

## Putative mechanisms of drought tolerance in maize (*Zea mays* L.) via root system architecture traits

Formatted: Space Before: 0 pt, After: 0 pt

Formatted: Space After: 0 pt

### ABSTRACT

Formatted: Indent: Left: 0"

Identifying maize genotypes with favorable root architecture traits for drought tolerance is prerequisite for initiating a successful breeding program for developing high yielding and drought tolerant varieties of maize. The objectives of the present investigation were: (i) to identify drought tolerant genotypes of maize at flowering and grain filling, (ii) to elucidate the relationships between the drought tolerance and root architecture traits and (iii) to identify the putative mechanisms of drought tolerance *via* root system traits. A two-year experiment was carried out using a split plot experiment with three replications. The main plots were devoted to ~~three~~ irrigation regimes, *i.e.* well watering (WW), water stress at flowering (WSF) and at grain filling (WSG), and sub plots to 22 maize cultivars and populations. Drought tolerance index (DTI) had strong and positive associations with crown root length (CRL), root circumference (RC) and root dry weight (DRW) under both WSF and WSG, a negative correlation with brace root whorls (BW), and positive correlations with crown root number (CN) under WSF and brace root branching (BB) and crown root branching (CB) under WSG. These root traits could be considered as putative mechanisms of drought tolerance. The cultivars Pioneer-3444, SC-128, Egaseed-77, SC-10 and TWC-324 showed the most drought tolerant and the highest yielding in a descending order; each had a number of such drought tolerance mechanisms. Further investigation should be conducted to determine the underlying root mechanisms contributing to the selection of water-efficient hybrids of maize.

**Key words:** Corn, Crown and Brace roots, Correlations, Drought tolerance index.

Formatted: Indent: Left: 0", Space Before: 0 pt

### INTRODUCTION

Formatted: Space Before: 0 pt

Maize (*Zea mays* L.) in Egypt is mainly used for poultry industry and animal feed. For acreage and production, it ranks second to wheat among cereal crops in Egypt. It is grown as a summer season crop and well irrigated by water coming from Nile River and its branches and canals. Current maize hybrids cultivated in Egypt are selected under well irrigation and therefore are subject to yield losses when grown under water deficit. The amount of water available for irrigation is reducing, especially at the ends of canals and due to expanding maize cultivation into the deserts, where sandy soils are of low water holding capacity. In order to stabilize maize production in Egypt, there is a need to develop drought tolerant maize hybrids.

Maize is very sensitive to water stress during the flowering and grain-filling periods (Bai *et al.* 2006) [1]. However, Witt *et al.* (2012) [2] reported that the most critical period for yield production goes approximately from 2 weeks before flowering time until 2 weeks after flowering time. Developing maize varieties that are tolerant to drought is, therefore considered

37 critical for increasing the maize production. Several investigations have been undertaken across  
38 the years to improve drought tolerance in breeding programs. Edmeades *et al.* (1993) [3]  
39 reported that germplasm developed from drought tolerant source populations performed  
40 significantly better under drought stress compared to conventional populations.

41 Root system architecture traits are important for plant productivity under drought stress  
42 (Lynch1995) [4]. Plants avoid dehydration by increasing water uptake in the soil profile and  
43 adapt to the chemical and physical soil constraints, particularly under drought conditions, *via* the  
44 morphological plasticity of their root system (Lynch 2007) [5]. The importance of a deep and  
45 vigorous root system for maintaining yield under drought stress has been reported in maize by  
46 Hund *et al.* (2011) [6]. Rauf and Sadaqat (2008) [7] stated that "drought tolerant genotypes  
47 generally increase the photosynthates allocation for root elongation under drought stress". Rauf  
48 *et al.* (2009) [8] reported that genetic variation for root elongation has been shown in maize. The  
49 effects of root architecture and size on maize yield also depend on the distribution of soil  
50 moisture and the competition for water resources within the plant community (King *et al.* 2009)  
51 [9].

52 Trait interrelationships in particular determine the degree of association among traits  
53 and how they may increase selection efficiency. It is useful if indirect selection for root traits  
54 gives greater response to selection for grain yield trait than direct selection for the same trait. The  
55 main criterion for drought tolerance selection is the association of each root trait with grain yield  
56 under stress conditions [9, 10] (King *et al.* 2009 and Trachsel *et al.*, 2011).

57 To start a successful breeding program for improving drought tolerance, available  
58 maize germplasm should be screened for related traits to drought tolerance; e.g. root architecture  
59 traits under deficit irrigation to identify the best ones for further use in extracting the best  
60 parental inbred lines for developing drought tolerant hybrids. The objectives of the present  
61 investigation were to: (i) characterize 22 maize genotypes for root architecture traits and  
62 tolerance to deficit irrigation at flowering and grain filling stages in order to identify drought  
63 tolerant ones, (ii) elucidate the relationships between the drought tolerance and root traits and  
64 (iii) identify the putative mechanisms of drought tolerance *via* root system architecture.

## 65 MATERIALS AND METHODS

66 This study was carried out in the two successive growing seasons 2016 and 2017 at the  
67 Agricultural Experiment and Research Station of the Faculty of Agriculture, Cairo University,

Formatted: Right: 0", Space Before: 0 pt

Comment [f1]: standardize

Formatted: Space After: 0 pt

68 Giza, Egypt (30° 02'N latitude and 31° 13'E longitude with an altitude of 22.50 meters above sea  
69 level).

70 **Plant materials**

71 Twenty two maize (*Zea mays* L.) genotypes were used, namely 15 Egyptian cultivars (10 single  
72 crosses and 5 three-way crosses) and 7 open-pollinated populations (Table 1). These materials  
73 were kindly provided by Hi-Tec Company (Hi-Tec-2031, Hi Tec-2066, Hi Tec 1100), DuPont  
74 Pioneer Company (P-30K09, P-3444, P-32D99), Fine Seeds Company (Fine-1005), Egaseed  
75 Company (Egaseed-77), Wataniya Company (Watania 11) and Agricultural Research Center-  
76 Egypt (the rest of genotypes). These genotypes were chosen to represent the available germplasm  
77 in Egypt and some of them could be considered sources for extracting drought tolerant inbred  
78 lines.

80 **Table 1.**— Designation, origin and grain color of studied maize genotypes.

Genotype No.	Designation	Origin	Genetic nature	Grain colour
1	Hi-Tec-2031	Hi-Tec, Egypt	Single cross	White
2	P-30K09	DuPont Pioneer	Single cross	White
3	Fine 1005	Fine Seeds, Egypt	Single cross	White
4	Egaseed-77	Egaseed Co., Egypt	Single cross	White
5	SC-10	ARC, Egypt	Single cross	White
6	SC-128	ARC, Egypt	Single cross	White
7	Hi-Tec- 2066	Hi-Tec, Egypt	Single cross	Yellow
8	P-3444	DuPont Pioneer	Single cross	Yellow
9	SC-166	ARC, Egypt	Single cross	Yellow
10	P-32D99	DuPont Pioneer	Single cross	Yellow
11	Hi-Tec 1100	Hi-Tec, Egypt	3-way cross	White
12	Watania 11	Watania Co., Egypt	3-way cross	White
13	TWC-324	ARC, Egypt	3-way cross	White
14	TWC-360	ARC, Egypt	3-way cross	Yellow
15	TWC-352	ARC, Egypt	3-way cross	Yellow
16	Giza Baladi	ARC, Egypt	Population	White
17	Population-45	ARC, Egypt	Population	Yellow
18	Nubaria	ARC, Egypt	Population	Yellow
19	Nebraska Midland	USA	Composite	Yellow
20	Midland_Cunningham	Eldorado,Kansas, USA	Population	Yellow
21	Golden Republic	Beltsville,Kansas, USA	Population	Yellow
22	Sweepstakes 5303	USA	Population	Yellow

81 ARC = Agricultural Research Center, SC = Single cross, TWC = Three-way cross

82 |

### 83 **Experimental procedures**

84 Sowing date was April 24<sup>th</sup> in the 1<sup>st</sup> season (2016) and April 30<sup>ht</sup> in the 2<sup>nd</sup> season (2017).  
85 Sowing was done in rows; each row was 4 m long and 0.7 m width. Seeds were over sown in  
86 hills 25 cm apart, thereafter (after 21 days from planting and before the first irrigation) were  
87 thinned to one plant/hill to achieve a plant density of 24,000 plants/fed. Each experimental plot  
88 included two rows (plot size = 5.6 m<sup>2</sup>).

