

EFFECTS OF GENETIC COMPONENTS BY ENVIRONMENTAL INTERACTIONS OF MAIZE HYBRIDS ACROSS THE DIFFERENT LOCATIONS OF KHYBER PAKHTUNKHWA, PAKISTAN

Izhar Hussain¹, Arshad Ali Kalari², Altaf Hussain Dawar³, Sajid Hussain Kalari⁴, Allah Wasayo Kalari⁵, Majid Hussain Kalari², Mohsin Khan⁶ and Irfan Ali Chandio²

¹Department of Agriculture, University of Haripur, Haripur, KPK, Pakistan.

²Department of Plant Breeding and Genetics Sindh Agriculture University, Tandojam.

³Department of Microbiology, Lasbela University of Agriculture, Water and Marine Science,

⁴Department of Soil Science, Sindh Agriculture University, Tandojam

⁵Department of Agriculture Extension, Sindh Pakistan

⁶Department of Biotechnology, Sindh Agriculture University, Tandojam

Correspondence author: Ali.breeder110@gmail.com

ABSTRACT

The effect on genetic constitution by the environment interaction was studied for 13 maize genotypes, during 2012 over four different location of Khyber Pakhtun Khwa i.e. KPK Agricultural University Peshawar, Agricultural Research Station Baffa (Mansehra), Agricultural Research Station, Jamra (Mardan) and Cereal Crops Research Institute, Pirsabak. These experiments were carried out in randomized complete block design with three replication at each location during (July-October) 2012. Data were recorded on different morphological and yield parameters. Analysis of variance indicated significant differences among the four locations for followings traits, such as days to 50% silking, days to 50% pollen shedding, anthesis silking interval (ASI), plant height, ear height, grain moisture at harvest and grain yield per hectare. All the genotypes showed significant differences for almost parameters except anthesis silking interval (ASI), which was non significant across the four different locations. The genotypes x locations interactions also showed significant differences for days to 50% silking, days to 50% pollen shedding, height of plant, height of ear, grain moisture at harvest and grain yield per hectare, while non significant difference was observed for anthesis silking interval (ASI). Based on yield performance of genotypes across the four locations, Baffa ranked first as compared to the other three locations. Hybrid WD-2x8 was the highest yielding across the four locations followed by hybrid Hi-corn, while the lowest yield was observed in hybrid CS-King at all the four locations.

Key-words: Maize, Genotype, Environment, Yield associated traits.

INTRODUCTION

Maize is one of the most important and high yielding cereal crop in the world, it is considered as a queen of cereal crop. Maize is more significant and importance food supplies those countries, where population is rapidly increasing available. Maize is one of the three important cereal crops after wheat and rice in Pakistan (Anonymous 2012a). It contributes about 5.44% of the total cultivated area and 3.5% of the value of agricultural output. It is grown on an average area of 1.02 million ha with an annual production of 3.1 million tons. The bulk (97%) of the total production comes from two provinces i.e., KPK, accounts for 57% of the total area and 68% of total production and Punjab contributes 38% acreage with 30% of total maize grain production., while on very small areas of maize is planted, which accounts for (2-3%) in the province of Sindh and Balochistan (Anonymous 2012b). The phenotype of an individual is determined by both the genotype and the environment. These two effects are not always additive which indicates that genotype x environment interactions (GEI) are present. The GEI results in inconsistent performances among the genotypes across environments. Significant GEI results from the changes in the magnitude of differences among genotypes in different environments or changes in the relative ranking of the genotypes (Crossa, 1990). The genotype x environment interaction is an important aspect of plant breeding programs. It may arise when certain genotypes are grown in diverse set of environments. A significant G x E interaction for a quantitative trait such as seed yield can seriously limit the efforts on selecting superior genotypes for both new crop production and improved cultivar development (Kang and Gorman, 1989). Genotype x environment interaction for any variety reduces the usefulness of genotype mean over all locations for selecting and advancing superior genotypes. Genotype by environment (G x E) interactions is almost unanimously considered to be among the major factors limiting response to selection and also decreases the efficiency of breeding programs. G x E interactions become important when the rank of breeding lines changes in different environments. This change

