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## GGE BIPLLOT ANALYSIS OF YIELD STABILITY FOR ANDEAN DRY BEAN ACCESSIONS GROWN UNDER DIFFERENT ABIOTIC STRESS REGIMES IN TANZANIA

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### ABSTRACT

Large seeded Andean dry beans (*Phaseolus vulgaris* L.) are most preferred in Africa. However, low soil fertility and increasing drought stress conditions due to climate change are among the challenges faced by farmers in this region. The purpose of this study was to identify Andean genotypes with yield stability and to identify the best environments for selection of this trait. GGE-Biplot analysis was used to examine 183 Andean bean genotypes in Tanzania. Results indicated significant environment (E), genotype (G) and genotype x environment (GE) effects for seed yield. The environment explained 46%, genotype 20%, and GE 34% of the total (G + E + GE) variation. Two principal components explained 41.21% (PC1) and 26.35% (PC2) of GGE sum of squares. The Andean genotypes, ADP-102 (Jessica)- purple mottle, large seed, bush plant habit, released by Selian Agriculture Research Institution (ARI) in Tanzania; ADP-220 (G5625)- red, large seed, vine plant habit, from Central America; ADP-276 (G13654) - brown, medium seed, vine, from Mexico; and ADP-648 ('Red Kloud') large red kidney, with bush plant habit, were stable across environments and can be recommended for general adaptation across environments. An individual location for selection for yield stability was not identified, thus justifying the continued need for multiple test sites.

**Key Words:** Drought stress, low soil fertility stress, *Phaseolus vulgaris*, yield stability

### RÉSUMÉ

Les haricots secs andins à grosses graines (*Phaseolus vulgaris* L.) sont les plus préférés en Afrique. Cependant, la faible fertilité du sol et le stress croissant de la sécheresse causé par le changement climatique font partie des défis auxquels sont confrontés les agriculteurs de cette région. L'objectif de cette étude était d'identifier les génotypes andins avec une stabilité de rendement et d'identifier les meilleurs environnements pour la sélection de ce trait. L'analyse GGE-Biplot a été utilisée pour examiner

183 génotypes de haricot andin en Tanzanie. Les résultats ont montré des effets significatifs sur l'environnement (E), le génotype (G) et le génotype x environnement (GE) sur le rendement en grain. L'environnement expliquait 46%, le génotype 20% et l'EG 34% de la variation totale (G + E + GE). Deux composantes principales expliquent 41,21% (PC1) et 26,35% (PC2) de la somme des carrés GGE. Les génotypes andins, ADP-102 (Jessica) - marron violet, grosse graine, habitude de plante de brousse, publiés par l'Institut de recherche sur l'agriculture Selian (ARI) en Tanzanie; ADP-220 (G5625) - Rouge, grosse graine, habitude de plante de la vigne, d'Amérique centrale; ADP-276 (G13654) - brun, graine moyenne, vigne, en provenance du Mexique; et le grand rein rouge ADP-648 («Red Kloud»), avec l'habitude des plantes de brousse, étaient stables dans tous les environnements et pouvaient être recommandés pour une adaptation générale dans tous les environnements. Un emplacement individuel pour la sélection de la stabilité du rendement n'a pas été identifié, ce qui justifie le besoin de la continuité de test multiples des sites.

*Mots Clés:* Stress de sécheresse, stress lié à la faible fertilité du sol, *Phaseolus vulgaris*, stabilité du rendement

## INTRODUCTION

Dry bean (*Phaseolus vulgaris* L.) productivity is highly influenced by climatic conditions and deviations from the optimal conditions (Vandemark *et al.*, 2014). Bean productivity becomes even more limited under stressful environmental conditions caused by low fertility or periodic drought. Dry bean cultivars with tolerance to these stresses will be needed for sustainable bean production (Ghassemi *et al.*, 1995; Pessaraki and Szabolcs, 1999). Progress to improve yield of common bean under low fertility or drought stress has been slow compared to research conducted to improve common bean for various agronomic traits (Kelly *et al.*, 1998).

The larger-seeded dry beans of Andean origin are preferred in Africa (Wortmann *et al.*, 1995). These beans are even less stress tolerant and less broadly adapted to different environments, compared to dry bean of Middle American origin (Singh, 1999; Beebe *et al.*, 2013). Contributing reasons can be the narrower genetic diversity (Beebe *et al.*, 2001) and determinate bush growth habit prominent in Andean beans. Determinacy usually results in more synchronised and shorter flowering duration, which results in reduced potential to recover from stressful events. Andean beans with vine growth habits exist primarily as landraces, which are less adapted to warmer

lowland environments (Beebe *et al.*, 2001). Therefore, it is important to study the performance of different Andean genotypes with different growth habits across different environments to help select the best genotypes for target regions.

The study of Genotype x Environment (GE) interaction is important because soil fertility and climate change are global concerns. Basically, GE interaction occurs when genotypes respond differently in different environments. It is well known that significant GE influences the selection of genotypes for either broad adaptation across environments, or narrow adaptation for specific environments. A significant GE interaction indicates that the genotype effect depends on the environment and each genotype responds to environmental variation differently (Julius *et al.*, 2015). The purpose of this study was to use GGE to better understand the G and GE effects of different environments for selection of Andean dry bean for improved yield stability.

