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## Elucidation of combining ability for earliness and popping expansion volume in popcorn (*Zea mays* var. *everta*) across different environments

**Munnesh Kumar, JP Shahi, Ashok Singhamsetti, Anima Mahato, Kumari Shikha and Kartik Madankar**

### Abstract

Maize (*Zea mays* L.,  $2n = 20$ ), an economic crop, is a widely cross-pollinated annual, monoecious Poaceae family member native to Mexico and Central Africa. All twenty characters studied showed dominance gene action, except for traits like ear diameter and moisture content. The number of gene groups varied from one to six. Gene groups were involved and showed partial or over-dominance. Based on GCA and SCA studies, two good general combiners are identified as HKIPC-7 and WINPOP-29. Specific combiners for yield and popping characteristics are the WINPOP-47×HKIPC-7 cross for yield per plant, the WINPOP-8×HKIPC-7 and WINPOP-3×HKIPC-7 crosses for earliness and also the WINPOP-47×HKIPC-5 cross for popping expansion. G×E interaction is clearly indicated as narrow sense heritability is less than unity. In either the pedigree method or the bulk method, it is best to choose parents with dominant alleles for traits that affect yield.

**Keywords:** Popcorn, combining ability, popping volume, genetic parameters

### Introduction

Maize (*Zea mays* L.,  $2n = 20$ ) is the most diversified and versatile crop among cereal crops. It is an annual, monoecious, cross-pollinated plant with a native range of Mexico and Central Africa. It was domesticated from a wild maize ancestor (*Teosinte*) and is referred to as the "queen of cereals" due to its vast genetic base and high yield potential. In India, a highly populous country in which rice and wheat are staple food crops, followed by maize, which is the 3<sup>rd</sup> in area and 7<sup>th</sup> in production among corn-cultivated countries, accounting for approximately 4% of the global area and 2% of production (DACNET-2020) [1]. India's maize cultivation surpassed 9.2 million hectares in 2018-19 [1]. In 1950-51, it yielded 1.73 million metric tonnes (MT), which has climbed to 27.8 million metric tonnes (MT) in 2018-19, an increase of almost 16-fold (Sandhu, *et al.*, 2007) [2].

Popcorn is an extremely unique form of flint corn with a very hard endosperm and a low portion of soft starch and is characterised by its popping ability on heating the kernel, which is a unique quality of the endosperm (Acquaah, 2006) [3]. Popcorn swells and puffs up when heated and is a popular recreational snack with a high nutritional content (Rakshit, *et al.*, 2003) [4]. It is a globally popular snack because it is an excellent source of carbohydrates, energy and fibre. It's a healthy, tasty food that helps digestion by giving the body the fibre it needs (Rodvalho, *et al.*) [5]. Indian popcorn cultivars have a lower popping ratio as compared to European and American cultivars. The high demand for popcorn has spurred research to discover traits that dictate its idiosyncratic popping ability and many attempts have been made to develop a popcorn industry, but it has been a hurdle because of the absence of stable and optimal hybrids with high popping volume and efficiency (Larish, and Brewbaker, 1999) [6] and also a lack of superior popcorn germplasm for popping characteristics, including popping volume, flake size, popping percent and high-yield (Robbins, and Ashman, 1984 and Dofing, 1999) [7, 8] due to narrow genetic background because most of the popcorn lines are descended from flint germplasm (Kantety, *et al.*, 1995) [9]. However, it has relatively poor agronomic, yield and popping quality and also more susceptible and prone to disease and pests. These research, focused on designing new single-cross popcorn hybrids that are stable, uniform, early-maturing, with higher yields and better popping quality than composite cultivars. They need to be bred for a long time. The development of single-cross popcorn hybrids requires a proper road map to understand the genetic architecture of popcorn based on

the underlying diallel mating design in the breeding programme. The main object of this manuscript "Elucidation of combining ability, genetic effects for controlling certain traits and deciphering genetic component variance for yield and popping quality traits across different environment".

### Materials and Methods

The experimental genetic material was comprised of all the available popcorn inbred lines. The selected promising nine inbred line were planted in a crossing block during Rabi 2017 and all possible single cross combination carried out and their successive progenies are maintained by selfing and sib mating techniques. The thirty -six crosses thus formed along with the 9 parents popcorn inbreds and one standard check (Amber popcorn) was planted in completely randomized block design with three replication and each entry was planted in single row plot of 4m length maintaining in each season the inter-row spacing at 60 cm and Plant to plant spacing of 25 cm during *rabi*, 2017-18, *kharif*-2018 and *rabi* 2018-19 seasons.

Sr. No.	Seasons and Year	Code No.
1.	Rabi, 2017-18	E1
2.	Khariif,2018	E2
3.	Rabi,2018-19	E3

### Phenotypic data collection

Five randomly selected plants from each entry in each replication are recorded based on underlying agro-morphological, yield and quality traits, whereas for kernel architectural and quality traits, data is recorded from one selfed plant from each entry in each replication. These data were recorded for twenty different agro-morphological, yield attributing and popping traits, viz.: number of leaves per plant, leaf area, plant height (cm), ear height (cm), primary branches per tassel, days to 50% anthesis, days to 50% silking, days to 75% maturity, number of ears per plant, ear length (cm), ear diameter (cm), number of kernels per ear, number of kernels per row, hundred seed weight (gm), grain yield per plant (g), moisture content, popping expansion volume, flake size, flake volume and popping rate (%). Mean over replications, in each season /environment, were used for the statistical analysis. Analysis of data was done with the help of statistical software Windostat V.8.5. The procedure for analysis of variance for the crosses and parents was followed as suggested (Panse and Sukhatme, 1978) [10]. The detailed procedure of both graphical and component analysis is done in according (Jinks, 1954 and Hayman, 1954a) [11, 12]. Statistical analysis in order to estimate the general and specific combining ability (GCA and SCA) variances and effects, as described (Griffing, 1956a) [13].

