



Association and Diversity Studies for Yield and Its Attributing Traits in Maize (*Zea mays* L.) Hybrids

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Authors' contributions

This work was carried out in collaboration among all authors. All authors read and approved the final manuscript.

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ABSTRACT

Present investigation was conducted in 144 maize hybrids of CIMMYT, Hyderabad along with 6 checks at ICRISAT, Hyderabad. The genotypes were evaluated for six quantitative traits that are plant height, ear height, anthesis days, silking days and grain yield. Correlation among the traits came to a conclusion that plant height and ear height are positively correlated with grain yield. Path analysis revealed there is highest direct association of plant height on grain yield and ear height

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had a highest indirect positive association with grain yield. Hence selection plant height and ear height days plays a vital role in increasing yield. Clustering of genotypes based on the six traits divide the germplasm into two clusters. 85 and 65 genotypes were found in each clusters. Clustering the genotypes based the quantitative traits revealed close relatedness between the genotypes. Knowledge on association and diversity identified in our study help in plant breeding process.

Keywords: Maize hybrids; genotypes; grain; yield.

1. INTRODUCTION

Maize is one among the most important cereal crops. It is cultivated in 190mha in 165 countries [1]. It is an important cereal crop as it is a main component of human feed, animal feed and industrial purpose. In India maize is grown in 9.9 mha and produce 28 m tones of maize with on an average 6t/ha [2]. Growing importance of maize has increased its demand. Hence increasing the production and productivity of maize is need of the hour. Hence its important analyse the association of traits and available diversity in the germplasm. Analysis of genetic diversity in the germplasm will help in development and utilisation of diverse parents in the hybrid development. Correlation and path analysis of the important yield attributing traits help in indirect selection of yield through the associated traits when the heritability of the yield trait is low as in stress conditions. Many studies have been conducted in maize hybrids to assess the diversity and association among the traits in hybrids [2-5]. In present study we have attempted to analyse the diversity and association of traits in the CIMMYT maize hybrids to understand and utilise the results in future breeding programme.

2. MATERIALS AND METHODS

144 maize hybrids from CIMMYT, Hyderabad with 6 checks were evaluated for yield and yield attributing traits at Attur, Tamilnadu in 2022 Rabi. 150 genotypes were evaluated for six quantitative traits i.e., plant height (PH)(cm), ear height (EH)(cm), days for 50% anthesis (AD), days for 50% silking (SD), ear per plant (EPP) and grain yield (GY)(t/ha). 150 genotypes are laid out in alpha lattice design with two replications. Suitable agronomic practices were carried out. The observations for plant height, ear height and ear per plant was taken on five plants per plot, whereas the grain yield was measured on plot the converted to tonnes per hectare. Genotypic and phenotypic correlations were

done using the method by Searle [6]. Path analysis study was carried out in METAN software in R studio [7]. The cluster analysis using hierarchical clustering and ward. D method was done in R software using the factoextra and ggplot2 package Kassambara, [8]; Wickham,2023.

3. RESULTS AND DISCUSSION

3.1 Correlation

Yield is the prime most important character during selection and correlation studies provide a better understanding of this trait [9,10]. Correlation was worked out for all the five traits at genotypic and phenotypic levels (Table 1). Phenotypic correlation consists of both genotypic and phenotypic effects, it can be directly observed and are influenced by environmental factors. Inherent association between traits is studied through genotypic correlation it may be due to pleiotropic action of genes or due to linkage or more likely both [11]. Correlation studies in the present investigation revealed significant phenotypic and genotypic correlation of plant height, ear height and ear per plant with grain yield. Correlation of plant height with grain yield was also reported in earlier studies [12,13]. Ear height and plant height are significantly correlated. Since the plant height and ear height are significantly correlated with yield improving anyone of the trait will significantly improve the yield too. Anthesis and silking days both have perfect correlation with each other explaining the interdependency. But non-significant positive genotypic and negative phenotypic correlation was observed with grain yield, similar results were also obtained by other researchers [14-16]. Ear per plant are significantly correlated with all the traits both phenotypically and genotypically.

3.2 Path Coefficient

Multiple traits in correlation makes understanding the association between the traits more complex and doesn't have a meaningful