### 89 **Experimental design**

90 A split-plot design in randomized complete block (RCB) arrangement with three replications  
91 was used. Main plots were allotted to three irrigation regimes, *i.e.* well watering (WW), water  
92 stress at flowering (WSF) and water stress at grain filling (WSG). Each main plot was  
93 surrounded with an alley (4m width), to avoid water leaching between plots. Sub plots were  
94 devoted to twenty-two maize genotypes.

### 95 **Water regimes**

96 **1. Well watering (WW):** Irrigation was applied by flooding, the second irrigation was given  
97 after three weeks and subsequent irrigations were applied every 12 days.

98 **2. Water stress flowering (WSF):** The irrigation regime was just like well watering, but the 4<sup>th</sup>  
99 and 5<sup>th</sup> irrigations were withheld, resulting in 24 days water stress just before and during  
100 flowering stage.

101 **3. Water stress grain filling (WSG):** The irrigation regime was just like well watering, but the  
102 6<sup>th</sup> and 7<sup>th</sup> irrigations were withheld, resulting in 24 days water stress during grain filling stage.

### 103 **Agricultural practices**

104 All other agricultural practices were followed according to the recommendations of ARC, Egypt.  
105 Nitrogen fertilization at the rate of 120 kg N/fed was added in two equal doses of Urea 46 %  
106 before the first and second irrigation. Triple Superphosphate Fertilizer (46% P<sub>2</sub>O<sub>5</sub>) at the rate of  
107 30 kg P<sub>2</sub>O<sub>5</sub>/fed, was added as soil application before sowing during preparation of the soil for  
108 planting. Weed control was performed chemically with Stomp herbicide just after sowing and  
109 before the planting irrigation and manually by hoeing twice, the first before the first irrigation  
110 (after 21 days from sowing) and the second before the second irrigation (after 33 days from  
111 sowing). Pest control was performed when required by spraying plants with Lannate (Methomyl)  
112 90% (manufactured by DuPont, USA) against corn borers.

113 **Soil analysis**

114 Physical and chemical soil analyses of the field experiments were performed at laboratories of  
115 Soil and Water Research Institute of ARC, Egypt. Across the two seasons, soil type was clay  
116 loam: Silt (36.4%), clay (35.3%), fine sand (22.8%) and coarse sand (5.5%), pH (7.92), EC (1.66  
117  $\text{dSm}^{-1}$ ), SP (62.5),  $\text{CaCO}_3$  (7.7 %), Soil bulk density ( $1.2 \text{ g cm}^{-3}$ ),  $\text{HCO}_3$  (0.71 mEq/l), Cl (13.37  
118 mEq/l),  $\text{SO}_4$  (0.92mEq/l),  $\text{Ca}^{++}$  (4.7mEq/l),  $\text{Mg}^{++}$  (2.2mEq/l),  $\text{Na}^+$  (8.0mEq/l),  $\text{K}^+$   
119 (0.1mEq/l), N, P, K, Zn, Mn and Fe (371, 0.4, 398, 4.34, 9.08 and 10.14 mg/kg, respectively).

120 **Data recorded:**

121 **Grain yield plant<sup>-1</sup> (GYPP) (g):** It was estimated by dividing the grain yield plot<sup>-1</sup> (adjusted at 15.5% grain  
122 moisture) on number of plants plot<sup>-1</sup> at harvest.

123 **Grain yield ha<sup>-1</sup> (GYPH) (ton):** It was estimated by adjusting grain yield plot<sup>-1</sup> at 15.5% grain  
124 moisture to grain yield ha<sup>-1</sup> (ton).

125 **Root traits:**

126 At the end of each water stress treatment (80 and 100 days from emergence for WSF and  
127 WSG, respectively) and just after irrigation, three plant roots from each experimental plot were  
128 excavated by removing a soil cylinder of 40 cm diameter and a depth of 40 cm with plant base as  
129 the horizontal centre of the soil cylinder. Excavation was carried out using standard shovels. The  
130 excavated root crowns were shaken briefly to remove a large fraction of the soil adhering to the  
131 root crown. Most of the remaining soil was then removed by soaking the root crown in running  
132 water. In a third step, remaining soil particles were removed from the root crown by vigorous  
133 rinsing at low pressure. The clean roots were measured or visually scored (Fig. 1) for the  
134 following traits:

135 **Number of above-ground whorls occupied with brace roots (BW).**

136 **Number of brace roots (BN).**

137 **Angle of 1<sup>st</sup> arm of the brace roots originating from whorl 1 (BA) (score).**

138 **Branching density of brace roots (BB) (score).**

139 **Number of crown roots (CN) (score).**

140 **Crown roots angle (CA) (score).**

141 **Branching density of crown roots (CB) (score).**

142 Traits from No. 5 to No. 9 were assigned values from one to nine according to Trachsel *et*  
143 *al.* (2011) [10], where one indicates shallow root angles ( $10^\circ$ ), low root numbers and a

Formatted: Space After: 0 pt

Formatted: Indent: Left: -0.25"

Formatted: Indent: Left: -0.25"

144 low branching density and nine indicates steep root angles (90°), high numbers and a  
145 high branching density (Fig.1).

146 **10. Crown root length (CRL) (cm).** The root length, measured as the distance between the last  
147 node to the end tip of the root.

148 **11. Root circumference (RC) (cm).** RC was measured from maximum root system width.

149 **12. Root (crown and brace) dry weight (RDW) (g).** The measured root was first spread out in the  
150 sun for partial drying and then put in an oven for total drying at 40°C for 24 hours. After drying  
151 the roots were weighed using an electronic scale.

#### 152 **Drought tolerance index (DTI):**

153 Drought tolerance index is the factor used to differentiate between the genotypes from tolerance  
154 point of view and it is calculated by the equation of Fageria (1992) [11] as follows:

$$155 \text{DTI} = (Y1/AY1) \times (Y2/AY2)$$

156 Where, Y1 = trait mean of a genotype at well watering. AY1 = average trait of all genotypes at  
157 well watering. Y2 = trait mean of a genotype at water stress. AY2 = average trait of all  
158 genotypes at water stress. When DTI is  $\geq 1$ , it indicates that genotype is tolerant (T) to drought.  
159 If DTI is  $< 1$ , it indicates that genotype is sensitive (S) to drought.

#### 160 **Biometrical analyses**

161 Analysis of variance of the split-split plot design in RCB arrangement was performed on the  
162 basis of individual plot observation using the MIXED procedure of MSTAT ®. Combined  
163 analysis of variance across the two growing seasons was also performed if the homogeneity test  
164 was non-significant. Moreover, combined analysis for each environment separately across  
165 seasons was performed as randomized complete block design. Least significant difference (LSD)  
166 values were calculated to test the significance of differences between means according to Steel *et*  
167 *al.* (1997) [12].

Formatted: Right: 0"

Formatted: Indent: Left: 0", Right: 0"

Formatted: Indent: Left: -0.25"

Formatted: Space After: 0 pt



168  
169  
170  
171  
172

**Figure 1.** Images of brace roots angle (BA), brace roots branching density (BB), crown roots number (CN), crown roots angle (CA) and crown roots branching (CB) displayed were scored with 1, 3, 5, 7 and 9.

173  
174  
175  
176

Simple correlation coefficients were calculated between pairs of studied traits under well watering (WW), water stress (WS), severe water stress (SWS) and combined across all irrigation treatments according to Singh and Narayanan (2000) [13]. Spearman's rank correlation coefficients calculated among studied root traits and other studied traits under studied environments. It was

Formatted: Indent: Left: 0", First line: 0"  
Formatted: Font: Not Bold  
Formatted: Indent: Left: -0.5", Line spacing: 1.5 lines  
Formatted: Space After: 0 pt

177 computed by using SPSS 17 computer software and the significance of the rank correlation  
 178 coefficient was tested according to Steel *et al.* (1997) [12].

## 180 RESULTS AND DISCUSSION

### 181 3.1. Analysis of variance

182 Combined analysis of variance across seasons (S) of the split-split plot design (Table 2)  
 183 indicated that mean squares due to seasons were significant ( $P \leq 0.05$  or  $P \leq 0.01$ ) for six out of  
 184 studied 12 traits, namely brace root whorls (BW), brace root angle (BA), crown root angle (CA),  
 185 crown root branching (CB), grain yield/plant and grain yield/ha. Mean squares due to irrigation  
 186 regime were significant ( $P \leq 0.05$  or  $P \leq 0.01$ ) for six out of studied 12 traits, namely crown root  
 187 number (CN), CB, root circumference (RC) and root dry weight (RDW), GYPP and GYPH.  
 188 Mean squares due to genotype were significant ( $P \leq 0.01$ ) for all studied root and grain yield  
 189 traits.

192 **Table -2:** Mean squares from combined analysis of variance across 2016 and 2017 years for studied root  
 193 traits of 22 maize genotypes under four irrigation regimes.

