

Evaluation of Open Pollinated Maize (*Zea mays* L.) Varieties for Mid Altitude Areas of Western Guji Zone, Southern Oromia, Ethiopia

Natol Bakala^{1,*}, Belda Idao², Ibsa Jibat²

¹Oromia Agricultural Research Institute, Bako Agricultural Research Center, Bako, Ethiopia

²Oromia Agricultural Research Institute, Yabello Pastoral and Dryland Agricultural Research Center, Yabello, Ethiopia

Email address:

natymartyko@gmail.com (N. Bakala)

*Corresponding author

To cite this article:

Natol Bakala, Belda Idao, Ibsa Jibat. Evaluation of Open Pollinated Maize (*Zea mays* L.) Varieties for Mid Altitude Areas of Western Guji Zone, Southern Oromia, Ethiopia. *Plant*. Vol. 10, No. 1, 2022, pp. 19-25. doi: 10.11648/j.plant.20221001.13

Received: December 22, 2021; **Accepted:** January 11, 2022; **Published:** January 28, 2022

Abstract: Maize is a major grain crop that is widely adaptable to many agro-ecologies across the world. The goal of the study was to see how adaptable and stable open pollinated maize varieties are in terms of grain yield and yield-related features in the Western Guji Zone's mid-altitude areas. Different genotypes perform differently in each location, which may be used to boost production. Six open pollinated maize varieties were transported from Bako National Maize Research Center and planted in RCBD with three replications at Yabello Pastoral and Dryland Agriculture Research Center's Galana and Abaya sub sites for three years. The results of the analysis of variance revealed that there was a substantial yield difference between genotypes. Gibe-2 had the greatest average grain production of 5.85 t/ha, followed by Kulani with 5.63 t/ha across years and locations, according to the combined analysis of variance. Kulani was found to be the most stable of all the varieties, whereas Gibe 2 was shown to be the most unstable. Kulani's and ABO-additive Bako's main and multiplicative interaction stability values (ASVs) were both near to zero (0.08 and 0.27, respectively), but Gibe 2's ASV was significantly higher (1.62) and deviated from zero. As a result, Kulani was stable and high yielding across settings, whereas Gibe 2 was high yielding in a single environment (unstable). So, and Kulani were recommended for cultivation in the Galana and Abaya districts of southern Oromia, as well as regions with comparable agro ecologies, while Gibe 2 was recommended for the Galana district.

Keywords: Adaptation, ASV, Genotypes, Stability, *Zea mays*

1. Introduction

Maize (*Zea mays* L.) ($2n=20$), popularly known as corn, belongs to the Poaceae family and is a versatile crop that adapts well to a broad range of production settings [1, 2]. In terms of growing area, output, and grain yield, maize is the world's third most significant crop, trailing only rice and wheat [3] and it is important basic crop of trade product and recurring ingredient for millions of people in sub-Saharan Africa [4]. Maize is a versatile crop that adapts well to a wide range of production circumstances [5]. Maize is an important commodity in the global economy, and it is frequently traded [6]. Maize is one of the staple foods in Ethiopia, whose importance in consumption as well as production has significantly increased [7].

In Ethiopia average maize grain yield is low due to problems like insect pest damage, lack of high yielding varieties and poor crop management practices. The most important problem reported by farmers in Western Guji Zone was the lack of adaptable maize varieties and majority of the farmers in the area are growing local varieties. So far, no effort has been made in the zone to introduce and adapt improved maize varieties.

Genotype by environment interactions is the most importance to the plant breeder in selecting appropriate variety for appropriate environmental condition. Different genotypes may perform differently in each location, which may be leveraged to increase production. Variability in grain yield is due to difference in genetic potential among genotypes and environment effect. Grain yield is quantitative in nature, which usually exhibits GEI, which necessitates evaluation in multi-

environment trials before doing advanced selection [8]. Due to cross interaction, the existence of genotype by environment interaction (GEI) commonly alters the rankings of varieties in various environments, making appropriate selection challenging. As a result, analyzing and conversing genotype by environment interactions is critical for obtaining knowledge on genotype adaptability and stability. The AMMI (Additive Main Effects and Multiplicative Interaction) approach is a frequently utilized way for analyzing GE interaction as a measure of stability and adaptability [9]. In multiplication varietal trials, the AMMI model is a better model for analyzing GxE interactions [10]. It not only gives an estimate of each

genotype's overall GxE interaction impact, but also divides it into environment-related interaction effects. As a result, the current study will use open pollinated maize cultivars to evaluate the adaptability and stability of grain yield and yield-related features in the Western Guji Zone's mid-altitude areas.

