Effect of moisture stress on combining ability and gene action for polygenic traits in maize

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Abstract
Present study was carried out to assess the general combining ability effects of parents and specific combining ability effects of hybrids for yield and yield related traits purely under rainfed situations. Incidentally, it mimicked mid season drought conditions i.e. moisture stress at flowering stage for a period of 22 days. Combining ability analysis using Line × Tester design was conducted in maize (\textit{Zea mays} L) inbred lines by growing 135 F\textsubscript{1}'s generated by crossing fifteen lines with nine testers at ARS, Karimnagar, PJTSAU during rainy season, 2011. Female lines were derived from the recurrent selection cycle carried out in the drought tolerant population «Tuxpeno Sequia», procured from CIMMYT in 1996. Males were developed using pedigree breeding. Analysis of variance showed highly significant genotypic differences for all the traits studied indicating wide range of variability among the genotypes. The ratio of gca /sca was less than unity indicating the preponderance of non-additive gene action in the expression of all the characters studied except flowering traits. Moderate narrow sense heritability was observed for majority of the characters. Three lines KMLD-3, KMLD-11, and KMLD-19 and one tester, KML-9 were good general combiners for grain yield and one or more yield contributing characters. The crosses KMLD-3 × KML-99, KMLD-5 × BML-7, KMLD-5 × KML-801, KMLD-6 × KML-36, and KMLD-11 × KML-29 showed significant positive sca, standard heterosis and high mean values for grain yield plant\textsuperscript{-1} and two or more yield components. These hybrids were found to be early in flowering with significant negative heterosis providing some clue about their usefulness under drought conditions. Therefore further testing of these new inbred lines for use in a crossing program is recommended to combine major yield components with high yield to derive climate resilient hybrids.

Keywords: maize, moisture stress, combining ability, line × tester, standard heterosis, narrow sense heritability

Introduction
Drought is estimated to cause average annual yield losses in maize of about 17\% in the tropics (Edmeades et al, 1989), and this figure is expected to increase in the future as a large population of the crop is being shifted to marginal lands. Indian agriculture is dominated by rain fed farming and so even after development of large irrigation infra structure, still 63\% area is rain fed. In India, Maize is grown on 6 mha area mainly as rain fed crop during rainy (kharif) season. Telangana state is one of the major maize producing states in the country and maize is cultivated mainly as a rain fed crop in kharif season.

Maize is particularly sensitive to water stress in the period one week before to two weeks after flowering (Grant et al, 1989). Drought during this period results in increased anthesis-silking interval (ASI) (Edmeades et al, 2000) and consequently in grain abortion (Boyle et al, 1991). Hence, development of maize cultivars with high and stable yields under drought is most important particularly in Northern Telangana Zone where mid season and terminal droughts during kharif are common. Keeping this constraint in view an attempt has been made in the present investigation to study the gca and sca effects of parents and hybrids respectively in crosses where drought tolerant lines were used.

Materials and Methods
The experimental material comprised of fifteen lines, viz. KMLD-3, KMLD-5, KMLD-6, KMLD-11, KMLD-18, KMLD-19, KMLD-21, KMLD-61, KMLD-65, KMLD-66, KMLD-68, KMLD-70, KMLD-71, KMLD-73, and KMLD-802. These lines were derived from the recurrent selection cycle carried out in the population «Tuxpeno Sequia», procured from CIMMYT in 1996. Nine testers viz. BML-7, KML-9, KML-29, KML-36, KML-55, KML-99, KML-224, KML-801, and KML-802 were developed using pedigree breeding. These parents were crossed in line × tester fashion during rabi season of 2011-12 at Agricultural Research Station, Karimnagar to generate 135 F\textsubscript{1}s. During kharif season of 2012, all the single crosses were evaluated in a randomized block design with two replications with row-to-row and plant-to-plant spacing of 75 cm × 20 cm, respectively purely under rain fed situation against DHM-117, the popular check. Crop was sown on July 25\textsuperscript{th} and at flowering stage crop was under
severe moisture stress in the month of September. Incidentally, it mimicked mid-season drought condition for a period of 22 days. Rainfall at vegetative stage i.e. in the month of August was 10.5% deficit (21.78 mm) over the normal rainfall (246.52 mm) and at flowering stage i.e. in the month of September was 103.52% deficit (97 mm) over the normal rainfall (172.66 mm). The data were recorded on five selected plants for 11 yield and yield contributing characters. Combining ability analysis was done according to the model given by Kempthorne (1957).