## MATERIALS AND METHODS

A panel of 183 dry bean accessions, representing bush and vine growth habit types were tested across three sites in Tanzania, for two years. The panel comprised of East African landraces, local checks, cultivars and advanced

breeding lines representative of the global genetic diversity for Andean beans grown in Africa and the Americas. These accessions represent a subset of the Andean Diversity Panel (ADP) described by Cichy *et al.* (2015). A description of individual accessions in the ADP can be found at [http://arsftfbean.uprm.edu/bean/?page\\_id=941](http://arsftfbean.uprm.edu/bean/?page_id=941).

**Experimental sites.** The field trials were conducted in three distinct regions of Tanzania, namely Arusha, Morogoro and Mbeya. The dry bean breeding programme based at the Sokoine University of Agriculture in Morogoro, Tanzania is located at 6°49.26122 S/37°39.67322 E, at altitude 526 meters above the sea level (masl). The soils are reddish brown, well-drained fine clay Ultisol. The area receives bimodal rainfall with lower precipitation (November to January) and higher precipitation (March to May) periods. Generally, low fertility soils is a common problem due to lack of fertiliser use. But recently rainfall has become unreliable, thus accentuating the effect of heat and terminal drought stress in the Morogoro region. Bean production is sparse in this area, compared to the Arusha, Kigoma, and Southern Highland (Mbeya) regions, where most of the bean production occurs. Nonetheless, the SUA research farm is a prominent environment used to evaluate and select dry bean breeding lines for production across Tanzania.

The other test environments used by the SUA bean breeding programme include the

Selian Research Station in Arusha region, which is located at 3°22.00022 S/36°40.99982 E, at an elevation of 1387 masl. The soil type is a volcanic ash Nitisol. Like Morogoro, the site receives bimodal rainfall with lower precipitation (November to January) and higher precipitation (March to May) periods.

The Uyole Research Station near Mbeya in the Southern Highlands is located at 8°542 003 S/33°272 003 E. The altitude at the site ranges from 1750 - 1850 masl. Uyole has a well-drained heavy soil which tends toward acidic with low to medium levels of soil nutrients. The area is characterised by unimodal rainfall starting from October/November and ending in April/May. Uyole and Selian Stations have separate national dry bean breeding programs that target cultivar development for their specific regions. The experiments conducted at these sites received rainfall only (precipitation not reported) as no supplemental irrigation was provided.

**Experimental design.** The experimental design was a randomised complete block, with two replications. Single row plots of 8 m length and with 40 plants per row with interrow spacing of 50 cm were used.

**Data collection and statistical analysis.** From each site, soil samples were taken prior to planting, to check for the nutrient status of the fields (Table 1). For GGE stability analysis, only seed yield ( $\text{kg ha}^{-1}$ ) data were used. Seed

TABLE 1. Soil mineral N and available P status for five Tanzania field test environments used to screen a subset of 183 genotypes from the Andean Diversity Panel for yield performance

Site	Year	Environment	Number of genotypes	Soil N = $\text{NO}_3\text{-N} + \text{NH}_4\text{-N}$ ( $\text{kg N ha}^{-1}$ )	Bray P 1 ( $\text{mg kg}^{-1}$ )
Mbeya	2013	Env 1	183	49	7
Morogoro	2013	Env 2	183	53	2
Morogoro	2014	Env 3	183	49	10
Mbeya	2014	Env 4	183	67	5
Arusha	2014	Env 5	183	71	21

yield was estimated from a row per plot harvested.

A combined analysis across the five environments, abbreviated as Env 1, Env 2, Env 3, Env 4 and Env 5 representing Mbeya 2013, Morogoro 2013, Morogoro 2014, Mbeya 2014 and Arusha 2014, respectively, for 183 genotypes was performed using the PROC GLM procedure in SAS version 9.4 (SAS Institute, Cary, NC, 2011). During the test of hypotheses for mixed model analysis of variance, environments and replications were treated as random variables; whereas genotypes were treated as a fixed variable. A single location with different years (growing seasons) were treated as different environments (E), because field conditions and weather varied each year. Partitioning of variances into different sources to test the effects of G, E and GE interactions was performed.

**Genotype main effect.** To perform GGE biplot analysis, least square means from each environment were analysed using the GGE BiplotGUI package from the statistical software R (Yan and Tinker, 2006). A generalised linear model was used to determine the variation in yield among E, G and GE. The GGE biplots show the first two principal components (PC1 and PC2), where axis 1 is for PC1 and axis 2 is for PC2.

**Heritability estimates.** Narrow sense heritability for yield across environments was estimated, using the variance components; which included genotypic (G) variance,

environmental (E) variance and genotype x environment (GE) variance across environments generated using the PROC GLM procedure in SAS. The calculation was done using the mean squares from the ANOVA Table 2 as described by Hallauer *et al.* (2010):

$$h^2 = \frac{\delta^2_G}{\left[ \delta^2_G + \frac{\delta^2_{GXE}}{e} + \frac{\delta^2_E}{re} \right]}$$

Where:

$\delta^2_G$  = genotypic variance;

$\delta^2_{GXE}$  = genotypic by environment variance;

$\delta^2_e$  = error variance; and

r = number of replications

## RESULTS AND DISCUSSION

There was significant variation ( $P < 0.005$ ) for all three sources of variation (E, G and GE) for seed yield (Table 2). Variation among genotypes evaluated in different environments could be explained by the fact that the genotypes were genetically diverse. Therefore, when genetic variation is high, there is greater potential for genetic gains through breeding and selection (Kelly *et al.*, 1998). When variation due to GE is high, then the genotypes are responding differently for yield in the different environments. In this case, closer inspection of the genotype ranking for yield

TABLE 2. Mean squares from analysis of variance for 183 Andean bean genotypes evaluated in five different environments in Tanzania

Traits	Source	DF	Mean square	Pr > F
Tanzania	Environment (Env)	4	38364618	<.0001
	Genotype	182	209813	<.0001
	Env x genotype	728	139993	<.0001

across environments is needed to identify stable yielding genotypes.

