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Heterosis for Grain Yield and Other Agronomic Traits in Foxtail Millet

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Heterosis for Grain Yield and Other Agronomic Traits in Foxtail Millet


ABSTRACT

Foxtail millet [Setaria italica (L.) P. Beauv.] is a largely self-pollinating species that is used as a warm-season annual in the USA. Nearly all cultivars of this species grown in the USA are selections from land races. This research was undertaken to determine whether sufficient high-parent heterosis is expressed in foxtail millet for grain yield and other key traits to justify the development and use of varietal crosses. Seven diverse parents and 21 F2s and 21 F3s produced from biparental crosses were evaluated in five environments in 1996. Genotype × environment interaction was highly significant for grain yield, but the highest yielding entries were high-yielding in each environment. High-parent heterosis for grain yield was detected in 18 of 21 F2s. On the basis of the estimate of average heterosis, which was highly significant in every environment, the expected yield of the F1 generation was 68% greater than the average yield of the parental cultivars. This high level of heterosis for grain yield suggested that varietal crosses or other types of cultivars in which there exists a relatively high amount of heterozygosity would provide a significant yield benefit over nonhybrid cultivars. Although significant heterotic effects were observed for each of the other traits, additive effects were more important. Significant correlations between traits of the estimates of additive and/or variety heterosis effects suggested that at least some of the genes controlling grain yield, plant height, and spike length were either the same or in coupling phase linkage.

In the great plains of the USA, foxtail millet is used primarily as a warm-season annual forage. The USDA has not released any estimates of land area planted to foxtail millet. However, it is often included in wheat–continuous crop rotations, which in some environments have shown to be superior to the more traditional wheat–fallow rotations (Senft 1998). The grain of foxtail millet also is harvested for pet birdseed, and in China, India, and other parts of East Asia this species has been an important food crop for centuries.

Foxtail millet is largely a self-pollinating species. Outcrossing rates have been estimated from 0.0 to only 1.4% for plants separated by 0.30 m (Till-Bottraud et al., 1992), although Li et al. (1935) reported rates as high as 5.6% for some varieties under certain conditions. Nearly all foxtail millet cultivars grown in the USA are selections from land races. A primary reason that varietal crosses or selfed selections from planned crosses have not been widely used is that foxtail millet is one of the most difficult species to cross-pollinate (Baltensperger, 1996). The flowers are small (about 1 mm in length), and anthesis generally occurs near midnight and in the morning but varies greatly with the environment (Malm and Rachie, 1971). However, Siles et al. (2001) described an artificial technique of hybridization that resulted in 67.5% hybrid seed set per flower crossed. Also, Wang (1991) reported the discovery of male-sterile varieties of foxtail millet. The ability to hybridize foxtail millet opens other options to breeders besides selecting from land races. If nonadditive gene action is important, then either mid- and/or high-parent heterosis may be sufficient to justify the production and use of varietal crosses.

Information on the inheritance of important agronomic traits of foxtail millet, including susceptibility to key diseases, is limited. Most of the previous work has focused on estimating broad-sense heritabilities and realized genetic gains, with little attention directed to measuring levels of heterosis or to assessing the relative importance of different types of gene action (Athwal and Singh, 1966; Singh and Athwal, 1966; Gill and Randhawa, 1975; Vishwanatha et al., 1981; Gurunadha Rao et al., 1984; Prasada Rao et al., 1985). Darmency et al. (1987) reported that most of 19 morphological and reproductive traits were probably under the control of nonadditive genetic components, but this research was conducted on an interspecific cross between foxtail millet and its wild relative S. viridis (L.) P. Veauv. Also, information on the importance of genotype × environment interaction for this species when grown in environments of the Great Plains is lacking.

Important agronomic traits in foxtail millet include not only grain yield, days to heading, days to maturity, and plant height, but also number of tillers and spike length. When foxtail millet is used as a forage, tillering is a desirable trait. However, non-tillering cultivars are preferred for use in producing birdseed because the seeds typically are larger. Spike length also is correlated with seed size. One of the more important diseases affecting foxtail millet is leaf spot, which is caused by at least three species of Helminthosporium spp., H. setariae Sawada, H. turcicum Pass., and H. carbonum Ullstrup (Haenseler, 1941; Robert, 1962).

