



Association Analysis over Seasons among Morphological, Physiological and Yield Components with Kernel Yield in Maize (*Zea mays* L.)

N Sabitha ^{a*}, D Mohan Reddy ^a, D Lokanatha Reddy ^a,
P Sudhakar ^a and B Ravindra Reddy ^a

^a S V Agricultural College, Acharya N G Ranga Agriculture University, Tirupati -517502, A.P., India.

Authors' contributions

This work was carried out in collaboration among all authors. Author NS designed the study, performed the statistical analysis, wrote the first draft of the manuscript and did literature searches. Authors DMR and BRR verified the review of literature and results of the experiment. Fourth author managed the analyses of the study. All authors read and approved the final manuscript.

Article Information

DOI: 10.9734/JABB/2024/v27i5774

Open Peer Review History:

This journal follows the Advanced Open Peer Review policy. Identity of the Reviewers, Editor(s) and additional Reviewers, peer review comments, different versions of the manuscript, comments of the editors, etc are available here: <https://www.sdiarticle5.com/review-history/114988>

Original Research Article

Received: 24/01/2024
Accepted: 29/03/2024
Published: 02/04/2024

ABSTRACT

Forty five single cross hybrids made from 10 inbred lines of maize through diallel mating design were evaluated for three seasons viz. *rabi*, *summer* and *kharif* from 2016-17 to 2017-18. Kernel yield had consistent significant and positive associations with SPAD meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index in *rabi*, *summer* and *kharif* seasons. Similar trend of positive and significant association of kernel yield with all the above characters were recorded at genotypic level indicating existence a close relationship among these characters. Days to 50% tasseling, days to 50% silking, days to maturity and specific leaf weight showed consistent negative and significant correlations with kernel yield both at phenotypic and genotypic level indicating yield penalty with increase in

*Corresponding author: E-mail: nsabitha84@gmail.com;

days to 50% tasselling, days to 50% silking and specific leaf weight because more of vegetative growth and less time for reproductive growth which consequently lead to less kernel yield. The associations of anthesis-silking interval with kernel yield were consistently negative and non-significant in all the three seasons at phenotypic level revealing that narrow interval of anthesis silking interval facilitates good seed setting. Plant height showed either negative and significant or negative but non-significant association with kernel yield across seasons suggesting that any increase in plant height may lead to reduction in kernel yield and thus medium plant height is desirable for recording higher kernel yields in Maize. Based on the results of character association analysis it was concluded that SPAD chlorophyll meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index might be given due importance, while formulating selection indices aimed at kernel yield improvement as these characters had showed consistently positive and significant associations with kernel yield. A plant with medium plant height and duration coupled with higher SPAD meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index is desirable for getting higher kernel yield in maize.

Keywords: Maize; association analysis; kernel yield; yield components; cereal crop; maize breeding; breeding programs.

1. INTRODUCTION

Maize (*Zea mays* L.) is the third most important productive potential cereal crop after wheat and rice and is considered one of the most versatile crops with greater adaptability in various agro-ecological regions [1]. It is a multipurpose crop used as food, feed, and industrial raw materials for diverse products [2]. Grain yield of maize is expected to increase due to combined effects of increased genetic potential of the hybrids and improved cultural practices and management. It is reported that half of the yield increase was contributed by improvements in genetic gain [3]. Therefore the principal goal of maize breeding programs is to develop new hybrids/ varieties that outperform existing hybrids/ varieties with respect to many characters. Maize yield had increased greatly with the extension of single cross hybrids throughout the world [4]. The kernel yield is a complex quantitative character influenced by several yield-contributing characters and controlled by multiple factors [5]. Hence, selection for kernel yield alone may not be much effective [6]. Knowledge of interrelationships between kernel yield and its contributing components significantly improves the efficiency of breeding programs through the use of appropriate selection indices [7], (Dan Singh et al., 2017). Correlation measures the degree of association: genetic or non-genetic between two or more characters and is measured by a correlation coefficient [8]. Phenotypic correlation involves both genetic and environmental effects, whereas genotypic correlation is the association of breeding values of the two characters [9,10,11]. Both measure

the extent to which degree the same genes or closely linked genes cause co-variation in two different characters [12]. Character association studies will help to assess the relationship between the yield and its contributing traits for enhancing efficiency selection [13]. The present study was conducted to determine the extent and nature of morpho-physiological and yield components relationship with kernel yield in maize through a simple coefficient correlation analysis.

