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# Impulsiveness Mediates the Association between *GABRA2* SNPs and Lifetime Alcohol Problems

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## Abstract

Genetic variants in *GABRA2* have previously been shown to be associated with alcohol measures, electroencephalography (EEG)  $\beta$  waves and impulsiveness-related traits. Impulsiveness is a behavioral risk factor for alcohol and other substance abuse. Here, we tested association between 11 variants in *GABRA2* with NEO-impulsiveness and problem drinking. Our sample of 295 unrelated adult subjects was from a community of families with at least one male with DSM-IV alcohol use diagnosis, and from a socioeconomically comparable control group. Ten *GABRA2* SNPs (single-nucleotide polymorphisms) were associated with the NEO-impulsiveness ( $P < 0.03$ ). The alleles associated with

higher impulsiveness correspond to the minor alleles identified in previous alcohol dependence studies. All ten SNPs are in linkage disequilibrium (LD) with each other and represent one effect on impulsiveness. Four SNPs and the corresponding haplotype from intron 3 to intron 4 were also associated with Lifetime Alcohol Problems Score (LAPS,  $P < 0.03$ ) (not corrected for multiple testing). Impulsiveness partially mediates (22.6% average) this relation between *GABRA2* and LAPS. Our results suggest that *GABRA2* variation in the region between introns 3 and 4 is associated with impulsiveness and this effect partially influences the development of alcohol problems, but a direct effect of *GABRA2* on problem drinking remains. A potential functional SNP rs279827, located next to a splice site, is located in the most significant region for both impulsiveness and LAPS. The high degree of LD among nine of these SNPs and the conditional analyses we have performed suggest that all variants represent one signal.

**Keywords:** alcohol problems, *GABRA2*, impulsiveness, mediation, single-nucleotide polymorphisms

Impulsivity has repeatedly been identified as an important determinant of alcohol use and problems (Dick et al. 2010; Lejuez et al. 2010; Zucker et al. 2011) (unless otherwise differentiated, we hereafter refer to “alcohol use and problems” as alcohol involvement). According to Whiteside and Lynam (2001) trait Impulsivity is a multidimensional behavioral construct which includes at least four different component traits: urgency (impulsiveness in NEO\_PI), sensation seeking, lack of premeditation, and lack of perseverance. These subscales show differential relations with alcohol involvement. Specific to alcohol problems, Magid and Colder (2007) found in a group of nonabstaining undergraduates that those with both high scores on the UPPS (Whiteside et al. 2005) urgency subscale (impulsiveness in the NEO) and low scores on perseverance had a higher level of alcohol problems. Both the urgency scale in the UPPS impulsivity scale and the NEO-impulsiveness (Costa & McCrae, 1992) measure impulsive behaviors under conditions of negative affect which can be motivated by a coping strategy involving use of alcohol to deal with emotional distress, often with disregard for negative consequences (Dawe et al. 2004).

It is possible that impulsiveness and alcohol involvement may share a common underlying biological pathway such that particular genetic variants are associated with increased risk for alcohol involvement and that this effect is mediated by impulsiveness. The identification of the genetic risk for both impulsive behavior and alcohol involvement may help to identify the biological mechanisms underlying these complex traits. We previously reported (Villafuerte et al. 2012) an association between *GABRA2* and impulsiveness in a sample enriched for alcoholism. Gamma-aminobutyric acid (GABA), an inhibitory neurotransmitter in the CNS, has a role in risk for developing alcohol use disorder (AUD) through the fast-acting receptor complex,  $GABA_A$  (Grobin et al. 1998; Krystal et al. 2006). Repeated alcohol exposure affects the GABA system (Grobin et al. 1998) through binding sites at the GABA receptors, reducing neural inhibitory action. Genetic variation in *GABRA2* has been reproducibly associated with both alcoholism (Agrawal et al. 2006; Bauer et al. 2007; Covault et al. 2004; Edenberg et al. 2004; Enoch et al. 2006; Fehr et al. 2006; Lappalainen et al. 2005; Lind et al. 2008a,b; Pierucci-Lagha et al. 2005; Soyka et al. 2008) and an electroencephalography (EEG) measure, the  $\beta$  frequency band (Edenberg et al. 2004). Alcoholics (Rangaswamy et al. 2002) and their at-risk offspring (Rangaswamy et al. 2004) have increased power in the  $\beta$  frequency band (13–28 Hz). Furthermore, *GABRA2*

