

*Full Length Research Paper*

# Evaluating natural infection of fungal, bacterial and viral pathogens to dry bean genotypes under field conditions

Papias H. Binagwa<sup>1,2\*</sup>, Guohao He<sup>2</sup>, Marceline Egnin<sup>2</sup>, Gregory C. Bernard<sup>2</sup>,  
Desmond Mortley<sup>2</sup> and Conrad K. Bonsi<sup>2</sup>

<sup>1</sup>Integrative Bioscience (IBS), Ph.D. Program, Tuskegee University, AL 36088, USA.

<sup>2</sup>Department of Agricultural and Environmental Sciences, Tuskegee University, AL 36088, USA.

Received 11 December, 2019; Accepted 3 February, 2020

**Fungal, bacterial and viral diseases are economic foliar diseases that cause yield losses, between 40 and 100%, in commonly grown dry bean cultivars in the world. Development of disease resistance genotypes is a complex interaction between genetic and environmental factors. This study focused on determining the natural infection of disease-causing pathogens of angular leaf spot, powdery mildew, bacterial blight and bean common mosaic virus in different agro-ecologies in relation to grain yield. Diversity of 211 bean genotypes were tested at two different disease hot spots areas under incomplete block design, with two replications for two cropping seasons in Tanzania. Diseases severity was significantly different ( $p < 0.001$ ) for genotypes and their interactions with the environment and season. Higher disease severity was observed at Lyamungo site than Selian site. Effects of genotypes by environment were observed with maximum yield of 2170 kg/ha to low yield of 398 kg/ha with the grand mean of 1151.54 kg/ha. High annual rainfall and relative humidity contributed to disease development among the tested environment. Five genotypes (FEB 189, A774, NUA 16, KG 71-4 and DOR 766) expressed trait of resistance to above diseases and are advised to be incorporated in breeding programs for enhancing dry bean productivity.**

**Key words:** Diseases, losses, productivity, G\*E interaction.

## INTRODUCTION

Dry bean (*Phaseolus vulgaris* L.) is the most widely grown and significantly consumed grain legume in the world (Broughton et al., 2003; FAOSTAT, 2018). There are two origin centers, Mesoamerican origin in southern Mexico and Guatemala, as well as Andean origin in Peru and Columbia (Landon, 2008; Kwak et al., 2009) where this crop originated from; before spreading across the world (Kwak and Gepts, 2009). Dry bean is the primary

food crop with the highest level of variation in adaptation, maturity, growth habit (habitat) and seed characteristics (size, shape and color) (Peters, 1993). Additionally, some geographic regions are favored with producing large size seeds, medium (25 to 40 g per 100 seeds) or large (>40 g per 100 seeds).

Global production is hindered by biotic and abiotic factors resulting in commercial varieties yielding lower

\*Corresponding author. E-mail: pbinagwa8224@tuskegee.edu. Tel: +255764951596.

than their potentiality (De Leque and Creamer, 2014). Based on the economic importance, fungal diseases cause higher losses followed by bacterial and viral diseases (Mahuku and Riascos, 2004). For instance, the percentage of damage caused by fungal diseases is 80% by Angular Leaf Spot (ALS), *Phaeoisariopsis griseola* (Sacc.); 100% by Anthracnose, *Colletotrichum lindemuthianum*; and more than 50% by Powdery mildew (PM), *Erysipelas polygoni*. Bean common blight (*Xanthomonas axonopodis* pv. *phaseoli*), which is a bacterial disease, causes 45% losses (Nkalubo et al., 2007). Viral diseases, such as *Bean common mosaic virus* (BCMV) and *Bean common mosaic necrosis virus* (BCMN), can cause 100% yield losses (Hillocks et al., 2006; Buruchara et al., 2010; Singh and Schwartz, 2010; Mwaipopo et al., 2016) with their major transmission and survival structures which includes seeds, wind and plant debris.

These diseases (ALS, CBB, PM and BCMV) severely affect farmers' field in Tanzania, which ranks number one in Africa and 6<sup>th</sup> in the world (FAOSTAT, 2018). The development of foliar disease resistance bean genotypes through understanding of the environmental factors and gene alleles interactions may help gain insight into disease etiology and sub-classification; also, management options would offer better strategies for bean breeding program (Wang et al., 2005). The marked recent improvement in bean breeding program is in the initial stage for biotechnology approaches (Harwell et al., 2011). It has been shown that field crop phenotyping under natural infection assists desirable trait assessments for genetic variability aimed at selecting genotypes with better traits for enhanced improvement (Sankaran et al., 2015). The objectives of this research study are to (a) evaluate the response of dry bean genotypes to ALS, CBB, PM and BCMV diseases at different environments and seasons; (b) screen best genotypes with high yield and resistance traits for breeding purposes to all diseases and their specificity; (c) compare disease occurrence in relation to cropping seasons for actual rainfall and temperature.

## MATERIALS AND METHODS

### Plant materials

A total of 211 genotypes were collected from different sources: Ethiopia (12), International Center for Tropical Agriculture (CIAT)-Kawanda (184), Kenya (10), Tanzania (3) and Rwanda (2) (Supplement 1) with various market classes evaluated under natural field conditions for 2016/17 and 2017/18 cropping season. From these materials, resistant checks of MEX 54 and G5686 (Mahuku et al., 2009) was used for ALS, Vax 1 (Singh et al., 2001) was used for CBB, MAZ 47 was used for BCMV; unfortunately, checks for PM disease was not included.

### Experimental area

The experiment was conducted at low to high altitudes, 1407 m

above sea level of S03°21.690' and E36°37.879' at Selian Agricultural Research Institute (SARI) and Tanzania Coffee Research Institute commonly called Lyamungo, with 992 m.a.s.l of 03°19.905' and E037°14.067', respectively. The characteristics of this soil area are: Eutrophic Brown Soils on volcanic and Alluvial sediments - Medium texture (loamy soils) (Brady and Weil, 2002; Landon, 1991). The soil contains organic carbon (0.53%), organic matter (0.92%), total nitrogen (0.079%), exchangeable potassium (0.17 cmol (+)/kg) and medium available phosphorous (8.0 mg/kg); this means that the soil fertility status is medium fertility which is moderately suitable for bean cultivation (Kiriba et al., 2020). These selected sites are close to the weather station, from whence respective weather data were collected.

### Disease evaluation and grain yield determination

Each disease (ALS, PM, CBB and BCMV) was rated using 1 to 9 scale as described by CIAT (1987) and CIAT-Kawanda (2013); where 1 to 3 refer to resistant, 4 to 6 intermediate, and 7 to 9 susceptible. Grain yield was measured from each plot using digital scale with 11lb (model No. SKS - 006, China). Final grain yield was extrapolated into kg/ha, using the following formula:

$$\text{Grain yield (kg/ha)} = (dy * 10) / dx$$

Where  $dy$  = plot weight and  $dx$  = plot area.

### Experimental design

In both locations, in all growing season, trials were laid out under incomplete block design with two replications. The experimental plot size was 4 rows, 3.2 m long and 50 cm apart and 20 cm within a row. The harvested net plot size was 3.2 m<sup>2</sup> of the centered two rows of each plot. Other practices were carried out as recommended by National *Phaseolus* Bean Research Program in Tanzania (Binagwa, 2017).

### Statistical analysis

The collected data were subjected to GenStat 16<sup>th</sup> Edition with the following linear model:

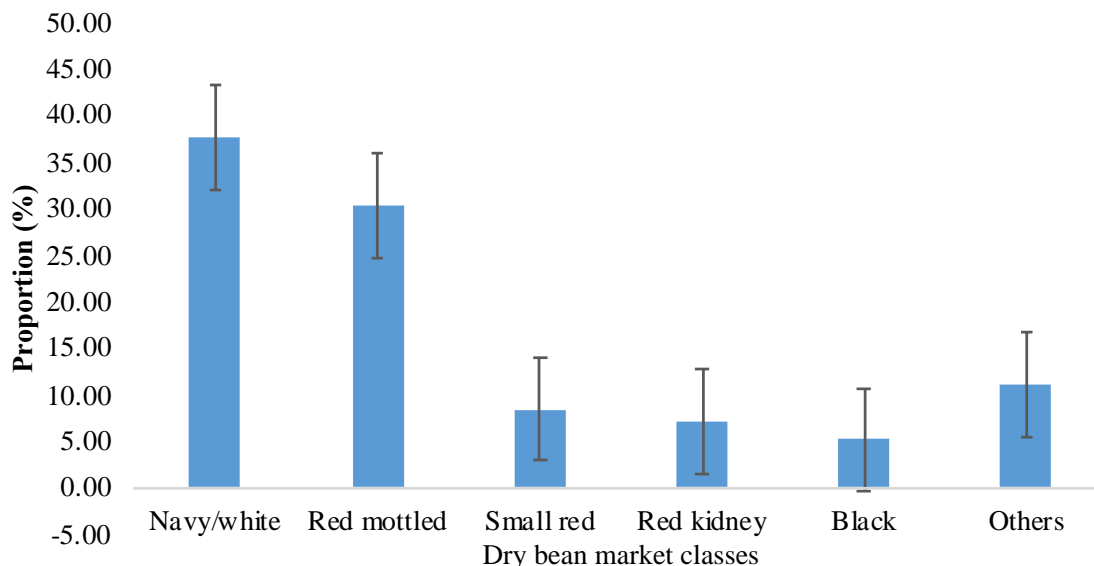
$$Y_{ijk} = \mu + G_i + y_j + G_i * y_j + 2k + G_i * y_j * 2k + e_{ijk}$$

Where  $Y_{ijk}$  = Response variable (Yield) for variety  $i$ , environment  $j$  and season  $k$ ;  $\mu$  = Overall mean for all the observed response;  $G_i$  = Fixed effect of variety;  $y_j$  = random environmental effect of the observed response;  $G_i * y_j$  = Interaction effects between variety and environment;  $2k$  = Random effect of replication within a season;  $G_i * y_j * 2k$  = Interaction effect of variety, environment and season;  $e_{ijk}$  = Random term error which is assumed to be normally distributed with 0 mean and variance  $\delta^2$  which were summarized in a given results. Data were tested using Analysis of Variance (ANOVA) for single and multiple treatment interactions. The protected Least Significant Differences (LSD) of ( $p=0.05$ ) were used to test for treatment comparison (Clewer and Scarisbrick, 2001; Yu 2008).

## RESULTS

### Classification of dry bean genotypes used

Through seed morphological description process, the



**Figure 1.** Proportional of dry bean seed market classes.

genotypes utilized were classified as 63.50% from Mesoamerican and 36.50% from Andean gene pools of origin. Based on market class, navy/white is dominant by 32.70% followed by red mottled, 30.33%. The remaining market classes were small red (8.53%), red kidney (7.10%), black, carioca, kablanquet and cream (Figure 1). Seed size was large, medium and small, proportional to 36.02, 9.95 and 54.03%, respectively.

### Effects of fungal disease-causing pathogens

The ANOVA analysis showed that the effects of genotype, environment, season and the interactions of genotype and environment (G\*E) were significantly different at  $p < 0.001$ ; while that of genotype and seasons were significant at  $p = 0.002$  for ALS infections. The effects of ALS disease caused by *P. griseola* were high at Lyamungo site with the severity of up to 5.00 scale; while 4.20 at SARI. The G\*E effects were observed between the range of 2.00 to 3.50 scale (Figure 2A). In relation to season, infection was high in the 2017/18 season ALS, with severity score  $> 5.00$ ; while that of 2016/17 reached  $\sim 2.80$  and most of the genotypes expressed none pathogen infection (Figure 2B). Overall, 38 genotypes had higher severity scores above the grand mean (Supplement 2). For PM infection caused by *E. polygoni*, disease severity was high at Lyamungo site with score  $> 6.00$ ; while  $\sim 4.50$  at SARI (Figure 2C and D). The effects of genotypes, environment, season, G\*E, G\*S and G\*E\*S were also significant ( $p < 0.001$ ) for PM infection. The PM infection was high at Lyamungo, followed by with SARI site during the 2017/2018 growing season, with severity scores between 2.00 to 7.00 and 1.00 to 5.00

(Figure 2E) (Supplement 2).