in rank is called crossover G x E interaction (Baker 1988). Genotypes by environment interactions (G x E) are a known fact in cultivar evaluations. The G x E is composed of G x Location, G x Year, and G x L x Y constituents. In general G x Y is larger than G x E, although both were observed to be nearly equivalent for switch grass biomass yield (Hopkin *et al.*, 1995). If the G x Y component is larger, then multiple year tests are needed. It is usually not considered during the early stages of testing. Later, as the cultivar/hybrid testing program becomes focused on few potential releases, testing in multiple years is essential for selecting the best and most stable entries. Therefore the aimed of this study was to evaluate 13 maize genotypes for their adaptability (G X E) of performance for yield and yield components at four different locations of KPK.

MATERIAL AND METHODS

A series of experiments were conducted at four locations of Khyber Pakhtun Khwa i.e. KPK Agricultural University (AU) Peshawar, Cereal Crops Research Institute (CCRI) Pirsabak, Agriculture Research Station (ARS) Baffa (Mansehra) and Agriculture Research Station (ARS) Jamra (Mardan). For this purpose 10 maize hybrids along an open pollinated variety (OPV) and two populations were used (Table 1).

Table 1. Maize genotypes, origin and source used in experiment during crop season 2008.

| S.No. | Genotypes | Origin | Source |
|-------|-------------|------------|---------|
| 1 | 974 (w) | Hybrid | Pioneer |
| 2 | P – 3025(w) | Hybrid | Pioneer |
| 3 | Opener (w) | Hybrid | Pioneer |
| 4 | CS – King | Hybrid | Ginocrm |
| 5 | Hi – Corn | Hybrid | Ginocrm |
| 6 | CS – 201 | Hybrid | Ginocrm |
| 7 | WD -2 x 8 | Hybrid | CCRI |
| 8 | WD – 3x 6 | Hybrid | CCRI |
| 9 | P – 30k08 | Hybrid | Pioneer |
| 10 | R – 4210 | Hybrid | Rafhan |
| 11 | SW C4 | Population | AUP |
| 12 | AZ C4 | Population | AUP |
| 13 | Jalal | OPV | CCRI |

The experiment at each location was conducted in a randomized complete block design (RCBD) and was replicated three times. Sowing was done on 20th June 2008 at KPK AU, Peshawar, 24th June 2008 at ARS Baffa, 30th June 2008 at ARS Jamra (Mardan) and on 5th July 2008 at CCRI Pirsabak. Each genotype was planted in a plot of four rows having five meters length. The plant to plant distance was 25 cm and row to row distance was of 75 cm. A basal fertilizer dose of 50 kg urea and 200 kg SSP per acre was applied. Before sowing, seeds were treated with Confidor and Imkan fungicides. Standard cultural practices essential for crop management were practiced at all locations through out the season. Data were recorded on the following parameters; Days to 50% silking, Days to 50% pollen shedding Anthesis silk interval, Plant height Ear height, Fresh ear weight at harvest Grain moisture at harvest

Grain yield per hectare . (Grain yield (kg ha⁻¹)= fresh ear wt. x (100 – grain moisture content) x 0.8 x 10000).

RESULTS

Flowering traits

Mean square values for days to 50 % silking, days to 50% pollen shedding and anthesis silking interval (ASI) are given in the Table 2, 3,4,5,6 indicated that highly significant differences were observed among the four locations for days to 50 % silking, days to 50% pollen shedding and anthesis silking interval (ASI). Highly significant differences were observed among hybrids as well across four locations for days to 50 % silking and days to 50% pollen shedding while differences were non significant for anthesis silking interval (ASI). The interaction between genotypes and locations was also highly significant for days to 50% silking and days to 50% pollen shedding. The mean values for days to 50% silking and days to 50% pollen shedding of maize genotypes at four locations are given in Table 2. It was noted that hybrid P-30K08 had the highest mean value of 58.5 days followed by Jalal (OPV) with 58.4 days for 50% silking. The shortest duration for days to 50% silking was observed for hybrid WD-3x6 with

mean value of 54.3 days. The highest mean value of 57.8 for days to 50% pollen shedding was observed for hybrid P-30K08, followed by Jalal (OPV) with mean value 57.5 days. The minimum value for days to 50% pollen shedding was observed for hybrid WD-3 x 6 i.e. 53.5 days.