### Result and Discussion

All the twenty characters studied showed dominance gene action as  $H_1$  is greater than D and leaf area, plant height, ear height, days to 50% silking, ear per plant, ear length, kernels per row, popping expansion volume, flake size, flake volume and popping rate characters contributed majorly by dominance gene action, hence selection lines based on these traits should be carried out in later generation rather than early segregating generation. Traits like ear diameter and moisture content are majorly contributed by additive gene action as D is greater than  $H_1$ . Number of gene groups varied around one to 6 gene groups are involved in controlling these twenty

characters except traits like traits like moisture content and popping expansion volume controlled by a single gene in all three environment, similarly days to 50% silking, physiological maturity, ear diameter and hundred seed weight in E2 and ear per plant in E1, E2 are controlled by single gene as these traits recorded  $h^2/H_2$  less than unity. Narrow sense heritability is measure of heritability as well as the contribution of genetic variance and environmental variance.

All the twenty characters studied recorded less than unity for narrow sense heritability indicating the phenotypic variance for these characters is mainly attributed by environmental influence rather than solely genotypic variance. This findings is congruity with (El-Moula, *et al.*, 2004) [14] for days to 50% silking and ear height, Kalla, *et al.*, (2001) [15] for ear length and hundred seed weight, (Kanagarasu, *et al.*, 2010) [16] for grain yield per plant, ear diameter, cob length, plant height, ear height, leaf length, 100-grain weight, grain rows per cob, days to 50 per cent tasseling and days to 50 per cent silking (Rameeh, *et al.*, 2000) [17] for number kernels per row, (Vijayabharathi, *et al.*, 2009a) [18] for all the characters studied reported non- additive gene action.

Among all the parents HKIPC-7, HKIPC-8-3 can exploited for the yield per plant, number of kernel row per ear, ear diameter and ear length, as both parents showed significant high positive GCA effect for respective characters. Parents HKIPC-7, WINPOP-29 can exploited for developing early maturing inbred lines as these parents showed highest negative GCA effect for days to 50% anthesis, days to 50% silking and physiological maturity. Parents HKIPC-7, WINPOP-47 can exploited for developing more number of leafs per plant and leaf area in a inbred lines as these parents showed highest positive GCA effect for days to 50% anthesis, days to 50% silking and physiological maturity. WINPOP-47 for popping expansion volume, WINPOP-29 and HKIPC-5 for popping rate, WINPOP-3 for flake volume, WINPOP-8 and HKIPC-8-3 for flake volume. WINPOP-13 and WINPOP-29 parents can be exploited for developing dwarf inbred lines as these parents showed highest negative GCA effect.

**SCA:** Based on SCA effect HKI PC-5×HKIPC-7 cross can utilized to obtain desirable segregants for yield per plant and number of leafs per plant.

The WINPOP-8×HKIPC-7, WINPOP-3×HKIPC-7 cross can utilized to obtain desirable segregants for hundred seed weight and earliness. WINPOP-3×HKIPC-8-3 cross can utilized to obtain desirable segregants for number of kernels per ear, ear diameter and ear length. WINPOP-3×WINPOP-29 cross can utilized to obtain desirable segregants for flake volume, ear per plant and earliness. WINPOP-47×HKIPC-5 cross can utilized to obtain desirable sergeants for popping expansion volume and number of leafs per plant. WINPOP-8×WINPOP-47 cross can utilized to obtain desirable segregants for ear per plant and primary branches per tassel. This findings are congruity with reported that a cross combination which were of SCA high x SCA high and high SCA x low GCA parents exhibited greater heterosis (Singh, *et al.*, 2002) [19] found out that two popcorn combinations with low GCA x high GCA (Moterle, *et al.* 2012) [20] reported one popcorn cross combination with high GCA x high GCA as good specific combiners with higher *per se* performance (Vieira, *et al.*, 2011) [21].

**Table 1:** ANOVA of 9x9 half diallel design (excluding reciprocal) for twenty different characters studied under *rabi* 2017-18 (E1), *kharif*: 2018 (E2) and *rabi*, 2018:19 (E3)

