

Combining ability and gene action studies in brinjal (*Solanum melongena* L.)

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Received: August 2017 / Accepted: November 2017

Abstract

Combining ability effects were estimated for fruit yield and its component characters in a line x tester analysis model comprising 24 crosses developed by using eight lines and three testers. On the basis of *gca* and *sca* variances, most of the characters under study indicated predominance of non additive gene action. The parents HABR-5, HABR-6, BL-3, Swarna Pratibha and HAB-899 were good general combiners for fruit yield, and fruit weight. The F_1 hybrid HABL-11 x HAB-381 was the combinations for fruit yield.

Keywords: Combining ability, Gene action, GCA, SCA, Predictability ratio

Introduction

Brinjal (*Solanum melongena* L.) is one of the important vegetable crops of the family Solanaceae being grown in India. Eggplant is widely consumed with distinct preferences for their size, shape and colour in different ethnic groups and geographical regions of the India and world. This crop possesses rich genetic diversity for various qualitative and quantitative traits and has a great potential for improvement as per preference of consumers. Yield being a complex character is influenced by a number of component characters and they are polygenically inherited. Hence it requires a better understanding of their gene action to formulate a sound breeding strategy. Line x tester analysis is one such approach to know the nature and magnitude of gene action and also identifies good general combiners and specific crosses for different economic traits. Thus combining ability analysis gives the estimates of combining ability effects and aids in selecting desirable parents and crosses for further exploitation. Hence the present study has been undertaken in a line x tester design

using eight lines and three testers to study gene action and combining ability of yield and its component traits.

Materials and Methods

Advanced breeding lines and released variety of brinjal from ICAR RCER Research Centre, Ranchi were used as parents in the crossing programme. Crosses were done in line x tester fashion using eight lines *viz.*, HABR-5, HABR-6, BL-22, HABL-11, HABL-3, BL-5, Swarna Pratibha and three testers *viz.*, HAB-898, HAB-899 and HAB-381. The trials were conducted during main season of 2008-09 and 2000-10 at Experimental Farm of ICAR RCER Research Centre, Ranchi (23.35° N and 85.33° E at 629m altitude). Total annual rainfall was 1430mm with 1100 mm during June to September and the average maximum and minimum temperatures 37°C and 40°C respectively. Eleven parents along with 24 F_1 s were evaluated in a randomized block design. The net plot size was 2.25m x 2.4m with 60 x 45 cm inter and intra spacing respectively. All the recommended agronomic practices were followed to raise the normal crop. Data was recorded on ten randomly selected plants in each treatment over each replication for seven characters *viz.*, fruit yield (q/ha), fruit weight (g), fruit length (cm), fruit breadth(cm), days to 50% flowering, number of branches/plant and plant height (cm). Combining ability analysis was done according to Kempthorne (1957) using SPAR 2.0. Relative importance of *gca* and *sca* was established by calculating predictability ratio $2\delta^2g/2\delta^2g + \delta^2s$ (Baker 1978). Average degree of dominance was estimated by using the formula VD / VA .

Results and Discussion

Analysis of variance indicated highly significant variance for all traits among parents and hybrids. The parents vs hybrids component was significant for most of the traits indicating superior performance of hybrids over parents (Table 1). The analysis of variance for combining ability, VA, VD, average degree of dominance, predictability

ratio, contribution of lines, testers and lines x testers are given in Table 2. The variance due to lines and testers (indicative of gca variance) were significant and higher in magnitude for fruit yield, fruit weight, fruit length, fruit breadth and plant height. This indicates that these characters are predominantly governed by additive gene action. Variance due to lines x testers which was related to dominance variance was significant for fruit yield, fruit weight, fruit length, fruit breadth and days to 50% flowering. Hence, both additive and dominance gene action were important for fruit yield, fruit weight, fruit length and fruit breadth. Similar results were reported by Aswani and Khandelwal (2005), Suneetha *et al.* (2008) and Sao and Mehta (2010). Predominance of non additive gene action for yield and yield components

was reported by Rai *et al.* (2005), Nalini *et al.* (2011) and Ramesh Kumar and Arumugam (2016). Choudhary (2001) established presence of both additive and non additive gene actions for inheritance of various characters. Predictability ratio was observed to be <0.5 for all the characters studied indicating the predominant role of non additive component of variance for improvement of these characters. Reddy and Patel (2014) and Prasad *et al.* (2010) revealed similar findings. Higher values of average degree of dominance for all these characters confirmed the results of above findings.