2. MATERIALS AND METHODS

Forty five single cross hybrids developed from 10 inbred lines of maize through diallel mating design (Method II and Model II) were evaluated for their performance over three seasons viz., *rabi*, *summer* and *kharif* from 2016-17 to 2017-18 at Agricultural Research Station, Perumallapalli, A.P. The experiment was laid out in a randomized block design with three replications with five meters row length with 75 × 20 cm in *kharif* and 60 × 20 cm in both *rabi* and *summer* between rows and between plant to plant spacing was adopted respectively. The two seeds per hill were dibbled and one week after germination thinning operation was carried out to maintain single plant per hill. All the recommended package of practices was followed in raising a healthy crop. Data were recorded for 15 morpho-physiological and yield contributing characters on five randomly selected plants in each replication. The mean values for different characters were analysed using online software "OPSTAT" developed by Chaudhary Charan Singh

Agricultural University, Hissar. Correlation coefficients for different characters were estimated by the method suggested by Panse and Sukhatme [14].

3. RESULTS AND DISCUSSION

Phenotypic and genotypic correlation coefficients worked out for all the 15 characters over three seasons *i.e* *rabi*, *summer* and *kharif* are presented in Table 1. Genotype correlations were in general greater in magnitude than the phenotypic correlations indicating the lesser influence of environments on the expression of the characters.

Kernel yield had consistent significant and positive associations with SPAD meter reading (0.39**, 0.50** and 0.26**), specific leaf area (0.44**, 0.27** and 0.22**), cob length (0.53**, 0.49** and 0.45**), cob girth (0.62**, 0.59** and 0.52**), number of kernel rows cob⁻¹ (0.59**, 0.69** and 0.56**), number of kernels row⁻¹ (0.58**, 0.57** and 0.57**), 100 kernel weight (0.65**, 0.57** and 0.52**) and harvest index (0.26**, 0.29** and 0.32**) in *rabi*, *summer* and *kharif* seasons, respectively at phenotypic level. Similar trend of positive and significant association of kernel yield with all the above characters was recorded at genotype level. The significant and positive associations of SPAD meter readings [15], cob length [16,17] (Ghirmire and Timsina 2015) [18-25], cob girth [15,18,19,20,22,21,23,24,26], number of kernel rows cob⁻¹ [18] number of kernels row⁻¹ [15,18,20,22,21,27,24,23,28,25], 100 kernel weight [15,18,20,22,21,24,27,29,25] were reported by several research workers in maize.

Days to 50% tasseling (-0.45**, -0.27** and -0.48**), days to 50% silking (-0.40**, -0.26** and -0.47**), days to maturity (-0.48**, -0.36** and -0.31**) and specific leaf weight (-0.42**, -0.30** and -0.29**) showed consistent negative and significant correlations with kernel yield at phenotypic level. Ghimire and Timsina [15] reported that more number of days to 50% tasseling and days to 50% silking will result in more vegetative growth and less time for reproductive growth and leads to less yield. Negative associations of days to tasseling, days to silking with days to maturity with kernel yield as recorded in the present study are desirable in maize breeding aimed at the development of high yielding hybrids/ varieties with earliness. These results are in agreement with the findings of Ghimire and Timsina [15] Vijay Kumar et al. [18] and Woldu Mogesse [29] in maize.

