

Original Research Article

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Combining Ability Analysis for Yield and Yield Contributing Traits in Maize (*Zea mays* L.) Hybrids

G. B. Bharat^{1*}, P. B. Wadikar¹, M. V. Dhuppe² and N. R. Thakur³

¹Department of Agricultural Botany, College of Agriculture, Latur, India

²Oilseed Research Station, Latur, India

³Department of Agricultural Botany, College of Agriculture, Parbhani, India

*Corresponding author

ABSTRACT

Study on combining ability effects were carried out in 28 hybrids obtained by crossing 8 x 8 genotypes in half diallel fashion and parents, hybrids evaluated with standard check Maharaja in RBD with three replication during *khari*2018-19 at research farm of Agriculture botany College of Agriculture, Latur for twelve traits *viz.*, days to 50 per cent tasseling, days to 50 per cent silking, days to maturity, plant height (cm), ear length (cm), ear girth (cm), kernel row number, number of grains per row, 100-grain weight (g), fodder yield per plant (g), grain yield per plant (g) and Harvest index (%). The analysis of variance revealed that significant differences among treatments, parents, hybrids and parents *v/s* hybrids for all characters except days to 50 per cent tasseling and days to 50 percent silking. This validated that, considerable amount of variability among experimental material. The combining ability analysis revealed the presence of higher magnitude of SCA than GCA variance for all characters under study. The ratio of additive to dominance variance was lower than unity for all traits, indicating higher non-additive variance than additive variance. Among parents, 231-A, Hyd.19051 and Hyd.19051 was found to good general combiner for grain yield per plant. While the parent 231-A exhibited good general combining ability for majority of characters. Best specific combinations were Hyd.18227 x Hyd.08R-740-2, Hyd.19051 x Hyd.12288, Hyd.18227 x Hyd.18169 and Hyd.18063 x Hyd.18169 for grain yield per plant.

Keywords

Maize, Combining ability, GCA and SCA

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Introduction

Maize is third most important cereal crop after rice and wheat in the world. It is referred as the “Queen of Cereals” and “Miracle Crop” due to its high productivity potential as compared to the other cereals. Being a potential field crop in India, maize occupies an important place as a source of human food (26%), animal feed (11%), poultry feed (43%), industrial products (12%) and three

per cent for seed and other purposes (FICCI). Many forms of maize are used i.e. Flour corn, Popcorn, Flint corn, Dent corn, Waxy corn, Sweet corn, Pod corn and Stripped maize.

India is the fifth largest producer of maize in the world contributing 3% of the total global production. It is cultivated on nearly 150 million ha in about 160 countries having wider diversity of soil, climate, biodiversity and management practices that contributes

36% (782 MT) of the global grain production. The United States of America (USA) is the largest producer of maize contributes nearly 35% of the total production in the world. The important maize growing countries are USA, China, Brazil, Mexico, India, Philippines, South Africa and Indonesia.

In India, major maize growing states are Karnataka, Madhya Pradesh, Maharashtra, Rajasthan, Bihar, Uttar Pradesh, Telangana, Gujarat and Tamil Nadu. In India, maize has been grown in an area of 9.2 million ha with production of 24.17 MT and average productivity of 2.56 t/ha.

Being a highly cross pollinated crop, it offers tremendous scope for the plant breeders for genetic improvement. It possesses one of the most well studied genetic systems among cereals which have motivated a rich history of research into the genetics of various traits in maize. Information on different traits of interest especially their genetic control is prerequisite for planning the genetic improvement strategies.

The main goal of maize breeding is to obtain new hybrids with high genetic potential for yield and positive features that exceed the existing commercial hybrids. The commercial production of hybrids however, depends upon two factors *viz.*, the behavior of the line itself and the behavior of line in hybrid combination. The behavior of a line in hybrid combination is assessed through the estimation of combining ability effects.

Sprague and Tatum (1942) coined combining ability to designate the average performance of line in hybrid combination and to define those cases in which certain combination do relatively better or worse than would be expected on the basis of the average performance of the lines involved. Diallel crossing program have been applied to

achieve this goal by providing a systematic approach for detection of suitable parents and crosses for the investigated characters.

Materials and Methods

Total eight maize inbred lines (10306, Hyd.19051, Hyd.12288, Hyd.18227, Hyd.18063, Hyd.18169, Hyd.08R-740-2, 231-A) were used as parents in 8 x 8 half diallel crossing scheme. All the possible crosses $n(n-1)/2$ excluding reciprocals were effected in a half diallel mating design to produce 28 hybrids during *summer* 2018 at Experimental Farm, Department of Agril. Botany, College of Agriculture, Latur. Seeds were sown in the field by keeping plant to plant and row to row distances 20 cm and 60 cm respectively.

