.9639	-0.0753	0.7061	-0.1048	0.5351	-0.1177	0.8494	-0.0070	0.9905	0.1107	0.7010
---	---	1636261_at	0.1308	0.5949	0.1124	0.5237	0.4392	0.0747	0.2290	0.5485	0.0284	0.9183	-0.2007	0.2587	-0.1218	0.8400	-0.1811	0.4743	-0.0593	0.8545
pall	pallbearer	1636262_at	-0.5542	0.0455	-0.5656	0.0162	-0.3950	0.0584	0.1891	0.6854	0.1898	0.3944	0.0007	0.9981	-0.1598	0.8594	0.0563	0.9209	0.2161	0.5767
---	---	1636263_s_at	0.2337	0.1455	-0.0565	0.6069	0.2256	0.3766	0.2209	0.5140	0.2250	0.2070	0.0041	0.9856	-0.0845	0.9405	-0.0758	0.8792	0.0086	0.9874
VAChT	VAChT	1636264_at	-0.0273	0.9167	-0.0619	0.6383	-0.1279	0.4796	-0.0307	0.9578	0.0633	0.7243	0.0941	0.5315	-0.0184	0.9885	-0.0559	0.8903	-0.0375	0.9220
Btk29A	fickle	1636265_s_at	-0.4152	0.3029	0.8793	0.0980	-0.1546	0.6392	-0.1399	0.7906	-0.2082	0.3409	-0.0683	0.7737	0.8238	0.6702	1.1954	0.1287	0.3715	0.6607
psh	persephone	1636266_at	-0.9568	0.1174	-2.0946	0.0142	-1.2902	0.0038	0.0465	0.9465	-0.0473	0.8522	-0.0937	0.6402	-0.7498	0.7215	-0.9766	0.2518	-0.2268	0.8412
CG13914	CG13914	1636267_at	0.4478	0.5308	0.0257	0.9233	-0.2626	0.5008	-0.0231	0.9727	0.3939	0.0269	0.4170	0.0133	-0.0426	0.9914	-0.1222	0.9315	-0.0796	0.9491
CG10570 /// DyakCG10570	CG10570	1636268_at	-0.9258	0.2534	0.4100	0.6292	-2.1398	0.0102	-1.6289	0.1853	-3.1990	0.0019	-1.5702	0.0263	0.7704	0.8108	-1.9390	0.1400	-2.7094	0.0790
CG33191	CG33191	1636269_at	0.2636	0.1557	0.0800	0.6048	0.2247	0.1483	-0.0268	0.9705	-0.0070	0.9804	0.0198	0.9313	-0.1513	0.7245	0.0087	0.9796	0.1600	0.3981
Prosalpha3T	Prosalpha3T	1636270_at	-0.0173	0.9343	0.0608	0.7037	0.2695	0.0749	0.1305	0.6958	0.1905	0.2147	0.0601	0.7134	-0.1247	0.8062	0.0067	0.9876	0.1315	0.5592
---	---	1636271_a_at	0.1906	0.2926	0.0682	0.7803	-0.0153	0.9432	0.3248	0.5128	0.5662	0.0373	0.2413	0.3110	0.3662	0.3587	0.3313	0.1237	-0.0348	0.9057
tey	teyrha-meyrha	1636272_at	0.0487	0.7752	-0.0146	0.9013	-0.3997	0.2804	-0.3208	0.6613	0.0525	0.9147	0.3733	0.2343	-0.0102	0.9914	-0.0702	0.7787	-0.0600	0.8119
Dpit47	DNA polymerase i	1636273_at	-0.6256	0.0157	0.2729	0.4570	0.4080	0.0758	0.2141	0.6088	-0.6109	0.0107	-0.8250	0.0017	0.0630	0.9657	0.4325	0.2573	0.3695	0.3644
wgn	Wengen	1636274_at	-2.2789	0.0004	-3.4152	0.0014	-3.7792	0.0000	-0.1732	0.8085	0.1241	0.7056	0.2972	0.2446	0.2733	0.7436	-0.8389	0.0391	-1.1122	0.0320
CG33528	CG33528	1636275_a_at	-0.3208	0.7950	0.8125	0.1774	-0.2956	0.3325	-0.5916	0.5008	-2.0308	0.0018	-1.4392	0.0049	0.4311	0.9238	-0.9651	0.5321	-1.3962	0.3614
Unr	Upstream of N-ras	1636276_at	-1.1115	0.0440	0.1527	0.8387	-0.2003	0.5765	-0.3988	0.5680	-0.3796	0.2920	0.0192	0.9663	0.0637	0.9884	0.9685	0.2982	0.9048	0.3569
---	---	1636277_at	0.2066	0.2368	0.0788	0.4922	0.2287	0.1851	0.0630	0.8960	0.0794	0.6588	0.0164	0.9340	-0.0061	0.9952	-0.0621	0.8445	-0.0560	0.8530
CG9850	Dlg-interacting me	1636278_at	0.0290	0.8973	0.2004	0.2808	-0.0628	0.6865	-0.1225	0.7825	-0.1683	0.3694	-0.0458	0.8264	0.1053	0.8882	0.0002	1.0000	-0.1051	0.7482
l(1)1Bi	lethal (1) 1Bi	1636279_at	0.4429	0.2814	-0.0236	0.9840	0.2815	0.1768	0.3471	0.5895	0.7922	0.0234	0.4451	0.1256	-0.1050	0.9752	0.4085	0.6533	0.5136	0.5615
CG12360	CG12360	1636280_s_at	-0.0418	0.9079	0.9226	0.0463	1.1356	0.0003	0.0657	0.9255	-0.1705	0.4383	-0.2361	0.2132	-0.1688	0.8767	0.8686	0.0622	1.0373	0.0523
Sema-1b	semiphorin	1636281_at	-0.3772	0.0708	-0.4710	0.2158	-0.7003	0.0041	-0.1043	0.8640	0.1440	0.5325	0.2483	0.1980	0.1918	0.7644	0.1035	0.7455	-0.0883	0.7838
Iwr	ubiquitin-conjugati	1636282_s_at	0.6104	0.0108	0.1061	0.5991	0.0011	0.9958	-0.0353	0.9508	0.3599	0.0309	0.3952	0.0132	0.1240	0.8661	-0.0229	0.9622	-0.1469	0.6414
CG14826	CG14826	1636283_at	0.0262	0.9220	0.2330	0.1797	0.1319	0.5774	-0.1413	0.7797	-0.2703	0.1960	-0.1290	0.5285	0.0153	0.9898	-0.0190	0.9649	-0.0343	0.9246
CG12432	CG12432	1636284_at	-0.0717	0.6640	0.2562	0.1141	0.3510	0.0493	-0.2234	0.5128	-0.3621	0.0500	-0.1386	0.4110	-0.1449	0.7644	-0.0894	0.6969	0.0555	0.8287
---	---	1636285_at	0.2866	0.1076	0.0363	0.7297	0.1968	0.2584	0.1389	0.6907	0.1807	0.2653	0.0418	0.8194	-0.0975	0.8439	-0.1236	0.5618	-0.0260	0.9261
CG7382	CG7382	1636286_at	-0.4224	0.1644	-0.6145	0.0375	-0.5518	0.0189	0.1500	0.7000	0.0713	0.7343	-0.0787	0.6728	-0.0363	0.9831	-0.1439	0.7578	-0.1076	0.8256
CG10592	CG10592	1636287_at	-0.0541	0.7640	0.0838	0.5120	0.0800	0.5879	-0.1259	0.7268	-0.2083	0.1947	-0.0824	0.6143	0.0474	0.9487	-0.0872	0.7346	-0.1346	0.5649

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
yuri	yuri gagarin	1636288_at	0.1510	0.5830	-0.1242	0.5112	-0.0613	0.8125	-0.1406	0.7031	0.0961	0.6064	0.2367	0.1177	-0.1648	0.7644	-0.1421	0.5636	0.0226	0.9449
DNApol-iota	DNApol-iota	1636289_s_at	-0.1334	0.5204	0.5620	0.0871	0.8110	0.0076	0.3360	0.4844	-0.3330	0.1916	-0.6690	0.0106	0.0291	0.9860	0.2118	0.5910	0.1826	0.6476
---	---	1636290_at	0.0634	0.6855	0.1446	0.3155	-0.0233	0.9176	-0.1315	0.7475	-0.0079	0.9775	0.1236	0.4645	0.0115	0.9914	0.1062	0.6949	0.0947	0.7307
Rae1	Rae1	1636291_at	-0.1929	0.1817	-0.4685	0.0628	-0.1703	0.3692	-0.0044	0.9956	0.1755	0.3671	0.1799	0.2973	-0.3385	0.5519	-0.0510	0.8967	0.2875	0.2879
Cyp313a4	Cyp313a4	1636292_at	0.3417	0.3627	0.2672	0.5910	0.4938	0.0091	0.0787	0.9465	0.0230	0.9637	-0.0557	0.8879	-0.1907	0.9246	-0.1100	0.9158	0.0808	0.9320
CG2217	CG2217	1636293_at	1.7539	0.1366	-0.2506	0.9041	0.7929	0.5148	1.5496	0.3660	2.8041	0.0112	1.2545	0.1484	0.5755	0.9467	0.8230	0.8101	0.2475	0.9469
---	---	1636294_at	0.1755	0.3869	-0.1414	0.3979	0.2352	0.2100	0.1022	0.8640	0.2007	0.3531	0.0985	0.6531	-0.1296	0.8680	-0.0663	0.8807	0.0633	0.8800
CG6796	CG6796	1636295_at	-0.0253	0.9083	-0.3710	0.1968	-0.5916	0.0058	-0.1129	0.7929	0.4708	0.0151	0.5837	0.0034	0.0318	0.9776	0.1972	0.4729	0.1655	0.5634
CG12945	CG12945	1636296_at	-0.3451	0.2282	0.6555	0.1455	0.5691	0.0512	-0.1478	0.8399	-0.7872	0.0112	-0.6394	0.0170	-0.3345	0.8141	0.0750	0.9363	0.4095	0.5023
CG3173	CG3173	1636297_at	0.2995	0.3255	0.0156	0.9413	-0.1061	0.5567	0.1678	0.6852	0.7843	0.0024	0.6165	0.0041	0.2528	0.8236	0.4376	0.3424	0.1849	0.7338
CG31093	CG31093	1636298_at	0.1345	0.3751	0.0642	0.6882	0.0736	0.6789	-0.1009	0.8578	0.0812	0.7312	0.1821	0.3224	-0.1241	0.8298	-0.1120	0.6764	0.0120	0.9752
CG4707	CG4707	1636299_at	0.0551	0.8693	-0.0415	0.7968	-0.0112	0.9522	0.1142	0.7973	0.1286	0.4985	0.0144	0.9499	0.0994	0.9365	0.1359	0.7787	0.0365	0.9471
CG14593	CG14593	1636300_at	0.0759	0.5934	0.4081	0.0330	0.3437	0.0802	0.0050	0.9956	-0.1848	0.3044	-0.1898	0.2342	0.0384	0.9516	0.1264	0.5232	0.0880	0.6731
larp	meteor	1636301_s_at	0.0897	0.8804	0.3674	0.3247	-0.1458	0.5963	-0.1937	0.6634	-0.2197	0.3072	-0.0259	0.9210	0.1597	0.9495	-0.0230	0.9884	-0.1827	0.8509
Su(dx)	Suppressor of del	1636302_s_at	-0.9919	0.0363	-1.4826	0.0057	-1.4519	0.0002	0.1386	0.8594	0.4595	0.1022	0.3209	0.2015	0.1472	0.8971	0.0659	0.9204	-0.0813	0.8889
dhd	thioredoxin	1636303_at	0.9696	0.6431	-2.6623	0.3085	-1.2549	0.2430	0.4032	0.6338	3.3786	0.0002	2.9754	0.0002	-1.0295	0.9296	-0.1864	0.9782	0.8431	0.8652
---	---	1636304_at	0.4764	0.0421	0.1019	0.5037	0.1644	0.4683	0.0332	0.9640	0.2184	0.2656	0.1851	0.2954	0.0428	0.9619	-0.0135	0.9752	-0.0562	0.8646
CG17327 /// CG9133 /// Dy	CG17327 /// CG9	1636305_a_at	-0.3499	0.1229	0.2875	0.1855	0.3938	0.0251	-0.1561	0.6325	-0.7661	0.0012	-0.6100	0.0019	-0.1516	0.8692	0.0901	0.8610	0.2417	0.5340
---	---	1636306_at	0.1489	0.3410	0.2134	0.3642	0.2057	0.2601	-0.0826	0.8578	-0.0852	0.6484	-0.0025	0.9904	-0.1471	0.7896	-0.1109	0.6694	0.0362	0.9110
---	---	1636307_at	0.0217	0.9161	0.1372	0.4204	0.1868	0.2833	-0.0820	0.8738	-0.1149	0.5525	-0.0329	0.8773	-0.1060	0.8486	-0.0357	0.9213	0.0703	0.8038
CG8605	CG8605	1636308_at	0.6463	0.0080	0.6196	0.0486	0.4512	0.0142	0.1834	0.5511	0.8889	0.0007	0.7055	0.0010	0.2867	0.6695	0.7733	0.0276	0.4866	0.1092
CG4089	CG4089	1636309_at	-2.8322	0.0005	-4.2681	0.0014	-3.9196	0.0000	0.3492	0.5854	1.1241	0.0050	0.7748	0.0160	0.1380	0.8718	-0.2279	0.5165	-0.3659	0.2960
Victoria	Turandot E	1636310_at	0.2546	0.1699	-0.1308	0.4212	-0.0043	0.9896	0.1732	0.6202	0.2712	0.1175	0.0980	0.5701	0.0006	0.9998	-0.1126	0.7490	-0.1133	0.7422
Gpdh	5' gene	1636311_at	0.7314	0.0098	-0.1194	0.7229	0.4552	0.0551	0.2008	0.6122	0.5046	0.0193	0.3039	0.0875	-0.4786	0.6483	-0.3562	0.3950	0.1224	0.8173
CG12038	CG12038	1636312_at	-0.2560	0.2056	0.0520	0.7321	-0.0739	0.7369	-0.1078	0.8063	-0.2639	0.1296	-0.1561	0.3282	0.1047	0.9174	0.1101	0.8061	0.0054	0.9929
CG4914	CG4914	1636313_at	0.0626	0.7292	-0.1804	0.4385	-0.1538	0.3221	0.1089	0.7941	0.1792	0.2968	0.0703	0.6941	-0.1216	0.8825	-0.2180	0.4994	-0.0964	0.8003
mtSSB	mitochondrial sing	1636314_at	0.1637	0.6039	-0.1422	0.7025	0.0714	0.7509	0.0673	0.8871	0.3127	0.0554	0.2454	0.0867	-0.1209	0.9387	-0.1442	0.8192	-0.0233	0.9761
CG1360	CG1360	1636315_at	0.3770	0.0717	0.2289	0.1985	0.2565	0.1814	0.0101	0.9883	0.0030	0.9899	-0.0071	0.9715	-0.0022	0.9985	-0.1803	0.4380	-0.1780	0.4597
CG1531	CG1531	1636316_a_at	-0.0822	0.9014	1.2015	0.1176	1.0602	0.0039	-0.1557	0.8824	-0.3180	0.3883	-0.1623	0.6634	-0.0296	0.9943	1.1468	0.2181	1.1764	0.2376
---	---	1636317_at	0.1466	0.5298	0.1494	0.3341	0.1771	0.3240	-0.0712	0.9024	-0.0184	0.9479	0.0528	0.8033	-0.0404	0.9568	0.1164	0.6093	0.1567	0.4771
---	---	1636318_at	0.0243	0.9040	0.0796	0.5814	0.2834	0.3585	0.1188	0.7992	-0.0017	0.9951	-0.1205	0.5081	-0.0041	0.9978	0.0839	0.8611	0.0879	0.8413
---	---	1636319_at	0.2011	0.2379	-0.9956	0.0172	-0.4566	0.2542	0.4536	0.6615	1.1349	0.0297	0.6813	0.1229	-0.2291	0.7120	-0.1994	0.4421	0.0298	0.9347
Arp11	Arp11	1636320_at	-0.6224	0.0049	-0.0160	0.8760	-0.0887	0.6534	0.0940	0.7929	-0.3510	0.0242	-0.4450	0.0052	0.1871	0.7464	0.3220	0.1749	0.1350	0.6093
CG13624	CG13624	1636321_s_at	0.1634	0.6109	0.7003	0.0502	0.4278	0.1217	-0.1331	0.7929	-0.2053	0.3282	-0.0722	0.7476	-0.0191	0.9939	0.1121	0.8877	0.1312	0.8529
CG11190	CG11190	1636322_at	-0.3910	0.2374	0.1281	0.6485	0.0382	0.8403	0.1475	0.6607	-0.0699	0.7068	-0.2174	0.1294	0.1655	0.9057	0.4941	0.3120	0.3286	0.5345
CG2249 /// DsimCG2249 ///	anon-fast-evolving	1636323_at	0.0499	0.8463	0.0620	0.8470	0.1967	0.4684	-0.2996	0.4979	-0.1379	0.5877	0.1616	0.4660	-0.4030	0.6955	-0.1580	0.7579	0.2449	0.5959
CG17623	CG17623	1636324_at	0.1464	0.4788	0.1168	0.4203	0.1685	0.5655	-0.1864	0.6929	-0.0688	0.7968	0.1176	0.5898	-0.0963	0.8815	0.0032	0.9953	0.0995	0.7240
Osi7	Osiris	1636325_at	0.2474	0.3198	0.5046	0.1092	0.5243	0.0120	-0.0385	0.9744	-0.2892	0.3258	-0.2506	0.3464	0.1039	0.9239	0.1230	0.7863	0.0190	0.9732
---	---	1636326_s_at	-0.1974	0.2508	-0.5107	0.0960	-0.5562	0.2173	-0.1557	0.9181	0.1606	0.7744	0.3163	0.4748	-0.3689	0.7152	-0.0465	0.9460	0.3224	0.4597
Tap42	Two A-associated	1636327_at	-0.4921	0.0420	-0.0976	0.6935	0.0557	0.7194	0.0299	0.9680	-0.2885	0.1374	-0.3183	0.0704	-0.0847	0.9507	0.1265	0.8105	0.2112	0.6306
CG12507	CG12507	1636328_at	0.1739	0.2757	-0.1658	0.3916	0.1123	0.4312	0.0323	0.9641	0.1700	0.3850	0.1377	0.4445	-0.1943	0.6749	-0.0648	0.7842	0.1295	0.5156
CG11333	CG11333	1636329_at	0.1363	0.3181	0.2469	0.1474	0.1626	0.3251	0.0194	0.9759	-0.0925	0.5919	-0.1119	0.4550	0.1946	0.6531	0.1716	0.3105	-0.0230	0.9234
CG17107	CG17107	1636330_at	0.2351	0.3193	0.2039	0.3466	0.0683	0.7393	-0.1377	0.7519	-0.1225	0.5469	0.0152	0.9494	0.0391	0.9816	-0.0029	0.9981	-0.0420	0.9360
Sox102F	Sox102F	1636331_at	1.0855	0.0466	1.7409	0.0350	2.0466	0.0015	-0.1835	0.8899	-0.4958	0.2669	-0.3123	0.4579	-0.4301	0.8339	0.0806	0.9531	0.5107	0.5744
Tm1	Tropomyosin	1636332_at	0.6881	0.0032	0.3326	0.1644	0.1904	0.2356	-0.0707	0.8791	0.3015	0.0610	0.3722	0.0167	0.1392	0.8298	0.0583	0.8812	-0.0810	0.8086
mthl2	Mth-like 2	1636333_a_at	-2.8050	0.0438	-4.4722	0.0100	-4.8373	0.0000	-0.4297	0.3660	0.4854	0.0748	0.9150	0.0032	-0.2429	0.9717	-1.3516	0.4342	-1.1087	0.5444
CG11913	CG11913	1636334_at	0.1904	0.4662	0.1035	0.6506	0.0982	0.5828	0.0751	0.8956	0.0850	0.6970	0.0098	0.9675	0.1929	0.8305	0.0083	0.9925	-0.1846	0.6491
CG12923	CG12923	1636335_at	-0.0651	0.7758	0.1810	0.2712	0.0823	0.6664	-0.0824	0.8860	-0.2071	0.2919	-0.1248	0.5065	0.0140	0.9923	0.0890	0.8478	0.0750	0.8682
CG14618	CG14618	1636336_at	-0.0728	0.6973	-0.1889	0.5318	-0.0473	0.8647	0.0325	0.9674	0.4663	0.0330	0.4338	0.0281	-0.1271	0.9076	0.4146	0.2707	0.5417	0.1897
pgant6	polypeptide GalN	1636337_s_at	1.4047	0.0044	1.5281	0.0162	1.6091	0.0000	0.2465	0.3357	0.5025	0.0053	0.2561	0.0568	0.0845	0.9588	0.4936	0.2599	0.4091	0.3807
CG15021	CG15021	1636338_at	-2.3390	0.0058	-3.5563	0.0014	-3.2895	0.0000	0.4642	0.5148	0.8272	0.0350	0.3630	0.2866	0.3296	0.8270	-0.3496	0.5995	-0.6793	0.2918
CG13334	CG13334	1636339_at	0.2722	0.2333	0.0335	0.7409	-0.0848	0.6240	0.0807	0.8942	0.1753	0.3970	0.0946	0.6489	0.1347	0.7733	0.0458	0.8750	-0.0889	0.6941
mtTFB2	mitochondrial tran	1636340_at	-0.0605	0.7042	-0.0129	0.9326	0.0063	0.9771	-0.1876	0.6282	0.1192	0.5626	0.3068	0.0740	-0.0994	0.8202	0.1333	0.4598	0.2328	0.2126
fzy	fizzy	1636341_at	-0.0977	0.8837	0.1144	0.8437	-0.7384	0.2485	-0.6344	0.5012	0.3178	0.5557	0.9522	0.0400	-0.0792	0.9875	0.2225	0.8960	0.3017	0.8336
---	---	1636342_at	0.2769	0.2529	-0.0358	0.7246	0.0596	0.7629	0.1416	0.7713	0.2374	0.2492	0.0958	0.6519	0.1248	0.8446	-0.0109	0.9828	-0.1357	0.6311
CG6283	CG6283	1636343_at	0.3211	0.1671	-0.0764	0.6855	0.2794	0.3037	0.3554	0.5073	0.1467	0.6464	-0.2087	0.4349	0.0837	0.9088	0.0688	0.8444	-0.0149	0.9701
chp	chaoptin	1636344_at	0.2843	0.2738	-0.2434	0.0944	0.1747	0.4915	0.1579	0.6514	0.3264	0.0576	0.1685	0.2685	-0.2649	0.7810	-0.2801	0.5090	-0.0152	0.9833
syt	Synaptotagmin	1636345_s_at	0.0235	0.9082	-0.1141	0.4402	-0.1600	0.3534	-0.1812	0.7511	-0.1182	0.6757	0.0629	0.8233	0.0374	0.9657	-0.0762	0.7951	-0.1136	0.6508
CG7839 /// DmirCG7839	anon-fast-evolving	1636346_at	0.5270	0.0868	0.0993	0.8940	0.1900	0.5504	0.1109	0.9255	0.9614	0.0151	0.8505	0.0159	-0.0205	0.9950	0.4312	0.5855	0.4516	0.5681
Hs6st	Heparan sulfate 6	1636347_at	0.0106	0.9804	0.3544	0.1482	0.3917	0.0573	0.0138	0.9893	0.0177	0.9600	0.0038	0.9898	-0.1526	0.8910	0.3185	0.4499	0.4711	0.2784
Act87E	Actin 87E	1636348_s_at	-1.4713	0.0540	-2.1313	0.0990	-3.6595	0.0008	-0.6361	0.0555	-0.7259	0.0033	-0.0898	0.6485	0.8453	0.8400	-1.3846	0.4203	-2.2298	0.2136
---	---	1636349_at	-0.4894	0.0758	0.0334	0.8266	-0.2275	0.3049	-0.0954	0.9029	-0.2577	0.3150	-0.1623	0.5076	-0.0500	0.9646	0.1187	0.7383	0.1687	0.6052
---	---	1636350_at	-0.3325	0.0462	0.2062	0.4194	0.0663	0.6795	-0.1282	0.7349	-0.3514	0.0430	-0.2232	0.1366	-0.0386	0.9653	0.1602	0.5184	0.1988	0.4136
CG17333 /// DyakCG17333	CG17333	1636351_at	0.6280	0.0238	0.9247	0.0513	1.4793	0.0003	0.0129	0.9883	-0.9532	0.0014	-0.9661	0.0008	-0.4477	0.6824	-0.4204	0.3383	0.0273	0.9682
CG15443	CG15443	1636352_at	-0.3256	0.0543	-0.0686	0.5185	-0.2786	0.1140	-0.0404	0.9515	0.0361	0.8834	0.0765	0.6910	0.1499	0.7707	0.2925	0.1684	0.1426	0.5360
CG8112	CG8112	1636353_s_at	1.9891	0.0644	-0.3659	0.6651	0.6151	0.0172	-0.1848	0.8589	0.2469	0.5380	0.4318	0.1962	-1.2481	0.6960	-1.9954	0.1382	-0.7472	0.6087
CG11107	CG11107	1636354_at	0.2126	0.5959	0.2434	0.2941	0.0871	0.6647	0.0423	0.9591	0.3793	0.0804	0.3370	0.0821	0.2275	0.8692	0.5000	0.3534	0.2726	0.6461
CG11260	CG11260	1636355_at	-0.0700	0.6522	-0.0792	0.4655	-0.0146	0.9473	0.0696	0.8589	0.0327	0.8596	-0.0369	0.8172	-0.0263	0.9816	0.0313	0.9410	0.0575	0.8699
CG3927	CG3927	1636356_at	0.0337	0.8343	-0.1312	0.3599	-0.1384	0.3537	-0.1059	0.8189	0.1487	0.4268	0.2546	0.1141	-0.0464	0.9409	-0.0745	0.7439	-0.0281	0.9168

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14834	CG14834	1636357_at	0.1078	0.5243	-0.1648	0.3280	-0.2995	0.2079	-0.0469	0.9507	0.1106	0.6381	0.1575	0.4231	-0.0664	0.9365	-0.1116	0.7066	-0.0452	0.9013
eIF-4B	Eukaryotic initiatio	1636358_a_at	0.3324	0.2728	1.3007	0.0236	0.5688	0.0423	-0.2308	0.7809	-0.6716	0.0557	-0.4408	0.1510	0.5012	0.6483	0.3631	0.4056	-0.1381	0.7947
CG31139	CG31139	1636359_at	-0.2029	0.3309	0.1041	0.4790	-0.0637	0.7492	-0.1603	0.7604	-0.1191	0.6377	0.0412	0.8776	0.1255	0.8837	0.2978	0.3486	0.1724	0.6231
---	---	1636360_at	0.2870	0.3020	0.0543	0.6838	-0.1844	0.2794	-0.1420	0.8553	0.1671	0.5848	0.3092	0.2198	-0.0056	0.9964	-0.0347	0.9460	-0.0291	0.9473
---	---	1636361_at	0.2738	0.1282	0.0374	0.9263	0.1923	0.2391	0.1370	0.8190	0.1570	0.5303	0.0200	0.9463	-0.1821	0.7743	-0.0928	0.7871	0.0893	0.7848
CG32169	CG32169	1636362_at	-0.1601	0.4856	-0.1746	0.5728	-0.0381	0.8959	0.1580	0.8358	0.3808	0.1836	0.2229	0.4066	-0.0454	0.9588	0.1708	0.4954	0.2162	0.3859
Mapmodulin	Mapmodulin	1636363_s_at	0.8160	0.0774	0.7921	0.0582	-0.2957	0.3624	-0.8364	0.0193	0.5874	0.0081	1.4238	0.0001	0.3267	0.8331	0.7600	0.2236	0.4334	0.5234
CG17376	CG17376	1636364_a_at	-0.1321	0.4774	-0.0356	0.8553	-0.1012	0.4941	0.0446	0.9319	0.0116	0.9604	-0.0330	0.8549	0.0350	0.9705	0.0661	0.8376	0.0311	0.9249
sick	sickie	1636365_at	-0.1966	0.3431	-0.1200	0.5373	-0.0286	0.9134	0.0228	0.9838	0.0503	0.8836	0.0276	0.9298	-0.0732	0.9246	0.1214	0.6749	0.1947	0.4697
---	---	1636366_at	0.1555	0.3611	0.2511	0.3626	0.3630	0.0305	0.1477	0.5993	-0.0845	0.5827	-0.2323	0.0694	0.1002	0.8494	-0.0143	0.9684	-0.1145	0.6280
CG11018	CG11018	1636367_at	0.1244	0.4053	-0.0348	0.7677	-0.1466	0.3973	-0.1063	0.8233	0.1464	0.4458	0.2527	0.1246	-0.0277	0.9816	0.0203	0.9622	0.0480	0.8926
CG14269	CG14269	1636368_at	0.1905	0.2589	0.0136	0.9034	0.2628	0.3626	0.0173	0.9838	0.0396	0.8802	0.0223	0.9261	-0.0874	0.8870	-0.1391	0.5644	-0.0517	0.8650
---	---	1636369_at	-0.0509	0.8061	0.0736	0.5701	0.2925	0.0939	-0.0174	0.9777	-0.0692	0.7051	-0.0518	0.7660	-0.1094	0.8191	0.0407	0.8852	0.1502	0.4572
---	---	1636370_at	0.0886	0.6692	0.0118	0.9101	0.0171	0.9453	0.0551	0.9116	-0.0490	0.7998	-0.1040	0.4907	0.1185	0.8472	-0.0207	0.9614	-0.1392	0.6140
CG2941	CG2941	1636371_at	-0.3999	0.2673	0.1284	0.5911	-0.2260	0.2248	-0.2970	0.4908	-0.2866	0.2104	0.0104	0.9711	-0.0372	0.9869	0.2059	0.7289	0.2431	0.6621
BBS4	BBS4	1636372_at	-0.0340	0.8231	0.0665	0.6845	0.0905	0.6480	-0.0570	0.9233	-0.0342	0.8872	0.0227	0.9173	-0.1481	0.8024	-0.1087	0.6958	0.0394	0.9082
CG5155	CG5155	1636373_at	0.3054	0.0917	0.1908	0.3003	-0.1260	0.4273	-0.0506	0.9345	0.0624	0.7752	0.1130	0.5203	0.2436	0.6955	-0.0508	0.8967	-0.2944	0.2769
CAH1	Carbonic anhydra	1636374_at	-1.4516	0.0094	-1.7752	0.0249	-1.6743	0.0000	-0.0044	0.9955	-0.1588	0.3098	-0.1544	0.2688	-0.0316	0.9884	-0.4061	0.3749	-0.3745	0.4340
CG30060	CG30060	1636375_at	0.0584	0.8060	0.0501	0.6930	0.2651	0.2437	0.1094	0.8663	-0.0827	0.7639	-0.1921	0.3632	-0.0132	0.9923	0.0328	0.9496	0.0460	0.9169
CG14947	CG14947	1636376_at	0.0280	0.8918	0.1501	0.4458	-0.0028	0.9905	-0.0988	0.8073	-0.1764	0.2757	-0.0776	0.6384	0.0970	0.8903	0.1764	0.5179	0.0794	0.8048
pHCl	pHCl	1636377_at	-0.9916	0.0345	-0.1439	0.3787	-1.2701	0.0003	-0.7652	0.1420	-0.6319	0.0502	0.1332	0.6873	0.0225	0.9894	-0.2976	0.3972	-0.3201	0.3803
Syn	synapsin	1636378_a_at	0.5163	0.3372	-0.1193	0.4615	0.3212	0.4150	-0.2805	0.8462	-0.6414	0.2266	-0.3609	0.4754	-0.3386	0.8379	-0.7472	0.2522	-0.4087	0.5709
DnaJ-60	DnaJ-like-60	1636379_a_at	-0.3298	0.2626	-0.4037	0.0668	-0.5788	0.0169	-0.0773	0.9182	-0.7822	0.0055	-0.7049	0.0053	0.0772	0.9657	-0.7609	0.1199	-0.8381	0.1179
dnt	Doughnut	1636380_at	-0.0560	0.9485	0.4209	0.2554	-0.2900	0.1876	0.0434	0.9375	-0.0211	0.9252	-0.0646	0.7038	0.7709	0.7046	0.5530	0.5289	-0.2180	0.8404
CG33156	CG33156	1636381_a_at	0.1180	0.5463	0.1153	0.3972	0.0755	0.7480	-0.0063	0.9948	0.1880	0.3545	0.1943	0.2804	-0.0558	0.9465	0.0002	1.0000	0.0560	0.8667
CG17574	CG17574	1636382_at	-1.1798	0.0135	-1.4226	0.0021	-1.4653	0.0002	-0.0997	0.8281	0.0413	0.8585	0.1410	0.3911	0.0857	0.9632	-0.2515	0.6483	-0.3372	0.5270
CG13856	CG13856	1636383_at	-1.0910	0.0257	-2.3269	0.0213	-2.2509	0.0050	0.2653	0.9099	0.7375	0.3292	0.4722	0.5122	0.0512	0.9862	-0.7030	0.2916	-0.7542	0.2855
CG32703	CG32703	1636384_at	0.0502	0.8314	0.0768	0.5992	-0.1040	0.5835	0.0630	0.9251	0.0344	0.9009	-0.0285	0.9067	0.0741	0.9449	-0.0320	0.9508	-0.1061	0.7836
CG11317	CG11317	1636385_at	-0.4241	0.1441	-0.1729	0.3458	-0.2347	0.2979	-0.0228	0.9777	-0.2189	0.2995	-0.1962	0.3009	-0.0721	0.9511	-0.0066	0.9935	0.0655	0.8914
CG8179	CG8179	1636386_s_at	-0.0062	0.9845	0.0326	0.7641	-0.0558	0.8140	-0.0324	0.9603	0.0139	0.9567	0.0463	0.8130	0.1615	0.8461	-0.0176	0.9768	-0.1791	0.6282
CG10300	CG10300	1636387_at	-0.0219	0.9684	-0.0349	0.7620	0.0452	0.8610	-0.1784	0.8768	-0.1829	0.6865	-0.0045	0.9928	-0.2799	0.8086	-0.4046	0.3941	-0.1248	0.8379
CG14220	CG14220	1636388_at	-0.4505	0.0644	0.3032	0.2974	0.1896	0.3188	-0.0551	0.9376	-0.5153	0.0207	-0.4602	0.0208	0.0139	0.9916	0.2805	0.3625	0.2666	0.4037
---	---	1636389_at	-0.0284	0.8958	0.0000	1.0000	-0.0197	0.9393	-0.0868	0.8915	-0.0899	0.7166	-0.0032	0.9906	0.0392	0.9701	-0.0096	0.9854	-0.0487	0.8946
CG14568	CG14568	1636390_at	0.0279	0.8999	0.1839	0.3409	0.0937	0.5906	0.0537	0.9254	-0.0508	0.8112	-0.1045	0.5345	0.1332	0.7953	0.0501	0.8701	-0.0832	0.7390
CG15609	CG15609	1636391_at	-0.1053	0.5286	-0.2118	0.4718	-0.4202	0.0547	-0.0869	0.8449	0.1257	0.4619	0.2126	0.1454	0.1086	0.9137	0.0377	0.9470	-0.0708	0.8818
CG15785	CG15785	1636392_at	0.1239	0.4613	0.2225	0.3493	-0.0149	0.9348	-0.0404	0.9559	-0.0651	0.7869	-0.0248	0.9187	0.2426	0.5754	0.1021	0.6280	-0.1405	0.4865
CG15005	CG15005	1636393_at	0.2470	0.2164	-0.0126	0.9148	0.0345	0.8614	0.1546	0.7293	0.1794	0.3804	0.0248	0.9193	0.0686	0.9635	-0.1564	0.7448	-0.2250	0.6094
---	---	1636394_a_at	-0.7663	0.0043	-0.8497	0.0713	-0.8679	0.0124	0.1558	0.8671	0.2474	0.4720	0.0916	0.8043	0.2529	0.8357	-0.0373	0.9646	-0.2902	0.5905
CG9468	CG9468	1636395_at	-0.2924	0.7661	-0.1822	0.6545	0.1294	0.6638	0.6188	0.6450	-0.2353	0.7634	-0.8541	0.1409	0.3123	0.8661	-0.1835	0.8571	-0.4958	0.5225
dod	dodo	1636396_at	0.1026	0.6818	-0.0375	0.8693	-0.2619	0.2702	-0.1629	0.8028	0.2470	0.3536	0.4099	0.0809	0.4088	0.6423	0.1667	0.6743	-0.2420	0.5163
CG13615	CG13615	1636397_at	0.0038	0.9844	-0.0736	0.6695	-0.0297	0.8710	0.0675	0.9071	-0.0285	0.9127	-0.0959	0.6106	-0.0210	0.9862	-0.0214	0.9633	-0.0004	0.9992
CG10086	CG10086	1636398_at	0.6889	0.5525	0.1004	0.5804	0.0960	0.7984	-0.2403	0.9269	-0.3782	0.6721	-0.1379	0.8818	-0.0726	0.9852	-0.7561	0.3909	-0.6835	0.4597
---	---	1636399_at	0.0706	0.8043	-0.1351	0.4185	-0.0469	0.7797	0.3304	0.4174	0.3714	0.1005	0.0410	0.8785	0.0594	0.9581	-0.0718	0.8738	-0.1312	0.7134
---	---	1636400_at	0.0702	0.6778	-0.1545	0.2322	0.0224	0.9179	0.0421	0.9466	0.3180	0.0756	0.2760	0.0832	-0.1730	0.7196	-0.0916	0.6819	0.0813	0.7199
Nmt	N-myristoyl transfe	1636401_a_at	0.5723	0.0157	0.4036	0.2892	0.1429	0.3624	-0.2096	0.5137	0.4001	0.0261	0.6097	0.0025	0.0548	0.9717	0.3759	0.3236	0.3211	0.4192
CG10081	CG10081	1636402_at	-0.1385	0.4680	0.1159	0.3782	0.2409	0.1023	-0.1557	0.6413	-0.2574	0.1135	-0.1017	0.5204	-0.1214	0.8439	0.0521	0.8867	0.1734	0.5092
CG5276	CG5276	1636403_at	-0.4745	0.0591	0.2655	0.4858	0.8843	0.0011	0.1104	0.8192	-0.3339	0.0794	-0.4444	0.0167	-0.3233	0.7550	0.5780	0.1750	0.9013	0.0772
Mst35Bb	protamine	1636404_at	0.2803	0.0737	0.1102	0.4397	-0.0651	0.7257	-0.0437	0.9422	0.0067	0.9800	0.0503	0.7932	0.0923	0.8222	-0.0164	0.9521	-0.1087	0.5477
CG4329	CG4329	1636405_a_at	-0.0259	0.9241	0.1415	0.4257	0.1390	0.4219	0.1455	0.7134	-0.0532	0.8112	-0.1987	0.2180	-0.1102	0.8940	0.0049	0.9941	0.1150	0.7484
CG10321 /// DereCG10321	CG10321	1636406_at	-0.0365	0.9144	0.2861	0.3168	0.0993	0.5603	0.1901	0.7039	0.1445	0.5602	-0.0456	0.8650	0.3169	0.7358	0.4131	0.2874	0.0962	0.8525
gol	goliath	1636407_at	0.1901	0.2645	0.1561	0.4640	0.2539	0.1471	-0.0012	0.9988	-0.2390	0.2767	-0.2378	0.2245	-0.1178	0.8222	-0.2496	0.2378	-0.1318	0.5691
---	---	1636408_at	0.2127	0.2451	0.0935	0.4615	0.0877	0.6760	0.0687	0.8776	0.1989	0.1914	0.1303	0.3545	-0.0246	0.9860	0.0692	0.8781	0.0937	0.8088
CG11034	CG11034	1636409_at	0.1753	0.4658	0.0406	0.6879	0.2021	0.5021	0.1724	0.6954	0.1023	0.6556	-0.0701	0.7514	0.0736	0.9589	0.1396	0.7824	0.0660	0.9067
CG3505	CG3505	1636410_at	-0.8145	0.0135	-2.2637	0.0100	-1.5978	0.0015	0.4933	0.4529	0.9386	0.0167	0.4454	0.1615	-0.2107	0.8814	-0.4750	0.3698	-0.2643	0.6484
---	---	1636411_at	0.2431	0.4074	0.0969	0.7040	0.1834	0.2803	0.1509	0.8158	0.1896	0.4759	0.0388	0.8992	-0.2053	0.8203	-0.0336	0.9563	0.1717	0.6771
MICAL	MICAL	1636412_at	0.4970	0.5740	-0.2669	0.4088	-1.0315	0.0242	-1.0026	0.0144	0.4900	0.0259	1.4927	0.0002	-0.1627	0.9681	-0.4256	0.7341	-0.2629	0.8495
---	---	1636413_at	0.1217	0.6111	0.3635	0.1886	0.4425	0.0370	-0.0145	0.9874	-0.1476	0.5284	-0.1331	0.5345	0.0839	0.9420	0.0848	0.8642	0.0008	0.9992
CG30004	CG30004	1636414_at	-1.1023	0.0677	-0.1213	0.8728	-1.3829	0.0003	-0.3052	0.6941	-0.5411	0.1299	-0.2359	0.4925	0.8993	0.6999	0.3185	0.7977	-0.5808	0.5786
CG14225	CG14225	1636415_at	-0.9290	0.0275	-0.2316	0.6286	-0.1177	0.6751	-0.0105	0.9909	-0.7265	0.0041	-0.7160	0.0026	-0.1967	0.9357	-0.0528	0.9668	0.1439	0.8923
---	---	1636416_at	0.4401	0.0443	0.1896	0.3448	0.5386	0.0490	0.1825	0.5261	0.1703	0.2606	-0.0122	0.9492	0.1387	0.9340	-0.1902	0.7707	-0.3289	0.5629
---	---	1636417_at	0.2299	0.3197	0.0491	0.6886	0.1867	0.2800	0.0966	0.7918	0.0363	0.8509	-0.0602	0.7010	-0.0290	0.9774	-0.1019	0.7090	-0.0729	0.8044
Hcf	Host cell factor	1636418_s_at	0.1182	0.6097	-0.0152	0.9444	0.1284	0.3790	-0.0566	0.9218	-0.0857	0.6644	-0.0291	0.8884	0.0086	0.9958	-0.0095	0.9911	-0.0181	0.9760
CG1647	CG1647	1636419_at	-0.1536	0.6870	0.0441	0.8645	-0.1425	0.4214	-0.0068	0.9941	0.4012	0.0403	0.4080	0.0240	0.1607	0.9064	0.5976	0.2086	0.4368	0.3830
CG16826 /// DyakCG16826	CG16826	1636420_at	0.6854	0.4937	-0.1116	0.5023	-0.1543	0.6606	-0.2058	0.8155	0.1480	0.7116	0.3537	0.2558	-0.1727	0.9734	-0.6898	0.6122	-0.5171	0.7199
CG14260	CG14260	1636421_at	0.2378	0.1857	0.2530	0.4135	0.4804	0.0203	0.0548	0.9589	0.0924	0.7851	0.0376	0.9112	0.0567	0.9589	-0.0342	0.9462	-0.0909	0.8162
---	---	1636422_at	0.0700	0.7425	0.0147	0.8893	0.1328	0.3438	0.0527	0.9165	0.2188	0.1602	0.1661	0.2362	0.0204	0.9862	0.0968	0.7573	0.0764	0.8154
CG8012	CG8012	1636423_at	0.3032	0.4331	0.3917	0.4955	0.2539	0.2919	0.0528	0.9744	-0.7057	0.0765	-0.7585	0.0385	0.1904	0.9101	-0.6158	0.2938	-0.8062	0.2039
---	---	1636424_s_at	0.0872	0.5766	0.0286	0.8101	0.0482	0.8287	-0.0553	0.9138	0.0198	0.9301	0.0751	0.6519	-0.0885	0.8692	0.1114	0.6312	0.1999	0.3649
Or33a	Olfactory receptor	1636425_at	0.2011	0.4219	0.0206	0.8471	0.1733	0.3666	0.0763	0.9112	0.0756	0.7718	-0.0007	0.9981	-0.0006	0.9998	-0.0305	0.9353	-0.0299	0.9269

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG6421	CG6421	1636426_at	-0.1569	0.8025	1.8880	0.0050	0.9747	0.0457	-1.5224	0.1279	-2.6689	0.0017	-1.1464	0.0394	-0.6163	0.7196	-0.5279	0.4640	0.0884	0.9291
Mt2	DNA(5-cytosine) r	1636427_a_at	0.1819	0.4690	0.0208	0.9678	0.1562	0.6667	0.0860	0.9029	0.3219	0.1572	0.2359	0.2503	-0.1549	0.9309	0.1813	0.8083	0.3362	0.5868
---	---	1636428_at	-0.1065	0.6541	0.2628	0.0975	0.0802	0.6924	-0.0102	0.9880	-0.1437	0.3767	-0.1335	0.3617	0.2053	0.7485	0.2016	0.4678	-0.0037	0.9940
CG5928	CG5928	1636429_at	0.3565	0.1524	-0.0140	0.9022	0.2876	0.2468	0.0000	0.9999	0.0661	0.7741	0.0661	0.7507	-0.0956	0.9342	-0.1153	0.8117	-0.0197	0.9727
---	---	1636430_at	-0.0798	0.6808	0.0996	0.5348	-0.2229	0.3191	-0.1158	0.8679	0.0335	0.9226	0.1493	0.5347	0.0127	0.9901	-0.0417	0.9027	-0.0545	0.8507
CG8768	CG8768	1636431_at	-0.5179	0.2461	-0.4251	0.0460	-0.6967	0.0053	-0.2557	0.5259	-0.3758	0.0773	-0.1201	0.5661	-0.1713	0.9092	-0.4758	0.3650	-0.3045	0.5958
---	---	1636432_at	0.1522	0.5960	-0.0138	0.9337	0.0268	0.8884	-0.0318	0.9669	-0.0198	0.9469	0.0120	0.9621	-0.0343	0.9816	-0.1030	0.8135	-0.0687	0.8812
CG2260	CG2260	1636433_at	-0.0491	0.9071	-0.0180	0.9524	0.3607	0.0577	0.3530	0.2701	0.3946	0.0430	0.0416	0.8493	-0.0088	0.9958	0.4809	0.1962	0.4897	0.2199
---	---	1636434_at	-0.2211	0.2057	-0.2222	0.5197	-0.5057	0.0680	-0.0470	0.9514	0.3078	0.1452	0.3548	0.0649	0.1847	0.8454	0.0548	0.9289	-0.1299	0.7770
CG11068	CG11068	1636435_at	0.0832	0.6598	-0.1524	0.3350	0.0800	0.6246	0.0623	0.9205	0.2017	0.2930	0.1394	0.4351	-0.1869	0.7230	-0.1154	0.6408	0.0715	0.7945
eff	Suppressor of GM	1636436_at	-0.0516	0.7484	0.7489	0.0878	0.4800	0.0137	-0.1398	0.7815	-0.4774	0.0298	-0.3376	0.0728	0.3090	0.7052	0.5185	0.1294	0.2096	0.5612
CG31638	CG31638	1636437_at	-0.4823	0.0815	0.1774	0.3800	0.0608	0.7499	0.0354	0.9653	-0.5177	0.0250	-0.5531	0.0119	0.0373	0.9618	0.1524	0.4937	0.1151	0.6222
---	---	1636438_s_at	0.0625	0.7100	0.1719	0.2139	0.0223	0.9048	-0.1442	0.6122	-0.1383	0.3410	0.0059	0.9736	0.0456	0.9500	-0.0236	0.9462	-0.0692	0.7894
CG34354	CG12425	1636439_at	-0.0154	0.9405	-0.1494	0.5009	0.0595	0.7859	0.0405	0.9649	0.1323	0.6268	0.0918	0.7254	-0.1002	0.9246	0.0062	0.9938	0.1064	0.8056
Sip1	SRY interacting pr	1636440_at	-1.3868	0.0018	-0.3169	0.5263	-1.2529	0.0008	-0.1653	0.8156	-0.4868	0.0817	-0.3215	0.1957	0.6795	0.5659	0.5508	0.2880	-0.1287	0.8524
---	---	1636441_at	0.2156	0.4417	0.1061	0.4870	-0.0700	0.6942	-0.0612	0.9406	0.0467	0.8828	0.1079	0.6588	0.1759	0.8400	0.0434	0.9404	-0.1325	0.7494
CG1999	CG1999	1636442_at	-0.0165	0.9395	0.1241	0.5067	0.0850	0.7155	0.0757	0.9413	0.0429	0.9173	-0.0327	0.9276	0.0390	0.9679	0.1567	0.5597	0.1178	0.6749
shep	alan shepard	1636443_s_at	0.1310	0.5880	0.1904	0.1267	0.1371	0.5620	-0.1445	0.7971	-0.0313	0.9220	0.1132	0.6260	0.0710	0.9421	-0.0036	0.9959	-0.0746	0.8482
CG13871	CG13871	1636444_at	-0.0153	0.9418	-0.0293	0.7951	-0.0525	0.8022	0.0834	0.8462	0.0262	0.9044	-0.0572	0.7363	-0.0302	0.9806	-0.0440	0.9167	-0.0137	0.9745
---	---	1636445_at	0.1551	0.4993	-0.0607	0.6885	0.1009	0.5828	0.1354	0.7722	0.1384	0.5056	0.0030	0.9904	-0.0197	0.9898	-0.1284	0.7448	-0.1087	0.7849
CSN6	Drosophila COP9	1636446_at	0.2127	0.1550	0.6544	0.0843	0.9448	0.0008	0.0752	0.8872	-0.3656	0.0466	-0.4408	0.0137	-0.0914	0.9056	0.2373	0.3815	0.3287	0.2525
CG14024	CG14024	1636447_at	0.1620	0.3223	0.2600	0.1509	0.5266	0.0370	-0.0280	0.9755	-0.2443	0.2717	-0.2164	0.2791	0.1828	0.7691	0.2133	0.4183	0.0304	0.9346
---	---	1636448_at	0.0956	0.7141	-0.1984	0.4063	0.1446	0.5931	0.0414	0.9672	0.0682	0.8422	0.0269	0.9348	-0.1417	0.8655	-0.0415	0.9390	0.1002	0.8061
CG32984	CG32984	1636449_at	0.1920	0.2843	-0.1615	0.4258	0.0795	0.6049	0.2562	0.4130	0.2629	0.1284	0.0067	0.9754	-0.0339	0.9734	-0.1388	0.6024	-0.1049	0.7089
---	---	1636450_at	-0.7684	0.0582	-1.4003	0.0095	-1.1861	0.0020	-0.0442	0.9722	0.4261	0.1747	0.4702	0.0952	-0.2931	0.8379	-0.0555	0.9531	0.2376	0.7224
Atf-2	Activating transcri	1636451_a_at	0.0652	0.7255	0.1258	0.6440	0.0204	0.9162	-0.0146	0.9857	-0.0580	0.8000	-0.0434	0.8405	0.0902	0.9239	-0.0673	0.8824	-0.1575	0.6423
CG14423	CG14423	1636452_at	-0.0204	0.9218	-0.0331	0.8721	-0.0783	0.7768	-0.0513	0.9218	-0.0250	0.9089	0.0262	0.8893	0.0184	0.9862	-0.0615	0.8539	-0.0799	0.7810
CG4988	CG4988	1636453_at	0.1156	0.4661	0.1315	0.5387	0.2775	0.1951	0.1284	0.7000	-0.0088	0.9703	-0.1373	0.3261	0.0567	0.9741	-0.0896	0.8849	-0.1464	0.7698
CG7956 /// DyakCG7956	CG7956	1636454_at	-0.1083	0.7122	0.0729	0.6682	-0.2808	0.1442	0.0012	0.9988	0.2474	0.2699	0.2463	0.2177	0.1488	0.8236	0.2154	0.4391	0.0665	0.8542
---	---	1636455_at	-0.2334	0.3040	0.1622	0.1896	-0.0170	0.9448	-0.1071	0.8539	-0.2635	0.2077	-0.1564	0.4269	0.1343	0.8348	0.0247	0.9541	-0.1095	0.7163
CG13053	CG13053	1636456_at	0.1524	0.3024	0.2482	0.1721	0.3782	0.0320	0.0840	0.8291	0.0924	0.5670	0.0083	0.9649	-0.1321	0.8257	0.1514	0.5618	0.2835	0.2707
---	---	1636457_at	-0.0256	0.9014	-0.1419	0.4204	-0.3012	0.1869	-0.0747	0.8967	-0.0473	0.8471	0.0273	0.9045	0.0546	0.9748	-0.0479	0.9451	-0.1025	0.8542
CG4702	CG4702	1636458_at	0.0303	0.8934	0.1477	0.4801	-0.0247	0.8979	0.0135	0.9909	-0.0577	0.8660	-0.0712	0.8067	0.1394	0.8472	-0.0079	0.9910	-0.1472	0.6498
CG18284 /// CG31872 /// D	CG31872 /// CG1(	1636459_s_at	0.2197	0.2024	0.0771	0.4966	0.0711	0.7332	-0.1527	0.6471	-0.1666	0.3072	-0.0139	0.9460	0.0157	0.9900	-0.1867	0.5074	-0.2024	0.4738
Jon65Ai	Jonah 65A	1636460_at	0.2234	0.6293	-0.0421	0.7815	0.1152	0.5672	0.3129	0.2939	0.1853	0.2944	-0.1277	0.4384	0.0986	0.9588	-0.1776	0.7897	-0.2761	0.6302
CG3630	CG3630	1636461_at	-2.3416	0.0107	-2.4709	0.0804	-2.9634	0.0000	-0.2444	0.8180	-0.1652	0.7363	0.0792	0.8716	0.2791	0.9411	-0.0749	0.9690	-0.3540	0.8044
---	---	1636462_at	0.0738	0.7807	-0.0239	0.9153	-0.0862	0.6724	0.0951	0.8636	0.1385	0.5048	0.0434	0.8490	-0.0023	0.9990	-0.0143	0.9788	-0.0120	0.9809
Ucp4B	Ucp4B	1636463_at	-0.1587	0.2802	0.0464	0.7042	-0.0719	0.7065	-0.0301	0.9639	-0.0706	0.7321	-0.0405	0.8417	0.0055	0.9952	0.0936	0.7273	0.0881	0.7425
Rrp6	Rrp6	1636464_at	0.4417	0.2750	-0.0373	0.9313	0.1718	0.3939	0.1896	0.7855	0.6819	0.0256	0.4923	0.0590	-0.0738	0.9735	0.1135	0.8857	0.1873	0.7705
CG9215	CG9215	1636465_at	-0.1588	0.6470	-0.0784	0.7721	0.2041	0.2154	0.1878	0.6099	0.0034	0.9899	-0.1844	0.2682	-0.0093	0.9964	0.1303	0.8380	0.1396	0.8123
Cdlc2	dynein	1636466_s_at	0.1672	0.2618	0.0667	0.5751	0.0282	0.9186	0.0504	0.9426	-0.0900	0.6944	-0.1404	0.4564	-0.0390	0.9641	-0.0509	0.8792	-0.0119	0.9743
---	---	1636467_at	0.0791	0.7151	0.2205	0.3428	0.4383	0.0517	-0.0393	0.9688	-0.2419	0.3653	-0.2026	0.4065	0.0649	0.9449	0.1208	0.7084	0.0559	0.8870
CG5033	CG5033	1636468_a_at	0.7129	0.0776	0.1045	0.8752	0.5159	0.0076	0.4367	0.5008	0.8500	0.0218	0.4133	0.1785	0.0745	0.9717	0.3315	0.5446	0.2570	0.6486
CG14363	CG14363	1636469_at	0.0221	0.8993	0.0990	0.6356	0.1612	0.3331	-0.0306	0.9538	-0.0681	0.6792	-0.0375	0.8185	-0.1351	0.8013	-0.0300	0.9333	0.1050	0.6659
CG34026	CG34026	1636470_at	0.2785	0.3276	0.1036	0.4088	0.2804	0.1328	-0.0024	0.9961	-0.0372	0.8531	-0.0348	0.8468	-0.0788	0.9460	-0.1565	0.6937	-0.0777	0.8689
mod(mdg4)	Modifier67.2	1636471_at	0.2004	0.4295	0.1647	0.4689	0.1624	0.4195	-0.2842	0.4511	-0.1730	0.4129	0.1113	0.5869	0.0206	0.9913	0.0303	0.9627	0.0096	0.9874
unc-4	Paired HomeoDor	1636472_at	-0.0782	0.7241	0.0054	0.9669	-0.0108	0.9608	0.0777	0.8895	-0.0326	0.8985	-0.1103	0.5480	-0.0272	0.9754	-0.0180	0.9599	0.0091	0.9794
CG12861	CG12861	1636473_at	0.1616	0.4675	0.0543	0.5704	0.0429	0.8205	-0.0735	0.8942	0.0256	0.9214	0.0990	0.5908	0.0405	0.9677	0.0001	1.0000	-0.0405	0.9109
CG12602	CG12602	1636474_at	0.6202	0.7234	0.1864	0.4018	-1.2470	0.0048	-1.2415	0.4408	-0.6528	0.4773	0.5887	0.4825	0.0039	0.9998	-0.9610	0.5886	-0.9650	0.5892
otp	Orthopedia	1636475_at	0.1476	0.3706	-0.0023	0.9864	-0.0958	0.7031	-0.0397	0.9493	0.1587	0.3726	0.1985	0.2021	0.0177	0.9862	0.0310	0.9358	0.0133	0.9716
CG11109	CG11109	1636476_a_at	0.6897	0.0057	0.7310	0.0952	0.7644	0.0007	0.2349	0.3743	0.4121	0.0136	0.1771	0.1838	0.3212	0.7070	0.6266	0.0900	0.3055	0.4016
CG31934	CG31934	1636477_at	0.0907	0.5780	0.0059	0.9672	-0.0248	0.8832	0.0111	0.9860	-0.0006	0.9979	-0.0117	0.9503	0.0428	0.9646	-0.0565	0.8792	-0.0993	0.7409
CG9596	CG9596	1636478_at	0.4709	0.0342	0.4645	0.1794	1.1195	0.0006	0.3567	0.2466	-0.0107	0.9699	-0.3674	0.0317	-0.1155	0.9126	0.0810	0.8786	0.1965	0.6225
---	---	1636479_at	-0.1569	0.5074	0.1997	0.1098	-0.0366	0.8846	-0.0593	0.9397	0.0417	0.8924	0.1010	0.6693	0.1369	0.8521	0.1717	0.5941	0.0348	0.9345
CG14532	CG14532	1636480_at	0.2623	0.0984	0.1794	0.1388	0.2471	0.1899	0.1865	0.5859	0.0937	0.6299	-0.0928	0.5955	0.1029	0.8331	0.1260	0.5543	0.0231	0.9353
---	---	1636481_s_at	0.1937	0.3230	0.1239	0.4312	0.1254	0.4086	0.0579	0.9196	0.0644	0.7595	0.0065	0.9766	0.2804	0.6557	0.1174	0.6789	-0.1630	0.5448
CG14005	CG14005	1636482_at	-0.2348	0.3410	-0.0092	0.9332	-0.0407	0.8360	0.1752	0.7857	-0.1619	0.5727	-0.3371	0.1549	0.2279	0.7248	-0.0212	0.9634	-0.2491	0.3847
CG6511	CG6511	1636483_at	-0.8357	0.0974	0.4898	0.3458	0.0822	0.7417	-0.2057	0.8605	-0.9847	0.0269	-0.7790	0.0441	0.1343	0.9400	0.5470	0.3022	0.4127	0.4653
RNaseX25	acid RNase	1636484_at	0.0169	0.9570	-0.0044	0.9864	0.1642	0.2729	-0.1764	0.6712	-0.5476	0.0131	-0.3712	0.0413	-0.3143	0.6955	-0.5129	0.1300	-0.1986	0.5811
CTCF	CTCF	1636485_at	0.2764	0.6351	-0.0074	0.9845	0.1173	0.7524	0.3529	0.5376	0.8926	0.0095	0.5397	0.0486	0.1575	0.9475	0.5715	0.4311	0.4140	0.5969
Gdi /// Taf11	TBP-associated fa	1636486_at	0.2716	0.1026	-0.0943	0.5362	-0.1014	0.5449	0.0436	0.9482	0.4300	0.0299	0.3863	0.0294	0.1360	0.7779	0.1373	0.5259	0.0014	0.9975
CG7079	CG7079	1636487_at	0.2677	0.7299	-0.0080	0.9539	-0.1282	0.6327	0.0101	0.9955	-0.0123	0.9816	-0.0224	0.9597	0.1186	0.9710	-0.3945	0.6614	-0.5131	0.5547
CG4468 /// DereCG4468 /// CG4468		1636488_at	0.0647	0.7949	0.1280	0.3914	0.1048	0.7277	-0.0178	0.9860	-0.1911	0.4339	-0.1733	0.4333	0.1209	0.9142	0.0631	0.9170	-0.0578	0.9157
CG5987	CG5987	1636489_at	-0.5173	0.1249	-0.3386	0.2156	-0.1217	0.5104	-0.0385	0.9603	-0.1467	0.5027	-0.1081	0.6056	-0.0428	0.9829	0.0477	0.9494	0.0905	0.8863
PGRP-SB1	PGRP-SB1	1636490_at	2.9298	0.0033	0.9030	0.6416	1.6091	0.0714	0.5220	0.6662	2.4937	0.0019	1.9718	0.0031	-0.3541	0.9657	0.3052	0.9312	0.6593	0.8088
CG3955	CG3955	1636491_at	-0.0681	0.8247	0.1550	0.3312	0.1331	0.5991	0.0325	0.9712	-0.1155	0.6540	-0.1480	0.5004	0.0376	0.9816	0.0794	0.8851	0.0418	0.9372
CR30029	CR30029	1636492_at	-1.2014	0.0115	-0.2932	0.0920	-0.5249	0.0166	-0.0412	0.9774	-0.8677	0.0226	-0.8265	0.0172	0.1237	0.8692	0.0401	0.9341	-0.0837	0.8247
dei	delilah	1636493_at	-2.1165	0.0028	-2.9423	0.0085	-2.5438	0.0001	0.7082	0.2789	0.9368	0.0232	0.2286	0.5452	0.2931	0.8465	0.2341	0.7547	-0.0590	0.9469
---	---	1636494_at	0.1191	0.4013	0.1913	0.4299	0.2084	0.3280	-0.0827	0.8172	0.0524	0.7545	0.1350	0.2879	0.0310	0.9717	0.0790	0.7699	0.0480	0.8719

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1636495_at	0.0832	0.7476	0.1167	0.4504	0.0418	0.8420	0.0788	0.9254	0.1789	0.5009	0.1001	0.7075	0.0709	0.9142	0.1542	0.5158	0.0833	0.7524
Nup153	Nup153	1636496_at	0.5071	0.4635	0.1127	0.6370	-0.2774	0.1664	0.1061	0.9068	1.1275	0.0031	1.0214	0.0028	0.4216	0.8479	0.7857	0.3806	0.3640	0.7266
---	---	1636497_at	0.0234	0.9086	0.0870	0.3983	0.1251	0.5709	-0.2321	0.5726	-0.2745	0.1890	-0.0424	0.8625	-0.0448	0.9503	0.0556	0.8481	0.1004	0.6652
CG6168	CG6168	1636498_at	-2.6952	0.0010	-0.2479	0.4185	-1.8757	0.0004	-1.5709	0.0458	-2.8045	0.0004	-1.2336	0.0082	0.2224	0.8292	-0.2892	0.5183	-0.5116	0.2536
Atg6	Autophagy-specifi	1636499_at	-0.4105	0.0797	-0.0443	0.8777	0.1950	0.1963	-0.0199	0.9774	-0.5820	0.0055	-0.5621	0.0038	-0.2820	0.7644	-0.1516	0.7453	0.1305	0.7810
CG32425	CG32425	1636500_at	0.0469	0.8377	0.3526	0.1476	0.4047	0.0158	-0.0247	0.9759	-0.0724	0.7636	-0.0477	0.8355	-0.0016	0.9993	0.1466	0.5338	0.1481	0.5355
CG32656	CG32656	1636501_at	-0.0923	0.9700	0.0644	0.8821	-0.1001	0.5071	-0.0723	0.9883	-1.2360	0.2832	-1.1637	0.2585	-0.0499	0.9943	-1.1741	0.4843	-1.1242	0.5100
CG7718 /// DyakCG7718	CG7718	1636502_at	-0.0011	0.9964	0.1505	0.5327	0.2090	0.3171	0.1186	0.8589	0.2016	0.4148	0.0830	0.7502	0.1364	0.8513	0.5027	0.1033	0.3663	0.2398
---	---	1636503_at	-0.0650	0.6914	-0.0365	0.8582	-0.0496	0.7986	0.2781	0.5502	0.1270	0.6377	-0.1511	0.5172	0.2285	0.7628	0.2153	0.5085	-0.0132	0.9798
CG13008	CG13008	1636504_a_at	0.1132	0.4827	0.3406	0.1578	0.1759	0.5279	-0.0499	0.9558	0.0148	0.9687	0.0647	0.8132	0.0690	0.9416	0.2704	0.3397	0.2014	0.5042
CG15083	CG15083	1636505_at	-0.6606	0.0915	0.5395	0.0628	0.1555	0.5305	0.1389	0.6869	-0.1581	0.3302	-0.2970	0.0440	0.6028	0.6749	1.1755	0.0637	0.5726	0.3348
CG40116	CG40116	1636506_at	-0.0902	0.5940	0.0046	0.9745	0.0219	0.9030	0.0136	0.9860	-0.0144	0.9562	-0.0280	0.8956	0.0370	0.9651	0.0245	0.9467	-0.0124	0.9716
vanin-like	vanin-like	1636507_at	0.2400	0.5576	0.8707	0.0149	1.8658	0.0001	0.1945	0.8436	0.2174	0.5825	0.0230	0.9597	-0.7258	0.4825	0.7413	0.1402	1.4671	0.0368
Nca	neurocalcin	1636508_at	-0.8901	0.0025	0.0605	0.7964	-0.2292	0.3686	-0.1532	0.8028	-0.8257	0.0057	-0.6725	0.0086	0.0779	0.9277	0.0724	0.8501	-0.0056	0.9907
---	---	1636509_at	0.1775	0.4188	0.0883	0.7320	0.1768	0.3719	0.0469	0.9567	-0.0304	0.9252	-0.0773	0.7612	-0.2200	0.7215	-0.2179	0.3909	0.0022	0.9965
Lsd-1	Lipid storage drop	1636510_a_at	2.5534	0.0007	1.4673	0.0431	2.7506	0.0001	1.5030	0.0359	1.0981	0.0134	-0.4048	0.2563	0.1991	0.9235	0.1759	0.8500	-0.0232	0.9841
---	---	1636511_at	0.0989	0.6303	0.0909	0.5028	0.4427	0.0183	0.0674	0.9011	0.0079	0.9772	-0.0595	0.7570	-0.1475	0.8023	0.1401	0.5907	0.2876	0.2578
---	---	1636512_s_at	0.1240	0.4704	-0.0708	0.5225	0.0372	0.8439	0.1314	0.7711	0.2654	0.1620	0.1340	0.4579	0.0242	0.9779	0.0307	0.9265	0.0065	0.9849
kis	kismet	1636513_a_at	0.5977	0.0411	0.5354	0.0945	0.3426	0.1394	-0.1146	0.8640	0.2529	0.2915	0.3675	0.0844	0.2201	0.8424	0.3442	0.4485	0.1242	0.8274
Tak1	TGF-beta activate	1636514_at	-0.5199	0.0154	0.1057	0.6071	-0.0659	0.7050	-0.1512	0.6622	-0.4009	0.0238	-0.2497	0.0922	-0.0350	0.9776	0.1632	0.6093	0.1981	0.5270
CG14354	CG14354	1636515_at	0.1244	0.4735	-0.0156	0.8969	0.2193	0.2237	0.0325	0.9639	0.1326	0.5098	0.1000	0.6010	0.0412	0.9679	0.0563	0.8858	0.0151	0.9714
CG32189	CG32189	1636516_at	0.2779	0.3524	0.2082	0.0935	0.5807	0.0236	-0.0075	0.9956	-0.2066	0.5207	-0.1991	0.4924	-0.2895	0.7036	-0.2581	0.4115	0.0314	0.9434
gprs	gprs	1636517_at	0.3581	0.0625	0.0955	0.4827	0.1749	0.3333	0.1836	0.5947	0.1745	0.3257	-0.0091	0.9674	-0.0320	0.9812	-0.1022	0.7727	-0.0702	0.8514
---	---	1636518_s_at	0.0170	0.9520	-0.0180	0.8677	0.2385	0.2179	0.0100	0.9907	0.0273	0.9167	0.0173	0.9404	-0.1847	0.7628	-0.0289	0.9445	0.1558	0.5637
CG13716	CG13716	1636519_at	-0.3040	0.2801	-0.0756	0.5554	-0.7399	0.0052	-0.6306	0.1341	-0.1156	0.6926	0.5150	0.0311	-0.0135	0.9914	-0.1312	0.6686	-0.1177	0.7060
CG16940	CG16940	1636520_s_at	-0.0774	0.7676	-0.0673	0.8777	-0.3706	0.0696	-0.0744	0.9108	0.4710	0.0321	0.5455	0.0108	0.3490	0.7726	0.4848	0.3350	0.1358	0.8354
Thd1	Thd1	1636521_at	-0.2517	0.2227	0.0864	0.8547	-0.4247	0.0325	-0.3228	0.3389	-0.1861	0.3478	0.1367	0.4583	0.2360	0.8461	0.1193	0.8665	-0.1167	0.8600
CG16850	CG16850	1636522_at	0.0307	0.8844	0.4537	0.0272	0.0695	0.7920	-0.0191	0.9857	-0.2912	0.2267	-0.2721	0.2070	0.2414	0.6955	0.2133	0.4077	-0.0281	0.9382
CG7276	CG7276	1636523_at	0.1577	0.5440	0.1743	0.4006	0.0891	0.6409	-0.0995	0.8174	-0.0780	0.6869	0.0214	0.9175	0.0738	0.9447	-0.0830	0.8500	-0.1568	0.6510
---	---	1636524_at	0.0660	0.6703	0.0212	0.9030	-0.0747	0.7160	-0.0014	0.9985	0.0568	0.7657	0.0582	0.7327	0.1174	0.8012	0.1115	0.5881	-0.0059	0.9849
CG13541	CG13541	1636525_at	-0.0811	0.7570	0.1128	0.4341	0.0671	0.8347	-0.0435	0.9744	-0.0247	0.9600	0.0188	0.9637	0.0429	0.9643	0.1521	0.5800	0.1092	0.7075
CG9451	CG9451	1636526_at	2.2794	0.0021	1.3052	0.0534	2.4781	0.0010	0.8553	0.4200	0.7931	0.1750	-0.0622	0.9314	-0.5618	0.7220	-0.3085	0.6772	0.2533	0.7426
por	porcupine	1636527_at	0.1542	0.6674	-0.2925	0.3553	-0.4218	0.1232	0.1813	0.7409	0.5896	0.0229	0.4083	0.0622	0.3963	0.7606	0.4452	0.4094	0.0490	0.9491
Gp93	Glycoprotein 93	1636528_at	1.1700	0.0071	1.1785	0.0172	1.6383	0.0000	0.3943	0.4102	0.7906	0.0096	0.3963	0.0949	-0.1277	0.9142	0.8469	0.0622	0.9745	0.0559
CG33483	CG33483	1636529_at	0.2705	0.0997	0.1975	0.2239	0.1327	0.3725	0.0139	0.9840	0.0927	0.6021	0.0788	0.6357	0.1092	0.8191	0.1199	0.5576	0.0106	0.9717
betaTub56D	beta tubulin	1636530_s_at	-0.6227	0.0251	-0.1315	0.2353	-0.0677	0.6698	0.1650	0.6496	0.2118	0.2266	0.0468	0.8132	0.0932	0.9056	0.6808	0.0391	0.5876	0.0741
---	---	1636531_s_at	0.1972	0.2663	-0.1215	0.5726	-0.0666	0.8150	0.3024	0.4860	0.0365	0.9096	-0.2659	0.1984	0.0843	0.9421	-0.1966	0.6080	-0.2809	0.4465
CG13958	CG13958	1636532_at	0.2548	0.1301	-0.0142	0.9010	-0.1556	0.5177	0.0255	0.9749	0.1000	0.6542	0.0746	0.7242	0.1353	0.8270	-0.0372	0.9275	-0.1725	0.5247
Lhr	Lethal hybrid resc	1636533_at	-0.5244	0.0557	-0.1416	0.7385	-0.2371	0.3480	0.2129	0.7427	0.5305	0.0663	0.3177	0.2132	0.2774	0.8141	0.8233	0.0999	0.5460	0.2784
CRMP	dihydropyrimidine	1636534_a_at	0.1453	0.5152	0.3886	0.1535	0.6997	0.0064	0.0459	0.9434	-0.3760	0.0450	-0.4219	0.0178	-0.1971	0.7956	-0.0373	0.9425	0.1598	0.6428
CG15912	CG15912	1636535_at	-0.0166	0.9594	0.0941	0.7681	0.4621	0.0316	0.1561	0.7253	-0.0439	0.8681	-0.1999	0.2669	-0.2845	0.7464	-0.1679	0.6876	0.1167	0.7954
Pglym87	Pglym87	1636536_at	0.0165	0.9561	-0.0147	0.8893	0.0120	0.9499	-0.0420	0.9517	-0.1264	0.5366	-0.0844	0.6716	-0.1662	0.8192	-0.1575	0.6229	0.0087	0.9852
CG30274	CG30274	1636537_at	0.7097	0.0285	0.8486	0.0118	0.9292	0.0012	-0.1207	0.8707	-0.2459	0.3519	-0.1252	0.6381	-0.0767	0.9291	-0.0753	0.8407	0.0014	0.9979
---	---	1636538_at	-0.0717	0.6217	-0.0535	0.8466	-0.0461	0.8469	-0.1408	0.8085	-0.0708	0.8031	0.0700	0.7868	-0.0394	0.9778	-0.0400	0.9409	-0.0007	0.9992
Dak1	Dak1	1636539_at	0.1415	0.4699	0.8690	0.0149	0.6122	0.0059	0.0818	0.8196	-0.1366	0.3394	-0.2183	0.0835	0.2346	0.7464	0.5470	0.0869	0.3124	0.3205
CG8165	CG8165	1636540_at	-0.1052	0.6268	-0.4917	0.2084	-1.2143	0.0020	-0.2698	0.5357	0.2913	0.1984	0.5612	0.0136	0.1818	0.8714	-0.0302	0.9670	-0.2120	0.6607
CG33052	CG33052	1636541_at	0.4274	0.0283	0.2345	0.4801	0.0985	0.6854	-0.0396	0.9463	0.5552	0.0058	0.5948	0.0025	0.1064	0.9238	0.3720	0.3000	0.2656	0.4863
CG15277	CG15277	1636542_at	0.0618	0.7505	-0.0774	0.5909	0.1212	0.5457	0.0937	0.8714	-0.1008	0.6586	-0.1945	0.2871	0.0505	0.9648	0.0758	0.8592	0.0253	0.9535
CG1969	CG1969	1636543_a_at	1.7456	0.0019	1.4011	0.0075	1.6342	0.0001	-0.0472	0.9477	-0.4704	0.0280	-0.4232	0.0275	0.0069	0.9963	-0.2336	0.4721	-0.2405	0.4677
CG7011	CG7011	1636544_at	0.7856	0.0046	1.1568	0.0260	1.2139	0.0003	0.0477	0.9345	0.2968	0.0835	0.2491	0.1022	-0.0749	0.9535	0.7237	0.0721	0.7986	0.0732
CheB38c	CheB38c	1636545_at	0.0767	0.6832	0.0283	0.8243	0.5165	0.0105	0.0827	0.8877	-0.2025	0.3105	-0.2852	0.1059	-0.2231	0.6272	-0.1492	0.4379	0.0739	0.7391
CG1640	CG1640	1636546_s_at	0.7897	0.0070	0.4388	0.1354	1.0563	0.0006	0.1664	0.7010	-0.3046	0.1241	-0.4711	0.0162	-0.4004	0.3517	-0.4973	0.0519	-0.0969	0.7073
---	---	1636547_at	-0.1141	0.4860	-0.9170	0.0198	-1.0534	0.0001	-0.0946	0.8578	0.4627	0.0225	0.5573	0.0060	0.1090	0.8442	-0.2182	0.3259	-0.3272	0.1752
blw	mitochondrial ATF	1636548_at	-0.1523	0.3466	-0.2297	0.3313	-0.3730	0.0183	-0.1474	0.5943	-0.1198	0.4060	0.0276	0.8662	-0.0521	0.9535	-0.2411	0.3526	-0.1890	0.4912
CG3996	CG3996	1636549_at	0.1172	0.4635	0.1022	0.6438	-0.1569	0.3049	0.0916	0.8738	0.1402	0.5099	0.0486	0.8326	0.1803	0.8062	0.1948	0.5425	0.0145	0.9764
Obp22a	Odorant-binding p	1636550_at	-0.0527	0.7756	0.0206	0.8399	0.2080	0.1680	0.1507	0.7149	0.0495	0.8383	-0.1013	0.5912	-0.0442	0.9588	-0.0278	0.9420	0.0165	0.9622
CG14683 /// DyakCG14683	CG14683	1636551_at	-0.4880	0.0341	0.4767	0.1694	0.7785	0.0212	0.0500	0.9322	-0.5548	0.0073	-0.6048	0.0029	-0.1270	0.9301	0.3902	0.3999	0.5172	0.2842
dys	Dysfusion	1636552_at	0.0546	0.7474	0.1321	0.4475	-0.0574	0.7558	-0.0768	0.8791	0.0147	0.9542	0.0915	0.5959	0.0747	0.9499	0.1287	0.7625	0.0540	0.9121
CG15766	CG15766	1636553_at	-0.1601	0.4547	-0.2697	0.1422	-0.1848	0.4926	0.2170	0.6533	0.2115	0.3739	-0.0055	0.9851	0.1197	0.8122	0.0339	0.9162	-0.0858	0.7154
---	---	1636554_at	-0.0528	0.7536	-0.0470	0.7724	0.2938	0.0941	0.2766	0.4630	0.1239	0.5741	-0.1527	0.4242	0.0332	0.9657	0.1264	0.5620	0.0933	0.6853
CG17784	CG17784	1636555_at	0.1555	0.3854	0.0567	0.6716	-0.1740	0.2798	-0.1387	0.7293	0.1008	0.6094	0.2395	0.1334	0.1314	0.8012	-0.0243	0.9451	-0.1557	0.4875
CG8141	CG8141	1636556_at	0.0361	0.9103	0.1548	0.3917	0.1488	0.3391	-0.1028	0.8470	-0.0484	0.8484	0.0544	0.8046	-0.0188	0.9913	0.1406	0.7340	0.1594	0.6848
epsin-like	epsin-2	1636557_a_at	0.5913	0.0912	0.7971	0.0238	0.6917	0.0014	-0.0634	0.8968	0.1807	0.2651	0.2441	0.0907	-0.0104	0.9952	0.4746	0.2217	0.4849	0.2454
abd-A	abdominal-A	1636558_a_at	-0.3903	0.5879	1.0387	0.0818	0.9287	0.0084	-0.2547	0.6820	-0.5916	0.0463	-0.3370	0.1916	0.1489	0.9775	1.1424	0.3458	0.9936	0.4367
Prosbeta5	20S proteasome	1636559_s_at	-0.1499	0.3920	0.6815	0.0128	0.8172	0.0012	0.0472	0.9339	-0.5266	0.0075	-0.5738	0.0030	-0.0825	0.8903	0.3129	0.1512	0.3954	0.1142
---	---	1636560_s_at	0.1812	0.3768	0.0510	0.7383	0.1713	0.3207	0.0906	0.8701	0.0817	0.7166	-0.0088	0.9710	-0.1234	0.8541	-0.1228	0.6926	0.0006	0.9992
CG9967	CG9967	1636561_at	-0.1189	0.7479	-0.2663	0.4457	-0.0026	0.9935	0.4859	0.1223	0.8382	0.0017	0.3523	0.0438	0.0683	0.9798	0.5494	0.3558	0.4810	0.4411
---	---	1636562_at	-0.0764	0.7025	-0.1931	0.4457	-0.2778	0.1476	-0.1257	0.8508	0.0460	0.8892	0.1717	0.4561	-0.1302	0.8909	-0.0838	0.8684	0.0464	0.9252
CG32399	CG32399	1636563_at	0.1198	0.5709	-0.0213	0.8444	-0.2348	0.1572	0.0845	0.8628	0.2716	0.1168	0.1871	0.2294	0.1689	0.7810	0.0715	0.8386	-0.0973	0.7483

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14757	CG14757	1636564_at	-0.9541	0.2625	-1.4619	0.0361	-2.0421	0.0002	-0.2923	0.6413	-1.3249	0.0019	-1.0326	0.0033	0.0706	0.9898	-2.0258	0.0943	-2.0964	0.1104
---	---	1636565_at	0.2804	0.1772	-0.1383	0.4102	0.0089	0.9715	0.1642	0.7115	0.1024	0.6514	-0.0618	0.7833	-0.0968	0.8960	-0.2837	0.2907	-0.1868	0.5172
CG32368	CG32368	1636566_at	-0.1105	0.9583	-1.3054	0.6031	-0.6326	0.6448	0.8401	0.6570	1.5317	0.0926	0.6916	0.4161	0.1680	0.9914	0.3213	0.9538	0.1533	0.9777
CG31199	CG31199	1636567_at	0.7403	0.5208	0.1390	0.4522	-0.4314	0.0495	-0.1866	0.8094	-0.2207	0.4947	-0.0341	0.9278	0.4743	0.8960	-0.7094	0.6270	-1.1838	0.3921
---	---	1636568_at	0.1742	0.4561	0.0995	0.4071	0.2400	0.2677	0.1006	0.8732	0.0363	0.9045	-0.0644	0.7932	0.0878	0.9400	-0.0692	0.8979	-0.1570	0.6933
CG11883	CG11883	1636569_a_at	-0.2500	0.2595	0.0613	0.7304	0.2102	0.2277	-0.3150	0.4166	-0.6624	0.0084	-0.3474	0.0729	-0.3328	0.7423	-0.1621	0.7466	0.1706	0.7253
mex1	midgut expression	1636570_at	0.0672	0.9811	-0.0735	0.6028	-0.0470	0.8486	0.1453	0.9777	-0.6301	0.6771	-0.7755	0.5525	-0.0106	0.9994	-1.0565	0.6443	-1.0459	0.6445
Rtnl2	tropomyosin-like	1636571_at	0.3350	0.0460	-0.1044	0.3196	0.0800	0.6318	0.3285	0.3016	0.3226	0.0856	-0.0059	0.9799	0.0609	0.9380	-0.2665	0.2566	-0.3273	0.2033
Duox	Dual oxidase	1636572_at	0.5075	0.2269	-0.1160	0.4925	-0.7002	0.0242	-0.8025	0.2326	1.5355	0.0031	2.3381	0.0003	-0.2523	0.8049	0.9060	0.0564	1.1583	0.0421
---	---	1636573_at	-0.0097	0.9704	-0.0098	0.9340	-0.3525	0.0574	-0.0873	0.9098	0.1502	0.5701	0.2375	0.2832	0.0151	0.9898	0.0455	0.9069	0.0305	0.9320
CG33056	CG33056	1636574_s_at	0.4832	0.0614	1.2244	0.0236	0.2762	0.1629	-0.0145	0.9872	-0.6610	0.0086	-0.6465	0.0057	0.8244	0.2892	-0.0425	0.9494	-0.8669	0.0710
---	---	1636575_s_at	-0.2137	0.2804	0.6424	0.0505	0.5102	0.0158	-0.2274	0.5917	-0.8464	0.0028	-0.6189	0.0068	-0.1344	0.8365	-0.0987	0.7584	0.0358	0.9231
norpA	phospholipase C	1636576_s_at	-0.7482	0.0144	0.7789	0.1149	0.1115	0.4568	-0.3429	0.2810	-0.2219	0.2368	0.1209	0.5026	0.5258	0.6660	1.2729	0.0335	0.7471	0.1551
---	---	1636577_at	0.0949	0.6374	-0.0709	0.6021	-0.0101	0.9744	0.2150	0.6380	0.1208	0.6279	-0.0942	0.6887	0.0757	0.9400	-0.0604	0.8963	-0.1360	0.6923
---	---	1636578_at	0.1595	0.3035	0.2021	0.2958	0.0983	0.6179	-0.0625	0.9285	-0.0703	0.7774	-0.0078	0.9761	-0.0415	0.9514	-0.0122	0.9709	0.0293	0.9161
mdy	midway	1636579_s_at	2.3578	0.0011	0.1964	0.7146	1.3176	0.0002	0.0583	0.9612	0.2653	0.4235	0.2071	0.5018	-0.8995	0.3738	-1.8627	0.0200	-0.9632	0.1184
CG17387	CG17387	1636580_at	0.0023	0.9904	0.1193	0.5427	0.0580	0.7377	-0.0494	0.9540	0.0024	0.9945	0.0518	0.8493	0.0581	0.9449	0.1485	0.5818	0.0904	0.7577
---	---	1636581_s_at	0.2269	0.2292	0.2497	0.0767	0.3704	0.0437	0.0505	0.9319	0.1795	0.3124	0.1290	0.4356	0.0358	0.9717	0.0640	0.8548	0.0281	0.9371
CG34011	CG34011	1636582_at	0.1040	0.4830	-0.0200	0.8437	0.1293	0.6220	0.0184	0.9777	-0.0549	0.7905	-0.0733	0.6782	-0.0217	0.9831	0.0090	0.9842	0.0307	0.9252
CG5932	CG5932	1636583_at	0.3420	0.2529	0.1520	0.5896	0.0670	0.8340	-0.0909	0.9447	-0.1629	0.7061	-0.0720	0.8704	0.1347	0.9340	-0.3543	0.5093	-0.4890	0.3601
---	---	1636584_at	0.0483	0.8478	0.1182	0.3492	-0.0193	0.9136	-0.0789	0.8897	-0.0907	0.6758	-0.0118	0.9605	0.0666	0.9174	-0.0512	0.8721	-0.1177	0.6233
hig	hikaru genki	1636585_a_at	-0.1535	0.3467	-0.1665	0.4004	-0.1279	0.4978	-0.1490	0.6990	-0.0298	0.9027	0.1192	0.4801	-0.1525	0.8202	-0.0506	0.9054	0.1020	0.7499
---	---	1636586_at	-0.0137	0.9633	-0.0039	0.9739	-0.1761	0.5255	0.1307	0.8189	0.1930	0.3989	0.0623	0.8039	-0.0427	0.9779	0.0244	0.9682	0.0670	0.8949
CG7288	CG7288	1636587_at	0.5374	0.0098	0.4867	0.2048	0.3877	0.0425	-0.0657	0.8735	0.3087	0.0379	0.3744	0.0106	0.1446	0.8859	0.2110	0.6019	0.0664	0.8985
---	---	1636588_at	0.2635	0.1184	0.1113	0.4007	-0.0793	0.6275	0.1686	0.7252	0.2457	0.2562	0.0771	0.7398	0.2396	0.6732	0.0659	0.8268	-0.1737	0.4635
CG31958 /// CG31960	CG31960 /// CG3	1636589_s_at	-0.0625	0.7799	0.0227	0.8389	-0.0194	0.9398	0.0540	0.9319	-0.0425	0.8637	-0.0965	0.6114	0.0629	0.9340	-0.0235	0.9519	-0.0864	0.7577
CG12423	CG12423	1636590_s_at	0.0224	0.9040	0.0177	0.9189	0.1059	0.6117	0.0176	0.9803	0.1225	0.5060	0.1049	0.5381	-0.0677	0.9328	0.0654	0.8531	0.1331	0.6311
skpB	skpB	1636591_at	0.0387	0.7974	-0.8060	0.0230	0.2563	0.5383	0.2406	0.8280	1.3912	0.0064	1.1506	0.0090	-0.7818	0.3362	0.4274	0.3066	1.2092	0.0390
---	---	1636592_at	0.0314	0.9409	0.1131	0.4626	0.2584	0.1838	0.1470	0.8738	0.0879	0.8309	-0.0592	0.8776	-0.1963	0.7506	0.0294	0.9457	0.2256	0.4033
---	---	1636593_s_at	0.0314	0.8518	0.0088	0.9683	-0.0458	0.8431	-0.0192	0.9814	-0.0807	0.7257	-0.0614	0.7792	0.0039	0.9964	-0.0985	0.7000	-0.1023	0.6816
CG15088	CG15088	1636594_at	1.3145	0.3721	0.2112	0.3232	0.4218	0.0474	0.1513	0.8265	-0.2526	0.3510	-0.4039	0.0897	0.2429	0.9734	-1.1844	0.5251	-1.4274	0.4402
CG13138	CG13138	1636595_a_at	0.1844	0.3542	0.0377	0.8224	0.6613	0.0871	0.1417	0.8074	0.5628	0.0232	0.4211	0.0474	-0.3507	0.7936	0.3948	0.4954	0.7455	0.2075
CG10075	CG10075	1636596_at	-0.2349	0.4320	-0.5014	0.1296	-0.4270	0.0526	-0.1069	0.8546	-0.0892	0.7153	0.0176	0.9473	-0.1743	0.8744	-0.3776	0.3749	-0.2033	0.6668
Osi13	Osiris	1636597_at	0.0371	0.9099	0.0163	0.8777	-0.0022	0.9944	0.1088	0.8908	0.0441	0.9045	-0.0648	0.8309	0.1225	0.8235	0.0198	0.9587	-0.1028	0.6842
Unc-76	Unc-76	1636598_s_at	-0.3994	0.0339	-0.5787	0.0432	-0.7063	0.0021	-0.2483	0.4404	-0.1746	0.3271	0.0737	0.6893	-0.1397	0.8706	-0.4501	0.1744	-0.3104	0.3764
CG1227	CG1227	1636599_at	0.1763	0.3413	0.0393	0.8415	-0.0867	0.6463	0.0254	0.9669	0.2670	0.0960	0.2416	0.0916	0.0095	0.9939	0.1465	0.6344	0.1370	0.6574
Zeelin1	Zeelin1	1636600_a_at	-1.0023	0.0218	-1.1475	0.0287	-2.2274	0.0032	-0.0678	0.9165	-0.2212	0.2738	-0.1534	0.4150	0.9256	0.6832	-0.4523	0.6600	-1.3779	0.1651
---	---	1636601_at	0.3121	0.1668	-0.0475	0.7803	-0.0900	0.6605	0.0439	0.9471	0.1837	0.3331	0.1398	0.4249	-0.0185	0.9913	-0.1607	0.6819	-0.1421	0.7216
CG11253	CG11253	1636602_at	0.3105	0.3181	0.1510	0.1739	0.4498	0.0334	0.2171	0.6673	0.1003	0.7208	-0.1168	0.6359	-0.1559	0.8134	-0.1882	0.5042	-0.0323	0.9330
CG9297	CG9297	1636603_a_at	-2.6004	0.0087	-2.0395	0.0324	-2.5406	0.0001	-0.2643	0.3553	-1.2363	0.0002	-0.9719	0.0003	0.2592	0.9467	-0.4906	0.7158	-0.7498	0.5481
CG12490	CG12490	1636604_at	-0.0318	0.8519	0.0936	0.4209	0.2563	0.1342	-0.0176	0.9777	-0.0520	0.7824	-0.0344	0.8483	-0.0757	0.9174	0.0870	0.7802	0.1626	0.5433
CG2816	CG2816	1636605_at	-0.2051	0.6838	-0.2193	0.2470	-0.7156	0.0347	-0.4314	0.3909	-0.1260	0.6998	0.3053	0.2291	-0.0401	0.9762	-0.2157	0.5199	-0.1756	0.6150
sll	slalom	1636606_at	0.7743	0.0050	0.3881	0.1176	0.3658	0.0765	-0.3186	0.4333	0.4376	0.0544	0.7561	0.0032	-0.0974	0.8995	0.0751	0.8454	0.1726	0.5669
Rab2	Rab-protein 2	1636607_at	0.2714	0.5113	0.4598	0.2350	-0.1048	0.6423	-0.1874	0.8483	0.0288	0.9566	0.2163	0.5361	0.2975	0.6955	0.1148	0.7603	-0.1827	0.5858
CG32843	CG32843	1636608_at	-0.2215	0.4582	0.0137	0.9061	-0.1091	0.5101	-0.2669	0.6652	-0.6026	0.0462	-0.3357	0.2013	-0.1685	0.7726	-0.4140	0.1067	-0.2455	0.3430
---	---	1636609_at	0.1064	0.5339	0.1505	0.3233	0.3777	0.1065	-0.0513	0.9515	0.0207	0.9525	0.0721	0.7809	-0.0377	0.9679	0.0106	0.9821	0.0483	0.8880
CG17359	CG17359	1636610_at	-0.1416	0.6682	0.1672	0.3716	-0.2205	0.3726	-0.0268	0.9834	0.0412	0.9204	0.0680	0.8407	0.3641	0.7215	0.3750	0.3760	0.0110	0.9874
Try29F	Trypsin 29F	1636611_at	-0.6205	0.7110	-0.1399	0.3769	0.0517	0.7907	0.2225	0.9635	-0.7979	0.5656	-1.0204	0.3935	-0.0419	0.9914	-0.5706	0.5338	-0.5287	0.5729
pk	prickle-spiny legs	1636612_a_at	-2.4520	0.0055	-3.6503	0.0242	-3.9389	0.0002	0.4923	0.7595	1.0851	0.1148	0.5927	0.3497	0.6647	0.8049	0.0631	0.9738	-0.6017	0.6186
CG31554	CG31554	1636613_at	0.1325	0.4875	0.2442	0.3798	0.1711	0.3226	0.1840	0.5900	-0.0449	0.8407	-0.2289	0.1377	0.0943	0.9063	0.1219	0.7068	0.0276	0.9449
vav	vav	1636614_a_at	0.2043	0.3200	0.4693	0.0445	0.6053	0.0379	-0.0420	0.9500	-0.1386	0.4759	-0.0966	0.6072	-0.3436	0.7498	0.0959	0.8852	0.4394	0.3538
elav	embryonic lethal,	1636615_at	0.3096	0.6606	-0.6362	0.0162	-1.9506	0.0132	-0.1813	0.8942	1.4221	0.0078	1.6034	0.0027	0.7929	0.7707	0.2038	0.9085	-0.5891	0.6388
Or63a	Odorant receptor	1636616_at	0.1805	0.3500	0.0376	0.7525	0.1664	0.4556	0.2097	0.6637	0.1169	0.6529	-0.0929	0.7027	0.0017	0.9990	0.0215	0.9506	0.0198	0.9479
---	---	1636617_a_at	0.1598	0.5228	0.4053	0.2258	-0.0511	0.8260	-0.5074	0.1659	-0.5276	0.0257	-0.0202	0.9419	-0.0686	0.9238	-0.1278	0.6154	-0.0592	0.8461
CG8664	CG8664	1636618_at	0.0956	0.7578	0.5355	0.1139	0.2908	0.1329	-0.0421	0.9451	-0.2030	0.2450	-0.1609	0.3098	0.1800	0.8400	0.1228	0.7896	-0.0572	0.9099
---	---	1636619_at	0.0248	0.8836	0.0140	0.8993	0.0448	0.8526	-0.1180	0.6999	-0.1100	0.4519	0.0080	0.9642	-0.1425	0.8650	-0.0624	0.9023	0.0801	0.8533
l(2)08717	lethal (2) 08717	1636620_s_at	0.3166	0.5179	0.0551	0.9403	0.6937	0.0064	0.2491	0.8174	0.1668	0.7383	-0.0824	0.8681	-0.5942	0.6955	-0.0736	0.9430	0.5206	0.4272
---	---	1636621_at	-0.0762	0.8226	-0.2224	0.5485	-0.3190	0.1215	0.1797	0.8677	0.5924	0.1190	0.4127	0.2271	0.2821	0.7070	0.3495	0.2505	0.0673	0.8743
Peritrophin-15a	Dm-peritrophin-15	1636622_at	0.3865	0.3593	-0.1922	0.4124	0.2927	0.1843	0.1896	0.5735	0.1289	0.4739	-0.0607	0.7441	-0.2987	0.8049	-0.5728	0.2415	-0.2741	0.6152
Ance-5	Ance-5	1636623_at	-1.5084	0.0088	-1.7377	0.0189	-1.5299	0.0027	-0.0968	0.9247	-0.6817	0.0334	-0.5850	0.0394	-0.4847	0.7751	-0.9210	0.1948	-0.4363	0.5767
---	---	1636624_at	0.2196	0.2894	0.0354	0.7042	0.2103	0.3136	0.0777	0.8975	0.1475	0.4768	0.0698	0.7445	0.0213	0.9831	0.1079	0.6830	0.0866	0.7513
mtc	montecristo	1636625_at	0.4178	0.4672	-1.1998	0.3138	-1.5696	0.0571	-0.6210	0.1125	1.6970	0.0003	2.3179	0.0000	-0.2635	0.9611	-0.1873	0.9401	0.0762	0.9749
---	---	1636626_s_at	-0.0440	0.8547	-0.1368	0.3941	-0.1798	0.2894	-0.0494	0.9223	0.0843	0.6179	0.1337	0.3392	0.0037	0.9980	-0.0144	0.9808	-0.0181	0.9709
CG18554	CG18554	1636627_at	0.2066	0.3224	0.2477	0.3147	0.3690	0.0418	0.1260	0.8738	-0.0141	0.9741	-0.1401	0.6182	0.0844	0.8956	0.0225	0.9531	-0.0618	0.8394
CG11866	CG11866	1636628_at	0.1570	0.3146	-0.3576	0.1007	-0.7633	0.0013	0.0087	0.9900	0.5249	0.0055	0.5162	0.0035	0.2302	0.7215	-0.0504	0.9027	-0.2806	0.3097
CG13792	CG13792	1636629_at	2.3406	0.0121	2.6280	0.0042	2.9341	0.0005	0.4198	0.5008	0.6171	0.0654	0.1973	0.5421	0.3529	0.9108	0.9491	0.3909	0.5962	0.6216
CG5091 /// CG5096	CG5096 /// CG50	1636630_s_at	-0.4260	0.2916	-0.1658	0.5048	-0.6632	0.1303	-0.5960	0.1341	0.1472	0.5702	0.7432	0.0049	-0.0451	0.9848	0.3814	0.4794	0.4265	0.4318
CG1139	CG1139	1636631_at	0.2949	0.6178	0.0638	0.7861	0.0777	0.7819	0.1597	0.9171	-0.4633	0.3369	-0.6230	0.1408	0.0708	0.9775	-0.4877	0.4089	-0.5585	0.3616
CG9784	CG9784	1636632_at	-0.6968	0.0801	0.4443	0.1784	0.7893	0.0014	0.1869	0.6041	-0.8801	0.0012	-1.0670	0.0003	-0.2438	0.8400	0.2911	0.5823	0.5349	0.2964