The associations of anthesis-silking interval with kernel yield were consistently negative but non-significant (-0.03, -0.10 and -0.09) and significantly negative at genotypic level with kernel yield in all the three seasons. Plant height showed either negative and significant or negative but non-significant (-0.03, -0.24** and -0.32**) association with kernel yield across seasons. Contrary to the present study results, Ghilmire and Timsina [15], Hailegebrial and Yemane [19] and Jemal et al. [9] reported positive and significant associations of plant height with kernel yield in maize. This may be due to variation in the study environment, experimental material and number of genotypes handled in the study.

Associations among the characters revealed that days to 50% tasseling had significant positive correlations with days to 50% silking, days to maturity, specific leaf weight and negative significant association with SPAD meter reading, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight, harvest index at both phenotypic and genotypic level. Days to 50% silking showed positive and significant associations with anthesis-silking interval. Days to maturity showed negative and significant associations with SPAD meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index.

Association of days to maturity with SPAD meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index were consistently significant and negative, while these same characters had positive and significant association with specific leaf weight in all the three seasons. Anthesis-silking interval showed inconsistent negative or positive associations with plant height, SPAD meter readings, specific leaf area, specific leaf weight, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index except with days to maturity.

In the present study plant height had consistent significant and negative associations with SPAD meter readings, specific leaf area and positive and significant associations with specific leaf weight. Contrary to these findings Woldu Mogesse [29] observed significant and positive correlations of plant height with ear height, days to maturity, ear length, ear diameter, number of kernels row⁻¹ and number of kernel row cob⁻¹ at

Table 1. Estimates of phenotypic and genotypic correlation coefficients among morpho-physiological and yield components and with kernel yield in maize over seasons