### Effect of bean bacterial blight pathogen

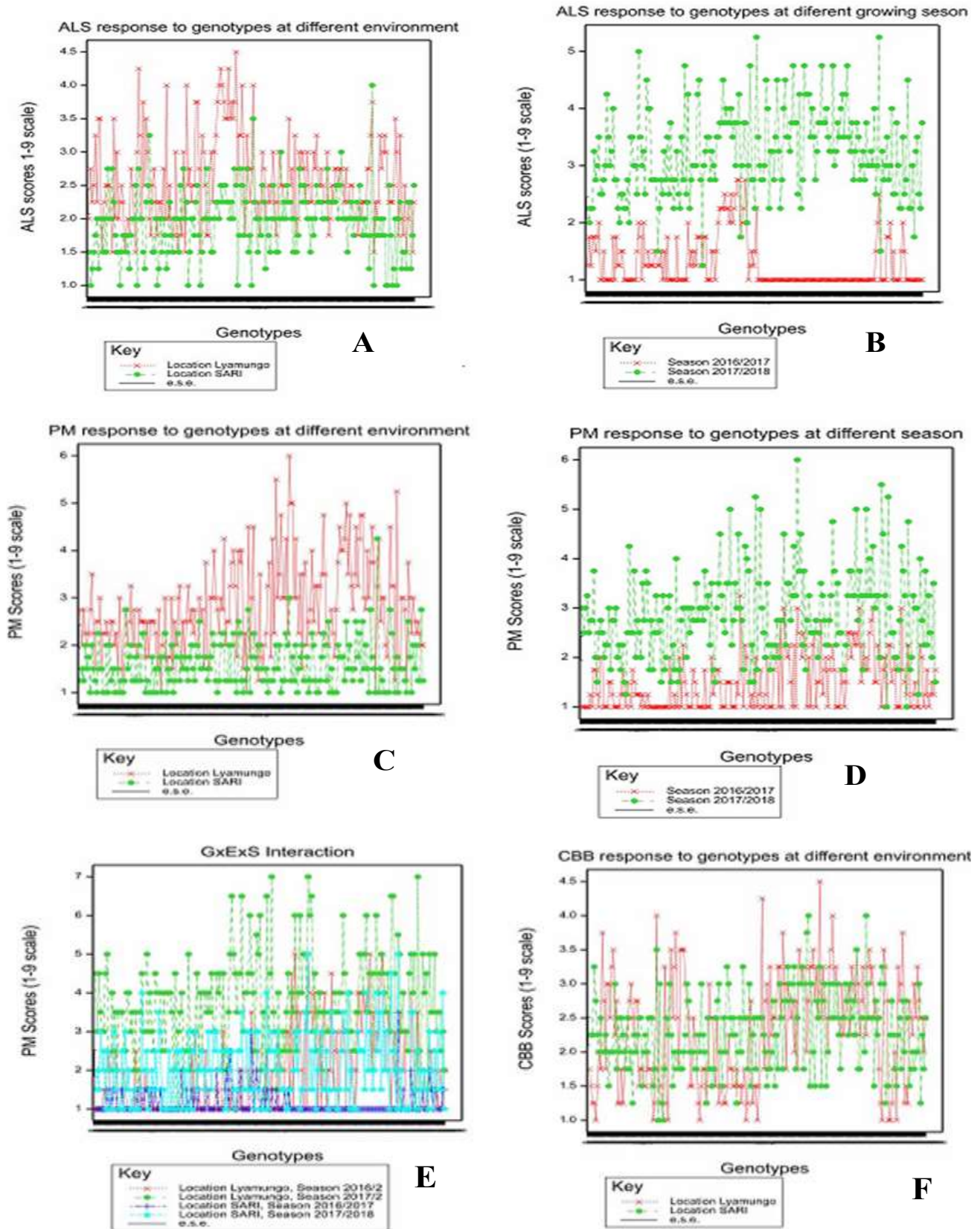
Two bean growing season results showed significant difference ( $p < 0.001$ ) between genotype, season, G\*E, G\*S and G\*E\*S for CBB reaction, and about 82 genotypes showed their disease severity scores were above the grand mean of  $> 3.00$  scores. The effect was high at Lyamungo,  $\sim 4.70$ ; while at SARI, severity scores was high up to 4.20 (Figure 2F). The growing season of 2017/2018 had more infections of CBB than that of 2016/2017 growing season, with more scatter points above 3.50 disease scores (Figure 3A and Supplement 2).

### Infection of bean common mosaic virus

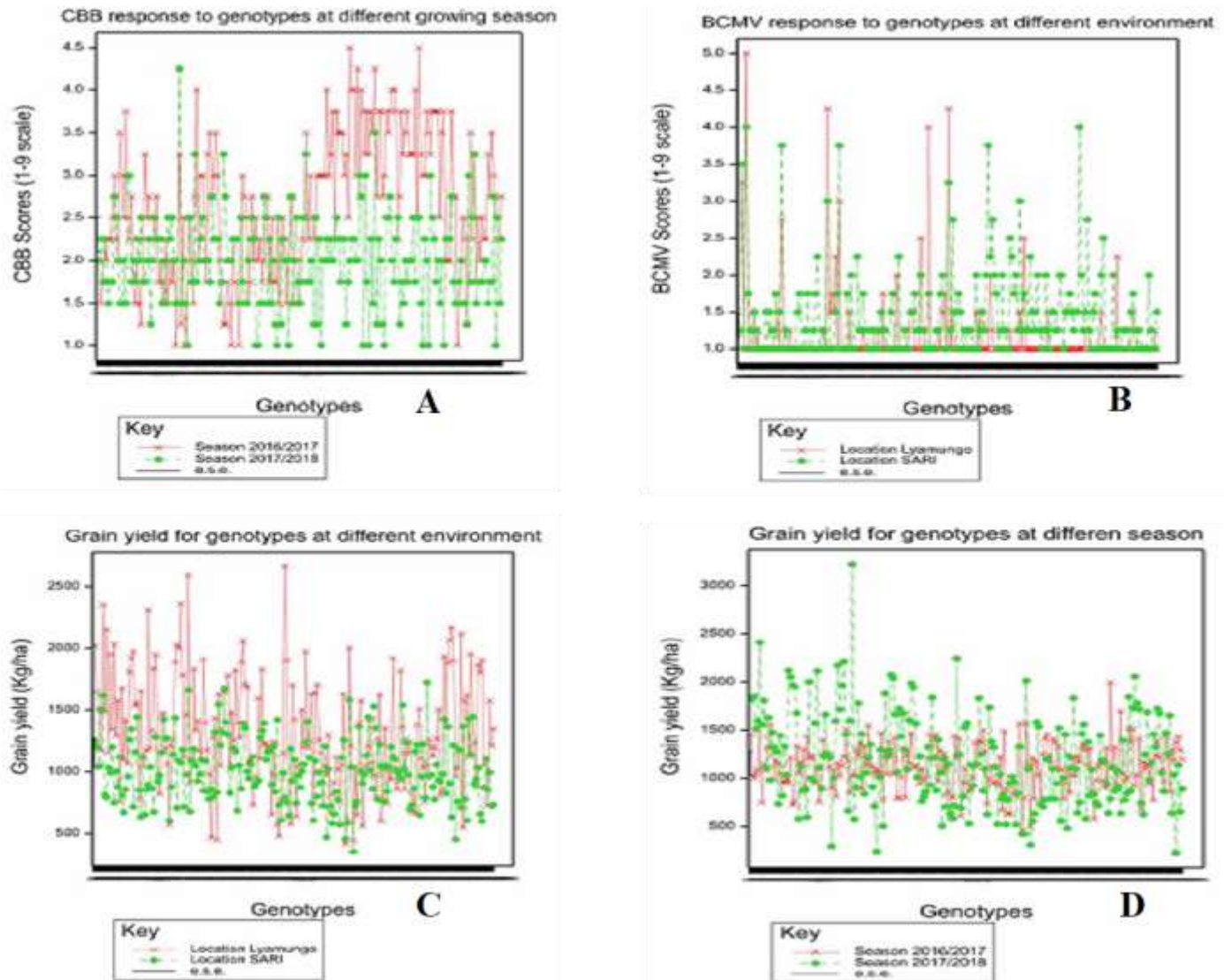
There was significant difference ( $p < 0.001$ ) between G\*E and G\*E\*S ( $p = 0.03$ ) for dry bean common mosaic virus. A few genotypes, namely CC 906, 222/1, Flor De Mayo, DOR 755, MLB48-89A and A 686, had severity scores within the ranges of 3.25 to 4.50. Narrow variation observed showed resistance scores across the tested sites due to most genotypes (Figure 3B and Supplement 2).

### Grain yield production across the environment and season

Genotype, environment, G\*E and G\*E\*S showed



**Figure 2.** Response of disease to dry bean genotypes (A) genotype and environment interaction for ALS; (B) Genotype and season interaction for ALS; (C) Genotype and environment interaction for PM; (D) Genotype and season interaction for PM; (E) Genotype, environment and season interaction for PM; (F) Genotype and environment interaction for CBB.

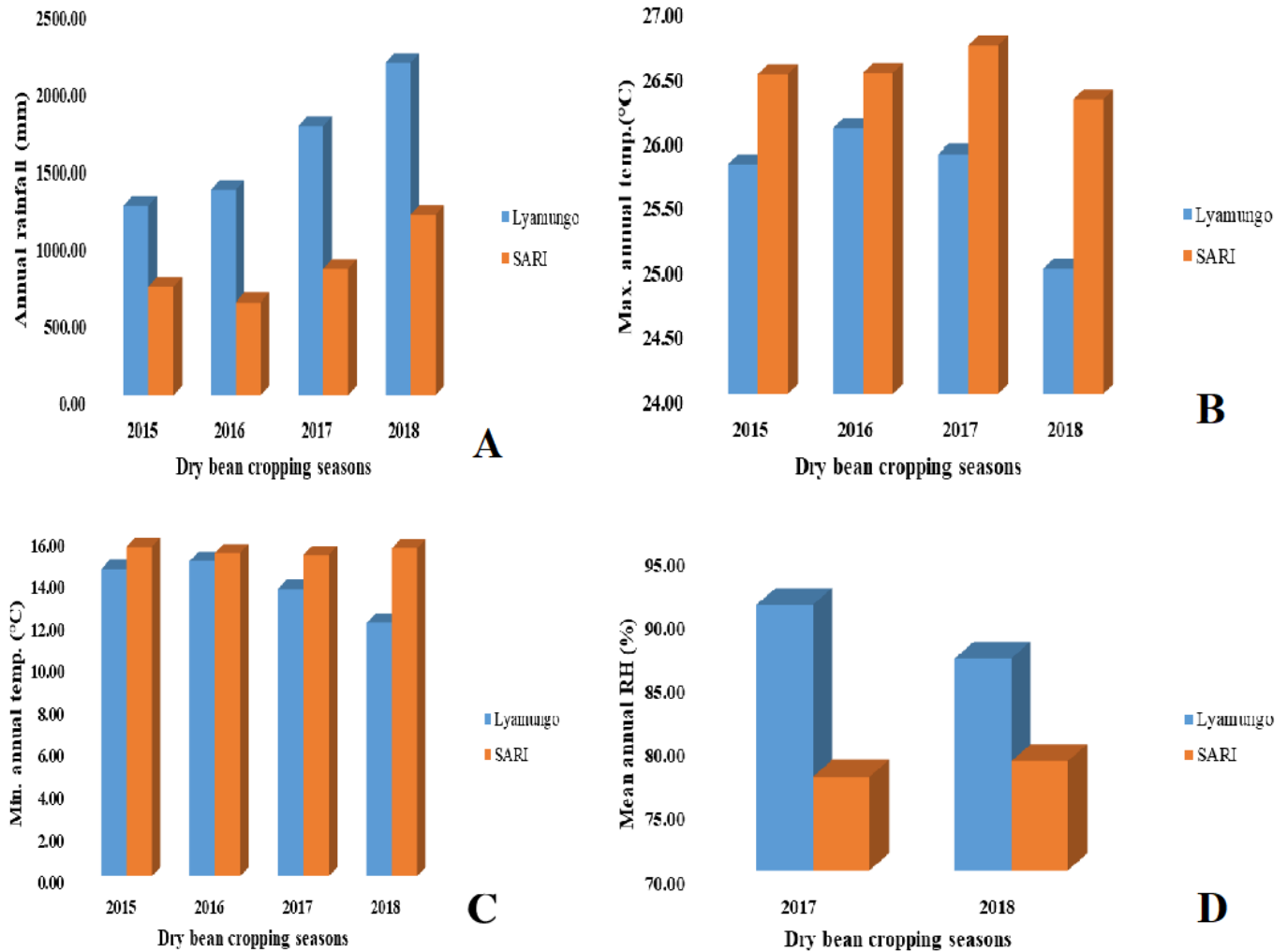


**Figure 3.** Response of disease to dry bean genotypes and grain yield; (A) Genotype and season interaction for CBB; (B) Genotype and environment interaction for BCMV; (C) Genotype and environment interaction for grain yield; (D) Genotype and season interaction for grain yield.