2. Materials and Methods

2.1. Description of the Study Areas

The experiment was conducted at Galana and Abaya districts of Western Guji Zone, southern Oromia (Table 1).

Table 1. Descriptions of the study area.

| Sites | PH | Altitude | Available P. in ppm | CEC meq/100g soil | Texture | | | Soil class | Mean annual rainfall (mm) | Mean annual temperature (°C) |
|--------|------|----------|---------------------|-------------------|---------|-------|-------|-----------------|---------------------------|------------------------------|
| | | | | | %sandy | %clay | %silt | | | |
| Abaya | 5.68 | 1480 | 1.42 | 20.40 | 50 | 30 | 20 | Sandy clay loam | 850 | 16-36 |
| Galana | 5.84 | 1670 | 1.84 | 60.60 | 48 | 20 | 32 | Clay loam | 950 | 14-34 |

Source: mereology station and soil analysis.

2.2. Descriptions of Experimental Materials and Design

Six open pollinated maize varieties (table 2) were imported from Bako National Maize Research Center and grown for three years at Galana and Abaya subsites of Yabello Pastoral and Dryland Agricultural Research Center. There were three

replications of a totally randomized block design. The plants were cultivated in accordance with agronomic recommendations. Each plot was 12.6 m² in size, with 6 rows of 3 m length and spacing of 25cm x 75cm. Each plot was 8.4 m² in size, with four (4) rows in the centre. For all genotypes at each location, the prescribed fertilizer rate was applied.

Table 2. Lists and description of materials used in an experiment.

| Variety | Year of release | Altitude (masl) | Rain fall (mm) | Maturity date | Yield on research field (kg/ha) | Production status |
|----------|-----------------|-----------------|----------------|---------------|---------------------------------|-------------------|
| Kulani | 1995 | 1700-1400 | 1000-1200 | 150 | 6000-7000 | Under production |
| ABO-Bako | 1985 | 500-1000 | 1000-1200 | 150 | 5000-7000 | Under production |
| Gutto | 1988 | 1000-1700 | 800-1200 | 126 | 3000-5000 | Under production |
| Gibe 1 | 2001 | 1000-1800 | 1000-1700 | 145 | 6000-7000 | Under production |
| Gambella | 2002 | 500-1000 | 1000-1200 | 110 | 6000-7050 | Under production |

Source: (EARO, 2004).

2.3. Collected Data

Plant height, ear length, ear height, number of kernels per row, and number of rows per cob were gathered in plots, whereas all agronomic data such as days to physiological maturity, hundred kernel weights, and grain yield were obtained in plants.

2.4. Data Analysis

The PROC ANOVA program in SAS software was used to perform analysis of variance for phenological, yield, and yield-related data, with genotypes treated as fixed effects and replication within environment as a random effect, as stated by Gomez and Gomez [11]. Least significant different (LSD) was used for mean separation ($P < 0.05$). Adaptability and stability of the genotypes was estimated using the Genstat 15th edition. G x E biplots were generated to evaluate the genotypes simultaneously for yield and stability. ASV (AMMI Stability Values) were estimated for both genotypes and environments. The G E interaction was studied using the AMMI (Additive Main Effect and Multiplicative Interaction)

model, which combines traditional analysis of variance and principal component analysis. The contribution of each genotype and environment to the G x E interaction in the AMMI model is measured using a biplot graph presentation in which yield means are displayed against IPCA (Interaction Principal Component Axis) scores) [12]. The AMMI model is:

$$Y_{ij} = \mu + g_i + e_j + \sum \lambda_k + \alpha_{ik} y_{jk} + R_{ij}$$

Where, Y_{ij} is the yield of i^{th} genotypes in j^{th} environment; μ the overall mean; g_i is the effect of the i^{th} genotype; e_j is the effect of the j^{th} environment; λ_k is the square root of the eigenvalue of the PCA axis k . Then α_{ik} and y_{jk} are the principal components scores for PCA (Principal Component Axis) k of the i^{th} genotype and j^{th} environment, respectively, and R_{ij} is the residual. Environment and genotype PCA scores are expressed as unit vector times the square root of λ_k (environment PCA score = $\lambda_k 0.05 y_{ik}$, genotype PCA score = $\lambda_k 0.05 \alpha_{ik}$) [10]. AMMI stability value (ASV) was computed for each genotype in order to rank the genotypes utilized in this study in terms of stability [13] as follows:

$$ASV = \sqrt{\left[\frac{(IPCA1\ scores) \times (IPCA1ss)}{IPCA1SS}\right]^2 + [IPCA2score]^2}$$