Character(s)	Season(s)	DT	DS	ASI	DM	PH	SPAD	SLA	SLW	CL	CG	NKRC	NKPR	KW	HI	KY
DT	Rabi	1.00	0.86**	0.10	0.46**	0.03	-0.24*	-0.23**	0.20*	-0.30**	-0.49**	-0.36**	-0.32**	-0.34**	-0.24**	-0.45**
	Summer	1.00	0.96**	0.33**	0.40**	0.11	-0.40**	-0.32**	0.32**	-0.26**	-0.32**	-0.24**	-0.28**	-0.24**	-0.30**	-0.27**
DS	Rabi	1.00	0.93**	0.09	0.46**	0.35**	-0.34**	-0.34**	0.34**	-0.37**	-0.46**	-0.37**	-0.52**	-0.42**	-0.40**	-0.48**
	Summer	1.00	1.00	0.56**	0.50**	-0.02	-0.19*	-0.16	0.11	-0.30**	-0.39**	-0.26**	-0.25**	-0.30**	-0.22*	-0.40**
ASI	Rabi	1.00	1.00	0.59**	0.39**	0.13	-0.38**	-0.33**	0.33**	-0.27**	-0.32**	-0.26**	-0.30**	-0.25**	-0.29**	-0.26**
	Summer	1.00	1.00	0.45**	0.46**	0.38**	-0.34**	-0.38**	0.37**	-0.39**	-0.46**	-0.42**	-0.54**	-0.45**	-0.38**	-0.47**
DM	Rabi	1.00	0.28**	0.61**	1.00	0.24**	-0.09	0.03	0.09	-0.13	0.02	0.06	0.05	-0.01	0.02	-0.03
	Summer	1.00	0.92**	0.94**	1.00	0.14	0.13	-0.14	-0.19*	0.18*	-0.15	-0.16	-0.17*	-0.16	-0.11	-0.10
PH	Rabi	1.00	1.62**	1.54**	1.00	0.19*	0.16	-0.11	-0.19*	0.17*	-0.15	-0.15	-0.23**	-0.20*	-0.20*	-0.09
	Summer	1.00	0.61**	0.65**	1.00	-0.11	-0.23**	-0.27**	0.23**	-0.35**	-0.44**	-0.41**	-0.42**	-0.38**	-0.31**	-0.48**
SPAD	Rabi	1.00	0.52**	0.53**	1.00	0.19*	-0.47**	-0.26**	0.28**	-0.40**	-0.42**	-0.40**	-0.53**	-0.39**	-0.34**	-0.36**
	Summer	1.00	0.56**	0.59**	1.00	0.27**	-0.23**	-0.31**	0.26**	-0.37**	-0.47**	-0.41**	-0.33**	-0.25**	-0.23**	-0.31**
SLA	Rabi	1.00	0.01	-0.04	-0.15	1.00	0.16	-0.16	0.16	-0.02	-0.01	0.16	-0.07	-0.01	0.05	-0.03
	Summer	1.00	0.14	0.16	0.35**	1.00	-0.26**	-0.30**	0.32**	-0.19*	-0.15	-0.16	-0.35**	-0.21*	-0.10	-0.24**
SLW	Rabi	1.00	0.49**	0.52**	1.42**	0.38**	1.00	-0.16	-0.29**	0.24**	-0.23**	-0.22*	-0.36**	-0.32**	-0.24**	-0.32**
	Summer	1.00	-0.30**	-0.23**	0.06	-0.27**	1.00	0.19*	1.00	0.31**	-0.23**	0.50**	0.48**	0.51**	0.36**	0.39**
CL	Rabi	1.00	-0.50**	-0.51**	-0.60**	-0.66**	-0.36**	1.00	0.44**	-0.45**	0.47**	0.60**	0.54**	0.65**	0.49**	0.50**
	Summer	1.00	-0.58**	-0.65**	-2.21**	-0.58**	-0.31**	1.00	0.36**	-0.32**	0.41**	0.39**	0.34**	0.44**	0.34**	0.26**
CG	Rabi	1.00	-0.38**	-0.24**	0.21*	-0.58**	-0.43**	0.69**	1.00	-0.90**	0.48**	0.43**	0.39**	0.53**	0.29**	0.44**
	Summer	1.00	-0.55**	-0.56**	-0.60**	-0.54**	-0.59**	0.77**	1.00	-0.98**	0.35**	0.30**	0.24**	0.50**	0.22*	0.24**
NKRC	Rabi	1.00	-0.66**	-0.62**	-0.10	-0.62**	-0.67**	0.95**	1.00	-0.91**	0.25**	0.35**	0.34**	0.26**	0.23**	0.30**
	Summer	1.00	0.33**	0.17*	-0.26**	0.54**	0.38**	-0.56**	-0.97**	1.00	-0.43**	-0.36**	-0.40**	-0.34**	-0.50**	-0.27**
NKPR	Rabi	1.00	0.53**	0.54**	0.59**	0.54**	0.61**	-0.76**	-1.00**	1.00	-0.36**	-0.31**	-0.25**	-0.52**	-0.23**	-0.30**
