

STUDIES OF GENETIC PARAMETERS AND VARIABILITY IN ELITE MAIZE (*Zea mays* L.) LINES

ABSTRACT

The present investigation was carried out with 30 maize inbred lines at Seed Research Technology Centre, Rajendranagar. Analysis of variance revealed significant differences among the genotypes for all the traits studied. Phenotypic coefficient of variance (PCV) was slightly higher than the genotypic coefficient of variance (GCV) indicating the role of experimental variance to the total variance. High PCV and GCV was observed for SVI-2 whereas cob yield per plant and grain yield per plant showed high PCV and moderate GCV. It indicated the presence of a high degree of variability. Ear height, 100- grain weight, seed vigour index - I, speed of germination had moderate PCV and GCV. High heritability coupled with high genetic advance was recorded in ear height, 100-seed weight, cob yield per plant, seed vigour index-1, seed vigour index-2, speed of germination and grain yield per plant indicated additive gene action in inheritance and simple selection would be effective for yield improvement. Based on GCV, PCV, heritability and genetic advance over mean were recorded for ear height, ear length, cob yield per plant, grain yield per plant, SVI-1, SVI-2 and speed of germination which indicates the importance of these traits in selection of inbred lines.

Keywords: *Maize, inbreds, PCV, GCV, heritability, genetic advance, genetic variability.*

INTRODUCTION

Maize (*Zea mays* L.) is the most important cereal crops next to wheat and rice where it is considered has “Queen of the cereals” and also best known as golden crop because every part of this crop is useful to man, animals and the industries. Maize is the third-largest cereal crop in the world with a production of 1148 million metric tonnes in 2019 (FAOSTAT). It is the third most important cereal in India, next to wheat and rice in terms of area (9.8 m.ha), production (31.6 million tonnes) and productivity (31.9 q ha⁻¹) (Indiastat, 2020-21). It also plays a key role in the Indian economy by contributing a significant share in global agricultural imports and exports.

Maize crop is majorly used as feed for animals and food for humans. Apart from food and feed it is also extensively used for corn oil, starch and biofuel production. It is processed into many food products such as tortillas, snacks, flour, flakes etc. It is enriched with abundant amount of macronutrients like starch, fibre, protein and fat along with micronutrients like vitamin B complex, β -carotene and essential minerals such as magnesium, zinc, phosphorus, copper, etc. Many hybrids or varieties are released by both public and private to cater to the various needs.

Development of high yielding maize varieties is the most fundamental goal of any maize breeder to increase yield. Grain yield is the collective product of inherited and environmental factors. Genetic variability in maize genotypes plays a vital role in grain yield variation. Maize production could be increased through development of improved genotypes capable of producing enhanced yield under different agro-climatic conditions. Genetic variability among maize genotypes can be estimated based on qualitative and quantitative traits.

The major breeding approach for increasing productivity rely on production of hybrids using heterosis breeding. In this context, the first step is to develop maize inbred lines and assess the extent of genetic variability. Genetic variability is the pre requisite for any crop improvement program. Genetic variability, which is a heritable difference among cultivars, is required at an appreciable level within a population to facilitate and sustain an effective long term plant breeding program. Progress from selection has been reported to be directly related to the magnitude of genetic variance in the population. Improvement in any trait depends solely on the amount of variability present in the base material for that trait. Therefore, variability is a key for crop improvement (Welsh, 1981).

The study of variability and genetic advance in the germplasm will help to ascertain the real potential of the genotype (Larik, 2000). Accordingly, study of genetic parameters like genotypic coefficient of variation, phenotypic coefficient of variation, heritability and genetic advance as per cent of mean provides a clear cut idea about the extent of variability present in a plant population and a relative measure of efficiency of selection of genotypes based on phenotype in a highly variable population (Bocanski, 2009). The estimates of genetic parameters like heritability and genetic advance helps in predicting the gain under selection. Therefore, this study was undertaken to study the genetic variability, heritability and genetic advance among the maize genotypes for yield and yield contributing traits.