Mean seed yields are presented in Table 3. The highest mean seed yield (1382 kg ha<sup>-1</sup>) was recorded in Mbeya 2014 (Env 4); while the minimum yield (499 kg ha<sup>-1</sup>) was recorded in Morogoro 2013 (Env 2). Seed yield for Mbeya 2013 (Env1), Morogoro 2014 (Env 3) and Arusha 2014 (Env 5) did not differ significantly. The five environments could be ranked in terms of their grain yield potential as follows: Mbeya 2014 > Arusha 2014 > Morogoro 2014 > Mbeya 2013 > Morogoro 2013. The percentage sum of squares for E, G, and GE was 46, 20 and 34%, respectively (Table 4). Van Eeuwijk *et al.* (1995); Beyene

*et al.* (2011) and Sserumaga *et al.* (2015), in studies of maize, reported higher genotypic variation in grain yield than the effect of GE interaction for yield. Similar results were reported by Hoyos-Villegas *et al.* (2016), who examined the performance and adaptability of Mesoamerican common bean genotypes under irrigation *versus* rain-fed conditions in Michigan. However, in the present study, E was higher than GE and G, suggesting that the environmental conditions differed. According to soil test results (Table 1), there was wide variability in soil fertility levels between test sites. This variability in soil fertility may have contributed to the variation in yield observed across the five environments, as well.

TABLE 3. Seed yield for 183 Andean bean genotypes evaluated in five environments in Tanzania

Environments	Location	Mean seed yield (kg ha <sup>-1</sup> )†
Environment 1	Mbeya 2013	778.5 <sup>b</sup>
Environment 2	Morogoro 2013	499.4 <sup>c</sup>
Environment 3	Morogoro 2014	783.0 <sup>b</sup>
Environment 4	Mbeya 2014	1382.0 <sup>a</sup>
Environment 5	Arusha 2014	789.4 <sup>b</sup>

† Means followed by the same letter within the same column are not significantly different (P = 0.05) Tukey comparisons of mean

TABLE 4. Effect of E, G and GE from total SS<sub>(G)</sub>, SS<sub>(E)</sub> and SS<sub>(GE)</sub> and heritability estimates for 183 common bean genotypes evaluated in five different environments in Tanzania

Source of variation	Tanzania (%)
Environment	46
Genotypic	20
Gen x Env	34
CV	28
Heritability	26

Abiotic stress tolerance measured by seed yield is a heritable trait in common bean and other crops (Clark and Duncan, 1991). In this study, heritability was low 26% indicating that seed yield was highly influenced by the environment (Table 4).

**Biplot analysis.** The significant GE interactions for yield observed in the present study indicated that these data were suitable for GGE Biplot analysis. To visualise the yields from genotypes and interactions between genotypes and environment, biplots were generated based on a table, “Tester-centered

(G+GE)” without scaling in row metric preserving (JK biplot) in R. Several biplots were generated to examine the relationships among G and GE, as described in Yan and Tinker (2006).

From the GGE biplot analysis, the first principal component captured 41.21% of the total sum of squares for G and GE; and the second principal component captured 26.35%.

Figure 1 was used to visualise relationships between test environments. The angle formed when two environmental axes meet explains if there is a correlation between the two environments. An acute angle  $> 90^\circ$  indicates