Results and Discussion

In the present investigation, the analysis of variance for all the yield and yield component traits showed that, variance due to hybrids was highly significant for all the traits studied indicating the manifestation of parental genetic variability in their crosses (Table 1). The mean squares for hybrids were partitioned into three components viz., due to lines, due to testers and due to line × tester interactions. The differences among hybrids due to the lines, testers and line × tester interaction were significant for all the characters except number of ears plant\(^{-1}\) in lines and testers, ear length, number of kernels row\(^{-1}\) and grain yield plant\(^{-1}\) in lines and ear height and ear girth in testers, suggesting that the experimental material possessed considerable variability and that both gca and sca were involved in genetic expression of these traits. A higher proportion of sca variance than gca variance was noticed for all the traits except days to 50% pollen shed and days to 50% silk emergence indicating significantly higher non-additive interactions among the hybrids, which could be exploited. Higher sca variance than the gca variance exhibiting preponderance of non-additive gene effects has also been earlier reported by Aminu and Izge (2013). However, the results contradict the view reported by Sharma et al (2004), who found the preponderance of additive genetic effects in the control of most traits in maize.

Estimates of gca effects (Table 2) indicated that the lines KMLD-3 KMLD-11, and KMLD-19 were good general combiners for grain yield plant\(^{-1}\). Lines KMLD-19 and KMLD-21 were good general combiners for 100 grain weight and number of kernels row\(^{-1}\), respectively. Lines KMLD-5 and KMLD-6 were good combiners for kernel rows and ear girth. For ear length, KMLD-71 and KMLD-73 were good general combiners and for days to 50% silking, KMLD-19, KMLD-65, KMLD-68 and KMLD-71 were good general combiners. KMLD-3, KMLD-5, KMLD-19, and KMLD-61 were good general combiners for plant height. Among testers, KML-9 was a good general combiner for grain yield plant\(^{-1}\), ear girth and number of kernels row\(^{-1}\). BML-7 was a good general combiner for plant height, ear height, ear girth and 100 grain weight and KML-224 was a good general combiner for plant height, ear length and 100 grain weight. KML-55, KML-801, and KML-802 were good general combiners for days to 50% pollen shed and days to 50% silking.

Among the hybrids, KMLD-5 × KML-801 was the best specific combiner for grain yield followed by KMLD-6 × KML-36 and KMLD-11 × KML-29 (Table 3). The lines KMLD-5, KMLD-18 and KMLD-68 with tester BML-7 were good specific combiners for grain yield plant\(^{-1}\). KMLD-70 × KML-801 was the only combination with significant positive sca for grain yield plant\(^{-1}\) coupled with significant negative sca for days to 50% silking desirable for early maturity. Line KMLD-5 with testers BML-7 and KML-9 showed significant positive sca for plant height, ear height, ear length, and ear girth, lines KMLD-19 and KMLD-65 with the same testers showed significant positive sca for plant height and ear girth. In most of the hybrids that had the highest sca effects for grain yield plant\(^{-1}\), one of the parents was KMLD-3 or BML-7 or KML-9. These are the best general combining parents either for grain yield plant\(^{-1}\) and yield attributing characters and therefore, the combination of favorable genes from the parents for the corresponding traits could have resulted in high and desirable sca effects. In the present study, some of the superior hybrids were from either one of the parents with high gca effect or parents that are low × low general combiners, indicating that the parents with either high gca or low gca would have a higher chance of having excellent complimentarity with other parents. This is in similarity with that of earlier findings of Premlatha and Kalamani (2010).

