

Genetic analysis for the improvement of grain yield, yield contributing traits and quality in Rabi sorghum (*Sorghum bicolor* (L.) Moench). Line x Tester Analysis In Rabi Sorghum For Yield, Yield Components And Quality Traits

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ABSTRACT:

An experiment study was conducted to assess the estimates of heterosis and combining ability of thirty F₁ hybrids of rabi sorghum (*Sorghum bicolor* (L.) Moench) developed by crossing three lines and ten testers in line x tester fashion with respect to quality traits, grain yield and its components. The hybrids and their parents were evaluated using thirty F₁ hybrids of rabi sorghum (*Sorghum bicolor* (L.) Moench) developed by crossing three lines and ten testers in line x tester fashion. for their combining ability and also to check the superiority of hybrids (heterosis) over parents with respect to quality traits, grain yield and its components. to estimate the heterosis and combining ability governing quantitative traits. Among thirty hybrids, two potential hybrids M-35-2A x RSV 2388 and 104A x Phule Vasudha were found to be good specific combiners with highest standard heterosis and *per se* performance for grain yield over checks CSH 15R. AKMS-66-2A and 104A were the good general combiners with males viz. Phule Vasudha, RSV 2388, Phule Revati and RSV 1910. The estimates of general combining ability and specific combining ability revealed the presence of both additive and non-additive genetic components for most of the traits under study.

KEY WORDS: Rabi Sorghum, L x T, Heterosis, Combining ability.

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INTRODUCTION:

Sorghum (*Sorghum bicolor* (L.) Moench) plays a vital role in providing nutrition to human race along with wheat, rice and maize. It is the source of food, fodder and predominantly cultivated in states of Maharashtra, Karnataka and Tamil Nadu. Rabi sorghum is cultivated over 4.24 m ha in India with an annual production of 4.78 mt. In Maharashtra, rabi sorghum are being cultivated over 16.6 lakh ha with an annual production of 20.36 lakh ton (Anonymous, 2022). The traits like grain yield and its components contributing traits are governed by polygenes with complex gene action and hence understanding the nature and magnitude of gene action help the breeder in selection of an appropriate breeding method. For improvement in such an important crop, the most important prerequisite is the selection of suitable parents, which could combine well and produce desirable hybrids. Combining

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ability analysis helps in identifying the better parents, and these parents can be used for hybridization program in order to produce superior hybrids (Ingle *et al.*, 2018). As a general rule, general combining ability (GCA) is the result of additive gene effects, while the specific combining ability (SCA) is the result of non-allelic interactions (Jinks, 1954; Ingle *et al.*, 2018). In the present study, an attempt has been made to estimate the heterosis in F_1 hybrids with respect yield, (what about quality traits?) the combining ability and gene action governing the quantitative and qualitative traits in sorghum, ~~using line x tester mating designs.~~

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MATERIAL AND METHODS:

Thirteen genotypes of sorghum ~~were studied in this experiment. Thirteen genotypes~~ consist of three lines namely 104A, AKMS-662 A and M-35-2A and ten testers namely, RSV 2388, RSV 2371, RSV 1910, RSV 1988, RSV 2394, RSV 2482, RSV 2595, Phule Vasudha, Phule Suchitra and Phule Revati were used as parental lines. These parents were crossed in a line ~~x~~ tester mating design and developed 30 hybrids. resultant thirty All the hybrids along with their parents and standard check CSH 15R were tested during *rabi* 2022-23. Heterosis was assessed over the better parent (heterobeltosis) and standard check (standard heterosis). Estimation of these two types heterosis were done for following characters *viz.* days to 50 per cent flowering, days to maturity, plant height, 1000 seed weight, grain yield per plot, dry fodder yield per plot, total sugar %, crude protein % and crude fiber %. The heterosis was estimated out as per the method suggested by Singh and Narayanan (1993). ~~Thirty~~ The hybrids and ~~thirteen~~ parents were also evaluated to study the combining ability and gene action for nine characters. The combining ability analysis was carried out as per the method suggested by Kempthorne (1957).