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1636633_at	0.2673	0.0806	0.0454	0.7426	-0.0426	0.8355	-0.0216	0.9722	-0.0317	0.8778	-0.0101	0.9589	-0.0221	0.9816	-0.1597	0.4754	-0.1377	0.5540
MRP	Multidrug-resistan	1636634_a_at	-0.3538	0.1077	-0.5005	0.0291	-1.0313	0.0004	-0.2455	0.4643	0.4073	0.0322	0.6528	0.0025	0.2541	0.6898	0.1728	0.5192	-0.0813	0.7943
CG12164	CG12164	1636635_at	0.0905	0.6050	-0.0051	0.9677	0.0584	0.8006	0.1218	0.7304	-0.0361	0.8619	-0.1579	0.2679	0.1032	0.8479	-0.0020	0.9975	-0.1053	0.6656
CG11771	CG11771	1636636_at	-0.4856	0.0159	-0.2012	0.1625	0.0458	0.8119	0.0214	0.9726	-0.2998	0.0564	-0.3212	0.0277	-0.2805	0.7142	-0.1547	0.6594	0.1259	0.7307
CG13731	CG13731	1636637_x_at	0.4599	0.1261	0.3924	0.3036	0.3365	0.2822	-0.0937	0.9300	-0.1806	0.5977	-0.0869	0.8049	0.1204	0.9333	-0.0808	0.9085	-0.2012	0.6942
DyakCG7217	CG7217	1636638_s_at	-0.1179	0.7143	0.2615	0.3245	0.4253	0.0311	-0.0362	0.9592	-1.0119	0.0007	-0.9757	0.0005	-0.1893	0.8846	-0.4344	0.3713	-0.2451	0.6452
Spp	shanti	1636639_at	0.9661	0.0017	0.7456	0.0231	1.0661	0.0003	0.1950	0.5179	0.5543	0.0048	0.3593	0.0197	-0.0258	0.9816	0.4549	0.0800	0.4807	0.0891
CG14811	CG14811	1636640_at	0.5556	0.0134	0.0679	0.7241	0.1983	0.2873	0.0105	0.9874	0.0562	0.7697	0.0457	0.7990	-0.1013	0.8425	-0.3053	0.1395	-0.2040	0.3427
CG40275	CG40275	1636641_at	0.0350	0.8869	0.0085	0.9515	-0.0638	0.8089	-0.1520	0.7711	0.0039	0.9901	0.1559	0.4545	-0.2090	0.6272	-0.0572	0.8040	0.1518	0.4069
CG13712	CG13712	1636642_at	0.0165	0.9173	0.0576	0.6594	-0.1548	0.4539	0.0163	0.9854	0.1091	0.6265	0.0928	0.6568	0.2621	0.6389	0.0769	0.7877	-0.1852	0.4296
CG6983	CG6983	1636643_at	-0.9797	0.0107	-0.9235	0.1408	-1.4939	0.0001	-0.2331	0.6506	-0.2565	0.3060	-0.0234	0.9395	0.2573	0.8609	-0.2716	0.6827	-0.5289	0.3851
RpII18	RNA polymerase	1636644_at	0.5471	0.0675	0.6005	0.0618	0.4250	0.0580	0.1806	0.6010	0.2930	0.0926	0.1124	0.5016	0.2499	0.7956	0.2548	0.5488	0.0048	0.9946
Nup75	CG5733	1636645_at	1.0712	0.0010	0.7253	0.1694	0.7004	0.0016	-0.0323	0.9603	0.3970	0.0298	0.4293	0.0136	-0.0832	0.9514	0.1213	0.8185	0.2045	0.6399
sna	snail	1636646_at	-0.0077	0.9868	-0.1228	0.3777	0.2094	0.4496	0.1447	0.8578	0.0236	0.9562	-0.1212	0.6914	-0.1592	0.8692	-0.0201	0.9763	0.1391	0.7534
CG40160	CG40160	1636647_s_at	-0.0780	0.7644	-0.2594	0.4685	-0.5904	0.0044	-0.2264	0.3863	0.1022	0.5138	0.3286	0.0212	-0.0121	0.9952	-0.1002	0.8792	-0.0881	0.8893
---	---	1636648_s_at	-0.3218	0.4980	-0.3491	0.4369	-0.3632	0.0279	-0.0684	0.8907	0.4658	0.0129	0.5342	0.0041	-0.0679	0.9816	0.4267	0.5258	0.4945	0.4615
CG10107 /// DmirCG10107	CG10107	1636649_a_at	0.0794	0.8657	0.2711	0.6029	0.4459	0.0267	0.2184	0.7031	-0.0659	0.8471	-0.2843	0.2290	0.0921	0.9734	0.2325	0.7810	0.1403	0.8788
CG10176	CG10176	1636650_at	-0.5400	0.1302	0.6146	0.2464	-0.2993	0.4346	0.0714	0.8905	-0.3980	0.0296	-0.4694	0.0091	0.8684	0.6272	0.6737	0.3534	-0.1947	0.8364
Csp	cysteine string prc	1636651_a_at	0.1887	0.3144	0.4408	0.0559	-0.2235	0.2670	-0.0709	0.8822	0.4349	0.0159	0.5058	0.0049	0.5733	0.3273	0.6460	0.0568	0.0727	0.8561
CG14955	CG14955	1636652_a_at	0.0299	0.8861	-0.3815	0.0255	-0.1952	0.3923	0.1655	0.6504	0.2735	0.1185	0.1080	0.5288	-0.0469	0.9491	-0.0747	0.7828	-0.0278	0.9271
nec	serpin43Ac	1636653_at	-0.6061	0.1145	-0.7905	0.3026	-0.1738	0.4014	0.2383	0.6506	-0.2617	0.3073	-0.5000	0.0348	-0.3674	0.8603	-0.2419	0.8262	0.1255	0.9135
CG15626	CG15626	1636654_s_at	0.6708	0.0250	0.8664	0.0309	0.8437	0.0037	-0.1837	0.6988	-0.5013	0.0291	-0.3176	0.1024	-0.1137	0.9199	-0.2952	0.4406	-0.1816	0.6622
---	---	1636655_s_at	0.2779	0.2296	-0.0060	0.9620	-0.0845	0.6733	0.4305	0.2876	0.3779	0.1121	-0.0526	0.8490	0.1865	0.7810	-0.2102	0.4757	-0.3967	0.1952
CG13379	CG13379	1636656_at	0.4249	0.0344	0.4319	0.0755	0.3059	0.1336	-0.0505	0.9263	-0.1085	0.5368	-0.0580	0.7434	0.0440	0.9653	-0.0105	0.9841	-0.0545	0.8831
CheA87a	CheA87a	1636657_at	0.1122	0.6396	-0.1844	0.4115	0.1658	0.3039	0.1343	0.8358	0.0970	0.7368	-0.0373	0.8992	-0.1678	0.7770	-0.1287	0.6447	0.0391	0.9121
Lcp65Ad	Lcp65Ad	1636658_at	0.0595	0.7444	0.0368	0.7124	0.0603	0.7414	-0.0212	0.9755	-0.0856	0.6522	-0.0644	0.7202	-0.0178	0.9862	0.0696	0.8168	0.0873	0.7434
Cpr65Ec	CG8634	1636659_at	-0.0931	0.5870	0.0359	0.7343	0.1571	0.3355	0.0756	0.8628	-0.0459	0.8112	-0.1215	0.4016	-0.0162	0.9848	0.0315	0.9164	0.0477	0.8477
CG14903	CG14903	1636660_at	0.1828	0.2632	-0.2450	0.4000	-0.3931	0.0261	0.0046	0.9956	0.3959	0.0248	0.3913	0.0163	0.1409	0.8650	-0.0446	0.9326	-0.1855	0.6016
---	---	1636661_at	0.2488	0.1290	0.0649	0.5185	0.1461	0.4587	-0.0331	0.9639	-0.0002	0.9994	0.0329	0.8864	-0.0337	0.9667	-0.0624	0.8247	-0.0287	0.9231
CG8945	CG8945	1636662_at	0.1436	0.4002	0.0377	0.8075	-0.0394	0.8618	0.0361	0.9500	0.0342	0.8718	-0.0019	0.9924	0.0466	0.9514	-0.0684	0.8185	-0.1150	0.6399
CG7126	CG7126	1636663_at	-0.7465	0.0034	-1.6978	0.0048	-0.7466	0.0527	0.9984	0.0976	0.9125	0.0183	-0.0859	0.8317	-0.1224	0.8906	-0.2747	0.4089	-0.1523	0.6800
CG1077	CG1077	1636664_at	-0.0921	0.7328	-0.0949	0.3828	-0.1991	0.1906	0.0229	0.9826	0.1160	0.6789	0.0931	0.7219	0.0130	0.9893	-0.0557	0.8356	-0.0687	0.7708
---	---	1636665_at	0.0768	0.7442	-0.1165	0.2810	0.2418	0.2673	0.0672	0.9020	0.0872	0.6648	0.0200	0.9264	-0.1923	0.8049	-0.0852	0.8472	0.1071	0.7800
CG32591	CG32591	1636666_at	0.2691	0.2312	-0.0342	0.7701	0.1285	0.6708	-0.0094	0.9917	0.0021	0.9941	0.0115	0.9614	0.0427	0.9741	-0.1805	0.6006	-0.2233	0.5050
---	---	1636667_at	-0.1676	0.3600	0.1523	0.5212	0.1247	0.4603	-0.0939	0.9017	-0.2259	0.3701	-0.1320	0.5934	0.0932	0.8543	0.0693	0.7921	-0.0239	0.9348
CG9972	CG9972	1636668_at	0.0192	0.9555	-0.1621	0.3981	-0.1493	0.3445	0.0405	0.9675	-0.0879	0.7824	-0.1285	0.6318	0.0957	0.9246	-0.1731	0.6344	-0.2689	0.4370
Ca-P60A	organellar-type Ca	1636669_at	-1.1259	0.0845	-2.0455	0.0190	-3.0353	0.0000	-0.3683	0.4704	0.2502	0.3728	0.6185	0.0201	0.4407	0.8486	-0.8207	0.3856	-1.2614	0.2094
---	---	1636670_at	0.1013	0.5706	-0.0631	0.5790	0.1292	0.4181	0.0085	0.9891	0.2140	0.1294	0.2055	0.1038	-0.1341	0.8283	0.0411	0.9185	0.1753	0.5176
CG13479	CG13479	1636671_at	-0.1644	0.5292	0.1872	0.2245	-0.0696	0.7277	-0.0932	0.8987	-0.1394	0.5940	-0.0461	0.8704	0.3664	0.3712	0.3239	0.1394	-0.0425	0.8889
Brd	Bearded	1636672_at	0.0066	0.9844	-0.1405	0.4047	-0.0357	0.8564	0.2212	0.6510	0.1597	0.5256	-0.0615	0.8184	-0.0945	0.9246	-0.0616	0.9046	0.0329	0.9435
ttk	tramtrack-69	1636673_s_at	-0.4560	0.3591	0.2920	0.3137	-0.6922	0.0093	0.0324	0.9612	0.2788	0.1168	0.2464	0.1211	0.9750	0.5754	0.9879	0.1948	0.0129	0.9924
CG11407	CG11407	1636674_at	-1.1483	0.3193	-1.1035	0.1097	-1.8072	0.0002	-0.5916	0.6120	-1.2321	0.0414	-0.6405	0.2209	-0.1636	0.9776	-1.6864	0.2110	-1.5228	0.2869
CG31145	CG31145	1636675_at	0.1934	0.3240	0.1399	0.4046	0.0705	0.6789	0.0041	0.9956	-0.1498	0.4205	-0.1539	0.3517	0.0235	0.9721	-0.1040	0.5609	-0.1275	0.4690
CG2150	CG2150	1636676_at	0.2346	0.3737	0.0104	0.9753	-0.0921	0.6795	0.0877	0.8908	0.2040	0.3491	0.1163	0.5869	0.2520	0.7686	0.0295	0.9614	-0.2225	0.5647
CG11876 /// DyakCG11876	CG11876	1636677_s_at	-0.1960	0.4731	0.2399	0.0624	0.3186	0.0740	-0.1054	0.7742	-0.7188	0.0014	-0.6134	0.0016	-0.1258	0.8973	-0.2145	0.5776	-0.0887	0.8501
CG14343	CG14343	1636678_at	0.0778	0.6800	0.2153	0.3949	-0.0625	0.8615	-0.0450	0.9375	0.0095	0.9708	0.0545	0.7668	0.2402	0.7810	0.1041	0.8353	-0.1361	0.7520
Mrtf	Myocardin-related	1636679_at	0.7138	0.1818	0.9783	0.1953	-0.0189	0.9756	-0.3328	0.4174	0.5895	0.0179	0.9222	0.0015	0.8568	0.7768	1.1234	0.3802	0.2666	0.8806
---	---	1636680_at	0.1696	0.2845	0.1372	0.4341	0.3289	0.0820	-0.0473	0.9451	-0.0656	0.7824	-0.0183	0.9397	0.0318	0.9721	-0.0318	0.9312	-0.0636	0.8256
CG14621	CG14621	1636681_at	0.1361	0.7583	1.2267	0.0247	1.2242	0.0013	-0.1414	0.8485	-0.2141	0.4469	-0.0726	0.8123	-0.2196	0.8906	0.8170	0.1617	1.0366	0.1187
CG31778	CG31778	1636682_at	2.0981	0.0048	1.1336	0.2912	1.5990	0.0019	0.0253	0.9893	-0.0585	0.9226	-0.0838	0.8682	-0.3442	0.9156	-0.8478	0.4563	-0.5036	0.6877
CG14610	CG14610	1636683_at	0.0702	0.6563	-0.0651	0.6100	0.1081	0.4994	0.0978	0.8623	-0.0834	0.7218	-0.1813	0.3234	0.0153	0.9862	-0.0829	0.7304	-0.0982	0.6622
CG31126	CG31126	1636684_at	-0.0397	0.8696	0.2653	0.4005	0.4041	0.1077	-0.0710	0.8738	-0.2875	0.0682	-0.2166	0.1191	-0.1393	0.9056	0.0602	0.9277	0.1995	0.6623
---	---	1636685_at	0.1633	0.4007	0.0442	0.7275	0.1524	0.4509	-0.0444	0.9502	0.0871	0.7014	0.1315	0.4843	-0.0184	0.9871	-0.0306	0.9420	-0.0122	0.9769
garz	gartenzwerg	1636686_at	0.1419	0.6507	0.1833	0.2726	0.5700	0.0241	0.3060	0.3625	0.2392	0.2099	-0.0668	0.7478	0.1497	0.8571	0.3422	0.3002	0.1925	0.5984
CG34366	CG13111	1636687_at	0.1762	0.5045	0.2524	0.0865	0.1955	0.4553	0.0549	0.9592	0.1069	0.7511	0.0520	0.8774	0.0393	0.9814	0.0525	0.9265	0.0132	0.9835
Cyp4s3	Cyp4s3	1636688_at	0.5743	0.2692	0.0891	0.8083	0.7098	0.0652	-0.1237	0.8211	0.0191	0.9521	0.1429	0.4835	-0.8128	0.7092	-0.4635	0.6384	0.3492	0.7394
CG4259	CG4259	1636689_at	0.7830	0.0868	0.2891	0.6479	0.5435	0.1079	-0.4663	0.5125	-0.3959	0.2968	0.0705	0.8740	-0.8135	0.6496	-0.9644	0.1700	-0.1509	0.8779
Or42a	Odorant receptor	1636690_at	-0.0883	0.6956	-0.0696	0.5697	0.1240	0.5405	0.0334	0.9610	-0.0442	0.8519	-0.0776	0.6849	-0.1900	0.7893	-0.0242	0.9619	0.1658	0.6129
Sfmbt	CG16975	1636691_a_at	0.0711	0.7975	-0.0236	0.9113	-0.2307	0.1488	-0.1750	0.5913	0.3119	0.0627	0.4869	0.0061	0.0790	0.9441	0.3033	0.3667	0.2243	0.5332
---	---	1636692_at	0.2279	0.2491	0.0816	0.6766	0.1354	0.4615	-0.0268	0.9672	-0.0427	0.8463	-0.0159	0.9397	-0.0427	0.9618	-0.2293	0.3446	-0.1865	0.4683
CG4330	CG4330	1636693_at	-0.4051	0.2162	0.0717	0.9047	0.1790	0.4796	-0.2198	0.6202	-0.4592	0.0437	-0.2394	0.2262	-0.4533	0.7644	-0.0045	0.9987	0.4488	0.4944
CG10673	CG10673	1636694_at	-0.2364	0.2587	-0.0707	0.7566	-0.0578	0.7805	0.0766	0.8987	-0.2566	0.1891	-0.3332	0.0616	0.1321	0.8541	0.0485	0.9157	-0.0836	0.8207
Pk17E	Bicoid interacting	1636695_at	0.0021	0.9941	0.4658	0.1389	-0.0374	0.8499	-0.2577	0.4861	-0.2327	0.2401	0.0251	0.9175	0.2379	0.8065	0.1881	0.6762	-0.0498	0.9300
CG32804	CG32804	1636696_at	0.1568	0.3914	-0.0067	0.9745	-0.2245	0.3823	-0.0101	0.9909	0.2318	0.2305	0.2420	0.1610	0.0886	0.9235	-0.1066	0.7737	-0.1953	0.5420
---	---	1636697_at	0.2419	0.1777	0.0651	0.6131	-0.0230	0.9361	-0.0239	0.9755	0.0391	0.8778	0.0630	0.7630	0.1650	0.8293	-0.0605	0.9009	-0.2255	0.4978
nxf4	nuclear export fac	1636698_at	-0.0593	0.7582	0.1771	0.1123	-0.1057	0.5516	-0.1592	0.7380	-0.1732	0.4262	-0.0140	0.9584	0.1878	0.8192	-0.0754	0.8754	-0.2632	0.4465
---	---	1636699_at	0.0452	0.8720	-0.0445	0.7273	0.1406	0.4587	0.0504	0.9385	-0.1617	0.4125	-0.2121	0.2155	-0.0339	0.9773	-0.2161	0.4388	-0.1823	0.5361
---	---	1636700_at	0.1775	0.3248	-0.1681	0.1798	-0.3083	0.1764	0.0840	0.9035	0.3538	0.1137	0.2698	0.1776	0.0019	0.9992	-0.0506	0.9009	-0.0525	0.8870
fz2	D-frizzled2	1636701_at	-0.0977	0.5164	0.0174	0.8982	-0.0923	0.6361	0.0141	0.9888	-0.2202	0.3536	-0.2343	0.2648	0.0817	0.9291	-0.1721	0.5847	-0.2538	0.4011

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG15343	CG15343	1636702_at	0.3328	0.1622	0.7635	0.0287	0.8616	0.0019	0.4930	0.1549	0.5224	0.0210	0.0295	0.9078	0.3163	0.6955	0.9127	0.0290	0.5964	0.1138
CG31817	CG31817	1636703_at	-0.3328	0.2991	-0.6443	0.1190	-0.7942	0.0032	-0.0909	0.9373	0.2466	0.4822	0.3375	0.2635	-0.1439	0.8425	-0.1393	0.6713	0.0046	0.9928
CG34104	CG34104	1636704_at	0.0153	0.9528	0.0067	0.9523	0.2017	0.1732	0.0005	0.9994	-0.0780	0.6792	-0.0785	0.6445	-0.0178	0.9816	0.0125	0.9649	0.0303	0.8984
mbo	members only	1636705_at	0.8594	0.0039	0.2336	0.4097	0.7085	0.0029	0.2994	0.2870	0.5066	0.0085	0.2072	0.1598	-0.1475	0.8222	-0.1781	0.5310	-0.0306	0.9362
Myd88	krapfen	1636706_at	-0.3900	0.0512	-0.8359	0.0902	-1.2393	0.0061	-0.3229	0.5735	-0.0057	0.9899	0.3173	0.2271	0.0377	0.9852	-0.4750	0.2938	-0.5127	0.2842
---	---	1636707_at	0.1179	0.5789	0.1036	0.4172	0.1431	0.5177	-0.0015	0.9987	-0.1743	0.3761	-0.1728	0.3258	0.0989	0.8461	-0.0556	0.8460	-0.1545	0.4791
CG3061	dnaJ like2	1636708_at	-0.3861	0.0291	-0.1256	0.5514	-0.2489	0.1025	-0.2088	0.4456	-0.1408	0.3553	0.0681	0.6604	-0.0842	0.9180	0.1346	0.6658	0.2188	0.4539
coilin	CG8710	1636709_at	-0.5122	0.1021	0.1141	0.8014	0.1996	0.2001	-0.1264	0.7093	-0.1437	0.3616	-0.0173	0.9268	-0.1772	0.8940	0.4588	0.3446	0.6360	0.2189
Neu2	Neu2	1636710_at	0.2191	0.2028	0.0357	0.7974	0.1367	0.3736	0.0420	0.9507	0.0161	0.9542	-0.0259	0.9095	-0.1581	0.7758	-0.1184	0.6497	0.0396	0.9055
CG31265	CG31265	1636711_at	-0.2581	0.8923	-0.3801	0.1709	0.0534	0.7660	0.5556	0.8518	-0.5786	0.6310	-1.1342	0.2426	-0.0755	0.9898	-0.7189	0.6012	-0.6434	0.6423
RpL21	Ribosomal protein	1636712_at	0.0409	0.7951	0.1530	0.2611	0.2424	0.1454	0.0172	0.9777	-0.1497	0.3356	-0.1669	0.2233	-0.0101	0.9914	0.0328	0.9156	0.0429	0.8729
CG14985	CG14985	1636713_at	-0.1724	0.3361	0.2706	0.3380	-0.1257	0.5476	-0.1571	0.8507	-0.1194	0.7396	0.0378	0.9193	0.2982	0.6749	0.1582	0.6191	-0.1400	0.6639
---	---	1636714_at	0.2795	0.0908	0.0788	0.7118	0.0973	0.6552	-0.0622	0.9293	0.0953	0.6844	0.1575	0.4115	0.1139	0.8917	0.0793	0.8546	-0.0346	0.9372
---	---	1636715_at	-0.0532	0.7667	0.0520	0.7696	0.1487	0.3309	0.1824	0.6872	0.0089	0.9784	-0.1734	0.3728	0.1106	0.8734	0.1482	0.6118	0.0376	0.9199
Cyp4c3	Cytochrome P450	1636716_at	0.5062	0.1027	-0.1638	0.2442	0.3036	0.1332	0.5546	0.1737	0.7165	0.0109	0.1619	0.4950	-0.0380	0.9816	-0.2084	0.6093	-0.1703	0.6867
CG6005	CG6005	1636717_at	-0.4572	0.0850	-0.9094	0.0164	-1.1258	0.0004	-0.1049	0.8362	0.3240	0.0914	0.4289	0.0205	0.2677	0.6607	0.0407	0.9157	-0.2270	0.3723
CG4502	CG4502	1636718_s_at	0.3949	0.0781	0.1896	0.5617	-0.1614	0.3232	-0.0573	0.9110	0.2316	0.1523	0.2889	0.0514	0.2560	0.7726	0.0918	0.8628	-0.1642	0.6990
CG5316	CG5316	1636719_s_at	0.1020	0.6531	0.0125	0.9041	0.1622	0.3834	-0.0916	0.8735	-0.2093	0.2995	-0.1177	0.5490	-0.0842	0.9309	-0.1728	0.6093	-0.0885	0.8237
---	---	1636720_at	0.1415	0.4671	0.1314	0.3222	0.1807	0.3166	-0.0867	0.9255	-0.1171	0.7190	-0.0304	0.9291	-0.1186	0.8145	-0.0304	0.9270	0.0882	0.7075
Pc	Polycomb	1636721_at	0.3589	0.0794	0.0605	0.8970	-0.2051	0.4205	-0.1195	0.8889	0.6264	0.0357	0.7459	0.0108	0.2475	0.8062	0.3803	0.3571	0.1329	0.7933
CG31917	CG31917	1636722_at	-0.0357	0.9132	0.2945	0.2415	0.4062	0.1083	-0.0386	0.9491	-0.1625	0.3427	-0.1239	0.4333	-0.0014	0.9998	0.3007	0.6032	0.3021	0.6045
---	---	1636723_at	0.5054	0.0584	0.0644	0.7239	0.1296	0.4683	-0.0172	0.9777	0.1443	0.3633	0.1615	0.2473	-0.2128	0.8097	-0.2848	0.4379	-0.0719	0.8870
CG8292	CG8292	1636724_at	-0.2693	0.2809	-0.1597	0.3848	-0.3829	0.1149	0.1251	0.8162	0.1771	0.4170	0.0520	0.8293	0.0876	0.9216	0.1341	0.6864	0.0464	0.9096
CG5334	CG5334	1636725_at	0.0400	0.8344	0.0576	0.5997	-0.0069	0.9771	-0.1322	0.7123	-0.1262	0.4577	0.0060	0.9766	-0.1228	0.8940	-0.1296	0.7489	-0.0067	0.9909
CG17180	CG17180	1636726_at	-0.1312	0.5547	0.2415	0.1494	0.2644	0.2377	0.2109	0.7190	-0.1485	0.6122	-0.3595	0.1287	0.2878	0.7739	0.3135	0.4794	0.0257	0.9694
CG4768	CG4768	1636727_at	-0.0057	0.9907	1.1329	0.0355	0.5810	0.0308	-0.6336	0.2848	-0.3304	0.3517	0.3032	0.3423	-0.1383	0.9246	0.7058	0.1418	0.8441	0.1184
Ptr	Ptc-related Disp-li	1636728_at	-1.8272	0.0084	-3.5180	0.0047	-3.2449	0.0000	0.4517	0.6324	1.0873	0.0264	0.6356	0.1221	0.1892	0.8719	-0.5265	0.2408	-0.7157	0.1498
---	---	1636729_at	0.0836	0.7673	0.1384	0.5244	0.1875	0.2237	0.0942	0.8707	0.0313	0.9114	-0.0629	0.7805	-0.0053	0.9965	0.0016	0.9991	0.0069	0.9894
CG13373	CG13373	1636730_at	0.0315	0.9284	-0.4783	0.2734	-0.4670	0.0339	-0.1626	0.7408	0.3733	0.0854	0.5359	0.0134	-0.2111	0.8686	-0.1444	0.8285	0.0667	0.9231
CG30357	CG30357	1636731_at	0.0460	0.8147	0.1226	0.4189	0.2104	0.1347	-0.0321	0.9592	-0.0900	0.6207	-0.0579	0.7438	0.0369	0.9542	0.0629	0.7897	0.0260	0.9198
CG11695	CG11695	1636732_at	-0.6209	0.0218	-0.0979	0.3180	-0.2229	0.3223	0.0672	0.9066	-0.4827	0.0170	-0.5500	0.0058	0.0945	0.8574	0.0154	0.9650	-0.0791	0.7492
CG32135	CG32135	1636733_at	1.2927	0.0568	0.6675	0.1450	-0.4129	0.4559	-0.2929	0.6361	0.9440	0.0081	1.2370	0.0015	0.9703	0.7202	0.8430	0.4556	-0.1273	0.9354
---	---	1636734_at	-0.0233	0.8921	0.0570	0.6149	0.1578	0.5279	0.0325	0.9647	-0.1001	0.6474	-0.1326	0.4736	-0.0548	0.9409	-0.0686	0.8182	-0.0137	0.9688
CG3065	CG3065	1636735_s_at	-0.1917	0.3101	0.1849	0.4522	0.0394	0.8070	0.1506	0.6720	0.1303	0.4577	-0.0203	0.9218	0.3334	0.6749	0.4895	0.1349	0.1561	0.6605
CG31274 /// MESK4	CG31274 /// Mise	1636736_s_at	-0.3098	0.2182	0.0985	0.3194	-0.1553	0.2867	-0.2191	0.4815	-0.3773	0.0309	-0.1581	0.2947	-0.0697	0.9514	-0.0783	0.8710	-0.0086	0.9874
---	---	1636737_at	-0.0737	0.7186	-0.0496	0.6416	0.0554	0.7460	0.1377	0.7857	0.0502	0.8529	-0.0874	0.6903	0.0208	0.9862	0.0438	0.9192	0.0229	0.9530
RpL37b	Ribosomal protein	1636738_at	0.1513	0.5571	0.0894	0.4982	0.2638	0.1426	-0.0299	0.9627	-0.1418	0.4188	-0.1119	0.4917	0.0033	0.9978	0.0638	0.8610	0.0605	0.8599
CG12551	CG12551	1636739_at	0.1447	0.5071	0.1617	0.2521	0.0626	0.7174	0.0247	0.9759	-0.0529	0.8373	-0.0776	0.7158	0.1522	0.8270	0.1427	0.6536	-0.0096	0.9846
Fdh	glutathione-depen	1636740_at	0.7829	0.0035	1.3581	0.0055	1.7886	0.0000	0.1897	0.5249	-0.8548	0.0007	-1.0446	0.0002	-0.1489	0.8379	-0.1496	0.6454	-0.0007	0.9992
Hsc70Cb	Hsc70Cb	1636741_s_at	-0.2951	0.4960	0.2200	0.3532	0.2978	0.2882	0.2288	0.7939	0.4683	0.1876	0.2396	0.4824	0.2728	0.8270	1.0345	0.0661	0.7617	0.1655
nord	nord	1636742_at	-3.0008	0.0005	-1.7289	0.0329	-1.9789	0.0042	-1.0365	0.1153	-2.2663	0.0006	-1.2298	0.0048	-0.8975	0.6724	-1.1839	0.1536	-0.2864	0.7764
---	---	1636743_s_at	-3.1290	0.0033	-1.2913	0.0207	-2.6360	0.0005	-1.1686	0.1332	-1.8178	0.0030	-0.6492	0.1205	-0.0847	0.9787	-0.3294	0.6931	-0.2447	0.7800
---	---	1636744_at	-0.0268	0.8847	0.0807	0.5606	-0.0352	0.8224	-0.1410	0.7138	-0.1932	0.2723	-0.0522	0.7900	0.0832	0.9174	0.1135	0.7202	0.0304	0.9373
Art4	Arginine methyltra	1636745_at	0.7129	0.0338	-0.0136	0.9618	0.9329	0.0004	-0.0268	0.9803	0.0992	0.7495	0.1260	0.6386	-0.8158	0.1628	-0.4718	0.0834	0.3439	0.2094
CG13826	CG13826	1636746_at	-0.2562	0.2549	-0.0494	0.7185	-0.2901	0.0835	-0.3257	0.4482	-0.2821	0.2269	0.0436	0.8748	-0.0791	0.9246	0.0358	0.9367	0.1149	0.7179
CG8051	CG8051	1636747_at	1.1667	0.0554	-0.2894	0.2535	-0.0600	0.8711	-0.2029	0.8676	0.0427	0.9465	0.2457	0.5629	-0.5400	0.7215	-1.4321	0.0449	-0.8921	0.1823
---	---	1636748_at	0.1487	0.3013	-0.1932	0.1954	-0.2188	0.2098	-0.1773	0.5079	0.2529	0.0769	0.4301	0.0054	-0.0570	0.9317	-0.0354	0.9156	0.0215	0.9433
---	---	1636749_at	0.1138	0.6550	-0.0684	0.5935	-0.3157	0.2197	-0.3526	0.2943	0.0592	0.8087	0.4117	0.0277	-0.1106	0.9420	0.1049	0.8754	0.2154	0.6734
CG13737	CG13737	1636750_s_at	1.3518	0.0237	0.1696	0.3761	0.0608	0.7938	-0.0863	0.9254	-0.0860	0.7998	0.0003	0.9995	0.2250	0.9036	-1.1948	0.0881	-1.4198	0.0729
---	---	1636751_x_at	0.2404	0.2319	0.1205	0.3599	0.1359	0.3448	0.1054	0.8244	0.1519	0.4246	0.0465	0.8241	0.0740	0.8870	0.1296	0.5214	0.0556	0.8190
---	---	1636752_at	0.4047	0.0860	0.0501	0.8630	0.4457	0.0155	0.0944	0.8816	0.0319	0.9166	-0.0625	0.7971	-0.1032	0.9246	-0.0788	0.8807	0.0244	0.9640
CG7094	CKIa-like	1636753_at	0.1447	0.3977	0.1267	0.4625	0.0820	0.7523	-0.0113	0.9921	0.0485	0.8739	0.0598	0.8194	-0.0880	0.9238	-0.0501	0.9157	0.0379	0.9293
---	---	1636754_at	-0.0778	0.7104	0.0062	0.9655	0.1648	0.2362	0.0908	0.8811	-0.0611	0.8112	-0.1518	0.4349	0.0285	0.9816	0.0883	0.8056	0.0598	0.8739
CG13966	CG13966	1636755_at	-0.8595	0.1414	0.7776	0.0980	0.7348	0.0218	-0.1931	0.6999	-1.0048	0.0018	-0.8117	0.0027	0.1312	0.9653	0.5832	0.4810	0.4521	0.6043
CG31627	CG31627	1636756_at	0.1548	0.4421	0.2460	0.4331	0.2983	0.2306	0.0226	0.9777	-0.0444	0.8725	-0.0670	0.7716	0.0021	0.9994	-0.0530	0.9167	-0.0551	0.9028
CG14107	CG14107	1636757_at	0.2311	0.2273	0.0317	0.8026	0.0991	0.6369	0.1264	0.8811	0.1917	0.5332	0.0654	0.8442	0.0558	0.9467	0.0977	0.7455	0.0419	0.9057
CG18853 /// phr	CG18853 /// photc	1636758_s_at	-0.2111	0.3630	-0.4090	0.2549	-0.4232	0.1144	-0.0959	0.9339	-0.0227	0.9653	0.0732	0.8526	-0.1695	0.8564	-0.1465	0.7454	0.0230	0.9688
CG8303	CG8303	1636759_at	-0.2544	0.7449	-2.3318	0.0111	-1.0266	0.0152	1.1852	0.1109	1.2343	0.0127	0.0491	0.9274	-0.0818	0.9848	-0.8111	0.3949	-0.7294	0.4653
CG13125	CG13125	1636760_a_at	-0.0955	0.6299	0.2048	0.2503	0.3337	0.1617	-0.1881	0.6338	-0.2761	0.1529	-0.0880	0.6618	-0.1129	0.8768	-0.0023	0.9980	0.1106	0.7346
---	---	1636761_at	0.4678	0.1269	0.3227	0.0575	0.0608	0.8458	-0.1070	0.9105	-0.1502	0.6587	-0.0432	0.9053	0.1823	0.8157	-0.0412	0.9363	-0.2235	0.5049
scf	supercoiling factor	1636762_a_at	-1.5267	0.0024	-1.4111	0.0536	-0.9448	0.0024	-0.3237	0.5179	-0.5133	0.0563	-0.1896	0.4459	-0.6166	0.6898	-0.2785	0.6971	0.3381	0.6216
CG33922 /// CG33923	CG33922 /// CG3	1636763_at	-0.0539	0.8296	-0.0295	0.8082	-0.0032	0.9885	0.0809	0.8732	0.1099	0.5626	0.0291	0.8904	-0.0416	0.9651	-0.0019	0.9985	0.0398	0.9096
CG31075	CG31075	1636764_at	0.1116	0.7392	0.3801	0.1350	-0.1349	0.6135	-0.9060	0.1663	-1.1837	0.0097	-0.2777	0.4615	-0.4124	0.5126	-0.9217	0.0222	-0.5093	0.1200
mod(mdg4)	Modifier67.2	1636765_at	-0.5739	0.1007	0.4039	0.4056	0.1086	0.6612	-0.0786	0.9441	-0.5674	0.0793	-0.4887	0.0892	0.3193	0.8270	0.4328	0.4854	0.1135	0.8918
Ent2	Equilibrative nucle	1636766_at	0.8999	0.0144	1.3038	0.0440	1.5098	0.0001	0.2223	0.5582	-0.0759	0.7423	-0.2982	0.0881	-0.0539	0.9816	0.1305	0.8604	0.1844	0.7668
LysP	lysozyme	1636767_at	0.2051	0.5309	0.0089	0.9379	0.1776	0.4253	0.1439	0.7409	0.0078	0.9792	-0.1361	0.4527	0.0672	0.9653	-0.0945	0.8734	-0.1617	0.7346
---	---	1636768_s_at	-0.8981	0.0035	-1.6846	0.0114	-2.4808	0.0000	-0.3330	0.4171	1.2525	0.0007	1.5855	0.0002	0.1336	0.9101	0.3883	0.3458	0.2547	0.5702
CG3651	CG3651	1636769_at	-0.1404	0.4051	-0.4928	0.1259	-0.3276	0.1186	-0.0762	0.8846	0.2959	0.0983	0.3720	0.0278	-0.1796	0.8235	-0.0572	0.9115	0.1224	0.7516
Arc-p20	p20 subunit of Arp	1636770_at	-0.0457	0.8406	-0.0638	0.6310	0.2095	0.3709	0.1188	0.7596	0.2485	0.1350	0.1297	0.4045	0.0061	0.9971	0.4581	0.2459	0.4520	0.2841

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CoRest	CG33525	1636771_at	-0.5463	0.2589	-0.1487	0.7547	-0.0335	0.8341	-0.2138	0.5407	-0.0204	0.9377	0.1933	0.2356	-0.2351	0.8903	0.3238	0.6414	0.5589	0.3921
CG32694	CG32694	1636772_s_at	-2.4343	0.0004	-2.8295	0.0023	-3.1886	0.0000	-0.1300	0.7862	-0.2375	0.2296	-0.1075	0.5869	0.1487	0.8202	-0.6562	0.0391	-0.8049	0.0333
CG10981	CG10981	1636773_a_at	0.4786	0.0209	-0.0112	0.9702	-0.2899	0.1005	-0.1048	0.7647	0.4701	0.0077	0.5749	0.0019	0.1237	0.8610	0.0455	0.9184	-0.0782	0.8287
cpsf	Drosophila cleava	1636774_at	0.4412	0.0440	-0.1520	0.6679	-0.0366	0.8123	-0.0660	0.8791	0.6768	0.0015	0.7428	0.0006	-0.0735	0.9677	0.0779	0.9167	0.1514	0.7933
CG18622	CG18622	1636775_at	0.0298	0.9402	0.9259	0.0122	0.7655	0.0597	0.0697	0.9108	-0.4371	0.0337	-0.5068	0.0114	0.2231	0.9092	0.4705	0.5093	0.2474	0.7558
CG40359	CG40359	1636776_at	0.0638	0.7137	-0.0307	0.7711	0.0636	0.8004	0.1146	0.8485	0.2607	0.2347	0.1461	0.4857	0.0434	0.9530	0.1302	0.5753	0.0867	0.7304
CG5968	CG5968	1636777_at	-0.5300	0.2812	-1.9124	0.0664	-1.6795	0.0020	0.0422	0.9834	1.1556	0.0208	1.1134	0.0152	-0.3834	0.8657	-0.1500	0.9152	0.2334	0.8402
CG13679	CG13679	1636778_at	0.6539	0.1237	0.5413	0.0413	0.4179	0.1844	-0.0475	0.9445	-0.1055	0.6275	-0.0580	0.7910	0.2543	0.8609	0.0335	0.9734	-0.2209	0.7442
---	---	1636779_at	0.1153	0.5648	0.1969	0.3167	0.0771	0.6500	-0.2098	0.5039	-0.1969	0.2371	0.0129	0.9511	-0.0417	0.9630	-0.0257	0.9492	0.0160	0.9655
meso18E	meso18E	1636780_at	-1.1623	0.0329	-0.4499	0.3745	-0.5928	0.1572	-0.0728	0.9200	-0.1237	0.6117	-0.0510	0.8415	-0.0213	0.9946	0.3462	0.6695	0.3676	0.6415
CG2144	CG2144	1636781_at	1.0324	0.0022	0.9171	0.1085	0.7902	0.0007	0.0556	0.9297	0.6518	0.0045	0.5962	0.0040	0.1940	0.8692	0.5198	0.2518	0.3258	0.5050
CG10035	CG10035	1636782_at	0.1454	0.5514	0.1525	0.3879	0.4317	0.0136	0.1100	0.8373	-0.0872	0.7070	-0.1972	0.2780	-0.0117	0.9928	0.0134	0.9819	0.0250	0.9548
nerfin-2	nervous fingers 2	1636783_at	0.2096	0.1674	0.1809	0.2285	0.1341	0.5448	-0.0046	0.9956	-0.0571	0.7951	-0.0525	0.7941	-0.0193	0.9829	-0.0623	0.8166	-0.0430	0.8779
Spn3	serpin 3	1636784_at	0.0696	0.8295	0.0333	0.8678	0.1397	0.4119	0.1833	0.7121	0.0536	0.8552	-0.1296	0.5624	-0.0240	0.9852	-0.0429	0.9274	-0.0189	0.9659
---	---	1636785_at	-0.1665	0.2956	-0.0271	0.8263	0.0291	0.8995	-0.0457	0.9467	-0.1422	0.4834	-0.0965	0.6248	-0.0082	0.9916	-0.0307	0.9111	-0.0225	0.9276
CG13474	CG13474	1636786_at	0.1325	0.4206	-0.0478	0.7873	-0.0592	0.7587	0.1928	0.6313	0.2492	0.2037	0.0564	0.7986	0.0587	0.9441	0.0250	0.9507	-0.0337	0.9231
th	Inhibitors of Apopt	1636787_s_at	-0.3795	0.1936	0.0625	0.8524	0.0356	0.9117	-0.0032	0.9970	-0.1730	0.5062	-0.1698	0.4677	0.0146	0.9950	0.2440	0.6762	0.2294	0.6954
loqs	loquacious	1636788_a_at	-1.4349	0.0011	-0.9836	0.0262	-0.9757	0.0003	0.0108	0.9860	-0.6844	0.0013	-0.6953	0.0007	-0.0258	0.9862	-0.1542	0.6769	-0.1284	0.7389
Tango11	Transport and Gol	1636789_s_at	-0.6651	0.0077	0.2039	0.3942	0.0439	0.7877	-0.2834	0.5280	-0.9904	0.0021	-0.7070	0.0055	-0.0560	0.9515	0.1680	0.5630	0.2240	0.4301
---	---	1636790_at	0.1523	0.3251	0.0041	0.9818	0.1024	0.5358	-0.0049	0.9956	0.0019	0.9945	0.0068	0.9783	0.0532	0.9142	0.0230	0.9341	-0.0301	0.8999
CG6718	CG6718	1636791_s_at	0.6247	0.0263	0.7332	0.0204	1.1078	0.0006	0.1869	0.5220	0.5082	0.0060	0.3214	0.0269	-0.3001	0.7588	0.3820	0.3397	0.6821	0.1289
CG13278	CG13278	1636792_at	0.0360	0.8527	-0.0008	0.9961	0.2961	0.3332	-0.0352	0.9620	0.1807	0.3659	0.2159	0.2176	-0.1921	0.8459	0.0778	0.8987	0.2699	0.5257
Cyp4d2	cytochrome P-450	1636793_at	-0.3176	0.3627	-0.5323	0.2899	-0.4431	0.1674	-0.1674	0.8096	-0.8512	0.0083	-0.6837	0.0135	-0.1738	0.9342	-0.7836	0.2329	-0.6098	0.3807
---	---	1636794_at	0.1269	0.5422	-0.0147	0.9010	0.1398	0.5888	0.0043	0.9956	0.0311	0.9084	0.0268	0.9095	0.0609	0.9646	-0.0151	0.9829	-0.0760	0.8800
Aats-ala-m	mitochondrial alar	1636795_at	-0.1759	0.3259	0.2974	0.0949	0.4120	0.0298	0.0812	0.8942	-0.2520	0.2147	-0.3333	0.0699	-0.0596	0.9514	0.2405	0.3978	0.3001	0.3079
CG15295	CG15295	1636796_at	0.1248	0.5096	-0.0182	0.9428	0.2420	0.1260	0.1977	0.5664	0.1062	0.5798	-0.0915	0.6082	-0.2188	0.7768	-0.1075	0.8008	0.1113	0.7774
Aats-ser	Seryl-tRNA synthe	1636797_at	0.1761	0.3897	0.3235	0.0907	0.5602	0.0194	0.0722	0.9037	-0.1303	0.5285	-0.2025	0.2448	-0.2653	0.6898	-0.0737	0.8402	0.1915	0.4978
CG13482	CG13482	1636798_at	0.0938	0.9492	-1.5990	0.0710	-0.8862	0.1494	1.0534	0.4128	1.6432	0.0289	0.5898	0.3703	-0.0698	0.9923	-0.0489	0.9890	0.0209	0.9945
CG33274	CG33274	1636799_at	0.0533	0.8422	0.1611	0.3522	0.2349	0.1125	0.0088	0.9937	-0.1139	0.6445	-0.1227	0.5751	-0.0824	0.8993	0.0531	0.8769	0.1355	0.5995
---	---	1636800_at	3.3425	0.0006	3.0095	0.0087	3.9393	0.0000	1.4858	0.0221	1.3594	0.0029	-0.1264	0.7349	0.2679	0.8049	0.8943	0.0667	0.6264	0.1911
CG8232 /// DsimCG8232	CG8232	1636801_at	0.2078	0.1577	0.0373	0.7322	-0.0590	0.8008	-0.0715	0.8864	-0.2793	0.0992	-0.2078	0.1692	-0.0032	0.9984	-0.4076	0.1925	-0.4044	0.2249
CG33279	CG33279	1636802_at	0.1329	0.4315	0.0732	0.5262	-0.1755	0.3511	0.0122	0.9883	0.0939	0.6789	0.0817	0.6960	0.0527	0.9447	0.0024	0.9969	-0.0504	0.8694
---	---	1636803_at	0.1460	0.5090	0.3197	0.2386	0.3989	0.0800	0.0403	0.9600	-0.1661	0.4489	-0.2064	0.2789	-0.0112	0.9933	0.0994	0.7967	0.1106	0.7527
CG14629 /// DsimCG14629	CG14629	1636804_at	0.0369	0.9365	-0.3260	0.6888	0.7276	0.0366	0.4006	0.5353	-0.0901	0.8345	-0.4907	0.1028	-0.5820	0.7721	-0.4247	0.6483	0.1573	0.8934
CG8944	CG8944	1636805_a_at	0.4811	0.3950	0.4566	0.2044	-0.0735	0.7329	0.0603	0.9172	0.5992	0.0058	0.5389	0.0056	0.5606	0.7822	0.5539	0.5447	-0.0066	0.9966
Alg10	Alpha 3 glucosyltr	1636806_at	0.1677	0.3442	0.5746	0.0283	0.5129	0.0064	0.0043	0.9956	-0.1502	0.3683	-0.1545	0.2964	0.0153	0.9901	0.1912	0.4856	0.1759	0.5335
Ogg1	Ogg1	1636807_at	0.0392	0.8848	-0.3651	0.0375	-0.3154	0.1582	-0.0642	0.9247	0.6161	0.0089	0.6803	0.0034	-0.1375	0.8744	0.1655	0.6601	0.3030	0.3867
---	---	1636808_at	0.1926	0.4023	0.1981	0.4317	0.0613	0.7646	-0.0997	0.9148	-0.0541	0.8907	0.0456	0.8968	0.1731	0.8246	-0.0104	0.9875	-0.1836	0.5995
Lam	intermediate filam	1636809_at	0.5046	0.3915	-0.1285	0.3624	-0.7423	0.0041	0.2133	0.7850	1.4765	0.0015	1.2632	0.0017	0.8009	0.6496	0.8169	0.2351	0.0160	0.9888
CG11872	CG11872	1636810_at	-0.6335	0.0877	-0.3034	0.3996	-0.5306	0.0167	-0.2975	0.6391	-0.5152	0.0973	-0.2177	0.4588	0.1367	0.9246	-0.0451	0.9538	-0.1818	0.7455
Oatp33Ea	Organic anion trar	1636811_at	-0.3548	0.7994	-0.0328	0.7505	-0.1794	0.3285	0.0165	0.9962	-0.7232	0.5622	-0.7396	0.5076	-0.1675	0.9238	-0.6118	0.2731	-0.4444	0.4572
CG10103	CG10103	1636812_at	-0.1234	0.4950	-0.7426	0.0674	-0.3338	0.3934	0.1827	0.6615	0.4824	0.0239	0.2997	0.0935	-0.2126	0.8972	0.0040	0.9988	0.2166	0.7604
CG3731	CG3731	1636813_s_at	-0.6974	0.0149	0.0652	0.9315	0.0237	0.9373	-0.0165	0.9893	-1.1348	0.0026	-1.1183	0.0016	0.0623	0.9829	-0.3310	0.6458	-0.3932	0.5778
---	---	1636814_at	0.1419	0.5409	-0.0447	0.8542	-0.0539	0.8474	-0.0910	0.9117	0.1196	0.6898	0.2106	0.3845	-0.1895	0.8270	-0.1554	0.7060	0.0341	0.9470
numb	Numb	1636815_a_at	-0.5102	0.0422	-0.2707	0.1666	-0.5646	0.0119	-0.0846	0.9011	0.0135	0.9687	0.0981	0.6697	0.1160	0.8850	0.1632	0.6156	0.0473	0.9099
CG33090	CG33090	1636816_s_at	1.1698	0.0200	1.2278	0.0217	1.7656	0.0004	-0.1315	0.7929	0.3841	0.0629	0.5156	0.0122	-0.6266	0.7070	0.3344	0.6625	0.9610	0.1911
---	---	1636817_s_at	0.2490	0.1751	-0.0021	0.9903	0.0262	0.8917	0.0813	0.8856	0.1258	0.5408	0.0444	0.8407	0.1141	0.8609	-0.0056	0.9925	-0.1197	0.6779
CG13133	CG13133	1636818_at	0.2070	0.3758	0.1526	0.4435	0.0415	0.8508	0.0794	0.9138	-0.0560	0.8521	-0.1355	0.5514	0.1683	0.8504	0.0763	0.8864	-0.0920	0.8461
---	---	1636819_at	0.0120	0.9631	0.0517	0.7042	0.0294	0.8583	0.0289	0.9608	0.0162	0.9438	-0.0127	0.9491	0.0385	0.9577	-0.1100	0.6154	-0.1485	0.4825
spo	spook	1636820_s_at	0.0398	0.9011	-0.0802	0.4737	-0.0913	0.6192	0.1808	0.5516	0.2332	0.1365	0.0524	0.7612	0.1072	0.9388	0.1084	0.8546	0.0013	0.9990
Arpc3A	Arpc3A	1636821_at	-0.0196	0.9150	0.0162	0.8972	0.2574	0.1249	0.0183	0.9808	0.0639	0.7747	0.0457	0.8291	-0.1267	0.8236	0.1476	0.5523	0.2743	0.2638
endos	dendos	1636822_s_at	0.0929	0.8347	0.3903	0.0302	0.2404	0.2200	-0.1536	0.7013	-0.5942	0.0067	-0.4406	0.0154	-0.0355	0.9848	-0.2150	0.6370	-0.1795	0.7027
CG30087	CG30087	1636823_at	-0.1530	0.3946	0.2139	0.2289	-0.0136	0.9527	-0.1985	0.5757	-0.3329	0.0702	-0.1343	0.4319	0.0704	0.9421	0.1494	0.6473	0.0790	0.8355
---	---	1636824_at	0.1498	0.4173	-0.0977	0.7126	-0.1710	0.4631	-0.0595	0.9343	0.2225	0.2981	0.2819	0.1350	-0.0178	0.9862	0.0150	0.9704	0.0328	0.9220
CG3492	CG3492	1636825_at	0.0746	0.7304	0.0986	0.4362	0.1760	0.4357	0.0292	0.9704	0.0373	0.8906	0.0081	0.9738	0.0950	0.9309	0.2545	0.4843	0.1595	0.6857
CG14072	CG14072	1636826_at	-0.0981	0.7640	0.0854	0.5117	0.0218	0.9262	-0.2916	0.3863	-0.2759	0.1460	0.0157	0.9483	0.0629	0.9260	0.1067	0.6675	0.0437	0.8880
CG32614	CG32614	1636827_at	0.1509	0.4494	-0.0110	0.9337	0.1779	0.3607	0.0441	0.9451	0.1060	0.5938	0.0619	0.7563	-0.0201	0.9831	-0.0336	0.9229	-0.0135	0.9681
CG15732	CG15732	1636828_at	-0.1938	0.2698	-0.0742	0.5755	-0.0752	0.7355	0.0001	0.9999	-0.1261	0.5038	-0.1262	0.4551	0.0563	0.9644	-0.1239	0.7618	-0.1802	0.6228
CG31902	CG31902	1636829_at	0.2903	0.2830	-0.2535	0.3599	0.2306	0.1394	0.2464	0.5932	0.3500	0.1336	0.1037	0.6703	-0.1157	0.8940	-0.1741	0.6154	-0.0584	0.8949
---	---	1636830_s_at	0.1595	0.3591	0.3392	0.1297	0.4101	0.1717	-0.0128	0.9860	-0.3298	0.0625	-0.3169	0.0476	0.0619	0.9238	-0.0513	0.8642	-0.1132	0.6238
brm	Complementation	1636831_s_at	-0.0383	0.9181	0.0423	0.8777	0.0203	0.9512	0.0343	0.9777	0.2023	0.5532	0.1679	0.5970	0.1056	0.9449	0.4051	0.3738	0.2995	0.5409
CG40188	CG40188	1636832_a_at	0.1500	0.3251	-0.0608	0.5653	-0.1140	0.5339	0.0653	0.9254	0.2193	0.3002	0.1540	0.4345	0.0268	0.9717	-0.0028	0.9942	-0.0296	0.9096
CG17364	CG17364	1636833_at	0.0182	0.9117	-0.1955	0.5692	0.1942	0.3393	0.2549	0.7134	0.1985	0.5600	-0.0564	0.8802	-0.2062	0.7644	-0.0770	0.8439	0.1292	0.6857
ms(3)K81	paternal-haploid	1636834_at	0.0850	0.5671	0.0087	0.9441	-0.2301	0.2650	-0.1092	0.8156	-0.0277	0.9144	0.0816	0.6702	0.0178	0.9846	-0.1156	0.5962	-0.1334	0.5363
CG16700	CG16700	1636835_at	-2.1828	0.0055	-2.7082	0.0154	-2.9953	0.0000	-0.1246	0.8736	-0.5850	0.0382	-0.4604	0.0621	0.2123	0.9428	-1.0251	0.2430	-1.2374	0.1990
CG8568	CG8568	1636836_at	-0.0127	0.9378	0.0349	0.8237	-0.0684	0.6830	0.0307	0.9705	-0.0314	0.9167	-0.0620	0.7962	0.0454	0.9405	-0.0266	0.9282	-0.0720	0.7393
CG32177	CG32177	1636837_at	0.3567	0.0571	0.5283	0.0286	0.7722	0.0048	0.0914	0.8189	-0.0581	0.7552	-0.1495	0.2906	-0.1261	0.8940	-0.0214	0.9707	0.1046	0.8082
n-syb	N-synaptobrevin	1636838_a_at	-1.0964	0.0195	-1.2103	0.0032	-1.0885	0.0064	0.1521	0.9117	0.2200	0.6551	0.0678	0.8974	-0.1809	0.8795	-0.0709	0.9208	0.1100	0.8524
Or71a	Odorant receptor	1636839_a_at	-0.0308	0.9267	-0.0145	0.9480	0.1636	0.3525	0.0256	0.9683	0.0347	0.8765	0.0091	0.9660	-0.0362	0.9779	-0.0820	0.8471	-0.0458	0.9161

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CBP	Ca2+ - binding pr	1636840_at	3.7307	0.0023	3.5720	0.0045	3.0827	0.0000	0.2342	0.9093	0.6497	0.3272	0.4155	0.5103	0.6966	0.5126	0.4854	0.3259	-0.2111	0.7131
CG12130 /// DereCG12130	CG12130 /// GA1	1636841_at	1.1057	0.0289	0.3005	0.2597	0.3577	0.0960	0.0559	0.9441	0.5120	0.0317	0.4561	0.0323	-0.0951	0.9514	0.0031	0.9988	0.0982	0.8758
CG8460	CG8460	1636842_at	0.6023	0.0072	0.0572	0.8605	0.2008	0.3482	0.1651	0.6823	0.6394	0.0053	0.4743	0.0124	0.0497	0.9643	0.1470	0.6524	0.0973	0.7848
CG10512	CG10512	1636843_a_at	-0.0841	0.7680	0.8788	0.1564	0.6508	0.0601	-0.0387	0.9715	-0.7790	0.0117	-0.7403	0.0089	0.2063	0.9142	0.1361	0.8884	-0.0702	0.9402
fu2	fu2	1636844_at	0.4640	0.0580	0.0744	0.6691	-0.1030	0.5088	-0.0093	0.9922	0.5832	0.0110	0.5925	0.0061	0.1493	0.8062	-0.0055	0.9925	-0.1548	0.5657
CG8885	CG8885	1636845_at	-0.6126	0.0360	-0.2697	0.2135	0.0180	0.9483	-0.0411	0.9345	-0.2922	0.0522	-0.2511	0.0605	-0.3825	0.7498	-0.0268	0.9762	0.3556	0.5032
---	---	1636846_at	-0.1666	0.6946	-0.2245	0.8401	-0.4471	0.1333	-0.1291	0.9098	0.6875	0.0615	0.8166	0.0199	-0.0285	0.9950	0.2561	0.8624	0.2846	0.8307
Mkrn1	Makorin 1	1636847_s_at	-0.4806	0.0702	-0.0924	0.7947	-0.6783	0.0236	-0.2977	0.5109	-0.3692	0.1214	-0.0715	0.7894	0.2818	0.7780	0.0463	0.9477	-0.2355	0.6152
CG6024	CG6024	1636848_at	-0.0889	0.5207	-0.8084	0.0259	-0.8119	0.0064	0.1770	0.8437	0.7832	0.0261	0.6062	0.0467	0.1051	0.8465	0.0036	0.9941	-0.1015	0.6836
CG8858	CG8858	1636849_at	0.2563	0.3445	0.2942	0.4100	0.8396	0.0044	0.1620	0.7971	0.2558	0.3239	0.0939	0.7322	-0.2502	0.8157	0.4266	0.3255	0.6769	0.1537
CG40338	CG40338	1636850_at	0.1958	0.4120	0.0957	0.5861	0.2775	0.1554	0.2085	0.5895	0.1290	0.5392	-0.0795	0.7002	-0.0749	0.9365	0.0425	0.9277	0.1174	0.7305
CG1722	CG1722	1636851_at	-0.7275	0.1422	-0.4579	0.0949	-0.6561	0.0055	0.0357	0.9768	-0.0540	0.8957	-0.0897	0.7908	-0.1990	0.8611	-0.2085	0.6862	-0.0095	0.9903
CG8939	CG8939	1636852_at	0.5929	0.0819	0.4329	0.4451	0.9025	0.0044	0.3816	0.5758	0.4616	0.1848	0.0799	0.8432	-0.0726	0.9779	0.3531	0.5972	0.4257	0.5156
Cpr47Ef	CG13214	1636853_at	0.2680	0.3322	-0.0187	0.8973	-0.2446	0.4148	-0.0842	0.8897	0.1331	0.5423	0.2173	0.2325	-0.2332	0.7893	-0.1256	0.7856	0.1077	0.8161
---	---	1636854_at	0.2277	0.2882	0.0272	0.8062	0.0839	0.5877	0.0656	0.8776	0.1675	0.2531	0.1020	0.4615	0.0510	0.9653	-0.0199	0.9693	-0.0709	0.8641
---	---	1636855_at	0.1275	0.5011	-0.1067	0.4436	-0.0165	0.9455	0.2293	0.5808	0.2610	0.2170	0.0317	0.9007	-0.0788	0.9201	-0.1577	0.5749	-0.0789	0.8102
Vhl	Von Hippel Linda	1636856_at	-0.3150	0.1688	-0.0478	0.8312	-0.0101	0.9564	-0.0192	0.9838	-0.2492	0.2631	-0.2300	0.2480	-0.0212	0.9851	0.0834	0.7994	0.1045	0.7221
CG15879	CG15879	1636857_at	0.4131	0.3783	0.0689	0.7593	0.0138	0.9742	-0.0846	0.9172	0.1362	0.6267	0.2207	0.3383	-0.1737	0.9092	-0.4460	0.4037	-0.2723	0.6389
cm	carmine	1636858_at	0.7305	0.0217	1.4317	0.0071	1.5970	0.0000	0.2892	0.5515	-0.2490	0.3278	-0.5382	0.0246	0.0986	0.9246	0.5797	0.1047	0.4811	0.1919
---	---	1636859_at	0.2183	0.3249	0.1948	0.3353	0.1786	0.3336	0.0494	0.9325	-0.0558	0.7927	-0.1051	0.5340	-0.0516	0.9543	-0.0606	0.8721	-0.0090	0.9841
---	---	1636860_at	0.0054	0.9762	-0.1140	0.5425	0.0584	0.7646	0.1471	0.6999	0.0436	0.8479	-0.1035	0.5473	-0.0084	0.9939	-0.0197	0.9601	-0.0113	0.9763
CG8617	CG8617	1636861_at	-0.1165	0.5656	-0.0471	0.7331	0.4634	0.0180	0.2360	0.4220	0.0451	0.8254	-0.1909	0.1881	-0.2309	0.6955	0.1525	0.5501	0.3834	0.1433
CG11322	CG11322	1636862_at	0.2086	0.1519	0.1637	0.3512	0.1512	0.3411	-0.1948	0.5839	-0.0587	0.7913	0.1361	0.4224	0.0320	0.9739	0.1128	0.6711	0.0808	0.7754
---	---	1636863_at	0.1792	0.3965	-0.0738	0.6469	-0.0132	0.9477	-0.0765	0.9015	0.1802	0.3794	0.2567	0.1509	-0.1552	0.7768	0.0079	0.9870	0.1630	0.5064
corto	Centrosomal and	1636864_at	-0.4732	0.5460	0.0208	0.9828	-1.6486	0.0095	-0.3256	0.2596	-0.0263	0.9121	0.2993	0.0548	1.3828	0.7041	0.5550	0.7596	-0.8278	0.6129
Os-C	Os-C	1636865_at	-0.0941	0.7043	0.0019	0.9954	-0.3297	0.1653	-0.2180	0.5539	-0.0380	0.8810	0.1799	0.2968	0.0347	0.9860	-0.0263	0.9719	-0.0610	0.9198
sec15	sec15	1636866_at	0.0306	0.8823	0.3465	0.3387	0.5264	0.0055	0.1228	0.8053	0.0678	0.7776	-0.0550	0.8053	-0.0546	0.9691	0.3254	0.3698	0.3800	0.3141
CG7987	CG7987	1636867_s_at	-0.4342	0.0832	-0.1186	0.4223	-0.3332	0.0387	0.0582	0.9061	-0.1828	0.2523	-0.2411	0.0905	0.2441	0.7536	0.0963	0.8248	-0.1478	0.6870
CG14939	CG14939	1636868_at	-0.4207	0.2944	-0.0405	0.8235	-0.4780	0.0096	-0.0579	0.9255	0.1873	0.3230	0.2452	0.1409	0.3042	0.8062	0.5893	0.2385	0.2852	0.6086
---	---	1636869_at	0.5622	0.1205	-0.1387	0.5811	-0.6452	0.0500	-0.1248	0.8857	0.8750	0.0098	0.9998	0.0032	0.3098	0.8157	0.1136	0.8864	-0.1962	0.7568
CG9793	CG9793	1636870_at	0.0825	0.7124	-0.1947	0.3063	-0.5638	0.0336	-0.5034	0.3743	0.3226	0.3193	0.8259	0.0112	-0.0287	0.9845	0.0831	0.8546	0.1118	0.7733
CG12943	CG12943	1636871_at	0.3105	0.0707	0.0897	0.5665	0.2354	0.2471	0.1257	0.7923	0.1537	0.4460	0.0280	0.9048	-0.0636	0.9168	-0.1764	0.3982	-0.1128	0.6195
Aats-phe	Phenylalanyl-tRN	1636872_at	0.0637	0.7030	0.2231	0.3279	0.1763	0.2396	-0.0245	0.9647	-0.1163	0.4435	-0.0918	0.5170	0.0875	0.9088	0.0477	0.9084	-0.0399	0.9157
CG9068	CG9068	1636873_at	-0.0089	0.9612	0.1645	0.4099	0.0623	0.7628	-0.2515	0.5504	-0.2355	0.2827	0.0160	0.9538	-0.0247	0.9705	0.0378	0.8765	0.0625	0.7495
---	---	1636874_at	-0.0829	0.6476	-0.1198	0.4475	0.1024	0.6029	0.2687	0.5610	0.1450	0.5711	-0.1237	0.6051	0.0039	0.9970	0.0243	0.9521	0.0205	0.9545
skpE	skpE	1636875_at	0.1207	0.5463	0.0369	0.7299	-0.0773	0.7486	0.0340	0.9683	0.0767	0.7761	0.0427	0.8713	0.0593	0.9296	0.0466	0.8807	-0.0127	0.9695
KP78a	KP78a	1636876_at	0.1876	0.4259	0.1062	0.5539	-0.0083	0.9729	-0.0279	0.9836	-0.1104	0.7667	-0.0825	0.8135	0.0786	0.8791	0.0156	0.9621	-0.0631	0.7918
CG1134	CG1134	1636877_at	-0.1030	0.6939	0.5863	0.0950	0.6697	0.0323	-0.1045	0.9013	-0.7262	0.0160	-0.6217	0.0192	-0.0130	0.9923	0.1833	0.5844	0.1963	0.5581
---	---	1636878_at	0.2221	0.2373	0.0718	0.7264	0.0494	0.7978	-0.0128	0.9852	0.0128	0.9552	0.0256	0.8886	0.0850	0.9357	-0.0497	0.9248	-0.1347	0.7228
CG14356	CG14356	1636879_at	0.2488	0.7138	0.7880	0.2828	-0.0692	0.8337	-0.0644	0.9584	-0.1406	0.7075	-0.0761	0.8388	0.7237	0.7644	0.2953	0.8284	-0.4284	0.7133
cpx	complexin	1636880_s_at	-2.1421	0.0028	-1.7153	0.0374	-3.2883	0.0001	-1.1178	0.0221	-1.2662	0.0012	-0.1484	0.5676	0.1374	0.9717	-0.8468	0.3802	-0.9842	0.3271
---	---	1636881_at	0.2089	0.3005	0.1300	0.4509	0.1899	0.3776	-0.0293	0.9759	-0.3976	0.0963	-0.3683	0.0853	-0.0690	0.9407	-0.1315	0.6803	-0.0626	0.8699
---	---	1636882_s_at	-0.0210	0.9226	-0.9540	0.1426	-1.0489	0.0305	-0.0447	0.9857	0.7779	0.1671	0.8225	0.1036	-0.0941	0.8882	0.0238	0.9531	0.1180	0.6668
HmgD	High mobility grou	1636883_s_at	-0.1277	0.8252	-0.4498	0.2482	-0.3501	0.0399	-0.3765	0.6597	-0.2580	0.5602	0.1185	0.7975	-0.4079	0.7324	-0.4890	0.3259	-0.0811	0.9086
CG4658	CG4658	1636884_s_at	-0.6966	0.0063	-0.1435	0.4422	-0.0726	0.6605	-0.1929	0.5140	-0.9451	0.0005	-0.7522	0.0007	-0.2074	0.7506	-0.3329	0.2174	-0.1254	0.6848
CG17560	CG17560	1636885_at	0.6870	0.6868	-2.8399	0.4290	1.6649	0.1018	2.8939	0.2420	1.6806	0.2518	-1.2133	0.3688	-1.4866	0.8692	-1.6764	0.6713	-0.1898	0.9727
spn-E	Spindle-E (homek	1636886_at	0.4939	0.2514	-0.1260	0.8608	-1.2127	0.0171	-1.0031	0.0104	1.0449	0.0008	2.0480	0.0000	-0.1573	0.9721	0.2176	0.8997	0.3748	0.7815
CG11247	CG11247	1636887_s_at	-0.1556	0.3735	-0.1140	0.7935	-0.7852	0.0013	-0.2612	0.5000	-0.0727	0.7670	0.1885	0.3111	0.1530	0.8611	-0.1404	0.7346	-0.2933	0.4167
Aph-4	Alkaline phosphat	1636888_a_at	-1.5391	0.0049	-0.0718	0.9010	-2.2376	0.0009	-1.5841	0.0970	-2.5140	0.0016	-0.9299	0.0663	0.5713	0.7220	-1.1370	0.1047	-1.7082	0.0500
---	---	1636889_at	-0.2226	0.2140	0.0219	0.8606	-0.0348	0.8429	-0.0018	0.9978	-0.0269	0.9084	-0.0251	0.9011	0.0115	0.9898	0.0588	0.8185	0.0473	0.8524
CG31029	CG31029	1636890_at	-0.2362	0.3815	-0.1049	0.6048	-0.1097	0.5966	0.0462	0.9558	-0.3178	0.1556	-0.3640	0.0722	-0.2164	0.7393	-0.2663	0.3236	-0.0499	0.8949
---	---	1636891_at	0.2216	0.3670	0.1363	0.4245	0.1105	0.4985	-0.1102	0.8551	-0.1011	0.6856	0.0091	0.9733	-0.1068	0.8494	-0.1132	0.6587	-0.0065	0.9861
upd3	unpaired 3	1636892_at	-0.0397	0.8186	0.3279	0.2911	0.4425	0.1240	-0.0452	0.9603	-0.1518	0.5603	-0.1066	0.6720	-0.2201	0.8122	0.0640	0.9144	0.2841	0.4741
CG32798	CG32798	1636893_at	-0.0823	0.6818	0.0316	0.7943	0.2129	0.2740	0.1819	0.6818	-0.0109	0.9732	-0.1928	0.3032	0.0934	0.9246	0.1345	0.7215	0.0411	0.9283
---	---	1636894_s_at	0.2149	0.3802	-0.1013	0.7168	-0.0321	0.9108	-0.0915	0.9149	0.3590	0.1778	0.4505	0.0635	-0.1967	0.8215	0.0099	0.9903	0.2067	0.5936
Or9a	Odorant receptor	1636895_at	-0.0126	0.9548	-0.0229	0.9024	0.3170	0.1646	0.1095	0.8507	-0.1413	0.5310	-0.2508	0.1837	-0.1770	0.7707	-0.0547	0.8845	0.1222	0.6653
CG12576	CG12576	1636896_a_at	-0.2247	0.3094	0.0361	0.7693	0.0473	0.8142	-0.5053	0.1321	-1.0687	0.0008	-0.5635	0.0078	-0.4821	0.3828	-0.9133	0.0246	-0.4312	0.1829
CG33003	CG33003	1636897_at	0.1025	0.7693	0.1580	0.3021	0.3897	0.0283	0.0554	0.9393	-0.0600	0.8210	-0.1154	0.5864	-0.0863	0.9092	0.2009	0.4539	0.2872	0.2964
CG14448	CG14448	1636898_at	0.2236	0.3251	0.0000	1.0000	-0.0727	0.6482	-0.0218	0.9755	0.0217	0.9292	0.0436	0.8257	-0.0651	0.9342	-0.1544	0.5613	-0.0892	0.7608
CG1623	CG1623	1636899_s_at	-0.1979	0.4894	0.4368	0.3313	0.4530	0.1158	-0.3335	0.5445	-0.8604	0.0092	-0.5269	0.0446	-0.3029	0.8243	-0.0933	0.9157	0.2095	0.7492
CG5390 /// DyakCG5390	CG5390	1636900_at	-0.3848	0.1023	-1.7977	0.0252	-1.4999	0.0001	0.1849	0.5929	-0.0993	0.6054	-0.2842	0.0718	-0.0399	0.9853	-1.4049	0.0276	-1.3650	0.0410
---	---	1636901_at	0.0734	0.6640	0.0003	1.0000	0.3403	0.0400	0.0618	0.9116	-0.0898	0.6500	-0.1516	0.3474	-0.1066	0.8222	-0.0323	0.9157	0.0743	0.7442
CG1840	CG1840	1636902_at	0.6152	0.0524	0.5156	0.0781	0.7190	0.0046	-0.1240	0.6869	0.9135	0.0004	1.0375	0.0002	-0.1713	0.8608	0.8867	0.0457	1.0581	0.0422
---	---	1636903_at	-0.0203	0.9288	0.1264	0.5199	0.0798	0.7133	0.1298	0.7929	0.1199	0.5846	-0.0100	0.9688	0.1761	0.7726	0.3108	0.2226	0.1347	0.6366
CG40235	CG40235	1636904_at	0.0795	0.6775	0.0123	0.9444	0.0516	0.7865	0.0318	0.9647	-0.0175	0.9509	-0.0493	0.8185	0.0137	0.9898	-0.0049	0.9925	-0.0186	0.9548
lea	Robo2	1636905_at	0.7339	0.4266	0.4248	0.6219	1.5037	0.0007	0.1911	0.9330	0.3599	0.6307	0.1688	0.8253	-0.6928	0.7781	0.2107	0.8948	0.9035	0.4067
Sans	Sans	1636906_s_at	0.5054	0.1089	-0.7988	0.0386	0.1467	0.5774	0.4886	0.3420	0.7154	0.0238	0.2268	0.4123	-0.4984	0.6717	-0.6564	0.1537	-0.1580	0.7774
CG32982	CG32982	1636907_at	0.2835	0.1443	0.0428	0.8874	-0.1492	0.5035	0.0720	0.9098	0.3925	0.0566	0.3205	0.0777	0.0560	0.9677	-0.1456	0.7317	-0.2016	0.6086
---	---	1636908_s_at	0.2066	0.4865	-0.0188	0.9362	0.3001	0.2434	0.2429	0.6354	0.1956	0.4553	-0.0473	0.8738	-0.1248	0.8940	-0.0425	0.9402	0.0823	0.8555

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG4078	CG4078	1636909_at	0.5429	0.1112	0.0365	0.7167	0.4582	0.0367	0.0338	0.9539	0.2166	0.1743	0.1828	0.2013	-0.2670	0.7726	-0.2107	0.6256	0.0562	0.9192
CG13704	CG13704	1636910_at	2.1016	0.0073	1.5045	0.0843	2.4821	0.0002	0.7458	0.4979	0.2412	0.7224	-0.5046	0.3452	-0.1694	0.9491	-0.4303	0.6117	-0.2608	0.7808
jim	jim	1636911_at	-0.4255	0.4993	-0.1198	0.8397	-0.3310	0.0857	-0.0983	0.9423	-0.1043	0.8325	-0.0060	0.9902	0.0923	0.9816	0.1086	0.9428	0.0163	0.9924
CG7745	CG7745	1636912_at	0.3334	0.1132	0.5474	0.1004	0.3825	0.0510	0.0513	0.9311	0.2040	0.2395	0.1527	0.3353	0.2128	0.7823	0.2582	0.4380	0.0455	0.9211
CG13120	CG13120	1636913_at	0.0285	0.8862	-0.0127	0.9432	0.0468	0.8101	0.0323	0.9733	-0.0119	0.9743	-0.0443	0.8757	0.0499	0.9644	-0.0736	0.8611	-0.1235	0.7168
Cdk5alpha	Cdk5 activator-like	1636914_at	-0.5561	0.2350	0.0692	0.6423	0.1661	0.4917	-0.2603	0.8532	-0.8704	0.0881	-0.6101	0.1789	-0.1047	0.8202	-0.1153	0.5591	-0.0105	0.9709
CG11380	CG11380	1636915_at	0.1775	0.4204	-0.1015	0.4617	0.0425	0.8453	0.0176	0.9803	0.0493	0.8202	0.0317	0.8776	-0.0557	0.9515	-0.1072	0.7406	-0.0516	0.8918
disco-r	disco-related	1636916_at	0.1128	0.5139	-0.4309	0.0564	0.0059	0.9806	0.2732	0.4013	0.3802	0.0430	0.1070	0.5496	-0.0328	0.9816	-0.1525	0.6660	-0.1197	0.7456
CG41056	CG41056	1636917_at	0.3582	0.0324	0.0434	0.6805	-0.0256	0.9061	-0.0641	0.9030	0.1104	0.5480	0.1745	0.2561	0.0000	0.9999	-0.1855	0.4448	-0.1856	0.4597
CG31169	CG31169	1636918_a_at	-0.0693	0.7724	-0.6071	0.0186	-0.5476	0.0667	0.0648	0.9300	0.4670	0.0374	0.4022	0.0435	-0.1383	0.9171	-0.0578	0.9381	0.0805	0.8986
mago	Mago nashi	1636919_at	0.6341	0.0277	0.6666	0.2098	0.2421	0.1609	-0.2019	0.7590	0.2113	0.4749	0.4132	0.1016	0.2482	0.8222	0.2479	0.6144	-0.0003	0.9997
CG12016	CG12016	1636920_s_at	0.3366	0.1488	-0.1198	0.6461	0.0373	0.8882	-0.2953	0.4690	-0.5126	0.0275	-0.2174	0.2738	-0.4016	0.7095	-1.0739	0.0391	-0.6723	0.1605
l(2)06496	dynactin, p24 subu	1636921_at	-0.9035	0.0018	-1.0002	0.0338	-1.1378	0.0001	0.1049	0.7556	0.2287	0.1146	0.1238	0.3539	0.0448	0.9514	0.0659	0.8180	0.0210	0.9457
CG12267	CG12267	1636922_at	0.0661	0.6852	-0.0014	1.0000	0.3636	0.1784	0.1826	0.5898	0.1124	0.5423	-0.0701	0.6981	-0.2065	0.8541	0.0824	0.9075	0.2889	0.5564
CG11198	Acetyl CoA carbox	1636923_a_at	0.9152	0.0493	1.0083	0.1578	1.8808	0.0001	0.2507	0.5507	-0.5810	0.0155	-0.8318	0.0019	-0.5665	0.7746	-0.4782	0.6011	0.0883	0.9414
---	---	1636924_at	-0.1576	0.3145	0.1335	0.5373	0.2397	0.2101	-0.1145	0.7658	-0.2760	0.0928	-0.1615	0.2755	-0.2765	0.7307	-0.0938	0.8393	0.1827	0.6225
CG31477	CG31477	1636925_at	-0.4210	0.0560	-0.3027	0.0351	-0.5178	0.0607	0.1352	0.8903	0.0309	0.9504	-0.1044	0.7742	0.0724	0.9309	-0.0297	0.9462	-0.1021	0.7442
cact	cactus	1636926_s_at	-0.2947	0.1112	-0.1857	0.3307	-0.3528	0.2169	0.0302	0.9677	0.1214	0.5671	0.0912	0.6531	0.1099	0.9291	0.2243	0.5972	0.1145	0.8178
CG30379	CG30379	1636927_at	-0.3036	0.3406	-0.2409	0.2631	-0.3863	0.0658	-0.2571	0.5492	-0.1112	0.6569	0.1459	0.4923	0.1208	0.9093	0.0296	0.9621	-0.0912	0.8514
CG1486	CG1486	1636928_s_at	0.1004	0.8652	0.9335	0.0917	0.4069	0.0623	-0.1139	0.8196	-0.1200	0.5704	-0.0061	0.9802	0.3277	0.8767	0.7135	0.3765	0.3858	0.6668
ave	aveugle	1636929_at	-0.3146	0.2807	0.0602	0.7074	-0.2540	0.1458	-0.1679	0.7228	-0.1576	0.4801	0.0103	0.9692	0.1464	0.8431	0.2480	0.4100	0.1016	0.7767
CG9121	CG9121	1636930_at	-0.1301	0.6756	0.1998	0.3589	-0.0308	0.9166	-0.1977	0.7149	-0.3491	0.1560	-0.1514	0.5300	-0.0966	0.8963	0.0512	0.9037	0.1478	0.6196
br	broad complex	1636931_at	0.0353	0.9003	0.0410	0.7016	0.0886	0.7004	-0.0856	0.8791	-0.1961	0.3117	-0.1105	0.5602	0.0044	0.9964	-0.0153	0.9701	-0.0197	0.9548
CG9886	CG9886	1636932_at	1.4563	0.0022	1.9759	0.0240	2.0553	0.0001	0.2476	0.7349	-0.9345	0.0113	-1.1821	0.0024	0.1866	0.9057	-0.3074	0.6117	-0.4940	0.3921
CG13562	CG13562	1636933_at	-0.6754	0.0264	-0.2164	0.3963	-0.3492	0.0430	-0.0918	0.9186	-0.6031	0.0362	-0.5113	0.0446	0.1133	0.8846	-0.1310	0.6949	-0.2444	0.4175
zormin	D-Titin	1636934_at	-0.2917	0.6335	1.1444	0.0451	-0.5661	0.0882	-0.0784	0.9619	-0.4917	0.2591	-0.4132	0.2933	1.5086	0.2926	1.0070	0.1718	-0.5016	0.5299
CG2219	CG2219	1636935_at	0.0830	0.7117	-0.3646	0.0909	-0.7709	0.0030	-0.2053	0.5046	0.2497	0.1247	0.4550	0.0079	0.1659	0.8404	-0.1531	0.6876	-0.3190	0.3622
CG15778	CG15778	1636936_at	0.1269	0.4712	0.2030	0.3609	0.1209	0.5849	0.1247	0.8550	0.0606	0.8495	-0.0641	0.8185	0.2265	0.7768	0.2364	0.5070	0.0099	0.9855
---	---	1636937_at	0.1905	0.1951	0.1215	0.5469	-0.1266	0.5140	-0.0949	0.8303	0.0483	0.8188	0.1432	0.3579	0.2780	0.6903	-0.0090	0.9877	-0.2870	0.3293
---	---	1636938_at	0.1031	0.4706	-0.1032	0.3583	0.0000	0.9998	0.1765	0.5932	0.1570	0.3583	-0.0195	0.9245	0.0112	0.9913	-0.1081	0.6332	-0.1193	0.5969
Tango6	Transport and Gol	1636939_at	0.3130	0.1525	-0.1008	0.7696	-0.0088	0.9692	-0.1353	0.7205	0.2680	0.1159	0.4033	0.0165	-0.2706	0.7720	-0.2802	0.4934	-0.0096	0.9876
---	---	1636940_x_at	-0.1562	0.3675	-0.0911	0.4742	-0.0071	0.9711	0.1394	0.7000	0.1734	0.3015	0.0340	0.8605	-0.0365	0.9618	0.1011	0.6662	0.1376	0.5393
CG9007 /// DmirCG9007	CG9007	1636941_at	0.5885	0.1668	0.5504	0.1026	0.0494	0.8643	-0.2571	0.7930	0.0970	0.8520	0.3541	0.3332	0.3552	0.7464	0.1404	0.8138	-0.2148	0.6677
CG8498 /// DyakCG8498	CG8498	1636942_at	-0.0556	0.8657	0.1405	0.7054	-0.1193	0.5692	-0.0368	0.9756	-0.0189	0.9675	0.0180	0.9611	0.1581	0.8479	0.1136	0.7914	-0.0445	0.9248
Spn5	serpin 5	1636943_s_at	-1.0552	0.0011	-1.3579	0.0695	-1.5070	0.0001	-0.0745	0.8578	-0.0892	0.5832	-0.0147	0.9372	-0.0380	0.9898	-0.3590	0.5855	-0.3210	0.6317
Edg84A	ecdysone-depend	1636944_at	0.1240	0.4492	0.1325	0.3414	0.0856	0.7189	0.0253	0.9803	-0.1198	0.6728	-0.1451	0.5535	0.1992	0.7046	0.0228	0.9492	-0.1764	0.4322
PpV	Protein phosphata	1636945_at	-0.0919	0.6258	0.7381	0.0441	0.4101	0.0578	-0.1463	0.7104	-0.2477	0.1645	-0.1013	0.5686	0.2112	0.8114	0.4816	0.1747	0.2704	0.4755
CG6687	CG6687	1636946_at	2.9827	0.0033	-0.0150	0.9907	2.2624	0.0024	1.7498	0.1840	2.4418	0.0085	0.6921	0.3438	-0.5714	0.8270	-0.6810	0.5471	-0.1096	0.9431
CG9598	CG9598	1636947_at	2.6130	0.0006	1.4183	0.0026	1.9716	0.0001	0.3665	0.5741	1.0846	0.0066	0.7181	0.0245	-0.1890	0.7823	-0.0624	0.8843	0.1266	0.7026
Aats-gly	Glycyl-tRNA synth	1636948_a_at	1.0397	0.0013	1.4962	0.0507	1.5923	0.0000	0.4746	0.1194	0.5317	0.0110	0.0571	0.7802	0.2915	0.7960	0.8428	0.0871	0.5513	0.2614
CG7843	CG7843	1636949_s_at	0.5112	0.0592	0.4931	0.0695	0.2067	0.2421	-0.1553	0.7149	0.4695	0.0237	0.6248	0.0040	0.2041	0.8086	0.5197	0.1374	0.3156	0.3803
CG11077	CG11077	1636950_at	-0.2703	0.4762	-1.5918	0.0356	-1.1724	0.0065	0.1499	0.8605	0.1267	0.7223	-0.0232	0.9523	-0.3395	0.8331	-1.0247	0.1297	-0.6853	0.3205
CG2857	CG2857	1636951_at	0.1695	0.3497	0.0376	0.7990	0.0619	0.7386	-0.0242	0.9777	0.0377	0.9024	0.0619	0.8047	-0.0543	0.9515	-0.0837	0.8101	-0.0294	0.9375
CG9308	CG9308	1636952_at	0.1350	0.4491	-0.0438	0.6852	-0.0097	0.9617	0.0448	0.9532	0.1338	0.5543	0.0890	0.6865	0.0699	0.9142	-0.0813	0.7669	-0.1512	0.5243
Lmpt	Limpet	1636953_a_at	-1.1005	0.0104	-0.4497	0.3662	-1.5200	0.0006	-0.6682	0.1316	-1.0512	0.0028	-0.3830	0.1064	0.4331	0.7893	-0.3320	0.6605	-0.7651	0.2830
NetB	NETRIN	1636954_at	-0.7743	0.1593	-3.4066	0.0078	-2.8003	0.0020	0.7046	0.6634	2.4766	0.0071	1.7721	0.0187	0.1621	0.9342	-0.1082	0.9111	-0.2702	0.7026
---	---	1636955_at	0.2844	0.1752	0.1722	0.6478	0.2849	0.2523	0.0626	0.9191	0.0920	0.6692	0.0294	0.8974	-0.1066	0.9320	0.0000	1.0000	0.1066	0.8348
Hsc70-1	Heat shock protein	1636956_at	-2.2223	0.0022	-1.0409	0.0618	-1.9086	0.0005	-0.4940	0.1787	-1.4264	0.0004	-0.9324	0.0013	-0.0462	0.9911	-0.5817	0.4985	-0.5356	0.5448
CG34012	CG34012	1636957_s_at	-0.1540	0.4163	-0.2662	0.0536	-0.6750	0.0065	-0.2380	0.5038	0.0898	0.6765	0.3278	0.0562	0.1814	0.7755	0.1711	0.5523	-0.0103	0.9831
l(2)k01209	lethal (2) k01209	1636958_s_at	0.2131	0.4229	-0.4846	0.2855	-0.5105	0.0358	-0.0512	0.9376	0.6523	0.0053	0.7035	0.0022	-0.0199	0.9939	0.0587	0.9505	0.0786	0.9231
CG12587	CG12587	1636959_at	0.0751	0.5914	0.0464	0.6636	0.0120	0.9470	0.0183	0.9745	0.0349	0.8487	0.0166	0.9235	0.0471	0.9306	-0.0453	0.8492	-0.0924	0.6228
micr	midlife-crisis	1636960_a_at	-0.3218	0.2849	-0.4920	0.4808	-0.0535	0.7762	-0.0131	0.9883	-0.2791	0.1775	-0.2661	0.1516	-0.5279	0.7478	-0.4457	0.5398	0.0822	0.9330
CG9027	CG9027	1636961_a_at	-0.0686	0.8524	0.3450	0.5254	0.9221	0.0023	0.0936	0.8738	-0.3471	0.0924	-0.4406	0.0248	-0.4735	0.7726	0.1682	0.8664	0.6417	0.3798
CG8349	CG8349	1636962_at	-0.0246	0.8852	0.0295	0.8152	0.0302	0.9074	0.0374	0.9697	-0.1225	0.6717	-0.1599	0.5155	0.0625	0.9177	-0.0768	0.7595	-0.1393	0.5270
CG12111 /// DsimCG12111	CG12111	1636963_at	0.2492	0.2150	0.0949	0.5221	0.0199	0.9043	-0.1459	0.7571	-0.0275	0.9226	0.1183	0.5532	-0.0730	0.9309	-0.0472	0.9096	0.0257	0.9464
---	---	1636964_at	0.1487	0.5429	0.2230	0.2310	0.1539	0.5444	-0.0409	0.9626	-0.1255	0.6247	-0.0846	0.7328	-0.0964	0.9168	-0.0499	0.9194	0.0465	0.9162
---	---	1636965_at	-0.0481	0.8664	0.2235	0.1317	0.0487	0.8082	-0.0337	0.9659	0.0449	0.8710	0.0786	0.7236	0.2353	0.6660	0.2265	0.2878	-0.0088	0.9809
---	---	1636966_s_at	0.1468	0.6258	0.0433	0.7359	0.1685	0.4149	0.0429	0.9675	0.0330	0.9324	-0.0099	0.9776	0.0157	0.9898	0.0939	0.7667	0.0782	0.8073
CG5315	CG5315	1636967_a_at	0.7802	0.0085	0.0801	0.5992	0.2990	0.0913	0.0180	0.9774	0.3583	0.0297	0.3403	0.0233	-0.1854	0.8513	-0.2890	0.4895	-0.1036	0.8430
CG5559	CG5559	1636968_at	0.2298	0.2585	0.0939	0.5803	0.3134	0.0756	-0.0747	0.9011	-0.0233	0.9321	0.0514	0.8151	-0.2640	0.4511	0.0215	0.9411	0.2855	0.1438
---	---	1636969_at	0.1580	0.2701	0.0912	0.5842	0.1273	0.5027	0.1650	0.6785	0.0630	0.7798	-0.1019	0.5836	0.0567	0.9400	0.0632	0.8386	0.0066	0.9856
CG9394	CG9394	1636970_at	-3.4648	0.0015	-1.7984	0.0145	-3.1370	0.0001	-1.2030	0.0458	-1.6513	0.0012	-0.4482	0.1413	0.1124	0.9740	0.0995	0.9414	-0.0129	0.9930
CG12464	CG12464	1636971_at	-0.0174	0.9521	-0.1153	0.5660	0.1043	0.5641	0.1951	0.7525	0.0379	0.9198	-0.1572	0.5504	-0.0042	0.9963	0.0258	0.9328	0.0300	0.9096
CG8003	CG8003	1636972_at	0.1842	0.5278	0.3092	0.5436	1.2572	0.0003	0.1075	0.8589	-0.4067	0.0650	-0.5142	0.0164	-0.3268	0.7644	0.2671	0.5855	0.5939	0.2148
smp-30	Drosophila cold ac	1636973_at	2.8664	0.0054	1.0449	0.2541	0.8729	0.0251	-0.1299	0.9518	-1.0460	0.0806	-0.9162	0.0861	0.0730	0.9862	-2.6044	0.0304	-2.6774	0.0421
dlp	Dally-like	1636974_at	-0.8229	0.0190	-0.8738	0.0686	-1.0111	0.0006	0.0287	0.9744	-0.1978	0.3705	-0.2265	0.2438	0.1268	0.9238	-0.2740	0.5523	-0.4009	0.3753
CG32236	CG32236	1636975_at	0.0373	0.8636	0.0581	0.5614	0.1266	0.4587	-0.0833	0.8546	-0.1696	0.3071	-0.0863	0.6019	0.1905	0.7726	0.0197	0.9664	-0.1709	0.5722
CG5322	CG5322	1636976_at	-1.4747	0.0027	-2.5460	0.0057	-2.8322	0.0000	-0.4691	0.2168	-0.0614	0.8327	0.4077	0.0503	-0.2005	0.8903	-1.1160	0.0625	-0.9155	0.1267
CG9318	CG9318	1636977_at	0.2566	0.3998	0.7693	0.1592	0.8157	0.0014	-0.2791	0.5039	-0.0512	0.8601	0.2278	0.2525	-0.3153	0.8236	0.5043	0.3842	0.8197	0.1897

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG17912	CG17912	1636978_at	0.7685	0.0380	-0.0076	0.9702	-0.5118	0.0431	-0.1516	0.8732	1.0967	0.0059	1.2483	0.0020	0.3921	0.6557	0.3252	0.3447	-0.0669	0.8905
CG7912	CG7912	1636979_at	0.5055	0.1756	-0.0145	0.9227	0.0131	0.9643	0.0764	0.9072	0.1933	0.3701	0.1168	0.5791	0.0445	0.9816	-0.3180	0.4502	-0.3625	0.3967
CG4643	CG4643	1636980_s_at	0.6220	0.0048	0.3792	0.1015	0.2100	0.4036	0.0562	0.9182	0.2704	0.1063	0.2142	0.1525	0.1884	0.8465	0.0671	0.9149	-0.1212	0.8069
---	---	1636981_at	0.1399	0.4453	0.0330	0.8459	-0.0008	0.9972	0.0532	0.9406	0.0403	0.8837	-0.0129	0.9597	-0.0709	0.9324	-0.1903	0.4881	-0.1194	0.6877
---	---	1636982_at	0.0940	0.6187	0.3061	0.1554	0.2493	0.1362	0.0189	0.9755	0.0172	0.9352	-0.0017	0.9928	0.1998	0.7324	0.2404	0.3259	0.0406	0.9067
---	---	1636983_at	0.1692	0.4767	0.0896	0.5335	-0.0325	0.8693	0.0206	0.9808	0.0916	0.7043	0.0710	0.7564	0.0336	0.9735	-0.0017	0.9988	-0.0352	0.9186
CG32092	CG32092	1636984_at	-0.1816	0.6334	0.0410	0.9226	0.1911	0.6004	-0.2410	0.7929	-0.3068	0.4269	-0.0658	0.8815	-0.3511	0.7893	-0.0998	0.9075	0.2513	0.6861
CG31666 /// chinmo	CG31666 /// chror	1636985_s_at	0.0422	0.7825	-0.0270	0.7901	-0.1148	0.4592	0.0331	0.9507	0.0850	0.5997	0.0519	0.7461	0.1647	0.7196	0.0475	0.8556	-0.1171	0.5615
l(2)35Df	lethal (2) 35Df	1636986_at	0.2260	0.6637	0.4981	0.1286	0.4269	0.0865	0.2614	0.6197	0.3502	0.1795	0.0888	0.7566	0.1774	0.9411	0.5184	0.4976	0.3410	0.6761
---	---	1636987_at	0.2521	0.1602	0.0249	0.8095	0.1356	0.3846	0.1399	0.6835	0.0004	0.9985	-0.1395	0.3400	0.0033	0.9970	0.0004	0.9998	-0.0029	0.9941
CG8476	CG8476	1636988_at	-0.0657	0.7423	0.1193	0.5074	-0.0508	0.7978	0.0414	0.9436	0.1164	0.5057	0.0750	0.6620	0.1935	0.7768	0.2007	0.5110	0.0072	0.9875
CG1399	Dcarmil	1636989_a_at	-0.6646	0.0711	0.0151	0.9733	0.6024	0.0195	0.4167	0.5617	-0.6051	0.1004	-1.0217	0.0081	-0.1295	0.9447	0.1693	0.8174	0.2988	0.6225
botv	Drosophila Ext-like	1636990_at	0.3695	0.2311	0.1112	0.6810	0.1203	0.6173	-0.1066	0.8358	0.1098	0.6067	0.2164	0.2128	-0.0664	0.9774	-0.1811	0.7932	-0.1146	0.8779
emp	epithelial membra	1636991_s_at	0.0570	0.6940	0.1653	0.5177	0.5885	0.0128	0.3349	0.2898	-0.2352	0.2046	-0.5701	0.0051	0.1592	0.8243	0.1159	0.7421	-0.0433	0.9168
---	---	1636992_at	0.1856	0.2412	0.2165	0.2158	0.1608	0.3876	-0.1073	0.8540	-0.1850	0.3908	-0.0778	0.7298	-0.0281	0.9717	-0.0078	0.9838	0.0203	0.9413
---	---	1636993_at	0.2560	0.2331	0.0000	1.0000	0.1696	0.4037	0.0367	0.9722	-0.0233	0.9528	-0.0600	0.8435	-0.0921	0.8825	-0.0683	0.8280	0.0237	0.9433
CG14712	CG14712	1636994_at	-0.2300	0.6872	-0.6058	0.3741	-1.0449	0.0133	-0.2451	0.8875	0.2727	0.6827	0.5178	0.3331	0.1273	0.9081	0.0221	0.9737	-0.1052	0.8307
tgy	twiggy	1636995_at	0.0650	0.7918	0.3896	0.0568	0.5831	0.0080	-0.0662	0.8791	-0.3256	0.0353	-0.2594	0.0549	-0.0565	0.9412	0.0237	0.9505	0.0802	0.7749
CG17329	CG17329	1636996_at	0.1909	0.5790	0.1662	0.3437	0.1472	0.5523	0.1525	0.8189	0.0918	0.7689	-0.0607	0.8402	0.1799	0.8480	0.1486	0.7475	-0.0313	0.9551
CG18528	CG18528	1636997_at	0.2797	0.4427	-0.1327	0.6374	-0.2864	0.3502	-0.2336	0.7838	0.3995	0.2570	0.6330	0.0489	-0.3251	0.6898	-0.1956	0.5776	0.1295	0.7338
sca	scabrous	1636998_at	-0.0795	0.6043	-0.2371	0.0863	0.1270	0.4704	0.1063	0.7825	0.2098	0.1850	0.1036	0.4987	-0.1235	0.8065	-0.0224	0.9478	0.1011	0.6577
MESK2	Misexpression suj	1636999_a_at	-1.5606	0.0023	-0.1092	0.8385	-0.2737	0.1703	-0.4045	0.5539	-1.7842	0.0010	-1.3797	0.0017	-0.1902	0.8825	-0.2582	0.6266	-0.0681	0.9197
Vha68-1	Vha68-1	1637000_at	-1.6708	0.0004	-2.2842	0.0056	-2.5137	0.0000	-0.0559	0.9311	0.1668	0.3914	0.2227	0.1887	0.2171	0.8331	-0.3986	0.3413	-0.6157	0.1764
---	---	1637001_at	-0.2121	0.2765	-1.0365	0.0790	-1.3713	0.0016	-0.0378	0.9857	0.9927	0.0479	1.0305	0.0266	-0.2593	0.7386	-0.0611	0.9035	0.1982	0.5700
CG14125	CG14125	1637002_at	0.3856	0.2149	0.0257	0.9132	0.1289	0.6116	-0.1206	0.8518	0.0490	0.8750	0.1696	0.4427	-0.0933	0.9029	-0.3178	0.2380	-0.2244	0.4319
CG6927	CG6927	1637003_at	0.2031	0.6567	-0.3677	0.5885	-0.7603	0.2285	-0.2044	0.7304	0.7685	0.0109	0.9728	0.0023	-0.2737	0.9400	0.0702	0.9696	0.3439	0.7982
pcs	poirot	1637004_at	-1.3644	0.0049	0.0106	0.9813	0.2224	0.2854	-0.1105	0.8817	-1.0852	0.0019	-0.9748	0.0019	-0.1558	0.8740	0.4891	0.1920	0.6449	0.1271
---	---	1637005_at	0.1755	0.3068	0.1325	0.5815	0.0410	0.8113	-0.2249	0.4861	-0.0057	0.9822	0.2193	0.1536	0.0721	0.9340	0.0974	0.7741	0.0252	0.9486
CG33174 /// DmirCG33174	CG33174	1637006_at	0.1076	0.4904	0.0214	0.8470	-0.0668	0.7442	0.0019	0.9978	-0.0273	0.9148	-0.0292	0.8936	0.0237	0.9816	-0.0113	0.9781	-0.0350	0.9107
CG5445 /// DyakCG5445	CG5445	1637007_s_at	-0.1405	0.3841	0.5722	0.0263	0.5686	0.0644	-0.0548	0.9389	-0.9153	0.0018	-0.8605	0.0014	-0.2100	0.8076	-0.2329	0.5336	-0.0229	0.9660
CG14687	CG14687	1637008_at	-2.6280	0.0283	-2.7563	0.0091	-3.4183	0.0000	0.5159	0.5363	0.1840	0.7158	-0.3319	0.4155	1.2368	0.6749	0.3005	0.8584	-0.9363	0.4569
CG17689	CG17689	1637009_at	0.1857	0.5571	0.0846	0.5103	0.2554	0.2719	-0.0869	0.8975	-0.0525	0.8556	0.0344	0.8974	-0.1052	0.9226	-0.0298	0.9600	0.0755	0.8772
---	---	1637010_at	0.2609	0.2806	0.1152	0.4088	0.1998	0.3004	-0.0309	0.9649	-0.2036	0.2833	-0.1727	0.3142	0.0951	0.8926	-0.0466	0.9080	-0.1416	0.6168
CG33287	CG33287	1637011_at	-0.0642	0.7985	-0.0403	0.6962	-0.0329	0.8526	0.0601	0.9542	0.2134	0.4767	0.1533	0.5934	-0.0542	0.9296	0.1213	0.5591	0.1755	0.3859
m2	E(spl) region trans	1637012_at	0.0764	0.7196	-0.0163	0.9208	0.1726	0.4817	-0.0047	0.9961	-0.3629	0.2502	-0.3581	0.2025	-0.0427	0.9764	-0.2606	0.4544	-0.2179	0.5523
CG34145	CG34145	1637013_at	0.1102	0.7661	0.2174	0.2221	0.2140	0.1758	0.0240	0.9774	-0.0116	0.9709	-0.0356	0.8836	0.2339	0.6903	0.2233	0.3441	-0.0107	0.9778
sinu	sinuous	1637014_at	-0.3446	0.1753	-0.2549	0.4967	-0.4006	0.0696	0.0752	0.9029	0.1978	0.3292	0.1226	0.5295	0.2889	0.7485	0.2620	0.5064	-0.0269	0.9618
ran	ran	1637015_s_at	0.2766	0.1688	0.3536	0.1648	0.1921	0.3034	-0.0278	0.9726	0.2529	0.2059	0.2807	0.1164	0.1754	0.8157	0.3793	0.2086	0.2039	0.5334
CG33324	CG33324	1637016_at	-1.0967	0.0977	0.0363	0.9282	0.0497	0.8569	0.2557	0.8166	-0.5934	0.1688	-0.8491	0.0356	0.3067	0.8424	0.3842	0.5622	0.0775	0.9300
Cont	contactin	1637017_at	-0.2924	0.1964	-0.0938	0.8396	-0.2837	0.0990	-0.1368	0.7770	0.1856	0.3677	0.3224	0.0755	0.0200	0.9922	0.4295	0.3358	0.4094	0.3825
CG5548 /// DyakCG5548	CG5548	1637018_at	0.1381	0.5703	0.4763	0.2202	0.3009	0.1761	-0.1648	0.7046	-0.6466	0.0067	-0.4818	0.0150	-0.0831	0.9523	-0.3655	0.3733	-0.2824	0.5175
---	---	1637019_at	0.2181	0.2846	-0.0058	0.9806	0.3084	0.0438	0.1104	0.7850	0.0973	0.5922	-0.0130	0.9507	-0.0075	0.9933	-0.0847	0.7158	-0.0772	0.7426
---	---	1637020_at	-0.0917	0.6539	0.0366	0.7793	0.0812	0.7144	0.1169	0.7735	0.1316	0.4585	0.0147	0.9458	-0.0179	0.9885	-0.0602	0.8764	-0.0424	0.9096
---	---	1637021_at	-0.0685	0.6783	0.0636	0.5684	0.1722	0.3358	0.0857	0.8942	0.0305	0.9197	-0.0552	0.8212	-0.0719	0.8888	0.0080	0.9829	0.0799	0.7150
CG3532	CG3532	1637022_at	-0.4546	0.2879	1.0422	0.0798	1.1816	0.0004	0.1505	0.9029	-0.7771	0.0560	-0.9276	0.0174	0.0582	0.9761	0.8213	0.0897	0.7631	0.1326
CG14370	CG14370	1637023_at	0.1045	0.5501	0.0090	0.9340	-0.0654	0.7973	0.0360	0.9545	0.0215	0.9292	-0.0144	0.9473	0.1469	0.8439	-0.0852	0.8353	-0.2321	0.4647
VhaPPA1-1	VhaPPA1-1	1637024_at	-1.1387	0.0030	-0.6262	0.1037	-0.8669	0.0005	-0.2368	0.3793	-0.6293	0.0023	-0.3925	0.0113	-0.1266	0.9238	-0.2488	0.5923	-0.1222	0.8239
---	---	1637025_at	-0.1219	0.6644	-0.0688	0.5431	-0.0643	0.7327	0.0029	0.9973	0.0373	0.9105	0.0344	0.9046	0.0451	0.9589	0.1069	0.7063	0.0618	0.8480
---	---	1637026_at	-0.2926	0.1769	0.0567	0.6097	-0.0180	0.9486	-0.1980	0.7753	-0.0921	0.7963	0.1059	0.7348	0.0161	0.9913	0.1141	0.7591	0.0980	0.7935
CG10165	CG10165	1637027_s_at	-0.0404	0.9008	1.1398	0.0108	0.9767	0.0014	-0.1318	0.7073	-0.1800	0.2641	-0.0482	0.7881	0.0173	0.9928	0.9328	0.0622	0.9154	0.0812
SIFR	SIFamide recepto	1637028_at	0.1466	0.6236	-0.4252	0.1498	-0.1872	0.5947	0.1406	0.8981	0.5453	0.1289	0.4047	0.2091	-0.2571	0.8202	-0.2026	0.7027	0.0545	0.9342
PQBP-1	Poly-glutamine tra	1637029_at	0.3430	0.1464	0.0950	0.6659	0.0010	0.9968	0.0134	0.9857	0.1950	0.2583	0.1816	0.2385	0.1654	0.8513	-0.0298	0.9613	-0.1951	0.6196
CG5948	CG5948	1637030_at	-0.0254	0.9235	0.0351	0.7990	0.1422	0.5658	0.1247	0.8822	0.1349	0.6800	0.0102	0.9776	-0.1478	0.7726	0.0320	0.9271	0.1798	0.4258
gsb-n	gooseberry neuro	1637031_at	-4.0208	0.0005	-0.7181	0.2476	-2.1065	0.0020	-1.2938	0.2789	-3.3849	0.0012	-2.0911	0.0059	-0.0875	0.9555	-0.3604	0.4297	-0.2730	0.5746
beta4GalNAcTB	beta4GalNAcTB	1637032_at	-0.4414	0.1249	0.8837	0.0323	0.7476	0.0117	-0.3477	0.5511	-1.5708	0.0009	-1.2231	0.0015	-0.1205	0.9063	0.0309	0.9590	0.1513	0.7148
sip1	septin interacting	1637033_at	0.1115	0.7939	0.0108	0.9901	0.3531	0.0818	0.0207	0.9760	0.0807	0.6768	0.0600	0.7436	-0.3600	0.8480	-0.0368	0.9796	0.3233	0.7152
CG32371	CG32371	1637034_at	0.1444	0.3548	0.1992	0.3522	0.0799	0.6770	-0.0305	0.9675	0.0359	0.8932	0.0664	0.7573	0.0524	0.9491	-0.0138	0.9737	-0.0663	0.8301
unc	section 5	1637035_at	-0.0440	0.8478	0.0351	0.7194	0.0030	0.9935	0.0257	0.9774	-0.1379	0.5602	-0.1635	0.4267	0.0809	0.8999	-0.0558	0.8678	-0.1367	0.5886
pr	purple	1637036_s_at	0.7507	0.0042	0.9942	0.0521	0.7184	0.0117	-0.0406	0.9507	-0.1065	0.5901	-0.0659	0.7361	0.2208	0.8461	0.2149	0.6827	-0.0059	0.9945
ade3	Complementation	1637037_at	2.1808	0.0020	0.8837	0.4523	2.1409	0.0001	0.4331	0.3733	0.3388	0.2160	-0.0943	0.7528	-0.8071	0.7644	-0.9818	0.3856	-0.1747	0.9121
tex	tex	1637038_at	0.5012	0.0110	0.1723	0.6437	0.4725	0.0236	-0.0361	0.9460	-0.0671	0.6990	-0.0310	0.8599	-0.2485	0.7677	-0.1498	0.7115	0.0987	0.8255
mus309	bloom	1637039_at	0.3273	0.1579	0.0455	0.7969	0.1070	0.5391	0.0775	0.8640	0.1355	0.4132	0.0580	0.7371	0.1153	0.8647	0.1261	0.6730	0.0108	0.9810
---	---	1637040_at	-0.1454	0.4469	0.0757	0.6657	0.1228	0.4711	0.1556	0.6656	0.0131	0.9595	-0.1426	0.3687	-0.0132	0.9913	0.1251	0.6381	0.1383	0.5995
CG3332 /// DtraCG3332	CG3332	1637041_a_at	1.3130	0.3656	0.0141	0.9113	0.1397	0.6100	0.0421	0.9677	-0.1047	0.7460	-0.1467	0.5906	-0.0959	0.9903	-1.3043	0.4459	-1.2084	0.4945
CG11961	CG11961	1637042_a_at	1.8156	0.0008	2.0694	0.0281	2.4325	0.0000	0.1142	0.7921	-0.5151	0.0105	-0.6293	0.0026	-0.3220	0.8521	-0.2745	0.7439	0.0474	0.9640
pip	pipe	1637043_at	0.1381	0.6880	0.1072	0.4024	0.0084	0.9729	-0.0071	0.9956	-0.2002	0.5496	-0.1931	0.5234	0.1057	0.8270	-0.1877	0.3372	-0.2934	0.1674
---	---	1637044_at	-0.3848	0.0708	-0.1790	0.2941	-0.4142	0.0269	0.1180	0.7967	-0.0709	0.7481	-0.1889	0.2592	0.0894	0.8940	-0.0827	0.7922	-0.1721	0.5078
---	---	1637045_s_at	0.1191	0.6656	0.2867	0.1652	0.3245	0.1676	0.1716	0.5931	-0.1694	0.3030	-0.3410	0.0271	0.2012	0.8145	0.1123	0.8053	-0.0889	0.8453
l(2)03659	Hsp related	1637046_at	0.2916	0.4139	0.0478	0.7112	-0.0959	0.6919	-0.0398	0.9493	-0.0111	0.9676	0.0287	0.8907	-0.0296	0.9901	-0.3580	0.5037	-0.3283	0.5493