Source of variation	Env.	DF	Traits									
			LP	LA	PBPT	PHT	EHT	DFA	DFS	PM	EPP	EL
Replicates	E1	2	2.107 *	1475.514 ***	16.647 ***	491.613 **	21.037 *	3.030	2.696	2.467	0.094 **	3.858
	E2	2	2.590 *	2088.553	26.912 ***	1686.489 ***	2092.368 ***	7.385	4.452	14.319 **	0.062 **	2.107
	E3	2	9.286 ***	677.896 *	101.775 ***	1140.234 ***	337.484 ***	4.289	4.356	6.200	0.257 ***	4.050
Treatments	E1	44	4.058 ***	4722.470 ***	22.972 ***	1420.262 ***	370.603 ***	28.954 ***	30.038 ***	30.294 ***	0.121 ***	10.668 ***
	E2	44	2.730 ***	5021.721 ***	18.357 ***	997.494 ***	314.309 ***	9.520 ***	13.030 ***	4.963 *	0.045 ***	13.374 ***
	E3	44	3.870 ***	4230.461 ***	26.621 ***	1604.018 ***	565.045 ***	26.355 ***	29.767 ***	32.418 ***	0.038 *	10.404 ***
Parents	E1	8	0.851	2291.461 ***	1.429	669.487 ***	122.302 ***	11.750 ***	11.426 ***	11.593 ***	0.052 *	3.111 *
	E2	8	2.127 ***	2912.433 *	12.455 ***	391.898 ***	176.141 **	0.954	3.500	1.870	0.018	5.059
	E3	8	1.092	1541.256 ***	6.416	797.579 ***	190.507 ***	8.787 **	9.917 **	11.759 ***	0.005	3.096 *
Hybrids	E1	35	3.516 ***	3003.657 ***	13.408 ***	571.720 ***	226.139 ***	12.580 ***	11.470 ***	11.470 ***	0.123 ***	3.263 **
	E2	35	1.741 ***	3302.508 ***	12.034 ***	565.659 ***	222.861 ***	10.092 ***	14.174 ***	5.588 **	0.050 ***	8.283 **
	E3	35	3.021 ***	2651.947 ***	15.525 ***	564.376 ***	382.947 ***	12.580 ***	11.686 ***	11.470 ***	0.035 *	3.202 **
Parent Vs. Hybrids	E1	1	48.672 ***	84329.010 ***	530.086 ***	37125.440 ***	7413.261 ***	739.674 ***	828.817 ***	838.757 ***	0.626 ***	330.317 ***
	E2	1	42.196 ***	82068.480 ***	286.891 ***	20956.470 ***	4620.330 ***	58.017 ***	49.202 ***	7.824	0.063 *	258.061 ***
	E3	1	55.809 ***	80992.100 ***	576.600 ***	44443.000 ***	9934.783 ***	649.007 ***	821.400 ***	930.891 ***	0.400 ***	320.937 ***
Error	E1	88	0.449	38.722	1.013	72.240	5.180	2.507	2.318	2.292	0.019	1.487
	E2	88	0.563	1315.545	3.010	91.012	64.447	2.537	1.793	2.788	0.010	4.055
	E3	88	0.743	169.145	5.581	102.734	24.729	2.827	2.787	2.745	0.022	1.449
Total	E1	134	1.659	1598.114	8.457	521.133	125.406	11.199	11.425	11.490	0.054	4.537
	E2	134	1.305	2544.035	8.406	412.476	176.759	4.902	5.522	3.674	0.022	7.086
	E3	134	1.897	1510.305	13.925	611.178	206.814	10.574	11.670	12.540	0.031	4.428
GCA	E1	8	2.418 ***	1798.982 ***	6.330 ***	534.881 ***	146.890 ***	9.629 ***	8.621 ***	8.722 ***	0.024 ***	1.401 **
	E2	8	1.028 ***	2091.958 ***	7.165 ***	304.048 ***	112.536 ***	4.631 ***	5.945 ***	1.414	0.017 ***	0.500
	E3	8	2.356 ***	1146.063 ***	10.722 ***	617.136 ***	251.258 ***	9.116 ***	8.760 ***	9.111 ***	0.008	1.342 **
SCA	E1	36	1.116 ***	1524.196 ***	7.952 ***	459.763 ***	118.344 ***	9.656 ***	10.322 ***	10.404 ***	0.044 ***	4.035 ***
	E2	36	0.884 ***	1581.007 ***	5.887 ***	338.820 ***	103.044 ***	2.849 ***	3.987 ***	1.708 *	0.014 ***	5.337 ***
	E3	36	1.053 ***	1468.841 ***	8.463 ***	516.348 ***	174.369 ***	8.711 ***	10.180 ***	11.183 ***	0.014 *	3.940 ***
Error	E1	88	0.150	12.907	0.338	24.080	1.727	0.836	0.773	0.764	0.006	0.496
	E2	88	0.188	438.515	1.003	30.337	21.482	0.846	0.598	0.929	0.003	1.352
	E3	88	0.248	56.382	1.860	34.245	8.243	0.942	0.929	0.915	0.007	0.483
GCA/SCA Ratio	E1		0.213	0.107	0.072	0.107	0.113	0.091	0.075	0.075	0.043	0.023
	E2		0.110	0.132	0.115	0.081	0.101	0.172	0.143	0.057	0.115	-0.019
	E3		0.238	0.070	0.122	0.110	0.133	0.096	0.077	0.073	0.007	0.023
Predictability Ratio	E1		0.299	0.177	0.125	0.176	0.185	0.153	0.130	0.131	0.079	0.044
	E2		0.180	0.208	0.187	0.139	0.169	0.256	0.223	0.102	0.187	-0.040
	E3		0.322	0.123	0.196	0.180	0.210	0.161	0.133	0.127	0.015	0.043

\* $\leq 0.05$ , \*\* $\leq 0.01$ , R-18- *rabi*, 2017-18, K-18- *kharif*-2018, R-19- *rabi*, 2018-19, DF- Degree of freedom, LPP-No. of leaves per plant; LA- Leaf area; PBPT- No. of primary branches per tassel; EHT-Ear height; DFA- Days to 50% anthesis; DFS- Days to 50% silking, DM-days to 75% maturity, EPP-No. of Ears per plant, EL- Ear length, ED-Ear diameter, KRPE- No. of kernels rows per ear, KPR- No. of kernels per row, YPP- Yield per plant, HSW- Hundred seed weight, MC- Moisture content, PEV-Popping expansion volume, FS- Flake size, FV- Flake volume, PR- Popping rate

**Table 2:** ANOVA of 9x9 half diallel design (excluding reciprocal) for twenty different characters studied under *rabi* 2017-18 (E1), *kharif*: 2018 (E2) and *rabi*, 2018:19 (E3)

