

## **Correlation and Clustering for Yield and Its Attributing Traits in Maize (*Zea mays* L.)hybrids**

### **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 anthesis days and silking are positively correlated with grain yield. Hence selection anthesis and silking days plays a vital role in increasing yield especially in stressful environments. 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 helps in understanding the relatedness between the genotypes.

### **Introduction:**

Maize is the third most important cereal crop after rice and wheat. It is not as popular as the rice and wheat as it is not used primarily for consumption. But the popularity of maize increasing gradually as its usage is increasing gradually in feed as well as biofuel production. Maize is cultivated in about 170 countries and in India, maize is cultivated in 9.26Mha (INDIASTAT,2020-21).The commercial hybrid production was feasible after the suggestion given by D. F. Jones in 1918 proposed the use of double cross hybrid. The double cross hybrids were developed by crossing two single cross hybrids. Later in 1960's development of superior inbreds took place after rigorous research by eminent scientists. Plant breeders started developing superior inbreds. Single cross hybrids became popular among growers after 1960's. Assessment and exploitation of available genetic variability that exist in working germplasm for traits of economic importance is a short term strategy of breeding any crop. The study of inheritance of various developmental and productive traits through the correlation and clustering of genotypes which give an insight about the contribution of different characters to grain yield which is helpful for framing an effective breeding programme. Yield is a complex trait understanding the factors contributing it is beneficial for breeding superior hybrids. Knowledge on the genetic diversity has been successfully applied in earlier studies(Al-Naggar et al., 2022; Fufa et al., 2005). Genetic diversity analysis also helps in heterotic grouping of genotypes. Our study evaluated the maize hybrids for association between the traits as well as grouping of those hybrids to different clusters to assess its diversity.

### **Material and method:**

144 maize hybrids from CIMMYT, Hyderabad with 6 checks were evaluated for yield and yield attributing traits at ICRISAT, Hyderabad 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(Searle, 1961). The cluster analysis using hierarchical clustering and ward.D method was done in R software using the factoextra and ggplot2 package (Kassambara, 2017; Wickham,2023).

## Results and discussion:

### Correlation analysis:

Phenotypic and genotypic correlation among all the traits under study is mentioned in the table 1 and 2. Correlation generally helps in identifying the traits which are mutually contributing to enhance each other or impair each other. Knowledge in this aspect is crucial to aid in the selection of genotype in the breeding process. In the study of 150 genotypes of maize its clearly visible that grain yield had a significant phenotypic correlation with plant height, ear height and ear per plant. Plant height and ear height both had significant negative phenotypic correlation with anthesis days and silking days. Highest significant correlation has been observed between the anthesis days and silking days as well as plant height and ear height. In the genotypic correlation ear per plant correlation was not estimated as there was negative covariance between the traits observed explaining the traits such as grain yield, anthesis days, silking days, plant height and ear height has the negative effect. Anthesis days and silking days has a significant positive genotypic correlation with the grain yield. Plant height had a significant negative correlation with anthesis days and silking days. Highest significant positive genotypic correlation was observed between the anthesis and silking days as well as plant height and ear height. Correlation between the anthesis and silking days had been observed in the earlier studies (Azad et al., 2012; Chase and Nanda, 1967). Positive correlation between the plant height silking and anthesis days was reported in the earlier studies, although we could see there was negative correlation between all the three traits(Nzuve et al., 2014). High correlation between plant height and ear height was observed in earlier studies(Hallauer et al., 2010; Nzuve et al., 2014; Salami, 2002). Positive contribution of plant height and ear height was reported in earlier studies(Nzuve et al., 2014). By observing the correlation study we can say the anthesis days and silking days has a significant contribution on grain yield. In stress breeding studies its important trait of selection.

Table 1: Phenotypic correlation among the traits

Traits	GY	AD	SD	PH	EH
AD	0.07				
SD	0.08	0.93**			
PH	0.18*	-0.33**	-0.33**		
EH	0.26**	-0.25**	-0.26**	0.69**	
EPP	0.27**	-0.14	-0.09	0.00	-0.02

\*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	0.21*				

SD	0.24**	0.99**			
PH	0.03	-0.25**	-0.22**		
EH	0.07	0.02	0	0.62**	
EPP	NA	NA	NA	NA	NA

\*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

**Cluster:**

The genotypes are divided into two clusters and further subdivide into four sub clusters. On keen observation of division of 150 genotypes into two main clusters it was observed that low yielding, late flowering, lower plant height, lower ear height and lower number of ears per plant genotypes are grouped in one cluster and higher yielding, early flowering, higher plant height and ear height and higher number of ears per plant are grouped in other cluster. The clusters are formed at the height of 25. Further these genotypes are divided into two more clusters in each cluster. Predicting the ideal number of cluster by hierarchical clustering using Euclidean distance and ward.D2 method it was found ideal clustering is 2 clusters. Intracluster and intercluster distance between two clusters are mentioned in the table. Intra cluster distance was lowest in cluster I. Inter cluster distance between the cluster I and cluster II was 3.62 higher than the intercluster distance. Total of 85 genotypes are in cluster I and 65 are in cluster II. Clustering of genotypes based on the quantitative traits will assist in selection of genotypes based on the traits. In earlier studies also the clustering based on the quantitative traits was performed to access the diversity between the genotypes (Sabitha et al., 2022; Zaman and Islam, 2013). Plausible explanation of lower genetic distance as well mean difference between the two clusters explain the closer relationship among the genotypes i.e., most of the hybrids from the parents who are closely related (Subramanian and Subbaraman, 2010).