Where, ASV=AMMI Stability Value; IPCA1SS = Interaction Principal Component Axis 1 sum of squares; IPCA1score = Interaction Principal Component Axis 1 score; IPCA2score = Interaction Principal Component Axis 2 score.

3. Results and Discussion

3.1. Days to Maturity

An analysis of variance revealed that there is a significant difference between types in days to maturity (P0.01) for three consecutive years for both sites. Kulani was late mature among all genotypes which took 153 days and 150.67 days in 2014 and 2015 cropping season respectively. Gambella matured earlier than other genotypes under study for Galana site (table 3). At Abaya site, Kulani was late maturing one among all genotypes under study, which took 150.67, 152.00 and 151.67 days in 2014, 2015 and 2016 cropping season respectively. Gambella variety took 103.67, 111.33 and

110.32 days to mature in 2014, 2015 and 2016 cropping seasons. (Table 3) Bakala et al. [14] found significant differences among genotypes in their study of high land maize evaluation.

3.2. Grain Yield

Grain yield differed significantly between genotypes in all cropping seasons, according to the analysis of variance. The higher grain yield was obtained from Kulani 6.48t/ha in 2014, 4.65t/ha in 2015 and 5.01t/ha in 2016 cropping season while, the lowest grain yield was obtained from Gutto LMS 4.31t/ha, 4.28t/ha and 4.28t/ha in 2014, 2015 and 2016 cropping seasons respectively (table 3). The yield variability observed among genotype, showed the potential of the variety and specific adaptability of the genotype. For Abaya site, maximum grain yield was obtained from Gibe 2 (5.08, 6.50 and 6.71t/ha in 2014, 2015 and 2016 cropping season respectively. Bassa and Goa [15] reported that different maize varieties produce significantly different grain yields at different locations over years. Taye et al. [16] also reported significant yield difference among diifferent maize genotypes.

Table 3. Mean performance of days to maturity and grain yield for Abaya and Galana site.

| Varieties | Days to maturity (days) | | | | | | Mean | Grain yield (t/ha) | | | | | | Mean |
|-----------|-------------------------|---------|---------|------------|---------|---------|---------|--------------------|---------|--------|------------|---------|---------|---------|
| | Galana site | | | Abaya Site | | | | Galana site | | | Abaya Site | | | |
| | 2014 | 2015 | 2016 | 2014 | 2015 | 2016 | | 2014 | 2015 | 2016 | 2014 | 2015 | 2016 | |
| Gibe-1 | 143b | 141.67b | 143.67c | 141.67b | 146b | 146.33b | 144.83b | 5.21b | 4.52ab | 4.35b | 4.52ab | 4.47c | 6.28a | 5.42c |
| Gibe-2 | 132c | 135.00c | 137.67d | 135.00c | 139c | 137.00c | 136.17c | 5.41b | 4.65a | 4.57b | 4.65a | 6.50a | 6.65a | 5.85a |
| ABO-Bako | 152a | 145.67b | 152.33a | 145.67b | 149ab | 150.67a | 149.83a | 5.44b | 4.25c | 4.68ab | 4.25c | 4.57c | 5.49b | 4.93e |
| Gutto-LMS | 124d | 127.33d | 126.67e | 127.33d | 125d | 126.00d | 125.83d | 4.31c | 4.28bc | 4.28b | 4.28bc | 4.26c | 4.21c | 4.03f |
| Gambella | 110e | 103.67e | 110.67f | 103.67e | 111.33e | 110.32e | 110.11e | 5.54b | 4.52ab | 4.67ab | 4.52ab | 4.15c | 6.20a | 5.16d |
| Kulani | 153a | 150.67a | 147.67b | 150.67a | 152a | 151.67a | 151.06a | 6.48a | 4.42abc | 5.01a | 5.08a | 5.47b | 6.71a | 5.63b |
| LSD | 7*** | 4.45*** | 3.48*** | 4.45*** | 4.13*** | 3.34*** | 1.45*** | 0.76*** | 0.24** | 0.42* | 0.24** | 0.42*** | 0.59*** | 0.22*** |
| CV (%) | 1.88 | 2.27 | 1.40 | 2.27 | 1.66 | 1.34 | 1.60 | 4.41 | 2.92 | 5.03 | 2.92 | 4.75 | 5.48 | 6.45 |

*, **, *** = significant at P < 0.05, at P < 0.01 and at P< 0.001, respectively, ns = non-significant. DM=days to maturity, Yld=grain yield, LSD=least significant difference, CV=coefficient of variance.

