HETEROSIS FOR YIELD, YIELD COMPONENTS AND QUALITY TRAITS IN RICE HYBRIDS (ORYZA SATIVA L.)

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Abstract

Twenty hybrids developed from crossing four CMS lines and five testers were evaluated for the extent of heterosis over standard parent for yield, yield components and quality traits in rice during *kharif*, 2013. Four crosses out of twenty crosses exhibited highly significant standard heterosis for grain yield per plant. Heterosis for grain yield was manifested due to the significant and positive heterosis for its components *viz.*, Ear bearing tillers, number of fertile grains per panicle, test weight, Harvest index, Hulling%, Milling%, Kernel length, L/B ratio, kernel length after cooking and Elongation ratio. The top four heterotic combinations identified for grain yield per plant were APMS 9A×RM 83-19-3, CMS 12A×RM 83-19-3, APMS 10A×RM 89-12-3, APMS 10A×RM 80-55-2 which exhibited more than 20% standard heterosis.

Key words: CMS lines, heterosis, quality traits, rice, yield components.

Introduction

Rice is a staple food crop in India providing 43% of calories requirement for more than 70% of Indian population. In India, rice is cultivated in an area of 43.77 m. ha. with a production of 95.32 mt. Still there is need for the increase in productivity to meet the needs of the growing population. Hybrid rice giving a yield advantage of about 20-30% over high yielding varieties is a better choice for increasing productivity and quality also play a major role for fetching higher remunerative price. A higher yield over high yielding check varieties and wider adaptability has been instrumental in rapid spread of hybrid rice in India. Heterosis refers to the increase (or) decrease in F₁ value over the mean parental value. From the view point of plant breeding, increased yield of F, over the better (or) best commercial variety is more relevant (Virmani et al., 1981). Commercial exploitation of heterosis has been made possible by the use of cytoplasmic genetic male sterility and fertility restoration system. Identification of locally adapted restorers, which show consistently high degree of restoration of CMS lines would be great value in commercial hybrid programme. Hence, the present study was carried out to study the performance of the experimental hybrid crosses in order to estimate the magnitude of standard heterosis. Hence, the crosses with high heterotic potential could be exploited for further evaluation and commercial cultivation.

Materials and Methods

Four CMS lines viz., APMS 6A, APMS 9A, CMS 12A and APMS 10A were crossed with five testers RM 89-12-3, RM 83-19-3, RM 80-55-2, RM 80-55-3 and RM 1-21-1 in a line × Tester fashion to synthesize 20 hybrids (table 1). The hybrids along with parents and checks (varietal check RGL 2537 and hybrid check MTUHR 2089) were grown in a randomized block design (RBD) with three replications at Regional Agricultural Research Station, Anakapalle during kharif 2013. Recommended package of practice were followed to raise the crop. Each plot consisted of three rows of 5 m length, single seedling was planted per hill with a spacing of $20 \text{ cm} \times 15 \text{ cm}$. Observations were recorded on ten randomly selected plants for plant height, days to 50% flowering, ear bearing tillers plant⁻¹, panicle length, number of fertile grains panicle⁻¹, number of unfilled grains panicle⁻¹, test weight, harvest index and grain yield plant⁻¹. Quality characters

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Table 1 : Experimental material.

	- · - · - · · ·		
S. no.	Hybrid	Line	Tester
1.	TCN12591	APMS 6A (A1)	RM 89-12-3(R1)
2.	TCN12575	APMS 6A (A1)	RM 83-19-3(R2)
3.	TCN12580	APMS 6A (A1)	RM 80-55-2(R3)
4.	TCN12581	APMS 6A (A1)	RM 80-55-3(R4)
5.	TCN12553	APMS 6A (A1)	RM 1-21-1(R5)
6.	TCN12667	APMS 9A (A2)	RM 89-12-3(R1)
7.	TCN12662	APMS 9A(A2)	RM 83-19-3(R2)
8.	TCN12649	APMS 9A(A2)	RM 80-55-2(R3)
9.	TCN 12650	APMS 9A(A2)	RM 80-55-3(R4)
10.	TCN 12635	APMS 9A(A2)	RM 1-21-1(R5)
11.	TCN 12536	CMS 12 A(A3)	RM 89-12-3(R1)
12.	TCN 12528	CMS 12 A(A3)	RM 83-19-3(R2)
13.	TCN 12761	CMS 12 A(A3)	RM 80-55-2(R3)
14.	TCN 12762	CMS 12 A(A3)	RM 80-55-3(R4)
15.	TCN 12746	CMS 12 A(A3)	RM 1-21-1(R5)
16.	TCN 12692	APMS 10A(A4)	RM 89-12-3(R1)
17.	TCN 12693	APMS 10 A(A4)	RM 83-19-3(R2)
18.	TCN 12680	APMS 10 A(A4)	RM 80-55-2(R3)
19.	TCN 12681	APMS 10 A(A4)	RM 80-55-3(R4)
20.	TCN 12673	APMS 10 A(A4)	RM 1-21-1(R5)
	Varietal Check	RGL2537	CRT145*CRT1014
	Hybrid Check	MTUHR 2089	APMS 9A /
			MTU II-156-13-11-2