The estimates of general combining ability effects indicated that HABR-5 was best general combiner for fruit weight and fruit breadth, HABR-6 for fruit weight and fruit length, BL-22 for plant height, HABL-11 for days to

Table 1: Analysis of variance with parents and crosses

| Source of variation | d.f | Fruit yield (q/ha) | Fruit weight (g) | Fruit length (cm) | Fruit breadth (cm) | Days to 50% flowering | No. of branches/ plant | Plant height (cm) |
|---------------------|-----|--------------------|------------------|-------------------|--------------------|-----------------------|------------------------|-------------------|
| Replications | 1 | 214275.39* | 6299.84** | 7.09* | 0.68 | 20.20 | 0.017 | 196.90 |
| Treatments | 34 | 77732.12** | 4121.35** | 36.41** | 2.77** | 31.21* | 0.641 | 174.46** |
| Parents | 10 | 23199.32 | 3583.38** | 15.88** | 0.33* | 14.90 | 0.869 | 33.97 |
| Crosses | 23 | 91787.79** | 4504.46** | 43.74** | 3.80** | 37.58* | 0.568 | 243.12** |
| P. vs. C | 1 | 299779.55** | 689.76 | 73.30** | 3.56** | 47.71** | 0.048 | 0.32 |
| Error | 34 | 29649.61 | 476.47 | 1.43 | 0.18 | 16.94 | 0.516 | 83.78 |

*P = 0.05, **P = 0.01

Table 2: ANOVA for Line x Tester analysis

| Source of variation | d.f | Fruit yield (q/ha) | Fruit weight (g) | Fruit length (cm) | Fruit breadth (cm) | Days to 50% flowering | No. of branches/ plant | Plant height (cm) |
|-------------------------------------|-----|--------------------|------------------|-------------------|--------------------|-----------------------|------------------------|-------------------|
| Lines | 7 | 156262.49* | 7484.87** | 60.46** | 7.35** | 47.54 | 0.445 | 649.67** |
| Testers | 2 | 10944.31 | 1753.81 | 123.97** | 2.65 | 29.28 | 0.543 | 35.15 |
| Lines x Testers | 14 | 71099.51** | 3407.20** | 23.91** | 2.19** | 33.79* | 0.633 | 69.55 |
| Error | 34 | 29649.61 | 476.47 | 1.43 | 0.18 | 16.94 | 0.516 | 83.78 |
| Genetic component | | | | | | | | |
| V_A | | 1428.92 | 75.78 | 1.38 | 0.12 | 0.26 | -0.004 | 11.98 |
| V_D | | 22868.61 | 1762.06 | 32.06 | 1.78 | 9.6 | 0.018 | 66.86 |
| V_D/V_A | | 16.00 | 23.25 | 23.23 | 14.83 | 36.92 | -4.5 | 5.58 |
| V_{gca} | | 714.46 | 37.89 | 0.69 | 0.06 | 0.13 | -0.002 | 5.99 |
| V_{scn} | | 22868.61 | 1762.06 | 32.06 | 1.78 | 9.6 | 0.018 | 66.86 |
| Predictability ratio | | 0.06 | 0.04 | 0.04 | 0.06 | 0.026 | -0.29 | 0.15 |
| Contribution of lines (%) | | 51.81 | 50.57 | 43.07 | 58.82 | 38.50 | 23.84 | 81.33 |
| Contribution of s (%) | | 1.04 | 3.39 | 24.65 | 6.07 | 6.77 | 8.31 | 1.26 |
| Contribution of lines x testers (%) | | 47.15 | 46.04 | 33.28 | 35.11 | 54.72 | 67.85 | 17.41 |

*P = 0.05, **P = 0.01

Table 3: GCA effects of 11 parents for seven characters in brinjal

| Source of variation | Fruit yield (q/ha) | Fruit weight (g) | Fruit length (cm) | Fruit breadth (cm) | Days to 50% flowering | No. of branches/ plant | Plant height (cm) |
|--------------------------|--------------------|------------------|-------------------|--------------------|-----------------------|------------------------|-------------------|
| <i>GCA effects Lines</i> | | | | | | | |
| HABR-5 | -185.81* | 56.68** | -4.38** | 2.20** | 0.25 | 0.16 | -2.53 |
| HABR-6 | -162.22* | 33.80** | 5.25** | -0.03 | 2.35 | -0.30 | -0.08 |
| BL-22 | -156.02* | 10.18 | -1.94** | 0.25 | -0.08 | 0.008 | -20.14** |
| HABL-11 | -50.94 | -45.49** | -1.61** | -0.99** | -5.85** | -0.26 | -5.28 |
| HABL-2 | 140.33 | -3.78 | 0.07 | 0.28 | -0.65 | -0.13 | -0.16 |
| BL-3 | 205.70** | 8.92 | -1.28** | 0.39* | -0.71 | 0.41 | 9.30 |
| BL-5 | 31.37 | -18.07 | -0.10 | -0.69** | 0.95 | 0.33 | 14.59** |
| Swarna Pratibha | 177.60* | -42.24** | 3.98** | -1.40** | 3.72* | -0.22 | 4.30 |