3.3. Combined Analysis of Variance

Combining analysis of variance (ANOVA) across sites for grain yield revealed a significant in genotype location interaction, indicating that genotype x environment interactions affected maize genotype yield performance. Similarly, Anley et al. [17] reported different genotypes perform differently for yield and yield related traits under different environmental conditions.

Combined analysis of variance showed that a very highly significant (P<0.0001) variation was observed between

genotypes, environment and the genotypes x environment interaction for plant height, ear height, Cob diameter, hundred seed weight and grain yield (Table 4). This indicated that the varieties and the test environments are variable, and the varieties performed differently across locations and years for almost all traits. Combined analysis of variance indicated that genotypes and environment showed significant effect (P<0.05) while G x E had non-significant effect on number of rows per cob and number of seeds per row. Traits less affected by environments are high heritability [18].

Table 4. Over all mean of maize genotypes for yield, yield related traits and phonological growths.

| Varieties | Traits mean | | | | | | | |
|-----------|-------------|---------|---------|-----------|---------|-----------|---------|------------|
| | DM (days) | PH (cm) | EH (cm) | NRPC (no) | CD (cm) | NSPR (no) | HSW (g) | YLD (t/ha) |
| Gibe-1 | 144.83b | 144.28c | 97.48ab | 13.69ab | 4.39c | 35.26ab | 31.72b | 5.42c |
| Gibe-2 | 136.17c | 178.39a | 90.41c | 13.33bc | 4.32c | 34.04abc | 31.67bc | 5.85a |
| ABO-Bako | 149.83a | 181.57a | 94.73bc | 13.39b | 4.34c | 35.56ab | 29.58cd | 4.93e |
| Gutto-LMS | 125.83d | 163.53b | 89.56c | 12.44c | 5.87b | 33.56bc | 29.03de | 4.03f |
| Gambella | 110.11e | 180.78a | 95.35bc | 14.44a | 4.27c | 35.83a | 27.64e | 5.16d |
| Kulani | 151.06a | 181.74a | 103.60a | 13.33bc | 6.33a | 32.97c | 34.00a | 5.63b |

| Varieties | Traits mean | | | | | | | |
|------------------------|-------------|------------|-----------|-----------|----------|-----------|----------|------------|
| | DM (days) | PH (cm) | EH (cm) | NRPC (no) | CD (cm) | NSPR (no) | HSW (g) | YLD (t/ha) |
| Significance level | | | | | | | | |
| Genotype | 4568.84*** | 4119.92*** | 470.81*** | 7.44** | 15.45*** | 24.88* | 9.36*** | 5.52*** |
| Environment | 27.55*** | 822.90*** | 296.06** | 19.04*** | 38.68*** | 307.08*** | 66.07*** | 7.87*** |
| Genotype x Environment | 12.84** | 327.37*** | 267.90*** | 1.85ns | 16.35*** | 13.11ns | 42.24*** | 1.20*** |
| Error | 4.83 | 113.91 | 85.13 | 1.96 | 0.15 | 9.36 | 6.85 | 0.09 |
| CV | 6.22 | 1.61 | 9.69 | 10.13 | 7.86 | 8.86 | 8.57 | 5.61 |

*, **, *** = significant at $P < 0.05$, at $P < 0.01$ and at $P < 0.001$, respectively, ns = non-significant. DM=days to maturity, PH=plant height, EH=ear height, NRPC=number of rows per cob, NSPR=number of seeds per row, HSW=hundred seed weight, CD=cob diameter and Yld=grain yield.