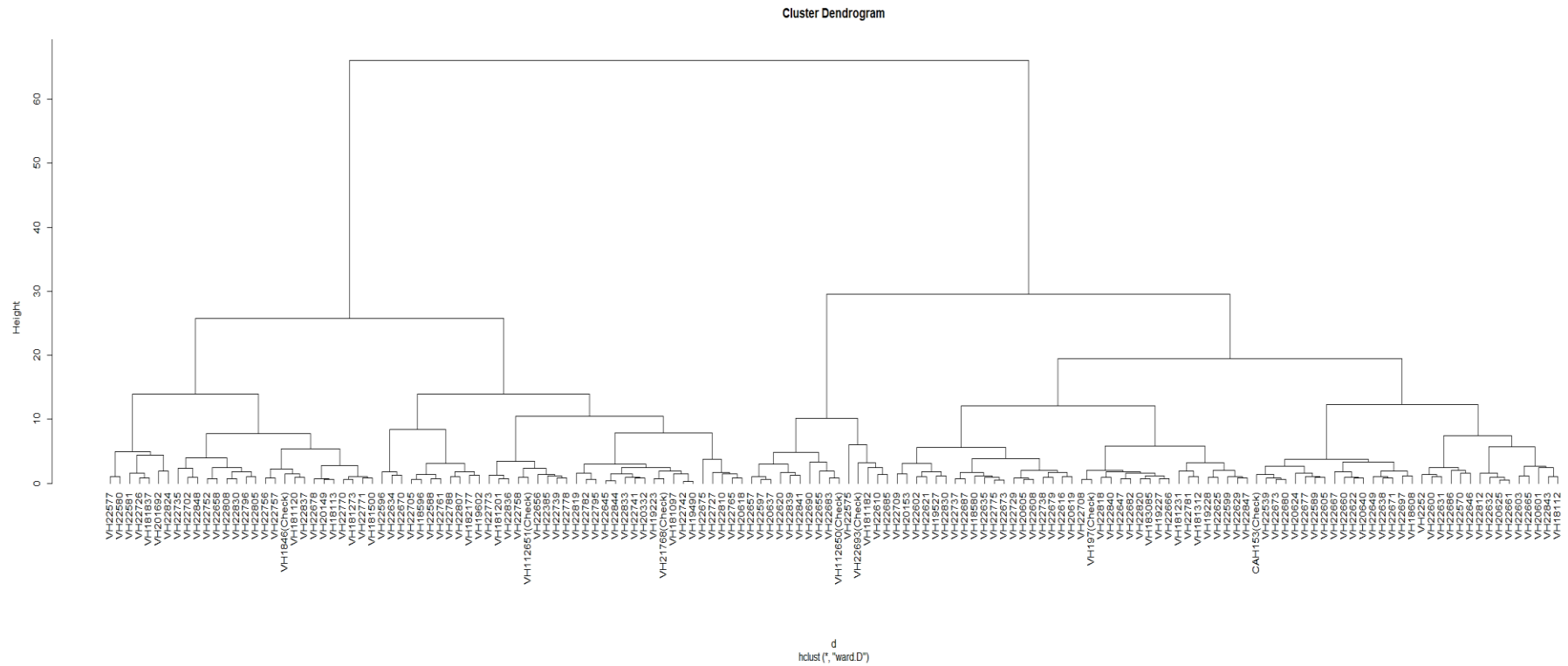


Fig. 1. Dendrogram representing clustering of 150 maize hybrids

Table 1. Phenotypic correlation among the traits

Traits	GY	AD	SD	PH	EH
AD	-0.09				
SD	-0.1	0.97**			
PH	0.52**	-0.14	-0.15		
EH	0.36**	0.13	0.11	0.73**	
EPP	0.24**	0.28**	0.24**	0.19*	0.21**

*Significant at 5% level, ** Significant at 1% level
GY, Grain Yield; AD, Anthesis Days; SD, Silking Days;
PH, Plant Height, EH, Ear Height; EPP Ear Per Plant

Table 2. Genotypic correlation among the traits

Traits	GYF	AD	SD	PH	EH
AD	0.05				
SD	0.03	1**			
PH	0.79**	-0.14	-0.16		
EH	0.55**	0.17*	0.15	0.78**	
EPP	0.22**	0.77**	0.78**	0.61**	0.47**

*Significant at 5% level, ** Significant at 1% level
GY, Grain Yield; AD, Anthesis Days; SD, Silking Days;
PH, Plant Height, EH, Ear Height; EPP Ear Per Plant

Table 3. Path coefficient study among five traits in maize hybrids

	PH	EH	AD	SD	EPP
PH	0.52	-0.11	-0.01	0.02	0.02
EH	0.42	-0.13	-0.003	0.02	0.02
AD	-0.10	0.01	0.03	-0.08	0.04
SD	-0.11	0.02	0.03	-0.08	0.03
EPP	0.04	-0.01	0.01	-0.01	0.19

*Bold numbers are direct effects
GY, Grain Yield; AD, Anthesis Days; SD, Silking Days;
PH, Plant Height, EH, Ear Height; EPP Ear Per Plant

Table 4. Contribution of traits in each cluster

Character	Cluster I	Cluster II
Grain yield	10.09	8.92
Ear height	105.83	92.56
Plant height	211.10	188.66
Anthesis days	61.61	64.12
Silking days	63.35	66.30
Ear per plant	0.96	0.97

Table 5. Inter and intra cluster distance

Cluster	I	II
I	2.75	3.63
II		2.80

interpretation. Hence partitioning the correlation to direct and indirect effects to specify the cause and their relative importance becomes important.

Highest positive direct effect was by plant height (0.52) for grain yield followed by ear per plant (0.19) on grain yield (Table 3). Ear height was giving negative direct effect on grain yield. Significant association of plant height with grain yield was also observed in earlier studies [17,16]. Silking days was negatively associated with the grain yield [18]. Indirect effect was highest by ear height (0.42) on grain yield through plant height rest all the traits had negligible contribution of indirect effects. With the obtained results we can conclude that increase in grain yield can be obtained by increasing the plant height and ear height and decreasing the days to silking.