viz., Hulling per cent, Milling per cent, Head rice recovery, L/B ratio, Water uptake, Kernel length after cooking, elongation ratio, Alkali spreading value and Amylose content were recorded for all hybrids. Estimation of heterosis over standard parent was computed as suggested by Liang et al. (1971).

Results and Discussion

The analysis of variance of Line \times Tester revealed that the variance due to genotypes were highly significant for yield, yield components and quality character except for hulling per cent and elongation ratio indicating considerable variation in the material under study (tables 2a & 2b).

The estimates of standard heterosis for yield, yield components and quality character were given in tables 3a and 3b, respectively. In the present investigation, considerable heterosis existed both in positive and

lable 2a: Line × Tester ANOVA for yield and yield components in rice hybrids.

Source of variation	fp	Plant height (cm)	Days to 50% flowering	Ear bearing tillers	Panicle length (cm)	Fertile grains/panicle	Unfilled grains /panicle	Test weight (g)	Harvest	Grain yield/ plant (g)
Replicates	2	35.389	13.117*	1.358	*685.2	44.927	0.301	0.323	14.307*	82.853
Crosses	19	48.336*	63.996**	29.048**	5.882**	3522.07**	72.368**	9.322**	27.253**	1269.468**
Line Effect	3	72.483	37.839	116.299**	4391	264.639	66.616	18.294	31.540	2813.100*
Tester Effect	4	38.694	50.208	19.077	5.474	2224.815	33.169	9.489	23.945	1528.886
Line * Tester Eff.	12	45.514*	75.131**	10.559*	**068.9	4768.861**	86.873**	7.023**	27.284**	**180.797
Error	3%	20.659	3.696	4578	0.577	202.066	1.552	0.256	1.891	40.164
Total	59	30.071	23.434	12.349	2.347	1265.896	24.315	3.178	10.479	437.489

Table 2b : Line \times Tester ANOVA for quality traits in rice hybrids.

Source of	df.	Hulling	Milling	Head rice	Kernal	Kernal	L/B	Water	Amylose	Alkali	Kernal	Elongation
variation		0%	o %	recovery	(mm)	(mm)	rano	пртаке	content	spreading value	spreading length after value cooking	Kano
Replicates	2	8.253**	23.546*	0.424	0.047*	0.001	0.034*	646.667*	0.324	1.017	*090:0	31088.860
Crosses	19	10.903	22.955**	18.067**	0.271**	0.087 **	0.398**	3384.737**	1.761**	3.701 **	0.226**	31094.820
Line Effect	3	11.026	45.180	3.993	0.376	0.133	0.489	5767.778	1.105	4594	0.287	31068.480
Tester Effect	4	8.021	15.774	5.477	0.285	0.077	0.464	3166.042	4.919**	3.642	0.182	31133.670
Line * Tester Eff.	12	11.834**	19.792**	25.783**	0.240**	** 6200	0.354**	2861.875**	0.873*	3.497 **	0.225**	31088.460
Error	38	2.916	5.899	3.242	0.008	0.002	0.008	91.842	0.387	9650	600:0	31084.190
Total	59	5.669	11.990	7921	0.094	0.029	0.135	1171.073	0.827	1.610	0.080	31087.770

^{*}significant at 5% level, **significant at 1% level.

Table 3a: Standard heterosis for yield, yield components in rice hybrids.