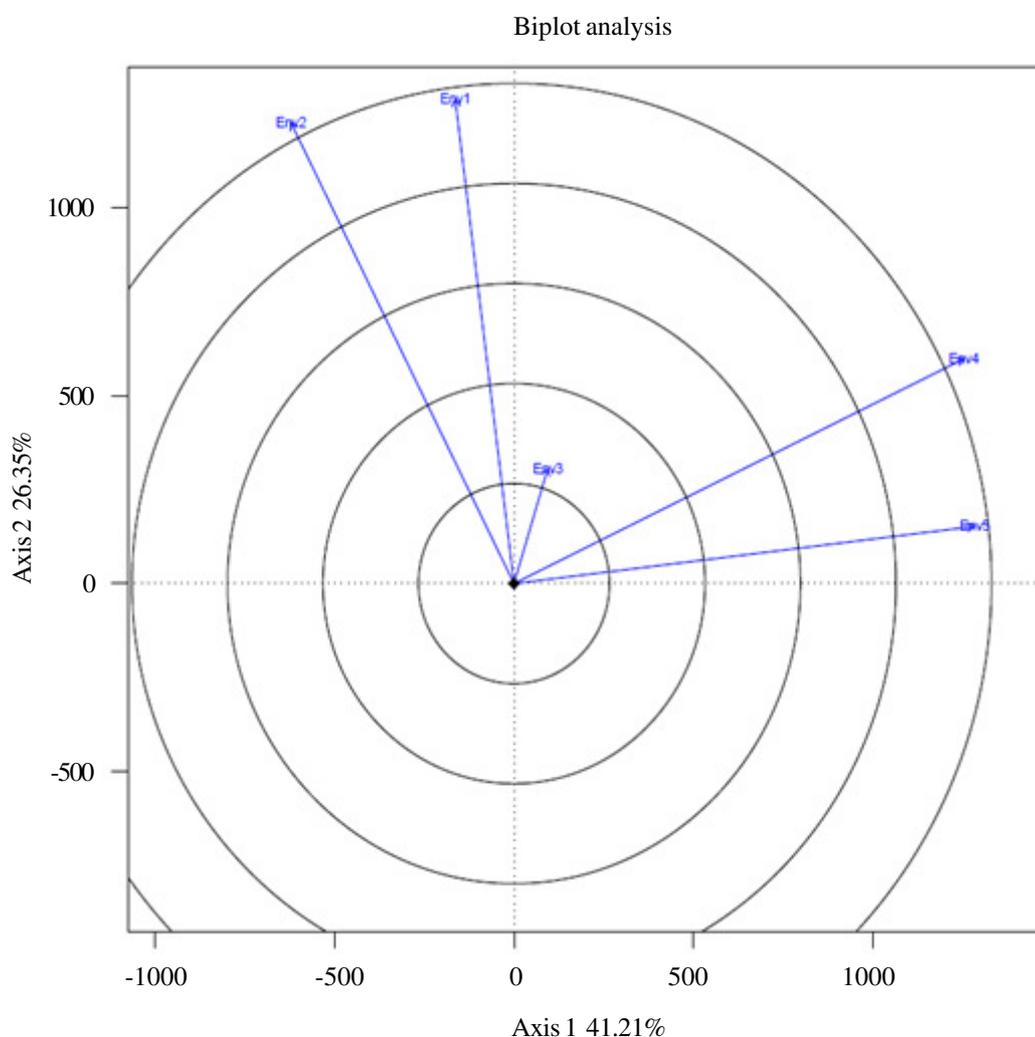


Figure 1. Relation among five environments in a genotype x environment (GE) interaction study for yield performance of Andean dry bean in Tanzania.

positive correlation, an obtuse angle  $< 90^\circ$  indicates a negative correlation and a right angle ( $= 90^\circ$ ) indicates no correlation (Yan and Tinker, 2006). Our results (Fig. 1) indicate a negative correlation between Morogoro 2013 and Arusha 2014, no correlation between Mbeya 2013 and Arusha 2014, and all other environments are positively correlated. A shorter distance between environments indicates similarity for discriminating genotypes, this is determined by the angle between the environment and the length of

their vectors. Thus, Mbeya 2013, Morogoro 2013 and Morogoro 2014 are similar and fall in one group; while Mbeya 2014 and Arusha 2014 are similar and fall in another group. The positive correlations among environments indicate that fewer test environments can be used to discriminate yield performance of the genotypes.

The length of environment vectors gives information about the discriminating ability of the environments (Yan and Tinker, 2006). From Figure 2, Morogoro 2013 and Mbeya 2014 are