significant difference ( $p=0.004$ ) in season ( $p=0.004$ ). Grain yield of all genotypes at the Lyamungo site was higher than that of SARI site across the testing seasons. For instance, maximum yield at Lyamungo was  $>2500$  kg/ha; while the highest yield at SARI was  $\sim 1800$  kg/ha and the lowest yield  $<500$  kg/ha, with the grand mean of 1151.54 kg/ha for both locations (Figure 3C). The 2017/2018 growing season resulted in higher yield of  $>3000$  kg/ha; while that of 2016/2017 gave rise to approximately 2100 kg/ha (Figure 3D). Generally, Lyamungo site during 2017/2018 season performed better in this study and the best top five bean genotypes, FEB 189, A774, NUA 16, KG 71-4 and DOR 766, produced yield of 2127, 1982, 1793, 1725 and 1715 kg/ha, respectively (Figure 2C and D and Supplement 2).

### Climate and diseases occurrence during this study period

Annual rainfall (mm), maximum temperature ( $^{\circ}\text{C}$ ) and mean temperature for consecutive four cropping seasons (2015-2018) were collected as well as annual relative humidity (%) for two cropping seasons (2017-2018). High annual rainfall was observed at Lyamungo site than that of SARI, across the cropping seasons. For instance, in 2018, annual rainfall was 2156.40 mm at Lyamungo but 1169.20 at SARI (Figure 4A). Mean and maximum temperature was higher at SARI compared to Lyamungo (Figure 4B and C); while relative humidity was reverse to temperature (Figure 4D). The negative correlation in both sites indicated that the higher the rainfall, the lower the



**Figure 4.** Annual climate statistics for experimental sites; (A) Annual rainfall, (B) Maximum average annual temperature; (C) Minimum average annual temperature; (D) Mean annual relative humidity.

temperature and the higher the relative humidity in these cropping seasons. Through comparative analysis, the high rainfall and relative humidity resulted in high severity of fungal, bacterial and viral diseases at Lyamungo; while at SARI, though disease severity was lower, productivity was also low, which may be attributed to terminal drought and abortion of flowers due to high temperature.

#### Dry bean market class variations with respect to diseases effects and grain yield

Small red genotypes dominated the higher grain yield (>1500 kg/ha) compared to all accessions; while navy and red mottled genotypes produced yield range between 1000 and 1490 kg/ha. Red mottled genotypes with large to medium size were more susceptible to ALS, CBB and PM, with lower yield compared to other market classes. Small seeded genotypes, especially black, red and khaki

striped such as A 686, MLB 48-89A, DOR 755, FLOR DE MAYO, 222/1 and CC 906, were more susceptible to BCMV disease and their yield was lower (Supplement 2). Although some genotypes were not infected by pathogens, their yields were lower (<1000 kg/ha), which may be due to poor germination and adaptability (Table 1). From the combined analysis that expressed the effects of GxE across the two bean growing seasons, ten resistant genotypes were identified for each diseases and were listed in Table 1. From this identification, small white and small red expressed trait of resistance under this field conditions than other market classes.

#### DISCUSSION

The responses of genotypes to diseases through genotype and environmental interactions led to yield variations. Infection caused by ALS disease-causing

**Table 1.** Identified top 10 genotypes with trait of resistance per specific disease, market type, severity and grain yield.

Economic disease	Genotypes	Market type	Score (1-9)	Yield (kg/ha)
Angular Leaf Spot (ALS)	KG 30-29	White	1.00	1257.00
	Ranjonomby	White	1.00	1679.00
	DOR 755	Small red	1.00	1545.00
	ZABR 16575-51F22	White	1.00	962.00
	CN Bunsu (64)	White	2.00	897.00
	CC814	Carioca	2.00	1345.00
	CN Bunsu (60)	White	2.00	815.00
	CN Bunsu (62)	White	2.00	1145.00
	G 31	White	2.00	1169.00
	G 78	White	2.00	1239.00
Powdery mildew (PM)	PAN 72	White	1.00	1219.00
	CN Bunsu (64)	White	1.00	897.00
	SM 133	White	1.00	1412.00
	ZABR 16573-78F22	White	1.00	1070.00
	F1Population	White	1.00	916.00
	G 30	White	1.00	1190.00
	G 60	White	1.00	1378.00
	MEX 54	Cream	1.00	1725.00
	MICHETTE	White	1.00	645.00
	Awash-1	White	1.00	1270.00
Common Bacterial Blight (CBB)	DOR 771	Small red	1.00	1481.00
	DOR 662	Carioca	1.00	1042.00
	DOR 766	Small red	1.00	1715.00
	KG 4-20	Small red	1.00	1223.00
	RRN 47	Small red	1.00	1485.00
	Selian 05	Khaki	1.00	1171.00
	DOR 711	Small red	1.00	1368.00
	MAZ 41	Medium red	2.00	927.00
	DOR 708	Red kidney	2.00	1199.00
	KG 114-185	Small red	2.00	1542.00
Bean Common Mosaic Virus (BCMV)	296/6	Carioca	1.00	1278.00
	ALS 3	Black	1.00	1472.00
	BAT 332	Cream	1.00	1391.00
	C.2014/Hu/11	White	1.00	1149.00
	C.2017/Hu/11	White	1.00	1224.00
	C.2019/Hu/11	White	1.00	1244.00
	CANPSULA	White	1.00	1014.00
	CIM 9313-1	Khaki	1.00	1500.00
	CN Bunsu (62)	White	1.00	1145.00
	CN Bunsu (64)	White	1.00	897.00

pathogen was observed during this study. Extreme rainfall and relative humidity in the field created an environment for diseases occurrence. Each bean genotype responded differently to pathogen infection in different environments, though phenotypic expression was strongly influenced by this pathogen. Some reports

also stated that this pathogen caused serious infection because of high level of moisture in dry bean production areas, such as excess moisture in the field in Kenya (Mwang'ombe et al., 2007) and great lake regions in Tanzania, Uganda and Ethiopia (Pastor-Corrales et al., 1998; Nkalubo et al., 2007). Some genotypes which

tolerated excess moisture identified in this study could be selected for production in these specific areas. Also, other effects of ALS to yield loss in dry bean genotypes was observed in a three-year field experiment conducted by Jesus et al. (2001) in the 1997 and 1999 experiments, whereby ALS reached higher disease levels than rust under rust and ALS inoculations pathogens. Same results were gotten by Cichy et al. (2015), Liebenberg et al. (1997) on Andean diversity study, which was attributed to G\*E and inappropriate good agricultural practices.

Powdery mildew severity was less considered as economic disease in Tanzania, but these experiments showed its economic impact, in that those plants attacked failed to produce even a single seed. The infection was high, up to 6.00 severity scores at Lyamungo site for most large seeded genotypes including NUA 137, NUA 145, NUA 15, SWAP 09 and NUA 244 which leads to poor grain yield of 1016.87, 758.28, 1056.09, 549.89 and 857.27kg/ha, respectively. Same infection was caused by *E. polygoni* pathogen of PM causes extensive damage with significant losses of up to 69% in Columbia, US; the infection occurred during flowering time (Steadman et al., 2005) and other losses of about 40 to 50% at Mexico's farmers' field. Effects of CBB range from leaves, pods to seeds and that's why it is regarded as seed borne disease (Lopez et al., 2006). The effects occurred both in the field through natural infection, which occurs under normal environmental conditions, and in greenhouse through inoculation procedures. High severity occurs under high rainfall and relative humidity as well as warm temperature conditions, between 25 and 35°C (Chaube et al., 1992). However, in the green house, high relative humidity does not influence the CBB, causing organisms to infect plants like that under field conditions (Akhavan et al., 2009).

This study shows the environmental condition for the diseases to occur across the tested locations. *Bean common mosaic virus* showed less variation in its occurrence in the testing sites, while few genotypes were affected by this virus. Despite not expressing its economic importance, but for those few genotypes affected, it really hit across the tested environments. The losses caused by BCMV and BCMNV impacted severely not only on commercial scale cultivation of this high-value crop but also on production by smallholder farmers in the developing world, where bean serves as a key source of dietary protein and mineral nutrition (Worrall et al., 2015). The tested resource materials in this study reflects a better source of Mesoamerican gene pool to improve the common bean growing cultivars, especially those succumbed by ALS, PM, CBB and BCMV diseases (Table 1). Even the identified Andean with moderate resistant for the above diseases could be improved via available resistant bean varieties with the aim to develop the preferred market classes based on region preferences. Additionally, the study shows that extreme rainfall and temperature affects bean grown with disease occurrence and flower abortion respectively, which can

cause poor yield. The genetic and environmental differences contribute to the phenotype and sometimes it is used to confirm the GxE interactions among treatment effects (Tabery and Griffiths, 2010). The future approach on these materials will be to carry genotypic sequencing for better correlation between field and laboratory data.

## Conclusion

From this research study, different genotypes expressed trait of resistance under field conditions for the four foliar economic diseases of ALS, CBB, PM and BCMV. Most of the small seeded genotypes dominated the identified genotypes for each trait of focus on the target diseases. This reflects the opportunities to improve the Mesoamerican gene pool across the dry bean research networks. Despite this fact, some of the genotypes expressed trait of resistance under the field but more work needs to be done under the controlled environment with focus placed on the identified genotypes. Apart from disease, other environmental factors like ambient temperature, rainfall variations and relative humidity affected the yield variations.

## CONFLICT OF INTEREST

The authors have not declared any conflict of interests.

## ACKNOWLEDGEMENT

The authors are grateful for the support rendered by Tuskegee University through the Integrative Biosciences (IBS) Ph.D. Program under the synergism of the College of Agriculture, Environment and Nutrition Sciences (CAENS), College of Arts and Sciences (CAS) and College of Veterinary Medicine (CVM). The authors also appreciate the Government of Tanzania (GoT) for the assistance rendered through Tanzania Agricultural Research Institute – Selian Centre and the International Center for Tropical Agriculture (CIAT) as well as Pan-Africa Bean Research Alliance (PABRA).