Hybrids	Plant	Days to 50%	Ear bearing	Panicle	Fertile grains/	Unfilled grains	Test	Harvest	Grain yield
,	height (cm)	flowering	tillers	length (cm)	panicle	/panicle	weight (g)	index	/plant (g)
AIRI	4.01	**06.6-	34.60**	-25.35**	-33.56**	**86'29	6.91**	-5.87*	-43.37**
A1R2	-1.69	-8.95**	76.78**	-21.99**	-39.54**	109.45**	7.87**	10.27**	-9.64
A1R3	3.94	-3.51*	37.44**	-19.13**	-9.71*	-38.06**	-11.80**	3.93	32.01*
A1R4	-3.10	-11.82**	50.24**	-17.70**	-32.57**	30.18**	14.36**	-3.56	-34.77*
A1R5	**86.6-	-8.63**	41.23**	-11.43**	-22.73**	9.71	0.99	0.92	-1.72
A2R1	-1.97	-10.86**	15.64	-22.35**	-28.71**	25.46**	11.30**	0.65	-11.36
A2R2	2.81	-2.24	9.95	-13.32**	-22.73**	36.48*8	-0.53	-2.37	59.90**
A2R3	-2.74	-6.07**	-18.96	-18.70**	-14.83**	43.83**	7.67**	-6.42*	47.85**
A2R4	-6.18*	-3.51*	7.58	-16.34**	-25.12**	2.36	5.57*	11.67**	-25.6
A2R5	-6.32**	-4.15**	00.6	-13.38**	-27.13**	-1.84	5.38*	-10.24**	-38.21**
A3R1	-5.34*	-11.18**	35.55**	-11.43**	-40.02**	30.71**	20.66**	3.95	38.90**
A3R2	-7.31**	-6.39**	12.80	-9.54**	-7.81	-12.34	16.46**	429	75.39**
A3R3	-8.15**	-1.92	-13.74	-20.15**	-5.34	-37.53**	3.67	-7.81**	75.22**
A3R4	-2.11	**06.6-	12.80	-27.80**	-38.76**	24.93**	16.27**	3.80	32.70**
A3R5	-8.92**	-3.51*	-6.16	-17.52**	-31.69**	11.81	32.49**	-5.02	51.64*
A4R1	-10.61**	2.56	22.27	-5.88*	10.47*	-70.08**	27.15**	13.95**	117.04**
A4R2	-5.62*	-14.06**	24.64	-14.19**	-37.69**	10.24	0.23	-030	-31.33**
A4R3	4.57	96:0-	26.54*	-15.93**	41.23**	30.18**	14.55**	*80.9	151.46*
A4R4	-13.14**	-2.24	21.33	-19.82**	-33.24**	62.0	3.67	5.23	39.41**
A4R5	-2.95	**06.6-	28.91*	-14.65**	-28.16**	29.66**	9.39**	0.74	22.55**

^{*}significant at 5% level, **significant at 1% level.

Table 3b: Standard heterosis for quality traits in rice hybrids.