*P = 0.05, **P = 0.01

Table 4: SCA effects in 8x3 (Line x Tester) crosses of brinjal

| Source of variation | Fruit yield (q/ha) | Fruit weight (g) | Fruit length (cm) | Fruit breadth (cm) | Days to 50% flowering | No. of branches/plant | Plant height (cm) |
|----------------------|--------------------|------------------|-------------------|--------------------|-----------------------|-----------------------|-------------------|
| HABR-5 x HAB-898 | -40.06 | -9.24 | -1.53 | -0.05 | -3.87 | 0.15 | -9.49 |
| HABR-5 x HAB-899 | 124.21 | -4.79 | -1.47 | 0.93** | 4.03 | 0.54 | 0.58 |
| HABR-5 x HAB-381 | -84.15 | 14.03 | 3.00** | -0.88** | -0.17 | -0.69 | 8.91 |
| HABR-6 x HAB-898 | 164.66 | -47.12** | 6.50** | -1.79** | -2.17 | -0.27 | 3.18 |
| HABR-6 x HAB-899 | -99.48 | 68.59** | -3.25** | 1.60** | 4.23 | -0.62 | 0.62 |
| HABR-6 x HAB-381 | -65.18 | -21.47 | -3.25** | 0.19 | -2.07 | 0.89 | -3.80 |
| BL-22 x HAB-898 | -111.05 | 5.51 | -4.33** | 0.97** | 2.87 | -0.03 | -3.71 |
| BL-22 x HAB-899 | 233.51 | 58.09** | 4.78** | 0.16 | -3.73 | 0.20 | 2.06 |
| BL-22 x HAB-381 | -122.47 | -63.60** | -0.46 | -1.13** | 0.87 | -0.16 | 1.64 |
| HABL-11 x HAB-898 | -216.13 | -9.20 | -0.99 | -0.29 | 1.13 | -0.31 | 4.64 |
| HABL-11 x HAB-899 | -142.08 | -38.00* | -0.08 | -0.78** | 0.33 | -0.04 | -4.17 |
| HABL-11 x HAB-381 | 358.21** | 47.20** | 1.06 | 1.07** | -1.47 | 0.35 | -0.47 |
| HABL-2 x HAB-898 | 105.89 | 22.84 | 0.62 | -0.10 | -4.47 | 0.19 | 8.89 |
| HABL-2 x HAB-899 | -135.02 | -19.45 | 0.73 | -0.70* | -1.87 | 0.59 | -1.92 |
| HABL-2 x HAB-381 | 29.13 | -3.39 | -1.35 | 0.80* | 6.33 | -0.77 | -6.97 |
| BL-3 x HAB-898 | 141.77 | 32.27* | 0.54 | 0.52 | 7.20* | -0.23 | 1.55 |
| BL-3 x HAB-899 | 108.73 | -25.53 | 0.32 | -0.62** | -4.80 | -0.08 | 0.50 |
| BL-3 x HAB-381 | -250.49* | -6.74 | -0.86 | 0.10 | -2.40 | 0.31 | -2.05 |
| BL-5 x HAB-898 | -94.64 | 19.88 | -2.14* | 0.81* | 0.43 | -0.14 | -7.36 |
| BL-5 x HAB-899 | -20.50 | -24.66 | -2.44** | -0.03 | 1.43 | 0.13 | 4.71 |
| BL-5 x HAB-381 | 115.14 | 4.78 | 4.59** | -0.78* | -1.87 | 0.18 | 2.69 |
| S.Pratibha x HAB-898 | 49.56 | -14.95 | 1.32 | -0.07 | -1.13 | 0.64 | 2.30 |
| S.Pratibha x HAB-899 | -69.37 | -14.25 | 1.40 | -0.56 | 0.37 | -0.71 | -2.38 |
| S.Pratibha x HAB-381 | 19.82 | 29.20 | -2.72** | 0.63* | 0.77 | 0.06 | 0.08 |
| S.E | 121.76 | 15.43 | 0.85 | 0.30 | 2.91 | 0.51 | 6.47 |

*P = 0.05, **P = 0.01

50% flowering, BL-3 for fruit yield, fruit breadth, Swarna Pratibha for fruit length and fruit yield, HAB-898 for fruit length, HAB-899 for fruit weight and fruit length and HAB-381 for fruit breadth. None of the parents displayed desirable general combining ability effects for all traits indicating scope for improving their general combining ability. In general, contribution of lines was higher for all traits except for days to 50% flowering and number of branches/plant where percentage contribution of lines x testers was highest.

