



Studies on combining ability and gene action for yield and quality traits in Baby corn (*Zea mays* L.)

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Abstract: An investigation was carried out to assess the combining ability and nature of gene action in baby corn (*Zea mays* L.) genotypes, by making crosses of eight inbred lines namely, HKI 3209, SML 1, EC 595979, CM 128, VQL 1, G 18 seq C5 F 76-2-2-1-1-2-BBB, HKI 209, in diallel mating design (without reciprocal crosses) by following Griffing Model-I and Method-II during *rabi* season 2013-14 at the experimental farm of BAC, Sabour, Bhagalpur (Bihar). The crosses were evaluated in a randomized complete block design (RCBD) to assess the combining ability and nature of gene action. Based on general combining ability, parents CM 128 and VQL 1 were best parent for breeding programme for yield and quality improvement. The specific combining ability also indicates that hybrid CM128 x VQL1 was best specific combiner for baby corn yield (4.11q/ha), fodder yield (50.91q/ha) as well as for better quality. The crosses CM 128 x HKI 209 recorded the high mean (7.35%) and SCA effects (2.57 °Brix) for quality traits. The gene action analysis also revealed preponderance of non-additive gene action for yield and its contributing characters.

Keywords: Baby corn, Combining ability, Gene action, Quality traits

INTRODUCTION

Maize (*Zea mays* L.), is the third most important cereal crop in the world. It has been cultivated for centuries as a grain crop and more recently it becoming very popular as vegetables, such as baby corn (*Zea mays* var. *saccharata*) (Muthukumar *et al.*, 2005; Mahajan *et al.*, 2007).

Baby corn is very delicious, decorative and nutritious vegetable rich in fibrous protein which is easy to digest and low calorific vegetable. It is grown for its young, fresh, finger like green ears, harvested at the time of silk emergence and before pollination and fertilization (Ramachandrapa *et al.*, 2004). Light yellow colour baby corn ears with regular row arrangement, 10 to 12 cm long and diameter of 1.0 to 1.5 cm are preferred in the market (Muthukumar *et al.*, 2005). India is emerging as one of the potential baby corn producing countries. Among the Indian states, Rajasthan is first in respect of area, where this crop occupies area, production and productivity 10.5 lakh ha, 19.5 lakh tones, 18.6 q ha⁻¹ respectively (Government of Rajasthan, 2010).

The production areas of baby corn (*Zea mays* L.) are still confined to a few countries, including Thailand, Indonesia, India, and Brazil. Baby corn production and improvement on the part of the breeders has resulted in the development of number of varieties. Even though, there was no much works had been done (Chauhan *et al.*, 2010). For exploring the potential yield and good

quality baby corn harvest, we need to explore the standard heterosis. The development of new hybrids for the exploration of heterosis depends on the combining ability of the lines or varieties involved in the production of these hybrids. It is a known fact that selection of parents on the basis of their mean performance does not necessarily lead to desired results (Rai and Asati, 2011). Therefore, devising a sound breeding strategy to improve the yield and quality traits of this crop is of paramount importance. Combining ability analysis help breeders in choosing suitable genotypes as parents for hybridization and superior cross combinations through GCA and SCA studies, respectively (Rodrigues and da Silva, 2002; Rai and Asati, 2011) and at the same time, it also elucidates the nature and magnitude of different types of gene action involved, which is essential for an effective breeding program. Hence, this investigation was undertaken to study the estimates of general and specific combining ability and gene action in baby corn (*Zea mays* L.) for yield components and quality characters.

MATERIALS AND METHODS

Experimental material: The present experiment was carried out at BAC Farm of Bihar Agricultural University, Sabour, Bhagalpur (Bihar), during *rabi* 2013-14 and *kharif* 2014. The eight genetically diverse baby corn inbreds viz., HKI 3209, SML 1, EC 595979, CM

128, VQL 1, G 18 seq C5 F76-2-2-1-1-2-BBB, HKI 209, were crossed in diallel mating design following Model-I, Method-II of Griffing (1956) during *rabi* 2013-14 (Table 1). The commonly used method of combining ability analysis that includes one way crosses and their parents without reciprocal crosses (Singh and Narayanan, 1993).

