



## Combining Ability and Heterosis for Yield and Quantitative Traits in Maize, *Zea mays* (L.)

K. Sumalini\*

Agricultural Research Station, ANGRAU, Karimnagar-505 001.

Combining ability analysis was conducted in late maturing maize (*Zea mays* L.) inbred lines for yield attributes. Both additive and non additive gene effects were present in the material under study. Variance due to *gca* was preponderant for days to 50% pollen shedding, days to 50% silk emergence, plant height, ear height, number of kernel rows, number of kernels per row and grain yield indicating importance of additive genes in controlling the traits. Three lines KML-70, KML-223 and KML-286 and two testers, KML-36 and KML-226 were good general combiners for grain yield. The crosses KML-57 x KML-226, KML-70 x KML-3, KML-70 x KML-29 and KML-286 x KML-29 are desirable with significant positive *sca*, heterosis and high mean values for grain yield. KML-223 x KML-3 recorded significant negative heterosis for days to 75 per cent dry husk, combined with significant negative *sca* and low mean performance. Hence, this hybrid could be suitable for medium maturity situations.

**Key words:** Maize, heterosis, combining ability, gene action, line x tester analysis

Maize (*Zea mays* L.) is the second most important cereal crop in the world and provides more human food than any other cereal. In India, maize is the third most important cereal after rice and wheat growing in an area of 8.26 MHa with the productivity of 2.02 t ha<sup>-1</sup> and production of 16.72 MT. Maize has many possible uses viz., food, feed for livestock and as raw material for industry. The knowledge of gene action and combining ability helps in identifying the best combiners which may be hybridized either to exploit heterosis or to accumulate genes through selection and in understanding the characters to choose the proper selection method to be followed in breeding programme. In the present study, an attempt has been made to estimate the heterosis in F<sub>1</sub> hybrids with respect to yield, combining ability and gene action governing the quantitative traits in maize, using line x tester mating design.

### Materials and Methods

Fifteen diverse late maturity genotypes of maize including seven lines namely, KML-57, KML-70, KML-141, KML-223, KML-234, KML-235 and KML-286 and eight testers namely, KML-3, KML-29, KML-32, KML-35, KML-36, KML-58, KML-99 and KML-226 were involved in crossing in a line x tester mating design in *Rabi*, 2009-10 and the resultant fifty six crosses along with popularly grown hybrid of Monsanto India Pvt Ltd. i.e. 900MGold as check were grown in Randomized Block Design with three replications during *Kharrif*, 2010 at Agricultural Research Station, Karimnagar. The seed material was sown in a two row plot of 4 m length, having an

inter and intra-row spacing of 75 cm and 20 cm, respectively. Two seeds per hill were planted and later on thinning was done at 2-leaf stage to have optimum plant population. Heterosis was assessed over the standard check (standard heterosis). Five plants from each plot were randomly selected for recording observations on plant height, ear height, ear length, ear girth, number of kernel rows and number of kernels per row. Observations on days to 50 per cent pollen shedding, days to 50 per cent silk emergence, days to 75 per cent dry husk, number of ears per plant, grain yield (q/ha) and 100 kernels weight were recorded on plot basis. Number of ears per plant was recorded on plot basis to know barrenness, an undesirable trait. Combining ability analysis was carried out as per the method suggested by Kempthorne (1957). Heterosis was estimated over the check as per the standard procedure.

### 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. The mean squares for hybrids were partitioned into three components viz., due to lines, due to testers and due to line x tester interactions. The differences among hybrids due to the lines, testers and line x tester interaction were significant for all the characters except plant height, ear height, number of ears per plant, number of kernel rows in line x testers thereby suggesting that the