## REFERENCES

- Akhavan A, Bahar M, Saeidi G, Lak M (2009). Factors affecting epiphytic population of *Xanthomonas axonopodis* pv. *phaseoli* in epidemiology of bean common blight. *Journal of Science and Technological Agriculture and Natural Resources* 13:265-277.
- Binagwa PH (2017). State of art: National report - Phaseolus bean research program - Tanzania. Arusha, Arusha, Tanzania.
- Brady N, Weil R (2002). *The nature and properties of soils* 3rd Edition. New Jersey: Prentice Hall.
- Broughton W, Hernandez G, Blair M, Beebe S, Gepts P, Vanderleyden J (2003). Beans (*Phaseolus vulgaris* spp.)-model food legumes. *Plant Soil* 252:55-128.
- Buruchara R, Mukankusi C, Ampofo K (2010). *Bean disease and pests identification management-Handbook for small-scale seed producer.*



- Kampala: CIAT.
- Cichy K, Porch T, Beaver J, Cregan DF, Glahn R, Grusak M, Kamfwa K, Katuramu DN, McClean P, Mndolwa E, Nchimbi-Msolla S, Pastor-Corrales MA, Miklas P (2015). A *Phaseolus vulgaris* Diversity Panel for Andean Bean Improvement. *Crop Science* 55:2149-2160.
- Chaube H, Kumar J, Mukhopadhyay A (1992). *Plant diseases of international importance*. New Jersey: Prentice Hall.
- CIAT (1987). *Standard system for the evaluation of bean germplasm*. Van Schonhoven (Compilers). Cali: CIAT.
- CIAT-Kawanda (2013). *Training manual on phenotyping diseases of common bean (*Phaseolus vulgaris* L.) and characterization of the disease pathogens*. Kampala: CIAT.
- Clewer AG, Scarisbrick DH (2001). *Practical statistics and experimental design for plant and crop science*. Chichester. New York: Wiley.
- De Leque J, Creamer B (2014). Major constraints and trends for common bean production and commercialization; establishing priorities for future research. *Agronomia Colombiana* 32(3):423-431.
- FAOSTAT (2018). *Food and Agriculture Organization*. Retrieved from [www.faostat.fao.org](http://www.faostat.fao.org)
- Harwell L, Hood L, Goldberg M, Reynolds A, Silver L (2011). *Genetics: From genes to genomes*. Boston: McGraw Hill.
- Steadman J, Hall R, Forster R (2005). *Compendium of Bean Diseases*. The American Phytopathological Society 20:209-2018.
- Hillocks R, Madata C, Chirwa R, Minja E, Msolla S (2006). *Phaseolus* Bean Improvement in Tanzania, 1959–2005. *Euphytica* 150(1):215-231
- Jesus W, Ribeiro do Vale F, Coelho R, Hau B, Zambolim L, Costa L, Bergamin A (2001). Effects of Angular Leaf Spot and Rust on Yield Loss of *Phaseolus vulgaris*. *Phytopathology* 91(11):1045-1053.
- Kiriba K, Msaky M, Mahenge N, Paul S, Kessy G, Binagwa PH (2020). Yield response of bush bean varieties (*Phaseolus vulgaris*) to different planting densities and rates of phosphorus fertilizer. *International Journal of Agricultural Research* 15(1)40-48.
- Kwak M, Gepts P (2009). Structure of genetic diversity in the two major gene pools of common bean (*Phaseolus vulgaris* L., Fabaceae). *Theoretical and Applied Genetics* 118(5):979-992.
- Kwak M, Kami J, Gepts P (2009). The putative Mesoamerican domestication center is in Lerma-Santiago Basin of Mexico. *Crop Science* 49(2):554-563.
- Landon A (2008). The "how" of the three sisters: The origin of agriculture in Mesoamerican and the human niche. *Nebraska Anthropology* 40:110-124.
- Landon J (1991). *Booker tropical soil manual. A handbook for soil survey and agricultural land evaluation in the tropics and sub-tropics*. New York: Longman Scientific and Technical.
- Liebenberg M, Pretorius Z (1997). A review of angular leaf spot of common bean (*Phaseolus vulgaris* L.). *African Plant Protection* 3:81-106.
- Lopez R, Asensio C, Gilbertson R (2006). Phenotypic and genetic diversity in strains of common bacterial blight. (*Xanthomonas campestris* pv. *phaseoli* and *X. campestris* pv. *phaseoli* var. *fuscans*) in a secondary center of diversity of the common bean host suggests multiple introduction events. *Phytopathology* 96:1204-1213
- Mahuku GS, Iglesias AM, Jara C (2009). Genetics of angular leaf spot resistance in the Andean common bean accession G5686 and identification of markers linked to the resistance genes. *Euphytica* 167:381-396
- Mahuku S, Riascos J (2004). Virulence and molecular diversity within *Colletotrichum lindemuthianum* isolates from Andean and Mesoamerican bean varieties and regions. *European Journal of Plant Pathology* 110:253-263.
- Mwaipopo B, Msolla S, Njau P, Tairo F, William W, Binagwa P, Mbanzibwa D (2016). Viruses infecting common bean (*Phaseolus vulgaris* L.) in Tanzania. A review on molecular characterization, detection and disease management options. *African Journal of Agricultural Research* 12(18):1486-1500.
- Mwang'ombe A, Wagara I, Kimenju J, Buruchara R (2007). Occurrence and Severity of Angular Leaf Spot of Common Bean in Kenya as Influenced by Geographical Location, Altitude and Agroecological Zones. *Plant Pathology Journal* 6:235-241.
- Nkalubo S, Melis R, Laing M, Opio F (2007). Yield loss associated with anthracnose disease on Ugandan market class dry bean cultivars. *African Crop Science Conference Proceedings* 8:869-874.
- Pastor-Corrales M, Jara C, Singh S (1998). Pathogenic variation in sources of and breeding for resistance to *Phaeoisariopsis griseola* causing angular leaf spot in common bean. *Euphytica* 103:161-171.
- Peters A (1993). China. *Michigan Dry Bean Digest* 4:18-20.
- Sankaran S, Helmut KDS, Okamuro J, Poland J, Kalcsits L (2015). *Advances in field based high throughput phenotyping and data management*. Washington DC: Grains and specialty crops.
- Singh S, Schwartz H (2010). *Breeding Common Bean for Resistance to Diseases: A Review*. *Crop Sciences* 50:2199-2223.
- Singh SP, Munoz CG, Teran H (2001). Registration of common bacterial blight resistant dry bean germplasm VAX 1, VAX 3 and VAX 4. *Crop Science* 41:275-276.
- Wang WB, Clayton D, Todd J (2005). Genome-wide association studies: Theoretical and practical concerns. *Nature Reviews Genetics* 6:109-116.
- Worrall E, Wamonje F, Mukeshimana G, Harvey J, Carr JM (2015). *Bean Common Mosaic Virus and Bean Common Mosaic Necrosis Virus: Relationships, Biology, and Prospects for Control*. Elsevier 93:34-39.
- Yu J, Holland J, McMullen M, Buckler E (2008). Genetic design and statistical power of nested association mapping in maize. *Genetics* 178:539-551.

## Supplement 1. Seed characterization.

S/N	Genotype	Gene pool	Origin	Market class	Seed color	Seed size
1	MAZ 37	Andean	CIAT breeding line	Red kidney	Red	Large
2	MAZ 41	Mesoamerican	CIAT breeding line	Red mottled	Red	Medium
3	MAZ 42	Mesoamerican	CIAT breeding line	Red mottled	Red mottled	Medium
4	MAZ 44	Andean	CIAT breeding line	Red mottled	Red mottled	Large
5	MAZ 46	Andean	CIAT breeding line	Red mottled	Red mottled	Large
6	MAZ 47	Andean	CIAT breeding line	Red mottled	Red mottled	Large
7	MAZ 48	Andean	CIAT breeding line	Red mottled	Red mottled	Large
8	MAZ 50	Mesoamerican	CIAT breeding line	Red mottled	Red mottled	Medium
9	MAZ 52	Mesoamerican	CIAT breeding line	Red mottled	Red mottled	Medium
10	MAZ 59	Mesoamerican	CIAT breeding line	Red mottled	Red	Medium
11	MAZ 56	Andean	CIAT breeding line	Red mottled	Red mottled	Large
12	MAZ 57	Andean	CIAT breeding line	Red mottled	Red mottled	Large
13	MAZ 49	Andean	CIAT breeding line	Red mottled	Red mottled	Large
14	MAZ 70	Mesoamerican	CIAT breeding line	Carioca	Brown striped	Small
15	MAZ 72	Andean	CIAT breeding line	Red kidney	Red	Large
16	MAZ 74	Andean	CIAT breeding line	Red kidney	Red	Large
17	MAZ 84	Andean	CIAT breeding line	Red mottled	Red mottled	Large
18	MAZ 255	Andean	CIAT breeding line	Red mottled	Red mottled	Large
19	DOR 662	Mesoamerican	CIAT breeding line	Carioca	Brown striped	Small
20	DOR 708	Mesoamerican	CIAT breeding line	Red kidney	Purple	Medium
21	DOR 710	Mesoamerican	CIAT breeding line	Small red	Red	Small
22	DOR 711	Mesoamerican	CIAT breeding line	Small red	Red	Small
23	DOR 755	Mesoamerican	CIAT breeding line	Small red	Red	Small
24	DOR 766	Mesoamerican	CIAT breeding line	Small red	Red	Small
25	CN Bunsí (60)	Mesoamerican	CIAT breeding line	Navy	White	Small
26	CN Bunsí (62)	Mesoamerican	CIAT breeding line	Navy	White	Small
27	CN Bunsí (63)	Mesoamerican	CIAT breeding line	Navy	White	Small
28	CN Bunsí (64)	Mesoamerican	CIAT breeding line	Navy	White	Small
29	CN Bunsí (65)	Mesoamerican	CIAT breeding line	Navy	White	Small
30	CN Bunsí (66)	Mesoamerican	CIAT breeding line	Navy	White	Small
31	CN Bunsí (67)	Mesoamerican	CIAT breeding line	Navy	White	Small
32	CN Bunsí (68)	Mesoamerican	CIAT breeding line	Navy	White	Small
33	NUA 209	Andean	CIAT breeding line	Red mottled	Red mottled	Large
34	NUA 210	Andean	CIAT breeding line	Red mottled	Red mottled	Large
35	NUA 272	Mesoamerican	CIAT breeding line	Yellow	Cream	Medium
36	NUA 231	Andean	CIAT breeding line	Red mottled	Red mottled	Large
37	NUA 244	Andean	CIAT breeding line	Red mottled	Red mottled	Large
38	NUA 40	Andean	CIAT breeding line	Red mottled	Red mottled	Large

## Supplement 1. Contd.

39	NUA 48	Andean	CIAT breeding line	Red mottled	Red mottled	Large
40	NUA 57	Andean	CIAT breeding line	Red mottled	Red mottled	Large
41	NUA 9	Andean	CIAT breeding line	Red mottled	Red mottled	Large
42	NUA 64	Andean	CIAT breeding line	Red mottled	Red mottled	Large
43	NUA 110	Andean	CIAT breeding line	Red kidney	Red	Large
44	NUA 117	Mesoamerican	CIAT breeding line	–	–	Medium
45	NUA 212	Andean	CIAT breeding line	Red mottled	Red mottled	Medium
46	NUA 213	Andean	CIAT breeding line	Red mottled	Red mottled	Large
47	NUA 125	Mesoamerican	CIAT breeding line	–	Red	Medium
48	NUA 232	Andean	CIAT breeding line	Red mottled	Red mottled	Large
49	NUA 130	Andean	CIAT breeding line	Red mottled	Red mottled	Large
50	NUA 137	Andean	CIAT breeding line	Red kidney	Red	Large
51	NUA 156	Mesoamerican	CIAT breeding line	–	Red	Medium
52	NUA 23	Andean	CIAT breeding line	Red mottled	Red mottled	Large
53	NUA 11	Andean	CIAT breeding line	Red mottled	Red mottled	Large
54	NUA 13	Andean	CIAT breeding line	Red mottled	Red mottled	Large
55	NUA 15	Andean	CIAT breeding line	Red mottled	Red mottled	Large
56	NUA 16	Mesoamerican	CIAT breeding line	Small red	Red	Small
57	NUA 17	Andean	CIAT breeding line	Red mottled	Red mottled	Large
58	NUA 18	Andean	CIAT breeding line	Red mottled	Red mottled	Large
59	NUA 19	Andean	CIAT breeding line	Red mottled	Red mottled	Large
60	NUA 256	Andean	CIAT breeding line	Speckled sugar	Purple striped	Large
61	NUA 30	Andean	CIAT breeding line	Red mottled	Red mottled	Large
62	NUA 31	Andean	CIAT breeding line	Red mottled	Red mottled	Large
63	NUA 134	Andean	CIAT breeding line	Red kidney	Red	Large
64	NUA 115	Andean	CIAT breeding line	Red mottled	Red mottled	Large
65	NUA 116	Andean	CIAT breeding line	Red mottled	Red mottled	Large
66	NUA 129	Mesoamerican	CIAT breeding line	–	Red	Medium
67	NUA 152	Andean	CIAT breeding line	Red kidney	Red	Large
68	NUA 158	Mesoamerican	CIAT breeding line	–	Black stripe	Medium
69	NUA 160	Mesoamerican	CIAT breeding line	Small red	Red	Small
70	NUA 161	Andean	CIAT breeding line	Red kidney	Red	Large
71	NUA 163	Mesoamerican	CIAT breeding line	Small red	Red	Small
72	NUA 165	Mesoamerican	CIAT breeding line	Red mottled	Red mottled	Medium
73	NUA 39	Andean	CIAT breeding line	Red mottled	Red mottled	Large
74	NUA 59	Andean	CIAT breeding line	Red mottled	Red mottled	Large
75	NUA 66	Mesoamerican	CIAT breeding line	Red mottled	Red mottled	Medium
76	NUA 67	Andean	CIAT breeding line	Speckled sugar	Purple striped	Large
77	NUA 79	Andean	CIAT breeding line	Red kidney	Red	Large