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1637047_at	0.1216	0.6890	0.2559	0.1345	0.2456	0.1775	0.0290	0.9753	-0.1739	0.4649	-0.2029	0.3301	0.0646	0.9239	0.0823	0.7625	0.0177	0.9566
CG1662	CG1662	1637048_at	-0.0417	0.8826	1.0271	0.0621	1.2974	0.0018	0.2171	0.5913	-1.0967	0.0008	-1.3138	0.0002	0.0634	0.9779	0.1550	0.8319	0.0916	0.9034
bi	optomotorblind	1637049_at	0.0477	0.8378	-0.0674	0.5332	0.0459	0.7859	0.0469	0.9602	0.1202	0.6683	0.0732	0.7908	0.0355	0.9535	0.0221	0.9402	-0.0134	0.9595
CG33054	CG33054	1637050_at	-0.1875	0.5365	0.0759	0.8777	-0.2912	0.1660	-0.2485	0.7929	-0.3974	0.3083	-0.1489	0.7172	0.0791	0.9458	-0.2244	0.5414	-0.3034	0.3973
RhoGAP71E	RhoGAP71E	1637051_at	1.1127	0.0017	0.7346	0.0287	0.7603	0.0399	0.4174	0.5756	0.9694	0.0194	0.5520	0.1055	0.1773	0.7677	0.6363	0.0365	0.4590	0.1063
---	---	1637052_at	0.2204	0.1755	0.0947	0.5666	0.1505	0.4965	-0.0130	0.9860	0.0604	0.7713	0.0735	0.6851	-0.0470	0.9421	-0.0455	0.8730	0.0015	0.9965
Cas	CAS/CSE1 segre	1637053_at	0.7951	0.0043	0.2018	0.5480	0.3762	0.0737	-0.2963	0.3374	0.4494	0.0199	0.7456	0.0013	-0.4160	0.6824	-0.1604	0.7439	0.2556	0.5642
RluA-1	multiple	1637054_s_at	-3.8491	0.0060	-5.0293	0.0026	-5.1990	0.0000	0.1411	0.9247	0.4733	0.2953	0.3322	0.4299	0.0765	0.9898	-1.0007	0.4382	-1.0771	0.4114
---	---	1637055_s_at	-0.2693	0.1470	-0.2876	0.1303	-0.3062	0.0718	-0.3286	0.3619	-0.5222	0.0192	-0.1935	0.2982	-0.2584	0.7947	-0.6174	0.1374	-0.3591	0.3972
CG11200	CG11200	1637056_s_at	1.5423	0.0049	0.6834	0.2656	1.1341	0.0201	0.1181	0.9423	-0.1968	0.7156	-0.3149	0.4776	-0.0879	0.9677	-0.8329	0.1395	-0.7449	0.2107
nrm	Neuromusculin	1637057_at	-2.6865	0.0014	-2.5395	0.0638	-3.3159	0.0000	-0.5229	0.1009	-0.6145	0.0064	-0.0916	0.6343	0.2574	0.9460	-0.4246	0.7567	-0.6820	0.5776
CG18110	CG18110	1637058_at	0.1886	0.4533	-0.0319	0.7938	0.1166	0.6664	0.0355	0.9693	0.1379	0.5947	0.1024	0.6806	-0.0636	0.9487	-0.0883	0.8185	-0.0247	0.9525
DnaJ-1	dnaJ like1	1637059_s_at	-0.7829	0.2445	-0.2849	0.5481	-0.3640	0.5135	0.2037	0.9311	-0.0514	0.9604	-0.2552	0.7297	0.2652	0.9061	0.5308	0.5251	0.2656	0.7804
Nopp140	Nopp140	1637060_a_at	0.6252	0.0910	0.4396	0.0461	0.6867	0.0630	0.1291	0.7942	0.5226	0.0179	0.3935	0.0361	0.0297	0.9914	0.5066	0.4134	0.4769	0.4597
CG6151	CG6151	1637061_at	0.0907	0.6824	0.0385	0.7035	-0.1106	0.6181	-0.0421	0.9598	-0.0234	0.9407	0.0187	0.9462	0.0678	0.9246	-0.0154	0.9688	-0.0833	0.7640
CG18745	CG18745	1637062_at	0.1483	0.8868	0.3965	0.3468	0.3205	0.1588	-0.0421	0.9893	-0.4803	0.5323	-0.4382	0.5333	0.0651	0.9829	-0.1990	0.8287	-0.2641	0.7423
CG33099	CG33099	1637063_at	1.3316	0.1457	1.1145	0.0455	0.2588	0.1562	-0.4762	0.3452	-0.1291	0.6999	0.3471	0.1804	0.3627	0.8953	-0.5713	0.6024	-0.9340	0.3764
Adk2	Adenylate kinase-	1637064_at	0.3189	0.3122	0.3349	0.1638	0.5336	0.0640	-0.0392	0.9688	-0.5830	0.0323	-0.5439	0.0273	-0.0944	0.9525	-0.4652	0.3146	-0.3708	0.4488
CG12022	CG12022	1637065_a_at	-0.4149	0.0431	-0.0156	0.9399	0.0293	0.8806	0.1381	0.8000	-0.1891	0.4024	-0.3272	0.0954	-0.0432	0.9648	0.0177	0.9674	0.0608	0.8601
CG8239	CG8239	1637066_at	-0.2698	0.4828	0.4976	0.1302	0.1134	0.5597	-0.1730	0.7680	0.0577	0.8596	0.2307	0.3083	0.1919	0.8768	0.7557	0.1211	0.5638	0.2594
CG6074	CG6074	1637067_at	-1.9236	0.2665	-2.8885	0.0101	-3.4713	0.0000	-0.2579	0.6160	-0.4458	0.0840	-0.1880	0.4365	0.1254	0.9898	-1.6925	0.4285	-1.8180	0.4045
CG6621	CG6621	1637068_at	-0.3222	0.2860	0.1914	0.5633	0.3388	0.0921	-0.1165	0.8005	-0.5137	0.0135	-0.3972	0.0249	-0.1220	0.9445	0.0329	0.9702	0.1549	0.8147
CG4622	CG4622	1637069_at	-0.0952	0.7572	0.1717	0.3433	0.6894	0.0027	0.1800	0.6445	-0.1960	0.3051	-0.3761	0.0335	-0.2425	0.8093	0.0939	0.8754	0.3363	0.4322
CG34340	CG10017	1637070_at	-0.0026	0.9915	0.0133	0.8979	0.1542	0.2836	0.1307	0.7688	0.0181	0.9492	-0.1126	0.5414	-0.0237	0.9816	-0.0163	0.9669	0.0075	0.9849
CG12851	CG12851	1637071_at	0.1595	0.5294	0.0982	0.5114	0.1514	0.4204	-0.0422	0.9507	0.1478	0.4510	0.1900	0.2632	-0.0845	0.9127	0.0680	0.8550	0.1525	0.6005
CG11459	CG11459	1637072_at	2.7707	0.0007	0.4633	0.3996	2.5673	0.0055	0.8882	0.6195	2.1127	0.0256	1.2245	0.1223	-1.2896	0.2884	-0.2481	0.7344	1.0415	0.1289
CG12995	CG12995	1637073_a_at	0.2168	0.4617	0.2092	0.3799	0.1973	0.2013	0.0573	0.9039	-0.0570	0.7578	-0.1144	0.4309	0.0595	0.9545	-0.0043	0.9949	-0.0637	0.8779
CG11035	CG11035	1637074_at	-0.4730	0.0774	-0.2117	0.3019	-0.3684	0.0573	-0.2556	0.6872	-0.4092	0.1634	-0.1536	0.6062	-0.1126	0.8609	-0.2153	0.4016	-0.1027	0.7287
CG15115	CG15115	1637075_at	0.0046	0.9812	-0.0934	0.5562	0.0321	0.8891	0.0957	0.8605	0.0929	0.6764	-0.0028	0.9909	-0.1260	0.8400	-0.1178	0.6816	0.0082	0.9847
CG8861	CG8861	1637076_at	-0.3031	0.5302	-0.4412	0.1051	-0.2265	0.1178	0.0714	0.9507	-0.1774	0.6151	-0.2488	0.4035	-0.2322	0.7733	-0.3465	0.3087	-0.1143	0.7831
CG3777 /// DsimCG3777	CG3777	1637077_s_at	0.5531	0.2455	1.0939	0.1380	1.3188	0.0034	0.2463	0.7805	0.1671	0.6918	-0.0792	0.8526	0.0927	0.9761	0.6243	0.3958	0.5316	0.4892
Poxm	pox meso	1637078_a_at	-0.0597	0.8723	0.4498	0.1579	0.3449	0.0759	-0.1198	0.9046	-0.4766	0.1407	-0.3569	0.2197	0.0534	0.9775	-0.0795	0.9081	-0.1329	0.8112
CG5098	CG5098	1637079_at	0.5052	0.2122	0.1284	0.7329	-0.3759	0.0722	-0.0549	0.9649	0.7803	0.0261	0.8352	0.0124	0.5443	0.6557	0.4150	0.3949	-0.1294	0.8364
CG9129 /// DyakCG9129	CG9129	1637080_at	-0.1148	0.6775	-0.0452	0.7758	0.3075	0.1927	0.1823	0.7857	-0.0442	0.9087	-0.2265	0.3774	-0.0758	0.9462	-0.1207	0.7710	-0.0449	0.9234
---	---	1637081_at	0.0663	0.6487	0.0791	0.6535	0.2095	0.2727	-0.0398	0.9610	0.0634	0.8137	0.1032	0.6439	-0.0630	0.9421	0.0352	0.9345	0.0982	0.7494
---	---	1637082_at	-0.0728	0.7562	0.1313	0.4857	0.2142	0.3723	0.0514	0.9496	-0.3766	0.0962	-0.4280	0.0406	0.0943	0.9441	-0.0670	0.9156	-0.1613	0.7287
---	---	1637083_at	-0.1417	0.3367	0.0059	0.9776	0.1848	0.2395	0.1590	0.5628	0.0037	0.9865	-0.1553	0.2180	0.0259	0.9816	0.1293	0.6573	0.1034	0.7351
CG14883	CG14883	1637084_at	-0.3890	0.2283	0.6083	0.0191	0.3289	0.2151	-0.1277	0.8899	-0.3391	0.2747	-0.2115	0.4719	0.1285	0.9142	0.5087	0.2107	0.3802	0.3766
Cdk5	Cyclin-dependent	1637085_at	-0.0613	0.7667	0.3434	0.3278	-0.0403	0.8458	-0.0643	0.9300	-0.1833	0.4076	-0.1189	0.5792	0.1783	0.7768	0.0781	0.8269	-0.1002	0.7492
CG11843	CG11843	1637086_at	0.1505	0.5919	-0.2506	0.4081	-0.0693	0.7704	0.1265	0.8190	0.2537	0.2420	0.1272	0.5499	0.0653	0.9635	-0.0097	0.9914	-0.0750	0.8880
---	---	1637087_at	0.0558	0.7484	-0.1118	0.2874	0.3145	0.1819	0.1946	0.6856	0.2138	0.3476	0.0192	0.9460	-0.1479	0.8461	-0.0936	0.8185	0.0543	0.9004
---	---	1637088_at	0.0178	0.9467	0.0147	0.8861	0.1569	0.3981	-0.0879	0.8473	-0.1089	0.5455	-0.0210	0.9186	-0.0738	0.9401	0.0800	0.8455	0.1539	0.6375
Syb	synaptobrevin	1637089_at	0.0065	0.9753	0.3415	0.0848	0.3460	0.0876	0.0362	0.9584	-0.1052	0.5999	-0.1414	0.4085	0.0350	0.9741	0.2639	0.3058	0.2289	0.3964
---	---	1637090_at	-0.0673	0.7880	0.3370	0.3181	0.0766	0.7293	-0.0716	0.9345	-0.2633	0.3109	-0.1916	0.4270	0.1429	0.8439	0.0383	0.9358	-0.1046	0.7625
CG9772 /// DyakCG9772	CG9772	1637091_a_at	0.2469	0.3606	0.1733	0.5334	-0.1233	0.7146	-0.5340	0.1140	0.0409	0.8800	0.5749	0.0073	-0.3079	0.8270	-0.0087	0.9942	0.2992	0.6328
Upf3	Upf3	1637092_a_at	-0.1101	0.5648	0.1887	0.1567	0.3041	0.1823	0.1007	0.8087	0.1111	0.5293	0.0104	0.9597	-0.0416	0.9653	0.2610	0.2992	0.3026	0.2565
CG34045	CG34045	1637093_at	0.1424	0.4313	0.1005	0.4199	0.3977	0.2309	0.0176	0.9866	0.0445	0.8983	0.0269	0.9314	-0.1284	0.8940	0.0416	0.9429	0.1700	0.6641
CG17564	CG17564	1637094_at	0.0610	0.7659	0.0821	0.4597	-0.0845	0.6113	0.0397	0.9584	0.1042	0.6439	0.0645	0.7714	0.1256	0.8510	0.1679	0.5629	0.0423	0.9118
CG15818 /// DsimCG1581	CG15818	1637095_at	0.5973	0.3807	-0.0945	0.4643	-2.5569	0.0018	-2.4583	0.0347	0.1060	0.9045	2.5643	0.0016	-0.1637	0.9441	-0.5671	0.4192	-0.4034	0.5940
---	---	1637096_s_at	0.2256	0.3341	-0.0111	0.9719	-0.0932	0.5792	0.1555	0.7887	0.1625	0.5202	0.0071	0.9810	0.3588	0.6898	0.0170	0.9797	-0.3417	0.3674
CG7497	CG7497	1637097_at	1.0833	0.0144	2.9086	0.0096	2.1758	0.0002	0.1822	0.7982	-0.6288	0.0363	-0.8110	0.0077	0.8881	0.5421	1.1137	0.1083	0.2256	0.7787
CG9101	CG9101	1637098_at	0.0015	0.9950	-0.0786	0.7770	0.1111	0.5018	0.2127	0.7605	0.1130	0.7519	-0.0997	0.7618	-0.0110	0.9914	0.0325	0.9277	0.0435	0.8891
CG17049	CG17049	1637099_at	-0.0378	0.8180	-0.1130	0.5547	0.1142	0.5974	-0.0233	0.9711	-0.0937	0.6006	-0.0703	0.6810	-0.2593	0.7478	-0.0091	0.9906	0.2502	0.4801
Bx42	Ski-interacting pro	1637100_at	-0.4034	0.1704	-0.0745	0.8888	0.2778	0.1626	0.2236	0.5896	-0.2153	0.3109	-0.4389	0.0275	-0.1067	0.9589	0.0742	0.9377	0.1809	0.8001
pasha	partner of drosha	1637101_at	0.0853	0.8117	0.2489	0.3330	0.2872	0.1194	-0.0963	0.9130	0.2800	0.3204	0.3763	0.1290	-0.2080	0.8062	0.3459	0.3152	0.5539	0.1443
---	---	1637102_at	-0.0365	0.8333	-0.1416	0.2632	-0.1526	0.2910	0.0542	0.9029	0.0188	0.9252	-0.0354	0.8293	0.1002	0.8513	0.0188	0.9590	-0.0814	0.7492
CG9640	CG9640	1637103_at	-0.7678	0.0102	-0.8424	0.0949	-0.8330	0.0038	-0.1629	0.6785	-0.0848	0.6873	0.0781	0.6851	-0.2251	0.8744	-0.3127	0.6012	-0.0876	0.9096
---	---	1637104_at	0.0771	0.6961	0.1669	0.1750	0.1562	0.4599	-0.0446	0.9592	-0.2075	0.3749	-0.1628	0.4509	0.1125	0.8837	0.0231	0.9620	-0.0893	0.8011
rgr	regular	1637105_at	0.0093	0.9753	-1.6210	0.0014	-1.5080	0.0033	0.0446	0.9777	0.8949	0.0365	0.8502	0.0286	0.0460	0.9742	-0.4227	0.2168	-0.4686	0.2072
CG7197 /// DereCG7197 ///	CG7197	1637106_at	-0.3091	0.2119	0.4757	0.0872	0.7945	0.0006	0.0268	0.9728	-0.7307	0.0035	-0.7575	0.0018	-0.5516	0.5821	0.0086	0.9935	0.5602	0.2287
CG10809	CG10809	1637107_at	0.1597	0.5620	-0.0563	0.6766	-0.2368	0.1742	0.0617	0.9314	0.2461	0.2517	0.1844	0.3482	0.2226	0.7997	-0.0800	0.8787	-0.3027	0.4153
---	---	1637108_at	0.2315	0.4395	-0.1113	0.7263	-0.0215	0.9190	0.1406	0.9093	0.2823	0.4929	0.1417	0.7370	0.0425	0.9463	0.0172	0.9558	-0.0253	0.9234
CG3961	CG3961	1637109_s_at	-1.8737	0.0005	-2.3992	0.0101	-3.4645	0.0000	-0.1793	0.8387	0.8109	0.0223	0.9902	0.0056	0.8005	0.3482	0.2309	0.6380	-0.5696	0.2231
CG3618	CG3618	1637110_at	-1.6967	0.0019	-1.3892	0.0103	-1.7646	0.0000	-0.1847	0.5680	-0.4768	0.0113	-0.2921	0.0533	0.0934	0.9296	-0.2036	0.5728	-0.2970	0.3939
PR2	focal adhesion kin	1637111_a_at	-0.1602	0.6063	-0.1445	0.6550	-0.0598	0.8414	0.0178	0.9835	0.1223	0.5649	0.1046	0.5961	0.0239	0.9913	0.1351	0.8132	0.1112	0.8439
---	---	1637112_at	0.0200	0.9066	-0.0658	0.5524	-0.0126	0.9477	0.0189	0.9790	0.1061	0.5856	0.0872	0.6341	-0.0163	0.9831	0.0279	0.9204	0.0442	0.8474
CG15105	CG15105	1637113_s_at	-2.6760	0.0022	-3.3468	0.0075	-4.2054	0.0000	0.0082	0.9932	0.2231	0.2397	0.2148	0.2044	1.0584	0.4584	-0.2245	0.8132	-1.2829	0.1156
Or47b	Olfactory receptor	1637114_at	0.1936	0.2071	0.1298	0.4048	-0.0065	0.9859	0.0010	0.9988	0.0452	0.8362	0.0442	0.8190	-0.1272	0.8270	-0.0322	0.9341	0.0950	0.7314
---	---	1637115_at	0.4699	0.1800	0.1564	0.4558	0.0222	0.9274	0.1279	0.8246	0.1392	0.5638	0.0113	0.9682	0.0864	0.9390	-0.0301	0.9587	-0.1165	0.7764

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1637116_at	0.4978	0.0112	-0.0516	0.6498	-0.0014	0.9963	0.1646	0.8671	0.5410	0.1182	0.3765	0.2268	-0.2018	0.8305	-0.1362	0.7787	0.0656	0.9046
CG9483	CG9483	1637117_at	0.0404	0.8774	-0.0204	0.8828	0.2449	0.2146	0.0982	0.8340	-0.0825	0.6846	-0.1807	0.2587	-0.1569	0.8122	-0.1600	0.5818	-0.0031	0.9948
CG30010	CG30010	1637118_at	0.5126	0.0463	0.0849	0.7985	0.0548	0.8097	0.0847	0.9036	0.0653	0.8163	-0.0194	0.9461	-0.1588	0.8513	-0.3743	0.2698	-0.2154	0.5630
CG16863	CG16863	1637119_at	0.0888	0.7821	-0.1814	0.3908	-0.0207	0.9432	-0.0505	0.9658	0.3044	0.3360	0.3549	0.2019	-0.3124	0.7707	0.0548	0.9407	0.3672	0.4357
CG30098	CG30098	1637120_at	1.7501	0.0650	-0.9916	0.4827	0.1410	0.8642	0.7626	0.5166	2.7376	0.0017	1.9751	0.0042	-0.3295	0.9589	-0.0762	0.9829	0.2533	0.9199
Rbp4	Testis specific RR	1637121_at	0.0603	0.7578	-0.0412	0.7758	-0.1177	0.5837	0.0329	0.9643	0.1941	0.3272	0.1612	0.3718	0.0160	0.9898	-0.2611	0.3135	-0.2771	0.3082
CG30436	CG30436	1637122_at	-0.0374	0.8335	-0.1478	0.3962	0.0018	0.9944	0.0714	0.8939	0.1281	0.4929	0.0566	0.7733	0.0248	0.9862	0.0843	0.8551	0.0594	0.8982
CG33281	CG33281	1637123_a_at	-0.8727	0.0030	-0.6104	0.0555	-0.8121	0.0159	0.2518	0.7478	0.0667	0.8846	-0.1851	0.5880	-0.0422	0.9589	-0.0049	0.9929	0.0374	0.9089
sec13	sec13	1637124_at	0.2518	0.1525	0.6705	0.0166	0.8265	0.0007	-0.0427	0.9253	-0.0016	0.9939	0.0412	0.7878	-0.0961	0.8609	0.3796	0.0955	0.4757	0.0702
CG2004	CG2004	1637125_at	0.8709	0.0332	1.0480	0.0114	1.6868	0.0001	0.2712	0.5680	0.7094	0.0106	0.4382	0.0489	-0.2494	0.7953	0.9718	0.0391	1.2212	0.0333
CG4983	CG4983	1637126_at	0.0536	0.7695	-0.2442	0.0772	-0.1512	0.3027	0.0496	0.9491	0.2995	0.1643	0.2499	0.1962	-0.2319	0.6749	-0.1495	0.5259	0.0825	0.7527
CG10958	CG10958	1637127_at	0.4884	0.1623	0.2274	0.3430	0.2460	0.1575	-0.1313	0.8721	0.2415	0.4087	0.3729	0.1405	-0.0970	0.9342	0.0052	0.9949	0.1022	0.8305
CG33340	CG33340	1637128_a_at	0.1802	0.4601	0.2407	0.2064	0.2834	0.1530	-0.0113	0.9883	-0.2220	0.2125	-0.2107	0.1861	0.0187	0.9892	-0.0602	0.8850	-0.0789	0.8271
GstE3	Glutathione S tran	1637129_at	0.9626	0.0166	0.7641	0.1354	1.0401	0.0115	0.4672	0.1787	0.1750	0.4202	-0.2922	0.1188	0.3189	0.8473	0.0680	0.9507	-0.2509	0.7524
CG30159	CG30159	1637130_at	0.3451	0.1779	0.3067	0.0679	0.5780	0.0029	-0.0288	0.9733	-0.3060	0.1478	-0.2772	0.1429	-0.2368	0.7116	-0.2629	0.3131	-0.0261	0.9440
---	---	1637131_at	0.1088	0.4681	-0.1111	0.4110	-0.0057	0.9770	0.2836	0.3463	0.2829	0.1025	-0.0007	0.9978	-0.0001	0.9999	-0.1260	0.6043	-0.1259	0.6081
---	---	1637132_at	0.1321	0.4394	-0.0215	0.8625	0.0880	0.7236	0.0583	0.9376	-0.0255	0.9338	-0.0838	0.7176	-0.0767	0.8888	-0.1021	0.6495	-0.0254	0.9297
CG11350	CG11350	1637133_x_at	0.3055	0.1777	0.1649	0.3382	0.2915	0.2071	0.1335	0.8379	0.0046	0.9899	-0.1289	0.5999	0.0121	0.9928	-0.0866	0.8353	-0.0987	0.7923
Or94a	Odorant receptor	1637134_at	0.4295	0.1132	-0.0410	0.7090	-0.2254	0.3300	-0.2302	0.6854	0.1290	0.6677	0.3592	0.1268	0.0214	0.9894	-0.1822	0.6156	-0.2036	0.5709
CG15321	CG15321	1637135_at	0.9010	0.0750	-0.0652	0.9196	-0.0982	0.7552	0.2473	0.8283	1.0514	0.0246	0.8042	0.0459	0.2739	0.8751	0.4427	0.5333	0.1689	0.8474
laf	labial associated f	1637136_at	0.3275	0.1924	0.2100	0.1574	0.0186	0.9548	-0.3193	0.5891	-0.0171	0.9705	0.3022	0.2635	-0.0344	0.9778	-0.1124	0.7455	-0.0780	0.8341
Dhc64C	dynein	1637137_a_at	0.2915	0.1056	0.4980	0.2687	0.4430	0.0182	-0.1253	0.8111	-0.2241	0.2845	-0.0988	0.6440	0.0609	0.9577	0.0175	0.9751	-0.0435	0.9224
CG13898	CG13898	1637138_at	-0.0975	0.6227	0.0473	0.6068	0.0369	0.8452	0.0010	0.9988	-0.1308	0.4845	-0.1319	0.4318	-0.0166	0.9839	-0.0330	0.9085	-0.0164	0.9507
nab	nab	1637139_at	0.2948	0.1955	0.1024	0.5952	0.1488	0.3972	0.2754	0.5735	0.1504	0.5812	-0.1250	0.6248	0.2336	0.7036	0.0866	0.7856	-0.1470	0.5906
CG32830	CG32830	1637140_at	0.0627	0.7619	0.0811	0.6110	0.1873	0.3562	-0.1291	0.6886	-0.0871	0.5926	0.0420	0.8030	-0.2026	0.7337	-0.0899	0.7727	0.1126	0.6892
CG4705	CG4705	1637141_at	0.3563	0.1217	0.6095	0.0562	0.6053	0.0136	-0.1680	0.7327	-0.2862	0.1890	-0.1182	0.5895	-0.0700	0.9460	-0.0845	0.8407	-0.0145	0.9769
CG9547	CG9547	1637142_at	1.0729	0.0011	0.7959	0.0386	0.9295	0.0009	0.0739	0.8856	-0.2907	0.0954	-0.3646	0.0272	0.0259	0.9848	-0.5184	0.0933	-0.5443	0.1039
CG9643	CG9643	1637143_at	0.1428	0.5419	0.5914	0.0902	0.5047	0.0188	0.0140	0.9857	-0.0769	0.7115	-0.0908	0.6183	0.0171	0.9923	0.3174	0.4333	0.3003	0.4747
Map205	Microtubule-assoc	1637144_a_at	-0.6756	0.0627	-1.2823	0.0075	-1.5764	0.0006	-0.1185	0.8462	0.4482	0.0507	0.5667	0.0124	0.1800	0.9092	-0.2103	0.7497	-0.3903	0.4994
CG14787	CG14787	1637145_at	0.1119	0.4607	0.3153	0.3828	0.0367	0.8680	-0.1553	0.7149	-0.1370	0.5037	0.0183	0.9394	0.1407	0.8461	0.0120	0.9832	-0.1286	0.7007
---	---	1637146_at	-0.9230	0.0070	-0.5877	0.0265	-0.6795	0.0227	-0.0231	0.9852	-0.3412	0.2177	-0.3182	0.1993	0.0311	0.9878	-0.2442	0.6166	-0.2752	0.5689
---	---	1637147_s_at	0.2341	0.4108	0.2509	0.3143	0.2367	0.3228	-0.0294	0.9777	-0.0701	0.8418	-0.0407	0.9007	0.0866	0.9342	-0.0111	0.9872	-0.0977	0.8173
CG13366 /// DmirCG13366	CG13366	1637148_at	-0.4731	0.1277	-0.4916	0.0624	-0.4591	0.0871	-0.1869	0.7673	0.1923	0.4960	0.3792	0.1133	-0.1587	0.8782	0.3500	0.3735	0.5087	0.2201
Fit1 /// ran	Fermitin 1 /// ran	1637149_s_at	-0.2440	0.2200	-0.2765	0.1028	-0.7392	0.0013	0.0964	0.7764	0.6256	0.0017	0.5292	0.0021	0.4393	0.5126	0.5288	0.1042	0.0895	0.8222
CG13928	CG13928	1637150_at	-1.3280	0.0635	0.0138	0.9308	-0.5096	0.0554	-0.2168	0.8968	-1.1027	0.0507	-0.8859	0.0742	0.1295	0.9447	-0.1486	0.8467	-0.2781	0.6475
mre11	meiotic recombina	1637151_at	-0.8173	0.0885	-0.8902	0.1149	-1.2632	0.0001	-0.4304	0.1183	-0.3451	0.0437	0.0853	0.6117	-0.1524	0.9449	-0.4677	0.4951	-0.3152	0.6630
---	---	1637152_at	0.0824	0.6408	-0.1466	0.3103	-0.1013	0.6408	0.1903	0.6327	0.3458	0.0783	0.1555	0.3916	-0.0426	0.9683	-0.0305	0.9473	0.0121	0.9788
Dis3	Dis3	1637153_at	0.2613	0.2150	0.0891	0.8337	-0.0009	0.9980	-0.1151	0.8402	0.3000	0.1573	0.4151	0.0362	-0.0172	0.9939	0.2322	0.6729	0.2495	0.6399
CG7470	CG7470	1637154_at	1.5939	0.0994	1.3488	0.3322	2.1668	0.0000	0.5379	0.6672	-0.2622	0.7027	-0.8001	0.1313	-0.3623	0.9457	-0.4540	0.8324	-0.0917	0.9704
CG32407	CG32407	1637155_at	0.8219	0.0859	0.2805	0.2120	0.7478	0.0009	0.0932	0.9068	0.4004	0.1169	0.3071	0.1793	-0.2605	0.8331	-0.0563	0.9463	0.2042	0.7299
CG14717	CG14717	1637156_at	0.0677	0.6412	-0.0066	0.9641	0.1652	0.4423	0.1160	0.7764	-0.0881	0.6464	-0.2040	0.1804	-0.1185	0.8541	-0.1337	0.6414	-0.0152	0.9702
---	---	1637157_at	0.1446	0.3882	0.0705	0.5632	0.0988	0.5523	-0.0546	0.9311	0.0437	0.8554	0.0983	0.5949	-0.1088	0.8472	-0.0539	0.8721	0.0548	0.8577
---	---	1637158_at	0.2780	0.1357	-0.0499	0.8200	0.1398	0.5524	-0.2252	0.5626	-0.1706	0.4050	0.0546	0.8075	-0.0023	0.9994	-0.1047	0.8153	-0.1024	0.8108
CG9065	CG9065	1637159_at	-0.2873	0.3542	0.1025	0.5103	0.1328	0.6115	-0.0290	0.9653	-0.6236	0.0047	-0.5945	0.0034	-0.0263	0.9914	-0.2020	0.7435	-0.1757	0.7764
CG14103 /// CG34116 /// D	CG14103 /// CG3	1637160_at	-0.1994	0.3589	-0.0906	0.7972	-0.0244	0.9086	0.0854	0.8310	0.0793	0.6438	-0.0061	0.9745	-0.0986	0.9400	0.1083	0.8427	0.2069	0.6328
myoglianin	Myoglianin	1637161_s_at	-0.5154	0.2224	0.2405	0.5485	0.5129	0.1397	-0.2637	0.7604	-0.9792	0.0166	-0.7156	0.0380	-0.2833	0.8692	-0.0027	0.9994	0.2806	0.7199
CG12054	CG12054	1637162_at	-0.9882	0.0067	-0.0561	0.9028	-0.1934	0.3534	0.0053	0.9952	-0.7814	0.0017	-0.7867	0.0010	0.0811	0.9689	-0.0034	0.9989	-0.0845	0.9103
CG7003 /// DmirCG7003	CG7003	1637163_at	-0.0107	0.9883	-0.5814	0.1997	-0.8121	0.0287	-0.4887	0.3171	0.7297	0.0187	1.2184	0.0012	-0.2765	0.9174	0.1255	0.9325	0.4019	0.6964
Tctp	Translationally cor	1637164_at	0.0641	0.6595	0.1698	0.2290	0.0874	0.5888	0.0532	0.9108	-0.0499	0.7847	-0.1031	0.4709	0.1086	0.8236	-0.0126	0.9703	-0.1211	0.5755
Cchl	CG6022	1637165_at	-0.9142	0.0127	-0.2084	0.4094	-0.0480	0.8508	0.0343	0.9647	-0.9303	0.0015	-0.9646	0.0008	-0.0015	0.9998	-0.0356	0.9617	-0.0341	0.9575
RfC3	RfC3	1637166_at	0.4385	0.0875	-0.2685	0.3751	-0.1421	0.5882	-0.1860	0.8089	0.8210	0.0154	1.0071	0.0037	-0.2686	0.8000	0.0616	0.9301	0.3302	0.4697
---	---	1637167_at	0.1785	0.3876	0.0754	0.4753	0.1862	0.4403	0.0198	0.9852	0.2078	0.3925	0.1880	0.3932	-0.0930	0.8823	0.0166	0.9664	0.1096	0.6770
---	---	1637168_at	0.1044	0.5310	-0.0715	0.5372	0.1476	0.3047	0.0133	0.9847	0.1236	0.4528	0.1103	0.4619	-0.1444	0.7893	-0.0977	0.7064	0.0467	0.8813
Nat1	Nat1	1637169_at	0.0101	0.9832	0.0194	0.9700	-0.1168	0.5211	0.0979	0.8667	0.4512	0.0352	0.3533	0.0587	0.2667	0.8465	0.4237	0.4619	0.1570	0.8260
---	---	1637170_at	0.0137	0.9323	0.0668	0.5372	-0.1659	0.3151	-0.0859	0.8196	-0.0454	0.8023	0.0405	0.8073	-0.0104	0.9914	-0.0393	0.9064	-0.0289	0.9231
---	---	1637171_at	-0.0974	0.6680	0.0395	0.7271	0.1803	0.2708	-0.0139	0.9857	-0.1502	0.4042	-0.1363	0.4039	-0.1317	0.8869	0.0224	0.9688	0.1541	0.6877
mRpS14	mitochondrial ribo	1637172_at	0.0940	0.8034	0.6505	0.1376	0.4013	0.1175	0.0124	0.9916	-0.1002	0.7279	-0.1125	0.6590	0.1649	0.9246	0.4456	0.4421	0.2807	0.6553
CG34010	CG34010	1637173_at	0.0386	0.9029	0.1462	0.4233	0.0741	0.7032	0.1103	0.8925	-0.1362	0.6586	-0.2465	0.3223	0.1660	0.7779	-0.0463	0.9058	-0.2123	0.4126
---	---	1637174_s_at	0.0444	0.8857	0.2450	0.4871	0.3583	0.0481	-0.2910	0.3484	-0.3129	0.0807	-0.0219	0.9201	-0.4296	0.6898	-0.0570	0.9358	0.3726	0.4093
CG12170	CG12170	1637175_at	-0.4799	0.0135	-0.3624	0.5773	-0.0338	0.8725	0.2371	0.3822	-0.4100	0.0155	-0.6470	0.0012	-0.2603	0.8767	-0.4413	0.5140	-0.1810	0.8253
Osi16	Osiris	1637176_at	0.1853	0.2024	0.0518	0.7861	0.0882	0.6600	-0.0754	0.9039	-0.0017	0.9952	0.0737	0.7361	0.0496	0.9499	-0.0213	0.9557	-0.0709	0.8046
CG15032	CG15032	1637177_at	0.5260	0.0401	0.0839	0.4455	0.4649	0.0124	-0.0766	0.8983	-0.1315	0.5301	-0.0549	0.8039	-0.0314	0.9691	-0.1188	0.6009	-0.0873	0.7154
---	---	1637178_at	0.0765	0.7546	0.1009	0.6031	0.1051	0.5225	-0.0298	0.9649	-0.0147	0.9557	0.0151	0.9472	0.0771	0.9206	0.1603	0.5578	0.0831	0.7907
CG14774	CG14774	1637179_at	0.0093	0.9589	0.1016	0.4542	0.1149	0.6496	-0.0008	0.9994	-0.0595	0.8229	-0.0587	0.8056	-0.1551	0.8222	-0.1265	0.6958	0.0286	0.9434
CG17977	CG17977	1637180_at	0.0110	0.9758	0.0861	0.4733	-0.2316	0.2988	-0.3487	0.5038	-0.2232	0.4361	0.1254	0.6617	0.0769	0.9238	0.0382	0.9279	-0.0387	0.9170
---	---	1637181_at	-0.0296	0.8759	-0.0259	0.7968	0.0391	0.8212	0.1301	0.7492	0.0288	0.9068	-0.1013	0.5629	-0.0742	0.8903	-0.0610	0.8165	0.0133	0.9646
CG9503	CG9503	1637182_at	0.2003	0.3834	-0.3257	0.0711	0.0076	0.9729	0.2819	0.4719	0.2962	0.1584	0.0143	0.9573	-0.0172	0.9914	-0.0377	0.9479	-0.0205	0.9701
---	---	1637183_at	0.1084	0.5473	-0.0978	0.7974	0.1698	0.5578	0.0616	0.9451	0.0926	0.7618	0.0310	0.9210	-0.2723	0.6557	-0.1460	0.5823	0.1263	0.6391
CG15029	CG15029	1637184_at	0.0409	0.8726	0.1696	0.5687	0.1613	0.5312	-0.0267	0.9697	-0.1155	0.5536	-0.0888	0.6310	0.0355	0.9717	-0.0455	0.9064	-0.0809	0.7877

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1637185_s_at	0.2192	0.3470	-0.0240	0.8135	0.2829	0.1669	0.0585	0.9412	0.0856	0.7511	0.0271	0.9226	-0.0689	0.9390	-0.0731	0.8461	-0.0043	0.9929
---	---	1637186_at	-0.1572	0.4674	-0.0219	0.8534	-0.0921	0.6603	0.1138	0.8794	0.0089	0.9816	-0.1050	0.6979	-0.0067	0.9946	0.0470	0.8893	0.0537	0.8599
CG9861	CG9861	1637187_at	0.3718	0.0588	0.2779	0.1547	0.1834	0.3182	-0.0060	0.9951	-0.0354	0.8965	-0.0293	0.9027	0.1256	0.8270	-0.0845	0.7707	-0.2102	0.3878
CG30492	CG30492	1637188_s_at	-0.2295	0.2407	0.1425	0.5350	-0.1671	0.2308	-0.6294	0.0492	-0.7604	0.0022	-0.1310	0.4467	-0.1959	0.7423	-0.3722	0.1375	-0.1763	0.4978
CG7146	CG7146	1637189_at	-0.2532	0.3341	-0.2323	0.2783	-0.3439	0.0923	-0.0787	0.8830	0.1322	0.4884	0.2109	0.1944	0.0320	0.9816	0.2802	0.3669	0.2483	0.4457
CG9149	CG9149	1637190_at	-1.5907	0.0024	-1.2256	0.0145	-0.8238	0.1083	-0.1199	0.8841	0.0698	0.8477	0.1897	0.4773	-0.4972	0.7707	0.4591	0.5457	0.9564	0.2094
Coprox	Coproporphyrinog	1637191_at	-0.5395	0.0210	-0.0796	0.4847	0.4470	0.0206	0.1258	0.7605	-0.5071	0.0116	-0.6329	0.0026	-0.3791	0.4511	-0.1454	0.6015	0.2337	0.3818
CG16732	CG16732	1637192_at	-0.0584	0.9395	0.0393	0.7622	-0.1275	0.4008	0.0166	0.9937	-0.3712	0.3971	-0.3878	0.3191	0.1818	0.9291	-0.3548	0.6154	-0.5366	0.4269
eIF3-S8	eIF3-S8	1637193_at	0.3836	0.1153	0.4763	0.0816	0.5227	0.0121	0.3579	0.2506	0.2496	0.1768	-0.1084	0.5532	0.3185	0.6955	0.3468	0.2941	0.0283	0.9525
wmd	wing morphogene	1637194_at	-0.0231	0.9116	-0.0892	0.5690	0.1291	0.6458	-0.0017	0.9985	0.2753	0.1434	0.2769	0.1005	-0.2078	0.8235	0.2353	0.5626	0.4431	0.2691
CG31321	CG31321	1637195_at	0.2390	0.6386	-0.0138	0.9069	0.0243	0.8832	-0.1896	0.7556	-0.2174	0.4224	-0.0278	0.9313	-0.1032	0.9275	-0.3241	0.3738	-0.2209	0.5758
CG41105	CG41105	1637196_at	0.1252	0.6956	0.0802	0.5601	0.0589	0.7733	-0.0119	0.9895	0.0752	0.7643	0.0871	0.6918	0.1114	0.9016	0.1850	0.5994	0.0735	0.8668
neb	tiovivo	1637197_at	0.9129	0.0103	-0.2870	0.3298	-0.2858	0.2550	0.1425	0.7596	0.4137	0.0446	0.2712	0.1278	0.0377	0.9816	-0.7582	0.0664	-0.7958	0.0741
CG10345	CG10345	1637198_at	0.1486	0.8021	0.0622	0.6915	-0.6294	0.0423	-0.4477	0.2917	0.0071	0.9846	0.4548	0.0471	0.0674	0.9816	-0.3102	0.6830	-0.3776	0.6056
---	---	1637199_at	0.1291	0.4978	-0.0087	0.9771	0.0550	0.7691	-0.0808	0.8817	0.0563	0.8044	0.1371	0.4306	-0.1586	0.8395	-0.1541	0.6636	0.0045	0.9937
---	---	1637200_at	0.1368	0.3339	-0.0300	0.7680	0.0879	0.6124	0.0026	0.9956	0.0200	0.9192	0.0175	0.9193	0.0010	0.9994	-0.0146	0.9593	-0.0156	0.9491
tinc	tincar	1637201_s_at	0.1200	0.4772	-0.0508	0.8702	0.1316	0.5320	0.1621	0.7922	0.1960	0.4511	0.0340	0.9105	0.0877	0.9238	0.0556	0.9054	-0.0322	0.9401
Med	Medea	1637202_a_at	0.1021	0.7145	-0.0122	0.9147	-0.2588	0.3756	0.0229	0.9790	0.2226	0.3074	0.1997	0.3092	-0.0471	0.9741	-0.1215	0.7833	-0.0743	0.8779
rev7	rev7	1637203_at	-0.3939	0.2033	-0.5432	0.0588	-0.3350	0.2144	0.0693	0.9098	0.0781	0.7306	0.0088	0.9710	-0.1445	0.8692	-0.1332	0.7459	0.0114	0.9846
Ptp99A	Protein tyrosine pl	1637204_at	-1.4806	0.0041	-2.3761	0.0200	-2.4973	0.0000	0.3273	0.4806	0.7794	0.0078	0.4521	0.0474	0.3633	0.8157	0.0527	0.9608	-0.3105	0.6567
CG11608	CG11608	1637205_at	0.2048	0.4448	0.1724	0.2977	0.0614	0.7625	-0.1864	0.5756	-0.1663	0.3376	0.0201	0.9235	-0.0355	0.9816	-0.1163	0.7782	-0.0808	0.8524
CG8405	CG8405	1637206_at	0.0616	0.9441	0.4368	0.2617	0.4019	0.0567	-0.0506	0.9744	-0.0802	0.8738	-0.0296	0.9511	-0.0879	0.9826	0.3443	0.7561	0.4322	0.6700
---	---	1637207_at	0.2202	0.1885	0.0304	0.7893	-0.0541	0.7668	-0.0362	0.9445	0.0673	0.6939	0.1036	0.4628	0.0364	0.9589	0.0097	0.9789	-0.0267	0.9234
---	---	1637208_at	0.1076	0.5544	0.0368	0.8266	0.1150	0.6287	-0.0634	0.9110	0.0171	0.9495	0.0806	0.6662	-0.0364	0.9739	0.0412	0.9231	0.0776	0.8202
CG11112	CG11112	1637209_at	0.1600	0.6045	0.1590	0.2348	0.0882	0.6611	-0.0356	0.9673	-0.2791	0.2202	-0.2435	0.2334	0.1189	0.8775	0.0670	0.8760	-0.0519	0.9004
---	---	1637210_at	0.2510	0.2728	0.0132	0.9591	0.0267	0.9094	-0.0622	0.9413	0.0592	0.8500	0.1214	0.6213	-0.0258	0.9816	0.0038	0.9941	0.0297	0.9330
CG13045	CG13045	1637211_at	0.1939	0.2524	0.1304	0.5902	0.3795	0.0710	0.0098	0.9931	0.0158	0.9623	0.0060	0.9831	-0.0863	0.9467	0.0568	0.9275	0.1432	0.7527
CG33275	CG33275	1637212_at	-0.5795	0.0200	-0.2180	0.3505	-0.1183	0.5044	0.1363	0.8403	-0.0741	0.8126	-0.2103	0.3668	-0.1311	0.8192	0.0636	0.8406	0.1947	0.4148
CG31115	CG31115	1637213_at	0.1465	0.3781	0.0010	0.9999	0.0828	0.6311	0.0142	0.9866	0.0269	0.9252	0.0127	0.9606	-0.0991	0.8906	-0.1438	0.6226	-0.0448	0.9056
scra	Anillin	1637214_a_at	0.2046	0.7043	-0.0159	0.9880	-1.1903	0.0160	-0.5885	0.4800	0.5412	0.2250	1.1297	0.0123	0.1844	0.9514	-0.0436	0.9787	-0.2281	0.8443
CG10881	CG10881	1637215_at	0.3430	0.1653	0.2422	0.3698	0.2023	0.2002	-0.0364	0.9558	-0.1868	0.2922	-0.1504	0.3512	0.0997	0.8972	0.0224	0.9624	-0.0774	0.8307
CG7328	CG7328	1637216_at	0.0063	0.9891	-0.3662	0.2150	-0.3424	0.0474	-0.1171	0.8591	0.2609	0.2742	0.3779	0.0767	-0.3031	0.7779	-0.3780	0.4147	-0.0748	0.9077
ecd	ecdysoneless	1637217_at	-0.7617	0.0686	-0.5954	0.1104	-0.1350	0.4713	0.4240	0.1279	-0.0708	0.7157	-0.4948	0.0060	0.0004	0.9999	0.2783	0.6146	0.2779	0.6168
---	---	1637218_at	-0.1580	0.4617	0.0000	1.0000	-0.0010	0.9969	0.1581	0.7579	-0.1246	0.6068	-0.2827	0.1502	0.0014	0.9994	-0.1359	0.5966	-0.1372	0.5963
CG1690	CG1690	1637219_at	0.1826	0.2833	0.0678	0.6951	0.0689	0.7334	-0.0304	0.9659	0.0157	0.9553	0.0460	0.8296	-0.0753	0.9101	0.0143	0.9714	0.0896	0.7440
CG11984	CG11984	1637220_s_at	-0.7506	0.0406	-0.6814	0.2339	-0.9582	0.0028	-0.2319	0.4706	-0.2420	0.1604	-0.0101	0.9632	0.0336	0.9913	-0.2045	0.8053	-0.2381	0.7492
CG5705	CG5705	1637221_at	-0.4860	0.0527	0.0726	0.4958	0.2199	0.4000	-0.2231	0.5307	-0.3000	0.1048	-0.0769	0.6934	-0.3435	0.7220	0.3557	0.3759	0.6992	0.1192
---	---	1637222_at	0.1083	0.4540	0.1447	0.3741	0.3711	0.0807	0.0171	0.9774	-0.0651	0.6999	-0.0822	0.5719	-0.1366	0.8653	-0.0630	0.8958	0.0736	0.8618
CG8929	CG8929	1637223_s_at	-0.0225	0.9395	-0.4710	0.3844	-0.5400	0.0146	-0.0704	0.9068	0.4244	0.0357	0.4949	0.0119	0.0099	0.9967	-0.0902	0.9263	-0.1001	0.9057
CG13905	CG13905	1637224_at	-0.0260	0.9761	-0.0474	0.6725	0.0885	0.6425	0.1390	0.9187	-0.3363	0.4395	-0.4753	0.2048	-0.0219	0.9911	-0.2393	0.5644	-0.2175	0.6109
bnl	fibroblast growth f	1637225_at	-0.0011	0.9970	-0.2103	0.0759	-0.2700	0.1314	-0.0753	0.8979	-0.0681	0.7688	0.0072	0.9761	-0.0449	0.9635	-0.3584	0.1658	-0.3136	0.2557
CG11885	CG11885	1637226_at	0.3513	0.2747	-0.2112	0.4734	0.1596	0.4648	0.5223	0.1109	0.7476	0.0031	0.2253	0.1945	0.0669	0.9737	0.1070	0.8821	0.0401	0.9549
---	---	1637227_at	0.0943	0.5345	0.1625	0.3269	-0.0199	0.9171	-0.2706	0.4713	-0.1782	0.3901	0.0924	0.6590	0.1569	0.7707	0.0735	0.8018	-0.0835	0.7520
CG10663	CG10663	1637228_at	-1.0814	0.0038	-0.5063	0.2211	-1.3157	0.0005	-0.4608	0.4239	-0.2787	0.3941	0.1822	0.5632	0.2703	0.7726	0.2271	0.6011	-0.0433	0.9394
CG11796	CG11796	1637229_a_at	0.6390	0.6658	0.0186	0.9176	0.0753	0.7530	-0.0032	0.9988	-0.5574	0.1803	-0.5542	0.1367	-0.1923	0.9816	-1.2158	0.5214	-1.0234	0.6036
---	---	1637230_at	0.0864	0.7241	0.2144	0.1746	0.1220	0.4245	0.0583	0.9413	0.1086	0.6743	0.0504	0.8482	0.2486	0.7230	0.2843	0.3340	0.0358	0.9330
---	---	1637231_s_at	-0.0039	0.9847	0.0554	0.6506	-0.0589	0.8088	-0.0400	0.9592	-0.1240	0.5756	-0.0840	0.6962	0.1282	0.8235	0.1222	0.6380	-0.0060	0.9874
CG17150 /// DmirCG17150	CG17150	1637232_at	0.1427	0.4184	0.0193	0.9115	0.0282	0.9089	0.1504	0.7130	0.1886	0.3155	0.0383	0.8599	0.0173	0.9894	-0.0112	0.9829	-0.0285	0.9410
CG13229	CG13229	1637233_at	-0.0912	0.6787	0.1416	0.5757	0.0510	0.8272	-0.1053	0.9132	-0.2992	0.3334	-0.1939	0.5098	0.0358	0.9751	0.0733	0.8487	0.0376	0.9231
---	---	1637234_s_at	0.1118	0.5329	0.0019	0.9940	0.2459	0.1317	-0.0626	0.9098	-0.0595	0.7793	0.0031	0.9887	-0.1516	0.8191	-0.2823	0.2811	-0.1308	0.6550
---	---	1637235_at	0.0127	0.9405	0.0141	0.8915	0.0875	0.5905	0.0169	0.9764	-0.0171	0.9318	-0.0340	0.8338	0.0672	0.9142	0.0512	0.8685	-0.0160	0.9601
CG41136	CG41136	1637236_at	-0.6116	0.0162	-0.6525	0.0137	-0.9456	0.0036	0.1557	0.8748	0.5921	0.0861	0.4364	0.1545	-0.0651	0.9550	-0.1621	0.6614	-0.0970	0.8178
---	---	1637237_at	-0.0438	0.8312	0.0000	1.0000	-0.0205	0.9098	-0.0784	0.8949	0.0255	0.9267	0.1039	0.6002	-0.0604	0.9112	0.0501	0.8481	0.1105	0.5910
---	---	1637238_at	-0.0436	0.7849	0.1771	0.4483	0.1727	0.5866	-0.1459	0.8159	-0.1504	0.5718	-0.0044	0.9883	-0.0415	0.9756	0.0605	0.9054	0.1020	0.8000
CG34034	CG34034	1637239_s_at	-1.4860	0.0010	0.3526	0.3471	-0.5695	0.1803	-0.9830	0.2032	-2.0764	0.0017	-1.0934	0.0167	0.0694	0.9666	0.0046	0.9969	-0.0649	0.9158
Atg7	Autophagy-specifi	1637240_a_at	0.4162	0.0461	0.6887	0.0389	0.6186	0.0050	-0.0806	0.8594	0.0350	0.8718	0.1156	0.4549	-0.0708	0.9424	0.2520	0.4006	0.3228	0.2993
---	---	1637241_at	0.1324	0.4657	0.1514	0.4195	-0.1009	0.6128	-0.2524	0.6564	-0.2483	0.3726	0.0040	0.9906	0.0325	0.9752	-0.0223	0.9587	-0.0548	0.8751
CG34353	CG12274	1637242_at	0.0663	0.6875	0.1210	0.3943	-0.0919	0.5977	-0.0081	0.9943	0.0529	0.8656	0.0610	0.8206	0.0936	0.8729	0.0707	0.8130	-0.0229	0.9434
CG31498	CG31498	1637243_at	0.0572	0.7262	-0.0141	0.8930	0.0342	0.8338	-0.0667	0.8698	-0.1135	0.4429	-0.0468	0.7650	-0.0834	0.8841	-0.2475	0.2385	-0.1641	0.4648
APC4	Anaphase Promot	1637244_at	0.1009	0.6857	0.1482	0.3446	0.5856	0.0089	-0.0574	0.9196	0.0157	0.9518	0.0730	0.6893	-0.4077	0.5757	0.1071	0.7971	0.5148	0.1443
CG15729	CG15729	1637245_at	-0.0234	0.9197	-0.1261	0.4572	-0.0855	0.6145	-0.0020	0.9985	0.0139	0.9690	0.0159	0.9571	-0.0228	0.9775	-0.0284	0.9263	-0.0056	0.9852
CG32627	CG32627	1637246_a_at	-2.5881	0.0015	-1.8805	0.1413	-3.0654	0.0001	-1.1518	0.1231	-1.8922	0.0021	-0.7404	0.0687	0.0347	0.9943	-1.0493	0.3404	-1.0840	0.3513
---	---	1637247_at	-0.0114	0.9507	0.0533	0.8207	-0.0376	0.8147	-0.0845	0.8156	-0.0251	0.8973	0.0594	0.6913	-0.0643	0.9305	-0.0677	0.8304	-0.0034	0.9936
CG30194 /// DereCG30194	CG30194	1637248_at	0.3642	0.0819	0.0926	0.4922	0.0338	0.8403	-0.0280	0.9763	0.1019	0.7056	0.1299	0.5769	0.1013	0.8541	-0.0398	0.9085	-0.1411	0.5579
CG10721	CG10721	1637249_at	-1.0505	0.0008	0.2970	0.3293	0.3328	0.0711	-0.2814	0.4062	-1.5709	0.0002	-1.2895	0.0002	-0.6297	0.2553	-0.0191	0.9670	0.6106	0.0679
CG17816	CG17816	1637250_at	0.3525	0.1495	0.2634	0.3629	0.1295	0.4944	-0.1019	0.8716	0.0440	0.8810	0.1459	0.4877	-0.0516	0.9677	0.0471	0.9296	0.0987	0.8138
Sucb	Sucb	1637251_a_at	-0.0061	0.9890	0.9191	0.0348	0.7890	0.0019	0.0047	0.9956	-0.5178	0.0504	-0.5225	0.0314	0.1849	0.8541	0.4669	0.2430	0.2820	0.5151
---	---	1637252_s_at	0.0676	0.6861	0.2055	0.2362	0.1012	0.5672	-0.1368	0.7121	-0.0449	0.8356	0.0919	0.5860	-0.0676	0.9449	0.0077	0.9911	0.0753	0.8418
CG17570	CG17570	1637253_s_at	0.3103	0.1692	0.3092	0.0792	-0.0323	0.8576	-0.2066	0.4504	-0.1949	0.1873	0.0117	0.9503	0.2745	0.7644	-0.1415	0.7572	-0.4160	0.2924

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
spdo	sanpodo	1637254_at	0.6739	0.1622	1.1362	0.1380	1.0462	0.0007	-0.0841	0.9247	-0.0612	0.8612	0.0228	0.9461	0.3005	0.8379	0.5835	0.3236	0.2830	0.6677
alpha-Est7	fragment H	1637255_a_at	1.3338	0.0288	1.4190	0.2161	1.5595	0.0002	-0.1817	0.8990	-1.0774	0.0290	-0.8958	0.0388	-0.3820	0.8940	-1.0873	0.2985	-0.7052	0.5332
CG34349	CG11819	1637256_at	-0.6875	0.0130	-1.6954	0.0086	-1.8150	0.0021	0.2867	0.3596	0.8974	0.0010	0.6107	0.0031	0.2594	0.8930	-0.0324	0.9807	-0.2918	0.7205
CG9009	CG9009	1637257_at	-1.6928	0.0031	-1.9874	0.0312	-1.0074	0.0024	-0.2428	0.5317	-1.5358	0.0002	-1.2930	0.0002	-1.0174	0.4981	-1.6444	0.0446	-0.6270	0.3939
---	---	1637258_at	0.4334	0.0230	0.1144	0.6151	0.0088	0.9732	0.0040	0.9956	0.1002	0.6825	0.0962	0.6659	0.1575	0.7287	-0.0864	0.6945	-0.2439	0.2263
---	---	1637259_at	0.0275	0.9413	0.2023	0.2164	-0.0351	0.8782	-0.1598	0.7845	-0.0210	0.9532	0.1388	0.5597	0.0967	0.8689	0.1014	0.6969	0.0047	0.9903
CG6066	CG6066	1637260_at	0.0411	0.8988	0.3063	0.1194	0.1489	0.5516	-0.0156	0.9866	-0.0045	0.9898	0.0111	0.9689	-0.0278	0.9816	0.1479	0.5897	0.1757	0.5153
Cp7Fc	Chorion protein c	1637261_at	0.0531	0.8166	0.0111	0.9686	-0.1691	0.3481	-0.2629	0.6338	-0.2379	0.3919	0.0250	0.9408	-0.1703	0.8714	-0.0928	0.8764	0.0775	0.8923
CG7698	CG7698	1637262_at	-0.1216	0.3841	-0.2457	0.3443	-0.1762	0.2151	-0.1316	0.6972	0.3714	0.0252	0.5031	0.0040	-0.1934	0.7423	0.1674	0.5140	0.3607	0.1691
mRpL45	mitochondrial ribo	1637263_at	-0.4981	0.0908	-0.8278	0.0306	-1.0802	0.0009	-0.0445	0.9491	-0.1298	0.5313	-0.0854	0.6738	-0.0747	0.9555	-0.2889	0.4655	-0.2142	0.6117
ncd	non-claret disjunct	1637264_at	0.1042	0.8301	-0.0337	0.9596	-0.1453	0.6758	-0.3522	0.5735	0.0163	0.9739	0.3685	0.1960	-0.3330	0.8716	-0.1456	0.9075	0.1874	0.8617
CG7550	CG7550	1637265_at	-0.1656	0.3508	-0.6203	0.0092	-0.5338	0.0063	0.2274	0.4455	0.7071	0.0018	0.4797	0.0060	0.0230	0.9816	0.0710	0.7978	0.0480	0.8694
---	---	1637266_at	-0.1429	0.5769	0.2217	0.2281	0.2062	0.2897	-0.0640	0.9102	-0.1947	0.2822	-0.1307	0.4397	-0.0060	0.9964	0.1615	0.6146	0.1675	0.6031
CG8138	CG8138	1637267_at	0.2128	0.2639	-0.0072	0.9493	-0.0831	0.5872	-0.0659	0.9185	0.1423	0.4936	0.2082	0.2409	-0.0738	0.8966	-0.1524	0.4806	-0.0786	0.7484
---	---	1637268_at	0.1561	0.3487	0.0988	0.4866	0.0960	0.5492	0.0359	0.9606	0.0215	0.9385	-0.0144	0.9534	0.2379	0.6152	0.0682	0.7854	-0.1696	0.4078
CG6310	CG6310	1637269_at	-2.6347	0.0090	-3.8676	0.0047	-3.8706	0.0000	-0.0332	0.9540	0.0543	0.7759	0.0875	0.5785	-0.0241	0.9952	-1.0293	0.2938	-1.0052	0.3293
D2R	Dopamine 2-like r	1637270_at	0.3650	0.1006	0.1302	0.5452	0.3159	0.1367	0.0157	0.9866	0.0724	0.7962	0.0568	0.8276	0.0080	0.9939	0.0750	0.8145	0.0669	0.8287
CG14605	CG14605	1637271_a_at	-0.0280	0.8700	-0.2900	0.2259	-0.0987	0.5442	-0.0312	0.9838	0.0446	0.9298	0.0758	0.8571	-0.1399	0.8486	-0.0091	0.9887	0.1308	0.7022
Obp58c	Odorant-binding p	1637272_at	0.0817	0.6886	0.0846	0.7284	0.1623	0.3723	-0.0937	0.8133	-0.1368	0.3940	-0.0432	0.8059	-0.1451	0.8033	-0.0662	0.8389	0.0790	0.7838
---	---	1637273_at	0.0507	0.8242	0.0921	0.5678	0.1732	0.2628	0.0680	0.9308	-0.0716	0.7988	-0.1397	0.5320	-0.1228	0.8425	-0.0648	0.8519	0.0580	0.8618
CG14488	CG14488	1637274_at	0.0453	0.8246	-0.0247	0.8201	0.2322	0.1648	0.1200	0.7351	0.0678	0.7098	-0.0521	0.7630	-0.0312	0.9742	-0.0547	0.8754	-0.0234	0.9452
CG13335 /// DyakCG13335	CG13335	1637275_a_at	2.1264	0.0098	0.4610	0.0177	1.3034	0.0061	0.7719	0.5128	1.4936	0.0245	0.7217	0.1939	0.0836	0.8692	-0.1325	0.5252	-0.2161	0.2964
Pen	Importin-a2	1637276_at	0.6879	0.7018	-1.8535	0.4639	-2.1688	0.0606	-0.5349	0.5233	2.8615	0.0004	3.3964	0.0001	-0.4141	0.9774	0.0462	0.9950	0.4603	0.9224
CG34130	CG34130	1637277_at	0.2546	0.4058	0.1612	0.3624	0.3507	0.0608	0.1139	0.8889	0.0845	0.8009	-0.0293	0.9302	0.0976	0.8692	0.1013	0.7020	0.0037	0.9924
tin	muscle-specific-ho	1637278_at	-0.0020	0.9934	0.0962	0.4206	0.2774	0.2186	-0.1453	0.7327	-0.3647	0.0587	-0.2194	0.1948	-0.1240	0.8472	-0.0503	0.9018	0.0737	0.8259
CG12951	CG12951	1637279_at	0.3507	0.0555	0.0208	0.8993	0.0580	0.7532	-0.0767	0.8738	-0.0874	0.6442	-0.0107	0.9597	-0.0779	0.9092	-0.2043	0.3941	-0.1264	0.6294
LpR1	yolkless-like	1637280_at	-1.1699	0.0210	-1.1135	0.3304	-1.6068	0.0093	-0.3428	0.7117	-0.7799	0.0670	-0.4371	0.2473	0.1395	0.9764	-0.8200	0.4753	-0.9595	0.4037
---	---	1637281_at	0.0663	0.7513	0.1230	0.5275	0.1076	0.4683	0.0773	0.8736	0.0066	0.9800	-0.0708	0.6900	0.0854	0.9168	0.0443	0.9198	-0.0412	0.9167
CG6171 /// DyakCG6171	Anon-becker2 /// (	1637282_at	-0.9047	0.0114	-0.0465	0.6897	0.3292	0.0987	0.2281	0.7014	-0.4297	0.1143	-0.6577	0.0151	-0.1898	0.8009	0.3868	0.2018	0.5766	0.1011
---	---	1637283_at	-0.0654	0.8295	0.0711	0.6645	0.0748	0.6935	-0.1009	0.8605	-0.1116	0.6256	-0.0107	0.9674	-0.0529	0.9717	0.0534	0.9276	0.1063	0.8183
Rlip	Rlip	1637284_at	-1.0221	0.0028	-0.8592	0.0557	-1.1754	0.0002	-0.3503	0.3645	0.2067	0.3564	0.5571	0.0122	-0.2121	0.8202	0.1216	0.8043	0.3337	0.3921
slmo	Kisir	1637285_s_at	-0.1261	0.6088	0.3158	0.2475	0.6857	0.0169	0.0343	0.9774	-0.6862	0.0283	-0.7204	0.0146	-0.2312	0.7779	-0.2104	0.5762	0.0208	0.9696
raw	cyrano	1637286_a_at	-0.0397	0.8809	0.0930	0.7631	-0.1822	0.2099	0.1363	0.7138	0.3785	0.0326	0.2422	0.1088	0.4311	0.5854	0.4650	0.1744	0.0339	0.9452
CG13260	CG13260	1637287_at	-0.0145	0.9688	0.0777	0.4475	-0.1202	0.4982	-0.2083	0.7441	-0.0531	0.8887	0.1552	0.5785	0.0989	0.8510	-0.0220	0.9504	-0.1208	0.6052
CG7548	CG7548	1637288_at	-0.0535	0.8201	-0.0698	0.6008	0.1595	0.4419	-0.0325	0.9712	-0.0230	0.9456	0.0095	0.9733	-0.0794	0.9503	0.1348	0.7707	0.2143	0.5995
---	---	1637289_at	0.1401	0.5820	-0.2170	0.2596	0.0339	0.8421	0.2105	0.5332	0.1977	0.2643	-0.0128	0.9547	0.0514	0.9683	-0.1184	0.7779	-0.1698	0.6431
Dsk	sulfakinin	1637290_at	0.2018	0.4078	0.1991	0.2635	0.2685	0.1002	0.1062	0.8073	0.2055	0.2349	0.0993	0.5617	0.0427	0.9740	0.1555	0.6593	0.1127	0.7621
CG18031	CG18031	1637291_at	-0.3056	0.2670	0.0367	0.9201	0.4122	0.1548	0.4057	0.4316	-0.1808	0.5531	-0.5864	0.0282	-0.1731	0.9237	-0.1791	0.8166	-0.0060	0.9953
Esp	Epidermal stripes	1637292_at	-2.4741	0.0005	-2.7185	0.0424	-3.0720	0.0000	-0.3266	0.1613	-0.1673	0.2370	0.1593	0.2076	0.0387	0.9928	-0.4092	0.7093	-0.4478	0.6728
---	---	1637293_at	-0.3158	0.3585	-0.0507	0.7123	0.0502	0.7868	0.0823	0.9253	-0.0246	0.9505	-0.1069	0.6999	-0.0991	0.9438	0.0426	0.9505	0.1417	0.7777
CG4641	CG4641	1637294_at	-0.2244	0.3343	-0.0101	0.9239	-0.1596	0.4806	-0.0714	0.9378	-0.2167	0.4352	-0.1453	0.5872	-0.0614	0.9306	0.0728	0.8028	0.1342	0.5807
---	---	1637295_at	0.1184	0.4405	0.3583	0.0575	0.4085	0.0370	-0.0758	0.8863	-0.0841	0.6813	-0.0082	0.9710	-0.1143	0.8270	-0.0224	0.9498	0.0919	0.7075
---	---	1637296_at	0.5393	0.0173	0.2017	0.1454	0.2779	0.2657	0.3696	0.1545	0.3844	0.0227	0.0149	0.9393	0.0374	0.9776	-0.0234	0.9647	-0.0609	0.8895
---	---	1637297_at	0.2385	0.2853	-0.0060	0.9642	0.1231	0.6688	0.3600	0.3521	0.3597	0.1040	-0.0003	0.9994	0.0662	0.9589	-0.1270	0.7738	-0.1932	0.6196
CG5567	CG5567	1637298_at	-0.6608	0.0113	-0.7118	0.0456	-0.7846	0.0030	-0.2036	0.6471	-0.2959	0.1670	-0.0924	0.6821	0.0781	0.9555	-0.3154	0.4380	-0.3935	0.3476
CG6928	CG6928	1637299_s_at	0.0195	0.9293	0.7520	0.0261	0.4003	0.1779	-0.4035	0.1897	0.0447	0.8529	0.4482	0.0152	-0.0941	0.9589	0.8181	0.1151	0.9122	0.1110
DyakCG8229	CG8229	1637300_at	0.0451	0.8098	0.5969	0.0978	0.3311	0.2531	0.0465	0.9734	-0.6835	0.0519	-0.7300	0.0258	0.2064	0.8297	-0.1188	0.8212	-0.3252	0.4301
slo	slowpoke	1637301_a_at	-2.1594	0.0019	-1.6304	0.0223	-1.6507	0.0007	-0.0867	0.8950	-0.3857	0.0793	-0.2990	0.1242	-0.3787	0.7886	-0.3950	0.5251	-0.0163	0.9865
flfl	falafel	1637302_s_at	0.1106	0.6244	0.0091	0.9864	-0.6127	0.0238	-0.0494	0.9314	0.4408	0.0181	0.4902	0.0069	0.4388	0.7230	0.3353	0.5469	-0.1035	0.8880
CG15494	CG15494	1637303_at	-0.0732	0.7741	0.1460	0.2570	0.1031	0.4935	-0.1057	0.8009	-0.2246	0.1807	-0.1189	0.4561	0.1424	0.8461	0.2075	0.5037	0.0651	0.8719
CG14911	CG14911	1637304_at	0.1146	0.6441	-0.0117	0.9108	-0.0235	0.9207	0.0883	0.8698	0.0979	0.6444	0.0096	0.9679	0.0382	0.9816	-0.0902	0.8678	-0.1284	0.7764
CG14471	CG14471	1637305_a_at	0.2734	0.2019	0.1198	0.3398	0.1270	0.4448	-0.0914	0.8535	0.0302	0.9026	0.1216	0.4754	-0.0041	0.9967	-0.1054	0.7200	-0.1012	0.7307
CG31720	CG31720	1637306_at	0.6120	0.6510	0.0107	0.9208	0.1242	0.5123	0.0816	0.9658	-0.2075	0.7230	-0.2892	0.5604	-0.1261	0.9816	-0.8143	0.5464	-0.6882	0.6214
CG31694	CG31694	1637307_at	0.2651	0.0945	0.7453	0.0242	0.7065	0.0032	-0.0279	0.9624	0.0023	0.9916	0.0302	0.8702	0.0820	0.9238	0.5474	0.0694	0.4654	0.1300
CG11417	CG11417	1637308_at	-0.1563	0.7161	-0.5056	0.0332	-0.1665	0.5015	0.2935	0.3698	0.8879	0.0012	0.5943	0.0041	-0.2292	0.8961	0.3963	0.5640	0.6255	0.3543
Cyp12e1	Cyp12e1	1637309_a_at	-0.0178	0.9630	-0.7206	0.0707	-1.0338	0.0007	-0.6228	0.2500	-0.6032	0.0636	0.0196	0.9611	-0.2898	0.7673	-1.3479	0.0204	-1.0582	0.0457
---	---	1637310_at	-0.2268	0.2358	-0.6589	0.0621	-0.5448	0.0071	0.1700	0.7576	0.5756	0.0232	0.4056	0.0592	-0.0214	0.9768	0.0096	0.9741	0.0310	0.8977
pburs	bursicon beta	1637311_at	0.0684	0.7108	0.0782	0.5994	-0.0389	0.8055	0.0752	0.8709	0.0179	0.9385	-0.0573	0.7428	0.2593	0.6587	0.0453	0.9009	-0.2140	0.3825
br	broad complex	1637312_a_at	-0.3679	0.0827	-1.1877	0.0500	-0.8311	0.0542	0.7342	0.3817	1.2956	0.0142	0.5613	0.1847	-0.0003	0.9999	-0.1336	0.7101	-0.1334	0.7087
CG32595	CG32595	1637313_at	-0.0420	0.8745	-0.0035	0.9817	-0.0070	0.9801	-0.1665	0.7255	-0.2480	0.2455	-0.0815	0.7198	-0.0209	0.9898	-0.1986	0.5896	-0.1777	0.6336
tub	tube	1637314_at	0.0187	0.9528	0.2253	0.2902	-0.1333	0.6218	-0.1750	0.7931	-0.2599	0.3489	-0.0849	0.7796	0.1916	0.8122	-0.0872	0.8478	-0.2788	0.4107
CG14065	CG14065	1637315_at	0.5299	0.0745	0.1047	0.5561	0.2593	0.2260	0.1593	0.7906	0.3986	0.1031	0.2393	0.2791	0.1057	0.9514	-0.0517	0.9505	-0.1574	0.8038
CG31697	CG31697	1637316_at	-0.1502	0.3883	0.0798	0.5495	-0.3709	0.1797	-0.2704	0.4270	0.0023	0.9935	0.2726	0.1036	0.1598	0.7997	-0.0136	0.9766	-0.1733	0.5299
Bub3	Bub3	1637317_at	0.0540	0.8430	0.0797	0.6688	0.2583	0.1561	0.0989	0.8377	0.1531	0.4119	0.0543	0.7885	-0.1148	0.8609	0.0267	0.9504	0.1415	0.6189
CG9270	CG9270	1637318_at	0.0492	0.8409	-0.0570	0.5564	-0.0665	0.7001	0.0747	0.8789	0.2750	0.0992	0.2004	0.1783	-0.0316	0.9689	0.0187	0.9567	0.0503	0.8542
Fancl	Fancl	1637319_at	-0.0595	0.9071	0.0612	0.5906	-0.4036	0.0461	-0.5566	0.3625	-0.3081	0.3875	0.2485	0.4490	-0.1744	0.8874	-0.1512	0.7941	0.0232	0.9746
CG18472	CG18472	1637320_at	0.0327	0.8692	0.1317	0.2320	0.2746	0.1720	0.1031	0.7939	0.0084	0.9733	-0.0946	0.5496	0.0018	0.9990	0.0213	0.9566	0.0195	0.9538
nAcRbeta-64B	AChR protein of D	1637321_at	-0.3397	0.0611	-0.7534	0.0145	-0.9044	0.0334	-0.0338	0.9842	0.6289	0.1108	0.6626	0.0643	-0.1687	0.7215	-0.0484	0.8593	0.1204	0.5637
pyd	tamou	1637322_at	1.9185	0.0166	-0.2175	0.7187	-0.8396	0.1086	-0.1915	0.9093	2.1033	0.0028	2.2947	0.0012	0.5498	0.8270	0.0285	0.9905	-0.5212	0.6469

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1637323_at	0.2080	0.2499	0.0190	0.8646	0.1969	0.2609	0.0491	0.9300	0.0582	0.7675	0.0091	0.9654	-0.1827	0.7697	-0.0497	0.9019	0.1330	0.6410
---	---	1637324_at	-0.0258	0.8831	0.0408	0.8535	0.0040	0.9878	-0.1382	0.7897	-0.1622	0.4633	-0.0240	0.9262	-0.0513	0.9677	-0.1224	0.7593	-0.0711	0.8745
Mst57Da	Male-specific RNA	1637325_at	-0.0582	0.7632	0.1678	0.3382	-0.0241	0.9036	-0.1892	0.5968	-0.3219	0.0771	-0.1327	0.4349	0.0737	0.8882	0.0708	0.7663	-0.0029	0.9933
CG15773	CG15773	1637326_at	0.5237	0.0224	0.7253	0.0118	0.2449	0.1205	-0.0310	0.9722	0.0936	0.7176	0.1246	0.5733	0.1597	0.7768	0.1504	0.5531	-0.0093	0.9821
CG11376	CG11376	1637327_at	0.2541	0.2662	0.0700	0.8815	-0.1438	0.5140	0.1226	0.7654	0.5011	0.0115	0.3785	0.0239	0.3207	0.7848	0.3230	0.5404	0.0023	0.9979
CG14782	CG14782	1637328_at	-0.3699	0.1293	-0.3855	0.4376	-0.3879	0.1535	0.1568	0.5724	0.1090	0.4614	-0.0479	0.7573	0.1476	0.9405	0.0413	0.9671	-0.1063	0.8997
tho2	tho2	1637329_a_at	-0.0505	0.8769	0.1402	0.6885	0.2120	0.2016	0.0155	0.9883	0.1923	0.4561	0.1768	0.4498	0.0411	0.9816	0.3816	0.3476	0.3404	0.4250
---	---	1637330_at	0.2452	0.2324	-0.0414	0.7019	0.1467	0.4014	0.1288	0.7138	0.1721	0.2860	0.0433	0.8102	-0.0426	0.9646	-0.0377	0.9265	0.0048	0.9915
---	---	1637331_at	0.2321	0.2640	-0.0274	0.9304	0.3483	0.1284	0.0275	0.9770	-0.1427	0.5702	-0.1702	0.4367	-0.1143	0.8889	-0.1358	0.6954	-0.0215	0.9622
imd	Immunodeficiency	1637332_at	0.2852	0.0830	-0.6500	0.0162	-0.2081	0.2876	0.2943	0.1883	0.5225	0.0031	0.2282	0.0627	-0.1582	0.7925	-0.4788	0.0743	-0.3206	0.2231
---	---	1637333_at	0.0431	0.9046	0.1306	0.4578	-0.1042	0.6625	-0.0092	0.9924	-0.0112	0.9715	-0.0020	0.9938	0.1120	0.9280	0.1739	0.7038	0.0619	0.9118
---	---	1637334_at	0.0419	0.8661	0.0599	0.7179	0.1352	0.4901	0.0180	0.9860	-0.0863	0.7672	-0.1043	0.6819	-0.0053	0.9962	-0.0381	0.9192	-0.0328	0.9215
fln	flightin	1637335_at	0.1935	0.3446	-0.1010	0.7236	0.1153	0.5922	0.0261	0.9808	-0.0379	0.9189	-0.0640	0.8326	-0.2107	0.7506	-0.1782	0.5488	0.0326	0.9347
CG5377	CG5377	1637336_at	-0.2725	0.1243	-0.2182	0.2612	0.1433	0.3873	0.0101	0.9921	0.0139	0.9652	0.0039	0.9880	-0.3503	0.3734	-0.0190	0.9577	0.3313	0.1512
---	---	1637337_at	0.2132	0.3766	0.2116	0.1114	-0.1654	0.3393	-0.0496	0.9627	0.1495	0.6376	0.1991	0.4577	0.1392	0.7644	0.0518	0.8450	-0.0874	0.6867
CG12517	CG12517	1637338_at	-0.6445	0.3387	1.9195	0.1253	0.4362	0.4760	-0.2174	0.9242	0.5895	0.4024	0.8068	0.1867	1.3189	0.6749	3.2679	0.0365	1.9490	0.1588
Ccap	Crustacean Cardi	1637339_at	0.0703	0.6618	0.2328	0.1963	0.2250	0.1687	-0.0540	0.9373	-0.2498	0.2082	-0.1958	0.2770	0.0285	0.9816	-0.0004	0.9998	-0.0289	0.9411
---	---	1637340_s_at	0.1757	0.2666	0.0681	0.5625	-0.0318	0.8561	0.0119	0.9857	-0.1001	0.5262	-0.1119	0.4197	0.0464	0.9640	-0.0661	0.8659	-0.1125	0.7223
Rpt3	Rpt3	1637341_at	-0.1991	0.2028	0.5299	0.0474	0.9798	0.0003	0.1099	0.8640	-0.5410	0.0256	-0.6509	0.0070	-0.2842	0.4603	0.1945	0.3104	0.4787	0.0540
l(2)03709	lethal (2) 03709	1637342_s_at	0.3180	0.1299	-0.0300	0.9101	-0.2018	0.2179	-0.1823	0.5336	0.2053	0.1773	0.3876	0.0122	-0.0774	0.9405	-0.0923	0.8272	-0.0149	0.9769
---	---	1637343_at	-0.1133	0.5612	-0.0221	0.8559	0.0529	0.7832	-0.0828	0.8679	-0.0345	0.8842	0.0483	0.8069	-0.0537	0.9642	0.0753	0.8685	0.1290	0.7259
CG31529	CG31529	1637344_at	-0.1191	0.5774	0.0499	0.7727	-0.0456	0.8009	-0.0429	0.9529	-0.1867	0.3616	-0.1438	0.4470	-0.0432	0.9641	-0.1355	0.6288	-0.0923	0.7578
CG31856	CG31856	1637345_at	0.1200	0.4362	-0.0120	0.9113	-0.0161	0.9367	0.0908	0.8899	0.3056	0.1623	0.2149	0.2791	-0.1037	0.8380	-0.0595	0.8303	0.0442	0.8761
CG33514	CG33514	1637346_at	-0.1160	0.9528	0.1444	0.3859	-0.0006	0.9982	-0.3520	0.9220	-1.3170	0.2239	-0.9650	0.3303	0.1483	0.9823	-0.8604	0.6028	-1.0087	0.5372
CG5317	CG5317	1637347_at	0.1160	0.5552	0.0875	0.4214	0.3147	0.0722	0.0848	0.8441	0.0385	0.8521	-0.0463	0.7961	0.1197	0.8494	0.1912	0.4729	0.0715	0.8284
---	---	1637348_at	-0.1173	0.5269	0.0303	0.8217	0.0305	0.8583	0.0675	0.8794	-0.0289	0.8883	-0.0964	0.5115	0.0860	0.8806	0.0625	0.8290	-0.0235	0.9387
Rpn1	Rpn1	1637349_at	0.4079	0.1743	0.6092	0.0395	0.7434	0.0009	0.2421	0.4897	0.2944	0.1138	0.0523	0.8035	0.0823	0.9503	0.4919	0.1948	0.4096	0.3071
CG31086	CG31086	1637350_at	-2.6681	0.0012	-2.0693	0.0388	-2.3883	0.0000	-0.4121	0.1329	-0.9474	0.0006	-0.5352	0.0039	-0.1646	0.9499	-0.4560	0.5818	-0.2915	0.7457
CG15631	CG15631	1637351_at	0.1087	0.5511	0.0149	0.9028	-0.0409	0.8596	-0.0973	0.7857	0.0490	0.7846	0.1463	0.2724	-0.0675	0.9460	-0.1541	0.6382	-0.0866	0.8179
CG18812	CG18812	1637352_a_at	-1.3126	0.0016	-0.8625	0.0772	-1.2947	0.0002	-0.0801	0.8735	-0.3286	0.0635	-0.2485	0.1104	0.2241	0.8424	-0.0184	0.9836	-0.2425	0.6263
Dat	Arylalkylamine N-	1637353_at	-1.4779	0.0028	1.5431	0.0290	0.9723	0.0554	-0.8069	0.3932	-3.4315	0.0004	-2.6245	0.0006	-0.2716	0.8554	-0.3878	0.5432	-0.1162	0.8904
CG8172	CG8172	1637354_at	0.1997	0.3891	-0.0244	0.8630	0.1121	0.6966	-0.0377	0.9630	0.0935	0.7077	0.1312	0.5344	-0.2115	0.7517	-0.2017	0.4925	0.0098	0.9839
CG33696	CG33696	1637355_at	0.0941	0.6280	-0.0221	0.8739	0.0305	0.9043	-0.0305	0.9641	0.0622	0.7739	0.0927	0.6098	-0.0073	0.9940	-0.0416	0.9050	-0.0343	0.9124
Su(var)2-HP2	Heterochromatin p	1637356_a_at	0.4162	0.2845	0.4012	0.4062	0.2756	0.2372	0.1155	0.9218	0.7666	0.0372	0.6511	0.0456	0.1435	0.9447	0.5988	0.3259	0.4552	0.4823
CG9463	CG9463	1637357_at	0.1820	0.4054	-0.1885	0.3775	0.0504	0.8576	0.3231	0.4902	0.2623	0.2973	-0.0608	0.8301	0.0460	0.9739	-0.2761	0.4228	-0.3221	0.3641
CG10232	CG10232	1637358_at	0.1710	0.4437	-0.0408	0.8160	-0.2135	0.2911	-0.1672	0.7130	0.0893	0.7082	0.2566	0.1643	0.0534	0.9666	0.0434	0.9363	-0.0101	0.9852
Lsd-2	Lipid storage drop	1637359_at	1.0203	0.0178	0.6415	0.1167	0.9797	0.0135	0.0031	0.9962	-0.0419	0.8892	-0.0450	0.8619	-0.3280	0.8446	-0.4492	0.5281	-0.1212	0.8982
CG11382	CG11382	1637360_at	0.1893	0.3770	0.1771	0.1551	0.4700	0.0071	0.0953	0.7857	-0.0228	0.9097	-0.1181	0.3786	-0.0260	0.9751	0.0330	0.9175	0.0590	0.8178
CG12179	CG12179	1637361_a_at	-0.1158	0.7304	-0.1454	0.5731	-0.4684	0.0326	-0.3116	0.4815	0.3675	0.1193	0.6792	0.0070	-0.3243	0.6955	0.0579	0.9151	0.3822	0.2879
CG12484	CG12484	1637362_at	-0.0558	0.8339	0.0672	0.6038	-0.0195	0.9539	0.1129	0.8233	-0.0731	0.7551	-0.1860	0.2960	0.0865	0.9017	-0.1962	0.4452	-0.2827	0.2855
CG5890	CG5890	1637363_at	-0.0434	0.8053	-0.0604	0.7145	-0.1736	0.3723	0.0399	0.9489	0.2933	0.0917	0.2535	0.1017	-0.0073	0.9964	0.1058	0.8307	0.1131	0.8040
CG13558	CG13558	1637364_at	0.2212	0.3334	0.1853	0.2338	0.2452	0.2056	-0.0187	0.9777	-0.1366	0.4386	-0.1179	0.4651	0.2295	0.7628	0.1123	0.7716	-0.1172	0.7492
CG31122	CG31122	1637365_at	-0.3706	0.3346	-0.5636	0.0752	-0.9119	0.0050	-0.2088	0.5556	0.2973	0.1035	0.5061	0.0082	0.0914	0.9626	0.1839	0.7829	0.0925	0.9020
CG16876	CG16876	1637366_at	-2.7158	0.1143	-1.9053	0.0140	-2.2603	0.0012	-0.3711	0.8336	-1.6795	0.0204	-1.3084	0.0357	0.0533	0.9939	-1.0649	0.5062	-1.1182	0.4856
CG10799 /// DyakCG10799	CG10799	1637367_at	2.7148	0.0058	1.5169	0.1746	3.1625	0.0001	1.2541	0.2248	0.5426	0.3909	-0.7116	0.1962	-0.4968	0.8472	-0.5709	0.6191	-0.0741	0.9624
CG9406	CG9406	1637368_at	-0.2690	0.1489	-0.2714	0.1961	-0.0650	0.7789	0.1022	0.8578	-0.1000	0.6690	-0.2022	0.2768	-0.1565	0.8320	-0.1248	0.7245	0.0317	0.9414
Hdac3	Hdac3	1637369_at	0.1314	0.4291	0.2126	0.3089	-0.1679	0.4342	-0.2133	0.5621	-0.1156	0.5707	0.0978	0.6082	-0.1214	0.8791	-0.0163	0.9759	0.1051	0.7703
yellow-f	yellow f	1637370_at	-0.8767	0.0507	-2.6108	0.0074	-2.2160	0.0001	-0.1811	0.8578	0.4754	0.1880	0.6565	0.0483	-0.4830	0.7095	-1.0825	0.0660	-0.5995	0.2834
---	---	1637371_at	0.0124	0.9594	0.1247	0.3141	0.0651	0.8113	-0.1172	0.8190	-0.1826	0.3731	-0.0654	0.7675	0.1449	0.8091	0.0520	0.8852	-0.0929	0.7464
CG33335	CG33335	1637372_at	0.1074	0.6053	-0.0542	0.7524	-0.5929	0.0513	-0.3908	0.2789	0.0483	0.8632	0.4391	0.0288	0.1261	0.8541	-0.0888	0.8069	-0.2148	0.4627
---	---	1637373_at	-0.1565	0.4293	-0.0282	0.8250	-0.2226	0.3243	-0.0415	0.9633	-0.0416	0.8990	-0.0001	0.9996	0.1756	0.6961	0.0447	0.8679	-0.1309	0.5092
CG33779	CG33779	1637374_at	0.1371	0.3914	0.0043	0.9884	-0.1739	0.3119	0.0403	0.9603	0.2881	0.1808	0.2478	0.1997	-0.0450	0.9588	-0.0307	0.9376	0.0143	0.9695
---	---	1637375_at	0.1571	0.4145	0.2052	0.1143	0.0021	0.9922	-0.0675	0.8942	0.0641	0.7485	0.1316	0.4016	0.0888	0.8953	-0.0275	0.9462	-0.1163	0.6719
CG30108	CG30108	1637376_at	0.2036	0.3358	0.3364	0.0788	0.0435	0.9091	0.1229	0.8132	0.3652	0.0770	0.2424	0.1854	0.3318	0.7893	0.4434	0.3972	0.1115	0.8764
---	---	1637377_x_at	-0.0870	0.6331	-0.3884	0.2186	-0.6679	0.0373	-0.3109	0.6113	0.1543	0.6556	0.4652	0.0897	-0.2797	0.7070	-0.0966	0.8117	0.1831	0.5817
ia2	ia2	1637378_s_at	-1.4599	0.0113	-3.8406	0.0048	-3.5458	0.0000	0.3073	0.7293	2.2578	0.0006	1.9505	0.0006	-0.0837	0.9296	-0.2841	0.3407	-0.2004	0.5341
CG14671	CG14671	1637379_at	-0.3433	0.1969	0.1381	0.4537	0.0932	0.6660	-0.1119	0.8642	-0.2256	0.3400	-0.1137	0.6322	0.0329	0.9816	0.3336	0.2861	0.3007	0.3642
CaMKI	calmodulin-depen	1637380_s_at	0.7341	0.0209	0.6950	0.0248	0.8926	0.0230	-0.0891	0.9309	-0.4302	0.1505	-0.3411	0.2039	-0.2521	0.8439	-0.5225	0.3053	-0.2703	0.6323
---	---	1637381_at	0.3090	0.2728	0.1783	0.3393	0.1947	0.3037	0.0623	0.9567	0.0019	0.9965	-0.0604	0.8675	-0.0649	0.9550	-0.1742	0.6312	-0.1094	0.7838
---	---	1637382_at	0.5276	0.0833	0.3340	0.0665	0.5858	0.0323	0.2019	0.5363	0.0707	0.7223	-0.1313	0.4098	-0.0283	0.9742	-0.0445	0.8893	-0.0163	0.9595
CG11168	CG11168	1637383_at	-0.1062	0.5339	0.0693	0.7886	-0.1148	0.4418	-0.0145	0.9838	-0.0445	0.8341	-0.0299	0.8791	0.1712	0.8086	0.1235	0.7106	-0.0477	0.9065
CG3638	CG3638	1637384_a_at	-0.2104	0.7583	-0.3148	0.5492	-0.0288	0.9052	-0.0802	0.9072	-0.2555	0.2501	-0.1753	0.3946	-0.3108	0.8954	-0.4108	0.6750	-0.0999	0.9353
CG9780	CG9780	1637385_a_at	0.3304	0.3891	1.2689	0.1884	1.2245	0.0126	-0.1596	0.9353	-1.6662	0.0115	-1.5065	0.0110	-0.0736	0.9871	-0.6351	0.5607	-0.5615	0.6166
rdgC	retinal degenerati	1637386_at	-2.5069	0.0006	-1.2383	0.0459	-2.4661	0.0000	-0.8035	0.0465	-1.0350	0.0016	-0.2315	0.2634	-0.0647	0.9734	-0.3322	0.4999	-0.2675	0.6040
Sap47	Synapse-associat	1637387_s_at	0.0972	0.8962	1.5223	0.0485	1.7918	0.0003	-0.0709	0.9247	-0.6913	0.0085	-0.6204	0.0083	-0.2907	0.9108	0.6737	0.4729	0.9644	0.3079
CG13511	CG13511	1637388_at	0.1076	0.7178	0.3183	0.1182	-0.1042	0.5608	0.0776	0.9154	-0.0721	0.7977	-0.1497	0.4965	0.4722	0.4128	0.2744	0.3667	-0.1978	0.5444
CG11575	CG11575	1637389_at	0.1404	0.4628	-0.0028	0.9878	0.2341	0.1592	0.0555	0.9345	0.0429	0.8703	-0.0126	0.9597	-0.0696	0.9142	0.0622	0.8386	0.1318	0.5869
CG4406	CG4406	1637390_at	-0.1826	0.5870	0.1187	0.7438	0.4151	0.0236	0.1793	0.6391	-0.0525	0.8228	-0.2318	0.1650	-0.0462	0.9839	0.2902	0.6015	0.3364	0.5403
Cpr76Bd	CG9299	1637391_at	0.2787	0.1264	0.2165	0.2202	0.3286	0.0595	-0.0202	0.9838	-0.1096	0.6755	-0.0894	0.7138	0.0434	0.9717	0.0321	0.9496	-0.0112	0.9835

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9263	CG9263	1637392_at	0.2023	0.3098	-0.1252	0.2482	0.0620	0.7785	0.2850	0.3991	0.3335	0.0798	0.0485	0.8214	-0.0495	0.9523	-0.0346	0.9301	0.0149	0.9688
CG33528	CG33528	1637393_at	-0.0028	0.9924	-0.0436	0.7175	-0.0499	0.8463	-0.1052	0.7753	-0.0919	0.5829	0.0132	0.9461	-0.1278	0.9007	-0.1896	0.6416	-0.0617	0.9056
CG31781	CG31781	1637394_s_at	-1.2848	0.0121	-0.3556	0.2802	-0.8189	0.0271	-0.2551	0.6673	-0.5832	0.0450	-0.3281	0.1937	0.3177	0.7644	0.0766	0.9120	-0.2411	0.6190
CG13084	CG13084	1637395_at	0.0302	0.8692	0.0807	0.7345	-0.1219	0.6759	0.0586	0.9435	0.0579	0.8485	-0.0007	0.9981	0.2299	0.8049	0.2134	0.6083	-0.0165	0.9793
---	---	1637396_at	0.1123	0.5432	0.0384	0.7165	0.1161	0.5484	-0.1042	0.7898	-0.0160	0.9459	0.0882	0.5802	-0.0129	0.9928	-0.0764	0.8642	-0.0636	0.8849
CG6631	CG6631	1637397_a_at	-0.0211	0.9151	-0.1313	0.5248	-0.3360	0.2987	-0.4244	0.1790	-0.5803	0.0090	-0.1560	0.3793	-0.3074	0.8202	-0.6453	0.2321	-0.3380	0.5679
capu	cappucino	1637398_a_at	-0.0654	0.8952	-1.3583	0.2594	-0.8472	0.0668	0.4640	0.3328	0.5759	0.0436	0.1118	0.7014	-0.0087	0.9991	-0.7119	0.5688	-0.7032	0.5788
CG16741	CG16741	1637399_at	-0.0423	0.8060	0.0236	0.8959	-0.0049	0.9848	0.0426	0.9540	0.0800	0.7380	0.0374	0.8768	0.0795	0.9116	0.0754	0.8196	-0.0042	0.9924
---	---	1637400_at	-0.0425	0.8307	0.0346	0.8599	0.2099	0.3248	0.0183	0.9847	-0.0809	0.7542	-0.0992	0.6586	-0.0080	0.9939	0.0811	0.7916	0.0891	0.7520
CG11095	CG11095	1637401_at	0.0650	0.7112	-0.1182	0.7292	0.3085	0.0461	-0.0055	0.9952	-0.1307	0.5063	-0.1252	0.4808	-0.5234	0.4207	-0.2643	0.4475	0.2591	0.4699
CG32109	CG32109	1637402_at	-0.3389	0.0825	-0.5098	0.0510	-0.3141	0.2212	-0.1098	0.8350	0.1296	0.5467	0.2394	0.1789	-0.4059	0.6483	-0.1507	0.7105	0.2552	0.4912
CG7953	CG7953	1637403_at	0.1435	0.8667	0.0010	1.0000	-0.1859	0.6325	0.0046	0.9985	0.1687	0.7962	0.1641	0.7820	0.2649	0.9294	-0.2896	0.8172	-0.5544	0.5900
CG10262	CG10262	1637404_at	-0.0698	0.6774	0.0845	0.4996	0.2435	0.3766	-0.1309	0.8014	-0.0310	0.9159	0.0999	0.6417	-0.2969	0.7305	0.0149	0.9829	0.3118	0.4056
CG32649	CG32649	1637405_at	-1.1226	0.0035	-0.5249	0.1315	-0.7461	0.0556	-0.2437	0.7169	-0.5746	0.0639	-0.3309	0.2271	0.0432	0.9853	0.0256	0.9791	-0.0177	0.9846
---	---	1637406_at	0.1577	0.4485	0.1866	0.2135	0.3571	0.0702	-0.0184	0.9835	-0.1214	0.5829	-0.1030	0.6182	-0.0407	0.9516	0.0085	0.9823	0.0493	0.8509
CG32243	CG32243	1637407_at	0.0259	0.9528	-0.4742	0.1853	-0.4156	0.0162	-0.3575	0.4561	0.1475	0.6050	0.5050	0.0366	-0.4536	0.6660	-0.3479	0.4094	0.1057	0.8467
---	---	1637408_at	0.2061	0.2708	-0.0868	0.6700	0.0036	0.9886	-0.0519	0.9311	0.1286	0.4876	0.1805	0.2561	-0.1664	0.7720	-0.1362	0.6024	0.0302	0.9297
CG12865	CG12865	1637409_at	-0.1209	0.6780	-0.0935	0.7069	0.3195	0.2075	0.2959	0.6828	0.0753	0.8656	-0.2206	0.4959	-0.0690	0.9717	0.0300	0.9708	0.0990	0.8831
l(2)gl	lethal giant larve	1637410_s_at	-0.0334	0.9110	0.2806	0.4192	-0.1453	0.2928	-0.2297	0.7187	-0.2190	0.4681	0.0106	0.9766	0.0729	0.9088	0.0579	0.8501	-0.0150	0.9640
Pur-alpha	Purine-rich bindin	1637411_s_at	0.1885	0.5895	1.0802	0.0203	0.4812	0.0263	-0.4749	0.3775	-0.4293	0.1566	0.0455	0.9007	0.0411	0.9816	0.3740	0.3261	0.3329	0.4037
sas	responsible for do	1637412_a_at	0.2399	0.5572	-0.2969	0.2306	-0.4944	0.0277	-0.1161	0.8796	0.6465	0.0222	0.7626	0.0065	0.0240	0.9914	0.1105	0.8792	0.0865	0.9023
CG13424	CG13424	1637413_at	0.2763	0.4548	-0.1702	0.4861	-0.4410	0.0439	-0.0467	0.9716	1.1128	0.0057	1.1595	0.0028	0.2415	0.8128	0.6063	0.1423	0.3648	0.3963
CG4455	CG4455	1637414_at	-0.2491	0.4424	-0.0186	0.9615	-0.0967	0.6301	-0.0085	0.9940	-0.0118	0.9735	-0.0034	0.9906	-0.0223	0.9914	0.1158	0.8586	0.1381	0.8128
CG6576	CG6576	1637415_at	0.3451	0.1162	0.2300	0.2437	0.3596	0.1483	0.0767	0.9248	0.1179	0.6710	0.0412	0.8870	-0.0589	0.9621	-0.0567	0.9157	0.0022	0.9976
CG17631	CG17631	1637416_at	0.1217	0.5710	0.1421	0.4744	0.2942	0.0609	-0.0924	0.8578	-0.1192	0.5497	-0.0268	0.9045	-0.0785	0.9030	0.1661	0.4881	0.2447	0.3079
CG6565	CG6565	1637417_at	-0.4314	0.1279	0.2058	0.2535	0.5659	0.0273	0.1170	0.8801	-0.5168	0.0562	-0.6338	0.0157	-0.1398	0.8870	0.2078	0.5971	0.3475	0.3597
---	---	1637418_at	0.0719	0.7603	0.1644	0.4358	0.5253	0.0082	0.1588	0.6546	-0.2736	0.1087	-0.4324	0.0123	-0.0595	0.9457	0.0633	0.8633	0.1228	0.6683
---	---	1637419_at	0.1301	0.3876	-0.1556	0.4606	-0.0622	0.7408	0.2478	0.5453	0.3625	0.0884	0.1147	0.5888	0.0648	0.9305	-0.0995	0.7166	-0.1643	0.5100
GABA-B-R2	metabotropic GAE	1637420_a_at	-0.1768	0.4203	-0.2966	0.4071	0.0000	0.9998	-0.4606	0.3051	-0.5793	0.0352	-0.1188	0.6645	-0.5334	0.3162	-0.2886	0.2835	0.2447	0.3884
Cht2	Chitinase 2	1637421_at	-0.5760	0.1511	-0.1723	0.6003	-0.2301	0.7302	0.2452	0.4915	-0.1861	0.3347	-0.4313	0.0193	-0.1545	0.9689	-0.4073	0.7362	-0.2528	0.8499
CG3246	CG3246	1637422_at	2.6234	0.0013	1.7050	0.0780	3.6526	0.0001	1.0642	0.1662	0.6923	0.1366	-0.3719	0.3916	-0.9211	0.6749	-0.2051	0.8698	0.7160	0.4301
CG6674	CG6674	1637423_at	-0.1382	0.6002	0.0749	0.5372	0.0820	0.7402	-0.0860	0.9234	-0.3351	0.2132	-0.2491	0.3101	0.0162	0.9898	-0.0182	0.9670	-0.0345	0.9263
sti	Citron	1637424_at	0.0065	0.9862	0.2059	0.5456	-0.7736	0.0066	-0.7945	0.0532	-0.2526	0.2888	0.5419	0.0184	0.1783	0.9112	0.0717	0.9376	-0.1066	0.8918
---	---	1637425_at	0.0456	0.8731	0.0178	0.8611	0.0998	0.6806	0.0075	0.9943	-0.0683	0.7998	-0.0757	0.7511	-0.0683	0.9495	0.0492	0.9221	0.1175	0.7513
CG11449	CG11449	1637426_at	0.0790	0.6451	0.3095	0.1433	0.1112	0.4985	-0.0308	0.9665	-0.2887	0.1307	-0.2579	0.1306	0.0808	0.8928	0.0329	0.9263	-0.0479	0.8737
---	---	1637427_at	-0.0067	0.9768	0.0000	1.0000	-0.0601	0.7138	-0.0195	0.9760	-0.1075	0.5296	-0.0880	0.5827	-0.0405	0.9487	-0.0820	0.7000	-0.0415	0.8684
pyd	tamou	1637428_a_at	0.1937	0.5846	-0.3714	0.5276	-0.1015	0.6406	0.0443	0.9600	0.3640	0.1151	0.3197	0.1214	-0.2340	0.8949	-0.1371	0.8858	0.0969	0.9161
CG8441	CG8441	1637429_at	-0.0573	0.7367	0.0856	0.6536	0.2684	0.1004	0.1980	0.5587	0.0497	0.8188	-0.1483	0.3540	0.0738	0.9246	0.1860	0.4881	0.1123	0.7027
CG1572	CG1572	1637430_s_at	-1.1971	0.0203	0.2646	0.2215	-0.2347	0.4619	0.2169	0.5086	0.3914	0.0317	0.1745	0.2658	0.7886	0.6332	1.9541	0.0276	1.1655	0.1192
CG34363	CG12904	1637431_at	0.1173	0.4684	-0.0005	1.0000	0.0681	0.6858	0.0376	0.9387	0.0012	0.9951	-0.0364	0.8233	0.1106	0.8387	0.0628	0.8341	-0.0478	0.8744
CG3097	CG3097	1637432_at	2.8732	0.0011	2.3227	0.0144	2.4070	0.0003	0.4121	0.1366	0.7047	0.0021	0.2926	0.0548	0.3457	0.8837	0.1865	0.8857	-0.1592	0.8977
CG30166	CG30166	1637433_at	0.1793	0.3686	0.0459	0.7962	0.0843	0.6333	0.0217	0.9803	-0.0668	0.7993	-0.0885	0.6940	0.0869	0.9092	0.0306	0.9445	-0.0563	0.8779
CG14563	CG14563	1637434_at	0.0571	0.8840	0.2150	0.2901	0.1082	0.7027	0.0004	0.9997	-0.0060	0.9899	-0.0064	0.9874	0.1082	0.9451	0.0140	0.9890	-0.0942	0.8858
kek4	kekkon4	1637435_at	1.0922	0.1827	1.9650	0.0395	0.8108	0.0568	0.1472	0.8449	0.0283	0.9447	-0.1188	0.6822	1.3431	0.6531	0.7938	0.5240	-0.5493	0.6763
---	---	1637436_at	-0.0765	0.6453	-0.0900	0.5651	-0.0175	0.9257	0.0316	0.9584	0.1031	0.5477	0.0715	0.6657	-0.0286	0.9742	0.0608	0.8382	0.0894	0.7231
CG6254	CG6254	1637437_at	-0.3438	0.2683	-0.2382	0.0870	0.0104	0.9729	0.0477	0.9462	0.0584	0.8128	0.0107	0.9664	-0.2866	0.7464	0.1019	0.8389	0.3885	0.3089
Wnt2	Wnt oncogene an	1637438_at	-2.0463	0.0067	-2.3041	0.0439	-1.6063	0.0027	0.4534	0.6506	0.4152	0.4025	-0.0381	0.9503	-0.0967	0.9816	0.2151	0.8680	0.3119	0.7734
CG14709	CG14709	1637439_at	-0.0279	0.8973	-0.2235	0.1215	-0.0966	0.6301	0.1282	0.7315	0.3569	0.0388	0.2287	0.1235	-0.2212	0.6955	-0.2760	0.2267	-0.0548	0.8600
CG17344	CG17344	1637440_at	-0.0049	0.9814	0.0645	0.5662	-0.1076	0.5774	-0.1798	0.6325	-0.1372	0.4757	0.0426	0.8410	-0.0074	0.9923	0.0893	0.6465	0.0968	0.6166
CG3982	CG3982	1637441_at	0.0601	0.7855	0.2319	0.1387	0.0665	0.7228	-0.2072	0.4350	-0.2138	0.1405	-0.0066	0.9711	0.0114	0.9914	0.0800	0.7891	0.0685	0.8178
ttv	tout velu	1637442_at	0.1982	0.1823	0.2388	0.4845	0.1217	0.6045	-0.2799	0.4145	-0.0078	0.9790	0.2721	0.1093	-0.1672	0.8521	-0.0274	0.9646	0.1398	0.7442
---	---	1637443_at	0.2090	0.2457	-0.0215	0.8423	0.1386	0.5295	0.1323	0.8336	0.1078	0.6942	-0.0244	0.9343	-0.1204	0.8016	-0.1675	0.3982	-0.0470	0.8575
pug	Pugilist	1637444_s_at	2.5278	0.0035	0.9677	0.4478	2.4775	0.0000	0.7211	0.3952	0.4694	0.3322	-0.2517	0.5993	-0.7951	0.8008	-1.1171	0.3866	-0.3220	0.8484
CG31513	CG31513	1637445_at	0.0750	0.6120	0.0247	0.8259	0.0185	0.9084	-0.0499	0.9222	0.0330	0.8718	0.0829	0.5908	-0.0411	0.9427	-0.0339	0.8983	0.0072	0.9806
---	---	1637446_at	0.2850	0.1171	0.2661	0.2152	0.1453	0.4226	-0.2417	0.4238	-0.2871	0.0865	-0.0455	0.8085	-0.0322	0.9751	-0.2866	0.2430	-0.2544	0.3293
CG12498	CG12498	1637447_at	0.1054	0.6163	0.0408	0.7840	0.0256	0.9053	-0.1371	0.7203	-0.1574	0.3735	-0.0203	0.9240	0.0272	0.9831	-0.0728	0.8625	-0.1000	0.7785
RpS19a	Ribosomal protein	1637448_s_at	0.4615	0.0377	0.8947	0.0119	0.9160	0.0010	0.1794	0.7221	-0.2958	0.1907	-0.4752	0.0273	0.1029	0.8655	0.0292	0.9404	-0.0737	0.8039
---	---	1637449_at	0.3207	0.3053	0.2422	0.4209	-0.0404	0.8082	-0.0151	0.9838	0.0971	0.6145	0.1122	0.5054	0.1881	0.8283	-0.0147	0.9829	-0.2028	0.6036
---	---	1637450_at	0.1167	0.6203	0.0113	0.9509	0.0558	0.7618	0.1278	0.7857	0.1795	0.3659	0.0518	0.8134	0.0159	0.9862	-0.0422	0.8910	-0.0582	0.8255
CG31239	CG31239	1637451_at	-0.1935	0.3634	-0.8899	0.0659	-0.9532	0.0119	0.0573	0.9441	0.6400	0.0139	0.5827	0.0130	-0.0983	0.8814	-0.1806	0.4816	-0.0823	0.7824
---	---	1637452_at	0.1413	0.3731	0.0843	0.4567	0.1418	0.3757	0.0609	0.9028	0.0277	0.9008	-0.0332	0.8599	0.0279	0.9741	0.0953	0.6843	0.0674	0.7878
CG3797 /// DmirCG3797	CG3797	1637453_at	0.0181	0.9609	0.0872	0.7025	0.0050	0.9852	-0.0748	0.9228	0.3502	0.1328	0.4251	0.0474	0.1413	0.8705	0.4191	0.2110	0.2778	0.4326
Obp19a	Odorant-binding p	1637454_a_at	0.0713	0.7512	0.2401	0.3492	0.0341	0.8740	-0.1013	0.8676	-0.1548	0.4929	-0.0534	0.8276	0.0168	0.9913	-0.0800	0.8487	-0.0968	0.7940
CG11085	CG11085	1637455_at	0.2042	0.2164	0.0734	0.4941	0.1110	0.5411	-0.0903	0.8384	0.0563	0.7793	0.1466	0.3335	-0.0315	0.9816	-0.0173	0.9724	0.0142	0.9762
CG8878	CG8878	1637456_at	-0.2473	0.4305	-0.0655	0.5409	0.0644	0.7294	0.1999	0.5544	0.3389	0.0569	0.1390	0.3923	0.1686	0.8097	0.3940	0.1624	0.2253	0.4528
CG7646	CG7646	1637457_at	-0.7186	0.0109	-0.5951	0.4626	-1.0068	0.0058	-0.3209	0.6582	-0.1092	0.7985	0.2117	0.5343	0.0560	0.9852	-0.0119	0.9938	-0.0680	0.9434
---	---	1637458_at	0.0402	0.8381	0.0857	0.7410	0.1731	0.4508	-0.1565	0.7138	-0.2341	0.2287	-0.0775	0.7065	-0.0312	0.9862	-0.0147	0.9865	0.0164	0.9820
---	---	1637459_at	0.1136	0.4145	0.0486	0.6518	0.0643	0.6779	0.1485	0.6144	0.0837	0.6063	-0.0648	0.6745	0.0521	0.9530	-0.0083	0.9877	-0.0604	0.8618
CG30190	CG30190	1637460_at	-2.1368	0.0009	-0.2818	0.1431	-0.4872	0.0652	-0.1819	0.8401	-1.8876	0.0007	-1.7057	0.0006	-0.0123	0.9939	0.0216	0.9710	0.0340	0.9461

