



## Assessment of heterobeltiosis and useful heterosis for seed yield and its components in (*Brassica juncea* L.)

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(Received: 22 July 2015; Revised: 7 Nov 2016; Accepted: 21 Nov 2016)

### Abstract

Ninety six crosses of Indian mustard [*Brassica juncea* (L.) Czern & Coss.] obtained by crossing of forty eight lines with two testers in a line x tester fashion were used to assess the possibility of estimating the amount of heterosis for seed yield, yield components and oil content. These parents and crosses were grown in randomized complete block design replicated thrice and observations were taken on days to maturity, plant height at maturity (cm), number of branches plant<sup>-1</sup>, number of siliquae plant<sup>-1</sup>, seed yield plant<sup>-1</sup> (g), and oil content (%). Heterosis of higher magnitude was expressed for seed yield plant<sup>-1</sup> and number of siliqua plant<sup>-1</sup>. The crosses ACN-9 x ACN-169 and Pusa bold x ACN-164 were identified as the superior crosses which can be utilized for development of high yielding genotypes.

**Key words:** Mustard, heterosis, heterobeltiosis and useful heterosis

### Introduction

Rapeseed-mustard is the third important oilseed crop in the world after soybean (*Glycine max*) and palm (*Elaeis guineensis* Jacq.) oil. Among the seven edible oilseeds cultivated in India, rapeseed-mustard (*Brassica spp.*) contributes 28.6% in the total production of oilseeds. In India, it is the second most important edible oilseed after groundnut sharing 27.8% in the India's oilseed economy. The share of oilseeds is 14.1% out of the total cropped area in India, rapeseed-mustard accounts for 3%. Indian mustard (*Brassica juncea*) is largely self pollinated crop but certain amount (5-18%) of cross pollination may take place (Labana and Banga, 1984).

Heterosis breeding could be potential alternative for achieving quantum jump in production and productivity. The heterosis component is largely dependent on diverse parents with good general combining ability (gca) high. Selection of desirable heterotic crosses at an early stage is very important in developing high-yielding genotypes. Hence, the present study was aimed to select cross F<sub>1</sub> generation which can be forwarded further generations.

### Materials and Methods

The study was conducted during the year 2013-14 at experiment farm of Agricultural Botany Section, College of Agriculture, Nagpur, India. During Rabi 2013-14, 96 crosses, where obtained by crossing 48 lines with two testers in L x T mating design. Crossed seeds of these 96 hybrids crosses, along with 50 parents were planted in randomized block design with three replications with plot size 0.45 x 0.3 m. Recommended package of practices were followed to raise a good crop. The data were recorded on five randomly selected plants from each genotype on following six characters viz., days to maturity, plant height at maturity (cm), number of branches plant<sup>-1</sup>, number of siliqua plant<sup>-1</sup>, seed yield plant<sup>-1</sup> (g) and oil content (%).

### Results and Discussion

Heterosis breeding has been extensively exposed and utilized for increasing yield in a number of crops particularly cross pollinated crops. The heterosis observed in these crosses justifies the development of commercial hybrid in Indian mustard. In order to explode the hybrid vigor at commercial level, attempt

Table 1: Analysis of variance for the experimental design

Sources	d.f	Days to 50% flowering	Days to maturity	Plant height at maturity(cm)	Number of branches plant <sup>-1</sup>	Number of siliqua plant <sup>-1</sup>	Seed yield plant <sup>-1</sup>
Replication	2	24.35	51.50	288.68	0.57	281.84	2.30
Treatments	145	21.96**	24.49**	433.67**	0.47**	1907.74**	30.02**
Parents	49	24.49**	19.94**	450.21**	0.56**	7733.76**	31.80**
Crosses	95	20.33**	21.13**	429.08**	0.37**	9128.42**	29.35**
Parents vs. crosses	1	53.36**	567.51**	59.35	5.90**	74467.93**	6.69**
Error	290	2.95	3.89	98.24	0.12	198.00	2.24

\*\* = significant at 1% level

Table 2: Heterobeltiosis (H<sub>1</sub>), and useful heterosis (H<sub>2</sub>) for various characters

