# Combining ability and heterosis for yield and its related traits in rice hybrids

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#### ABSTRACT

Combining ability analysis and heterosis for yield and its attributing traits was carried out in rice through line x tester analysis of 30 hybrids developed by crossing three females with ten male lines during wet season 2010 at Mandya, Karnatak. The hybrids along with parents and a standard check KRH-2 grown in a RCBD with two replications were evaluated for grain yield and yield contributing traits during dry season 2011. The estimates of gca effects indicated that, among females, KCMS 49A and among males Thanu are good general combiners for grain yield and most of the traits studied. High sca effects were observed in the crosses, KCMS 47A × KMR 4, KCMS 48A × MSN 75 and KCMS 47A × MSN 98 and they were found to be the best combinations for grain yield and its traits. The crosses KCMS 47A × KMR 4 and KCMS 49A × MSN 93 exhibited high mean seed yield and high standard heterosis over standard check KRH-2.

Key words: hybrid rice, yield, CMS, combining ability, heterosis

Success of any plant breeding programme depends on the choice of appropriate genotypes as parents in the hybridization programme. Among different methods available, combining ability analysis is one of the effective approaches available for estimating the combining ability effects that help in selecting desirable parents and crosses for the exploitation of heterosis. To exploit maximum heterosis using male sterility system in hybrid breeding programme, knowledge on the combining ability of different male sterile and restorer lines is essential. From a practical point of view, standard heterosis is the most important of the two levels of heterosis because it is aimed at developing desirable hybrids superior to the existing high yielding commercial varieties. Significant heterosis and standard heterosis have been reported in rice by a number of workers such as Umakantha et al., (2002), Nadali (2010) and Tiwary et al., (2011) etc. Accordingly, the present investigation was undertaken to study combining ability for yield and its traits with a view to identify good combiners and also to identify high yielding non aromatic rice hybrids.

### **MATERIAL AND METHODS**

The experimental material for this study comprised of three newly developed non-aromatic CMS lines KCMS 47A, KCMS 48A and KCMS 49A having WA type of cytoplasm} and ten testers (Thanu, KMR-3, KMR-4, KMR-12, MSN-36, MSN-75, MSN-91, MSN-93, MSN-98 and MSN-99) by validating SSR tightly linked to fertility restoration. These were crossed in 3 line  $\times$  10 tester fashion during wet season 2010 to produce thirty F<sub>1</sub> hybrids by adopting clipping method of crossing. These F, hybrids, parents along with standard check variety KRH-2 were evaluated during 2011 in RCBD with two replications. The hybrids and their parents and checks were transplanted one seedling hill<sup>-1</sup> with a spacing of 15 cm  $\times$  15 cm. All the recommended package of practices were followed to ensure good crop growth and development. Five competitive plants were randomly selected to record the observations on grain yield and yield contributing characters viz., days to 50% flowering, plant height,

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No. of tillers, plant<sup>-1</sup>, No. of Panicles plant<sup>-1</sup>, Panicle length (cm), Yield plant<sup>-1</sup>, 1000 grain weight and L/B ratio. The mean values of these five plants were used for combining ability analysis as suggested by Kemtphorne (1957). The variances for general combining ability (gca) and specific combining ability (sca) were tested against their respective error variances derived from ANOVA reduced to mean level. Significance test for gca and sca effects were performed using t-test. The aroma of the new hybrids was tested using standard KOH test using IR58025A (highly aromatic) as check. The per cent increase or decrease of  $F_1$  hybrids over mid parent and standard check was calculated to estimate heterotic effects of yield and its related traits.

The overall gca status of parents was calculated based on the methods of Arunachalam and Bandyopadyay (1979) with slight modification as suggested by Mohan Rao (2001).

## **RESULTS AND DISCUSSION**

Analysis of variance revealed presence of significant difference among the genotypes studied. Mean squares of parents and crosses were significantly different for all the traits indicated that they are suitable for genetic studies. The mean squares due to females (lines) and males (testers) were significant most of the traits studied. The variance due to hybrids differed significantly for all the characters. Mean sum of squares due to lines x testers interaction were highly significant for all the characters indicating that the lines differed significantly from testers in respect of all the characters studied. The variance due to hybrids differed significantly for all the characters except no. of tillers plant<sup>-1</sup>. Thus, suggesting the importance of heterosis breeding for improvement of these traits. The analysis of variance for combining ability revealed that the estimates of sca variances were predominant for all the characters studied as revealed by the ratio of gca and sca variances. This indicates predominance of nonadditive gene action in respect of all the traits studied. These results are in agreement with earlier findings of Jagadeesan and Ganesan (2006), Rahimi et al. (2010), Saravanan et al. (2006), Anandkumar et al. (2004).

