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Screening Selected Maize Single Crosses for Tolerance to Low P in Acidic Soils of Bumala and Maseno.

Eugene O. Olung'ati^{1*}, S. O. Gudu², E. Ouma², J. Ochuodgo¹ & O. K. Kiplagat¹

¹ University of Eldoret, P. O. Box 1125-30100 Eldoret- Kenya.

² Rongo University, P.O. Box 103-404040 Rongo, Kenya.

* Author for Correspondence ORCID ID: <https://orcid.org/0000-0001-6640-9376>; Email: ojukuolungati@yahoo.com.

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Generally, 13% of Kenya's arable land mass (7.5 million ha) is acidic and prone to poor phosphorus (P) availability and soil acidity. This results in crop yield losses due to the direct adverse effects Aluminium toxicity and P deficiency due to fixation of this element in the soil. The objective of this study was to develop and select P efficient maize single crosses developed by crossing as per North Carolina II mating design. Sixty maize genotypes, among them 34 single crosses were screened under acidic soils in Bumala and Maseno in a randomised complete block design. Sixty-seven percent of these single crosses were efficient, while 33% were inefficient. Two percent were efficient and responsive, 14% were inefficient but responsive, and the 79% were efficient but non-responsive. Generally, GY had a positive correlation with EH (0.45) and PH (0.61), while PH and EH had a positive correlation ($r = 0.86$) for the single crosses. The addition of P had significant effect on the grain yield, plant height, ear height and flowering of the genotypes at Bumala and Maseno. However, the effect of 26 kgP/ha was marginal at Maseno as compared to Bumala. The sites and genotypes varied significantly with regard to soil analysis and grain yield respectively, with the efficient and responsive genotypes selected for use in low input farming systems. Also, some of the efficient but non-responsive lines can also be selected for low input farming.

Keywords:

Maize,
Single Crosses,
Aluminium Toxicity,
P Efficiency

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INTRODUCTION

Globally, maize is considered as the third in importance after wheat and rice among consumed cereals (Ali et al., 2013; Onasanya et al., 2009). In Sub Saharan Africa and Kenya, the cereal is ascribed a unique importance because of its value as a staple food crop, an industrial raw material, and an animal feed (Magenya et al., 2008). Produced by both small- and large-scale farmers, maize provides Kenyans with a source for approximately 35.7% of consumed calories due to its staple status (Gichuru, 2013). However, its production is hampered by both biotic and abiotic constraints (pests, diseases, poor weather and soil conditions, poor seed quality, etc.). Soils that are categorised as being low in available P are however considered major challenge to maize productivity (Ouma et al., 2012). Also, according to Magenya et al (2008) and Gichuru (2013), P sorption in acid soils leads to phosphorous (P) deficiency, poor soil fertility, significantly lowers maize productivity. In Kenya, approximately 7.5 million ha of maize producing as well as agriculturally viable land is acidic (Kisinyo et al., 2014). In such soils, applied P becomes fixed due to acidity and concurrent Aluminium (Al) toxicity, thus making chemical amelioration less effective (Mumtaz et al., 2014). In western Kenya, soil acidity is a common occurrence and is commonly associated with Aluminium toxicity as well as P deficiency. Such areas are reported as expressing an available soil P of between 2-5 mg/kg, whereas the minimum threshold for maize production is set at 10 mg/kg (Brink & Bellay, 2006; Yang et al., 2013; Kisinyo et al., 2014; Ouma et al., 2012). This

situation has resulted in yield losses of between 20%-58%, while its accompanying Al toxicity has resulted in yield losses of between 16%-45% (Kisinyo, 2011; Ouma et al., 2012). Addressing the issue of low P availability therefore becomes vital in Western Kenya.