MATERIALS AND METHODS

The present pursuit entails study of genetic parameters and variability in a set of 30 inbreds of maize developed by the Maize Research Centre, Rajendranagar for grain yield and yield contributing characters including seed vigour traits. The experiment was conducted with standard agronomical package of practices at Seed Research Technology Centre, Rajendranagar, Professor Jayashankar Telangana State Agricultural University (PJTSAU) during the *Rabi* 2020 in Randomized Block Design with three replications and the plot size was 4 rows of 6 metres length for each genotype. The data subjected to INDOSTAT software to estimate genetic coefficient of variation (%), phenotypic coefficient of variation (%), heritability (%) (Broadsense) and genetic advance as percent of mean. The estimates for variability treated as per the categorization proposed by Siva Subramanian and Madhavamenon (1973), heritability and genetic advance as percent of mean estimates according to criteria proposed by Johnson *et al.* (1955). From the heritability estimates the genetic advance was calculated by the following formula given by Burton, (1952).

RESULTS AND DISCUSSION

The major breeding approach for increasing productivity rely on production of hybrids using heterosis breeding. In this context, the first step is to develop maize inbred lines and assess the extent of genetic variability. Selection of genetically superior genotypes as well as inbred lines for hybrid breeding requires sufficient genetic variability and high heritability in the base population. Analysis of variance (Table. 1) revealed highly significant differences among genotypes in respect of almost all the characters under studied at 1% and 5% level. This indicates that a sufficient range of variability in all the traits exists among the populations. Significant difference among the genotypes for studied characters have also been reported by Bisen *et al.* (2018), Bartaula *et al.* (2019), Prakash *et al.* (2019) and Taiwo *et al.* (2020).

The mean values, genotypic and phenotypic variances, heritability, and the genetic advance of traits studied are some of the key parameters which determine the efficiency of a breeding program.

MEAN PERFORMANCE

The mean performance of the inbred lines for eighteen yield, yield contributing characters and seedling vigour traits is presented in Table 2. Among the 30 maize inbreds, days to tasseling was recorded highest in PFSR- 56 (66.0) and lowest in PFSR-12 (57.6), days to silking(68.0,59.6) and days to maturity (112.0,111.6) were highest in PFSR-12 and lowest in MGC- 137. Plant height was highest in PFSR- 104 (168.7) and lowest in GP-311 (116.3). Highest ear height was observed in PFSR-198 (93.60), while the lowest was observed in the inbred line PFSR-151 (61.76). Ear length was highest in PFSR-135 (14.63), lowest in PFSR-56 (10.23). Ear diameter was highest in MGC- 137 (4.06) and lowest in PFSR-49 (3.03).

Number of kernels rows per ear and speed of germination were in highest in PFSR-56 (14.6) and lowest in MGC-137 (9.3). MGC-137 (32) had the highest number of kernels per row and GP-170 (17.6) had lowest. 100- Grain weight was highest in PFSR-29 (33.4) and lowest in MGC-7 (16.0). Shelling percentage was highest in MGC-7 (84.2%) and lowest in PFSR-204 (75.1%). Cob yield per plant was highest in MGC-137 (122.3) and lowest in GP-19 (46.0).

Grain yield per plant (98.6,38.3) and germination first count (100,78) were highest in PFSR-17 and lowest in MGC-137. Germination final count (100,84), seed vigour index-1 (3940.0,2127.6)and seed vigour index- 2 (102.0,30.2) were highest in GP-311 and lowest in MGC-137. In other words the performances of the genotypes with respect to these characters were statistically different, suggesting scope for improvement. These results are in agreement with the results of (Dar *et al.*, 2018), (Gazal *et al.*, 2017), (Kumar *et al.*, 2018) and (Asghar and Khan, 2005).

Genetic Variability Parameters

In crop breeding, variability is very significant. The degree of variability present in crop species is critical because it serves as the basis for selection. Genetic and environmental influences contribute to the overall variation in a population. The presence of significant genetic variability in the breeding materials is critical for a successful plant breeding program exploitation. The computed values of phenotypic, genotypic coefficient of variation, genetic advance and heritability are presented in Table 3 and 4.

The traits studied were at three levels *i.e.*, high, medium, and low. PCV showed a wide range of variation from 2.72 to 23.10 percent while the GCV ranged 1.30 to 22.83 percent for the different

traits studied. Phenotypic Coefficient of variance is slightly higher than the Genotypic coefficient of variance which indicating the role of experimental variance to the total variance. Both PCV and GCV were high for SVI-2 (23.10, 22.83), whereas cob yield per plant (20.21, 19.10) and grain yield per plant (20.36, 19.23) had high PCV and moderate GCV. It implies that presence of high variability among the genotypes for this trait and the possibility of selection of these traits for improving maize yield. Ear height (12.03, 11.81), 100- grain weight (18.82, 18.48), seed vigour index – I (13.34, 12.86), speed of germination (18.49, 18.33) had the moderate PCV and GCV.