Plant height (cm)

Statistical analysis of the data regarding plant height indicated highly significant differences among the four locations used in this study. The hybrids also exhibited significant variations across the four locations. The interaction between genotype and locations (G X L) was significant (Table 7). The mean values for plant height given in Table 7, indicated that hybrid P-30K08 showed highest mean value for this trait i.e. 213.9 cm while lowest mean value was expressed by hybrid WD-3X6 having a value 150.3 cm. Other hybrids showed their mean values for plant height with in this range.

Table 2. Mean values for days to 50% silking, and pollen shedding plant height and ear height of maize genotypes across four locations of KPK.

| Maize genotypes | Days to 50% Silking | Days to 50% Pollen shedding | Plant height (cm) | Ear height (cm) |
|-----------------|---------------------|-----------------------------|-------------------|-----------------|
| 974 (W) | 55.83 d-g | 55.92 bc | 180.36 b | 85.03 de |
| P-3025(W) | 56.83 b-d | 56.08 bc | 173.44 b-e | 77.39 f |
| Opener (W) | 54.83 fg | 54.42 de | 176.97 bc | 82.31 ef |
| CS King | 57.42 a-c | 57.17 ab | 163.55 ef | 81.09 ef |
| Hi Corn | 57.83 ab | 56.75 a-c | 178.90 bc | 94.41 b |
| CS-201 | 57.75 ab | 56.75 a-c | 168.79 c-e | 85.80 c-e |
| WD-2X8 | 55.00 e-g | 54.00 e | 177.16 bc | 92.62 bc |
| WD-3X6 | 54.33 g | 53.50 e | 150.34 g | 79.64 ef |
| P-30K08 | 58.50 a | 57.75 a | 213.91 a | 107.60 a |
| R-4210 | 57.50 a-c | 56.58 a-c | 173.38 b-e | 84.90 de |
| SW-C4 | 56.17 c-f | 55.42 dc | 164.52 d-f | 76.24 f |
| AZ-C4 | 56.42 c-f | 55.42 dc | 157.81 fg | 77.17 f |
| JALAL | 58.42 a | 57.50 a | 175.39 b-d | 91.03 b-d |
| LSD $p < 0.05$ | 1.527 | 1.392 | 10.93 | 7.432 |

Table 3. Means values of maize genotypes evaluated over four locations of KPK for grain moisture% at harvest and grain yield ha^{-1} during year 2008.

| Maize Hybrid | Anthesis Silking Interval | | Grain yield (kg ha^{-1}) |
|----------------|---------------------------|-----------|-------------------------------------|
| 974 (W) | 2.50 b | 27.55 c-e | 7733.5 ab |
| P-3025(W) | 2.33 b | 28.38 b-e | 7519.1 bc |
| Opener (W) | 2.42 b | 29.03 bc | 6454.2 d |
| CS King | 2.83 ab | 26.88 de | 5060.0 f |
| Hi Corn | 2.42 b | 28.58 b-d | 7972.2 a |
| CS-201 | 2.75 ab | 28.21 b-e | 6484.6 cd |
| WD-2X8 | 2.50 b | 26.74 e | 8261.5 a |
| WD-3X6 | 2.50 b | 27.26 c-e | 6232.2 e |
| P-30K08 | 2.58 ab | 30.88 a | 7466.5 bc |
| R-4210 | 3.00 a | 28.15 b-e | 6183.5 e |
| SW-C4 | 2.58 ab | 28.48 b-e | 5805.5 e |
| AZ-C4 | 2.67 ab | 27.41 c-e | 5770.9 e |
| JALAL | 2.42 b | 29.78 ab | 5934.4 e |
| LSD $p < 0.05$ | 0.5601 | 1.816 | 1256 |

Ear height (cm)

Analysis of variance for ear height indicated highly significant differences among the four locations. The genotypes also showed highly significant variations across the four locations, while the interaction between genotype and locations (G X L) was also highly significant. The mean values for ear height are given in Table 8. Highest mean value of 107.6 cm was observed for hybrid P-30K08, followed by hybrid Hi Corn with mean value of 94.4 cm, while minimum mean value for ear height was observed for genotype SW-C4 i.e. 76.24 cm. Significant differences were observed among 13 genotypes across four locations for anthesis silking interval (ASI). Therefore, the means for hybrids were not subjected to LSD test.