Phosphorus is among the major crop nutrients required for growth and development of crops, and deficiencies of this mineral leads to negative effects on the crop's development process and eventually reducing yield (Ward et al., 2011). As a major nutrient, this element is required in significantly larger amounts, similar to Nitrogen and Potassium. It is an important molecule in the Adenosine triphosphate molecular structure as well as in phospholipids and nucleic acids and is an essential requirement in photosynthesis (Obura, 2008). In addition to these roles, Phosphorus availability is important in minimizing Aluminium induced root damage, and ameliorating accumulation of the toxin in root tips cells (Sun et al., 2008). In terms of importance, the fully oxidised and inorganic form of this mineral element is considered as the most important form for plant use due to its function in the above-mentioned roles (Satyaprakash et al., 2017). In addition to soil acidity and Al toxicity, P availability is also influenced by lack of or presence of organic matter in soil as well as the continual use of acidic fertilisers (Sharma et al., 2013; Ware et al., 2007). Crops exposed to low P conditions express symptoms such as the progressive purpling of leaves from tips to the margins and eventually the whole leaf, necrosis of the stem, poor flowering and

seed fill, poor yields, and the complete elimination of young susceptible plants (Fageria, 2009).

Traditionally, P deficiency can be ameliorated by application of lime or organic and inorganic fertilisers (Kisinyo, 2011). However, in most P deficient soils, only 20% of applied P remains available for plant use and acquisition in a majority of soil ecosystems due to fixation of the remaining 80% (Balemi & Ngeshio 2012; Mumtaz et al., 2014). This then, due to the diverse nature of soil, leads to the development of P depleted pockets supplying between 10-15% of supplied P (Obura, 2008). A process that may eventually lead to lack of economic sustainability. In addition, liming or the use of other soil amendments, is considerably expensive for small scale farmers and it is therefore prudent to explore genetically conditioned tolerance (Kisinyo et al., 2014; Ouma et al., 2012). According to Kisinyo et al. (2014), maize genotypes with adequate phosphorus utilization potential (PUE), or those adopted to enhanced acquisition of P (PAE) become an indispensable tool in dealing with the constrain of low available P, especially for low-income small-scale farmers. This study therefore aimed at selecting P efficient single crosses developed by classical breeding.

MATERIALS

Study Sites

The study was carried out in three different locations; Rongo University, Maseno University, and Bumala. Rongo University farm was the crossing site for development of single cross hybrids, Maseno University farm and the farmer's field in Bumala were the sites for evaluation of the F1 hybrids. Rongo University is located at between 1300-1500 m above sea level, receives an average of 1600 mm of rainfall pa, and experiences a temperature of between 20-21.7 °C (Low et al., 1997; MoA, 2014). Bumala is located between 1135-1500 m above sea level, experiences a temperature of between 20.5-22.7 °C, receives an annual rainfall of approximately 900-1700 mm, and is reported as having acidic soil with ah pH of 4.5-4.6 and available soil P of 2.74 mg/kg (Ouma et al., 2012). Maseno is located approximately 1500 m above sea level, experiencing a rainfall average of 1750 mm pa, a temperature of 28.7 °C, and is has

acidic soil of pH 4.5-5.4 with an available soil P of 4.5 mg/kg (Gichimu et al., 2008). While soils in Bumala is classified as orthicferralsols (Ouma et al., 2012), Maseno soils are classified as dystricnitisols (Mwai et al., 2001).

Genetic Material

The germplasm used in the study was sourced from Rongo University and the University of Kwazulu Natal. From Rongo University was sourced the 14 Aluminium tolerant and P efficient maize genotypes, while the 9 Maize streak virus tolerant genotypes were sourced from the University of Kwazulu Natal. The material from the University of Kwazulu Natal were all inbred lines, while from Rongo University were a mixture of inbred lines, and populations from Brazilian introductions (Ouma et al., 2012).

METHODOLOGY

Development of Single Cross Hybrids

Crossing was done in Rongo, at the University collage field in the short rains of 2015, using the North Carolina II mating design with the Kenyan inbreds as female and South African inbreds as male. Pollen was transferred in pollen bags set up overnight from the tassel of the male plants to the silks of the female pants at 9 am in the morning to 10 am during the two-week flowering period. It resulted in 34 single cross genotypes for screening. The crossing block consisted of two plots staggered by a week and a half, each consisting of two plots (male and female), with each male and female having two rows for each of the genotypes.