Among the CMS lines, KCMS 49A was the best general combiner as it showed highly significant *gca* effects for all the traits in desirable (positive)

direction except for days to 50 % flowering, Plant height and no. of tillers plant<sup>-1</sup>. Among the testers, Thanu was found superior general combiner for no. of panicles plant<sup>-1</sup>, and yield plant<sup>-1</sup> (Table 1). For Panicle length KMR-4 was found to be superior. Testers KMR-12 and MSN-36 were found good general combiners for spikelet fertility and no. of spikelets panicle<sup>-1</sup>, respectively. The tester MSN-91 was good general combiner for L/B ratio. It was evident from the results that the lines KCMS 48A and KCMS 49A had high (H) overall *gca* status. Among the testers, Thanu, KMR-3, KMR-4, KMR-12, MSN-36, MSN-93 and MSN-98 posses high (H) overall *gca* status (Table 2). Similar results were reported by Swamy *et al.* (2003) and Saidaiah *et al.*, (2010).

The estimate of *sca* effects with their respective standard error for each character in thirty cross combinations were presented in Table 2. None of the crosses exhibited high sca effects for all the characters studied. The majority of the crosses showed significant sca effects, which involved atleast one parent having high gca effects. Only six out of forty crosses, showed positively significant sca effects for grain yield plant<sup>-1</sup> of which highest being KCMS  $49A \times MSN$  75. This hybrid also had highly significant sca values for panicle length, spikelet fertility and yield hectare<sup>-1</sup> in positive direction. The hybrid KCMS 48A × MSN 75 exhibited highest positive significant sca for spikelet fertility and 1000 grain weight. For yield plant<sup>-1</sup>, the cross KCMS  $47A \times KMR 4$  was found superior. Another hybrid KCMS 47A × MSN 98 was found superior for earliness as it exhibited lowest negative sca for days to 50% flowering. This cross also exhibited superior L/B ratio. For panicle weight, hybrid KCMS 48A×MSN 36 was found superior. The hybrids KCMS  $49A \times KMR$  12 and KCMS 49A × MSN 75 were found good specific combiners for no. of spikelets panicle<sup>-1</sup> and plant height respectively. These results are in corroboration with the earlier findings of Jagadeeasan and Ganesan (2006) and Saidaiah et al. (2010). Seventeen out of thirty hybrids studied exhibited high (H) overall gca status and thirteen crosses exhibited low (L) overall gca status as estimated by the method suggested by Arunachalam and Bandyopadyay (1979) with slight modification as suggested by Mohan Rao (2001). All the twenty crosses with high overall sca effects have parents with all types of combination of gca effect viz., H x H, H x L and L x L. These results are in agreement with the earlier