\*1Corresponding author email: sumalinikatragadda@gmail.com

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 for days to 75% dry husk, ear length, ear girth and equal proportion of *sca* variance to *gca* variance for number of ears per plant, 100 kernel weight indicates that the both the additive and non-additive interactions were significantly higher among the hybrids, which would be important for their exploitation. Higher *sca* variance than the *gca* variance exhibiting

preponderance of non-additive gene effects has also been earlier reported by Joshi *et al.* (2002) and Gautam (2003). However, the component of variation due to *gca* was higher than that due to *sca* for days to 50% pollen shedding, days to 50% silk emergence, plant height, ear height, number of ears per plant, number of kernel rows, number of kernels per row, 100 kernels weight and grain yield, revealing the preponderance of additive gene actions for major attributes (Table 1). It infers that the experimental materials may be more efficiently

**Table 1. Analysis of variance for combining ability.**

Source	df	Grain yield	Days to 50% pollen shedding	Days to 50% silk emergence	Days to 75% dry husk	Plant height	Ear height	Ear length	Ear girth	No. of ears/plant	No. of kernel rows	No. of kernels/row	100 kernels weight
Replications	2	183.789	14.024**	16.685**	10.661	994.667**	1164.875**	8.046**	0.662	0.004	1.256	31.327	0.554
Hybrids	55	1011.06**	13.339**	11.395**	9.549**	875.628**	255.975**	6.442**	3.294**	0.015**	3.415**	43.16**	53.066**
Lines	6	4788.548**	38.534**	34.206**	37.496**	3994.927**	1427.913**	26.69**	7.646**	0.064**	13.774**	241.635**	231.387**
Testers	7	1771.208**	53.04**	44.595**	7.067*	2136.149**	428.5**	4.81**	7.334**	0.02**	5.707**	20.395**	104.142**
Lines x testers	42	344.731**	3.123**	2.603*	5.97**	219.927	59.80	3.821**	1.999**	0.007	1.554	18.601**	19.079**
Error	110	48.969	1.612	1.648	2.436	177.273	81.80	1.039	0.419	0.006	1.104	8.546	1.057
<i>gca</i> variance		130.451	1.896	1.635	0.725	126.472	38.60	0.53	0.244	0.002	0.364	4.996	6.608
<i>sca</i> variance		98.587	0.504	0.318	1.178	14.218	-7.33	0.927	0.527	0.000	0.15	3.352	6.007
Additive variance (VA)		521.804	7.584	6.540	2.900	505.888	154.384	2.120	0.976	0.008	1.456	19.984	26.432
Dominance Variance (VD)		394.348	2.016	1.272	4.712	56.872	-29.336	3.708	2.108	0.000	0.600	13.408	24.028
<i>gca</i> variance/ <i>sca</i> variance		1.323	3.762	5.142	0.615	8.895	-5.263	0.572	0.463	0.000	2.427	1.490	1.100

Significant at P<0.05; \*\*Significant at P<0.01

exploited by adopting any population improvement method for composite development to derive new inbred lines. There exists great potential for improving these traits through recurrent selection. These findings were in close agreement with those of Vasal *et al.* (1993) and Singh *et al.* (1998) for days to silking and plant and ear heights.

Estimates of *gca* effects (Table 2) indicated that the lines KML-70, KML-223 and KML-286 were good

general combiners for grain yield. Line KML-223 was a good general combiner for all the yield contributing traits viz., ear girth, number of ears per plant, number of kernel rows and number of kernels per row. Lines KML-70 and KML-286 were found to be good general combiners for plant height, ear height and 100 kernels weight, where as KML-70 was good combiner for number of ears per plant and KML-286 was a good combiner for days to 50% silk emergence and ear length.