Source of variation	DF	traits										
		ED	KRPE	KPR	YPP	HSW	MOI	PEV	FZ	FV	PR	
Replicates	E1	2	0.021	0.647	25.697	63.379	0.087	1.369	0.291	0.036	438.337	15.814
	E2	2	0.292	3.941	17.783	27.282	2.987	4.298 *	0.431	0.038	58.454	15.814
	E3	2	0.022	1.443	30.561	74.658	2.681	4.241 **	0.128	0.022	51.526	27.528
Treatments	E1	44	0.249 ***	3.951 ***	74.112 ***	2292.279 ***	11.069 ***	1.264 ***	12.876 ***	0.514 ***	5558.506 ***	110.186 ***
	E2	44	0.375 ***	3.848 *	30.873 ***	1821.063 ***	3.416 ***	3.636 ***	14.791 ***	0.875 ***	8761.368 ***	110.186 ***
	E3	44	0.218 ***	4.033 ***	76.874 ***	2310.482 ***	15.059 ***	1.790 **	14.547 ***	0.713 ***	7835.882 ***	107.879 ***
Parents	E1	8	0.403 ***	6.196 ***	23.318	128.659 **	3.198 *	0.975	4.944 ***	0.227 ***	676.957 ***	70.055 ***
	E2	8	0.027	1.148	13.794 *	96.329 **	1.536	2.494 *	19.429 ***	0.300 ***	5168.207 ***	70.055 ***
	E3	8	0.377 ***	6.193 ***	22.378	260.347 **	3.972 **	5.008 ***	6.616 ***	0.408 ***	575.006	59.929
Hybrids	E1	35	0.205 ***	1.755	12.947	1600.284 ***	8.021 ***	1.339 ***	12.244 ***	0.234 ***	3307.802 ***	37.698 ***
	E2	35	0.276 ***	3.838 *	20.359 ***	1190.019 ***	3.799 ***	3.921 ***	11.136 ***	0.702 ***	6263.460 ***	37.698 ***
	E3	35	0.179 ***	1.806	16.462	1608.156 ***	8.108 ***	1.101	14.939 ***	0.406 ***	4630.131 ***	68.767 **
Parent Vs. Hybrids	E1	1	0.574 ***	62.853 ***	2621.262 ***	43821.040 ***	180.706 ***	0.963	98.475 ***	12.607 ***	123385.600 ***	2968.301 ***
	E2	1	6.609 ***	25.785 **	535.489 ***	37705.540 ***	5.069 *	2.788	105.638 ***	11.508 ***	124933.400 ***	2968.301 ***
	E3	1	0.348 *	64.688 ***	2627.258 ***	43292.950 ***	347.042 ***	0.160	64.274 ***	13.888 ***	178124.200 ***	1860.380 ***
Error	E1	88	0.082	1.342	12.131	41.959	1.238	0.527	0.626	0.018	181.913	13.311
	E2	88	0.104	2.456	6.625	27.875	1.186	0.963	0.483	0.026	145.508	13.311
	E3	88	0.076	1.396	12.210	71.888	1.445	0.852	0.566	0.032	283.934	32.085
Total	E1	134	0.136	2.188	32.685	781.190	4.449	0.782	4.643	0.181	1951.189	45.158
	E2	134	0.196	2.935	14.753	616.674	1.945	1.890	5.180	0.305	2973.297	45.158
	E3	134	0.122	2.262	33.717	806.990	5.933	1.210	5.150	0.255	2760.209	56.904
GCA	E1	8	0.331 ***	1.891 ***	12.167 **	939.305 ***	4.611 ***	1.207 ***	3.165 ***	0.053 ***	572.545 ***	31.402 ***
	E2	8	0.133 ***	1.180	7.985 **	545.521 ***	1.158 **	1.747 ***	5.253 ***	0.052 ***	1567.462 ***	31.402 ***

	E3	8	0.261 ***	1.773 ***	9.486 *	924.469 ***	4.358 ***	2.033 ***	3.433 ***	0.238 ***	378.864 ***	29.881 **
SCA	E1	36	0.028	1.190 ***	27.490 ***	725.157 ***	3.485 ***	0.247	4.543 ***	0.198 ***	2137.344 ***	37.912 ***
	E2	36	0.123 ***	1.305 *	10.803 ***	620.687 ***	1.134 ***	1.093 ***	4.859 ***	0.345 ***	3221.121 ***	37.912 ***
	E3	36	0.031	1.249 ***	29.211 ***	735.870 ***	5.167 ***	0.277	5.164 ***	0.238 ***	3108.205 ***	37.310 ***
Error	E1	88	0.027	0.447	4.044	13.986	0.413	0.176	0.209	0.006	60.638	4.437
	E2	88	0.035	0.819	2.208	9.292	0.395	0.321	0.161	0.009	48.503	4.437
	E3	88	0.025	0.465	4.070	23.963	0.482	0.284	0.189	0.011	94.645	10.695
GCA/SCA Ratio	E1		42.908	0.177	0.031	0.118	0.124	1.318	0.062	0.022	0.022	0.073
	E2		0.102	0.067	0.061	0.080	0.094	0.168	0.099	0.012	0.044	0.073
	E3		3.888	0.152	0.020	0.115	0.075	3.498	0.059	0.091	0.009	0.066
Predictability Ratio	E1		0.988	0.261	0.059	0.191	0.199	0.725	0.110	0.043	0.043	0.128
	E2		0.169	0.119	0.109	0.138	0.158	0.251	0.165	0.023	0.080	0.128
	E3		0.886	0.233	0.038	0.187	0.131	1.022	0.106	0.154	0.017	0.116