	Days to 50% flowering	Plant height (cm)	No. of tillers plant <sup>-1</sup>	No. of panicles plant <sup>-1</sup>	Panicle length (cm)	Spikelet fertility (%)	No. of spikelets panicle <sup>-1</sup>	Yield plant <sup>-1</sup> (g)	1000 grain weight (g	L/B ratio )	overall gca status
Lines											
KCMS 47A	-1.48	0.03	0.37	-0.44 **	1.49	1.70 **	-6.77	-3.15 **	-0.65 **	-0.26 **	L
KCMS 48A	0.46	0.37	0.09	-0.27 **	1.36	1.15 **	-14.03 **	-1.74 *	-0.06	-0.01	Н
KCMS 49A	1.02	-0.40	-0.46	0.71 **	-2.85 *	-2.85 **	20.80 **	4.89 **	0.71 **	0.27 **	Н
SEm±	1.084	0.5241	0.6101	0.0489	1.2294	0.0824	4.8169	0.6912	0.0475	0.0358	
Testers											
Thanu	5.64 **	-0.05	0.04	0.69 **	6.51 **	1.97 **	43.89 **	5.72 **	-0.85 **	0.08	Н
KMR-3	-2.69	2.06 *	1.76	0.03	9.19 **	-4.53 **	-36.02 **	3.56 **	2.90 **	-0.08	Н
KMR-4	-1.08	0.34	0.93	0.02	10.90 **	0.63 **	-29.93 **	3.33 *	0.77 **	0.13 *	Н
KMR-12	7.92 **	0.67	0.98	0.32 **	-2.06	4.97 **	21.85 *	-2.28	-0.17	-0.05	Н
MSN-36	5.26 *	-0.61	0.37	0.37 **	1.50	0.47 **	45.39 **	5.06 **	-0.82 **	-0.13	Н
MSN-75	-8.69 **	-1.77	-1.41	-1.30 **	-45.68 **	1.13 **	-37.82 **	-18.74 **	-0.33 **	-0.49 **	L
MSN-91	6.64 **	-0.39	-0.63	-0.04	10.60 **	4.80 **	-3.11	0.40	-1.34 **	0.34 **	L
MSN-93	4.26 *	-1.33	-1.30	0.24 *	2.10	0.97 **	14.79	4.72 **	0.49 **	0.20 **	Н
MSN-98	-7.97 **	1.45	-0.24	0.24 *	4.08	-3.53 **	-1.65	1.21	0.93 **	-0.11	Н
MSN-99	-9.30 **	-0.36	-0.52	-0.58 **	2.87	-6.87 **	-17.38	-2.99 *	-1.57 **	0.10	L
SEm±	1.9792	0.9568	1.1139	0.0892	2.2446	0.1504	8.7944	1.262	0.0868	0.0653	

Table.1 Estimates of general combining ability effects of lines and testers for yield and yield contributing character

\*Significance at 0.05, \*\* Significance at 0.01

reports of Mohan Rao (2001), Saidaiah *et al.* (2010). The proportional contribution of lines, testers and their interactions to total variances showed that testers played an important role toward difficult traits, indicating influence of testers on these traits. The smaller contribution of interactions of the line x tester than testers, indicating higher estimates of variances due to general combining ability. Nadali (2010) observed higher estimates of gca variances due to testers in rice. Contribution of interactions of line x tester was higher than lines for all the traits except for spikelet fertility and L/B ratio indicating higher estimates of gca variances for available to tester than testers.

In the present study among the lines KCMS 49A and among the testers Thanu was the good general combiner for yield and its majority of the traits since they had high positive *gca* effects for yield and majority of yield attributing characters. These best combiners could be utilized in hybrid development breeding programme. The crosses KCMS 47A × KMR 4, KCMS 48A × MSN 75 and KCMS 47A × MSN 98 were identified as most promising for yield based on *sca* effects. Hence these could be used for the exploitation of heterosis for yield and related characters.

Heterosis was computed as increase or decrease in F<sub>1</sub> value over mid parent and over best commercial variety (standard heterosis). The relative magnitude of heterosis was expressed as heterosis over mid parent and standard checks for yield and other characters (Table. 3). Significant heterosis for days to 50 % flowering in negative direction over mid parent (-15.99%) and over the check KRH-2 (-5.38%) was recorded by the hybrid KCMS  $49A \times MSN$  99 (-16.43%). For plant height the cross KCMS 47A × MSN 98 showed significant mid parent heterosis (-15.35%) followed by the hybrid KCMS  $47A \times MSN$  99 (-11.43%) while none of the hybrids showed significant negative heterosis over the check KRH-2. Earlier workers including Umakantha et al. (2002), Nadali (2010), Saidaiah et al., (2010), Tiwary et al., (2011) also noticed significant negative heterosis for early flowering and plant height. With respect to no. of tillers plant<sup>-1</sup> the hybrid KCMS  $48A \times KMR 4$  recorded maximum heterosis over mid parent (30.86%) while none of the crosses showed positive significant heterosis over the check for this trait. For no. of panicles plant<sup>-1</sup> the cross KCMS 48A × KMR 4 showed highest significant mid parent heterosis (44.44%) followed by