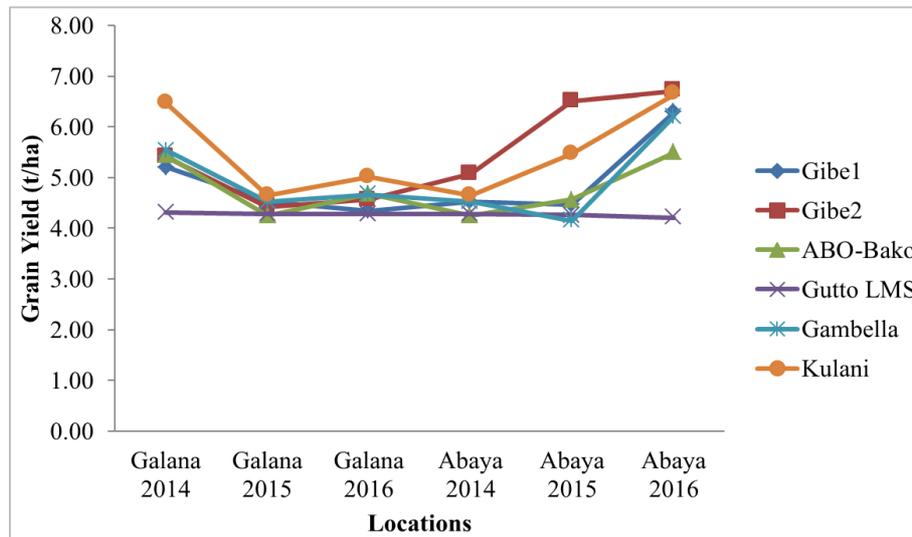


Figure 1. The performance of maize varieties across three years at Abaya and Galana sub sites.

GEI exists when the response standards for various genotypes are not parallel [19]. Cross-over interaction is a sort of GEI in which the genotypes' ranking varies depending on the environment [20]. There was a rank change of varieties across years which may mean the presence of crossover interaction. Kulani, Gibe 2 and Gambella were relatively well performed and high yielder across years, while Gutto LMS showed poor but consistent yield performance across years and locations (Figure 1). There was a change in rank of genotypes across years which may suggest the presence of crossover interaction. Similarly, Akbar *et al.* [21], Rehman *et al.* [22] reported significant differences among maize cultivars for grain yield under different environmental condition.

3.4. Stability Analysis

The AMMI analysis gives a graphical representation of

the main impact and interaction effect information for genotypes and environments on the same graph. Variance in yield data revealed that all three components genotype (G), environment (E), and G x E interaction were very significant, showing a wide range of variety occurred across varieties, location, and seasonal fluctuations (Table 5). Further, the mean squares from AMMI analysis indicated variation among G, E and G x E interaction showed highly significant different level at ($P < 0.01$) (Table 6). G x E interaction was further partitioned into two principal component analysis axis (IPCA) interactions. This variability was may be due to larger dissimilarity in rainfall, number of rainy days in each environment and high variation in mean sunshine hours among the environments. Several authors also reported supportive results [4, 23, 24].

Table 5. Combined analyses of variance using AMMI Model.

| Source of variation | df | Sum Square | Mean Square | Sum Squares Explained | |
|------------------------|-----|------------|-------------|-----------------------|---------|
| | | | | % total | % G x E |
| Total | 107 | 101.25 | 0.5 | | |
| Genotypes | 5 | 27.6 | 5.52** | 27.26 | |
| Environments | 5 | 39.36 | 7.87** | 38.87 | |
| Genotype x environment | 25 | 26.22 | 1.05** | 25.90 | |
| IPCAI | 9 | 13.63 | 1.51** | | 51.98 |
| IPCAC | 7 | 7.47 | 1.07** | | 28.49 |
| Residuals | 9 | 5.13 | 0.57** | | |
| Error | 60 | 5.13 | 0.09 | | |

** $P < 0.01$.

Table 6. IPCA 1 and IPCA 2 scores, genotypes mean and six open pollinated maize varieties tested at six locations.

| Genotypes | Graph ID | Genotype mean | IPCAG [1] | IPCAG [2] | ASV |
|------------|----------|---------------|-----------|-----------|------|
| ABO-Bako | ABO | 4.93 | 0.46 | 0.01 | 0.29 |
| Gambella | Gambella | 5.16 | 0.40 | 0.52 | 0.49 |
| Gibe-1 | Gibe1 | 5.42 | -0.61 | 0.54 | 0.79 |
| Gibe-2 | Gibe2 | 5.85 | -0.92 | -0.69 | 1.62 |
| Gutto-LMS | Gutto | 4.30 | 0.74 | -0.68 | 1.20 |
| Kulani | Kulani | 5.63 | -0.06 | 0.28 | 0.08 |
| Grand mean | | 5.22 | | | |

IPCA=Interaction Principal Component Axis, ASV=AMMI stability value.