\*≤0.05, \*\*≤0.01, R-18- rabi, 2017-18, K-18- kharif-2018, R-19- rabi,2018-19, DF- Degree of freedom, LPP-No. of leafs per plant; LA- Leaf area; PBPT- No. of primary branches per tassel; EHT-Ear height; DFA- Days to 50% anthesis; DFS- Days to 50% silking, DM-days to75% maturity, EPP-No. of Ears per plant, EL- Ear length, ED-Ear diameter, KRPE- No. of kernels rows per ear, KPR- No. of kernels per row, YPP- Yield per plant, HSW- Hundred seed weight, MC- Moisture content, PEV-Popping expansion volume, FS- Flake size, FV- Flake volume, PR- Popping rate

**Table 3:** General Combining Ability of nine parents studied under rabi, 2017:18 (E1), kharif- 2018 (E2) and rabi, 2018-19 (E3)

Traits / Parents	Number of leafs per plant						Leaf area					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	12.90	-0.371 **	11.00	-0.270 *	12.90	-0.313 *	204.76	-8.972 ***	221.38	-6.108	211.47	-7.304 ***
2 Parent	13.23	-0.538 ***	12.17	0.224	11.90	-0.701 ***	206.99	2.904 **	210.67	2.781	203.37	1.008
3 Parent	12.63	-0.289 *	12.33	-0.094	12.63	-0.250	246.89	16.164 ***	260.33	17.897 **	230.40	9.623 ***
4 Parent	12.16	-0.175	11.00	-0.170	12.17	-0.098	195.18	-8.510 ***	204.75	-8.803	195.43	-6.480 **
5 Parent	12.00	-0.520 ***	11.00	-0.280 *	12.00	-0.429 **	203.78	-6.802 ***	206.00	-7.473	203.77	-4.635 *
6 Parent	13.00	0.547 ***	12.50	0.242	13.00	0.477 **	243.67	7.569 ***	250.17	6.777	235.50	10.299 ***
7 Parent	13.55	0.404 ***	10.63	-0.377 **	13.57	0.368 *	155.47	-22.183 ***	155.50	-25.307 ***	158.13	-18.783 ***
8 Parent	13.45	0.610 ***	12.67	0.512 ***	13.47	0.641 ***	222.53	16.751 ***	232.00	18.231 **	220.23	12.123 ***
9 Parent	12.90	0.333 **	10.67	0.212	12.87	0.305 *	195.27	3.079 **	195.00	2.005	203.87	4.150
Gi <math>\hat{C}</math> 0 at 95%	0.254 ***		0.284 ***		0.326 ***		2.355 ***		13.727 ***		4.922 ***	
Gi--Gj at 95%	0.380 ***		0.426 ***		0.489 ***		3.533 ***		20.591 ***		7.383 ***	
GCA/SCA Ratio	0.213		0.110		0.238		0.107		0.132		0.070	
Predictability Ratio	0.299		0.180		0.322		0.177		0.208		0.123	

**Table 4:** General Combining Ability of nine parents studied under rabi, 2017:18 (E1), kharif- 2018 (E2) and rabi, 2018-19 (E3)

Traits / Parents	Number of Primary branches per tassel						Plant height					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	15.17	-0.231	11.67	0.265	14.10	-0.624	89.67	0.153	83.33	-3.465 *	78.90	-1.323
2 Parent	14.62	-0.654 ***	12.80	0.744 *	14.90	-0.757	94.00	-3.288 *	96.67	3.232 *	82.00	-4.687 **
3 Parent	15.30	-1.302 ***	15.33	0.144	13.00	-1.727 ***	72.00	-8.905 ***	108.33	0.263	62.00	-9.935 ***
4 Parent	13.92	0.118	10.07	0.935 **	13.90	0.152	69.82	-4.370 **	88.33	1.505	66.70	-4.156 *
5 Parent	14.68	-0.370 *	11.00	-0.856 **	14.20	-0.487	65.67	-8.502 ***	85.00	-6.253 ***	65.67	-7.696 ***
6 Parent	16.05	0.599 ***	12.67	-0.129	15.90	0.858 *	84.11	3.968 **	95.00	-0.859	84.10	4.589 **
7 Parent	15.94	1.339 ***	9.67	-0.977 ***	16.00	1.349 ***	94.05	5.697 ***	75.00	-7.525 ***	94.07	6.353 ***
8 Parent	15.67	0.255	15.33	0.387	15.67	0.246	113.02	12.263 ***	110.00	9.202 ***	113.03	13.247 ***
9 Parent	15.41	0.244	13.00	1.356 ***	17.87	0.989 *	92.17	2.984 *	91.67	3.899 **	92.17	3.607 *
Gi <math>\hat{C}</math> 0 at 95%	0.381 ***		0.657 ***		0.894 ***		3.217 ***		3.611 ***		3.836 ***	
Gi--Gj at 95%	0.571 ***		0.985 ***		1.341 ***		4.825 ***		5.416 ***		5.754 ***	
GCA/SCA Ratio	0.072		0.115		0.122		0.107		0.081		0.110	
Predictability Ratio	0.125		0.187		0.196		0.176		0.139		0.180	

**Table 5:** General Combining Ability of nine parents studied under rabi, 2017:18 (E1), kharif- 2018 (E2) and rabi, 2018-19 (E3)