Experimental methods: The parents and their resulting 28 F₁s were raised in a Randomized Complete Block Design (RCBD) with 3 replications during *kharif* 2014. Each plot consisted 2 rows of each 5 m in length and spacing between rows and plants were 60 cm and 20 cm, respectively. Two seed per hill were planted to ensure the optimum population and recommended package of practices were followed to raise a healthy crop (Crop production guide, 2012). The twelve observations were taken such as 1. Randomly 10 plants basis- Plant height (PHT), Number of leaves per plant (NOL), Number of Baby corn /plant (NBC), Baby corn length (BCL), 2. Plant basis- Days to 50% tasseling (DFT), Baby corn yield (BCY), Fodder yield (FY). 3. Bulk baby corn sample of each entry and each replication basis- Total soluble solid (TSS) was estimated by hand refractometer in Brix, Total sugar (TS), Reducing sugar (RS) and Non reducing sugar (NRS) was determined by Nelson and Somoygi (1944), Crude protein (CP) was estimated by the Micro-Kjeldhal method (AOAC, 1970). Mean data were subjected for analysis of general combining ability (GCA), specific combining ability (SCA) and gene action as per method given by Griffing (1956) (method II and model I) using the software WINDOSTAT(version 7.1).

RESULTS AND DISCUSSION

Analysis of variance for combining ability revealed that mean sum of squares of GCA & SCA for yield and yield contributing trait and quality characters were highly significant for all the characters presented in **Table 1**. Parental lines used for diallel mating and its sources.

S. N.	Parents	Name of the parents	Sources
1	P1	HKI 3209	KARNAL, HISAR
2	P2	SML 1	BAU, SABOUR
3	P3	EC 595979	CIMMYT, MEXICO
4	P4	CM 128	DMR, NEW DELHI
5	P5	VQL 1	ALMORA, UT-TRAKHAND
6	P6	G 18 seq C5 F76-2-2-1-1-2-BBB	CIMMYT, MEXICO
7	P7	HKI 209	KARNAL, HISAR
8	P8	DTPYC-F38-5-2-1-1-2-2-1-B4	CIMMYT, MEXICO
9	C ₁ (Check)	VL Baby corn 1	ALMORA, UT-TRAKHAND
10.	C ₂ (Check)	G5414	ADVANTA INDIA LTD.
11.	C ₃ (Check)	HM4	KARNAL, HISAR

Table 2. Analysis of variance for combining ability analysis for baby corn yield and quality traits.

Sources	df	DFT	PHT	NOL	NBC	BCL	BCY	FY	TS	RS	NRS	TSS	CP
GCA	7	1893.02**	8183.30**	158.11**	6.68**	55.43**	179.39**	66252.99**	101.37**	28.92**	24.79**	66.57**	5.40**
SCA	28	4608.48**	20266.85**	387.19**	17.27**	127.32**	447.15**	187673.88**	225.99**	67.05**	48.71**	177.63**	8.35**
Error	14	0.30	25.42	0.01	0.00	0.12	0.39	767.12	0.006	0.003	0.010	0.01	0.004
Estimate of variance component													
σ²gca		271.55	1208.36	22.91	1.06	37.37	26.78	12142.09	12.46	3.81	2.39	11.10	2.39
σ²sca		4608.18	20241.43	387.18	17.27	650.22	446.76	186906.75	225.98	67.05	48.70	177.62	48.70
σ²e		0.30	25.42	0.01	0.00	0.45	0.39	767.12	0.006	0.002	0.01	0.01	0.01
σ²gca/σ²sca		-0.06	-0.06	-0.06	-0.06	-0.06	-0.06	-0.07	-0.05	-0.05	-0.04	-0.06	-0.04

* Significant at P=0.05

** Significant at P= 0.01

Table 3. Estimates of mean and general combining ability effects (GCA) of the parents for baby corn yield and quality traits.