## Supplement 1. Contd.

78	NUA 145	Andean	CIAT breeding line	–	Purple	Large
79	NUA 200	Andean	CIAT breeding line	Red mottled	Red mottled	Large
80	NUA 204	Mesoamerican	CIAT breeding line	Red mottled	Red mottled	Medium
81	NUA 207	Mesoamerican	CIAT breeding line	Speckled sugar	Purple striped	Medium
82	NUA 209	Andean	CIAT breeding line	Red mottled	Red mottled	Large
83	NUA 211	Andean	CIAT breeding line	Speckled sugar	Purple striped	Large
84	NUA 224	Andean	CIAT breeding line	Red mottled	Red mottled	Large
85	NUA 225	Andean	CIAT breeding line	Red mottled	Red mottled	Large
86	NUA 226	Andean	CIAT breeding line	Red mottled	Red mottled	Large
87	NUA 229	Andean	CIAT breeding line	Red mottled	Red mottled	Large
88	NUA 233	Andean	CIAT breeding line	Red mottled	Red mottled	Large
89	NUA 235	Andean	CIAT breeding line	Red mottled	Red mottled	Large
90	NUA 236	Andean	CIAT breeding line	Red mottled	Red mottled	Large
91	NUA 238	Andean	CIAT breeding line	Red mottled	Red mottled	Large
92	NUA 239	Andean	CIAT breeding line	Red mottled	Red mottled	Large
93	NUA 240	Andean	CIAT breeding line	Red mottled	Red mottled	Large
94	NUA 245	Andean	CIAT breeding line	Red mottled	Red mottled	Large
95	NUA 257	Andean	CIAT breeding line	Red mottled	Red mottled	Large
96	NUA 273	Andean	CIAT breeding line	Speckled sugar	Purple striped	Large
97	G 5	Mesoamerican	Ethiopia	Navy	White	Small
98	G 23	Mesoamerican	Ethiopia	Navy	White	Small
99	G 30	Mesoamerican	Ethiopia	Navy	White	Small
100	G 31	Mesoamerican	Ethiopia	Navy	White	Small
101	G 60	Mesoamerican	Ethiopia	Navy	White	Small
102	G 78	Mesoamerican	Ethiopia	Navy	White	Small
103	G 79	Mesoamerican	Ethiopia	Navy	White	Small
104	G 87	Mesoamerican	Ethiopia	Navy	White	Small
105	G 90	Mesoamerican	Ethiopia	Navy	White	Small
106	G 100	Mesoamerican	Ethiopia	Navy	White	Small
107	CZ 114-8	Mesoamerican	Kenya	Small red	Red	Small
108	CZ 102-24	Andean	Kenya	Red kidney	Red	Large
109	CZ 102-29	Andean	Kenya	Red kidney	Red	Large
110	CZ 108-27	Mesoamerican	Kenya	Navy	White	Small
111	CZ 114-46	Mesoamerican	Kenya	Small red	Red	Small
112	CZ 114-50	Mesoamerican	Kenya	Small red	Red	Small
113	CZ 114-51	Mesoamerican	Kenya	Small red	Red	Small
114	KG 114-177	Mesoamerican	CIAT breeding line	–	Purple	Small
115	KG 114-178	Mesoamerican	CIAT breeding line	–	Purple	Small
116	KG 114-179	Mesoamerican	CIAT breeding line	Black	Black	Small

## Supplement 1. Contd.

117	KG 114-182	Mesoamerican	CIAT breeding line	Black	Black	Small
118	KG 114-185	Mesoamerican	CIAT breeding line	Small red	Red	Small
119	KG 4-3	Mesoamerican	CIAT breeding line	—	Khak	Small
120	KG 4-20	Mesoamerican	CIAT breeding line	Small red	Red	Small
121	KG 15-6	Andean	CIAT breeding line	Kablanket	Dotted purple	Large
122	KG 30-29	Mesoamerican	CIAT breeding line	Navy	White	Small
123	KG 24-43	Mesoamerican	CIAT breeding line	Navy	White	Small
124	KG 65-5	Mesoamerican	CIAT breeding line	Navy	White	Small
125	KG 67-10	Mesoamerican	CIAT breeding line	Navy	White	Small
126	KG 67-11	Mesoamerican	CIAT breeding line	Navy	White	Small
127	KG 71-4	Mesoamerican	CIAT breeding line	—	Purple	Small
128	KG 71-5	Mesoamerican	CIAT breeding line	—	Purple	Small
129	KG 97-11	Andean	CIAT breeding line	Speckled sugar	Brown striped	Large
130	ZABR 16575-17F22	Mesoamerican	CIAT breeding line	Navy	White	Small
131	ZABR 16575-24F22	Mesoamerican	CIAT breeding line	Navy	White	Small
132	ZABR 16575-39F22	Mesoamerican	CIAT breeding line	Navy	White	Small
133	ZABR 16575-51F22	Mesoamerican	CIAT breeding line	Navy	White	Small
134	ZABR 16575-60F22	Mesoamerican	CIAT breeding line	Navy	White	Small
135	ZABR 16575-86F22	Mesoamerican	CIAT breeding line	Navy	White	Small
136	ZABR 16573-78F22	Mesoamerican	CIAT breeding line	Navy	White	Small
137	ZABR 16574-46F22	Mesoamerican	CIAT breeding line	Navy	White	Small
138	ZABR 16576-11F22	Mesoamerican	CIAT breeding line	Navy	White	Small
139	ZABR 16577-51F22	Mesoamerican	CIAT breeding line	Navy	White	Small
140	RANJONOMBY	Mesoamerican	CIAT breeding line	Navy	White	Small
141	Navy line 5	Mesoamerican	CIAT breeding line	Navy	White	Small
142	Navy line 19	Mesoamerican	CIAT breeding line	Navy	White	Small
143	Navy line 15	Mesoamerican	CIAT breeding line	Navy	White	Small
144	Navy line 22	Mesoamerican	CIAT breeding line	Navy	White	Small
145	Navy line 25	Mesoamerican	CIAT breeding line	Navy	White	Small
146	Navy line 38	Mesoamerican	CIAT breeding line	Navy	White	Small
147	Navy line 40	Mesoamerican	CIAT breeding line	Navy	White	Small
148	Navy line 48	Mesoamerican	CIAT breeding line	Navy	White	Small
149	Navy line 43	Mesoamerican	CIAT breeding line	Navy	White	Small
150	Navy line 51	Mesoamerican	CIAT breeding line	Navy	White	Small
151	Navy line 52	Mesoamerican	CIAT breeding line	Navy	White	Small
152	Navy line 54	Mesoamerican	CIAT breeding line	Navy	White	Small
153	RWR 2075	Andean	Rwanda	Red kidney	Red	Large
154	RWR 1059	Mesoamerican	Rwanda	Red mottled	Red mottled	Medium
155	KABABALA	Mesoamerican	CIAT breeding line	Navy	White	Small

## Supplement 1. Contd.

156	SWP 12	Mesoamerican	Kenya	Navy	White	Small
157	SWP 09	Andean	Kenya	White kidney	White	Large
158	SWP 10	Mesoamerican	Kenya	Navy	White	Small
159	MEXICO 54	Mesoamerican	CIAT breeding line	Carioca	Cream	Medium
160	PAN 72	Mesoamerican	CIAT breeding line	Navy	White	Small
161	JESCA	Andean	Tanzania	Kablanket	Dotted purple	Large
162	Selian 05	Mesoamerican	Tanzania	–	Khak	Small
163	CC 13	Mesoamerican	CIAT breeding line	Small red	Red	Small
164	CC 547	Mesoamerican	CIAT breeding line	Black	Black	Small
165	CC 814	Mesoamerican	CIAT breeding line	Carioca	Brown striped	Small
166	CC 960	Mesoamerican	CIAT breeding line	Black	Black	Small
167	RRN 47	Mesoamerican	CIAT breeding line	Small red	Red	Small
168	RRN 48	Andean	CIAT breeding line	Red mottled	Red mottled	Large
169	217/7	Mesoamerican	CIAT breeding line	–	Khak	Small
170	217/2	Mesoamerican	CIAT breeding line	–	Khak	Small
171	222/1	Mesoamerican	CIAT breeding line	Black	Black	Small
172	296/6	Mesoamerican	CIAT breeding line	Carioca	Brown striped	Small
173	RAZ 36	Mesoamerican	CIAT breeding line	Navy	White	Small
174	RAZ 44	Mesoamerican	CIAT breeding line	Navy	White	Small
175	CIM 9313-1	Mesoamerican	CIAT breeding line	–	Khak	Small
176	Lyamungo 90	Andean	Tanzania	Red mottled	Red mottled	Large
177	ALS 3	Mesoamerican	CIAT breeding line	Black	Black	Small
178	BAT 332	Mesoamerican	CIAT breeding line	Cream	Cream	Small
179	IBC 2	Mesoamerican	CIAT breeding line	Navy	White	Small
180	Awash-1	Mesoamerican	Ethiopia	Navy	White	Small
181	Awash Meka	Mesoamerican	Ethiopia	Navy	White	Small
182	PI 207262	Mesoamerican	CIAT breeding line	–	Khak	Small
183	A 686	Mesoamerican	CIAT breeding line	Black	Black	Small
184	A 774	Mesoamerican	CIAT breeding line	Cream	Cream	Small
185	A 797	Andean	CIAT breeding line	Red mottled	Red mottled	Medium
186	G 5686	Mesoamerican	CIAT breeding line	Cream	Cream	Large
187	CANPSULA	Mesoamerican	CIAT breeding line	Navy	White	Small
188	TU	Mesoamerican	CIAT breeding line	Black	Black	Small
189	CAL 113	Andean	CIAT breeding line	Red mottled	Red mottled	Large
190	C.202/Hu/3	Mesoamerican	CIAT breeding line	Navy	White	Small
191	C.202/Hu/11	Mesoamerican	CIAT breeding line	Navy	White	Small
192	C.2014/Hu/11	Mesoamerican	CIAT breeding line	Navy	White	Small
193	C.2019/Hu/11	Mesoamerican	CIAT breeding line	Navy	White	Small
194	C.2017/Hu/11	Mesoamerican	CIAT breeding line	Navy	White	Small

## Supplement 1. Contd.

195	C.2018/Hu/11	Mesoamerican	CIAT breeding line	Navy	White	Small
196	MLB 17-89A	Andean	CIAT breeding line	Red mottled	Red mottled	Large
197	MLB 40-89A	Mesoamerican	CIAT breeding line	–	Khak	Small
198	MLB 48-89A	Mesoamerican	CIAT breeding line	–	Purple	Small
199	Amendon	Mesoamerican	CIAT breeding line	–	Dotted purple	Medium
200	Vax 1	Mesoamerican	CIAT breeding line	Carioca	Brown striped	Small
201	Vax 2	Mesoamerican	CIAT breeding line	Carioca	Brown striped	Small
202	FLORDEMAYO	Mesoamerican	CIAT breeding line	–	Dotted purple	Small
203	FEB 181	Mesoamerican	CIAT breeding line	Small red	Red	Small
204	FEB 189	Mesoamerican	CIAT breeding line	Small red	Red	Small
205	SAB 662	Mesoamerican	CIAT breeding line	Navy	White	Small
206	CORNELL 49822	Mesoamerican	CIAT breeding line	black	Black	Small
207	SM 133	Mesoamerican	CIAT breeding line	Navy	White	Small
208	MONT-CALM	Andean	CIAT breeding line	Red kidney	Red	Large
209	MICHETTE	Mesoamerican	CIAT breeding line	Navy	White	Small
210	MEXICAN 142	Mesoamerican	CIAT breeding line	Navy	White	Small
211	R.K. MICHIGA	Andean	CIAT breeding line	Red kidney	Red	Large
212	DONTIMOTEA	Mesoamerican	CIAT breeding line	small red	Red	Small
213	F1 Population	Mesoamerican	CIAT breeding line	Navy	White	Small