Hybrids Hulling Milling Head rice Kernel L/B Water Amylee Alkail Kernel Frending Prodigh alter Frending Prodigh alter Prodigh alter	THE COUNT	Statistate Hotels	tore death	table of a ballidate necession quality times in the hyprides	CITAB:							
4.11* 8.55*** 797* 538 10.96*** 9.10** 1.80 25.86*** -6.55 13.15 4.71* 8.55*** 133 3.79* 6.70** 9.76** -15.52** -8.85** -6.25 0.59 2.20 1.35 5.43* 2.83 8.23** -10.02** -15.23** -6.85** 1.50 0.59 1.77 1.35 108** 10.02** -10.03** -6.85** 1.50 0.59 0.90 0.40 562* 1.39 781** -7.03** -10.24* -2.14 -2.84 6.25 5.87* 0.90 0.40 562* -1.39 -7.81** -14.23** -16.39 -11.5 -3.14* -3.04* -3.16*	Hybrids	Hulling %	Milling %	Head rice recovery	Kernel length (mm)	Kernel breadth (mm)	L/B ratio	Water uptake	Amylose	Alkali preading value	Kernel length after cooking	Elongation ratio
4,71% 8,55%** -1,33 -3,79% 6,70%** -9,76%* -15,29** -6,25 -6,25 0,59 5,33%** 2,43 -2,83 8,23*** -10,00*** -10,03*** -6,85*** 15.0 191 2,0 1,3 -1,08*** 1,9 6,46*** 4,17 4,31** -6,85** 15.0 191 0,0 1,7 1,3 -3,68 -3,10* 13,02** -10,34* -2,84 6,25 5,87 0,0 0,40 -5,62* -1,39 -1,425** -10,34* -2,16 1,40** -10,34* -10,49* -10,49** -1,42** -1,425** -1,34* -1,34* -1,43* -1,43* -1,63* -1,43* -1,43* -1,63* -1,43* -1,54* -1,56* -1,40** -1,16** -1,40** -1,16** -1,40** -1,16** -1,40** -1,16** -1,40** -1,16** -1,40** -1,16** -1,40** -1,16** -1,50** -1,40** -1,16** -1,50** -1,40*	A1R1	-3.42	*76.7-	-3.58	10.96**	9.10**	1.80	25.86**	-4.85*	-6.25	13.15	5.86**
5.33** 2.48 -5.43* -2.83 8.23** -10.02** -12.93** -6.85** 15.0 -10.1 2.20 1.35 -1.085** 1.96 6.46** 4.17 4.31** -6.84 6.5 5.87 1.77 2.13 -3.68 -3.10* 1.302** -14.25** -13.79** 1.80 -3.12* 5.74** 0.90 -0.40 -5.62* -1.39 -7.81** 7.08* -10.34* -2.16 1.37* 2.14** 2.16** 1.11 4.30 2.16* 1.11 4.30 2.16* 2.15** 2.14** 2.16** 4.14** 2.24** 2.24* 2.24<	A1R2	4.71*	8.55**	-1.33	-3.79*	6.70**	**92.6-	-15.52**	-8.85**	-625	0.59	6.61**
220 1.35 -1.088** 1.96 6.46** 4.17 4.31** -2.84 6.55 5.87 1.77 2.13 -3.68 -3.10* 13.02** -14.25** -13.9* 1.80 -31.25* 2.74** 0.90 -0.40 -5.62* -1.39 -7.81** 7.08* -10.34 -2.16 -18.75 4.30** 0.36 -0.46 -2.07 -10.45** -7.81** 7.08* -10.34 -2.16 -18.75 4.30** 0.36 -1.15 -3.11 -3.81* 8.33** -11.68** -1.56 -18.75 1.47** 0.38 -1.15 -3.11 -3.81* 8.33** -11.16** -1.56 -1.87 1.47** 1.88 -1.24 -4.78 3.56* -1.16** 4.13** -2.48 -1.87 4.69** 1.53 0.74 -1.46 2.74 -4.13* -1.25 -2.48 -1.87 -1.87** 1.53 0.75 -1.53 -1.10 -1	A1R3	5.33**	2.48	-5.43*	-2.83	8.23**	-10.02**	-12.93**	-6.85**	12.50	1.91	9.14**
1.77 2.13 -3.68 -3.10* 13.02** -14.25** -13.79** 1.80 -31.25* 2.74*** 0.90 -0.40 -5.62* -1.39 -7.81** 7.08* -10.34* -2.16 -18.75 4.30*** 0.46 -2.07 -10.45** -3.19* 1.11 4.30 -16.38** -6.01** -18.75 4.30*** 0.36 -1.68 -9.24** 4.03** 7.40** -16.38** -15.6 -18.75 1.47** 0.78 -1.15 -3.11 -3.81* 8.33** -11.16** -15.2** -2.48 43.75** 0.64 1.58 -1.24 -4.78 3.56* -3.72 7.58** 2.59 2.50 12.50 1.40** 1.59 1.59 -1.46 -7.4 -4.13* 7.28** 4.31 -3.21 -3.12** 4.69** 1.59 -1.50 1.58* 3.19 -1.19 1.12 -1.22 -1.87* 1.87** 1.53** 2.53	A1R4	2.20	1.35	-10.85**	1.96	6.46**	4.17	4.31**	-2.84	6.25	5.87	8.56**
090 040 -5.62* -1.39 -7.81*** 7.08* -10.34* -2.16 -18.75 4.30*** 0.46 -2.07 -10.45*** -3.19* 1.11 4.30 -16.38** -6.01** -18.75 4.30*** 0.36 -1.68 -9.24** -4.03** -10.63** -16.38** -1.56 -18.75 2.05* 2.88 -1.15 -3.11 -3.81* 8.33** -11.16** -15.28* -1.56 1.47** 0.64 2.88 -1.15 -1.14 -3.81 8.33** -11.16** -15.29* 2.48 43.75** 0.64 1.53 0.74 -1.46 2.74 -4.13* 7.28** 4.31 -3.21 1.27** 1.15** 1.46** 1.53 0.74 -1.63 1.83 3.19 -1.19 1.72 -1.22 -1.87 8.26** 1.53 0.53 -6.00* 4.89** -12.19** 1.56** -5.24* -5.00** 7.59** 2.54	A1R5	1.77	2.13	-3.68	-3.10*	13.02**	-14.25**	-13.79**	1.80	-31.25*	2.74**	15.43**