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	Days to 50%	Plant height	No. of tillers	No. of panicles	Panicle length	Spikelet fertility	No. of spikelets	Yield plant <sup>-1</sup>	1000 grain	L/B ratio	overall
	flowering	(cm)	plant <sup>-1</sup>	plant <sup>-1</sup>	(cm)	(%)	panicle <sup>-1</sup>	(g)	weight	Tatio	sca status
	nowening	(em)	plant	pium	(em)	(70)	puniere	(5)	(g)		status
KCMS 47A×Thanu	-1.24	0.30	0.63	-0.05	4.99	-0.87 **	-8.43	2.59	0.22	-0.17	Н
KCMS 47A× KMR 3	0.42	0.50	1.07	0.03	1.07	-2.37 **	-3.63	-0.55	0.22	0.09	Н
KCMS $47A \times KMR 4$	4.14	-1.58	-1.93	0.52 **	0.58	1.47 **	21.74		-0.66 **	-0.03	Н
KCMS $47A \times KMR 4$ KCMS $47A \times KMR 12$	5.31	1.75	1.18	-0.16	-2.47	3.13 **	-4.20		-0.00 · -0.49 **	-0.02	L
KCMS $47A \times MSN 36$	-5.02	0.36	-0.21	0.08	-5.37	-0.37	7.37	2.68	-0.18	-0.14	H
KCMS $47A \times MSN 30$ KCMS $47A \times MSN 75$	-0.24	0.03	0.07	-0.44 **		-3.03 **	-21.43	-2.06	0.71 **	-0.02	L
KCMS $47A \times MSN 91$	2.26	0.05	0.63	0.26	-0.37	1.30 **	22.97	3.16	0.40 *	0.19	H
KCMS $47A \times MSN 93$	4.81	-0.25	-0.70	0.02	2.38	-0.87 **	-18.73		• -0.31 *	-0.14	L
KCMS $47A \times MSN 98$ KCMS $47A \times MSN 98$	-9.30 *	-1.20	0.24	-0.26	-4.81	2.63 **	10.90	0.71	0.26	-0.14 0.34 **	L L
KCMS $47A \times MSN 99$	-1.14	-0.89	-0.98	-0.02	3.82	-1.03 **	-6.56	0.06	-0.80 **	-0.11	L
KCMS $48A \times$ Thanu	2.66	-0.71	-0.26	0.26	-3.29	-0.32	9.93	-0.60	-0.24	-0.06	L
KCMS $48A \times KMR 3$	4.65	-1.98	-1.48	0.49 **	-3.51	1.18 **	11.93	3.24	-0.17	0.13	H
KCMS $48A \times KMR 4$	-0.62	2.90	2.86	0.00	-3.16	-2.48 **	2.54	0.89	0.27	-0.22	Н
KCMS 48A × KMR 12	-5.79	-1.93	-2.03	0.01	8.13 *	-1.32 **	-30.54	2.21	-0.04	-0.06	Н
KCMS 48A × MSN 36	-0.29	-0.49	-0.25	0.05	9.78 *	1.18 **	-0.98	-3.04	-0.46 **	-0.15	L
KCMS 48A × MSN 75	-3.68	4.01 *	3.86	-0.68 **	-2.68	5.52 **	19.33	-4.35	1.03 **	0.32 **	Н
KCMS 48A × MSN 91	1.16	-2.87	-2.59	-0.14	1.57	0.85 **	-31.08	-2.90	-0.34 *	-0.05	L
KCMS 48A × MSN 93	-2.29	-1.09	-0.26	-0.24	-4.76	-1.32 **	25.73	2.82	-0.36 *	-0.11	Н
KCMS 48A × MSN 98	1.43	-0.68	-1.48	0.07	3.76	-2.82 **	-20.14	-1.72	-0.25	-0.16	L
KCMS 48A × MSN 99	2.77	2.84	1.63	0.18	-5.86	-0.48	13.29	3.46	0.56 **	0.36 **	Н
KCMS 49A × Thanu	-1.41	0.42	-0.37	-0.21	-1.70	1.18 **	-1.50	-1.99	0.03	0.23	L
KCMS 49A × KMR 3	-5.08	1.46	0.41	-0.52 **	2.44	1.18 **	-8.30	-2.69	-0.69 **	-0.21	L
KCMS 49A $\times$ KMR 4	-3.52	-1.32	-0.93	-0.52 **	2.57	1.02 **	-24.28	-8.03 **	• 0.39 *	0.24 *	L
KCMS 49A × KMR 12	0.48	0.18	0.85	0.15	-5.66	-1.82 **	34.74 *	4.04	0.53 **	0.08	Н
KCMS 49A × MSN 36	5.31	0.13	0.46	-0.13	-4.41	-0.82 **	-6.39	0.36	0.64 **	0.29 *	Н
KCMS 49A × MSN 75	3.92	-4.04 *	-3.93	1.11 **	2.51	-2.48 **	2.10	6.42 **	-1.74 **	-0.30 *	Н
KCMS 49A × MSN 91	-3.41	1.90	1.96	-0.12	-1.21	-2.15 **	8.11	-0.27	-0.06	-0.14	Н
KCMS 49A × MSN 93	-2.52	1.35	0.96	0.21	2.38	2.18 **	-6.99	4.67 *	0.67 **	0.25 *	Н
KCMS 49A × MSN 98	7.87 *	1.88	1.24	0.19	1.05	0.18	9.24	1.01	-0.01	-0.18	Н
KCMS 49A × MSN 99	-1.63	-1.95	-0.65	-0.16	2.03	1.52 **	-6.73	-3.51	0.24	-0.26 *	L
SEm±	3.428	1.6573	1.9293	0.1545	3.8878	0.2604	15.2323	2.1859	0.1503	0.1131	