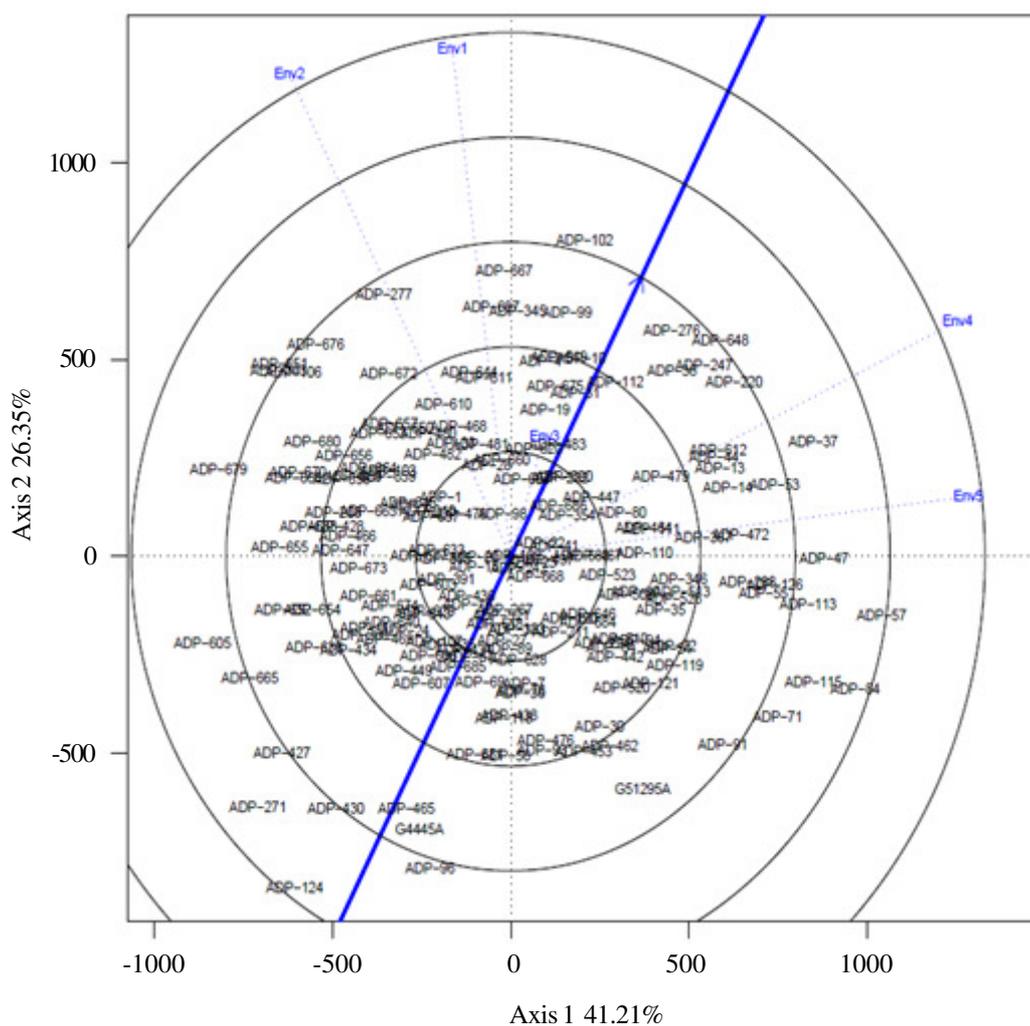


Figure 2. Discriminativeness vs representativeness in a genotype x environment (GE) interaction study for yield performance of Andean dry bean in Tanzania. A description of ADP accessions is available at [http://arsfiftbean.uprm.edu/bean/?page\\_id=941](http://arsfiftbean.uprm.edu/bean/?page_id=941).

the most discriminating; while Morogoro 2014 is the least discriminating environment. Less discriminating environments give less information on genotypic performance, and thus should be eliminated as a test environment. The ideal environment should be located at or close to the environment axis with an arrow. An environment with a smaller angle to environmental axis is more representative. In this case, Morogoro 2014 was the most representative, but was the least discriminating, so has little value. Morogoro 2013 and Arusha 2014 are the least representative. The most

discriminating, but least representative environments (Morogoro 2013 and Arusha 2014) can be used to select genotypes adapted to these specific environments. The most discriminating and most representative environments (Mbeya 2013 and Mbeya 2014) can be used to select genotypes with wider adaptation.

The ideal environment for selecting genotypes for general adaptation should be located at the centre of concentric circles (Fig. 3). This is the most informative and most discriminating environment. None of the five

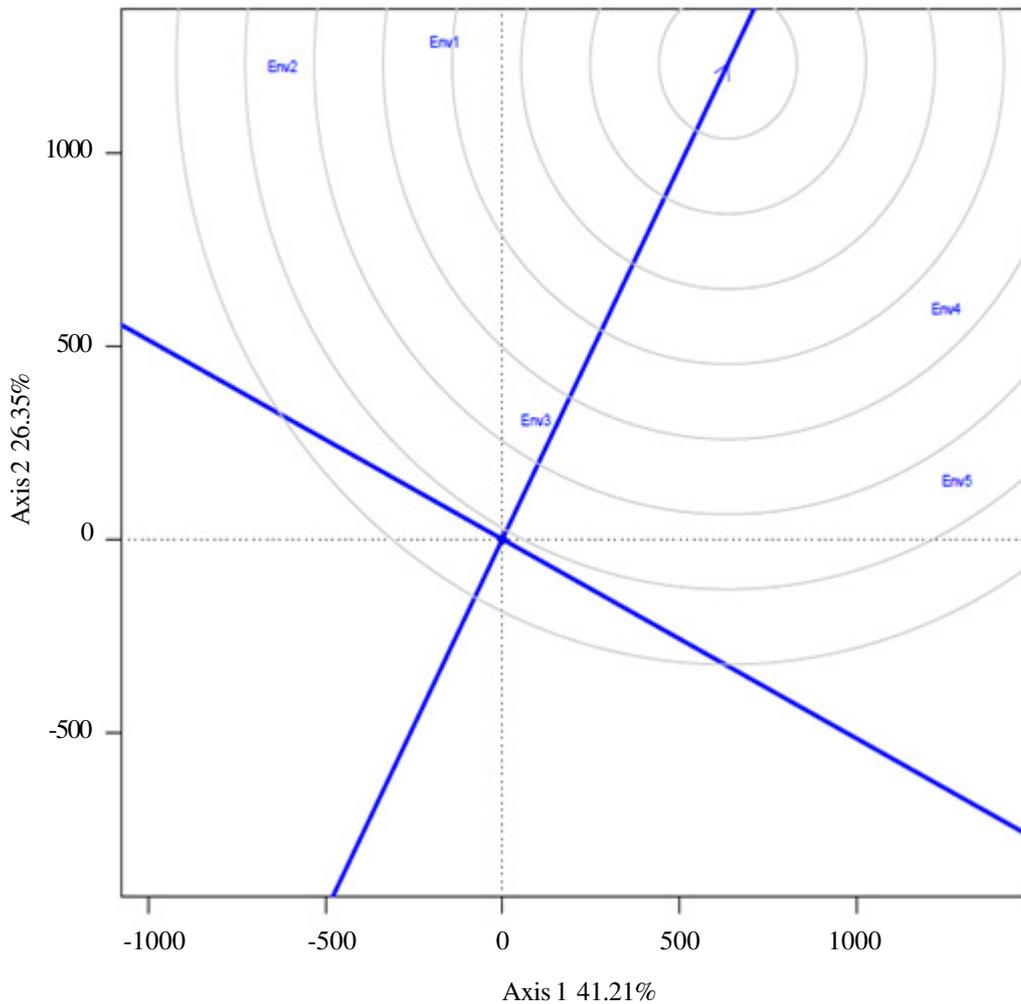


Figure 3. Ranking of the environments relative to the ideal environment for Andean dry bean yield performance in Tanzania.