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

Line / Testers	Grain yield	Days to 50% pollen shedding	Days to 50% silk emergence	Days to 75% dry husk	Plant height	Ear height	Ear length	Ear girth	No. of ears/plant	No. of kernel rows	No. of kernels/row	100 kernels weight
KML-57	0.882	0.994**	1.268**	1.304**	-7.488**	-1.78	-1.604**	-0.421**	-0.074**	0.321	-4.220**	-0.149
KML-70	18.952**	0.327	-0.065	-0.03	11.804**	6.637**	0.146	-0.408**	0.028*	-1.054**	-2.637**	3.601**
KML-141	-12.743**	-1.506**	-1.274**	-1.988**	2.762	1.637	1.271**	-0.754**	0.053**	-0.095	5.071**	-3.232**
KML-223	9.873**	0.411	0.143	-0.071	-1.571	-1.238	-0.812**	0.617**	0.060**	1.030**	1.988**	-3.065**
KML-234	-10.273**	-1.673**	-1.065**	-0.946**	-16.821**	-8.738**	-0.633**	0.271*	-0.030*	-0.304	-1.304*	-2.774**
KML-235	-18.335**	1.786**	1.851**	1.679**	-9.280**	-8.821**	1.009**	0.738**	-0.047**	0.780**	1.905**	1.935**
KML-286	11.644**	-0.339	-0.857**	0.054	20.595**	12.304**	0.621**	-0.042	0.01	-0.679**	-0.804	3.685**
SE (Lines) ±	1.322	0.240	0.243	0.295	2.516	1.709	0.193	0.122	0.014	0.199	0.552	0.194
KML-3	-3.152*	2.613**	2.321**	0.089	17.125**	1.798	0.08	0.804**	-0.024	-0.565**	0.262	2.339**
KML-29	-1.462	0.423	0.321	0.899**	-3.256	-4.583*	-0.162	0.004	0.021	0.911**	0.643	-0.28
KML-32	1.452	-0.815**	-1.250**	-0.577	-9.399**	-3.917*	0.261	-0.082	-0.002	-0.423	1.119	-0.375
KML-35	-1.205	-2.673**	-2.393**	-0.625	-14.446**	-4.488*	-0.539*	0.466**	0.023	0.339	-0.881	1.815**
KML-36	5.276**	-0.815**	-0.536*	0.280	-1.256	-1.726	0.857**	0.280*	0.007	0.244	0.833	0.958**
KML-58	-11.381**	-0.625*	-0.298	-0.720*	-2.113	6.893**	-0.672**	-0.939**	0.012	-0.613**	0.262	-4.661**
KML-99	-8.229**	0.851**	1.036**	0.28	3.839	0.321	0.009	-0.753**	-0.064	0.149	-1.833**	-1.137**
KML-226	18.700**	1.042**	0.798**	0.375	9.506**	5.702**	0.166	0.218	0.027	-0.042	-0.405	1.339**
SE (Testers) ±	1.428	0.259	0.262	0.319	2.718	1.846	0.208	0.132	0.016	0.215	0.597	0.210

Significant at P<0.05; \*\*Significant at P<0.01

Lines KML-141 and KML-234 were found to be good general combiners for earliness i.e. days to 50% pollen shedding, days to 50% silk emergence and days to 75% dry husk, Line KML-141 for number of ears per plant and number of kernels per row, KML-234 for ear girth and KML-235 for ear length, ear girth, number of kernel rows, number of kernels per row and 100 kernels weight.

Among testers, KML-36 and KML-226 were good general combiners for grain yield. For 100 kernels weight, KML-3, KML-35, KML-36 and KML-226 were good general combiners and KML-3, KML-35 and KML-36 for ear girth. For ear length KML-36 was good general combiner. For earliness i.e. for days to 50% pollen shedding and days to 50% silk emergence, KML-32, KML-35 and KML-36 were good general combiners. For days to 50% pollen

shedding and days to 75% dry husk, KML-58 was good general combiner.