Table 2. Estimates of specific combining ability effects in crosses for yield and yield contributing characters

\*Significance at 0.05, \*\* Significance at 0.01

KCMS 47A × KMR 3 (37.77%), for this trait also none of the hybrids have recorded positive significant heterosis over KRH-2. Maximum mid parent heterosis for panicle length was exhibited by the hybrid KCMS 49A × MSN 98 (14.26%) followed by the hybrid KCMS 49A × KMR 12 (14.08%) while no positively significant standard heterosis was recorded for this trait also. Joshi (2000), Umakantha *et al.*, (2002), Nadali (2010) and Tiwary *et al.* (2011) reported similar results in their respective studies. For Spikelet fertility out of thirty hybrids studied no one exhibited significant positive mid parent heterosis and standard heterosis over the check.

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The hybrid KCMS  $49A \times KMR$  12 recorded highest heterosis over mid parent (64.20%), over the check KRH-2 (39.86%) for no. of spikelets panicle<sup>-1</sup>.

Seed yield is a complex trait and it is the multiplicative end product of several yield components. Many hybrids showed positive significant heterosis for this trait. For yield plant<sup>-1</sup> the range of mid parent heterosis was recorded from -37.41% (KCMS 48A × MSN 75) to 118.88% (KCMS 47A × KMR 4). However only one hybrid *viz.*, KCMS 49A × MSN 93 manifested significant positive heterosis over the check KRH-2 (16.31%). Joshi (2000), Umakantha *et al.*, (2002),