-046 -2.07 -1.045** -3.19* 1.11 4.30 -16.38** -6.01** -18.75 2.05* 0.36 -1.68 -9.24** 4.03** 7.40** -16.38** -15.5 -1.56 -18.75 1.47** -1.64 -15.24** -0.64 -1.87 -1.87** -1.63** -1.63** -1.10** -1.52** -2.48 -43.75** 0.64 -1.47** -1.66** -1.52** -2.48 -43.75** 0.64 -1.47** -1.65** -2.48 -43.75** 0.64 -1.47** -1.10** -1.52** -2.48 -43.75** 0.64 -1.47** 0.64** -1.65 1.40** 0.64** -1.65 1.40** 0.64** -1.65 1.40** 0.64** -1.65 1.40** 0.64** -1.65** 1.40** 0.64** -1.69** -1.19** -1.19** -1.19** -1.19** -1.19** -1.15** -1.20** -1.20** -1.20** -1.20** -1.20** -1.20** -1.20** -1.20** -1.20** -1.20**	A2R1	0.90	-0.40	-5.62*	-1.39	-7.81**	7.08*	-10.34*	-2.16	-18.75	4.30**	16.05**
0.36 -1.68 -9.24** -4.03** 740** -10.63** -16.58** -1.56 -1.875 -1.875 1.47** -0.78 -1.15 -3.11 -3.81* 8.33*** -11.16** -15.52** -2.48 43.75** 0.64 2.88 -1.24 4.78 3.56* -3.72 7.65** 2.50 12.50 5.82** 1.53 0.74 -1.46 2.74 -4.13* 7.28** 4.31 -3.21 -3.21 -3.125* 4.69** 1.53 0.74 -1.46 2.74 -4.13* 7.28** 4.31 -3.21 -1.25 1.25 8.26** 3.24** 1.53 0.75 0.75 0.59 2.91 5.17 -3.21 -1.875 8.26** 3.28** 1.29 0.55 -6.00* 4.89** -12.19** 19.68** -45.69** -50.00** 7.59** 2.34 3.88* 3.82 -8.03** -7.22** -11.56** -7.15* -5.42* 1.50*	A2R2	-0.46	-2.07	-10.45**	-3.19*	1.11	430	-16.38**	-6.01**	-18.75	2.05*	11.81**
-0.78 -1.15 -3.81* 8.33** -11.16** -15.52** -2.48 -43.75** 0.64 2.88 -1.24 -4.78 3.56* -3.72 7.65** 2.59 2.50 12.50 5.82** 1.53 0.74 -1.46 2.74 -4.13* 7.28** 4.31 -3.21 -1.25 4.69** 5.82** 1.97 1.50 -1.63 1.83 3.19 -1.19 1.72 -1.22 -1.875 8.26** 7.59** 2.53 -3.23 -9.32** 3.43* 0.59 2.91 5.17 -3.21 -18.75 8.26** 1.29 0.95 -6.00* 4.89** -12.19** 19.68** 9.48 -6.37** 7.59** 7.59** -5.39** 4.72 10.36** 4.41* 5.74* 45.69** -18.75 12.1** 3.88* 3.82 -8.03** 7.22** 7.22** -11.56** -4.97* -12.50 6.60** 3.00 9.41 3.	A2R3	0.36	-1.68	-9.24**	-4.03**	7.40**	-10.63**	-16.38**	-1.56	-18.75	1.47**	12.95**
2.88 -1.24 -4.78 3.56* -3.72 7.65** 2.59 2.50 12.50 5.82*** 1.53 0.74 -1.46 2.74 -4.13* 7.28** 4.31 -3.21 -3.12* 4.69** 1.57 1.50 -1.63 1.83 3.19 -1.19 1.72 -1.22 -18.75 8.26** 2.53 -3.23 -9.32** 3.43* 0.59 2.91 5.17 -3.21 -18.75 8.26** 1.29 0.95 -6.00* 4.89** -12.19** 19.68** 948 -6.37** -50.00** 7.59** 5.39** 4.58 4.72 10.36** 4.41* 5.74* 45.69** 1.96 -18.75 12.71** 3.88* 3.82 -8.03** 3.02* 7.22** 7.12** -4.97* 1.25* 4.65** 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 3.11 3.35**	A2R4	-0.78	-1.15	-3.11	-3.81*	8.33**	-11.16**	-15.52**	-2.48	-43.75**	0.64	6.47**
1.53 0.74 -1.46 2.74 -4.13* 7.28** 4.31 -3.21 -3.21 -3.21 4.69** 4.69** 1.97 1.50 -1.63 1.83 3.19 -1.19 1.72 -1.22 -18.75 8.26** 2.53 -3.23 -9.32** 3.43* 0.59 2.91 5.17 -3.21 -18.75 8.26** 1.29 0.95 -6.00* 4.89** -12.19** 19.68** 9.48 -6.37** -50.00** 7.59** -5.39** 4.58 4.72 10.36** 4.41* 5.74* 45.69** 1.96 -18.75 12.1** 3.88* 3.82 -8.03** 7.22** 7.22** -11.56** -11.56** -5.42* 12.50 4.65** 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -5.50* 6.60** 2.11 3.35** 4.11 10.76** -15.52** 10.94** -10.34* -1.08 -5.32*	A2R5	2.88	-124	4.78	3.56*	-3.72	7.65**	2.59	2.50	12.50	5.82**	-1.08