Among hybrids, KML-286 x KML-29 was the best specific combiner for grain yield followed by KML-57 x KML-35, KML-235 x KML-32, KML-235 x KML-58, KML-57 x KML-226 and KML-223 x KML-3 (Table 3). The cross KML-223 x KML-3 is desirable for early maturity with significant negative *sca*, heterosis and low mean values for number of days to 75 per cent dry husk. The cross KML-57 x KML-226 was desirable with significant positive *sca*, heterosis and high mean values for grain yield and ear girth, significant *sca* and heterotic effects for ear length, ear girth and 100 kernels weight. The crosses KML-57 x KML-226, KML-70 x KML-3, KML-70 x KML-29, KML-57 x KML-35, KML-223 x KML-3 and KML-286 x KML-29 are desirable with significant positive *sca*,

**Table 3. Estimates of *sca* effects, standard heterosis and *per se* performance of best performing crosses.**

Character		Cross combinations								SE (m) ±
		KML-286 x KML-29	KML-57 x KML-35	KML-235 x KML-32	KML-235 x KML-58	KML-57 x KML-226	KML-223 x KML-3	KML-70 x KML-3	KML-70 x KML-29	
Grain yield (q/ha)	<i>Sca</i>	16.870**	16.676**	16.035**	14.302**	13.871**	13.865**	12.852**	9.729**	3.499
	het.	32.757**	17.709**	-6.480	-26.966**	41.757**	23.662**	35.007**	32.992**	5.714
	<i>per se</i>	94.400	83.700	66.50	51.933	100.80	87.933	96.00	94.567	-
Days to 50 per cent pollen shedding	<i>Sca</i>	-0.423	-0.327	0.024	-0.500	-1.042	-2.030**	0.720	-0.089	0.635
	het.	-5.78**	-8.67**	-3.47**	-4.05**	-3.47**	-3.47**	1.16	-4.05**	1.037
	<i>per se</i>	54.333	52.667	55.667	55.333**	55.667	55.667	58.333	55.333	-
Days to 50 per cent silk emergence	<i>Sca</i>	-0.571	0.018	0.292	-0.661	-1.173	-1.571*	0.637	0.304	0.642
	het.	-7.73**	-7.73**	-4.42**	-4.42**	-4.42**	-4.42**	-1.10	-4.97**	1.048
	<i>per se</i>	55.667	55.667	57.667*	57.667*	57.667	57.667	59.667	57.333	-
Days to 75 per cent dry husk	<i>Sca</i>	-0.482	1.125	-0.631	-0.488	-0.542	-5.548**	0.411	-0.399	0.780
	het.	-0.71	2.13	0.71	0.71	1.42	-5.67**	0.71	0.71	1.274
	<i>per se</i>	94.667	96.000	94.667	94.667	95.333	88.667	94.667	94.667	-
Plant height	<i>Sca</i>	-6.786	0.821	-0.768	-7.054	6.869	13.000	-6.042	-2.661	6.657
	het.	8.78	-14.39	-13.17	-12.44	7.56	21.95	17.80	5.37	10.871
	<i>per se</i>	148.667	117.000	118.667	119.667	147.000	166.667	161.000	144.000	-
Ear height	<i>Sca</i>	-4.875	3.113	-1.083	1.107	7.589	3.619	0.411	-1.875	4.522
	het.	8.52	-1.70	-19.886*	2.27	23.295**	10.795	18.715*	3.98	7.385
	<i>per se</i>	63.667	57.667	47.000	60.000	72.333	65.000	69.667	61.000	-
Ear length	<i>sca</i>	-0.917	-0.382	-0.228	0.205	1.646**	0.14	-0.151	-1.575**	0.510
	het.	4.615*	-8.105**	13.846**	10.769**	8.720**	3.797**	7.895**	-2.357**	0.832
	<i>per se</i>	17.000	14.933	18.500**	18.100*	17.667	16.867	17.533	15.867	-
Ear girth	<i>sca</i>	-0.258	-0.641	0.115	0.972**	1.64**	-1.317**	-0.492	-0.525	0.324
	het.	-2.733**	-4.624**	3.992**	3.992**	8.195**	-0.212	-1.472**	-6.723**	0.529
	<i>per se</i>	15.433	15.133	16.500	16.500	17.167	15.833	15.633	14.800	-
No. of ears/plant	<i>sca</i>	-0.003	0.002	0.04	-0.047	0.036	0.019	0.031	0.002	0.038
	het.	3.236**	-4.793**	-0.622	-8.234**	-0.935	6.052**	3.966**	5.635**	0.063
	<i>per se</i>	0.990	0.913	0.953	0.880	0.950	1.017	0.997	1.013	-
No. of kernel rows	<i>sca</i>	-0.369	0.202	-0.827	0.030	0.583	-0.268	0.149	-0.994	0.525
	het.	2.379**	9.521**	0.000	4.764**	9.521**	4.764**	-7.143**	-4.764**	0.858
	<i>per se</i>	14.333	15.333	14.000	14.667	15.333	14.667	13.000	13.333	-
No. of kernels/row	<i>sca</i>	0.232	-2.494	0.381	-2.429	2.363	0.155	0.780	-1.935	1.462
	het.	-3.001	-25.999**	7.001**	-4.000	-10.000**	4.001	-7.999**	-15.001**	2.387
	<i>per se</i>	32.333	24.667	35.667	32.000	30.000	34.667	30.667	28.333	-
100 kernel weight	<i>sca</i>	-1.304*	1.101*	-0.125	0.161	1.577**	-3.173**	-2.506**	4.113**	0.514
	het.	1.922*	3.846**	0.001	-11.538**	3.846**	-15.386**	5.770**	17.309**	0.839
	<i>per se</i>	35.333	36.000	34.667	30.667	36.000	29.333	36.667	40.667	-