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG9527	CG9527	1637461_at	0.4772	0.0788	0.2516	0.1587	0.1369	0.4639	0.0764	0.8796	0.3258	0.0629	0.2494	0.1056	0.1925	0.8424	0.1295	0.7962	-0.0630	0.9086
CG13833	CG13833	1637462_at	-1.8253	0.0008	-1.6967	0.0583	-1.9792	0.0001	-0.3902	0.3381	-0.7314	0.0080	-0.3412	0.1036	-0.0836	0.9737	-0.6176	0.3068	-0.5340	0.3989
Nrg	neuroglian	1637463_a_at	-0.6522	0.0040	0.6822	0.2666	0.3023	0.1842	0.0699	0.9254	-0.6697	0.0096	-0.7396	0.0037	0.4100	0.7644	0.7146	0.1995	0.3046	0.6231
---	---	1637464_at	-0.2207	0.3515	-0.2578	0.2543	-0.0938	0.5412	0.1035	0.8578	0.1887	0.3736	0.0852	0.6962	-0.0572	0.9589	0.0070	0.9925	0.0642	0.8834
CG31157	CG31157	1637465_at	0.0632	0.7642	-0.0650	0.8223	0.3059	0.0911	0.0529	0.9125	0.0453	0.8101	-0.0076	0.9685	-0.2026	0.8042	-0.0563	0.9149	0.1463	0.7000
Ric	Ras-related which	1637466_at	-0.2913	0.2312	-0.8006	0.0157	-0.9439	0.0024	0.1116	0.8196	0.4481	0.0272	0.3365	0.0540	0.2302	0.8014	-0.0628	0.9156	-0.2930	0.4573
CG5245	CG5245	1637467_at	0.3150	0.1578	-0.0671	0.5638	0.1783	0.3090	0.0306	0.9649	0.4275	0.0293	0.3968	0.0253	-0.1321	0.8472	0.1541	0.6142	0.2862	0.3271
CG7516	CG7516	1637468_at	-0.1963	0.4379	-0.1264	0.4901	0.3900	0.0438	0.3108	0.5461	0.1662	0.5638	-0.1446	0.5880	-0.1510	0.8725	0.2490	0.5176	0.4000	0.2960
pen-2	presenilin enhanc	1637469_at	-0.0523	0.7878	0.2256	0.3991	0.3223	0.2193	-0.0958	0.8872	-0.0162	0.9626	0.0796	0.7476	-0.1688	0.8439	0.2199	0.5501	0.3887	0.2837
CG4420 /// DsimCG4420	CG4420	1637470_at	-0.0078	0.9844	0.8809	0.0074	0.4778	0.0588	-0.2982	0.5433	-0.8535	0.0057	-0.5553	0.0230	0.2366	0.8182	0.0516	0.9402	-0.1850	0.6925
---	---	1637471_at	-0.0481	0.8744	0.0554	0.5985	0.2241	0.2039	0.0121	0.9883	-0.1702	0.3983	-0.1823	0.3064	-0.0499	0.9589	0.0498	0.9054	0.0998	0.7516
CG32570	CG32570	1637472_at	0.1328	0.5605	0.4022	0.0459	0.2361	0.2650	0.0255	0.9722	-0.2031	0.2680	-0.2286	0.1601	0.1377	0.8235	0.0568	0.8773	-0.0809	0.7936
CG18787 /// CG18789	CG18789 /// CG1	1637473_s_at	0.3440	0.1886	-0.5422	0.2111	-0.5047	0.0108	-0.0496	0.9550	0.9705	0.0024	1.0201	0.0012	-0.0025	0.9993	0.0942	0.8500	0.0967	0.8345
CG34104	CG34104	1637474_at	0.1348	0.6334	-0.0698	0.5727	-0.0656	0.7372	0.1186	0.8820	0.3081	0.2606	0.1895	0.4637	0.0619	0.9589	-0.0328	0.9531	-0.0947	0.8287
feo	fascetto	1637475_at	0.5656	0.5121	-1.1869	0.2143	-1.2234	0.0194	0.1408	0.7929	1.5949	0.0003	1.4541	0.0002	-0.0517	0.9922	0.0606	0.9781	0.1122	0.9471
CG15579	CG15579	1637476_at	0.0161	0.9633	-0.1197	0.4737	-0.2347	0.3964	0.1103	0.8738	0.1674	0.5138	0.0570	0.8376	-0.0257	0.9862	0.0685	0.8879	0.0942	0.8221
CG9447	CG9447	1637477_at	0.0846	0.6677	0.1386	0.4099	0.1454	0.5399	-0.0353	0.9689	-0.1525	0.5469	-0.1173	0.6246	-0.0956	0.8811	-0.1022	0.7119	-0.0066	0.9865
heph	ectopic margin	1637478_s_at	0.8227	0.0512	0.5652	0.3632	-0.6854	0.0131	-0.4171	0.4652	0.6894	0.0328	1.1065	0.0025	0.9161	0.5089	0.6234	0.3288	-0.2927	0.6873
pn	prune	1637479_at	1.1818	0.0035	0.7905	0.1112	1.5518	0.0000	0.0339	0.9668	-0.2601	0.2216	-0.2940	0.1215	-0.7439	0.3712	-0.6950	0.1293	0.0489	0.9387
---	---	1637480_at	0.1276	0.4823	0.2211	0.3044	-0.0139	0.9653	-0.1204	0.7104	0.0260	0.8974	0.1464	0.2710	0.1630	0.8553	0.2284	0.5523	0.0654	0.8967
Tollo	Tollo	1637481_at	-4.5109	0.0010	-5.9678	0.0014	-5.5282	0.0000	0.2700	0.8192	0.8297	0.0754	0.5596	0.1753	-0.0748	0.9826	-0.5114	0.5329	-0.4367	0.6067
oxt	peptide O-xylosylt	1637482_at	-0.2300	0.1249	0.1373	0.4966	0.2319	0.2867	-0.2513	0.4117	-0.3246	0.0598	-0.0733	0.6798	-0.0557	0.9623	0.1158	0.7722	0.1715	0.6279
CG8736	CG8736	1637483_at	0.1793	0.4704	0.4217	0.3190	0.4055	0.1670	-0.0343	0.9639	0.0818	0.7262	0.1161	0.5560	-0.0597	0.9737	0.1841	0.7198	0.2438	0.6130
CG6985	CG6985	1637484_at	-0.4455	0.0401	-0.7884	0.0181	-0.9094	0.0014	-0.0752	0.9098	0.4295	0.0474	0.5047	0.0156	0.0716	0.9357	0.0846	0.8172	0.0131	0.9769
CG10827	CG10827	1637485_at	-0.0978	0.9442	0.1446	0.5984	-0.1972	0.2792	-0.3222	0.9182	-0.8424	0.3944	-0.5202	0.5895	0.1800	0.9365	-0.2742	0.7439	-0.4542	0.5476
CG31698	CG31698	1637486_at	0.2162	0.3134	0.0493	0.7891	0.1821	0.3281	0.3045	0.4979	0.3779	0.1112	0.0734	0.7820	0.0131	0.9913	0.0152	0.9701	0.0020	0.9965
Dcp-1	caspase	1637487_at	1.0693	0.2418	0.2829	0.3952	0.7314	0.0187	0.4663	0.7538	0.5925	0.3662	0.1262	0.8662	0.1453	0.9460	-0.0599	0.9547	-0.2053	0.7935
Cad88C	Cad88C	1637488_at	0.2864	0.2571	0.0895	0.6571	0.3807	0.1479	0.1083	0.8507	0.0618	0.8112	-0.0466	0.8478	-0.1041	0.9421	-0.1606	0.7621	-0.0564	0.9264
CG6133	CG6133	1637489_at	1.0313	0.0391	0.3847	0.0975	0.8168	0.0037	0.4914	0.3542	1.2564	0.0023	0.7650	0.0124	-0.0184	0.9928	0.5718	0.2254	0.5902	0.2439
CG13314	CG13314	1637490_at	-0.2438	0.4006	0.0387	0.8012	-0.2111	0.2902	-0.1243	0.7922	-0.1773	0.3654	-0.0530	0.8054	-0.0036	0.9977	-0.1014	0.7626	-0.0978	0.7627
CG3835	CG3835	1637491_s_at	0.7699	0.0043	0.6262	0.4229	1.2372	0.0007	0.1011	0.8864	-0.3433	0.1498	-0.4444	0.0439	-0.4764	0.7768	-0.4720	0.5321	0.0044	0.9979
epsilonTry	epsilonTrypsin	1637492_at	0.6880	0.2756	0.0645	0.6350	0.2906	0.1442	0.1541	0.7409	0.0069	0.9820	-0.1472	0.4479	-0.0716	0.9823	-0.3716	0.6473	-0.3000	0.7243
---	---	1637493_at	0.3353	0.1164	0.2183	0.2138	0.0170	0.9440	-0.2253	0.6096	0.0340	0.9126	0.2593	0.1896	0.1091	0.8427	0.0393	0.9093	-0.0698	0.7986
CG3071	CG3071	1637494_at	0.5528	0.0547	-0.4167	0.3986	0.0261	0.9186	0.3060	0.4710	1.2210	0.0008	0.9150	0.0016	-0.1105	0.9514	0.3676	0.5126	0.4780	0.3884
CG18735	CG18735	1637495_at	0.1542	0.5411	0.2402	0.1012	0.0599	0.8024	-0.0452	0.9610	-0.0297	0.9321	0.0155	0.9599	0.2218	0.8222	0.0803	0.8962	-0.1415	0.7686
CG30219	CG30219	1637496_at	-1.6706	0.0009	-0.5124	0.0520	-1.1838	0.0007	-0.5077	0.1526	-1.4472	0.0004	-0.9395	0.0011	0.1513	0.8903	-0.2911	0.4903	-0.4424	0.2964
CG6418	CG6418	1637497_at	-0.0075	0.9757	0.2971	0.1336	0.6328	0.0233	0.0608	0.9441	-0.1161	0.6821	-0.1769	0.4509	-0.2154	0.7770	0.1817	0.6063	0.3971	0.2456
CG41106	CG41106	1637498_at	0.3741	0.0617	0.1113	0.5566	0.3105	0.0747	-0.1404	0.8531	-0.0153	0.9711	0.1251	0.6598	-0.2156	0.7595	-0.2200	0.4598	-0.0044	0.9932
CG5953	CG5953	1637499_s_at	-0.0175	0.9664	-1.1640	0.2450	-0.9224	0.2116	0.4745	0.6015	0.9553	0.0434	0.4808	0.2429	0.3802	0.9287	-0.2543	0.9039	-0.6345	0.6732
---	---	1637500_at	-0.1359	0.5887	0.0611	0.6814	-0.2051	0.1883	-0.1793	0.7081	0.0115	0.9733	0.1908	0.3406	0.0348	0.9626	0.0747	0.7628	0.0398	0.8880
CG1835	CG1835	1637501_a_at	-0.0787	0.7096	-0.1869	0.5205	-0.1654	0.4987	0.2734	0.5757	0.2688	0.2872	-0.0046	0.9880	0.3285	0.7070	0.3075	0.3973	-0.0210	0.9703
CG13296	CG13296	1637502_at	-0.0117	0.9710	0.2068	0.2130	-0.0464	0.8427	-0.0851	0.9218	-0.2310	0.3894	-0.1459	0.5755	0.0895	0.8875	0.1176	0.6490	0.0280	0.9331
ear	ENL/AF9-related	1637503_at	0.0536	0.8142	0.4159	0.0925	0.7184	0.0113	0.0441	0.9518	-0.0923	0.6906	-0.1364	0.4788	-0.2001	0.8521	0.1615	0.7618	0.3617	0.4258
CG17754	CG17754	1637504_at	0.3130	0.4601	0.5507	0.3490	-0.0815	0.7952	-0.4812	0.5724	0.0382	0.9531	0.5194	0.1809	0.0449	0.9885	0.0206	0.9885	-0.0243	0.9835
---	---	1637505_at	0.0227	0.8968	0.1762	0.4262	0.1265	0.5135	0.1257	0.7764	0.0813	0.7034	-0.0445	0.8338	0.1707	0.7997	0.1212	0.7020	-0.0495	0.8982
CG32758	CG32758	1637506_at	-1.2448	0.0028	-0.6250	0.1402	-0.4458	0.2417	-0.1816	0.8073	-0.7216	0.0229	-0.5400	0.0466	-0.4028	0.7478	-0.1551	0.8242	0.2477	0.6684
CG32835	CG32835	1637507_at	0.1543	0.3339	-0.1630	0.3205	0.1906	0.3158	0.2308	0.5357	0.2967	0.1256	0.0659	0.7565	-0.1975	0.7686	-0.1099	0.7439	0.0876	0.8004
slv	saliva	1637508_at	0.8488	0.0049	0.6662	0.0944	0.6941	0.0031	-0.0971	0.8358	0.6052	0.0052	0.7023	0.0016	-0.0920	0.9421	0.4347	0.2509	0.5268	0.2037
Tim10	Tim10	1637509_s_at	-0.2425	0.3174	0.2992	0.0921	0.0833	0.7211	0.0175	0.9803	-0.4418	0.0183	-0.4593	0.0095	0.3323	0.7464	0.3410	0.4317	0.0088	0.9905
Hexo1	Hexosaminidase	1637510_s_at	-0.7427	0.0074	-1.1239	0.0412	-0.8469	0.0050	0.1276	0.8309	0.1902	0.4196	0.0626	0.8069	-0.2091	0.8283	-0.2629	0.5300	-0.0538	0.9231
CG12521	CG12521	1637511_at	-0.0490	0.8468	0.0462	0.8250	0.1830	0.3648	0.1031	0.8189	0.0670	0.7485	-0.0361	0.8605	-0.1309	0.8861	0.0611	0.9075	0.1920	0.6036
CG9294	CG9294	1637512_at	0.0261	0.9043	-0.1441	0.4023	0.0262	0.9052	0.1289	0.6933	0.1557	0.3071	0.0268	0.8806	0.0695	0.9004	0.0624	0.8132	-0.0071	0.9838
CG11275	CG11275	1637513_at	-0.7563	0.0230	-0.7173	0.0723	-0.4818	0.0084	0.0297	0.9664	-0.3620	0.0551	-0.3917	0.0260	-0.1637	0.8882	-0.2889	0.5231	-0.1252	0.8178
CG14142	CG14142	1637514_at	0.2663	0.1658	0.2901	0.2484	0.2061	0.2192	-0.0657	0.9066	-0.1557	0.4006	-0.0900	0.6241	0.2502	0.7478	-0.0342	0.9494	-0.2844	0.3972
cul-2	cul-2	1637515_s_at	0.0142	0.9523	0.2117	0.4075	0.0694	0.7441	-0.0922	0.8376	-0.1739	0.3073	-0.0818	0.6364	0.0792	0.9294	0.1155	0.7293	0.0364	0.9273
CG15449	CG15449	1637516_at	0.6671	0.0489	0.1679	0.3531	0.2384	0.2040	0.1746	0.5863	0.0668	0.7266	-0.1078	0.4914	0.0532	0.9640	-0.2860	0.3615	-0.3392	0.2995
KrT95D	Krueppel target at	1637517_s_at	1.4940	0.0260	1.3290	0.1821	2.1079	0.0000	0.2033	0.7235	-0.1703	0.5401	-0.3736	0.1061	-0.5635	0.8305	-0.3366	0.8137	0.2269	0.8800
CG9314	CG9314	1637518_at	0.2996	0.1686	0.7926	0.0793	0.1293	0.4650	0.1729	0.6099	0.2460	0.1463	0.0731	0.6814	0.8298	0.3712	0.7040	0.1517	-0.1257	0.8467
Pvf3	CG13783	1637519_at	0.1483	0.4201	0.2430	0.4068	0.1022	0.7219	-0.2123	0.7982	-0.1421	0.7158	0.0702	0.8583	-0.0705	0.9589	-0.0997	0.8481	-0.0292	0.9578
CG1518	CG1518	1637520_at	0.4802	0.0425	0.9452	0.0411	1.1843	0.0004	0.0617	0.9339	-0.2511	0.2512	-0.3128	0.1081	-0.2148	0.8235	0.3094	0.4397	0.5241	0.2107
CG13449	CG13449	1637521_at	-0.3186	0.7052	-0.0657	0.6492	-0.6928	0.0299	-0.2488	0.6338	-0.6732	0.0168	-0.4244	0.0672	-0.0294	0.9933	-0.4216	0.6294	-0.3922	0.6540
CG5043	CG5043	1637522_at	0.0878	0.5484	-0.0610	0.5928	0.3355	0.0353	0.3129	0.3206	0.3397	0.0675	0.0267	0.9037	0.0595	0.9409	0.2397	0.3240	0.1802	0.4851
Nek2	Nek2	1637523_at	0.1494	0.4558	0.2901	0.3795	0.1014	0.6718	-0.2691	0.5008	0.0576	0.8308	0.3267	0.0851	-0.1280	0.9246	0.1191	0.8478	0.2471	0.6150
CG40270	CG40270	1637524_at	0.2627	0.2129	-0.1584	0.3849	-0.1416	0.5456	0.1113	0.8578	0.2607	0.2408	0.1495	0.4804	0.0851	0.9374	-0.1173	0.7790	-0.2024	0.5767
Mpcp	Mitochondrial pho	1637525_s_at	-0.1739	0.2645	0.3026	0.0799	0.0761	0.6664	-0.0380	0.9477	-0.4425	0.0145	-0.4046	0.0133	-0.0216	0.9845	-0.1581	0.5414	-0.1365	0.6100
Ggamma30A	G protein gamma	1637526_s_at	1.9211	0.3532	-1.2722	0.0100	-0.2040	0.4078	1.6415	0.0038	1.1922	0.0013	-0.4493	0.0528	0.0565	0.9963	-2.0879	0.4302	-2.1444	0.4309
CG33140	CG33140	1637527_at	0.1547	0.4651	-0.1525	0.4870	0.1768	0.4848	0.1126	0.8822	0.3352	0.1948	0.2226	0.3509	-0.0393	0.9717	-0.0993	0.7735	-0.0601	0.8746
---	---	1637528_at	-0.0174	0.9422	-0.0522	0.6597	0.0149	0.9459	-0.0245	0.9744	0.0254	0.9215	0.0500	0.8087	-0.0083	0.9929	0.1070	0.6614	0.1153	0.6311
---	---	1637529_at	-0.1495	0.5150	0.0508	0.6243	0.0485	0.8841	0.0113	0.9872	-0.0073	0.9780	-0.0185	0.9278	-0.0143	0.9939	0.0880	0.8868	0.1023	0.8538

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1637530_at	0.0967	0.7121	-0.0455	0.8212	-0.1131	0.4814	0.0025	0.9978	-0.0532	0.8619	-0.0557	0.8346	0.2014	0.7893	-0.0043	0.9950	-0.2057	0.5444
CG1440 /// DsimCG1440	CG1440	1637531_at	0.1286	0.6536	1.1058	0.0457	1.4926	0.0001	-0.2082	0.7768	-0.8748	0.0117	-0.6666	0.0233	-0.4396	0.6702	0.1530	0.7591	0.5927	0.1806
DppIII	Dipeptidyl aminop	1637532_s_at	0.6463	0.0170	0.9232	0.1118	0.9941	0.0005	0.2289	0.5998	0.3278	0.1341	0.0989	0.6640	0.0853	0.9611	0.4533	0.3446	0.3681	0.4690
---	---	1637533_at	0.1169	0.5773	0.1302	0.4645	0.1646	0.3195	-0.0168	0.9852	-0.0001	0.9997	0.0167	0.9482	-0.0904	0.8810	0.1207	0.6270	0.2111	0.3719
olf186-F	lethal (2) k11505	1637534_at	-1.3237	0.0047	-2.4140	0.0027	-1.9140	0.0054	0.4050	0.7595	1.1748	0.0444	0.7698	0.1274	-0.0173	0.9928	0.0804	0.9036	0.0977	0.8646
---	---	1637535_at	-0.5353	0.2587	-0.6597	0.0843	0.0921	0.7597	-0.2561	0.8034	-0.0527	0.9274	0.2035	0.6276	-0.9997	0.3381	-0.4122	0.4619	0.5874	0.3019
CG14303	CG14303	1637536_at	0.3045	0.1343	-0.2813	0.2152	-0.2647	0.0972	-0.1105	0.8358	0.4281	0.0400	0.5386	0.0095	-0.0688	0.9516	-0.0356	0.9492	0.0331	0.9444
Lar	bypass	1637537_at	0.1173	0.8738	0.4012	0.2085	-0.0290	0.9483	-0.3751	0.7117	-0.0426	0.9509	0.3324	0.4432	0.0323	0.9908	0.3610	0.5523	0.3287	0.5988
CG9485	CG9485	1637538_s_at	1.8610	0.0025	0.5546	0.6512	1.0189	0.0034	0.7217	0.1811	-0.3118	0.3489	-1.0335	0.0044	0.2087	0.9620	-1.6147	0.1739	-1.8234	0.1603
hbs	hibris	1637539_a_at	-0.1503	0.7781	-0.4790	0.1386	-0.6586	0.0863	-0.0307	0.9879	0.0182	0.9801	0.0489	0.9341	-0.2254	0.9095	-0.5365	0.4475	-0.3111	0.6896
CG3123	CG3123	1637540_at	0.1906	0.3332	-0.0150	0.8875	-0.0182	0.9358	0.0245	0.9745	0.0257	0.9220	0.0013	0.9958	-0.1229	0.7758	-0.1338	0.4828	-0.0109	0.9696
Fs	dFollistatin	1637541_at	0.7507	0.0318	2.3274	0.0427	1.6253	0.0195	0.5400	0.2705	0.2954	0.3122	-0.2446	0.3580	1.7017	0.3660	2.0351	0.0637	0.3334	0.7780
---	---	1637542_s_at	0.5640	0.0201	0.3421	0.3733	0.0403	0.8865	-0.2091	0.7091	0.9793	0.0031	1.1884	0.0008	0.1739	0.8755	0.6393	0.1389	0.4654	0.2967
CG30036	CG30036	1637543_at	0.1024	0.5642	0.0074	0.9488	0.0120	0.9533	-0.2085	0.6060	-0.1902	0.3587	0.0182	0.9425	-0.0821	0.9030	-0.1302	0.6214	-0.0481	0.8870
---	---	1637544_s_at	0.0620	0.7680	0.0173	0.9327	0.1012	0.7173	0.0365	0.9563	0.2427	0.1757	0.2062	0.1998	-0.1475	0.8235	0.0984	0.7692	0.2459	0.3828
CG1837	CG1837	1637545_at	-0.2951	0.1089	-0.0701	0.5401	-0.0469	0.7802	-0.0259	0.9672	-0.2186	0.1814	-0.1928	0.1891	-0.1774	0.7997	0.0245	0.9591	0.2019	0.5042
RpLP0	Apurinic endonucl	1637546_at	-0.0051	0.9777	0.1366	0.3591	0.1516	0.3014	0.0448	0.9228	-0.1182	0.4117	-0.1629	0.1911	0.0175	0.9860	-0.0750	0.7897	-0.0925	0.7154
CG8199	CG8199	1637547_at	-0.0357	0.8530	0.2160	0.3849	0.2912	0.1764	-0.1495	0.6919	-0.2548	0.1430	-0.1052	0.5373	-0.3145	0.7106	-0.0824	0.8686	0.2321	0.5335
arg	arginase	1637548_at	3.2583	0.0012	1.5223	0.1917	3.3499	0.0001	1.3570	0.1378	0.6847	0.2144	-0.6722	0.1733	-0.4454	0.8628	-1.1124	0.2668	-0.6669	0.5430
CG12680	CG12680	1637549_at	0.4360	0.0613	-0.0125	0.9486	0.0331	0.8772	-0.1570	0.6576	0.0911	0.6307	0.2481	0.1021	-0.1468	0.8141	-0.1798	0.4983	-0.0330	0.9264
desat2	desaturase2	1637550_at	0.3656	0.4044	0.3037	0.2630	0.3465	0.1009	-0.0712	0.9517	-0.1351	0.7203	-0.0638	0.8658	0.1894	0.8331	-0.1469	0.7361	-0.3364	0.3803
CG1887	CG1887	1637551_at	0.1425	0.5936	0.1250	0.5779	0.2247	0.2589	0.0523	0.9311	-0.1302	0.4852	-0.1825	0.2544	0.0042	0.9984	0.0139	0.9865	0.0096	0.9878
shep	alan shepard	1637552_s_at	0.1987	0.7145	-0.7548	0.0100	-1.6140	0.0015	-0.2983	0.6010	1.5630	0.0007	1.8613	0.0002	0.4692	0.8122	0.6390	0.4323	0.1698	0.8779
CG34408	CG2967	1637553_at	-0.0222	0.9210	0.1139	0.4332	0.1714	0.2603	0.0550	0.9325	-0.0552	0.8174	-0.1102	0.5629	-0.2214	0.7324	-0.0156	0.9738	0.2058	0.4730
Dp	lethal(2)49Fk	1637554_a_at	0.3947	0.1419	1.5451	0.0258	1.2074	0.0001	-0.0770	0.9154	-0.5540	0.0230	-0.4770	0.0269	0.1217	0.9277	0.5882	0.1668	0.4666	0.2993
mtTFB1	Mitochondrial Trar	1637555_at	-0.4526	0.0366	-0.2939	0.1926	-0.4856	0.0111	-0.0029	0.9956	0.0687	0.6807	0.0716	0.6321	0.1040	0.8906	0.1167	0.7205	0.0127	0.9777
CG11857	CG11857	1637556_at	0.6611	0.0086	1.3585	0.0142	1.2665	0.0001	0.0374	0.9353	0.0080	0.9699	-0.0294	0.8525	0.0441	0.9804	0.6718	0.0966	0.6277	0.1399
CG10724	CG10724	1637557_a_at	-1.1179	0.0065	-0.0520	0.7932	-0.2557	0.1339	-0.0283	0.9675	-0.8708	0.0012	-0.8425	0.0009	0.1377	0.8736	0.2634	0.4382	0.1256	0.7492
CG8594	CG8594	1637558_at	0.2189	0.2252	-0.5152	0.0829	-0.4252	0.0597	-0.1080	0.8493	0.7586	0.0039	0.8666	0.0013	-0.1867	0.8270	0.0197	0.9739	0.2065	0.5867
CG13601	CG13601	1637559_at	-0.3533	0.1319	-0.7667	0.0100	-0.7688	0.0014	-0.0629	0.9043	0.2067	0.2239	0.2695	0.0780	-0.0722	0.9487	-0.2572	0.4421	-0.1850	0.6084
eIF-4G	translation initiatio	1637560_at	0.5451	0.0508	0.6485	0.0964	0.0530	0.7834	-0.2099	0.5987	0.4394	0.0358	0.6493	0.0040	0.2553	0.8049	0.5369	0.1957	0.2816	0.5334
lectin-33A	lectin-33A	1637561_at	1.9389	0.1103	2.3376	0.0095	2.0943	0.0105	-0.6273	0.7631	-1.0398	0.2449	-0.4125	0.6551	-0.1358	0.9816	-0.4744	0.7562	-0.3386	0.8348
GCR(ich)	GCR(ich)	1637562_at	-0.3716	0.6258	-0.0600	0.7386	-1.1719	0.0227	-1.2588	0.1119	-1.4121	0.0095	-0.1533	0.7684	-0.3838	0.8461	-1.2579	0.1248	-0.8741	0.2960
---	---	1637563_at	0.0992	0.6217	0.1917	0.1674	-0.0772	0.6774	-0.0381	0.9558	-0.0595	0.7940	-0.0214	0.9252	0.2393	0.6955	-0.0154	0.9710	-0.2547	0.3248
CG5382	CG5382	1637564_a_at	0.3484	0.1398	0.3886	0.0625	0.1986	0.2465	0.1028	0.7596	0.5632	0.0029	0.4604	0.0042	0.2404	0.7726	0.5585	0.1226	0.3181	0.3848
CG12645	CG12645	1637565_at	0.0017	0.9934	-0.0559	0.7146	0.1283	0.5592	0.1243	0.8216	0.1072	0.6587	-0.0172	0.9497	-0.0495	0.9421	0.0109	0.9765	0.0605	0.8178
---	---	1637566_at	0.0097	0.9612	0.0477	0.6033	-0.0280	0.9138	-0.1056	0.7768	-0.1213	0.4539	-0.0157	0.9351	-0.0329	0.9737	0.0011	0.9991	0.0341	0.9197
CG30360	CG30360	1637567_at	0.2197	0.3830	-0.1007	0.6533	0.0487	0.7626	0.4038	0.5500	0.2861	0.4286	-0.1178	0.7575	0.0863	0.9168	-0.1934	0.5141	-0.2797	0.3427
CG4060	CG4060	1637568_at	0.0686	0.8190	0.1856	0.3341	0.1005	0.5824	-0.0949	0.8604	-0.0344	0.8965	0.0606	0.7781	-0.0304	0.9677	-0.0027	0.9950	0.0277	0.9194
BobA /// CG13465	CG13465 /// Broth	1637569_s_at	-0.0895	0.7608	0.1483	0.4986	0.1802	0.3146	0.0544	0.9569	-0.1122	0.7223	-0.1666	0.5275	0.0523	0.9515	0.0440	0.9120	-0.0083	0.9847
CG4679	CG4679	1637570_at	0.2501	0.1112	0.2821	0.5836	0.2861	0.1018	0.0108	0.9894	0.2215	0.2367	0.2107	0.2077	0.0052	0.9984	0.2699	0.6408	0.2647	0.6443
CG11577	CG11577	1637571_at	0.7194	0.0156	0.7235	0.1202	0.2983	0.1881	-0.2062	0.7753	0.6376	0.0412	0.8438	0.0078	0.2929	0.8062	0.5597	0.2437	0.2668	0.6189
---	---	1637572_at	0.1662	0.4886	0.0020	0.9899	0.1240	0.6316	0.2471	0.5357	0.1272	0.5720	-0.1199	0.5602	-0.0180	0.9875	-0.1280	0.6536	-0.1100	0.7087
---	---	1637573_at	0.1600	0.4786	0.0934	0.5990	0.4008	0.0312	0.2246	0.4704	0.0380	0.8635	-0.1866	0.2132	-0.0335	0.9816	0.0475	0.9259	0.0810	0.8443
Acf1	Chromatin Access	1637574_at	0.0639	0.8218	-0.1413	0.6286	-0.2912	0.0697	-0.1878	0.6790	0.4447	0.0423	0.6325	0.0058	-0.1256	0.8940	0.1783	0.6408	0.3039	0.3967
CG8028	CG8028	1637575_at	0.0809	0.7695	-0.0682	0.6913	0.0257	0.9281	0.0505	0.9558	-0.1544	0.5614	-0.2049	0.3680	0.1062	0.8692	-0.0650	0.8533	-0.1712	0.5232
CG8839	CG8839	1637576_s_at	0.2567	0.3778	1.0638	0.0286	1.1888	0.0019	0.0771	0.9257	-0.5793	0.0278	-0.6564	0.0102	-0.0045	0.9984	0.3092	0.4861	0.3137	0.4851
Zip3	Zinc/iron regulate	1637577_at	-0.6158	0.5655	-2.3829	0.0188	-1.9535	0.0062	0.0104	0.9938	2.4319	0.0001	2.4215	0.0001	-0.2393	0.9672	0.7961	0.6310	1.0354	0.5175
Set	Set	1637578_at	-0.0380	0.8709	0.6682	0.0792	0.7589	0.0021	0.0498	0.9507	-0.2998	0.1777	-0.3496	0.0818	0.0955	0.9342	0.4916	0.1744	0.3962	0.3006
CG32792	CG32792	1637579_at	-0.1083	0.4531	0.0938	0.5557	0.0346	0.8468	-0.0195	0.9760	-0.0634	0.7372	-0.0439	0.8075	0.0705	0.9243	0.0535	0.8803	-0.0170	0.9628
CG14000	CG14000	1637580_at	0.0397	0.7994	-0.1541	0.5102	0.1024	0.6601	0.1596	0.7630	0.1308	0.5973	-0.0288	0.9175	-0.0966	0.8991	-0.0161	0.9732	0.0804	0.8182
lola	longitudinals abse	1637581_at	-0.3968	0.2439	-0.0330	0.9508	-0.2053	0.4724	-0.2246	0.8115	-0.2793	0.4739	-0.0548	0.9030	-0.0346	0.9898	0.0025	0.9994	0.0371	0.9666
Nedd4	Nedd4	1637582_a_at	-0.7363	0.1535	0.1790	0.5508	-0.8147	0.0027	-0.2821	0.6990	-0.1160	0.7739	0.1662	0.6262	0.6950	0.6695	0.6906	0.2748	-0.0044	0.9974
CG17385 /// DyakCG17385	CG17385	1637583_at	-0.3771	0.0971	0.4965	0.0165	0.3278	0.1990	0.2464	0.4191	-0.7576	0.0016	-1.0040	0.0003	0.2601	0.6960	-0.0146	0.9766	-0.2746	0.3439
pgant3	polypeptide GalN/	1637584_at	0.9716	0.0031	0.5200	0.1337	0.5564	0.1630	0.1750	0.5859	0.8611	0.0008	0.6860	0.0013	0.1885	0.9246	0.3618	0.6152	0.1733	0.8393
CG12278	CG12278	1637585_at	0.0341	0.8864	0.0475	0.7957	0.0014	0.9964	0.1192	0.8294	0.0863	0.7279	-0.0329	0.8976	0.2353	0.8270	0.0677	0.9230	-0.1676	0.7456
---	---	1637586_at	0.1618	0.2991	0.2954	0.1703	0.3062	0.0850	0.0674	0.9029	-0.0071	0.9795	-0.0745	0.6920	0.0757	0.9411	0.1927	0.5625	0.1170	0.7492
CG16716	CG16716	1637587_at	0.1857	0.5117	0.0439	0.7701	-0.0360	0.9098	0.0007	0.9994	0.0416	0.8765	0.0409	0.8616	-0.0239	0.9841	-0.0664	0.8611	-0.0425	0.9096
kl-3	male fertility factor	1637588_at	-0.1495	0.4361	-0.0089	0.9337	-0.0275	0.8952	0.1975	0.6380	0.0895	0.7061	-0.1080	0.6016	-0.0160	0.9877	0.0903	0.7421	0.1063	0.6781
CG15423	CG15423	1637589_at	0.2738	0.2721	0.0454	0.7840	0.2388	0.3530	0.2604	0.6327	0.2289	0.4044	-0.0315	0.9235	0.0559	0.9657	-0.0589	0.9124	-0.1148	0.7802
CG5156	CG5156	1637590_at	-0.5518	0.4393	1.0714	0.3525	0.2555	0.5989	-0.1176	0.9141	0.1806	0.6385	0.2982	0.3433	0.6399	0.8672	1.8440	0.2106	1.2041	0.4405
---	---	1637591_at	0.0152	0.9355	-0.1717	0.3650	-0.0434	0.8086	0.1395	0.7492	0.1691	0.3846	0.0296	0.8972	0.1164	0.8222	0.1125	0.6298	-0.0039	0.9921
CG10073	CG10073	1637592_at	-0.6849	0.0039	-0.1102	0.5185	-0.8523	0.0014	-0.6604	0.0855	-0.8618	0.0031	-0.2014	0.3245	-0.0543	0.9452	-0.2254	0.3340	-0.1711	0.4892
CG15537	CG15537	1637593_at	0.5074	0.3929	0.0501	0.8707	0.7751	0.0031	0.7028	0.3056	0.0002	0.9996	-0.7026	0.0558	0.0816	0.9701	-0.3225	0.5842	-0.4040	0.4825
CG15364	CG15364	1637594_at	-0.8252	0.1628	-0.2007	0.8120	-1.0509	0.0032	-0.3138	0.7631	-0.0704	0.9096	0.2434	0.5851	0.4340	0.8424	0.5191	0.5833	0.0851	0.9460
---	---	1637595_at	-0.1514	0.3888	0.1537	0.2554	0.2487	0.1042	0.0061	0.9941	-0.1066	0.5624	-0.1127	0.4903	0.0438	0.9521	0.2842	0.1801	0.2404	0.2863
---	---	1637596_at	0.0894	0.6227	0.0714	0.5825	0.1095	0.5319	-0.0945	0.8776	-0.1205	0.6010	-0.0260	0.9195	-0.0203	0.9846	-0.0697	0.8189	-0.0493	0.8764
CG11980	CG11980	1637597_a_at	0.3621	0.1403	0.4133	0.4100	0.4430	0.0127	0.0714	0.8930	0.0504	0.8195	-0.0211	0.9228	0.1139	0.9462	0.3243	0.5431	0.2104	0.7154
CG15715	CG15715	1637598_at	-0.2161	0.4976	0.6430	0.0373	0.1002	0.6382	-0.1435	0.7135	-0.4448	0.0208	-0.3013	0.0621	0.4887	0.6749	0.4765	0.3152	-0.0122	0.9874

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
l(3)mbt	tumor-supressor	1637599_at	-0.0044	0.9886	-0.1424	0.4399	-0.0429	0.8411	-0.0259	0.9805	0.1286	0.6621	0.1545	0.5436	0.0198	0.9898	-0.0912	0.8372	-0.1110	0.7761
CG14926	CG14926	1637600_at	0.2239	0.1551	0.0322	0.8932	-0.0221	0.9388	0.0320	0.9673	0.0679	0.7861	0.0358	0.8829	0.0562	0.9340	-0.1172	0.6118	-0.1733	0.4334
CG7739	CG7739	1637601_at	0.0274	0.9277	-0.2694	0.5737	-0.3252	0.0639	0.0233	0.9759	0.4131	0.0351	0.3899	0.0281	0.0683	0.9746	0.1180	0.8779	0.0497	0.9471
Edem1	CG3810	1637602_at	0.1476	0.5885	0.3515	0.2286	0.0366	0.8761	-0.2004	0.7031	0.1914	0.4468	0.3919	0.0725	0.0368	0.9831	0.3461	0.3802	0.3093	0.4568
---	---	1637603_at	0.1830	0.3954	-0.0407	0.7167	-0.1438	0.3357	0.0400	0.9312	0.0770	0.6092	0.0369	0.8107	0.1607	0.8236	-0.0952	0.8053	-0.2558	0.4037
CG18356	CG18356	1637604_at	0.1446	0.4215	0.1084	0.4200	0.1914	0.2558	-0.0071	0.9937	-0.0527	0.8197	-0.0456	0.8293	-0.0742	0.9514	-0.0674	0.9019	0.0068	0.9914
CG1146	CG1146	1637605_s_at	-1.3617	0.0023	-0.1470	0.8259	-1.1771	0.0243	-0.6069	0.2401	-1.2716	0.0022	-0.6647	0.0233	0.4653	0.8206	-0.0717	0.9590	-0.5370	0.5511
CG12795	CG12795	1637606_at	0.8106	0.0188	0.8269	0.0242	0.5271	0.0541	0.4411	0.2674	0.6869	0.0105	0.2458	0.2441	0.6456	0.3712	0.6707	0.0957	0.0251	0.9659
CG8027	CG8027	1637607_at	0.5499	0.0313	0.3624	0.1578	0.1984	0.2089	-0.1392	0.7850	0.3713	0.0793	0.5105	0.0146	0.0298	0.9775	0.1522	0.5597	0.1224	0.6476
Ppt1	Palmitoyl-protein t	1637608_at	0.1426	0.3146	0.1209	0.6610	0.1411	0.5719	-0.0160	0.9860	0.1189	0.6247	0.1349	0.5264	-0.1014	0.9246	-0.0891	0.8571	0.0124	0.9838
---	---	1637609_at	0.4080	0.0192	0.2460	0.1695	0.0341	0.9036	-0.0939	0.8650	-0.0077	0.9798	0.0862	0.6728	0.0521	0.9554	-0.0039	0.9943	-0.0560	0.8788
Chrac-16	Chrac-16	1637610_at	0.2410	0.1120	-0.1041	0.3495	0.1093	0.7197	0.2169	0.7599	0.4055	0.1811	0.1887	0.5249	-0.0208	0.9816	-0.0680	0.7753	-0.0472	0.8509
Yippee	Yippee	1637611_at	-0.0789	0.6789	-0.2458	0.0752	-0.0042	0.9880	-0.1256	0.7596	-0.4362	0.0215	-0.3107	0.0529	-0.3068	0.6272	-0.5437	0.0606	-0.2369	0.3800
CG6017	ankyrin domain	1637612_at	-0.9168	0.0093	-0.5870	0.1722	-0.8083	0.0031	-0.4574	0.1078	-0.3195	0.0619	0.1379	0.3768	-0.0781	0.9666	0.0065	0.9952	0.0846	0.9023
---	---	1637613_at	0.2464	0.3346	-0.0095	0.9606	0.3172	0.0462	0.0756	0.8507	0.0155	0.9418	-0.0601	0.6962	-0.1950	0.7220	-0.0807	0.7781	0.1143	0.6484
CG12413 /// CG40068	CG12413 /// CG4(	1637614_s_at	-0.0392	0.8987	0.3541	0.0461	0.4910	0.1206	-0.0397	0.9688	-0.4656	0.0803	-0.4259	0.0739	-0.1243	0.8298	0.0439	0.9062	0.1682	0.5034
CG13192	CG13192	1637615_at	0.5144	0.0716	0.5210	0.1325	0.8651	0.0010	0.0764	0.8738	0.0048	0.9846	-0.0717	0.6835	-0.1448	0.8745	0.0096	0.9906	0.1545	0.7026
---	---	1637616_at	0.0218	0.9509	0.1508	0.4523	-0.0562	0.7325	-0.0701	0.9507	-0.0990	0.7977	-0.0290	0.9409	0.3033	0.5519	0.1426	0.5636	-0.1606	0.5137
CG18190	CG18190	1637617_at	0.3456	0.4492	-0.2612	0.7726	-0.6420	0.3066	-0.8572	0.0437	0.8939	0.0036	1.7511	0.0002	-0.5545	0.9088	0.3097	0.9062	0.8642	0.6389
Hs2st	heparan-sulfate-2	1637618_at	0.8757	0.0019	0.5800	0.1110	0.5139	0.0200	0.0268	0.9622	0.4802	0.0071	0.4534	0.0055	0.0851	0.9515	0.2526	0.5679	0.1675	0.7272
Dscam	Down syndrome c	1637619_s_at	-1.6665	0.0009	-2.6040	0.0088	-2.8543	0.0000	0.2339	0.7190	0.7113	0.0239	0.4774	0.0722	0.2776	0.8202	-0.2945	0.5810	-0.5721	0.2728
CG9186	CG9186	1637620_s_at	0.5526	0.0718	0.7586	0.1882	0.8043	0.0012	0.0896	0.8544	0.1573	0.3826	0.0678	0.7176	0.0089	0.9967	0.3159	0.5805	0.3070	0.5967
CG15196	CG15196	1637621_at	0.0160	0.9346	0.0078	0.9653	-0.2082	0.1894	0.0806	0.8516	0.1562	0.3261	0.0756	0.6382	0.0363	0.9705	-0.0058	0.9924	-0.0421	0.9026
Cyp12a4	Cyp12a4	1637622_s_at	0.0523	0.9328	-1.8747	0.0121	-1.4367	0.0282	0.2049	0.9345	1.5205	0.0443	1.3156	0.0500	-0.2439	0.8650	-0.3956	0.5018	-0.1517	0.8338
CG14270	CG14270	1637623_at	-0.0418	0.9083	0.1124	0.8176	0.1955	0.4452	0.1312	0.7126	0.3912	0.0239	0.2599	0.0746	0.1051	0.9653	0.5232	0.4180	0.4181	0.5416
---	---	1637624_a_at	0.0328	0.9083	0.0796	0.5540	0.0360	0.8614	-0.0979	0.8293	-0.2410	0.1640	-0.1431	0.3737	0.0049	0.9970	0.0501	0.9197	0.0452	0.9186
CG32813	CG32813	1637625_s_at	-0.4658	0.4409	-0.0824	0.9102	-1.1040	0.0027	-0.4707	0.1119	-0.5303	0.0094	-0.0596	0.7583	0.6426	0.7953	-0.0317	0.9885	-0.6743	0.5399
CG15412	CG15412	1637626_at	-2.0215	0.0004	-1.7028	0.0959	-2.0090	0.0005	0.3639	0.4596	-0.6062	0.0305	-0.9701	0.0024	0.6529	0.7464	-0.3162	0.7563	-0.9691	0.2738
Ptp4E	Protein tyrosine pl	1637627_at	-1.3549	0.0016	-1.6015	0.0031	-1.5495	0.0001	-0.2757	0.3775	-0.2418	0.1688	0.0339	0.8706	-0.2980	0.7187	-0.5528	0.1126	-0.2548	0.4730
CG6550	CG6550	1637628_at	0.2429	0.3435	0.0629	0.8846	0.4522	0.0102	0.1128	0.8791	0.1322	0.6409	0.0194	0.9517	-0.2264	0.8012	0.0153	0.9829	0.2418	0.5403
ssh	MKP-like	1637629_a_at	-0.0211	0.9528	-0.5829	0.4758	-0.8230	0.0085	-0.1626	0.7923	0.4009	0.1092	0.5635	0.0203	-0.1000	0.9813	-0.2418	0.8444	-0.1417	0.9096
---	---	1637630_at	0.0457	0.8527	0.0009	0.9953	0.0075	0.9724	0.1296	0.8248	0.0413	0.8933	-0.0883	0.7136	-0.0886	0.8655	-0.1340	0.5398	-0.0454	0.8716
tutl	turtle	1637631_at	0.2713	0.1756	-0.1385	0.4126	0.2182	0.3042	0.2211	0.5793	0.3132	0.1230	0.0921	0.6630	-0.2442	0.7204	-0.2692	0.3285	-0.0250	0.9499
CG12194	CG12194	1637632_at	0.9112	0.3101	0.1593	0.4429	1.0926	0.0002	0.5928	0.5156	-0.1503	0.7995	-0.7430	0.0852	-0.2675	0.9235	-0.8409	0.3467	-0.5734	0.5547
CG4095	CG4095	1637633_at	-0.0094	0.9620	0.3412	0.0667	-0.0245	0.9263	-0.3409	0.4420	-0.4099	0.0911	-0.0690	0.7996	0.1435	0.8400	0.0434	0.9259	-0.1002	0.7722
CG10953	CG10953	1637634_at	0.2068	0.3075	0.0838	0.8292	0.5828	0.0155	0.1349	0.8776	0.0006	0.9988	-0.1343	0.6669	-0.1395	0.8270	-0.0541	0.8879	0.0854	0.7860
CG4766	CG4766	1637635_at	0.2207	0.3765	0.0850	0.5606	0.0314	0.8414	-0.1042	0.8310	0.0177	0.9498	0.1219	0.4962	0.0988	0.8846	0.0133	0.9766	-0.0855	0.7788
DopEcR	DopEcR	1637636_at	0.3288	0.1461	-0.2122	0.4217	-0.4948	0.0184	-0.1888	0.5932	0.3820	0.0393	0.5708	0.0042	0.0215	0.9831	-0.0050	0.9925	-0.0264	0.9360
shi	dynamin	1637637_at	-0.2600	0.2486	0.3957	0.3322	0.3882	0.0666	0.0556	0.9538	-0.2379	0.3742	-0.2934	0.2104	-0.0122	0.9939	0.3900	0.2686	0.4022	0.2855
CG10702 /// DyakCG10702	CG10702	1637638_s_at	0.3970	0.7365	-0.8691	0.0227	-0.6021	0.0104	0.2185	0.7000	0.0421	0.9075	-0.1764	0.4772	0.0209	0.9971	-1.2896	0.3549	-1.3105	0.3705
gsb	gooseberry	1637639_at	-2.5045	0.0013	-0.2755	0.2470	-1.9053	0.0024	-1.2783	0.1033	-2.1317	0.0015	-0.8534	0.0456	0.2046	0.8956	0.0960	0.9152	-0.1086	0.8918
CG32320	CG32320	1637640_at	-0.0023	0.9951	-0.1243	0.5308	0.1287	0.6078	-0.0549	0.9518	-0.0542	0.8703	0.0007	0.9981	-0.3344	0.6749	-0.1702	0.6239	0.1642	0.6366
Mkp3	MKP-like	1637641_at	-0.8276	0.0508	-1.3903	0.0095	-2.1698	0.0003	0.0636	0.9641	0.5679	0.1290	0.5043	0.1311	0.6060	0.5461	0.1536	0.7932	-0.4524	0.3468
CG15047	CG15047	1637642_at	0.4811	0.0094	0.0359	0.7803	0.2464	0.4650	0.0185	0.9774	-0.0174	0.9398	-0.0359	0.8462	-0.2145	0.8897	-0.1834	0.8055	0.0312	0.9724
FucTB	alpha1,3-fucosyltr	1637643_at	-0.1306	0.4539	-0.2189	0.5205	-0.4569	0.0723	-0.3898	0.2466	0.0216	0.9401	0.4114	0.0287	-0.2082	0.8320	-0.3105	0.4479	-0.1023	0.8452
CG13813	CG13813	1637644_at	1.0668	0.5467	2.9092	0.0323	4.9113	0.0000	-0.2298	0.8244	-0.5215	0.1912	-0.2918	0.4393	-1.9414	0.7307	1.3840	0.5897	3.3254	0.1932
miple2	miple2	1637645_at	-2.5843	0.0054	-3.4780	0.0014	-2.5305	0.0033	0.2618	0.8568	-0.1728	0.7845	-0.4346	0.3653	-0.0678	0.9848	-0.6299	0.4404	-0.5621	0.5083
---	---	1637646_at	0.0576	0.7256	0.0350	0.7194	0.0249	0.9008	-0.0212	0.9803	0.0469	0.8670	0.0681	0.7710	0.0288	0.9717	0.0486	0.8642	0.0198	0.9443
CG31148	CG31148	1637647_at	0.0020	0.9938	-0.0909	0.4741	0.0593	0.8596	0.0671	0.9029	0.1333	0.4736	0.0662	0.7276	-0.2005	0.8222	-0.1090	0.8225	0.0915	0.8480
CG11191	CG11191	1637648_at	-0.4210	0.0479	-0.0942	0.7750	0.0636	0.7978	-0.0168	0.9880	-0.5056	0.0567	-0.4888	0.0421	-0.1174	0.9092	-0.0515	0.9288	0.0659	0.8949
CG1707	CG1707	1637649_at	0.2665	0.2794	0.3379	0.1721	0.3600	0.2147	0.2205	0.7490	-0.0410	0.9222	-0.2615	0.3411	0.1136	0.9235	0.0308	0.9620	-0.0829	0.8746
CG4164	CG4164	1637650_at	1.1191	0.0067	0.4529	0.0447	0.8289	0.0022	0.2605	0.5757	1.5778	0.0004	1.3173	0.0004	-0.0603	0.9679	0.9458	0.0391	1.0061	0.0457
CG4004	CG4004	1637651_at	-0.1915	0.2711	-0.2183	0.2496	-0.0591	0.7851	0.0701	0.9110	0.0705	0.7669	0.0005	0.9986	-0.0450	0.9588	-0.0721	0.8185	-0.0271	0.9360
CG30222	CG30222	1637652_a_at	0.2370	0.1631	-0.2310	0.3433	0.0835	0.6824	0.1266	0.8162	0.2358	0.2724	0.1092	0.6143	-0.1680	0.8192	-0.0660	0.8785	0.1020	0.7744
---	---	1637653_at	0.1936	0.3500	0.0604	0.6809	0.1218	0.5300	0.0645	0.9314	0.1035	0.6824	0.0390	0.8815	-0.0245	0.9775	-0.0696	0.7870	-0.0450	0.8699
CG10566	CG10566	1637654_at	-0.4774	0.1045	-0.6865	0.0444	-0.5021	0.0612	0.4196	0.4751	0.3389	0.2867	-0.0806	0.8219	0.1399	0.8650	-0.0781	0.8657	-0.2180	0.5271
CG40341	CG40341	1637655_s_at	-0.0720	0.6608	0.0689	0.6276	0.0423	0.8600	0.0199	0.9777	-0.1090	0.5671	-0.1289	0.4369	0.1214	0.8564	0.0820	0.8185	-0.0395	0.9173
---	---	1637656_at	-0.0580	0.8129	-0.0767	0.6303	0.1306	0.5538	0.0014	0.9988	-0.0980	0.6800	-0.0994	0.6423	-0.1662	0.8270	0.0246	0.9624	0.1908	0.5693
---	---	1637657_at	0.0471	0.7949	-0.0214	0.9099	0.3210	0.1226	0.0574	0.9009	-0.0152	0.9452	-0.0725	0.6343	-0.0947	0.9076	-0.0032	0.9961	0.0915	0.7933
mthl3	mth-like 3	1637658_at	-0.5521	0.0151	0.2787	0.5677	0.0923	0.5844	-0.2792	0.4500	-0.2300	0.2531	0.0492	0.8296	0.0151	0.9939	0.7043	0.1135	0.6892	0.1433
CG18146	CG18146	1637659_a_at	-1.3348	0.0027	-0.7245	0.0678	-1.1423	0.0034	-0.1816	0.8337	-0.7143	0.0362	-0.5327	0.0716	-0.1307	0.8740	-0.2068	0.5404	-0.0761	0.8570
CG5150	CG5150	1637660_at	0.3488	0.3827	0.1266	0.3372	0.0295	0.9046	0.0617	0.9386	0.0911	0.7422	0.0293	0.9187	0.0154	0.9939	-0.1503	0.7932	-0.1657	0.7526
---	---	1637661_at	0.2400	0.3276	0.0089	0.9475	0.3227	0.0525	-0.1552	0.7409	-0.1668	0.4362	-0.0116	0.9648	-0.1986	0.7707	-0.2663	0.3496	-0.0677	0.8599
msta	msta	1637662_at	-2.4630	0.0018	-1.2801	0.0207	-1.8566	0.0000	-0.2761	0.5242	-1.5805	0.0003	-1.3043	0.0004	-0.0691	0.9741	-0.7245	0.1512	-0.6553	0.2235
Tbp	42kDa polypeptide	1637663_at	0.2982	0.1006	0.5260	0.1309	0.4120	0.0279	-0.0894	0.8794	-0.1741	0.4015	-0.0848	0.6893	0.1620	0.8395	0.1752	0.6209	0.0132	0.9810
CG1340	CG1340	1637664_at	-0.0355	0.8748	0.2434	0.0755	0.1048	0.5404	-0.0920	0.8680	-0.2467	0.2064	-0.1547	0.3959	0.0423	0.9489	0.1151	0.5818	0.0729	0.7492
CG31367	CG31367	1637665_at	-0.4225	0.3358	-0.1047	0.6512	-0.1598	0.4906	-0.1320	0.8478	0.0841	0.7831	0.2160	0.3497	-0.1772	0.8940	0.4064	0.4094	0.5836	0.2561
CG15651	CG15651	1637666_at	-0.1721	0.4528	0.0272	0.8164	-0.0532	0.8182	-0.0088	0.9943	-0.1381	0.6477	-0.1293	0.6390	-0.0922	0.8890	-0.0459	0.9044	0.0463	0.8921
CG17295	CG17295	1637667_at	0.5022	0.0293	0.6620	0.1272	0.2508	0.4387	-0.2043	0.7654	0.0944	0.7917	0.2987	0.2555	0.3030	0.7644	0.2659	0.5529	-0.0371	0.9512

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG31101	CG31101	1637668_at	0.0926	0.6508	-0.0173	0.8643	0.0532	0.8061	-0.0875	0.8073	-0.0193	0.9228	0.0683	0.6414	-0.0832	0.9011	-0.0734	0.8185	0.0098	0.9810
CG7442	CG7442	1637669_at	-1.6334	0.0008	-0.1704	0.6399	-0.9869	0.0069	-1.0308	0.0654	-1.5806	0.0011	-0.5499	0.0585	-0.2822	0.8141	-0.1717	0.7787	0.1105	0.8669
CHES-1-like	Checkpoint suppr	1637670_s_at	0.2651	0.2189	0.2272	0.3735	0.1712	0.4307	-0.0486	0.9584	-0.0904	0.7578	-0.0419	0.8859	0.1060	0.9174	-0.0564	0.9182	-0.1624	0.6761
CG14767	CG14767	1637671_a_at	0.0481	0.7578	-0.1161	0.5392	-0.0980	0.5074	-0.0724	0.8498	-0.0885	0.5549	-0.0161	0.9249	-0.1786	0.7464	-0.3619	0.1212	-0.1833	0.4411
CG15919	CG15919	1637672_at	-0.0629	0.7603	0.2737	0.1852	0.2725	0.3770	-0.1995	0.7581	-0.2226	0.4407	-0.0231	0.9479	-0.0755	0.9225	0.1716	0.5199	0.2471	0.3508
Nup58	Nup58	1637673_at	-0.4968	0.0619	-0.4615	0.2176	-0.3678	0.2087	0.0582	0.9252	0.1771	0.3499	0.1189	0.5079	-0.1740	0.9095	0.1465	0.8426	0.3204	0.5809
Est-P	Esterase P	1637674_at	-0.0618	0.7453	-0.0439	0.7193	-0.0061	0.9779	-0.2613	0.4456	-0.2473	0.1834	0.0139	0.9534	-0.1990	0.7436	-0.2681	0.2807	-0.0691	0.8307
---	---	1637675_at	-0.0215	0.9314	-0.0434	0.7623	0.1536	0.3040	0.0562	0.9228	-0.0441	0.8455	-0.1002	0.5617	-0.0068	0.9959	0.0913	0.8180	0.0981	0.7872
CG5482	CG5482	1637676_at	0.3346	0.0887	0.5602	0.1930	0.3114	0.1095	-0.1375	0.7213	-0.2063	0.2362	-0.0688	0.7091	0.1341	0.8973	0.0828	0.8821	-0.0513	0.9231
ftz	fushi-tarazu	1637677_at	-0.0392	0.8890	-0.0462	0.7722	0.0964	0.6624	0.1551	0.7838	0.1694	0.4881	0.0144	0.9606	-0.0179	0.9853	0.0354	0.9191	0.0533	0.8547
CG10340	anon-fast-evolvin	1637678_at	-0.2043	0.3521	0.1726	0.3799	-0.1220	0.5705	-0.0541	0.9311	-0.1879	0.3085	-0.1338	0.4362	0.2834	0.7485	0.1398	0.7542	-0.1436	0.7409
UbcD2	Ubiquitin conjugat	1637679_s_at	0.3200	0.2585	0.0824	0.5537	-0.1597	0.4061	-0.0508	0.9375	0.3372	0.0766	0.3880	0.0293	0.3082	0.7464	0.1547	0.7421	-0.1535	0.7392
Tsp42Eq	tetraspanin 42E	1637680_at	0.5616	0.6884	0.4547	0.4347	1.2212	0.0108	-0.0986	0.9794	-1.2246	0.1897	-1.1261	0.1787	-0.5671	0.8427	-1.0503	0.3623	-0.4832	0.7154
CG3253	L-gene	1637681_at	-0.0680	0.7583	-0.0190	0.8530	0.0828	0.6052	-0.0409	0.9504	-0.0494	0.8309	-0.0085	0.9704	0.0219	0.9823	0.0902	0.7351	0.0683	0.8083
Traf2	TNF-receptor-ass	1637682_at	-0.2283	0.2428	-0.1198	0.5207	-0.3203	0.0450	0.0331	0.9603	0.1973	0.2680	0.1642	0.3082	0.1606	0.7810	0.2800	0.2430	0.1194	0.6577
---	---	1637683_at	0.0195	0.9297	0.0429	0.8413	0.0532	0.8003	-0.1037	0.9149	-0.0375	0.9298	0.0662	0.8507	-0.0742	0.8680	0.0022	0.9955	0.0764	0.6981
unc-104	unc-104	1637684_at	-0.3622	0.1441	0.1018	0.6305	-0.2363	0.3025	-0.2417	0.6041	-0.2474	0.2963	-0.0057	0.9850	-0.0060	0.9970	0.1488	0.7439	0.1548	0.7253
---	---	1637685_at	0.4114	0.0394	-0.0168	0.8703	0.1656	0.3339	0.2295	0.4690	0.2659	0.1181	0.0364	0.8545	0.0757	0.9309	-0.0245	0.9587	-0.1002	0.7577
---	---	1637686_at	0.0365	0.8923	-0.1255	0.3640	0.1916	0.2628	0.1383	0.7424	-0.0662	0.7654	-0.2045	0.2170	-0.0134	0.9935	-0.0508	0.9325	-0.0373	0.9433
ct	cut	1637687_at	-1.2842	0.0045	-1.2498	0.0698	-1.7661	0.0007	-0.5579	0.4420	-0.0983	0.8482	0.4596	0.1932	-0.2038	0.7485	-0.0670	0.8580	0.1368	0.6389
---	---	1637688_at	0.0032	0.9897	0.0044	0.9713	0.2889	0.1767	0.0195	0.9833	-0.0227	0.9407	-0.0422	0.8643	-0.1890	0.7220	-0.0164	0.9648	0.1726	0.4597
CG5317 /// DyakCG5317	CG5317	1637689_at	0.1659	0.4969	0.2382	0.4271	0.4386	0.0158	0.1440	0.7333	-0.4207	0.0327	-0.5647	0.0056	-0.0004	0.9999	-0.4159	0.2430	-0.4155	0.2774
CG10357	CG10357	1637690_at	0.0554	0.7886	-0.0135	0.8975	-0.0827	0.6305	-0.0709	0.8735	0.0055	0.9809	0.0764	0.6280	0.0587	0.9499	-0.0628	0.8750	-0.1215	0.6955
CG4570	CG4570	1637691_at	0.0459	0.9284	-0.5915	0.6038	-1.0238	0.2211	-0.8008	0.0657	1.3348	0.0008	2.1356	0.0001	-0.4390	0.9460	0.5176	0.8466	0.9566	0.6524
---	---	1637692_at	-0.0680	0.7179	-0.0607	0.5728	-0.1899	0.3572	0.0522	0.9345	0.1931	0.3072	0.1410	0.4216	0.0792	0.8906	0.0267	0.9402	-0.0525	0.8515
CG34409	CG31411	1637693_at	0.3483	0.0602	0.1809	0.2266	0.1952	0.2374	-0.0542	0.9005	-0.0437	0.7998	0.0105	0.9533	0.1149	0.8097	0.0655	0.7925	-0.0494	0.8461
---	---	1637694_at	0.0320	0.8728	0.1532	0.5164	0.0908	0.5774	-0.0605	0.9011	-0.0502	0.7947	0.0103	0.9589	-0.1191	0.8751	0.0186	0.9699	0.1377	0.6719
---	---	1637695_at	0.1050	0.5491	-0.0082	0.9771	-0.0648	0.7604	-0.0929	0.8590	0.0745	0.7372	0.1674	0.3307	0.0584	0.9589	0.0162	0.9776	-0.0422	0.9237
---	---	1637696_at	0.1475	0.5471	0.1509	0.3407	0.3464	0.0940	0.2229	0.4847	0.1329	0.4531	-0.0900	0.5984	0.0637	0.9460	0.1426	0.6450	0.0789	0.8255
---	---	1637697_at	0.0042	0.9883	-0.1534	0.4679	-0.3061	0.2832	0.0614	0.9532	0.1493	0.6445	0.0879	0.7851	0.0648	0.9635	0.0387	0.9505	-0.0261	0.9633
CG30339	CG30339	1637698_at	-0.1971	0.4038	0.0647	0.7127	0.0740	0.7608	-0.0997	0.8999	-0.2009	0.4537	-0.1012	0.7116	-0.0832	0.8940	-0.0758	0.7963	0.0074	0.9847
---	---	1637699_at	0.0716	0.7403	0.1399	0.4924	0.1810	0.4222	-0.1314	0.7349	-0.2240	0.1917	-0.0926	0.5917	0.1637	0.7979	0.1187	0.6929	-0.0450	0.9028
eIF2B-delta	eIF2B-delta	1637700_s_at	0.2608	0.4437	0.4757	0.2804	0.7711	0.0029	0.3503	0.2067	0.3293	0.0532	-0.0210	0.9185	0.0277	0.9918	0.4380	0.4976	0.4103	0.5358
Cpr49Ab	CG30042	1637701_at	0.1681	0.8827	0.0962	0.8313	0.1217	0.4150	0.1505	0.9494	-0.7197	0.2787	-0.8702	0.1392	0.1301	0.9752	-0.7350	0.4868	-0.8651	0.4098
CG13307	CG13307	1637702_at	-0.0749	0.6469	0.1581	0.4009	0.0653	0.8003	-0.0152	0.9805	-0.0782	0.6495	-0.0629	0.6950	0.0149	0.9914	0.1763	0.5712	0.1614	0.6129
Socs36E	Suppressor of cyt	1637703_a_at	-0.6089	0.4185	-0.5122	0.5175	-0.7889	0.2366	0.5348	0.4908	1.3964	0.0055	0.8616	0.0277	0.7355	0.8386	1.5112	0.2974	0.7757	0.6294
---	---	1637704_at	0.1995	0.3661	-0.1108	0.3808	0.0829	0.5755	0.2906	0.3166	0.3022	0.0765	0.0116	0.9572	0.0214	0.9816	-0.0707	0.7719	-0.0921	0.6734
ewg	erect wing	1637705_at	0.2771	0.3758	0.4962	0.1652	-0.1087	0.5714	-0.2461	0.5242	-0.0728	0.7641	0.1734	0.3510	0.2953	0.7810	0.0228	0.9782	-0.2726	0.5767
CG7330	CG7330	1637706_at	0.2084	0.4528	0.1759	0.1601	0.1782	0.5330	-0.0576	0.9413	-0.0832	0.7552	-0.0256	0.9253	0.0715	0.9405	0.0515	0.9087	-0.0201	0.9630
CG4009	CG4009	1637707_at	0.1049	0.4708	-0.1496	0.4493	0.1667	0.2430	0.1089	0.8424	0.2120	0.3015	0.1032	0.6167	-0.1636	0.7633	-0.1018	0.6883	0.0618	0.8287
CG17672	CG17672	1637708_a_at	0.0741	0.6314	-0.0435	0.6908	0.2181	0.2269	0.2245	0.3836	0.2317	0.1102	0.0072	0.9688	-0.1082	0.8202	0.0995	0.6414	0.2077	0.3027
CG31797	CG31797	1637709_at	0.0273	0.9393	0.1728	0.3598	0.2557	0.3620	0.1003	0.8350	-0.0832	0.6904	-0.1835	0.2629	0.1231	0.9092	0.0590	0.9214	-0.0640	0.9032
kst	beta[[H]]-spectrin	1637710_at	-0.0406	0.9410	0.0244	0.9582	-0.1660	0.4178	0.7688	0.1976	0.7182	0.0515	-0.0506	0.9078	1.0861	0.2740	0.9477	0.0761	-0.1384	0.8287
CG13592	CG13592	1637711_at	0.2124	0.1983	0.0814	0.7077	0.0674	0.7557	-0.0549	0.9380	0.0703	0.7795	0.1252	0.5361	-0.1536	0.8076	-0.1794	0.5093	-0.0258	0.9435
CG30007	CG30007	1637712_at	0.2446	0.3607	-0.3894	0.3507	0.0017	0.9950	0.2267	0.7023	0.7035	0.0181	0.4768	0.0548	-0.2479	0.8222	-0.0548	0.9409	0.1931	0.7095
PH4alphaNE3	prolyl-4-hydroxyla	1637713_at	0.1253	0.5240	-0.0362	0.8608	0.0938	0.5950	0.0826	0.9314	0.2065	0.4910	0.1239	0.6787	-0.0941	0.9081	-0.0061	0.9925	0.0879	0.8032
Ance-3	Ance-3	1637714_a_at	-0.9057	0.0513	0.7455	0.1669	0.3615	0.5052	-0.5359	0.5933	-1.2386	0.0232	-0.7028	0.1212	-0.1595	0.9523	0.4461	0.6021	0.6056	0.4655
metl	methyltransferase	1637715_a_at	0.3107	0.2005	0.9076	0.0155	1.0256	0.0019	0.1679	0.7415	-0.2031	0.3756	-0.3711	0.0674	-0.0035	0.9984	0.4179	0.1948	0.4214	0.2217
sqd	RNA-binding prote	1637716_a_at	0.8646	0.0161	0.1307	0.3690	-0.3568	0.0699	-0.1296	0.7028	0.8794	0.0007	1.0090	0.0002	0.4489	0.7070	0.3093	0.5538	-0.1396	0.8238
sev	sevenless	1637717_at	-1.3187	0.0472	-0.4518	0.1549	-0.3321	0.2875	0.3139	0.6736	-0.9438	0.0153	-1.2577	0.0026	-0.0414	0.9916	-0.0686	0.9646	-0.0271	0.9852
CG4820	CG4820	1637718_at	-0.2706	0.1836	0.2447	0.2377	-0.2224	0.3153	0.0105	0.9922	-0.2870	0.1933	-0.2974	0.1317	0.2825	0.7686	0.1779	0.6987	-0.1046	0.8407
CG16793	CG16793	1637719_at	0.1181	0.4574	0.1878	0.4126	0.0909	0.5305	-0.0549	0.9191	-0.0776	0.6821	-0.0227	0.9100	0.2370	0.6695	0.0422	0.9009	-0.1948	0.3921
CG6856	CG6856	1637720_at	0.1936	0.3613	0.5975	0.1934	0.3909	0.0340	-0.0505	0.9586	-0.0049	0.9899	0.0457	0.8804	0.1042	0.9206	0.3996	0.2404	0.2954	0.4093
CG3301	CG3301	1637721_at	0.8333	0.2314	-0.1516	0.3952	-0.2038	0.3762	-0.1058	0.8938	0.2359	0.3811	0.3417	0.1454	-0.2336	0.9330	-0.8635	0.3187	-0.6299	0.4938
---	---	1637722_at	0.3715	0.0970	0.4198	0.0447	0.4127	0.0606	-0.1757	0.5778	-0.1698	0.2980	0.0058	0.9773	-0.0545	0.9495	-0.1494	0.5823	-0.0949	0.7484
mod(mdg4)	Modifier67.2	1637723_at	-0.0772	0.7051	-0.1336	0.2150	-0.4983	0.0286	-0.3295	0.2640	0.0096	0.9717	0.3391	0.0365	0.0328	0.9816	-0.2123	0.4996	-0.2452	0.4351
CG11905	CG11905	1637724_s_at	0.0846	0.8056	0.2033	0.5025	0.0368	0.8965	-0.1396	0.8546	-0.2287	0.4216	-0.0891	0.7692	0.0711	0.9644	0.0320	0.9643	-0.0391	0.9486
CG1756	CG1756	1637725_at	-1.5128	0.0062	-0.0758	0.8118	-0.3002	0.3561	-0.1368	0.9311	-1.3706	0.0102	-1.2338	0.0099	-0.0537	0.9761	0.0015	0.9995	0.0552	0.9264
Mmp2	Matrix metalloprot	1637726_at	0.0020	0.9936	-0.0298	0.7660	0.1274	0.5475	-0.0217	0.9766	0.0709	0.7419	0.0926	0.6166	-0.1231	0.8454	0.0112	0.9817	0.1343	0.6310
CG7049	CG7049	1637727_at	-0.4188	0.3132	0.2006	0.4192	0.3297	0.0738	0.0667	0.8810	-0.5090	0.0057	-0.5757	0.0020	-0.1322	0.9238	0.2190	0.6673	0.3512	0.4620
CG8508	CG8508	1637728_at	0.3990	0.0782	-0.0507	0.7680	-0.0486	0.8523	-0.0676	0.9375	0.1216	0.6708	0.1892	0.4211	0.0178	0.9908	-0.2127	0.5232	-0.2305	0.4870
nompC	no mechanorecep	1637729_at	0.2698	0.1329	-0.1796	0.4063	0.1311	0.4556	0.2490	0.5341	0.5373	0.0184	0.2883	0.1215	-0.3652	0.5929	-0.2010	0.5199	0.1642	0.6140
CG17361	CG17361	1637730_at	0.2907	0.3463	0.2754	0.4129	0.1811	0.5983	-0.3618	0.6615	0.2445	0.5671	0.6063	0.0869	-0.4157	0.6897	0.0515	0.9402	0.4672	0.2841
---	---	1637731_at	0.3524	0.1078	0.4265	0.0373	0.3905	0.0606	-0.2194	0.6338	-0.2874	0.2025	-0.0679	0.7882	-0.0605	0.9530	0.0200	0.9684	0.0805	0.8354
Gr98a	Gustatory recepto	1637732_at	0.0793	0.6407	-0.0482	0.8217	0.1643	0.3553	0.2333	0.4716	0.2588	0.1365	0.0255	0.9034	-0.1046	0.8661	-0.0518	0.8850	0.0527	0.8745
CG31063	CG31063	1637733_at	-0.6284	0.0197	0.9209	0.0435	1.0651	0.0005	-0.0872	0.8589	-1.2158	0.0003	-1.1286	0.0002	-0.3468	0.7644	0.2309	0.6679	0.5778	0.2532
Tepl	Thiolester contain	1637734_at	0.0738	0.7573	0.0533	0.6130	0.0042	0.9848	-0.0592	0.9314	-0.0736	0.7622	-0.0144	0.9557	-0.0461	0.9589	-0.0300	0.9409	0.0162	0.9659
MrgBP	MrgBP	1637735_at	0.3835	0.0415	0.4121	0.0500	0.3539	0.0861	0.0581	0.9098	-0.1391	0.4064	-0.1972	0.1751	0.1274	0.8305	-0.0427	0.9111	-0.1700	0.5131
GalNAc-T2	polypeptide GalN/	1637736_a_at	-0.5148	0.1183	-0.0682	0.9101	0.0359	0.8686	-0.2019	0.6558	0.0881	0.7299	0.2900	0.1351	-0.2934	0.8378	0.6399	0.2591	0.9334	0.1409

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	**Gene Title**	**Affy ID**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**	**FC**	**FDR**
l(1)G0230	lethal (1) G0230	1637737_at	-0.1371	0.5797	0.7094	0.0752	0.8237	0.0032	0.0958	0.8915	-1.2547	0.0008	-1.3505	0.0004	0.0527	0.9780	-0.3360	0.4694	-0.3887	0.4037
Srp14	Srp14	1637738_at	0.2569	0.2697	0.6117	0.0540	0.8961	0.0045	0.2816	0.4221	0.0824	0.7136	-0.1991	0.2534	0.1057	0.9499	0.5682	0.2408	0.4624	0.3695
CG16721	CG16721	1637739_at	-0.4764	0.1114	0.2289	0.4918	0.3588	0.4716	-0.1750	0.8274	-1.8373	0.0005	-1.6623	0.0005	-0.2625	0.8953	-0.8898	0.2167	-0.6274	0.4069
CG11404	CG11404	1637740_at	0.0372	0.8836	-0.0332	0.9230	-0.0599	0.8676	0.3105	0.4174	0.2293	0.2841	-0.0812	0.7199	0.1637	0.8609	0.2041	0.6152	0.0404	0.9394
CG30284	CG30284	1637741_a_at	-0.1862	0.3224	-0.0035	0.9885	-0.0101	0.9603	-0.0095	0.9915	-0.0766	0.7272	-0.0671	0.7403	0.0103	0.9939	0.0768	0.8588	0.0666	0.8743
---	---	1637742_at	0.0581	0.7529	0.0430	0.7631	0.1530	0.3693	0.0173	0.9846	0.0801	0.7391	0.0628	0.7821	-0.1643	0.7726	-0.1156	0.6729	0.0486	0.8864
CG9461	CG9461	1637743_at	1.0737	0.0812	0.6774	0.0597	0.1150	0.5928	-0.2670	0.7419	0.9380	0.0167	1.2050	0.0033	0.3187	0.8806	0.5071	0.5501	0.1885	0.8591
CG30365	CG30365	1637744_at	0.0177	0.9212	0.0492	0.7492	0.1674	0.2651	0.0448	0.9441	-0.0138	0.9598	-0.0587	0.7712	-0.0580	0.9277	-0.0694	0.7916	-0.0114	0.9720
CG14891	CG14891	1637745_a_at	0.0145	0.9410	0.0000	1.0000	0.0478	0.8231	-0.1447	0.7773	0.0630	0.8112	0.2078	0.2814	-0.0211	0.9831	-0.0163	0.9658	0.0048	0.9903
CG18545	CG18545	1637746_at	0.2305	0.1159	0.2310	0.2995	0.3499	0.0517	-0.0148	0.9860	-0.1147	0.5877	-0.0999	0.6112	0.0123	0.9901	-0.0541	0.8591	-0.0664	0.8044
CG8841	CG8841	1637747_s_at	0.2910	0.2754	0.5015	0.0273	0.3946	0.0406	-0.1855	0.6006	-0.0588	0.7867	0.1267	0.4516	0.1442	0.8472	0.3419	0.2518	0.1978	0.5464
---	---	1637748_at	0.1054	0.5462	0.0679	0.6631	0.1281	0.3835	-0.0307	0.9603	-0.1102	0.5319	-0.0795	0.6386	0.0145	0.9914	-0.0111	0.9841	-0.0256	0.9494
D2R	Dopamine 2-like r	1637749_at	-0.2581	0.3012	-0.4550	0.0515	-0.2720	0.2395	0.2659	0.4798	0.2057	0.3124	-0.0603	0.7894	-0.1872	0.7220	-0.1743	0.4379	0.0128	0.9704
Sox21b	sox-like	1637750_at	-2.8752	0.0004	-0.2857	0.5285	-1.3633	0.0228	-1.4916	0.1081	-2.8980	0.0008	-1.4064	0.0117	-0.0876	0.9396	-0.1583	0.6908	-0.0707	0.8846
CG32563	CG32563	1637751_at	-0.0147	0.9418	0.0146	0.9722	0.3802	0.0492	0.1201	0.8605	-0.1175	0.6743	-0.2376	0.2840	-0.2396	0.7485	0.0034	0.9965	0.2430	0.4606
CG9363	CG9363	1637752_a_at	0.0854	0.7159	-0.0551	0.5862	0.1140	0.5491	0.1254	0.8045	-0.2107	0.3021	-0.3362	0.0663	0.0949	0.9125	-0.1282	0.7093	-0.2231	0.4783
CG14748	CG14748	1637753_at	-1.3519	0.0032	1.2494	0.3935	-0.9219	0.1022	-0.8401	0.2789	-0.6772	0.1368	0.1629	0.7406	1.3921	0.7215	1.8167	0.2489	0.4246	0.8375
CG9458	CG9458	1637754_at	0.6250	0.6857	-0.8091	0.6748	1.6683	0.0595	1.6073	0.3710	-0.0562	0.9715	-1.6634	0.0696	-0.4552	0.9520	-1.3077	0.5860	-0.8526	0.7442
CG10987	CG10987	1637755_at	0.0383	0.8636	-0.6371	0.1503	-0.4416	0.0764	0.1508	0.9075	0.6759	0.1046	0.5251	0.1586	-0.0312	0.9851	-0.0405	0.9478	-0.0093	0.9875
CG12960	CG12960	1637756_at	0.1517	0.3791	-0.1134	0.4612	-0.1198	0.4209	0.1701	0.7064	0.3778	0.0704	0.2077	0.2648	-0.0412	0.9729	-0.0981	0.7971	-0.0570	0.8913
CG10280	CG10280	1637757_at	0.1002	0.5147	0.2595	0.3716	0.2326	0.1705	-0.1124	0.7803	-0.1010	0.5783	0.0115	0.9573	0.0546	0.9643	0.0598	0.9054	0.0053	0.9924
CG7737	CG7737	1637758_at	-0.0451	0.8828	-0.4756	0.1570	-0.4656	0.2199	0.4073	0.4596	0.8539	0.0115	0.4466	0.0940	0.1819	0.9340	0.2509	0.7692	0.0690	0.9435
---	---	1637759_at	-0.0252	0.8810	0.0501	0.5978	-0.0182	0.9417	0.0262	0.9703	-0.0244	0.9220	-0.0506	0.8007	0.2567	0.6960	0.1588	0.5901	-0.0979	0.7614
---	---	1637760_at	-0.0810	0.5953	0.2416	0.3385	0.0176	0.9195	-0.2636	0.4126	-0.1678	0.3558	0.0958	0.5912	0.1877	0.7644	0.3542	0.1722	0.1666	0.5578
CG31878	CG31878	1637761_at	0.0170	0.9289	0.0554	0.7631	-0.0142	0.9539	0.0060	0.9940	0.0905	0.6142	0.0845	0.6072	-0.1226	0.7990	-0.0459	0.8724	0.0766	0.7410
---	---	1637762_s_at	0.3814	0.0636	-0.1099	0.4558	-0.0097	0.9628	-0.0391	0.9656	0.1791	0.4759	0.2181	0.3191	-0.0463	0.9390	-0.1601	0.3820	-0.1137	0.5649
CG8298 /// DyakCG8298	CG8298	1637763_a_at	-0.1137	0.6166	-0.3788	0.0463	-0.2833	0.2137	-0.1876	0.6364	0.5330	0.0141	0.7207	0.0022	-0.3472	0.6749	0.1600	0.6691	0.5071	0.1588
CG9806	CG9806	1637764_at	0.0750	0.6406	0.0804	0.7833	0.1355	0.5103	0.4074	0.5419	0.2038	0.5895	-0.2035	0.5509	0.1467	0.7707	0.1043	0.6576	-0.0425	0.8857
CG14147	CG14147	1637765_at	0.0195	0.9444	0.0974	0.4287	0.1969	0.2387	0.1145	0.7803	-0.0619	0.7620	-0.1764	0.2491	0.1566	0.7597	0.0810	0.7499	-0.0756	0.7615
CG10252 /// DyakCG10252	CG10252	1637766_at	-0.0466	0.8566	-0.0436	0.7241	-0.1222	0.7225	0.1716	0.7389	0.2195	0.3426	0.0479	0.8565	0.0075	0.9946	-0.0576	0.8754	-0.0651	0.8427
CG3305 /// DyakCG3305	CG3305	1637767_at	-0.0943	0.5678	0.0454	0.7324	-0.1117	0.4297	-0.1081	0.7327	-0.0963	0.5206	0.0118	0.9476	0.0309	0.9734	0.0014	0.9989	-0.0294	0.9256
CG30100	CG30100	1637768_at	-0.0524	0.8734	0.3203	0.1216	0.4400	0.0910	0.1593	0.7815	-0.3868	0.1036	-0.5461	0.0185	0.1688	0.8202	0.0259	0.9590	-0.1429	0.6669
CkIIbeta	casein kinase 2	1637769_s_at	0.4424	0.0694	0.8186	0.0882	0.6821	0.0029	0.0963	0.9011	-0.2010	0.4403	-0.2973	0.1848	0.3056	0.7230	0.2765	0.4622	-0.0291	0.9566
CG11505	CG11505	1637770_a_at	0.1533	0.4892	0.2421	0.2445	-0.0140	0.9610	-0.3363	0.5068	0.0051	0.9899	0.3414	0.1539	0.0160	0.9914	-0.0530	0.9129	-0.0691	0.8694
CG7220	CG7220	1637771_s_at	0.4411	0.1198	-0.4539	0.0798	-0.7788	0.0047	-0.2595	0.4081	0.1245	0.4968	0.3840	0.0220	0.0444	0.9816	-0.7854	0.0912	-0.8298	0.1011
CG4726	CG4726	1637772_at	-0.8074	0.3691	-1.7313	0.0185	-1.8948	0.0056	-0.2047	0.8611	-0.5678	0.1736	-0.3631	0.3452	-0.0380	0.9946	-1.6139	0.2086	-1.5759	0.2523
Baldspot	baldspot	1637773_s_at	-0.0341	0.9124	0.0051	0.9882	-0.0993	0.7382	0.0151	0.9858	0.5719	0.0107	0.5568	0.0073	0.1435	0.9016	0.4972	0.2254	0.3537	0.4120
CG8389	CG8389	1637774_s_at	0.5895	0.0185	0.8428	0.0395	1.4523	0.0020	-0.1535	0.6921	-0.7359	0.0024	-0.5824	0.0039	-0.8449	0.3564	-0.4887	0.3034	0.3563	0.4822
PpD5	protein phosphata	1637775_at	0.1164	0.4399	0.1032	0.3823	0.0882	0.6212	-0.0395	0.9479	-0.1098	0.5480	-0.0703	0.6940	0.0517	0.9296	0.1272	0.5141	0.0755	0.7249
CG6444	CG6444	1637776_at	0.0096	0.9596	0.0190	0.8630	-0.1355	0.5516	0.1468	0.6972	0.1459	0.4157	-0.0009	0.9968	0.1552	0.8689	0.2348	0.5439	0.0796	0.8736
CG13982	CG13982	1637777_at	0.0002	0.9994	0.0586	0.5632	0.1388	0.4325	-0.0123	0.9880	-0.2710	0.1548	-0.2587	0.1287	-0.0281	0.9816	-0.0426	0.9221	-0.0145	0.9732
CG1969	CG1969	1637778_a_at	2.4618	0.0005	1.7479	0.0014	1.8661	0.0000	0.1583	0.8250	0.5966	0.0380	0.4383	0.0784	-0.0756	0.9195	-0.1389	0.6093	-0.0633	0.8462
CG17341	CG17341	1637779_at	0.3303	0.2379	0.2009	0.2502	0.1441	0.4318	-0.0104	0.9884	0.0578	0.7795	0.0682	0.7046	0.0334	0.9831	-0.0908	0.8550	-0.1242	0.7699
CG4836	CG4836	1637780_a_at	0.0643	0.6991	-0.0341	0.7721	0.1004	0.5877	0.0484	0.9311	0.0589	0.7664	0.0105	0.9597	0.0610	0.9112	0.0443	0.8734	-0.0167	0.9513
ald	altered disjunction	1637781_at	-0.1995	0.4321	-0.0724	0.7159	-0.5145	0.0800	-0.7020	0.1345	-0.1260	0.7000	0.5761	0.0307	-0.2685	0.8270	0.0029	0.9989	0.2713	0.6260
CG12020	CG12020	1637782_at	0.1647	0.4762	0.0256	0.8128	0.0743	0.7687	-0.0985	0.8676	0.0698	0.7833	0.1684	0.3854	-0.2005	0.8062	-0.1357	0.7331	0.0648	0.8905
CG12077	CG12077	1637783_at	0.1663	0.3369	0.0048	0.9666	0.2160	0.1687	-0.0549	0.9218	0.0627	0.7545	0.1176	0.4601	-0.2793	0.5765	-0.1500	0.5246	0.1293	0.5969
CG7669	CG7669	1637784_at	0.0926	0.6975	0.1451	0.3605	0.2678	0.1818	0.0885	0.9138	0.0412	0.9084	-0.0473	0.8773	-0.1983	0.7230	-0.0269	0.9451	0.1714	0.4913
L	Lobe	1637785_at	-0.4396	0.0898	0.9513	0.0221	0.7980	0.0062	-0.2955	0.5552	-1.2014	0.0014	-0.9059	0.0028	0.0123	0.9946	0.2416	0.5875	0.2293	0.6129
CG40323	CG40323	1637786_s_at	0.4435	0.0263	-0.7782	0.0717	-0.6848	0.0889	0.0352	0.9838	1.0487	0.0179	1.0135	0.0130	-0.2338	0.7644	-0.2358	0.4875	-0.0021	0.9977
CG8233	CG8233	1637787_a_at	-0.0134	0.9528	-0.0130	0.9781	-0.1865	0.3241	-0.0110	0.9893	0.0982	0.6423	0.1091	0.5568	0.1495	0.8651	0.1016	0.8255	-0.0480	0.9211
CG31176	CG31176	1637788_at	-2.3987	0.0012	-4.2577	0.0048	-4.2126	0.0001	0.4081	0.6720	1.0016	0.0353	0.5934	0.1450	0.2039	0.9277	-0.4640	0.5448	-0.6679	0.3762
CG31772	CG31772	1637789_at	-0.1978	0.3533	0.1988	0.1001	-0.0936	0.7019	-0.1318	0.8589	-0.2510	0.3564	-0.1192	0.6671	0.0750	0.8825	0.0359	0.9036	-0.0391	0.8831
CG12250	CG12250	1637790_a_at	0.1606	0.4472	0.0197	0.8961	0.3010	0.1705	0.1776	0.7136	0.2196	0.3264	0.0419	0.8718	-0.1453	0.8395	-0.0236	0.9617	0.1216	0.7154
CG31922	CG31922	1637791_at	-0.7173	0.0139	-0.2821	0.4200	-0.1958	0.4769	-0.0090	0.9956	-0.0401	0.9298	-0.0312	0.9385	-0.0430	0.9816	0.4684	0.2735	0.5115	0.2651
A3-3	A3-3	1637792_at	-0.8994	0.0041	-0.9695	0.0480	-1.2736	0.0004	-0.0176	0.9838	0.0457	0.8638	0.0633	0.7778	0.2722	0.7822	0.0433	0.9499	-0.2288	0.6171
CG9791	CG9791	1637793_at	0.3206	0.0614	0.2468	0.5162	0.3124	0.0409	0.0972	0.8074	0.4182	0.0166	0.3210	0.0313	-0.0588	0.9636	0.2930	0.3992	0.3518	0.3271
CG4927	CG4927	1637794_at	-1.2092	0.0157	-1.1009	0.0052	-1.3087	0.0001	-0.0029	0.9962	-0.2148	0.2953	-0.2119	0.2457	-0.0796	0.9589	-0.4361	0.3098	-0.3566	0.4317
CG7778 /// DyakCG7778	CG7778	1637795_at	-2.2824	0.0127	-1.7935	0.1502	-2.9709	0.0017	-0.9645	0.1412	-1.4677	0.0037	-0.5032	0.1551	0.3932	0.9390	-0.8873	0.5988	-1.2806	0.4301
---	---	1637796_at	0.0932	0.5389	-0.1045	0.5764	-0.0528	0.8582	-0.0706	0.9248	0.0168	0.9604	0.0874	0.7138	-0.0747	0.9365	-0.2070	0.4954	-0.1323	0.6861
CG14400	CG14400	1637797_at	2.2921	0.0012	2.3982	0.0677	3.4781	0.0000	0.0571	0.9695	-1.0128	0.0167	-1.0699	0.0080	-0.6513	0.7823	-0.3765	0.7573	0.2748	0.8307
CG15717	CG15717	1637798_at	1.4051	0.0006	1.9520	0.0067	2.5589	0.0000	0.3011	0.3571	-0.4729	0.0196	-0.7739	0.0014	-0.3337	0.6660	0.1119	0.7684	0.4457	0.1792
r-l	UMP synthase	1637799_at	0.3502	0.1219	0.3959	0.2338	-0.0144	0.9645	0.2134	0.7333	0.3118	0.2665	0.0984	0.7444	0.5578	0.5967	0.1190	0.8511	-0.4388	0.3576
CG3347	CG3347	1637800_at	-0.0439	0.8175	0.1536	0.3528	-0.0603	0.8054	-0.0478	0.9619	-0.0845	0.7949	-0.0367	0.9089	0.1735	0.8270	0.0946	0.8304	-0.0789	0.8555
se	sepia	1637801_at	0.0428	0.8676	0.0566	0.7708	0.0454	0.8178	-0.0067	0.9937	0.1402	0.4439	0.1469	0.3667	-0.1745	0.8206	-0.1309	0.7205	0.0436	0.9211
---	---	1637802_at	0.2087	0.3117	-0.0231	0.8185	0.1955	0.2787	-0.0260	0.9639	0.0848	0.6115	0.1108	0.4365	-0.0328	0.9800	0.0166	0.9734	0.0494	0.9025
---	---	1637803_s_at	0.1362	0.4772	0.1128	0.4322	0.1090	0.4835	-0.0268	0.9610	-0.0963	0.5385	-0.0695	0.6436	0.0418	0.9653	-0.0544	0.8850	-0.0963	0.7484
CG18673	CG18673	1637804_at	0.9729	0.4257	0.4465	0.1587	0.1395	0.7615	-0.2608	0.6597	0.1398	0.6653	0.4006	0.1140	0.1107	0.9862	-0.4418	0.8232	-0.5525	0.7513
---	---	1637805_at	0.0940	0.7186	0.0049	0.9660	-0.0436	0.7826	-0.0603	0.9220	-0.0020	0.9941	0.0583	0.7776	-0.0245	0.9848	-0.0925	0.8086	-0.0680	0.8613