environments tested, fell within or even near the circle. Therefore, all environments are suitable for selecting genotypes for specific adaptation but not for general adaptation.

Results to identify the best genotype in each environment are presented in Figure 4 (which-won-where). All genotypes located at the vertices in each sector of a polygon are the best for the environment in the sector where the vertices are located (Yan *et al.*, 2000; Yan

and Tinker, 2006; Frutos *et al.*, 2013; Makumbi *et al.*, 2015). ADP-102 is the best in Mbeya 2013 and Morogoro 2014, ADP-277 is the best in Morogoro 2013, ADP-648 is the best in Mbeya 2014; whereas ADP-37, ADP-57 and ADP-84 are the best in Arusha 2014. The worst genotypes (low yield) are located at the vertices of the sectors in the polygon with no environments; thus, ADP-676, ADP-651, ADP-679, ADP-605, ADP-271, ADP-124.

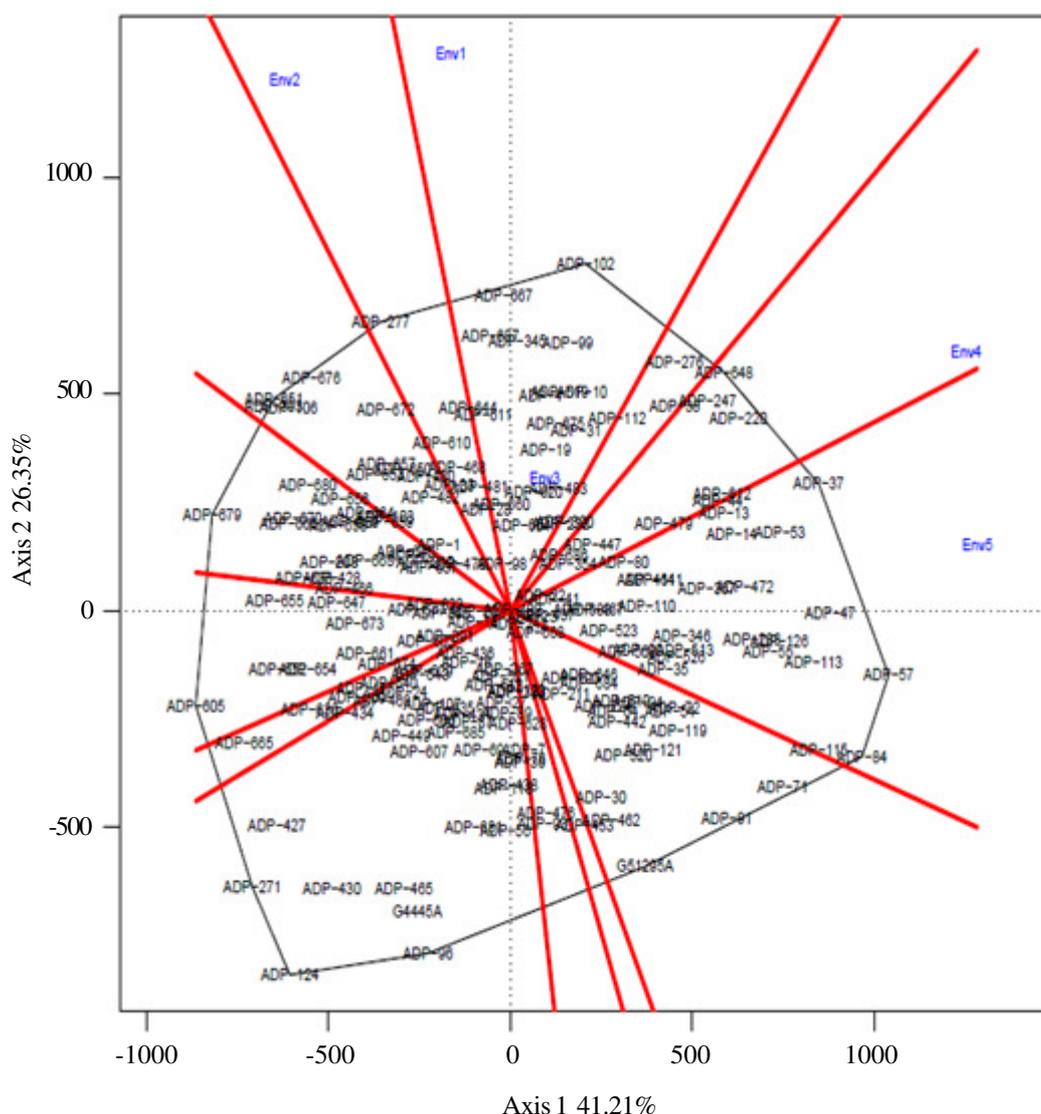


Figure 4. Identification of the best Andean dry bean genotypes for yield in five different environments (which-won-where) in Tanzania.