	Summer	1.00	0.65**	0.61**	0.13	0.57**	0.59**	-0.94**	-0.83**	1.00	-0.25**	-0.37**	-0.35**	-0.25**	-0.24**	-0.29**
KW	Rabi	1.00	-0.44**	-0.34**	0.07	-0.47**	-0.07	0.79**	0.86**	-0.80**	1.00	0.64**	0.60**	0.50**	0.62**	0.46**
	Summer	1.00	-0.55**	-0.55**	-0.53**	-0.76**	-0.32**	0.91**	0.73**	-0.74**	1.00	0.50**	0.37**	0.41**	0.43**	0.28*
HI	Rabi	1.00	-0.55**	-0.60**	-1.85**	-0.52**	-0.41**	0.91**	0.62**	-0.71**	1.00	0.63**	0.53**	0.42**	0.49**	0.45**
	Summer	1.00	-0.68**	-0.51**	0.09	-0.54**	-0.03	0.65**	0.92**	-0.88**	0.96**	1.00	0.67**	0.53**	0.66**	0.32**
KY	Rabi	1.00	-0.40**	-0.39**	-0.29**	-0.52**	-0.22**	0.77**	0.50**	-0.50**	0.86**	1.00	0.58**	0.59**	0.60**	0.33**
	Summer	1.00	-0.62**	-0.62**	-1.01**	-0.59**	-0.33**	0.77**	0.69**	-0.74**	0.89**	1.00	0.62**	0.49**	0.56**	0.37**
NKRC	Rabi	1.00	-0.53**	-0.41**	0.07	-0.62**	0.15	0.76**	0.90**	-0.87**	0.98**	1.00	0.52**	0.69**	0.40**	0.59**
	Summer	1.00	-0.33**	-0.35**	-0.44**	-0.69**	-0.28**	0.79**	0.57**	-0.57**	1.01**	0.85**	1.00	0.56**	0.67**	0.29**
NKPR	Rabi	1.00	-0.68**	-0.68**	-1.04**	-0.68**	-0.50**	0.94**	0.79**	-0.85**	1.03**	0.95**	1.00	0.53**	0.45**	0.19*
	Summer	1.00	-0.62**	-0.52**	0.03	-0.62**	-0.07	0.60**	1.01**	-0.91**	0.99**	0.87**	1.00	0.56**	0.30**	0.58**
KW	Rabi	1.00	-0.39**	-0.42**	-0.55**	-0.72**	-0.53**	0.92**	0.84**	-0.84**	0.98**	0.80**	0.93**	1.00	0.57**	0.36**
	Summer	1.00	-0.75**	-0.76**	-1.29**	-0.59**	-0.51**	0.85**	0.64**	-0.67**	0.80**	0.76**	0.92**	1.00	0.47**	0.34**
HI	Rabi	1.00	-0.47**	-0.41**	-0.05	-0.52**	-0.01	0.66**	0.91**	-0.93**	0.91**	0.83**	0.93**	0.89**	1.00	0.37**
	Summer	1.00	-0.25**	-0.28**	-0.41**	-0.53**	-0.34**	0.64**	0.45**	-0.46**	0.81**	0.81**	1.05**	0.91**	1.00	0.26**
KY	Rabi	1.00	-0.56**	-0.63**	-2.38**	-0.32**	-0.41**	0.56**	0.45**	-0.46**	0.73**	0.73**	0.72**	0.70**	1.00	0.46**
	Summer	1.00	-0.31**	-0.27**	0.03	-0.47**	0.15	0.51**	0.67**	-0.56**	0.65**	0.46**	0.71**	0.57**	0.55**	1.00
KY	Rabi	1.00	-0.49**	-0.52**	-0.66**	-0.47**	-0.25**	0.50**	0.46**	-0.44**	0.67**	0.55**	0.63**	0.55**	0.61**	1.00
	Summer	1.00	-0.53**	-0.56**	-1.39**	-0.30**	-0.43**	0.69**	0.77**	-0.61**	0.49**	0.54**	0.45**	0.63**	0.58**	1.00
KY	Rabi	1.00	-0.56**	-0.50**	-0.07	-0.66**	0.01	0.47**	0.84**	-0.81**	0.78**	0.84**	0.83**	1.00**	0.81**	1.00
	Summer	1.00	-0.34**	-0.34**	-0.26**	-0.59**	-0.25**	0.76**	0.68**	-0.70**	0.93**	0.79**	0.87**	0.87**	0.86**	1.00
KY	Rabi	1.00	-0.64**	-0.71**	-2.31**	-0.44**	-0.49**	0.48**	0.54**	-0.67**	0.80**	0.69**	0.83**	0.87**	0.70**	1.00
	Summer	1.00	-0.64**	-0.71**	-2.31**	-0.44**	-0.49**	0.48**	0.54**	-0.67**	0.80**	0.69**	0.83**	0.87**	0.70**	1.00