Parents	DFT		PHT (cm)		NOL		NBC		BCL(cm)		BCY (q/ha)	
	Mean	GCA	Mean	GCA	Mean	GCA	Mean	GCA	Mean	GCA	Mean	GCA
HKI 3209	46.33	-0.21	100.33	0.52	12.37	-0.28**	2.03	-0.09**	6.87	-0.23	6.30	-0.25
SML - 1	47.00	0.96**	81.00	1.28	13.43	0.25**	2.10	-0.04**	8.27	0.26	7.92	0.18
EC 595979	44.33	-0.18	91.33	-0.05	11.92	0.07**	3.10	0.12**	6.18	-0.40	3.93	-0.33
CM 128	45.67	0.36*	94.00	6.32**	13.12	0.21**	3.10	0.12**	8.15	0.42	6.37	0.27
VQL - 1	44.67	-0.04	91.67	2.65**	13.66	0.37**	3.03	0.31**	6.78	-0.04	5.53	1.16**
G 18	45.33	-0.14	71.67	-7.95**	11.23	-0.44**	2.07	-0.18**	7.11	-0.03	4.93	-0.17
HKI 209	43.67	-0.38*	79.67	-1.58	12.24	-0.06*	2.10	-0.27**	7.04	0.36	5.10	-0.53**
DTPYC	43.67	-0.38*	74.67	-1.18	12.10	-0.12**	2.03	0.02	6.30	-0.32	4.90	-0.33
Mean	45.08		85.54		12.51		2.45		7.09		5.62	0.18
SEm±	0.5		5.1		13.34		0.0		0.4		0.4	
CD at 5%	1.4		12.0		0.1		0.09		1.07		1.2	
SE(g) _i ±		0.16		1.49		0.02		0.01		0.10		0.08

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Parents	FY (q/ha)		TS (%)		RS (%)		NRS (%)		TSS (%Brix)		CP (%)	
	Mean	GCA	Mean	GCA	Mean	GCA	Mean	GCA	Mean	GCA	Mean	GCA
HKI 3209	163.33	-13.44	6.82	-2.34**	3.28	-1.10**	3.54	-1.23**	6.82	-1.88**	2.29	-0.01
SML - 1	156.00	-6.17	10.71	0.84**	5.53	-0.29**	5.18	1.14**	10.71	0.09**	2.51	0.15**
EC 595979	182.67	-5.84	9.38	1.01**	4.34	0.36**	5.04	0.66**	9.38	-0.57**	2.66	0.14**
CM 128	176.00	6.76	12.00	0.76**	6.82	0.57**	5.18	0.19**	12.00	0.00	2.54	0.09**
VQL - 1	197.33	-1.01	9.62	-0.06*	5.00	0.08**	4.62	-0.14**	9.62	1.56**	2.01	-0.02
G 18	181.67	4.49	10.87	0.28**	5.32	-0.12**	5.55	0.14**	10.87	0.03	2.35	-0.07**
HKI 209	172.67	4.99	8.33	-0.36**	6.58	0.44**	1.75	-0.18**	8.33	0.29**	2.68	0.09**
DTPYC	222.67	10.23	8.62	-0.14**	5.68	0.08**	2.94	-0.22**	8.62	0.59**	2.29	-0.38**
Mean	181.54		9.54		5.32		4.23		9.54		2.42	
SEm±	9.9		0.078		0.050		0.099		0.166		0.062	
CD at 5%	28.5		0.22		0.14		0.28		0.47		0.17	
SE(g) _i		8.19		0.02		0.01		0.02		0.03		0.01

Table 4. Estimates of mean and specific combining ability effects (SCA) of the hybrids for baby corn yield and quality traits.