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RESULTS AND DISCUSSION:

The analysis of variance for quality, yield and yield components traits studied are presented in Table 1. Variance due to parents was highly significant for all the traits ~~studied~~—indicating good amount of genetic differences among the parents. Variance due to hybrids was also significant for all the nine traits. Variance due to lines was significant for all traits except days to 50% flowering and crude protein. Variance due to tester was significant for all traits except days to 50% flowering. Variance due to line ~~x~~ testers ~~tester~~ was significant for days to 50% flowering, days to maturity, plant height and total sugar content and remaining are found non-significant. The relative estimation of variance due to general combining ability

and specific combining ability revealed the presence of both additive and non-additive genetic components for most of the traits (Ingle, *et al* 2018). The predominance of non-additive gene action were recorded in traits *viz.* Thousand grain weight, grain yield total sugar and crude protein. These findings were confirmed by Chaudhary and Narkhede (2004) [\(Also try to mention recent references for confirmation your findings\)](#). Whereas predominance of additive gene action were shown by the traits *viz.* days to 50 % flowering, days to maturity, plant height, dry fodder yield and crude protein. Similar findings were reported by Jain and Patel (2014) and Aru *et al* (2019).

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The analysis of variance for combining ability for quality, yield and yield components traits studied are presented in Table2. Harer and Bapat (1982) stated that the *per se* performance of the parents with the nature of combining ability provide the criteria for the choice of parents-. On the basis-, those parents which performed well for *per se* and *gca* effects can be consider as good combiner.

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Among the lines, AKMS--66 2A recorded low mean value and negative *gca* effects for days to 50% flowering [\(Please mention the mean values and *gca* values in parentheses after every trait\)](#) and maturity coupled with high mean value -as well as positive *gca* effects for grain yield, total sugar and crude protein content which might be useful earliness, high productive with more nutritional properties. The line 104A shows high mean value for plant height and dry ~~fodder~~fodder yield with desirable *gca* effects indicating, it is suitable for more fodder yield. Among the testers, RSV 2388, RSV 1910, RSV 2371, Phule Vasudha and Phule Revati were found to be good combiner for quality and yield components traits. A perusal of mean performance and general combining abilities of parent exhibited relationship amongst them, this finding corroborates with that of Nandanwankar (2010).

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Among the hybrids M-35-2A(low) ~~×~~ RSV 2388(low) and 104A (low) ~~×~~ Phule Vasudha (high) recorded maximum *sca* effects for grain yield. The hybrids M-35-2A(low) ~~×~~ Phule Vasudha (high) and 104A(high) ~~×~~ RSV 2371(low) showed significant *sca* effects for dry fodder yield. The hybrid M-35-2A (high) ~~×~~ RSV 2371(high) earliness for flowering with more *sca* effects. The hybrid M-35-2A (low) ~~×~~ RSV 2394 (high) matures early with more *sca* effects. The hybrid 104A (average) ~~×~~ RSV 2595 (high) recorded more plant height with maximum *sca* effects. Similar results were also observed by —Chaudhary and Narkhede (2004) and Takalkar (2010).

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The hybrid 104A (average) ~~xx~~ Phule Revati (high) recorded more thousand grain with maximum *sca* effects. ~~_AKM66-2A(high) ~~xx~~ RSV 2388(average)~~ recorded high total sugar and crude fibre content with more *sca* effects. The hybrid 104A (low) x RSV 1988 (average) recorded more crude ~~proten~~protein content with high *sca* effects. The derivatives of low ~~xx~~ low, low ~~xx~~ high, low ~~xx~~ average, average ~~xx~~ low and high ~~xx~~ low parental combinations shows non additive gene action. The derivatives of high ~~xx~~ high parental combinations show additive gene action. In case of non-additive gen action later generation selection is suitable whereas for additive gene action desirable segregants can be selected in early generations.

In the estimation of heterosis, Sharma (1994) opinioned that heterosis over the best check could be considered as better criteria for evaluation of hybrids. The present study revealed the distribution of heterosis in both positive and negative directions for all the traits. The ~~potentially?~~ of hybrid might be judged by comparing the per se performance and heterotic vigour. Close association per se performance of hybrids and heterosis was observed in most of the traits.