Variance Source	Mean Squares						
	BW	BN	BA	BB	CN	CA	
Season (S)	5.32*	487.8	33.5**	5.5	0.4	103.2**	
Irrigation regime(I)	2.78	2139.6**	3.2	12.9	32.5*	5.4	
I x S	4.9*	615.6	3.3	15.1	4.3	10.4	
Genotype (G)	2.91**	1014.5**	6.1**	16.6**	12.3**	9**	
G x S	0.218	85.9	2.2	10.8**	4*	1.7	
G x I	0.449	146.8	1.5	3.7	2.5	1.6	
G x S x I	0.362	122.6	1.2	5.2*	2.3	1.1	
	CB	CRL	RC	RDW	GYPP	GYPH	
Season (S)	28.2**	243.5	107.5	94.5	26041.5*	124.7**	
Irrigation regime(I)	26**	115.7	618.1**	1336.5**	47158.4**	2041.1**	
I x S	3.8	201.9	232.9*	1278.1**	3864.3	225.5**	
Genotype (G)	13.1**	59.4**	263.2**	955.5**	12428.3**	707.3**	
G x S	4.7**	13.6	26.9	234.1**	3439.6**	46.4**	
G x I	2.5	17.2	26.7	132.9	1335.8**	34.8**	
G x S x I	1.8	23.1	32.2	142.4	1383.5**	19.6**	

193 BW= Number of above-ground whorls occupied with brace roots, BN= Number of brace roots, BA= Brace root  
 194 angle, BB= Branching density of brace roots, CN= Number of crown roots, CA=Crown roots angle, CB=Branching

Formatted: Indent: Left: -0.49"

Formatted: Font: Not Bold

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt



195 density of crown roots, CRL= Crown root length, RC=Root circumference, RDW= Roots dry weight, GYPP= Grain  
196 yield/plant, GYPH= grain yield/ha, \* and \*\* indicate significance at 0.05 and 0.01 probability levels, respectively.

197 |  
198 Mean squares due to the 1<sup>st</sup> order interaction were significant ( $P \leq 0.05$  or 0.01) for four  
199 traits (BN, RC, RDW and GYPH) due to I×S, for six traits (BB, CN, CB, RDW, GYPP and  
200 GYPH) due to G×S and two traits (GYPP and GYPH) due to G× I. Mean squares due to the 2<sup>nd</sup>  
201 | order interaction, *i.e.* G×S× I, were significant ( $P \leq 0.01$ )—for three traits, namely BB, GYPP and  
202 GYPH (Table 2).

203 Combined analysis of variance of a randomized complete blocks design (RCBD) (data  
204 not presented) under four environments, *i.e.* well watering at flowering (WWF), well watering at  
205 grain filling (WWG), water stress at flowering (WSF) and water stress at grain filling (WSG)  
206 across two seasons indicated that mean squares due to genotypes under all environments were  
207 significant ( $P \leq 0.05$  or 0.01) for 35 out of 46 studied cases (76.1%).

208 | Root system architecture is important for plant productivity under drought stress\*  
209 conditions [4] (Lynch, 1995). In order to improve plant performance, breeders need to select  
210 genotypes with a root architecture adapted to the conditions of the target environment. Results of  
211 the present study indicated that climatic conditions had a significant effect on BW, BA, CA, CB,  
212 GYPP and GYPH and that irrigation regime had a significant effect on CN, CB, RC, RDW,  
213 GYPP and GYPH. Moreover, genotype had an obvious effect on all studied traits. The role of  
214 maize genotype is in accordance with the finding of Trachsel et al. [10] (2011) for maize root  
215 traits and Al-Naggar et al. (2016a) [14, 15] for grain yield. Mean squares due to the the 1<sup>st</sup> and  
216 2<sup>nd</sup> order interaction were significant for some root and yield traits, indicating that for such traits,  
217 the rank of maize genotypes differ from irrigation regime to another, and from one year to  
218 another and the possibility of selection for improved root and grain yield under a specific water  
219 stressed environment as proposed by Al-Naggar et al. (2009, 2011, 2016 b, 2017 a,b) [16-20].  
220 Combined analysis of variance of RCBD under each of the four environments indicated the  
221 significance of differences among studied genotypes for the majority of studied root traits and  
222 grain yield under each irrigation regime.

### 223 3.3. The effect of genotype

224 Average, minimum and maximum values of all studied traits of 22 genotypes across all  
225 irrigation treatments combined across two seasons are presented in Table (3).

226 |

Formatted: Space After: 0 pt

Formatted: Indent: First line: 0"

227 **Table 3:** Average, minimum (Min) and maximum (Max) values of all studied traits of each genotype  
 228 combined across all irrigation regimes and across 2016 and 2017 seasons.

Formatted: Font: Not Bold

Parameter	Traits					
	BW (No.)	BN (No.)	BA (score)	BB (score)	CN (score)	CA (score)
Average	2.5	37.1	6.7	4.9	3.2	6.7
Min	1.9 (8)	25.6 (21)	5.5 (1)	3.4 (18)	1.9 (21)	5.6 (7)
Max	3.0 (10,11,17)	49.0(10)	7.7(19)	6.2(9)	4.5(6)	8.1(10)
LSD <sub>.05</sub>	0.36	6.8	0.74	1.09	0.86	0.76
	CB (score)	CRL (cm)	RC (cm)	RDW (g)	GYPP (g)	GYPH (ton)
Average	4.2	22.8	32.7	22.3	107.3	7.18
Min	3.0 (21)	20.4 (18)	25.9 (21)	11.2 (20)	62.5(22)	2.69(22)
Max	6.5 (8)	26.1 (5)	38.1 (8)	36.8(8)	158.5(6)	13.03(8)
LSD <sub>.05</sub>	0.91	2.57	2.85	6.05	9.72	0.39

229 Means of minimum and maximum are followed by genotype No. (Between brackets). BW= Number of  
 230 aboveground whorls occupied with brace roots, BN= Number of brace roots, BA= Brace root angle, BB= Branching  
 231 density of brace roots, CN= Number of crown roots, CA=Crown roots angle, CB=Branching density of crown roots,  
 232 CRL= Crown root length, RC=Root circumference, RDW= Roots dry weight, GYPP= Grain yield/plant, GYPH=  
 233 grain yield/ha.

Formatted: Space After: 0 pt

234  
 235 Genotypes varied for grain yield/fed from 13.03 ton (genotype No. 8) to 2.69 ton  
 236 (genotype No. 22), grain yield/plant from 158.5 g (genotype No. 6) to 62.5 g (genotype No. 22),  
 237 number of above-ground whorls occupied with brace roots from 3.0 from (genotype No. 17) to  
 238 1.9 (genotype No. 8), number of brace roots from 49.0 (genotype No. 10) to 25.6 (genotype  
 239 No. 21), angle of 1<sup>st</sup> arm of the brace roots originating from whorl 1 from 7.7 (genotype No. 19)  
 240 to 5.5 (genotype No. 1), branching density of brace roots from 6.2 (genotype No. 9) to 3.4  
 241 (genotype No. 18), number of crown roots from 4.5 (genotype No. 6) to 1.9 (genotype No.  
 242 21), crown roots angle from 8.1 (genotype No. 10) to 5.6 (genotype No. 7), branching density  
 243 of crown roots from 6.5 (genotype No. 8) to 3.0 (genotype No. 21), crown root length from  
 244 26.1 cm (genotype No. 5) to 20.4 cm (genotype No. 18), root circumference from 38.1 cm  
 245 (genotype No. 7) to 25.9 cm (genotype No. 21) and roots dry weight from 36.8 g (genotype  
 246 No. 8) to 11.2 g (genotype No. 20).

247 The genotype No. 8 (Pioneer-3444) exhibited the highest mean values for four traits  
248 [GYPH, root circumference (RC), crown root branching (CB) and roots dry weight (RDW)] and  
249 second highest for GYPP, brace root branching (BB), number of crown roots (CN), crown root  
250 length (CRL), *i.e.* most important yield and root traits. The genotype No. 6 (SC-128) developed  
251 by ARC-Egypt was the highest in GYPP and number of crown roots and second highest in crown  
252 root branching. The genotype No. 4 (Egaseed 77) developed by Fine Seed Co. showed the third  
253 highest in grain yield and the highest in brace root angle (BA). The genotype No. 5 (SC-10)  
254 developed by ARC-Egypt showed the highest means for one trait (crown root length; CRL); it  
255 gave the fourth highest grain yield per plant and per hectare.