ADP-96, G51295A, ADP-91 and ADP-71 are the worst genotypes (Fig. 4).

The average environmental axis (an axis with an arrow) is pointing to higher mean yield across environments (Fig. 5). Therefore, ADP-102 had the highest mean seed yield, followed by ADP-648, ADP-276 then ADP-220. However, ADP-124 had the lowest mean seed yield.

The line perpendicular to the average environmental axis points to greater variability in either direction (poorer stability). Thus, ADP-57 and ADP-84 were highly unstable; while ADP-112 was highly stable. A genotype can be very unstable if it has higher above average yield in one environment and below average yield in another environment. Moreover, a genotype can be highly stable,

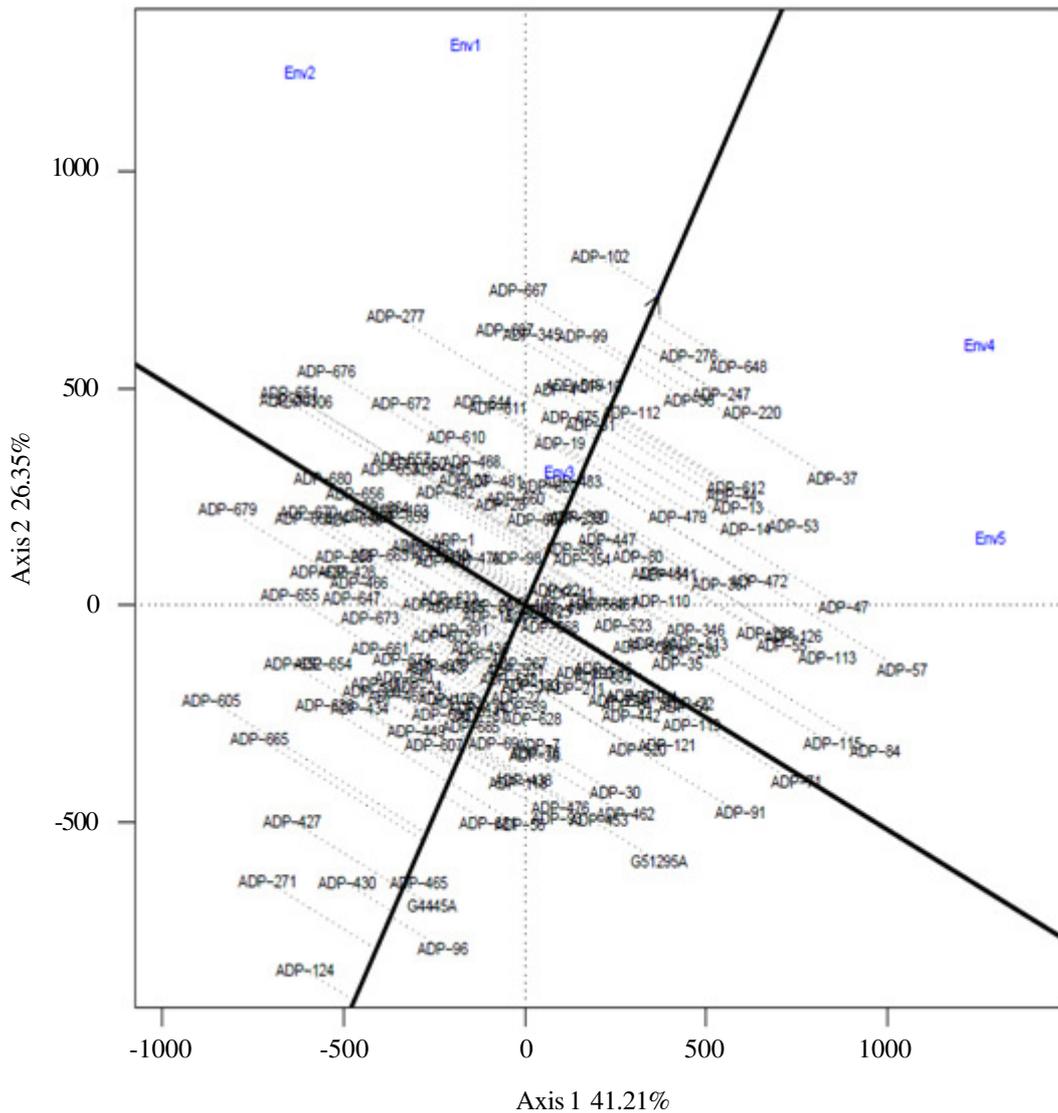


Figure 5. Mean seed yield and Andean dry bean genotype stability across five environments in Tanzania.