Table 4. Days to 50% silking comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK AU Peshawar | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|-----------------|-----------|-----------|---------------|-----------|
| 974 (W) | 60 | 51 | 55 | 57 | 55.83 d-g |
| P-3025(W) | 62 | 55 | 54 | 57 | 56.83 b-d |
| Opener (W) | 58 | 51 | 52 | 58 | 54.83 fg |
| CS King | 62 | 52 | 55 | 60 | 57.42 a-c |
| Hi Corn | 61 | 55 | 57 | 59 | 57.83 ab |
| CS-201 | 63 | 54 | 56 | 58 | 57.75 ab |
| WD-2X8 | 59 | 52 | 52 | 58 | 55.00 e-g |
| WD-3X6 | 59 | 48 | 54 | 56 | 54.33 g |
| P-30K08 | 65 | 55 | 54 | 59 | 58.50 a |
| R-4210 | 63 | 54 | 52 | 60 | 57.50 a-c |
| SW-C4 | 62 | 50 | 54 | 59 | 56.17 c-f |
| AZ-C4 | 61 | 49 | 55 | 60 | 56.42 c-f |
| JALAL | 65 | 54 | 55 | 60 | 58.42 a |
| Means | 61.46 a | 52.46 d | 54.26 c | 58.54 b | 56.67 |

Table 5. Days to 50% pollen shedding comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK AU Peshawar | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|-----------------|-----------|-----------|---------------|-----------|
| 974 (W) | 60 | 49 | 60 | 55 | 55.92 bc |
| P-3025(W) | 61 | 53 | 55 | 55 | 56.08 bc |
| Opener (W) | 57 | 49 | 54 | 57 | 54.42 de |
| CS King | 61 | 50 | 59 | 59 | 57.17 ab |
| Hi Corn | 60 | 53 | 56 | 58 | 56.75 a-c |
| CS-201 | 61 | 52 | 57 | 57 | 56.75 a-c |
| WD-2X8 | 58 | 49 | 53 | 56 | 54.00 e |
| WD-3X6 | 58 | 46 | 56 | 54 | 53.50 e |
| P-30K08 | 64 | 53 | 56 | 58 | 57.75 a |
| R-4210 | 61 | 51 | 55 | 59 | 56.58 a-c |
| SW-C4 | 60 | 48 | 56 | 58 | 55.42 dc |
| AZ-C4 | 59 | 47 | 57 | 58 | 55.42 dc |
| JALAL | 63 | 51 | 56 | 59 | 57.50 a |
| Means | 60.26 a | 50.23 d | 56.00 c | 57.28 b | 55.94 |

Grain moisture at harvest (%)

Highly significant differences were observed among the four locations for grain moisture percentage at harvest. Genotypes also showed significant differences for grain moisture at harvest. Genotype x location interaction was also significant for grain moisture at harvest. Means values of 13 maize genotypes evaluated over four locations for

the parameter of grain moisture at harvest is given in Table 9. The mean values for grain moisture percentage at harvest indicated that all the maize genotypes were not having similar grain moisture percentage and had significant differences. The maximum grain moisture percentage was observed in hybrid P-30K08 with value of 30.9 %.The minimum grain moisture percentage was observed for hybrid WD-2X8, with value of 26.7 %.

Table 6. Anthesis silking interval comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK AU Peshawar | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|-----------------|-----------|-----------|---------------|---------|
| 974 (W) | 1 | 3 | 3 | 2 | 2.50 b |
| P-3025(W) | 2 | 3 | 2 | 2 | 2.33 b |
| Opener (W) | 2 | 3 | 3 | 2 | 2.42 b |
| CS King | 3 | 3 | 3 | 2 | 2.83 ab |
| Hi Corn | 1 | 3 | 3 | 2 | 2.42 b |
| CS-201 | 3 | 3 | 3 | 2 | 2.75 ab |
| WD-2X8 | 2 | 3 | 2 | 3 | 2.50 b |
| WD-3X6 | 2 | 3 | 2 | 3 | 2.50 b |
| P-30K08 | 2 | 3 | 3 | 2 | 2.58 ab |
| R-4210 | 3 | 4 | 3 | 2 | 3.00 a |
| SW-C4 | 3 | 3 | 3 | 2 | 2.58 ab |
| AZ-C4 | 3 | 3 | 2 | 3 | 2.67 ab |
| JALAL | 2 | 3 | 2 | 2 | 2.42 b |
| Means | 2.21 d | 3.23 a | 2.67 b | 2.23 c | 2.58 |