Crosses	Days to maturity		Plant height at maturity plant <sup>-1</sup>		Number of branches plant <sup>-1</sup>		Number of siliqua plant <sup>-1</sup>		Seed yield plant <sup>-1</sup>	
	H <sub>1</sub>	H <sub>2</sub>	H <sub>1</sub>	H <sub>2</sub>	H <sub>1</sub>	H <sub>2</sub>	H <sub>1</sub>	H <sub>2</sub>	H <sub>1</sub>	H <sub>2</sub>
ACN -9 X ACN-141	0.96	1.61	17.30**	17.30**	-1.01	-2.00	37.61**	78.01**	-28.73	-36.92**
ACN -9 X ACN-142	0.00	0.65	18.64**	18.64**	3.03	2.00	-14.41**	100.59**	77.95**	33.46*
ACN -9 X ACN-143	-1.92	-1.28	0.92	0.92	31.48**	-26.00**	-30.47**	6.89	-41.84**	-32.69*
ACN -9 X ACN-144	-0.32	0.33	-2.49	-2.49	-30.08**	-14.00	-13.06*	28.35**	10.98	5.00
ACN -9 X ACN-145	3.52*	4.19**	2.99	2.99	-11.11	-12.00	-7.77	40.20**	28.72	-3.46
ACN -9 X ACN-146	0.32	0.96	12.61*	12.61*	-11.67	6.00	-5.91	65.84**	-2.85	3.46
ACN -9 X ACN-147	0.00	0.65	3.61	3.91	-0.95	4.00	5.81	48.72**	-18.39	-18.08
ACN -9 X ACN-148	-1.27	-0.64	-0.97	-0.97	1.96	4.00	-2.26	48.45**	2.23	7.69
ACN -9 X ACN-149	0.00	0.65	2.07	2.07	2.86	8.00	-37.58	36.23**	-22.34	-18.46
ACN -9 X ACN-150	1.27	1.93	-0.88	-0.87	-24.32**	-16.00	-30.19	52.55**	-39.25**	-20.58
ACN -9 X ACN-151	-2.23	-1.60	19.46**	19.47**	-6.67	12.00	12.20**	113.79**	45.08**	75.77**
ACN -9 X ACN-152	2.23	2.89	12.88**	12.89*	-3.60	7.00	-19.30**	74.40**	-11.25	18.08
ACN -9 X ACN-153	-0.32	0.33	5.66	5.66	-6.67	-2.00	12.73**	117.85**	-32.13**	-13.08
ACN -9 X ACN-154	1.60	2.26	15.78**	15.78**	-1.01	-2.00	38.86**	66.97**	6.78	-14.62
ACN -9 X ACN-155	0.64	1.29	2.44	2.44	-18.70**	0.00	50.05**	112.44**	-5.50	26.15
ACN -9 X ACN-156	-2.56	-1.93	8.88	8.88	15.15	14.00	-0.43	46.78**	4.76	1.54
ACN -9 X ACN-157	-1.60	0.96	9.29	9.30	12.61	31.00**	15.27**	75.12**	32.21*	2.69