Moderate PCV and low GCV were noticed for plant height (12.87, 9.98), cob length (10.22, 8.56), number of kernels row per cob (12.40, 8.67), and number of kernels per row (13.50, 9.80). Low phenotypic coefficient of variation and genotypic coefficient of variation were observed for traits days to tasseling (3.92, 3.27), days to silking (3.76, 3.06), days to maturity (2.72, 1.30), ear diameter (6.65, 5.31), shelling % (3.48, 2.72), germination 1st count (5.26, 5.00) and germination 2nd count (3.28, 2.98) indicating the absence of variation for these traits in the genotypes studied and improvement can't be achieved through the traditional methods and new methods such as mutation breeding and DNA technologies can be used to create variation.

These findings are in agreement with the findings of Ahmed *et al.* (2020), Chavan *et al.* (2020), Chandel *et al.* (2019), Prakash *et al.* (2019), Bartaula *et al.* (2019), Ramyashree *et al.* (2015) and Adebisi *et al.* (2013). Thus, indicating presence of sufficient inherent genetic variance over which selection could be effective. These characters with low magnitude of genetic variability may have limited utility in a programme of selection for their improvement.

Heritability and Genetic Advance

High heritability coupled with high genetic advance was noticed for ear height (96.4%, 23.9), 100-seed weight (96.4%, 37.4), cob yield per plant (89.3%, 37.18), seed vigour index-1(93.1%, 25.6), seed vigour index-2 (97.7%, 46.50), speed of germination (98.2%,37.42) and grain yield per plant (89.2%, 37.4) which suggest that these characters are controlled by the additive type of gene action in the inheritance and simple selection for these traits would be fruitful. Thus, heritability estimates coupled with high genetic advance would be more reliable (Reddy *et al.*, 2012).

High heritability coupled with low genetic advance was observed for days to 50% tasseling (69.7%, 5.63), days to 50% silking (65.9%, 5.11), ear diameter (63.9%, 8.75), shelling percentage

(61.1%, 4.39), germination 1st count (90.5%, 9.80), germination 2nd count (82.4%, 5.57) indicating that they are controlled by the non-additive type of gene action which is influenced by the environment. For the traits number of rows per ear and number of kernels per row which showed moderate heritability and moderate genetic advance suggested that these characters are controlled by joint action of genetic and non genetic factors.

High heritability and moderate genetic advance for plant height (60.1%, 15.9) and cob length (70.1%, 14.8) suggested that these traits are highly heritable and selection on the basis of this trait will be effective. Low heritability and genetic advance for days to maturity (22.8%, 1.27) indicated that the need of creation of variability either by hybridization or mutation followed by selection.

The findings of this study are in broadly agreement with the findings of Ahmed *et al.* (2020), Chavan *et al.* (2020), Singh *et al.* (2019), Prakash *et al.* (2019), Bartaula *et al.* (2019), Ramyashree *et al.* (2015), Ali *et al.* (2013) and Adebisi *et al.* (2013). The characters that showed high heritability coupled with high genetic advance could be improved through simple selection.

Conclusion

Based on PCV, GCV, heritability, along with high genetic advance as percent of mean traits namely ear height, ear length, cob yield per plant, grain yield per plant, SVI-1, SVI-2 and speed of germination) were found to be crucial in selection process for the improvement of yield on maize.

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Table. 1. Analysis of variance for yield, yield components and seed vigour traits in Maize

S.No.	Character	Mean sum of squares		
		Replications (d.f.=2)	Treatments (d.f.=29)	Error (d.f.=58)
1	Days to 50% tasseling	1.944	14.10**	1.784
2	Days to 50% silking	3.033	13.295**	1.953
3	Days to maturity	24.044	14.619*	7.757
4	Plant height (cm)	428.796	787.911**	142.840
5	Ear height (cm)	5.097	251.797**	3.105
6	Ear length (cm)	0.558	4.393**	0.546
7	Ear diameter (cm)	0.014	0.123**	0.020
8	No of rows per cob	1.111	3.617**	0.927