<table>
<thead>
<tr>
<th>Source</th>
<th>df</th>
<th>Grain yield plant(^{-1})</th>
<th>Days to 50% pollen shed</th>
<th>Days to 50% silking</th>
<th>Plant height</th>
<th>Ear height</th>
<th>Number of ears plant(^{-1})</th>
<th>Ear length</th>
<th>Ear girth</th>
<th>Kernel rows</th>
<th>Number of kernels row(^{-1})</th>
<th>100 grain weight</th>
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<td>0.37</td>
<td>967.50**</td>
<td>0.53</td>
<td>0.01</td>
<td>2.60</td>
<td>8.22**</td>
<td>0.03</td>
<td>529.20**</td>
<td>44.69**</td>
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<td>7.35**</td>
<td>7.31**</td>
<td>378.74**</td>
<td>187.65**</td>
<td>0.03**</td>
<td>2.67**</td>
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<td>16.18**</td>
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<td>3.82**</td>
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</tr>
<tr>
<td>Testers 8</td>
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<td>42.77**</td>
<td>36.58**</td>
<td>902.46**</td>
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<td>0.03</td>
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<td>65.60**</td>
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<td>217.16**</td>
<td>92.43**</td>
<td>0.03</td>
<td>2.26***</td>
<td>1.24**</td>
<td>2.38**</td>
<td>12.53**</td>
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<td>Error 134</td>
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<td>0.37</td>
<td>607.50**</td>
<td>0.37</td>
<td>0.37</td>
<td>0.37</td>
<td>0.37</td>
<td>0.37</td>
<td>0.37</td>
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<td>0.37</td>
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*Significant at p<0.05; **Significant at p<0.01
Hybrids with good specific combining ability and per se performance could be selected to recover transgressive segregants. The superiority of high × low or average × low combiners as parents could be explained on the basis of interaction between positive alleles from good/average combiners and negative alleles for the poor combiners as parents. The high yield of such crosses or hybrids would be non-fixable and thus could be exploited for heterosis breeding. The superior cross combinations involving low × low or average × low combiners as parents could be used as sources of new desirable inbred lines. For the exploitation of hybrid vigour, per se performance, sca effects and the extent of heterosis among hybrids could be important. Selection based on one of the aforementioned criteria alone may not be effective. The hybrid with high per se performance need not always reveal high sca effect and vice versa. So selection must be based on significant sca and high heterotic effects coupled with high mean performance.

The present study identified hybrids KMLD-3 × KML-99, KMLD-5 × BML-7, KMLD-5 × KML-801, KMLD-6 × KML-36, and KMLD-11 × KML-29 with significant positive sca, standard heterosis and high mean values for grain yield plant−1 and two or more yield components. These hybrids were found to be early in flowering with significant negative heterosis suitable to drought prone situations. Further testing of the new inbred lines viz. KMLD-3, KMLD-5, KMLD-6, KMLD-11, KMLD-29, KMLD-36, and KMLD-99 for use in a crossing program is recommended to combine major yield components with high yield under moisture stress conditions.

**Implications**

Dry matter yield and composition of maize whole-plant for silage can be controlled by multiple management factors. Despite this, uncontrollable environmental factors, such as drought stress and heat

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**Table 2 - Estimates of general combining ability effects of inbred lines of maize.**