Traits / Parents	Ear height						Days to 50% anthesis					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	51.40	-1.691 ***	53.33	-2.111	51.57	-2.337 **	106.67	-0.502	49.67	-1.175 ***	102.67	-0.434
2 Parent	58.11	-2.168 ***	57.83	1.343	56.10	-2.992 ***	111.00	1.074 ***	49.33	0.704 **	107.00	1.141 ***
3 Parent	49.33	-4.411 ***	63.77	-0.457	44.67	-5.361 ***	109.67	0.771 **	49.33	0.401	103.67	0.475
4 Parent	45.00	-3.046 ***	48.33	0.919	38.67	-4.176 ***	109.67	0.620 *	49.67	-0.024	105.67	0.687 *
5 Parent	43.11	-2.619 ***	51.67	-3.884 **	39.20	-3.179 ***	106.33	-0.168	48.67	-0.024	102.33	-0.101
6 Parent	58.00	2.846 ***	53.33	-0.505	55.67	4.563 ***	107.33	0.377	48.00	0.158	103.33	0.444
7 Parent	56.67	2.727 ***	43.33	-4.248 **	54.33	2.793 ***	111.00	-0.077	49.33	0.582 *	105.67	-0.253
8 Parent	58.40	6.748 ***	68.33	5.229 ***	58.77	8.969 ***	106.33	-2.108 ***	48.67	-0.933 ***	102.33	-2.040 ***
9 Parent	60.48	1.613 ***	51.67	3.714 **	58.57	1.721 *	107.00	0.013	48.67	0.310	103.00	0.081
Gi <math>\hat{C}</math> 0 at 95%	0.861 ***		3.038 ***		1.882 ***		0.599 ***		0.603 ***		0.636 ***	
Gi--Gj at 95%	1.292 ***		4.557 ***		2.823 ***		0.899 ***		0.904 ***		0.954 ***	
GCA/SCA Ratio	0.113		0.101		0.133		0.091		0.172		0.096	
Predictability Ratio	0.185		0.169		0.210		0.153		0.256		0.161	

**Table 6:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits / Parents	Days to 50% silking						Physiological maturity					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	109.00	-0.448	53.00	-1.135 ***	106.33	-0.253	141.00	-0.455	82.33	0.290	141.33	-0.273
2 Parent	113.67	1.007 ***	50.67	0.744 **	110.33	1.141 ***	145.67	1.000 ***	81.33	-0.013	145.33	1.061 ***
3 Parent	111.67	0.582 *	50.00	0.259	106.00	0.263	144.00	0.636 *	81.00	0.350	142.33	0.455
4 Parent	112.00	0.522 *	52.33	-0.347	108.33	0.596 *	144.00	0.515 *	81.67	-0.438	144.00	0.636 *
5 Parent	109.33	-0.266	52.00	0.077	105.00	-0.313	141.33	-0.273	80.67	0.168	140.33	-0.333
6 Parent	110.67	0.522 *	50.00	-0.044	107.33	0.657 *	142.67	0.515 *	82.67	0.138	142.33	0.576 *
7 Parent	114.33	0.067	51.67	0.956 ***	109.00	-0.101	146.33	0.061	80.67	0.320	145.33	0.000
8 Parent	109.00	-2.024 ***	50.33	-1.044 ***	105.00	-2.010 ***	141.00	-2.030 ***	81.00	-0.135	140.00	-2.091 ***
9 Parent	110.67	0.037	51.00	0.532 *	106.67	0.020	142.67	0.030	80.33	-0.680 *	141.67	-0.030
Gi <math>\times</math> 0 at 95%	0.576 ***		0.507 ***		0.632 ***		0.573 ***		0.632 ***		0.627 ***	
Gi--Gj at 95%	0.864 ***		0.760 ***		0.948 ***		0.860 ***		0.948 ***		0.941 ***	
GCA/SCA Ratio	0.075		0.143		0.077		0.075		0.057		0.073	
Predictability Ratio	0.130		0.223		0.133		0.131		0.102		0.127	

**Table 7:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits / Parents	Ears per plant						Ear length					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	1.30	0.049 *	1.23	0.055 **	1.17	0.014	14.09	-0.025	13.33	0.006	14.10	-0.100
2 Parent	1.11	-0.064 **	0.96	-0.028	1.10	-0.007	13.77	-0.356	11.42	-0.328	13.77	-0.342
3 Parent	1.12	-0.016	1.04	-0.072 ***	1.07	-0.016	13.27	-0.376	15.33	0.157	13.27	-0.360
4 Parent	1.10	-0.005	1.00	-0.002	1.07	-0.035	13.48	0.252	12.77	0.063	13.70	0.306
5 Parent	1.08	0.027	1.08	-0.022	1.03	-0.029	14.93	-0.166	12.87	-0.140	14.93	-0.151
6 Parent	0.92	0.000	1.14	0.030	1.07	-0.010	12.17	-0.096	14.17	-0.064	12.20	-0.075
7 Parent	0.90	-0.067 **	1.10	-0.004	1.10	0.017	11.67	-0.289	13.50	0.427	11.67	-0.275
8 Parent	1.04	0.075 **	1.07	0.047 **	1.10	0.017	14.17	0.466 *	11.07	-0.146	14.17	0.391
9 Parent	0.89	0.001	1.08	-0.004	1.13	0.050 *	13.93	0.591 **	12.87	0.024	13.93	0.606 **
Gi <math>\times</math> 0 at 95%	0.053 ***		0.038 ***		0.056 ***		0.462 ***		0.762 ***		0.456 ***	
Gi--Gj at 95%	0.079 ***		0.056 ***		0.084 ***		0.692 ***		1.143 ***		0.683 ***	
GCA/SCA Ratio	0.043		0.115		0.007		0.023		-0.019		0.023	
Predictability Ratio	0.079		0.187		0.015		0.044		-0.040		0.043	