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Ada	Adenosine deamir	1637806_at	0.4497	0.1349	0.6181	0.0941	0.8961	0.0009	0.0295	0.9759	-0.3351	0.1590	-0.3646	0.0881	-0.1178	0.9238	0.0551	0.9325	0.1730	0.7032
inaC	protein kinase C	1637807_at	0.1490	0.4730	0.2678	0.2257	0.1455	0.4307	0.0008	0.9990	0.0244	0.9175	0.0236	0.9078	0.1789	0.8134	0.1033	0.7927	-0.0756	0.8523
CG6151	CG6151	1637808_a_at	0.1308	0.8231	1.3156	0.0906	1.7216	0.0003	-0.1229	0.8822	-0.9293	0.0063	-0.8064	0.0072	-0.3879	0.8461	0.4168	0.6421	0.8048	0.3446
CG18341	CG18341	1637809_at	-0.0556	0.7824	-0.0117	0.9147	0.2441	0.2743	-0.0410	0.9538	-0.2043	0.2969	-0.1633	0.3605	-0.0431	0.9555	0.0341	0.9230	0.0772	0.7708
CG5780	CG5780	1637810_at	0.5999	0.0072	0.3334	0.2941	0.4674	0.0256	-0.1104	0.8856	0.3176	0.2217	0.4280	0.0689	-0.1565	0.7893	0.0121	0.9784	0.1686	0.5161
CG5273	CG5273	1637811_s_at	-2.9410	0.0049	-1.1146	0.2496	-1.1878	0.0077	0.0009	0.9994	-1.9567	0.0009	-1.9576	0.0005	0.2053	0.9646	-0.0350	0.9900	-0.2403	0.8886
CG30430	CG30430	1637812_at	0.0302	0.8478	-0.0691	0.5846	0.1695	0.3669	-0.0968	0.8735	0.0001	0.9996	0.0969	0.6578	-0.1599	0.8065	-0.0563	0.8866	0.1037	0.7422
abd-A	abdominal-A	1637813_at	-0.7259	0.0186	-0.1005	0.7685	-0.4711	0.0618	-0.4646	0.2642	-0.4134	0.0934	0.0512	0.8585	0.0271	0.9898	0.2841	0.5515	0.2569	0.5995
Jafrac2	jafrac2	1637814_s_at	0.4466	0.0619	3.1275	0.0021	2.4369	0.0001	0.4876	0.3443	-0.4747	0.1091	-0.9623	0.0040	1.0330	0.1628	2.1487	0.0018	1.1156	0.0333
Cka	connector of kinas	1637815_s_at	0.3193	0.2723	0.3675	0.3280	0.2672	0.1320	-0.0144	0.9893	0.3453	0.1613	0.3596	0.1036	0.0872	0.9514	0.3778	0.3604	0.2906	0.5050
---	---	1637816_s_at	-0.1970	0.2458	0.0981	0.8624	0.1974	0.5485	-0.1223	0.9029	-0.9837	0.0091	-0.8614	0.0100	-0.2888	0.8427	-0.9452	0.1197	-0.6564	0.2869
---	---	1637817_at	0.0370	0.8838	0.2170	0.3312	0.1842	0.4347	0.0858	0.8748	0.0286	0.9140	-0.0572	0.7871	0.0257	0.9816	-0.0059	0.9925	-0.0316	0.9321
CG6902	CG6902	1637818_at	0.0668	0.8161	0.0104	0.9396	0.0925	0.7774	0.0153	0.9901	0.1078	0.7399	0.0926	0.7583	-0.2200	0.7220	-0.0295	0.9450	0.1905	0.4825
CG6192	CG6192	1637819_at	-0.0763	0.8975	0.0258	0.9360	-0.5743	0.0106	-0.3959	0.4094	0.0443	0.9044	0.4401	0.0675	0.0949	0.9653	0.1711	0.8244	0.0762	0.9249
X11Lbeta	X11Lbeta	1637820_at	0.0644	0.8730	-0.4665	0.1795	-0.5884	0.0269	-0.0041	0.9961	0.3759	0.1699	0.3801	0.1211	-0.1694	0.9036	-0.3275	0.5363	-0.1581	0.7967
CG32023	CG32023	1637821_at	0.2046	0.6243	0.0030	0.9810	-0.1043	0.5603	0.0692	0.9488	0.0014	0.9976	-0.0678	0.8456	0.2709	0.6955	-0.1802	0.5457	-0.4512	0.1422
CG17904	CG17904	1637822_at	0.2685	0.1072	-0.3797	0.1937	-0.1841	0.4200	0.1042	0.8595	0.5409	0.0189	0.4366	0.0288	-0.1404	0.8608	-0.0097	0.9886	0.1308	0.7192
CG1434	CG1434	1637823_at	-0.6412	0.0230	-0.1044	0.8258	0.3428	0.4942	0.3498	0.3443	-0.2811	0.1848	-0.6309	0.0060	-0.2435	0.9064	-0.0069	0.9973	0.2366	0.7907
CG13653	CG13653	1637824_at	0.0414	0.8249	0.2095	0.1947	0.2452	0.1544	-0.0616	0.9029	-0.0816	0.6595	-0.0200	0.9209	-0.0911	0.8909	0.0212	0.9590	0.1123	0.6848
CG11127	CG11127	1637825_at	-0.0088	0.9683	0.5190	0.0230	0.7080	0.0112	0.1500	0.7271	-0.4279	0.0335	-0.5780	0.0056	-0.0643	0.9619	0.0267	0.9652	0.0910	0.8514
CG10623	selenocysteine me	1637826_at	-0.0264	0.9365	0.0768	0.8132	0.2078	0.3146	-0.5216	0.1883	-1.7602	0.0003	-1.2386	0.0006	-0.6681	0.5461	-1.4820	0.0276	-0.8139	0.1428
GlcAT-I	GlcAT-I	1637827_at	0.5693	0.0164	0.8247	0.0329	0.5663	0.0135	-0.1397	0.8016	-0.1919	0.4029	-0.0522	0.8396	0.0633	0.9363	-0.0136	0.9747	-0.0769	0.8001
---	---	1637828_a_at	0.1266	0.5444	-0.0457	0.8233	-0.1845	0.2166	0.0790	0.9116	0.3260	0.1454	0.2471	0.2198	0.0475	0.9653	0.0126	0.9821	-0.0349	0.9330
CG4020	CG4020	1637829_at	0.8130	0.5510	-2.1181	0.3811	1.8869	0.0394	2.5171	0.1884	1.3509	0.2448	-1.1662	0.2643	-1.5340	0.8049	-1.6871	0.5329	-0.1531	0.9695
CG31145	CG31145	1637830_s_at	-1.4048	0.0011	-1.0749	0.1359	-1.1142	0.0001	-0.2395	0.4553	-0.2179	0.2083	0.0216	0.9193	-0.1023	0.9643	0.1299	0.8838	0.2322	0.7439
CG30071	CG30071	1637831_at	0.1753	0.3856	0.1954	0.3389	0.2028	0.2354	-0.0332	0.9649	-0.1440	0.4940	-0.1108	0.5796	-0.0391	0.9717	0.0631	0.8718	0.1022	0.7463
---	---	1637832_at	0.0181	0.9509	-0.0498	0.6485	-0.1767	0.4731	0.0032	0.9956	0.1626	0.4063	0.1594	0.3637	0.1082	0.8875	0.1059	0.7567	-0.0024	0.9965
Jhe	JH-esterase	1637833_at	-0.2348	0.8947	1.4428	0.2345	1.9739	0.0227	1.6658	0.5255	2.0260	0.1392	0.3602	0.8172	1.0434	0.7707	3.8870	0.0365	2.8436	0.1000
exex	extra-extra	1637834_at	0.0308	0.8897	0.0125	0.9545	0.2949	0.1343	0.2182	0.5744	0.1023	0.6477	-0.1159	0.5542	-0.1963	0.8215	-0.1149	0.8045	0.0814	0.8646
CG12177	CG12177	1637835_at	-0.3038	0.3481	-0.3007	0.5250	-0.5859	0.0056	-0.2292	0.7028	-0.5670	0.0457	-0.3378	0.1703	-0.0468	0.9831	-0.7478	0.1398	-0.7010	0.1975
---	---	1637836_at	0.2138	0.1623	0.3038	0.1365	0.0158	0.9405	-0.1594	0.6462	-0.2057	0.2204	-0.0463	0.8063	0.0796	0.8692	-0.0062	0.9875	-0.0858	0.6877
CG14115	CG14115	1637837_at	-0.0236	0.8908	0.0398	0.8546	-0.0767	0.7350	0.0823	0.9182	0.0255	0.9438	-0.0568	0.8414	0.0540	0.9441	0.0773	0.7906	0.0233	0.9431
CG34380	CG13988	1637838_at	-0.5712	0.0440	0.0407	0.6962	-0.4453	0.0180	-0.4027	0.3237	-0.8247	0.0048	-0.4220	0.0513	0.1250	0.8541	-0.0403	0.9275	-0.1653	0.5794
Spase25	Spase 25-subunit	1637839_at	0.8778	0.0038	1.6736	0.0078	1.6669	0.0001	0.2347	0.4356	-0.0544	0.7867	-0.2892	0.0546	-0.0718	0.9503	0.3933	0.2367	0.4651	0.1990
---	---	1637840_at	-0.0202	0.9156	0.0083	0.9414	0.0982	0.6590	0.0676	0.8915	-0.0235	0.9200	-0.0911	0.5837	0.0053	0.9952	-0.0117	0.9739	-0.0169	0.9548
CG17985	CG17985	1637841_at	0.1080	0.5610	0.6129	0.0521	0.6141	0.0020	-0.0200	0.9777	-0.2147	0.2239	-0.1947	0.2187	0.0819	0.9066	0.3481	0.1512	0.2662	0.2993
CG14531	CG14531	1637842_at	-0.0722	0.7283	-0.0286	0.7880	0.0260	0.8843	0.0302	0.9633	-0.0636	0.7595	-0.0938	0.5889	0.0068	0.9952	-0.0649	0.8721	-0.0717	0.8430
RfaBp	retinoid-fatty acid	1637843_at	1.2020	0.0089	0.3464	0.0987	0.8121	0.0008	0.1107	0.9011	0.8820	0.0085	0.7713	0.0094	-0.3655	0.6557	0.0021	0.9989	0.3676	0.2822
CG11584	CG11584	1637844_at	0.1943	0.2273	0.0768	0.6028	-0.0640	0.7405	-0.1634	0.6576	0.0361	0.8784	0.1995	0.2072	0.3132	0.7493	0.0871	0.8853	-0.2261	0.6166
CG31230	CG31230	1637845_at	-0.0544	0.7987	-0.0433	0.7802	-0.0226	0.9030	0.1167	0.8143	0.2459	0.2074	0.1292	0.4908	0.1188	0.7945	0.1405	0.4694	0.0217	0.9351
---	---	1637846_at	0.2285	0.3487	-0.0375	0.8880	-0.0978	0.7201	0.0948	0.9376	0.2116	0.5825	0.1169	0.7639	-0.0012	0.9996	0.0497	0.9058	0.0510	0.8921
CG5362	CG5362	1637847_at	0.2077	0.2529	0.3837	0.1374	0.5278	0.0266	-0.1486	0.8156	-0.7108	0.0120	-0.5622	0.0203	-0.2841	0.7230	-0.6251	0.0809	-0.3410	0.3332
CG6873	Adf/cofilin-like	1637848_at	-0.3583	0.1357	0.0186	0.9515	-0.0756	0.6914	0.0822	0.9254	-0.0871	0.7864	-0.1692	0.5064	0.2389	0.7462	0.1554	0.6429	-0.0835	0.8305
OdsH	Ods-site homeobc	1637849_at	0.2629	0.3454	0.0941	0.4841	0.1730	0.4212	0.0988	0.8735	0.1854	0.4025	0.0867	0.7031	0.2147	0.7215	0.0421	0.9151	-0.1726	0.5153
Su(H)	suppressor of hair	1637850_at	0.2783	0.2514	0.0719	0.6523	0.1824	0.3122	0.1003	0.8446	0.2502	0.1819	0.1499	0.3922	0.0282	0.9816	0.0920	0.7824	0.0638	0.8555
CG7422	CG7422	1637851_at	0.5034	0.4950	-0.3477	0.0667	-0.2109	0.2250	0.2119	0.6628	0.2466	0.2935	0.0348	0.9008	-0.0225	0.9952	-0.6462	0.4729	-0.6237	0.4970
CG12442	CG12442	1637852_at	0.1266	0.5444	-0.0186	0.9318	-0.0614	0.8275	0.0051	0.9956	0.2198	0.3418	0.2147	0.2972	-0.0718	0.9400	0.1171	0.7319	0.1889	0.5431
Hk	Hyperkinetic	1637853_a_at	-1.9203	0.0007	-1.9169	0.0028	-2.1235	0.0002	-0.0765	0.9507	0.1634	0.6767	0.2399	0.4612	0.0348	0.9726	-0.0787	0.8117	-0.1135	0.6872
CG18335	CG18335	1637854_at	-0.1539	0.5080	0.1103	0.3258	-0.0290	0.8832	0.0252	0.9778	-0.1463	0.5618	-0.1716	0.4349	0.2546	0.7361	0.0680	0.8836	-0.1865	0.5868
CG15720	CG15720	1637855_at	-0.0581	0.7547	0.0307	0.8016	0.1067	0.6296	-0.0378	0.9603	-0.1264	0.5616	-0.0886	0.6744	-0.0171	0.9922	-0.0442	0.9451	-0.0271	0.9629
CG34388 /// DereCG14842	CG14842	1637856_at	0.0714	0.6770	-0.2053	0.5664	-0.1419	0.4181	0.0484	0.9526	0.1035	0.6882	0.0551	0.8298	-0.0010	0.9996	-0.1569	0.5208	-0.1559	0.5299
lectin-24A	Lectin24A	1637857_at	0.0522	0.8067	-0.1622	0.3389	-1.1862	0.3911	-1.0486	0.7680	0.4343	0.8169	1.4829	0.2774	-0.0203	0.9933	0.1534	0.8376	0.1737	0.7964
Cyp313b1	Cyp313b1	1637858_at	0.1240	0.8295	0.1568	0.7626	0.3931	0.3034	0.1929	0.8251	-0.5110	0.1293	-0.7040	0.0277	0.0076	0.9990	-0.4709	0.6294	-0.4784	0.6228
ATbp	(A+T)-stretch bind	1637859_at	-0.4079	0.2512	-0.6599	0.1149	0.0373	0.9263	0.3307	0.5900	0.0657	0.8732	-0.2650	0.3540	-0.4320	0.8202	-0.1623	0.8858	0.2697	0.7683
tam	DNA polymerase	1637860_at	0.1700	0.4387	0.3710	0.2808	0.5490	0.0353	-0.1556	0.6673	0.1330	0.4597	0.2886	0.0648	-0.2766	0.7644	0.1402	0.7693	0.4168	0.3011
CG7786	CG7786	1637861_at	0.0122	0.9533	0.0315	0.8979	0.0233	0.9332	0.0037	0.9956	0.0083	0.9800	0.0046	0.9863	0.0949	0.9168	-0.0375	0.9408	-0.1324	0.7077
CG6654	CG6654	1637862_at	-0.7072	0.0115	0.1832	0.2911	0.0912	0.6099	-0.2099	0.6041	-0.9734	0.0012	-0.7636	0.0021	-0.0985	0.8903	-0.0846	0.8059	0.0139	0.9733
---	---	1637863_x_at	-0.0548	0.7909	0.1537	0.4658	0.0635	0.7625	0.0076	0.9940	-0.0939	0.6939	-0.1015	0.6323	0.0542	0.9452	0.1304	0.6097	0.0762	0.7907
---	---	1637864_at	0.2102	0.2827	0.0302	0.7903	0.1466	0.4963	-0.1143	0.8304	0.0272	0.9241	0.1415	0.4652	-0.1113	0.9011	-0.1921	0.5800	-0.0809	0.8474
CG40381	CG40381	1637865_at	0.0500	0.8530	-0.0369	0.8777	0.1238	0.5510	-0.1046	0.8875	-0.0706	0.8210	0.0340	0.9101	0.0203	0.9898	-0.0063	0.9935	-0.0267	0.9563
CG9246	CG9246	1637866_at	0.3470	0.3396	0.7929	0.1434	1.1001	0.0077	0.3128	0.5568	0.3267	0.2340	0.0139	0.9681	-0.0796	0.9742	0.7165	0.2318	0.7961	0.2148
mid	neuromancer 2	1637867_at	-3.6489	0.0033	-4.8111	0.0014	-5.4268	0.0000	-0.1461	0.9269	0.8771	0.0688	1.0231	0.0246	-0.1871	0.9298	-0.4540	0.5246	-0.2669	0.7355
---	---	1637868_at	0.1793	0.2675	0.1147	0.3453	0.1325	0.3686	-0.0734	0.8974	-0.0522	0.8240	0.0212	0.9262	-0.0004	0.9998	0.0225	0.9326	0.0228	0.9199
PFE	Pray For Elves	1637869_at	-2.8152	0.0005	-0.7335	0.0990	-1.9499	0.0054	-1.3154	0.1140	-1.8233	0.0039	-0.5078	0.2585	-0.2815	0.8692	-0.1796	0.8445	0.1020	0.9124
dpr5	dpr5	1637870_a_at	0.0650	0.8132	-0.0006	1.0000	-0.0072	0.9762	0.2410	0.5983	0.2035	0.3898	-0.0375	0.8921	0.0349	0.9644	0.0246	0.9425	-0.0104	0.9756
CG11423	CG11423	1637871_at	0.2029	0.2346	0.0189	0.8503	0.0562	0.7978	-0.0556	0.9346	-0.0040	0.9896	0.0516	0.8212	0.0305	0.9775	0.1120	0.6955	0.0815	0.7884
Spat	serine pyruvate ar	1637872_at	2.5540	0.0041	1.8729	0.0833	3.4544	0.0000	0.8278	0.4612	-0.4448	0.4856	-1.2725	0.0273	-0.7485	0.7220	-1.0981	0.2026	-0.3495	0.7323
scpr-C	SCP-containing pr	1637873_at	0.1128	0.4882	0.1277	0.2752	0.2947	0.0509	0.0374	0.9518	0.1900	0.2701	0.1526	0.3302	0.0356	0.9779	0.2614	0.3856	0.2258	0.4759
Lasp	Lasp	1637874_at	-0.9573	0.1187	0.7102	0.2650	-0.0659	0.7475	-0.0771	0.9661	-0.3717	0.4546	-0.2946	0.5259	0.5460	0.7588	1.2278	0.1070	0.6818	0.3743

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1637875_a_at	-0.0108	0.9528	0.4340	0.0483	0.5559	0.0045	0.0712	0.8671	-0.4491	0.0094	-0.5202	0.0029	0.1245	0.8243	0.1413	0.5644	0.0168	0.9612
CG5976	CG5976	1637876_a_at	0.3132	0.3576	1.1667	0.0436	1.1925	0.0001	-0.1488	0.8158	-0.3763	0.1323	-0.2275	0.3198	-0.0118	0.9952	0.6656	0.1692	0.6774	0.1990
---	---	1637877_at	0.0945	0.6177	-0.1462	0.3380	0.0198	0.9159	0.0587	0.9048	0.1661	0.3019	0.1074	0.4797	-0.1136	0.8326	-0.0746	0.7898	0.0390	0.8994
---	---	1637878_at	-0.0232	0.9046	-0.3314	0.2164	0.4116	0.0649	0.3860	0.2729	0.7079	0.0053	0.3218	0.0839	-0.4244	0.5765	0.3523	0.2942	0.7767	0.0607
CG12720 /// DsimCG12720	CG12720	1637879_at	-0.0869	0.5631	-0.2096	0.0798	-0.3298	0.0828	0.0587	0.9445	0.1100	0.6918	0.0513	0.8553	-0.0497	0.9221	-0.0215	0.9368	0.0282	0.9033
CG13606	CG13606	1637880_at	0.7310	0.0033	0.4893	0.2669	0.3305	0.1463	-0.0497	0.9404	0.0745	0.7417	0.1242	0.4998	0.1180	0.9174	-0.0269	0.9665	-0.1449	0.7492
CG10373	CG10373	1637881_at	0.2302	0.2521	0.4469	0.0978	0.6035	0.0038	0.1093	0.7809	-0.1823	0.2663	-0.2916	0.0504	0.0078	0.9952	0.2591	0.4194	0.2513	0.4498
---	---	1637882_at	0.1169	0.5428	0.2210	0.2697	0.0878	0.5770	-0.0531	0.9300	-0.0785	0.7008	-0.0254	0.9053	0.0014	0.9994	0.1020	0.7371	0.1005	0.7371
---	---	1637883_at	0.0991	0.6220	0.2659	0.1726	0.1688	0.4600	-0.0068	0.9937	-0.1402	0.4441	-0.1334	0.4192	0.1450	0.8297	0.0898	0.8040	-0.0552	0.8871
CG40378	CG40378	1637884_at	0.1006	0.5089	0.0138	0.9299	0.0162	0.9243	-0.0214	0.9745	0.0866	0.6407	0.1080	0.4965	-0.0528	0.9226	-0.1229	0.5085	-0.0701	0.7346
---	---	1637885_at	-0.0543	0.7678	-0.0882	0.5061	0.1664	0.2684	0.0540	0.9362	-0.0388	0.8838	-0.0928	0.6487	-0.0112	0.9913	-0.0029	0.9950	0.0083	0.9821
CG10339	CG10339	1637886_at	0.0625	0.8471	0.0495	0.6430	0.0680	0.7327	-0.0627	0.8987	-0.1279	0.4459	-0.0651	0.7020	-0.0096	0.9950	-0.1999	0.6007	-0.1904	0.6216
CG1703	CG1703	1637887_at	0.8041	0.0078	0.1659	0.6832	0.2257	0.2169	0.1309	0.7979	1.2082	0.0006	1.0773	0.0005	0.2113	0.8521	0.6791	0.1413	0.4678	0.3300
CG7991	CG7991	1637888_at	-0.4230	0.1131	0.2224	0.2985	0.0583	0.7617	-0.0458	0.9639	-0.3593	0.1774	-0.3135	0.1896	-0.0170	0.9908	0.1201	0.7378	0.1371	0.6861
CG5883	CG5883	1637889_at	-0.0684	0.7609	-0.0571	0.8628	-0.0220	0.8989	-0.0248	0.9857	0.1606	0.6424	0.1854	0.5384	-0.2383	0.7464	-0.0363	0.9425	0.2020	0.5403
---	---	1637890_at	0.0570	0.6947	-0.1644	0.1996	-0.1160	0.5475	-0.0329	0.9538	0.0922	0.5869	0.1251	0.3878	-0.0341	0.9835	-0.0227	0.9719	0.0114	0.9849
---	---	1637891_at	0.0781	0.6643	-0.0101	0.9444	0.0799	0.7211	-0.0478	0.9496	-0.1361	0.5476	-0.0883	0.6893	-0.1043	0.8331	-0.1489	0.4812	-0.0446	0.8729
CG12594	CG12594	1637892_at	-0.0876	0.6708	0.0506	0.7880	-0.1312	0.3868	-0.0321	0.9637	0.0172	0.9509	0.0494	0.8141	0.1225	0.7726	0.0220	0.9404	-0.1004	0.6097
Vps28	Vps28	1637893_at	0.0806	0.7304	0.4539	0.0527	0.4700	0.0293	0.2730	0.5680	-0.2241	0.3695	-0.4972	0.0305	0.0872	0.9400	0.1863	0.6281	0.0992	0.8247
CG14509	CG14509	1637894_at	0.2903	0.2125	-0.2456	0.3095	-0.2901	0.2705	0.0328	0.9672	0.2945	0.1584	0.2617	0.1619	-0.1327	0.9199	-0.3320	0.4619	-0.1992	0.6875
Dhc16F	dynein-related hea	1637895_at	-0.1293	0.3655	0.0454	0.7930	0.0783	0.7027	-0.0354	0.9619	-0.0881	0.6962	-0.0527	0.8118	0.0019	0.9984	-0.0569	0.8190	-0.0588	0.8002
Dhc62B	dynein-related hea	1637896_at	0.3851	0.1005	0.1450	0.5373	0.2609	0.1818	-0.0217	0.9777	-0.0288	0.9183	-0.0072	0.9776	-0.2712	0.7464	-0.3433	0.3158	-0.0721	0.8801
CycT	cyclin T2	1637897_at	1.0834	0.0179	1.8121	0.0299	0.3753	0.2938	-1.1791	0.0444	-0.3598	0.2817	0.8193	0.0136	0.1375	0.9514	0.3564	0.6157	0.2189	0.7803
---	---	1637898_at	0.1858	0.4013	0.0847	0.7003	-0.0792	0.7496	-0.0539	0.9666	-0.0473	0.9204	0.0067	0.9874	0.0542	0.9646	-0.0225	0.9666	-0.0767	0.8578
CG13176	CG13176	1637899_at	-0.5688	0.1237	0.0418	0.9564	0.0199	0.9161	-0.0478	0.9368	-0.4639	0.0163	-0.4161	0.0161	-0.0525	0.9875	0.1638	0.8807	0.2163	0.8194
CG11852 /// DyakCG11852	CG11852	1637900_at	0.8445	0.1579	0.7987	0.0471	0.5623	0.2214	-0.0653	0.9761	-0.9011	0.0941	-0.8358	0.0832	0.0341	0.9921	-1.0244	0.1745	-1.0585	0.1990
CG14151	CG14151	1637901_at	0.1201	0.4528	0.2066	0.1402	0.2938	0.1376	-0.0778	0.9098	-0.0201	0.9513	0.0578	0.8144	-0.0809	0.8558	0.0210	0.9429	0.1019	0.6036
CG10918	CG10918	1637902_at	-0.0622	0.7903	0.4113	0.1237	0.3499	0.1442	-0.2273	0.6893	-0.3953	0.1328	-0.1680	0.5098	-0.0448	0.9742	0.2152	0.5523	0.2600	0.4677
---	---	1637903_at	0.2623	0.1955	0.5625	0.2500	0.3740	0.1287	-0.0741	0.9518	-0.2438	0.4917	-0.1697	0.6220	-0.0226	0.9898	-0.1041	0.8386	-0.0815	0.8739
CG31787	CG31787	1637904_at	0.0407	0.8547	0.0336	0.7239	0.0672	0.6938	0.1179	0.7931	0.0298	0.9070	-0.0881	0.6402	0.1546	0.7726	0.1346	0.5832	-0.0199	0.9512
CG2924	CG2924	1637905_s_at	0.4247	0.1062	1.0327	0.0266	0.4633	0.0324	-0.4753	0.2309	-0.1769	0.4732	0.2984	0.1559	0.0788	0.9340	0.4478	0.1388	0.3691	0.2409
---	---	1637906_at	0.0148	0.9380	-0.0248	0.8730	0.0945	0.5594	0.2040	0.5515	0.1336	0.4709	-0.0704	0.7074	0.0541	0.9405	0.1029	0.6786	0.0488	0.8699
CG10476	CG10476	1637907_at	0.0363	0.8760	0.1064	0.4655	0.1907	0.2915	-0.0339	0.9583	-0.1047	0.5696	-0.0708	0.6924	-0.0874	0.9298	0.0509	0.9188	0.1383	0.7001
---	---	1637908_at	0.0978	0.6805	0.0107	0.9208	0.1410	0.4504	0.0333	0.9602	-0.0170	0.9485	-0.0503	0.7981	-0.1244	0.8689	-0.0209	0.9658	0.1035	0.7655
Pp1-Y2	Pp1-Y2	1637909_at	0.1817	0.3254	0.0017	0.9899	-0.2006	0.3300	0.0572	0.9479	0.1580	0.5503	0.1008	0.6964	0.1473	0.7979	-0.0419	0.9088	-0.1893	0.4449
CG30125	CG30125	1637910_at	0.2099	0.3492	-0.1049	0.5054	-0.0869	0.6216	0.0214	0.9803	0.2669	0.2010	0.2455	0.1893	0.0713	0.9499	-0.0013	0.9994	-0.0726	0.8736
CG16892	CG16892	1637911_at	-0.4928	0.0922	-0.2553	0.3473	-0.0120	0.9736	0.2074	0.5330	0.0124	0.9623	-0.1950	0.2091	-0.2082	0.8940	0.1823	0.8083	0.3904	0.5232
---	---	1637912_at	-0.7993	0.0120	-0.5017	0.0270	-0.9283	0.0090	-0.5429	0.1787	-0.3843	0.1151	0.1586	0.5002	-0.1820	0.8270	-0.0459	0.9341	0.1361	0.7333
CG11638	CG11638	1637913_at	0.4945	0.0868	0.2232	0.4675	0.3450	0.2049	-0.1212	0.8678	0.3780	0.1408	0.4992	0.0370	-0.1594	0.9267	0.3241	0.5935	0.4834	0.4058
Obp83g	Odorant-binding p	1637914_at	-0.6095	0.0080	-0.1786	0.2184	-0.0580	0.8516	0.1867	0.6763	-0.4104	0.0554	-0.5971	0.0072	-0.1211	0.8940	-0.1307	0.7424	-0.0095	0.9856
---	---	1637915_at	0.1397	0.5038	-0.0262	0.9363	-0.1446	0.3923	-0.0459	0.9380	0.1249	0.4910	0.1709	0.2741	0.0254	0.9877	-0.1693	0.6830	-0.1947	0.6271
CG33013	CG33013	1637916_at	0.3003	0.1912	0.1650	0.2672	0.1692	0.4363	-0.0934	0.8967	0.0314	0.9243	0.1248	0.6000	-0.1440	0.8609	-0.0946	0.8302	0.0494	0.9147
ttk	tramtrack-69	1637917_s_at	-0.3912	0.4201	-0.6068	0.0218	-1.1592	0.0002	-0.0276	0.9866	0.4202	0.3019	0.4478	0.2144	0.5794	0.1628	0.1737	0.3820	-0.4056	0.0828
CG16786	CG16786	1637918_s_at	0.1389	0.6168	0.0070	0.9511	0.0930	0.7194	0.0076	0.9952	-0.0819	0.7955	-0.0894	0.7492	-0.0590	0.9277	-0.0483	0.8737	0.0106	0.9751
CG15233	CG15233	1637919_at	0.1384	0.5661	-0.0058	0.9596	-0.0335	0.8662	0.1081	0.8427	0.1141	0.6075	0.0060	0.9809	0.0044	0.9964	0.0355	0.9129	0.0311	0.9157
---	---	1637920_at	0.0344	0.8243	0.0784	0.5658	0.0806	0.6903	-0.2234	0.6086	-0.1596	0.4873	0.0638	0.7958	0.0340	0.9657	0.0262	0.9402	-0.0078	0.9835
Gr23a	Gustatory recepto	1637921_a_at	-0.0477	0.7453	-0.0315	0.8135	0.0019	0.9943	-0.0637	0.9110	-0.0251	0.9214	0.0387	0.8540	-0.1267	0.8222	-0.0169	0.9650	0.1099	0.6677
CG5592	CG5592	1637922_at	0.0291	0.8868	0.0903	0.5524	0.2791	0.1516	0.1471	0.7949	0.2187	0.3517	0.0715	0.7805	-0.1189	0.8270	-0.0044	0.9935	0.1145	0.6403
CG31938	CG31938	1637923_at	0.5531	0.1252	0.6466	0.2381	1.1263	0.0003	0.2598	0.5311	0.2655	0.2185	0.0057	0.9835	-0.2756	0.8472	0.3926	0.5246	0.6682	0.2778
---	---	1637924_at	0.2331	0.1234	0.0167	0.8848	0.0467	0.8149	-0.0919	0.8068	0.0015	0.9945	0.0934	0.5195	-0.0390	0.9521	-0.0488	0.8551	-0.0098	0.9748
CG32790	CG32790	1637925_at	0.1496	0.3792	0.0674	0.5978	0.1227	0.5666	0.1400	0.7857	0.0902	0.7146	-0.0497	0.8397	0.1655	0.8145	0.0330	0.9447	-0.1325	0.6800
CG14084	CG14084	1637926_s_at	0.5132	0.0129	0.7937	0.0323	1.0175	0.0003	0.2206	0.6214	0.3118	0.1587	0.0911	0.6964	0.2077	0.7644	0.7422	0.0365	0.5344	0.1049
---	---	1637927_at	0.0028	0.9897	0.1485	0.1781	-0.0455	0.7797	-0.1570	0.7432	-0.1816	0.4019	-0.0246	0.9240	0.0428	0.9506	0.0044	0.9925	-0.0384	0.8918
Oatp58Db	Organic anion trar	1637928_at	-0.5222	0.6234	0.1687	0.3576	0.3624	0.0365	0.0334	0.9937	-0.6390	0.4553	-0.6724	0.3769	0.0013	0.9998	-0.0950	0.8632	-0.0963	0.8503
CG32602	CG32602	1637929_a_at	0.0686	0.7560	-0.1101	0.4261	-0.0429	0.8067	0.1129	0.8796	0.2239	0.3925	0.1110	0.6776	-0.0036	0.9976	-0.0072	0.9906	-0.0036	0.9941
CG14089	CG14089	1637930_at	0.2649	0.0702	0.1814	0.1104	0.1973	0.4460	0.1003	0.8838	0.0588	0.8463	-0.0415	0.8807	0.0746	0.9092	0.0394	0.9121	-0.0352	0.9121
CG15036	CG15036	1637931_at	0.0093	0.9687	0.0591	0.5843	0.1425	0.3968	0.1299	0.7556	0.1613	0.3814	0.0314	0.8831	-0.0603	0.9421	0.1807	0.4951	0.2411	0.3629
---	---	1637932_at	0.1335	0.5697	-0.2191	0.0818	0.1454	0.5003	0.1068	0.8612	0.1287	0.5890	0.0220	0.9357	-0.1896	0.7423	-0.0926	0.7466	0.0970	0.7271
Pdi	protein disulfide is	1637933_a_at	-0.9156	0.0821	-1.3186	0.0160	-0.8396	0.0428	0.0937	0.9489	0.7745	0.0614	0.6808	0.0651	-0.3935	0.7644	0.1333	0.8642	0.5268	0.3583
---	---	1637934_at	-0.2327	0.1497	-0.1121	0.6027	-0.5329	0.0300	-0.3771	0.1601	-0.1056	0.5460	0.2715	0.0651	0.0604	0.9752	-0.0916	0.8995	-0.1520	0.7919
ARY	CG40064	1637935_at	0.1936	0.2259	0.0003	1.0000	0.0398	0.8055	-0.1243	0.7131	0.0447	0.8144	0.1689	0.2197	-0.0603	0.9305	-0.1058	0.6673	-0.0455	0.8812
CG32512	CG32512	1637936_at	-1.3027	0.0233	-0.3615	0.0996	-0.4073	0.2505	-0.1778	0.8479	-1.9113	0.0007	-1.7334	0.0006	-0.0660	0.9816	-0.9126	0.1420	-0.8465	0.2051
---	---	1637937_at	0.3980	0.0390	0.0050	0.9692	0.2538	0.0899	-0.0248	0.9696	0.1000	0.5818	0.1247	0.4260	-0.0943	0.8737	0.0280	0.9404	0.1223	0.6263
CG18636	CG18636	1637938_at	0.0501	0.8639	0.0567	0.6084	0.1080	0.6106	0.0062	0.9937	0.0127	0.9569	0.0066	0.9736	0.0199	0.9885	0.1867	0.5523	0.1668	0.6051
Spn43Aa	Serine protease ir	1637939_at	-0.3569	0.7234	0.2620	0.2630	0.5042	0.0469	0.0362	0.9912	-0.6457	0.3715	-0.6818	0.2870	-0.2158	0.8202	-0.1063	0.8397	0.1095	0.8237
CG4452	CG4452	1637940_s_at	0.4621	0.2415	0.3287	0.1402	0.6593	0.0350	-0.0168	0.9872	-0.4416	0.0793	-0.4249	0.0614	-0.3744	0.8024	-0.6721	0.2645	-0.2978	0.6604
RpS10a	Ribosomal protein	1637941_at	0.3497	0.0840	0.4668	0.0173	0.1500	0.3762	-0.0339	0.9592	-0.0189	0.9402	0.0150	0.9461	0.2752	0.6660	0.0077	0.9887	-0.2675	0.3094
CG8538	CG8538	1637942_at	0.1273	0.7177	0.2972	0.3381	0.0925	0.6343	-0.1073	0.8544	0.0724	0.7763	0.1797	0.3539	0.1952	0.8823	0.6908	0.1577	0.4956	0.3362
how	struthio	1637943_at	-0.7260	0.0233	-1.0162	0.0235	-1.6418	0.0007	-0.1149	0.7591	0.2752	0.0880	0.3901	0.0147	0.3275	0.7779	0.0406	0.9617	-0.2869	0.5969

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG10841	CG10841	1637944_at	0.0242	0.9076	0.0639	0.5937	0.1269	0.4308	0.0753	0.8671	0.0503	0.7963	-0.0251	0.8955	0.0197	0.9848	0.0082	0.9858	-0.0115	0.9751
---	---	1637945_at	0.0325	0.9245	-0.6813	0.1041	-1.2689	0.0023	-0.3408	0.5608	0.9514	0.0074	1.2922	0.0012	-0.1801	0.8810	0.0560	0.9402	0.2361	0.6328
CG5068	CG5068	1637946_at	-0.9170	0.0030	-0.6283	0.0405	-0.4717	0.0224	0.0260	0.9666	-0.0694	0.7121	-0.0954	0.5504	-0.2135	0.7743	0.1379	0.7054	0.3514	0.2917
CG30389	CG30389	1637947_s_at	0.1725	0.6779	0.4014	0.2906	0.0777	0.7014	-0.1208	0.8844	0.0434	0.9127	0.1642	0.5568	0.2059	0.8869	0.2435	0.6876	0.0376	0.9622
---	---	1637948_at	0.1910	0.3735	0.0514	0.6248	0.0245	0.9535	0.0824	0.9311	0.2280	0.4326	0.1456	0.6073	-0.0779	0.9495	0.0101	0.9903	0.0880	0.8526
---	---	1637949_at	-0.0855	0.7386	0.0911	0.5244	-0.3981	0.0874	-0.0587	0.9441	-0.0098	0.9790	0.0489	0.8605	0.1259	0.8192	-0.0243	0.9473	-0.1502	0.5267
---	---	1637950_at	0.0905	0.5650	-0.0377	0.8901	0.1052	0.5359	0.1239	0.8409	0.0460	0.8810	-0.0779	0.7570	-0.0769	0.9298	-0.1599	0.5959	-0.0830	0.8133
CG8461	CG8461	1637951_at	0.1159	0.6735	-0.5228	0.1784	-0.8017	0.0025	0.0354	0.9688	0.7021	0.0094	0.6668	0.0072	0.2193	0.8222	0.0771	0.9002	-0.1422	0.7626
---	---	1637952_s_at	0.2229	0.2390	0.0003	1.0000	0.0908	0.6962	-0.0360	0.9688	0.0312	0.9257	0.0673	0.8039	-0.0947	0.8480	-0.0795	0.7435	0.0153	0.9596
CG6324	CG6324	1637953_at	-0.0386	0.8769	0.0058	0.9712	-0.0976	0.7113	0.0057	0.9955	0.0735	0.7562	0.0678	0.7540	0.1402	0.8192	0.2456	0.3158	0.1053	0.7099
CG5844 /// DyakCG5844	Enoyl-CoA hydrat	1637954_at	-0.5796	0.0334	0.5854	0.0667	0.9099	0.0008	-0.0804	0.8822	-1.1765	0.0004	-1.0962	0.0003	-0.2498	0.7485	0.1857	0.5995	0.4355	0.2094
CG1827	CG1827	1637955_a_at	-0.1627	0.4021	-0.0761	0.8438	0.5309	0.0067	0.4193	0.1106	-0.2190	0.1573	-0.6383	0.0014	-0.2033	0.8424	-0.1227	0.8231	0.0806	0.8889
btz	barentsz	1637956_s_at	-0.3345	0.5633	-0.4807	0.5994	-0.3535	0.2431	0.0048	0.9956	0.0092	0.9746	0.0044	0.9852	0.0211	0.9964	-0.0603	0.9751	-0.0814	0.9598
Rbp9	female sterile(2)Bi	1637957_s_at	-2.1095	0.0011	0.5091	0.3955	-1.7477	0.0013	-1.1728	0.1140	-2.1671	0.0011	-0.9943	0.0205	1.3138	0.3493	0.7144	0.3247	-0.5994	0.4318
CG33510	CG33510	1637958_at	-0.0369	0.9112	0.3910	0.1272	0.5847	0.0210	0.1630	0.7805	-0.0139	0.9705	-0.1769	0.4388	0.1669	0.8828	0.3971	0.3458	0.2302	0.6216
CG4334 /// DyakCG4334	CG4334	1637959_at	0.6919	0.0113	0.2501	0.3450	0.5516	0.0057	-0.1631	0.8285	0.0538	0.8907	0.2169	0.4239	-0.1917	0.7726	-0.1465	0.6325	0.0452	0.9084
CG14839	CG14839	1637960_at	0.0943	0.6889	0.0998	0.4432	0.2100	0.2495	0.0540	0.9349	-0.1715	0.3918	-0.2255	0.1957	0.0749	0.9467	-0.0607	0.9073	-0.1356	0.7253
zen	zerknult	1637961_at	0.1324	0.4247	-0.2046	0.3916	-0.0057	0.9868	0.0130	0.9922	0.0954	0.7795	0.0824	0.7943	-0.0652	0.9088	-0.1290	0.5398	-0.0638	0.7919
spir	spire	1637962_at	-0.1679	0.7191	-1.4417	0.0895	-0.9517	0.0641	0.3768	0.7225	1.1310	0.0259	0.7543	0.0791	-0.3071	0.8999	-0.3099	0.7787	-0.0028	0.9989
CG14191	CG14191	1637963_at	-5.2736	0.0065	-3.9643	0.0465	-5.3006	0.0002	-0.2728	0.5008	-0.9227	0.0019	-0.6500	0.0053	1.1841	0.8380	0.3464	0.9273	-0.8377	0.7625
Eig71Eg	Gene VI	1637964_at	0.0885	0.5945	-0.0492	0.8423	-0.0025	0.9916	-0.0717	0.8601	0.1241	0.4090	0.1958	0.1317	-0.0837	0.9063	-0.0296	0.9425	0.0542	0.8746
CG31397	CG31397	1637965_at	-0.0995	0.6541	-0.0396	0.8056	-0.1207	0.6496	0.2024	0.5735	0.2108	0.2534	0.0084	0.9710	0.0836	0.9168	-0.0121	0.9814	-0.0956	0.7663
slpr	slipper	1637966_at	-0.0815	0.8138	0.2745	0.3495	0.1895	0.4032	-0.1828	0.7982	-0.0045	0.9913	0.1783	0.5276	-0.1067	0.9404	0.2011	0.6812	0.3078	0.4978
Tim8	Tim8	1637967_at	0.4394	0.0382	0.8016	0.0537	0.7281	0.0172	0.1339	0.7577	-0.0429	0.8611	-0.1768	0.2957	0.1783	0.8609	0.3647	0.3685	0.1864	0.6827
CG40315	CG40315	1637968_at	0.1335	0.4027	0.0021	0.9875	0.0974	0.6049	-0.0204	0.9777	0.0961	0.6416	0.1165	0.5126	-0.0335	0.9589	0.0563	0.8133	0.0898	0.6509
CG16984	CG16984	1637969_at	0.0292	0.8881	-0.0707	0.5422	0.3009	0.2814	0.4108	0.1119	0.1764	0.2522	-0.2344	0.0881	-0.1408	0.9092	0.0735	0.9129	0.2143	0.6484
---	---	1637970_at	0.1229	0.5491	0.2603	0.1238	0.2258	0.1522	0.0047	0.9956	0.0256	0.9278	0.0209	0.9328	0.2979	0.7220	0.3081	0.3769	0.0101	0.9856
CG8129	CG8129	1637971_a_at	3.0457	0.0279	0.9866	0.5285	2.1736	0.0006	-0.1745	0.9624	-0.0258	0.9865	0.1487	0.9000	-1.4020	0.7230	-2.1627	0.1920	-0.7606	0.6875
Umbrea	Umbrea	1637972_at	-0.0042	0.9891	-0.0807	0.6767	0.2805	0.1422	0.2994	0.4586	0.0628	0.8171	-0.2365	0.2271	-0.1638	0.8541	-0.0745	0.8893	0.0893	0.8507
CG4407	CG4407	1637973_a_at	-1.3788	0.0015	-1.1177	0.0393	-1.4408	0.0000	-0.0045	0.9956	-0.2328	0.2239	-0.2282	0.1818	0.4259	0.6538	0.0647	0.9121	-0.3612	0.3569
CG31246	CG31246	1637974_at	0.2268	0.2309	-0.2311	0.1081	-0.0112	0.9560	0.2201	0.5144	0.3111	0.0825	0.0910	0.6146	-0.1032	0.8760	-0.2058	0.4246	-0.1026	0.7284
Dhc36C	dynein-related hea	1637975_at	0.0729	0.6553	0.0247	0.9188	0.3995	0.0938	0.0013	0.9986	-0.0137	0.9513	-0.0150	0.9354	-0.2793	0.7230	-0.0828	0.8619	0.1965	0.5900
---	---	1637976_at	-0.0664	0.6469	0.0218	0.9376	-0.0205	0.9091	0.0208	0.9803	-0.0071	0.9816	-0.0278	0.9111	-0.0375	0.9811	0.0366	0.9470	0.0741	0.8699
Notum	wingful	1637977_a_at	0.0983	0.5789	0.2050	0.2848	0.1523	0.3380	-0.0048	0.9956	-0.1078	0.6159	-0.1030	0.5975	0.1102	0.8465	0.1233	0.6295	0.0131	0.9714
CG10194	CG10194	1637978_at	0.4691	0.0301	0.9468	0.0214	1.1814	0.0001	0.0333	0.9540	-0.2928	0.0682	-0.3261	0.0293	-0.2688	0.5905	0.2139	0.3217	0.4827	0.0679
CG13567	CG13567	1637979_at	-0.1509	0.6838	-0.1069	0.6327	0.2689	0.2287	-0.1354	0.9036	-0.3267	0.3781	-0.1913	0.5996	-0.2425	0.8215	-0.2208	0.6497	0.0217	0.9759
awd	killer of prune	1637980_at	0.4073	0.0492	1.3186	0.0057	1.6921	0.0004	0.0888	0.8942	-1.0801	0.0011	-1.1689	0.0005	-0.2019	0.7500	0.1285	0.6669	0.3303	0.2370
---	---	1637981_at	0.2566	0.2247	0.2169	0.3525	0.3304	0.1036	0.0196	0.9777	-0.0776	0.7101	-0.0973	0.5906	0.1642	0.8270	0.1101	0.7731	-0.0542	0.9004
Nurf-38	pyrophosphatase	1637982_at	0.3479	0.0473	0.6490	0.0121	0.8053	0.0077	0.1359	0.8399	-0.6160	0.0225	-0.7518	0.0057	0.1443	0.8128	0.0231	0.9552	-0.1212	0.6577
CG10365	CG10365	1637983_s_at	-0.1671	0.3594	1.2822	0.0085	1.1463	0.0013	-0.3099	0.3793	-1.8007	0.0001	-1.4908	0.0001	-0.1169	0.8882	-0.2789	0.3647	-0.1620	0.6311
---	---	1637984_at	-0.1196	0.5414	-0.1486	0.4010	0.1970	0.1829	0.3520	0.4374	0.1691	0.5216	-0.1829	0.4329	-0.0258	0.9816	0.0951	0.7484	0.1209	0.6574
CG32087	CG32087	1637985_at	0.1670	0.4424	0.1613	0.3133	0.1684	0.2562	-0.1516	0.7742	-0.1212	0.6217	0.0305	0.9095	0.1041	0.8744	0.0563	0.8792	-0.0477	0.8928
CG15253	CG15253	1637986_at	-0.1917	0.5920	0.0181	0.8678	-0.0524	0.7606	-0.1159	0.8244	-0.3173	0.1157	-0.2014	0.2701	0.0339	0.9816	-0.2260	0.4899	-0.2599	0.4280
ofs	eIF-4G-like proteir	1637987_at	-0.3484	0.3670	-0.4791	0.1382	-0.7429	0.0128	-0.0135	0.9937	0.8620	0.0177	0.8755	0.0102	0.2997	0.7768	0.6240	0.1599	0.3243	0.4930
CG14483	CG14483	1637988_at	0.0293	0.9117	-0.1881	0.6311	-0.1464	0.3943	0.0631	0.9254	-0.0179	0.9529	-0.0810	0.7049	0.0344	0.9860	-0.2285	0.6280	-0.2629	0.5709
CG4810	CG4810	1637989_at	0.0563	0.8062	0.1000	0.5014	0.0927	0.6435	0.0635	0.8811	0.0870	0.5801	0.0235	0.8924	0.0394	0.9659	0.0887	0.7692	0.0493	0.8856
fbl	fumble	1637990_s_at	0.6717	0.0265	0.8750	0.0145	1.3432	0.0003	0.2347	0.5735	-0.0941	0.7030	-0.3288	0.0862	-0.1759	0.8461	0.1626	0.7017	0.3385	0.3803
---	---	1637991_at	0.0751	0.6703	-0.0006	0.9979	0.1603	0.3256	-0.0260	0.9639	-0.0594	0.7402	-0.0334	0.8495	-0.1398	0.8692	-0.0752	0.8773	0.0646	0.8904
CG6607	CG6607	1637992_at	-0.4722	0.1279	0.3202	0.1096	0.3820	0.0476	0.0703	0.9241	-0.3974	0.0748	-0.4677	0.0261	0.1163	0.9142	0.4645	0.2032	0.3482	0.3688
MED11	Mediator complex	1637993_at	0.0309	0.9173	-0.2969	0.1656	-0.2141	0.2111	0.2183	0.6138	0.3093	0.1522	0.0910	0.6896	-0.0411	0.9653	-0.0693	0.8389	-0.0282	0.9369
mars	mars	1637994_at	-0.1176	0.5312	-0.1598	0.7583	-0.3790	0.1439	-0.1650	0.7714	0.0855	0.7663	0.2505	0.2443	-0.2166	0.8609	-0.0943	0.9009	0.1224	0.8493
CG15404	CG15404	1637995_at	0.1004	0.5137	0.1582	0.5303	0.2398	0.2140	0.1055	0.8446	0.0690	0.7724	-0.0365	0.8771	-0.0768	0.9401	0.0175	0.9739	0.0942	0.8073
---	---	1637996_at	0.2235	0.3335	0.4095	0.0387	0.3669	0.0243	0.1340	0.7229	0.0360	0.8732	-0.0980	0.5608	0.0856	0.9340	0.1365	0.7191	0.0509	0.9109
CG32155	CG32155	1637997_at	0.9836	0.0046	1.1032	0.0290	1.7699	0.0003	-0.1194	0.8578	-0.5081	0.0406	-0.3886	0.0722	-0.8080	0.3926	-0.4199	0.4134	0.3881	0.4683
Cpr65Az	CG12330	1637998_at	0.0906	0.5225	0.0732	0.5641	0.0553	0.7369	0.0874	0.8671	-0.0559	0.8053	-0.1433	0.4045	0.1520	0.7464	0.1045	0.6279	-0.0475	0.8547
CG31079	CG31079	1637999_at	-0.0023	0.9895	-0.0396	0.7793	-0.0889	0.7496	0.0540	0.9508	0.1887	0.4559	0.1347	0.5783	0.0661	0.9142	0.0652	0.8153	-0.0009	0.9982
sala	spalt	1638000_at	-0.9319	0.7470	-0.7043	0.0674	-0.4733	0.0888	0.1153	0.9900	-1.2657	0.5713	-1.3810	0.4848	-0.0305	0.9964	-1.0970	0.5102	-1.0665	0.5299
CG8889	CG8889	1638001_a_at	0.1835	0.3711	0.0851	0.8323	0.0139	0.9467	-0.2545	0.5459	0.1380	0.5578	0.3925	0.0495	-0.1771	0.8558	0.0008	0.9998	0.1780	0.6877
CG32808	CG32808	1638002_at	0.0062	0.9854	0.0104	0.9702	0.3370	0.2202	0.2226	0.7511	0.0487	0.9084	-0.1739	0.5645	-0.1939	0.8243	-0.1279	0.7734	0.0660	0.8949
CG14204	CG14204	1638003_at	0.1482	0.4147	-0.0466	0.8606	0.0029	0.9907	-0.0847	0.8738	0.1040	0.6115	0.1887	0.2591	-0.0302	0.9816	-0.2120	0.4677	-0.1819	0.5499
Cbp53E	calbindin	1638004_at	-1.2037	0.0548	0.0253	0.9713	-1.3883	0.0216	-0.7387	0.4797	-1.9348	0.0050	-1.1962	0.0251	0.7204	0.7576	-0.6552	0.5199	-1.3755	0.1875
smg	Smaug	1638005_s_at	0.5250	0.2160	-0.5575	0.4334	-1.2203	0.0006	-0.6730	0.1280	1.0636	0.0026	1.7366	0.0002	-0.0223	0.9952	0.0681	0.9629	0.0904	0.9428
CG10211	CG10211	1638006_at	0.0937	0.8031	-0.0263	0.9702	-0.1058	0.8053	-0.3695	0.6082	0.1120	0.8006	0.4815	0.1345	-0.1740	0.9449	0.0409	0.9757	0.2149	0.8223
CG1542	CG1542	1638007_at	0.1775	0.6478	-0.6748	0.2062	-0.0923	0.7158	0.5747	0.3301	1.1541	0.0055	0.5794	0.0616	-0.0869	0.9717	0.2630	0.7011	0.3500	0.5886
CG2003	CG2003	1638008_s_at	0.7179	0.0041	1.6958	0.0083	1.5661	0.0000	0.4743	0.0976	-0.0388	0.8605	-0.5131	0.0052	0.5104	0.5236	0.8871	0.0419	0.3767	0.3404
CG9399	CG9399	1638009_s_at	-0.1022	0.6577	0.2392	0.2532	-0.1937	0.4889	-0.4818	0.2596	-0.7375	0.0106	-0.2556	0.2623	-0.1159	0.9095	-0.3542	0.3158	-0.2383	0.5332
cngl	CNG channel-like	1638010_at	0.0810	0.7502	-0.0063	0.9832	0.2196	0.1577	0.2013	0.6197	-0.0356	0.8974	-0.2369	0.1889	0.0895	0.9101	-0.0810	0.8287	-0.1705	0.5702
Past1	EH domain contai	1638011_a_at	-0.0423	0.8601	0.2156	0.3755	0.0350	0.9091	0.1127	0.8550	0.3892	0.0821	0.2765	0.1643	0.2156	0.8461	0.5641	0.2042	0.3485	0.4631
rt	Drosophila Proteir	1638012_at	0.2033	0.4290	0.0357	0.8486	-0.5817	0.0448	-0.5804	0.0681	0.1883	0.3119	0.7687	0.0013	-0.0088	0.9964	-0.0679	0.9180	-0.0591	0.9197

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG30289	CG30289	1638013_at	0.3992	0.2293	0.3516	0.2766	0.6999	0.0046	0.0547	0.9228	0.1840	0.2853	0.1293	0.4192	-0.2277	0.7770	-0.0610	0.9085	0.1667	0.6593
CG5808	CG5808	1638014_at	-0.2825	0.5619	0.0151	0.9564	-0.0110	0.9625	-0.0136	0.9922	-0.0043	0.9914	0.0092	0.9794	-0.0885	0.9748	0.4118	0.5762	0.5003	0.4869
---	---	1638015_at	-0.0645	0.6575	-0.0585	0.6096	-0.1128	0.6067	-0.0526	0.9074	0.0572	0.7394	0.1098	0.4204	-0.0339	0.9816	0.0587	0.9064	0.0926	0.8200
CG2993	CG2993	1638016_at	-0.1696	0.2859	-0.3667	0.0752	-0.2538	0.1789	0.1444	0.8099	0.3291	0.1656	0.1847	0.4066	-0.1083	0.8513	-0.0942	0.7354	0.0141	0.9695
Cpr78E	CG7160	1638017_at	-0.4743	0.0661	-0.7616	0.0462	-0.9917	0.0029	-0.2606	0.5418	0.0067	0.9837	0.2673	0.1774	-0.1204	0.8874	-0.1299	0.7215	-0.0096	0.9849
---	---	1638018_at	0.0007	0.9971	0.1719	0.2273	-0.0138	0.9579	-0.0110	0.9863	-0.0832	0.6295	-0.0722	0.6519	0.1212	0.8655	0.0969	0.7854	-0.0243	0.9517
---	---	1638019_at	-0.0131	0.9429	-0.0065	0.9609	-0.0631	0.6827	-0.1134	0.7425	-0.0078	0.9735	0.1056	0.4630	-0.0402	0.9653	0.0324	0.9350	0.0726	0.8129
CG10467	CG10467	1638020_at	0.9067	0.0104	0.0288	0.8551	1.0676	0.0151	0.5080	0.2747	0.5730	0.0426	0.0651	0.8358	-0.5621	0.6955	-0.3189	0.6259	0.2432	0.7243
CG4757	CG4757	1638021_at	-2.6856	0.0190	-0.1998	0.6775	-2.6913	0.0386	-2.0133	0.0925	-0.5093	0.4881	1.5039	0.0246	0.2514	0.9775	1.7444	0.4083	1.4930	0.4978
CG7971	anon-fast-evolving	1638022_at	-0.0648	0.8844	-0.7615	0.2632	-0.8043	0.0125	-0.2506	0.7349	0.4723	0.1490	0.7229	0.0222	-0.1117	0.9643	-0.1565	0.8690	-0.0448	0.9640
CG14880	CG14880	1638023_s_at	-0.0490	0.7388	-0.0552	0.5584	-0.0980	0.6339	-0.0605	0.9342	-0.2555	0.2346	-0.1950	0.3198	-0.0092	0.9943	-0.1016	0.7784	-0.0924	0.7938
---	---	1638024_at	0.0792	0.7365	-0.0157	0.9173	-0.1275	0.4399	0.1998	0.6338	0.3348	0.1034	0.1350	0.4947	0.3414	0.7215	0.3542	0.3669	0.0127	0.9847
---	---	1638025_at	0.0017	0.9936	0.2304	0.2728	0.3124	0.1480	0.0137	0.9857	-0.1019	0.5901	-0.1156	0.4863	-0.0437	0.9767	0.0718	0.8903	0.1155	0.7849
CG15121	CG15121	1638026_at	-0.0813	0.7367	-0.1810	0.3546	0.1436	0.5035	0.1454	0.7409	-0.0294	0.9136	-0.1748	0.3226	-0.0750	0.9056	-0.1279	0.5963	-0.0528	0.8578
CG17429	CG17429	1638027_at	0.0584	0.7828	-0.1543	0.3910	0.1816	0.3306	0.1442	0.7326	0.2717	0.1465	0.1275	0.4763	-0.0826	0.9400	0.0519	0.9211	0.1345	0.7247
CG30485	CG30485	1638028_at	0.0915	0.5064	-0.1500	0.4103	-0.0024	0.9910	0.1035	0.8248	0.1160	0.5519	0.0125	0.9570	0.0211	0.9787	-0.1257	0.5119	-0.1469	0.4407
---	---	1638029_at	0.2218	0.2885	0.1148	0.4399	0.0875	0.7155	-0.0194	0.9838	0.1019	0.6844	0.1213	0.5794	-0.0663	0.9340	-0.0338	0.9341	0.0324	0.9266
CG15237	CG15237	1638030_at	-0.0571	0.7443	-0.1588	0.3764	-0.2619	0.1914	0.2101	0.5249	0.1725	0.3257	-0.0376	0.8516	0.2124	0.7500	0.0541	0.9003	-0.1584	0.6052
CG14512	CG14512	1638031_at	-0.1465	0.4257	-0.1994	0.5485	-0.3164	0.0651	-0.1421	0.7699	0.1801	0.3935	0.3222	0.0811	0.1652	0.7932	0.1107	0.7151	-0.0545	0.8800
CG14921	CG14921	1638032_at	0.0513	0.8519	0.2107	0.2763	0.2284	0.2913	0.1327	0.7828	0.0608	0.8035	-0.0719	0.7369	0.2481	0.7324	0.2519	0.4151	0.0038	0.9945
CG6055	CG6055	1638033_at	-3.2804	0.0031	-4.6177	0.0028	-3.3616	0.0001	0.7004	0.4233	0.5069	0.2988	-0.1935	0.7046	-0.4096	0.8442	-0.5667	0.5232	-0.1572	0.8934
---	---	1638034_at	0.1601	0.4222	0.1069	0.5152	0.1354	0.4557	0.0831	0.8794	-0.0871	0.6827	-0.1702	0.3174	0.0769	0.9441	-0.1232	0.7547	-0.2001	0.5688
CG31841	CG31841	1638035_at	0.1169	0.6077	0.1869	0.1015	-0.0208	0.9228	-0.0777	0.9029	-0.0828	0.7322	-0.0051	0.9849	0.1605	0.7644	-0.1008	0.6937	-0.2613	0.2668
Cyp1	Cyclophilin 1	1638036_at	-0.0244	0.8887	0.3102	0.1808	0.3015	0.0821	0.0663	0.9017	-0.0662	0.7485	-0.1325	0.4161	0.1252	0.7845	0.2405	0.2084	0.1154	0.5820
dpr17	dpr17	1638037_a_at	0.0491	0.8667	-0.4932	0.1998	-0.0712	0.7049	-0.1353	0.8321	0.0172	0.9628	0.1525	0.5168	-0.3245	0.7052	-0.3264	0.3484	-0.0019	0.9979
CG4335	CG4335	1638038_at	2.5353	0.0068	0.9462	0.4189	2.9319	0.0000	1.1392	0.0674	0.8700	0.0239	-0.2692	0.4257	-0.5134	0.8689	-0.3360	0.8386	0.1774	0.9171
CG6048	CG6048	1638039_at	0.1495	0.6314	0.1904	0.3413	0.1275	0.6316	-0.0525	0.9649	-0.1948	0.5731	-0.1423	0.6690	0.0843	0.9246	0.0018	0.9989	-0.0825	0.8256
shd	shade	1638040_at	-0.3025	0.2899	-0.3128	0.3605	-0.1194	0.7069	-0.3124	0.5066	-0.2345	0.3554	0.0779	0.7781	-0.3205	0.7726	-0.3122	0.5327	0.0082	0.9924
mod(mdg4)	Modifier67.2	1638041_at	-0.3580	0.2699	0.0670	0.8489	-0.6598	0.0647	-0.4527	0.4908	-0.3675	0.2983	0.0852	0.8305	0.0890	0.9717	-0.0279	0.9810	-0.1169	0.8921
CG32083	CG32083	1638042_at	0.1729	0.3703	-0.0950	0.5361	0.0278	0.8948	0.1302	0.8170	0.0738	0.7832	-0.0564	0.8219	-0.0356	0.9643	-0.0993	0.6772	-0.0636	0.8111
CG3530	MYOTUBULARIN	1638043_a_at	-0.5445	0.0650	-0.7109	0.0737	-0.6550	0.0158	0.1539	0.7664	0.2502	0.2591	0.0963	0.6769	0.0907	0.9588	0.1083	0.8767	0.0176	0.9835
CG5902 /// DyakCG5902	CG5902	1638044_a_at	0.2192	0.1384	0.6823	0.0121	0.8473	0.0045	0.0148	0.9860	-0.4685	0.0273	-0.4832	0.0151	-0.2915	0.6272	-0.1925	0.4406	0.0990	0.7290
CG14326	CG14326	1638045_at	0.2721	0.3187	0.2078	0.3388	0.0218	0.9191	-0.2241	0.6908	-0.2394	0.3694	-0.0154	0.9628	0.0683	0.9474	0.0254	0.9617	-0.0429	0.9211
Adgf-B	Adenosine deamir	1638046_at	0.1010	0.6812	0.1884	0.2098	0.1321	0.5827	-0.0813	0.8732	-0.0788	0.6994	0.0025	0.9910	0.0170	0.9892	0.0679	0.8487	0.0509	0.8878
CG17262 /// cnir	CG17262 /// cornii	1638047_at	0.1869	0.4331	0.1393	0.4987	0.1906	0.3037	0.2281	0.6822	0.3271	0.2058	0.0989	0.7204	0.1900	0.7779	0.3342	0.2385	0.1442	0.6475
CG31119	CG31119	1638048_at	-1.6234	0.0024	-0.7143	0.0088	-0.9035	0.0030	0.0219	0.9819	-0.9004	0.0029	-0.9222	0.0015	0.0281	0.9848	-0.4645	0.1579	-0.4926	0.1695
CG30494	CG30494	1638049_at	-0.1551	0.6130	0.1251	0.6625	-0.0273	0.8773	-0.6508	0.1194	-0.4408	0.0814	0.2100	0.3661	-0.4341	0.6496	-0.0678	0.9085	0.3663	0.3558
Sox14	Sox box protein 1	1638050_s_at	0.2548	0.4860	0.1603	0.7863	0.6106	0.1479	0.2161	0.7168	-0.4589	0.0916	-0.6750	0.0132	-0.3041	0.8972	-0.5414	0.5541	-0.2373	0.8298
CG17323	CG17323	1638051_at	-1.9704	0.0008	-2.8679	0.0101	-2.3695	0.0000	0.1850	0.5832	0.5265	0.0085	0.3415	0.0334	-0.2086	0.8903	-0.3223	0.5946	-0.1137	0.8844
CG41128	CG41128	1638052_at	-0.2963	0.3158	0.2213	0.6212	-0.2436	0.1592	-0.1553	0.8578	-0.4358	0.1604	-0.2806	0.3241	0.2260	0.8099	0.0127	0.9873	-0.2133	0.6130
Cyp4p1	Cytochrome P450	1638053_at	0.7693	0.0065	-1.1980	0.0162	-0.4395	0.0512	0.6220	0.1697	0.9749	0.0044	0.3529	0.1501	-0.0945	0.9277	-0.6518	0.0714	-0.5573	0.1313
CG13056	CG13056	1638054_at	0.4403	0.0136	0.0264	0.7886	0.1216	0.5418	-0.0551	0.9463	-0.0005	0.9987	0.0546	0.8364	-0.0033	0.9964	-0.1296	0.5129	-0.1263	0.5303
Cpr72Ea	CG4818	1638055_at	0.1807	0.4207	0.0425	0.6480	0.1972	0.3304	0.2011	0.7023	0.0881	0.7610	-0.1129	0.6511	-0.0413	0.9589	0.0403	0.9075	0.0816	0.7527
Gr39a	Gustatory recepto	1638056_at	-0.0584	0.8197	-0.0085	0.9775	0.3035	0.0574	-0.1155	0.6998	-0.2393	0.0830	-0.1238	0.3238	-0.2222	0.7644	0.0382	0.9402	0.2604	0.4148
CG18765	CG18765	1638057_at	-0.1673	0.4501	-0.1009	0.3133	0.0976	0.5079	0.0654	0.9223	-0.0601	0.8099	-0.1255	0.5234	-0.2253	0.7070	-0.1561	0.5523	0.0692	0.8256
---	---	1638058_at	0.0731	0.5980	0.0207	0.8796	-0.0126	0.9680	-0.1286	0.7225	-0.2214	0.1729	-0.0928	0.5660	-0.0162	0.9877	-0.0035	0.9942	0.0127	0.9724
Dmn	dynamitin	1638059_at	-0.6905	0.0195	0.2628	0.2647	0.1896	0.2226	0.0072	0.9937	-0.3541	0.0622	-0.3612	0.0375	0.0170	0.9914	0.5416	0.1078	0.5245	0.1409
CG10077	CG10077	1638060_at	-0.0483	0.8892	-0.2977	0.5977	-0.8334	0.0321	-0.5392	0.3053	0.3738	0.2265	0.9130	0.0060	-0.0708	0.9816	0.1244	0.9111	0.1951	0.8310
---	---	1638061_at	0.1707	0.4396	0.0056	0.9660	0.3585	0.0731	0.1047	0.8315	0.0856	0.6913	-0.0192	0.9344	-0.0797	0.9441	0.0339	0.9507	0.1136	0.7798
---	---	1638062_at	0.1158	0.4491	-0.1980	0.4737	-0.1816	0.3564	-0.1752	0.6010	0.1444	0.4054	0.3196	0.0406	-0.1595	0.8814	-0.2357	0.5860	-0.0763	0.8919
CG34417	CG3950	1638063_at	-0.4345	0.5801	0.1496	0.8437	0.0295	0.9500	0.1867	0.9311	0.3698	0.5926	0.1831	0.7975	0.1586	0.9616	0.7218	0.4309	0.5631	0.5634
CG12342 /// dgo	CG12342 /// Diego	1638064_a_at	-1.0156	0.0068	-0.3259	0.6094	-0.2809	0.1740	0.1420	0.8271	-0.4783	0.0602	-0.6203	0.0134	-0.1016	0.9398	0.0675	0.9157	0.1691	0.7154
---	---	1638065_at	0.1421	0.4436	0.1177	0.5197	-0.0433	0.8573	-0.0067	0.9941	-0.0902	0.6711	-0.0835	0.6662	0.1889	0.7215	-0.0356	0.9175	-0.2245	0.3215
Os-E	Olfactory-specific	1638066_at	0.2200	0.4858	0.0412	0.8500	0.1167	0.5048	-0.0629	0.9252	-0.1848	0.3686	-0.1219	0.5345	-0.0071	0.9964	-0.2123	0.6382	-0.2051	0.6473
CG5261	CG5261	1638067_a_at	-0.3480	0.2663	0.8932	0.1330	1.0424	0.0058	-0.0520	0.9641	-1.4102	0.0015	-1.3581	0.0010	-0.2261	0.9092	-0.2099	0.8180	0.0162	0.9884
RpS11	Ribosomal protein	1638068_a_at	0.0619	0.6635	0.5206	0.0314	0.8138	0.0112	0.1179	0.8094	-0.3426	0.0797	-0.4605	0.0162	-0.0655	0.9204	0.0166	0.9639	0.0821	0.7484
CG13740	CG13740	1638069_at	0.0830	0.6684	0.3391	0.0586	0.1304	0.4094	-0.1859	0.6062	-0.1444	0.4429	0.0416	0.8426	0.0320	0.9774	-0.0301	0.9445	-0.0621	0.8577
CG10418	CG10418	1638070_at	0.1497	0.5039	0.5787	0.1121	0.6633	0.0099	0.2562	0.4779	-0.0894	0.6868	-0.3456	0.0507	0.2650	0.7877	0.3407	0.4124	0.0757	0.8949
---	---	1638071_at	-0.0301	0.8665	-0.0746	0.5597	-0.0010	0.9958	0.0537	0.9094	0.0087	0.9702	-0.0450	0.7882	0.0814	0.8692	-0.0433	0.8779	-0.1247	0.5488
CG15225	CG15225	1638072_at	0.0456	0.8184	0.2262	0.1957	0.2759	0.0910	-0.1452	0.7475	-0.1797	0.3715	-0.0344	0.8823	-0.0793	0.9174	0.0604	0.8735	0.1396	0.6257
mRpL38	mitochondrial ribo	1638073_at	-0.0335	0.9144	-0.0854	0.4191	-0.3355	0.1011	-0.1113	0.8271	-0.0793	0.7302	0.0320	0.8915	0.0930	0.9137	-0.1152	0.7439	-0.2082	0.5032
CG6673	CG6673	1638074_at	-0.0215	0.9706	0.6799	0.1018	0.8268	0.0023	-0.1975	0.6327	-1.2283	0.0005	-1.0308	0.0006	-0.4093	0.7877	-0.3499	0.6142	0.0594	0.9479
CG10494	CG10494	1638075_a_at	-0.1424	0.5469	-0.0467	0.7833	0.2457	0.2979	0.2207	0.6327	-0.0491	0.8701	-0.2697	0.1811	0.0184	0.9898	0.1050	0.7897	0.0867	0.8256
---	---	1638076_at	0.2427	0.1047	0.0940	0.4672	0.3693	0.1336	0.0953	0.8776	0.1342	0.5577	0.0388	0.8776	-0.1638	0.8479	0.0099	0.9901	0.1737	0.6498
CG7634	CG7634	1638077_at	0.0137	0.9392	0.6782	0.1274	0.5489	0.1121	0.1616	0.7023	0.0726	0.7539	-0.0890	0.6578	0.4456	0.7644	0.7867	0.1957	0.3411	0.6145
Side	similar to Deadpar	1638078_at	-0.0614	0.7632	-0.0977	0.5598	-0.1562	0.5198	0.2345	0.6822	-0.0221	0.9555	-0.2566	0.2874	0.1266	0.8541	-0.0338	0.9409	-0.1604	0.5989
l(2)05510	lethal (2) 05510	1638079_at	-1.6563	0.0032	-2.2540	0.0098	-2.5367	0.0001	-0.1010	0.8822	-0.2011	0.4012	-0.1000	0.6815	0.0641	0.9816	-0.8009	0.2026	-0.8651	0.2054
CG13658	CG13658	1638080_at	-0.3262	0.3396	0.0868	0.8988	-0.6103	0.0808	-0.5758	0.3166	0.0616	0.8924	0.6374	0.0400	0.3341	0.8465	0.5646	0.4303	0.2305	0.7878
CG3119	CG3119	1638081_at	-0.1530	0.6397	-0.0370	0.7066	-0.2327	0.3480	-0.0600	0.9567	-0.0651	0.8687	-0.0051	0.9890	0.1033	0.8510	0.0300	0.9341	-0.0733	0.7861

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG6678	CG6678	1638082_at	0.2733	0.1531	0.0662	0.6087	0.2008	0.4149	0.1824	0.7293	0.1717	0.4876	-0.0107	0.9711	0.0708	0.9302	0.0264	0.9504	-0.0444	0.9024
---	---	1638083_at	0.1604	0.4222	0.1108	0.3878	0.0494	0.7639	-0.0202	0.9705	0.0530	0.7510	0.0732	0.6056	-0.1114	0.8270	-0.1191	0.5972	-0.0076	0.9835
---	---	1638084_at	-0.2715	0.0967	-0.1138	0.4390	-0.4241	0.0347	-0.0154	0.9860	-0.1608	0.4529	-0.1453	0.4554	0.0328	0.9589	0.0429	0.8664	0.0100	0.9714
ran-like	ran-like	1638085_at	-0.0278	0.8694	0.0000	1.0000	-0.0304	0.8517	-0.0926	0.7949	-0.0651	0.6967	0.0276	0.8722	-0.0244	0.9760	-0.0022	0.9961	0.0222	0.9350
CG3746	CG3746	1638086_at	-2.7482	0.0026	-0.4174	0.3260	-1.2889	0.0020	-0.4608	0.4596	-2.1039	0.0004	-1.6431	0.0007	0.2991	0.8400	-0.3754	0.5591	-0.6745	0.2855
CG33523	MSP protein	1638087_at	-0.1970	0.6840	0.7308	0.1340	0.9273	0.0175	0.2308	0.7809	-0.5640	0.1014	-0.7948	0.0179	0.1196	0.9649	0.4438	0.5621	0.3241	0.6877
CG30172	CG30172	1638088_at	-0.1306	0.5609	-0.0491	0.8611	0.1541	0.4814	0.1534	0.7556	-0.0318	0.9149	-0.1852	0.3418	-0.0601	0.9545	-0.0601	0.8967	0.0000	1.0000
---	---	1638089_at	0.0444	0.7985	0.1209	0.3047	0.0495	0.7819	-0.0411	0.9373	0.0179	0.9331	0.0590	0.7136	0.0950	0.8400	0.0227	0.9425	-0.0723	0.7470
CG4592	CG4592	1638090_at	0.3256	0.2637	0.5114	0.0713	0.5316	0.0270	-0.1502	0.7937	-0.5625	0.0245	-0.4123	0.0536	-0.1385	0.8894	-0.3049	0.4110	-0.1664	0.6877
CG15143	CG15143	1638091_at	0.1880	0.3970	-0.0812	0.4829	0.0219	0.9181	-0.0094	0.9932	0.2097	0.3427	0.2190	0.2629	-0.0360	0.9717	0.0674	0.8437	0.1033	0.7156
Arpc3A	Arpc3A	1638092_a_at	-0.1887	0.3650	0.3090	0.1074	0.5371	0.0574	0.3222	0.3013	0.2539	0.1633	-0.0683	0.7269	0.1291	0.9238	0.7581	0.0966	0.6290	0.1823
---	---	1638093_at	0.1754	0.2408	-0.2175	0.3178	0.0176	0.9343	0.2068	0.5953	0.5220	0.0163	0.3151	0.0758	-0.0304	0.9701	0.0392	0.9009	0.0697	0.7764
CG6255	CG6255	1638094_at	0.2072	0.1896	0.2039	0.4682	0.4850	0.0128	-0.0321	0.9745	-0.3373	0.1727	-0.3052	0.1687	-0.1800	0.7979	-0.0961	0.7949	0.0838	0.8187
CG3709	CG3709	1638095_at	0.8035	0.0156	0.1585	0.7705	0.2749	0.1957	0.0637	0.9441	1.0329	0.0027	0.9692	0.0021	-0.1079	0.9598	0.2516	0.7184	0.3595	0.5792
CG14315	CG14315	1638096_at	-0.0042	0.9844	0.1521	0.1849	0.2892	0.1454	-0.1048	0.8084	-0.0763	0.7003	0.0285	0.8899	-0.0522	0.9309	0.0997	0.6389	0.1520	0.4528
CG33543	CG33543	1638097_at	0.0276	0.9049	0.0805	0.6681	0.0244	0.9358	-0.1989	0.6893	-0.2039	0.3884	-0.0051	0.9861	0.0763	0.9398	-0.0681	0.8779	-0.1443	0.6677
CG31954	CG31954	1638098_at	0.1642	0.3777	0.5271	0.0223	0.4272	0.0436	-0.1242	0.7762	-0.1637	0.3810	-0.0395	0.8525	-0.0552	0.9492	0.0769	0.8187	0.1320	0.6345
---	---	1638099_at	0.0887	0.6590	0.0034	0.9813	0.4494	0.0438	0.0174	0.9803	0.0700	0.7299	0.0525	0.7861	-0.1305	0.8692	-0.0177	0.9739	0.1128	0.7577
Ptpmeg	split central compl	1638100_s_at	1.2528	0.0128	0.9472	0.2255	1.1453	0.0045	0.1227	0.8070	0.1691	0.4110	0.0464	0.8404	-0.0581	0.9878	-0.1959	0.8721	-0.1377	0.9067
CG15528	MKP-like	1638101_at	-0.6521	0.0471	1.1672	0.0185	0.3596	0.3683	-0.7662	0.2992	-1.9610	0.0016	-1.1948	0.0083	-0.0337	0.9862	-0.1411	0.7989	-0.1073	0.8484
---	---	1638102_s_at	0.0382	0.8651	-0.0192	0.8523	-0.0072	0.9718	0.1123	0.7929	0.1778	0.3125	0.0656	0.7246	0.0130	0.9875	-0.0297	0.9188	-0.0427	0.8633
---	---	1638103_at	0.1264	0.3927	-0.2447	0.3749	0.0877	0.7227	0.4451	0.0899	0.5444	0.0045	0.0993	0.5019	-0.1713	0.8331	-0.0079	0.9924	0.1634	0.6562
---	---	1638104_a_at	0.2478	0.1522	0.5038	0.0625	0.6914	0.0085	0.0810	0.8903	-0.1687	0.4038	-0.2497	0.1545	0.0201	0.9892	0.1702	0.6058	0.1501	0.6528
Dys	Dystrophin-like pr	1638105_at	0.2122	0.2609	0.0815	0.5686	0.0296	0.9052	0.0119	0.9934	0.0449	0.9081	0.0329	0.9235	-0.0288	0.9705	-0.0257	0.9353	0.0031	0.9926
---	---	1638106_at	0.0736	0.6676	0.1105	0.3450	0.2780	0.1637	-0.0262	0.9761	-0.0544	0.8471	-0.0282	0.9155	-0.1265	0.8331	-0.0157	0.9696	0.1107	0.6881
CG5762 /// DmauCG5762 /	CG5762	1638107_at	0.4667	0.2994	0.0012	0.9960	-0.0337	0.8880	0.0079	0.9937	-0.0219	0.9394	-0.0298	0.9011	0.1802	0.9112	-0.3511	0.5591	-0.5313	0.3676
Sgs8	group II	1638108_at	0.0682	0.6697	0.0009	0.9950	0.1149	0.5383	-0.0258	0.9684	-0.0053	0.9830	0.0205	0.9215	-0.1349	0.8207	-0.0502	0.8893	0.0847	0.7704
CG4692 /// DyakCG4692	ATP synthase /// (	1638109_s_at	-0.0998	0.6964	0.4863	0.1353	0.2733	0.1828	-0.1117	0.8732	-0.7173	0.0100	-0.6056	0.0131	0.0740	0.9590	-0.2901	0.4969	-0.3641	0.3914
CG40467	CG40467	1638110_at	-0.2870	0.1416	-0.9482	0.0783	-0.4024	0.0860	0.3372	0.6122	0.6634	0.0518	0.3262	0.2795	-0.3628	0.7230	-0.1218	0.8380	0.2410	0.6144
fru	fru-satori	1638111_at	0.5203	0.1003	0.5419	0.0108	0.3372	0.1035	-0.0206	0.9838	0.1144	0.6611	0.1350	0.5509	0.1164	0.8845	0.0471	0.9205	-0.0693	0.8621
CG1908 /// DmirCG1908	CG1908	1638112_at	-0.1547	0.4333	0.0435	0.7996	-0.2082	0.2405	-0.1920	0.4860	-0.0937	0.5541	0.0983	0.4838	0.0231	0.9852	0.1054	0.7554	0.0823	0.8164
beta4GalNAcTA	beta4GalNAcTA	1638113_at	-0.6938	0.0078	-0.1812	0.5764	-0.3433	0.0621	-0.2369	0.5336	-0.3203	0.1061	-0.0833	0.6896	-0.2056	0.8378	0.1019	0.8583	0.3075	0.4699
CG13022	CG13022	1638114_at	0.1659	0.4075	0.0153	0.9146	0.1258	0.4421	0.0693	0.8943	-0.0964	0.6162	-0.1657	0.2931	-0.0186	0.9841	-0.0826	0.7399	-0.0640	0.8044
---	---	1638115_at	0.1168	0.5293	0.1963	0.3715	0.1125	0.5623	0.0222	0.9777	-0.0120	0.9696	-0.0342	0.8869	0.1103	0.8153	-0.0163	0.9593	-0.1266	0.5378
---	---	1638116_at	-0.1337	0.5281	0.0072	0.9491	-0.1393	0.4202	0.0826	0.8444	0.0088	0.9709	-0.0738	0.6435	0.1933	0.7673	0.1770	0.5404	-0.0164	0.9694
CG14451	CG14451	1638117_at	0.0016	0.9934	-0.0047	0.9821	0.0581	0.8283	-0.0182	0.9803	-0.0729	0.7316	-0.0547	0.7872	0.0235	0.9848	0.0367	0.9349	0.0132	0.9766
CG4103	CG4103	1638118_at	-0.2297	0.5015	0.4126	0.0878	0.7233	0.0097	-0.0748	0.9110	-0.5186	0.0228	-0.4438	0.0273	-0.3999	0.7230	0.1523	0.8103	0.5522	0.2757
PGRP-LE	Peptidoglycan rec	1638119_at	-0.2985	0.0958	-0.0421	0.7908	0.1129	0.6404	0.2387	0.6046	-0.0952	0.7251	-0.3339	0.1069	0.0903	0.8903	0.1186	0.6601	0.0283	0.9344
CG1927	CG1927	1638120_at	-0.6128	0.0090	-0.0807	0.6701	-0.3359	0.0524	-0.0809	0.8822	-0.5022	0.0150	-0.4213	0.0196	0.2830	0.6749	0.0109	0.9841	-0.2720	0.3485
CG9961	CG9961	1638121_at	0.6230	0.0198	0.5998	0.0827	1.2167	0.0033	0.3177	0.3487	-0.0122	0.9678	-0.3300	0.0618	-0.3475	0.7215	-0.0367	0.9557	0.3108	0.4569
CG10874	CG10874	1638122_a_at	0.0705	0.8250	0.2762	0.1743	0.6130	0.0493	0.1121	0.8871	-0.2032	0.4652	-0.3153	0.1849	-0.1017	0.9449	-0.0028	0.9989	0.0990	0.8667
---	---	1638123_at	0.1521	0.2937	-0.4296	0.1026	0.1902	0.3358	0.3949	0.1411	0.5909	0.0039	0.1961	0.1760	-0.1838	0.8424	0.0548	0.9274	0.2386	0.5506
CG30121	CG30121	1638124_at	0.4203	0.0190	0.0242	0.8879	0.1937	0.3883	0.0289	0.9647	0.1240	0.4939	0.0951	0.5809	-0.1320	0.8744	-0.2581	0.4307	-0.1260	0.7398
msi	musashi	1638125_a_at	-0.8538	0.3509	-3.0124	0.0230	-2.3387	0.0002	0.5658	0.2397	1.8322	0.0004	1.2664	0.0010	-0.2906	0.9514	-0.3381	0.8635	-0.0475	0.9842
---	---	1638126_at	-0.5912	0.0094	-0.2165	0.1589	-0.2083	0.4108	0.0702	0.9110	-0.3849	0.0578	-0.4551	0.0188	0.0126	0.9942	-0.1071	0.8399	-0.1197	0.8041
Dlic2	Dlic2	1638127_s_at	0.2610	0.4295	0.2620	0.1642	-0.0260	0.9091	-0.0223	0.9863	0.4067	0.1866	0.4290	0.1200	0.2365	0.7956	0.4126	0.2664	0.1761	0.6763
CG30280	CG30280	1638128_at	2.8252	0.0039	0.6359	0.2583	1.7497	0.0001	1.2103	0.1714	0.2532	0.6792	-0.9571	0.0509	-0.0513	0.9862	-1.6234	0.0391	-1.5720	0.0559
Or74a	Odorant receptor	1638129_at	0.0835	0.5727	0.0696	0.6766	0.1197	0.5193	0.1398	0.7929	0.0499	0.8611	-0.0899	0.6937	0.0032	0.9970	-0.0399	0.8967	-0.0431	0.8774
CG7379	CG7379	1638130_at	-0.1084	0.7213	0.5230	0.0902	0.6963	0.0023	0.1388	0.7904	-0.5455	0.0187	-0.6844	0.0041	0.0086	0.9952	0.0985	0.8356	0.0900	0.8430
5-HT1A	serotonin-recepto	1638131_s_at	-0.0745	0.6619	-0.0663	0.5926	0.1589	0.4711	-0.0306	0.9629	0.1272	0.4895	0.1578	0.3246	-0.2349	0.7422	-0.0013	0.9994	0.2336	0.4523
CG10184	CG10184	1638132_at	0.4026	0.1560	0.5579	0.0314	0.8351	0.0845	-0.1760	0.5735	0.0881	0.6201	0.2641	0.0684	-0.4219	0.8270	0.1625	0.8884	0.5844	0.4786
qkr58E-3	KH domain encon	1638133_at	-0.1368	0.4845	-0.3774	0.2553	-0.2327	0.2236	-0.0605	0.9302	-0.3030	0.1337	-0.2425	0.1799	-0.1276	0.9142	-0.5818	0.1498	-0.4542	0.2868
---	---	1638134_at	0.0661	0.6457	0.1068	0.4441	0.0278	0.9277	0.1145	0.8265	0.0434	0.8712	-0.0711	0.7437	0.2417	0.7633	0.0435	0.9353	-0.1982	0.5807
CG10555 /// DsimCG10555	CG10555	1638135_at	-0.2659	0.6205	-0.0550	0.9536	-0.3198	0.2643	-0.1001	0.7956	-0.0771	0.6663	0.0230	0.9034	0.0097	0.9987	0.1190	0.9451	0.1093	0.9417
CG15765 /// DanaCG1576	CG15765	1638136_at	0.4380	0.0327	0.0550	0.7581	0.1279	0.5137	0.2148	0.5798	0.1912	0.3433	-0.0236	0.9229	-0.0214	0.9816	-0.3174	0.1420	-0.2960	0.2034
CG13310	CG13310	1638137_at	-0.3091	0.3299	0.0012	0.9940	-0.3886	0.0561	-0.3796	0.5093	-0.5113	0.0936	-0.1317	0.6794	0.0469	0.9520	-0.0594	0.8531	-0.1063	0.6794
CG9014	CG9014	1638138_at	0.0706	0.6818	0.0730	0.7543	0.2681	0.2065	-0.0338	0.9471	-0.0675	0.6814	-0.0337	0.8393	-0.1824	0.8192	-0.0972	0.8189	0.0852	0.8372
CG31664	CG31664	1638139_at	-3.3948	0.0008	-4.0182	0.0085	-3.6328	0.0004	0.6197	0.5735	0.9196	0.1020	0.2999	0.5946	-0.0040	0.9997	0.0752	0.9631	0.0791	0.9548
CG14043 /// DyakCG14043	CG14043	1638140_at	0.0165	0.9242	0.0915	0.7319	-0.0494	0.7656	-0.1173	0.7880	0.1904	0.2923	0.3077	0.0587	0.1393	0.8564	0.2755	0.3737	0.1361	0.6990
---	---	1638141_x_at	0.1293	0.5719	0.1671	0.2788	0.2995	0.0635	0.0179	0.9860	0.0492	0.8759	0.0313	0.9136	-0.1414	0.8023	-0.0027	0.9961	0.1387	0.5808
CG3288	CG3288	1638142_at	0.0483	0.7410	0.0487	0.8056	0.2041	0.4199	0.1111	0.7809	-0.0750	0.6940	-0.1861	0.2094	-0.0579	0.9545	-0.0717	0.8625	-0.0138	0.9769
Pvr	VEGF receptor	1638143_a_at	-0.9137	0.0023	-0.8164	0.0402	-0.8704	0.0039	-0.1738	0.5363	-0.2431	0.0983	-0.0692	0.6474	0.0475	0.9764	-0.0140	0.9849	-0.0615	0.9080
CG40267	CG40267	1638144_at	-0.1130	0.5431	0.1011	0.3353	-0.0204	0.9388	-0.1268	0.8192	-0.0417	0.8871	0.0851	0.7129	0.0951	0.9238	0.2177	0.5234	0.1226	0.7464
---	---	1638145_at	0.1354	0.6088	0.1759	0.2937	0.2897	0.1172	0.0240	0.9819	0.1373	0.6184	0.1134	0.6605	-0.0231	0.9884	0.1827	0.6191	0.2058	0.5719
Rtc1	Rtc1	1638146_at	0.7793	0.0307	0.8407	0.0456	1.4841	0.0010	0.3507	0.5128	0.0468	0.9045	-0.3039	0.2306	-0.3831	0.7697	0.1192	0.8820	0.5023	0.3803
CG7772	CG7772	1638147_at	-1.3727	0.0083	-1.0620	0.0521	-1.1689	0.0006	-0.3261	0.4631	-0.5155	0.0386	-0.1894	0.3975	-0.2576	0.8541	-0.4749	0.4073	-0.2174	0.7442
CG33137	CG33137	1638148_at	0.1335	0.5664	0.0374	0.6963	-0.3402	0.1814	-0.2065	0.5039	-0.1280	0.4532	0.0785	0.6410	0.0000	0.9999	-0.5083	0.2246	-0.5083	0.2574
---	---	1638149_at	0.1031	0.4802	-0.1600	0.5370	-0.2271	0.2306	-0.0535	0.9470	0.2993	0.1863	0.3528	0.0841	0.1309	0.8940	0.1588	0.7032	0.0279	0.9584
---	---	1638150_at	-0.0589	0.7878	0.0129	0.9620	0.0707	0.6519	0.0033	0.9960	-0.2136	0.3043	-0.2169	0.2400	-0.0592	0.9421	0.0216	0.9587	0.0808	0.7865

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG14837	CG14837	1638151_a_at	0.0340	0.9008	0.0389	0.7827	0.1665	0.3155	0.2476	0.4690	0.1508	0.4287	-0.0968	0.6010	0.0586	0.9558	0.1227	0.7315	0.0641	0.8779
---	---	1638152_s_at	0.2846	0.1758	0.1816	0.3748	0.2328	0.2243	-0.1205	0.8162	-0.1059	0.6431	0.0147	0.9549	0.0886	0.9093	-0.1420	0.6345	-0.2306	0.4114
CG7907	CG7907	1638153_at	-0.0314	0.9095	-0.1935	0.3712	-0.0627	0.6969	0.2530	0.5149	0.2430	0.2368	-0.0100	0.9689	0.0037	0.9970	0.0252	0.9494	0.0215	0.9495
CG2187	CG2187	1638154_at	-1.9716	0.0134	-0.3112	0.4889	-1.0846	0.0065	-0.5898	0.5068	-1.6911	0.0042	-1.1013	0.0171	-0.1225	0.9717	-0.2841	0.7932	-0.1616	0.8918
l(3)neo43	lethal (3) neo43	1638155_at	-0.2066	0.2220	0.0592	0.9115	-0.2701	0.3466	-0.0108	0.9937	-0.1220	0.6952	-0.1112	0.6961	0.2195	0.7707	0.0441	0.9313	-0.1753	0.6124
fy	fuzzy	1638156_at	0.4457	0.0596	0.5591	0.2789	0.2006	0.4497	-0.0586	0.9488	0.3683	0.1463	0.4269	0.0642	0.2726	0.8014	0.4531	0.3018	0.1805	0.7257
CG18012	CG18012	1638157_at	0.7335	0.0129	0.6492	0.0692	0.6242	0.0053	-0.0595	0.9295	-0.0808	0.7247	-0.0213	0.9295	-0.0687	0.9571	-0.2224	0.5591	-0.1537	0.7060
CG4845	CG4845	1638158_at	0.0489	0.8281	-0.0052	0.9752	0.1105	0.5977	0.1456	0.7827	0.5977	0.0140	0.4522	0.0282	0.0886	0.9398	0.5893	0.1070	0.5008	0.1883
---	---	1638159_at	0.0859	0.6397	-0.0452	0.7767	-0.0239	0.8924	-0.1537	0.6823	0.0355	0.8791	0.1892	0.2271	0.0072	0.9939	-0.1441	0.5214	-0.1513	0.5007
CG12538	CG12538	1638160_at	0.0971	0.6987	0.3101	0.2350	0.1601	0.4307	0.1339	0.7817	0.0520	0.8405	-0.0818	0.6976	0.1956	0.7046	0.2028	0.3322	0.0073	0.9841
Ssl1	Ssl1	1638161_at	-0.5909	0.0717	0.1813	0.3343	-0.0113	0.9563	-0.1762	0.6537	-0.4512	0.0253	-0.2750	0.1037	0.0443	0.9816	0.4119	0.3101	0.3676	0.3887
CG1244	CG1244	1638162_s_at	-0.4496	0.1778	0.2867	0.5093	0.5337	0.1199	0.2075	0.8162	-0.3030	0.4000	-0.5105	0.1025	-0.0316	0.9914	0.5155	0.4460	0.5471	0.4301
trr	trithorax-related	1638163_a_at	-0.1609	0.7261	0.3282	0.4924	0.4502	0.0240	0.3637	0.6615	-0.0477	0.9320	-0.4114	0.2491	0.0210	0.9935	0.3238	0.6049	0.3027	0.6318
CG4289	CG4289	1638164_at	0.4809	0.0307	0.2166	0.3704	0.1535	0.5265	-0.1263	0.7313	-0.1098	0.5313	0.0165	0.9354	-0.0683	0.9589	-0.4273	0.2385	-0.3590	0.3519
---	---	1638165_s_at	0.9832	0.3557	0.9964	0.0586	-0.6564	0.1221	-0.1252	0.9269	1.0384	0.0192	1.1636	0.0071	1.5112	0.6659	0.9830	0.4958	-0.5282	0.7442
Rpb10	Rpb10	1638166_at	0.0164	0.9612	0.5233	0.0640	0.2641	0.2719	0.0915	0.8671	-0.1971	0.3115	-0.2886	0.0940	0.3883	0.6955	0.3143	0.4565	-0.0740	0.8982
RabX6	RabX6	1638167_at	-0.2172	0.4578	0.4776	0.1301	0.5244	0.0234	0.0113	0.9931	-0.3123	0.2179	-0.3235	0.1533	0.0079	0.9964	0.3626	0.3963	0.3547	0.4214
---	---	1638168_at	0.2066	0.3664	0.0892	0.4918	0.3381	0.1797	0.0665	0.9375	0.0158	0.9675	-0.0507	0.8600	-0.0698	0.9506	-0.0422	0.9376	0.0276	0.9533
---	---	1638169_s_at	-0.0988	0.5576	0.5415	0.3075	1.0507	0.0034	-0.0215	0.9857	-0.4442	0.0985	-0.4227	0.0800	-0.4295	0.7204	0.2775	0.6014	0.7070	0.1808
CG40130	CG40130	1638170_at	0.1502	0.4799	0.1298	0.4107	0.0936	0.5578	-0.0815	0.8937	-0.0017	0.9951	0.0798	0.7109	-0.0015	0.9994	0.0367	0.9158	0.0382	0.9023
CG1753	CG1753	1638171_s_at	2.4082	0.0004	2.8742	0.0040	2.3324	0.0003	0.7172	0.0417	-0.1278	0.5477	-0.8451	0.0012	1.3755	0.2884	0.4360	0.5373	-0.9395	0.1906
---	---	1638172_at	0.3559	0.0888	0.1448	0.3436	0.2737	0.2297	-0.0203	0.9819	-0.1886	0.3849	-0.1684	0.3930	0.0307	0.9751	-0.1040	0.7043	-0.1347	0.6052
CG2950 /// DyakCG2950	CG2950	1638173_s_at	-0.8315	0.0915	-0.2201	0.6893	-0.6457	0.0628	-0.2105	0.6010	-0.4651	0.0294	-0.2546	0.1592	0.2438	0.9239	0.1891	0.8767	-0.0547	0.9659
CG1246	CG1246	1638174_at	0.0631	0.9719	-0.3036	0.0589	-0.4904	0.1399	0.0285	0.9952	-0.3199	0.7869	-0.3485	0.7391	0.0023	0.9999	-0.8655	0.5654	-0.8678	0.5691
mst	lethal(1)19Fg	1638175_at	-0.2275	0.3970	0.0914	0.8739	0.9677	0.0122	0.3646	0.3576	0.2369	0.2995	-0.1277	0.5691	-0.3612	0.8016	0.6646	0.2507	1.0258	0.1192
Magi	Magi	1638176_at	-1.8553	0.0052	-2.3420	0.0048	-2.2493	0.0002	0.6812	0.4034	0.9158	0.0493	0.2346	0.6100	0.5498	0.5228	0.4192	0.2974	-0.1306	0.7925
---	---	1638177_at	0.0566	0.7664	0.0242	0.8101	0.2662	0.1499	0.0014	0.9988	-0.1314	0.5436	-0.1327	0.4909	-0.0704	0.9545	-0.0961	0.8445	-0.0257	0.9614
CG31949	CG31949	1638178_at	0.1355	0.5870	-0.1570	0.3110	-0.0121	0.9552	0.0796	0.8598	0.1873	0.2483	0.1077	0.4861	-0.1220	0.8465	-0.1405	0.6156	-0.0185	0.9614
CG12107	CG12107	1638179_at	-0.4009	0.2824	0.1874	0.4015	0.6371	0.0137	-0.1093	0.8189	-0.7321	0.0029	-0.6227	0.0035	-0.4288	0.7220	-0.0393	0.9630	0.3895	0.4634
---	---	1638180_at	0.1071	0.6045	-0.0934	0.4699	-0.0969	0.6016	-0.0012	0.9988	0.3052	0.1759	0.3064	0.1290	-0.0323	0.9764	0.0450	0.9111	0.0773	0.8123
CG14743	CG14743	1638181_at	0.0391	0.8651	0.1394	0.2270	-0.0308	0.9181	-0.3180	0.5255	-0.2148	0.4287	0.1032	0.7109	-0.0344	0.9831	0.1014	0.8430	0.1359	0.7549
CG5999	CG5999	1638182_at	3.9192	0.0014	2.1206	0.0576	3.7194	0.0002	0.1699	0.8156	0.1387	0.6706	-0.0312	0.9291	-0.7279	0.8013	-1.2332	0.2930	-0.5053	0.7097
CG13868	CG13868	1638183_at	0.8433	0.2936	0.1495	0.8646	0.5130	0.0882	0.5650	0.2688	0.2123	0.5054	-0.3527	0.1900	0.2314	0.9588	-0.5360	0.7030	-0.7674	0.5612
CG13936	CG13936	1638184_at	-2.4892	0.0015	1.0728	0.2084	-1.3476	0.0148	-2.2000	0.0249	-3.9311	0.0003	-1.7312	0.0038	0.3300	0.8940	-0.2319	0.8551	-0.5620	0.5672
CG14613	CG14613	1638185_at	-0.3679	0.0499	0.0552	0.6288	0.2465	0.1247	0.0957	0.8074	-0.6780	0.0018	-0.7737	0.0006	-0.0223	0.9826	-0.1418	0.5654	-0.1195	0.6385
CG9025 /// DmirCG9025 ///	CG9025 /// GA21	1638186_a_at	-0.5312	0.0208	-0.3140	0.0363	-0.6152	0.0154	-0.3110	0.3839	-0.6433	0.0072	-0.3323	0.0687	0.0447	0.9689	-0.3790	0.1948	-0.4237	0.1860
CG13819	CG13819	1638187_at	0.5773	0.0571	0.3662	0.2314	-0.0438	0.8872	0.0089	0.9951	0.2234	0.4553	0.2145	0.4262	0.3201	0.7142	-0.0235	0.9689	-0.3436	0.3601
---	---	1638188_at	0.0029	0.9907	0.0556	0.6000	0.0598	0.7673	-0.0887	0.8594	-0.0353	0.8839	0.0534	0.7911	-0.0225	0.9816	-0.0392	0.9023	-0.0167	0.9555
S	star	1638189_s_at	0.1834	0.5666	-0.2820	0.2944	-0.2873	0.2876	0.2428	0.7304	1.2313	0.0029	0.9885	0.0046	0.4148	0.7230	0.7332	0.1414	0.3184	0.5523
CG9667	CG9667	1638190_at	0.2822	0.1631	0.1030	0.7185	0.3510	0.0504	0.2577	0.4786	0.4851	0.0212	0.2274	0.1916	-0.0284	0.9862	0.2621	0.4848	0.2905	0.4411
cv-c	crossveinless-c	1638191_at	0.0481	0.8493	0.0796	0.4968	0.2615	0.1787	0.1400	0.8671	-0.0793	0.8337	-0.2193	0.4289	0.0341	0.9816	0.1628	0.6382	0.1287	0.7224
Wsck	Wsck	1638192_a_at	-0.4328	0.1734	-0.6676	0.0082	-0.2923	0.3865	0.4973	0.1649	0.8559	0.0028	0.3586	0.0678	0.1298	0.9447	0.6232	0.2518	0.4934	0.3921
CG5612	CG5612	1638193_at	-2.6781	0.0011	-1.7131	0.0314	-2.2568	0.0000	-0.3222	0.2330	-1.3535	0.0002	-1.0313	0.0002	0.2011	0.9286	-0.2509	0.7788	-0.4520	0.5584
CG4409	CG4409	1638194_at	-1.0277	0.0073	-0.8933	0.0205	-0.7790	0.0768	-0.1081	0.9375	-0.2185	0.6239	-0.1104	0.8062	-0.0057	0.9959	-0.1370	0.6139	-0.1314	0.6311
lwf	lots wife	1638195_at	1.3997	0.0511	-0.7870	0.0052	-0.5320	0.0070	0.5010	0.2132	1.0774	0.0017	0.5764	0.0159	0.1803	0.9226	-1.1225	0.0797	-1.3028	0.0710
alphaTry	alphaTrypsin	1638196_at	1.0161	0.6466	-0.7893	0.0521	-0.6629	0.0943	0.1633	0.9445	0.4442	0.5353	0.2809	0.6896	-0.0432	0.9967	-1.6806	0.5459	-1.6375	0.5629
CG3573	inositol 5-phospha	1638197_at	-1.0169	0.0109	-0.7925	0.1230	-0.4287	0.0602	0.1560	0.8640	-0.1687	0.6425	-0.3247	0.2663	-0.2092	0.8202	0.1287	0.7824	0.3380	0.3812
CG12768	CG12768	1638198_at	-1.0408	0.0471	-0.4384	0.1426	-1.3105	0.0004	-0.6397	0.2501	-0.5871	0.0780	0.0526	0.8948	0.1285	0.8194	0.0080	0.9863	-0.1205	0.6289
RhoGAPp190	Rp190RhoGAP	1638199_s_at	0.1031	0.8031	-0.4810	0.0710	-0.6183	0.0118	-0.0326	0.9688	0.4503	0.0440	0.4829	0.0211	-0.0406	0.9816	-0.1397	0.7703	-0.0992	0.8439
pcx	pecanex	1638200_at	-0.2686	0.5906	-0.1306	0.8350	-0.1323	0.5018	0.1088	0.8738	0.2454	0.3078	0.1366	0.5624	0.0731	0.9816	0.3748	0.6059	0.3017	0.6881
26-29-p	26-29kD-proteina	1638201_at	-0.0048	0.9872	0.2616	0.3452	0.7803	0.0011	0.3411	0.3291	-0.4267	0.0412	-0.7678	0.0019	-0.1880	0.8086	-0.1555	0.6605	0.0325	0.9431
CG34356	CG12524	1638202_at	0.0858	0.6780	0.3322	0.1141	0.2569	0.0865	-0.1659	0.7507	-0.3248	0.1504	-0.1589	0.4593	0.0484	0.9653	0.1188	0.7340	0.0705	0.8575
CG7339	CG7339	1638203_at	0.1012	0.7390	0.1105	0.7524	0.7777	0.0141	0.2035	0.5495	0.0184	0.9429	-0.1851	0.2402	-0.4451	0.7352	0.0807	0.9275	0.5258	0.3626
cav	anonymous fast e	1638204_s_at	-2.2570	0.0039	-0.9050	0.0802	-1.6557	0.0042	-0.8755	0.1483	-2.0452	0.0007	-1.1697	0.0042	-0.2591	0.8953	-0.7901	0.2642	-0.5310	0.4848
---	---	1638205_s_at	-0.0011	0.9964	-0.0242	0.9171	-0.2857	0.2230	-0.2378	0.7104	-0.0033	0.9939	0.2345	0.3849	0.0032	0.9978	-0.0618	0.8721	-0.0650	0.8515
CG30458	CG30458	1638206_at	0.2336	0.1687	0.2762	0.1301	0.3164	0.0723	0.0988	0.8373	0.0575	0.7954	-0.0413	0.8433	0.0713	0.9309	0.0186	0.9664	-0.0527	0.8866
CG15617	CG15617	1638207_at	-0.0138	0.9690	1.4873	0.0595	1.1621	0.0091	-0.4932	0.5777	-1.8238	0.0026	-1.3306	0.0062	-0.1260	0.9638	-0.2202	0.8235	-0.0942	0.9282
CG17019	CG17019	1638208_at	-0.2034	0.5473	0.1005	0.6536	0.3704	0.0601	0.2052	0.5680	-0.0245	0.9242	-0.2297	0.1627	-0.0302	0.9884	0.3590	0.4242	0.3892	0.3953
---	---	1638209_at	0.2216	0.2524	0.0123	0.9108	0.0944	0.6566	0.0597	0.9345	-0.0116	0.9720	-0.0713	0.7609	0.0867	0.9101	-0.0478	0.9089	-0.1345	0.6484
---	---	1638210_at	0.0968	0.5429	0.0215	0.8323	-0.0561	0.8136	-0.1375	0.7803	0.1068	0.6422	0.2443	0.1809	-0.0684	0.9221	0.0096	0.9829	0.0780	0.7761
CG10933	CG10933	1638211_at	-1.4887	0.0076	-2.0703	0.0063	-2.0826	0.0008	0.3095	0.8155	0.6137	0.2393	0.3042	0.5528	0.0328	0.9862	-0.1344	0.8117	-0.1672	0.7381
---	---	1638212_at	-0.1539	0.4869	0.1400	0.3878	0.1535	0.3049	-0.0137	0.9863	-0.2210	0.2570	-0.2073	0.2336	0.0583	0.9445	0.1020	0.7264	0.0437	0.9004
CG40484	CG40484	1638213_at	-0.0962	0.5593	-0.0742	0.6285	-0.0052	0.9820	0.0538	0.9269	-0.0206	0.9338	-0.0744	0.6864	-0.0578	0.9342	-0.0744	0.7919	-0.0166	0.9596
CG34342	CG10629	1638214_at	0.1626	0.3767	0.1972	0.4472	0.0713	0.6519	0.2614	0.6005	0.2318	0.3662	-0.0296	0.9235	0.0281	0.9816	-0.0725	0.8450	-0.1006	0.7495
CG8043 /// DyakCG8043	CG8043	1638215_at	-0.1303	0.5907	0.3587	0.0263	0.8004	0.0051	0.1355	0.8280	-0.3873	0.1051	-0.5229	0.0230	-0.3628	0.6440	0.0768	0.8624	0.4395	0.1910
CG13623	CG13623	1638216_at	-0.0554	0.8816	0.5623	0.1495	0.3878	0.1618	0.0379	0.9463	-0.1995	0.2072	-0.2373	0.0941	0.0426	0.9852	0.1918	0.7604	0.1493	0.8201
CG10166	CG10166	1638217_at	0.9013	0.0062	1.2358	0.0067	1.3873	0.0002	0.2491	0.4551	0.3734	0.0446	0.1244	0.4683	0.1400	0.8692	0.5994	0.0881	0.4594	0.1990
CG10909	CG10909	1638218_at	0.3446	0.1064	0.0982	0.4593	0.0146	0.9505	0.1079	0.7753	0.1929	0.2220	0.0850	0.5917	0.0264	0.9823	-0.0177	0.9689	-0.0441	0.9080
---	---	1638219_at	0.2659	0.2699	0.0034	0.9800	0.2057	0.2456	0.0869	0.8233	0.0672	0.6994	-0.0197	0.9150	-0.1150	0.8243	-0.0555	0.8509	0.0595	0.8256