<table>
<thead>
<tr>
<th>Lines/Testers</th>
<th>Grain yield plant−1</th>
<th>Days to 50% pollen shed</th>
<th>Days to 50% silking</th>
<th>Plant height</th>
<th>Ear length</th>
<th>Number of ears plant−1</th>
<th>Ear girth</th>
<th>Kernel rows</th>
<th>Number of kernels row−1</th>
<th>100 grain weight</th>
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<td>9.06**</td>
<td>9.45**</td>
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<td>0.13</td>
<td>0.03**</td>
<td>0.47</td>
<td>-0.76</td>
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<td>KMLD-5</td>
<td>0.63</td>
<td>0.58</td>
<td>0.64</td>
<td>16.67**</td>
<td>18.05**</td>
<td>0.02</td>
<td>0.12</td>
<td>0.64**</td>
<td>1.14**</td>
<td>-0.81</td>
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<td>KMLD-6</td>
<td>3.13</td>
<td>1.15**</td>
<td>1.37**</td>
<td>3.84</td>
<td>3.12**</td>
<td>0.03</td>
<td>0.24</td>
<td>0.61**</td>
<td>1.20**</td>
<td>0.58</td>
</tr>
<tr>
<td>KMLD-11</td>
<td>4.54**</td>
<td>0.20</td>
<td>0.64</td>
<td>6.44**</td>
<td>6.11**</td>
<td>-0.04</td>
<td>0.37</td>
<td>0.03</td>
<td>0.03</td>
<td>-0.31</td>
</tr>
<tr>
<td>KMLD-18</td>
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<td>0.31</td>
<td>-5.72**</td>
<td>-3.11**</td>
<td>-0.10**</td>
<td>0.49</td>
<td>0.10</td>
<td>1.29**</td>
<td>1.53**</td>
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<td>KMLD-19</td>
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<td>-1.02**</td>
<td>8.44**</td>
<td>6.62**</td>
<td>0.05**</td>
<td>0.78</td>
<td>-0.19</td>
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<td>3.33**</td>
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<tr>
<td>KMLD-21</td>
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<td>0.32</td>
<td>0.75**</td>
<td>-2.72</td>
<td>-4.50**</td>
<td>0.04</td>
<td>0.01</td>
<td>-0.81**</td>
<td>-1.14**</td>
<td>1.58**</td>
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<tr>
<td>KMLD-61</td>
<td>1.77</td>
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<td>2.50</td>
<td>0.03</td>
<td>0.40**</td>
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<td>0.36</td>
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<td>1.90</td>
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<td>0.07</td>
<td>0.14</td>
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<td>1.38</td>
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<td>0.05</td>
<td>-0.46**</td>
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<td>0.07</td>
<td>0.57**</td>
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<td>-0.25**</td>
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Significant at p<0.05; **Significant at p<0.01
Table 3 - Estimates of sca effects, standard heterosis and per se performance of best performing crosses.

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<td>Kernel rows</td>
<td>0.04</td>
<td>0.04</td>
<td>0.04</td>
<td>0.04</td>
<td>0.04</td>
<td>0.04</td>
</tr>
<tr>
<td>Number of kernels per row</td>
<td>1.37</td>
<td>1.37</td>
<td>1.37</td>
<td>1.37</td>
<td>1.37</td>
<td>1.37</td>
</tr>
<tr>
<td>Grain yield plant (g)</td>
<td>11.16**</td>
<td>11.16**</td>
<td>11.16**</td>
<td>11.16**</td>
<td>11.16**</td>
<td>11.16**</td>
</tr>
<tr>
<td>Days to 50% silking</td>
<td>77.22**</td>
<td>77.22**</td>
<td>77.22**</td>
<td>77.22**</td>
<td>77.22**</td>
<td>77.22**</td>
</tr>
<tr>
<td>Number of ears per plant</td>
<td>74.00</td>
<td>74.00</td>
<td>74.00</td>
<td>74.00</td>
<td>74.00</td>
<td>74.00</td>
</tr>
<tr>
<td>Ear girth (cm)</td>
<td>49.30**</td>
<td>49.30**</td>
<td>49.30**</td>
<td>49.30**</td>
<td>49.30**</td>
<td>49.30**</td>
</tr>
<tr>
<td>100 grain weight (g)</td>
<td>54.00</td>
<td>54.00</td>
<td>54.00</td>
<td>54.00</td>
<td>54.00</td>
<td>54.00</td>
</tr>
</tbody>
</table>

Significant at p<0.05; **Significant at p<0.01

stress, can have major effects on DM yield and composition of maize whole-plant for silage. Results from this study show that low DM yields and poor quality of maize whole-plant for silage are beyond drought stress. Daily maximum temperatures should be considered when planning strategies to insure good quality forage supply and reduce risk in dairy farming systems.

References