**Table 8:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits Parents	Ear diameter						Number of kernels rows per ear					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	3.00	-0.053	2.70	-0.048	3.00	-0.108 *	10.73	-0.170	11.33	-0.377	10.73	-0.217
2 Parent	2.80	-0.091	2.53	-0.070	2.80	-0.084	10.67	-0.261	12.00	-0.013	10.67	-0.275
3 Parent	2.82	-0.170 ***	2.63	-0.085	3.00	-0.120 **	11.33	-0.023	11.33	-0.013	11.33	-0.038
4 Parent	3.00	-0.037	2.53	-0.070	3.00	-0.029	11.33	-0.196	11.33	0.471	11.33	-0.105
5 Parent	3.00	-0.067	2.60	-0.073	3.00	-0.059	11.33	-0.086	11.33	-0.074	11.33	-0.105
6 Parent	3.07	-0.062	2.40	-0.003	3.07	-0.044	9.33	-0.485 *	10.67	-0.559 *	9.33	-0.502 *
7 Parent	2.87	-0.061	2.57	0.017	2.87	-0.053	11.78	-0.092	11.33	-0.013	11.77	-0.005
8 Parent	4.00	0.413 ***	2.53	0.257 ***	4.00	0.359 ***	14.73	0.909 ***	10.67	0.229	14.73	0.862 ***
9 Parent	3.20	0.129 **	2.70	0.076	3.20	0.138 **	11.47	0.403 *	12.67	0.350	11.47	0.386 *
Gi <math>\times</math> 0 at 95%	0.108 ***		0.122 ***		0.105 ***		0.438 ***		0.593 ***		0.447 ***	
Gi--Gj at 95%	0.162 ***		0.183 ***		0.157 ***		0.658 ***		0.890 ***		0.671 ***	
GCA/SCA Ratio	42.908		0.102		3.888		0.177		0.067		0.152	
Predictability Ratio	0.988		0.169		0.886		0.261		0.119		0.233	

**Table 9:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits / Parents	Number of kernels per row						Yield per plant					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	30.33	0.598	28.67	-0.537	32.33	0.570	36.33	-0.122	27.11	-6.045 ***	39.80	-2.594
2 Parent	26.67	-1.582 **	24.33	0.206	26.00	-1.548 **	29.76	-4.045 ***	22.84	4.965 ***	29.23	-0.542
3 Parent	31.33	-0.244	28.00	0.479	29.73	-0.381	32.95	-9.432 ***	21.73	-6.451 ***	40.93	-5.742 ***
4 Parent	30.17	0.695	24.00	0.085	30.17	0.846	37.70	-3.483 **	32.50	-1.752 *	32.50	-3.706 **
5 Parent	29.73	-0.813	28.15	0.854 *	29.73	-0.660	42.00	-6.886 ***	21.65	-6.855 ***	38.10	-9.954 ***
6 Parent	28.33	-0.329	25.67	-1.340 **	28.33	-0.175	38.51	-0.044	34.07	0.262	30.13	-1.615
7 Parent	24.33	-0.923	26.33	-0.461	24.33	-0.769	33.15	-1.810	17.32	-0.749	29.53	-3.503 *
8 Parent	34.07	1.774 **	30.33	1.418 **	32.73	1.140 *	52.33	22.439 ***	22.57	15.559 ***	58.63	21.646 ***
9 Parent	30.33	0.824	25.33	-0.703	30.33	0.979	36.44	3.382 **	19.95	1.067	32.70	6.009 ***
Gi <math>\times</math> 0 at 95%	1.318 ***		0.974 ***		1.322 ***		2.452 ***		1.998 ***		3.209 ***	
Gi--Gj at 95%	1.977 ***		1.461 ***		1.984 ***		3.677 ***		2.997 ***		4.813 ***	

GCA/SCA Ratio	0.031	0.061	0.020	0.118	0.080	0.115
Predictability Ratio	0.059	0.109	0.038	0.191	0.138	0.187

**Table 10:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits / Parents	Hundred seed weight						Moisture content					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	12.96	0.140	21.93	0.265	11.70	0.125	14.50	0.246 *	13.37	-0.542 **	13.77	0.160
2 Parent	12.82	-0.233	20.63	-0.038	12.20	-0.499 *	13.93	0.334 **	14.30	-0.297	13.93	0.327 *
3 Parent	13.89	0.350	20.97	0.498 **	11.73	0.001	12.72	-0.196	12.10	-0.240	12.17	-0.249
4 Parent	12.56	-0.429 *	20.70	-0.308	11.17	-0.708 ***	12.95	-0.499 ***	13.35	-0.101	12.37	-0.558 ***
5 Parent	10.40	-0.855 ***	20.13	-0.219	10.70	-0.420 *	13.30	-0.499 ***	14.25	0.309	11.87	-0.661 ***
6 Parent	13.20	-0.942 ***	21.20	0.061	13.40	-0.593 **	13.72	0.122	13.62	-0.317	13.53	0.133
7 Parent	13.53	0.719 ***	19.53	-0.550 **	14.43	0.868 ***	13.60	0.015	13.74	0.307	13.23	-0.043
8 Parent	13.14	0.789 ***	21.47	0.272	12.83	1.043 ***	14.05	0.380 **	14.53	0.707 ***	15.97	0.691 ***
9 Parent	13.69	0.462 *	21.13	0.020	12.13	0.183	14.03	0.096	15.33	0.174	14.60	0.200
Gi <math>\hat{<math>0</math> at 95%</math>	0.421 ***		0.412 ***		0.455 ***		0.275 ***		0.371 ***		0.349 ***	
Gi--Gj at 95%	0.632 ***		0.618 ***		0.682 ***		0.412 ***		0.557 ***		0.524 ***	
GCA/SCA Ratio	0.124		0.094		0.075		1.318		0.168		-23.498	
Predictability Ratio	0.199		0.158		0.131		0.725		0.251		1.022	

