Heterotic grouping of maize inbred lines on the yield basis of combining ability: Efficiencies and prospects

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Abstract

Founding the perfect fit of inbreds into heterotic groups is a pivotal to the development of triumphant maize hybrids. Here the objective of this study were to (i) classifying inbred line(s) into known heterotic groups with unknown testers, HGCA, HGY and HGCA methods (ii) analyze breeding efficiencies of this method(s) and (iii) future prospects of heterotic grouping research. These results demonstrate the usefulness of combining ability in classifying inbreds into useful heterotic group(s) for further studies.

Keywords: maize genotypes, heterotic groups, combing ability, line × tester mating design, grain yield

Introduction

In general, as we all know that maize (Zea mays L.) is a miracle crop and queen of cereals, because of its highest genetic yield potential among the cereals. The global maize production is about 1.09 billion metric tonnes from 153.0 million hectares and USA has highest productivity (10.57 t ha⁻¹), which is double than the global average (4.92 t ha⁻¹). Whereas, the average productivity in India is about 2.68 t ha⁻¹ with production of 24.26 million tonnes from 9.3 million hectares, the country lags far behind in productivity against world average. However, in Telangana State maize is grown in almost all the districts in an area of 0.64 million hectares, with a production of about 2.60 million tonnes [1].

Is that heterotic grouping is desirable? because of highly declining in yield reduction and increasing of population growth, we have to boost the grain yield production, considering the above points we have to select the best combination of inbreds in breeding programs for desired traits. Strength of a breeding program depends on the genetic variability in the base populations and development of superior inbreds, for that ‘n’ numbers of breeding methods are available, but they are time consuming. Hence, heterotic grouping is the best way to select the desired combination of inbreds in a short period of time and by using these researchers have great influence on how a maize inbred line is assigned to a heterotic group.

Heterosis and combining ability plays a significant role in crop improvement, as it helps the breeder to study and compare the performance of the new lines in hybrid combinations. It provides the basis for selecting good combiners and also for understanding the nature of gene action. Apart from selection of superior lines and analysis of their combining ability, placing them in well-defined heterotic groups is essential to increase the probability of success in heterosis breeding. Identification and utilization of heterotic groups and their patterns is essential in maize heterosis breeding [2]. A heterotic group is a group of related or unrelated genotypes displaying similar combining ability and giving a heterotic response when crossed to opposite or other genetically distinct germplasm group and establishment of the best combination of inbreds among the heterotic groups is crucial to the development of successful maize hybrids [3, 4, 5, 6].

Classification of inbreds into heterotic groups based on sca effects of grain yield is influenced by interaction between two inbred lines, hybrids and environment, which leads to classification of same inbred into different groups in different studies. This issue can be resolved by using heterotic group’s specific and general combining ability (HSGCA) method of heterotic grouping [7, 8, 9]. Here we classify inbreds into heterotic groups with unidentified testers on combining ability yield basis by studying their interactions in all way in line × tester mating fashion.
Materials and Methods
Genetic materials
The study material involved seeds of 20 maize inbred lines collected from the Maize Research Center (MRC), Hyderabad. Among this 15 lines different heterotic pattern are identified with five testers based on combining ability grain yield in a line x tester (L x T) mating fashion. The pedigree and genetic background of the inbred lines used for this study are listed in table 1.

Experimental Design
Heterotic groupings
GCA effects for grain yield of all the five testers were evaluated to identify a pair of testers with significantly highly divergent GCA to allocate the inbred lines and to commence heterotic groups (Legesse). Moreover, three presumed heterotic groups were also commenced among the rest of the testers which were not identified on the bases of their GCA values. In all circumstances, the inbred lines based on the SCA values enumerate from their crossing performance with the testers in line x tester mating design were allocated toward any one or both of the heterotic tester groups. An inbred line which possessed negative SCA with any one of a heterotic tester was grouped with the tester with which it revealed negative effects. Another inbred line revealing positive SCA with both testers was assigned towards both groups and, the one which revealed negative SCA with the two testers was discarded (vasal et al., 1992; Parenonti et al., 2001).