256 On the contrary, the genotype No. 22 (Pop. Sweepstakes 5303) exhibited the lowest  
257 means for two traits, namely GYPP, GYPH. The genotype No. 21 (Pop. Golden Republic)  
258 exhibited the lowest means for two traits, namely BN and CN. The genotype No. 18 (Pop.  
259 Nubaria) showed the lowest means for two traits (BB and CRL).

260 Means of the 22 maize genotypes showed wide ranges of performance (difference  
261 between minimum and maximum values) for all studied root and yield traits across all irrigation  
262 treatments. Three commercial varieties showing the highest grain yield showed also the highest  
263 means for a number of root traits. The superiority of these three commercial varieties in six root  
264 traits (RC, CB, RDW, BB, CN and CRL) for Pioneer-3444, two traits (CN and CB) for SC-128,  
265 one trait (BA) for Egaseed 77 and one trait (CRL) for SC-10 might be the reason of their  
266 superiority in grain yield, because good roots may help the plants to uptake more water and  
267 nutrients from the soil for their biological activities, especially under drought conditions [4, 21,  
268 22] (Wright and Nageswara, 1994; Lynch, 1995; Henry et al., 2011).

269 In general, the commercial varieties P-3444, SC-128, Egaseed-77 and SC-10 were the  
270 best genotypes in our experiment; they showed the highest grain yield and the best root  
271 architectural traits across all studied irrigation treatments; they could be recommended for  
272 farmers use under a range of different environments as well as for maize breeding programs. On  
273 the contrary, it is observed that most of root and yield traits with undesirable mean values were  
274 exhibited by populations and the *vice versa* for traits with desirable means, which were mostly  
275 shown by the single crosses.

276 **Genotype × water stress interaction**

277 For root traits (Table 4), data were measured under WWF, WWG, WSF and WSG.  
 278 Under WWF, WWG, WSF and WSG, for BW the lowest mean was exhibited by genotypes No.  
 279 2, 13, 17 and 21 and the highest mean was shown by genotypes No. 17, 19, 4 and 10, for BN the  
 280 lowest mean by genotypes No. 21, 12, 4 and 21 and the highest mean by genotypes No. 11, 11,  
 281 10 and 10, for BA the lowest by genotypes No. 1, 9, 14 and 1 and the highest mean was shown  
 282 by genotypes No. 19, 21, 21 and 19, for BB the lowest by genotypes No. 18, 18, 13 and 20 and  
 283 the highest mean was shown by genotypes No. 5, 15, 6 and 9, for CN the lowest by genotypes  
 284 No. 18, 19, 13 and 13 and the highest mean was shown by genotypes No. 12, 8, 6 and 3, for CA  
 285 the lowest by genotypes No. 2, 5, 7 and 1 and the highest mean was shown by genotypes No. 10,  
 286 10, 21 and 10, for CB the lowest by genotypes No. 21, 17, 19 and 19 and the highest by  
 287 genotypes No. 8, 8, 6 and 8, for CRL the lowest by genotypes No. 14, 18, 22 and 22 and the  
 288 highest mean by genotypes No. 8, 5, 9 and 4, for RC the lowest by genotypes No. 18, 19, 19 and  
 289 21 and the highest by genotypes No. 7, 8, 7 and 8 and for RDW the lowest by genotypes No. 20,  
 290 18, 19 and 21 and the highest by genotypes No. 8, 8, 5 and 8, respectively.

Formatted: Space After: 0 pt

291 **Table 4.** Average, minimum (Min) and maximum (Max) values under each irrigation treatment  
 292 for all studied root traits and grain yield across two seasons.

Formatted: Font: Not Bold

Parameter	WWF	WWG	WSF	WSG	WWF	WWG	WSF	WSG
	<b>Brace Root Whorls No.</b>				<b>Brace Root No.</b>			
<b>Aver.</b>	2.52	2.48	2.29	2.64	39	37.1	31.5	40.8
<b>Min</b>	2 (2)	1.66 (13)	1.8 (17)	1.5 (21)	27.3 (21)	22.7 (12)	23 (4)	25.2 (21)
<b>Max</b>	3.1(17)	3.33(19)	2.9 (4)	3.3(10)	47(11)	54.7(11)	43.3(10)	59(10)
<b>LSD<sub>05</sub></b>	0.7	0.81	0.57	0.81	16.58	14.5	7.3	14.76
	<b>Brace Root Angle (Score)</b>				<b>Brace Root Branching (Score)</b>			
<b>Aver.</b>	6.7	6.7	6.9	6.5	5.3	4.7	4.9	4.7
<b>Min</b>	5 (1)	5 (9)	5.8 (14)	4.7 (1)	3.3 (18)	2 (18)	3 (13)	2.3 (20)
<b>Max</b>	8.3 (19)	7.3 (21)	7.5 (21)	7.5 (19)	7 (5)	7 (15)	6.8 (6)	6.2 (9)
<b>LSD<sub>05</sub></b>	1.62	1.88	1.02	1.25	2.38	2.66	1.66	2.02
	<b>Crown Root Number (Score)</b>				<b>Crown Root Angle (Score)</b>			
<b>Aver.</b>	3.82	2.66	3.38	3.05	6.8	6.5	6.9	6.5
<b>Min</b>	1.7 (18)	1(19)	1.8 (13)	1.8 (13)	5.7 (2)	5.3 (5)	5 (7)	5.2 (1)
<b>Max</b>	6 (12)	4 (8)	5.3 (6)	5 (3)	8 (10)	8 (10)	8 (21)	8.5 (10)
<b>LSD<sub>05</sub></b>	2.2	1.8	1.3	1.47	1.6	1.92	1.2	1.25
	<b>Crown Root Branching (Score)</b>				<b>Crown Root Length (cm)</b>			
<b>Aver.</b>	4.6	4.1	4.6	3.7	22.4	23.2	23.9	21.76
<b>Min</b>	3 (2)	2 (17)	3.2 (19)	2.2 (19)	18.6 (14)	18.8 (18)	21.2 (22)	16.9 (22)

<b>Max</b>	6 (8)	7.3 (8)	6.3 (6)	6.5 (8)	25.9 (8)	28.1(5)	26.2 (9)	26 (4)
<b>LSD<sub>05</sub></b>	1.95	2.35	1.49	1.54	6.67	5.1	4.1	4.4
	<b>Root Circumference (cm)</b>				<b>Root Dry Weight (g)</b>			
<b>Aver.</b>	34.7	30.7	34.4	30.9	26.2	21	18.8	23.3
<b>Min</b>	28.1(18)	23.3 (19)	26.5(19)	23.3(21)	8.2 (20)	8.2 (18)	9.8 (19)	9.9 (21)
<b>Max</b>	40.4(7)	41(8)	42.5(7)	36.6(8)	40.7 (8)	44.9 (8)	33.6 (5)	40.1(8)
<b>LSD<sub>05</sub></b>	6.48	6.5	4.97	4.95	14.36	12.96	9.53	11.53
	<b>Grain Yield/Plant (g)</b>				<b>Grain Yield/ha(ton)</b>			
	<b>WW</b>	<b>WSF</b>	<b>WSG</b>		<b>WW</b>	<b>WSF</b>	<b>WSG</b>	
<b>Aver.</b>	128.2	91.4	102.2		9.03	5.8	6.72	
<b>Min.</b>	82.9 (19)	31.8 (22)	58.9 (15)		3.91 (22)	1.39 (22)	2.77 (22)	
<b>Max.</b>	168.1(1,5)	156.4(6,4)	179.7(8,6,4)		15.25(8,5,6)	10.55(4,8,6)	13.45(8,6)	
<b>LSD<sub>05</sub></b>	23	13.3	12.7		0.75	0.63	0.71	

Means of minimum and maximum are followed by genotype No. (Between brackets).

Formatted: Indent: Left: -0.56"

Formatted: Indent: Left: -0.56", Line spacing: 1.5 lines

Formatted: Space After: 0 pt

For grain yield (Tables 5 and 6), data were measured under WW, WSF and WSG. The lowest mean GYPP was shown by genotypes No. 19, 22 and 15 and the highest by genotypes No. 1, 6 and 8 under WW, WSF and WSG, respectively. For GYPH, the lowest mean was exhibited by Genotypes No. 22, 22 and 22 and the highest mean was shown by Genotypes No. 8, 4 and 8 under WW, WSF and WSG, respectively.

Formatted: Indent: Left: 0", First line: 0"

Formatted: Font: Not Bold

**Table 5.** Means of grain yield/plant and grain yield/ha for each genotype under each irrigation regime (well watering; WW, water stress at flowering; WSF and water stress at grain filling; WSG) across 2016 and 2017 seasons.