*Significant at 5% level of significance; **Significant at 1% level of significance; Values upper diagonal indicates phenotypic correlations; Values lower diagonal indicates genotypic correlations

DT=Days to tasseling; DS=Days to silking; ASI=Anthesis-silking interval; DM=Days to maturity; PH=Plant height; SPAD=SPAD meter readings; SLA=Specific leaf area; SLW=Specific leaf weight; CL=Cob length; CG=Cob girth; NKRC=Number of kernel rows cob⁻¹; NKPR=Number of kernels row⁻¹; KW=100 kernel weight; HI=Harvest index; KY=Kernel yield plant⁻¹

phenotypic and genotypic level. This could be due to variation in the experimental material, sample size and environment.

SPAD meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index showed consistent significant and positive associations among the characters at phenotypic and genotypic level across three seasons. However, specific leaf weight had consistently showed significant and negative associations over three seasons both at phenotype and genotype level with SCMR, specific leaf area, specific leaf weight, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index.

4. CONCLUSION

From the present investigation it is evident that SPAD meter readings, specific leaf area, cob length, cob girth, number of kernel rows cob⁻¹, number of kernels row⁻¹, 100 kernel weight and harvest index had consistent positive and significant associations with kernel yield. Selection for any one of these characters would bring in simultaneous improvement of other characters and also finally improvement in kernel yield. Hence, simultaneous selection of these characters might be given due importance, while formulating selection indices or making selections for high yielding hybrids in maize. Negative associations of anthesis- silking interval with kernel yield suggests that narrow interval of anthesis- silking interval is desirable for good seed setting.

ACKNOWLEDGEMENTS

Funding for the experiment was provided by ANGRAU. Inbred lines of maize were shared from Agriculture Research Station, Peddapuram.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

1. Muhammad S, Muhammad A, Muhammad A, M. Atif. Comparative evaluation and correlation estimates for grain yield and quality attributes in Maize. Pakistan Journal of Botany. 2008;40(6):2361-2367.

2. Sumalini K, Shoba Rani T. Heterosis and combining ability for polygenic traits in late maturity hybrids of Maize (*Zea mays* L.). Madras Agricultural Journal. 2010;97(10-12):340-343.
3. Duvick, DN. Genetic progress on yield of united states maize (*Zea mays* L.) Maydica. 2005;50:193-202.
4. Quin X, Feng Xu YLS, Siddique KHM, YC Liao. Maize yield improvements in China. Past trends and Future directions. Plant breeding. 2016;135:166-176.
5. Kashiani G, Saleh G. Estimation of genetic correlation on sweet corn inbred lines using SAS mixed model. American Journal of Agricultural and Biological Sciences, New York. 2010;5:309-314.
6. Talebi R, Fayaz F, Jeloder NAB. Correlation and path coefficient analysis of yield and yield components of Chickpea (*Cicer arietinum* L.) under dry condition in the west of Iran. Asian Journal of Plant Science. 2007;6:1151-1154.
7. Pavlov J, Delic N, Markovic K, Crevar M, Camdzija, Stevanovic M. Path analysis for morphological traits in maize (*Zea mays* L.). Genetika. 2015;47(1):295-301.
8. Gomez KA, Gomez AA. Statistical Procedure for Agricultural Research. International Rice Research Institute, Manila, Philippines. 1984;305.
9. Falconer JS. Introduced to quantitative genetics 2nd edition. Longman, New York USA; 1981.
10. Muhammad Y, Muhammad S. Correlation analysis of S1 families of maize for grain yield and its components. International Journal of Agriculture and Biology. 2001;387-388.
11. Menkir A. Genetic variation for grain mineral content in tropical-adapted maize inbred lines. Food Chemistry, Science World Journal. 2008;110(1):454-464.
12. Hallauer AR, Miranda FJB. Quantitative genetics in maize breeding. 2nd edition. Iowa State University Press, Iowa, Ames. USA; 1988.
13. Pranay G, Shashibhushan D, Rani KJ, Bhadru D, Kumar CVS. Correlation and path analysis in elite maize (*Zea mays* L.) lines. International Journal of Plant & Soil Science. 2022;34(24):414-422.
14. Panse VG, Sukhatme PV. Statistical methods for agricultural workers. Published by ICAR, New Delhi; 1967.
15. Ghimire B, Timsina D. Analysis of yield and yield attributing traits of maize