Crosses	DFT		PHT		NBL		NBC		BCL		BCY		FY	
	Mean	SCA	Mean	SCA	Mean	SCA	Mean	SCA	Mean	SCA	Mean	SCA	Mean	SCA
HKI 3209 x SML 1	44.00	-1.72**	91.33	-3.15	12.79	0.15*	2.10	-0.38**	7.90	0.24	13.26	-0.42	266.67	21.27
HKI 3209 x EC 595979	44.33	-0.26	98.67	5.51	14.68	1.92**	3.10	0.46**	7.00	0.00	15.89	2.72**	256.67	10.94
HKI3206 x CM128	46.33	1.21**	97.00	-2.52	12.77	-0.14*	2.07	-0.58**	8.78	0.96	13.60	-0.17	289.33	31.01
HKI 3209 x VQL 1	44.33	-0.39	88.00	-7.85*	12.14	-0.95**	4.03	1.20**	5.68	-7.50	17.37	2.71**	267.33	16.77
HKI 3209 x G 18	43.00	-1.62**	81.00	-4.25	12.33	0.08	2.03	-0.31**	7.53	0.16	14.37	0.94*	284.67	28.61
HKI 3209 x HKI 209	44.33	-0.06	98.67	7.05	11.86	-0.77**	2.10	-0.116**	8.18	0.42	13.34	0.36	293.33	36.77
HKI 3209 x DTPYC	43.67	-0.72	84.00	-8.02*	12.68	0.11	3.10	0.56**	7.58	0.50	14.45	1.28**	266.00	4.21
SML 1 x EC 595979	47.00	1.24**	102.67	8.75*	13.27	-0.02	3.13	0.44**	7.19	-0.30	13.51	-0.09	291.33	38.34
SML 1 x CM 128	46.33	0.04	116.33	16.05**	14.93	1.49**	4.10	1.41**	7.38	-0.93	17.28	3.08**	287.33	21.74
SML 1 x VQL 1	46.67	0.78	94.33	-2.29	12.53	-1.07**	2.13	-0.75**	8.34	-5.32	13.59	-1.50**	266.00	8.17
SML 1 x G 18	45.33	-0.46	102.67	16.65**	13.76	0.97**	3.13	0.74**	7.32	-0.54	15.77	2.01**	292.67	29.34
SML 1 x HKI 209	45.67	0.11	87.00	-5.39	12.53	-0.63**	2.10	-0.20**	8.35	0.10	13.44	0.04	298.00	34.17
SML 1 x DTPYC	45.33	-0.22	90.67	-2.12	12.61	-0.49**	2.17	-0.42**	8.27	0.69	13.60	0.00	309.33	40.27
EC 595979 x CM 128	43.33	-1.82**	97.00	-1.95	13.19	-0.06	2.03	-0.82**	7.76	0.11	14.70	1.01*	283.00	17.07
EC 595979 x VQL 1	44.67	-0.09	101.00	5.71	12.22	-1.20**	2.07	-0.98**	8.33	-4.68	13.56	-1.01*	292.67	34.51
EC 595979 x G 18	45.33	0.68	76.33	-8.35*	12.47	-0.13	2.03	-0.52**	7.33	0.13	14.52	1.28**	265.00	1.34
EC 595979 x HKI 209	44.33	-0.09	88.67	-2.39	14.39	1.42**	3.10	0.63**	7.95	0.36	14.54	1.65**	291.00	26.84
EC 595979 x DTPYC	45.33	0.91*	86.67	-4.79	13.39	0.47**	3.03	0.28**	6.79	-0.13	15.01	1.39**	281.67	12.27
CM 128 x VQL 1	44.33	-0.96*	101.00	-0.65	14.65	1.09**	3.10	0.05	9.21	-4.61	19.59	4.41**	321.67	50.91*
CM 128 x G 18	44.67	-0.52	81.33	-9.72*	12.28	-0.46**	2.03	-0.52**	8.20	0.18	13.42	-0.43	310.00	33.74
CM 128 x HKI 209	46.00	1.04	101.33	3.91	12.18	-0.94**	2.10	-0.37**	7.86	-0.55	13.14	-0.35	306.67	29.91
CM 128 x DTPYC	46.00	1.04	115.33	17.51**	12.64	-0.42**	3.10	0.35**	7.40	-0.33	13.45	-0.24	302.67	20.67
VQL 1 x G 18	45.00	0.21	99.00	11.61**	13.49	0.58**	3.10	0.35**	7.14	-0.43	15.87	1.14*	299.33	30.84
VQL 1 x HKI 209	45.67	1.11	92.33	-1.42	14.83	1.54**	3.10	0.44**	8.32	0.36	16.75	3.37**	283.33	14.34
VQL 1 x DTPYC	44.33	-0.22	101.67	7.51	13.32	0.10	3.03	0.09**	7.79	0.51	14.54	-0.03	250.00	-24.23
G 18 x HKI 209	44.00	-0.46	83.67	0.51	12.37	-0.10	2.10	-0.07*	9.55	1.59	13.58	-0.54	300.00	25.51
G 18 x DTPYC	45.33	0.88*	87.33	3.78	13.22	0.80**	3.13	0.68**	7.11	-0.18	15.44	2.20**	315.00	35.27
HKI 209 x DTPYC	43.67	-0.56	107.33	17.41**	13.50	0.71**	2.03	-0.33**	8.01	0.33	14.20	1.31**	317.33	37.11
Mean	44.94		95.94		13.11		2.66		7.79		14.71		288.86	
SEm ±	0.5		5.1		0.1		0.0		0.4		0.6		9.9	
CD at 5%	1.4		12.0		0.2		0.09		1.07		1.5		28.5	
SOij		0.43		3.97		0.06		0.03		0.28		0.49		21.85