has been associated with childhood conduct disorder symptoms (Dick et al. 2006) and trajectories of externalizing behavior (Dick et al. 2009). The single-nucleotide polymorphisms (SNPs) reported in these studies expand a region with two haplotype blocks in high linkage disequilibrium (LD) facilitating allele comparison across studies. The majority of studies on alcohol dependence or abuse (Agrawal et al. 2006; Bauer et al. 2007; Covault et al. 2004; Edenberg et al. 2004; Enoch et al. 2006; Fehr et al. 2006; Lappalainen et al. 2005; Lind et al. 2008a,b; Pierucci-Lagha et al. 2005; Soyka et al. 2008) reported the minor allele associated with the disorder.

Here we report genetic associations of eleven SNPs in *GABRA2* with both impulsiveness and the Lifetime Alcohol Problems Score (LAPS), a composite index of problem alcohol use over the life course, and carry out an analysis of the role of impulsiveness as a mediator of the relationship between *GABRA2* SNPs and LAPS. The study was carried out on the adult subsample of a population-based study of families which had at least one male with DSM-IV AUD diagnosis, or were from a comparable group of control families (Zucker et al. 2000).

## Materials and methods

### *Subjects and assessment*

The sample consisted of 295 (167 females) biologically unrelated adult subjects from the Michigan Longitudinal Study (MLS) who were genotyped for eleven *GABRA2* SNPs and for whom NEO-PI-R and the LAPS data were available. This is an ongoing multiwave, community-recruited prospective study of families of men with a drunk-driving conviction and AUD diagnosis who were living with a 3- to 5-year-old son/daughter and the biological mother at time of recruitment (mean age 32; range 22–46 at baseline). The study began recruitment in 1985. In addition, control families without a history of substance abuse were recruited from the same or socioeconomically comparable neighborhoods. Families identified during the community canvass for controls who also had an AUD diagnosis were recruited as well (Zucker et al. 1996). For this study only the parents were selected, as both personality traits are more stable in adulthood and alcohol problems are more evident compared to the youth population. One hundred twenty-five subjects (51 females) had a DSM-IV lifetime alcohol dependence/abuse diagnosis and 170 (116 females) did not have this diagnosis. All subjects were unrelated and represent parents and/or partners. The great majority were of Caucasian origin, with 3.0% (9) of other ethnicity (1 African American, 2 Native American, and 6 Hispanic-Caucasian). All subjects were extensively assessed at 3-year intervals with behavioral and alcohol measures appropriate for age. Written informed consent was obtained from all participants after the nature of the study had been explained to them; the protocol was approved by the Institutional Review Board at the University of Michigan.

The LAPS (Zucker 1991; Zucker et al. 1997) is a time-based, multidimensional measure of problem alcohol involvement which scales the extent to which alcoholism-related symptomatology has been salient over the life course. Lifetime Alcohol Problems Score is constructed of three component subscores which assess onset of first symptom (age of first drunkenness), variety of symptomatology (number of different alcohol related difficulties

at any time), and life-invasiveness of the symptoms (index of life course duration of all drinking-related problems from onset to present, corrected for period of risk exposure). Lifetime Alcohol Problems Score is the sum of these three (standardized) component scores. For the number of alcohol-related difficulties, questions included missing school, lost friends, divorce or separation, getting fired or laid off, received ticket for drunk driving, had a car accident, kept drinking after promising to stay sober, number of times being admitted at the hospital among others. The index of life course duration of drinking related problems is the age at which these items occurred for the first and most recent times. Lifetime Alcohol Problems Score is an effective indicator to scale the extent of alcoholic risk load in preclinical stages of development for offspring, and for adults it is a metric of chronicity and severity. Measures of validity included discriminant analyses which revealed expected associations between LAPS and measures of cognitive functioning, family relationships, self-concept, and temperament (Zucker 1990; Zucker 1991; Zucker et al. 1997). We used the maximum LAPS score across waves T1–T6 and used this variable for analysis.

Personality traits were assessed using the NEO-PI-R questionnaire (Costa & McCrae, 1992) at all waves within a period of approximately 15 years. Impulsiveness is a facet from the neuroticism domain that measures the tendency to act on cravings and urges in response to distress along with later regret. As the scores on the impulsiveness facet from the neuroticism domain did not differ significantly across assessment waves, we composed a new dependent variable for the impulsiveness facet by averaging the individual scores across the different data waves (T4–T6). Subjects with Lifetime AUD diagnosis have higher impulsiveness score ( $16.5 \pm 4.0$ ;  $P = 0.0003$ ) than those without AUD diagnosis ( $14.8 \pm 3.9$ ).