Table 1: List of maize inbred lines used in this study for heterotic grouping

<table>
<thead>
<tr>
<th>Genotypes</th>
<th>Pedigree</th>
<th>Source</th>
<th>Grain texture</th>
<th>Silking Colour</th>
</tr>
</thead>
<tbody>
<tr>
<td>MGC-6</td>
<td>(CML451-B*7/(CML451/CL-RCY016)-B-18-1-1-1-BBB)-B-11-BB</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>MGC-9</td>
<td>((CML161 x CML451)-B18-1-BBB/CML161-B)-B13-BB(NonQ)-BBB/CML395/MBRCSBF114-1.2-3-3-4-2-B)-DH-3007-B*4)-B-8-BB</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>MGC-15</td>
<td>(AMDROUTI( DT-Tester)Cl32-36-b*5/(POP502C5#18/GEWMN-0145)-B-21-2-1-1-B)-3-BB</td>
<td>MRC, AR(R)</td>
<td>Semi-flint</td>
<td>Orange</td>
</tr>
<tr>
<td>MGC-32</td>
<td>AMDROUTI(5x6)C3F2-B2-15-1-BB</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>sMG-92</td>
<td>PT9633001-B<em>4-1-B</em>6-1-BBB-#-BB</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>MGC-137</td>
<td>(MARSSYN-155)-4-1-1-BB</td>
<td>MRC, AR(R)</td>
<td>Dent</td>
<td>Orange</td>
</tr>
<tr>
<td>MGC-230</td>
<td>CML452=Acb3228BNC-166-1-1-1-B*15-#-BB</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow with cap</td>
</tr>
<tr>
<td>MGC-237</td>
<td>(POP501C5#8/GEWMN-0039)-B-10-1-1-1-BBB</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>MGC-238</td>
<td>(CML451/CA00360/P3011F2-3-5-6-1-B*10)-B-4</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Orange</td>
</tr>
<tr>
<td>MGC-239</td>
<td>(CML161 x CLQ-RCYQ49=) (CML176 CL-G2501-B)-B-21-2-1-1-B)-B-9-B*11</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Orange</td>
</tr>
<tr>
<td>MGC-242</td>
<td>CML227-B*12</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>MGC-248</td>
<td>DTPYC9-F-46-3-6-1-2-2-1-2-B*7-B-B</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Pinkish orange</td>
</tr>
<tr>
<td>MGC-252</td>
<td>NE8008-B*9</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow with cap</td>
</tr>
<tr>
<td>MGC-254</td>
<td>CLQ-RCYQ36-B-1-B*8-B-B</td>
<td>MRC, AR(R)</td>
<td>Semi-dent</td>
<td>Yellow</td>
</tr>
<tr>
<td>MGC-256</td>
<td>CA00360/P3011F2-3-5-6-1-B*12</td>
<td>MRC, AR(R)</td>
<td>Flint</td>
<td>Yellow with cap</td>
</tr>
<tr>
<td>BML-6</td>
<td>BML-6(SRRL65-896-1-1-2-#2-1-1-1-1</td>
<td>MRC, AR(R)</td>
<td>Semi-flint</td>
<td>Yellow</td>
</tr>
<tr>
<td>BML-7</td>
<td>BML-7(</td>
<td>x</td>
<td>x pollen - CML226-B9R1-1-1-1-xb-xb-xb-xb</td>
<td>MRC, AR(R)</td>
</tr>
<tr>
<td>GP-170</td>
<td>Selected from CIMMYT lines</td>
<td>CIMMYT</td>
<td>Dent</td>
<td>Yellow with cap</td>
</tr>
<tr>
<td>GP-311</td>
<td>Selected from CIMMYT lines</td>
<td>CIMMYT</td>
<td>Dent</td>
<td>Yellow with cap</td>
</tr>
</tbody>
</table>

ARI(R), Agricultural Research Institute (Rajendranagar); BML, Code for Hyderabad Maize Line; CIMMYT, International Maize and Wheat Improvement Center; GP, Germplasm; MGC, Maize Germplasm selected from CIMMYT; MRC, Maize Research Center.

Abbreviations: ARI(R), Agricultural Research Institute (Rajendranagar); BML, Code for Hyderabad Maize Line; CIMMYT, International Maize and Wheat Improvement Center; GCA, General Combining Ability; GP, Germplasm; HSGCA, Heterotic group’s Specific and General Combining Ability; MGC, Maize Germplasm selected from CIMMYT; MRC, Maize Research Center; SCA, Specific Combining Ability; SCA_GY, Specific Combining Ability and Grain Yield; USA, United States of America.

Heterotic group’s specific and general combining ability computation
Heterotic group’s specific and general combining ability (HSGCA) method proposed by Fan et al. (2009) was computed as follows:

\[
\text{HSGCA} = \left( \text{Cross mean (X)} \right) - \text{Line mean (X)} - \text{Tester mean (X)} + \text{Overall mean (X)}
\]

\[
\text{GCA} = \text{Line mean (X)} - \text{Overall mean (X)}
\]

Where,

\[
X_{ij} = \text{mean yield of the cross between } i^{th} \text{ tester and } j^{th} \text{ line}
\]

\[
X_{i} = \text{mean yield of the } i^{th} \text{ tester}
\]

\[
X_{j} = \text{mean yield of the } j^{th} \text{ line}
\]

Identification of testers: The key for HSGCA method is to identify representative tester lines (Fan et al. 2009). Grain yield GCA effects of all the testers are to be estimated to identify a pair of testers with significantly contrasting GCA to

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assign the inbred lines and to initiate heterotic groups (Legesse et al. 2009) [7].

**Formation of heterotic groups:** With the representative testers identified based on the GCA effects and divergence, heterotic grouping was done following the HSGCA method developed by Fan et al. (2009). The following steps were followed for heterotic grouping of the inbred lines.

**Step 1:** Placed all inbred lines with negative HSGCA effects into the same heterotic groups as their tester. At this step, a line might be assigned to more than one heterotic group.

**Step 2:** If an inbred line was assigned to more than one heterotic group in Step 1, we kept the line in the heterotic group if its HSGCA had the smallest value (or largest negative value) and removed it from other heterotic groups.

**Step 3:** If a line had positive HSGCA effect with all representative testers, we must be cautious to assign that line to any heterotic group because the line might belong to a heterotic group different from all the tester. In conclusion, In order to classify inbred lines into heterotic grouping, evaluated the 15 maize inbred lines with two highly divergent testers out of five testers and assigned the 13 lines into two heterotic groups A and B. Eight and five lines were belonged to tester group BML-7 (Group-A) and GP-311 (Group-B) respectively. This study demonstrates the usefulness of combining ability effects in classifying inbred lines into useful heterotic groups for further studies.

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**References**
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