ACN -9 X ACN-158	0.31	0.96	8.28	2.02	-3.42	-2.00	48.75*	126.09**	100.51**	50.38**
ACN -9 X ACN-159	6.07**	6.75**	23.03**	20.11**	4.90	16.00	33.57**	74.76**	76.87**	63.46**
ACN -9 X ACN-160	3.19*	3.86*	20.32**	14.27*	9.91	10.00	119.12**	163.99**	-21.08	-4.62
ACN -9 X ACN-161	1.60	2.26	29.06**	21.63**	15.15	14.00	22.65**	54.30**	-15.36	-0.38
ACN -9 X ACN-162	-2.88	-2.25	25.82**	19.97**	27.27**	26.00**	67.80**	88.73**	18.78	60.77**
ACN -9 X ACN-163	-2.23	-1.60	17.21**	17.21**	-11.71	-2.00	20.39**	66.83**	46.10**	46.15**
ACN -9 X ACN-164	-1.92	-1.28	6.92	6.72	15.15	14.00	86.44**	107.98**	192.53**	46.15**
ACN -9 X ACN-165	-1.27	-0.64	6.16	6.17	8.57	14.00	51.73**	149.03**	55.70**	66.54**
ACN -9 X ACN-166	-1.92	-1.28	6.58	6.58	-2.86	2.00	34.04**	69.90**	8.85	23.46
ACN -9 X ACN-167	0.96	1.61	10.21	10.22	13.13	12.00	-25.33**	30.87**	0.40	6.92
ACN -9 X ACN-168	0.96	1.61	6.12	6.12	-14.29	-10.00	-16.35**	29.47**	4.48	-19.23
ACN -9 X ACN-169	-2.23	-1.60	8.05	8.05	-17.07*	2.00	15.80**	85.13**	-20.46**	79.23**
ACN -9 X ACN-170	-0.96	-0.32	-5.98	-5.98	-11.71	-2.00	-24.62**	33.57**	24.84	14.23
ACN -9 X ACN-171	-0.64	0.00	13.53*	13.53*	12.38	15.52*	-4.01	83.19**	55.76**	56.47**
ACN -9 X ACN-172	-2.23	-1.60	17.38**	12.52	19.61*	22.00	20.02**	63.36**	95.67**	53.08**
ACN -9 X ACN-173	-0.64	0.00	32.49**	24.62**	9.09	8.00	18.22*	42.00**	-21.07	-3.46
ACN -9 X ACN-174	0.32	0.96	5.15	5.15	36.36**	35.00**	47.25**	101.17**	91.77**	104.23**
ACN -9 X ACN-175	0.32	0.96	31.43**	14.91*	17.17*	16.00	70.86**	102.03**	-14.92	-1.54
ACN -9 X ACN-176	0.96	1.61	15.78**	15.78**	15.15	14.00	15.97*	60.93**	-0.73	9.62
ACN -9 X ACN-177	0.00	0.65	8.74	8.74	13.98	6.00	20.55**	42.09**	36.25**	47.31**
ACN -9 X ACN-178	4.20**	3.54*	7.54	7.55	-4.76	0.00	15.94**	74.00**	2.71	47.31**