	Days to 50% flowering		Plant height (cm)		No. of tillers plant <sup>-1</sup>		No. of panicles plant <sup>-1</sup>		Panicle length (cm)	
Crosses	MP Heterosis	SH over KRH-2	MP Heterosis	SH over KRH-2	MP Heterosis	SH over KRH-2	MP Heterosis	SH over KRH-2	MP Heterosis	SH over KRH-2
KCMS 47A × Thanu	-0.50	6.45 **	5.19	10.95 **	11.35	-0.95	29.90 *	6.38	4.79	-10.72 **
KCMS 47A × KMR 3	-9.90 **	-2.15 **	-3.88	4.69	21.89 *	12.52	37.77 **	20.17	2.62	-8.87 **
KCMS 47A× KMR 4	1.01 **	7.53 **	9.36 **	9.70 *	-1.54	-9.58	11.79	-4.28	7.93 **	-5.66 *
KCMS 47A × KMR 12	6.53 **	13.98 **	14.67 **	19.25 **	13.19	11.57	28.99 *	15.92	6.72 *	-5.73 *
KCMS 47A × MSN 36	-3.45 **	5.38 **	-1.51	7.04	-5.22	-3.84	14.77	3.16	6.18 *	-7.58 **
KCMS 47A × MSN 75	-7.02 **	3.23 **	-7.43 *	-1.57	-6.16	-12.46	7.95	-6.41	-11.18 *	* -20.89 *
KCMS 47A × MSN 91	0.48	11.83 **	6.13	15.18 **	-1.86	0.98	6.67	2.11	5.62 *	-5.14
KCMS 47A × MSN 93	0.51	5.38 **	9.27 **	15.34 **	-7.09	-11.54	-0.59	-10.66	7.60 **	-6.84 *
KCMS 47A × MSN 98	-2.02 **	4.30 **	-15.35 **	-9.39 *	8.98	-0.95	15.65	2.11	0.33	-11.09 *
KCMS 47A × MSN 99	-14.08 **	-3.23 **	-11.43 **	-2.98	-10.49	-9.61	-1.14	-7.47	0.87	-12.20 *
KCMS 48A × Thanu	-1.98 **	6.45 **	12.56 **	16.43 **	8.80	-4.79	20.02	-1.08	12.50 **	-5.18
KCMS 48A × KMR 3	-8.29 **	1.08 *	3.37	10.48 *	10.08	0.03	16.36	2.11	9.86 **	-3.44
KCMS 48A × KMR 4	-4.98 **	2.69 **	8.92 *	7.04	30.86 **	18.29	44.44 **	24.44	9.62 **	-5.18
KCMS 48A × KMR 12	0.00	8.60 **	8.52 *	10.64 *	-4.95	-7.67	3.53	-6.41	9.94 **	-3.88
KCMS 48A × MSN 36	-3.88 **	6.45 **	6.24	13.30 **	-6.74	-6.72	11.77	1.05	5.15	-9.43 **
KCMS 48A × MSN 75	-0.72 *	11.83 **	-6.98 *	-2.98	22.53 *	12.52	32.94 **	15.95	1.05	-10.91 *
KCMS 48A × MSN 91	-1.90 **	10.75 **	8.89 **	15.96 **	-20.38 *	-19.22	-17.12	-20.23	3.53	-7.95 **
KCMS 48A × MSN 93	-2.02 **	4.30 **	6.73 *	10.48 *	-8.71	-14.40	0.00	-9.60	11.97 **	
KCMS 48A × MSN 98	-9.45 **	-2.15 **	-2.39	2.50	16.32	4.04	0.59	-10.66	6.22 *	-6.84 *
KCMS 48A × MSN 99	-15.29 **	-3.23 **	-4.65	2.50	14.43	13.91	14.14	7.43	1.93	-12.20 **
KCMS 49A × Thanu	-3.02 **	3.76 **	6.63 *	13.15 **	12.99	-2.74	17.86	-5.36	13.66 **	
KCMS 49A $\times$ KMR 3	-10.89 **	-3.23 **	-7.00 *	1.88	29.03 **	15.41	29.20 *	10.63	5.69 *	-3.11
KCMS 49A $\times$ KMR 5 KCMS 49A $\times$ KMR 4	-4.04 **	2.15 **	3.87	4.85	0.54	-10.56	15.17	-3.22	12.59 **	
KCMS 49A $\times$ KMR 4 KCMS 49A $\times$ KMR 12	-3.02 **	2.15 3.76 **	11.89 **	4.85	4.52	0.03	22.90	-3.22 8.49	14.08 **	