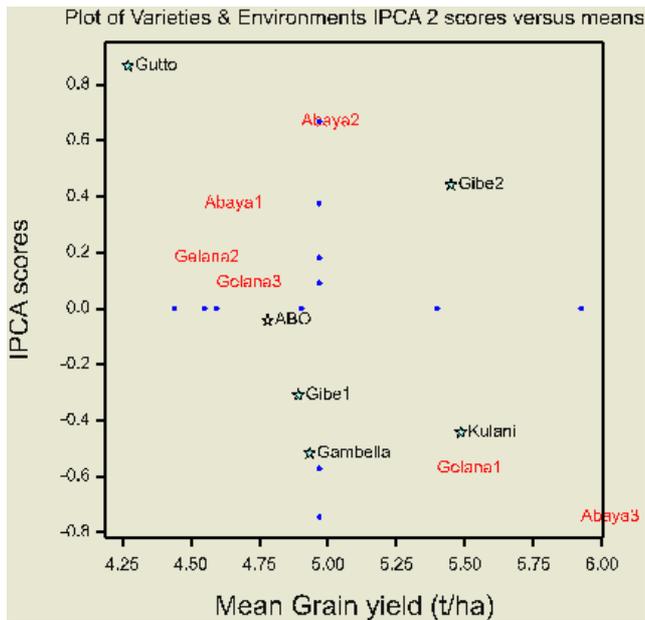


Figure 2. AMMI-1 model for grain yield (t/ha) showing the means of genotypes (numbers) and environments (upper case letter followed by number).

By plotting both the genotypes and the environments on the same graph, the relationship between genotypes and the environments can be seen clearly. The larger the IPCA scores, either positive or negative, as it is a comparative value, the better specifically a genotype is adapted to certain environments (Table 6). The more IPCA scores approximate to zero, the more stable the genotype is to overall environments. Accordingly, Kulani has IPCA value close to zero, high yielder and stable genotypes across environments (Table 6). ABO-Bako was a variety IPCA value relatively close to zero (stable) but gave lower yield below the average. Whereas, Gibe 1 and Gibe 2 were high yielder but, relatively higher IPCA score value deviating from zero, indicating that these varieties were not stable and thus adaptable for specific environment. The genotypes with high ASV were most unstable and while genotypes with low ASV were stable. The ASV indicated that the genotypes Kulani and ABO were most stable across environments and the genotypes, Gibe 2 and Gutto-LMS performed superiorly in certain environments (table 6). The sign of the scores indicate the pattern of interaction of the genotypes across environments and the reaction of environments for the different genotypes. Genotypes and environments with similar sign of IPCA1 scores interact positively for yield (tone/ha). But,

if they have opposite sign of IPCA1 scores, their interaction is negative and the environment is not favorable for the genotype [10, 20]. Similar results were reported by Souza et al [26], Anley et al [17], Abera et al [25].

AMMI 2 bi-plot: the AMMI 2 bi plot with IPC1 in X-axis and IPC2 in the Y-axis were plotted below (figure 3). The first interaction principal component (IPC1) explained 73.19% and the second interaction principal component (IPC2) explained 14.79% of the sum of square of GEI. The two IPC's cumulatively explained about 87.98% of the sum of squares of GEI (figure 3). Purchase [27] stated that, genotypes close to origin are stable while those far from origin are considered to as unstable genotypes. In terms of adaptability, the genotype closest to a given vector in any environment is more adaptable to that environment, whereas the genotype furthest from a given vector in any environment is less adaptable to that environment [28]. Inline to the following principle Gibe 2 was adaptable to Abaya2 and Abaya1 environments while Kulani and Gibe 1 were adaptable to Abaya3, Galana1 and Galana3 environments (figure 3).