ACN -9 X ACN-179	0.96	1.61	-2.80	-3.08	-8.11	2.00	63.87**	100.18**	30.80	3.08
ACN -9 X ACN-180	3.19*	3.86*	7.18	7.18	-13.51	-4.00	-30.01**	43.44**	-21.04	-32.31*
ACN -9 X ACN-181	5.11**	5.79**	5.93	5.94	-8.77	4.00	106.96**	174.58**	98.97**	49.23**
ACN -9 X ACN-182	1.27	1.93	7.48	7.13	14.29	20.00*	91.04**	112.57**	148.72**	86.54**
ACN -9 X ACN-183	1.28	1.61	15.25**	7.50	12.96	22.00*	22.27**	118.97**	102.05**	51.54**
ACN -9 X ACN-184	0.64	1.29	28.83**	19.70**	13.13	12.00	98.25**	119.78**	103.12**	55.38**
ACN -9 X ACN-185	-1.60	-0.96	18.64**	16.01**	-5.05	-6.00	61.61**	83.15**	74.87**	31.15*
ACN -9 X ACN-186	1.60	2.23	19.46**	19.47**	-4.27	12.00	39.13**	130.96**	21.56*	95.38**
ACN -9 X ACN-187	3.83**	4.51**	18.31**	18.32**	-18.10*	-14.00	17.30**	68.77**	36.92*	2.69
ACN -9 X ACN-188	1.92	2.58	14.13*	14.13*	5.88	8.00	24.38**	71.47**	53.75**	39.62**
Pusa bold X ACN-141	3.86*	3.86**	13.94*	13.99*	-1.01	-2.00	13.28*	50.34**	48.02**	48.08**
Pusa bold X ACN-142	4.82**	4.82**	8.47	8.51	11.11	10.00	-25.46**	77.47**	28.41*	28.46*
Pusa bold X ACN-143	3.22*	3.22*	14.12*	14.17*	-5.56	2.00	23.83**	73.55**	100.73**	132.31**
Pusa bold X ACN-144	3.86*	3.86*	18.72**	18.78**	-23.58**	-6.00	22.20**	71.65**	9.57	9.62
Pusa bold X ACN-145	3.86*	3.86*	15.37**	15.42**	19.19*	18.00*	11.67*	65.21**	48.79**	48.85**
Pusa bold X ACN-146	2.58	2.58	15.69**	15.74**	-1.67	18.00*	8.90*	88.55**	28.93*	37.31**
Pusa bold X ACN-147	1.61	1.61	22.59**	22.64**	20.00**	26.00**	39.18**	88.82**	37.55**	38.08**
Pusa bold X ACN-148	4.82**	4.82**	14.03*	14.08*	19.61*	22.00*	1.70	53.58**	25.96	32.69*
Pusa bold X ACN-149	4.51**	4.51**	9.98	10.03	-6.67	-2.00	-37.70**	36.01**	-15.75	-11.54
Pusa bold X ACN-150	6.12**	6.12**	22.13**	22.18**	-15.32*	-6.00	-42.36**	29.70**	91.82**	150.77**
Pusa bold X ACN-151	4.82**	4.82**	16.15**	16.20**	-21.90**	-18.00*	-17.47**	62.96**	-22.86*	-6.54
Pusa bold X ACN-152	4.51**	4.51*	16.88**	16.94**	-2.70	8.00	34.24**	175.80**	30.67**	73.85**
Pusa bold X ACN-153	3.22*	3.22*	8.28	8.33	-20.00	-16.00	-50.23**	8.20	-7.21	18.85
Pusa bold X ACN-154	1.93	1.93	8.60	8.65	-1.01	-2.00	-13.25	12.39	-6.57	-6.54
Pusa bold X ACN-155	8.36**	8.36**	18.68**	18.73**	-18.70**	0.00	-5.77	41.42**	-28.26**	-4.23
Pusa bold X ACN-156	6.75**	6.75**	14.81**	14.86**	-5.05	-6.00	11.57	61.87**	-20.42	-20.38
Pusa bold X ACN-157	3.22*	3.22*	2.44	2.49	-18.92*	-10.00	-7.43	31.64**	-9.27	-9.23
Pusa bold X ACN-158	0.00	0.00	4.78	3.96	17.95*	-4.00	-44.58**	16.72**	-23.11	-23.08
Pusa bold X ACN-159	2.89	2.89	4.52	-2.02	-1.96	0.00	-5.93	35.20**	0.73	0.77
Pusa bold X ACN-160	0.00	0.00	19.15**	17.40**	-11.71	-2.00	15.52**	51.69**	37.04**	45.00**
