Heterosis and Inbreeding Depression for Seed Cotton Yield and Yield Attributing Traits in Intrahirsutum 
(G. hirsutum L. X G. hirsutum L.) Hybrids of Cotton

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ABSTRACT

The present investigation was undertaken to estimate the extent of heterosis for yield and its quantitative traits in crosses of G. hirsutum. For this purpose, two hybrids were developed by using four parents during kharif 2014. These hybrids along with four parents were planted in kharif 2015. The hybrids DSMR10 x DRCR4 and DRGR2572 x M5 exhibited significant positive heterosis over mid parent and better parent for seed cotton yield and lint yield. These hybrids also showed significant heterosis for plant height, number of bolls per plant, boll weight, lint yield, number of monopodia and sympodia per plant.

Keywords: Heterobeltiosis, Mid-parent heterosis, Non-additive gene action, Inbreeding depression.

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Introduction

Cotton, the “king of fibre”, is an important cash crop having profound influence on economics and social affairs of the country. In India, cotton is the most important commercial crop, belonging to the family Malvaceae and the genus Gossypium. It has diversity in categories of cultivated species such as diploids and tetraploids, of which only four species (Gossypium arboreum L., Gossypium herbaceum L., Gossypium hirsutum L. and Gossypium barbadense L.) are cultivated. Among the four cultivated species, upland cotton (Gossypium hirsutum L.) is known for its production potential, while Egyptian cotton (Gossypium barbadense L.) is known for high fibre quality. The intraspecific hybrids of G. hirsutum L. have good fibre properties as well as higher yields.

The presence of non-additive gene action or overdominance is the primary justification for initiating a hybrid breeding program.

The present study was undertaken to find out the extent of heterosis for the seed cotton yield and its component characters in upland cotton (Gossypium hirsutum L.).

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Materials and Methods

The present study was conducted at Main Agricultural Research Station, Dharwad Farm, University of Agricultural Sciences Dharwad during Kharif 2014-15. The parent material used in the study comprised of two hybrids involving four parents. The resulting two F1s and four parents were grown in randomized block rows with two replication, having two rows in each replication of 5.4 m length, along with populations used for other studies during kharif 2015-16. Here, spacing in between rows and plant to plant was 90 cm and 45 cm respectively. Recommended agronomic practices were followed to raise the crop. Data were recorded for seed cotton yield, number of bolls per plant, boll weight, ginning outturn lint index, seed index, number of sympodia per plant and plant height. Population size of P1, P2 and F1 was 30 and 400 for F2 in both cross combination. Calculation were carried out according to the formulae given by Fehr (1987).

Results and Discussion

Results on mid-parent and better parent heterosis for eight characters are presented in Table 1. The trait seed cotton yield per plant showed high significant positive heterosis over mid-parent (54.26 %) and over better parent (40.79 %) in cross DSMR10 x DRCR4, also combination DRGR2572 X M5 also showed high significant positive heterosis over mid-parent (55.33 %) and over better parent (40.27 %). Similar results were reported by Ahuja and Tuteja (2000), Tuteja et al., (2004), Maisuria et al., (2007) and Wankhade et al., (2009). Significant inbreeding depression was observed from F1 to F2 for cross DSMR10 x DRCR4 (37.21 %) and DRGR2572 X M5 (29.82 %). These results are in conformity with Pradeep, et al., (2003), Shastri (2003) and Kumari et al., (2009).

Results for boll weight showed significance for both mid-parent heterosis (3.60 %) and heterobeltiosis (-3.42 %) were found to be significant in negative direction. In cross DSMR10 x DRCR4 and in cross DRGR2572 X M5 mid-parent heterosis (-5.65 %) and heterobeltiosis (-8.13 %) were significant. Inbreeding depression from F1 to F2 was highly significant in cross DSMR10 x DRCR4 (6.98 %) and in cross DRGR2572 X M5 it was non-significant. Inbreeding depression was significant in cross DSMR10 x DRCR4 the generations indicating the predominance of dominant gene action. Similar results were obtained by Kumari et al., (2009).

For ginning out-turn both mid-parent heterosis (9.89 %) and heterobeltiosis (9.62 %) were found to be highly significant in cross DSMR10 x DRCR4. In cross DRGR2572 X M5 mid-parent heterosis (7.35 %) and heterobeltiosis (8.50 %) were also
significant. Patel and Pethani (1998) and Tuteja et al., (2000) reported similar results. Significant and similar level of inbreeding depression was observed in both crosses 7.51 % in cross DSMR10 x DRCR4 and 7.49 % in DRGR2572 X M5. Similar results were noticed by Shastri (2003) and Kumari et al., (2009).