## Supplement 2. Combined data analysis.

S/N	Accessions	DFF days		DM days		ALS score		PM score		CBB score		BCMV score		Yield (kg/ha)	
		LYA	SARI	LYA	SARI	LYA	SARI	LYA	SARI	LYA	SARI	LYA	SARI	LYA	SARI
1	217/2	49.25	41.50	99.75	97.75	2.00	1.50	2.75	1.25	1.75	2.25	1.00	1.25	2021.48	1244.30
2	222/1	49.00	43.00	103.50	102.50	2.75	1.00	2.00	1.50	1.50	2.25	3.25	3.50	1644.77	1201.25
3	296/6	48.25	43.50	99.75	97.25	2.25	1.25	2.75	1.25	1.25	2.25	1.00	1.00	1513.12	1042.66
4	A 686	48.25	43.00	101.75	102.50	2.50	1.50	2.25	2.00	1.25	3.25	5.00	4.00	1175.31	1502.73
5	A 774	45.25	40.00	99.25	95.00	3.25	1.75	2.00	1.50	1.00	2.75	1.00	1.75	2350.23	1613.05
6	A 797	47.75	40.50	101.75	100.00	2.25	2.00	2.25	1.75	1.50	2.25	1.00	1.25	1488.75	813.20
7	ALS 3	46.25	38.50	100.50	98.75	3.50	1.25	2.75	1.00	2.00	2.00	1.00	1.00	2152.27	790.97
8	AMENDON	44.75	39.75	100.00	98.50	3.50	2.00	3.50	2.00	1.75	2.50	1.50	1.50	1158.44	1030.86
9	Awash Meka	49.25	44.25	102.50	100.50	1.50	2.00	2.25	1.50	3.75	2.00	1.00	1.25	1952.03	992.27
10	Awash-1	47.75	41.00	101.50	99.75	2.00	1.50	1.50	1.25	2.75	2.00	1.25	1.00	1339.06	1201.70
11	BAT 332	49.75	43.25	101.50	99.75	2.25	2.00	2.50	1.00	1.75	2.25	1.00	1.00	2033.44	749.06
12	C.2014/Hu/11	44.50	37.75	99.75	101.50	2.50	1.50	2.50	2.25	3.00	2.00	1.00	1.00	1292.58	1005.00

## Supplement 2. Contd.

13	C.2017/Hu/11	49.00	43.00	103.25	102.75	2.50	2.75	3.00	1.50	3.00	1.50	1.00	1.00	1576.25	872.73
14	C.2018/Hu/11	45.00	39.50	101.00	101.75	2.50	2.00	1.50	1.50	2.50	2.00	1.00	1.50	759.06	1117.89
15	C.2019/Hu/11	47.25	42.50	101.75	105.50	2.00	2.00	2.25	1.00	3.25	2.00	1.00	1.00	1678.75	808.52
16	C.202/Hu/3	45.00	38.75	99.00	101.00	1.50	2.75	2.25	1.00	3.50	2.00	1.00	1.50	1093.28	672.03
17	CAL 113	42.25	39.00	97.00	96.00	3.50	2.25	2.50	2.00	2.25	3.00	1.00	1.25	1418.37	856.68
18	CANPSULA	46.00	38.75	102.25	100.75	2.00	1.75	2.25	1.00	2.75	1.75	1.00	1.00	1065.31	961.72
19	CC 13	49.25	43.00	103.00	98.50	3.00	1.50	2.00	2.00	1.25	2.50	1.75	1.75	1806.56	1259.92
20	CC 547	49.50	43.00	102.00	98.50	2.00	1.50	2.50	1.25	1.25	2.00	1.25	1.00	1918.91	1342.34
21	CC 814	47.50	42.00	97.75	98.75	2.25	1.00	2.50	1.50	1.75	2.50	1.00	1.50	1974.53	715.78
22	CC 906	47.50	44.50	104.00	97.50	2.50	1.50	2.00	1.00	1.25	2.25	2.75	3.75	1535.39	1160.86
23	CIM 9313-1	49.75	44.25	100.50	96.75	2.25	1.75	2.25	1.00	1.75	2.00	1.00	1.00	1557.81	1442.42
24	CN Bunsu (60)	49.00	42.00	103.00	100.25	1.50	1.75	1.50	1.50	2.25	1.75	1.00	1.25	776.87	852.66
25	CN Bunsu (62)	49.25	42.75	104.50	100.75	1.75	1.50	3.00	1.00	2.50	2.50	1.00	1.00	1653.05	636.41
26	CN Bunsu (63)	44.50	39.50	98.00	99.75	2.00	1.75	1.50	1.25	2.75	1.75	1.25	1.00	1040.94	1121.56
27	CN Bunsu (64)	46.75	40.25	98.75	99.25	1.75	1.25	1.25	1.00	3.00	2.00	1.00	1.00	1105.08	653.75
28	CN Bunsu (65)	45.00	42.75	97.25	99.25	2.75	1.75	2.75	1.25	2.50	1.25	1.00	1.00	1174.30	855.94
29	CN Bunsu (66)	48.50	41.00	99.75	95.00	2.50	2.00	2.75	2.75	2.25	2.25	1.00	1.00	2309.30	895.56
30	CN Bunsu (67)	44.75	39.50	98.25	95.25	2.00	2.00	2.00	1.50	2.75	2.00	1.00	1.50	1192.03	1037.69
31	CN Bunsu (68)	50.25	42.25	104.50	100.00	1.50	2.50	2.50	1.75	2.75	2.25	1.00	1.75	1330.00	862.97
32	CORNELL 49242	48.25	43.00	103.75	102.75	3.00	1.00	3.25	2.00	1.50	2.50	1.00	1.00	1833.12	675.70
33	CZ 102-24	39.75	34.50	95.75	94.75	4.25	2.75	2.00	1.25	1.75	2.50	1.00	1.00	1950.43	1281.78
34	CZ 102-29	42.25	36.00	97.25	99.50	3.25	2.00	2.75	1.25	1.75	1.50	1.00	1.25	931.73	922.55
35	CZ 108-27	47.00	39.25	100.00	98.00	2.00	2.00	2.00	1.75	1.50	2.00	1.00	1.75	817.60	971.16
36	CZ 114-46	40.25	35.25	93.50	98.75	3.75	1.50	1.75	1.25	1.50	2.25	1.00	1.00	1462.84	1158.22
37	CZ 114-50	40.25	34.75	94.75	99.00	2.50	1.25	2.75	2.00	2.00	2.00	1.00	1.00	1474.09	1425.10
38	CZ 114-51	41.00	36.75	95.75	72.25	3.50	2.25	2.50	1.25	1.75	2.50	1.00	1.25	1194.47	1268.41
39	CZ 114-8	43.00	38.25	97.25	98.25	3.00	2.50	2.50	2.25	1.75	2.50	1.00	1.75	1057.69	1271.36
40	DONTIMOTEO	47.25	38.00	103.50	99.50	2.00	3.25	2.50	2.25	1.50	2.50	1.00	2.25	568.61	604.92
41	DOR 662	47.00	41.75	100.75	98.50	1.75	2.25	1.75	1.00	1.00	1.50	1.00	1.00	1209.06	875.70
42	DOR 708	47.50	42.00	100.25	102.25	2.50	1.50	2.50	1.25	1.25	1.75	1.00	1.00	1257.27	1141.02
43	DOR 710	45.75	40.50	101.00	98.25	2.75	1.50	1.75	1.00	4.00	3.50	1.25	1.25	1888.98	1437.03
44	DOR 711	49.00	41.00	101.25	103.00	2.25	2.00	2.50	1.00	1.75	1.00	1.00	1.00	2027.27	707.89
45	DOR 755	49.50	44.00	105.25	103.25	1.75	1.00	2.50	1.75	1.50	3.00	4.25	3.00	2001.87	1088.67
46	DOR 766	45.75	39.75	100.25	99.75	2.25	1.25	2.50	1.00	1.25	1.25	1.50	1.00	2359.37	1070.47
47	DOR 771	48.25	41.75	101.25	101.25	2.25	2.00	2.25	1.50	1.00	1.00	1.75	1.00	1780.78	1180.47
48	F1POPULATION	44.75	41.50	100.00	99.00	2.00	1.50	1.50	1.00	3.25	1.75	1.50	1.00	1117.73	714.87
49	FEB 181	47.50	41.75	101.25	100.00	2.75	2.00	2.75	1.50	1.25	2.50	2.25	1.50	1459.30	977.95