Significant at P<0.05; \*\*Significant at P<0.01

heterosis and high mean values for grain yield. These results are in similarity with the earlier findings of Lata *et al.* (2008).

In the estimation of heterosis, Sharma (1994) opined that heterosis over the best check or the local variety could be considered as the better criteria for evaluation of hybrids. The present study revealed that the distribution of heterosis in negative direction for days to 50% pollen shedding, days to 50% silk emergence and in both positive and negative directions for the remaining traits.

The hybrid combination KML-223 x KML-3 recorded significant negative heterosis for days to 75 per cent dry husk, combined with significant negative *sca* and low mean performance. Hence, this hybrid could be suitable for medium maturity situations.

For ear length, hybrids KML-70 x KML-36, KML-141 x KML-3, KML-235 x KML-29, KML-235 x KML-99 and KML-286 x KML-32 recorded significant standard heterosis combined with good *sca* and mean performance. For ear girth, KML-57 x KML-3 and KML-223 x KML-35 recorded significant standard heterosis with positive *sca* effects and good *per se* performance. KML-70 x KML-58 recorded significant *sca*, heterosis for number of ears per plant, but failed to express *per se* performance.

The crosses KML-223 x KML-35 and KML-223 x KML-36 recorded significant *sca*, heterosis and high mean values for number of kernel rows.

For number of kernels per row, crosses KML-223 x KML-36, KML-235 x KML-35 and KML-235 x KML-99 recorded significant *sca* effects coupled with heterosis but failed in *per se* performance.

For 100 kernels weight the crosses KML-57 x KML-3, KML-70 x KML-32, KML-70 x KML-35 and KML-70 x KML-36 recorded significant *sca* with heterosis and good *per se* performance.

Based on *per se* performance, KML-29 and KML-226 lines involving as one of the parents in cross combinations *i.e.*, KML-57 x KML-226, KML-223 x KML-226, KML-70 x KML-29 and KML-286 x KML-29

recorded high grain yields. These results are in similarity with the earlier findings of Srivastava and Singh (2003) and Jyoti *et al.* (2006). The lines KML-70, KML-223 and KML-286 and the testers KML-36 and KML-226 had significant *gca* effect for grain yield and its contributing characters and thus could be used as parents for the development of high yielding single cross hybrids. The high yielding hybrids KML-57 x KML-226, KML-223 x KML-226 and KML-70 x KML-3 with medium maturity could be tested for its performance over the locations for exploitation on commercial scale. The hybrids KML-57 x KML-35, KML-223 x KML-3 and KML-286 x KML-29 showed high *sca* effect, *per se* performance for grain yield and were identified for further multilocation evaluation.

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