Field Screening for Tolerance to Maize Streak Virus, Aluminium Toxic/ P Efficient

The field experiment was laid out in Randomised Completed Blocks with the genotypes and two P levels, P0, and P1, as treatments. Each treatment was replicated twice. A total of 34 single crosses, 2 repeats, 23 parietals, and 1 check were evaluated. Each plot had ten genotypes and each block has six plots, with each genotype represented in singular rows (3 m long, with 0.75 m inter-row spacing and 0.25 m intra-row spacing) (Gichuru, 2013; Ngwira & Khonje, 2005; Scott *et al.*, 2009).

The genotypes under screening were subjected to two rates of P, (P0) 0 kgP/ha and (P1) 26 Kg P/ha) while N was applied at 30 kg N/ha at planting and at knee height 45 Kg N/ Ha (NAFIS 2016; Gudu *et al.*, 2011; Gudu *et al.*, 2005). Due to its capacity to supply N at 18%, which is equivalent to 6.48 kg of N per 100 kg DAP, the treatment with -P (P0) was supplied slightly more CAN than +P (26 kgP/ha) to balance the 18% N supplied by DAP.

P-Deficiency to Lrance Assessment

P efficiency was evaluated on basis of grain yield (GY), plant height (PH), ear height (EH), days to 50% silking and 50% anthesis. Data was collected for all these traits for P efficiency assessment (Jiang *et al.*, 2010; Too *et al.*, 2014), with plant height and ear height being collected at plant maturity.

Model: $Y_{ijk} = \mu + b_i + P_j + G_k + PG_{jk} + \epsilon_{ijk}$

Where; μ -general mean, b- blocking effect, G- Genotype effect, P- Treatment effect, PG- treatment genotype interaction, ϵ -error term.

% Response to P application of P for the various inbred lines and single cross hybrids was calculated according to Ouma *et al.* (2012) as:

$$\%R = \frac{P1}{P0}$$

Data Analysis

The data for analysis was subjected to ANOVA in GENSTAT version 14 and mean separation done using DMRT test at 5% level of significance.

Soil Sampling

Soil sampling was done using systemic quadrates and the zigzag method (Midwest Laboratories, 2004; Okalebo *et al.*, 2002). The method used nested the zigzag systems within the quadrates. The sub-samples were mixed thoroughly and approximately 1.2 kg composite samples were packed in a black polythene bag and transported to the laboratory for air-drying, grinding, and sieving via a 2 mm sieve. The samples were then tested for pH using the HANNA soil analysis tool kit (Vanek, 2017), texture and organic carbon according to Okalebo *et al.* (2002), and available P according to Olsen (1972).

RESULTS

Soil Testing Results

Results of the soil analysis showed that the two sites were generally low in fertility, and had low pH and available P. Bumala had a pH 4.6, while Maseno had a pH was 5.2. These values indicated that the two sites have strongly acidic soils (*Table 1*), with Maseno having a slightly higher available P. The two sites also had low C level but Maseno had a slight advantage over Bumala for this trait as well.

Table 1: Soil analysis results for the experimental plots used in the study at Maseno and Bumala

| Site | Organic Carbon | P | pH | Textural Class | Soil Type |
|--------|----------------|-----|-----|-----------------|--------------------|
| Bumala | 2.66 | 3.3 | 4.6 | Sandy clay loam | Orthicferasols |
| Maseno | 3.18 | 4.8 | 5.2 | Clay loam | Dystric Nitrilsols |

Response in P deficient/ low P conditions in Acidic soils of Maseno and Bumala

The genotypes used in the study varied significantly in terms of PH, EH, GY, days to 50% silking and days to 50% tasselling ($p < 0.05$), with the single cross hybrids outperforming the parental inbred lines under both low P (0kgP/ha) and high P (26 kgP/ha) conditions. The interaction between the genotypes and sites was significant ($p < 0.05$) for

GY, PH, EH, days to 50% tasselling and days to 50% silking. The interaction between sites and P was significant ($p < 0.05$) for GY, PH, EH, and days to 50% tasselling but not for days to 50% silking, while the interaction between genotypes and P was significant ($p < 0.05$) for GY, PH, EH. Only PH and EH were significantly affected by the interaction between site, genotype, and P (*Table 2*).