At maturity all the parents were crossed following half diallel mating design, resultantly 28 F₁s were generated. Seeds harvested from F₁s, parents and standard check (Maharaja) were sown by following randomized block design (RBD) with three replications. All the agronomic practices were taken for establishment of proper crop stand in field.

Data were recorded for yield components i.e. days to 50 per cent tasseling, days to 50 per cent silking, days to maturity, plant height (cm), ear length (cm), ear girth (cm), kernel row number, number of grains per row, 100-grain weight (g), fodder yield per plant (g), grain yield per plant (g) and harvest index (%). The observations were recorded on randomly selected five plants and subjected to statistical analysis. Analysis of variance technique suggested by Panse and Sukhatme (1967) was followed to test the differences among the genotypes for all the characters. Combining ability analysis was performed with the data obtained for parents and hybrids according to Model-I, Method-II proposed by Griffing (1956).

Results and Discussion

Analysis of variance

The analysis of variance for randomized block design was carried out (Table. 1) for the character of yield and its components to see the ignorance of differences among the treatments, replications and other partitioning source of variation. The analysis of variance revealed that, the all characters i.e. days to 50% tasseling, days to 50% silking, days to maturity, plant height (cm), ear length (cm), ear girth (cm), kernel row number, no. of grains per row, 100 grain weight, fodder yield per plant (g), grain yield per plant (g), harvesting index (%) had showed significant differences between treatments as well as parents. Parents *v/s* F₁s, were significant all the characters except days to 50% tasseling and days to 50% silking. Similar results reported by Jawahar and Kumar(2012) and Ratna *et al.*, (2014).

Combining ability effects

The knowledge on combining ability assists in the selection of suitable parental line. Among various biometrical techniques available, combining ability analysis proposed by Griffing (1956) had been extensively used by breeders. It provides information on the performance of genotypes in hybrid combination and also nature of gene action involved in the control of metric traits.

General combining ability effects

The general combining ability effects of eight parents for twelve traits were presented in Table 3. Among the eight parents studied, Hyd.12288 (-1.621), Hyd.10306 (-0.654) and Hyd.19051 (-0.454) expressed significant and desirable negative GCA effects and proved to be good general combiner for days to tasseling while parents, Hyd.12288 (-1.258),

Hyd.08R-740-2 (-0.942), Hyd.10306 (-0.792) and Hyd.19051 (-0.492) expressed significant and desirable negative GCA effects and proved to be good general combiner for days to silking. For days to maturity the parents Hyd.10306 (-1.207), Hyd.18063 (-1.117) and Hyd.12288 (-0.150) shows negatively significant GCA effects.

The negative significant GCA effects was registered for ear height by three parents Hyd.10306 (-5.71), 231-A (-3.58) and Hyd.18169 (-2.11). While two parents *viz.*, Hyd.10306 (0.352) and 231-A (0.795) depicted significant and positive GCA effects for ear length. For ear girth parent Hyd.19051 (0.421) exhibited highest positively significant GCA effect.

The trait kernel row number was recorded significant positive GCA effects in two parents 231-A (0.439) and Hyd.12288 (0.366) while parent, 231-A (1.545) explored highest positive significant GCA effect for number of grains per row. Among the parents, the parent Hyd.19051 (2.30) exhibited highest positively significant GCA effect for 100 grain weight. For fodder yield per plant parents, Hyd.18169 (5.792) and Hyd.18063 (2.225) were depicted positive significant GCA effects. The parent 231-A manifested highest, significant and positive GCA effects (7.408) followed by Hyd.19051 (6.342) and Hyd.19051 (4.842) respectively for grain yield per plant. Among the eight parents, three parents showed positively significant GCA effects for harvesting index. Out of eight parents, 231-A was found to good general combiner for eight characters including grain yield per plant and the parents, Hyd.19051 and Hyd.12288 for seven characters. The high positive value of GCA effect of these parents indicates that their contribution in transferring those traits to their hybrids is high. This is in accordance with findings of Gosai *et al.*, (2017), Begum *et al.*, (2018) and Karim *et al.*, (2018).