Genotype	WW	WSF	Ch%	WSG	Ch%	WW	WSF	Ch%	WSG	Ch%
	<b>Grain yield/plant</b>					<b>Grain yield/ha</b>				
<b>1</b>	168.1	78.0	53.6	102.7	38.9	9.95	4.40	55.8	6.30	36.7
<b>2</b>	131.7	73.3	44.3	92.0	30.1	8.51	3.79	55.5	5.51	35.2
<b>3</b>	124.0	75.6	39.1	109.0	12.2	7.98	4.29	46.3	6.29	21.2
<b>4</b>	151.6	147.9	2.5	132.5	12.6	9.56	8.35	12.7	6.36	33.5
<b>5</b>	166.3	123.2	25.9	126.0	24.2	10.22	5.96	41.7	6.65	34.9
<b>6</b>	150.4	156.4	-4.0	168.7	-12.2	10.05	8.14	19.1	8.38	16.6
<b>7</b>	128.5	131.2	-2.1	106.8	16.9	7.34	6.41	12.6	4.76	35.2
<b>8</b>	150.4	137.6	8.5	179.7	-19.5	12.11	8.21	32.2	10.67	11.9
<b>9</b>	134.4	105.6	21.4	121.0	9.9	8.12	5.64	30.6	6.69	17.7
<b>10</b>	134.3	98.9	26.4	117.7	12.3	8.32	5.31	36.2	6.43	22.8
<b>11</b>	125.5	78.5	37.4	84.7	32.5	7.61	4.02	47.2	4.50	40.9
<b>12</b>	119.4	91.0	23.8	111.5	6.6	7.79	5.12	34.2	6.09	21.8
<b>13</b>	149.4	111.1	25.6	120.7	19.2	9.28	5.96	35.8	7.16	22.8

14	133.6	89.7	32.9	81.9	38.7	5.65	4.15	26.5	3.86	31.7
15	125.4	84.7	32.5	58.9	53.1	4.96	3.79	23.6	3.05	38.5
16	118.6	56.2	52.6	81.9	30.9	4.30	2.84	33.9	4.12	4.1
17	110.9	65.0	41.4	70.8	36.2	4.86	2.80	42.4	3.62	25.6
18	110.5	74.2	32.9	85.8	22.4	5.37	3.22	40.1	4.54	15.4
19	82.9	59.4	28.4	75.8	8.5	3.83	2.33	39.1	3.38	11.9
20	106.6	79.7	25.2	91.4	14.3	4.64	3.00	35.4	3.63	21.9
21	100.8	61.8	38.7	70.4	30.2	3.79	2.60	31.5	3.04	19.8
22	96.9	31.8	67.2	58.9	39.3	3.10	1.11	64.2	2.19	29.4
<b>Average</b>	128.2	91.4	28.7	102.2	20.3	7.15	4.61	35.5	5.33	25.5
<b>Min.</b>	82.9	31.8		58.9		3.10	1.11		2.19	
<b>Max.</b>	168.1	156.4		179.7		12.11	8.35		10.6	
<b>LSD<sub>.05</sub></b>	23	13.3		12.7		0.6	0.5		0.6	
<b>LSD<sub>.01</sub></b>	30.5	17.6		16.8		0.8	0.7		0.8	

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

304 Ch% = 100(WW-WSF or WSG)/WW

Formatted: Space After: 0 pt

305  
306 On the contrary, the worst genotypes were No. 22 (Sweepstakes) in 3 traits (GYPP,  
307 GYPH, CRL) under WSG, 3 traits (GYPP, GYPH, CRL) under WSF and one trait (GYPH)  
308 under WW, the genotype No. 21 (Golden Republic) in 4 traits (BW, BN, RC, RDW) under  
309 WSG, two traits (BN,CB) under WWF, the genotype No. 19 (Nebraska) in one trait (CB) under  
310 WSG, and 3 traits (CB, RC, RDW) under WWG and the genotype No. 18 (Nubaria) in two traits  
311 (CN, RC) under WWG and one trait (GYPP) under WW.

312 The four highest and the four lowest performing genotypes under water stress at  
313 flowering (WSF) and grain filling (WSG) across seasons are presented in Table (6). Under WSF  
314 conditions, the highest mean grain yield/ha was achieved by the single cross Egaseed-77  
315 (developed by Egaseed Co.), followed by P-3444 (developed by Pioneer Co.), SC 128  
316 (developed by ARC, Egypt) and HT-2066 (developed by Hi Tec Co.) in a descending order. The  
317 single cross Egaseed-77 was amongst the four highest genotypes under WSF for GYPH, GYPP,  
318 BA and CRL. The single cross P-3444 was amongst the four highest genotypes under WSF for  
319 GYPH, GYPP, CN, CB and CRL. The single cross SC-128 was amongst the four highest  
320 genotypes under WSF for GYPH, GYPP, BB, CN, CB, RC, and RDW. The single cross HT-  
321 2066 was amongst the four highest genotypes under WSF for GYPH, GYPP, CN and RC.

Formatted: Space After: 0 pt

322 **Table 6.** The four highest and the four lowest genotypes for studied traits under water stress at flowering  
323 (WSF) and grain filling (WSG) across seasons.

Formatted: Font: Not Bold

WSF	WSG	WSF	WSG	WSF	WSG
Brace root whorls No.		Brace root No. Highest		Brace root angle (score)	

Formatted: Space After: 0 pt

Pop-45	32D99	32D99	32D99	Nebraska	Nebraska
HT-1100	HT-1100	TWC-352	TWC-352	Golden	SC-10
32D99	TWC-360	Pop-45	HT-1100	Fine 1005	Golden
TWC-360	Pop-45	HT-1100	TWC-360	Eg-77	Sweep
<b>Lowest</b>					
Fine 1005	Eg-77	Fine 1005	P-3444	SC-128	TWC-352
SC-128	P-3444	Midland	Eg-77	HT-2066	Giza
Eg-77	30K09	Golden	30K09	SC-166	TWC-324
P-3444	Golden	Eg-77	Golden	TWC-360	HT-2031
<b>Brace root branching (score)</b>		<b>Crown root number (score)</b>		<b>Crown root angle (score)</b>	
<b>Highest</b>					
SC-128	SC-166	SC-128	Fine 1005	Golden	32D99
TWC-352	SC-128	P-3444	HT-2031	32D99	Nebraska
SC-166	P-3444	HT-2066	SC-128	Midland	Midland
32D99	SC-10	TWC-352	HT-1100	TWC-324	Golden
<b>Lowest</b>					
Golden	Nubaria	Eg-77	SC-166	TWC-360	P-3444
Giza	Wat- 11	Sweep	Midland	P-3444	HT-1100
Nebraska	Golden	TWC-324	TWC-324	HT-2031	HT-2031
TWC-324	Midland	Golden	Golden	HT-2066	HT-2066
<b>Crown root branching (score)</b>		<b>Crown root length (cm)</b>		<b>Root circumference (cm)</b>	
<b>Highest</b>					
SC-128	P-3444	P-3444	Eg-77	HT-2066	P-3444
P-3444	HT-1100	SC-166	P-3444	TWC-352	30K09
TWC-352	HT-2066	SC-10	HT-1100	TWC-352	TWC-352
SC-166	SC-128	Eg-77	SC-10	SC-128	HT-2031
<b>Lowest</b>					
Fine 1005	Golden	Pop-45	Nubaria	Nubaria	Nebraska
Eg-77	32D99	HT-2066	Golden	Midland	Midland
TWC-324	TWC-324	Midland	Giza	Golden	Nubaria
Nebraska	Nebraska	Sweep	Sweep	Nebraska	Golden
<b>Root dry weight (g)</b>		<b>Grain yield/plant (g)</b>		<b>Grain yield/ha</b>	
<b>Highest</b>					
SC-10	P-3444	SC-128	P-3444	Eg-77	P-3444
Fine 1005	HT-1100	Eg-77	SC-128	P-3444	SC-128
SC-128	SC-128	P-3444	Eg-77	SC-128	TWC-324
TWC-352	HT-2031	HT-2066	SC-10	HT-2066	SC-166
<b>Lowest</b>					
Midland	Nebraska	Golden	Pop-45	Pop-45	Nebraska
TWC-324	Midland	Nebraska	Golden	Golden	TWC-352
Golden	Nubaria	Giza	TWC-352	Nebraska	Golden
Nebraska	Golden	Sweep	Sweep	Sweep	Sweep

324

325

326

327

328

329

330

331

332

Under WSG conditions, the highest mean grain yield/ha was achieved by the single cross P-3444 (developed by Pioneer) followed by SC-128 (developed by ARC), TWC-324 (developed by ARC) and SC-166 (developed by ARC) in a descending order. The single cross P-3444 was amongst the four highest genotypes in GYPH, GYPP, BB, CB, CRL, RC and RDW, i.e. most important grain yield and root architecture traits. The single cross SC-128 was amongst the four highest genotypes in GYPH, GYPP, BB, CN, CB and RDW (the most important grain yield and root architecture traits). The single cross SC-166 was amongst the four highest genotypes in GYPH and BB.