Table 5. Mean performance of crosses for quality traits.

Crosses	Total sugar (%)			Reducing sugar (%)			Non reducing sugar (%)			Crude protein (%)			Total soluble solid (°brix)		
	Mean	SCA	SE	Mean	SCA	SE	Mean	SCA	SE	Mean	SCA	SE	Mean	SCA	SE
HKI 3209 x SML 1	7.32	1.05**	0.05	2.99	-0.93**	0.06	4.33	-0.13	0.05	2.06	-0.45**	0.05	4.63	-1.48**	0.05
HKI 3209 x EC 595979	10.87	2.33**	0.07	6.53	1.97**	0.06	4.34	0.35**	0.05	2.41	-0.06	0.05	9.80	-5.43**	0.05
HKI3206 x CM 128	6.58	-1.71**	0.05	3.81	-0.96**	0.05	2.77	-0.74**	0.05	2.43	0.01	0.05	6.07	0.20*	0.05
HKI 3209 x VQL 1	6.56	-0.91**	0.05	4.26	-0.02	0.05	2.29	-0.89**	0.05	2.97	0.66**	0.05	8.63	1.01**	0.05
HKI 3209 x G 18	7.14	-0.67**	0.05	4.07	-0.01	0.05	3.07	-0.66**	0.05	2.41	0.14**	0.05	7.90	1.88**	0.05
HKI 3209 x HKI 209	6.87	-0.29**	0.05	4.46	-0.18**	0.05	2.41	-0.11	0.05	2.39	-0.14**	0.05	8.73	2.47**	0.05
HKI 3209 x DTPYC	6.43	-0.96**	0.05	4.06	-0.23**	0.05	2.38	-0.73**	0.05	1.74	-0.21**	0.05	4.70	-1.89**	0.05
SML 1 x EC 595979	10.00	-1.72**	0.05	5.43	0.06	0.05	4.57	-1.78**	0.05	2.58	-0.06	0.05	9.73	2.34**	0.05
SML 1 x CM 128	13.16	1.69**	0.05	4.31	-1.28**	0.05	8.85	2.97**	0.05	2.95	0.37**	0.05	7.33	-0.52**	0.05
SML 1 x VQL 1	11.36	0.72**	0.05	5.95	0.86**	0.05	5.41	-0.14	0.05	2.41	-0.07	0.05	9.60	0.07	0.05
SML 1 x G 18	12.50	1.51**	0.05	5.21	0.32**	0.05	7.29	1.20**	0.05	2.45	0.02	0.05	7.13	-0.87**	0.05
SML 1 x HKI 209	10.57	0.22**	0.05	4.46	-0.99**	0.05	6.10	1.22**	0.05	2.49	-0.10*	0.05	7.40	-0.86**	0.05
SML 1 x DTPYC	10.87	0.30**	0.05	5.43	0.34**	0.05	5.43	-0.04	0.05	2.64	0.52**	0.05	10.87	2.31**	0.05
EC 595979 x CM 128	11.28	-0.36**	0.05	6.94	0.71**	0.05	4.34	-1.07**	0.05	2.58	0.01	0.05	7.43	0.24**	0.05