Table.1 Analysis of variance for different characters in maize

Source	D.F.	Days to 50% tasseling	Days to 50% silking	Days to maturity	Plant height (cm)	Ear length (cm)	Ear girth (cm)	Kernel row number	No. of grains per row	100 grain weight (g)	Fodder yield per plant (g)	Grain yield per plant (g)	Harvesting index (%)
Replications	2	0.210	0.009	1.009	6.953	0.035	0.015	1.313	0.826	0.720	0.481	11.861	0.335
Treatments	35	14.575**	13.831**	38.716**	519.433**	9.547**	2.853**	3.076**	44.066**	39.461**	256.879**	1251.188**	77.029**
Parents	7	16.946**	18.755**	34.089**	706.755**	5.997**	1.943**	2.272**	18.372**	9.787**	103.184**	518.547**	48.377**
F ₁ 's	27	14.465**	13.067**	40.791**	458.55**	9.524**	2.396**	2.277**	46.348**	48.525**	299.070**	1321.373**	76.458**
Parents v/s F ₁ 's	1	0.955	0.001	15.080**	852.000**	35.017**	21.559**	30.260**	162.315**	69.711**	193.572**	4484.666**	293.004**
Error	70	1.348	1.409	1.685	30.134	0.628	0.093	0.530	1.118	1.009	17.34	53.880	2.85

* Significant at 5 % level, ** Significant at 1 % level

Table.2 Analysis of variance of combining ability for different characters in maize

Source	D. F.	Days to 50% tasseling	Days to 50% silking	Days to maturity	Plant height (cm)	Ear length (cm)	Ear girth (cm)	Kernel row number	No. of grains per row	100 grain weight (g)	Fodder yield per plant (g)	Grain yield per plant (g)	Harvesting index (%)
GCA	7	7.704**	8.802**	7.994**	191.302**	2.496**	1.187**	0.781**	13.798**	26.955**	79.872**	389.485**	26.388**
SCA	28	4.147**	3.563**	14.133**	168.605**	3.354**	0.892**	1.086**	14.912**	9.704**	87.065**	423.957**	25.499**
Error	70	0.450	0.470	0.562	10.045	0.209	0.031	0.177	0.373	0.336	5.783	17.960	0.951
δ^2 gca/sca		0.096	0.169	0.045	0.014	0.027	0.034	0.033	0.007	0.184	0.008	0.008	0.003

Table.3 The general combining ability effects of eight parents for twelve traits

Sr. No.	Parents/ crosses	Days to 50% tasseling	Days to 50% silking	Days to maturity	Plant height (cm)	Ear length (cm)	Ear girth (cm)	Kernel row number	No. of grains per row	100 grain weight (g)	Fodder yield per plant (g)	Grain yield per plant (g)	Harvesting index (%)
	Parents												
1	Hyd.10306	-0.654**	-0.792**	-1.207**	-5.717**	0.352*	-0.142**	-0.218	1.525**	-1.186**	-1.775*	-0.558	0.253
2	Hyd.19051	-0.454**	-0.492*	-0.017	3.983**	0.028	0.421**	-0.024	-0.275	2.301**	-2.875**	6.342**	1.903**
3	Hyd.12288	-1.621**	-1.258**	-0.150	-1.550	-0.095	0.408**	0.366**	0.568**	-0.236	0.592	4.842**	1.034**
4	Hyd.18227	0.412**	-0.058	0.983**	6.883**	-0.925**	-0.009	-0.247*	0.072	0.041	-0.808	1.108	0.134
5	Hyd.18063	0.863**	1.675**	-1.117**	3.883**	0.048	-0.195**	-0.157	-0.888**	-2.266**	2.225**	-10.258**	-2.553**
6	Hyd.18169	0.062	-0.092	-0.083	-2.117**	-0.328*	-0.112*	0.112	-1.778**	0.421*	5.792**	-2.458	-1.672**
7	Hyd.08R-740-2	1.046**	-0.942**	0.050	-1.783	0.125	-0.602**	-0.271*	-0.768**	-1.323**	-1.442*	-6.425**	-0.960**
8	231-A	0.346	0.075	1.450**	-3.583**	0.795**	0.231**	0.439**	1.545**	2.248**	-1.708*	7.408**	1.862**
	S.E.gi	0.395	0.404	0.442	1.869	0.270	0.104	0.248	0.360	0.342	1.418	2.500	0.575
	S.E.gi-gj	0.598	0.611	0.668	2.826	0.408	0.157	0.375	0.544	0.517	2.144	3.779	0.869
	Crosses												
9	Hyd.10306 x Hyd.19051	3.159**	2.070**	-0.441	11.863**	2.759**	1.170**	0.489	3.081**	-0.210	0.863	14.744**	2.652
10	Hyd.10306 x Hyd.12288	0.326	-0.163	3.026**	5.396	1.349**	0.684**	0.632	2.438**	-2.674**	-12.270**	3.911	3.252*
11	Hyd.10306 x Hyd.18227	-3.374**	-3.363**	-4.441**	10.296**	-1.448**	-1.066**	-1.321**	-4.065**	-4.217**	10.463**	-39.689**	-11.166**
12	Hyd.10306 x Hyd.18063	-0.157	0.237	-4.341**	-16.704**	0.506	0.187	0.189	2.628**	-2.477**	11.430**	-5.322	-2.325**
13	Hyd.10306 x Hyd.18169	-3.357**	-1.663*	1.959**	1.963	0.049	-0.230	0.119	-0.815	-1.364*	-1.137	2.211	0.871
14	Hyd.10306 x Hyd.08R-740-2	2.326**	1.304*	5.826**	-6.370*	-0.504	0.660**	0.402	-4.959**	1.746**	13.763**	-8.156*	-3.025**
15	Hyd.10306 x 231-A	-0.307	-0.496	1.093	4.430	0.459	1.260**	0.526	5.128**	0.510	-15.304**	11.678**	3.817*
16	Hyd.19051 x Hyd.12288	-1.874**	-0.796	-4.074**	-4.304	2.072**	0.120	-0.528	4.738**	2.840**	10.496**	29.678**	4.025**
17	Hyd.19051 x Hyd.18227	-1.907**	-1.663*	4.126**	-17.404**	0.002	-0.130	-0.814*	-3.432**	2.563**	-3.104	-6.589	-0.353
18	Hyd.19051 x	-0.024	-0.396	-4.441**	5.263	-0.304	1.190**	1.029*	1.661**	1.370*	0.863	14.444**	2.851