The objective of this research was to determine the presence and importance of heterosis for grain yield and other important agronomic traits of foxtail millet. Also, the relative importance of genotype × environmental effects and the interaction of heterosis with these effects were measured. This information is needed by breeders to determine the best types of cultivars to develop and to design testing programs with appropriate numbers and types of environments.
MATERIALS AND METHODS

Two cultivars (Red Siberian and Golden German) and five plant introductions (PI614814, PI614815, PI614816, PI614817, and PI614818) were selected as parents for this study. The two cultivars were randomly chosen from many early cultivars that were introduced into and cultivated in the USA. The five plant introductions were selections from PI458628, PI531445, PI473598, NESE2, and PI464233, respectively, that matured in western Nebraska and on the basis of evaluations in the Nebraska Panhandle from 1991 to 1994 showed above average grain yield and resistance to Wheat streak mosaic virus (Siles et al., 2001).

In the summer of 1994, the parents were crossed in a half-diallel arrangement to produce 21 F1 progenies. Seed of the F1 and F2 generations of each cross was produced in a greenhouse at the University of Nebraska-Lincoln in the winter of 1994 and summer of 1995, respectively. The F1 generation of each cross was obtained by mixing equal quantities of seed from each of 200 F1 plants. In the same summer, the F1 and F2 generations from each of the 21 crosses were evaluated in five environments in western Nebraska, in a split-plot design with two replications per environment. The main plots were the generations, and the subplots were the crosses. The seven parents were included in all replications of each main plot. Seed of a parent was produced by self-pollinating a single plant in the same environment as seed of the generation (F1 or F2) of the main plot in which the parent was grown.

Four environments (E1 through E4) were located at the High Plains Agricultural Laboratory at Sidney, NE, and one environment (E5) was at the Panhandle Research and Extension Center at Scottsbluff, NE. E1 through E3 were dryland sites, whereas E4 and E5 were irrigated. The soil types were keith loam (fine-silty, mixed, mesic Argid Argustolls) at E1 and E4, duroc loam (fine-silty, mixed, mesic Pachic Haplustolls) at E2 and E3, and tripp fine sandy loam (coarse-silty, mixed, mesic Argid Arupolls) at E5. Sunflower (Helianthus annuus L.), fallow, wheat (Triticum aestivum L.), and amaranth (Amaranthus spp.) were the previous crops at E1, E2, E3, E4, and E5, respectively. Sowing dates were 3 June, 22 May, 5 June, 5 June, and 4 June at E1, E2, E3, E4, and E5, respectively. The plot size was four rows by 2.1 m in length at E1, E2, and E3 or four rows by 1.5 m in length at E4 and E5. Adjacent rows within and between adjacent plots were spaced 0.3 m apart. All entries were planted at an average rate of 5.7 kg of seed ha\(^{-1}\).

Days to heading and maturity, grain yield, plant height, spike length, and number of tillers per plant were recorded at each environment. Days to heading and maturity were recorded on a plot basis and were counted from the date of planting until 50% of the spikes emerged from the flag leaf and until the spikes turned pale yellow, respectively. Plant height, spike length, and number of tillers per plant were recorded as the mean of 10 randomly selected plants from the central two rows of each plot. Plant height and spike length were evaluated on the main culm, whereas the number of tillers per plant was recorded as the number of seed-bearing tillers per plant. The two central rows from each plot were harvested to evaluate grain yield. At E1, E2, and E3 only the middle 1.5 m section of each row was harvested; at E4 and E5 the middle 2.1 m of each row was harvested. Evaluations of resistance to Helminthosporium leaf spot were performed only at E3 and E4, where natural levels of infection were sufficiently high to discern differences among entries. The disease reaction was rated subjectively on a plot basis by a 10-class scale (0 = no lesions at all or traces; 9 = lesions on 90% or more of leaf surface on all plants). Disease reaction was scored at the flowering stage.