Table 7. Plant height (cm) comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK AU Peshawar | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|-----------------|-----------|-----------|---------------|------------|
| 974 (W) | 188 | 229 | 159 | 146 | 180.36 b |
| P-3025(W) | 186 | 202 | 148 | 158 | 173.44 b-e |
| Opener (W) | 188 | 221 | 138 | 161 | 176.97 bc |
| CS King | 174 | 198 | 136 | 147 | 163.55 ef |
| Hi Corn | 195 | 212 | 149 | 160 | 178.90 bc |
| CS-201 | 173 | 203 | 144 | 156 | 168.79 c-e |
| WD-2X8 | 170 | 222 | 157 | 160 | 177.16 bc |
| WD-3X6 | 149 | 178 | 143 | 131 | 150.34 g |
| P-30K08 | 225 | 268 | 177 | 186 | 213.91 a |
| R-4210 | 182 | 206 | 148 | 157 | 173.38 b-e |
| SW-C4 | 168 | 217 | 118 | 156 | 164.52 d-f |
| AZ-C4 | 168 | 199 | 116 | 148 | 157.81 fg |
| JALAL | 190 | 220 | 143 | 148 | 175.39 b-d |
| Means | 181.16 b | 213.51 a | 144.28 d | 154.74 c | 173.422 |

Grain yield kg ha⁻¹

Mean square values for grain yield kg ha⁻¹ are given in Table 10. Highly significant differences were observed among the four locations for grain yield. Like (Charles *et al.*, 2013). genotypes also showed significant differences for grain yield. Similarly the genotype x location interaction was also significant for this parameter. Mean values for grain yield kg ha⁻¹, indicated that hybrid WD-2X8 had the maximum grain yield of 8261.5 kg ha⁻¹, followed by hybrid Hi-corn hybrid with grain yield of 7972.2 kg ha⁻¹. The minimum grain yield kg ha⁻¹ was observed for hybrid CS King with average value of 5060 kg ha⁻¹. Comparison of average grain yield ha⁻¹.of hybrids for the four

locations is given in Table 10. The data in Table 10 reveals that hybrid Hi-Corn ranked first with average grain yield of 12531 kg ha⁻¹ at Baffa (location 2), and hybrid P-30k08 ranked second with average grain yield 12141 kg ha⁻¹ at Pirsabak (location 4) followed by hybrid WD-2x8 which was third in rank with average yield of 11580 kg ha⁻¹. Maximum number of genotypes produced high yield in ARS Baffa as compared to the other three locations.

The overall means for grain yield ha⁻¹ given in table. 10 showed that maximum grain yield of 10210.15 kg ha⁻¹ was produced at ARS Baffa (Mansehra) followed by 9333 kg ha⁻¹ and 5451 kg ha⁻¹ CCRI Pirsabak and KPK AU, Peshawar, respectively with grain yield. The minimum grain yield was of 2697 kg ha⁻¹ was produced at Jamra. LSD results revealed significant differences among these values. Mean square values for grain yield ha⁻¹ also showed that all genotypes were significantly different from each other at all locations. The interaction between genotypes and locations were also highly significant for grain yield ha⁻¹.