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG16888	CG16888	1638220_at	0.0624	0.6926	0.0209	0.9523	-0.1139	0.5468	-0.1040	0.8337	0.0415	0.8673	0.1455	0.4066	0.1294	0.8461	-0.0328	0.9404	-0.1623	0.5781
Obp83cd	Odorant-binding p	1638221_at	0.2775	0.6987	1.3271	0.2135	0.2357	0.5778	0.6141	0.2905	0.2277	0.5327	-0.3864	0.2058	1.6410	0.5924	1.2842	0.3299	-0.3568	0.8354
CG8252	CG8252	1638222_at	-0.1083	0.5121	-0.0955	0.6225	0.3022	0.2993	-0.0728	0.9284	-0.0780	0.7899	-0.0052	0.9862	-0.2202	0.8141	0.0344	0.9569	0.2545	0.5331
CG13361	CG13361	1638223_at	0.0476	0.8026	0.0239	0.9219	-0.0530	0.7761	-0.0011	0.9988	0.1982	0.2680	0.1993	0.2108	-0.0257	0.9856	-0.0489	0.9229	-0.0232	0.9612
CG32250	CG32250	1638224_at	0.0946	0.5974	-0.0030	0.9844	0.4943	0.0231	0.1825	0.5766	-0.0479	0.8183	-0.2304	0.1224	-0.2964	0.5604	-0.0051	0.9925	0.2913	0.2237
inx7	innexin7	1638225_a_at	0.1689	0.8862	-0.3034	0.0809	-0.0919	0.7131	0.1919	0.9318	-0.0354	0.9732	-0.2274	0.7551	-0.0760	0.9869	-0.6176	0.5800	-0.5415	0.6328
CG10562	CG10562	1638226_at	-1.7082	0.1053	-3.0406	0.0074	-3.1196	0.0000	-0.4247	0.5902	-0.4579	0.2534	-0.0332	0.9478	-0.3310	0.9340	-1.6120	0.1925	-1.2809	0.3261
SP1070	SP1070	1638227_at	-0.8844	0.1117	-0.8362	0.4044	-0.8512	0.0944	0.3792	0.8432	0.5178	0.4869	0.1386	0.8675	0.3870	0.9012	0.5679	0.6479	0.1808	0.9089
---	---	1638228_s_at	0.1418	0.5022	-0.0516	0.7117	0.0339	0.8645	0.1063	0.8156	0.1302	0.4879	0.0239	0.9124	-0.0144	0.9923	-0.0614	0.9081	-0.0469	0.9217
l(3)01239	lethal (3) 01239	1638229_at	-0.5945	0.0282	0.3164	0.2329	0.5096	0.0578	0.0681	0.8841	-0.6860	0.0019	-0.7541	0.0008	-0.0662	0.9665	0.3560	0.3885	0.4222	0.3247
CG3160	CG3160	1638230_at	-0.4512	0.0571	-0.1604	0.5929	0.0526	0.8842	-0.2076	0.7327	-0.3711	0.1686	-0.1635	0.5372	-0.4765	0.6557	-0.1079	0.8584	0.3686	0.4076
CG4393	CG4393	1638231_at	-0.7672	0.0207	-0.2414	0.2038	-1.1222	0.0029	-0.3174	0.4822	-0.2592	0.2872	0.0582	0.8332	0.2909	0.7601	-0.0413	0.9496	-0.3322	0.4119
---	---	1638232_at	0.2136	0.4860	-0.0113	0.9153	0.3107	0.1369	0.1326	0.7608	0.1551	0.4240	0.0226	0.9218	-0.1611	0.8122	-0.0538	0.8988	0.1073	0.7409
CG4646	CG4646	1638233_at	-0.0312	0.9226	0.4152	0.1210	0.4474	0.1052	-0.0812	0.9102	-0.4640	0.0475	-0.3828	0.0641	-0.0494	0.9816	0.0775	0.9167	0.1270	0.8332
---	---	1638234_s_at	0.1513	0.4277	-0.0313	0.7935	0.2108	0.1653	-0.0350	0.9592	0.0392	0.8701	0.0742	0.6961	-0.0942	0.8700	-0.0734	0.8029	0.0208	0.9487
DptB	diptericin-like prot	1638235_at	2.1952	0.2210	-0.3221	0.8859	-0.6260	0.6955	0.2127	0.9409	3.9270	0.0012	3.7143	0.0009	0.7262	0.9527	1.3538	0.7567	0.6276	0.9020
CG9302	CG9302	1638236_at	0.3283	0.0794	0.1605	0.7052	0.3061	0.0847	0.1686	0.6197	1.0500	0.0004	0.8815	0.0005	0.0400	0.9829	0.8520	0.0661	0.8120	0.0922
CG5412	CG5412	1638237_at	-0.0574	0.8034	0.4334	0.1112	0.8267	0.0012	0.1982	0.5680	0.0283	0.9089	-0.1699	0.2922	-0.1169	0.8609	0.4589	0.0961	0.5758	0.0702
CG5778	CG5778	1638238_at	0.6093	0.1635	1.4213	0.0758	0.6746	0.0263	-0.9070	0.0393	-0.6279	0.0181	0.2791	0.1987	-0.3055	0.8940	-0.1220	0.9277	0.1834	0.8736
---	---	1638239_at	0.1897	0.2699	-0.1082	0.5205	0.0010	0.9972	-0.1016	0.8265	-0.0088	0.9746	0.0928	0.6073	-0.1141	0.8461	-0.1411	0.5855	-0.0270	0.9371
CG8785	CG8785	1638240_s_at	-2.1443	0.0030	-0.6753	0.4761	-2.2384	0.0016	-1.8886	0.0180	-1.8901	0.0015	-0.0014	0.9981	-0.2490	0.9445	-0.4205	0.7403	-0.1715	0.9086
---	---	1638241_at	0.2962	0.1616	0.0081	0.9534	-0.0778	0.6727	-0.1471	0.5946	0.0100	0.9622	0.1571	0.2097	0.0118	0.9922	-0.2047	0.4570	-0.2165	0.4421
---	---	1638242_at	0.2499	0.2107	0.1766	0.4239	0.1409	0.3454	-0.0227	0.9733	0.0296	0.8966	0.0523	0.7781	-0.0143	0.9898	-0.1387	0.5667	-0.1244	0.6166
CG34402	CG17793	1638243_at	0.1030	0.7139	0.4663	0.0132	0.0641	0.7684	-0.1273	0.8738	-0.1957	0.5060	-0.0684	0.8295	0.1534	0.8461	0.0809	0.8583	-0.0725	0.8674
CG8177 /// DmirCG8177	CG8177	1638244_s_at	-0.9619	0.0021	-0.9753	0.0775	-2.5221	0.0001	-0.7521	0.0799	-0.1803	0.4888	0.5718	0.0182	0.7407	0.4415	0.0146	0.9895	-0.7261	0.1744
CG13766	CG13766	1638245_at	-0.1197	0.7392	0.4893	0.0266	0.5375	0.0848	-0.3091	0.6513	-0.5832	0.0793	-0.2741	0.3686	-0.2316	0.8159	0.0401	0.9515	0.2717	0.5299
CG5804	CG5804	1638246_at	-0.7658	0.8314	-1.2173	0.0945	-0.7670	0.0552	0.7398	0.8999	-1.7996	0.3557	-2.5394	0.1358	0.0365	0.9982	-2.7600	0.4037	-2.7965	0.4096
CG17065	CG17065	1638247_at	-1.4334	0.0079	-0.9235	0.2257	-1.3840	0.0070	0.1376	0.7389	-0.4727	0.0177	-0.6103	0.0034	0.4419	0.8608	-0.0914	0.9557	-0.5333	0.6254
CG32350	CG32350	1638248_at	0.3201	0.3353	0.4485	0.2778	-0.0556	0.7743	-0.2446	0.5280	0.1984	0.3347	0.4430	0.0227	0.1821	0.8609	0.3352	0.4223	0.1532	0.7513
regucalcin	regucalcin	1638249_at	0.3338	0.1208	0.2291	0.1569	0.2150	0.1746	-0.0933	0.8844	-0.1293	0.5863	-0.0360	0.8906	0.0108	0.9893	0.0445	0.8461	0.0338	0.8844
---	---	1638250_at	0.2307	0.2115	0.1035	0.3599	0.0606	0.7704	0.1054	0.8174	-0.0440	0.8497	-0.1494	0.3658	0.2049	0.7220	0.0682	0.8358	-0.1367	0.6053
CG5549	CG5549	1638251_at	-0.1259	0.6767	0.0897	0.3835	0.0047	0.9843	-0.0314	0.9592	-0.0816	0.6525	-0.0502	0.7789	0.2746	0.7726	0.3321	0.4094	0.0575	0.9184
pll	pelle	1638252_at	0.0996	0.5743	0.1951	0.3137	0.5987	0.0582	0.1726	0.8089	-0.0837	0.8113	-0.2563	0.3228	-0.0441	0.9616	0.1961	0.4429	0.2402	0.3616
CG5361	CG5361	1638253_at	0.0103	0.9915	0.1278	0.4860	-0.8718	0.0513	-0.9278	0.3444	-0.7117	0.2069	0.2161	0.7199	0.0409	0.9926	-0.5008	0.6467	-0.5417	0.6169
---	---	1638254_at	-0.0754	0.6791	-0.0234	0.8174	0.0275	0.8647	0.2264	0.4626	0.0099	0.9696	-0.2165	0.1454	0.0971	0.8191	0.0169	0.9515	-0.0802	0.6719
---	---	1638255_at	0.1928	0.3946	-0.0645	0.5957	0.0888	0.7201	0.0575	0.9491	0.0517	0.8766	-0.0058	0.9856	-0.1433	0.8814	-0.3059	0.3999	-0.1626	0.6883
---	---	1638256_at	-0.0232	0.9203	-0.0945	0.8841	0.1548	0.3236	-0.4339	0.1956	-0.2540	0.2086	0.1799	0.3319	-0.7334	0.4415	-0.3523	0.4861	0.3811	0.4567
CG2931	CG2931	1638257_at	-0.3229	0.0813	-0.3955	0.1340	-0.3678	0.0368	0.0595	0.8976	0.4560	0.0093	0.3966	0.0106	-0.0182	0.9892	0.4382	0.1199	0.4564	0.1313
---	---	1638258_at	0.1880	0.3906	0.0105	0.9390	0.2782	0.0879	0.0318	0.9626	-0.1202	0.5308	-0.1521	0.3577	-0.1072	0.8454	-0.1095	0.6586	-0.0023	0.9953
Aats-val	Valyl-tRNA synthe	1638259_s_at	0.2531	0.4888	0.9773	0.0611	1.4197	0.0005	0.5122	0.2909	0.1579	0.6165	-0.3542	0.1638	0.1479	0.9301	0.9830	0.0831	0.8351	0.1517
CG2269 /// DmirCG2269	CG2269	1638260_s_at	-0.6531	0.0140	0.0060	0.9759	-0.2547	0.1855	-0.2348	0.4887	-0.7426	0.0024	-0.5079	0.0076	0.0042	0.9964	-0.1079	0.6879	-0.1121	0.6680
Osi22	Osiris	1638261_at	-0.1288	0.5943	0.1719	0.4972	-0.0108	0.9691	-0.0649	0.9038	-0.0990	0.6056	-0.0341	0.8676	-0.0400	0.9742	0.1347	0.6949	0.1746	0.5930
CG17819	CG17819	1638262_at	0.1426	0.3930	0.0563	0.5978	0.0779	0.7159	0.1484	0.6844	0.1146	0.5263	-0.0339	0.8639	0.0899	0.8609	0.0379	0.9047	-0.0520	0.8443
---	---	1638263_at	-0.0859	0.6553	0.0886	0.7031	-0.0368	0.9031	0.1217	0.7441	0.0034	0.9892	-0.1183	0.4427	0.0817	0.9238	0.0345	0.9404	-0.0473	0.9053
CG1304	CG1304	1638264_at	0.1321	0.5401	0.0645	0.5325	-0.0096	0.9577	0.1118	0.7293	-0.0254	0.8979	-0.1372	0.2948	0.1573	0.8222	-0.0644	0.8778	-0.2217	0.4631
---	---	1638265_s_at	0.0613	0.7710	0.4633	0.0705	0.3847	0.0401	0.0639	0.9036	-0.2443	0.1517	-0.3082	0.0491	0.0165	0.9901	-0.0137	0.9796	-0.0303	0.9414
---	---	1638266_at	0.0564	0.7632	0.2490	0.1463	0.2751	0.1420	-0.0402	0.9647	-0.2229	0.3653	-0.1827	0.4179	0.0801	0.9243	0.0396	0.9289	-0.0405	0.9171
kermit	lethal (2) 02045	1638267_s_at	0.0333	0.9323	-0.2299	0.3742	0.0218	0.9331	0.2312	0.7251	0.4119	0.1615	0.1807	0.5310	-0.0905	0.9405	0.0280	0.9643	0.1185	0.7933
TBPH	TBPH	1638268_at	-0.5232	0.1423	-0.3106	0.2848	-0.7519	0.0041	-0.1540	0.8578	-0.0981	0.7963	0.0558	0.8791	0.2957	0.7758	0.1547	0.7788	-0.1410	0.7944
---	---	1638269_at	0.0060	0.9807	0.0744	0.5649	0.2913	0.1538	0.0394	0.9371	0.0129	0.9524	-0.0264	0.8776	-0.1482	0.8042	0.0195	0.9621	0.1677	0.5176
CSN4	COP9 signalosom	1638270_at	0.2860	0.3198	0.5605	0.1296	0.3323	0.0957	-0.4012	0.3276	-0.6550	0.0136	-0.2538	0.2325	-0.1890	0.8270	-0.3681	0.2973	-0.1791	0.6463
---	---	1638271_at	0.1280	0.5885	0.0369	0.7272	0.2347	0.2412	-0.0522	0.9438	-0.0699	0.7863	-0.0177	0.9473	-0.0836	0.9234	0.0943	0.7927	0.1779	0.5579
Srp19	Signal recognition	1638272_at	0.5565	0.0528	0.8096	0.0632	0.5930	0.0081	-0.0028	0.9960	0.2298	0.1733	0.2326	0.1235	0.2409	0.8202	0.4871	0.2507	0.2462	0.6030
CG10000	CG10000	1638273_at	0.1721	0.5679	-0.0252	0.8929	0.1124	0.4421	0.1924	0.5357	0.2484	0.1237	0.0560	0.7508	0.0100	0.9939	0.0137	0.9796	0.0037	0.9945
pix	pixie	1638274_s_at	0.2564	0.1273	0.2659	0.2937	0.4951	0.0071	0.2063	0.5258	0.0160	0.9504	-0.1903	0.2118	0.0234	0.9829	0.0624	0.8550	0.0390	0.9095
CG11699	CG11699	1638275_at	-0.4389	0.0615	0.0517	0.8266	0.1544	0.4399	0.0658	0.9098	-0.6146	0.0059	-0.6805	0.0022	0.1004	0.9400	-0.0229	0.9738	-0.1233	0.8044
Obp56e	Odorant-binding p	1638276_at	0.1857	0.5506	0.0931	0.3790	0.2519	0.1586	0.0332	0.9587	-0.1115	0.5377	-0.1446	0.3509	-0.0887	0.8609	0.0197	0.9523	0.1084	0.6237
CG11550	CG11550	1638277_at	-1.9482	0.0054	-2.8040	0.0034	-1.5139	0.0029	1.0043	0.3166	0.6915	0.2371	-0.3128	0.5931	-0.1171	0.8692	-0.1007	0.7707	0.0164	0.9701
Myo61F	myosin61F	1638278_s_at	0.0555	0.9454	-2.0773	0.0386	-1.1003	0.0025	0.8708	0.3452	1.5204	0.0115	0.6495	0.1707	-0.1366	0.9677	-0.6411	0.4772	-0.5045	0.5959
CG13840	CG13840	1638279_at	-0.0084	0.9750	0.1810	0.3966	0.3923	0.0881	-0.0344	0.9639	-0.1945	0.3399	-0.1601	0.3900	-0.0656	0.9545	0.1372	0.7229	0.2028	0.5702
---	---	1638280_at	-1.2275	0.0055	-1.1685	0.0114	-1.5147	0.0006	-0.1638	0.8386	-0.2623	0.3969	-0.0984	0.7669	0.1684	0.8270	-0.1082	0.7886	-0.2765	0.4004
CG7448	CG7448	1638281_at	0.1407	0.3869	-0.1466	0.3634	-0.0291	0.9053	0.0358	0.9745	0.1595	0.5926	0.1237	0.6628	-0.1046	0.9036	-0.1743	0.6028	-0.0697	0.8672
CG14196	CG14196	1638282_at	-1.0003	0.0279	-0.7173	0.1523	-0.6675	0.1372	-0.1721	0.8908	-1.3926	0.0060	-1.2205	0.0064	-0.1644	0.9342	-1.0780	0.1012	-0.9136	0.1796
CG9799	CG9799	1638283_at	0.1082	0.7567	0.5130	0.0775	1.1149	0.0032	0.2687	0.6656	-0.1414	0.6757	-0.4100	0.1206	-0.1605	0.9016	0.4171	0.3725	0.5776	0.2403
CG3199 /// DyakCG3199	CG3199	1638284_at	-0.0449	0.7951	-0.0148	0.9010	0.2969	0.0904	0.1763	0.7081	-0.0024	0.9939	-0.1788	0.3676	-0.1141	0.8145	-0.0989	0.6512	0.0151	0.9588
Gyk	Glycerol kinase	1638285_a_at	0.3289	0.1472	0.7872	0.0142	1.3477	0.0004	-0.2617	0.5317	-1.0746	0.0011	-0.8129	0.0021	-0.8031	0.1902	-0.5690	0.0795	0.2341	0.4648
pie	maternal effect ea	1638286_at	-0.0125	0.9735	-0.2674	0.0658	-0.2746	0.2400	-0.3016	0.5280	0.5627	0.0316	0.8643	0.0029	-0.1934	0.8331	0.2399	0.5496	0.4333	0.2750
CG5756	CG5756	1638287_a_at	0.0778	0.7490	-0.0293	0.8001	0.0113	0.9552	0.0935	0.8485	0.1513	0.4114	0.0578	0.7696	0.0556	0.9523	-0.0544	0.8963	-0.1100	0.7307
CG32184	CG32184	1638288_at	0.0890	0.6557	0.1085	0.4063	0.0302	0.8754	-0.2204	0.5220	-0.0208	0.9352	0.1996	0.2171	-0.0372	0.9635	-0.0846	0.7454	-0.0474	0.8737

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1638289_at	0.1691	0.4475	-0.0379	0.8240	-0.0608	0.7050	-0.0748	0.8687	-0.0361	0.8639	0.0388	0.8309	-0.0485	0.9515	0.0280	0.9425	0.0765	0.7909
CG32528	CG32528	1638290_at	-1.0187	0.0052	-0.1930	0.2516	-0.2380	0.1500	-0.1402	0.7270	-0.5903	0.0062	-0.4502	0.0128	-0.1020	0.9168	0.2598	0.4424	0.3618	0.2982
---	---	1638291_at	-0.0449	0.8161	-0.0689	0.7051	-0.1917	0.2917	0.0033	0.9956	0.0897	0.6839	0.0864	0.6662	0.0822	0.8461	0.0113	0.9687	-0.0709	0.7189
Rrp46	Rrp46	1638292_at	0.4587	0.3209	0.2170	0.6127	0.5365	0.0132	0.2676	0.6371	0.4328	0.1183	0.1653	0.5447	-0.2127	0.8973	0.1461	0.8658	0.3588	0.5841
---	---	1638293_at	0.1779	0.3555	0.1276	0.5642	0.2260	0.3264	0.1700	0.6471	-0.0524	0.8137	-0.2224	0.1655	0.1301	0.8461	-0.0364	0.9347	-0.1665	0.5702
mus301	spindle-C	1638294_at	0.3505	0.0910	0.0717	0.4914	0.0715	0.8127	-0.2840	0.6113	0.4081	0.1435	0.6921	0.0135	-0.2066	0.8235	0.3213	0.4003	0.5279	0.1964
CG1129	CG1129	1638295_s_at	1.8967	0.0006	0.9259	0.0035	1.2026	0.0002	0.2766	0.3855	1.4624	0.0002	1.1859	0.0002	-0.1757	0.8097	0.5276	0.0912	0.7032	0.0559
CG32064 /// DsimCG32064	CG32064	1638296_at	-0.1329	0.4417	0.0337	0.7911	0.0417	0.8221	0.0310	0.9633	-0.0803	0.6959	-0.1113	0.5208	-0.0164	0.9894	0.0650	0.8546	0.0814	0.7933
CG12056	CG12056	1638297_at	-1.0746	0.0130	-0.5425	0.1086	-0.2951	0.1899	-0.0135	0.9865	-0.6058	0.0077	-0.5923	0.0051	-0.3101	0.7899	-0.1309	0.8461	0.1792	0.7541
obst-J	CG7348	1638298_at	-0.1205	0.4711	0.0863	0.4901	-0.0467	0.7718	-0.1958	0.4694	-0.2006	0.1669	-0.0049	0.9788	0.0690	0.9396	0.0172	0.9703	-0.0518	0.8924
CG7323	CG7323	1638299_at	-0.8473	0.3155	-0.2996	0.4726	-0.5097	0.1323	0.1755	0.8439	-0.0077	0.9879	-0.1833	0.5764	0.4202	0.8940	0.4675	0.7339	0.0472	0.9810
---	---	1638300_at	0.1340	0.5533	-0.0790	0.5249	-0.0693	0.7439	-0.0184	0.9838	0.2562	0.2239	0.2746	0.1444	0.0953	0.8828	0.1251	0.6408	0.0298	0.9316
CG8157	CG8157	1638301_at	0.1645	0.5777	-0.5702	0.1304	-0.2840	0.1148	-0.1581	0.7590	0.3564	0.1063	0.5145	0.0174	-0.2959	0.7324	-0.3199	0.3802	-0.0240	0.9658
CG1090	CG1090	1638302_a_at	-0.2561	0.1139	-0.0023	0.9860	-0.0935	0.6146	0.0774	0.8817	-0.1627	0.3695	-0.2401	0.1286	0.0570	0.9404	0.1338	0.5897	0.0768	0.7816
CG33160	CG33160	1638303_at	0.6233	0.0236	0.6775	0.0217	0.6625	0.0102	0.0238	0.9838	-0.3083	0.2519	-0.3321	0.1654	0.0321	0.9816	-0.1822	0.5715	-0.2143	0.4984
---	---	1638304_at	0.0930	0.6601	-0.1147	0.4402	0.2857	0.1784	0.2436	0.5038	0.1504	0.4540	-0.0932	0.6381	-0.3333	0.5754	-0.1929	0.4729	0.1404	0.6222
Mip	drostatin-B2	1638305_at	0.1302	0.5200	0.1545	0.3590	0.1362	0.4734	0.1108	0.8143	-0.0837	0.6954	-0.1945	0.2438	0.1712	0.8270	0.0558	0.9111	-0.1154	0.7578
CG5038	CG5038	1638306_at	-0.8058	0.0590	-0.4631	0.0485	-0.7411	0.0094	-0.2482	0.5350	-0.2449	0.2408	0.0033	0.9901	0.0747	0.9589	0.1342	0.8018	0.0596	0.9178
---	---	1638307_s_at	0.3819	0.0415	0.2743	0.0801	0.2616	0.0923	-0.1241	0.7604	-0.0227	0.9257	0.1014	0.5563	-0.1395	0.7644	-0.1033	0.6252	0.0362	0.8926
His3:CG31613 /// His3:CG3	His3:CG31613 ///	1638308_s_at	-0.5028	0.5137	-3.1232	0.0415	-1.7108	0.0095	0.2249	0.8671	1.7336	0.0039	1.5087	0.0042	-1.0140	0.7595	-0.7430	0.6177	0.2710	0.8874
CG16733	CG16733	1638309_at	-0.1141	0.4662	-0.1176	0.3911	-0.3359	0.1454	-0.0756	0.9218	0.3143	0.1723	0.3899	0.0630	-0.0092	0.9939	0.2303	0.3778	0.2395	0.3807
Hlc	RNA helicase gen	1638310_at	-0.6695	0.0161	-0.3870	0.1937	-0.0549	0.7717	0.1449	0.6580	-0.6996	0.0017	-0.8445	0.0005	-0.3430	0.5765	-0.4616	0.1050	-0.1186	0.7045
CG12817	CG12817	1638311_at	0.0631	0.7245	0.1201	0.5757	0.2383	0.3036	-0.0065	0.9956	-0.0165	0.9682	-0.0100	0.9761	-0.0455	0.9743	0.0753	0.8831	0.1207	0.7733
CG13189	CG13189	1638312_at	0.3399	0.3321	-0.0343	0.9350	0.1843	0.5096	0.0858	0.8449	0.0951	0.5893	0.0093	0.9639	-0.2686	0.8692	-0.3829	0.5800	-0.1143	0.8985
CG8321	CG8321	1638313_at	-0.8776	0.0047	-0.7500	0.0398	-1.1239	0.0014	-0.1709	0.6587	-0.6443	0.0049	-0.4734	0.0119	0.2016	0.8400	-0.5395	0.1769	-0.7410	0.1080
CG12418	CG12418	1638314_at	1.0101	0.0627	0.3896	0.3011	-0.4651	0.0883	-0.2597	0.4199	0.9272	0.0009	1.1869	0.0002	0.5863	0.7057	0.2227	0.7787	-0.3636	0.6001
CG8493	CG8493	1638315_s_at	-0.6216	0.0238	0.1423	0.4562	-0.3480	0.0596	-0.4837	0.2704	-0.8587	0.0060	-0.3751	0.1049	0.1837	0.6780	0.0421	0.8725	-0.1417	0.4567
---	---	1638316_at	-0.0883	0.5439	-0.0121	0.9540	0.1896	0.3455	0.1439	0.6869	-0.0181	0.9397	-0.1621	0.2794	0.0499	0.9467	0.1271	0.6028	0.0772	0.7749
CG8097 /// DyakCG8097	CG8097	1638317_at	0.2377	0.2988	-0.4275	0.1383	-0.1227	0.6578	-0.0360	0.9749	0.8336	0.0098	0.8696	0.0049	-0.3925	0.6824	0.0676	0.9111	0.4601	0.2597
CG7544	CG7544	1638318_at	0.5089	0.1243	-0.5399	0.1116	-0.3231	0.0934	-0.0009	0.9994	0.6591	0.0256	0.6600	0.0160	-0.2322	0.7418	-0.2298	0.4388	0.0024	0.9965
inaD	inactivation no aft	1638319_at	0.0330	0.8727	0.1442	0.2884	0.2535	0.0908	0.1145	0.7556	0.0069	0.9778	-0.1076	0.4770	-0.0537	0.9467	0.0436	0.9075	0.0973	0.7280
---	---	1638320_at	0.2298	0.3098	-0.0132	0.9078	-0.0915	0.7185	0.0452	0.9540	0.2272	0.2958	0.1820	0.3579	0.1909	0.7644	0.0782	0.8279	-0.1127	0.7138
CG4847	CG4847	1638321_s_at	2.2986	0.0017	1.4080	0.1248	2.4238	0.0002	0.1883	0.8796	0.0942	0.8670	-0.0941	0.8488	-0.7222	0.7092	-0.7802	0.3236	-0.0580	0.9612
CG10168	CG10168	1638322_at	-0.0467	0.7855	0.0898	0.5419	-0.0790	0.6404	-0.0334	0.9613	-0.0185	0.9459	0.0149	0.9492	0.0746	0.9137	0.1300	0.6138	0.0554	0.8604
CG8116	CG8116	1638323_at	0.3056	0.2373	1.1024	0.0144	0.8336	0.0014	-0.1583	0.7149	-0.3539	0.0760	-0.1956	0.2737	0.2669	0.7780	0.5457	0.1720	0.2788	0.5167
CG6921 /// DyakCG6921	CG6921	1638324_s_at	1.5238	0.0072	1.5285	0.0181	2.3150	0.0002	0.4988	0.3631	0.0594	0.8884	-0.4394	0.1146	-0.3323	0.8558	0.1026	0.9320	0.4349	0.5862
Hn	Henna	1638325_at	2.8118	0.0016	1.1472	0.2443	2.5594	0.0001	0.6899	0.4586	0.1204	0.8540	-0.5696	0.2065	-0.6954	0.7644	-1.4441	0.1395	-0.7487	0.4647
CG7685	CG7685	1638326_at	0.4947	0.0424	0.2605	0.1538	0.1211	0.4624	0.2283	0.5461	0.6643	0.0054	0.4360	0.0211	0.2470	0.6955	0.3538	0.1718	0.1068	0.7253
CG17754	CG17754	1638327_a_at	-0.0123	0.9857	1.3735	0.0578	0.9125	0.0127	-0.1775	0.5735	-0.9262	0.0006	-0.7487	0.0008	0.1962	0.9506	0.2351	0.8552	0.0390	0.9803
---	---	1638328_at	0.1716	0.2834	0.0595	0.5884	0.2663	0.0966	-0.0028	0.9956	0.0364	0.8611	0.0392	0.8284	-0.1172	0.7997	-0.0514	0.8439	0.0658	0.7717
png	pan gu	1638329_at	0.4890	0.3005	-0.0357	0.9099	0.0169	0.9626	-0.2594	0.5249	0.3108	0.1440	0.5702	0.0096	-0.3082	0.8692	-0.0633	0.9590	0.2449	0.7815
Dot	Dorothy	1638330_at	0.0915	0.7265	-1.6426	0.0108	-1.2020	0.0074	-0.1023	0.9373	1.2549	0.0059	1.3572	0.0025	-0.4968	0.6272	-0.4621	0.2634	0.0346	0.9548
CG31054 /// CG4849	CG31054 /// CG4	1638331_s_at	0.2118	0.3819	-0.0024	0.9884	0.3057	0.1235	0.0503	0.9445	-0.0582	0.8210	-0.1084	0.6002	-0.0906	0.8608	-0.1148	0.6041	-0.0243	0.9340
CG31926	CG31926	1638332_at	0.0155	0.9426	0.3134	0.2084	0.0582	0.8363	0.0203	0.9866	-0.0093	0.9827	-0.0297	0.9354	0.2378	0.7506	0.3732	0.2269	0.1354	0.7067
pip	pipe	1638333_at	0.0858	0.6187	0.1004	0.6506	0.1864	0.1914	-0.0632	0.8819	0.0346	0.8546	0.0977	0.4806	-0.0986	0.8657	-0.0737	0.8053	0.0249	0.9394
CG13168	CG13168	1638334_at	-0.1185	0.4755	0.0837	0.7637	0.1752	0.2536	-0.0508	0.9228	-0.2199	0.1629	-0.1691	0.2333	-0.1263	0.8875	0.0193	0.9729	0.1456	0.6969
CG18493	CG18493	1638335_at	-0.0244	0.9934	-0.2197	0.3353	-0.0309	0.8998	0.1068	0.9857	-1.0728	0.4480	-1.1796	0.3447	-0.1712	0.9848	-1.3321	0.5199	-1.1608	0.5862
CG8422	CG8422	1638336_at	-0.1257	0.6080	-0.0074	0.9810	0.0235	0.9176	0.1214	0.7929	0.0236	0.9289	-0.0977	0.6072	0.0993	0.8940	0.1168	0.7090	0.0175	0.9661
---	---	1638337_at	0.0483	0.7549	0.0042	0.9762	0.1194	0.4736	0.0846	0.8350	0.2088	0.1723	0.1242	0.3834	-0.0495	0.9421	0.0226	0.9473	0.0722	0.7695
Fbp1	Protein-1	1638338_a_at	0.3426	0.4277	0.2558	0.4037	0.1944	0.3340	-0.1519	0.8764	-0.4811	0.1507	-0.3292	0.2786	0.2799	0.8326	-0.0833	0.9230	-0.3632	0.5314
mip40	Myb-interacting pr	1638339_at	-0.3880	0.0964	0.2143	0.3036	0.1189	0.4404	-0.1299	0.7330	-0.7110	0.0022	-0.5811	0.0031	-0.0790	0.9148	-0.0047	0.9935	0.0743	0.8178
---	---	1638340_at	0.0422	0.8042	0.1459	0.4819	0.0631	0.7058	-0.1115	0.7791	-0.0969	0.5898	0.0146	0.9438	-0.0229	0.9862	-0.0188	0.9704	0.0041	0.9941
CG18331	CG18331	1638341_s_at	0.4466	0.0669	0.5092	0.1068	0.3211	0.1302	-0.1062	0.8281	0.1565	0.4209	0.2627	0.1172	0.2084	0.8157	0.4064	0.2518	0.1980	0.6186
CG2046 /// DyakCG2046	CG2046	1638342_at	-0.2632	0.2869	0.2616	0.2611	0.7936	0.0045	-0.0406	0.9436	-0.5616	0.0051	-0.5210	0.0042	-0.4956	0.5461	0.0274	0.9650	0.5230	0.1972
CG1143	CG1143	1638343_at	1.4319	0.3993	-0.0794	0.7194	0.2364	0.2743	0.7038	0.2864	0.0809	0.8752	-0.6229	0.0742	0.1232	0.9914	-2.1896	0.3710	-2.3128	0.3670
CG5287	CG5287	1638344_at	1.0345	0.0021	1.1995	0.0325	1.8270	0.0000	0.0836	0.8815	0.2243	0.2408	0.1407	0.4331	-0.5374	0.3712	0.4776	0.1345	1.0150	0.0320
fog	folded gastrulatior	1638345_at	-0.2506	0.2649	0.0102	0.9813	-0.7818	0.0014	-0.5574	0.1787	-0.3378	0.1765	0.2196	0.3400	0.2169	0.8097	-0.0728	0.8963	-0.2897	0.4533
---	---	1638346_at	-0.0380	0.9056	-0.0659	0.6748	-0.5027	0.1751	-0.0583	0.9380	0.1190	0.6251	0.1773	0.3826	0.4236	0.7062	0.0607	0.9353	-0.3629	0.4533
---	---	1638347_at	0.2003	0.2452	0.3366	0.1940	0.2633	0.3677	-0.1123	0.8942	-0.2195	0.4523	-0.1072	0.7200	0.1826	0.8049	0.0478	0.9213	-0.1348	0.6963
CG31658	CG31658	1638348_at	0.1986	0.3841	0.0812	0.4579	-0.1760	0.4322	-0.2146	0.4850	-0.0772	0.6797	0.1374	0.3631	0.0773	0.9317	-0.1305	0.6897	-0.2079	0.4912
eIF-5A	anon-fast-evolving	1638349_s_at	0.0604	0.7178	0.4094	0.0179	0.4589	0.0092	0.0354	0.9518	-0.1565	0.3452	-0.1920	0.1867	-0.0381	0.9550	0.0618	0.8081	0.0999	0.6389
CG6712	CG6712	1638350_at	0.5434	0.1890	-0.0941	0.8791	0.1460	0.6180	0.2542	0.8140	0.9463	0.0334	0.6921	0.0714	0.0425	0.9893	0.3580	0.6156	0.3154	0.6626
Ef1alpha48D	elongation factor	1638351_s_at	0.0943	0.6056	0.0781	0.5499	-0.0045	0.9815	-0.0211	0.9733	-0.0242	0.9113	-0.0031	0.9874	0.0954	0.8594	0.0280	0.9354	-0.0674	0.7976
CG16896	CG16896	1638352_at	-0.5289	0.0593	-0.3701	0.2385	-0.5183	0.0101	0.0001	0.9999	0.1904	0.3402	0.1903	0.2834	-0.0393	0.9816	0.4002	0.2790	0.4395	0.2666
---	---	1638353_at	0.2522	0.1489	0.0596	0.5661	0.3038	0.1002	-0.0314	0.9622	-0.1174	0.5319	-0.0861	0.6322	-0.0408	0.9515	-0.0317	0.9193	0.0091	0.9774
CG12007	CG12007	1638354_at	0.2319	0.1314	-0.0497	0.6370	-0.0108	0.9562	0.0535	0.9118	0.5555	0.0040	0.5020	0.0038	-0.0092	0.9928	0.1551	0.5434	0.1643	0.5214
---	---	1638355_at	-0.0868	0.6214	-0.0829	0.5089	-0.0822	0.6361	0.0579	0.9361	0.0379	0.8957	-0.0201	0.9393	-0.0486	0.9330	-0.0564	0.8144	-0.0078	0.9794
---	---	1638356_at	0.1126	0.4616	-0.0782	0.5894	0.1150	0.6017	0.1512	0.6663	0.0606	0.7625	-0.0906	0.5879	0.0550	0.9611	0.0900	0.8281	0.0350	0.9365
---	---	1638357_at	0.1771	0.3194	0.1477	0.3428	0.0936	0.6358	0.0255	0.9666	-0.1293	0.4324	-0.1547	0.2810	0.0881	0.8692	0.0434	0.8879	-0.0447	0.8769

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Bx	rhombotin	1638358_s_at	0.1020	0.5027	0.0639	0.6700	0.0196	0.9348	0.0033	0.9956	0.0046	0.9846	0.0013	0.9952	0.0709	0.9095	0.0854	0.7439	0.0144	0.9655
CG14478	CG14478	1638359_s_at	-1.1042	0.0065	-1.7711	0.0067	-1.8617	0.0001	0.0384	0.9624	0.5399	0.0208	0.5015	0.0176	0.2269	0.8270	-0.0779	0.9065	-0.3048	0.4975
Oatp26F	Organic anion trar	1638360_at	-0.0451	0.7967	0.1024	0.5175	-0.1743	0.4659	-0.2455	0.5453	-0.1031	0.6662	0.1424	0.4774	-0.0583	0.9588	0.0333	0.9478	0.0916	0.8138
CG12374 /// DyakCG12374	CG12374	1638361_at	0.5328	0.6507	-0.4859	0.2789	-0.0028	0.9959	0.0790	0.9386	0.2219	0.4795	0.1429	0.6429	-0.4135	0.9246	-0.8419	0.5888	-0.4284	0.8149
---	---	1638362_at	0.0577	0.7519	0.0570	0.7356	-0.3269	0.0748	-0.2609	0.5008	-0.1135	0.6172	0.1474	0.4463	0.0159	0.9898	-0.0723	0.8356	-0.0882	0.7745
---	---	1638363_at	0.3013	0.1541	0.1629	0.3678	0.2079	0.1865	-0.1649	0.7469	-0.0970	0.7064	0.0679	0.7861	-0.0524	0.9238	-0.0847	0.6756	-0.0323	0.8977
---	---	1638364_at	0.1744	0.4833	0.0392	0.8718	0.0275	0.9081	-0.0768	0.9065	-0.0886	0.7168	-0.0118	0.9648	0.0247	0.9779	-0.0629	0.8223	-0.0876	0.7170
CG3244	CG3244	1638365_at	-2.0923	0.0114	-2.5847	0.0236	-1.9217	0.0001	0.1865	0.7293	0.3759	0.1173	0.1894	0.3985	-0.1250	0.9689	-0.0427	0.9774	0.0822	0.9440
---	---	1638366_at	-0.6272	0.3627	-1.0108	0.4274	-1.0659	0.0100	-0.3783	0.3836	0.5157	0.0417	0.8939	0.0023	-0.1585	0.9775	-0.0618	0.9797	0.0967	0.9602
CG7686	CG7686	1638367_at	-0.6196	0.0106	-0.2013	0.1962	0.0434	0.8033	0.2236	0.5335	-0.6504	0.0049	-0.8740	0.0008	-0.0205	0.9816	-0.2929	0.1489	-0.2724	0.2094
CG31674	CG31674	1638368_at	0.6055	0.1747	0.6608	0.0455	0.3284	0.0785	-0.3423	0.2842	-0.1357	0.4912	0.2065	0.2198	0.0129	0.9962	-0.1073	0.9058	-0.1202	0.8812
---	---	1638369_at	0.0474	0.8374	0.0476	0.6431	-0.0061	0.9860	0.0200	0.9866	0.0414	0.9198	0.0213	0.9544	0.2274	0.7769	0.1930	0.6028	-0.0344	0.9434
CG12701	CG12701	1638370_s_at	0.4474	0.7990	-1.2229	0.2074	-1.7541	0.0123	-0.7716	0.5363	1.5951	0.0227	2.3667	0.0024	-0.2994	0.9682	-0.5645	0.8299	-0.2650	0.9231
CG31849	CG31849	1638371_at	-1.3861	0.0042	-0.7925	0.0236	-1.6303	0.0004	-0.3438	0.3776	-0.0392	0.8965	0.3047	0.1208	0.0917	0.9534	-0.0406	0.9585	-0.1324	0.8200
---	---	1638372_at	0.0489	0.8153	-0.0326	0.7456	0.2649	0.1709	0.0489	0.9255	-0.0148	0.9505	-0.0637	0.7019	-0.1200	0.8609	-0.0339	0.9402	0.0861	0.7972
CG32161	CG32161	1638373_at	0.1135	0.4716	-0.1200	0.4343	-0.0755	0.7453	0.1655	0.7608	0.1852	0.4466	0.0197	0.9473	-0.1817	0.7823	-0.1604	0.5971	0.0213	0.9589
Obp19b	Odorant-binding p	1638374_at	-0.1240	0.6403	0.2721	0.0668	0.0774	0.6669	0.0139	0.9860	-0.1904	0.3073	-0.2043	0.2165	0.1125	0.8888	-0.0026	0.9979	-0.1151	0.7422
Tim17b2	Translocase inner	1638375_at	-0.0528	0.7560	-0.2451	0.0537	-0.1180	0.4349	0.0398	0.9434	0.1565	0.3292	0.1167	0.4324	-0.1141	0.8400	-0.0764	0.7949	0.0377	0.9067
Osi15	Osiris	1638376_at	0.3730	0.0718	0.0534	0.6107	0.2622	0.1402	-0.2249	0.5793	0.2159	0.3030	0.4408	0.0257	-0.3046	0.6092	-0.2319	0.3551	0.0728	0.8200
CG30025 /// CG30031 /// dr	CG30031 /// CG3(	1638377_x_at	0.3489	0.1103	-0.2392	0.3279	-0.0875	0.7065	0.1542	0.8155	0.4310	0.0972	0.2768	0.2339	-0.0301	0.9831	-0.1892	0.5907	-0.1591	0.6580
CG13116	CG13116	1638378_at	-0.3276	0.2345	-0.5994	0.1308	-0.7909	0.0310	0.2128	0.8164	0.0715	0.8819	-0.1413	0.7116	0.5124	0.5259	0.0936	0.8628	-0.4188	0.2956
---	---	1638379_at	0.2309	0.3669	0.3201	0.0694	0.3357	0.0625	-0.0610	0.8796	-0.1317	0.3481	-0.0707	0.6122	0.0602	0.9545	0.1168	0.7499	0.0566	0.8941
CG41433	CG41433	1638380_at	0.1508	0.4362	0.0369	0.7595	0.2265	0.3768	-0.0762	0.9086	-0.0369	0.8988	0.0393	0.8757	-0.1989	0.7822	-0.0294	0.9531	0.1695	0.6134
CG1695 /// CG32506	CG32506 /// CG1(	1638381_s_at	-0.2333	0.2646	-0.1530	0.3683	0.0017	0.9948	0.1677	0.7121	-0.1553	0.4714	-0.3230	0.0811	0.1454	0.8554	0.0211	0.9689	-0.1243	0.7425
CG30391	CG30391	1638382_at	0.3786	0.0533	0.0168	0.8985	0.1154	0.5694	-0.0908	0.9060	-0.0833	0.7820	0.0076	0.9804	-0.1424	0.8303	-0.1587	0.5906	-0.0163	0.9696
CG8949	CG8949	1638383_at	0.0205	0.9548	0.1211	0.3059	0.1791	0.2645	-0.1053	0.8717	-0.0495	0.8696	0.0557	0.8296	-0.0633	0.9530	0.1035	0.7985	0.1668	0.6291
---	---	1638384_at	0.1336	0.4761	0.0396	0.6976	-0.0081	0.9674	0.0265	0.9705	0.0070	0.9801	-0.0195	0.9308	0.0608	0.9309	0.1042	0.6827	0.0434	0.8904
---	---	1638385_at	0.0442	0.7990	-0.0119	0.9377	0.2673	0.1749	0.0232	0.9757	0.0336	0.8948	0.0104	0.9649	-0.1995	0.7423	-0.1084	0.7090	0.0912	0.7578
HLH3B	Helix loop helix pr	1638386_at	0.2790	0.2818	0.0344	0.9130	0.1577	0.4960	0.2620	0.5109	0.2709	0.1965	0.0089	0.9730	0.1051	0.9396	0.0582	0.9320	-0.0469	0.9371
CG15107	CG15107	1638387_at	0.0431	0.8885	0.0878	0.7487	0.0833	0.6584	-0.2483	0.5255	0.0536	0.8391	0.3019	0.0991	-0.2703	0.7768	0.1475	0.7676	0.4178	0.3205
CG10559	CG10559	1638388_at	0.3436	0.3522	0.5625	0.1501	0.5533	0.0634	0.0251	0.9759	-0.7579	0.0035	-0.7830	0.0018	0.1475	0.9365	-0.3428	0.5818	-0.4902	0.4112
---	---	1638389_at	-0.0301	0.8530	0.0885	0.5145	0.1983	0.2376	0.1271	0.7760	0.0202	0.9403	-0.1069	0.5631	-0.0843	0.8867	-0.0022	0.9971	0.0821	0.7484
---	---	1638390_at	0.1944	0.4115	0.2034	0.2141	0.2055	0.4337	0.0024	0.9962	0.0064	0.9808	0.0040	0.9856	-0.0942	0.9309	-0.1110	0.8061	-0.0168	0.9762
---	---	1638391_at	-0.1483	0.3343	0.0000	1.0000	0.0807	0.6955	-0.0481	0.9165	0.0363	0.8439	0.0844	0.5461	-0.2343	0.7956	-0.0419	0.9461	0.1924	0.6389
Acp98AB	Accessory gland p	1638392_at	-0.0725	0.7445	0.0643	0.5407	0.0810	0.7813	-0.1264	0.6698	-0.0311	0.8670	0.0953	0.4772	-0.1289	0.8870	0.0644	0.9005	0.1933	0.5969
---	---	1638393_at	-0.6158	0.0814	-0.1085	0.5537	0.0071	0.9783	-0.1575	0.8337	-0.5021	0.0793	-0.3446	0.1745	-0.3068	0.6955	0.0069	0.9925	0.3136	0.3420
CG3499	CG3499	1638394_at	0.1035	0.6676	0.5748	0.0882	1.0229	0.0006	0.2101	0.6615	-0.2818	0.2194	-0.4918	0.0249	-0.1301	0.8940	0.3625	0.3033	0.4925	0.1990
CG31802	CG31802	1638395_at	0.1455	0.3024	0.1093	0.4672	0.0361	0.8543	-0.0815	0.8836	-0.0460	0.8520	0.0356	0.8738	0.0544	0.9365	0.0165	0.9630	-0.0379	0.8982
---	---	1638396_at	0.0449	0.8426	0.0142	0.9487	0.3110	0.0962	0.0877	0.8251	0.0306	0.8813	-0.0571	0.7281	-0.1735	0.8206	0.0534	0.9133	0.2269	0.4900
CG11388	CG11388	1638397_at	-0.2767	0.0795	-0.2585	0.0817	-0.2339	0.3141	0.0524	0.9036	-0.1975	0.1575	-0.2499	0.0511	0.0562	0.9652	-0.1069	0.8109	-0.1631	0.6621
Shal	Shaker cognate 1	1638398_at	-3.3433	0.0014	-3.6209	0.0020	-3.9942	0.0000	-0.7081	0.0189	-0.8206	0.0009	-0.1125	0.4526	-0.1918	0.9101	-0.9522	0.1199	-0.7603	0.2245
Gr64a	Gustatory recepto	1638399_at	0.0188	0.9192	0.1565	0.1605	0.0917	0.6889	0.0103	0.9882	-0.1645	0.3111	-0.1748	0.2251	-0.1925	0.7990	-0.2028	0.5373	-0.0102	0.9846
CG8503	CG8503	1638400_at	-1.8337	0.0024	-1.9277	0.0202	-2.6075	0.0000	-0.6199	0.2311	-0.4069	0.1840	0.2130	0.4660	0.2035	0.8906	-0.3769	0.5135	-0.5804	0.3082
CG13589	CG13589	1638401_at	0.0890	0.7178	0.0325	0.8136	-0.0601	0.7783	-0.0484	0.9435	0.0654	0.7793	0.1138	0.5472	0.0425	0.9527	-0.0707	0.7922	-0.1132	0.6231
CG40195	CG40195	1638402_at	0.1508	0.3895	0.0017	0.9952	0.3609	0.1188	0.1251	0.6854	0.1969	0.1675	0.0718	0.6230	-0.0659	0.9421	0.1215	0.7019	0.1874	0.5252
CG17856	CG17856	1638403_at	-0.1330	0.4414	-0.1647	0.4150	-0.0101	0.9591	0.0743	0.8665	-0.0646	0.7224	-0.1389	0.3296	-0.1364	0.8513	-0.1362	0.6847	0.0002	0.9997
---	---	1638404_at	0.2010	0.4201	0.1445	0.4789	0.0805	0.7425	-0.0133	0.9872	0.0436	0.8680	0.0569	0.7981	-0.0024	0.9990	-0.0825	0.8460	-0.0802	0.8409
CG9338 /// DsimCG9338	CG9338	1638405_at	-1.2401	0.0135	-0.9136	0.0554	-1.5408	0.0017	-0.1112	0.9196	-0.8013	0.0255	-0.6901	0.0298	0.4379	0.7812	-0.3750	0.6075	-0.8129	0.2523
CG32606	CG32606	1638406_at	0.4189	0.1619	0.1676	0.3389	0.1711	0.3018	0.1927	0.6006	0.0585	0.7966	-0.1341	0.4415	0.0108	0.9924	-0.0278	0.9485	-0.0387	0.9161
CG31244	CG31244	1638407_at	0.0923	0.7438	0.0005	0.9986	0.1510	0.6231	0.0283	0.9777	0.0525	0.8787	0.0242	0.9397	-0.1782	0.7707	-0.1121	0.7048	0.0661	0.8434
CG11297	CG11297	1638408_at	-0.0183	0.9218	0.0489	0.6464	0.0115	0.9625	-0.1123	0.8350	-0.0452	0.8667	0.0671	0.7647	-0.1052	0.8521	-0.1041	0.6908	0.0011	0.9979
CG12030	CG12030	1638409_at	1.0312	0.0077	2.0555	0.0057	2.1677	0.0000	0.1898	0.7425	-0.5622	0.0341	-0.7519	0.0060	0.0010	0.9998	0.2922	0.4032	0.2912	0.4172
CG5131	CG5131	1638410_at	0.1461	0.5708	0.2145	0.0827	0.0030	0.9916	-0.1635	0.6823	0.1259	0.5246	0.2894	0.0829	0.0039	0.9982	0.0962	0.8389	0.0923	0.8366
CG6897	CG6897	1638411_at	0.0735	0.8533	0.0079	0.9887	0.0208	0.9575	-0.3773	0.4128	0.1247	0.6676	0.5020	0.0349	-0.3452	0.8543	0.0168	0.9925	0.3620	0.6677
px	plexus	1638412_at	-0.5456	0.2699	-0.5994	0.4953	-0.8313	0.0016	0.0156	0.9857	0.0145	0.9606	-0.0011	0.9965	0.2312	0.9515	-0.0599	0.9762	-0.2912	0.8418
CG18180 /// DyakCG1818(	CG18180	1638413_at	0.4381	0.0606	0.0997	0.6888	0.1875	0.3529	0.1853	0.5863	0.1806	0.3030	-0.0048	0.9826	0.1442	0.8609	0.1422	0.7056	-0.0020	0.9979
Bmcp	Bmcp	1638414_a_at	0.0350	0.9344	-0.0423	0.7328	0.1191	0.5037	-0.2555	0.7039	-0.4156	0.1765	-0.1601	0.6082	-0.3521	0.7230	-0.4252	0.3098	-0.0731	0.9025
FK506-bp2	FK506-binding pr(	1638415_at	0.6295	0.0066	1.1031	0.0112	0.9116	0.0022	0.0700	0.9216	-0.2245	0.2990	-0.2945	0.1224	0.2296	0.6660	0.1584	0.4665	-0.0712	0.7787
cype	cyclope	1638416_at	-0.2084	0.4601	0.1050	0.6776	-0.0005	0.9980	-0.1869	0.6533	-0.6410	0.0071	-0.4541	0.0196	-0.0637	0.9717	-0.3759	0.3935	-0.3122	0.4995
CG31373	CG31373	1638417_at	1.7740	0.0027	0.7038	0.3484	2.0719	0.0004	0.7071	0.2438	0.1099	0.8031	-0.5972	0.0661	-0.6782	0.7423	-1.0170	0.2254	-0.3388	0.7346
FucTD	FucTD	1638418_at	0.1547	0.4804	0.2084	0.1727	0.1349	0.5180	-0.0440	0.9538	-0.1498	0.4902	-0.1058	0.6138	0.0424	0.9589	0.1114	0.6605	0.0690	0.8082
Sgs1	Salivary gland sec	1638419_at	-0.1458	0.4964	0.0769	0.5557	-0.0518	0.7892	-0.1158	0.7891	-0.1468	0.4235	-0.0310	0.8823	-0.0533	0.9611	0.0713	0.8703	0.1246	0.7220
CG12523	CG12523	1638420_at	-0.0942	0.6228	-0.1487	0.3191	0.1147	0.4894	0.4255	0.1680	0.2638	0.1580	-0.1618	0.3476	0.0776	0.8909	0.0539	0.8546	-0.0237	0.9369
CG33124	CG33124	1638421_at	0.0057	0.9816	0.0838	0.7242	0.1631	0.2439	-0.0241	0.9777	0.1002	0.6841	0.1244	0.5595	-0.0147	0.9915	0.0882	0.8379	0.1029	0.7882
---	---	1638422_at	0.0094	0.9674	0.1067	0.6728	0.2816	0.3074	0.0499	0.9345	-0.1108	0.5649	-0.1607	0.3237	0.0590	0.9330	0.0120	0.9758	-0.0471	0.8774
---	---	1638423_at	0.2205	0.2113	0.1385	0.4019	-0.2235	0.1758	0.0047	0.9956	0.1012	0.6839	0.0965	0.6702	0.1064	0.8222	-0.0780	0.7339	-0.1845	0.3621
CG10514	CG10514	1638424_at	-2.3281	0.0014	-0.9416	0.2968	-3.1201	0.0011	-1.0724	0.0943	-0.8854	0.0263	0.1870	0.6250	1.1537	0.6904	0.5032	0.7105	-0.6505	0.6141
CG15643	CG15643	1638425_at	-0.1201	0.5862	0.4118	0.2095	0.4816	0.0593	0.2508	0.6615	0.0600	0.8691	-0.1908	0.4639	0.2965	0.7387	0.5990	0.1173	0.3026	0.4351
CG14035	CG14035	1638426_at	0.1763	0.3068	-0.0882	0.5008	-0.2580	0.2326	-0.1301	0.7850	0.1275	0.5469	0.2576	0.1415	0.0310	0.9816	-0.1922	0.5617	-0.2232	0.4944

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
Su(fu)	suppressor of fus	1638427_at	0.0210	0.9220	0.4132	0.0453	0.4572	0.0344	0.0441	0.9538	-0.1730	0.4204	-0.2171	0.2445	0.0231	0.9884	0.3207	0.3358	0.2976	0.3921
---	---	1638428_at	-0.0129	0.9442	-0.0629	0.8378	-0.4846	0.0519	-0.0024	0.9960	0.2143	0.1517	0.2167	0.1058	0.2362	0.8292	0.2537	0.6041	0.0176	0.9821
CG33475	CG33475	1638429_at	-0.0356	0.8962	-0.0266	0.8819	-0.1424	0.3941	-0.1049	0.7902	0.0165	0.9447	0.1214	0.4206	-0.0327	0.9717	0.0117	0.9778	0.0444	0.8891
---	---	1638430_at	0.0125	0.9510	-0.0267	0.8796	0.0801	0.6599	-0.0389	0.9676	-0.0054	0.9889	0.0335	0.9130	-0.2548	0.5905	-0.0814	0.7413	0.1734	0.4162
CG2911 /// DereCG2911 ///	CG2911	1638431_at	0.2543	0.3303	-0.2007	0.2451	-0.3479	0.0881	0.1643	0.7104	0.7080	0.0046	0.5437	0.0091	0.2012	0.7697	0.2855	0.3160	0.0844	0.8177
CG10082	CG10082	1638432_a_at	-0.2055	0.7761	-0.8936	0.0815	-1.0303	0.0108	0.4999	0.6338	1.2677	0.0218	0.7678	0.0944	0.6732	0.7215	0.6005	0.4514	-0.0727	0.9482
---	---	1638433_at	0.2366	0.2534	-0.1518	0.4149	-0.0493	0.8228	0.1298	0.8384	0.1822	0.4642	0.0524	0.8495	0.1716	0.7997	-0.0653	0.8695	-0.2368	0.4078
---	---	1638434_at	0.0756	0.6782	0.2129	0.1429	0.1818	0.3248	-0.0594	0.9026	-0.0033	0.9892	0.0561	0.7403	-0.0739	0.9177	0.0947	0.7454	0.1686	0.5144
---	---	1638435_at	0.2060	0.2235	-0.0074	0.9515	0.1302	0.4612	0.0288	0.9558	0.0301	0.8718	0.0013	0.9943	0.0123	0.9913	-0.0713	0.8153	-0.0836	0.7578
hug	pyrokinin	1638436_at	-0.0779	0.6081	-0.0300	0.9046	0.0138	0.9431	-0.0264	0.9755	-0.0265	0.9289	-0.0001	0.9995	0.0141	0.9893	0.0318	0.9275	0.0177	0.9550
CR14638	CR14638	1638437_at	3.2572	0.0005	2.5038	0.0039	2.6218	0.0001	-0.0456	0.9401	-0.0141	0.9567	0.0315	0.8795	-0.6087	0.6557	-1.1739	0.0567	-0.5652	0.3205
Ptp61F	protein tyrosine pt	1638438_a_at	1.2623	0.0049	0.6187	0.0975	0.4444	0.2248	0.0189	0.9814	0.1236	0.5537	0.1047	0.5888	-0.2541	0.8878	-0.3118	0.6774	-0.0577	0.9516
RpS3A	ribosomal protein	1638439_a_at	0.1199	0.3791	0.0488	0.6618	0.0939	0.5096	-0.0028	0.9956	0.0384	0.8228	0.0411	0.7880	-0.0090	0.9913	-0.0293	0.9095	-0.0203	0.9319
Ca-beta	calcium channel b	1638440_at	0.1802	0.3665	0.0385	0.7133	0.3644	0.0710	0.2240	0.4902	0.0982	0.6051	-0.1259	0.4391	-0.0711	0.8904	0.0770	0.7340	0.1481	0.4616
CG18519	CG18519	1638441_a_at	-1.1073	0.0359	-1.3875	0.0145	-1.0590	0.0035	0.6979	0.3425	0.8954	0.0407	0.1975	0.6517	-0.1411	0.8461	-0.1499	0.6450	-0.0088	0.9852
CG4655	CG4655	1638442_at	-0.3188	0.1037	-0.6540	0.0412	-0.9989	0.0112	-0.1220	0.9029	0.6320	0.0557	0.7540	0.0174	-0.0971	0.9152	-0.0568	0.9075	0.0403	0.9273
CG3062	CG3062	1638443_at	-0.1015	0.6775	0.2049	0.2562	0.2472	0.2358	0.0520	0.9048	-0.2223	0.1145	-0.2743	0.0368	-0.0243	0.9862	0.0768	0.8678	0.1011	0.7980
---	---	1638444_s_at	0.1644	0.3876	-0.1444	0.4414	-0.2181	0.2510	-0.0220	0.9778	0.1760	0.4002	0.1980	0.2817	0.2687	0.7215	-0.0286	0.9558	-0.2973	0.3629
SPoCk	Secretory Pathwa	1638445_a_at	0.7502	0.0899	0.4872	0.0760	0.4146	0.1489	0.0663	0.9653	0.4514	0.2694	0.3851	0.2962	0.1933	0.8882	0.1817	0.7741	-0.0115	0.9890
CG3542	CG3542	1638446_a_at	0.0400	0.9090	0.3651	0.2412	0.6731	0.0456	0.0509	0.9376	0.1954	0.3077	0.1445	0.4158	-0.2232	0.8689	0.4759	0.3651	0.6991	0.2100
CG11141	CG11141	1638447_s_at	-0.7505	0.0446	-0.3129	0.4660	-0.1011	0.7604	0.3065	0.6321	0.1676	0.6310	-0.1389	0.6697	0.0713	0.9816	0.5708	0.3823	0.4995	0.4683
CG13211	CG13211	1638448_at	0.2395	0.8661	0.4439	0.0367	0.5681	0.0505	-0.1776	0.9422	-0.6983	0.3236	-0.5207	0.4260	-0.1047	0.9848	-0.2543	0.8845	-0.1496	0.9293
CG13953	CG13953	1638449_at	0.0274	0.8892	-0.0051	0.9686	-0.1565	0.5429	0.0524	0.9626	0.0949	0.7940	0.0426	0.9053	0.2496	0.6477	0.0165	0.9634	-0.2331	0.2993
---	---	1638450_at	-0.0129	0.9535	-0.0588	0.5754	0.0407	0.8403	-0.0474	0.9305	0.0429	0.8324	0.0903	0.5666	-0.0812	0.9246	-0.0644	0.8760	0.0168	0.9696
---	---	1638451_at	0.3259	0.0793	0.3444	0.0925	0.3075	0.1205	0.2459	0.4509	0.1210	0.5201	-0.1248	0.4561	0.0674	0.9238	0.0483	0.8867	-0.0191	0.9541
per	period clock prote	1638452_at	-0.1657	0.8335	-0.3258	0.3583	-0.3890	0.1624	-0.0820	0.9412	-0.1076	0.7798	-0.0256	0.9494	0.2146	0.9457	-0.3999	0.7101	-0.6145	0.5398
CG9849 /// DyakCG9849	CG9849	1638453_at	1.1023	0.0043	0.9680	0.0220	1.1084	0.0002	0.1454	0.7556	0.5184	0.0184	0.3730	0.0444	0.0174	0.9916	0.4031	0.2814	0.3856	0.3293
---	---	1638454_at	-0.3320	0.3181	-0.3055	0.1345	-0.8124	0.0100	-0.0850	0.9311	0.6160	0.0391	0.7010	0.0146	0.1045	0.9405	0.1753	0.7225	0.0708	0.9056
---	---	1638455_at	0.2219	0.2185	0.2367	0.0597	0.1153	0.4643	0.1195	0.7927	0.1944	0.2968	0.0750	0.6999	0.1948	0.7726	0.0713	0.8595	-0.1236	0.7045
CG8531	CG8531	1638456_at	-0.1205	0.4049	0.1572	0.1716	0.3536	0.0365	0.0519	0.9186	-0.2365	0.1292	-0.2884	0.0449	-0.1425	0.7707	0.0919	0.6957	0.2344	0.2781
CG18347 /// DyakCG18347	CG18347	1638457_at	-0.8989	0.3752	-0.4965	0.1721	-0.8229	0.0008	-0.7273	0.0840	-1.2548	0.0009	-0.5275	0.0239	-0.4143	0.9137	-0.8037	0.5654	-0.3894	0.8149
---	---	1638458_at	0.1159	0.6849	0.3363	0.1388	0.5188	0.0075	-0.0027	0.9962	-0.2023	0.2960	-0.1996	0.2463	0.0227	0.9816	0.0109	0.9795	-0.0119	0.9739
---	---	1638459_at	-0.0063	0.9861	-0.0601	0.6629	0.1211	0.4320	0.0894	0.8816	-0.0298	0.9172	-0.1193	0.5539	0.0038	0.9982	-0.1253	0.7646	-0.1292	0.7465
CG18809	CG18809	1638460_s_at	-0.2349	0.5266	-0.0337	0.7421	-0.0091	0.9795	0.0479	0.9629	0.0735	0.8341	0.0256	0.9400	0.0421	0.9800	0.1631	0.7003	0.1210	0.7865
CG7271 /// term	CG7271 /// termin	1638461_s_at	0.0555	0.8344	0.4715	0.0981	0.2871	0.1811	-0.0983	0.8578	-0.4318	0.0356	-0.3335	0.0623	0.1216	0.8795	0.1354	0.6955	0.0138	0.9776
CG18088	CG18088	1638462_at	4.1721	0.0004	2.6047	0.0052	5.5746	0.0001	3.0091	0.0001	2.6411	0.0001	-0.3679	0.0900	0.9755	0.5421	1.7814	0.0391	0.8059	0.2960
CG11986	CG11986	1638463_at	-1.3601	0.0032	-0.3863	0.3503	-0.7267	0.0130	-0.5470	0.2592	-0.8810	0.0085	-0.3339	0.1925	-0.1168	0.9291	0.1413	0.7897	0.2580	0.5649
qtc	quick-to-court	1638464_a_at	2.4761	0.0016	3.3598	0.0044	4.0245	0.0000	0.3795	0.5008	-0.3361	0.2606	-0.7156	0.0152	-0.3271	0.8689	0.4104	0.6266	0.7376	0.3594
CG4848	CG4848	1638465_at	0.1465	0.5121	0.7330	0.1746	1.0482	0.0002	0.1395	0.8154	-0.2203	0.3565	-0.3598	0.0864	-0.1395	0.8909	0.4754	0.1998	0.6150	0.1389
Irk2	Inwardly rectifying	1638466_a_at	0.2870	0.8821	0.0828	0.7286	-1.1101	0.0171	-0.8736	0.3955	-2.5912	0.0016	-1.7177	0.0059	0.2467	0.9792	-2.7975	0.1838	-3.0442	0.1865
---	---	1638467_at	-0.1658	0.3255	-0.0969	0.4243	-0.0868	0.7304	0.0484	0.9300	0.0976	0.5767	0.0491	0.7855	-0.1381	0.8439	0.0308	0.9476	0.1689	0.5814
CG30051	CG30051	1638468_at	0.2093	0.4169	-0.0037	0.9960	0.2693	0.1357	-0.0028	0.9973	-0.1587	0.5064	-0.1560	0.4675	-0.0843	0.9611	-0.3132	0.5300	-0.2288	0.6607
CG3857	CG3857	1638469_s_at	-0.1315	0.6476	-0.9133	0.0257	-0.8918	0.0008	-0.3147	0.2990	0.1760	0.3302	0.4908	0.0083	-0.4487	0.7070	-0.4800	0.3259	-0.0313	0.9665
---	---	1638470_at	0.2447	0.3428	-0.0290	0.9067	0.2310	0.1966	0.2598	0.5350	0.2291	0.2985	-0.0307	0.9078	0.0832	0.9095	-0.0272	0.9494	-0.1104	0.7072
---	---	1638471_at	0.1802	0.3928	-0.1385	0.4540	-0.1520	0.4475	0.0723	0.9293	0.2115	0.3863	0.1392	0.5527	-0.0272	0.9717	-0.0644	0.7883	-0.0372	0.8880
Cpr62Bb	CG13935	1638472_at	0.9952	0.2453	1.9431	0.0269	1.6272	0.0008	-0.4850	0.5948	-1.9269	0.0022	-1.4419	0.0047	-0.1039	0.9829	-0.8014	0.4794	-0.6975	0.5520
CG4386	CG4386	1638473_at	-0.0847	0.6782	-0.0044	0.9836	0.0974	0.6455	-0.0487	0.9346	-0.1296	0.4801	-0.0809	0.6558	-0.0292	0.9676	0.0449	0.8643	0.0741	0.7307
---	---	1638474_at	0.0438	0.8601	0.3043	0.0688	0.2154	0.1402	-0.1076	0.8211	-0.4670	0.0203	-0.3594	0.0373	0.0530	0.9599	-0.0763	0.8530	-0.1293	0.6993
CG16885	CG16885	1638475_a_at	0.2143	0.2959	0.2349	0.2482	0.0131	0.9572	0.0106	0.9900	0.0141	0.9604	0.0035	0.9880	0.0486	0.9555	-0.0239	0.9531	-0.0726	0.8190
mnb	Minibrain	1638476_s_at	-0.0213	0.9753	-0.5777	0.2315	-1.3116	0.0020	-0.5641	0.4962	0.6444	0.1408	1.2086	0.0084	0.1880	0.9174	0.0339	0.9752	-0.1541	0.8488
CG13403	CG13403	1638477_at	-2.5548	0.0203	0.0494	0.6452	-1.5399	0.0097	-1.2121	0.4033	-2.7914	0.0050	-1.5793	0.0360	0.2043	0.9555	-0.4286	0.7273	-0.6328	0.5764
CG1231	CG1231	1638478_at	0.1845	0.5668	-0.0697	0.5993	-0.0696	0.8272	0.0803	0.9500	0.3258	0.3693	0.2455	0.4660	0.0887	0.8940	0.0840	0.7838	-0.0048	0.9913
---	---	1638479_at	-0.0049	0.9861	0.1680	0.4160	0.3595	0.0454	-0.1152	0.8449	-0.0652	0.8083	0.0501	0.8416	-0.2492	0.7215	0.1271	0.7031	0.3763	0.2189
Rlb1	Rlb1	1638480_at	-0.1280	0.5634	0.5090	0.1740	0.7479	0.0107	0.0011	0.9988	-0.0821	0.6514	-0.0832	0.6100	-0.1328	0.9246	0.4748	0.3000	0.6076	0.2136
CG2736	CG2736	1638481_at	3.0391	0.0014	1.4750	0.0768	3.0907	0.0001	0.2976	0.7825	0.8861	0.0518	0.5885	0.1384	-1.2305	0.5228	-0.5523	0.5652	0.6782	0.4746
---	---	1638482_at	0.0884	0.6243	-0.0429	0.7322	0.2038	0.2537	0.1520	0.8056	0.1589	0.5480	0.0069	0.9825	-0.1629	0.7230	-0.0666	0.7897	0.0963	0.6553
CG10920	CG10920	1638483_at	0.1310	0.4099	-0.0406	0.8535	-0.0860	0.6462	0.1161	0.8140	0.0274	0.9200	-0.0887	0.6595	0.0775	0.9128	-0.1396	0.5989	-0.2172	0.3921
Hsp67Bc	Gene 3	1638484_at	-0.0277	0.9535	-2.0629	0.0049	-0.6901	0.0709	1.2998	0.2596	2.1305	0.0080	0.8307	0.1737	-0.1329	0.8940	-0.1355	0.7620	-0.0026	0.9973
SP2637	SP2637	1638485_s_at	-0.1827	0.2145	0.6360	0.0760	0.5631	0.0621	-0.0194	0.9838	-0.5437	0.0216	-0.5243	0.0157	-0.0963	0.9467	0.2199	0.6429	0.3163	0.4851
Rbp1	RNA-binding prote	1638486_at	0.2901	0.4548	-0.4166	0.5741	-0.7194	0.0253	-0.6356	0.3433	-0.2351	0.5731	0.4005	0.2483	-0.2659	0.9142	-0.8371	0.3217	-0.5712	0.5310
CG13510	CG13510	1638487_at	0.3070	0.4023	0.2586	0.2312	0.1452	0.4200	-0.1460	0.7595	0.0175	0.9542	0.1635	0.3900	0.0682	0.9535	0.0390	0.9451	-0.0293	0.9523
---	---	1638488_at	0.0657	0.8054	0.0425	0.8345	0.1870	0.3907	-0.1333	0.8076	-0.1692	0.4562	-0.0359	0.8907	-0.1778	0.8077	-0.1112	0.7612	0.0667	0.8699
---	---	1638489_at	0.6670	0.0631	0.2014	0.3653	0.3205	0.3396	0.1103	0.8663	0.2665	0.2503	0.1561	0.4785	0.0871	0.9589	-0.0973	0.8903	-0.1844	0.7410
CG15365	CG15365	1638490_at	-1.2668	0.0388	0.1926	0.2453	0.0943	0.6335	-0.1276	0.9386	-1.6255	0.0057	-1.4980	0.0049	0.0754	0.9011	-0.0893	0.7284	-0.1647	0.4741
CG15831	CG15831	1638491_at	0.0537	0.7318	-0.0476	0.7078	0.0221	0.8997	0.0178	0.9807	-0.0496	0.8261	-0.0675	0.7245	0.0030	0.9975	-0.0107	0.9773	-0.0136	0.9655
CG13913	CG13913	1638492_at	0.1601	0.4674	-0.1372	0.4825	0.1400	0.6452	0.1663	0.7409	0.2148	0.3408	0.0486	0.8502	-0.2797	0.7726	-0.2316	0.6024	0.0482	0.9344
CG30470	CG30470	1638493_at	0.2982	0.0940	-0.0369	0.7796	0.1581	0.4982	0.1609	0.7937	0.1359	0.6259	-0.0249	0.9362	0.0164	0.9848	-0.1742	0.3728	-0.1906	0.3518
CG3124	CG3124	1638494_at	0.3085	0.1407	0.0470	0.8185	-0.0040	0.9903	0.0526	0.9343	0.0014	0.9960	-0.0512	0.8077	0.0899	0.9289	-0.1270	0.7395	-0.2170	0.5257
Ngp	Ngp	1638495_at	0.0283	0.9392	-0.1083	0.8062	-0.2847	0.1619	-0.1891	0.7796	0.3145	0.2648	0.5036	0.0498	0.0709	0.9689	0.0628	0.9341	-0.0081	0.9924

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
---	---	1638496_at	1.4734	0.0010	0.8788	0.0077	1.3530	0.0017	0.3611	0.5259	0.5125	0.0869	0.1513	0.6170	0.0350	0.9611	0.0457	0.8725	0.0107	0.9726
---	---	1638497_at	-0.0170	0.9407	0.0093	0.9340	-0.0842	0.5875	0.0062	0.9937	-0.1845	0.2589	-0.1907	0.1893	-0.0745	0.9374	-0.1545	0.6312	-0.0800	0.8307
CG1600	CG1600	1638498_s_at	-0.9471	0.0036	-0.9031	0.0930	-1.5092	0.0048	-0.2774	0.3887	-0.3627	0.0510	-0.0853	0.6487	0.3314	0.8767	-0.3760	0.6858	-0.7074	0.4037
Tfb1	Tfb1	1638499_s_at	-0.1139	0.5574	0.2057	0.1578	0.4806	0.0387	0.1369	0.7121	-0.2304	0.1704	-0.3673	0.0232	-0.0752	0.9238	0.1519	0.5800	0.2270	0.3898
DMAP1	DMAP1	1638500_at	0.1187	0.4147	0.2262	0.3955	0.2550	0.1090	0.0890	0.8046	-0.0473	0.7869	-0.1363	0.2950	0.1188	0.8743	0.1577	0.6173	0.0389	0.9234
CG6259	CG6259	1638501_at	-0.2183	0.1547	-0.2973	0.0360	0.0240	0.9382	0.2841	0.2397	0.3381	0.0259	0.0541	0.7224	-0.0020	0.9994	0.3445	0.4094	0.3465	0.4199
---	---	1638502_at	0.1093	0.5045	0.0183	0.8992	-0.1823	0.2854	-0.0980	0.8217	0.1400	0.4253	0.2380	0.1154	0.2085	0.6717	0.0486	0.8602	-0.1599	0.4322
CG10969	CG10969	1638503_at	0.0595	0.7550	0.1885	0.1493	0.0986	0.7068	-0.0339	0.9687	-0.1478	0.5360	-0.1139	0.6143	-0.0795	0.8884	-0.0563	0.8461	0.0232	0.9382
---	---	1638504_s_at	-0.2302	0.3869	-0.0956	0.6251	0.0903	0.6821	0.0063	0.9956	0.3387	0.2614	0.3323	0.2164	-0.1452	0.8833	0.4748	0.1938	0.6199	0.1303
t	tan	1638505_at	-1.9582	0.0044	0.8814	0.0899	-0.2155	0.6466	-0.9292	0.2092	-3.2002	0.0003	-2.2710	0.0006	0.1448	0.9535	-0.1506	0.8879	-0.2954	0.7243
CG9430	CG9430	1638506_at	0.2224	0.2837	0.1354	0.4551	0.2750	0.1406	-0.1378	0.7081	-0.1551	0.3662	-0.0173	0.9341	-0.0858	0.9092	-0.0784	0.8221	0.0073	0.9863
mRpL46	mitochondrial ribo	1638507_at	-0.4081	0.2669	-0.4451	0.1221	-0.5489	0.0048	0.0414	0.9451	0.0031	0.9899	-0.0383	0.8471	0.0661	0.9742	-0.0712	0.9288	-0.1373	0.8287
CG14934	CG14934	1638508_at	0.4283	0.2028	0.3020	0.4284	0.3074	0.1105	-0.3645	0.3405	-0.5325	0.0238	-0.1680	0.4151	-0.0300	0.9862	-0.4158	0.2853	-0.3858	0.3503
CG31110	CG31110	1638509_at	-1.0414	0.0012	-0.9631	0.0356	-1.2312	0.0002	-0.0806	0.8835	0.0975	0.6424	0.1781	0.2935	0.0778	0.8971	0.0901	0.7273	0.0124	0.9715
CG7744 /// DmirCG7744	CG7744	1638510_at	0.7112	0.0158	0.1829	0.6239	0.1247	0.5813	-0.3424	0.2217	0.8309	0.0011	1.1733	0.0002	-0.3441	0.7464	0.2783	0.5571	0.6225	0.1919
Aats-trp	Tryptophanyl-tRN	1638511_at	-0.1678	0.3492	0.4248	0.1106	0.0813	0.7185	0.0706	0.8899	0.3297	0.0579	0.2592	0.0898	0.4293	0.5619	1.1070	0.0204	0.6778	0.0812
Sb	stubbloid	1638512_at	-0.3786	0.0625	-1.2793	0.0149	-1.2551	0.0124	0.1256	0.9223	0.9344	0.0242	0.8088	0.0277	-0.1304	0.8379	0.0950	0.7634	0.2254	0.3981
atl	D-atlastin	1638513_s_at	0.2612	0.1321	0.0987	0.4861	0.1189	0.5215	-0.0689	0.9009	0.2603	0.1441	0.3292	0.0453	-0.0991	0.8407	0.0493	0.8619	0.1484	0.4815
CG40497	CG40497	1638514_a_at	0.1770	0.2845	0.0786	0.5081	0.0532	0.7570	-0.0665	0.9098	-0.0027	0.9919	0.0638	0.7537	-0.0264	0.9716	-0.0205	0.9451	0.0058	0.9847
CG13155 /// DyakCG13155	CG13155	1638515_at	0.3277	0.4535	0.0972	0.3380	0.0367	0.8665	0.0122	0.9893	-0.0872	0.7143	-0.0994	0.6372	-0.0236	0.9898	-0.1959	0.6363	-0.1723	0.6837
CG5708	CG5708	1638516_s_at	-0.0778	0.6832	0.1930	0.1194	0.0214	0.9065	-0.0199	0.9819	-0.0735	0.7701	-0.0536	0.8214	0.2276	0.5954	0.2190	0.2288	-0.0086	0.9769
CG7543	CG7543	1638517_at	0.1540	0.4006	-0.0764	0.5835	-0.1632	0.5572	-0.1936	0.7854	0.1517	0.6451	0.3453	0.1867	-0.1746	0.7768	-0.1658	0.5523	0.0088	0.9846
---	---	1638518_at	0.2105	0.2754	0.2700	0.1601	0.2861	0.0956	-0.0244	0.9727	-0.1214	0.5173	-0.0970	0.5824	0.0679	0.9132	-0.0078	0.9854	-0.0757	0.7660
CG32599	unknown protein	1638519_at	-2.5457	0.0008	-0.7999	0.0141	-1.3478	0.0093	-0.1374	0.9382	-1.3008	0.0208	-1.1634	0.0207	-0.0254	0.9816	-0.0746	0.8053	-0.0493	0.8779
nudE	nudE	1638520_at	0.0642	0.8589	0.7297	0.0354	0.3651	0.2718	-0.1697	0.7927	-0.8377	0.0071	-0.6680	0.0119	0.0418	0.9860	-0.2052	0.7425	-0.2470	0.6706
CG1193	CG1193	1638521_a_at	0.7582	0.0037	0.6807	0.0113	0.9282	0.0019	0.2011	0.6659	0.4821	0.0367	0.2810	0.1558	-0.1707	0.7519	0.5641	0.0391	0.7347	0.0320
CG12123 /// DsimCG12123	CG12123	1638522_at	-0.3718	0.0428	-0.7144	0.0437	-0.6267	0.0047	0.1374	0.7815	0.3538	0.0861	0.2164	0.2400	0.0837	0.9260	0.0725	0.8623	-0.0112	0.9832
CG32633	CG32633	1638523_at	-0.5732	0.8515	0.1529	0.4190	-1.1653	0.0139	-1.1275	0.8221	-2.0731	0.2934	-0.9456	0.6372	0.1124	0.9914	-1.2858	0.5777	-1.3982	0.5444
---	---	1638524_s_at	0.0101	0.9732	0.0262	0.8736	0.1693	0.3493	0.1383	0.8189	0.0361	0.9128	-0.1022	0.6796	0.0934	0.8692	0.1339	0.5750	0.0406	0.8955
Or67c	Odorant receptor	1638525_at	0.0525	0.7578	0.0143	0.8923	-0.0533	0.7257	-0.0079	0.9912	-0.0080	0.9734	-0.0001	0.9995	0.0633	0.8981	0.0235	0.9363	-0.0399	0.8715
CG3744	CG3744	1638526_a_at	-0.0794	0.8618	0.2323	0.5879	0.3528	0.1281	0.0571	0.9627	0.2774	0.4076	0.2204	0.4774	0.0151	0.9950	0.7098	0.1845	0.6947	0.2234
CG40196	CG40196	1638527_a_at	-0.1064	0.7072	0.4222	0.2947	-0.1406	0.4682	-0.5145	0.2442	-0.4250	0.1040	0.0895	0.7545	-0.0148	0.9928	0.0778	0.8903	0.0925	0.8533
CG15531	CG15531	1638528_at	0.4769	0.7180	-0.9509	0.6698	1.3999	0.1285	1.7778	0.2753	0.8260	0.4024	-0.9518	0.2718	-0.6835	0.9309	-0.8150	0.8048	-0.1315	0.9740
Scsalpha	succinyl-CoA synt	1638529_at	0.0629	0.6936	0.2061	0.6177	0.1810	0.4173	-0.1774	0.6634	-0.8056	0.0023	-0.6282	0.0040	-0.2661	0.8222	-0.6296	0.1874	-0.3635	0.4753
---	---	1638530_at	0.0851	0.7126	0.1302	0.3485	-0.1027	0.5578	0.0662	0.9293	-0.0464	0.8725	-0.1126	0.6100	0.0301	0.9841	0.0350	0.9498	0.0049	0.9935
CG40096	CG40096	1638531_at	0.2196	0.3181	0.0040	0.9810	-0.0289	0.9021	-0.0646	0.8955	-0.0830	0.6510	-0.0184	0.9263	-0.0287	0.9774	-0.0705	0.8218	-0.0417	0.8993
CG32376	CG32376	1638532_at	0.2297	0.3518	0.0242	0.8553	0.2055	0.3221	0.3138	0.4690	0.3489	0.1337	0.0351	0.9004	0.2058	0.7644	0.1761	0.5650	-0.0297	0.9429
l(3)neo38	lethal (3) neo38	1638533_at	0.2183	0.1666	0.2446	0.1502	0.1144	0.5857	-0.0889	0.8604	-0.0203	0.9378	0.0686	0.7220	0.1528	0.8534	0.1109	0.7964	-0.0418	0.9299
---	---	1638534_at	0.1494	0.4262	-0.0197	0.9564	-0.0657	0.7471	-0.0519	0.9515	0.0315	0.9235	0.0834	0.7427	-0.0094	0.9943	-0.0839	0.8428	-0.0745	0.8543
Shab	Shaker cognate b	1638535_a_at	-0.2285	0.4183	-0.1934	0.3783	-0.3943	0.0561	0.0680	0.9039	0.1098	0.5817	0.0418	0.8434	0.0306	0.9816	-0.1537	0.6109	-0.1843	0.5361
---	---	1638536_at	0.2237	0.4066	0.0031	0.9832	0.1058	0.6452	0.0085	0.9941	0.0381	0.9077	0.0296	0.9187	-0.1316	0.9004	-0.1293	0.7892	0.0023	0.9979
mun	munin	1638537_at	0.0885	0.5746	0.1093	0.6920	0.2244	0.1481	0.2248	0.4239	0.2554	0.1005	0.0307	0.8658	0.1308	0.8517	0.1379	0.6658	0.0071	0.9874
CG16970	CG16970	1638538_at	0.0019	0.9936	-0.0475	0.8738	0.0484	0.8153	0.1320	0.8459	0.0279	0.9378	-0.1041	0.6914	0.0382	0.9816	0.1253	0.8023	0.0871	0.8669
CG7137	Gene 3	1638539_at	0.3853	0.2790	0.1818	0.4479	0.5346	0.0621	0.0646	0.9529	0.4020	0.1811	0.3373	0.2118	-0.2714	0.8202	0.2000	0.7241	0.4714	0.3519
---	---	1638540_at	-0.0398	0.8285	-0.0305	0.7999	-0.0115	0.9641	0.0314	0.9677	0.1755	0.4034	0.1441	0.4561	-0.0886	0.8705	0.0385	0.9075	0.1271	0.5814
mRpL32	mitochondrial ribo	1638541_at	0.1517	0.4681	-0.0928	0.7971	-0.8217	0.0180	-0.4120	0.4895	0.5732	0.0738	0.9852	0.0048	0.2853	0.8049	0.4152	0.3815	0.1299	0.8308
CG15685	CG15685	1638542_at	0.0897	0.7329	0.0909	0.5390	-0.0139	0.9485	-0.0923	0.8796	0.0559	0.8342	0.1483	0.4525	-0.2305	0.7485	-0.0890	0.8248	0.1415	0.6724
CG33775	CG33775	1638543_at	-0.6580	0.0077	-1.0056	0.0360	-1.3484	0.0076	-0.2709	0.7507	0.4941	0.1799	0.7650	0.0286	-0.1034	0.9056	0.0042	0.9950	0.1076	0.7691
CG12929	CG12929	1638544_at	-0.3734	0.3255	-0.3176	0.1477	-0.1239	0.5003	0.1254	0.7663	-0.0976	0.6253	-0.2230	0.1645	-0.1083	0.9474	-0.0705	0.9289	0.0378	0.9585
CG30409	CG30409	1638545_at	0.1602	0.3457	0.0674	0.7533	0.0240	0.9213	-0.0233	0.9777	-0.1024	0.6731	-0.0791	0.7283	0.0570	0.9550	-0.1518	0.6344	-0.2088	0.4944
CG13218	CG13218	1638546_at	0.0134	0.9528	0.2348	0.3194	0.3223	0.0329	-0.1076	0.7929	-0.1570	0.3566	-0.0494	0.7915	-0.2292	0.7307	0.0395	0.9310	0.2687	0.3614
CG7955 /// DmirCG7955	CG7955	1638547_s_at	-0.2980	0.1203	0.0060	0.9606	0.1706	0.3039	-0.0442	0.9356	-0.3848	0.0253	-0.3405	0.0264	-0.3017	0.6832	0.0284	0.9538	0.3300	0.2938
CG15452	CG15452	1638548_at	0.0421	0.8161	-0.0036	0.9840	-0.0554	0.7944	0.0633	0.9116	0.1540	0.4089	0.0907	0.6230	0.1186	0.8706	0.0861	0.8196	-0.0324	0.9362
CG31460	CG31460	1638549_at	0.4004	0.2529	0.9444	0.0099	1.0669	0.0107	-0.1305	0.8667	-0.3575	0.1901	-0.2270	0.3703	-0.2472	0.8439	0.2263	0.6969	0.4734	0.3713
Pgi	glucose phosphat	1638550_s_at	0.0592	0.6853	-0.2068	0.6000	0.0905	0.7058	0.2699	0.4908	-0.4408	0.0410	-0.7106	0.0032	-0.1154	0.9405	-0.8712	0.0795	-0.7558	0.1409
CG13171	CG13171	1638551_at	0.0874	0.6835	0.0607	0.5534	0.3116	0.1165	0.0935	0.8498	-0.1255	0.5096	-0.2189	0.1728	0.0385	0.9689	-0.0734	0.8303	-0.1119	0.6974
dpr	defective proboscl	1638552_at	0.1842	0.3982	0.0432	0.6844	-0.0236	0.8836	0.1742	0.5932	0.2291	0.1640	0.0549	0.7618	0.0393	0.9514	0.0074	0.9841	-0.0319	0.9055
---	---	1638553_at	-0.0184	0.9487	0.0242	0.8101	-0.0309	0.9255	0.0903	0.8271	0.0753	0.6783	-0.0150	0.9395	0.1484	0.9031	0.1300	0.8228	-0.0183	0.9803
CG4115	CG4115	1638554_at	-0.7646	0.1142	-0.2386	0.5321	-1.2911	0.0001	-0.4847	0.1807	-1.2242	0.0007	-0.7395	0.0033	0.4357	0.7726	-0.8077	0.1974	-1.2433	0.0893
---	---	1638555_at	0.0820	0.7700	-0.0447	0.7552	-0.0160	0.9432	0.0492	0.9646	-0.0103	0.9813	-0.0596	0.8615	-0.0408	0.9717	-0.0241	0.9605	0.0167	0.9704
Oamb	octopamine recep	1638556_s_at	-2.3645	0.0009	-2.0853	0.0252	-2.1272	0.0005	-0.5939	0.1663	-1.3214	0.0010	-0.7274	0.0079	-0.4673	0.8128	-0.9145	0.2466	-0.4471	0.6126
CG34411 /// DsimCG32637	CG32637	1638557_at	0.0705	0.7834	0.0365	0.7589	0.2963	0.1984	0.1150	0.8908	0.0530	0.8892	-0.0620	0.8490	-0.0748	0.9400	-0.0226	0.9643	0.0522	0.9016
Snap25	Synapse protein 2	1638558_a_at	-0.7460	0.0662	-0.2020	0.2223	-0.5026	0.0739	-0.0546	0.9683	-0.2481	0.5138	-0.1934	0.5900	-0.0989	0.9142	0.0107	0.9870	0.1096	0.7710
---	---	1638559_at	0.0542	0.8301	-0.0210	0.9173	-0.0069	0.9750	-0.0249	0.9786	0.0838	0.7640	0.1087	0.6519	0.0219	0.9816	0.0338	0.9123	0.0120	0.9688
---	---	1638560_a_at	-0.3745	0.3155	0.6611	0.0413	0.8561	0.0025	0.0235	0.9838	-0.5293	0.0562	-0.5528	0.0306	-0.1808	0.8521	0.4382	0.2550	0.6190	0.1490
---	---	1638561_x_at	0.0585	0.8387	-0.3405	0.0598	-0.5195	0.1436	0.0539	0.9763	0.4991	0.2694	0.4452	0.2728	-0.0373	0.9611	0.0056	0.9908	0.0430	0.8837
Cyp6d5	Cyp6d5	1638562_a_at	0.9732	0.1175	-0.1661	0.8948	0.4595	0.0372	-0.1691	0.8578	-0.6413	0.0626	-0.4723	0.1186	-0.5085	0.8802	-1.4550	0.2430	-0.9465	0.4791
CG6490	CG6490	1638563_at	-1.2184	0.0437	-0.2671	0.8316	-1.7750	0.0036	-1.2154	0.2063	-1.3515	0.0272	-0.1361	0.8376	0.1775	0.9717	-0.2985	0.8633	-0.4760	0.7408
CG12279	CG12279	1638564_at	0.1708	0.5389	0.6745	0.0487	0.7944	0.0049	-0.0099	0.9937	-0.2461	0.3582	-0.2362	0.3242	-0.1124	0.9101	0.1397	0.7371	0.2521	0.4938

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
bap	bagpipe	1638565_at	0.4813	0.1205	-0.0226	0.8523	0.2213	0.2903	0.0794	0.9068	0.0837	0.7469	0.0043	0.9874	-0.1053	0.9246	-0.1791	0.6611	-0.0738	0.8856
---	---	1638566_at	-0.0624	0.7473	-0.0674	0.7185	0.1348	0.5333	0.1762	0.7487	0.0582	0.8514	-0.1180	0.6290	-0.1061	0.8767	-0.0036	0.9950	0.1025	0.7391
CG1092	CG1092	1638567_at	-0.1106	0.7189	0.1061	0.5518	-0.3502	0.0470	-0.3376	0.4653	-0.4272	0.0884	-0.0896	0.7392	0.0866	0.9168	-0.1508	0.6259	-0.2374	0.4148
H	hairless	1638568_s_at	0.5188	0.5936	0.3910	0.6016	-0.6007	0.1957	-0.2745	0.8532	0.7955	0.1367	1.0701	0.0331	0.6964	0.8465	0.7534	0.6425	0.0570	0.9824
---	---	1638569_at	-0.0651	0.6618	0.0126	0.9046	-0.1682	0.4568	-0.2223	0.5932	-0.0448	0.8729	0.1775	0.3607	0.1035	0.8965	0.0363	0.9402	-0.0672	0.8636
RpL10Ab	Ribosomal protein	1638570_a_at	0.1980	0.1957	0.3351	0.0368	0.5282	0.0193	0.1893	0.6280	-0.1019	0.6363	-0.2911	0.0906	-0.0479	0.9352	-0.0234	0.9390	0.0245	0.9246
CG32588	CG32588	1638571_at	0.1096	0.5601	0.0088	0.9359	0.1259	0.4287	0.0770	0.8942	0.1877	0.3390	0.1107	0.5622	0.0009	0.9994	-0.0368	0.8988	-0.0377	0.8864
---	---	1638572_s_at	0.3105	0.1025	0.1507	0.3906	-0.0626	0.7418	-0.0982	0.8337	-0.1199	0.5258	-0.0217	0.9209	0.2598	0.6955	-0.0258	0.9537	-0.2856	0.3110
CG7140	CG7140	1638573_at	0.1686	0.4422	-0.0489	0.6556	0.2879	0.1264	0.1212	0.7380	0.1113	0.5112	-0.0099	0.9605	-0.2495	0.6728	-0.0327	0.9316	0.2168	0.3762
---	---	1638574_at	-0.0687	0.7710	0.0874	0.6286	0.1523	0.4654	-0.2059	0.5720	-0.2526	0.1725	-0.0467	0.8254	-0.1611	0.8122	0.0212	0.9634	0.1823	0.5387
Mcm6	Minichromosome	1638575_at	0.5732	0.0489	-0.5048	0.4578	-0.2537	0.3718	-0.4253	0.3744	0.5188	0.0594	0.9442	0.0028	-0.5912	0.7677	-0.3539	0.7149	0.2373	0.8237
Rpb11	Rpb11	1638576_at	-0.0678	0.7474	0.4369	0.0470	0.0949	0.7185	-0.0908	0.8299	-0.1578	0.3375	-0.0670	0.6939	0.2704	0.7726	0.3006	0.4523	0.0302	0.9584
---	---	1638577_s_at	0.2040	0.3330	0.1294	0.4904	0.2678	0.1054	0.0971	0.8865	0.0944	0.7250	-0.0026	0.9927	0.0221	0.9816	-0.0047	0.9925	-0.0268	0.9316
CG15245	CG15245	1638578_at	0.1885	0.5466	0.1183	0.3808	0.1552	0.5301	-0.0224	0.9803	-0.1260	0.6054	-0.1036	0.6519	0.1978	0.7644	0.0248	0.9561	-0.1730	0.5525
---	---	1638579_at	0.0150	0.9612	-0.0042	0.9901	0.1393	0.3284	0.2335	0.6854	0.0510	0.8880	-0.1825	0.4741	-0.0542	0.9474	-0.0177	0.9655	0.0366	0.9161
caps	capricious	1638580_at	-1.1591	0.1007	0.2918	0.5452	0.0937	0.8463	0.1299	0.9300	-0.7698	0.0776	-0.8997	0.0280	0.4333	0.8453	0.6521	0.4833	0.2188	0.8525
CG14224	CG14224	1638581_at	-0.4828	0.2921	0.1014	0.8988	0.3462	0.1172	-0.0056	0.9943	-0.0372	0.8632	-0.0316	0.8700	-0.2730	0.9231	0.4384	0.6713	0.7114	0.4616
CG5321	CG5321	1638582_at	2.0262	0.0066	1.4590	0.2384	2.7269	0.0001	0.5571	0.2390	0.3116	0.2674	-0.2455	0.3378	-0.7460	0.8181	-0.3386	0.8527	0.4074	0.8007
CG31103	CG31103	1638583_at	-2.6351	0.0037	0.3175	0.1264	-0.6433	0.0866	-1.2964	0.1341	-3.4058	0.0004	-2.1094	0.0015	-0.3485	0.7953	-0.4761	0.3915	-0.1277	0.8636
CG9988	CG9988	1638584_at	0.0517	0.7785	-0.0925	0.5237	-0.1680	0.3944	0.1957	0.6354	0.2510	0.2147	0.0553	0.8086	0.0352	0.9816	-0.1467	0.6853	-0.1819	0.6022
CG4721	CG4721	1638585_at	0.4607	0.2587	0.9473	0.0949	1.1249	0.0094	-0.6766	0.4690	-1.2863	0.0189	-0.6097	0.1747	-0.6710	0.5461	-0.7690	0.1374	-0.0979	0.8908
hkb	huckebein	1638586_at	0.0842	0.7443	-0.0826	0.5601	0.0380	0.8761	0.1118	0.8671	0.1240	0.6390	0.0122	0.9676	0.0355	0.9679	-0.0918	0.7371	-0.1273	0.6140
tor	torso	1638587_at	0.4320	0.3887	-0.7347	0.2753	-1.1028	0.0365	-0.3701	0.4245	1.2417	0.0012	1.6118	0.0003	-0.2921	0.9390	0.0137	0.9961	0.3058	0.8375
CG15522	CG15522	1638588_at	0.1431	0.6113	0.0456	0.6830	-0.0984	0.5584	-0.1903	0.5915	-0.1724	0.3460	0.0179	0.9362	0.1704	0.7498	0.1141	0.6449	-0.0563	0.8474
CG18538	CG18538	1638589_at	0.1140	0.6326	0.0567	0.6159	0.0681	0.6647	-0.0258	0.9763	-0.0787	0.7607	-0.0529	0.8296	-0.0376	0.9557	0.0232	0.9414	0.0608	0.7989
CG30467	CG30467	1638590_at	-0.0101	0.9649	0.4324	0.0625	0.7040	0.0041	-0.0150	0.9907	-0.1845	0.5385	-0.1695	0.5350	-0.2758	0.7133	0.2685	0.3820	0.5443	0.1171
Tif-IA	Tif-IA	1638591_at	0.5279	0.0526	0.1886	0.6301	0.0800	0.6638	-0.2565	0.4073	0.1874	0.2789	0.4439	0.0112	0.0411	0.9848	0.0245	0.9769	-0.0166	0.9838
CG6560	CG6560	1638592_at	1.0727	0.1495	0.6140	0.1376	0.6127	0.0946	-0.0626	0.9786	-1.4250	0.0234	-1.3624	0.0176	-0.1479	0.9555	-1.4729	0.0714	-1.3250	0.1190
ND75	NADH:ubiquinone	1638593_a_at	-1.5698	0.0160	0.5956	0.5884	0.8392	0.0215	0.0860	0.8350	-1.9571	0.0000	-2.0432	0.0000	0.0301	0.9952	0.2148	0.9085	0.1846	0.9125
Gr28b	Gustatory recepto	1638594_at	0.1909	0.3525	0.1120	0.5256	-0.0683	0.7002	-0.0851	0.8822	0.0319	0.9068	0.1170	0.5427	0.0119	0.9914	-0.1329	0.6312	-0.1448	0.5991
---	---	1638595_at	0.2045	0.3522	-0.0673	0.6025	-0.0472	0.8015	-0.0722	0.8796	0.1175	0.4929	0.1896	0.1930	-0.2120	0.7726	-0.0761	0.8631	0.1360	0.7004
ac	achaete	1638596_at	0.0670	0.7645	-0.0060	0.9583	-0.2490	0.3090	-0.2960	0.4717	0.0108	0.9749	0.3068	0.1200	-0.0326	0.9701	-0.0097	0.9814	0.0228	0.9416
CG32733 /// CG32797	CG32733 /// CG3	1638597_s_at	-0.1689	0.5258	-0.2789	0.2810	-0.5729	0.0143	0.0383	0.9620	0.1675	0.4506	0.1292	0.5351	0.2743	0.6955	0.2028	0.5075	-0.0715	0.8524
MED15	Mediator complex	1638598_at	0.2052	0.5949	0.3365	0.0772	0.2652	0.1070	-0.0198	0.9803	0.3489	0.0754	0.3687	0.0404	0.0506	0.9816	0.4628	0.3158	0.4122	0.3921
---	---	1638599_at	0.0023	0.9907	0.0258	0.9153	0.0519	0.7457	-0.0114	0.9878	0.0758	0.7070	0.0871	0.6246	-0.0748	0.9239	-0.1038	0.7339	-0.0290	0.9369
ebi	beta transducin-lik	1638600_at	0.5243	0.0349	0.1945	0.5164	0.1002	0.6690	-0.0897	0.9068	0.2567	0.3041	0.3465	0.1160	0.0830	0.9421	-0.0145	0.9829	-0.0975	0.8255
spo2	spook 2	1638601_at	-0.0193	0.9245	0.0569	0.7054	0.0181	0.9355	-0.0775	0.8678	-0.1462	0.3862	-0.0687	0.6914	-0.0096	0.9928	0.0744	0.8244	0.0840	0.7812
---	---	1638602_at	0.0008	0.9972	0.0134	0.9130	0.2323	0.2390	0.0241	0.9803	0.1117	0.6783	0.0876	0.7287	-0.1514	0.7644	0.1073	0.6462	0.2586	0.2470
CG31909	CG31909	1638603_at	0.5145	0.3070	0.0184	0.9536	0.2044	0.2047	-0.2147	0.5008	0.1012	0.5817	0.3159	0.0429	-0.4427	0.7215	-0.3889	0.4600	0.0538	0.9416
CG8117	CG8117	1638604_at	0.0712	0.6575	0.0821	0.5902	-0.0599	0.7700	0.0039	0.9956	0.0663	0.6834	0.0623	0.6752	0.1200	0.8706	0.0805	0.8381	-0.0395	0.9224
CG4495	CG4495	1638605_at	-0.2830	0.2414	-0.2557	0.4934	0.0088	0.9641	-0.1561	0.6929	-0.2123	0.2458	-0.0562	0.7825	-0.3746	0.7149	-0.2490	0.5825	0.1256	0.8129
---	---	1638606_at	0.0450	0.8526	0.1957	0.1184	-0.0192	0.9062	-0.1042	0.8140	-0.0877	0.6561	0.0165	0.9394	0.1953	0.7324	0.1271	0.6312	-0.0682	0.8246
Ef1gamma	Ef1gamma	1638607_s_at	0.1365	0.4226	0.4579	0.0999	0.6603	0.0163	0.2038	0.7252	-0.2830	0.2804	-0.4869	0.0429	-0.0541	0.9632	0.0234	0.9647	0.0776	0.8514
---	---	1638608_at	0.0427	0.8366	-0.0213	0.8522	0.0184	0.9239	0.0437	0.9433	-0.1450	0.4185	-0.1888	0.2245	-0.0171	0.9898	-0.0439	0.9222	-0.0268	0.9471
MBD-R2	MBD-R2	1638609_s_at	-0.6311	0.1810	0.4502	0.1673	0.7111	0.0034	0.2158	0.7143	-0.6433	0.0249	-0.8591	0.0042	-0.0310	0.9914	0.4927	0.4138	0.5238	0.3969
CG7802	CG7802	1638610_at	0.0721	0.6920	-0.1303	0.4324	0.1396	0.3338	0.1640	0.7062	0.2050	0.3078	0.0411	0.8599	-0.0854	0.8653	-0.0343	0.9112	0.0511	0.8418
---	---	1638611_at	-0.0299	0.9629	-0.3575	0.5937	-0.9505	0.0304	-0.4380	0.6578	-0.2628	0.6170	0.1752	0.7307	0.2126	0.9342	-0.4667	0.5972	-0.6792	0.4220
obst-E	CG11142	1638612_at	-3.1567	0.0025	-2.4743	0.0030	-4.1987	0.0001	-0.8918	0.2789	-1.6430	0.0055	-0.7512	0.0847	0.6873	0.7726	-1.0096	0.3068	-1.6969	0.1271
CG32794	CG32794	1638613_at	0.2029	0.4539	-0.1587	0.4522	0.0231	0.9539	0.2043	0.7929	0.2853	0.3787	0.0809	0.8219	-0.1943	0.8202	-0.1124	0.8018	0.0818	0.8578
CG18273	CG18273	1638614_at	-2.4132	0.0004	-0.8537	0.0963	-1.8600	0.0020	-0.8773	0.2492	-1.4387	0.0072	-0.5615	0.1627	0.1051	0.9495	0.2168	0.6962	0.1116	0.8637
RpS7	Ribosomal protein	1638615_s_at	0.2939	0.1772	1.1677	0.0087	1.4549	0.0006	0.2170	0.7136	-1.0293	0.0032	-1.2463	0.0008	-0.0226	0.9845	-0.0471	0.9055	-0.0245	0.9463
mthl9	Mth-like 9	1638616_at	2.0995	0.0008	1.6752	0.0418	2.3685	0.0000	0.0596	0.9116	0.4721	0.0130	0.4126	0.0144	-0.6037	0.7018	-0.0129	0.9935	0.5908	0.3813
CG33725	CG33725	1638617_at	0.1962	0.3609	-0.0858	0.7090	0.0480	0.8091	-0.0343	0.9761	-0.0412	0.9175	-0.0069	0.9850	-0.0954	0.8513	-0.0144	0.9666	0.0810	0.7391
CG5877	CG5877	1638618_a_at	-0.3996	0.1151	0.3508	0.4808	0.5544	0.0082	0.1277	0.8387	-0.2406	0.3111	-0.3684	0.0812	-0.0696	0.9741	0.4463	0.3892	0.5159	0.3388
CG17698	CG17698	1638619_s_at	-0.5925	0.2786	0.4727	0.2122	0.0521	0.8883	-0.1660	0.8721	-0.8824	0.0222	-0.7164	0.0329	0.1116	0.9611	0.0803	0.9387	-0.0313	0.9757
---	---	1638620_x_at	-0.0587	0.7781	-0.4520	0.1298	-1.0049	0.0097	-0.0573	0.9558	0.5324	0.0596	0.5897	0.0258	0.0608	0.9742	-0.1368	0.8220	-0.1977	0.7053
CG31648	CG31648	1638621_at	-0.4142	0.0783	0.0476	0.7480	0.2125	0.4011	-0.1966	0.6325	-0.6334	0.0076	-0.4368	0.0234	-0.1348	0.9250	-0.0193	0.9834	0.1155	0.8525
inaF	inaF	1638622_at	0.0558	0.7563	-0.1417	0.3373	-0.2075	0.2569	0.0200	0.9790	0.0741	0.7396	0.0541	0.7989	-0.0502	0.9503	-0.2492	0.2814	-0.1990	0.4114
---	---	1638623_at	0.1092	0.5923	0.2440	0.1233	0.0817	0.6640	-0.2052	0.4927	-0.3071	0.0581	-0.1019	0.5073	0.0133	0.9898	0.0491	0.8721	0.0358	0.9031
P5cr	Pyrroline 5-carboy	1638624_at	0.2154	0.5851	0.3335	0.0373	0.6139	0.0295	-0.0315	0.9633	-0.2996	0.0993	-0.2680	0.0988	-0.1849	0.8972	-0.2270	0.7065	-0.0420	0.9563
---	---	1638625_at	0.3441	0.0979	0.2760	0.3659	0.1992	0.2787	-0.1315	0.7782	0.1796	0.3659	0.3111	0.0753	-0.0046	0.9964	0.0313	0.9411	0.0359	0.9211
CG17377	CG17377	1638626_a_at	-0.0022	0.9907	0.0264	0.7934	0.1601	0.4557	0.0496	0.9466	-0.0673	0.7932	-0.1169	0.5755	-0.1321	0.7726	-0.0081	0.9829	0.1240	0.5506
---	---	1638627_at	0.0159	0.9555	-0.0296	0.7814	-0.0806	0.6677	0.1110	0.8349	-0.0775	0.7451	-0.1885	0.3021	0.1666	0.8400	-0.0139	0.9829	-0.1805	0.6234
---	---	1638628_at	-0.0558	0.7717	-0.0140	0.8923	-0.0871	0.6195	0.0682	0.9029	0.1130	0.5631	0.0448	0.8284	0.0571	0.9237	-0.0348	0.9075	-0.0919	0.6668
CG15918	CG15918	1638629_at	-0.0216	0.9177	-0.0174	0.8696	0.0358	0.8909	-0.0163	0.9860	0.1395	0.5532	0.1558	0.4494	-0.0347	0.9779	0.0116	0.9841	0.0463	0.9109
CG1983	CG1983	1638630_at	-0.1369	0.5558	-0.2612	0.3845	0.0429	0.8082	0.0761	0.8794	0.1865	0.2789	0.1104	0.5028	-0.1483	0.8814	0.2007	0.6233	0.3490	0.3708
---	---	1638631_at	0.2036	0.2144	0.0131	0.9399	0.0896	0.6768	0.1334	0.7937	0.3646	0.0814	0.2311	0.2148	0.1743	0.7917	0.1986	0.4875	0.0243	0.9510
---	---	1638632_at	0.0887	0.6831	0.0544	0.6110	0.0444	0.8191	-0.0937	0.8471	-0.1317	0.4796	-0.0380	0.8545	0.0955	0.8692	-0.1075	0.6762	-0.2030	0.3898
ppk7	pickpocket 7	1638633_at	0.1897	0.3271	0.1569	0.4055	0.2817	0.3096	0.0917	0.8967	0.0183	0.9577	-0.0734	0.7760	0.0020	0.9994	0.0637	0.8831	0.0617	0.8800