Pusa bold X ACN-161	1.29	1.29	26.03**	18.78**	3.03	2.00	18.65**	49.98**	13.73	33.85*
Pusa bold X ACN-162	0.00	0.00	12.94	7.69	1.01	0.00	10.49	31.64**	-49.08**	-31.08*
Pusa bold X ACN-163	0.33	0.33	10.81	10.86	-11.71	-2.00	-21.77**	15.95*	-2.35	-2.31
Pusa bold X ACN-164	0.00	0.00	11.30*	11.09*	7.53	0.00	26.51**	30.15**	43.79**	43.85**
Pusa bold X ACN-165	8.36**	8.36**	12.42*	12.47*	-4.76	0.00	-30.00	26.50**	-9.03	-2.69
Pusa bold X ACN-166	4.19**	4.19**	8.97	9.02	-4.76	0.00	-6.88	24.61**	-2.68	10.38
Pusa bold X ACN-167	8.36**	8.36**	8.47	8.51	-21.21*	-22.00*	-35.67**	15.73*	-1.77	4.62
Pusa bold X ACN-168	5.47**	5.47**	3.04	3.08	-8.57	-4.00	-13.55*	33.08**	-22.34	-22.31
Pusa bold X ACN-169	7.40**	7.40**	16.70	16.75**	-18.70**	0.00	-45.05**	28.30**	-68.56**	-27.69*
Pusa bold X ACN-170	7.40**	7.40**	2.65	2.69	-26.13**	-18.00*	-40.48**	10.00	-19.65	-19.62
Pusa bold X ACN-171	8.05**	8.05**	-2.80	-2.76	-31.43**	-28.00**	-55.74**	-3.92	-34.11**	-18.65
Pusa bold X ACN-172	7.08**	7.08**	-4.70	-8.65	-17.65	-16.00	-25.40**	9.69	4.77	4.81
Pusa bold X ACN-173	6.12**	6.12**	7.78	1.38	43.43**	42.00**	19.99**	43.80**	-25.47	-8.85
Pusa bold X ACN-174	5.47**	5.47**	5.71	5.75	-9.09	-10.00	-24.88**	13.38	-37.88**	-33.85*
Pusa bold X ACN-175	6.75**	6.75**	-0.15	-12.70	9.09	8.00	-27.39**	-1.76	-25.56**	-13.85
Pusa bold X ACN-176	6.75**	6.75**	5.84	5.89	8.08	7.00	-7.65	32.54**	-18.84	-10.38
Pusa bold X ACN-177	7.40**	7.40**	8.05	8.10	5.05	4.00	84.57**	83.24**	-21.74	-15.38
Pusa bold X ACN-178	8.74**	8.05**	8.56	8.61	-4.76	0.00	-10.93	38.67**	-54.41**	-34.62*
Pusa bold X ACN-179	4.19**	4.19**	-4.89	-5.20	-13.51	-4.00	38.97**	73.32**	40.72**	40.77**
Pusa bold X ACN-180	5.47**	5.47**	5.02	5.06	-2.70	8.00	-1.27	93.51**	5.73	5.77
Pusa bold X ACN-181	5.79**	5.79**	-0.83	-0.78	-22.81**	-12.00	6.67	51.96**	-28.10*	-28.08*
Pusa bold X ACN-182	6.43**	6.43**	-3.60	-3.91	-21.90	-18.00*	115.05**	132.81**	19.95	20.00
Pusa bold X ACN-183	4.82**	4.82**	1.93	-4.92	-16.67**	-10.00	-29.89**	34.75**	-15.42	-15.38
Pusa bold X ACN-184	6.43**	6.43**	18.33**	9.94	-19.19*	-20.00*	25.77**	28.66**	-2.35	-2.31
Pusa bold X ACN-185	6.75**	6.75**	4.33	1.98	-11.11	-12.00	29.95**	29.65**	-9.65	-9.62
Pusa bold X ACN-186	8.05**	7.45**	-7.73	-7.69	-12.82	2.78	-20.78**	40.78**	-46.16**	-16.55
Pusa bold X ACN-187	7.08**	7.08**	7.78	7.82	-21.90**	-18.00*	-32.57	6.17	-18.88	-18.85
Pusa bold X ACN-188	7.08**	7.08**	14.58**	14.63**	-5.88	-4.00	40.04**	90.36**	-3.50	-3.46
C.D. at 1%	4.17	4.17	20.97	20.97	0.74	0.74	29.78	29.78	3.16	3.16
C.D.at5%	3.17	3.17	15.92	15.92	0.56	0.56	22.61	22.61	2.40	2.40
SE(m)	2.59	2.59	65.47	65.47	0.08	0.08	132.02	132.02	1.49	1.49