3.3 Clustering of Genotypes

150 genotypes are divided at the height of 25 into two clusters. 65 genotypes are placed in one cluster and 85 genotypes are placed in other cluster (Fig. 1). Cluster I has the genotypes with greater yield, plant height, ear height, lesser anthesis and silking days and cluster II has the genotypes opposite of cluster I (Table 4). Both the clusters have the similar ear per plant. Intra cluster distances are lower than inter cluster distance. Both the clusters had the similar intercluster distance (Table 5). Distribution of 150 maize hybrids into only two clusters suggest that the genotypes are closely related. Clustering of hybrids into two clusters say that these hybrids are divergent [19]. This is an important observation as this knowledge would assist in developing inbreds. Similar kind of experiment was also conducted by various other researchers and they two obtained similar results [3-5].

4. CONCLUSION

144 hybrids evaluated along with check concluded that plant height and ear height play an important role in increasing the grain yield. Negative association of days to silking suggest that decreasing the silking days play a vital role in increasing the yield. Clustering of genotypes into only two clusters suggest that most of the genotypes are closely related. The results of this study helps in planning the future breeding programme.

DISCLAIMER (ARTIFICIAL INTELLIGENCE)

Author(s) hereby declare that NO generative AI technologies such as Large Language Models (ChatGPT, COPILOT, etc) and text-to-image generators have been used during writing or editing of manuscripts.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

1. FAO. World Food and Agriculture - Statistical Yearbook. Rome; 2021.
2. Angrau. Maize outlook report January to May; 2021.
3. Rigon JPG, Capuani S, Rigon CAG. Genetic divergence among maize hybrids by morphological descriptors. *Bragantia* 2015;74(2):156–160. Available: <https://doi.org/10.1590/1678-4499.0246>
4. Sabitha N, Mohan Reddy D, Lokanadha Reddy D, Hemanth Kumar M, Sudhakar P, Ravindra Reddy B, Mallikarjuna SJ. Genetic divergence analysis over seasons in single cross hybrids of maize (*Zea mays* L.). *Acta Botanica Plantae*. 2022;1(2):12–18.
5. Zaman MA, AAA. Genetic diversity in exotic maize. *Bangladesh J. Agril. Res.* 2017;38(7):335–341.
6. Searle, S. R. Phenotypic, Genetic and Environmental Correlations. *Biometrics*. 1961;17(3):474–480.
7. Olivoto T, Lúcio AD. Metan: An R package for multi-environment trial analysis. *Methods in Ecology and Evolution*. 2020;11(6):783-789.
8. Kassambara A, Mundt F. Factoextra: Extract and visualize the results of multivariate data analyses. 2020;R Package Version 1.0.7.
9. Robinson, HF., Comstock, RE. And Harvey PH. Genotypic and phenotypic correlations in corn and their implications in selection. *Agron. J.* 1951;43:262-267.
10. Johnson, HW., Robinson, HF. and Comstock, RE. Estimates of genetic and environmental variability in soybean. *Agron. J.* 1955;47:314-318.
11. Mehra, R. Tickle, AN. Saxena, A. Munjal A. Rekhakhandia. and Singh, M. Correlation, path-coefficient and genetic diversity in blackgram [*Vigna mungo*(L) hepper]. *Int. Res. J. Pl. Sci.* 2016;7(1):001-011.
12. Nemati A, Sedghi M, Sharifi RS, Seiedi MN. Investigation of correlation between traits and path analysis of corn (*Zea mays* L.) grain yield at the climate of Ardabil region (Northwest Iran). *Notulae Botanicae Horti. Agrobotanici Cluj-Napoca*. 2009;37(1):194–198.
13. Yahaya MS, Bello I, Unguwanrimi AY. Correlation and path-coefficient analysis for grain yield and agronomic traits of maize (*Zea mays* L.). *Science World Journal*. 2021;16(1):10–13.
14. Akbar M, Shabbir S, Amer Hussain, Mohammad S. Evaluation of maize three way crosses through genetic variability, broad sense heritability, character association and path analysis. *J. Agric. Res.* 2008;46(1):39-45.
15. Atnafua B, Nageshwar Rao. Estimates of heritability, genetic advance and correlation study for yield and its attributes in maize (*Zea mays* L.). *J. Pl. Sci.* 2014: 2(1):1-4
16. Soumya, H., and Kamatar, M. Correlation and path analysis for yield and yield components in single cross maize hybrids (*Zea mays* L.). *J. Farm Sci.* 2017;30(2):153–156.
17. Patel DA, Patel JS, Bhatt MM, Bhatt HM. Correlation and path analysis in Forage maize (*Zea mays* L.). *Res. Crops*. 2005; 6(3):502-504.
18. Kumar S, Shashi JP, Singh J, Singh SP. Correlation and path analysis in early generation inbreds of maize (*Zeamays* L.). *Crop Improv.* 2006;33(2): 156-160
19. Wickham H. ggplot2: Elegant Graphics for Data Analysis. Springer-Verlag New York; 2016.

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