1.57 1.50 -1.63 1.83 3.19 -1.19 1.72 -1.22 -1.875 8.26** 2.53 -3.23 -9.32** 3.43* 0.59 2.91 5.17 -3.21 -18.75 5.33** 1.29 0.95 -6.00* 4.89** -12.19** 19.68** 9.48 -6.37** -50.00** 7.59** -5.39** 4.58 4.72 10.36** 4.41* 5.74* 45.69** 1.96 -18.75 12.71** 3.88* 3.82 -8.03** 3.02* 19.58** -13.77** 1.72 -4.97* -12.50 4.65** 3.88* 3.88 -14.82** -5.23** 7.22** -11.56** -21.55** -5.42* 12.50 6.60** 3.00 9.41 3.98* 4.79** 14.17** -8.13** 9.48 -1.22 -5.50* 6.60** 2.11 3.35** 4.11 10.76** 10.35** -10.36** -10.34* -1.03* -5.22** 3.91** </td <td>A3R1</td> <td>1.53</td> <td>0.74</td> <td>-1.46</td> <td>2.74</td> <td>-4.13*</td> <td>7.28**</td> <td>4.31</td> <td>-3.21</td> <td>-31.25*</td> <td>4.69**</td> <td>-2.51</td>	A3R1	1.53	0.74	-1.46	2.74	-4.13*	7.28**	4.31	-3.21	-31.25*	4.69**	-2.51
2.53 -3.23 -9.32** 3.43* 0.59 2.91 5.17 -3.21 -18.75 5.33** 1.29 0.95 -6.00* 4.89** -12.19** 19.68** 9.48 -6.37** -50.00** 7.59** -5.39** 4.58 4.72 10.36** 4.41* 5.74* 45.69** 19.6 -18.75 12.1** 3.88* 3.88* -8.03** 3.02* 19.58** -13.77** 1.72 4.97* -12.50 4.65** 2.94 3.88 -14.82** 7.22** 7.22** -11.56** -21.55** -5.42* 12.50 6.60** 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 2.11 3.35** 4.11 10.76** 15.52** 31.32** -5.32* -50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.94* -10.34* -1.08 -5.52** 3.91**	A3R2	1.97	1.50	-1.63	1.83	3.19	-1.19	1.72	-1.22	-18.75	8.26**	24.58**
1.29 0.95 -6.00* 4.89** -12.19** 19.68** 948 -6.37** -50.00** 7.59** -5.39** 4.58 4.41* 5.74* 45.69** 1.96 -18.75 12.71** 3.88* 3.88* 3.02* 19.58** -13.77** 1.72 4.97* -12.50 4.65** 2.94 3.88 -14.82** 5.23** 7.22** -11.56** -5.42* 1.25 0.93 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 2.11 3.35** 4.11 10.76** -15.52** 31.32** 55.86** -5.32* 50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.96** -10.34* -10.8 -5.62.5** 3.91**	A3R3	2.53	-3.23	-9.32**	3.43*	0.59	2.91	5.17	-3.21	-18.75	5.33**	-2.08
-5.39** 4.58 -4.72 10.36** 4.41* 5.74* 45.69** 1.96 -18.75 12.71** 3.88* 3.82 -8.03** 3.02* 19.58** -13.77** 1.72 -4.97* -12.50 4.65** 2.94 3.88 -14.82** -5.23** 7.22** -11.56** -21.55** -5.42* 12.50 0.93 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 2.11 3.35** 4.11 10.76** -15.52** 31.32** -5.32* -50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.96** -10.34* -1.08 -56.25** 3.91**	A3R4	1.29	0.95	-6.00*	4.89**	-12.19**	19.68**	9.48	-6.37**	-50.00**	7.59**	3.74*
3.88* 3.88* -8.03** 3.02* 19.58** -13.77** 1.72 -4.97* -12.50 4.65** 2.94 3.88 -14.82** -5.23** 7.22** -11.56** -21.55** -5.42* 12.50 0.93 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 2.11 3.35** 4.11 10.76** -15.52** 31.32** 25.86** -5.32* -50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.96** -10.34* -1.08 -56.25** 3.91**	A3R5	-5.39**	4.58	4.72	10.36**	4.41*	5.74*	45.69**	1.96	-18.75	12.71**	13.97**
2.94 3.88 -14.82** -5.23** 7.22** -11.56** -51.55** -5.42* 12.50 0.93 3.00 9.41 3.98 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 2.11 3.35** -4.11 10.76** -15.52** 31.32** 25.86** -5.32* -50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.96** -1.034* -1.08 -56.25** 3.91**	A4R1	3.88*	3.82	-8.03**	3.02*	19.58**	-13.77**	1.72	-4.97*	-12.50	4.65**	-3.85*
3.00 9.41 3.58* 4.79** 14.17** -8.13** 9.48 -1.22 -25.00* 6.60** 2.11 3.35** 4.11 10.76** -15.52** 31.32** 25.86** -5.32* -50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.96** -10.34* -1.08 -56.25** 3.91**	A4R2	2.94	3.88	-14.82**	-5.23**	7.22**	-11.56**	-21.55**	-5.42*	12.50	0.93	15.72**
2.11 3.35** 4.11 10.76** -15.52** 31.32** 25.86** -5.32* -50.00** 13.35** 1.53 0.59 -3.52 -1.88 10.35** -10.96** -10.34* -1.08 -56.25** 3.91**	A4R3	3.00	9.41	3.98	4.79**	14.17**	-8.13**	9.48	-1.22	-25.00*	**09'9	-1.38
1.53 0.59 -3.52 -1.88 10.35** -10.96** -10.34* -1.08 -56.25** 3.91**	A4R4	2.11	3.35**	4.11	10.76**	-15.52**	31.32**	25.86**	-5.32*	-50.00**	13.35**	7.97
	A4R5	1.53	0.59	-3.52	-1.88	10.35**	-10.96**	-10.34*	-1.08	-56.25**	3.91**	16.22**