Sorghum hybrids, such a 104A ~~xx~~ RSV 1910 —and 104A ~~xx~~ RSV 2371 ~~are-were~~ recorded the —significant negative standard heterosis over the check CSH-15-R were recorded for flowering and maturity. Similar results also reported by Umakanth *et al.* (2006) and Madhusudhan *et al.*, (2013). Totre and *et al* (2021). The hybrids M-35-2A ~~xx~~ P. Suchitra and M-35-2A ~~xx~~ RSV 1988- recorded the maximum significant standard heterosis in the positive direction for plant height. These results confirmed by Umakanth *et al.* (2006) and Nirde (2014). Among the hybrids 104A ~~xx~~ P. Revati (19.36 %) recorded significantly superior standard heterosis for 1000 seed weight. —A positive heterosis for this trait was also reported by Prabhakar *et al.* (2013). The hybrid 104A ~~xx~~ P. Vasudha showed the highest significant standard heterosis for grain yield. —Similar results were obtained by Umakanth *et al.* (2006) and Kulkarni and Patil (2004). The hybrids M-35-2A ~~xx~~ P. Vasudha M-35-2A ~~xx~~ RSV 1910 and M-35-2A ~~xx~~ RSV 1910 displayed the maximum standard heterosis in the positive direction compared to the check CSH-15R. Similar results obtained by Chavan *et al.*, (2016), Prakash *et al.* (2010) and Ghorade *et al.* (2014). In case of quality trait, the hybrid 104A ~~xx~~ P. Vasudha showed the highest significant standard heterosis for total sugar and crude protein. These results confirms the earlier findings of Chavan *et al.* (2016)-. [\(Please write some discussion in connection with your findings in brief way\).](#)

[You should include the tables for gca, sca and heterosis for all the parents and hybrids\).](#)

CONCLUSION:

The present study revealed that hybrids exhibited heterosis for grain yield were not heterotic for all the traits. The - general combining ability of the female 104B and AKMS-66-2B were the best-, while the male parent P. Vasudha and RSV 2388 were the best general combiners and could be exploited for future breeding programme of *rabi* sorghum. The results indicate that exploitation of the heterosis or hybrid vigour might be one of the promising methods to effect crop improvement in sorghum for grain purpose. These results also indicated that the heterosis for grain yield can be exploited commercially. [\(What about the potential hybrids you developed? Conclude about those best hybrids for further breeding or someting\)](#)

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UNDER PEER REVIEW

Table 1. Analysis of variance for parents and hybrids for yield, yield components and quality traits in sorghum

Sources of variation	d.f.	Mean sum of squares								
		Days to 50% flowering	Days to maturity	Plant height	1000 seed weight	Grain yield	Dry fodder yield	Total sugar	Crude protein	Crude fiber
		1	2	3	4	5	6	7	8	9
Parents	12	3.250**	5.371**	625.153**	15.068**	0.145**	0.477**	0.559**	0.531**	0.286**
Line	2	0.166	3.166**	2076.500**	11.727**	0.018	0.354**	0.488**	0.181	0.329**
Testers	9	1.383	4.827**	73.005**	17.473**	0.184**	0.557**	0.361**	0.667**	0.308**
Line vs Tester	1	26.216**	14.678**	2691.796**	0.102	0.043	0.001	2.478**	0.001	0.001
Parents vs Crosses	1	14.693**	2.737	6498.096**	47.780**	0.362**	0.309**	0.026	0.171	0.041
Crosses	29	4.048**	4.867**	1434.315**	20.500**	0.048**	0.233**	0.205**	0.319**	0.193**
Error	42	0.821	1.403	10.121	3.920	0.014	0.006	0.047	0.088	0.017

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

Table2: Analysis of variance for combing ability for yield, yield components and quality traits in sorghum

Sources of variation	d.f.	MSS								
		Days to 50 % flowering	Days to maturity	Plant height (cm)	1000 seed weight (g)	Grain yield (kg/ plot)	Dry fodder yield (Kg/ plot)	Total sugar content (%)	Crude protein content (%)	Crude fibre content (%)
Line	2	25.550**	24.050**	19861.40* *	10.55**	0.025*	1.778**	0.385**	0.740**	0.171**
Tester	9	2.489**	2.906**	103.24**	21.65**	0.059**	0.092**	0.144**	0.419**	0.093**
Line \times Tester	18	2.439**	3.717*	52.40**	21.03**	0.047**	0.133**	0.217**	0.222**	0.246**
Error	29	0.667	1.419	8.59	4.50	0.016	0.006	0.031	0.080	0.015
<i>gca</i>		2.03	1.86	1534.18	1.87	0.004	0.14	0.03	0.08	0.02
<i>sca</i>		0.81	1.15	21.14	8.55	0.016	0.06	0.09	0.07	0.11
<i>gca/sca</i>		2.51	1.61	72.57	0.22	0.256	2.26	0.39	1.12	0.15

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

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