EC 595979 x VQL 1	10.72	-0.10	0.05	5.95	0.21**	0.05	4.77	-0.31**	0.05	2.18	-0.28**	0.05	8.70	-0.17*	0.05
EC 595979 x G 18	11.20	0.04	0.05	6.41	0.87**	0.05	4.79	-0.83**	0.05	3.04	0.62**	0.05	5.90	-1.43**	0.05
EC 595979 x HKI 209	12.94	2.42**	0.05	5.95	-0.15**	0.05	6.99	2.57**	0.05	3.04	0.46**	0.05	7.63	0.03	0.05
EC 595979 x DTPYC	13.16	2.42**	0.05	5.43	-0.31**	0.05	7.72	2.72**	0.05	1.33	-0.78**	0.05	11.90	3.71**	0.05
CM 128 x VQL 1	10.57	0.00	0.05	6.25	0.29**	0.05	4.32	-0.29**	0.05	2.35	-0.07	0.05	11.40	2.07**	0.05
CM 128 x G 18	10.87	-0.03	0.05	4.46	-1.29**	0.05	6.41	1.26**	0.05	1.85	-0.52**	0.05	7.47	-0.33**	0.05
CM 128 x HKI 209	9.04	-1.22**	0.05	7.35	1.03**	0.05	1.69	-2.26**	0.05	2.43	-0.09	0.05	10.63	2.57**	0.05
CM 128 x DTPYC	10.87	0.39**	0.05	6.70	0.74**	0.05	4.17	-0.36**	0.05	2.29	0.23**	0.05	6.63	-1.71**	0.05
VQL 1 x G 18	10.87	0.78**	0.05	5.10	-0.16**	0.05	5.77	0.95**	0.05	2.24	-0.02	0.05	7.20	-2.28**	0.05
VQL 1 x HKI 209	9.26	-0.18**	0.05	6.25	0.42**	0.05	3.01	-0.60**	0.05	2.45	0.03	0.05	6.10	-3.64**	0.05
VQL 1 x DTPYC	9.62	-0.05	0.05	4.81	-0.66**	0.05	4.81	0.61**	0.05	2.29	0.34**	0.05	8.97	-1.06**	0.05
G 18 x HKI 209	8.93	-0.85**	0.05	5.21	-0.42**	0.05	3.72	-0.44**	0.05	2.58	-2.81**	0.05	5.07	-3.14**	0.05
G 18 x DTPYC	8.33	-1.67**	0.05	5.43	0.17**	0.05	2.90	-1.85**	0.05	1.17	-0.75**	0.05	8.53	0.04	0.05
HKI 209 x DTPYC	10.87	1.51**	0.05	5.32	-0.50**	0.05	5.55	2.01**	0.05	1.29	-0.77**	0.05	10.63	1.87**	0.05
Mean	9.96			5.31			4.65			2.32			8.10		
SEm ±	0.07			0.05			0.09			0.06			0.16		
CD at 5%	0.22			0.14			0.28			0.17			0.47		
SE(ij)		0.06			0.04			0.08			0.05				0.09