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Tab stops: Not at 3.25" + 6.5"

Formatted: Indent: Left: -0.63"

Formatted: Space After: 0 pt

333 Results from Tables (4 and 5) concluded that the best genotypes were No. 8 (P-3444) in  
 334 5 traits (GYPP, GYPH, CB, RC, RDW) under WSG, 4 traits (CN, CB, RC, RDW) under WWG,  
 335 3 traits (CA, CRL, RDW) under WWF and one trait (GYPH) under WW, the genotype No. 6  
 336 (SC 128) in 4 traits (GYPP, BB, CA, CB) under WSF, the genotype No.5 (SC 10) in two traits  
 337 (BB and CRL) under WWF and WWG, respectively, the genotype No. 7 (Hi-Tec 2066) in one  
 338 trait (RC) under WSF and RC under WWF, the genotype No. 4 (Egaseed 77) in one trait  
 339 (GYPH) under WSF, and the genotype No. 2 (30K09) in one trait (GYPH) under WSF.

340 The best genotypes in grain yield under drought at either flowering or grain filling were  
 341 characterized by one or more desirable root architecture traits. Accumulating genes of more  
 342 desirable root characteristics in one genotype might help plants to search water and nutrients in  
 343 the soil and consequently help plant to accomplish its biological activities and achieve almost its  
 344 potential grain yield under drought stress at flowering or grain filling stages [4, 10, 21-24]  
 345 (Wright and Nageswara, 1994; Lynch,1995; Hund et al.,2009 b;Hund,2010; Henry et al.,  
 346 2011;Trachsel et al. (2011). The studied single-cross hybrids P-3444, Egaseed-77 and SC-128  
 347 were considered drought tolerant genotypes under drought stress at flowering and/or grain filling  
 348 stages and would be offered to future breeding programs to utilize their genes of desirable root  
 349 architecture and grain yield traits in improving maize drought tolerance under Egyptian  
 350 conditions. It should be mentioned that the hybrid P-3444 was characterized in this experiment  
 351 by its ability to stay green even under water stress, which might help it to tolerate water stress at  
 352 grain filling stage in a way much better than other tested hybrids and populations.

### 333D) Drought tolerance index

354 Drought tolerance index (DTI) values of studied genotypes under the stressed  
 355 environments WSF and WSG are presented in Table (7). According to our scale, when DTI is  
 356  $\geq 1.0$ , it indicates that genotype is tolerant (T), if DTI is 1.0, it indicates that genotype is  
 357 moderately tolerant (MT) and if DTI is  $< 1.0$ , it indicates that genotype is sensitive (S).

358 Based on DTI values, the 22 studied maize genotypes were grouped into three categories  
 359 under water stress at flowering, namely tolerant (10 genotypes), moderately tolerant (two  
 360 genotypes) and sensitive (10 genotypes) (Table 7). Under water stress conditions at grain filling,  
 361 number of tolerant (T), and sensitive (S) genotypes were 11, and 11, respectively.

362 **Table 7.** Drought tolerance index (DTI) of each genotype under WSF and WSG environments.  
 363

Genotype No.	Designation	WSF	WSG	Genotype No.	Designation	WSF	WSG
--------------	-------------	-----	-----	--------------	-------------	-----	-----

Formatted: Space After: 0 pt

Formatted: Indent: Left: -0.56", Space After: 0 pt

Formatted: Space After: 0 pt

Formatted: Indent: Left: -0.69"

Formatted: Indent: Left: 0", First line: 0"

Formatted: Font: Not Bold

Formatted: Space After: 0 pt, Line spacing: single



1	Hi-Tec-2031	1.3	1.6	12	Watania -11	1.2	1.2
2	P-30K09	1.0	1.2	13	TWC-324	1.7	1.7
3	Fine 1005	1.0	1.3	14	TWC-360	0.7	0.6
4	Egaseed-77	2.4	1.6	15	TWC-352	0.6	0.4
5	SC-10	1.8	1.8	16	Giza Baladi	0.4	0.5
6	SC-128	2.5	2.2	17	Population-45	0.4	0.5
7	Hi-Tec-2066	1.4	0.9	18	Nubaria	0.5	0.6
8	P-3444	3.0	3.4	19	Nebraska Midland	0.3	0.3
9	SC-166	1.4	1.4	20	Midland_Cunningham	0.4	0.4
10	P-32D99	1.3	1.4	21	Golden Republic	0.3	0.3
11	Hi-Tec-1100	0.9	0.9	22	Sweepstakes 5303	0.1	0.2

364  
365 The highest DTI under both the two stressed environments (WSF and WSG) was  
366 exhibited by the genotype No. 8 (P-3444). The 2<sup>nd</sup> and 3<sup>rd</sup> highest genotypes in DTI were SC-128  
367 and Egaseed-77 under WSF and SC-128 and SC-10 under WSG. For productivity (grain  
368 yield/plant) under WSF, the genotype Egaseed-77 ranked 1<sup>st</sup>, but P-3444 and SC-128 ranked 3<sup>rd</sup>.  
369 Under WSG, P-3444, SC-128 and SC-10 ranked 1<sup>st</sup>, 2<sup>nd</sup> and 3<sup>rd</sup>, for productivity as well as  
370 drought tolerance index.

371 -On the contrary, the most drought sensitive genotypes were the open-pollinated  
372 populations Sweepstakes 5303, Golden Republic and Nebraska Midland under both water stress  
373 environments (WSF and WSG); their grain yield were the lowest.

### 374 3.3. Superiority of drought tolerant (T) to sensitive (S) genotypes

375 Based on grain yield/plant and drought tolerance index (DTI) the best three genotypes  
376 were the single cross hybrids P-3444, SC-128 and Egaseed-77 under WSF and P-3444, SC-128  
377 and SC-10 under WSG, while the most drought sensitive and lowest yielding genotypes were the  
378 populations Sweepstakes, Golden Republic and Nebraska Midland under both water stress  
379 environments (WSF and WSG). Data averaged for each of the two groups (T and S) under WSF  
380 and under WSG indicated that GYPP of drought tolerant (T) was greater than that of the  
381 sensitive (S) genotypes by 189.0 and 131.3 % under drought at flowering (WSF) and grain filling  
382 (WSG), respectively (Table 8).

383  
384 **Table 8.** Superiority (Sup.%) of the three most tolerant (T) to the three most sensitive (S) genotypes for  
385 selected traits under the stressed environments WSF and WSG, combined across 2016 and 2017 seasons.

Trait	WSF	WSG
-------	-----	-----

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt

Formatted: Indent: Left: 0", First line: 0"

Formatted: Font: Not Bold

	T	S	Sup. %	T	S	Sup. %
<b>Grain yield/plant</b>	147.3	51.0	189.0**	158.1	68.3	131.3**
<b>Crown root number</b>	4.2	2.4	76.7**	3.4	2.3	45.2*
<b>Crown root branching</b>	5.4	3.8	42.6*	4.6	2.5	84.4**
<b>Crown root length</b>	25.6	22.9	11.3*	23.3	18.6	25.4*
<b>Root circumference</b>	35.6	28.4	25.4**	32.6	26.4	23.6*
<b>Root dry weight</b>	20.1	10.7	86.7*	33.1	14.6	126.3**

386 \* and \*\* indicate significance at 0.05 and 0.01 probability levels, respectively.

387  
388 Significant superiority of drought tolerant (T) over sensitive (S) genotypes in GYPP  
389 under drought at flowering and grain filling was associated with significant superiority in higher  
390 CN (76.7 and 45.2%), CB (42.6 and 84.4%), higher CRL (11.3 and 25.4 %), higher RC (25.4  
391 and 23.6%) and higher RDW (86.7 and 126.3%), respectively.

### 3.4. Correlations between drought tolerance and root traits

393 Drought tolerance index had a strong significant ( $p \leq 0.01$ ) and positive correlation with  
394 grain yield/plant ( $r = 0.912^{**}$  and  $0.941^{**}$ ) under WSF and WSG conditions, respectively (Table  
395 9). Drought tolerance had a significant and positive correlation coefficient, with crown root  
396 length ( $r = 0.693^{**}$  and  $0.561^{**}$ ), root circumference ( $0.440^*$  and  $0.499^*$ ) crown root dry weight  
397 ( $r = 0.410^*$  and  $0.592^{**}$ ) under WSF and WSG conditions, respectively.