For each trait, an analysis of variance over all entries and across all environments was used to assess the relative importance of the environmental main effect and the interaction between environments and entries. Entries were treated as a fixed effect, whereas environments were treated as a random variable. The reference environmental space included both irrigated and dryland production sites in the high plains region of the Nebraska Panhandle and adjacent areas of Colorado and Wyoming with similar soil types, climate, and production practices. Additive (\(a\)) and heterotic (\(h\)) effects were estimated for each variety (\(i\)) and variety cross (\(ij\)) and the heterotic effects were partitioned into average (\(h\)), variety (\(hi\)), and environment (\(E\)) specific (\(sij\)) heterosis as described by Gardner and Eberhart (1966). The F1 crosses were not grown and evaluated due to a shortage of seed, but an estimate of the performance of the F1 generation across all crosses was obtained as \(\mu + F\), where \(\mu\) is the mean of the parental varieties. An estimate of the \(F1\) generation between the \(i\)th and \(j\)th parents was obtained as \(\mu + a_i + a_j + h_{ij} + h_i + h_j + s_{ij}\). Significance of the microenvironmental difference among environments between F1; and F1 whole plots was determined by an F test in an analysis of variance of parental data only.

RESULTS

The environmental difference between whole plots within environments was not statistically significant (\(p > 0.05\)); consequently, no corrections were made for this effect. Across environments, highly significant (\(p < 0.01\)) variation occurred among environments and entries for each trait (Table 1). For grain yield, the highest yielding

Table 1. Mean squares from an analysis of variance of data on seven agronomic traits measured on seven cultivars of foxtail millet and their half-diallel derived 21 F1 and 21 F2 progenies. Reaction to leaf spot was evaluated in two environments in 1996, and all other traits were evaluated in five environments in 1996.

<table>
<thead>
<tr>
<th>Source of Variation</th>
<th>df</th>
<th>Grain yield (Mg ha(^{-1}))^2</th>
<th>Days to heading</th>
<th>Days to maturity</th>
<th>Plant height (cm)</th>
<th>Spike length</th>
<th>Tillers per plant</th>
<th>Reaction to leaf spot disease (score)†</th>
</tr>
</thead>
<tbody>
<tr>
<td>Environments (Env)</td>
<td>4</td>
<td>139.7**</td>
<td>1 308.1**</td>
<td>1 818.3**</td>
<td>43 032**</td>
<td>136.5**</td>
<td>18.7**</td>
<td>10.2**</td>
</tr>
<tr>
<td>Entries (Ent)</td>
<td>48</td>
<td>2.6**</td>
<td>180.3**</td>
<td>112.4**</td>
<td>777**</td>
<td>90.7**</td>
<td>10.0**</td>
<td>20.4**</td>
</tr>
<tr>
<td>Additive effect</td>
<td>6</td>
<td>4.7**</td>
<td>1 010.5**</td>
<td>667.8**</td>
<td>4 589**</td>
<td>434.6**</td>
<td>76.7**</td>
<td>145.7**</td>
</tr>
<tr>
<td>Average heterosis</td>
<td>1</td>
<td>59.4**</td>
<td>645.9**</td>
<td>150.3**</td>
<td>2 101**</td>
<td>887.7**</td>
<td>6.9**</td>
<td>&lt;0.1</td>
</tr>
<tr>
<td>Specific heterosis</td>
<td>14</td>
<td>0.3</td>
<td>63.6**</td>
<td>42.1**</td>
<td>87</td>
<td>20.0**</td>
<td>&lt;0.1</td>
<td>4.3**</td>
</tr>
<tr>
<td>Residual</td>
<td>21</td>
<td>0.3</td>
<td>17.5**</td>
<td>10.9</td>
<td>166**</td>
<td>4.0**</td>
<td>0.7**</td>
<td>21.4</td>
</tr>
<tr>
<td>Ent × Env</td>
<td>192</td>
<td>0.4**</td>
<td>9.3**</td>
<td>9.0**</td>
<td>66**</td>
<td>2.4**</td>
<td>0.2**</td>
<td>48.9</td>
</tr>
<tr>
<td>Error</td>
<td>240</td>
<td>0.2</td>
<td>2.3</td>
<td>37</td>
<td>1.3</td>
<td>&lt;0.1</td>
<td>96.7</td>
<td>0.7</td>
</tr>
</tbody>
</table>

* Significant at the 0.05 level of probability.
** Significant at the 0.01 level of probability.
† A 0-to-9 visual rating scale was used, with a score of 0 indicating no disease lesions and a score of 9 indicating lesions covering at least 90% of the leaf surface.
Table 2. Means of seven foxtail millet cultivars for seven agronomic traits. Values for disease scores are averages across two environments in 1996 and for all other traits are averages across five environments in 1996.