Table 2 under studied. The mean squares of SCA were larger than those of GCA in all the characters. It was also revealed that preponderance of non-additive gene action involved in control of the characters. The importance of non additive gene actions were also confirmed by the findings of Anantha (2004) and Selvarani (2007) for days to tasseling, Geetha and Jayaraman (2000), Anantha (2004) and Prakash and Ganguli (2004) for plant height, Jayakumar and Sundaram (2007) for number of leaves per plant, Rodrigues and da Silva (2002) for baby corn length and Suneetha *et al.* (2000) for non reducing sugars in baby corn which is in agreement with our finding.

The high positive estimates of specific combining ability in absolute values indicates that hybrid performance is relatively superior or inferior to parent lines general combining ability, showing the importance of additive interactions resulting from the complementation degree among parent lines in relation to frequency of alleles in loci with some dominance, while low estimates of specific combining ability in absolute value indicates that hybrids behave as expected in relation to general combining ability of parent lines (Vencovsky and Barriga, 1992). In the selection of parent lines used to produce hybrids, the effect of a specific combining ability analyzed in an isolated way has a limiting value. Thus, other parameters should be considered such as and general combining ability of the respective parent lines (Oliveira *et al.*, 1998). The parent CM 128 and VQL 1 under present investigation were the best among eight parents shows desirable mean and GCA effect for yield, yield attributing traits and quality traits. So, these parents could be used extensively in hybrid breeding programme to increase baby corn yield and quality. Similar to present investigation were also reported by Dhasarathan *et al.* (2012) in baby corn for selection of suitable parental line for exploiting heterosis.

Therefore, superior hybrid combinations, which are important for breeding, are involved with at least one parental line which has the most favorable effects of general combining ability (Cruz and Regazzi, 1997). Thus, it is possible to analyze the hybrids that showed high performance for most of the yield (baby corn yield and fodder yield) and quality traits (T.S.S, T.S) such as CM128 x VQL1 and VQL1 x HKI209 (Table 4). Baby corn yield per plot in the hybrid CM128 x VQL1 is associated with high effects of general combining ability of both the parent lines. Therefore, in this case, the high productivity is not due to dominant genetic effects of inbred but due to additive effects. In the VQL1 x HKI209 hybrid is associated with high effect of the general combining ability of the VQL1 (BCY) and HKI209 (FY) inbred with one of highest effect of the estimated specific combining ability, since HKI209 inbred showed lower general combining ability. In this case, the participation of specific combining ability is significant for yield, contribution almost to

the general combining ability from both inbreds, regarding the dominance and epistasis effect (Gardner, 1963). However, the hybrid G18 x HKI209 and SML1 x CM128 exhibited highest specific combining ability for baby corn length and number of baby corns per plant and CM 128 x HKI 209 (reducing suger and total soluble solid) showed highest specific combining ability effect. About 5 hybrids exhibited higher mean performance and SCA effects for baby corn yield per plot. In respect of baby corn superiority, it is decided by its quality. Hence, the crosses CM128 x VQL1 and CM 128 x HKI 209 recorded the high mean and SCA effects for yield and quality traits of baby corn.

Conclusion

The present study concluded that CM128 & VQL1 were the best among the all eight parents as it showed desirable mean and GCA effects for most of the yield, yield contributing traits and quality traits of baby corn ear, respectively. Therefore these genotypes could be used extensively in hybrid breeding program with a view to increase baby corn yield with quality traits. Furthermore, based on mean and SCA effects, the one hybrid CM128 x VQL1 was found suitable for baby corn yield with better quality. For varietal improvement, these crosses could also be utilized for exploiting promising recombinants and it could be useful towards enhancing baby corn yield and quality.

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