Table 2: Mean square tale for maize genotypes tested for P efficiency under field condition in Bumala and Maseno

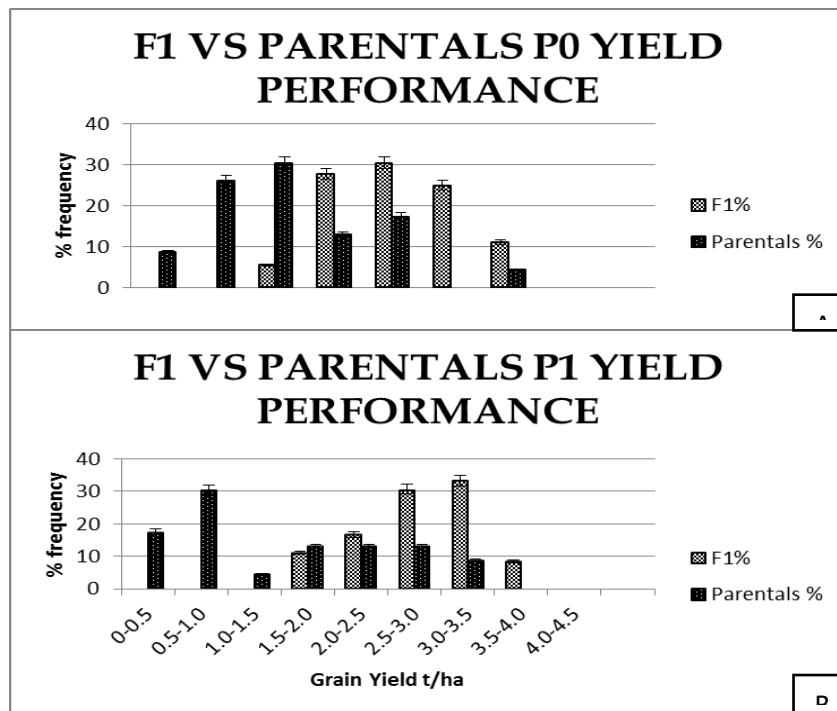
| Source of variation | df | GY | PH | EH | SILK | TASS |
|---------------------|-----|------------|-----------|-------------|------------|------------|
| Site | 1 | 109.712*** | 911121*** | 273537.4*** | 3.98 | 150.1** |
| Genotype | 59 | 5.5897*** | 305881*** | 24345.9*** | 244.31*** | 196.08*** |
| P | 1 | 16.3042*** | 157049*** | 85166.5*** | 1520.22*** | 1535.74*** |
| Site. Genotype | 59 | 1.7929*** | 326699*** | 3643.7*** | 65.19*** | 52.36*** |
| Site.P | 1 | 4.3532*** | 12386*** | 182261.1*** | 49.31 | 186.74** |
| Genotype.P | 59 | 1.0631*** | 5337*** | 2142.7*** | 30.77 | 32.62 |
| Site.Genotype.P | 59 | 0.5285 | 9449*** | 3679.4*** | 37.05 | 23.42 |
| Residual | 239 | 0.3874 | 1218 | 475.3 | 5.7 | 27.46 |
| Total | 478 | | | | | |
| Mean | | 2.119 | 148.03 | 50.07 | 73.54 | 68.42 |
| SE | | 0.6224 | 34.904 | 21.801 | 5.694 | 5.241 |
| CV | | 29.4 | 4.2 | 13.5 | 7.7 | 7.7 |

Note *, ** and *** indicates significance at $P \leq 0.05$, $P \leq 0.01$, $P \leq 0.001$ levels, respectively.