<table>
<thead>
<tr>
<th>Parental cultivar</th>
<th>Grain yield</th>
<th>Days to heading</th>
<th>Days to maturity</th>
<th>Plant height</th>
<th>Spike length</th>
<th>Tillers per plant</th>
<th>Leaf spot disease</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mg ha⁻¹</td>
<td>cm</td>
<td></td>
<td>number</td>
<td></td>
<td></td>
<td>score †</td>
</tr>
<tr>
<td>Golden German</td>
<td>2.54</td>
<td>64.2</td>
<td>94.5</td>
<td>98.4</td>
<td>14.4</td>
<td>2.5</td>
<td>1.4</td>
</tr>
<tr>
<td>PI614814</td>
<td>2.75</td>
<td>62.2</td>
<td>90.2</td>
<td>99.6</td>
<td>9.4</td>
<td>0.0</td>
<td>2.0</td>
</tr>
<tr>
<td>PI614815</td>
<td>3.79</td>
<td>51.5</td>
<td>81.6</td>
<td>96.4</td>
<td>9.9</td>
<td>2.7</td>
<td>1.4</td>
</tr>
<tr>
<td>PI614816</td>
<td>3.05</td>
<td>50.3</td>
<td>82.6</td>
<td>79.3</td>
<td>7.8</td>
<td>3.5</td>
<td>3.0</td>
</tr>
<tr>
<td>PI614817</td>
<td>2.35</td>
<td>53.6</td>
<td>81.7</td>
<td>95.0</td>
<td>13.3</td>
<td>0.0</td>
<td>9.0</td>
</tr>
<tr>
<td>Red Siberian</td>
<td>3.76</td>
<td>48.9</td>
<td>77.6</td>
<td>88.5</td>
<td>10.6</td>
<td>2.5</td>
<td>2.4</td>
</tr>
<tr>
<td>PI614818</td>
<td>2.58</td>
<td>49.2</td>
<td>81.7</td>
<td>95.0</td>
<td>13.3</td>
<td>0.0</td>
<td>9.0</td>
</tr>
<tr>
<td>LSD 0.05</td>
<td>0.57</td>
<td>2.8</td>
<td>7.1</td>
<td>1.4</td>
<td>0.4</td>
<td>1.4</td>
<td></td>
</tr>
</tbody>
</table>

† A 0- to 9-visual rating scale was used, with a score of 0 indicating no disease lesions and a score of 9 indicating lesions covering at least 90% of the leaf surface.

The parental cultivar was PI614815 at 3.79 Mg ha⁻¹ and the lowest yielding was PI614817 at 2.35 Mg ha⁻¹ (Table 2). The ranges for days to heading and maturity among the parental cultivars were 13.3 and 16.9 d, respectively; Red Siberian was the earliest cultivar for both maturity traits and Golden German the latest. Thus, most of the varietal differences in maturity were predictable from differences in days to heading. Plant height ranged from 79 to 100 cm and spike length from 7.8 to 14.4 cm. Two of the parental cultivars did not tiller, whereas the others ranged from 2.5 to 3.5 tillers per plant. One cultivar, PI614817, exhibited an extremely sensitive response to Helminthosporium leaf spot, scoring a 9 in each replication of both environments. All other parental cultivars had low to moderately low disease scores.

Heterosis, Grain Yield

In the across environmental analysis, the additive effect, average heterosis, and variety heterosis were highly significant, whereas specific heterosis was not significant. The deviation from a model with only the additive effect, average heterosis, and variety heterosis was not significant.

The ranking of the cultivars by the values of \( a_i \) (Table 3) was similar but not identical to the ranking based on per se yields. PI614815, the cultivar with the highest per se yield, also had the most positive value of \( a_i \). The most noticeable discrepancy between per se yields and \( a_i \) values was observed for the Red Siberian, Golden German, and PI614814 cultivars. Red Siberian had a highly significantly greater per se grain yield than either of the other two cultivars (3.76 Mg ha⁻¹ compared to 2.54 and 2.75 Mg ha⁻¹), but all three varieties had identical and significantly positive values of \( a_i \).