	Hyd.18063												
19	Hyd.19051 x Hyd.18169	0.443	0.704	-0.807	4.596	0.206	0.140	-0.408	-1.049	5.383**	-1.370	13.644**	3.007*
20	Hyd.19051 x Hyd.08R-740-2	-0.541	-1.330*	4.393**	-1.737	-2.981**	-0.470*	1.042*	-1.692**	-2.840**	-4.470	-14.722**	-2.282**
21	Hyd.19051 x 231-A	-1.841**	-0.130	2.326**	-3.270	1.816**	0.297*	-0.268	3.261**	-0.110	-12.537**	-0.222	2.563
22	Hyd.12288 x Hyd.18227	0.593	0.104	0.926	-5.204	-1.174*	-1.216**	-0.404	-0.342	1.533**	1.430	-3.089	-0.700
23	Hyd.12288 x Hyd.18063	4.809**	5.370**	5.026**	-17.537**	-1.348**	0.304	-0.194	0.618	-2.707**	-3.604	-9.389*	-1.443
24	Hyd.12288 x Hyd.18169	1.276*	2.470**	1.659*	-5.537	0.762*	0.687**	1.269**	-1.592**	0.753	10.830**	-1.856	-1.500
25	Hyd.12288 x Hyd.08R-740-2	1.293*	0.104	0.526	11.796**	-0.858	0.110	-0.181	0.731	-1.570**	-12.604**	-6.889**	1.034
26	Hyd.12288 x 231-A	-3.007**	-1.696*	2.126**	-6.404*	1.339**	0.310*	0.909*	0.618	2.460**	4.330	15.278**	1.706
27	Hyd.18227 x Hyd.18063	-0.891	-1.496*	-2.441**	2.363	-1.551**	-0.080	0.586	-2.785**	-2.037**	5.130*	-7.656	-2.387**
28	Hyd.18227 x Hyd.18169	1.909**	0.604	-3.474**	11.363**	1.426**	1.304**	0.982*	6.238**	-0.690	-3.770	26.878**	6.069**
29	Hyd.18227 x Hyd.08R-740-2	-1.407*	-0.096	-1.941**	0.030	5.572**	2.027**	2.299**	9.628**	2.920**	-15.537**	54.178**	12.990**
30	Hyd.18227 x 231-A	0.626	0.104	0.993	-24.170**	-1.098*	-2.240**	-0.478	-4.019**	2.283**	2.396	-0.656	-0.218
31	Hyd.18063 x Hyd.18169	-1.707*	-1.463*	-1.707*	13.696**	1.652**	0.157	0.226	1.531**	3.583**	-7.804*	19.578**	5.723**
32	Hyd.18063 x Hyd.08R-740-2	0.476	1.837**	-6.841**	-0.970	-1.268**	0.780**	1.276**	-0.745	-2.207**	-16.904**	-2.789	2.591
33	Hyd.18063 x 231-A	-1.157	-1.630*	3.426**	11.830**	0.429	-0.853**	0.232	2.608**	1.590**	1.363	11.711**	2.256
34	Hyd.18169 x Hyd.08R-740-2	-0.057	-0.396	3.459**	5.696	0.209	-1.136**	-0.694	-0.589	-4.894**	7.863*	-28.589**	-7.920**
35	Hyd.18169 x 231-A	1.643*	1.804**	1.726*	-3.837	0.506	0.997**	1.396**	1.098	3.036**	-3.302	17.911**	4.502**
36	Hyd.08R-740-2 x 231-A	1.326*	0.104	2.074**	-29.170**	-0.014	-0.280	-0.388	-3.579**	-2.254**	12.363**	-3.789	-1.934
	S.E sij	1.212	1.239	1.355	5.731	0.827	0.319	0.760	1.104	1.048	4.348	7.664	1.763
	S.E sij - sik	1.794	1.833	2.005	8.48	1.224	0.472	1.125	1.663	1.551	6.434	11.339	2.609