In summary, our sample consists of 295 subjects for which genotype, LAPS, and impulsiveness data were available. The correlation between impulsiveness and LAPS is significant ( $0.193$ ,  $P = 0.001$ ). Also, the sample includes 110 couples (220 subjects). We found no significant correlation between husband and wife variables for both impulsiveness and LAPS. The presence of dyads does not affect standard errors in the model.

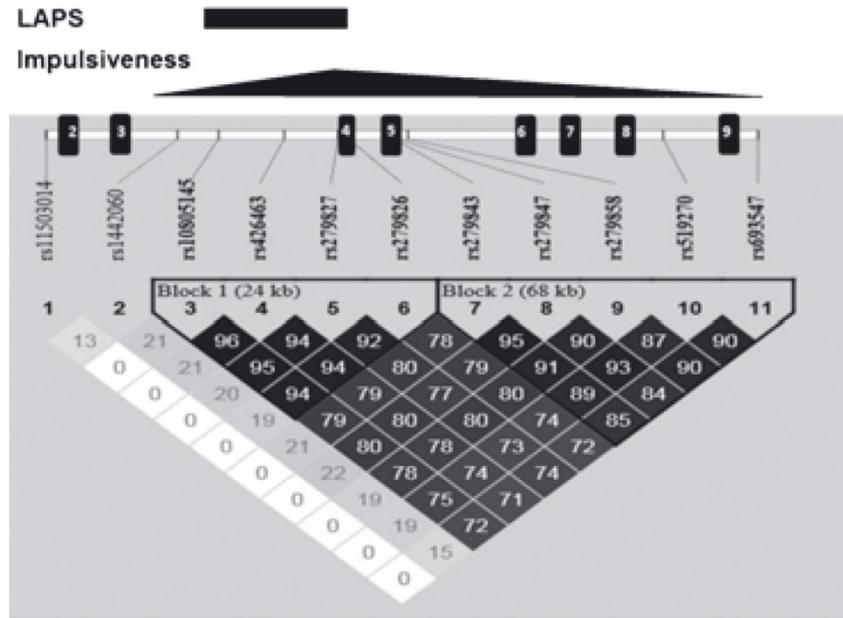
### *Single-nucleotide polymorphism genotyping*

We used the Illumina Addiction biology SNP array designed by Hodgkinson et al. (Hodgkinson et al. 2008). The panel includes SNPs from 130 candidate genes for alcoholism, addictions, and disorders of mood and anxiety and is genotyped using the Illumina GoldenGate platform (Illumina, Inc., San Diego, California, USA). Twelve SNPs from *GABRA2* were included in the panel. Three SNPs (rs10008315, rs9291283, and rs7678520) that were rare or had low call rates were excluded. In addition, we included two *GABRA2* SNPs, rs279826 (intron 4) and rs279858 (exon 5 and K132K), genotyped by Taqman (Villafuerte et al. 2012) previously associated with both alcohol measures and impulsiveness (Villafuerte et al. 2012). We included duplicates (78 for the array and 12 for the Taqman assay), and no discrepancies were observed. In summary, 11 SNPs are reported in the analyses.

Linkage disequilibrium (LD) between markers was calculated with Haploview (Barrett et al. 2005). All SNPs were in Hardy-Weinberg equilibrium.

Haplotypes were constructed for four SNPs (rs10805145, rs426463, rs279827, and rs279826) in high LD ( $LD > 0.93$ ) located in the proximal block (Figure 1). Because of the high LD among these SNPs, haplotypes were estimated manually and without ambiguity

for 288 subjects out of the 295 participants. Two major haplotypes, AAAA (54.3%) and the complementary CCGG (43.3%), were estimated without ambiguity for 288 subjects and were coded with 0, 1, and 2 if they have 0, 1, or 2 copies of the minor (risk) haplotype (CCGG), respectively.



**Figure 1.** Location and LD of eleven genotyped SNPs within the *GABRA2* gene (not drawn to scale). The black boxes depict the exons. The thicker areas in the triangles indicate the strongest association signal for both impulsiveness and LAPS.