\*,\*\* = significant at 5% and 1% level respectively

should be made to convert high yielding heterotic parents into cytoplasmic male sterile lines and search for fertility restorer lines in the germplasm to develop mustard hybrid.

Data in table 1 revealed that the mean squares due to replication were non significant for plant height at maturity, number of siliqua plant<sup>-1</sup> and seed yield plant<sup>-1</sup>. Treatment differences were significant for all the traits studied. Kang *et al.* (2013) reported that analysis of variance for various characters showed significant differences in different parents for all the traits. Similar results of significant mean sum of square of parents and crosses for all the traits studied were also reported by Rameeh (2012). Significant differences among parents and hybrid in F<sub>1</sub> for all the characters except 1000 seed weight were also reported by Sincik *et al.* (2011). Vaghela *et al.* (2011) also reported significant differences among genotypes for all the characters studied.

The cross Pusa bold X ACN-150 was identified to exhibit maximum useful heterosis of (150.77%) and (91.82%) heterobeltiosis for seed yield plant<sup>-1</sup>. The same cross also possessed useful heterosis of (29.70%) for number of siliqua plant<sup>-1</sup>. Another cross Pusa bold X ACN-143 recorded significant useful heterosis (132.31%) and heterobeltiosis (100.31%) for seed yield plant<sup>-1</sup>. The same cross also possessed significant useful heterosis (73.55%) and heterobeltiosis (23.83%) for number of siliqua plant<sup>-1</sup>. Another cross exhibiting significant useful heterosis for seed yield plant<sup>-1</sup> are ACN-9 X ACN-174 (H<sub>1</sub> 91.77 % H<sub>2</sub> 104.23%). The same cross also showed significant useful heterosis and heterobeltiosis for number of siliqua plant<sup>-1</sup> (H<sub>2</sub> 101.17%, H<sub>1</sub> 47.25%) and number of branches plant<sup>-1</sup> (H<sub>2</sub> 35.00%, H<sub>1</sub> 36.36%). ACN-9 X ACN-159 had significant useful heterosis and heterobeltiosis for seed yield plant<sup>-1</sup> (H<sub>2</sub> 63.46%, H<sub>1</sub> 76.87%), and number of siliqua plant<sup>-1</sup> (H<sub>2</sub> 74.76%, H<sub>1</sub> 33.57%). This was followed by ACN-9 X ACN-182 which possessed useful heterosis and heterobeltiosis for seed yield plant<sup>-1</sup> (H<sub>2</sub> 86.54%, H<sub>1</sub> 148.72%), Number of siliqua plant<sup>-1</sup> (H<sub>2</sub> 112.57%, H<sub>1</sub> 91.04), number of branches plant<sup>-1</sup> (H<sub>2</sub> 20.00%, H<sub>1</sub> 14.29%). Significant useful heterosis and heterobeltiosis were exhibited by the cross ACN-9 X ACN-183 (H<sub>2</sub> 51.54%, H<sub>1</sub> 102.05%), number of siliqua plant<sup>-1</sup> (H<sub>2</sub> 118.97%, H<sub>1</sub> 22.27%). Notable amount of heterosis for seed yield plant<sup>-1</sup>

(104.3%) and number of siliqua plant<sup>-1</sup> (39.03) were also reported by Joshi (2001), Katiyar *et al.* (2005) reported heterosis upto (91.6%) over better parents (90.5%) over check varieties for seed yield. Wassu (2011) reported standard heterosis upto (191.57%) for number of pod plant<sup>-1</sup> while, (16.64 to 66.09%) for seed yield plant<sup>-1</sup>. Results of high heterosis for seed yield plant<sup>-1</sup> contributing traits were also reported by Ghosh *et al.* (2002). The level of heterosis observed in these crosses justifies the development of commercial hybrid in Indian mustard. Commercial exploitation of hybrid in *Brassica juncea* has been reported by many mustard workers like Meena *et al.* (2014, 2015). Akbar *et al.* (2007) also reported that crosses with positive heterosis and heterobeltiosis may be utilized for hybrid seed production by cms and restorer lines.

Heterosis has positive association with the sca effects. Sca effects of crosses revealed that the crosses ACN-9 X ACN-169 and Pusa bold X ACN-164 exhibited significant positive sca effects seed yield plant<sup>-1</sup> and number of branches plant<sup>-1</sup> while, the crosses Pusa bold x ACN-169, ACN-9 X ACN-184 and ACN-9 X ACN-164 exhibited significant negative sca effects seed yield plant<sup>-1</sup> and number of siliqua plant<sup>-1</sup>. Maha (2008) also concluded that sca effects showed wide variation in the level of significant for different character studied.

Heterotic crosses showing substantial and significant sca effects for seed yield and number of siliqua plant<sup>-1</sup> over mid, better parent and over check varieties are indicated table 3. Dixit *et al.* (2005) reported that the crosses showing heterobeltiosis for seed yield plant<sup>-1</sup> possessed significant and positives sca effects. Similar results of significant correlation between heterosis for seed yield and mean performance of the crosses were reported by Rameeh (2012). Aher *et al.* (2009) found that the promising hybrid also had high *per se* performance and significant desirable sca effects for various characters.

From data of heterobeltiosis and useful heterosis, it was observed that crosses with high magnitude of heterosis had higher magnitude of sca effects and better *per se* performance. Hence selection of superior crosses for development of hybrids should necessarily base not only on the magnitude of heterosis, but also high mean performance and high

Table 3: Superior crosses showing high heterosis ( $H_1$ ,  $H_2$  and  $H_3$ ), high sca and high mean for number of seed yield plant<sup>-1</sup> and number of siliqua plant<sup>-1</sup>

Crosses	Seed yield plant <sup>-1</sup>				Number of siliqua plant <sup>-1</sup>			
	Mean	sca	$H_1$	$H_2$	Mean	sca	$H_1$	$H_2$
ACN-9 X ACN-169	15.53**	6.09**	—	79.23**	242.00**	79.26**	15.80**	85.13
Pusa bold X ACN-164	12.47**	3.84**	43.79**	43.85**	160.67	43.73	26.51**	30.15**