The significance of \( h_i \) was a result of the higher average grain yield of the F₂ and F₃ generations compared to the average yield of the parental cultivars (Table 4). The yield superiority of the F₂ over the parental generation was 32%, and in 18 of 21 crosses the F₂ exhibited high-parent heterosis for yield. In half of these 18 crosses, over-dominance was observed even at the F₁ generation. The expected yield of the F₂ generation was 4.98 Mg ha⁻¹. This was 68% greater than the average grain yield of the parental cultivars and 31% greater than the yield of the highest yielding parent.

Varieties Golden German, PI614814, and PI614817 had \( h_i \) values that were highly significantly greater than 0 (Table 3). The cross between Golden German and PI614814 had F₂ high-parent heterosis of 78%, and this F₂ was the highest yielding entry in the test at 4.90 Mg ha⁻¹. The estimated yield of the F₂ of this cross was 7.29 Mg ha⁻¹, which was more than twice the yield of the higher parent. The two varieties with the highest per se grain yields, PI614815 and Red Siberian, both had \( h_i \) values that were highly significantly less than 0. The F₂ derived from the crosses of these two varieties actually had less grain yield than either parent. The relationship between the level of heterosis observed for grain yield and the coancestry of the parents could not be determined because the phylogenetic relationships among the parents used in this research is not known.

Heterosis, Other Traits

For all other traits, the relative importance of additive effects was much greater than observed for grain yield.
Table 4. Generational means and estimate of average heterosis ($h$) for seven agronomic traits of foxtail millet. Reaction to leaf spot disease was evaluated in two environments in 1996, and all other traits were evaluated in five environments in 1996.

<table>
<thead>
<tr>
<th>Generational mean</th>
<th>Grain yield</th>
<th>Days to heading</th>
<th>Days to maturity</th>
<th>Plant height</th>
<th>Spike length</th>
<th>Tiller per plant</th>
<th>Reaction to leaf spot disease</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mg ha$^{-1}$</td>
<td></td>
<td>d</td>
<td></td>
<td>cm</td>
<td>number</td>
<td>score†</td>
</tr>
<tr>
<td>Parental</td>
<td>2.97</td>
<td>54.3</td>
<td>84.2</td>
<td>91.9</td>
<td>10.7</td>
<td>2.1</td>
<td>2.7</td>
</tr>
<tr>
<td>F$_1$</td>
<td>3.94</td>
<td>50.6</td>
<td>82.9</td>
<td>100.3</td>
<td>14.6</td>
<td>2.3</td>
<td>2.9</td>
</tr>
<tr>
<td>F$_2$</td>
<td>3.39</td>
<td>51.7</td>
<td>84.2</td>
<td>101.1</td>
<td>12.7</td>
<td>1.9</td>
<td>3.3</td>
</tr>
<tr>
<td>LSD 0.05(Parental vs. F2 or F3)</td>
<td>0.18</td>
<td>0.8</td>
<td>0.8</td>
<td>2.2</td>
<td>0.6</td>
<td>0.3</td>
<td>0.4</td>
</tr>
<tr>
<td>$h$</td>
<td>2.01**</td>
<td>-6.9**</td>
<td>-3.4**</td>
<td>12.3**</td>
<td>7.5**</td>
<td>0.7**</td>
<td>-0.1</td>
</tr>
</tbody>
</table>

** Significant at the 0.01 level of probability.
† A 0-to-9 visual rating scale was used, with a score of 0 indicating no disease lesions and a score of 9 indicating lesions covering at least 90% of the leaf surface.

Only 23% of the variation among entries was attributable to the additive effect for yield, whereas the minimal value for this same percentage for the other traits was 60% for spike length and the maximal value was 95% for leaf spot ratings (Table 1).

Although additivity was the most important effect for each of these other traits, average heterosis was statistically highly significant for each trait except leaf spot rating (nonsignificant), variety heterosis was highly significant for each trait except tiller number and leaf spot rating (nonsignificant), and specific heterosis was highly significant for heading and maturity date, spike length, and leaf spot rating and nonsignificant for plant height and tiller number (Table 1). On the basis of estimates of $h$ (Table 4), the expected $F_1$ heterosis was 70% for increased spike length, 33% for more tillers, 12 and 4% for fewer days to heading and maturity, and 13% for increased plant height. The residual from the full genetic model ($a_i$, $h$, $h_i$, and $s_i$ effects) was statistically significant for each of the other traits except days to maturity and reaction to Helminthosporium leaf spot (Table 1). The significance of this residual indicated that linkage and/or epistasis were involved in the inheritance of these traits in this material.