The trait lint index showed significant positive heterosis over mid-parent (31.38 %), heterobeltiosis (28.54 %) in cross DSMR10 x DRCR4 and in cross DRGR2572 X M5 both heterosis over mid-parent (32.53 %) and heterobeltiosis (33.81 %) were highly significant. Similar results were observed by Dukre et al., (2009). The inbreeding depression was highly significant in positive direction indicating the presence of non-additive gene action in both crosses DSMR10 x DRCR4 (30.85 %) and DRGR2572 X M5 (32.78 %).

Table 1 Percentage heterosis and inbreeding depression of eight quantitative characters

| Character                  | Cross                  | Heterosis (%) | Inbreeding depression (%) |
|----------------------------|------------------------|---------------|---------------------------|
|                            |                        | Hmp           | Hbp           | F2               |
| Seed cotton yield per plant| DSMR10 X DRCR4         | 54.26**       | 40.79**       | 37.21**         |
|                            | DRGR2572 X M5          | 55.33**       | 40.27**       | 29.82**         |
| Number of bolls per plant  | DSMR10 X DRCR4         | 35.45**       | 45.56**       | 35.45**         |
|                            | DRGR2572 X M5          | 54.15**       | 46.34**       | 20.31**         |
| Boll weight                | DSMR10 X DRCR4         | 3.60**        | -3.42*        | 6.98**          |
|                            | DRGR2572 X M5          | -5.65**       | -8.13**       | 1.43            |
| Ginning out turn           | DSMR10 X DRCR4         | 9.89**        | 9.62**        | 7.51**          |
|                            | DRGR2572 X M5          | 7.35**        | 8.50**        | 7.49**          |
| Lint index                 | DSMR10 X DRCR4         | 31.38**       | 28.54**       | 30.85**         |
|                            | DRGR2572 X M5          | 32.53**       | 33.81**       | 32.78**         |
| Seed index                 | DSMR10 X DRCR4         | 13.19**       | 11.15**       | 21.71**         |
|                            | DRGR2572 X M5          | 18.52**       | 17.72**       | 24.03**         |
| No. of Sympodia per plant  | DSMR10 X DRCR4         | -0.75         | -2.21         | 12.49*          |
|                            | DRGR2572 X M5          | 23.1**        | 36.16**       | 18.92**         |
| Plant height               | DSMR10 X DRCR4         | 5.82*         | 6.92          | 16.56**         |
|                            | DRGR2572 X M5          | 25.74**       | 26.10**       | 27.8**          |

*, ** - Significance at 5% and 1% probability level, respectively

For seed index mid-parent heterosis (13.19 %) and heterobeltiosis (11.15 %) were both highly significant in cross DSMR10 x DRCR4, in cross DRGR2572 X M5 both mid-parent heterosis (18.52 %) and heterobeltiosis (17.72 %) were also found to be highly significant. Similar results were observed by Shinde and Mehetre (2002) and Shastri (2003). Inbreeding depression in F2 was significant in both crosses, 21.71 % in DRCR4 X DSMR10 and 24.03 % in cross DRGR2572 X M5.

Both mid-parent heterosis and better parent
heterosis for number of sympodia per plant were non-significant in nature in cross DSMR10 x DRCR4 and significant mid-parent heterosis (23.10 %), better parent heterosis (36.16 %) was observed in cross DRGR2572 X M5. Similar observations were made by Patil et al., (2009) and Wankhade et al., (2009). Significant inbreeding depression was observed in both cross F₁ to F₂, 12.49 % in cross DSMR10 x DRCR4 and 18.92 % in cross DRGR2572 X M5. Similar results were obtained by Anuradha (1997), Shinde and Mehetre (2002) and Mahantesh Shastri (2003).

Plant height showed significant positive heterosis over mid-parent (25.74 %) and heterobeltiosis (26.10 %) in cross DRGR2572 X M5. Similar results were noticed by Maisuria et al., (2006) and Wankhade et al., (2009). Mid parent heterosis was significant while better parent heterosis was non-significant in cross DSMR10 x DRCR4. Inbreeding depression from F₁ to F₂ was 16.56 % in cross DSMR10 x DRCR4 and 27.80 % in cross DRGR2572 X M5. Anuradha (1997) and Shinde and Mehetre (2002) also observed positive inbreeding depression.

The two cross combinations showed significant positive heterosis for seed cotton yield and its component traits viz., number bolls/plant, number of sympodia and ginning outturn. Seth and Singh (1984) suggested that improvement in seed cotton yield could be achieved by giving due weightage to number of branches and number of bolls.

Significant heterosis and inbreeding depression was observed for all the characters studied in both crosses revealing that, characters are controlled by non-additive gene action and can be exploited for hybrid development. For varietal development, selection should be delayed upto F₆ generation, because values are not reliable in early generations.

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