\*, \*\*= significant at 5% and 1% level, respectively.

sca effects. The first cross ACN-9 X ACN-169 exhibited significant useful heterosis ( $H_2$ ) for seed yield plant<sup>-1</sup>. The same cross also exhibited significant sca and mean for seed yield plant<sup>-1</sup> and number of siliqua plant<sup>-1</sup> and significant heterosis in to ( $H_1$  and  $H_2$ ) for number siliqua plant<sup>-1</sup>. Another cross Pusa bold x ACN-164 also exhibited significant desirable heterosis ( $H_1$  and  $H_2$ ) for seed yield as well as number of siliqua plant<sup>-1</sup>, significant sca for seed yield and number of siliqua plant<sup>-1</sup> and significant mean for seed yield plant<sup>-1</sup>. These crosses after their evaluation in trials can be used for development of hybrids after conversion of female line in to CMS background and male parents as restorers.

## References

- Aher CD, Chinchane VN, Shelke LT, Bograonkar SB and Gaikwad AR. 2009. Genetic study in Indian mustard [*B. juncea* (L.) Czern & Coss.]. *Int J Plant Sci* **4**: 83-85.
- Akbar M, Tahira and Hussain M. 2007. Heterosis for seed yield and it's componants in rapeseed *J Agric Res* **45**: 95-104.
- Dixit S, Kumar K and Chauhan Y S. 2005. Heterosis in Indian mustard [*B. juncea* (L.) Czern & Coss.]. *Agric Sci Digest* **25**:19-22.
- Ghosh SK, Gulati SK and Raman R. 2002. Combining ability and heterosis for seed yield and it's components in Indian mustard [*B. juncea* (L.) Czern & Coss]. *Indian J Genet* **62**: 29-33.
- Joshi SV. 2001. Studies on heterosis and Combining ability analysis for yield components in Indian mustard (*B. juncea* L.) thesis (unpub.) submitted to Dr. PDKV., Akola.
- Kang SA, Saeed F and Riaz M. 2013. Breeding for improving the seed yield and yield contributing traits in *B. napus* L. by using line x tester analysis. *J Plant Breed Genet* **1**: 111-116.
- Katiyar RK, Chomola R, Banerjee PK and Singh HB. 2005. Assessment of heterosis and combining ability for seed yield in Indian mustard [*B. juncea* (L.) Czern. & Coss]. *J Oilseed Brassica* **7**: 33-37.
- Labana KS and Banga SK. 1984. Floral Biology in Indian mustard (*B. juncea* (L.)). *J Genet Agric* **38**: 131-138.
- Maha DC. 2008. Line x tester analysis for combining ability in mustard (*B. juncea*). Thesis (Unpub.) submitted to Dr. PDKV. Akola.
- Meena HS, Ram B, Kumar A, Singh BK, Meena PD, Singh VV and Singh D. 2014. Heterosis and standard heterosis for yield and important traits in *B. juncea*. *J Oilseed Brassica* **5**: 134-140.
- Meena HS, Kumar A, Ram B, Singh VV, Singh BK, Meena PD and Singh D. 2015. Combining ability and heterosis for seed yield and its components in Indian mustard (*B. juncea*). *J Agri Sci Tech* **17**: 1861-1871.
- Rameeh V. 2012. Study on combining ability and heterobelyosis on agronomic traits in spring type of rapeseed varieties. *J Oilseed Brassica* **3**: 43-50.
- Sincik MA, Goksoy T and Turan ZM. 2011. The Heterosis and Combining Ability of Diallel Crosses of Rapeseed Inbred Lines. *Not Bot Horti Agrobo* **39**: 242-248.
- Vaghela PO, Thakhar HS, Sutariya DA, Parmer SK and Prajapati DV. 2011. Heterosis and combining ability for yield and it's component traits in Indian mustard [*B. juncea* (L.) Czern & Coss.]. *J Oilseed Brassica* **2**: 39-43.
- Wassu. 2011. Combining ability and potential heterosis in Ethiopian mustard (*B. juncea* A. Braun). *East African J Sci* **5**: 99-107.