KCMS 49A $\times$ MSN 36	-3.02	0.00	8.95 **	19.09 **	-6.35	-7.67	15.65	2.11	13.05 **	
KCMS $49A \times MSN 30$ KCMS $49A \times MSN 75$	-10.90 **	-1.08 *	-2.12	4.69	-31.93 **	-38.46 **	-26.28 *	-37.27 *		-7.58 **
KCMS 49A × MSN 75 KCMS 49A × MSN 91	-7.25 **	3.23 **	2.79	4.09	3.84	3.87	11.86	5.30	-1.10	-8.32 **
KCMS 49A × MSN 91 KCMS 49A × MSN 93	-1.03 **	3.76 **	4.35	12.21 **	3.11	-4.79	7.21	-5.36	-1.10 12.19 **	
KCMS 49A × MSN 95 KCMS 49A × MSN 98	-9.09 **	-3.23 **	4.33 1.31	9.08 *	29.85 **	-4.79 14.28	18.99	-3.30 3.16	12.19	
KCMS 49A × MSN 98 KCMS 49A × MSN 99	-9.09 **	-5.38 **	-10.23 **		-16.66	-18.23	-2.90	-10.66	7.50 **	-3.33
	Spikelet fertility (%)		No. of spikelets panicle <sup>-1</sup>		Yield per plant (g)		1000 grain weight(g)		L/B ratio	
Crosses	MP	SH over	MP	SH	MP	SH	MP	SH	MP	SH
	Heterosis	KRH-2								
KCMS $47A \times$ Thanu	13.72	4.32	41.60 **	16.94	75.09 **	-4.54	-5.23 **	-21.30 *	*-12.19 *	*-13.53 **
KCMS 47A × KMR 3	12.15	2.90	-4.25	-18.40 *	61.98 **	-16.66 *	4.79 **	-1.96 *	2.30	-11.18 *
KCMS 47A× KMR 4	10.79	4.31	32.33 **	-3.60	118.88 **		-10.98 **	-18.02 *		-8.82 *
KCMS 47A × KMR 12	-2.66	-14.00	30.30 **	8.56	0.21	-43.06 **	-9.58 **	-21.39 *	*-13.90 *	*-12.94 *
KCMS 47A × MSN 36	-4.52	-13.24	16.91 *	25.08 **	40.27 **	-5.85	-6.09 **	-22.89 *	*-14.32 *	*-17.65 *
KCMS 47A × MSN 75	-54.82 **	-60.83 **	-8.34	-27.62 **	-27.95	-71.14 **	1.93	-16.83 *	* -25.88 *	*-23.53 *
KCMS 47A × MSN 91	11.28	2.87	30.15 **	9.60	92.40 **	-15.40 *	-4.48 **			1.18
KCMS 47A × MSN 93	14.59	-3.70	23.14 *	-1.60	35.93 **	-29.87 **	-5.42 **	-17.74 *	*-13.07 *	* -10.00 *
KCMS 47A × MSN 98	4.60	-9.65	16.66	4.61	46.01 **	-19.17 **	-5.07 **	-13.28 *		-5.88
KCMS 47A × MSN 99	11.23	-1.17	-1.09	-11.01	45.62 **	-30.26 **	-19.60 **		*-13.89 *	
KCMS 48A × Thanu	1.03	-5.28	47.29 **	22.16 *	59.36 **	-8.63	0.15	-20.71 *		-5.29
KCMS 48A × KMR 3	4.02	-2.47	-0.08	-14.49	75.45 **	-4.78	7.20 **		15.28 **	
KCMS 48A × KMR 4	3.88	-0.11	14.69	-16.05	83.48 **	-10.67	0.56	-11.28 *		-7.65
KCMS $48A \times KMR 12$	8.42	-2.02	10.85	-7.25	33.29 **	-20.50 **	0.17	-16.79 *		-8.24
KCMS $48A \times MSN 36$	11.90	3.93	9.69	17.74	20.48 *	-15.73 *	0.30	-21.52 *		-12.35 *
KCMS 48A × MSN 75	-59.67 **	-64.22 **		-11.86	-37.41 **	-73.16 **	12.13 **	-12.84 *		-9.65 *
KCMS 48A × MSN 91	11.11	4.95	-4.51	-19.25 *	58.05 **	-26.05 **	-0.51	-23.29 *		1.18
KCMS 48A × MSN 91 KCMS 48A × MSN 93	2.26	-12.00	44.42 **	15.91	78.10 **	-3.09	1.84	-15.36 *		-3.53
	2.20	12.00	TT.T2	15.71	/0.10	5.07	1.07	15.50	5.02	5.55