- genotypes in Chitwan, Nepal. World Journal of Agricultural Research. 2015;3(5):153-162.
16. Rafique M, Hussain A, Mahaood T, Alivi AW, Alvi MB. Heritability and interrelationships among grain yield and yield components in Maize (*Zea mays* L.). International Journal of Agriculture and Biology. 2004;6(6):1113-1134.
 17. Rigon JPG, Rigon CAG. Quantitative descriptors and their direct and indirect effects on corn yield. Bioscience Journal Uberland. 2014;30(2):356-362.
 18. Vijay Kumar SK. Singh PK. Bhati, Amita Sharma SK. Sharma, Vinay Mahajan. Correlation, path and genetic diversity analysis in Maize (*Zea mays* L.). Environment & Ecology. 2015;33(2A): 971-975.
 19. Hailegebrail K, Yemane T. Studies of heritability, genetic parameters, correlation and path coefficient in elite Maize (*Zea mays* L.) hybrids. Academic Research Journal of Agricultural Science and Research. 2015;3(10):396-303.
 20. Kandel M, Ghimire SK, Ojha BR, Shrestha J. Correlation and path coefficient analysis for grain yield and its attributing traits of maize inbred lines (*Zea mays* L.) under heat stress condition. International Journal of Agriculture Environment and Food Sciences. 2018;2(4):124-130.
 21. Prakash R, Ravikesavan R, Vinodhana NK, Senthil A. Genetic variability, character association and path analysis for yield and yield component traits in maize (*Zea mays* L.). Electronic Journal of Plant Breeding. 2019;10 (2):518-524.
 22. Godwin MV, Maria BO EK. Success. Path coefficient analysis, character association studies in selected Maize (*Zea maya* L.) genotypes grown in Southern Nigeria. Annual Research and Review in Biology. 2019;33(3):1-6.
 23. Singh D, Kumar A, Kumar R, Singh SK, Kushwaha N, Mohanty TA. Correlation and path coefficient analysis for 'yield contributing' traits in quality protein Maize (*Zea mays* L.). Current Journal of Applied Science and Technology. 2020;39(25):91-99.
 24. Ahmed N, Chowdhury AK, Uddin MS, Rashad MMI. Genetic variability, correlation and path analysis of exotic and local hybrid maize (*Zea mays* L.) genotypes. Asian Journal of Medical and Biological Research. 2020;6(1):8-15.
 25. Prasanna Rani M, Anusha CR, Roopa Lavanya G, Sai Kiran M. Genetic variability and inter relationship in Maize (*Zea mays* L.) genotypes for grain yield and yield component characters. International Journal of Environment and Climate Change. 2022;12(11):3264-3271.
 26. Satish babu K, gabrial ML, Mohan Krishna N. Studies on genetic association and p[ath analysis for yield and yield attributing characters in Maize (*Zea mays* L.) International Journal of Plant and Soil Science. 2022;34(22):49-54.
 27. Jemal Aman, Kassahun Bantte, Sentayehu Alamerew, Desta Berhe Sbhatu. Correlation and path coefficient analysis of yield and yield components of quality protein Maize (*Zea mays* L.) hybrids at Jimma, Western Ethiopia. International Journal of Agronomy. 2021;2020, Article ID 9651537, 7. Available: <https://doi.org/10.1155/2020/9651537>
 28. Rajasekhar D, Naveenkumar KL, Pandey PK, Sen D. Analysis of morphological variation, grouping and path coefficient studies in a set of maize inbred lines local to North East Hill Region of India. International Journal of Plant and Soil Science. 2022;34(17):105-113.
 29. Woldu Mogesse. Correlation and path coefficient analysis for grain yield and yield related trait of Maize (*Zea mays* L.) inbred lines. International Journal of Current Research Academy Reiew. 2021;9(2): 138-150.

© Copyright (2024): Author(s). The licensee is the journal publisher. This is an Open Access article distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Peer-review history:

The peer review history for this paper can be accessed here:
<https://www.sdiarticle5.com/review-history/114988>