*significant at 5% level, **significant at 1% level.

Character	Hybrids	Per se performance	SCA effect	Standard heterosis
Plant height	A_4R_4	82.40	-4.30 **	-13.14**
	A_4R_1	84.80	-2.51	-10.61**
	A_1R_5	85.40	-4.62**	-9.98**
Days to 50% flowering	A_4R_2	89.66	-7.92**	-14.06**
	A_3R_1	92.66	-3.77**	-11.82**
	A_1R_4	92.00	-2.87*	-11.18**
Ear bearing tillers plant ⁻¹	A_1R_2	24.86	2.69*	76.78**
	A_1R_4	21.13	0.09	50.24**
	A_1R_5	19.86	-0.51	41.23**
No. of fertile grains panicle ⁻¹	A_4R_1	274.80	84.31**	10.47**
No. of unfilled grains per panicle	A_4R_1	3.80	-8.71**	-70.08**
	A_1R_3	7.86	-7.41**	-38.06**
	A_3R_3	7.93	-3.23**	-37.53**
Test weight	A_3R_5	23.13	2.12**	32.49**
	A_3R_1	21.06	-0.72*	20.66**
	A_3R_2	20.33	0.38	16.46**
Harvest index	A_4R_1	51.66	2.90**	13.95**

50.63

50.00 97.40

84.06

61.93

Table 4a: The best heterotic hybrids identified for yield and yield components based on overall performance.