### Statistical analyses

#### Association analyses

General Linear Models (univariate) in SPSS was used to predict both impulsiveness and LAPS from *GABRA2* SNPs. Covariates in the model were gender, age, race, and DSM-IV diagnosis of AUD. To test the independent effect of SNP rs1442060, not in LD with the other SNPs, rs279827 was included as a covariate. Independent *t*-tests were used to compare impulsiveness and LAPS differences between subjects with and without lifetime AUD diagnosis.

#### Multiple testing

We used the Single-Nucleotide Polymorphism Spectral Decomposition (SNPSpD) test, a simple correction for multiple testing of SNPs in LD with each other, on the basis of the spectral decomposition (SpD) of matrices of pairwise LD between SNPs. The user-friendly Web interface (<http://gump.qimr.edu.au/general/daleN/SNPSpD/>) (Nyholt, 2004) provides the effective number of independent marker loci and experiment-wide significance

threshold required to keep type I error rate at 5%. For this study, including 11 SNPs, the effective number of independent marker loci is 5 and the experiment-wide significance threshold was 0.01.

#### *Mediation analyses*

To test the indirect effect, bootstrapping procedures in AMOS were used to examine whether impulsiveness mediated the relation between GABRA2 variants and LAPS. Bootstrapping (MacKinnon et al. 2007) is a nonparametric method based on resampling with replacement done 2,000 times. We used a 90% confidence interval. We tested the indirect effect of the four SNPs that were associated with both LAPS and impulsiveness.

## **Results**

### ***GABRA2 variants and the impulsiveness facet of the neuroticism domain (NEO-PI)***

Eleven SNPs from the GABRA2 gene spanning from intron 1 to intron 9 were analyzed with both the NEO-impulsiveness and the LAPSs in the 295 adult subjects. We found 10 SNPs associated with the NEO-impulsiveness ( $P < 0.03$ ) (Table 1). Nine of these associated SNPs were in LD with each other ( $r^2$  from 0.71 to 0.96) while one SNP, rs1442060, showed LD ( $r^2$ ) values of less than 0.21 ( $D'$  of 0.51) with the other nine SNPs. These polymorphisms explain between 2.6% and 4.8% of the total variance of impulsiveness. One of the strongest associated SNPs, rs279827 (impulsiveness ( $P = 0.001$ ), is located next to a splice acceptor site (Figure 1). Similar results were obtained for the Caucasian group ( $N = 286$ ) providing additional evidence that race does not confound our results. Four out of the 10 associated SNPs did not pass multiple correction. These SNPs are located between intron 4 and intron 9.

**Table 1.** Means for impulsiveness by *GABRA2* SNPs and haplotypes

Markers	Position	Minor: major allele/MAF	Number of minor alleles* mean $\pm$ SEM (N)			Effect size %	Signif- icance
			0	1	2		
rs11503014	Intron 1	C:G/0.30	15.73 $\pm$ 0.34 (141)	15.09 $\pm$ 0.35 (131)	16.43 $\pm$ 0.83 (23)		0.211
rs1442060	Intron 3	A:G/0.45	14.68 $\pm$ 0.51 (90)	15.13 $\pm$ 0.33 (144)	16.63 $\pm$ 0.42 (61)	3.7	0.004
rs10805145	Intron 3	C:T/0.46	14.7 $\pm$ 0.43 (83)	15.33 $\pm$ 0.31 (151)	17.0 $\pm$ 0.50 (61)	4.2	0.002
rs426463	Intron 3	C:A/0.46	14.62 $\pm$ 0.43 (84)	15.36 $\pm$ 0.32 (153)	17.14 $\pm$ 0.51 (58)	4.8	0.001
rs279827	Intron 3	G:A/0.46	14.58 $\pm$ 0.42 (86)	15.41 $\pm$ 0.32 (148)	17.01 $\pm$ 0.5 (61)	4.5	0.001
rs279826	Intron 4	G:A/0.46	14.62 $\pm$ 0.43 (84)	15.35 $\pm$ 0.32 (150)	17.06 $\pm$ 0.50 (61)	4.6	0.001
rs279843	Intron 4	T:G/0.44	14.64 $\pm$ 0.41 (91)	15.54 $\pm$ 0.32 (149)	16.81 $\pm$ 0.53 (55)	3.5	0.006
rs279847	Intron 4	A:C/0.44	14.75 $\pm$ 0.411 (92)	15.47 $\pm$ 0.33 (146)	16.78 $\pm$ 0.52 (57)	3.1	0.010
rs279858	Exon 5	G:A/0.43	14.75 $\pm$ 0.41 (94)	15.50 $\pm$ 0.33 (147)	16.78 $\pm$ 0.54 (54)	3.0	0.012
rs519270	Intron 8	T:C/0.43	14.85 $\pm$ 0.40 (96)	15.46 $\pm$ 0.33 (144)	16.72 $\pm$ 0.53 (55)	2.6	0.021
rs693547	Intron 9	T:A/0.43	14.83 $\pm$ 0.40 (97)	15.46 $\pm$ 0.33 (143)	16.78 $\pm$ 0.53 (55)	2.8	0.015
rs10805145- rs279826	Intron 3- Intron 4	CCGG/TAAA	14.70 $\pm$ 0.43 (83)	15.41 $\pm$ 0.32 (147)	17.14 $\pm$ 0.51 (58)	4.6	0.001