With regard to the performance of the single crosses across the sites in P0 5.6% expressed a GY of 1.0-1.49 t/ha, 27.8% expressed a GY of 1.5-2.0 t/ha, 30.6% expressed a GY of 2.0-2.49 t/ha, 25% expressed a GY of 2.5-3.0 t/ha, 11.1% expressed a GY of 3.0-3.49 t/ha. This performance was better than the parental lines that had 8.7% expressing 0-0.49 t/ha, 26.1% expressing 0.5-0.99 t/ha, 30.4% expressing 1.0-1.49 t/ha, 13.0% expressed a GY of

1.5-2.0 t/ha, 17.4% expressed a GY of 2.0-2.49 t/ha, 0% a GY of 2.5-3.0 t/ha, and 4.3% expressed a GY of 3.0-3.49 t/ha. On the other hand, all the single cross hybrids expressed a GY of greater than 1 t/ha under 26 kg/ha P, while only 47.82% of the parental inbred lines expressed the same grain yield under 26 kg/ha P, and majority of the parental inbred lines (52.18%) expressed a grain yield of below 1 t/ha under 26 kg/ha P (Figure 1).

Figure 1: Performance of F1 and Parental lines across P treatments; A (0kg P/ha), B (26kg P/ha)



The best single cross in Maseno under low P (P0) was 4BXSYNAL with a GY of 5.4 t/ha, and the worst was 13MAK-13BXKML036 with a GY of 0.94 t/ha. Under high P (P1), the best single cross in Maseno was 4BXSYNAL with a GY of 6.23 t/ha, and the worst was 13MAK-13BXKML036 with a GY of 1.90 t/ha. The best single cross in Bumala under low P was 203B-9X54B with a GY of 1.2 t/ha, while the worst was 203B-9X4B with a GY of 0.09 t/ha. Under high P the best single cross in Bumala was 1BXBRC with a GY of 2.1 t/ha and the worst was 9BXCON5 with a GY of 0.46 t/ha. Across the two sites, the best single cross under low P was 4BXSYNAL with a GY of 3.3 t/ha, while the worst single cross was 13MAK-13BXKML036 with a grain yield of 1.10. Under high P, the best single cross across the sites was 4BXSYNAL with a GY of 4.42 t/ha and the worst was 13MAK-13BXKML036 with a grain yield of 1.75 t/ha (Table

6). Comparatively, addition of P had a 21.1% incremental effect on the grain yield of the single cross F1 used in the study, with a 7.1% increase of grain yield in Maseno and a 29.1% increase of grain yield in Bumala due to addition of P. Also, the overall grain yield for the F1 hybrids in Maseno was 38.5% higher than in Bumala. In addition, the addition of P had a significant effect in improving ear height (EH) and plant height (PH), as well as reducing the days to 50% tasselling, and reducing the days to 50% silking. The single cross hybrids were further classified into different categories (tolerance to low P soils, and responsiveness to P) based on their grain yield at the different P levels. The classification resulted in the conclusion that 97.1% were efficient, 17.65% were responsive to P, and 14.71% were both P efficient and responsive (Table 6).

Table 3: Means for traits under low P and High P in acidic soils of Maseno and Bumala. Means across the same row with different letters differed significantly

| Trait Measured | P0 | P1 | MEAN | SED |
|------------------------|---------------------|--------------------|--------|--------|
| Grain yield | 0.421 ^a | 1.054 ^b | 0.737 | 0.369 |
| Plant height | 114.12 ^a | 151.6 ^b | 132.86 | 13.817 |
| Ear height | 28.31 ^a | 51.72 ^b | 40.01 | 8.118 |
| Days to 50% tasselling | 69.82 ^a | 65.14 ^b | 67.48 | 3.98 |
| Days to 50% silking | 75.56 ^a | 71.31 ^a | 73.43 | 4.644 |

Table 4: Means for traits across sites (Maseno and Bumala). Means across the same row with different letters differed significantly

| | Maseno | Bumala | MEAN | SED |
|------------------------|---------------------|---------------------|--------|--------|
| GY | 2.597 ^a | 1.641 ^b | 2.119 | 0.6224 |
| PH | 165.81 ^a | 130.24 ^b | 148.03 | 34.904 |
| EH | 59.82 ^a | 40.32 ^b | 50.07 | 21.801 |
| Days to 50% tasselling | 68.98 ^a | 67.86 ^a | 68.42 | 3.98 |
| Days to 50% silking | 73.45 ^a | 73.63 ^a | 73.54 | 4.644 |