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG32448	CG32448	1638634_at	0.1221	0.6868	0.2247	0.3848	0.5482	0.0244	0.1892	0.6854	-0.3458	0.1104	-0.5350	0.0137	-0.0799	0.9511	-0.1982	0.6331	-0.1183	0.8000
exo70	exo70	1638635_at	-0.0945	0.5471	0.2069	0.3902	0.4833	0.0073	-0.1487	0.6903	-0.1983	0.2524	-0.0496	0.7981	-0.3371	0.6749	0.0926	0.8297	0.4297	0.2075
CG12512	CG12512	1638636_at	-0.8697	0.0070	0.8521	0.2906	0.9181	0.0035	-0.1626	0.7805	-1.3179	0.0007	-1.1553	0.0008	-0.2099	0.9388	0.1905	0.8749	0.4004	0.6607
---	---	1638637_at	-0.2762	0.0980	0.2925	0.2583	0.4740	0.0217	-0.0039	0.9956	-0.4668	0.0265	-0.4629	0.0172	-0.1507	0.8270	0.1530	0.6216	0.3037	0.3037
CG12525	CG12525	1638638_at	0.1235	0.4970	0.0664	0.6512	0.1894	0.4503	-0.0490	0.9353	0.0197	0.9392	0.0687	0.7179	-0.0160	0.9922	-0.0392	0.9476	-0.0232	0.9661
CG18854 /// how	CG18854 /// strutt	1638639_a_at	0.5078	0.3024	-0.1529	0.5741	-0.9510	0.1079	-0.1991	0.6998	0.6913	0.0106	0.8904	0.0020	0.4203	0.8806	0.1322	0.9396	-0.2881	0.8307
CG15576	CG15576	1638640_at	0.1038	0.7039	0.0536	0.8459	-0.0510	0.7743	0.0590	0.9504	0.0193	0.9606	-0.0396	0.9000	-0.0866	0.8609	-0.1019	0.6380	-0.0153	0.9575
---	---	1638641_at	0.1743	0.2800	0.1124	0.4639	-0.0315	0.9127	-0.1256	0.7121	-0.0308	0.8811	0.0948	0.5287	0.0813	0.9445	-0.0584	0.9152	-0.1397	0.7298
CG17259	CG17259	1638642_at	0.7986	0.0207	1.1121	0.0195	1.2135	0.0001	0.2595	0.4568	0.3612	0.0602	0.1017	0.5912	0.1277	0.9142	0.6538	0.1206	0.5261	0.2230
---	---	1638643_at	0.1546	0.3137	-0.1335	0.2353	0.0739	0.6827	0.0995	0.7982	0.2537	0.1068	0.1542	0.2792	-0.1440	0.7707	-0.1196	0.5909	0.0244	0.9342
CG12479	CG12479	1638644_at	0.0724	0.7644	0.0128	0.9343	0.1759	0.2704	0.1490	0.7409	0.0173	0.9531	-0.1317	0.4884	0.0532	0.9589	-0.0829	0.8287	-0.1362	0.6660
---	---	1638645_at	0.1086	0.6808	0.1403	0.4567	0.4960	0.0130	0.3510	0.2501	-0.0243	0.9234	-0.3753	0.0284	0.1058	0.8960	0.0732	0.8619	-0.0326	0.9386
---	---	1638646_at	-0.0351	0.8723	0.0884	0.4698	0.1388	0.3486	-0.0393	0.9518	-0.1090	0.5807	-0.0697	0.7183	-0.0281	0.9777	-0.0797	0.7910	-0.0516	0.8729
Adgf-E	Adenosine deamir	1638647_at	-0.0087	0.9633	0.0983	0.4360	0.0253	0.9207	0.0858	0.8496	-0.0280	0.9020	-0.1138	0.4667	0.1911	0.7853	0.0869	0.8262	-0.1042	0.7674
CG8160	CG8160	1638648_at	0.0328	0.8530	-0.1340	0.4343	-0.0977	0.6391	-0.0206	0.9833	0.0699	0.7990	0.0905	0.7010	-0.1531	0.7686	-0.1320	0.5711	0.0211	0.9460
CG11340	CG11340	1638649_at	0.0383	0.8693	0.0581	0.8113	0.2496	0.1807	0.1987	0.6409	0.1905	0.3700	-0.0082	0.9747	0.1304	0.8330	0.0411	0.9175	-0.0893	0.7654
---	---	1638650_at	0.1330	0.4561	0.0226	0.8354	0.0583	0.7556	-0.0833	0.8894	-0.0987	0.6648	-0.0154	0.9517	0.0547	0.9246	-0.0250	0.9353	-0.0797	0.7154
CG3508	CG3508	1638651_a_at	-0.3356	0.1302	-0.0046	0.9682	-0.1094	0.5411	-0.0942	0.8327	-0.1382	0.4305	-0.0440	0.8187	-0.0325	0.9721	0.1378	0.5818	0.1702	0.4852
CG40006	CG40006	1638652_at	-0.5854	0.2665	-0.4059	0.5525	-1.2074	0.0017	-0.7801	0.2561	-1.2635	0.0081	-0.4834	0.1838	0.1634	0.9507	-0.9004	0.2361	-1.0638	0.1990
CG9932	CG9932	1638653_a_at	-1.6896	0.0068	-1.0866	0.0723	-1.0565	0.0045	0.0819	0.9074	-0.3169	0.1600	-0.3987	0.0534	0.1354	0.9592	0.1932	0.8539	0.0578	0.9582
CG14645 /// DyakCG14645	CG14645	1638654_at	1.0556	0.3811	-0.2979	0.2604	0.4472	0.2611	0.4060	0.3831	0.4513	0.0869	0.0454	0.8841	-0.1753	0.9779	-0.8807	0.5823	-0.7053	0.6677
CG14692	CG14692	1638655_at	-0.0561	0.8024	-0.0164	0.9159	0.1301	0.5048	0.1029	0.8631	0.1305	0.5692	0.0276	0.9150	-0.1498	0.7500	0.0585	0.8244	0.2083	0.3089
CG8646	CG8646	1638656_at	-4.7996	0.0008	-3.8380	0.0016	-4.4858	0.0000	-0.7677	0.1040	-1.4380	0.0009	-0.6704	0.0155	-0.3452	0.8494	-0.7272	0.3247	-0.3819	0.6389
CG33640	CG33640	1638657_at	0.0575	0.7303	0.0473	0.6879	0.1448	0.3190	0.1525	0.6998	-0.0167	0.9513	-0.1691	0.3047	-0.0792	0.8465	-0.0173	0.9499	0.0619	0.7513
Gos28	Gos28	1638658_at	0.2694	0.2550	0.1329	0.6546	0.2314	0.2887	0.2466	0.5008	0.5040	0.0176	0.2575	0.1362	0.2223	0.8033	0.3354	0.3571	0.1131	0.8032
CG11703	CG11703	1638659_at	0.2749	0.1450	0.0326	0.7599	0.1390	0.5430	0.0914	0.8220	0.1066	0.5282	0.0152	0.9386	0.0726	0.9132	0.0619	0.8445	-0.0106	0.9769
---	---	1638660_at	0.3613	0.0702	-0.0792	0.5498	0.0463	0.7986	0.0367	0.9448	0.1247	0.4260	0.0880	0.5552	-0.0458	0.9677	-0.3113	0.2840	-0.2655	0.3860
yip7	Machete	1638661_at	0.8218	0.5991	-1.2115	0.0093	0.2384	0.3481	1.1467	0.0104	1.0781	0.0011	-0.0686	0.7809	-0.3455	0.9611	-0.9918	0.6425	-0.6463	0.7819
---	---	1638662_at	0.0677	0.6839	-0.0148	0.9249	0.0659	0.6835	0.0276	0.9624	-0.0069	0.9775	-0.0345	0.8483	0.0150	0.9860	-0.1017	0.6227	-0.1167	0.5670
---	---	1638663_at	0.7521	0.0054	1.3587	0.0223	1.7595	0.0000	0.1355	0.7225	0.2309	0.1770	0.0953	0.5780	-0.2080	0.8122	0.8884	0.0388	1.0964	0.0333
eIF-4a	Eukaryotic initiatio	1638664_s_at	-0.0844	0.5514	-0.0197	0.8477	-0.1323	0.3607	0.0353	0.9466	-0.0399	0.8341	-0.0752	0.6193	0.1492	0.7387	0.0849	0.6904	-0.0643	0.7747
Antp	Sex comb extra	1638665_at	0.0074	0.9709	0.0581	0.8329	-0.0310	0.8999	-0.1258	0.8546	-0.0215	0.9532	0.1042	0.6890	0.0105	0.9922	0.1025	0.7128	0.0920	0.7447
CG13358	CG13358	1638666_at	0.0670	0.8314	0.3372	0.3228	0.2173	0.1939	-0.0968	0.8883	-0.1682	0.4885	-0.0714	0.7820	-0.0494	0.9514	0.0906	0.7543	0.1400	0.5892
CG6359	CG6359	1638667_a_at	-0.4271	0.0421	-0.1532	0.1989	0.0486	0.8067	-0.0657	0.8927	-0.0397	0.8510	0.0260	0.8939	-0.1701	0.8049	0.1748	0.5608	0.3449	0.2454
---	---	1638668_at	-0.0145	0.9418	-0.0216	0.8923	0.0764	0.6582	0.0860	0.9120	0.0461	0.8906	-0.0400	0.8926	0.0362	0.9514	-0.0187	0.9479	-0.0549	0.7967
CG11395	CG11395	1638669_at	2.3832	0.0016	0.8282	0.2468	1.7051	0.0015	0.4756	0.6247	0.5136	0.2901	0.0380	0.9503	-0.3536	0.8461	-1.1048	0.1382	-0.7512	0.3271
CG5484	CG5484	1638670_a_at	1.0397	0.0023	0.9105	0.0562	0.7485	0.0014	-0.0650	0.9053	0.7020	0.0027	0.7670	0.0011	-0.0242	0.9898	0.5711	0.1505	0.5953	0.1682
CG6196	CG6196	1638671_at	-0.1553	0.4466	0.0522	0.8100	-0.0146	0.9358	-0.1082	0.8479	0.0416	0.8800	0.1498	0.4400	0.1259	0.8222	0.2120	0.3518	0.0861	0.7492
---	---	1638672_at	-0.1296	0.4210	0.0328	0.7596	-0.1680	0.4081	-0.0571	0.9232	-0.1662	0.3626	-0.1091	0.5318	0.1451	0.7731	0.0755	0.7787	-0.0696	0.7907
Bap	beta-adaptin	1638673_at	-0.2671	0.5314	-0.1442	0.7329	-0.0832	0.6263	0.1532	0.7123	0.2852	0.1301	0.1321	0.4608	0.0892	0.9705	0.4209	0.5050	0.3317	0.6150
---	---	1638674_at	0.1456	0.4226	0.2903	0.2543	0.1976	0.2390	-0.1060	0.8498	-0.1532	0.4721	-0.0472	0.8407	0.0608	0.9407	0.1598	0.5453	0.0990	0.7333
CG10680	CG10680	1638675_at	2.2242	0.0038	1.1554	0.0830	2.2781	0.0005	0.6170	0.5317	0.5745	0.2670	-0.0425	0.9482	-0.5011	0.7628	-0.4744	0.5062	0.0267	0.9821
gom	gomdanji	1638676_at	-0.0145	0.9634	0.1314	0.4135	0.1311	0.4186	0.0825	0.9036	-0.1000	0.6930	-0.1825	0.3699	0.0306	0.9804	0.0575	0.8850	0.0269	0.9435
CG12691	CG12691	1638677_at	0.1730	0.3255	0.0125	0.9061	0.1259	0.4684	0.1136	0.7949	0.0763	0.7114	-0.0373	0.8577	-0.0041	0.9967	0.0192	0.9628	0.0232	0.9471
CG6522	CG6522	1638678_at	-1.1567	0.0014	-1.7518	0.0019	-1.9552	0.0000	0.1883	0.7293	0.6371	0.0169	0.4489	0.0446	0.3605	0.5519	0.0799	0.8297	-0.2806	0.3271
Obp57d	Odorant-binding p	1638679_at	-0.4122	0.1369	-0.1176	0.3923	0.0363	0.8642	-0.0698	0.9120	-0.3995	0.0515	-0.3297	0.0689	-0.1605	0.8156	-0.1584	0.6010	0.0020	0.9972
CG17282	CG17282	1638680_at	-0.4571	0.0365	-0.0708	0.8715	0.1850	0.2861	-0.0755	0.8863	-0.2576	0.1477	-0.1821	0.2574	-0.1734	0.8594	0.1361	0.7788	0.3094	0.4463
pio	PioPio	1638681_at	-2.3446	0.0015	-1.3282	0.1047	-2.1443	0.0005	-0.5450	0.2090	-0.9875	0.0033	-0.4426	0.0612	0.3440	0.8692	0.0978	0.9414	-0.2462	0.8103
neur	Neuralised	1638682_a_at	-0.7301	0.0977	-0.2056	0.5387	-1.4703	0.0004	-0.3105	0.7289	-0.3594	0.3799	-0.0489	0.9209	0.7187	0.5128	0.0380	0.9650	-0.6806	0.2136
---	---	1638683_at	-0.1272	0.6304	0.0341	0.8325	-0.0626	0.6944	-0.1922	0.7121	-0.2986	0.2064	-0.1064	0.6656	-0.0489	0.9589	0.1105	0.7200	0.1594	0.5781
CG34372	CG13552	1638684_at	0.2306	0.1875	0.2425	0.3228	0.2182	0.1777	0.0722	0.8794	-0.0157	0.9491	-0.0879	0.5900	0.0624	0.9467	-0.0129	0.9806	-0.0753	0.8336
CG40127	CG40127	1638685_at	-0.3567	0.2246	0.1911	0.3380	-0.0882	0.7597	-0.2608	0.5544	-0.5141	0.0315	-0.2533	0.2145	-0.0479	0.9816	0.0355	0.9628	0.0834	0.8949
---	---	1638686_at	0.2115	0.3637	0.0819	0.7336	0.3472	0.0487	-0.0322	0.9749	-0.2932	0.2448	-0.2610	0.2479	-0.1448	0.8461	-0.3027	0.3133	-0.1580	0.6334
CG40298	CG40298	1638687_at	-0.8529	0.0106	0.0189	0.9745	-0.2363	0.4030	-0.3862	0.5539	-0.9919	0.0102	-0.6057	0.0495	-0.1701	0.8846	-0.1844	0.7165	-0.0143	0.9847
CG15269	CG15269	1638688_at	0.0710	0.7523	0.0390	0.8328	0.0877	0.6079	-0.0251	0.9715	-0.0088	0.9746	0.0163	0.9415	0.1465	0.8202	0.0867	0.7972	-0.0597	0.8656
---	---	1638689_s_at	0.1248	0.5548	0.1301	0.5827	-0.0838	0.7508	-0.1122	0.8818	-0.1168	0.6939	-0.0045	0.9887	0.0851	0.8960	0.0176	0.9649	-0.0675	0.8247
CG5537	CG5537	1638690_at	0.2485	0.2996	0.1875	0.2788	0.0492	0.8139	0.2386	0.4259	0.4724	0.0116	0.2338	0.1129	0.4136	0.6557	0.3952	0.2775	-0.0183	0.9757
CG32202	CG32202	1638691_at	0.1178	0.4650	-0.0426	0.8065	0.2301	0.1811	0.1894	0.6461	0.4429	0.0339	0.2534	0.1551	-0.1257	0.8141	0.2546	0.2349	0.3803	0.1190
CG34118	CG34118	1638692_at	0.2272	0.2434	0.0709	0.5394	0.2007	0.3495	0.1663	0.6202	0.1611	0.3435	-0.0052	0.9799	-0.0398	0.9816	-0.0530	0.9312	-0.0132	0.9838
CG1882	CG1882	1638693_s_at	-0.7672	0.0201	0.2257	0.4804	0.5720	0.0099	0.1194	0.7335	-0.9838	0.0004	-1.1033	0.0002	-0.3520	0.7588	-0.1740	0.7652	0.1779	0.7492
---	---	1638694_at	0.2884	0.1500	0.3799	0.1000	0.2866	0.1950	-0.0262	0.9760	-0.1131	0.6385	-0.0869	0.7013	0.1536	0.7896	0.1198	0.6560	-0.0338	0.9199
CG3415	CG3415	1638695_at	0.7098	0.0623	0.2269	0.6871	0.5222	0.0117	0.3010	0.2501	0.1428	0.3694	-0.1582	0.2589	-0.1795	0.9296	-0.3089	0.6716	-0.1294	0.8868
CG3552	CG3552	1638696_a_at	0.0613	0.8294	0.0924	0.5978	0.2982	0.1111	-0.0893	0.8816	-0.2856	0.1606	-0.1963	0.2891	-0.1328	0.8472	-0.1006	0.7741	0.0321	0.9361
CG5597	CG5597	1638697_at	-2.2725	0.0048	-1.6042	0.0348	-2.3177	0.0003	-0.3487	0.5552	-1.4462	0.0013	-1.0975	0.0025	0.5726	0.8016	-0.6983	0.4691	-1.2709	0.2039
---	---	1638698_at	0.0601	0.7659	0.1079	0.3394	0.1168	0.6544	-0.0570	0.9349	-0.0733	0.7666	-0.0164	0.9500	0.1092	0.8480	-0.0305	0.9363	-0.1397	0.5814
CG6470	CG6470	1638699_at	0.0401	0.8228	0.0002	1.0000	0.1255	0.4929	0.0658	0.9004	0.0508	0.8094	-0.0150	0.9438	-0.0206	0.9831	0.0394	0.9089	0.0600	0.8338
Alk	Alk	1638700_at	-0.2100	0.3914	-0.1937	0.0948	-0.5247	0.0420	-0.2036	0.7104	0.2985	0.2306	0.5021	0.0317	0.0518	0.9653	0.0979	0.8123	0.0461	0.9168
CG3825	CG3825	1638701_at	-0.4185	0.0946	0.0818	0.4614	0.1651	0.3930	0.1399	0.6705	-0.2131	0.1701	-0.3530	0.0197	0.0334	0.9824	0.2734	0.4355	0.2400	0.5121
achi	achintya	1638702_a_at	-0.3348	0.0894	-0.0869	0.4808	-0.0982	0.5751	-0.0124	0.9863	-0.0830	0.6741	-0.0706	0.6983	0.0248	0.9869	0.1977	0.5897	0.1729	0.6416

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG32048	CG32048	1638703_a_at	-2.2288	0.0022	-2.5762	0.0130	-2.8785	0.0000	-0.3029	0.5680	0.0584	0.8718	0.3613	0.1360	0.1515	0.9555	-0.1856	0.8679	-0.3370	0.7040
Kap3	Kinesin II accesso	1638704_at	0.2637	0.4712	-0.2707	0.3036	-0.3980	0.0927	0.0077	0.9937	0.5438	0.0155	0.5360	0.0101	0.1149	0.9462	0.0286	0.9739	-0.0862	0.9034
CG14069	CG14069	1638705_at	0.1771	0.4325	0.2193	0.0721	0.1994	0.2673	0.0166	0.9824	-0.1603	0.3739	-0.1770	0.2662	0.1069	0.8815	0.0644	0.8678	-0.0424	0.9108
---	---	1638706_at	0.0888	0.6674	-0.0548	0.6935	-0.2315	0.2414	0.0329	0.9643	0.1289	0.5378	0.0960	0.6302	0.0913	0.9168	-0.0004	0.9998	-0.0917	0.8044
---	---	1638707_at	0.0945	0.5919	-0.0004	1.0000	0.2918	0.1652	-0.0163	0.9803	0.0656	0.7276	0.0819	0.6165	-0.1136	0.8439	0.0062	0.9908	0.1198	0.6389
gpp	grappa	1638708_s_at	-0.9510	0.0203	-0.7583	0.0186	-1.0473	0.0004	-0.1826	0.7479	0.0845	0.7781	0.2671	0.2287	-0.0426	0.9816	0.1418	0.7856	0.1843	0.6912
Ark	Apaf-1 related kill	1638709_at	-0.2401	0.3972	0.1335	0.6232	0.1064	0.5843	-0.4163	0.3051	-0.3882	0.1033	0.0281	0.9235	-0.1773	0.8439	0.1337	0.7642	0.3109	0.4078
CG14367	CG14367	1638710_at	-0.1331	0.6528	-0.3494	0.3986	-0.7548	0.0180	-0.4036	0.5008	-0.0482	0.9139	0.3554	0.2119	-0.0745	0.9530	-0.1770	0.6711	-0.1026	0.8287
CG5706	CG5706	1638711_at	0.3133	0.1113	0.6780	0.1772	0.6900	0.0107	0.0870	0.8796	0.4059	0.0451	0.3189	0.0723	0.1567	0.9174	0.8107	0.1243	0.6540	0.2255
---	---	1638712_at	0.2196	0.3973	0.0425	0.7329	0.0136	0.9358	-0.1102	0.7588	-0.0369	0.8531	0.0732	0.6437	0.0189	0.9913	-0.0385	0.9462	-0.0574	0.9057
CG33472	CG33472	1638713_at	0.0939	0.6169	0.0548	0.6141	0.0099	0.9703	0.0089	0.9937	0.0341	0.9149	0.0252	0.9280	-0.0928	0.9016	-0.1491	0.6122	-0.0564	0.8800
Acp26Aa	ovulin	1638714_at	-0.1969	0.2950	0.0459	0.7641	0.0807	0.6535	0.1105	0.8000	-0.0154	0.9531	-0.1259	0.4477	-0.0772	0.9365	0.1471	0.6630	0.2243	0.4815
---	---	1638715_s_at	0.1127	0.5923	0.0857	0.5478	-0.1680	0.3332	-0.2079	0.6197	-0.0078	0.9804	0.2002	0.2896	0.2035	0.6955	0.0723	0.7883	-0.1312	0.5649
CG10725	CG10725	1638716_a_at	0.2131	0.3112	0.1750	0.1339	0.4140	0.0852	0.0708	0.8877	-0.1234	0.4871	-0.1942	0.1987	0.0342	0.9775	-0.0011	0.9994	-0.0353	0.9311
CG10932	CG10932	1638717_at	0.9860	0.0070	1.0033	0.0113	1.4002	0.0006	0.1944	0.5969	-0.3145	0.0915	-0.5090	0.0085	-0.1712	0.8882	-0.2141	0.6732	-0.0429	0.9469
---	---	1638718_at	0.2162	0.3710	0.0356	0.7590	0.2033	0.1886	0.0555	0.9136	-0.0603	0.7541	-0.1158	0.4466	-0.1638	0.7644	-0.1996	0.3738	-0.0358	0.9095
CG14011	CG14011	1638719_s_at	0.1886	0.4026	0.1526	0.3716	0.1613	0.2643	0.0316	0.9610	0.0334	0.8862	0.0018	0.9933	0.0378	0.9611	0.0544	0.8550	0.0166	0.9575
CG40322	CG40322	1638720_x_at	0.2431	0.3126	0.1504	0.3823	-0.4151	0.0932	-0.1496	0.8196	0.2264	0.3906	0.3760	0.1015	0.1322	0.8521	0.0155	0.9760	-0.1167	0.7279
CG18624 /// DyakCG18624	CG18624	1638721_s_at	-0.1841	0.6577	-0.0330	0.9225	-0.2594	0.2977	-0.1062	0.8923	-0.8025	0.0086	-0.6963	0.0099	0.1713	0.9111	-0.5889	0.2618	-0.7602	0.1880
roq	roquin	1638722_at	0.3766	0.2121	0.7503	0.1498	0.4666	0.0325	-0.2043	0.6850	0.2634	0.2619	0.4676	0.0326	0.1736	0.8956	0.5510	0.2471	0.3774	0.4597
---	---	1638723_at	0.1790	0.4412	0.0433	0.8217	0.0682	0.7426	0.1108	0.8676	0.2130	0.3750	0.1022	0.6776	0.1480	0.8062	0.0476	0.9009	-0.1004	0.7266
CG18507	CG18507	1638724_at	-1.5208	0.0007	-1.1587	0.0410	-1.2876	0.0004	-0.1610	0.6533	-1.0452	0.0005	-0.8842	0.0006	-0.0286	0.9901	-0.6768	0.1657	-0.6482	0.2137
kek3	Kekkon-3	1638725_at	0.0191	0.9297	-0.0362	0.7957	0.0886	0.6974	0.2596	0.5357	0.3180	0.1445	0.0584	0.8132	-0.0440	0.9619	-0.0578	0.8754	-0.0138	0.9725
l(3)03670	lethal (3) 03670	1638726_at	-0.5345	0.0205	0.5131	0.0518	0.4806	0.0579	-0.0564	0.9491	-1.1250	0.0015	-1.0686	0.0011	-0.0482	0.9588	-0.0513	0.8939	-0.0032	0.9945
---	---	1638727_at	0.0121	0.9699	-0.0823	0.7390	-0.6697	0.0431	-0.1846	0.5317	0.1741	0.2607	0.3586	0.0174	0.4065	0.7506	0.0714	0.9353	-0.3351	0.5637
CG17994	CG17994	1638728_at	0.0620	0.7644	0.1528	0.3196	0.0634	0.7940	0.0851	0.9223	-0.0550	0.8749	-0.1401	0.5950	0.1421	0.8284	0.0335	0.9402	-0.1086	0.7305
CG8589	anon-fast-evolving	1638729_at	0.4388	0.1785	-0.7850	0.4491	-0.8580	0.1703	-0.3424	0.6338	1.3147	0.0035	1.6572	0.0008	-0.2526	0.9555	0.0835	0.9694	0.3361	0.8414
Hsp67Ba	Gene 1	1638730_at	0.2069	0.3013	0.0749	0.4932	0.1614	0.4683	-0.0014	0.9987	0.0847	0.6657	0.0860	0.6246	0.0147	0.9860	-0.0186	0.9505	-0.0334	0.8964
CG10898	CG10898	1638731_at	-0.0388	0.8512	0.1015	0.6492	0.1406	0.3225	-0.1346	0.6908	-0.0603	0.7460	0.0743	0.6449	-0.0998	0.8960	0.2159	0.4495	0.3156	0.2833
CG34131 /// CG7071	CG7071 /// CG34	1638732_at	-0.6813	0.0279	-0.2746	0.3054	-0.2682	0.1430	-0.0733	0.9017	-0.3466	0.0741	-0.2733	0.1108	0.0197	0.9913	0.0670	0.9075	0.0473	0.9279
CG3984	CG3984	1638733_at	2.9581	0.0050	0.7152	0.4737	2.6164	0.0001	1.3475	0.1119	0.9517	0.0635	-0.3959	0.4004	-0.3529	0.9246	-0.8812	0.4917	-0.5283	0.7068
CG4907	CG4907	1638734_at	-0.1306	0.5638	-0.0822	0.4967	0.2711	0.2175	0.1617	0.6937	0.0232	0.9303	-0.1385	0.4382	-0.2007	0.7726	0.0278	0.9538	0.2286	0.4597
prel	preli-like	1638735_at	0.0606	0.7179	0.6567	0.0232	0.8381	0.0008	-0.0803	0.8723	-0.9576	0.0007	-0.8773	0.0006	-0.2755	0.6660	-0.3369	0.1786	-0.0614	0.8548
CG11458	CG11458	1638736_at	0.2598	0.1223	0.0270	0.8230	-0.0500	0.8643	-0.0807	0.8589	0.1152	0.5042	0.1959	0.1788	-0.0400	0.9742	-0.0011	0.9994	0.0389	0.9282
CG17209	CG17209	1638737_at	0.7001	0.0298	0.3407	0.0858	0.5045	0.0127	0.1211	0.8256	0.8429	0.0027	0.7218	0.0032	-0.0274	0.9862	0.4424	0.2195	0.4698	0.2235
ced-6	ced-6	1638738_s_at	-0.3173	0.2464	0.0548	0.5795	-0.3096	0.4613	-0.3086	0.5586	-0.8561	0.0076	-0.5475	0.0315	0.1107	0.9587	-0.4885	0.4046	-0.5992	0.3215
Cyp4ac3	Cyp4ac3	1638739_at	-0.0867	0.9154	0.7484	0.4894	1.1874	0.0090	-0.3041	0.7333	-1.2202	0.0085	-0.9161	0.0181	-0.6500	0.8339	-0.2696	0.8852	0.3804	0.8123
---	---	1638740_at	0.2184	0.1457	0.1142	0.6575	-0.0330	0.8808	-0.0011	0.9988	0.0124	0.9619	0.0135	0.9504	-0.0527	0.9357	-0.1140	0.6075	-0.0613	0.8114
---	---	1638741_at	0.2580	0.3193	0.1200	0.6149	0.4704	0.0576	-0.0303	0.9733	-0.1055	0.6792	-0.0752	0.7584	-0.0618	0.9721	-0.1778	0.7339	-0.1160	0.8385
Cpr67Fb	CG18348	1638742_at	0.3363	0.1035	0.3908	0.3430	0.3043	0.0889	0.0852	0.9228	-0.0992	0.7542	-0.1843	0.4640	0.3154	0.7694	0.1806	0.7345	-0.1348	0.8117
CG31745	CG31745	1638743_at	-0.0197	0.9232	-0.2999	0.3811	-0.3106	0.1217	0.0572	0.9540	0.2798	0.3091	0.2226	0.3773	0.0287	0.9764	0.0790	0.7855	0.0503	0.8726
CG32439 /// CG34410	CG32439	1638744_at	0.0202	0.9071	0.2589	0.3179	-0.0390	0.8342	0.1113	0.8738	0.0691	0.8202	-0.0422	0.8838	0.2252	0.7478	0.2190	0.4669	-0.0062	0.9902
---	---	1638745_at	0.0308	0.8495	-0.1809	0.2092	-0.0074	0.9788	0.1246	0.7149	0.1300	0.4191	0.0054	0.9783	-0.0964	0.8903	-0.0211	0.9617	0.0753	0.8178
---	---	1638746_at	-1.4634	0.0070	-2.7311	0.0061	-2.6701	0.0001	-0.0653	0.9777	1.3174	0.0312	1.3827	0.0161	-0.1441	0.8454	0.1897	0.5503	0.3338	0.2858
---	---	1638747_at	0.3249	0.0920	0.0345	0.8699	0.0865	0.6630	0.1126	0.8609	0.0377	0.9058	-0.0749	0.7682	0.1685	0.7997	0.0151	0.9747	-0.1534	0.6052
emc	extramacrochaeta	1638748_at	0.1842	0.4853	0.5357	0.1550	-0.1339	0.6647	-0.0830	0.8266	-0.0663	0.6930	0.0167	0.9253	0.5761	0.6898	0.3066	0.6312	-0.2696	0.6782
CG34346	CG11072	1638749_at	-0.1473	0.8961	0.3050	0.7261	-0.0921	0.8917	-0.0469	0.9771	0.4685	0.2407	0.5154	0.1471	0.4543	0.9204	0.9542	0.5523	0.4999	0.7836
CG5440	CG5440	1638750_at	0.3245	0.1515	0.2134	0.2787	0.3131	0.0911	0.1574	0.7140	0.1924	0.3338	0.0350	0.8804	-0.0822	0.9246	-0.0685	0.8678	0.0138	0.9764
CG4004	CG4004	1638751_a_at	-0.0088	0.9812	-0.0361	0.8828	-0.1314	0.5663	-0.0248	0.9803	-0.0048	0.9898	0.0200	0.9484	0.0618	0.9717	0.0174	0.9834	-0.0444	0.9427
CG31961	CG31961	1638752_s_at	-0.0573	0.8356	-0.3541	0.2115	-0.2535	0.3286	0.0732	0.9218	0.3242	0.1479	0.2510	0.2124	-0.0902	0.9447	-0.0675	0.9111	0.0227	0.9701
Hs3st-B	CG7890	1638753_at	0.1196	0.5692	0.0020	0.9882	-0.3578	0.0405	-0.1340	0.8085	-0.0382	0.8990	0.0958	0.6774	0.1400	0.8368	-0.0657	0.8678	-0.2057	0.4776
CG8445	CG8445	1638754_s_at	0.0986	0.6700	-0.0396	0.7011	0.0507	0.8150	-0.0354	0.9558	0.1216	0.5025	0.1570	0.3158	-0.1213	0.8461	0.2119	0.4056	0.3332	0.2132
---	---	1638755_at	-0.1370	0.5035	0.1020	0.3862	0.0469	0.8281	-0.2045	0.5415	-0.1563	0.3803	0.0482	0.8053	-0.0570	0.9246	0.0337	0.9140	0.0907	0.6840
CG17279	CG17279	1638756_at	-0.0521	0.7758	-0.1747	0.1940	-0.1202	0.4528	0.2451	0.5117	0.3090	0.1160	0.0639	0.7689	-0.2199	0.7036	-0.0581	0.8629	0.1618	0.5211
CG17780	CG17780	1638757_a_at	-0.2081	0.6299	-0.2982	0.3102	-0.6154	0.0397	-0.1620	0.8984	0.3318	0.4413	0.4938	0.1825	-0.1011	0.9457	-0.1885	0.7098	-0.0874	0.8874
CG3016	CG3016	1638758_at	0.0457	0.7995	-0.1130	0.7054	-0.1113	0.5202	-0.1662	0.6425	0.0536	0.8023	0.2198	0.1576	-0.1665	0.8379	-0.1746	0.6288	-0.0081	0.9877
vnd	ventral nerve cord	1638759_at	-0.0145	0.9504	-0.0800	0.4044	0.0816	0.5794	0.0951	0.8129	0.1721	0.2804	0.0770	0.6352	-0.0175	0.9853	0.0426	0.8962	0.0601	0.8256
CG15545	CG15545	1638760_at	0.0871	0.6927	-0.5540	0.0442	-0.4768	0.0912	0.3378	0.5044	0.8149	0.0089	0.4771	0.0515	0.0865	0.8513	-0.0061	0.9875	-0.0926	0.6539
CG32151	CG32151	1638761_at	0.0624	0.7369	0.0283	0.8919	0.1303	0.4873	0.0666	0.9116	0.0726	0.7482	0.0060	0.9799	-0.1825	0.7726	0.0966	0.7722	0.2791	0.3175
RpL10Aa	Ribosomal protein	1638762_at	0.3820	0.0919	-0.0613	0.6179	-0.0004	0.9984	-0.0524	0.9230	0.0172	0.9426	0.0697	0.6854	-0.1064	0.9235	-0.2915	0.4234	-0.1851	0.6386
CG32030	CG32030	1638763_at	0.4836	0.2650	0.4180	0.1286	0.4183	0.0459	-0.0176	0.9873	-0.1445	0.6245	-0.1269	0.6418	0.1565	0.9309	0.1466	0.8551	-0.0100	0.9924
CG13516	CG13516	1638764_at	-0.1578	0.3823	0.4149	0.0534	0.3520	0.0609	-0.0693	0.9117	-0.2985	0.1320	-0.2293	0.1965	0.0859	0.8972	0.0948	0.7439	0.0090	0.9835
CG7818	CG7818	1638765_at	-0.0420	0.7857	0.1796	0.4246	0.2235	0.1806	0.0438	0.9311	0.2157	0.1437	0.1719	0.1937	-0.0926	0.9400	0.3043	0.4166	0.3969	0.3032
CG34006	CG34006	1638766_at	0.1000	0.5547	0.1016	0.4800	0.0155	0.9473	0.0021	0.9973	-0.0642	0.7542	-0.0663	0.7171	0.0504	0.9515	-0.0953	0.7459	-0.1457	0.5853
CG15059	CG15059	1638767_at	0.2544	0.1969	0.3247	0.0638	0.1096	0.4999	-0.0303	0.9745	-0.0486	0.8751	-0.0182	0.9505	0.2837	0.6955	0.0247	0.9613	-0.2590	0.4019
Cpr11A	CG2560	1638768_at	0.2013	0.2271	-0.0043	0.9710	-0.0168	0.9263	-0.0178	0.9757	0.1375	0.3442	0.1553	0.2250	-0.1937	0.6483	-0.1651	0.3206	0.0286	0.9031
CG15059	CG15059	1638769_a_at	-0.0433	0.7971	-0.1478	0.4755	-0.1588	0.3558	0.0187	0.9777	0.0110	0.9674	-0.0077	0.9711	0.0085	0.9928	-0.1085	0.6497	-0.1170	0.6216
cenB1A	centaurin beta 1A	1638770_a_at	-0.0822	0.6042	-0.0127	0.9673	0.0427	0.8757	-0.0198	0.9819	0.3051	0.1379	0.3249	0.0800	-0.1564	0.8760	0.3644	0.3397	0.5208	0.2056
---	---	1638771_at	-0.0828	0.6647	0.0557	0.6822	0.0555	0.7965	-0.1391	0.7582	-0.1758	0.3760	-0.0367	0.8729	-0.1584	0.8378	-0.0271	0.9590	0.1313	0.7148

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG18107	CG18107	1638772_at	2.1361	0.0005	1.5314	0.1841	2.7724	0.0000	0.4042	0.4341	-0.1466	0.6459	-0.5508	0.0368	-0.5439	0.8157	-0.7365	0.4446	-0.1926	0.8842
CG31353	CG31353	1638773_at	0.0634	0.7563	-0.0639	0.6518	0.1279	0.4820	0.1434	0.6539	0.1286	0.4195	-0.0148	0.9386	-0.1282	0.8704	-0.0629	0.8910	0.0653	0.8779
CG16904	CG16904	1638774_at	0.4944	0.8142	-2.2705	0.4810	1.8587	0.1063	2.2131	0.3739	0.1914	0.9223	-2.0217	0.1064	-1.9888	0.8270	-2.4593	0.5303	-0.4705	0.9285
Mdr50	Multi drug resistar	1638775_at	-0.1272	0.9578	0.1556	0.1651	-0.1659	0.6552	-0.5712	0.8794	-1.1618	0.3781	-0.5906	0.6585	-0.1620	0.9831	-1.0038	0.5994	-0.8418	0.6659
---	---	1638776_at	0.1012	0.4957	0.0781	0.5170	0.0082	0.9756	-0.0670	0.9083	-0.0618	0.7845	0.0052	0.9825	-0.0039	0.9964	-0.0288	0.9325	-0.0249	0.9333
Rim	Rim	1638777_at	-0.0449	0.8530	0.2400	0.1852	0.1096	0.5867	-0.0927	0.8732	-0.1325	0.5392	-0.0398	0.8656	0.1188	0.8292	0.2001	0.3761	0.0814	0.7608
CG10352	CG10352	1638778_at	0.4964	0.3787	0.0843	0.9318	0.3675	0.1421	0.1235	0.7003	0.1805	0.2195	0.0570	0.7162	-0.0934	0.9841	-0.0781	0.9647	0.0153	0.9935
---	---	1638779_at	0.4364	0.1986	0.3703	0.1777	0.0128	0.9607	-0.4633	0.3048	0.2997	0.2605	0.7630	0.0068	-0.1764	0.8802	0.2245	0.6437	0.4009	0.3813
GluRIIE	Glutamate recepto	1638780_at	-2.1387	0.0010	-2.4734	0.0176	-2.9027	0.0000	-0.0029	0.9961	-0.1213	0.5441	-0.1184	0.5098	0.2493	0.9030	-0.4692	0.5464	-0.7185	0.3485
CG31913	CG31913	1638781_at	0.1241	0.5890	-0.0581	0.6130	-0.1853	0.5391	-0.1101	0.8578	-0.0095	0.9781	0.1006	0.6615	-0.0916	0.9142	-0.0383	0.9358	0.0534	0.8962
CG8195	CG8195	1638782_at	0.0290	0.8979	0.0152	0.9659	-0.0670	0.7201	-0.2197	0.5233	0.2080	0.2501	0.4277	0.0161	-0.1167	0.8846	0.1996	0.5272	0.3163	0.3089
mRpL48	mitochondrial ribo	1638783_at	0.0175	0.9393	0.7487	0.0295	0.4946	0.0419	-0.0284	0.9757	-0.3990	0.0821	-0.3706	0.0719	0.2065	0.7596	0.3512	0.1995	0.1447	0.6350
---	---	1638784_s_at	0.0456	0.8119	-0.0650	0.6518	0.2647	0.1376	0.0614	0.9445	-0.0043	0.9903	-0.0657	0.8178	-0.1788	0.7266	-0.1103	0.6449	0.0684	0.7978
CG14585	CG14585	1638785_at	-0.0623	0.7232	0.2975	0.0616	0.2476	0.2322	0.0960	0.8159	-0.1974	0.2216	-0.2934	0.0480	0.1164	0.8795	0.0923	0.8080	-0.0242	0.9535
CG13036	CG13036	1638786_at	0.1174	0.6598	0.5097	0.3004	0.5027	0.0160	-0.0064	0.9951	-0.2994	0.1578	-0.2930	0.1222	0.0069	0.9956	0.1075	0.7618	0.1006	0.7724
CG40207	CG40207	1638787_at	-0.0746	0.7658	-0.1673	0.4479	-0.4055	0.0498	-0.1417	0.8248	0.2809	0.2589	0.4226	0.0604	-0.2248	0.8262	0.1780	0.7105	0.4028	0.3552
CG32574	CG32574	1638788_at	0.1639	0.3546	-0.0733	0.6506	0.2238	0.2011	0.1564	0.6238	-0.0619	0.7399	-0.2184	0.1211	0.1095	0.8837	-0.1418	0.6458	-0.2512	0.3870
fau	fau	1638789_at	-1.8982	0.0134	-1.6439	0.0775	-2.8560	0.0019	-0.6037	0.3206	-1.2198	0.0050	-0.6161	0.0558	0.6824	0.8270	-1.0053	0.4336	-1.6877	0.2094
CG31386	CG31386	1638790_at	0.1709	0.3602	-0.0523	0.8365	-0.0275	0.8701	0.0052	0.9956	0.2202	0.2422	0.2149	0.2003	-0.0394	0.9737	-0.1118	0.7457	-0.0724	0.8474
---	---	1638791_at	0.2346	0.2056	-0.0012	0.9970	0.0882	0.5539	0.3055	0.3443	0.5173	0.0131	0.2117	0.2024	0.0463	0.9589	0.1776	0.4931	0.1313	0.6296
CG32098	CG32098	1638792_at	0.1407	0.5447	-0.1300	0.4524	0.1089	0.6412	0.1989	0.6591	0.2404	0.2695	0.0414	0.8713	0.0945	0.9304	0.1661	0.6658	0.0716	0.8800
CG31406	CG31406	1638793_at	0.1560	0.3048	-0.0020	0.9882	0.2239	0.2739	0.0717	0.8500	-0.0053	0.9801	-0.0770	0.5755	-0.2069	0.7464	-0.1110	0.7200	0.0960	0.7577
CG32247	CG32247	1638794_at	0.1596	0.2657	0.0664	0.5662	0.0393	0.8362	-0.1316	0.7278	0.0048	0.9848	0.1364	0.3853	0.0105	0.9913	-0.0570	0.8307	-0.0676	0.7758
sec6	sec6	1638795_at	0.4189	0.0720	0.5709	0.1748	0.4217	0.0165	0.1056	0.8180	0.4508	0.0204	0.3452	0.0383	0.3487	0.7154	0.5385	0.1656	0.1898	0.6615
CG6195	CG6195	1638796_at	0.0715	0.6908	1.1116	0.0085	1.0202	0.0002	0.1904	0.5943	-0.2954	0.1027	-0.4858	0.0094	0.3569	0.3712	0.8585	0.0108	0.5016	0.0559
---	---	1638797_a_at	-0.1028	0.5444	-0.4664	0.0496	-0.0815	0.7467	0.2410	0.4024	-0.1819	0.2598	-0.4228	0.0102	-0.1415	0.8906	-0.5580	0.1420	-0.4164	0.2964
---	---	1638798_at	0.2521	0.2393	-0.1830	0.2362	0.0282	0.8604	0.2452	0.5369	0.2729	0.1856	0.0277	0.9124	-0.0365	0.9717	-0.1559	0.5644	-0.1194	0.6719
---	---	1638799_at	0.2241	0.3491	-0.4062	0.2568	-0.2606	0.3695	0.1914	0.7850	0.5629	0.0560	0.3715	0.1493	-0.1517	0.8157	-0.2108	0.4323	-0.0590	0.8694
CG14154	CG14154	1638800_a_at	-0.6354	0.0334	-0.3288	0.1161	-0.8119	0.0015	-0.1316	0.8473	-0.1884	0.4723	-0.0567	0.8445	-0.0082	0.9939	-0.0948	0.7439	-0.0866	0.7626
CG14229	CG14229	1638801_at	0.0937	0.6003	-0.1285	0.3679	-0.0455	0.8069	0.1007	0.7664	0.4478	0.0081	0.3471	0.0153	-0.0091	0.9939	0.1067	0.7248	0.1159	0.6906
CG31415	CG31415	1638802_at	0.1851	0.4364	0.2840	0.2669	0.2435	0.5401	0.0525	0.9446	-0.2280	0.2963	-0.2806	0.1456	0.1697	0.8858	0.0105	0.9924	-0.1592	0.7562
Rnp4F	RNA-binding prote	1638803_at	-0.0552	0.8856	0.1677	0.3913	0.3980	0.0842	-0.0387	0.9584	0.0959	0.6658	0.1345	0.4683	-0.2347	0.8192	0.2415	0.5888	0.4762	0.2750
Stlk	Ste20-like kinase	1638804_a_at	-0.6383	0.0602	0.5672	0.0137	0.4706	0.0865	0.0741	0.9154	-0.6320	0.0110	-0.7061	0.0040	0.1650	0.8940	0.5091	0.2501	0.3441	0.4683
CG10068 /// DyakCG10068	CG10068	1638805_at	-0.0156	0.9660	-0.1087	0.4129	-0.2550	0.1168	0.1865	0.6506	0.3594	0.0730	0.1729	0.3420	0.1539	0.7464	0.1102	0.6102	-0.0437	0.8721
Rbf2	Retinoblastoma-fa	1638806_at	0.1275	0.7097	-0.3902	0.4103	-0.9004	0.0060	-0.7941	0.1787	0.5648	0.1141	1.3589	0.0020	-0.1863	0.9467	0.0354	0.9828	0.2216	0.8372
CG4829	CG4829	1638807_s_at	-1.3089	0.0023	-1.4811	0.0191	-1.4129	0.0007	-0.0073	0.9953	-0.4084	0.1043	-0.4011	0.0768	-0.1414	0.8837	-0.5357	0.1395	-0.3943	0.2964
Syx4	syntaxin	1638808_at	-1.0529	0.0175	-1.6043	0.0094	-1.6367	0.0001	-0.1924	0.7235	-0.3476	0.1548	-0.1552	0.5131	-0.0813	0.9711	-0.8712	0.1250	-0.7899	0.1867
---	---	1638809_at	-0.8288	0.2947	-0.0704	0.7043	-0.4101	0.0152	0.1463	0.9573	-0.9607	0.1858	-1.1069	0.0901	0.4394	0.7230	0.1121	0.8839	-0.3273	0.5612
Faa	Fumarylacetoacet	1638810_at	0.3726	0.1997	0.5185	0.0458	0.9073	0.0116	0.2448	0.7121	-0.5355	0.0773	-0.7802	0.0110	0.0542	0.9717	-0.2027	0.6156	-0.2569	0.5140
Sug	Sugar-baby	1638811_at	-0.0275	0.9393	-0.1122	0.7582	0.1885	0.3965	-0.2084	0.6533	-0.0636	0.8193	0.1448	0.5002	-0.2699	0.8424	0.0873	0.9193	0.3572	0.5421
CG11501	CG11501	1638812_at	0.7918	0.0298	-0.1930	0.6502	0.0604	0.9645	0.2606	0.9488	1.3007	0.2518	1.0401	0.3123	-0.1080	0.9776	0.1984	0.8807	0.3064	0.7777
CG15499	CG15499	1638813_at	0.2744	0.3383	0.1820	0.3486	0.1563	0.3901	-0.0087	0.9937	0.1492	0.5527	0.1580	0.4772	0.0819	0.9296	-0.0505	0.9125	-0.1324	0.6886
---	---	1638814_at	0.1211	0.5563	0.1319	0.6870	0.0767	0.7799	-0.2257	0.7073	-0.0075	0.9855	0.2182	0.3928	0.0620	0.9618	0.1890	0.6270	0.1270	0.7614
CG14850	CG14850	1638815_at	0.0737	0.6305	0.1337	0.4351	0.2076	0.1861	0.0742	0.8815	0.0212	0.9292	-0.0530	0.7781	0.0246	0.9816	0.1051	0.7064	0.0806	0.7835
CG3884	CG3884	1638816_at	0.0960	0.8769	0.2789	0.4047	0.7091	0.0865	0.1506	0.9208	-0.4815	0.2995	-0.6322	0.1224	-0.1694	0.9365	-0.0967	0.9275	0.0727	0.9382
CG15594	CG15594	1638817_at	0.0594	0.6812	0.0077	0.9801	0.3035	0.1476	0.0330	0.9621	-0.0326	0.8973	-0.0655	0.7415	-0.0386	0.9717	0.0488	0.9075	0.0874	0.7894
Mlp60A	Muscle-specific LI	1638818_at	-2.0036	0.0588	-2.4963	0.0314	-2.8081	0.0000	-0.4096	0.1658	-0.4961	0.0134	-0.0865	0.6382	-0.1090	0.9862	-1.0068	0.4870	-0.8978	0.5488
CG31712	CG31712	1638819_at	-0.5967	0.0339	-0.3034	0.3579	-0.4404	0.0810	0.0083	0.9909	0.1160	0.4852	0.1077	0.4755	0.0186	0.9929	0.2948	0.5749	0.2762	0.6052
CG34362	CG12870	1638820_at	0.0252	0.8856	-0.1331	0.2445	-0.1043	0.6520	-0.0463	0.9426	-0.1226	0.5306	-0.0762	0.6918	-0.0481	0.9515	-0.1746	0.4703	-0.1265	0.6225
yuri	yuri gagarin	1638821_a_at	-0.0316	0.9144	-0.7492	0.0100	-1.2642	0.0068	-0.0472	0.9466	0.3513	0.0795	0.3985	0.0326	0.4211	0.7485	-0.3847	0.5064	-0.8058	0.1752
CG6859	CG6859	1638822_at	0.5197	0.0419	1.5116	0.0101	1.7507	0.0000	-0.0533	0.9529	-1.0092	0.0025	-0.9559	0.0019	-0.2329	0.7305	-0.0281	0.9519	0.2048	0.4944
CG30259 /// DmirCG30259	CG30259	1638823_at	0.2948	0.1069	0.1875	0.3992	0.1829	0.2159	0.0062	0.9937	-0.1315	0.4236	-0.1378	0.3448	0.0283	0.9742	-0.0096	0.9812	-0.0379	0.8991
nonA	no-on-transient A	1638824_a_at	-0.3012	0.3419	0.1835	0.6242	-0.0734	0.7445	-0.1925	0.7042	-0.3562	0.1252	-0.1637	0.4583	0.0899	0.9589	0.2264	0.6868	0.1365	0.8289
CG16813	CG16813	1638825_at	0.0678	0.6748	0.0404	0.8770	-0.1084	0.6491	-0.0101	0.9883	-0.0324	0.8810	-0.0223	0.9095	-0.0001	0.9999	-0.0340	0.9425	-0.0339	0.9342
---	---	1638826_at	0.2468	0.4914	0.0761	0.6715	0.2602	0.2196	-0.0465	0.9621	-0.1827	0.5062	-0.1362	0.6034	-0.0103	0.9946	-0.1870	0.6270	-0.1767	0.6458
CG3640	CG3640	1638827_at	0.1523	0.3420	0.1851	0.3704	0.0256	0.8981	-0.1482	0.6936	0.0901	0.6445	0.2383	0.1236	0.1615	0.7644	0.3270	0.1418	0.1656	0.4815
CG17378	CG17378	1638828_a_at	-0.1538	0.3572	0.0370	0.8311	0.1613	0.5192	-0.0429	0.9506	-0.0879	0.6880	-0.0450	0.8376	-0.2871	0.6955	0.0163	0.9759	0.3033	0.3343
mod(mdg4)	Modifier67.2	1638829_s_at	0.0974	0.5806	0.0033	0.9950	-0.3601	0.0580	-0.2154	0.5793	0.1300	0.5428	0.3454	0.0560	0.0930	0.9137	-0.0441	0.9263	-0.1372	0.6763
alien	d-Alien	1638830_at	-0.1399	0.4544	0.3074	0.2935	0.2190	0.2018	0.0303	0.9704	-0.1350	0.5461	-0.1653	0.3922	0.1480	0.8454	0.3914	0.1948	0.2434	0.4478
---	---	1638831_at	0.0816	0.7445	0.0798	0.6313	-0.0053	0.9826	-0.1107	0.8578	-0.0304	0.9223	0.0803	0.7379	-0.0430	0.9775	0.0768	0.8813	0.1198	0.7764
RpS23	Ribosomal protein	1638832_at	0.2265	0.1344	0.3452	0.0444	0.3212	0.0521	0.0362	0.9436	0.0054	0.9804	-0.0308	0.8551	0.0407	0.9587	0.0319	0.9264	-0.0087	0.9809
FucT6	FucT6	1638833_at	0.7454	0.0368	0.5879	0.0730	0.4075	0.0183	-0.0560	0.9163	0.5900	0.0044	0.6460	0.0018	0.0724	0.9710	0.4411	0.3698	0.3687	0.4764
MED18	Mediator complex	1638834_a_at	0.1870	0.4558	0.3296	0.1648	0.4950	0.0143	-0.0041	0.9956	-0.0707	0.6925	-0.0665	0.6828	0.0337	0.9816	0.1131	0.7870	0.0794	0.8555
---	---	1638835_at	0.0791	0.5905	-0.0973	0.6348	0.1544	0.4306	0.1058	0.8310	0.1975	0.3013	0.0917	0.6360	-0.0590	0.9365	-0.0181	0.9628	0.0408	0.8994
CG34133	CG34133	1638836_at	0.1912	0.2371	0.2127	0.0986	0.3914	0.0573	0.0835	0.8903	0.0309	0.9133	-0.0526	0.8191	-0.0942	0.8767	0.0030	0.9956	0.0971	0.7154
---	---	1638837_a_at	0.2155	0.2427	-0.3762	0.3503	-0.2549	0.2405	0.0612	0.9528	0.7084	0.0213	0.6472	0.0194	0.2071	0.8270	0.2737	0.5046	0.0666	0.9031
CG13030 /// DereCG13030	CG13030 /// GA1	1638838_at	0.1329	0.5969	0.2228	0.3793	0.0131	0.9676	0.0145	0.9922	-0.0464	0.9143	-0.0609	0.8656	0.1787	0.8016	0.0007	0.9996	-0.1780	0.5722
CG8092	CG8092	1638839_at	0.1571	0.5444	0.1695	0.5917	0.1481	0.5491	0.2387	0.6046	0.3582	0.1214	0.1195	0.6111	0.1987	0.8465	0.2451	0.5878	0.0464	0.9380
CG14641	CG14641	1638840_at	-0.2526	0.2104	-0.1744	0.5757	-0.0936	0.5991	0.1052	0.7579	0.2362	0.1063	0.1309	0.3283	-0.0388	0.9829	0.2580	0.5501	0.2967	0.4865

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
eIF2B-gamma	eIF2B-gamma	1638841_at	-0.5166	0.0389	0.1495	0.6948	0.1336	0.3964	-0.1593	0.6084	-0.5696	0.0039	-0.4103	0.0102	-0.1089	0.9400	0.1027	0.8716	0.2116	0.6607
---	---	1638842_at	0.0555	0.8112	0.0562	0.6028	-0.0123	0.9524	0.1133	0.8327	0.1499	0.4817	0.0366	0.8795	0.1863	0.7464	0.0329	0.9333	-0.1534	0.5532
---	---	1638843_at	0.0573	0.8033	-0.0003	1.0000	-0.0197	0.9550	-0.0154	0.9893	0.0084	0.9826	0.0238	0.9418	-0.0414	0.9814	-0.0250	0.9682	0.0164	0.9784
CG3714	CG3714	1638844_s_at	-0.1660	0.2711	0.0927	0.7272	0.5713	0.0087	-0.2785	0.3056	-0.6424	0.0026	-0.3639	0.0191	-0.6029	0.3712	-0.4128	0.2430	0.1901	0.6311
CG18643	CG18643	1638845_at	-0.3260	0.1003	0.9362	0.0157	0.6780	0.0264	-0.3387	0.5811	-1.0553	0.0056	-0.7166	0.0187	-0.0180	0.9923	0.2663	0.5523	0.2843	0.5271
CG3402	CG3402	1638846_at	-0.0442	0.8827	-0.7431	0.0621	-0.9219	0.0009	-0.0474	0.9314	0.5060	0.0084	0.5534	0.0033	0.2206	0.8331	-0.1697	0.7408	-0.3903	0.3825
sta	Stubarista	1638847_s_at	0.1124	0.4866	0.6864	0.0473	1.1410	0.0004	0.2868	0.4739	-0.4961	0.0289	-0.7829	0.0024	-0.0309	0.9808	0.0278	0.9505	0.0587	0.8761
CG6241	CG6241	1638848_at	0.6544	0.0094	0.1967	0.2073	0.4458	0.0262	-0.2228	0.5470	0.2752	0.1491	0.4980	0.0108	-0.4307	0.3712	-0.0810	0.8106	0.3497	0.2031
CG2875	CG2875	1638849_a_at	0.8155	0.0216	0.2327	0.5728	0.4412	0.0462	0.1605	0.8196	0.8399	0.0088	0.6793	0.0138	-0.1006	0.9474	0.1147	0.8584	0.2153	0.6717
---	---	1638850_at	0.0075	0.9732	0.0717	0.5124	0.1029	0.6298	-0.0770	0.9205	-0.1958	0.4191	-0.1189	0.6176	0.0461	0.9677	0.0461	0.9225	-0.0001	0.9999
CG5280	CG5280	1638851_at	-0.2580	0.1743	-0.0161	0.8976	0.3705	0.1026	-0.1987	0.5931	-0.3537	0.0640	-0.1550	0.3735	-0.3894	0.5869	0.0927	0.8248	0.4822	0.1562
CHKov2	CHKov2	1638852_at	-2.0594	0.0074	-2.0581	0.0142	-2.0592	0.0000	-0.2999	0.5664	-0.4915	0.0697	-0.1915	0.4486	-0.2996	0.8692	-0.4046	0.6012	-0.1050	0.9169
CG32971	CG32971	1638853_at	0.1332	0.5595	-0.1161	0.4696	0.0196	0.9042	0.0656	0.9297	0.1697	0.4536	0.1041	0.6413	0.0203	0.9848	-0.0112	0.9797	-0.0315	0.9249
CG17667 /// DlitCG17667	CG17667	1638854_s_at	-0.7647	0.0080	-1.2838	0.0341	-1.3836	0.0004	0.0067	0.9951	0.5093	0.0260	0.5026	0.0172	0.0044	0.9984	-0.1064	0.8537	-0.1107	0.8348
---	---	1638855_at	-0.0542	0.8024	-0.0022	0.9907	-0.2675	0.1519	-0.1917	0.6202	0.0249	0.9265	0.2166	0.2091	0.1007	0.8940	0.0212	0.9634	-0.0795	0.8209
CG2650 /// DsimCG2650	0.9kb transcript ///	1638856_at	0.2656	0.1417	0.1857	0.4530	0.1685	0.4440	-0.0136	0.9894	-0.0211	0.9528	-0.0074	0.9799	-0.1647	0.8122	-0.1831	0.5404	-0.0183	0.9659
CG3605	CG3605	1638857_at	0.0690	0.7828	-0.0946	0.7623	-0.1297	0.5231	-0.1243	0.8251	0.4126	0.0622	0.5369	0.0137	-0.1226	0.9238	0.2110	0.6473	0.3336	0.4457
CG8934	CG8934	1638858_at	-0.3563	0.5106	0.0236	0.8641	0.3583	0.0318	0.2296	0.8578	-0.4781	0.3028	-0.7077	0.0855	-0.0432	0.9514	0.0136	0.9690	0.0568	0.8321
CG14741	CG14741	1638859_at	0.6622	0.1036	0.6113	0.0535	0.8300	0.0009	-0.1896	0.7732	0.0176	0.9675	0.2073	0.4228	-0.1136	0.9333	0.3199	0.4677	0.4336	0.3293
---	---	1638860_s_at	0.2199	0.3248	0.2265	0.3560	0.0782	0.6638	-0.0215	0.9857	-0.0738	0.8294	-0.0523	0.8685	0.1437	0.8424	-0.0589	0.8929	-0.2026	0.5115
CG32457	CG32457	1638861_at	-0.0608	0.8367	-0.2472	0.0555	-0.3843	0.0443	-0.1542	0.8162	0.1532	0.5887	0.3074	0.1860	-0.1154	0.8454	-0.0348	0.9277	0.0806	0.7787
CG15553	CG15553	1638862_at	2.5278	0.0041	0.6916	0.4402	1.5170	0.0093	-0.1702	0.8897	0.2084	0.6525	0.3785	0.3110	-1.1325	0.6898	-1.4814	0.1874	-0.3489	0.8048
---	---	1638863_at	0.1421	0.4847	0.0216	0.8447	0.1816	0.2867	0.2494	0.4128	0.0412	0.8510	-0.2082	0.1666	-0.0388	0.9657	0.0532	0.8782	0.0920	0.7422
Rab10	Rab-protein 10	1638864_at	0.1341	0.3448	0.0382	0.7796	-0.0957	0.6436	-0.1617	0.6650	0.2762	0.1196	0.4379	0.0139	-0.0220	0.9848	0.1107	0.7065	0.1326	0.6366
---	---	1638865_x_at	0.1019	0.6705	-0.1302	0.6625	-0.0370	0.9183	0.1021	0.9228	0.0191	0.9693	-0.0830	0.8178	0.0127	0.9939	-0.2094	0.6093	-0.2221	0.5869
CG9542	CG9542	1638866_at	0.3378	0.1560	0.3525	0.2161	0.2095	0.3959	0.1668	0.8296	-0.4973	0.0946	-0.6641	0.0207	0.1299	0.8940	-0.3793	0.2825	-0.5092	0.1880
CG16800	CG16800	1638867_at	-0.0861	0.7433	-0.1335	0.4435	-0.2675	0.1937	0.2645	0.5664	0.2392	0.3165	-0.0252	0.9306	0.1859	0.8461	0.0509	0.9353	-0.1350	0.7704
Mcm10	Sensitized chrom	1638868_at	0.2392	0.5571	-0.8063	0.2731	-0.9443	0.0554	-0.3053	0.4908	1.3237	0.0007	1.6290	0.0002	-0.2594	0.9516	0.0491	0.9842	0.3085	0.8540
Cpr51A	CG10112	1638869_at	-2.5944	0.0017	-4.3398	0.0011	-3.9803	0.0000	0.7369	0.3773	1.2484	0.0163	0.5114	0.2262	0.2931	0.8202	-0.3001	0.5959	-0.5933	0.2798
CG1958	CG1958	1638870_at	0.1320	0.8225	-0.4065	0.3629	-0.5930	0.1464	-0.0209	0.9922	0.5450	0.2093	0.5659	0.1451	-0.1020	0.9717	-0.0877	0.9404	0.0143	0.9909
SCAR	SCAR	1638871_at	-0.1945	0.6692	0.0334	0.8535	-0.4476	0.0808	0.0622	0.9228	0.1404	0.4901	0.0781	0.7025	0.4708	0.7697	0.3916	0.5897	-0.0792	0.9343
Hsp68	Heat shock protei	1638872_at	-1.4411	0.3042	-0.5516	0.4368	-0.5766	0.5628	0.4805	0.8671	1.0515	0.3043	0.5710	0.5701	0.4757	0.9411	1.6559	0.4012	1.1802	0.5781
CG6175	CG6175	1638873_at	-0.4442	0.3938	0.2857	0.3956	0.0965	0.7772	0.0659	0.9412	-0.3342	0.1890	-0.4002	0.0815	0.2341	0.9032	0.2194	0.8083	-0.0148	0.9902
CG4585	undefined 2	1638874_at	-0.0918	0.5802	-0.0250	0.8777	0.1262	0.6147	0.0242	0.9672	0.0533	0.7751	0.0292	0.8733	-0.0672	0.9588	0.2157	0.5715	0.2829	0.4475
EDTP	Egg-derived tyros	1638875_at	0.6886	0.0081	0.2115	0.1518	0.5170	0.0573	-0.0663	0.9263	-0.3953	0.0699	-0.3290	0.0882	-0.1103	0.9260	-0.7592	0.0711	-0.6489	0.1313
Rpn2	Rpn2	1638876_at	0.0507	0.8503	0.1508	0.2753	0.8054	0.0023	0.2944	0.2655	0.1067	0.5214	-0.1877	0.1787	-0.2848	0.7196	0.3177	0.3225	0.6025	0.1013
AdoR	Adenosine recept	1638877_at	-0.4023	0.3060	0.1576	0.5916	0.1988	0.3560	0.0792	0.9205	-0.4321	0.0758	-0.5113	0.0260	-0.2095	0.8732	-0.0110	0.9925	0.1985	0.7392
CG17302	CG17302	1638878_at	0.0789	0.5983	-0.0900	0.5019	0.1379	0.5402	-0.0067	0.9937	-0.0114	0.9638	-0.0047	0.9816	-0.0319	0.9746	-0.1822	0.4743	-0.1503	0.5720
CG33012	CG33012	1638879_a_at	0.2866	0.6863	0.1199	0.3916	-0.4117	0.0398	-0.7269	0.2067	-0.2253	0.5458	0.5017	0.1046	-0.0567	0.9829	-0.2206	0.7599	-0.1639	0.8288
CG32220	CG32220	1638880_at	0.1580	0.4720	0.0748	0.5431	0.1400	0.4214	0.0163	0.9812	0.0454	0.8308	0.0290	0.8831	-0.1932	0.8191	-0.0591	0.9111	0.1341	0.7350
CG9550	CG9550	1638881_at	0.2639	0.1179	0.2138	0.3448	0.3688	0.1016	0.0649	0.9314	0.0679	0.8057	0.0030	0.9915	-0.1392	0.8655	0.0325	0.9507	0.1716	0.6279
ry	Xanthine DH	1638882_at	0.0998	0.8224	-0.7983	0.4189	-0.1585	0.4999	0.0268	0.9777	0.5671	0.0291	0.5403	0.0225	-0.5586	0.8114	-0.2444	0.8546	0.3142	0.7847
CG8636	CG8636	1638883_at	0.1643	0.2521	0.2501	0.1392	0.3992	0.0199	0.1614	0.5912	-0.0441	0.8147	-0.2055	0.1293	0.0364	0.9647	-0.0357	0.9170	-0.0721	0.7838
CG4238	CG4238	1638884_at	0.3258	0.2152	0.7539	0.0811	0.8769	0.0123	0.1213	0.8822	-0.1322	0.6790	-0.2535	0.3209	0.0180	0.9914	0.2934	0.4141	0.2754	0.4620
---	---	1638885_at	1.5164	0.0054	1.5656	0.0108	2.2018	0.0001	0.3831	0.1642	0.3555	0.0396	-0.0277	0.8898	-0.0970	0.9677	0.3973	0.5501	0.4944	0.4504
CG8319 /// DwilCG8319	CG8319	1638886_at	0.1627	0.5934	-0.1237	0.5573	-0.2030	0.1811	0.0017	0.9986	0.3653	0.0759	0.3636	0.0511	0.0219	0.9913	0.0372	0.9557	0.0153	0.9832
rad50	rad50	1638887_a_at	-0.0913	0.6943	-0.7225	0.0419	-0.3475	0.1594	-0.2662	0.3030	-0.3788	0.0209	-0.1126	0.4294	-0.6403	0.3985	-0.9371	0.0449	-0.2968	0.4867
unpg	unplugged	1638888_at	0.6764	0.1714	0.4416	0.2880	0.5993	0.0036	-0.0642	0.9288	-0.0602	0.8193	0.0041	0.9875	0.1022	0.9611	-0.2710	0.6789	-0.3732	0.5488
CG3918	CG3918	1638889_at	-0.1008	0.4950	-0.0450	0.7524	0.0436	0.8309	0.0240	0.9761	-0.2174	0.2775	-0.2414	0.1743	-0.1612	0.7506	-0.1270	0.5825	0.0342	0.9095
Tapdelta	Translocon-assoc	1638890_at	0.6701	0.0150	1.6088	0.0074	1.9242	0.0000	0.0587	0.9057	-0.3187	0.0514	-0.3775	0.0165	-0.1931	0.8395	0.6429	0.1077	0.8360	0.0710
CG31031	CG31031	1638891_at	-0.0777	0.6623	-0.1174	0.3721	-0.2594	0.2752	-0.0959	0.9029	0.0623	0.8506	0.1583	0.5235	-0.0726	0.8723	-0.0827	0.6765	-0.0101	0.9706
CG5397	CG5397	1638892_at	-1.7972	0.0015	-0.7287	0.2748	-1.3188	0.0062	-0.3561	0.5778	-1.4299	0.0018	-1.0738	0.0038	0.0433	0.9898	-0.2787	0.7439	-0.3220	0.6877
CG7029	CG7029	1638893_at	0.3909	0.5769	0.8662	0.1648	0.5912	0.0632	-0.0252	0.9888	-0.4722	0.2578	-0.4470	0.2299	0.3239	0.9011	0.2624	0.8380	-0.0614	0.9659
---	---	1638894_at	-0.0978	0.6404	-0.1040	0.5517	0.1495	0.3132	0.2454	0.5769	0.0065	0.9845	-0.2390	0.2372	0.0163	0.9913	0.0994	0.7963	0.0830	0.8287
---	---	1638895_at	0.0787	0.7347	0.0941	0.4843	0.2256	0.1985	-0.0539	0.9507	0.0929	0.7468	0.1467	0.5351	-0.0165	0.9877	0.1516	0.5469	0.1681	0.5007
Cpr62Bc	CG1919	1638896_at	1.0038	0.0027	0.6100	0.1784	0.6263	0.0086	0.0445	0.9627	0.0851	0.7803	0.0407	0.8928	0.3878	0.7293	-0.0317	0.9682	-0.4196	0.3921
CG3511	CG3511	1638897_at	-0.3312	0.0520	0.3572	0.2749	0.6009	0.0082	0.0242	0.9738	-0.4678	0.0176	-0.4920	0.0086	-0.1288	0.8534	0.2550	0.3667	0.3838	0.2042
CG13197	CG13197	1638898_at	0.1630	0.4897	-0.1422	0.4810	-0.2322	0.2647	-0.0579	0.9432	0.1185	0.6462	0.1764	0.4108	0.0265	0.9848	-0.1338	0.7124	-0.1603	0.6410
CG15312	CG15312	1638899_s_at	-0.0288	0.9422	-0.8760	0.0412	-0.4364	0.0444	0.0669	0.9375	0.7477	0.0093	0.6808	0.0085	-0.1408	0.9235	0.1213	0.8550	0.2621	0.6149
---	---	1638900_at	0.0248	0.8987	0.0463	0.6370	0.1607	0.3127	-0.0309	0.9592	-0.1337	0.4131	-0.1028	0.4998	0.0450	0.9411	-0.0684	0.7642	-0.1134	0.5720
CG10014	CG10014	1638901_at	0.1180	0.6017	-0.1219	0.5103	-0.0133	0.9539	-0.0504	0.9346	-0.0092	0.9743	0.0412	0.8444	-0.2039	0.8400	-0.2477	0.5724	-0.0438	0.9402
Or47a	Olfactory receptor	1638902_at	-4.5474	0.0017	-7.1177	0.0007	-6.1896	0.0000	0.6550	0.6327	1.7790	0.0162	1.1240	0.0645	-0.1663	0.9626	-0.7879	0.4200	-0.6216	0.5488
CG5724	CG5724	1638903_at	0.0454	0.8244	0.0242	0.8521	-0.0004	0.9981	-0.2164	0.6197	-0.2411	0.2688	-0.0248	0.9257	-0.1792	0.7464	-0.1472	0.5488	0.0320	0.9211
---	---	1638904_at	-0.1502	0.3854	-0.0861	0.4566	0.0957	0.7325	0.0859	0.9068	-0.0157	0.9657	-0.1016	0.6814	-0.2194	0.7220	0.0041	0.9941	0.2235	0.4058
CG4839	PKG-like	1638905_s_at	-3.2430	0.0004	-2.9112	0.0021	-2.5712	0.0000	-0.1942	0.7774	0.1655	0.5965	0.3598	0.1594	-0.4985	0.4207	0.4127	0.1957	0.9113	0.0410
CG32052	CG32052	1638906_at	0.0575	0.7311	0.0268	0.8058	0.2872	0.1206	0.0859	0.8327	0.0341	0.8673	-0.0518	0.7589	-0.2096	0.6749	-0.1277	0.5531	0.0820	0.7269
---	---	1638907_at	0.0306	0.9008	0.0213	0.8669	-0.1124	0.6801	-0.0549	0.9529	0.1009	0.7388	0.1558	0.5340	-0.0454	0.9576	-0.0208	0.9587	0.0246	0.9425
CG15635	CG15635	1638908_at	-0.0992	0.5312	-0.1904	0.4124	-0.2370	0.1495	0.0415	0.9558	0.0753	0.7553	0.0338	0.8889	0.1897	0.7644	0.1627	0.5620	-0.0270	0.9431
---	---	1638909_at	0.2571	0.2051	0.1035	0.4659	0.0536	0.8260	-0.0336	0.9649	0.0335	0.9045	0.0671	0.7623	-0.0999	0.8960	-0.1076	0.7484	-0.0078	0.9863

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG5561	CG5561	1638910_at	0.5075	0.0382	0.2221	0.1301	0.3720	0.0310	0.1516	0.6354	0.0333	0.8729	-0.1183	0.4214	-0.0020	0.9992	-0.1393	0.6425	-0.1374	0.6449
CG14551	CG14551	1638911_at	0.2505	0.2103	-0.0797	0.5731	0.0334	0.8531	0.0243	0.9744	0.1068	0.5920	0.0825	0.6636	0.1159	0.8427	-0.2234	0.3397	-0.3392	0.1823
Hrb98DE	hnRNA-binding pr	1638912_s_at	0.1925	0.4062	-0.2582	0.5463	-0.2989	0.2568	-0.3966	0.2309	0.2563	0.1905	0.6529	0.0035	-0.4036	0.7644	-0.1468	0.8491	0.2568	0.6800
CG15603	CG15603	1638913_at	-0.0228	0.9109	0.0142	0.9445	-0.1502	0.4692	0.1655	0.6662	-0.0088	0.9753	-0.1743	0.2930	0.1032	0.9011	0.0842	0.8356	-0.0190	0.9666
CG34366	CG13111	1638914_at	0.5322	0.1712	0.2734	0.1247	0.1929	0.3503	-0.1491	0.7857	-0.0366	0.9075	0.1125	0.6250	0.0835	0.9499	0.0481	0.9402	-0.0354	0.9491
Ets21C	Ets at 21C	1638915_at	-0.1225	0.7323	0.3839	0.5842	0.2905	0.6664	0.2289	0.7556	0.3266	0.3071	0.0977	0.7820	0.4307	0.8973	1.0521	0.3922	0.6214	0.6426
CG12688	CG12688	1638916_at	0.0658	0.7632	0.0275	0.8923	0.0182	0.9373	0.0576	0.9218	0.0310	0.8983	-0.0266	0.9003	0.0783	0.9301	0.1379	0.6643	0.0596	0.8790
---	---	1638917_at	0.0670	0.7600	-0.1911	0.3542	0.0288	0.8782	0.1351	0.7248	0.1113	0.5489	-0.0237	0.9095	-0.1900	0.7707	-0.1620	0.5825	0.0280	0.9431
---	---	1638918_at	-0.1416	0.5878	0.1258	0.3353	0.2535	0.2531	0.0628	0.9592	-0.1150	0.7675	-0.1778	0.5831	0.0700	0.9371	0.1776	0.5415	0.1077	0.7380
A16	A16	1638919_at	0.0110	0.9597	-0.0911	0.3881	-0.0452	0.8662	0.1280	0.7949	0.2156	0.2859	0.0876	0.6763	0.0789	0.9500	0.1689	0.6865	0.0901	0.8524
---	---	1638920_at	0.1272	0.6831	-0.0292	0.7713	0.1994	0.1894	0.0453	0.9641	0.0621	0.8594	0.0168	0.9597	-0.0735	0.9342	-0.0636	0.8764	0.0099	0.9838
---	---	1638921_at	0.1590	0.4357	-0.0258	0.8215	0.0603	0.7916	-0.1851	0.6084	-0.0025	0.9922	0.1826	0.2635	0.0921	0.9342	0.0180	0.9781	-0.0741	0.8800
---	---	1638922_at	0.2375	0.2671	0.1151	0.6264	0.2250	0.3123	0.0063	0.9956	-0.0599	0.8472	-0.0662	0.8061	-0.0337	0.9742	-0.0742	0.8271	-0.0405	0.9095
CG3566	CG3566	1638923_at	0.0843	0.6535	-0.0285	0.9472	0.1173	0.4882	0.1829	0.6114	0.1928	0.2891	0.0099	0.9658	-0.0548	0.9705	0.0134	0.9855	0.0682	0.8949
CG31845	CG31845	1638924_at	-0.0177	0.9633	0.0129	0.9355	0.1411	0.4784	-0.1653	0.7949	-0.0909	0.7705	0.0744	0.7983	-0.1014	0.8825	0.0423	0.9170	0.1437	0.6109
CG34024	CG34024	1638925_at	0.1457	0.5065	-0.0329	0.7760	0.1384	0.4114	0.1301	0.7187	0.0109	0.9656	-0.1192	0.4389	-0.0882	0.9056	-0.0992	0.7550	-0.0110	0.9804
---	---	1638926_at	0.1268	0.5388	0.2212	0.3566	-0.0915	0.5680	-0.1650	0.8034	-0.2547	0.3457	-0.0897	0.7570	0.1485	0.8379	-0.1376	0.6783	-0.2860	0.3518
CG3279	CG3279	1638927_at	0.2451	0.2299	0.1148	0.5304	0.0345	0.9019	-0.1649	0.6558	-0.0332	0.8906	0.1317	0.4309	-0.0757	0.9235	-0.0453	0.9086	0.0304	0.9338
Or23a	Olfactory receptor	1638928_at	0.0054	0.9785	0.1791	0.1587	0.0718	0.6519	-0.1017	0.7929	-0.1394	0.3898	-0.0376	0.8355	0.0059	0.9943	0.0269	0.9303	0.0211	0.9382
CG8920	CG8920	1638929_at	0.7141	0.1971	-0.3663	0.4305	-0.8894	0.0181	-0.5426	0.2438	1.0395	0.0033	1.5822	0.0004	-0.0033	0.9998	-0.1106	0.9402	-0.1073	0.9330
CG11248	CG11248	1638930_s_at	-0.0704	0.7538	0.1910	0.2342	0.2402	0.1416	-0.1951	0.4861	-0.1677	0.2643	0.0274	0.8776	-0.0420	0.9621	0.2328	0.3322	0.2747	0.2798
Pcf11	lethal (2) k08015	1638931_at	0.2309	0.4578	0.5413	0.2652	0.7593	0.0018	0.0048	0.9960	-0.3126	0.2882	-0.3174	0.2249	-0.0578	0.9773	0.0709	0.9248	0.1287	0.8305
---	---	1638932_at	0.1152	0.5623	0.1512	0.4533	-0.0509	0.7385	-0.0189	0.9839	-0.1527	0.5092	-0.1338	0.5306	0.0764	0.9355	0.0690	0.8721	-0.0074	0.9874
mus304	mutagen-sensitive	1638933_at	0.1915	0.4711	-0.3441	0.0588	-0.0075	0.9791	-0.3575	0.4998	0.4437	0.1119	0.8012	0.0070	-0.5396	0.3517	-0.1280	0.7162	0.4116	0.2050
---	---	1638934_at	0.0229	0.9059	0.2134	0.3021	0.0803	0.6736	-0.0500	0.9297	-0.1883	0.2578	-0.1383	0.3660	0.1496	0.7726	0.0647	0.8275	-0.0849	0.7439
CG7091	CG7091	1638935_at	-1.2573	0.0015	-1.6456	0.0057	-0.8675	0.0167	0.1469	0.8327	-0.2592	0.3335	-0.4061	0.0860	-0.7316	0.1628	-0.7993	0.0261	-0.0677	0.8459
CG40169	CG40169	1638936_s_at	-0.0326	0.8800	0.2308	0.3407	0.1699	0.4453	-0.0551	0.9307	0.0302	0.9062	0.0853	0.6564	-0.0893	0.9124	0.0721	0.8546	0.1613	0.5995
CG3488	CG3488	1638937_at	0.1632	0.3882	-0.1912	0.4332	-0.1328	0.4697	0.1894	0.5779	0.3539	0.0478	0.1645	0.2982	0.1069	0.8963	-0.0511	0.9141	-0.1580	0.6317
CG9418	CG9418	1638938_at	-0.0600	0.8929	-0.2904	0.2329	-0.8318	0.0011	0.2333	0.6013	1.0534	0.0014	0.8201	0.0023	0.7585	0.4415	0.7198	0.1504	-0.0388	0.9585
CalpA	Calpain A	1638939_at	-1.4624	0.0004	-0.8545	0.0401	-1.3877	0.0002	-0.2917	0.4867	-0.5860	0.0171	-0.2943	0.1408	0.1429	0.8655	-0.2018	0.5700	-0.3448	0.3175
RhoGAP15B	RhoGAP15B	1638940_at	-0.8827	0.0406	-1.0610	0.1894	-1.0124	0.0234	-0.0229	0.9909	0.0681	0.9114	0.0910	0.8592	-0.0756	0.9829	-0.0514	0.9692	0.0243	0.9849
---	---	1638941_at	-0.0874	0.6185	0.1129	0.3430	0.0667	0.7282	0.0748	0.8507	-0.1045	0.4903	-0.1793	0.1642	0.0271	0.9764	0.0021	0.9973	-0.0250	0.9350
CG30499	CG30499	1638942_at	0.3969	0.0967	1.3199	0.0108	1.6382	0.0001	0.1354	0.8076	-1.0110	0.0015	-1.1464	0.0005	-0.2239	0.7485	-0.0719	0.8622	0.1519	0.6345
Z600	frustart	1638943_at	0.2111	0.2665	0.0479	0.7044	-0.1904	0.2880	-0.1528	0.6354	0.0293	0.8910	0.1821	0.1969	0.1470	0.8284	-0.0448	0.9193	-0.1918	0.5194
CG15780 /// CG34434 /// C	CG15780	1638944_at	0.0661	0.7508	0.0986	0.4558	0.1228	0.4494	-0.0284	0.9705	-0.1828	0.3644	-0.1544	0.4005	0.0664	0.9487	0.1593	0.6331	0.0929	0.8057
CG1162	CG1162	1638945_at	0.0097	0.9684	0.1950	0.4870	0.1938	0.3432	-0.0323	0.9704	-0.1130	0.6502	-0.0807	0.7348	-0.1264	0.8850	-0.0022	0.9988	0.1242	0.7443
---	---	1638946_at	-0.0249	0.9104	-0.0466	0.7599	0.3935	0.1815	-0.1432	0.7115	-0.1742	0.3282	-0.0310	0.8815	-0.2239	0.7779	-0.0102	0.9887	0.2137	0.5587
mRpL24	mitochondrial ribo	1638947_at	-0.1700	0.6994	0.0137	0.9674	-0.0731	0.8129	-0.2415	0.5019	-0.0918	0.6732	0.1498	0.3991	-0.0830	0.9742	0.0515	0.9614	0.1345	0.8757
slam	slow as molasses	1638948_at	0.5649	0.1120	-0.0456	0.9387	-0.2046	0.4421	-0.0682	0.9386	0.5304	0.0464	0.5986	0.0179	0.0684	0.9816	-0.1268	0.9009	-0.1952	0.8162
CG31830	CG31830	1638949_at	0.1389	0.5047	0.0291	0.7932	0.1837	0.3462	0.1068	0.8640	-0.0332	0.9147	-0.1399	0.5132	-0.2052	0.7116	-0.1830	0.4293	0.0222	0.9453
dpr13	dpr13	1638950_at	-0.0100	0.9612	0.0054	0.9629	0.0314	0.8821	-0.0657	0.8955	0.0389	0.8576	0.1046	0.5146	-0.1616	0.7241	-0.1094	0.6061	0.0522	0.8354
CG1695	CG1695	1638951_at	0.0210	0.9222	-0.0136	0.8960	-0.0703	0.6832	0.1261	0.7415	0.1918	0.2563	0.0658	0.7129	0.1480	0.8270	0.1002	0.7692	-0.0478	0.9023
---	---	1638952_at	0.0589	0.8426	0.5214	0.0471	0.4336	0.0231	-0.0885	0.8394	-0.4595	0.0124	-0.3709	0.0193	0.0201	0.9914	0.1379	0.7892	0.1178	0.8184
drpr	draper	1638953_a_at	-0.4545	0.0356	-0.6596	0.0596	-1.2801	0.0003	-0.0995	0.7596	0.2373	0.0904	0.3369	0.0151	0.5082	0.4670	-0.0055	0.9941	-0.5137	0.1702
CG33639	CG33639	1638954_at	0.0046	0.9822	0.0259	0.8144	-0.1801	0.1973	-0.1958	0.5039	-0.1673	0.2846	0.0284	0.8776	0.1339	0.8815	-0.0530	0.9214	-0.1870	0.6145
CG32277	CG32277	1638955_at	3.8349	0.0047	2.4606	0.0022	6.3416	0.0000	3.7431	0.0262	1.4836	0.1038	-2.2595	0.0135	-0.0496	0.9405	-0.0655	0.8018	-0.0159	0.9566
Fas2	Fasciclin II	1638956_at	-0.0374	0.9245	-0.6755	0.1716	-1.1081	0.0040	-0.0326	0.9777	0.4741	0.1022	0.5066	0.0552	0.4123	0.7768	-0.0868	0.9309	-0.4991	0.4387
---	---	1638957_at	0.0646	0.7758	-0.0321	0.7796	-0.0279	0.8931	0.1540	0.7379	0.1088	0.6310	-0.0452	0.8482	0.3658	0.4415	0.1379	0.6032	-0.2279	0.3720
CG31644	CG31644	1638958_at	0.2435	0.2509	-0.1081	0.4088	0.0056	0.9830	0.0379	0.9562	0.1557	0.4186	0.1178	0.5133	-0.1649	0.7230	-0.1136	0.5923	0.0513	0.8408
CG8501	CG8501	1638959_at	-1.3774	0.1311	-1.5179	0.0018	-1.7774	0.0000	-0.3455	0.2949	-0.4854	0.0206	-0.1400	0.4432	-0.2165	0.9405	-0.6720	0.4565	-0.4555	0.6385
---	---	1638960_at	0.2693	0.2617	0.0608	0.6612	0.3410	0.0377	0.0124	0.9922	-0.1594	0.5665	-0.1718	0.4848	-0.3449	0.6749	-0.1302	0.7466	0.2147	0.5535
CG14264	CG14264	1638961_at	-0.1927	0.3611	-0.1033	0.7229	-0.2857	0.2708	-0.3443	0.6564	0.0074	0.9897	0.3517	0.2972	-0.0634	0.9421	0.0329	0.9402	0.0963	0.7568
CG6227	CG6227	1638962_at	0.2589	0.7141	-0.0151	0.9907	0.2710	0.4306	-0.0209	0.9852	0.6155	0.0233	0.6364	0.0127	-0.1260	0.9816	0.5101	0.7225	0.6360	0.6375
mrj	mrj	1638963_s_at	0.8605	0.0091	0.8779	0.0247	0.1066	0.6713	-0.1779	0.7281	0.3181	0.1629	0.4959	0.0236	0.5425	0.3734	0.3178	0.3284	-0.2247	0.5200
CG7800	CG7800	1638964_at	-0.2390	0.3796	-0.0711	0.7150	-0.2300	0.3498	-0.3064	0.4586	-0.1643	0.4837	0.1420	0.5118	-0.1385	0.9032	0.0416	0.9505	0.1801	0.6973
CG5859	CG5859	1638965_at	0.2373	0.3503	0.0966	0.6072	0.0836	0.6279	0.0573	0.9311	0.3916	0.0504	0.3343	0.0598	0.1233	0.8901	0.3386	0.2982	0.2153	0.5411
bin	Biniou	1638966_at	0.2528	0.2215	0.0478	0.7193	-0.0167	0.9348	0.1173	0.7425	0.1733	0.2719	0.0561	0.7403	0.1924	0.7395	0.0356	0.9275	-0.1568	0.5460
CG32224	CG32224	1638967_at	0.1721	0.4107	-0.0106	0.9579	0.0767	0.7688	0.0411	0.9705	0.1399	0.6675	0.0988	0.7515	-0.0272	0.9829	-0.1300	0.6909	-0.1029	0.7605
CG18171	CG18171	1638968_at	-0.1752	0.2512	0.1918	0.1920	0.0979	0.5633	-0.0155	0.9860	-0.1335	0.5485	-0.1179	0.5646	0.1090	0.8814	0.1215	0.7002	0.0125	0.9777
CG6279	CG6279	1638969_at	0.6936	0.0091	0.3670	0.2015	0.5201	0.0428	-0.1624	0.7048	0.1441	0.4849	0.3065	0.0829	-0.0046	0.9974	-0.1964	0.5625	-0.1917	0.5777
CG14543	CG14543	1638970_at	0.3929	0.0754	0.4793	0.1431	0.5898	0.0382	0.1417	0.7149	0.3138	0.0783	0.1721	0.2821	0.1242	0.9238	0.4892	0.2380	0.3650	0.4021
---	---	1638971_at	0.2290	0.1789	-0.0641	0.7238	0.1047	0.5283	0.0342	0.9488	0.1531	0.3074	0.1189	0.3891	-0.1413	0.8235	-0.1136	0.7028	0.0278	0.9401
Best3	Bestrophin 3	1638972_at	0.3735	0.0747	0.2251	0.2543	0.2934	0.1408	-0.0137	0.9866	0.0091	0.9764	0.0228	0.9252	0.0013	0.9994	0.0095	0.9829	0.0081	0.9837
CG11149	CG11149	1638973_s_at	-0.1787	0.8994	-0.0969	0.3910	-0.3398	0.1451	-0.1771	0.9507	-0.8139	0.3098	-0.6368	0.3873	-0.0904	0.9848	-0.6850	0.5404	-0.5947	0.6055
CG14963	CG14963	1638974_at	-0.0434	0.7805	0.0364	0.7620	0.0623	0.8377	-0.1055	0.8244	-0.0544	0.8112	0.0511	0.8047	-0.0812	0.9095	-0.0520	0.8884	0.0292	0.9347
CG13876	CG13876	1638975_at	-0.2955	0.3809	0.2143	0.3354	-0.0486	0.8661	-0.0285	0.9847	-0.1297	0.7460	-0.1012	0.7886	-0.0929	0.9294	0.1511	0.6904	0.2440	0.4854
---	---	1638976_at	-0.0600	0.7668	-0.0883	0.4907	-0.0701	0.7043	0.1962	0.5871	0.1600	0.3971	-0.0362	0.8662	-0.0445	0.9677	0.0634	0.8783	0.1079	0.7460
---	---	1638977_at	-0.1557	0.4170	0.0979	0.4735	0.1697	0.4777	-0.0431	0.9603	-0.1284	0.6164	-0.0854	0.7312	-0.1109	0.8650	0.0573	0.8773	0.1682	0.5391
---	---	1638978_at	0.2398	0.1078	0.3921	0.0893	0.4010	0.0677	-0.0247	0.9803	-0.1880	0.4523	-0.1633	0.4755	0.0164	0.9914	0.0906	0.8468	0.0743	0.8729

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
mu2	mutator 2	1638979_a_at	0.6555	0.2509	-0.4710	0.2270	-0.6147	0.0640	-0.1397	0.9154	1.1575	0.0126	1.2972	0.0045	-0.1741	0.9309	-0.0393	0.9710	0.1348	0.8779
CG3611	CG3611	1638980_at	0.1484	0.3950	0.0032	0.9825	0.0309	0.8516	0.1464	0.6086	0.0791	0.6191	-0.0673	0.6493	0.0687	0.9467	-0.0372	0.9411	-0.1059	0.7749
Ugt86Dh	Ugt86Dh	1638981_at	-0.3133	0.8797	0.0795	0.4914	-0.1814	0.2507	-0.3645	0.9380	-1.1750	0.4048	-0.8105	0.5473	0.0024	0.9998	-0.7016	0.5595	-0.7039	0.5629
---	---	1638982_at	0.0587	0.7871	-0.1177	0.4901	-0.0823	0.6316	0.0986	0.8942	0.2771	0.2633	0.1786	0.4416	0.0502	0.9449	-0.0291	0.9341	-0.0793	0.7543
btv	beethoven	1638983_at	0.1856	0.3392	-0.0674	0.7493	0.0394	0.8561	0.0777	0.9036	-0.0085	0.9792	-0.0861	0.6940	-0.0655	0.8999	-0.1694	0.3685	-0.1039	0.6150
CG17816	CG17816	1638984_s_at	-0.0735	0.8847	-0.0080	0.9913	-0.4388	0.1589	-0.2487	0.7023	0.3799	0.2022	0.6285	0.0268	0.0695	0.9833	0.3792	0.6543	0.3096	0.7264
CG12626	CG12626	1638985_at	0.1430	0.5244	0.2829	0.3197	0.1988	0.2733	0.0687	0.9105	-0.0746	0.7446	-0.1433	0.4304	0.0378	0.9667	-0.1065	0.6948	-0.1444	0.5724
CG11905	CG11905	1638986_s_at	0.2599	0.2314	0.1697	0.3657	0.2502	0.1189	-0.0251	0.9760	-0.0128	0.9682	0.0123	0.9621	-0.0668	0.9421	0.0684	0.8632	0.1352	0.6607
---	---	1638987_at	0.0529	0.7971	-0.1352	0.3567	0.0768	0.6390	0.1240	0.8074	0.1425	0.5057	0.0185	0.9416	-0.0435	0.9545	-0.0965	0.7027	-0.0530	0.8555
CG12914	CG12914	1638988_at	0.1608	0.3196	-0.0394	0.7138	-0.1747	0.3437	0.0629	0.9228	0.1686	0.4015	0.1058	0.5885	0.0022	0.9986	-0.1123	0.6669	-0.1146	0.6541
---	---	1638989_at	0.0540	0.7924	-0.0099	0.9273	0.0073	0.9801	0.1500	0.7850	0.1017	0.6965	-0.0483	0.8545	0.0777	0.9095	0.0128	0.9760	-0.0649	0.8321
---	---	1638990_at	0.5288	0.1451	0.1609	0.8001	0.2648	0.3445	0.1379	0.7936	0.5933	0.0136	0.4554	0.0260	-0.1108	0.9734	0.1389	0.9120	0.2497	0.8017
CG5506	CG5506	1638991_at	0.4503	0.8734	-0.5560	0.0146	-0.8131	0.0129	0.1794	0.9744	-0.0684	0.9743	-0.2478	0.8776	0.0741	0.9943	-1.2025	0.6472	-1.2766	0.6238
---	---	1638992_at	0.3313	0.0640	0.1260	0.3823	0.3206	0.1314	0.2480	0.5515	0.1859	0.4016	-0.0621	0.7976	0.0701	0.9515	-0.1693	0.6552	-0.2395	0.5061
---	---	1638993_at	-0.2051	0.3346	-0.2998	0.2126	-0.0811	0.6284	0.1646	0.6854	0.2027	0.2872	0.0381	0.8624	-0.0550	0.9416	-0.0169	0.9648	0.0381	0.9056
CG10627	CG10627	1638994_at	0.8324	0.0060	1.0781	0.0348	1.0643	0.0006	0.0411	0.9515	0.1558	0.4188	0.1147	0.5276	0.0882	0.9495	0.4888	0.2224	0.4005	0.3488
CG2292	CG2292	1638995_at	0.4256	0.0527	1.0324	0.1284	0.7262	0.0038	-0.1341	0.8667	0.0006	0.9988	0.1347	0.6411	0.1811	0.9092	0.5689	0.2978	0.3878	0.5049
CG11727 /// DsimCG11727	CG11727	1638996_at	-0.1995	0.6655	-0.0308	0.8835	0.3596	0.1403	0.0972	0.9482	0.2551	0.5737	0.1580	0.7244	-0.2958	0.7768	0.4164	0.3433	0.7121	0.1422
CG30342	CG30342	1638997_at	-0.1453	0.4601	0.3806	0.3433	0.5425	0.0079	0.1190	0.7293	-0.2650	0.0859	-0.3840	0.0130	0.0575	0.9759	0.2462	0.6166	0.1888	0.7154
cul-5	cullin-5	1638998_at	0.3469	0.2910	0.2534	0.6009	0.3058	0.0669	-0.2640	0.4209	-0.0676	0.7552	0.1964	0.2271	-0.2716	0.8439	-0.1085	0.8988	0.1631	0.8174
rno	rhinoceros	1638999_at	0.4525	0.3926	1.1958	0.0375	1.0043	0.0088	-0.2311	0.7803	-0.3127	0.3736	-0.0816	0.8358	0.1551	0.9340	0.4722	0.4294	0.3172	0.6225
CG40175	CG40175	1639000_at	-0.0494	0.8001	0.0892	0.4932	-0.0335	0.8752	0.0476	0.9376	-0.0287	0.9077	-0.0764	0.6825	0.1901	0.7677	0.0298	0.9462	-0.1603	0.5811
alpha-Est9	fragment J	1639001_a_at	-1.8317	0.0015	-0.1962	0.7966	-0.5747	0.0056	-0.6995	0.2051	-1.7656	0.0008	-1.0661	0.0041	-0.1034	0.9717	0.0277	0.9844	0.1311	0.8982
CG14801	CG14801	1639002_s_at	1.5840	0.0349	0.9280	0.2439	1.6192	0.0012	-0.3368	0.3887	-0.1058	0.6741	0.2310	0.2437	-0.8791	0.6584	-0.8921	0.2522	-0.0130	0.9924
CG13137	CG13137	1639003_at	0.0742	0.6597	-0.1344	0.4224	-0.0814	0.6599	0.0848	0.8671	0.2066	0.2489	0.1218	0.4740	0.0753	0.8909	0.0984	0.6660	0.0230	0.9365
tutl	turtle	1639004_at	0.1816	0.3069	0.2052	0.5827	-0.0860	0.5672	-0.1172	0.8000	0.2145	0.2513	0.3317	0.0513	0.0665	0.9412	-0.0136	0.9784	-0.0801	0.8178
CG33272	inverted repeat 1	1639005_x_at	-0.1576	0.6604	-0.3747	0.0818	-0.1106	0.7516	-0.0050	0.9978	0.0686	0.9167	0.0736	0.8955	-0.2897	0.7070	-0.1921	0.5708	0.0976	0.8039
mod(mdg4)	Modifier67.2	1639006_at	-0.2071	0.3269	0.0100	0.9328	0.0673	0.7727	0.2697	0.4501	0.2350	0.2250	-0.0347	0.8800	-0.0746	0.9142	0.0996	0.7200	0.1742	0.4869
---	---	1639007_at	0.1988	0.1895	0.0124	0.9323	0.0662	0.6713	-0.0167	0.9777	0.1645	0.2701	0.1812	0.1714	0.0317	0.9721	0.0709	0.8123	0.0391	0.9023
CG15260	CG15260	1639008_at	0.0811	0.6643	0.0367	0.7542	-0.0095	0.9770	-0.0719	0.9149	0.1342	0.5541	0.2061	0.2789	0.0126	0.9898	-0.0086	0.9829	-0.0213	0.9431
CG10962	CG10962	1639009_at	0.0211	0.8999	-0.0575	0.5518	-0.0978	0.5638	0.1072	0.8350	0.2102	0.2838	0.1030	0.5982	0.0634	0.9168	0.0451	0.8807	-0.0182	0.9510
---	---	1639010_at	0.2169	0.2740	-0.2068	0.1202	-0.0253	0.9123	0.0047	0.9956	0.1544	0.5541	0.1497	0.5253	-0.2142	0.6898	-0.1740	0.4270	0.0402	0.8935
CG12162	CG12162	1639011_a_at	-0.3509	0.3312	-0.4540	0.4258	-0.1413	0.5145	0.2835	0.4110	0.0495	0.8411	-0.2341	0.1699	0.0643	0.9835	0.1551	0.8807	0.0908	0.9276
---	---	1639012_at	-0.1617	0.4206	0.1499	0.2598	0.0556	0.8189	0.0264	0.9745	-0.1477	0.4908	-0.1742	0.3537	0.0940	0.9142	0.1760	0.5852	0.0820	0.8305
CG5027	CG5027	1639013_at	-0.0927	0.6249	-0.0696	0.8823	0.0972	0.5976	0.1784	0.5900	0.4451	0.0162	0.2667	0.0773	-0.0018	0.9997	0.3614	0.4167	0.3632	0.4285
CG5404	CG5404	1639014_at	0.8657	0.3334	-0.0672	0.6014	0.0397	0.8070	0.0826	0.8686	-0.0560	0.7952	-0.1386	0.3975	-0.0603	0.9914	-1.0686	0.3796	-1.0082	0.4272
G-salpha60A	G-alpha s	1639015_s_at	-0.6600	0.0098	-0.4702	0.0655	-0.8145	0.0007	-0.1277	0.7753	-0.1807	0.3427	-0.0529	0.8013	0.1470	0.8222	0.0020	0.9985	-0.1450	0.6202
CG8206	CG8206	1639016_at	-0.2977	0.2692	-0.0992	0.6819	-0.0353	0.8922	-0.0684	0.9220	-0.5480	0.0176	-0.4796	0.0192	-0.1374	0.9174	-0.4655	0.2975	-0.3281	0.4888
---	---	1639017_at	0.1457	0.3522	0.0043	0.9845	0.2493	0.2121	0.2557	0.3034	0.1323	0.3754	-0.1234	0.3580	-0.0634	0.9672	-0.2458	0.5644	-0.1824	0.6854
---	---	1639018_at	-0.0247	0.9173	0.3705	0.1065	-0.0686	0.6792	-0.0219	0.9777	0.0026	0.9930	0.0244	0.9201	0.1843	0.8049	0.1990	0.5378	0.0146	0.9766
CG33470 /// IM10	Immune induced r	1639019_s_at	0.2104	0.8102	-1.4584	0.0309	-0.5392	0.1921	0.4742	0.6326	1.6657	0.0051	1.1916	0.0139	-0.4011	0.8906	0.0538	0.9784	0.4549	0.7135
---	---	1639020_at	-0.0912	0.5910	0.0772	0.5308	0.1086	0.5013	-0.0564	0.9017	-0.1771	0.2310	-0.1207	0.3786	-0.0226	0.9824	0.0192	0.9614	0.0419	0.8982
---	---	1639021_at	0.1218	0.4421	0.0162	0.8841	0.2704	0.1366	0.0110	0.9880	-0.0213	0.9289	-0.0323	0.8718	-0.0493	0.9414	0.0011	0.9991	0.0503	0.8509
CG12836	CG12836	1639022_at	0.0070	0.9715	0.1063	0.4810	-0.2672	0.1313	-0.0969	0.8609	-0.0956	0.6723	0.0013	0.9958	0.0610	0.9309	0.0118	0.9764	-0.0492	0.8726
Nsf2	NEM-sensitive fus	1639023_at	-0.6197	0.1124	-0.1893	0.2931	-0.2952	0.2102	-0.2256	0.6854	0.2354	0.3728	0.4610	0.0509	-0.1709	0.8344	0.6000	0.0890	0.7710	0.0607
CG9630	CG9630	1639024_at	0.2139	0.4388	0.2816	0.3787	0.2577	0.1459	0.0754	0.9319	0.4530	0.0854	0.3776	0.1066	0.1044	0.9342	0.4112	0.2984	0.3068	0.4652
Trc8	Trc8	1639025_at	-0.5211	0.4821	-0.1317	0.8628	-0.6056	0.0081	-0.1339	0.8940	-0.2249	0.5281	-0.0910	0.8087	0.2671	0.9421	0.2382	0.8850	-0.0289	0.9874
---	---	1639026_at	0.0286	0.9328	0.0450	0.7239	0.0787	0.6538	-0.0569	0.9120	-0.0900	0.6196	-0.0331	0.8621	0.0267	0.9822	-0.1032	0.7554	-0.1299	0.6677
---	---	1639027_at	0.1254	0.4557	0.0915	0.5527	0.0192	0.9252	-0.1768	0.7589	0.0685	0.8258	0.2453	0.2718	-0.0411	0.9665	0.0375	0.9275	0.0786	0.8046
CG30296	CG30296	1639028_a_at	-2.0893	0.0033	-2.2417	0.0234	-3.4780	0.0000	-0.5278	0.1605	-0.8156	0.0043	-0.2878	0.1569	0.7030	0.7485	-0.9190	0.3137	-1.6219	0.1184
mei-S332	meiotic from via S	1639029_at	-0.5592	0.0502	-0.1942	0.4603	-0.3667	0.0689	-0.2873	0.5360	0.3188	0.1857	0.6060	0.0129	-0.1955	0.8626	0.1819	0.7305	0.3775	0.4142
---	---	1639030_at	0.0293	0.8675	0.0203	0.9048	0.0967	0.5250	0.0529	0.9345	-0.0039	0.9894	-0.0568	0.7884	0.0184	0.9831	0.0047	0.9924	-0.0137	0.9633
---	---	1639031_at	0.2836	0.1643	0.1409	0.4072	0.1156	0.5574	-0.0567	0.9376	0.0047	0.9884	0.0614	0.7967	0.1274	0.8609	0.0170	0.9726	-0.1104	0.7445
CG14786	CG14786	1639032_at	-0.7288	0.2796	-0.0677	0.9626	0.1335	0.7492	0.3898	0.2358	-0.6655	0.0052	-1.0553	0.0005	0.3970	0.9309	0.1191	0.9617	-0.2779	0.8918
l(2)01289	Fusion2	1639033_at	-1.8990	0.0014	-2.0783	0.0198	-2.5083	0.0000	-0.2379	0.5008	-0.0960	0.6476	0.1419	0.4170	0.2942	0.8357	-0.2899	0.6497	-0.5840	0.3307
CG7593	CG7593	1639034_at	0.3216	0.1566	0.0300	0.8120	0.2804	0.5159	-0.2948	0.7143	0.5198	0.1556	0.8145	0.0214	-0.2603	0.7783	0.2759	0.5068	0.5361	0.2051
Rad51D	Rad51D	1639035_at	-0.0399	0.8847	0.1611	0.3218	0.0248	0.9410	-0.1150	0.8657	-0.0765	0.7951	0.0384	0.8939	-0.1091	0.9296	0.0354	0.9567	0.1445	0.7526
RpS11	Ribosomal protein	1639036_at	0.2745	0.2656	-0.2554	0.3700	-0.2275	0.1341	0.1151	0.8155	0.4338	0.0323	0.3187	0.0684	0.1414	0.8608	0.0672	0.8857	-0.0742	0.8618
CG13588	CG13588	1639037_at	0.0179	0.9392	0.2369	0.2768	0.1368	0.4484	-0.0456	0.9620	-0.2210	0.3973	-0.1753	0.4675	-0.0769	0.9449	0.0705	0.8850	0.1474	0.6939
CG9213	CG9213	1639038_at	0.1300	0.6507	-0.3533	0.3388	-0.5235	0.0076	0.0084	0.9937	0.6857	0.0077	0.6773	0.0048	0.0338	0.9875	0.0421	0.9587	0.0083	0.9924
CG32852	CG32852	1639039_at	-0.1482	0.5364	-0.0388	0.7421	-0.0223	0.9314	0.0049	0.9956	-0.0708	0.7978	-0.0757	0.7583	-0.0606	0.9491	-0.1478	0.6291	-0.0872	0.8003
CG8291	CG8291	1639040_s_at	-0.2068	0.5227	0.1093	0.7775	0.5193	0.0630	0.2273	0.4861	-0.1136	0.5457	-0.3409	0.0360	-0.1433	0.9174	0.0400	0.9594	0.1833	0.7398
---	---	1639041_at	0.0574	0.7262	-0.0247	0.8982	0.1246	0.4525	-0.0531	0.9295	-0.0179	0.9452	0.0352	0.8630	0.0036	0.9970	-0.0843	0.7715	-0.0879	0.7492
CG6414	CG6414	1639042_at	0.0939	0.6640	-0.0243	0.8240	0.0336	0.8895	0.0641	0.9017	0.0225	0.9234	-0.0415	0.8292	0.0443	0.9738	0.0022	0.9988	-0.0421	0.9271
CG34350	CG11824	1639043_at	0.2118	0.3668	-0.0078	0.9607	-0.2299	0.3330	0.1586	0.7018	0.3677	0.0576	0.2092	0.2192	0.2223	0.8292	0.0468	0.9467	-0.1754	0.7199
Cad96Cb	Cad96Cb	1639044_at	-0.0217	0.9237	0.1905	0.4129	0.0819	0.7509	-0.1350	0.8053	-0.1339	0.5698	0.0010	0.9973	-0.0224	0.9862	0.0667	0.8792	0.0891	0.8157
mthl5	Mth-like 5	1639045_at	-0.1577	0.5339	-0.0206	0.9678	0.1084	0.5603	0.0406	0.9689	0.0295	0.9398	-0.0110	0.9737	-0.2046	0.8400	0.1312	0.8081	0.3358	0.4356
CG5454	CG5454	1639046_at	0.5372	0.0309	0.0497	0.8434	-0.1295	0.5057	-0.0766	0.8942	0.3007	0.1154	0.3773	0.0348	0.2068	0.8192	-0.1019	0.8372	-0.3087	0.4114
mask	multiple ankyrin re	1639047_at	-0.3928	0.4210	-0.4691	0.3887	-1.2876	0.0006	-0.3793	0.5311	0.4541	0.1492	0.8334	0.0102	0.5157	0.7772	0.3731	0.6695	-0.1426	0.8951