Grain yield plant⁻¹

negative directions for all the traits. The hybrid A4R2 showed significant negative, but desirable heterosis for days to 50% flowering as also reported by Yolanda and Vijendradas (1996). High number of ear bearing tillers plant⁻¹ was observed in A₁R₂, however presence of high heterosis for this trait could not result in higher yield, especially in hybrids where spikelet sterility was greatly affected due to varying levels of fertility restoration. Therefore due consideration is required for both panicles plant⁻¹ and spikelet fertility simultaneously (Virmani *et al.*, 1981).

 $\frac{A_4R_1}{A_5R_4}$

 A_1R_2

 A_4R_3

 $A_{1}R_{1}$

 $A_{2}R_{2}$

Fourteen hybrids exhibited significant positive heterosis for test weight (Yolanda and Vijendradas, 1996). Most of the hybrids exhibited positive and significant heterosis for kernel length after cooking and elongation ratio.

The best heterotic hybrids selected for grain yield, yield components and quality traits presented in the tables 4a & 4b. Based on *per se* performance, significant *sca* effects and heterosis for yield in the crosses APMS 6A × RM 83-19-3, CMS 12A × RM 83-19-3, APMS 10A ×

RM 89-12-3 and APMS $10A \times RM$ 80-55-2 recorded high *per se* performance, significant *sca* effects and standard heterosis over the best check MTUHR 2089 for grain yield plant⁻¹ were identified as promising heterotic hybrids.

11.67**

10.27**

151.46**

117.04**

59.90**

4.30**

4.11**

16.42**

22.97**

22.16**

A perusal of the results on hybrid vigour over standard check for the different traits studied revealed maximum expression of heterosis for grain yield per plant followed by ear bearing tillers. Hybrid vigour to an extent of 151.46 per cent over standard check in the hybrid A_4R_3 followed by extent of 117.04 per cent in the hybrid A_4R_1 . Similar results were reported by Singh *et al.* (2007), Tiwari *et al.* (2011) and Venkata Subbaiah *et al.* (2013).

All the high yielding hybrids also manifested significant and useful heterosis over standard check for traits like plant height, test weight, harvest index, kernel length after cooking and elongation ratio (table 4). The present study confirmed that high heterotic combinations were realized in the cross combinations APMS $6A \times RM$ 83-19-3, CMS $12A \times RM$ 83-19-3, APMS $10A \times RM$ 89-12-3 and APMS $10A \times RM$ 80-55-2. Parental gca

^{**}significant at 1% level, *significant at 5% level.

Table 4b: The best heterotic hybrids identified for quality traits based on overall performance.

Character	Hybrids	Per se performance	SCA effect	Standard heterosis
Hulling %	A_1R_3	79.42	1.40	5.33**
	A_1R_2	78.95	1.31	4.71*
Milling %	A_1R_2	74.27	3.49*	8.55**
Kernel length	A_1R_1	6.49	0.46**	10.96**
	A_4R_4	6.47	0.38**	10.76**
	A_3R_5	6.45	0.29 **	10.36**
Kernel breadth	A_4R_4	1.62	-0.30**	-15.52**
	A_3R_4	1.68	-0.07*	-12.19**
	A_2R_1	1.77	-0.18**	-7.81**
L/b ratio	A_4R_4	3.99	0.72**	31.32**
	A_3R_4	3.64	0.08	19.68**
	A_2R_5	3.27	0.35**	7.65**
Water uptake	A_4R_2	151.66	-18.33**	-21.55**
	A_2R_3	161.66	-2.58	-16.38**
	A_2R_4	163.33	-19.67**	-15.52**
Amylose content	A_1R_2	22.75	-0.56	-8.85**
	A_1R_3	23.25	-0.60	-6.85**
	A_3R_4	23.37	-0.69	-6.37**
Alkali spreading value	A_4R_5	2.33	-1.37**	-56.25**
	A_4R_4	2.66	-0.45	-50.00**
	A_2R_4	3.00	-0.58	-43.75**
Kernel length after cooking	A_4R_4	7.72	0.40**	13.35**
	A_3R_5	7.68	0.27**	12.71**
	A_3R_2	7.38	0.20**	8.26**
Elongation ratio	A_3R_2	1.44	0.12**	24.58**
	A_4R_5	1.34	0.07**	16.22**
	A_2R_1	1.34	0.13**	16.05**

^{**}significant at 1% level,

status for grain yield, yield components and quality traits indicating the importance of both additive and non-additive gene effects in realization of heterosis for grain yield. The heterosis potential realized in these four hybrids could be further evaluated across the locations and environments for testing their feasibility of commercialization in due course.

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^{*}significant at 5% level.