MAF, minor allele frequency.

\*Impulsiveness: NEO-PI impulsiveness facet from the neuroticism domain, raw scores.

To determine the interdependence of the SNPs, we repeated the association test using one of the most significantly, rs279827, as a conditional factor (covariate). None of the SNPs were longer significant indicating that there are not two or more independent SNPs influencing the association of *GABRA2* on impulsiveness. Figure 1 depicts the location of the 10 associated SNPs highlighting the strength of the association with impulsiveness around SNP rs279827.

Covariates in the model (age, gender and race) were not significant and were not included in the final analysis. However, as expected, subjects with an AUD diagnosis had significantly higher impulsiveness scores (16.5  $\pm$  0.40, mean  $\pm$  SEM;  $P = 0.0003$ ) compared to subjects with no AUD diagnosis (14.8  $\pm$  3.9, mean  $\pm$  SEM). In the full genetic model, lifetime alcohol use diagnosis as a binary covariate (diagnosis vs. no diagnosis) was significant ( $P < 0.002$ ).

Homozygotes for the minor haplotype (CCGG/CCGG) showed significantly higher scores of impulsiveness ( $P = 0.001$ ) (Table 2) but the haplotype was not a better variable than the most significant SNP alone.

**Table 2.** Means for Lifetime Alcohol Problems Score (LAPS) by *GABRA2* SNPs and haplotypes

Markers	Position	Minor allele	Number of minor alleles*			Effect size %	Significance
			mean $\pm$ SEM (N)				
			0	1	2		
rs11503014	Intron 1	C	10.55 $\pm$ 0.55 (141)	10.53 $\pm$ 0.23 (131)	10.30 $\pm$ 0.23 (23)		0.746
rs1442060	Intron 3	A	10.39 $\pm$ 0.28 (90)	10.57 $\pm$ 0.22 (144)	10.10 $\pm$ 0.34 (61)		0.487
rs10805145	Intron 3	C	9.80 $\pm$ 0.29 (83)	10.55 $\pm$ 0.21 (151)	10.93 $\pm$ 0.34 (61)	2.4	0.027
rs426463	Intron 3	C	9.82 $\pm$ 0.29 (84)	10.53 $\pm$ 0.21 (153)	11.00 $\pm$ 0.35 (58)	2.5	0.025
rs279827	Intron 3	G	9.82 $\pm$ 0.28 (86)	10.56 $\pm$ 0.22 (148)	10.93 $\pm$ 0.34 (61)	2.4	0.029
rs279826	Intron 4	G	9.82 $\pm$ 0.29 (84)	10.53 $\pm$ 0.21 (150)	11.00 $\pm$ 0.34 (61)	2.5	0.027
rs279843	Intron 4	T	11.0 $\pm$ 0.36 (91)	10.50 $\pm$ 0.22 (149)	10.0 $\pm$ 0.28 (55)		0.072
rs279847	Intron 4	A	11.02 $\pm$ 0.35 (92)	10.40 $\pm$ 0.22 (146)	10.07 $\pm$ 0.28 (57)		0.105
rs279858	Exon 5	G	10.10 $\pm$ 0.27 (94)	10.42 $\pm$ 0.22 (147)	11.00 $\pm$ 0.36 (54)		0.182
rs519270	Intron 8	T	10.14 $\pm$ 0.27 (96)	10.43 $\pm$ 0.22 (144)	10.87 $\pm$ 0.36 (55)		0.268
rs693547	Intron 9	T	10.16 $\pm$ 0.27 (97)	10.46 $\pm$ 0.22 (143)	10.76 $\pm$ 0.38 (55)		0.387
rs10805145- rs279826	Intron 3- Intron 4	CCGG	9.80 $\pm$ 0.29 (83)	10.51 $\pm$ 0.22 (147)	11.0 $\pm$ 0.35 (58)	2.6	0.024