9	No of seeds per row	12.878	22.684**	5.234
10	100-grain weight (g)	0.417	73.240**	0.898
11	Shelling %	3.830	17.504**	3.060
12	Cob yield per plant (g)	13.253	649.217**	24.868
13	Germination 1 st count	6.011	71.396**	2.425
14	Germination 2 nd count	5.700	27.318**	1.815
15	Seed vigour index-I	33527.04	465850.0**	11284.29
16	Seed vigour index-II (g)	17.359	723.044**	5.666
17	Speed of germination	0.139	7.401**	0.045
18	Grain yield per plant (g)	7.286	428.998**	16.562

** Significant at 1 per cent level * Significant at 5 per cent level

UNDER PEER REVIEW

Table. 2. Mean performance of maize inbreds for grain yield, yield components and seed vigour traits

S.No.	Genotype	Days to (50%) tasseling	Days to (50%) silking	Days to Maturity	Plant height (cm)	Ear height (cm)	Ear length (cm)	Ear diameter (cm)	No. of rows per cob	No. of kernel per rows
1	PFSR-9	64.0	66.0	119.0	124.7	64.76	13.6	3.4	10.0	23.3
2	PFSR-12	66.0	68.0	121.0	127.5	63.03	14.0	3.2	10.0	27.0
3	PFSR-17	58.0	60.0	113.0	134.5	65.0	10.6	3.5	12.0	22.3
4	PFSR-19	62.3	64.0	117.3	152.4	77.0	12.3	3.4	10.6	25.6
5	PFSR-29	61.3	63.0	115.6	163.5	77.4	13.4	3.4	10.6	23.0
6	PFSR-30	60.3	62.3	115.3	155.6	78.2	13.9	3.4	10.0	22.6
7	PFSR-32	62.0	63.6	116.6	147.4	78.0	13.2	3.5	10.6	21.6
8	PFSR-46	62.0	62.6	115.3	150.0	78.2	13.8	3.4	11.3	26.0
9	PFSR-49	60.6	62.6	115.6	125.4	66.8	13.8	3.0	10.0	23.6
10	PFSR-56	57.6	60.3	113.3	121.0	79.4	14.6	3.1	9.3	27.3
11	PFSR-70	59.6	61.3	114.3	144.4	80.2	11.9	3.6	11.3	21.6
12	PFSR-71	61.3	63.0	116.3	138.2	66.0	11.9	3.3	11.3	23.0
13	PFSR-84	63.6	65.3	118.3	157.5	82.0	13.2	3.5	10.6	25.0
14	PFSR-90	63.3	65.0	118.0	161.6	78.6	14.1	3.6	10.0	25.0
15	PFSR-92	62.6	64.3	117.3	165.1	83.6	14.6	3.5	10.0	26.6
16	PFSR-95	62.0	63.0	116.0	166.5	86.8	13.7	3.6	10.6	25.3
17	PFSR-104	61.3	63.3	116.3	168.7	91.4	14.2	3.6	10.0	26.3
18	PFSR-127	61.0	63.0	116.0	157.8	92.0	14.2	3.5	11.3	23.0
19	PFSR-130	62.0	63.3	116.3	163.0	89.2	14.2	3.6	10.0	25.6
20	PFSR-132	61.6	63.6	116.6	159.8	66.0	12.9	3.4	10.6	27.6
21	PFSR-135	58.6	61.0	113.6	119.2	80.2	10.2	3.7	11.3	23.3

Table. 2. (Contd.) Mean performance of maize inbreds for grain yield, yield components and seed vigour traits

S.No.	Genotype	Days to (50%) tasseling	Days to (50%) silking	Days to maturity	Plant height (cm)	Ear height (cm)	Ear length (cm)	Ear diameter (cm)	No. of kernels per row	No. of kernel rows
22	PFSR-151	61.0	62.6	115.3	160.7	61.7	13.5	3.6	11.3	22.3
23	PFSR-198	62.3	63.6	116.6	146.9	93.6	12.9	3.3	12.0	28.6
24	PFSR-204	62.6	63.6	116.6	167.8	84.7	13.9	3.3	10.0	25.6
25	GP-16	63.0	64.3	117.3	156.1	77.8	13.8	3.3	10.0	24.0
26	GP-19	65.0	66.6	119.6	149.1	84.6	14.4	3.3	10.6	24.0
27	GP-170	63.0	64.3	117.3	141.0	67.4	10.6	3.5	10.6	17.6
28	GP-311	65.6	67.6	120.6	116.3	68.5	11.8	3.9	12.6	21.3
29	MGC-7	65.3	67.3	120.3	128.4	71.4	12.6	3.5	12.6	27.6
30	MGC-137	58.0	59.6	111.6	136.1	78.4	14.1	4.0	14.6	32.0
General Mean		61.92	63.63	116.57	146.9	77.08	13.22	3.49	10.8	24.62
C. V. %		2.15	2.19	2.38	8.13	2.28	5.58	3.9	8.84	9.29
Highest Range		66.0	68.0	112.0	168.7	93.60	14.63	4.06	14.6	32.0
Lowest Range		57.6	59.6	111.6	116.3	61.76	10.23	3.03	9.3	17.6
Standard Error		0.7	0.80	1.60	6.90	1.01	0.42	0.08	0.5	1.32
C.D 5%		2.18	2.28	4.5	19.5	2.8	1.20	0.2	1.57	3.73
C.D 1%		2.90	3.03	6.05	25.9	3.83	1.60	0.30	2.09	4.97