## Supplement 2. Contd.

50	FEB 189	46.50	39.00	100.25	98.25	2.00	1.75	1.00	1.75	1.00	2.25	1.00	1.75	2590.31	1662.94
51	FLOR DE MAYO	46.00	39.25	103.75	103.75	4.00	1.25	2.00	1.00	3.00	3.00	3.00	3.75	915.87	678.75
52	G 100	44.25	39.50	96.75	93.75	2.00	1.50	2.25	1.75	3.50	2.25	1.00	1.00	1103.05	1084.05
53	G 23	48.50	40.50	101.50	98.75	2.50	2.25	3.00	2.25	2.50	1.50	1.00	1.00	1830.78	1185.62
54	G 30	44.75	40.25	96.75	100.25	2.00	1.75	1.50	1.00	3.25	2.00	1.00	1.00	1334.45	1045.55
55	G 31	46.25	44.00	105.50	101.50	1.75	1.50	1.75	1.50	3.75	1.75	1.00	1.75	1412.50	925.23
56	G 5	46.00	39.75	100.00	100.25	1.75	2.00	1.75	1.25	2.50	2.00	1.50	1.00	1174.61	981.09
57	G 5686	41.00	37.00	98.00	97.75	3.00	2.00	3.00	2.00	1.75	2.00	1.00	2.00	1095.23	1092.78
58	G 60	46.00	39.50	99.50	101.00	1.50	2.00	1.50	1.00	3.50	1.75	1.00	1.00	1910.31	846.33
59	G 78	49.00	43.00	103.25	99.25	1.75	1.50	2.50	1.25	3.50	1.75	1.00	1.00	1394.84	1084.06
60	G 79	46.50	42.00	101.75	102.00	2.00	2.50	2.00	1.75	3.50	2.00	1.00	2.25	1178.44	842.75
61	G 87	48.00	41.50	102.75	99.00	1.75	2.00	3.00	2.25	3.00	2.25	1.00	1.25	978.44	778.91
62	G 90	42.75	35.75	99.00	99.50	3.00	2.75	3.25	1.25	2.75	3.00	1.00	1.25	467.97	857.97
63	IBC 2	45.00	42.00	104.25	98.50	1.50	2.00	1.50	1.75	2.50	2.75	1.25	1.75	1277.19	793.81
64	JESCA	35.00	34.50	93.50	92.25	4.00	2.25	2.50	1.50	1.50	2.00	1.00	1.25	1435.00	1319.42
65	KABALABALA	47.75	40.00	101.00	97.00	2.50	1.00	1.75	1.75	1.50	2.00	1.00	1.00	448.94	830.00
66	KG 114-177	42.75	37.25	97.25	95.75	2.75	1.50	2.50	1.50	2.50	2.00	1.00	1.00	1628.46	1545.24
67	KG 114-178	44.75	40.00	97.25	96.00	2.25	2.00	3.25	1.25	2.25	1.75	1.00	1.25	1047.98	1214.76
68	KG 114-179	43.25	38.00	98.50	97.00	2.50	2.00	2.50	1.75	2.00	1.75	1.00	1.25	1341.87	1318.12
69	KG 114-182	42.25	37.25	97.25	72.75	2.50	1.75	2.75	2.00	1.25	2.25	1.25	1.00	1639.62	1673.37
70	KG 114-185	43.50	38.25	95.75	97.25	3.75	2.25	2.25	1.50	1.00	2.00	1.25	1.00	1683.41	1400.67
71	KG 15-6	43.00	38.00	99.25	98.00	3.75	2.50	2.75	1.25	1.50	3.00	1.00	1.25	1777.93	1298.85
72	KG 24-43	46.00	43.00	103.75	102.00	2.75	1.75	2.00	2.00	1.75	2.25	1.00	1.50	1020.29	826.88
73	KG 30-29	49.00	43.00	103.25	103.00	1.50	1.00	2.00	1.75	1.75	2.00	1.75	1.25	1477.69	1035.77
74	KG 4-20	42.25	37.25	96.50	100.50	3.25	1.50	3.00	2.50	1.25	1.25	1.00	1.25	1250.38	1194.90
75	KG 4-3	44.75	39.75	97.75	99.50	3.00	2.00	2.50	1.50	1.50	2.50	1.00	1.00	1820.14	1166.20
76	KG 67-10	48.00	39.50	101.25	101.75	2.50	2.00	2.75	1.50	3.00	1.75	1.00	1.00	1394.69	686.14
77	KG 67-11	45.25	38.75	98.50	100.75	1.75	2.75	1.50	1.75	2.50	1.75	1.00	1.50	1278.67	1049.61
78	KG 67-5	44.25	41.50	99.50	98.75	1.75	2.00	3.75	1.75	2.50	1.50	1.00	1.75	1887.89	861.48
79	KG 71-4	42.50	37.25	96.25	95.25	2.75	2.00	1.75	1.25	1.75	1.75	1.00	1.00	2056.49	1392.88
80	KG 71-5	42.75	37.25	95.75	96.50	3.00	1.50	1.50	1.50	1.50	2.50	2.00	1.00	1709.81	1351.73
81	KG 97-11	47.50	42.75	103.00	101.00	2.25	1.50	3.00	1.75	2.50	2.75	1.00	2.25	1686.72	1001.41
82	LYAMUNGU 90	38.00	33.50	94.50	93.00	3.00	2.00	3.00	2.25	1.25	3.00	1.00	1.00	1078.27	947.21
83	MAZ 37	39.50	37.00	96.50	97.75	3.25	2.25	4.00	2.00	1.50	2.00	1.00	1.00	1100.23	1057.27
84	MAZ 41	40.25	36.25	97.00	99.50	3.75	2.25	3.00	1.25	1.50	1.50	1.00	1.25	732.50	1122.19
85	MAZ 42	42.00	37.00	99.25	98.25	4.00	2.00	2.75	1.00	1.25	2.50	1.00	1.00	884.14	885.94
86	MAZ 44	40.00	33.25	99.00	96.50	4.25	2.50	2.25	1.25	1.25	2.00	1.25	1.00	960.08	911.72

## Supplement 2. Contd.

87	MAZ 46	43.25	37.00	98.50	99.25	4.00	2.25	3.00	2.00	1.50	3.25	1.50	1.50	1596.09	929.92
88	MAZ 47	43.25	38.50	98.25	96.25	3.75	2.50	2.75	2.25	2.25	3.00	1.50	1.50	1102.19	1275.86
89	MAZ 48	39.50	33.75	98.75	98.25	3.50	2.50	4.25	2.25	1.75	2.50	1.00	1.75	1829.61	1330.86
90	MAZ 49	41.25	35.00	99.50	97.50	3.50	2.25	2.00	1.25	1.75	2.25	1.00	1.25	1229.69	1392.50
91	MAZ 50	41.25	35.00	98.75	99.75	4.25	2.25	2.50	1.25	1.50	2.50	1.00	1.25	1227.42	1095.00
92	MAZ 52	41.50	38.25	100.00	101.00	3.50	2.00	2.50	2.00	1.50	2.25	2.50	2.00	1201.17	1160.16
93	MAZ 55	40.25	33.25	95.25	97.50	3.75	2.25	3.25	1.25	1.75	1.25	1.25	1.25	1166.87	1173.94
94	MAZ 56	42.00	40.75	98.25	99.50	3.50	2.00	3.75	2.00	1.50	2.00	1.00	1.00	655.70	897.19
95	MAZ 57	41.00	35.25	97.00	100.75	3.75	2.75	4.00	2.25	1.50	2.50	1.00	1.00	1242.58	1203.98
96	MAZ 59	40.00	33.50	95.50	94.00	4.50	2.50	3.25	1.25	1.25	2.00	4.00	1.75	971.87	836.48
97	MAZ 70	45.75	40.75	102.25	101.75	2.50	1.00	1.75	1.00	1.50	3.25	1.00	1.00	572.42	1421.95
98	MAZ 72	41.50	35.50	97.75	96.50	3.25	1.75	4.00	2.50	1.75	2.50	1.00	1.00	479.61	731.02
99	MAZ 74	41.25	37.00	97.25	96.75	3.25	2.75	3.75	2.25	1.00	2.50	1.00	1.00	771.17	607.11
100	MAZ 84	40.50	35.00	95.50	95.50	4.00	2.50	4.00	1.25	1.25	1.75	1.00	1.25	1151.33	991.33
101	MEX 54	46.00	41.50	101.50	99.75	2.25	1.75	1.50	1.00	2.50	3.00	1.25	1.25	2660.62	789.61
102	MEXICAN 142	48.50	41.50	104.50	103.50	1.75	2.25	2.25	1.25	2.75	1.50	1.00	1.75	1901.64	750.70
103	MICHETTE	45.00	38.25	99.25	97.50	3.25	2.75	1.00	1.50	1.25	2.25	1.50	2.00	644.53	645.94
104	MLB 17-89A	46.50	40.25	100.25	97.75	2.75	1.50	4.50	2.75	1.50	2.50	1.75	1.00	576.80	1186.80
105	MLB 40-89A	49.00	45.75	100.75	96.75	2.25	1.75	1.75	1.00	1.50	1.75	1.00	1.00	1702.81	926.80
106	MLB 48-89A	45.75	42.00	102.25	101.75	2.75	1.00	3.00	1.75	1.00	2.50	4.25	3.25	1423.36	1096.95
107	MONT-CALM	39.00	34.00	95.00	94.00	4.00	3.50	4.50	2.25	1.50	2.50	1.00	1.25	638.83	744.77
108	Navy line 15	42.25	41.00	102.00	104.50	2.25	2.25	2.25	1.75	2.50	2.50	1.50	2.75	1043.59	969.53
109	Navy line 19	45.50	41.75	100.25	100.75	2.00	2.00	2.00	1.50	4.25	2.50	1.00	1.50	1054.61	872.86
110	Navy line 22	44.50	40.25	96.75	96.00	2.25	1.50	1.25	2.00	1.75	2.25	1.00	1.00	1498.67	1005.48
111	Navy line 25	50.50	43.25	104.25	99.75	1.75	2.00	3.50	2.00	2.75	2.00	1.00	1.75	1202.19	1058.31
112	Navy line 38	42.00	38.25	100.25	98.25	2.25	1.75	1.75	1.50	3.00	1.50	1.00	1.50	1973.75	1241.09
113	Navy line 40	40.00	34.00	95.75	85.50	3.00	2.50	1.50	1.25	1.75	1.75	1.00	1.00	1204.61	1406.87
114	Navy line 43	44.75	41.50	97.00	96.75	2.75	2.25	2.00	1.25	3.25	1.75	1.00	1.25	1248.44	1115.00
115	Navy line 48	46.50	42.50	101.75	103.00	2.00	1.25	2.25	1.50	2.25	2.00	1.00	1.00	1629.53	743.98
116	Navy line 5	45.00	39.00	104.50	104.50	2.25	2.00	3.25	1.00	2.50	2.75	1.00	1.25	748.91	605.62
117	Navy line 51	45.50	42.50	101.75	98.25	2.50	1.75	3.75	1.25	2.50	1.50	1.00	1.25	1639.37	1086.95
118	Navy line 52	48.00	44.50	102.25	101.75	2.25	1.50	1.75	1.25	2.75	2.25	1.00	1.00	1703.52	798.41
119	Navy line 54	45.00	37.00	100.00	97.75	2.50	2.00	4.25	1.25	3.25	1.75	1.00	1.00	1257.03	996.84
120	NUA 11	42.50	40.00	98.75	100.75	3.00	2.50	2.25	1.50	3.25	3.00	1.50	2.00	1155.62	914.87
121	NUA 110	39.00	35.75	96.00	98.00	2.25	2.00	5.50	2.00	2.50	2.50	1.00	1.00	727.50	724.06
122	NUA 116	45.00	40.50	100.75	101.25	1.75	1.50	3.00	1.50	3.75	1.75	1.50	1.25	1051.02	1206.56
123	NUA 117	46.00	39.00	102.00	100.25	2.25	1.50	3.50	2.00	3.00	2.75	1.00	1.50	976.09	469.30