Table 3. Heterosis (%) of new hybrids for yield and various yield related traits

Table 3....contd...

#### Combining ability and heterosis in rice hybrids

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	Spikelet fertility (%)		No. of spikelets panicle <sup>-1</sup>		Yield per plant (g)		1000 grain weight(g)		g) L/B	L/B ratio	
Crosses	MP Heterosis	SH over KRH-2	MP	SH	MP	SH	MP	SH	MP	SH	
KCMS 48A × MSN 98	13.12	-0.01	-3.82	-13.41	34.90 **	-21.52 **	-0.47	-12.93	** -3.61	-12.00 **	
KCMS 48A × MSN 99	-3.57	-12.38	5.09	-5.08	59.20 **	-19.26 **	-5.72 **	-20.33	** 6.55	5.29	
KCMS 49A × Thanu	-10.32	-8.28	57.70 **	33.18 **	67.02 **	3.36	10.03 **	-16.17	** 22.67 *	* 8.24	
KCMS 49A × KMR 3	-2.71	-0.49	6.08	-7.62	64.61 **	-3.18	12.11 **	-2.80 *	* 22.89 *	* -5.88	
KCMS 49A × KMR 4	-2.92	1.62	17.42	-12.28	57.97 **	-15.91 *	8.61 **	-7.42 *	* 33.14 *	* 9.88 *	
KCMS 49A × KMR 12	-21.73 **	-22.60 **	64.20 **	39.86 **	54.01 **	-1.14	11.20 **	-10.95	**12.10 *	* 1.88	
KCMS 49A × MSN 36	-18.24 **	-17.11 *	20.88 **	31.58 **	43.96 **	7.24	15.19 **	-13.31	** 21.92 *	* 4.71	
KCMS 49A × MSN 75	-62.05 **	-63.10 **	19.30	-3.58	40.54 **	-33.36 **	4.79 **	-21.68	** -11.39 *	** -17.65 **	
KCMS 49A × MSN 91	-5.85	-3.05	34.28 **	15.58	85.32 **	-4.86	9.67 **	-18.70	**7.53 *	5.88	
KCMS 49A × MSN 93	-3.41	-8.66	42.96 **	16.89	97.26 **	16.31 *	15.53 **	-7.42 *	* 19.55 *	* 11.53 **	
KCMS 49A × MSN 98	-4.98	-7.92	27.61 **	16.80	59.28 **	-0.09	8.32 **	-8.48 *	* 10.57 *	-5.88	
KCMS 49A $\times$ MSN 99	-7.60	-8.17	10.95	1.88	44.71 **	-20.03 **	0.08	-18.42	** 5.14	-2.59	

\*Significance at 0.05, \*\* Significance at 0.01

Nadali (2010) and Tiwary *et al.* (2011) in their respective studies have reported different levels of heterosis for seed yield plant<sup>-1</sup>. The hybrid KCMS 49A × MSN 36 showed highest mid parent heterosis (15.19%) for 1000 grain weight. No hybrids out of thirty hybrids showed positively significant heterosis over the check KRH-2. Joshi (2000) and Tiwary *et al.*, (2011) reported significant heterosis for test weight. For L/B ratio the cross KCMS 49A × KMR 4 showed highest mid parent heterosis of 33.14%. Another hybrid KCMS 49A × MSN 93 recorded significant heterosis (11.53%) over the check. All the thirty hybrids studied were non aromatic compared to check KRH-2 which is slightly aromatic hybrid.

The present study resulted in identification of promising non- aromatic rice hybrids viz., KCMS 47A× KMR 4 and KCMS 49A × MSN 93 based on high mean seed yield and high heterosis over standard check KRH-2. Hence these hybrids could be considered for commercial cultivation.

### REFERENCES

- Anandkumar Singh NK and Chaudhary VK 2004. Line x tester analysis for grain yield and related characters in rice. *Madras Agricultural Journal*, 91 (4-6): 211-214.
- Jagadeesan S and Ganesan J 2006. Combining ability in rice (*Oryza sativa* L.). *Indian J. Agric. Res.*, 40 (2): 139– 142.
- Kempthorne O 1957. An Introduction to Genetic Statistics.

John Wiley and Sons, New York, pp.397-515.

- Mohan Rao A 2001. Heterosis is a function of genetic divergence in sunflower (*Helianthus annuus* L.).
  *Ph.D Thesis*, Acharya N. G. Ranga Agricultural University, Hyderabad, 208pp.
- Nadali BJ 2010. Heterosis and combining ability analysis for yield and related traits in hybrid rice. *Int. J. Biol.*, 2(2): 222-231.
- Rahimi, M., Rabiei, B., Samizadeh, H. and Kafighasemi, A., (2010), Combining ability and heterosis in rice (*Oryza sativa* L.) cultivars. J. Agri. Sci. Tech., 12: 223-231.
- Saidaiah P, Sudheer Kumar S and Ramesha MS 2010. Combining Ability Studies for Development of New Hybrids in Rice over Environments. *Journal of Agricultural Science*, 2 (2): 225-233.
- Saravanan K, Ramya B, Satheesh Kumar P and Sabesan T 2006. Combining ability for yield and quality characters in rice (*Oryza sativa* L.). *Oryza*, 43 (4): 274-277.
- Tiwary DK, Pandey P, Giri SP and Dwivedi JL 2011. Heterosis studies for yield and its components in rice hybrids using CMS system. *Asian Journal of plant sciences*, 10(1): 29-42.
- Umakanta S, Biswas PS, Prasad B and Khaleque Mian MA 2002. Heterosis and genetic analysis in rice hybrids. *Pakistan J. Biol. Sci.*, 5(1): 1-5.