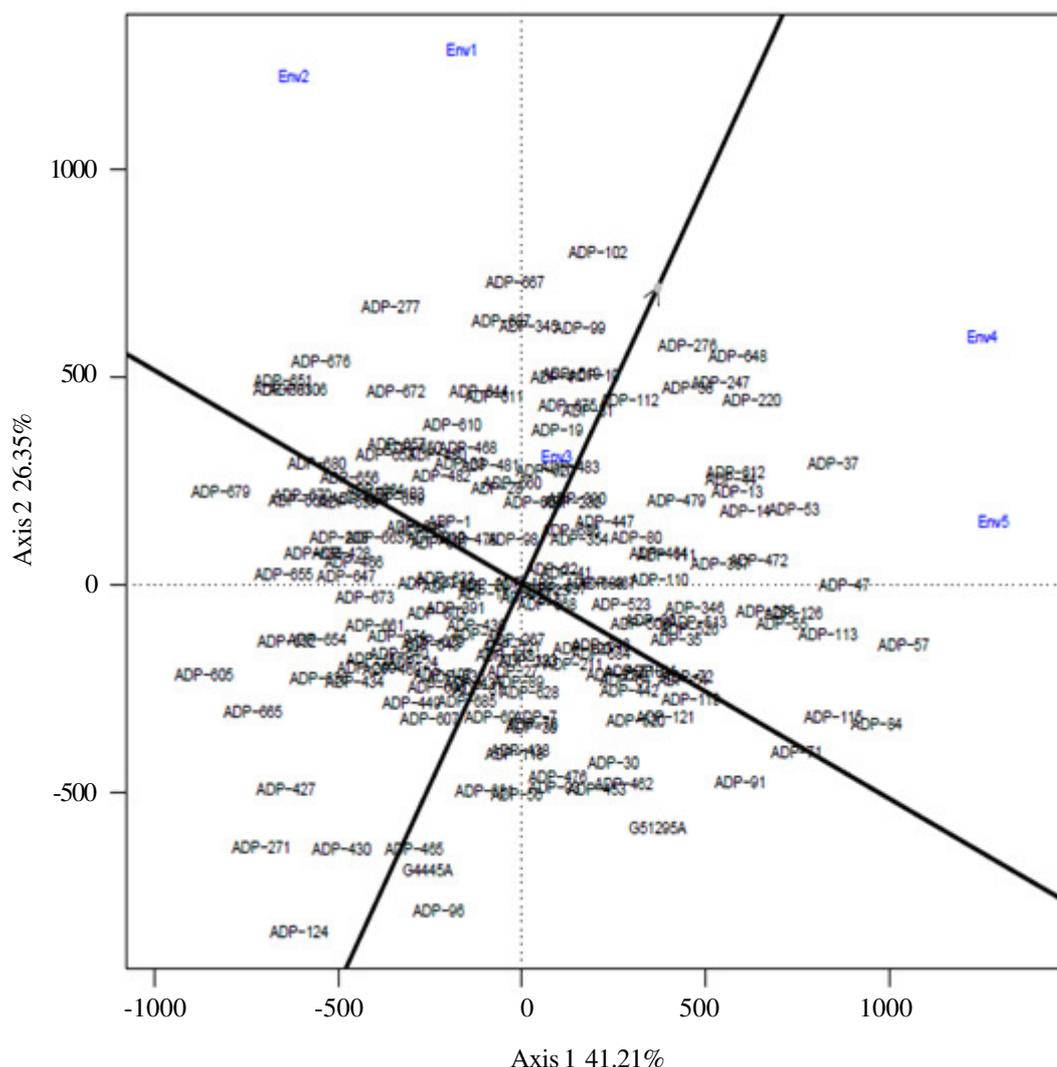


Figure 6. Ranking of the Andean dry bean genotypes relative to ideal genotype for yield performance in five test environments in Tanzania.

with low mean yield, provided that the yield is consistently low across environments. Therefore, stable genotypes are preferred only if they have higher mean yield across environments.

An ideal genotype has high mean yield performance, and higher stability across environments. The arrow inside the blue circle is where an ideal genotype should be. All genotypes located close to where an ideal genotype is, are more desirable. In this case,

ADP-102 was most desirable followed by ADP-276, ADP-667, ADP-99, ADP-684 and the poorest genotype was ADP-124 because it is at the bottom of environmental axis (Fig. 6).

### CONCLUSION

The study revealed significant G, E and GE variance for seed yield in Andean dry bean genotypes tested in different environments in

Tanzania. The Andean genotypes in this study exhibit a broad range of variation in yield under contrasting environments, which differed for water stress, soil fertility and general climatic conditions.

A few stable high yielding genotypes were found, including ADP-102 (Jessica)- purple mottle, large seed, bush plant habit, released by Selian Agriculture Research Institution (ARI) in Tanzania; ADP-220 (G5625)- red, large seed, vine plant habit from Central America; ADP-276 (G13654) - brown, medium seed, vine, from Mexico; and ADP-648 ('Red Kloud') large red kidney, with bush plant habit, from North America. These represent suitable materials for breeding for wide adaptation. The lines which performed well under low fertility and or water stress environments may possess abiotic stress tolerance and may be useful in breeding for improved yield stability for such conditions. An ideal environment was not identified for selecting genotypes with wide adaptation, indicating the continued need to maintain separate breeding and selection programs in the Northern (Arusha) and Southern Highlands (Mbeya) production regions of Tanzania.

Biplot analysis gave a good indication of genotypic performance across environments. An ideal genotype should have higher mean yield than other genotype and an ideal environment should have higher mean yield compared to other environments. GGE analysis can be helpful for making better decisions about which genotypes should be recommended for release in specific environments or across environments.

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