### ***GABRA2* variants and LAPSs**

Next, as impulsiveness has been associated with alcohol problems, we also tested the association of these SNPs with the LAPS. Four SNPs, including the most strongly associated SNP mentioned earlier, spanning the region from intron 3 to intron 4 were associated with LAPS (Table 2). Although, none of these SNPs survives multiple testing, we consider to report this association for the mediation analysis that follows. As expected, the association was in the same direction; that is, alleles associated with higher scores of impulsiveness were also associated with higher scores of LAPS (Table 2). The effect size of these SNPs varies between 2.4% and 2.5%, somewhat lower than the effect size for impulsiveness. Again SNP rs279827 showed one of the strongest effects on LAPS (2.5%). Gender, age, and race as covariates were not significant and were not included in the final model. However, as expected, LAPS scores were significantly higher in the AUD group ( $12.5 \pm 2.1$ ) compared

to the non-AUD subjects ( $8.9 \pm 1.9$ ) ( $P < 0.00001$ ). Figure 1 depicts the location of the four associated SNPs with LAPS, highlighting the strength of the association around SNP rs279827. Similar results were obtained in the Caucasian sample when we checked for possible stratification. Homozygotes for the minor haplotype (CCGG/CCGG) showed significantly higher scores of LAPS ( $P = 0.024$ ) (Table 3), but the haplotype was not a better variable than the most significant SNP alone.

**Table 3.** Standardized values of bootstrap mediation analyses of impulsiveness on the effect of *GABRA2* SNPs on Lifetime Alcohol Problems Score (LAPS)

	Total effects LAPS			Direct effects			Indirect effects			Mediated effect %	
	Estimate	SE	P-value	Estimate	SE	P-value	Estimate	SE	Confidence interval 90% P-value		
rs10805145	0.152	0.058	0.010	0.119	0.061	0.047	0.033	0.016	0.011–0.064	0.006	22.4
rs426463	0.156	0.058	0.007	0.121	0.062	0.053	0.035	0.016	0.012–0.066	0.006	22.4
rs279827	0.151	0.058	0.011	0.116	0.061	0.051	0.035	0.016	0.012–0.067	0.006	23.2
rs279826	0.155	0.059	0.011	0.120	0.062	0.048	0.035	0.016	0.012–0.066	0.006	22.5
rs10805145–rs279826	0.160	0.058	0.004	0.126	0.061	0.033	0.034	0.017	0.011–0.065	0.006	21.3

Of the three LAPS components (age of first drunkenness, number of different alcohol-related difficulties at any time, and index of life course duration of all drinking-related problems from onset to present, corrected for risk exposure), the association with *GABRA2* SNPs was driven mainly by number of alcohol related difficulties ( $P < 0.04$ ) and index of life course duration of all drinking related problems ( $P < 0.05$ ), and not by age of first drunkenness.

### Mediation analyses

Given that four *GABRA2* SNPs were associated with both impulsiveness and LAPS, we tested whether impulsiveness may mediate the effect of *GABRA2* on LAPS. We selected the four SNPs that showed association with both impulsiveness and LAPS (rs10805145, rs426463, rs279827, and rs279826) to test for mediation. Table 3 reports the standardized results of the model-based bootstrap that directly tests the significance of mediation effects using 2,000 samples from the original data. Evidence for partial mediation was observed for all four SNPs. Specifically, the indirect effect of *GABRA2* on LAPS was significant for all four SNPs and the direct effect of *GABRA2* on LAPS remained significant. On average, 22.6% of the relation of *GABRA2* SNPs was due to impulsiveness, with SNP rs279827 showing the highest mediated effect, of 23.2%.