Table. 2. (Contd.) Mean performance of maize inbreds for grain yield, yield components and seed vigour traits

S.No.	Genotype	100 seed weight (g)	Shelling (%)	Cob yield per plant (g)	Germination 1 st count	Germination 2 nd count	SVI -1	SVI-2	Speed of germination	Grain yield per plant (g)
1	PFSR-9	26.4	82.2	76.0	92.0	96.0	3278.3	68.9	6.8	62.5
2	PFSR-12	24.7	78.1	73.0	99.0	100.0	2920.0	64.3	8.1	57.0
3	PFSR-17	17.6	80.1	47.8	99.6	100.0	3380.0	61.6	8.7	38.3
4	PFSR-19	22.0	80.5	72.6	98.0	98.6	3536.6	71.3	8.1	58.5
5	PFSR-29	33.4	80.6	76.3	98.0	98.0	2861.6	58.8	8.5	61.5
6	PFSR-30	31.9	83.4	72.3	98.0	98.0	2702.3	58.8	6.9	60.3
7	PFSR-32	29.9	79.0	71.3	92.0	97.6	2628.6	82.0	11.5	56.3
8	PFSR-46	27.2	81.5	77.6	98.0	99.0	2677.0	75.2	8.4	63.3
9	PFSR-49	23.8	83.0	59.6	94.0	98.0	2996.3	78.4	8.4	49.5
10	PFSR-56	22.3	84.1	65.3	97.6	99.0	3396.6	77.2	5.5	55.0
11	PFSR-70	25.4	82.1	70.0	99.3	100.0	2676.6	64.0	7.1	47.5
12	PFSR-71	24.9	81.4	59.0	86.0	97.6	2204.3	70.3	7.9	48.0
13	PFSR-84	29.2	81.3	72.3	94.3	94.0	3235.0	75.2	7.8	59.0
14	PFSR-90	31.1	82.6	90.0	99.6	99.6	3020.0	65.7	8.5	74.3
15	PFSR-92	28.9	81.4	91.3	90.3	96.3	2829.3	69.0	9.4	74.3
16	PFSR-95	30.7	80.1	84.0	92.0	98.0	2982.6	74.1	8.8	67.1
17	PFSR-104	30.3	81.4	88.0	94.0	96.0	3027.6	74.8	6.3	71.6
18	PFSR-127	30.3	80.2	80.3	99.6	100.0	3200.0	58.0	8.4	64.5
19	PFSR-130	29.8	78.6	90.3	100.0	100.0	3153.3	82.0	8.5	71.0
20	PFSR-132	24.5	82.7	83.0	100.0	100.0	3023.3	61.6	8.0	68.6
21	PFSR-135	31.5	81.3	75.3	98.0	100.0	2806.3	49.6	6.7	61.3

Table. 2. (Contd.) Mean performance of maize inbreds for grain yield, yield components and seed vigour traits