## Supplement 2. Contd.

124	NUA 125	40.75	34.25	99.00	100.00	3.00	2.00	4.75	1.00	1.75	2.75	1.00	1.25	1294.61	1020.78
125	NUA 129	38.75	35.25	95.75	96.25	2.50	3.00	3.00	1.25	3.00	3.25	1.00	2.00	677.89	723.58
126	NUA 13	44.25	35.75	101.50	94.00	2.50	2.00	3.00	1.25	2.75	3.00	1.00	3.75	819.45	580.94
127	NUA 130	41.50	35.25	95.50	100.75	2.25	2.25	4.25	2.25	2.50	2.75	2.00	2.25	915.08	658.05
128	NUA 134	42.25	37.00	100.00	102.75	2.25	2.00	2.25	2.00	2.75	3.00	1.25	2.75	1121.72	840.94
129	NUA 137	36.50	37.75	97.25	97.25	2.25	2.50	6.00	3.00	2.00	3.25	1.00	2.00	1016.87	772.16
130	NUA 145	43.25	40.00	99.50	96.50	3.50	2.25	5.00	1.25	2.75	1.75	1.00	1.75	758.28	569.22
131	NUA 15	35.75	34.75	96.75	95.50	2.50	2.25	5.00	1.75	1.75	2.50	1.00	1.00	1056.09	1111.72
132	NUA 152	43.75	38.25	100.25	99.00	2.75	2.00	3.00	1.25	3.50	3.00	1.00	1.25	1629.69	1257.89
133	NUA 156	43.25	40.50	99.25	100.50	2.00	1.50	4.25	2.00	3.25	3.00	1.00	2.00	416.80	455.70
134	NUA 158	46.50	41.00	98.25	100.00	3.25	2.00	3.00	1.00	3.00	3.25	1.00	1.25	616.95	584.83
135	NUA 16	42.75	40.00	98.25	99.25	1.75	1.50	2.25	1.50	3.25	1.75	1.00	1.00	2003.67	1581.41
136	NUA 160	43.75	38.00	98.00	100.50	2.00	1.75	3.50	1.75	3.25	3.00	1.00	1.00	1210.70	1036.09
137	NUA 161	43.25	38.50	102.00	99.00	3.00	2.75	1.50	2.00	2.00	3.75	1.25	2.50	444.22	352.66
138	NUA 163	44.00	38.25	99.50	101.25	2.75	2.50	3.50	1.25	2.75	4.00	1.00	2.25	1042.03	723.83
139	NUA 165	42.75	36.50	97.50	96.75	3.00	2.25	3.75	1.25	3.25	1.50	1.00	1.75	651.64	758.12
140	NUA 17	43.50	38.50	98.75	97.75	2.75	2.50	2.00	1.75	3.25	3.00	1.00	2.00	1392.19	1360.39
141	NUA 18	43.25	35.50	97.25	97.50	2.25	2.25	2.50	1.00	3.50	2.00	1.00	1.25	1574.37	1002.50
142	NUA 19	44.75	38.00	101.00	97.00	3.00	2.00	2.75	1.25	2.75	1.50	1.50	3.00	559.30	907.73
143	NUA 200	43.00	36.75	97.75	95.50	2.50	2.00	4.00	2.25	3.25	2.75	1.00	1.25	967.81	978.12
144	NUA 204	42.00	34.25	100.00	102.50	2.25	1.50	2.50	2.00	2.50	2.50	2.50	1.75	1048.28	1131.87
145	NUA 207	44.25	36.25	101.25	100.25	2.25	2.00	3.25	1.25	4.50	3.25	1.00	1.25	1207.81	1442.03
146	NUA 209	46.50	39.00	100.00	99.00	3.00	1.75	3.25	1.00	2.50	1.50	1.00	1.25	1424.53	1012.66
147	NUA 210	43.00	40.50	99.50	97.50	2.75	2.00	2.25	1.00	3.25	2.75	1.00	2.25	1074.37	861.09
148	NUA 211	40.25	35.00	98.25	100.25	3.25	2.50	3.25	1.25	2.75	3.00	1.00	1.50	972.34	1527.66
149	NUA 212	42.50	36.25	96.00	97.25	2.50	2.25	3.50	1.75	2.50	1.50	1.00	1.25	795.47	1351.09
150	NUA 213	40.75	34.50	98.25	76.75	2.50	2.25	4.75	2.25	2.00	2.00	1.00	1.50	1264.45	771.80
151	NUA 224	42.75	38.00	101.25	100.50	2.75	2.25	3.50	2.00	3.25	3.00	1.00	2.00	1622.50	1219.92
152	NUA 225	43.00	39.50	97.50	99.50	2.50	2.00	2.00	1.25	3.50	2.50	1.00	1.25	601.41	1040.78
153	NUA 226	46.25	40.25	100.75	97.50	2.25	2.00	2.00	2.50	4.00	2.50	1.00	1.00	1025.00	759.45
154	NUA 229	46.00	40.00	101.25	101.00	2.25	2.00	2.50	1.00	3.00	3.00	1.00	1.25	1349.14	1047.27
155	NUA 23	41.25	37.00	98.00	97.25	3.00	2.75	2.75	1.50	3.25	2.50	1.00	2.00	744.84	840.94
156	NUA 231	47.00	41.75	100.75	101.25	1.75	2.25	2.25	1.00	3.00	3.00	1.50	1.25	1033.36	1037.81
157	NUA 232	46.25	38.50	102.25	100.25	2.00	2.25	1.75	1.00	2.50	2.25	1.00	1.00	937.42	1203.59
158	NUA 233	44.25	39.00	101.25	103.25	2.50	2.25	2.75	1.00	2.25	1.75	1.00	1.50	1917.66	1122.42
159	NUA 235	40.75	34.75	96.50	94.75	2.25	2.25	3.75	2.00	3.00	2.50	1.00	1.50	1020.62	1098.98
160	NUA 236	42.75	35.50	98.00	101.00	2.75	2.25	4.50	1.75	2.25	2.50	1.00	1.25	856.56	1007.03

## Supplement 2. Contd.

161	NUA 238	42.00	36.50	96.75	98.25	2.75	2.50	4.00	1.75	2.50	3.00	1.00	1.25	927.19	983.20
162	NUA 239	41.75	39.25	97.50	93.75	2.50	2.00	4.00	1.75	2.75	2.75	1.00	2.00	1817.11	945.23
163	NUA 240	45.00	37.25	98.25	97.50	2.75	2.50	4.25	1.25	3.00	2.25	1.00	2.00	1368.98	1537.89
164	NUA 244	45.25	40.25	98.75	97.25	2.75	3.00	5.00	2.50	2.25	3.00	1.00	1.00	857.27	1018.59
165	NUA 245	41.50	37.25	97.50	96.50	2.50	2.00	3.50	1.50	3.25	2.50	1.00	1.50	1267.11	1239.30
166	NUA 256	43.00	39.00	101.50	100.00	2.25	2.00	4.75	2.00	3.00	2.75	1.00	1.50	1150.78	975.08
167	NUA 257	40.75	35.75	98.25	97.75	2.75	2.00	3.25	1.50	2.75	2.25	1.00	1.75	1244.53	696.72
168	NUA 272	48.00	41.25	98.25	99.25	2.25	1.50	2.75	1.25	3.50	3.50	1.00	1.50	674.14	731.56
169	NUA 273	42.75	37.50	96.25	98.25	2.25	2.00	3.50	1.50	3.00	1.75	1.00	1.25	855.70	867.97
170	NUA 30	39.75	36.50	97.75	99.50	2.50	2.25	4.50	2.50	2.25	1.75	1.00	1.50	670.62	1165.31
171	NUA 31	42.50	36.00	96.50	98.50	2.50	2.00	3.25	1.75	2.75	2.50	1.00	1.50	1379.69	1217.89
172	NUA 39	45.25	39.75	97.25	98.00	2.00	1.75	4.25	2.25	3.25	2.50	1.00	4.00	1507.66	649.84
173	NUA 40	42.00	38.75	102.00	96.75	2.00	2.00	4.75	1.25	2.50	2.00	1.00	2.00	1108.36	960.62
174	NUA 48	46.00	41.75	99.50	95.00	2.00	2.25	4.75	2.50	2.25	4.00	1.00	1.25	1036.64	1223.73
175	NUA 57	46.75	40.25	98.00	96.25	1.75	2.00	3.00	1.50	3.25	3.00	1.25	1.50	758.44	788.05
176	NUA 59	45.75	40.75	102.75	103.75	2.25	2.50	2.75	2.00	3.50	2.50	1.25	2.75	1053.28	1722.67
177	NUA 64	43.75	39.25	98.75	98.25	2.00	2.00	4.00	1.50	3.00	2.50	1.00	1.25	1262.19	964.14
178	NUA 66	42.75	38.25	97.75	100.75	2.25	1.75	3.00	1.00	3.25	2.50	1.00	1.00	1205.23	975.86
179	NUA 67	46.75	41.25	98.50	98.00	2.25	2.00	3.75	2.75	2.50	1.50	1.00	1.50	1127.11	880.00
180	NUA 79	39.50	34.75	96.50	99.25	2.25	1.75	4.50	1.75	2.75	2.75	1.00	1.75	1195.00	773.28
181	NUA 9	43.50	38.50	99.25	99.25	2.75	2.25	3.75	1.50	2.50	2.25	1.00	1.00	1268.83	1055.23
182	PAN 72	45.25	40.75	99.00	98.00	2.75	1.25	1.00	1.00	1.75	2.50	1.00	1.00	1505.00	932.19
183	PI 207262	43.50	36.25	96.25	98.25	3.25	1.75	3.25	4.25	1.25	3.00	1.50	2.00	806.56	920.70
184	R.K. MICHIGA	35.50	35.00	95.25	95.50	3.75	4.00	4.25	1.00	1.00	3.25	1.00	2.50	855.16	982.62
185	RANJONOMBY	48.00	39.50	102.75	101.00	1.50	1.00	2.50	1.50	3.50	3.00	1.00	1.00	1933.36	1425.61
186	RAZ 36	47.25	39.00	100.25	100.00	2.25	2.00	3.00	1.25	3.25	1.25	1.00	1.00	1403.20	824.53
187	RAZ 44	49.50	44.25	103.00	98.25	2.00	1.75	3.00	1.00	2.75	1.50	1.00	1.00	1880.00	1359.45
188	RRN 47	47.50	42.50	99.00	99.75	2.25	1.75	2.00	1.00	1.00	1.50	1.00	1.00	2062.89	906.25
189	RRN 48	45.00	37.50	101.00	99.75	3.25	2.00	1.75	1.00	1.00	2.25	1.00	2.00	2168.36	633.05
190	RWR 1059	45.75	40.75	102.25	103.00	2.75	1.75	2.75	1.00	1.25	2.75	1.00	1.00	1899.37	1215.47
191	RWR 2075	41.50	38.25	99.25	100.50	3.00	2.00	4.50	2.75	1.75	2.25	2.25	1.25	961.17	453.28
192	SAB 662	47.25	42.00	98.00	98.75	3.25	1.75	3.25	2.25	2.00	1.75	1.00	1.00	1029.06	757.11
193	SELIAN 05	43.00	39.00	95.75	93.00	3.00	1.00	3.25	1.50	1.00	1.50	1.00	1.00	1160.67	1181.97
194	SM 133	44.00	41.25	99.50	96.75	2.50	1.00	1.25	1.00	3.25	2.25	1.00	1.25	2112.89	711.17
195	SWP 09	39.00	34.00	96.00	97.50	2.25	2.50	5.25	1.75	2.75	2.50	1.00	1.25	547.89	1367.11
196	SWP 10	46.75	40.25	100.75	100.50	2.25	1.75	1.75	1.00	3.00	1.75	1.00	1.25	1567.34	774.61
197	SWP 12	45.75	39.50	98.00	96.50	2.25	1.00	1.50	1.50	3.75	2.75	1.00	1.00	1616.72	605.16

## Supplement 2. Contd.

198	TU	45.00	39.50	98.50	100.75	2.50	1.25	3.00	1.50	1.25	2.00	1.00	1.50	1121.72	1432.42
199	VAX 1	43.00	36.00	100.00	100.50	3.50	2.00	3.00	1.50	1.50	2.75	1.50	1.75	1953.67	1139.45
200	VAX 2	47.50	39.50	100.50	99.50	3.00	1.00	2.75	1.25	1.25	2.50	1.25	1.25	1445.55	1446.64
201	ZABR 16573-78F22	46.75	40.50	100.50	99.75	1.75	2.00	1.25	1.00	2.75	1.50	1.00	1.00	1236.87	902.66
202	ZABR 16574-46F22	45.50	38.75	99.25	98.50	3.25	2.25	3.75	2.50	2.50	1.75	1.00	1.25	879.53	940.86
203	ZABR 16575-17F22	45.75	41.00	90.75	89.25	2.00	1.25	3.00	1.75	2.25	2.00	1.00	1.00	1862.11	1033.05
204	ZABR 16575-24F22	46.75	44.25	100.50	99.50	1.75	1.50	2.25	1.00	3.00	1.75	1.00	1.00	1802.73	659.22
205	ZABR 16575-39F22	47.75	41.75	99.50	99.50	2.50	1.50	3.00	2.00	2.50	2.00	1.00	1.00	1905.47	600.56
206	ZABR 16575-51F22	49.50	40.50	102.25	102.50	1.50	1.25	2.50	1.50	3.25	2.00	1.00	1.00	1050.23	873.67
207	ZABR 16575-52F22	45.50	38.25	101.25	101.00	2.00	2.25	2.25	2.00	2.75	3.00	1.00	2.00	1118.20	1100.48
208	ZABR 16575-60F22	42.25	36.25	96.75	101.25	2.25	2.25	2.50	1.25	2.50	1.25	1.00	1.25	727.42	856.58
209	ZABR 16575-86F22	48.00	42.25	100.50	101.00	2.25	1.25	2.25	1.25	2.50	2.25	1.00	1.00	1573.05	995.08
210	ZABR 16576-11F22	47.75	42.00	102.75	99.25	1.50	1.75	2.00	2.75	2.00	1.75	1.25	1.00	1214.69	724.77
211	ZABR 16577-51F22	45.00	42.50	99.25	98.25	2.25	2.50	2.00	1.25	2.50	2.50	1.00	1.50	1346.17	730.70
	Mean	44.50	39.16	99.45	98.53	2.75	1.95	2.80	1.58	2.33	2.28	1.19	1.45	1298.07	1005.00
	CV (%)		7.80		6.30		33.60		44.30		35.60		45.90		42.00
	SGE (5%)		ns		ns		<0.001		<0.001		<0.001		<0.001		<0.001

DFF: Days to 50% flowering; MD: days to maturity; ALS: Angular leaf spot; PM: powdery mildew; CBB: Common bacterial blight; BCMV: Bean common mosaic virus; CV: Coefficient variation; SGE: Significant effects; ns: Not significant.