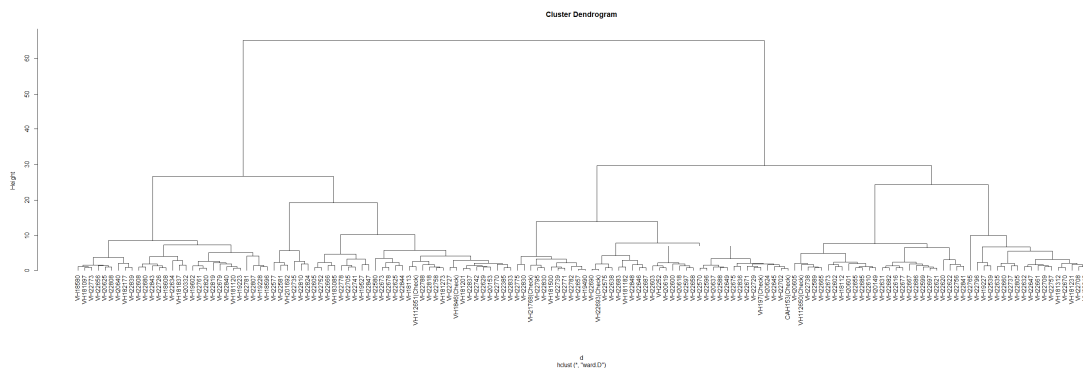


Figure 1: Dendrogram representing clustering of 150 maize hybrids

Table 3: Contribution of traits in each cluster

Character	Cluster I	Cluster II
Grain yield	7.37	6.83
Ear height	117.26	106.81
Plant height	221.62	206.88
Anthesis days	73.25	77.39

Silking days	74.82	79.25
Ear per plant	0.91	0.84

Table 4: Inter and intra cluster distance

Cluster	I	II
I	<b>2.71</b>	3.62
II		<b>2.85</b>

\*Bold values are intra cluster distance

### Reference:

Al-Naggar, A. M., Soliman, A., Hussien, M., and Mohaned, E. 2022. Genetic diversity of maize inbred lines based on morphological traits and its association with heterosis. *SABRAO Journal of Breeding and Genetics*. **54**:589–597.

Azad, M., Teixeira da Silva, J., and Biswas, B. 2012. Genetic Correlation among Various Quantitative Characters in Maize (*Zea mays* L.) Inbred Lines. *International Journal of Plant Breeding*. **6**:144–146.

Chase, S. S., and Nanda, D. K. 1967. Number of Leaves and Maturity Classification in *Zea mays* L.1. *Crop Science*. **7**(5).

Fufa, H., Baenziger, P. S., Beecher, B. S., Dweikat, I., Graybosch, R. A., and Eskridge, K. M. 2005. Comparison of phenotypic and molecular marker-based classifications of hard red winter wheat cultivars. *Euphytica*. **145**(1):133–146.

Hallauer, A. R., Carena, M. J., and Miranda Filho, J. B. de. 2010. Quantitative genetics in maize breeding. *Springer Science & Business Media*. **6**.

Kassambara, A. and Mundt, F. 2020. Factoextra: Extract and Visualize the Results of Multivariate Data Analyses. R Package Version 1.0.7.

Nzuve, F., Githiri, S., Mukunya, D. M., and Gethi, J. 2014. Genetic variability and correlation studies of grain yield and related agronomic traits in maize. *Journal of Agricultural Sciences*. **6**(9).

Sabitha, N., Mohan Reddy, D., Lokanadha Reddy, D., Hemanth Kumar, M., Sudhakar, P., Ravindra Reddy, B., and Mallikarjuna, S. J. 2022. Genetic divergence analysis over seasons in single cross hybrids of maize (*Zea mays* L.). *Acta Botanica Plantae*. **1**(2):12–18.

Salami, A. E. 2002. Responses to reciprocal recurrent selection in two early maturing complementary maize populations. PhD. Thesis, Univ. of Ibadan.

Searle, S. R. 1961. Phenotypic, Genetic and Environmental Correlations. *Biometrics*. **17**(3):474–480.

Subramanian, A., and Subbaraman, N. 2010. Hierarchical cluster analysis of genetic diversity in maize germplasm. *Electronic Journal of Plant Breeding*. **1**(4):431–436.

Wickham, H. *ggplot2: Elegant Graphics for Data Analysis*. 2016. Springer-Verlag New York,

Zaman, M. A., and Islam, M. A. 2013. Genetic diversity in exotic maize (*Zea mays* L.) hybrids. *Bangladesh Journal of Agricultural Research*. **38**(2):335–341.

UNDER PEER REVIEW