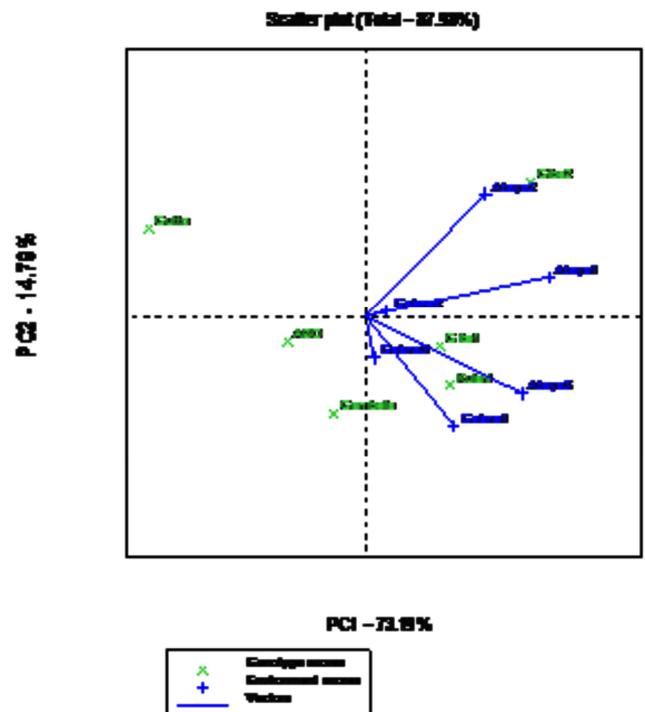


Figure 3. AMMI-2 model for grain yield (t/ha) showing the IPCA scores of open pollinated maize genotypes (numbers) planted across environments (upper cases followed by numbers).

4. Conclusion

In the West Guji zone, six open pollinated maize cultivars were evaluated for adaptation and grain yield performance. An analysis of variance for phenological, yield, and yield-related characteristics across locations and years revealed a significant difference across genotypes. The results indicated that the Kulani variety produced the highest yield at Abaya, followed by the Gambella variety. Gibe 2 was high yielder at Galana followed by Kulani. The result from stability analysis revealed that Kulani was high yielder and stable across test environments relative to other genotypes. ABO-Bako was also relatively stable but gave lower yield below the average. Gibe-1 and Gibe-2 varieties were high yielder but adaptable for specific environment. ASV analysis showed that Kulani and ABO-Bako were most stable across environments but Gibe 2 performed superiorly in certain environments. Generally, Kulani was recommended for wider adaptability, but Gibe 2 showed specific adaptability and recommended for specific area.