Table 8. Ear height (cm) comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK AU Peshawar | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|-----------------|-----------|-----------|---------------|-----------|
| 974 (W) | 82 | 112 | 78 | 68 | 85.03 de |
| P-3025(W) | 69 | 101 | 68 | 72 | 77.39 f |
| Opener (W) | 81 | 106 | 68 | 73 | 82.31 ef |
| CS King | 85 | 94 | 73 | 72 | 81.09 ef |
| Hi Corn | 97 | 118 | 79 | 84 | 94.41 b |
| CS-201 | 82 | 108 | 73 | 81 | 85.80 c-e |
| WD-2X8 | 79 | 119 | 85 | 87 | 92.62 bc |
| WD-3X6 | 68 | 96 | 86 | 69 | 79.64 ef |
| P-30K08 | 112 | 142 | 81 | 95 | 107.60 a |
| R-4210 | 87 | 102 | 68 | 82 | 84.90 de |
| SW-C4 | 71 | 102 | 61 | 72 | 76.24 f |
| AZ-C4 | 72 | 104 | 56 | 76 | 77.17 f |
| JALAL | 95 | 121 | 73 | 75 | 91.03 b-d |
| Means | 83.17 b | 109.68 a | 73.05 d | 77.25 c | 85.78 |

Table 9. Percent grain moisture comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK AU Peshawar | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|-----------------|-----------|-----------|---------------|-----------|
| 974 (W) | 26 | 32 | 31 | 21 | 27.55 c-e |
| P-3025(W) | 30 | 37 | 28 | 18 | 28.38 b-e |
| Opener (W) | 28 | 34 | 31 | 23 | 29.03 bc |
| CS King | 29 | 32 | 29 | 18 | 26.88 de |
| Hi Corn | 29 | 34 | 31 | 21 | 28.58 b-d |
| CS-201 | 30 | 33 | 30 | 20 | 28.21 b-e |
| WD-2X8 | 28 | 32 | 26 | 21 | 26.74 e |
| WD-3X6 | 29 | 30 | 30 | 21 | 27.26 c-e |
| P-30K08 | 29 | 40 | 32 | 23 | 30.88 a |
| R-4210 | 29 | 33 | 29 | 22 | 28.15 b-e |
| SW-C4 | 30 | 36 | 27 | 21 | 28.48 b-e |
| AZ-C4 | 28 | 32 | 28 | 21 | 27.41 c-e |
| JALAL | 29 | 38 | 32 | 21 | 29.78 ab |
| Means | 28.84 C | 33.96 A | 29.42 B | 20.80 D | 28.25 |

Table 10. Grain yield (kg ha⁻¹) comparison of 13 maize genotypes across four different locations of KPK.

| Maize Genotypes | KPK Peshawar | AU | ARS Baffa | ARS Jamra | CCRI Pirsabak | Means |
|-----------------|--------------|----|-----------|-----------|---------------|-----------|
| 974 (W) | 6966 | | 11244 | 3352 | 9371 | 7733.5 ab |
| P-3025(W) | 4924 | | 10736 | 2837 | 11579 | 7519.1 bc |
| Opener (W) | 5028 | | 10396 | 866 | 9526 | 6454.2 d |
| CS King | 3134 | | 9179 | 2143 | 5784 | 5060.0 f |
| Hi Corn | 7372 | | 12531 | 2589 | 9397 | 7972.2 a |
| CS-201 | 5597 | | 9699 | 2430 | 8212 | 6484.6 cd |
| WD-2X8 | 5597 | | 11580 | 4859 | 11009 | 8261.5 a |
| WD-3X6 | 5454 | | 8236 | 3387 | 7852 | 6232.2 e |
| P-30K08 | 8440 | | 6456 | 2829 | 12141 | 7466.5 bc |
| R-4210 | 5499 | | 8397 | 2649 | 8188 | 6183.5 e |
| SW-C4 | 4126 | | 9280 | 1763 | 8053 | 5805.5 e |
| AZ-C4 | 4200 | | 10040 | 1549 | 7294 | 5770.9 e |
| JALAL | 4526 | | 8291 | 2807 | 8114 | 5934.4 e |
| Means | 5451.1 c | | 10210.1 a | 2620.1 d | 8963.08 b | 6811.09 |