**Table 11:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits / Parents	Popping expansion volume						Flake size					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	7.98	-0.229	9.99	0.292 *	7.83	-0.206	1.51	-0.010	1.73	-0.031	1.67	0.059 *
2 Parent	10.81	0.458 ***	11.41	0.003	10.40	0.775 ***	2.08	0.119 ***	2.03	-0.087 **	2.23	0.280 ***
3 Parent	11.89	0.462 ***	7.83	-0.260 *	12.83	0.754 ***	1.91	-0.102 ***	2.04	-0.031	1.80	-0.016
4 Parent	8.90	-0.639 ***	8.48	-0.448 ***	9.70	-0.591 ***	1.61	-0.046 *	1.96	-0.110 ***	1.17	-0.271 ***
5 Parent	9.42	-0.051	8.64	-0.275 *	9.77	-0.412 **	2.13	-0.033	2.27	0.059 *	2.13	0.008
6 Parent	9.91	0.425 **	14.88	1.428 ***	9.87	0.551 ***	1.59	-0.009	2.77	0.073 **	1.50	-0.059 *
7 Parent	9.39	0.133	6.72	-1.036 ***	9.43	-0.461 ***	1.65	0.012	1.77	0.050	1.77	0.065 *
8 Parent	10.43	-1.008 ***	12.52	0.490 ***	11.30	-0.491 ***	1.80	-0.030	2.11	0.008	1.63	-0.092 **
9 Parent	11.67	0.449 ***	9.70	-0.194	11.90	0.081	2.28	0.098 ***	2.32	0.068 *	2.30	0.026
Gi <math>\hat{<math>0</math> at 95%</math>	0.299 ***		0.263 ***		0.285 ***		0.050 ***		0.061 ***		0.068 ***	
Gi--Gj at 95%	0.449 ***		0.394 ***		0.427 ***		0.076 ***		0.092 ***		0.101 ***	
GCA/SCA Ratio	0.062		0.099		0.059		0.022		0.012		0.091	
Predictability Ratio	0.110		0.165		0.106		0.043		0.023		0.154	

**Table 12:** General Combining Ability of nine parents studied under *rabi*, 2017:18 (E1), *kharif*- 2018 (E2) and *rabi*, 2018-19 (E3)

Traits / Parents	Flake volume						Popping rate					
	Mean	E1	Mean	E2	Mean	E3	Mean	E1	Mean	E2	Mean	E3
1 Parent	174.33	16.509 ***	174.07	10.087 ***	180.00	5.644 *	86.84	2.844 ***	86.84	2.844 ***	75.33	0.468
2 Parent	178.67	-1.622	197.53	-4.002 *	182.00	1.290	81.27	1.575 *	81.27	1.575 *	78.93	0.932
3 Parent	193.33	-8.119 ***	178.03	-17.939 ***	189.20	-11.486 ***	72.68	-0.248	72.68	-0.248	83.30	-0.123
4 Parent	190.17	-3.022	174.80	-12.258 ***	208.67	0.360	76.70	-0.116	76.70	-0.116	84.27	0.153
5 Parent	191.74	-6.226 **	164.77	13.984 ***	205.17	6.144 *	75.33	-1.515 *	75.33	-1.515 *	85.70	1.171
6 Parent	193.33	-2.783	288.47	17.413 ***	181.67	-7.571 **	83.75	1.339 *	83.75	1.339 *	79.40	-0.377
7 Parent	183.50	3.078	158.43	-7.120 ***	181.10	0.181	75.20	-1.444 *	75.20	-1.444 *	80.73	-0.620
8 Parent	197.88	-0.883	224.83	0.493	195.50	2.911	75.95	0.017	75.95	0.017	74.13	-3.753 ***
9 Parent	226.67	3.069	224.43	-0.657	216.67	2.526	74.03	-2.453 ***	74.03	-2.453 ***	86.97	2.150 *
Gi <math>\hat{<math>0</math> at 95%</math>	5.105 ***		4.565 ***		6.377 ***		1.381 ***		1.381 ***		2.144 ***	
Gi--Gj at 95%	7.657 ***		6.848 ***		9.566 ***		2.071 ***		2.071 ***		3.216 ***	
GCA/SCA Ratio	0.022		0.044		0.009		0.073		0.073		0.066	
Predictability Ratio	0.043		0.080		0.017		0.128		0.128		0.116	

## Conclusion

G×E interaction is clearly indicated as narrow sense heritability is less than unity. Selection of parents with dominant alleles in yield attributing characters is desirable in pedigree method or bulk method is to be followed. As all twenty characters studied recorded partial or over dominance selection of inbred lines in early generation is not suggestible, it is to be done in later generation to obtain better inbred line. Two good general combiners are recorded, HKIPC-7 can be exploited for yield per and earliness, WINPOP-29 can be

exploited for earliness and dwarfier lines. Specific combiners for yield and popping characters are WINPOP-47×HKIPC-7 cross for yield per plant, WINPOP-8×HKIPC-7 and WINPOP-3×HKIPC-7 cross for earliness and WINPOP-47×HKIPC-5cross for popping expansion

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