### Discussion

The evidence here indicates that a small part of the effect of *GABRA2* on alcoholism is mediated through a specific facet of behavior, namely, impulsiveness. The extent to which this behavioral trait has an underlying genetic component is just starting to be understood. Previously, we reported an association of two *GABRA2* SNPs with impulsiveness, a facet from the neuroticism domain in the NEO-PI-R. Here, we tested these two SNPs and nine additional SNPs with the NEO-impulsiveness, and also with a developmentally constructed,

life course measure of alcohol problems, LAPS. Ten SNPs were associated with impulsiveness and four of these were also associated with LAPS, before correcting for multiple testing. The alleles associated with higher scores for both impulsiveness and LAPS correspond to the same haplotype (minor alleles) previously described in other studies reporting an association of *GABRA2* variation with both alcoholism (Edenberg & Foroud, 2006; Fehr et al. 2006) and brain oscillations (Edenberg et al. 2004). Motivated by our previous finding (Villafuerte et al. 2012) and by a recent report in a sample of college students where higher scores on the urgency (impulsiveness) scale predicted alcohol problems but not alcohol use (Magid & Colder, 2007), we conducted association analyses of *GABRA2* variation with both impulsiveness and alcohol problems. Previously, we reported moderate associations of two of these *GABRA2* SNPs with higher percentage of alcoholic symptoms (Villafuerte et al. 2012). Some of the items on that measure overlap to a small degree with items comprising one component of LAPS, thus forecasting the association we report here. Our results suggest that genetic variation in *GABRA2* in the region comprising intron 3 to intron 4 may influence both level of impulsiveness and also level of alcohol problems. Notably, the association of *GABRA2* is stronger with impulsiveness than with alcohol problems, showing that an intermediate trait such as a personality trait is closer to the genetic influence, hence to the biology, than the manifestation of alcohol problems which are the result of several factors.

To further examine the casual relation of these associations, we performed mediation analysis where impulsiveness mediated the relation between *GABRA2* and LAPS. Indeed, bootstrap analyses indicated that the effect of *GABRA2* on LAPS was partially mediated by impulsiveness. On average, 22.6% of the effect of *GABRA2* on LAPS is accounted for by the impulsiveness-mediating effect. The remaining effect (77.4%) may be mediated by other behavioral risk factors not yet identified, the direct effect of *GABRA2* on LAPS and measurement error. It is also possible that *GABRA2* affects some factor that influences both impulsiveness and alcohol problem.

Furthermore, the associated gene region with both impulsiveness and LAPS comprises intron 3 to intron 4. No coding variation has been reported for exon 4. However, isoforms of human *GABRA2* mRNA where exon 4 (68 bp) has been spliced out were discovered in many brain regions. The product of this isoform would be a truncated protein (nonfunctional) of only 66 amino acids due to the creation of a stop codon (Tian et al. 2005). Notably, one of the associated SNPs rs279827 is located next to the acceptor splice site (Tian et al. 2005). It is not known if SNP rs279827 would have an effect on splicing. But it is intriguing to note that of 11 SNPs in the *GABRA2* gene tested, the SNP near this splice site is the most strongly associated. The haplotype analysis comprising the SNPs in the proximal block (Figure 1) shows similar results as the individual SNPs supporting the notion that in this study the haplotype does not unveil a better not genotyped SNP in the region or independent contribution of the genotyped SNPs.

*GABRA2* has been extensively investigated for its role in alcoholism, alcohol sensitivity, anxiolytic effects of benzodiazepines and its effect on EEG  $\beta$  patterns. The evidence suggests that this gene may have a pleiotropic effect on brain function involving overlapping mechanisms. The presence of alcohol and benzodiazepine binding sites in this subunit reveals the role of *GABRA2* on alcohol sensitivity and anxiety (Low et al. 2000; Uhart et al.

2013), while the effect of *GABRA2* genetic variation on the EEG  $\beta$  wave suggests a different mechanism, involving the excitation-inhibition homeostatic balance and impulsivity. Converging lines of evidence point to the role of *GABRA2* on impulsive-related behavior. Both in ADHD children and alcoholics with impulsive behaviors show increase in the EEG  $\beta$  activity (Bauer & Hesselbrock, 1993; Clarke et al. 2001). The frontal region is associated with self-regulation and inhibition/control behavior. It appears that genetic variation in *GABRA2* that influences the  $\beta$  activity in EEG may also influence inhibitory control behaviors such as impulsiveness. The moody and temper tantrum behavior may be related to the kinds of impulsiveness we describe here, involving lack of control under distress.

These results contribute to a refined understanding of the genetic role of *GABRA2* on impulsiveness, a major precursive behavioral risk characteristic (Masten et al. 2008; Zucker et al. 2008) and on LAPS that measures a facet of subsequent alcohol problems that is specific to problems due to consumption.

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