398 Moreover, drought tolerance index had a significant and negative correlation coefficient  
399 with brace root whorls; BW ( $-0.598^{**}$ ) and a significant and positive correlation coefficient with  
400 brace root branching; BB ( $0.506^*$ ) and crown root branching ( $0.489^*$ ) under WSG.

401 **Table 9.** Correlation coefficients between drought tolerance index (DTI) and means of studied traits of all genotypes  
402 under water stress at flowering (WSF) and at grain filling (WSG) across seasons.

Trait	WSF	WSG	Trait	WSF	WSG
Grain yield/plant	.912**	.941**	Crown root angle	-.319	-.203
Brace root whorls number	-.598**	-.288	Crown root branching	.381	.489*
Brace root Number	-.250	-.231	Crown root length	.693**	.561**
Brace root angle	-.183	-.193	Root circumference	.440*	.499*
Brace root Branching	.169	.506*	Root dry weight	.410*	.592**
Crown root number	.469*	.320			

403 \* and \*\* indicate significance at 0.05 and 0.01 probability levels, respectively.

### 3.5. Correlations between grain yield and root traits

404  
405 Estimates of rank correlation coefficients among grain yield/plant and all studied root  
406 traits across the two seasons under well watering, water stress at flowering (WSF) and grain  
407 filling (WSG) were calculated across all genotypes and presented in Table (10). Under well  
408

Formatted: Space After: 0 pt

Formatted: Space After: 0 pt

Formatted: Font: Not Bold

Formatted: Indent: Left: 0", First line: 0"

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Right: 0", Space After: 0 pt, Line spacing: single

Formatted: Indent: Left: 0", Right: 0", Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Right: 0", Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Right: 0", Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Space After: 0 pt, Line spacing: single

Formatted: Font: Not Bold

Formatted: Space After: 0 pt

Formatted: Indent: Left: 0"

409 watering, grain yield/plant had a significant ( $p \leq 0.01$ ) and positive association with the root dry  
 410 weight (RDW) (0.42), root circumference (RC) (0.43), crown root length (0.26), crown root  
 411 branching (CB) (0.27), number of crown roots (CN) (0.23) and brace root branching (BB) (0.34).

412 Data in Table (10) showed that under WSF, grain yield/plant was significantly ( $P \leq$   
 413 0.01) and positively correlated with each of RC ( $r=0.33$ ) and CN ( $r=0.27$ ). Under water stress at  
 414 grain filling (WSG), grain yield/plant had a significant and positive correlation ( $p \leq 0.01$  or  
 415  $p \leq 0.05$ ) with CRL ( $r=0.33$ ), CB ( $r=0.25$ ), RDW ( $r=0.23$ ), BB ( $r=0.18$ ) and RC ( $r=0.17$ ).

416 **Table 10.** Correlation coefficients between grain yield/plant and each of studied root traits of maize under  
 417 well watering (WW), water stress at flowering (WSF) and water stress at grain filling (WSG) across two  
 418 years.  
 419

Environment	BW	BN	BA	BB	CN	CA	CB	CRL	RC	RDW
WW	-0.2	-0.07	-0.09	0.34**	0.23**	-0.14	0.27**	0.26**	0.43**	0.42**
WSF	-0.07	0.01	-0.2	0.13	0.27**	-0.03	0.08	-0.03	0.33**	0.13
WSG	-0.14	-0.12	-0.02	0.18*	0.21**	-0.08	0.25**	0.33**	0.17*	0.23**

420 \* and \*\* indicate significance at 0.05 and 0.01 probability levels, respectively. GYPP = grain yield per plant, BW= Number  
 421 of above-ground whorls occupied with brace roots, BN= Number of brace roots, BA= Angle of 1st arm of the brace  
 422 roots originating from whorl 1, BB= Branching density of brace roots, CN= Number of crown roots, CA= Crown  
 423 roots angle, CB= Branching density of crown roots, CRL= Crown root length, RC= Root circumference, RDW=  
 424 Roots dry weight.  
 425

## 426 Grouping genotypes

### 427 Based on drought tolerance and grain yield

428 Mean grain yield/fed of studied genotypes under water stress at flowering (WSF) and  
 429 grain filling (WSG), was plotted against drought tolerance index of the same genotypes under  
 430 WSF and WSG, respectively (Fig. 2), which made it possible to distinguish between four groups,  
 431 namely tolerant and high- yielding, tolerant and low-yielding, sensitive and high-yielding and  
 432 sensitive and low-yielding according to Sattelmacher *et al.*, 1994 [25], Worku *et al.* (2007) [26]  
 433 and Al-Naggar *et al.* (2015) [27].

434

Formatted: Indent: Left: -0.5"

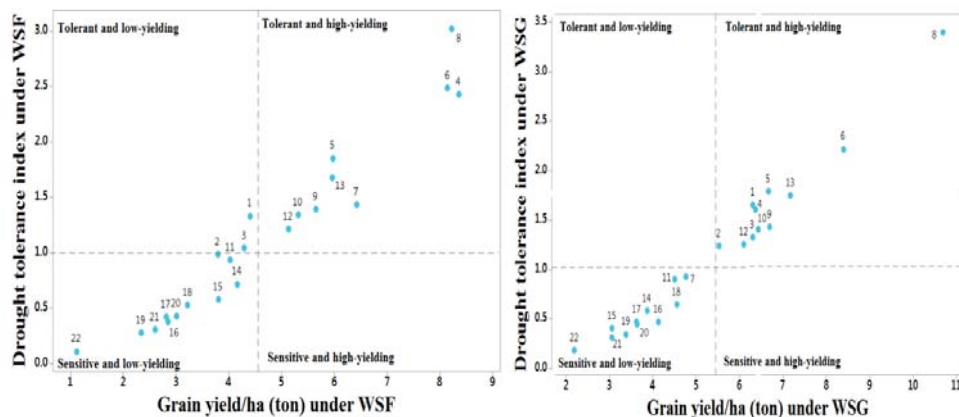
Formatted: Indent: Left: 0", First line: 0"

Formatted: Font: Not Bold

Formatted: Font: Not Bold

Formatted: Line spacing: 1.5 lines

Formatted: Space After: 0 pt



435  
436 **Figure. 2.** Relationships between drought tolerance index (DTI) and means of GYPH of genotypes (from No.1 to  
437 No.22) under water stress at flowering (WSF) and grain filling (WSG) combined across seasons. Broken lines  
438 represent mean grain yield/fed and DTI.  
439

440 Under water stress at flowering (WSF), the genotypes No 8 followed by No. 4, 6, 5, 7,  
441 13, 9, 10 and 12 were classified as the drought tolerant and high yielding genotypes, *i.e.* they  
442 could be considered as the most water stress tolerant and the most responsive genotypes to water  
443 stress at flowering in this study (Fig. 2). There was no genotype belonging to the group of  
444 sensitive and high yielding genotypes under WSF. The genotypes No. 1 and 3 occupied the  
445 group of tolerant and low yielding under WSF. The genotypes No 22, 19, 21, 16, 17, 20, 18, 15,  
446 14, 11 and 2 were classified as water stress sensitive and low yielding and therefore could be  
447 considered sensitive and low yielding.

448 Under water stress at grain filling (WSG), the genotypes No. 8 followed by 6, 13, 5, 1, 4,  
449 9, 10, 3, 12 and 2 were classified as drought tolerant and high yielding, they could be considered  
450 as the most water stress tolerant and the most responsive genotypes to water stress at grain filling  
451 in this study (Fig. 3). On the contrary, genotypes No. 22, 21, 15, 19, 20, 17, 16, 14, 18, 11 and 7  
452 were classified as water stress sensitive and low yielding (Fig. 2).

453 According to Fageria and Baligar (1994 and 1997a and b) [28-30] genotypes belonging to  
454 the 1<sup>st</sup> group "tolerant and high yielding" (above all) and 2<sup>nd</sup> group "tolerant and low yielding" (to  
455 a lesser extent) (we did not have) appear to be the most desirable materials for breeding  
456 programs that deal with adaptation to water stress. It was observed that the genotypes No. 8, 6, 4,  
457 13, 5, 9, 10 and 12 occupied the first group (E-R) under both WSF and WSG conditions; they

Formatted: Indent: Left: -0.19", Space Before: 0 pt

Formatted: Font: Not Bold

Formatted: Space After: 0 pt

458 had genes of high water efficiency; i.e. drought tolerance to both WSF and WSG stages and  
459 genes for high yield under well watering conditions.