References

- [1] Gerpacio R, Prabhu P. 2007. Tropical and subtropical maize in Asia: production systems, constraints, and research priorities. CIMMYT.
- [2] Riedelshemer C, Czedik-Eysenberg A, Grieder C, Lisek J, Technow F, Sulpice R, Altmann T, Stitt M, Willmitzer L, Melchinger AE. 2012. Genomic and metabolic prediction of complex heterotic traits in hybrid maize. *Nature genetics*. 1; 44 (2): 217-220.
- [3] Shiri MR. 2013. Grain yield stability analysis of maize (*Zea mays* L.) hybrids under different drought stress conditions using GGE biplot analysis." (2013): 107-112.
- [4] Nzuve F, Githiri S, Mukunya DM, Gethi J. 2013. Analysis of genotype x environment interaction for grain yield in maize hybrids. *Journal of Agricultural Science*, 5 (11), 75.
- [5] FAO (Food and Agriculture Organization of the United Nations).2015. FAOSTAT, Retrieved November 26, 2016 from <http://faostat3.fao.org/download/Q/QC/E>.
- [6] FAO (World Food Organization).2016. The State of Food Insecurity in the World: Undernourishment around the World in 2016. <http://faostat.fao.org/default.aspx>. Accessed 20, September 2016.
- [7] Jaleta, Moti, Menale Kassie, Paswel Marennya, Chilot Yirga, and Olaf Erenstein. "Impact of improved maize adoption on household food security of maize producing smallholder farmers in Ethiopia." *Food Security* 10, no. 1 (2018): 81-93.
- [8] Khalil I, Shah S, Ahmad H. 2010. Stability analysis of maize hybrids across North West of Pakistan. *Pak. J. Bot*, 42 (2), 1083-1091.
- [9] Annicchiarico, P., 1997. Additive main effects and multiplicative interaction (AMMI) analysis of genotype-location interaction in variety trials repeated over years. *Theoretical and applied genetics*, 94 (8), pp. 1072-1077.
- [10] Zobel, Richard W., Madison J. Wright, and Hugh G. Gauch Jr. "Statistical analysis of a yield trial." *Agronomy journal* 80, no. 3 (1988): 388-393.
- [11] Gomez K, and Arturo G. 1984. Statistical procedures for agricultural research. John Wiley & Sons.
- [12] Pretorius M, Allemann J. Smith M. 2015. Use of the AMMI model to analyse cultivar-environment interaction in cotton under irrigation in South Africa. 2 (2): 76–80.
- [13] Purchase JL. 1997. Parametric analysis to describe genotype x environment interaction and yield stability in winter wheat." PhD diss., Ph. D. Thesis, Bloemfontein, South Africa.
- [14] Bakala N, Abate B, Nigusie M. "Standard Heterosis of Maize (*Zea mays* L.) Inbred Lines for Grain Yield and Yield Related Traits at Southern Ethiopia, Hawassa". *American-Eurasian J. Agric. and Environ. Sci.*, 17: 2017. 257-264.
- [15] Bassa D, Goa Y. 2016. Performance Evaluation and Adaptation of Improved Maize (*Zea mays* L) Varieties for Highland of Alichu, Silti and Analemo Districts of Southern Ethiopia. *Journal of Natural Sciences Research*.
- [16] Taye T, Bekele N, Shimalis Y. 2016. Evaluation of highland maize at Bule hora District of Southern Oromia, Southern Ethiopia. *African Journal of Agricultural Research*. 25; 11 (34): 3178-3181. <http://www.academicjournals.org/AJAR>
- [17] Anley W, Zeleke H, Dessalegn Y. "Genotype X environment interaction of maize (*Zea mays* L.) across North Western Ethiopia". *Journal of Plant Breeding and Crop Science*, 5: 2013. 171-181.
- [18] Epinat-Le Signor C, Dousse S, Lorgeou J, Denis JB, Bonhomme R, Carolo P, Charcosset A. 2001. Interpretation genotype - environment interaction for early maize hybrids over 12 years. *Crop Sci*. 41: 663–669.
- [19] van Eeuwijk FA, Malosetti M, Yin X, Struik PC, Stam P. 2005. Statistical models for genotype by environment data: from conventional ANOVA models to eco-physiological QTL models. *Australian Journal of Agricultural Research*, 56 (9): 883-894.
- [20] Crossa I. 1990. Statistical analysis of multi-location trials. *Adv. in Agron*. 44, 55-85.
- [21] Akbar, Mohammad Muzahid, and Noorjahan Parvez. "Impact of service quality, trust, and customer satisfaction on customers loyalty." *ABAC journal* 29, no. 1 (2009).
- [22] Rehman, Abdul, M. Farrukh Saleem, Muhammad Ehsan Safdar, Safdar Hussain, and Naeem Akhtar. "Grain quality, nutrient use efficiency, and bioeconomics of maize under different sowing methods and NPK levels." *Chilean journal of agricultural research* 71, no. 4 (2011): 586.
- [23] Kumar P, Singh NK. 2015. Determining behavior of maize genotypes and growing environments using AMMI statistics. *SAARC Journal of Agriculture*. 15; 13 (1): 162-173.
- [24] Miah MA, Ahmed S, Uddin MS. 2016. Assessment of yield stability of maize inbred lines in multi-environment trials. *Bangladesh Journal of Scientific and Industrial Research*. 51: 61-68.
- [25] Abera W, van Rensburg JB, Labuschagne MT, Maartens H. "Genotype-environment interactions and yield stability analyses of maize in Ethiopia", *South African Journal of Plant and Soil*, 21: 2004. 251-254.

- [26] Souza FR., Ribeiro PH, Veloso CA, Corrêa LA. 2002. Yielding and phenotypic stability of corn cultivars in three municipal districts of Para State, Brazil. *Pesquisa Agropecuária Brasileira*, 37: 1269-1274.
- [27] Purchase, J. L., 1997. *Parametric analysis to describe genotype x environment interaction and yield stability in winter wheat* (Doctoral dissertation, University of the Free State).
- [28] Senguttuvel, P., Sravanraju, N., Jaldhani, V., Divya, B., Beulah, P., Nagaraju, P., Manasa, Y., Prasad, A. S., Brajendra, P., Gireesh, C. and Anantha, M. S., 2021. Evaluation of genotype by environment interaction and adaptability in lowland irrigated rice hybrids for grain yield under high temperature. *Scientific Reports*, 11 (1), pp. 1-13.