Phenotypic Relationships among Maize Performance Indicator Traits Across Sites

GY had a strong and positive correlation with both EH ($r = 0.82$) and PH ($r = 0.80$) for the all genotypes in the study. Also, PH and EH had strong positive correlation ($r = 0.82$) for all the genotypes in the study. For the inbred lines, correlation between GY and EH, GY and PH, and EH and PH was strong and positive at ($r = 0.87$), ($r = 0.93$), and

($r = 0.96$) respectively. For the single cross hybrids, correlation between PH and EH was positive and strong ($r = 0.86$), correlation between PH and GY was moderate and positive (0.61), while correlation between EH and GY was weak and positive (0.45). For all the genotypes in the study, GY had a positive and insignificant correlation with days to 50% tasselling ($r = 0.01$) and with days to 50% silking ($r = 0.09$), this was similar to the F1, where GY had a positive yet insignificant correlation with days to 50% tasselling ($r = -0.06$) and with days to

50% silking (0.17). For the inbred lines correlation and GY and days to 50% silking ($r = 0.50$) was moderate and positive (Table 4).

Table 5: Correlation between GY, EH, PH and days to 50% flowering of all genotypes tested for P-efficiency in Maseno and Bumala.

| | Grain Yield | Plant height | Ear Height | Days to 50% Tasseling | Days to 50% Silking |
|------------------------|-------------|--------------|------------|-----------------------|---------------------|
| Grain Yield | 1 | | | | |
| Plant height | 0.8** | 1 | | | |
| Ear Height | 0.82** | 0.82** | 1 | | |
| Days to 50% Tasselling | 0.01 | 0.15 | 0.25* | 1 | |
| Days to 50% Silking | 0.09 | 0.3 | 0.39* | 0.92** | 1 |

* and **; Significance at 5 and 1%, respectively.

Table 6: Table with Selected F1s for grouped for P efficiency and response to P based on Grain yield means.

| GENOTYPE | Grain Yield | | Response to P | |
|-------------|-------------|------|---------------|------------|
| | P0 | P1 | P1/P0 (R) | Categories |
| 4BXSYNAL | 3.34 | 4.42 | 1.33 | EN |
| 54BXBRC | 3.20 | 2.88 | 0.90 | EN |
| 9BXSYNAL | 3.18 | 3.44 | 1.08 | EN |
| 203B-9X54B | 3.14 | 3.01 | 0.96 | EN |
| 44BX203B-9 | 2.89 | 3.06 | 1.06 | EN |
| 44BX203B-14 | 2.85 | 2.73 | 0.96 | EN |
| 41BX203B | 2.69 | 2.36 | 0.88 | EN |
| 4BX203B-1 | 2.65 | 2.28 | 0.86 | EN |
| 41BXCON5 | 2.31 | 2.74 | 1.19 | EN |
| 13BX203B-14 | 2.17 | 3.67 | 1.69 | ER |
| 203B-9X4B | 2.13 | 3.24 | 1.52 | ER |
| 41BXBRC | 1.59 | 3.76 | 2.36 | IR |
| 1BXBRC | 1.57 | 3.54 | 2.25 | IR |
| 1BXAO809 | 1.53 | 1.96 | 1.28 | IN |
| 13BXKML036 | 1.10 | 1.75 | 1.59 | IR |

KEY: I, Inefficient; R, Responsive, E, efficient; N, non-responsive; IR, inefficient and responsive; ER, efficient and responsive; EN, efficient and nonresponsive; IN, inefficient and nonresponsive.

DISCUSSION

Variation in agronomic traits due to P addition, genotypic differences, and site.

The genotypes and P treatments had a significant effect on the plant height, ear height, grain yield, and days to flowering of the genotypes in the study. At Bumala the addition of P led to an overall increase in grain yield by 29.1%, plant height by 24.7% and ear height by 45.8% but reduced days to 50% tasselling by 7.2% and days to 50% silking by