S.No.	Genotype	100 seed weight (g)	Shelling (%)	Cob yield per plant (g)	Germinat ion 1 st count	Germinat ion 2 nd count	SVI -1	SVI-2	Speed of germination	Grain yield per plant (g)
22	PFSR-151	32.1	77.5	88.3	98.0	99.6	2787.6	71.4	11.3	68.5
23	PFSR-198	16.5	79.2	58.0	94.0	98.0	3287.3	64.3	9.6	46.0
24	PFSR-204	32.5	84.2	87.0	98.0	98.3	3499.6	89.8	10.0	73.3
25	GP-16	29.0	79.3	77.3	92.0	98.0	3215.3	68.6	6.6	61.3
26	GP-19	18.1	83.4	46.0	99.0	99.0	3627.3	88.7	11.1	38.5
27	GP-170	27.4	80.1	72.0	98.0	98.0	2672.6	43.1	7.6	57.6
28	GP-311	25.3	75.5	75.0	78.0	84.0	2127.6	30.2	8.8	56.6
29	MGC-7	16.0	75.1	64.3	98.0	98.0	3074.0	31.3	10.1	48.3
30	MGC-137	23.4	75.8	122.3	100.0	100.0	3940.0	102.0	11.8	98.6
General Mean		26.56	80.56	75.53	95.82	97.96	3025.60	67.71	8.54	60.96
C. V. %		3.56	2.17	6.60	1.62	1.37	3.51	3.51.	2.27	6.67
Highest Range		33.43	84.26	122.33	100.0	100.0	3940.0	102.0	11.82	98.6
Lowest Range		16.03	75.16	46.0	78.0	84.0	2127.6	30.28	5.50	38.3
Standard Error		0.54	1.01	2.87	0.89	0.7	61.3	1.37	0.12	2.34
C.D 5%		1.54	2.85	8.15	2.54	2.20	173.61	3.89	0.34	6.65
C.D 1%		2.06	3.80	10.84	3.38	2.92	230.9	5.17	0.45	8.84

Table. 3. Magnitude of variability, heritability and genetic advance for grain yield, yield components and seed vigour traits in Maize

S.no	Character	General Mean	Range		Phenotypic Coefficient of Variation (PCV)	Genotypic Coefficient of Variation (GCV)	Heritability in broad sense (%) (H_{bs})	Genetic Advance as per cent of mean (5%) (GAM)
			Maximum	Minimum				
1	Days to 50% tasseling	61.9	66.0	57.66	3.92	3.27	69.7	5.63
2	Days to 50% silking	63.63	68.0	59.66	3.76	3.06	65.9	5.11
3	Days to maturity	116.57	121.0	111.66	2.72	1.30	22.8	1.27
4	Plant height (m)	146.9	168.67	116.3	12.87	9.98	60.1	15.94
5	Ear height(cm)	77.08	93.6	61.76	12.03	11.8	96.4	23.89
6	Cob length	13.23	14.63	10.23	10.22	8.56	70.1	14.77
7	Cob diameter	3.49	4.06	3.03	6.65	5.31	63.9	8.75
8	No of rows per cob	10.88	14.0	9.33	12.40	8.67	49.2	12.56
9	No of seeds per row	24.62	32.0	17.66	13.50	9.80	52.6	14.64
10	100-grain weight (g)	26.56	33.43	16.03	18.82	18.48	96.4	37.39
11	Shelling %	80.56	84.26	75.16	3.48	2.72	61.1	4.39
12	Cob yield per plant (g)	75.53	122.33	46.0	20.21	19.10	89.3	37.18
13	Germination 1 st count	95.82	100.0	78.0	5.26	5.00	90.5	9.80
14	Germination 2 nd count	97.96	100.0	84.0	3.28	2.98	82.4	5.57
15	Seed vigour index-I	3025.60	3940.0	2127.66	13.34	12.86	93.1	25.57
16	Seed vigour index-II (g)	67.71	102.0	30.28	23.10	22.83	97.7	46.50
17	Speed of germination	8.54	11.82	5.05	18.49	18.33	98.2	37.42
18	Grain yield per plant (g)	60.96	98.66	38.33	20.36	19.23	89.2	37.43

Table 4. Estimates of genetic parameters for grain yield, yield components and seed vigour traits in Maize

S. NO	Character	Genetic Parameter	Gene effects	Influence of environment
1	Ear height 100-seed weight Cob yield per plant Seed vigour index-1 Seed vigour index-2 Speed of germination Grain yield per plant	High heritability with high GAM	Additive	Low
2	Plant height Cob length	High heritability with moderate GAM	Additive	Low
3	Days to tasseling Days to silking Ear diameter Shelling percentage Germination 1 st count Germination 2 nd count	Moderate heritability with low GAM	Additive and non-additive	Low
4	Number of rows per ear Number of kernels per row	Moderate heritability with moderate GAM	Additive and non-additive	Medium
5	Days to maturity	Low heritability with low GAM	Non-additive	High