DISCUSSION

New introduced maize crop strains are tested across the different locations in order to assess the pattern and the magnitudes of genotype x environments interactions. These genotypes x environments interactions if present for a certain trait of breeder's interest can reduce the correlation between phenotypic and genotypic values and will ultimately reduce progress from selection. Moreover, if the genotype x environment interactions is not prevailing, a single genotype can be recommended for the wider geographical area. This approach will not only lead to increase productivity, but can also considerably reduce the input cost by developing a single variety for a wider agro-ecological zone. To investigate the behavior of genotype at different locations, replicated data were recorded on yield and other agronomic traits. The data were analyzed by standard analysis of variance combined over locations and finally stability analysis was made by SAS (statistical analysis system) program (SAS, 1999). Mean square data for different parameters i.e. days to 50% silking, days to 50% pollen shedding, ASI, height of plant, height of ear, grain moisture at harvest and grain yield kg ha⁻¹ of 13 maize genotypes were used. Analysis of variance showed significant differences among the four locations for all parameters. All genotypes were significantly different from each other at four locations for days to 50% silking, days to 50% pollen shedding, anthesis silking interval, plant height, ear height, grain moisture at harvest and grain yield kg ha⁻¹. The genotype x location interactions was also highly significant for all parameters except anthesis silking interval which was not significantly different from each other at all locations. Maximum days to 50% silking and 50% pollen shedding was observed in Peshawar (location 1) while the lowest number of days to 50% silking and 50% pollen shedding were observed at Baffa (location 2). Based on yield performance of all genotypes across four locations, Baffa ranked first as compared to the other two locations. This may be due to the favorable environment at Baffa which makes it highly conducive for increased maize production. Hybrid WD-2x8 was high yielding across all four locations, followed by hybrid Hi-corn, while the lowest grain yield was observed for hybrid CS-King at all four locations. Non significant differences among the four locations were observed for anthesis silking interval by all hybrids. The maximum value for ASI was observed in Baffa (location 2) while the minimum value for ASI was observed at Peshawar (location 1). The maize hybrids used in this study indicated highly significant differences for plant height. Maximum plant height value was observed for hybrid P-30K08 while the minimum value for this parameter was shown by hybrid WD-3X6. The hybrids also showed highly significant differences for ear height. Maximum ear height was shown by hybrid P-30K08 and the lowest value was observed for the genotype SW-C4. Similar to Munawar *et al.*, (2013). highly significant differences were found among 13 genotypes for grain moisture percentage at harvest across four locations of KPK. Maximum grain moisture percentage was shown by hybrid P-30K08 and minimum grain moisture

was observed for hybrid WD-2X8.

CONCLUSION

Hence it is concluded from this study that Genotypes SW-C4, WD-2X8, 974 (W) and CS-210 had positive response to improved environmental conditions and therefore, it is recommended that these genotypes could give better performance under suitable environment.

REFERENCES

- Anonymous (2012a). *Maize, Sorghum and Millet*. Pakistan Agriculture Research Council.
- Anonymous (2012b). *Agricultural Statistics of Pakistan*. 2012. Ministry of Food, Agriculture and Livestock. Govt. Pak., Islamabad. 192.
- Baker, R.J. (1988). Tests for crossover genotype x environment interactions. *Canadian J. of P. Sci.*, 68: 405-410.
- Charles, M.K., N. M. Fredrick, O.A. George, and O.M. Odongo. (2013). genetic variability analysis for growth and yield parameters in double cross maize (*Zea mays* l.) genotypes in Kitale county of Kenya. *J. Plant Breed. Genet.* 1: 07-11.
- Crossa, J. (1990). Statistical analysis of multi-location trials. *Adv. Argon.*, 44: 55-85.
- Hopkins, A.A., K.P. Vogel, K.J. Moore, K.D. Johnson and I.T. Carlson (1995). Genotype effects and genotype by environment interactions for traits of elite switch grass populations. *Crop Sci.*, 35: 125-132.
- <http://old.parc.gov.pk/1SubDivisions/NARCCSI/CSI/msm.html>
- Kang, M.S. and D.P. Gorman (1989). Genotype x environment interactions in maize. *Agron. J.*, 81(4): 662-664.
- Munawar. M., G. Hammad and M. Shahbaz (2013). Evaluation of Maize (*Zea mays* L.) Hybrids under Different Environments by GGE Biplot Analysis. *American-Eurasian. J. Agric. & Environ. Sci.*, 13 (9): 1252-1257.
- SAS (1999). SAS/STAT User's guide. 8. Version. SAS Institute Inc. Cary. NC

(Accepted for publication October 2016)