5.6%. In Maseno, grain yield was increased by 7.1%, plant height by 0.42% and ear height by 8.8%, days to 50% tasselling and days to 50% silking reduced by 3.4% and 4.1% respectively due to the addition of P. This effect of P in improving agronomic crop traits identified in this study relates well with other studies. Fosu-Mensah & Mensah (2016), Ouma *et al.* (2013), and Umeri *et al.* (2016) reported that the addition of P resulted in a reduction of days to flowering (tasselling and silking), as well as an increase in grain yield. According to Ouma *et al.* (2013), P addition at 26 kgP/ha resulted in a grain yield increase by a margin of 73.5%, as well as a

corresponding increase in ear and plant height. Besides these findings, Lokhande *et al* (2015) concluded that there was an increase in wet and dry biomass, as well as increased plant height for Coriander due to application of Pat 45 kgP/ha. Also, Alias *et al* (2003) concluded a 15% increase in plant height and a 53.76% increase in grain yield due to P supply at 150 kgP/ha for maize. On the other hand, Dai *et al* (2013) reported a yield loss of 560 kg/ha/year for maize due to lack of P fertiliser thus indicating the mineral's importance to improve grain yields for the crop.

In addition, the different genotypes expressed varied grain yield response due to P addition. This could have been due to inherent genetic differences in acquiring P from the soil. These results are similar to those reported by Fosu-Mensah & Mensah (2016) as well as Ouma *et al* (2013) and Umeri *et al* (2016), who reported that genetic differences among genotypes contributed to variance in acquisition and utilization of soil P. Although the grain yield at Maseno was better than Bumala, the effect of P addition was marginal at Maseno as compared to Bumala. This difference may be due to the higher soil P, variation in response to added P at 26 kgP/ha at the two sites, and higher soil pH at Maseno as compared to Bumala. These results are comparable to those of Kihara *et al*. (2016) who reported variation in the effect of NPK (Nitrogen, Phosphorous, Potassium) fertiliser on grain yield across sites. The soils in these two regions also expressed a variation in organic carbon content with Maseno having a slightly higher level. High organic carbon is known to increase the soil Cation-exchange capacity (CEC) which affects retention of cation nutrients, as well as improving water infiltration and retention (Noellemeyer & Six, 2015). In addition to the soil issues, the variation in rain fall received by these two regions may also have affected overall grain yield.

The reported improvements in plant grain yield and yield components due to the addition of P can be ascribed to its role in the development of plant roots as these are the main water and mineral absorption organs. In addition, P is essential in photosynthesis, cell division and elongation, and being synergistic to Nitrogen absorption (Ouma *et al.*, 2013; Salehi & Anampanah, 2015). Despite the improved yields due to P addition, the overall yields of the genotypes

in the study at Bumala neither met the 3 t/ha threshold set by Gudu *et al.* (2011), nor did they meet 3.41-8.7 t/ha threshold set by Ouma *et al.* (2012). These results can be attributed to the prolonged drought at Bumala during the experimental season that also interacted with late flowering for some of the genotypes. According to Halindu *et al.* (2015), prolonged drought can cause yield losses of between 50-100% depending on the length of the scourge. At the study site in Bumala, nearby fields and the farmer's field had visible effects of the drought with losses of up to 100% for some of the neighbouring farms.

Phenotypic relationships among maize performance indicator traits across sites

PH and EH had a positive and significant correlation to each other as well as to GY. These results are in agreement with those of Appiah *et al.* (2014). Such positive correlation presents that these yield components are associated positively with GY and can be used effectively in selection purposes for GY (Akongwubel *et al.*, 2012). Between grain yield and flowering the general correlation was positive but weak and insignificant for all the genotypes in the study and for the F1s. These findings are similar to those reported by Yousuf and Saleem (2001). However, the correlation between grain yield and flowering dates was positive and moderate for parental inbred lines, where GY and days to 50% tasselling ($r = 0.41$), and GY and 50/5 silking ($r = 0.50$).

CONCLUSION

The study identified 23 P efficient single crosses and 11 inefficient single crosses. Of the efficient single crosses, only two were responsive to P addition, while of the inefficient three were responsive to P addition. These genotypes can therefore be utilized in further development of three way and top crosses for P efficiency as well as gene pyramiding to develop multiple tolerant maize genotypes. Also, genotypic variability for P efficiency and response to P was identified among the maize genotypes in study and was further enhance by soil fertility variation among the sites.

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