The correlation among the $a_i$ values between pairs of traits was significant for heading and maturity dates (0.93), plant height and spike length (0.77), and spike length and leaf spot rating (−0.86). Among $h_i$ values, significance was observed between heading and maturity dates (0.78), grain yield and plant height (0.76), grain yield and spike length (0.87), and plant height and spike length (0.93). The highest yielding entry, the $F_2$ between PI614814 × Golden German, exhibited high- or early-parent heterosis not only for yield, but also for each of the other traits except leaf spot rating.

**Entry × Environmental Interaction**

Among the six traits measured at all environments, the percentage of the total sums of squares for environments, entries, and their interaction that was attributable to the environmental main effect ranged from 77 for plant height to 10 for spike length (Table 1). For ratings of resistance to Helminthosporium leaf spot (measured only at E3 and E4), only 3% of this total of sums of squares was due to the environmental main effect. Over 98% of the variation among environments for grain yield was attributable to the difference between the irrigated and dryland environments. The average grain yield of the irrigated and dryland environments was 4.86 and 2.70 Mg ha$^{-1}$, respectively. For the other five traits measured at all environments, the comparison between irrigated and dryland environments was highly significant but was considerably less important than observed for grain yield.

The entry × environmental interaction was highly significant for all traits except reaction to leaf spot disease (Table 1). However, for every trait the interaction was less significant than either main effect. For those traits with a highly significant interaction, the sum of squares attributable to this interaction expressed as a percentage of the sum of squares among entries was greatest for grain yield at 62. The values of this percentage for days to heading, days to flowering, plant height, spike length, and tillers per plant were 21, 32, 41, 10, and 9, respectively.

The most variable of the parental cultivars for grain yield was PI614814, which ranked best among the parents in E4, an irrigated environment, but worst in E3, a dryland environment. The cause of this differential response may not have been water, however, because in the other irrigated environment, E5, this cultivar ranked only fifth best among the parents for grain yield. The best parental cultivar for grain yield across environments, PI614815, ranked either best or second best among the parents for grain yield in all environments (Table 3). The highest average grain yield, the $F_2$ from the Golden German × PI614814 cross, also ranked no worse than third best among all the entries in any environment.

A similar consistency across environments was observed for the grain yield values of the additive and heterotic effects. For example, PI614815, the parental cultivar with the most positive $a_i$ value across environments (Table 3), had $a_i$ values that were significantly greater than zero in three environments and positive in the other two environments. In only one instance did a parental cultivar have an $a_i$ value that was significantly positive in one environment and significantly negative in another environment (PI614814 in E4[+] and E5[−]). Average heterosis was highly significant and positive in every environment. In no instance was a variety heterosis effect for grain yield significantly greater than zero in one environment and significantly less than zero in another environment.
DISCUSSION

A primary objective of this research was to determine whether heterosis for grain yield and other important agronomic traits in foxtail millet is of sufficient magnitude to be of practical commercial value. This issue has become more relevant recently because of improved procedures for producing F2 seed of this species. The greatest amount of heterosis was observed for grain yield (67%) and spike length (68%). In 18 of 21 crosses, high-parent heterosis for grain yield was observed at the F2 generation. The correlation between pairs of traits among the estimated q, and h values suggested that at least some of the genes controlling grain yield in foxtail millet are the same as or linked in coupling phase with the genes controlling plant height and spike length.

Our predicted advantage of F1 grain yield over parental yield (2.01 Mg ha–1) may be biased upward if the size of seed from which a plant is produced has an effect on the grain yield of that plant. Seed size was not measured in this research. However, if seed size is dependent on the vigor of the parent plant, then on the basis of the observed heterosis for maturity and plant height, the seed from which the F2 generation was grown would have been the largest seed followed by the F1 generation and then the parents. Seed from which the F1 generation would be grown would have the same size as for the parents. The F2 generation had significantly lower seedling emergence than either the F1 generation and then the parents. Seed from which the F1 generation would be grown would have the same size as for the parents. Although the entry H11002 may be biased upward if the Heterosis in sorghum and pearl millet. p. 375–386. In J.G. Coors and S. Pandey (ed.) The genetics and exploitation of heterosis in crops. ASA, CSSA, and SSSA, Madison, WI.