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG8229 /// DyakCG8229	CG8229	1639048_a_at	-0.3371	0.0710	0.9565	0.0175	1.0168	0.0065	0.0220	0.9865	-1.5250	0.0012	-1.5470	0.0006	0.0273	0.9848	-0.0933	0.8221	-0.1206	0.7423
CG17145	CG17145	1639049_at	0.0915	0.6264	0.3571	0.0555	0.2431	0.2173	0.0061	0.9943	-0.0530	0.8147	-0.0591	0.7698	0.1596	0.7756	0.2879	0.2181	0.1283	0.6225
---	---	1639050_at	-0.0110	0.9687	0.2685	0.1592	0.2749	0.1205	-0.1209	0.8527	-0.1792	0.4673	-0.0583	0.8287	0.1277	0.8513	0.1372	0.6576	0.0095	0.9841
CG12938	CG12938	1639051_at	-0.6023	0.0849	-0.3392	0.0908	-0.7083	0.0256	-0.3751	0.5317	-0.2695	0.4027	0.1056	0.7580	-0.1321	0.8650	-0.1178	0.7466	0.0143	0.9769
CG12024	CG12024	1639052_a_at	-0.0224	0.9596	-0.1487	0.4185	-0.4150	0.1000	-0.1121	0.7807	-0.3144	0.0635	-0.2023	0.1748	0.1205	0.9515	-0.4569	0.4446	-0.5774	0.3475
---	---	1639053_at	0.1162	0.3974	0.1428	0.4160	0.1733	0.3559	0.0194	0.9834	0.0286	0.9232	0.0092	0.9725	-0.0610	0.9095	-0.0303	0.9188	0.0306	0.9080
---	---	1639054_s_at	0.2424	0.6657	-0.2873	0.4203	-0.4023	0.0283	-0.1365	0.9352	0.5021	0.3159	0.6385	0.1481	0.0363	0.9852	0.0100	0.9925	-0.0263	0.9695
CG9586	CG9586	1639055_at	-0.1823	0.2165	0.3856	0.1252	0.0609	0.7400	-0.0451	0.9411	-0.0721	0.7228	-0.0270	0.8976	0.1546	0.7853	0.4847	0.0661	0.3301	0.1993
---	---	1639056_at	0.0372	0.8973	0.1022	0.4996	0.3828	0.0959	0.0771	0.9234	0.0010	0.9978	-0.0761	0.7771	-0.1229	0.8270	-0.0432	0.9052	0.0797	0.7749
mRpS31	mitochondrial ribo	1639057_at	-0.3322	0.1741	-0.4147	0.1988	-0.2244	0.3097	-0.0926	0.8663	0.0383	0.8833	0.1309	0.4790	-0.2253	0.8206	-0.0977	0.8664	0.1276	0.7983
CG33958	CG33958	1639058_at	0.2712	0.1536	0.0596	0.6888	0.0532	0.8246	0.0271	0.9711	-0.0456	0.8529	-0.0727	0.7184	0.0551	0.9589	-0.0828	0.8380	-0.1378	0.6771
exu	exuperantia	1639059_s_at	1.2053	0.1183	-0.7830	0.6151	-0.7787	0.3481	-0.7253	0.0840	2.3491	0.0001	3.0744	0.0000	-0.8602	0.8692	0.1533	0.9647	1.0134	0.6536
l(1)10Bb	lethal(1)10Bb	1639060_at	-0.0003	0.9991	-0.1216	0.6416	-0.4111	0.0251	-0.0230	0.9672	0.2754	0.0630	0.2984	0.0299	0.2369	0.7779	-0.0364	0.9505	-0.2733	0.4683
enok	rotkehlchen	1639061_at	0.0430	0.9262	0.1946	0.4435	-0.0026	0.9916	0.2212	0.6088	0.2444	0.2625	0.0233	0.9305	0.3044	0.7953	0.3151	0.5404	0.0107	0.9895
CG31464	CG31464	1639062_at	0.2056	0.4023	0.0583	0.7311	0.2157	0.1648	0.0394	0.9461	-0.0731	0.6997	-0.1125	0.4719	-0.0544	0.9636	-0.0658	0.8893	-0.0114	0.9837
CG16969	CG16969	1639063_at	0.2583	0.3792	-0.0068	0.9832	0.3391	0.1081	0.2301	0.4376	0.3402	0.0423	0.1101	0.4762	-0.1352	0.9296	-0.0218	0.9812	0.1134	0.8618
Akt1	Related to PKA to	1639064_s_at	0.9380	0.0392	0.8126	0.0734	0.1477	0.5932	-0.2363	0.6622	0.3329	0.1972	0.5692	0.0223	0.4965	0.7464	0.2143	0.7900	-0.2822	0.6925
---	---	1639065_at	0.0442	0.8086	0.5199	0.1580	0.1030	0.5421	-0.1372	0.7929	-0.1736	0.4305	-0.0363	0.8859	0.0420	0.9800	0.1218	0.7971	0.0798	0.8745
CG34381	CG14004	1639066_at	-0.2951	0.2632	0.0623	0.5825	0.0402	0.8461	-0.0767	0.9105	-0.4776	0.0341	-0.4009	0.0438	0.0852	0.8513	0.0342	0.9054	-0.0509	0.8299
VhaM9.7-1	VhaM9.7-1	1639067_at	-0.1081	0.5471	-0.0407	0.7273	-0.2169	0.2804	-0.0499	0.9375	-0.1855	0.3205	-0.1356	0.4339	0.0995	0.8882	-0.1979	0.4639	-0.2974	0.2816
---	---	1639068_at	0.2379	0.2848	0.0349	0.8090	0.1187	0.5461	0.0283	0.9674	0.0402	0.8675	0.0119	0.9589	-0.0360	0.9677	-0.1783	0.4500	-0.1423	0.5689
Cyp12d1-d	Cyp12d1-d	1639069_at	2.3669	0.0009	2.1245	0.0060	2.8765	0.0005	0.9853	0.0482	-1.2596	0.0017	-2.2449	0.0001	0.2151	0.9260	-1.5064	0.0683	-1.7216	0.0613
CG17944	CG17944	1639070_a_at	-0.0167	0.9413	0.0500	0.8120	0.0983	0.6737	0.0438	0.9346	0.0644	0.7262	0.0206	0.9146	-0.0113	0.9916	0.0699	0.8362	0.0811	0.7875
Dredd	caspase	1639071_a_at	0.1651	0.2922	-0.1540	0.6190	0.1326	0.5544	0.0163	0.9819	0.3114	0.0723	0.2951	0.0583	-0.2136	0.8270	-0.0466	0.9438	0.1670	0.7170
hop	Hopskotch	1639072_at	0.2648	0.2754	0.1017	0.4542	-0.3419	0.1149	-0.0526	0.9052	0.2969	0.0441	0.3495	0.0143	0.4499	0.6927	0.2549	0.6101	-0.1951	0.7122
CG10104	CG10104	1639073_at	-0.0471	0.8114	-0.0744	0.5569	0.0245	0.9303	0.0091	0.9943	0.1245	0.6910	0.1154	0.6870	-0.1200	0.9092	-0.0087	0.9924	0.1113	0.8083
CG33275	CG33275	1639074_at	-1.5491	0.0069	-2.4773	0.0155	-2.4032	0.0002	0.3573	0.6869	0.4899	0.2344	0.1326	0.7710	0.3732	0.8331	-0.3157	0.7043	-0.6889	0.3626
Syx6	syntaxin	1639075_a_at	0.4338	0.0215	-0.0135	0.9493	0.4131	0.0327	-0.0240	0.9732	0.3211	0.0711	0.3450	0.0355	-0.3427	0.4128	-0.0729	0.7971	0.2698	0.2459
---	---	1639076_at	0.1082	0.6849	0.1548	0.4116	0.0665	0.7375	0.0413	0.9380	0.0900	0.5964	0.0487	0.7781	0.1122	0.9340	-0.0228	0.9760	-0.1350	0.7964
---	---	1639077_at	0.3032	0.0819	0.0011	0.9956	-0.0708	0.6427	0.2909	0.3837	0.2745	0.1441	-0.0164	0.9450	0.3748	0.5109	0.0098	0.9848	-0.3649	0.1990
CG32094	CG32094	1639078_at	0.1376	0.4394	0.0885	0.4807	0.0248	0.9098	0.0298	0.9696	-0.0335	0.9044	-0.0632	0.7775	0.1012	0.8472	-0.0409	0.9021	-0.1421	0.5376
Adgf-D	Adenosine deamir	1639079_at	3.0004	0.0015	1.6399	0.0570	3.3380	0.0001	0.9935	0.3386	0.2487	0.7216	-0.7448	0.1633	-0.6191	0.7215	-1.1304	0.1211	-0.5113	0.4938
fd3F	forkhead domain (	1639080_at	0.0627	0.8061	0.2735	0.0870	0.0936	0.5890	-0.1532	0.6327	-0.2417	0.1220	-0.0885	0.5701	-0.0051	0.9964	0.0726	0.8254	0.0777	0.7980
---	---	1639081_at	-0.0602	0.7601	0.0052	0.9664	-0.5024	0.0163	-0.2764	0.3972	0.0429	0.8581	0.3193	0.0556	0.1084	0.8400	0.1494	0.5154	0.0410	0.8926
CG14247 /// DmadCG1424	CG14247	1639082_at	0.1918	0.2648	-0.1006	0.4909	0.2407	0.3647	0.3064	0.7031	0.2958	0.4410	-0.0106	0.9819	0.0345	0.9768	0.0582	0.8867	0.0237	0.9524
faf	fat facets	1639083_at	0.6251	0.0106	0.4229	0.1343	0.6204	0.0017	0.2758	0.3855	0.5821	0.0067	0.3063	0.0605	0.0115	0.9914	0.3615	0.1251	0.3500	0.1591
---	---	1639084_s_at	0.0540	0.7810	-0.2621	0.3946	-0.1134	0.5709	0.1634	0.7608	0.2101	0.3734	0.0467	0.8623	0.1104	0.8740	-0.1102	0.7273	-0.2206	0.4318
Wwox	Wwox	1639085_at	-0.1050	0.5825	-0.2180	0.5391	-0.7113	0.0099	-0.2535	0.3663	0.3346	0.0422	0.5881	0.0022	0.1882	0.8541	0.0136	0.9873	-0.1746	0.7134
CG8863	CG8863	1639086_s_at	-0.2352	0.3191	0.1834	0.2496	-0.1905	0.2508	0.0225	0.9754	-0.2050	0.2505	-0.2275	0.1515	0.2164	0.7215	0.1643	0.5398	-0.0522	0.8813
His2Av	Histone H2A varia	1639087_at	-0.0185	0.9507	0.6170	0.0291	0.3931	0.0730	0.0637	0.9017	-0.1575	0.3554	-0.2211	0.1374	0.0980	0.9474	0.4141	0.3477	0.3161	0.5003
CG5620	CG5620	1639088_at	0.0534	0.7434	0.0910	0.5058	0.0348	0.8911	-0.1265	0.8028	-0.0292	0.9183	0.0973	0.6404	-0.1152	0.8206	-0.0060	0.9898	0.1092	0.6313
---	---	1639089_at	0.1513	0.4869	0.1756	0.3606	-0.2541	0.1079	-0.0969	0.8604	0.0443	0.8652	0.1412	0.4482	0.1811	0.7683	0.0545	0.8850	-0.1267	0.6533
CG40354	CG40354	1639090_a_at	-0.1688	0.4491	0.2488	0.3802	0.3319	0.1299	0.1438	0.8211	-0.1199	0.6715	-0.2637	0.2356	0.1499	0.7726	0.2657	0.2168	0.1158	0.6296
Rpn5	Rpn5	1639091_at	0.3363	0.0816	0.2526	0.2439	0.4427	0.0117	0.0597	0.9116	0.2740	0.1065	0.2143	0.1582	-0.0749	0.9177	0.2504	0.3042	0.3253	0.2130
CG5880	CG5880	1639092_at	0.1982	0.2478	0.3118	0.4552	0.1904	0.1941	-0.1790	0.6265	0.0466	0.8432	0.2256	0.1640	-0.0560	0.9738	0.1984	0.6686	0.2544	0.5676
Tom70	Translocase of ou	1639093_s_at	0.2268	0.3757	0.4417	0.2999	1.1106	0.0004	0.2016	0.7507	-0.0059	0.9890	-0.2074	0.4219	-0.3892	0.7070	0.3691	0.3872	0.7583	0.1156
CG10795	CG10795	1639094_at	-0.0525	0.7632	-0.2009	0.3180	-0.1290	0.4695	-0.0354	0.9599	0.2717	0.1390	0.3070	0.0658	-0.0141	0.9901	0.1570	0.5415	0.1711	0.5041
CG10426	pharbin-like	1639095_at	0.0561	0.8583	-0.2025	0.3257	-0.1640	0.2723	-0.0641	0.9029	0.1246	0.4856	0.1888	0.2163	-0.0449	0.9804	-0.1457	0.7626	-0.1008	0.8444
CG14096	CG14096	1639096_at	0.4653	0.0303	-0.2933	0.3781	-0.0278	0.9335	-0.0022	0.9986	0.4132	0.1024	0.4154	0.0692	-0.1816	0.7633	-0.2857	0.2404	-0.1040	0.7148
fs(1)Ya	young arrest	1639097_at	0.9588	0.2294	-1.0081	0.1068	-0.6342	0.2097	0.1591	0.6257	2.4202	0.0000	2.2611	0.0000	-0.4454	0.8692	-0.0570	0.9760	0.3884	0.7558
CG2837	CG2837	1639098_s_at	-1.2336	0.0252	0.2480	0.4814	0.0726	0.7965	-0.1064	0.9529	-1.1629	0.0271	-1.0565	0.0256	0.0377	0.9849	0.0183	0.9836	-0.0195	0.9792
CG32100	CG32100	1639099_at	-0.5083	0.0910	0.0168	0.9517	0.1987	0.3241	0.0942	0.8908	-0.4355	0.0617	-0.5297	0.0180	-0.1775	0.8164	0.1730	0.6080	0.3505	0.2829
---	---	1639100_at	-0.0137	0.9528	0.2185	0.4360	0.1764	0.4458	-0.2409	0.6933	-0.1500	0.6339	0.0909	0.7705	-0.0783	0.9246	-0.0223	0.9617	0.0560	0.8835
CG9837	CG9837	1639101_at	3.1362	0.0095	2.0838	0.0150	3.1841	0.0003	0.3296	0.8589	0.7071	0.2936	0.3775	0.5686	-0.5992	0.7932	-0.2357	0.8605	0.3635	0.7439
Nelf-E	Negative elongatic	1639102_at	-0.3581	0.2670	0.4545	0.0436	0.7059	0.0112	0.1223	0.8140	-0.6406	0.0078	-0.7629	0.0021	-0.1437	0.8949	0.2479	0.5609	0.3915	0.3488
---	---	1639103_s_at	0.3373	0.1166	0.0102	0.9229	0.1162	0.5122	0.2389	0.5586	0.0899	0.7114	-0.1490	0.4526	-0.0083	0.9922	-0.1453	0.4542	-0.1371	0.4938
---	---	1639104_at	0.0116	0.9751	0.1536	0.2681	0.0438	0.8148	-0.1028	0.8738	0.0099	0.9777	0.1127	0.6246	0.1039	0.9011	0.0468	0.9222	-0.0571	0.8919
ppk10	pickpocket 10	1639105_at	0.0062	0.9761	-0.2839	0.1776	-0.0479	0.8905	0.0725	0.9356	0.3304	0.2100	0.2580	0.2803	-0.1181	0.8609	0.0473	0.9093	0.1654	0.5670
Grip	Glutamate recepto	1639106_at	-2.3697	0.0011	-2.0018	0.0112	-2.5755	0.0001	0.0842	0.9436	-0.4819	0.1517	-0.5661	0.0642	0.6499	0.4464	0.0783	0.9085	-0.5716	0.2151
---	---	1639107_at	0.0646	0.6818	-0.0508	0.6369	0.0317	0.8729	-0.0917	0.8500	-0.0755	0.7118	0.0162	0.9412	-0.1803	0.6955	-0.0696	0.7625	0.1107	0.5876
CG31184	CG31184	1639108_at	-0.0046	0.9905	0.2822	0.3311	0.4089	0.2493	-0.1425	0.7937	0.0089	0.9795	0.1514	0.4748	-0.1582	0.9221	0.3796	0.4899	0.5378	0.3276
CRMP	dihydropyrimidine	1639109_a_at	0.0763	0.8320	-0.7141	0.2451	-0.5976	0.0568	-0.1393	0.8281	0.2950	0.2315	0.4342	0.0535	-0.2233	0.9117	-0.4440	0.5523	-0.2207	0.7982
CG4484	CG4484	1639110_at	0.1992	0.7151	0.5491	0.5051	0.9598	0.0218	0.1499	0.8968	-1.2552	0.0062	-1.4051	0.0022	-0.1832	0.9542	-0.7517	0.4202	-0.5685	0.5670
CG15362	CG15362	1639111_at	0.7551	0.0362	2.1123	0.0039	2.5225	0.0000	0.2834	0.5757	-0.2021	0.4548	-0.4855	0.0429	-0.1582	0.9011	1.0852	0.0415	1.2435	0.0422
---	---	1639112_s_at	0.1486	0.4198	0.0160	0.9578	0.1901	0.2339	0.1081	0.8056	0.0940	0.6294	-0.0141	0.9494	-0.0783	0.9310	-0.0642	0.8792	0.0141	0.9762
CG7705	CG7705	1639113_at	0.4119	0.0282	0.4625	0.1226	0.4625	0.0151	-0.0104	0.9915	-0.1175	0.6027	-0.1071	0.6070	0.0791	0.9270	-0.0148	0.9763	-0.0939	0.7791
---	---	1639114_at	-0.1801	0.2959	0.3372	0.1350	0.1529	0.3512	-0.1542	0.7225	-0.3666	0.0645	-0.2124	0.2258	0.0430	0.9588	0.1828	0.4280	0.1398	0.5684
rtGEF	Drosophila rho-typ	1639115_s_at	-1.3450	0.0251	-0.8271	0.1927	-1.2416	0.0002	-0.4125	0.2309	-0.3707	0.0730	0.0418	0.8608	0.0392	0.9914	0.1653	0.8880	0.1261	0.9097
beta-glycoprotein-hormone	beta-glycoprotein-	1639116_at	0.2328	0.4595	-0.1837	0.5177	-0.2967	0.1357	-0.1435	0.8076	0.2928	0.2105	0.4362	0.0436	0.0377	0.9816	-0.0965	0.8547	-0.1342	0.7632

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG33096	CG33096	1639117_a_at	0.2683	0.3683	0.4969	0.2793	0.5206	0.0153	0.0949	0.9185	0.0833	0.8128	-0.0115	0.9738	-0.0030	0.9989	0.3287	0.3240	0.3317	0.3458
Hrb87F	Heterogeneous nu	1639118_a_at	0.2049	0.5519	0.5308	0.0574	0.1296	0.5680	-0.0995	0.8578	0.2248	0.2638	0.3242	0.0723	0.3281	0.7726	0.4907	0.3034	0.1626	0.7787
---	---	1639119_at	-0.0377	0.7994	0.0295	0.8323	-0.0596	0.7677	0.0051	0.9956	0.0387	0.8877	0.0336	0.8899	-0.1539	0.7992	0.0755	0.8178	0.2295	0.3759
Gr58a	Gustatory recepto	1639120_at	-0.0556	0.8409	-0.0772	0.6725	0.0644	0.6703	0.0852	0.8908	-0.0834	0.7334	-0.1686	0.3826	-0.1035	0.9029	-0.0941	0.8139	0.0094	0.9849
Arp87C	gridlock	1639121_at	-0.2246	0.2585	0.0542	0.7153	-0.0472	0.8257	-0.0925	0.8336	0.0861	0.6460	0.1786	0.2318	0.0061	0.9964	0.4718	0.1251	0.4657	0.1537
CG31790	CG31790	1639122_at	0.1393	0.5389	0.2032	0.3795	0.1160	0.5138	-0.0369	0.9627	0.0476	0.8637	0.0844	0.7035	0.0411	0.9734	0.1328	0.6977	0.0917	0.8044
CG4882	undefined 1	1639123_at	-0.3989	0.0428	0.0749	0.7775	-0.0454	0.8216	0.0907	0.8344	0.0983	0.5817	0.0077	0.9705	0.1788	0.8378	0.5054	0.1497	0.3266	0.3757
CG31283	CG31283	1639124_at	0.1484	0.5409	-0.1431	0.3967	-0.3857	0.0398	0.0157	0.9838	0.7032	0.0027	0.6875	0.0018	-0.1338	0.8740	0.0336	0.9501	0.1674	0.6389
CG31630	CG31630	1639125_at	0.0047	0.9895	0.0371	0.8128	0.1345	0.4270	0.1780	0.7929	0.0729	0.8374	-0.1051	0.7214	0.0208	0.9816	0.0493	0.8509	0.0285	0.9151
CG31874	CG31874	1639126_at	0.1834	0.4042	0.0676	0.5500	0.0190	0.9522	0.0519	0.9435	-0.0639	0.8010	-0.1157	0.5724	0.0664	0.9444	-0.1649	0.5882	-0.2313	0.4322
CG1295	CG1295	1639127_at	0.1496	0.6231	0.2525	0.2612	0.2735	0.1571	0.0011	0.9988	-0.0522	0.8467	-0.0533	0.8216	0.0566	0.9445	0.1054	0.7043	0.0488	0.8849
CG32988	CG32988	1639128_at	0.1350	0.5311	0.0346	0.7831	0.0805	0.6146	0.1536	0.6510	0.0636	0.7424	-0.0900	0.5828	-0.0303	0.9717	-0.0649	0.8129	-0.0346	0.9057
Gr66a	Gustatory recepto	1639129_at	0.1157	0.4504	0.0260	0.9099	0.0782	0.6465	0.1405	0.7760	0.1800	0.3961	0.0394	0.8714	-0.0357	0.9814	0.0084	0.9911	0.0441	0.9211
kkv	Chitin Synthase B	1639130_a_at	-0.1161	0.6034	0.1873	0.3300	-0.0670	0.7922	-0.1108	0.8698	-0.3045	0.1966	-0.1938	0.3768	0.0208	0.9856	0.0016	0.9989	-0.0192	0.9596
CG14773	CG14773	1639131_at	-0.1252	0.5153	-0.0624	0.6416	0.1308	0.4706	0.1375	0.7967	0.1262	0.5931	-0.0113	0.9674	0.0179	0.9862	0.1626	0.5199	0.1446	0.5781
---	---	1639132_at	-0.1580	0.4548	0.0633	0.6302	0.1202	0.5857	0.0920	0.9218	-0.0228	0.9566	-0.1148	0.6999	0.1160	0.8379	0.0870	0.7544	-0.0289	0.9294
CG1939 /// DyakCG1939	CG1939	1639133_at	-1.0788	0.0144	-0.1574	0.3962	-0.4888	0.0810	-0.3675	0.5093	-1.2822	0.0018	-0.9146	0.0048	-0.1274	0.9277	-0.4226	0.3446	-0.2952	0.5420
---	---	1639134_at	0.2236	0.4178	0.1331	0.5475	0.1310	0.6125	0.1074	0.9132	0.3109	0.3240	0.2035	0.4953	0.1934	0.7726	0.1980	0.5050	0.0047	0.9924
Gtp-bp	GTP-binding prote	1639135_at	0.3931	0.1714	0.9779	0.0217	1.2710	0.0001	0.3231	0.3944	0.1419	0.5337	-0.1811	0.3559	0.0801	0.9340	0.6971	0.0461	0.6170	0.0833
psq	pipsqueak	1639136_s_at	-0.2735	0.5288	0.8502	0.1494	0.5823	0.0732	-0.5076	0.1899	-0.9507	0.0025	-0.4432	0.0408	-0.2373	0.8903	0.2846	0.6955	0.5220	0.4312
---	---	1639137_at	0.1808	0.2948	0.1119	0.3407	-0.0881	0.7178	-0.0381	0.9523	0.0658	0.7552	0.1039	0.5498	0.1112	0.8988	0.0875	0.8404	-0.0237	0.9597
CG5524	CG5524	1639138_at	0.0592	0.8182	-0.2692	0.4848	-0.3746	0.1262	-0.1105	0.8485	0.6639	0.0078	0.7744	0.0023	0.0196	0.9898	0.2639	0.4094	0.2443	0.4648
Fps85D	Fps oncogene ana	1639139_a_at	-1.4694	0.0015	-0.8139	0.1059	-1.9116	0.0000	-0.5775	0.1403	-0.9674	0.0024	-0.3900	0.0686	0.4325	0.7116	-0.3399	0.4977	-0.7724	0.1413
CG8400	CG8400	1639140_at	0.1503	0.4144	0.1296	0.5237	0.2448	0.2084	0.0448	0.9466	-0.0556	0.8122	-0.1004	0.5966	0.0199	0.9870	0.0561	0.8918	0.0361	0.9254
---	---	1639141_at	0.1638	0.3788	0.1456	0.3955	-0.1563	0.3423	-0.0921	0.8676	0.0430	0.8680	0.1350	0.4665	0.1351	0.7953	-0.0099	0.9806	-0.1450	0.5284
---	---	1639142_s_at	0.3718	0.1314	0.2025	0.4088	0.1334	0.7032	0.0783	0.9563	0.1884	0.6680	0.1101	0.7996	0.1172	0.9342	-0.0276	0.9714	-0.1448	0.7931
CG32302	CG32302	1639143_at	0.0376	0.8326	0.0887	0.4006	0.0414	0.8500	-0.0083	0.9937	-0.1824	0.3806	-0.1741	0.3514	-0.0269	0.9816	-0.1375	0.6124	-0.1106	0.6956
Tace	Tace	1639144_a_at	-0.2989	0.1665	-0.0194	0.8599	-0.1138	0.4947	-0.1630	0.8129	-0.2071	0.4658	-0.0440	0.8924	-0.0015	0.9994	0.0170	0.9711	0.0185	0.9646
Atpalpha	Na,K ATPase alph	1639145_s_at	-1.1767	0.0094	-0.2274	0.7087	-0.4724	0.0621	-0.1846	0.8189	-0.9868	0.0078	-0.8022	0.0119	0.0660	0.9816	0.0213	0.9871	-0.0448	0.9624
---	---	1639146_at	0.1165	0.5727	0.1152	0.3391	0.0504	0.8617	-0.0763	0.8895	0.1321	0.4933	0.2084	0.2040	0.0318	0.9816	0.0366	0.9402	0.0048	0.9924
CG30438	CG30438	1639147_s_at	-0.2000	0.6220	-0.8182	0.0541	-1.1291	0.0071	0.0930	0.9466	0.7175	0.0707	0.6246	0.0770	0.2412	0.8481	-0.0003	1.0000	-0.2415	0.6749
CG5325	Peroxisomal farne	1639148_at	-0.3884	0.0544	-0.1351	0.4844	-0.1817	0.3844	0.0046	0.9956	-0.2855	0.1156	-0.2901	0.0764	-0.0086	0.9959	-0.1686	0.6969	-0.1600	0.7130
CG15313	CG15313	1639149_at	0.0493	0.8310	-0.3815	0.2713	-0.1147	0.5569	0.3889	0.4669	0.5311	0.0688	0.1422	0.6293	0.0475	0.9761	0.0270	0.9655	-0.0205	0.9725
CG32071	CG32071	1639150_at	-0.1329	0.5718	0.0242	0.8378	0.2671	0.3399	0.0786	0.9130	-0.1779	0.4514	-0.2565	0.2062	-0.0054	0.9952	0.1101	0.6605	0.1155	0.6389
CG4272	CG4272	1639151_a_at	0.2074	0.2990	0.0823	0.5248	-0.5632	0.0538	-0.2459	0.4174	0.4104	0.0229	0.6563	0.0018	0.3763	0.7215	0.2433	0.6021	-0.1330	0.8035
Ptp52F	Ptp52F	1639152_at	3.0512	0.0009	2.8927	0.0069	4.1944	0.0000	0.6280	0.4234	-0.3230	0.4762	-0.9510	0.0224	-0.8134	0.5003	-0.5194	0.3606	0.2940	0.6375
CG13343	CG13343	1639153_at	-0.3584	0.0518	0.4456	0.0157	0.7347	0.0018	0.0071	0.9931	-0.7001	0.0018	-0.7072	0.0011	-0.2172	0.6159	0.0775	0.7105	0.2948	0.1390
CG31765	CG31765	1639154_at	0.1904	0.2529	0.2009	0.3821	0.2140	0.2382	0.0322	0.9665	0.1899	0.3517	0.1578	0.3972	0.0999	0.8833	0.1083	0.7127	0.0084	0.9846
---	---	1639155_at	0.1865	0.5260	0.1210	0.4632	0.0232	0.9200	-0.3144	0.4979	-0.2847	0.2480	0.0298	0.9214	0.0946	0.9445	-0.1083	0.8464	-0.2028	0.6469
---	---	1639156_at	0.0229	0.9425	0.1629	0.3763	0.0591	0.7587	-0.1478	0.7857	-0.1062	0.6790	0.0416	0.8762	0.0482	0.9474	0.1697	0.4466	0.1214	0.6130
CG6765	CG6765	1639157_at	0.0040	0.9869	-0.1113	0.6497	-0.0329	0.8604	-0.0407	0.9482	0.0615	0.7711	0.1022	0.5545	-0.2477	0.6955	-0.1008	0.7454	0.1469	0.6050
CG13473	CG13473	1639158_at	0.0663	0.7678	0.1853	0.2667	0.2093	0.3597	-0.0502	0.9330	-0.3519	0.0511	-0.3017	0.0597	-0.0362	0.9800	-0.1607	0.6465	-0.1244	0.7381
CG4069	CG4069	1639159_at	0.3324	0.2876	0.1275	0.7273	0.0115	0.9666	0.1898	0.6791	0.8454	0.0028	0.6555	0.0052	0.1981	0.9057	0.5558	0.3421	0.3577	0.5758
Toll-7	Toll-like	1639160_at	0.0722	0.7302	0.4019	0.1269	0.1426	0.5161	-0.0965	0.8796	-0.1020	0.6821	-0.0055	0.9847	0.2397	0.7215	0.1897	0.5183	-0.0500	0.8982
---	---	1639161_at	0.2144	0.3670	0.0162	0.9216	-0.0410	0.8504	0.0872	0.8905	0.1426	0.5284	0.0554	0.8177	0.0549	0.9672	-0.0104	0.9890	-0.0653	0.8924
---	---	1639162_at	-0.3587	0.2262	-0.2219	0.4949	-0.1671	0.5425	0.1397	0.8125	-0.3746	0.1065	-0.5144	0.0217	0.1480	0.9316	-0.2058	0.7573	-0.3538	0.5488
---	---	1639163_at	-0.0945	0.6547	0.0448	0.7159	0.1902	0.3756	0.0752	0.9116	-0.0530	0.8489	-0.1282	0.5422	-0.0334	0.9816	-0.0401	0.9409	-0.0067	0.9907
CG9486	CG9486	1639164_at	-3.6997	0.0005	-0.4117	0.5127	-2.5001	0.0004	-2.1896	0.0189	-3.1911	0.0004	-1.0015	0.0302	0.1663	0.9409	0.2180	0.8070	0.0517	0.9593
Mst84Dd	Male-specific RNA	1639165_at	0.1729	0.3690	-0.0409	0.6811	0.0642	0.7892	0.1387	0.8532	0.1228	0.6936	-0.0158	0.9631	-0.1241	0.8235	-0.0406	0.9085	0.0835	0.7541
CG10677	CG10677	1639166_at	0.0250	0.9248	-0.8399	0.0395	-1.0658	0.0010	0.1597	0.7845	1.3143	0.0007	1.1546	0.0007	0.1733	0.7220	0.1867	0.3612	0.0134	0.9659
CG14487	CG14487	1639167_at	0.3396	0.1658	0.0590	0.5667	0.0524	0.7831	-0.0016	0.9975	-0.0152	0.9404	-0.0136	0.9384	0.1486	0.8087	0.0097	0.9842	-0.1389	0.6140
CG1218	CG1218	1639168_at	-0.6606	0.0074	0.0524	0.9153	0.1334	0.4557	0.1038	0.8164	-0.3784	0.0381	-0.4822	0.0085	0.0454	0.9831	0.3662	0.4600	0.3208	0.5361
CG18754	CG18754	1639169_at	0.2882	0.2503	0.0009	0.9993	-0.0826	0.6756	0.0304	0.9744	0.2971	0.1930	0.2667	0.1930	0.0795	0.9357	0.0680	0.8792	-0.0115	0.9835
Cyp4d21	Cyp4d21	1639170_at	-0.1174	0.8182	-0.0099	0.9242	0.0258	0.8869	-0.0579	0.9610	-0.2806	0.3898	-0.2227	0.4601	0.2030	0.8202	0.1528	0.7198	-0.0503	0.9215
mRpL14	mitochondrial ribo	1639171_at	-0.0953	0.7644	0.0295	0.9153	-0.1947	0.6668	-0.1205	0.8611	0.4448	0.0742	0.5652	0.0186	0.0899	0.9742	0.4997	0.4794	0.4098	0.5785
---	---	1639172_at	-0.0343	0.8558	0.0804	0.5667	0.1830	0.3184	0.0238	0.9671	-0.1243	0.4255	-0.1482	0.2770	-0.0709	0.8999	-0.0093	0.9808	0.0616	0.8095
---	---	1639173_at	0.1360	0.4699	0.0108	0.9275	-0.0260	0.8884	-0.1265	0.7327	0.1067	0.5479	0.2332	0.1128	-0.1698	0.7116	-0.1399	0.4729	0.0300	0.9096
CG32552	CG32552	1639174_at	0.2457	0.1395	-0.0472	0.6108	0.2088	0.2326	0.3018	0.2631	0.2196	0.1663	-0.0823	0.6100	-0.1320	0.8445	-0.0866	0.8075	0.0455	0.9056
---	---	1639175_s_at	0.0044	0.9833	-0.1891	0.1268	-0.1282	0.4666	0.1569	0.6130	0.0803	0.6464	-0.0766	0.6300	-0.1452	0.8215	-0.1108	0.7169	0.0344	0.9256
CG1545	CG1545	1639176_at	0.0914	0.6749	-0.0483	0.8447	-0.0961	0.6121	-0.0790	0.8889	-0.0548	0.8157	0.0242	0.9167	-0.0036	0.9970	-0.0690	0.8185	-0.0654	0.8209
IFa	SIFamide	1639177_at	0.1324	0.4188	0.0756	0.4771	0.2384	0.4436	-0.0695	0.9300	0.1340	0.5964	0.2035	0.3377	-0.0946	0.9467	-0.0481	0.9451	0.0465	0.9384
mael	maelstrom	1639178_s_at	0.4470	0.2992	-0.2812	0.5229	-0.8207	0.0319	-0.4132	0.2977	0.7718	0.0060	1.1850	0.0006	-0.0678	0.9870	0.1030	0.9467	0.1708	0.8951
CG5103	CG5103	1639179_at	0.5056	0.0114	0.1677	0.1468	0.2661	0.2711	-0.0659	0.9339	0.0972	0.7209	0.1631	0.4607	0.0071	0.9939	-0.0603	0.8185	-0.0674	0.7774
CG12505	CG12505	1639180_at	0.1750	0.8449	-0.3077	0.6241	0.0203	0.9695	0.3571	0.6202	1.6634	0.0014	1.3063	0.0024	0.0810	0.9898	1.3908	0.2707	1.3098	0.3271
CG14598	CG14598	1639181_at	-0.2078	0.4452	-0.5457	0.1335	-0.1360	0.3868	0.3190	0.4757	0.3271	0.1714	0.0081	0.9786	-0.1696	0.8202	-0.1367	0.6948	0.0330	0.9394
CG11360	CG11360	1639182_at	-0.1693	0.4857	-0.1332	0.3583	0.0261	0.9148	-0.1482	0.7328	0.4366	0.0316	0.5848	0.0055	-0.1778	0.7485	0.5247	0.0483	0.7025	0.0336
mira	Miranda	1639183_a_at	-0.1274	0.8052	0.0715	0.5451	0.0220	0.8961	0.0161	0.9943	-0.4978	0.2917	-0.5140	0.2197	-0.1096	0.8903	-0.2087	0.4943	-0.0991	0.7774
CG16787	CG16787	1639184_at	-0.4616	0.0625	0.6336	0.0608	0.6644	0.0088	0.0768	0.8979	-0.1950	0.3282	-0.2718	0.1206	0.1762	0.8541	0.8994	0.0446	0.7231	0.1017
mip130	Myb-interacting pr	1639185_at	0.2651	0.4424	0.7848	0.1837	0.8312	0.0053	0.1092	0.9116	-0.1145	0.7584	-0.2237	0.4459	0.1034	0.9514	0.4859	0.3259	0.3825	0.4654

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG7918	CG7918	1639186_at	0.0557	0.7482	-0.0003	1.0000	-0.0597	0.6938	0.0379	0.9507	0.0260	0.9123	-0.0119	0.9565	-0.0292	0.9679	-0.0018	0.9975	0.0275	0.9176
CG40103	CG40103	1639187_at	-0.0061	0.9748	-0.0881	0.5103	-0.0343	0.8780	0.0891	0.8640	0.1614	0.3945	0.0723	0.7129	0.1986	0.8283	0.2928	0.4446	0.0942	0.8482
CG3308	CG3308	1639188_at	-0.3145	0.0708	-0.1606	0.2151	-0.1469	0.5578	-0.1404	0.7271	-0.0037	0.9893	0.1367	0.4177	-0.2201	0.7230	0.0668	0.8551	0.2869	0.2960
Lap1	Lap1	1639189_at	-0.7771	0.0473	-0.4477	0.0470	-1.2620	0.0001	-0.3234	0.4690	0.0915	0.7486	0.4149	0.0580	0.4307	0.6898	0.2922	0.5209	-0.1386	0.7933
mas	masquerade	1639190_at	0.0712	0.7680	0.0529	0.7740	-0.0347	0.8317	0.0085	0.9932	-0.1610	0.4285	-0.1694	0.3474	0.0816	0.9089	0.0243	0.9521	-0.0573	0.8635
CG18155	CG18155	1639191_at	0.4076	0.0910	0.2904	0.5406	-0.1082	0.5694	-0.4857	0.3619	-0.4116	0.1726	0.0741	0.8300	-0.0028	0.9994	-0.3044	0.6276	-0.3016	0.6311
gammaTub37C	gamma tubulin	1639192_at	0.2959	0.3816	0.0122	0.9803	-0.1734	0.5777	-0.0007	0.9994	0.2874	0.3051	0.2882	0.2475	-0.0005	0.9999	-0.0605	0.9541	-0.0600	0.9479
CG1971	CG1971	1639193_a_at	-0.4251	0.1557	0.0396	0.8178	-0.1555	0.4054	-0.3680	0.4532	-0.5874	0.0345	-0.2194	0.3728	-0.2814	0.7215	-0.3383	0.2901	-0.0569	0.9006
---	---	1639194_at	-0.0287	0.8606	-0.0341	0.8415	0.1475	0.3822	0.0079	0.9934	0.0013	0.9960	-0.0066	0.9776	-0.1530	0.7953	-0.0667	0.8422	0.0863	0.7651
CycA	cyclin A	1639195_a_at	0.3730	0.8096	-2.1039	0.3331	-1.8576	0.1494	0.0071	0.9973	2.6384	0.0017	2.6312	0.0010	-0.2624	0.9848	0.2947	0.9532	0.5571	0.8977
Jon25Bii	Jonah 25B	1639196_at	0.5120	0.1760	-0.0679	0.5032	0.3590	0.0612	0.2104	0.5117	0.1994	0.2396	-0.0110	0.9591	-0.1440	0.9345	-0.3001	0.6205	-0.1561	0.8255
CG18128	CG18128	1639197_at	-0.0052	0.9886	0.1482	0.5442	0.0326	0.9038	-0.0946	0.9314	-0.1379	0.7165	-0.0434	0.9132	-0.0706	0.9291	-0.1384	0.6144	-0.0678	0.8347
CG8635	CG8635	1639198_at	0.1233	0.5661	0.7252	0.0536	1.2752	0.0015	0.2293	0.5913	-0.2132	0.3316	-0.4425	0.0298	-0.2228	0.8203	0.4362	0.2675	0.6590	0.1359
---	---	1639199_at	-0.1002	0.4996	-0.0558	0.6434	0.1071	0.4795	0.1029	0.8496	-0.0189	0.9492	-0.1218	0.5265	-0.3111	0.7220	0.0622	0.9148	0.3733	0.3293
CG14333	CG14333	1639200_at	-0.4543	0.0541	-0.0232	0.8630	-0.2732	0.1063	-0.0659	0.8801	-0.1695	0.2591	-0.1036	0.4657	-0.0112	0.9916	-0.0838	0.7921	-0.0726	0.8178
---	---	1639201_at	0.1035	0.6055	0.0000	1.0000	0.1629	0.3018	-0.0506	0.9518	0.0286	0.9298	0.0793	0.7552	-0.0249	0.9862	0.0380	0.9438	0.0629	0.8908
CG7023	CG7023	1639202_a_at	-0.2306	0.1663	0.1168	0.5588	0.3798	0.1202	0.0416	0.9620	-0.6334	0.0140	-0.6749	0.0063	-0.0748	0.9426	0.0646	0.8903	0.1394	0.6964
CG32556	CG32556	1639203_at	-1.8172	0.0227	-1.0288	0.3600	-1.0255	0.0017	0.1481	0.8156	-0.5255	0.0416	-0.6736	0.0091	0.2830	0.9457	0.0272	0.9925	-0.2558	0.8813
ppk14	pickpocket 14	1639204_at	0.0570	0.8228	-0.0274	0.9132	0.2771	0.1005	0.1472	0.7307	0.1347	0.5051	-0.0126	0.9587	-0.1102	0.9142	-0.0587	0.9162	0.0515	0.9176
tra2	transformer-2	1639205_s_at	0.2943	0.2137	0.6819	0.0419	0.6378	0.0015	0.0588	0.9308	-0.1813	0.3697	-0.2402	0.1741	-0.0232	0.9848	0.0895	0.8018	0.1127	0.7224
CG15914	CG15914	1639206_at	0.2085	0.3923	-0.2421	0.4655	-0.6534	0.0233	-0.3398	0.2500	0.5722	0.0064	0.9120	0.0005	-0.0195	0.9914	-0.0630	0.9235	-0.0435	0.9412
CG14216	CG14216	1639207_at	-0.1124	0.4637	-0.5283	0.1770	-0.6524	0.0197	-0.1900	0.6845	0.0558	0.8428	0.2458	0.2054	-0.2564	0.7142	-0.5142	0.0890	-0.2578	0.3878
Mtor	Megator	1639208_at	0.0882	0.8195	0.3755	0.3880	0.9934	0.0022	0.1652	0.7840	0.0141	0.9707	-0.1511	0.5338	-0.3657	0.7956	0.3308	0.6019	0.6965	0.2574
CG32143	CG32143	1639209_at	0.1432	0.3876	-0.2190	0.1876	0.0741	0.6522	0.0749	0.9015	0.1884	0.3457	0.1135	0.5584	-0.2497	0.5524	-0.1808	0.3397	0.0689	0.7582
Fcp3C	Follicle cell proteir	1639210_at	-0.2447	0.3586	-0.2342	0.3956	-0.3341	0.1245	-0.0188	0.9857	0.0839	0.7679	0.1027	0.6787	0.0457	0.9829	-0.0942	0.8966	-0.1399	0.8172
---	---	1639211_at	-2.3447	0.0004	-1.4789	0.1574	-2.8051	0.0007	-0.4616	0.3863	-1.4449	0.0012	-0.9833	0.0039	0.7730	0.7758	-0.5274	0.6876	-1.3003	0.2833
CG8783	CG8783	1639212_a_at	-0.2300	0.2663	0.0445	0.7253	0.1780	0.2352	0.0205	0.9777	0.0421	0.8612	0.0216	0.9235	-0.1301	0.8400	0.1696	0.5410	0.2996	0.2757
CG34019	CG34019	1639213_s_at	0.1834	0.3689	-0.0133	0.8995	0.1767	0.2472	-0.0191	0.9745	0.0637	0.7053	0.0829	0.5663	0.0548	0.9199	0.0239	0.9353	-0.0310	0.9032
---	---	1639214_at	0.0430	0.8557	0.0579	0.6190	0.0749	0.6644	0.1835	0.6936	-0.0106	0.9747	-0.1940	0.3228	0.1394	0.7712	0.0339	0.9152	-0.1055	0.6328
Csat	D.melanogaster n	1639215_at	1.3834	0.0054	1.8210	0.0049	2.0331	0.0000	0.0583	0.9387	-0.1128	0.6522	-0.1711	0.4099	-0.1417	0.9011	0.4070	0.3178	0.5487	0.2095
CG31647	CG31647	1639216_a_at	0.1128	0.4462	0.0752	0.5046	0.0924	0.5376	-0.1398	0.7303	-0.1479	0.4307	-0.0081	0.9714	-0.0136	0.9898	-0.1091	0.6605	-0.0955	0.7072
brat	Brain Tumor	1639217_s_at	1.5912	0.0199	0.7209	0.0882	0.3339	0.4003	-0.3689	0.7949	1.2047	0.0436	1.5736	0.0088	0.2598	0.8786	0.6133	0.3372	0.3534	0.6168
CG15530	CG15530	1639218_s_at	-0.0592	0.8910	-0.4691	0.0293	-0.4372	0.0592	0.0706	0.9441	0.2264	0.4475	0.1559	0.5869	-0.1920	0.8331	-0.3319	0.3733	-0.1398	0.7492
Csl4	Csl4	1639219_at	0.3190	0.2464	-0.0236	0.9687	-0.1373	0.6124	0.2438	0.6986	0.5981	0.0444	0.3543	0.1694	0.2975	0.8298	0.3056	0.6253	0.0080	0.9937
CG7716	CG7716	1639220_at	-0.0969	0.6166	-0.0130	0.9010	0.1128	0.5336	0.1071	0.7451	0.0386	0.8346	-0.0685	0.6449	-0.0481	0.9387	0.0856	0.6935	0.1337	0.5041
dan	Tintin	1639221_at	-0.1348	0.4062	0.0132	0.8983	0.2616	0.1175	-0.0340	0.9620	-0.1147	0.5750	-0.0807	0.6838	-0.0850	0.8753	0.0043	0.9931	0.0893	0.7085
CG32191	CG32191	1639222_at	-1.1437	0.0227	-2.0347	0.0050	-1.6697	0.0033	-0.2969	0.5832	-0.2968	0.2844	0.0000	0.9999	-0.5794	0.7506	-1.0515	0.1647	-0.4721	0.5676
sr	Stripe	1639223_a_at	-1.3618	0.0548	-2.4523	0.0093	-2.6445	0.0002	0.3434	0.8399	1.5519	0.0228	1.2085	0.0397	0.3651	0.6955	0.3895	0.3005	0.0244	0.9661
DsimCG9273 /// RPA2	CG9273	1639224_at	0.4097	0.3231	-0.2279	0.4343	-0.3213	0.3089	-0.2617	0.6557	0.8726	0.0082	1.1343	0.0015	-0.0938	0.9762	0.4125	0.6104	0.5063	0.5239
CG13711	CG13711	1639225_at	0.1474	0.4003	-2.0181	0.0077	-1.2924	0.0276	0.6041	0.6247	2.0633	0.0055	1.4592	0.0157	0.0274	0.9816	-0.0383	0.9225	-0.0657	0.8375
---	---	1639226_at	0.0824	0.6870	0.1053	0.5089	0.0795	0.6093	-0.0978	0.8388	-0.0287	0.9085	0.0690	0.7187	-0.0887	0.9246	-0.1491	0.6656	-0.0604	0.8884
---	---	1639227_at	-0.1099	0.5470	0.1255	0.7724	0.1702	0.4310	0.2283	0.5720	0.1000	0.6702	-0.1283	0.5225	0.0915	0.9487	0.1483	0.7760	0.0568	0.9231
Actr13E	actin-related prote	1639228_at	-0.1340	0.6015	0.4268	0.0288	0.4875	0.0231	0.0062	0.9931	-0.3387	0.0251	-0.3449	0.0146	-0.1542	0.8571	0.0833	0.8649	0.2375	0.5154
vkg	type IV collagen, a	1639229_at	-1.7017	0.0285	-2.1608	0.1609	-2.1818	0.0000	-0.1410	0.6673	0.0050	0.9826	0.1460	0.3032	-0.1234	0.9853	-0.4733	0.8121	-0.3499	0.8622
v(2)k05816	v(2)k05816	1639230_at	-0.0113	0.9927	-1.4411	0.0964	1.2197	0.1074	2.3101	0.1495	1.0949	0.2606	-1.2153	0.1599	-0.4291	0.9056	-0.4750	0.7592	-0.0459	0.9837
---	---	1639231_at	0.1994	0.3176	-0.3491	0.1092	-0.0484	0.8252	-0.0251	0.9777	0.1833	0.4303	0.2084	0.3065	-0.2598	0.5461	-0.2119	0.2748	0.0479	0.8544
SP1029	SP1029	1639232_s_at	-2.7917	0.0020	0.3004	0.4786	-2.3120	0.0007	-2.1782	0.0201	-4.1229	0.0002	-1.9447	0.0017	0.5031	0.7726	-1.0419	0.1555	-1.5451	0.0775
c12.2	c12.2	1639233_at	-0.0371	0.9522	0.1784	0.8606	0.9697	0.0011	0.1964	0.7313	-0.1111	0.7061	-0.3076	0.1774	-0.4583	0.8513	0.2066	0.8880	0.6649	0.5326
---	---	1639234_at	0.0578	0.8313	0.1096	0.4549	0.1285	0.5056	-0.0563	0.9311	-0.0440	0.8611	0.0123	0.9593	-0.0293	0.9816	0.0482	0.9144	0.0775	0.8310
trn	anon-fast-evolving	1639235_at	0.1528	0.7323	1.5744	0.0737	0.5154	0.1304	-0.1638	0.9186	-0.8294	0.0929	-0.6655	0.1292	0.9051	0.4583	0.4214	0.5158	-0.4837	0.4557
CG13946	CG13946	1639236_at	-0.2239	0.2240	0.0022	0.9950	0.2819	0.1918	0.1221	0.8483	-0.0337	0.9180	-0.1558	0.4840	-0.1123	0.8270	-0.0085	0.9838	0.1038	0.6567
CG32814	CG32814	1639237_at	-0.0961	0.6334	0.1598	0.5344	0.0680	0.7404	-0.1739	0.8466	-0.1990	0.5773	-0.0251	0.9522	0.0411	0.9717	0.0435	0.9234	0.0024	0.9965
---	---	1639238_at	0.1068	0.7511	-0.0258	0.8012	0.1798	0.3586	0.2051	0.6233	0.1701	0.4275	-0.0350	0.8873	-0.0525	0.9467	-0.0341	0.9275	0.0184	0.9570
bsh	brain-specific hom	1639239_at	0.0760	0.6911	0.1523	0.2161	0.0407	0.8513	0.2390	0.3928	0.0346	0.8684	-0.2044	0.1455	0.0804	0.9031	0.0418	0.9093	-0.0386	0.9077
---	---	1639240_a_at	0.1267	0.4494	-0.0110	0.9188	0.0093	0.9621	0.0978	0.8449	0.1508	0.4298	0.0530	0.7981	-0.0364	0.9653	-0.0259	0.9430	0.0106	0.9766
---	---	1639241_at	-0.3118	0.1298	-0.2401	0.0986	-0.2771	0.1075	0.0242	0.9761	0.0911	0.6931	0.0669	0.7612	-0.1648	0.8386	-0.0009	0.9996	0.1639	0.6476
CG6455	Motor protein	1639242_a_at	-0.6469	0.0183	-0.2901	0.4268	-0.3244	0.0505	-0.1406	0.6354	-0.5502	0.0033	-0.4096	0.0075	-0.1052	0.9405	-0.1516	0.7778	-0.0465	0.9394
Spred	Sprouty-related pr	1639243_at	-0.7968	0.1926	-0.4048	0.2176	-0.4110	0.0824	-0.2528	0.7000	-0.1934	0.5509	0.0594	0.8657	0.0070	0.9984	0.2578	0.7573	0.2509	0.7557
---	---	1639244_at	0.0771	0.7121	0.1140	0.4751	0.1609	0.4146	0.0522	0.9422	-0.1084	0.6359	-0.1606	0.3990	-0.0579	0.9545	0.0504	0.9121	0.1083	0.7520
mys	lethal myospheroi	1639245_x_at	-0.6694	0.0217	-0.1895	0.2833	-0.8927	0.0017	-0.3264	0.4335	-0.1216	0.6339	0.2048	0.3267	0.1608	0.8077	0.1922	0.4976	0.0314	0.9350
CG4714	CG4714	1639246_at	-0.2749	0.1314	-0.2201	0.1353	-0.1507	0.4546	0.0731	0.8822	0.2467	0.1414	0.1736	0.2510	-0.0164	0.9901	0.1100	0.7490	0.1264	0.6974
PHDP	Putative homeodo	1639247_at	0.0242	0.9353	-0.1859	0.2765	0.0357	0.8272	0.0823	0.9228	0.0855	0.7838	0.0032	0.9919	-0.0624	0.9489	-0.0622	0.8821	0.0002	0.9997
CG15701	CG15701	1639248_at	0.1369	0.4639	0.0162	0.8779	0.3442	0.2269	0.0256	0.9755	-0.0822	0.7303	-0.1078	0.5976	-0.0850	0.9516	-0.1192	0.8329	-0.0341	0.9548
CG7587	CG7587	1639249_at	-0.0033	0.9889	0.1163	0.4058	0.2361	0.1678	-0.0731	0.9218	-0.1433	0.5527	-0.0702	0.7777	0.0020	0.9990	-0.0916	0.7734	-0.0936	0.7554
---	---	1639250_at	-0.0078	0.9787	0.0901	0.7275	0.1382	0.5244	0.0899	0.8099	-0.0306	0.8766	-0.1205	0.3810	0.1102	0.9328	0.1231	0.8218	0.0129	0.9849
CG17462	CG17462	1639251_at	0.3890	0.3652	0.1764	0.6879	-0.4522	0.1358	-0.4726	0.2125	0.3816	0.0946	0.8542	0.0021	0.0659	0.9775	0.1696	0.8132	0.1037	0.8918
CG11876 /// DyakCG11876	CG11876	1639252_s_at	-0.0865	0.6017	0.9085	0.1576	1.0801	0.0004	-0.1247	0.8231	-1.6294	0.0002	-1.5047	0.0002	-0.3429	0.8028	-0.5365	0.3358	-0.1936	0.7745
CG11242	CG11242	1639253_at	0.0063	0.9735	-0.1178	0.4326	0.1633	0.4505	0.1054	0.7749	0.4112	0.0157	0.3058	0.0337	-0.1282	0.8833	0.3171	0.3268	0.4453	0.2037
CG32036	CG32036	1639254_at	0.0030	0.9944	-0.2355	0.2906	-0.0391	0.8390	0.1023	0.8588	0.0375	0.8941	-0.0649	0.7775	-0.1577	0.8686	-0.1181	0.8071	0.0395	0.9402

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG1516	GH06348	1639255_s_at	0.8426	0.0683	-0.5020	0.5374	0.2118	0.5523	0.2251	0.8115	1.2526	0.0057	1.0275	0.0084	-0.5212	0.8185	-0.0912	0.9508	0.4300	0.6719
CG5676	CG5676	1639256_at	0.1896	0.3501	0.4910	0.1414	0.6312	0.0119	-0.0213	0.9803	-0.2080	0.3331	-0.1867	0.3347	-0.1146	0.8903	-0.0265	0.9590	0.0881	0.8209
CG11739	CG11739	1639257_s_at	1.8512	0.0021	1.0242	0.1415	1.6011	0.0015	0.3538	0.6718	0.1576	0.7349	-0.1962	0.6262	-0.3139	0.8465	-0.7274	0.2625	-0.4136	0.5626
CG7335	CG7335	1639258_at	0.1183	0.6033	0.0212	0.8490	0.1970	0.2874	0.1176	0.8028	0.0680	0.7640	-0.0496	0.8165	-0.0082	0.9939	-0.0599	0.8593	-0.0517	0.8751
---	---	1639259_at	-0.2541	0.1066	-0.0241	0.8788	0.1511	0.5018	0.1085	0.7950	-0.2455	0.1477	-0.3540	0.0279	-0.0144	0.9895	0.0100	0.9815	0.0244	0.9395
---	---	1639260_at	0.3055	0.1316	0.4413	0.1507	0.4990	0.0217	0.1877	0.5419	0.1831	0.2542	-0.0047	0.9812	0.3016	0.6702	0.4022	0.1498	0.1006	0.7619
---	---	1639261_at	0.1331	0.4139	0.0279	0.8844	0.1388	0.5027	0.1340	0.7550	0.1212	0.5420	-0.0128	0.9570	0.0244	0.9775	0.0224	0.9462	-0.0020	0.9956
CG11669	CG11669	1639262_at	0.2927	0.4864	-0.0114	0.9147	-0.0419	0.8624	0.3209	0.3685	0.2183	0.2838	-0.1026	0.6170	0.0307	0.9913	-0.3017	0.6239	-0.3324	0.5856
---	---	1639263_at	0.0814	0.6571	0.2277	0.3738	0.1336	0.4434	-0.0846	0.8671	-0.1908	0.2897	-0.1062	0.5444	0.0835	0.8909	-0.1611	0.4952	-0.2446	0.2993
Ddx1	Dead-box-1	1639264_at	0.1410	0.4421	0.3603	0.0962	0.3542	0.0530	-0.2095	0.5735	0.0273	0.9185	0.2368	0.1632	-0.2260	0.7220	0.2550	0.3340	0.4809	0.1116
CG15536	CG15536	1639265_at	-0.0322	0.8866	0.1289	0.6882	-0.1124	0.6357	-0.0941	0.8815	0.3556	0.0982	0.4497	0.0272	-0.1062	0.8882	0.2653	0.3358	0.3716	0.2095
CG15744	CG15744	1639266_at	0.1575	0.6908	-0.4820	0.3769	-0.4674	0.0356	0.1221	0.8578	0.8834	0.0040	0.7613	0.0047	0.1289	0.9545	0.2383	0.7656	0.1094	0.9051
CG30114	CG30114	1639267_at	-0.0240	0.9220	-0.0335	0.7410	0.0590	0.7500	0.1330	0.7327	-0.0053	0.9834	-0.1383	0.3916	-0.0175	0.9893	-0.0895	0.7959	-0.0720	0.8356
CG13324 /// DyakCG13324	CG13324	1639268_at	0.2029	0.3366	-0.0230	0.8814	0.0676	0.6935	0.0033	0.9956	0.0777	0.7146	0.0744	0.7001	-0.0243	0.9831	-0.0886	0.7882	-0.0642	0.8501
Pdh	pigment cell dehyd	1639269_a_at	-2.4474	0.0050	-2.1306	0.0478	-1.8375	0.0000	0.1651	0.8281	-0.0126	0.9780	-0.1777	0.5332	-0.2263	0.9365	0.2681	0.8177	0.4944	0.6075
RpS16	Ribosomal protein	1639270_at	0.2116	0.1957	0.3686	0.1375	0.4902	0.0110	0.0590	0.8967	-0.1393	0.3616	-0.1983	0.1366	-0.0132	0.9891	0.0205	0.9501	0.0337	0.9033
---	---	1639271_at	0.0546	0.7145	0.0742	0.6594	0.2703	0.1647	-0.1031	0.8266	-0.1714	0.3536	-0.0683	0.7241	-0.1204	0.8150	-0.0962	0.6853	0.0242	0.9350
CG3739	CG3739	1639272_at	0.0235	0.9245	0.1200	0.4603	0.1258	0.5305	-0.0952	0.8556	-0.0885	0.6811	0.0066	0.9776	0.0589	0.9589	0.0348	0.9483	-0.0241	0.9597
vri	vrille	1639273_s_at	0.2153	0.6438	-0.6127	0.2482	-0.5549	0.0219	-0.0266	0.9833	0.6488	0.0325	0.6753	0.0173	-0.0333	0.9922	-0.2570	0.7923	-0.2237	0.8178
CG15859	CG15859	1639274_at	-0.0637	0.7423	0.1972	0.2814	0.0758	0.6997	0.1040	0.7964	-0.2298	0.1585	-0.3338	0.0302	0.0825	0.8890	0.0282	0.9376	-0.0544	0.8502
in	inturned	1639275_at	-1.3552	0.0021	-1.1074	0.0055	-1.3743	0.0012	0.1589	0.8757	-0.0714	0.8789	-0.2303	0.4947	0.1950	0.8049	-0.0922	0.8342	-0.2871	0.3921
CG4680	CG4680	1639276_at	0.8732	0.3531	-0.8084	0.4878	-0.5309	0.1590	0.8677	0.1162	1.7359	0.0009	0.8682	0.0107	0.6134	0.8904	-0.0733	0.9814	-0.6867	0.7161
CG31067	CG31067	1639277_at	0.1523	0.4520	-0.0140	0.9002	-0.0885	0.7281	0.0100	0.9883	0.0265	0.9045	0.0165	0.9341	0.1005	0.9199	-0.0243	0.9650	-0.1248	0.7492
ETH	ecdysis triggering	1639278_at	-0.8016	0.4427	0.1758	0.5280	0.5949	0.3058	-0.2155	0.9412	-1.4791	0.0783	-1.2636	0.0905	-0.5859	0.8395	-0.3703	0.8114	0.2156	0.8949
Rlc1	Rlc1	1639279_a_at	-0.3223	0.2119	0.0967	0.8170	-0.0419	0.8782	-0.1370	0.6927	-0.2642	0.0992	-0.1272	0.3932	-0.1008	0.9514	0.0893	0.9062	0.1901	0.7381
CG6322	CG6322	1639280_at	-0.0770	0.8717	0.1264	0.6991	0.2818	0.1446	-0.0290	0.9715	0.1774	0.4025	0.2064	0.2669	-0.1978	0.8898	0.3565	0.5199	0.5543	0.3099
Fer2	48 related 2	1639281_at	-0.0263	0.9297	-0.1495	0.5737	0.0772	0.6739	0.0095	0.9942	-0.1477	0.6239	-0.1572	0.5561	-0.0132	0.9901	-0.1263	0.6157	-0.1131	0.6576
CG9220	CG9220	1639282_at	0.1335	0.4691	0.1715	0.3869	0.0933	0.5866	-0.0449	0.9563	-0.1428	0.5490	-0.0979	0.6710	0.0557	0.9515	-0.1336	0.6598	-0.1893	0.5099
---	---	1639283_at	0.1652	0.5580	0.0315	0.7483	0.0820	0.6255	0.0207	0.9825	-0.0796	0.7642	-0.1003	0.6621	0.0169	0.9893	-0.1074	0.7200	-0.1244	0.6607
---	---	1639284_at	0.0380	0.8638	0.0355	0.7402	-0.0878	0.6114	-0.0393	0.9538	-0.0345	0.8906	0.0048	0.9840	-0.0062	0.9948	-0.1453	0.5249	-0.1391	0.5506
CG14177	CG14177	1639285_at	-0.1936	0.3301	0.0127	0.9130	0.0495	0.7973	-0.0248	0.9704	-0.1050	0.5676	-0.0802	0.6473	-0.0997	0.9342	0.1123	0.8248	0.2120	0.6111
CG31689	CG31689	1639286_s_at	0.2005	0.6819	-0.3790	0.6179	-0.2825	0.2148	-0.4223	0.1787	-0.3415	0.0749	0.0808	0.6851	-0.4513	0.8379	-0.9854	0.2610	-0.5341	0.5814
CG9877 /// DereCG9877	CG9877	1639287_at	-0.1358	0.5320	-0.0301	0.8437	0.0533	0.8053	0.0152	0.9865	0.0104	0.9756	-0.0048	0.9861	-0.0682	0.9134	-0.0580	0.8457	0.0102	0.9769
Mys45A	Mystery 45A	1639288_at	0.5454	0.0444	0.3020	0.5365	0.4444	0.0576	0.0631	0.9270	0.7253	0.0046	0.6622	0.0041	-0.1279	0.9409	0.4652	0.3749	0.5931	0.2830
---	---	1639289_at	0.9904	0.0213	0.4175	0.1268	0.7389	0.0026	-0.1736	0.8219	-0.2792	0.3631	-0.1056	0.7452	-0.3970	0.5259	-0.5835	0.0694	-0.1865	0.5585
sdt	anon-fast-evolving	1639290_at	0.3070	0.2872	0.5074	0.0223	0.2316	0.2792	-0.2299	0.4846	-0.2122	0.2261	0.0177	0.9353	0.2782	0.7220	0.1912	0.5907	-0.0870	0.8379
---	---	1639291_at	0.0255	0.9220	0.1533	0.4223	0.2209	0.4270	-0.0797	0.9445	-0.0037	0.9939	0.0760	0.8404	-0.0183	0.9898	0.1140	0.7424	0.1323	0.6853
Frq1	frequenin	1639292_at	-0.0923	0.6035	-0.1538	0.3011	-0.6169	0.0087	-0.3295	0.2097	-0.0299	0.8887	0.2996	0.0401	-0.0052	0.9964	-0.1771	0.5404	-0.1718	0.5587
---	---	1639293_at	0.1758	0.2242	-0.1290	0.4216	0.1490	0.4683	0.1525	0.6988	0.3701	0.0464	0.2177	0.1777	-0.0876	0.8521	-0.0610	0.8073	0.0266	0.9211
---	---	1639294_at	0.1255	0.3970	0.2197	0.2579	0.2494	0.1237	0.0297	0.9626	-0.0285	0.9022	-0.0582	0.7502	0.0240	0.9805	0.1136	0.6227	0.0896	0.7118
CG4734	CG4734	1639295_at	0.3881	0.3802	0.2916	0.3613	0.4327	0.0917	0.0306	0.9679	-0.1720	0.4021	-0.2026	0.2598	0.1266	0.9246	-0.1475	0.7922	-0.2741	0.5629
CG3862	RCC GEF-related	1639296_at	-0.1693	0.4173	-0.2045	0.2819	-0.1773	0.3010	-0.2356	0.4568	-0.0458	0.8348	0.1898	0.2162	-0.3583	0.6749	-0.1887	0.6100	0.1696	0.6486
CG9568 /// DyakCG9568	CG9568	1639297_at	0.1618	0.9629	-1.3496	0.0230	-2.1373	0.0043	0.0431	0.9956	-0.1096	0.9690	-0.1527	0.9471	0.2087	0.9862	-1.9090	0.4899	-2.1176	0.4464
CG3213	CG3213	1639298_at	0.2207	0.2607	0.1778	0.3737	0.4054	0.0417	0.0077	0.9943	0.0324	0.9197	0.0247	0.9294	-0.1395	0.8655	0.1525	0.6762	0.2920	0.3851
Ugt37c1	UDP-glycosyltrans	1639299_at	1.0154	0.0243	0.5326	0.0751	0.8634	0.0200	-0.0341	0.9518	-0.1060	0.5235	-0.0718	0.6566	-0.1454	0.9238	-0.4700	0.3350	-0.3246	0.5391
---	---	1639300_at	-0.0910	0.7600	-0.0107	0.9642	0.0579	0.7010	-0.0328	0.9745	-0.0822	0.7904	-0.0494	0.8675	-0.0227	0.9764	-0.0423	0.8721	-0.0196	0.9394
CG4395	CG4395	1639301_at	0.1427	0.3071	0.2632	0.1181	0.3187	0.1350	0.0000	0.9999	-0.1471	0.5804	-0.1471	0.5393	0.0001	0.9999	0.0377	0.9064	0.0376	0.8951
CG5645	CG5645	1639302_at	0.2509	0.6744	0.0963	0.8378	-0.2208	0.5527	0.0228	0.9893	0.8052	0.0431	0.7824	0.0309	0.4226	0.8461	0.5875	0.5283	0.1649	0.8934
CG9001 /// DmirCG9001	CG9001	1639303_at	0.1185	0.7079	0.1767	0.5836	-0.1297	0.7421	-0.5407	0.3433	0.1638	0.6587	0.7045	0.0246	-0.2198	0.8917	0.2461	0.7250	0.4659	0.4578
---	---	1639304_at	0.2800	0.3190	0.0346	0.8874	0.0097	0.9769	0.1213	0.8217	0.2861	0.1706	0.1648	0.3988	0.0149	0.9916	0.1350	0.7260	0.1201	0.7548
Khc-73	kinesin-73	1639305_a_at	-0.2566	0.4006	-0.3670	0.1247	-0.3410	0.0883	0.1578	0.7605	0.3818	0.0877	0.2239	0.2650	0.2008	0.8427	0.2911	0.4962	0.0903	0.8717
CG17090	CG17090	1639306_s_at	0.4319	0.5235	-0.0741	0.7554	-0.7997	0.2006	-0.3434	0.5680	0.7863	0.0190	1.1297	0.0023	0.4712	0.8806	0.3571	0.8187	-0.1141	0.9461
CG9288 /// DyakCG9288	CG9288	1639307_at	-0.3433	0.1480	0.1659	0.5456	0.0437	0.8259	-0.0378	0.9627	-0.4288	0.0516	-0.3910	0.0479	0.0505	0.9689	0.0131	0.9841	-0.0374	0.9372
mod(mdg4)	Modifier67.2	1639308_at	-0.1378	0.6390	-0.0925	0.4824	-0.0840	0.7257	0.0042	0.9956	0.0681	0.7530	0.0639	0.7446	0.0591	0.9794	0.1920	0.7588	0.1329	0.8424
CG12333	CG12333	1639309_at	-1.1099	0.0020	-1.0344	0.0882	-1.2331	0.0003	-0.1696	0.7604	-0.0116	0.9750	0.1580	0.4909	-0.0480	0.9816	0.0564	0.9411	0.1045	0.8699
CG7101	CG7101	1639310_at	0.4204	0.0369	0.0748	0.5915	0.1443	0.3962	-0.1269	0.7111	0.5457	0.0047	0.6726	0.0011	-0.0496	0.9611	0.2435	0.3878	0.2931	0.3168
---	---	1639311_at	0.0525	0.7536	0.0621	0.6681	-0.0161	0.9440	0.1088	0.8577	0.1828	0.4140	0.0740	0.7551	0.0923	0.8815	0.1224	0.6324	0.0301	0.9271
RpL22	PARP-binding pro	1639312_at	0.0569	0.7169	-0.1293	0.5883	-0.0686	0.6941	0.0507	0.9039	0.0583	0.7121	0.0076	0.9649	0.0674	0.9460	-0.0799	0.8450	-0.1473	0.6498
CG17650	CG17650	1639313_at	0.0420	0.7865	-0.0313	0.8375	0.0078	0.9647	0.0813	0.8304	0.1336	0.3679	0.0523	0.7377	0.0115	0.9895	0.0569	0.8117	0.0454	0.8474
---	---	1639314_at	0.0447	0.8864	-0.0681	0.7296	-0.0021	0.9932	0.0580	0.9295	0.0806	0.7181	0.0226	0.9235	0.0712	0.9352	0.0387	0.9307	-0.0325	0.9342
---	---	1639315_at	0.0717	0.7196	0.2840	0.1365	0.1847	0.2786	-0.0462	0.9375	-0.3288	0.0589	-0.2826	0.0677	0.0738	0.9441	0.0722	0.8738	-0.0015	0.9979
---	---	1639316_at	0.1610	0.2663	0.1026	0.3902	0.2105	0.1698	-0.1264	0.7121	-0.1839	0.2378	-0.0574	0.7305	-0.2175	0.6897	-0.0786	0.7707	0.1389	0.5523
CG7896	CG7896	1639317_at	-0.1077	0.6187	0.0049	0.9680	-0.0577	0.7427	-0.1048	0.8409	-0.2747	0.1567	-0.1700	0.3411	-0.0436	0.9689	-0.1625	0.6011	-0.1188	0.7179
Su(Tpl)	Suppressor of GM	1639318_at	-0.0850	0.8295	0.4226	0.3766	0.4277	0.0275	0.1030	0.9262	-0.0173	0.9734	-0.1203	0.7381	0.1544	0.8951	0.5563	0.1894	0.4018	0.3705
skpF	skpF	1639319_at	0.2801	0.1245	0.1785	0.4658	0.0873	0.5952	0.1661	0.6533	0.1192	0.5315	-0.0469	0.8163	0.1158	0.8940	0.0731	0.8751	-0.0427	0.9235
Ddc	dopa decarboxyla	1639320_a_at	-1.5808	0.0016	-3.0595	0.0045	-1.8808	0.0143	0.8346	0.3084	0.8118	0.0923	-0.0229	0.9697	-0.1240	0.9616	-0.4221	0.5800	-0.2981	0.7138
Tl	toll	1639321_s_at	0.1003	0.7474	-0.8191	0.0179	-0.7858	0.0073	0.1628	0.8299	0.7210	0.0212	0.5583	0.0381	0.0011	0.9996	-0.2073	0.4512	-0.2084	0.4631
etaTry	etaTrypsin	1639322_at	-0.1392	0.9366	-0.0769	0.6594	0.1570	0.3195	0.2065	0.9538	-0.8126	0.4222	-1.0191	0.2463	-0.1310	0.9816	-0.9596	0.4875	-0.8286	0.5634
TotC	Turandot C	1639323_at	3.1884	0.0272	-0.5147	0.6988	2.6558	0.0677	3.5150	0.2266	2.3172	0.1785	-1.1978	0.4660	0.2839	0.9717	-1.3803	0.5045	-1.6643	0.4132

Prokupek et al., IMB (2009), Supplementary table 2 : Differential expression data from pairwise comparisons.

			STV-SRV		ST3-SR3		ST6-SR6		SR3-SR6		SR3-SRV		SR6-SRV		ST3-ST6		ST3-STV		ST6-STV	
Gene Symbol	Gene Title	Affy ID	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR	FC	FDR
CG12684	CG12684	1639324_at	0.1016	0.5515	0.0409	0.6588	0.0014	0.9949	0.0736	0.8738	0.0433	0.8341	-0.0303	0.8740	0.0084	0.9914	-0.0305	0.9124	-0.0389	0.8726
---	---	1639325_at	0.0424	0.8314	0.0092	0.9629	0.0425	0.8091	-0.1500	0.7425	-0.3410	0.0904	-0.1910	0.2939	-0.0138	0.9901	-0.1547	0.5415	-0.1409	0.5868
kat80	katanin 80	1639326_s_at	0.1263	0.4145	-0.4370	0.2303	-0.7039	0.0084	0.0102	0.9902	0.8878	0.0012	0.8776	0.0008	0.3547	0.7266	0.2465	0.5972	-0.1082	0.8472
CG5478	CG5478	1639327_at	-0.1724	0.5085	-0.1687	0.1845	-0.0451	0.8366	-0.0199	0.9777	0.0810	0.7011	0.1010	0.5802	-0.1953	0.7953	0.0726	0.8721	0.2679	0.4051
CG2453	CG2453	1639328_at	0.0355	0.9100	0.1870	0.3556	0.4113	0.0270	-0.0797	0.9228	-0.0916	0.7570	-0.0119	0.9692	-0.2842	0.7381	-0.0152	0.9823	0.2690	0.4699
---	---	1639329_at	-0.0486	0.8295	0.0353	0.7242	-0.0694	0.6965	-0.0849	0.8738	-0.0477	0.8428	0.0371	0.8637	-0.0359	0.9620	0.0357	0.9120	0.0716	0.7728
Tao-1	Tao-1	1639330_s_at	0.8114	0.1561	0.2614	0.7118	-1.0035	0.0368	-0.3199	0.3328	1.2866	0.0004	1.6065	0.0001	0.9397	0.7230	0.6789	0.5762	-0.2608	0.8635
Gr47a	Gustatory recepto	1639331_at	-0.3436	0.0504	-0.1695	0.3942	0.1110	0.5746	0.1842	0.7161	-0.0227	0.9471	-0.2069	0.3255	-0.0255	0.9825	0.1133	0.7090	0.1388	0.6311
Rap2l	Ras-associated pr	1639332_at	-0.1589	0.3609	-0.2601	0.1791	-0.3682	0.0345	-0.1152	0.7688	-0.0002	0.9994	0.1151	0.4656	-0.0576	0.9515	-0.1089	0.7452	-0.0514	0.8949
al	aristaless	1639333_at	-0.7313	0.1097	-0.6391	0.1336	-0.1141	0.5960	0.2735	0.7134	0.2608	0.4604	-0.0128	0.9761	0.0844	0.9181	0.1312	0.6762	0.0468	0.9053
Or92a	Odorant receptor	1639334_at	0.1924	0.4345	-0.0249	0.8207	-0.1176	0.4803	-0.0822	0.9030	0.0152	0.9636	0.0975	0.6684	-0.0777	0.9238	-0.1113	0.7185	-0.0336	0.9283
---	---	1639335_at	-1.6296	0.0134	-0.8136	0.1630	-2.6579	0.0000	-0.3236	0.4551	0.2033	0.3989	0.5268	0.0203	1.4668	0.2926	0.8778	0.2190	-0.5889	0.4374
CG12522	CG12522	1639336_at	0.1355	0.5146	0.1885	0.1099	0.0071	0.9860	-0.0142	0.9838	-0.1813	0.2638	-0.1671	0.2491	-0.0081	0.9964	0.0943	0.8785	0.1024	0.8547
CG9346	CG9346	1639337_at	0.0875	0.8062	-0.0512	0.8325	0.1399	0.5511	0.2437	0.3198	0.4909	0.0051	0.2473	0.0569	0.0638	0.9689	0.2915	0.5129	0.2277	0.6231
CG11912	CG11912	1639338_at	0.1086	0.7563	-0.4038	0.2202	-0.3146	0.3836	0.1846	0.7104	0.1968	0.3961	0.0121	0.9663	-0.2252	0.8122	-0.3378	0.3802	-0.1126	0.8178
---	---	1639339_at	0.0904	0.5574	0.0452	0.7151	0.2420	0.1512	0.1217	0.8084	0.1041	0.6422	-0.0176	0.9441	-0.2283	0.6955	-0.1317	0.6189	0.0966	0.7333
---	---	1639340_at	0.1658	0.2692	0.1569	0.5270	0.0923	0.7171	-0.0416	0.9567	0.0025	0.9934	0.0441	0.8541	-0.1675	0.8122	-0.0893	0.8131	0.0782	0.8309
CG14971	CG14971	1639341_at	1.9803	0.0039	1.3806	0.0228	1.4706	0.0003	0.0943	0.9160	0.3552	0.2010	0.2609	0.3026	0.0915	0.9717	-0.1941	0.8203	-0.2857	0.6958
Ccp84Af	cuticle cluster 3	1639342_at	0.1722	0.4531	0.2339	0.3196	0.0329	0.9055	-0.1024	0.9034	-0.0743	0.8308	0.0281	0.9341	0.1763	0.7726	-0.0377	0.9275	-0.2140	0.4206
---	---	1639343_at	0.2351	0.1404	0.0674	0.7218	0.1614	0.4195	-0.0668	0.8822	0.0245	0.9089	0.0913	0.5459	0.0844	0.9365	-0.0192	0.9732	-0.1037	0.7981
CG32651	CG32651	1639344_at	0.1425	0.3925	-0.1211	0.4513	0.1306	0.3956	0.0370	0.9518	0.2634	0.1175	0.2264	0.1321	-0.1523	0.8609	-0.0421	0.9407	0.1102	0.7956
CG14669	CG14669	1639345_at	0.1239	0.4485	-0.0356	0.7349	0.0608	0.7486	0.2551	0.5249	0.2913	0.1634	0.0362	0.8846	0.1191	0.8192	0.0549	0.8501	-0.0642	0.8050
CG9526	CG9526	1639346_at	-0.2216	0.2632	0.3608	0.0675	0.5333	0.0083	0.0042	0.9956	-0.6210	0.0038	-0.6252	0.0022	-0.2773	0.7092	-0.0625	0.8906	0.2148	0.5050
Cdk4	Protein kinase-like	1639347_s_at	-1.3375	0.0021	-2.4830	0.0018	-2.1261	0.0000	0.2100	0.7368	0.4096	0.1373	0.1996	0.4439	-0.1949	0.8270	-0.6237	0.1012	-0.4288	0.2651
---	---	1639348_at	0.2154	0.2300	0.0854	0.5902	0.3586	0.1005	0.0479	0.9300	0.0296	0.8915	-0.0184	0.9252	-0.1666	0.8133	0.0957	0.7943	0.2623	0.3807
CG8173	CG8173	1639349_at	0.4488	0.4057	-0.6127	0.3565	-1.3524	0.0319	-0.6636	0.1760	0.9938	0.0058	1.6574	0.0004	-0.1863	0.9717	0.0386	0.9889	0.2250	0.9011
---	---	1639350_at	0.1360	0.7903	0.0487	0.7836	-0.0077	0.9737	0.0094	0.9937	-0.1838	0.4848	-0.1931	0.4101	0.0573	0.9816	-0.1800	0.8055	-0.2373	0.7128
Cpr97Eb	CG15884	1639351_at	0.1311	0.5689	0.0879	0.7047	-0.0319	0.8727	-0.0022	0.9979	-0.1667	0.4559	-0.1645	0.4126	0.1002	0.8692	-0.0220	0.9557	-0.1222	0.6402
---	---	1639352_s_at	0.0738	0.7300	0.0388	0.7531	-0.0398	0.8469	-0.0900	0.8949	0.0055	0.9877	0.0955	0.6877	0.0245	0.9775	0.0198	0.9526	-0.0047	0.9886
CG17032	CG17032	1639353_at	-0.4744	0.1511	0.7578	0.3524	1.9738	0.0012	-0.6910	0.2413	-2.2706	0.0004	-1.5796	0.0009	-1.9054	0.2884	-0.9607	0.2825	0.9447	0.3170
CG9797	CG9797	1639354_at	0.2323	0.2011	0.3217	0.2490	0.2932	0.0630	-0.0326	0.9603	0.0315	0.8943	0.0641	0.7311	-0.0090	0.9943	0.0486	0.9157	0.0576	0.8880
---	---	1639355_s_at	-0.0137	0.9616	0.1381	0.7055	-0.0243	0.9065	-0.1347	0.7589	0.2081	0.2719	0.3428	0.0474	-0.0058	0.9976	0.3598	0.3878	0.3657	0.3953
---	---	1639356_at	0.2274	0.1342	0.0050	0.9669	0.1502	0.3748	0.0609	0.9036	0.0842	0.6459	0.0233	0.9057	-0.0409	0.9562	-0.1792	0.4008	-0.1382	0.5437
Cypl	Cyclophilin-like	1639357_at	0.1554	0.3315	0.2939	0.3697	0.5812	0.0376	0.0458	0.9346	0.0486	0.8119	0.0028	0.9893	-0.0166	0.9898	0.3323	0.2300	0.3489	0.2397
CG8207	CG8207	1639358_at	0.3297	0.1707	0.9079	0.0258	0.9534	0.0003	-0.0800	0.8568	-0.2649	0.0963	-0.1849	0.1931	-0.0261	0.9869	0.4425	0.2103	0.4687	0.2141
CG31974	CG31974	1639359_at	2.8632	0.0008	2.2401	0.0045	3.6336	0.0001	0.8899	0.3773	0.0490	0.9538	-0.8409	0.0985	-0.4624	0.6749	-0.4458	0.3031	0.0166	0.9832
---	---	1639360_at	0.0028	0.9895	0.0519	0.6937	-0.1219	0.6779	-0.1300	0.7915	-0.0530	0.8367	0.0770	0.7176	0.1217	0.8904	0.0259	0.9624	-0.0958	0.8177
garz	gartenzwerg	1639361_a_at	0.4908	0.4478	2.6401	0.0359	2.6235	0.0003	0.1767	0.8891	-0.5269	0.2135	-0.7036	0.0670	0.4447	0.8714	1.6589	0.1299	1.2141	0.2816

Supplementary table 3: Differential expression results for microarray compared to qRT-PCR

	CG9897		CG18628		CG17239	
	Microarray	qRT-PCR	Microarray	qRT-PCR	Microarray	qRT-PCR
ST3 - STV	-0.15	-0.04	-1.57	-3.86	0.50	3.63
ST6 - STV	-0.07	0.30	-0.48	-0.12	0.44	3.06
SR3 - SRV	0.75	-1.15	0.07	0.85	1.94	2.35
SR6 - SRV	-2.86	-1.40	0.03	-0.26	-1.65	1.45
STV - SRV	3.85	3.75	-3.72	-3.42	2.29	4.97
ST3 - SR3	3.12	4.86	-4.54	-8.14	0.74	6.27
ST6 - SR6	6.74	5.44	-3.40	-3.29	4.92	6.59

The resulting log2 fold change for differential gene expression calculated using microarray and RT-PCR data. Gene expression data from RT-PCR was normalized using rp49 prior to the calculation of fold change. FC is reported as relative to the first category in the listed comparison, for example in the comparison of ST3 - STV, CG9897 is down-regulated in ST3 relative to STV. ST = spermathecae, SR = Seminal receptacle, V = virgin, 3 = 3 hours post-mating, 6 = 6 hours post-mating.

Supplementary table 4: Categories of differentially regulated genes identified by DAVID

ST3 - STV	P-Value	FDR
Downregulated in STV		
Lipid metabolic proc.	9.7E-07	1.6E-03
Electron transport	1.1E-07	1.6E-04
Upregulated in ST3		
None Significant		

ST3- ST6	P-Value	FDR
Downregulated in ST3		
None Significant		
Upregulated in ST6		
None Significant		

Sp6- SpV
Downregulated in STV
None Significant
Upregulated in ST6
None Significant

SR3 - SRV	P-Value	FDR
Downregulated in SRV		
Electron transport	6.4E-08	9.6E-05
Oxidative phosphorylation	1.9E-05	2.9E-02
Co-enzyme metabolic proc.	2.7E-05	4.4E-02
Upregulated in SR3		
Defense response to bacterium	3.2E-11	4.8E-08
Protein kinase regulator activity	1.9E-10	2.7E-07
Gamete generat.	3.6E-08	5.9E-05
Cell cycle proc.	2.9E-07	4.3E-04
Fertilization	2.2E-06	3.3E-03
Regulation of Translation	7.3E-07	1.1E-03
Peptidoglycan metabolic proc.	2.8E-05	4.5E-02

SR3 - SR6	P-Value	FDR
Downregulated in SR3		
Serine-type peptidase activity	9.4E-06	1.7E-02
Upregulated in SR6		
None significant		

SR6 - SRV	P-Value	FDR
Downregulated in SRV		
Electron Transport	2.2E-08	3.2E-05
Upregulated in SR6		
Gamete generation	8.7E-18	1.3E-14
Cell cycle proc.	7.2E-15	1.1E-11
Protein kinase regulator activity	2.3E-12	3.0E-09
Asymmetric cell divison	7.1E-10	1.1E-06
Defense response to bacterium	6.1E-07	9.2E-04
Regulation of translation	1.0E-08	1.5E-05
DNA metabolic proc.	1.7E-09	2.8E-06
Fertilization	1.2E-05	1.8E-02
Embryonic development	7.6E-07	1.1E-03
Nervous system development	1.4E-05	2.2E-02
Organelle localization	7.9E-05	1.2E-01
Chromatin assembly	5.0E-05	8.2E-02

STV - SRV	P-Value	FDR
Upregulated in STV		
Carboxylic acid metabolic proc.	2.3E-09	3.1E-06
Electron transport	3.0E-05	4.5E-02
Lipid metabolic proc.	1.0E-06	1.5E-03
Heterocycle metabolic proc.	9.2E-06	1.4E-02
Arginine metabolic proc.	8.5E-05	1.4E-01
Monosaccharide metabolic proc.	1.4E-04	2.3E-01
Upregulated in SRV		
Organ development	1.0E-11	1.7E-08
Transmission of nerve impulse	2.2E-08	3.7E-05
Ion transport	8.8E-08	1.4E-04
Nervous system development	3.8E-07	6.3E-04
Regulation of developmental proc.	1.1E-04	1.6E-01
Embryonic development	3.4E-06	5.1E-03

ST3 - SR3	P-Value	FDR
Upregulated in ST3		
Macromolecule biosynthetic proc.	1.2E-05	1.7E-02
Protein targeting to ER	4.1E-05	6.6E-02
Upregulated in SR3		
Ion transmembrane transporter activity	7.3E-07	1.0E-03
Protein kinase CK2 regulator activity	1.1E-08	1.6E-05
Tissue development	1.6E-11	2.6E-08
Cell motility	2.5E-05	3.8E-02
Embryonic development	9.5E-05	1.4E-01

ST6 - SR6	P-Value	FDR
Upregulated in ST6		
Carboxylic acid metabolic proc.	8.8E-17	1.8E-13
Lipid metabolic proc.	8.3E-07	1.3E-03
Monosaccharide metabolic proc.	3.6E-07	5.9E-04
Cellular biosynthetic proc.	4.7E-07	7.1E-04
Coenzyme metabolic proc.	2.2E-05	3.5E-02
Heme binding	3.4E-05	4.5E-02
Peptidase activity	8.5E-05	1.1E-01
Upregulated in SR6		
Cell fate commitment	3.7E-08	6.1E-05
Transmission of nerve impulse	6.8E-07	1.1E-03
Regulation of developmental proc.	7.3E-09	1.1E-05
Open tracheal system development	8.0E-08	1.3E-04
Chemosensory behavior	1.0E-06	1.5E-03
Ion transport	1.0E-05	1.5E-02
Protein kinase CK2 regulator activity	1.9E-06	2.7E-03
Regulation of signal transduction	4.6E-05	6.9E-02
Embryonic development	6.5E-12	9.7E-09

DAVID categorization of differentially expressed genes (abs (FC) ≥ 2, FDR ≤ 0.05). Categories listed are considered to be overrepresented compared to chance (p ≤ 0.05, FDR ≤ .25). ST = spermathecae, SR = Seminal receptacle, V = virgin, 3 = 3 hours post-mating, 6 = 6 hours post-mating.

Supplementary table 5: Signifcantly upregulated pathways analyzed by Gene Set Enrichmen Analysis (GSEA).

			ST3_STV		ST6_STV		ST6_ST3		SR3_SRV		SR6_SRV		SR6_SR3		STV_SRV		ST3_SR3		ST6_SR6	
KEGG ID	Specific Pathway	General Pathway	down	up	down	up	down	up	down	up	down	up	down	up	SR up	ST up	SR up	ST up	SR up	ST up
DME00252	Alanine and aspartate metabolism	amino acid metabolism	X								X									X
DME00330	Arginine and proline metabolism	amino acid metabolism	X		X															X
DME00251	Glutamate metabolism	amino acid metabolism	X																	
DME00260	Glycine, serine and threonine metabolism	amino acid metabolism	X					X			X		X					X		X
DME00340	Histidine metabolism	amino acid metabolism	X					X			X									
DME00310	Lysine degradation	amino acid metabolism																		X
DME00271	Methionine metabolism	amino acid metabolism											X							
DME00360	Phenylalanine metabolism	amino acid metabolism			X			X		X	X		X				X			X
DME00380	Tryptophan metabolism	amino acid metabolism	X					X			X		X							X
DME00350	Tyrosine metabolism	amino acid metabolism	X		X			X		X	X		X				X			X
DME00220	Urea cycle and metabolism of amino groups	amino acid metabolism						X			X		X							X
DME00290	Valine, leucine and isoleucine biosynthesis	amino acid metabolism										X								
DME00280	Valine, leucine and isoleucine degradation	amino acid metabolism						X												X
DME00960	Alkaloid biosynthesis II	biosynthesis of secondary metabolites	X		X			X		X	X		X							X
DME00903	Limonene and pinene degradation	biosynthesis of secondary metabolites	X		X			X		X										X
DME00530	Aminosugars metabolism	carbohydrate metabolism																		X
DME00650	Butanoate metabolism	carbohydrate metabolism	X										X							X
DME00020	Citrate cycle (TCA cycle)	carbohydrate metabolism	X		X				X		X									X
DME00052	Galactose metabolism	carbohydrate metabolism	X								X		X							X
DME00010	Glycolysis / Gluconeogenesis	carbohydrate metabolism	X					X			X		X							X
DME00630	Glyoxylate and dicarboxylate metabolism	carbohydrate metabolism	X		X															X
DME00562	Inositol phosphate metabolism	carbohydrate metabolism													X					
DME00040	Pentose and glucuronate interconversions	carbohydrate metabolism	X																	
DME00030	Pentose phosphate pathway	carbohydrate metabolism	X																	X
DME00640	Propanoate metabolism	carbohydrate metabolism	X		X			X												X
DME00620	Pyruvate metabolism	carbohydrate metabolism	X								X		X							X
DME00500	Starch and sucrose metabolism	carbohydrate metabolism	X		X			X												X
DME04110	Cell cycle	cell growth and death										X								
DME00910	Nitrogen metabolism	energy metabolism	X		X															X
DME00190	Oxidative phosphorylation	energy metabolism	X		X				X		X		X		X		X			X
DME00720	Reductive carboxylate cycle (CO2 fixation)	energy metabolism	X		X						X									
DME03050	Proteasome	folding, sorting and degradation						X												X
DME03060	Protein export	folding, sorting and degradation				X				X										X
DME04130	SNARE interactions in vesicular transport	folding, sorting and degradation								X		X								
DME04120	Ubiquitin mediated proteolysis	folding, sorting and degradation				X		X												
DME01032	Glycan structures - degradation	glycan biosynthesis and metabolism	X		X			X												
DME00531	Glycosaminoglycan degradation	glycan biosynthesis and metabolism	X		X			X							X		X			
DME00563	Glycosylphosphatidylinositol(GPI)-anchor biosynthesis	glycan biosynthesis and metabolism						X												
DME00510	N-Glycan biosynthesis	glycan biosynthesis and metabolism				X		X		X										X
DME00511	N-Glycan degradation	glycan biosynthesis and metabolism	X		X															
DME00592	alpha-Linolenic acid metabolism	lipid metabolism								X		X								
DME00150	Androgen and estrogen metabolism	lipid metabolism	X		X			X												
DME00120	Bile acid biosynthesis	lipid metabolism	X					X												
DME01040	Biosynthesis of unsaturated fatty acids	lipid metabolism																		
DME00565	Ether lipid metabolism	lipid metabolism						X		X										X
DME00061	Fatty acid biosynthesis	lipid metabolism											X							
DME00071	Fatty acid metabolism	lipid metabolism	X		X															X
DME00561	Glycerolipid metabolism	lipid metabolism	X										X							X
DME00564	Glycerophospholipid metabolism	lipid metabolism						X		X										X

			ST3_STV		ST6_STV		ST6_ST3		SR3_SRV		SR6_SRV		SR6_SR3		STV_SRV		ST3_SR3		ST6_SR6	
KEGG ID	**Specific Pathway**	**General Pathway**	**down**	**up**	**down**	**up**	**down**	**up**	**down**	**up**	**down**	**up**	**down**	**up**	**SR up**	**ST up**	**SR up**	**ST up**	**SR up**	**ST up**
DME00591	Linoleic acid metabolism	lipid metabolism								X	X	X	X							
DME00600	Sphingolipid metabolism	lipid metabolism																X		X
DME00072	Synthesis and degradation of ketone bodies	lipid metabolism											X							X
DME00790	Folate biosynthesis	metabolism of cofactors and vitamins	X								X		X							X
DME00760	Nicotinate and nicotinamide metabolism	metabolism of cofactors and vitamins	X						X		X									
DME00670	One carbon pool by folate	metabolism of cofactors and vitamins																X		X
DME00860	Porphyrin and chlorophyll metabolism	metabolism of cofactors and vitamins	X					X												
DME00440	Aminophosphonate metabolism	metabolism of other amino acids						X												
DME00410	beta-Alanine metabolism	metabolism of other amino acids	X																	
DME00460	Cyanoamino acid metabolism	metabolism of other amino acids			X										X					X
DME00480	Glutathione metabolism	metabolism of other amino acids	X		X															
DME00450	Selenoamino acid metabolism	metabolism of other amino acids			X										X		X			
DME00430	Taurine and hypotaurine metabolism	metabolism of other amino acids			X										X					
DME00230	Purine metabolism	nucleotide metabolism				X		X		X		X								
DME00240	Pyrimidine metabolism	nucleotide metabolism				X							X							X
DME03410	Base excision repair	replication and repair				X				X		X								
DME03030	DNA replication	replication and repair		X		X		X		X		X		X						
DME03430	Mismatch repair	replication and repair				X		X		X		X								
DME03420	Nucleotide excision repair	replication and repair				X		X		X		X								
DME04070	Phosphatidylinositol signaling system	signal transduction													X					
DME03020	RNA polymerase	transcription				X		X		X			X							X
DME00970	Aminoacyl-tRNA biosynthesis	translation				X		X				X								
DME00624	1- and 2-Methylnaphthalene degradation	xenobiotics biodegradation and metabolism	X		X			X		X	X		X							X
DME00632	Benzoate degradation via CoA ligation	xenobiotics biodegradation and metabolism	X		X			X		X	X		X							X
DME00930	Caprolactam degradation	xenobiotics biodegradation and metabolism	X		X															X
DME00982	Drug metabolism - cytochrome P450	xenobiotics biodegradation and metabolism	X		X			X												
DME00983	Drug metabolism - other enzymes	xenobiotics biodegradation and metabolism	X																	
DME00642	Ethylbenzene degradation	xenobiotics biodegradation and metabolism	X		X			X		X	X		X							X
DME00361	gamma-Hexachlorocyclohexane degradation	xenobiotics biodegradation and metabolism	X								X		X							X
DME00980	Metabolism of xenobiotics by cytochrome P450	xenobiotics biodegradation and metabolism	X		X															
DME00626	Naphthalene and anthracene degradation	xenobiotics biodegradation and metabolism	X								X									X
DME00643	Styrene degradation	xenobiotics biodegradation and metabolism			X															X
DME00710	Carbon fixation		X		X			X	X		X									X
DME00940	Phenylpropanoid biosynthesis									X										

Pathways identified using GSEA as significantly differentially expressed (FDR ≤ 0.25). The direction of regulation is relative to the first category in each comparison, for example in the comparison of ST3_STV the pathway of amino acid metabolism is down-regulated in ST3. down = significantly down-regulated; up = significantly up-regulated; ST = spermathecae, SR = Seminal receptacle, V = virgin, 3 = 3 hours post-mating, 6 = 6 hours post-mating.