Highly significant specific combining ability effects (Table 4) were observed for the crosses HABR-5 x HAB-899 for fruit breadth, HABR-5 x HAB-381, HABR-6 x HAB-898 for fruit length, HABR-6 x HAB-899 for fruit weight and fruit breadth, BL-22 x HAB-898 for fruit breadth, BL-22 x HAB-899 for fruit weight and fruit length, HABL-11 x HAB-381 for fruit yield, fruit weight and fruit breadth, HABL-2 x HAB-381 for fruit breadth, BL-3 x HAB-898 for fruit weight, BL-5 x HAB-898 and Swarna Pratibha x HAB-381 for fruit breadth. These hybrids with higher specific combining ability effects are useful to derive high performing hybrids. Fifteen crosses had significant specific combining ability effects in addition to high *per se* performance for fruit yield, fruit weight, fruit length and fruit breadth (Table 5). These crosses involved parents with high x high, high x low and low x low general combining ability effects indicating presence of additive, dominance and epistatic

Table 5: GCA Status and *per se* performance of crosses with significant SCA effects

| Character | Crosses | GCA effects | Mean |
|--------------------|---------------------------|-------------|--------|
| Fruit yield (q/ha) | HABL-11 x HAB-381 | L x L | 47.71 |
| Fruit weight (g) | HABR-6 x HAB-899 | H x H | 165.25 |
| | BL-22 x HAB-899 | L x H | 125.88 |
| | HABL-11 x HAB-381 | L x L | 82.13 |
| | BL-3 x HAB-898 | L x L | 103.13 |
| Fruit length (cm) | HABR-5 x HAB-381 | L x L | 16.90 |
| | HABR-6 x HAB-898 | H x H | 10.63 |
| | BL-22 x HAB-899 | L x H | 14.93 |
| Fruit breadth (cm) | HABR-5 x HAB-899 | H x L | 4.91 |
| | HABR-6 x HAB-899 | L x L | 5.65 |
| | BL-22 x HAB-898 | L x L | 6.04 |
| | HABL-11 x HAB-381 | L x H | 3.18 |
| | HABL-2 x HAB-381 | L x H | 4.25 |
| | BL-5 x HAB-898 | L x L | 5.06 |
| | Swarna Pratibha x HAB-381 | L x H | 3.80 |

gene actions for controlling these characters. Similar results were obtained by Nalini et al. (2011), Deshmukh et al. (2015) and Prasad et al. (2015). High x low general combining ability combinations are suitable for heterosis breeding. High x high general combining ability combinations can be considered for developing superior variants through pedigree method.

LKjka k

बैंगन में फल उपज तथा इसके सम्बन्धित कारकों के लिए संयोजन क्षमता के प्रभावों का आंकलन लाइन x टेस्टर प्रजनन विधि द्वारा किया गया जिसमें 8 लाइनों एवं 3 टेस्टरों के प्रयोग से 24 संकरों

को विकसित कर मूल्यांकन किया गया। अध्ययन के अन्तर्गत अधिकांश विशेषतायें, सामान्य संयोजन क्षमता तथा विशिष्ट संयोजन क्षमता विविधता के आधार पर, असंयोजी जीन क्रिया हेतु प्रभावी पाया गया। पितृ एच.ए.बी.आर.-5, एच.ए.बी.आर.-6, बी.एल.-3, स्वर्ण प्रतिभा एवं एच.ए.बी.-899 को फल उपज एवं फल वजन का सामान्य संयोजन क्षमता उत्तम पाया गया। प्रथम पीढ़ी के संकर एच.ए.बी.एल.-11 x एच.ए.बी.-381 तथा एच.ए.बी.आर.-6 x एच.ए.बी.-899 को फल उपज एवं इसके सम्बन्धित घटकों के लिए सर्वश्रेष्ठ विशिष्ट संयोजी पाया गया।

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