Specific combining ability effects

The specific combining ability effects of 28 direct crosses with respect to twelve traits presented in Table 3. The estimates of the SCA effects of 28 hybrids ranged from -39.689 (Hyd.10306 x Hyd.18227) to 54.178 (Hyd.18227 x Hyd.08R-740-2) for grain yield per plant. The cross, Hyd.10306 x Hyd.18227 (-3.374) expressed highest significant and negative SCA effect for days to tasseling and days to silking. For days to maturity highest significant negative SCA recorded by the cross, Hyd.18063 x Hyd.08R-740-2 (-6.841) followed by Hyd.10306 x Hyd.18227 (-4.441) and Hyd.19051 x Hyd.18063 (-4.441).

Highest significant positive SCA recorded by the cross, Hyd.18063 x Hyd.18169 (13.696) for plant height followed by Hyd.10306 x Hyd.19051 (11.863) and Hyd.18063 x 231-A (11.830). For ear length and ear girth highest SCA effect recorded by the cross, Hyd.18227 x Hyd.08R-740-2 (5.572). For kernel row number out of 28 crosses, eight showed positively significant SCA effect.

The cross, Hyd.18227 x Hyd.08R-740-2 displayed maximum SCA effect (2.299) followed by Hyd.18169 x 231-A (1.396) and Hyd.18063 x Hyd.08R-740-2 (1.276). For number of grains per row Hyd.18227 x Hyd.08R-740-2 (9.628) had exhibited highest positively significant SCA effect. Among the hybrids, the cross Hyd.19051 x Hyd.18169 (5.383) was showed highest positively significant SCA effects for 100 grain weight followed by Hyd.18063 x Hyd.18169 (3.583) and Hyd.18169 x 231-A (3.036) respectively. For fodder yield per plant out of 28 crosses, eight showed positively significant SCA effect. The cross, Hyd.10306 x Hyd.08R-740-2 (13.763) displayed maximum SCA effect. For grain yield per plant, SCA effect fluctuated between -39.689 (Hyd.10306 x Hyd.18227) to 54.178 (Hyd.18227 x

Hyd.08R-740-2).

Among the 28 crosses, eleven crosses depicted significant and positive SCA effect under study. The cross, Hyd.18227 x Hyd.08R-740-2 (54.178) registered highest significant positive SCA followed by Hyd.19051 x Hyd.12288 (29.678), Hyd.18227 x Hyd.18169 (26.878) and Hyd.18063 x Hyd.18169 (19.578) respectively. For harvesting index Hyd.18227 x Hyd.08R-740-2 (12.990) had exhibited highest positively significant SCA effect followed by Hyd.18227 x Hyd.18169 (6.069) and Hyd.18063 x Hyd.18169 (5.723).

Good general combining inbred parents have not always showed high SCA effects in their cross combination. Thus it may be concluded that the information on GCA effects alone may not be sufficient to predict the extent of hybrid vigour by a particular cross combination. In many crosses significant SCA effects for yield per plant were associated with negative SCA effects of days to maturity.

This is desirable because we want hybrids having high yield and early in crop duration (Yerva *et al.*, 2016). In general, crosses involving both good general combiners as well as one good and other poor combiner showed high SCA effects, which is due to additive x additive and dominance x dominance gene action. Evaluation of parental inbreds together with *per se* and GCA effects resulted in the identification of 231-A as the best parent. The estimates of SCA effects of hybrids revealed that the hybrids Hyd.18227 x Hyd.08R-740-2 (54.178), Hyd.19051 x Hyd.12288 (29.678), Hyd.18227 x Hyd.18169 (26.878) and Hyd.18063 x Hyd.18169 (19.578) registered significant and favorable SCA effects for majority of traits including grain yield per plant.

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