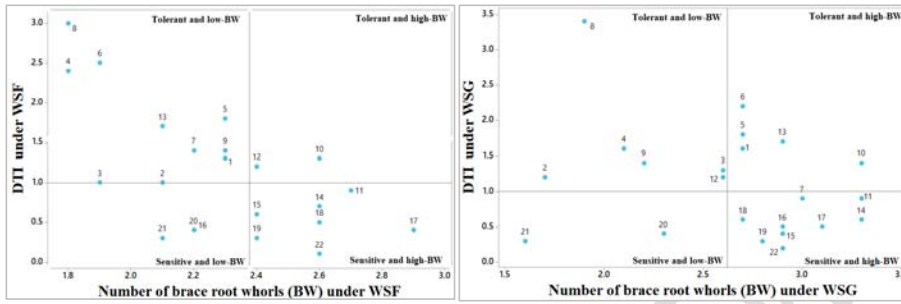
460 Summarizing the above-mentioned classifications, it is apparent that the genotypes No. 8  
461 (P-3444) followed by 6 (SC-128), 4 (Egaseed-77), 5 (SC-10), 13 (TWC-324), 7 (Hi Tec-2066), 9  
462 (SC-166), 10 (P-32D99) and 12 (Watania 11) were the best genotypes that occupied the first  
463 group (best one) in both classifications; they are the most efficient, most drought tolerant, the  
464 highest yielder under WSF—as well as WW. The genotypes No. 8 (P-3444) followed by 6 (SC-  
465 128), 13 (TWC-324), 5 (SC-10), 1 (Hi Tec-2031), 4 (Egaseed-77), 9 (SC-166), 10 (P-32D99), 3  
466 (Fine 1005), 12 (Watania 11) and 2 (P-30K09) were the best genotypes that occupied the first  
467 group (best one) in both classifications; they are the most efficient, most drought tolerant, the  
468 highest yielder under WSG—as well as WW.

469 It was observed that the genotypes No 8 (P-3444) followed by 6 (SC-128), 4 (Egaseed-  
470 77), 5 (SC-10), 13 (TWC-324), 7 (Hi Tec-2066), 9 (SC-166), 10 (P-32D99) and 12 (Watania 11)  
471 were the best in the first group for both stresses WSF and WSG; they are the most efficient, most  
472 drought tolerant and the highest yielders under WSF and WSG as well as WW. In accordance to  
473 these results, a previous study by Al-Naggar *et al.* (2011) [17], proved that the single cross  
474 hybrid SC-128 (genotype No. 6 in the present study) was the most water efficient (drought  
475 tolerant) under WSF and the most responsive to WW based on grain yield, ears/plant, kernels/  
476 plant, ASI and leaf senescence.

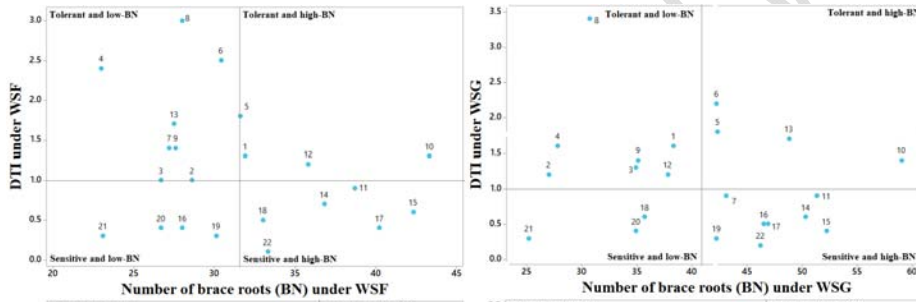
#### 477 **Based on drought tolerance and root traits**

478 Means of root traits of studied genotypes under water stress at flowering (WSF) and grain  
479 filling (WSG), were plotted against drought tolerance index (DTI) of the same genotypes under  
480 WSF and WSG; respectively (Fig. 3), which made it possible to distinguish between four groups,  
481 namely tolerant and high value of root trait, tolerant and low value of root trait, sensitive and  
482 high value of root trait and sensitive and low value of root trait.—According to Fageria and  
483 Baligar [29] (1997a), genotypes belonging to the 1<sup>st</sup> group "tolerant and high value of root trait"  
484 (above all) appear to be the most desirable materials for breeding programs.

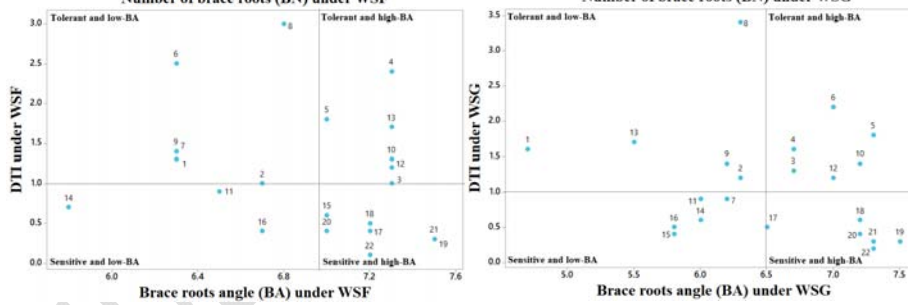
485



486

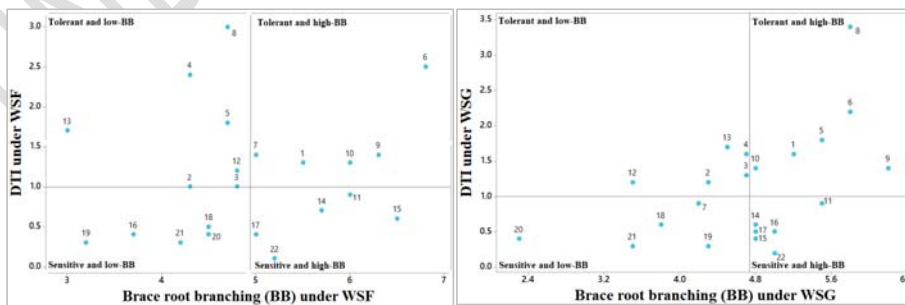


487



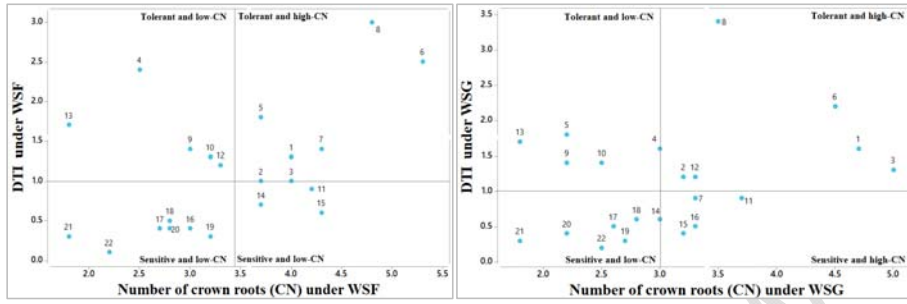
488

489

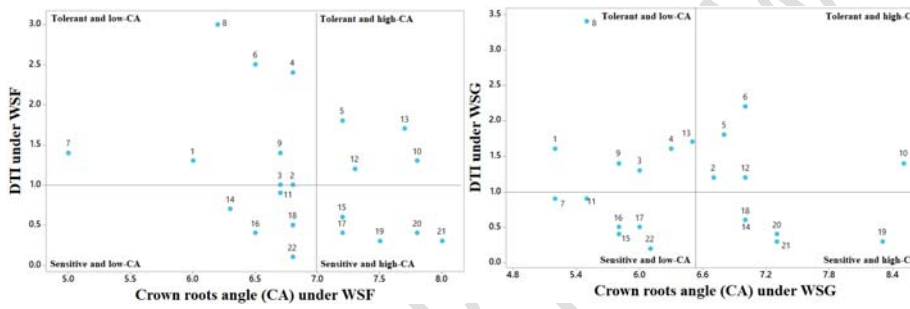


490

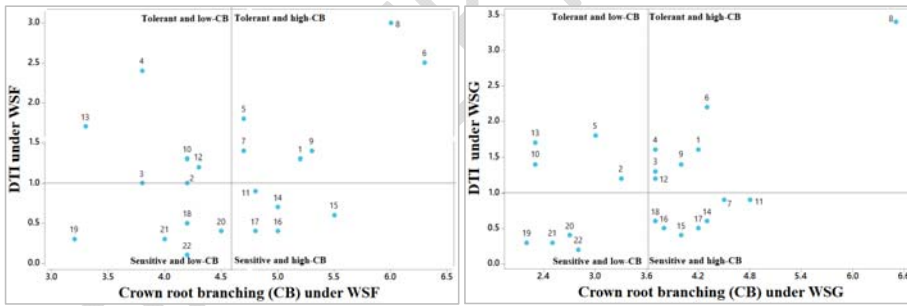
491



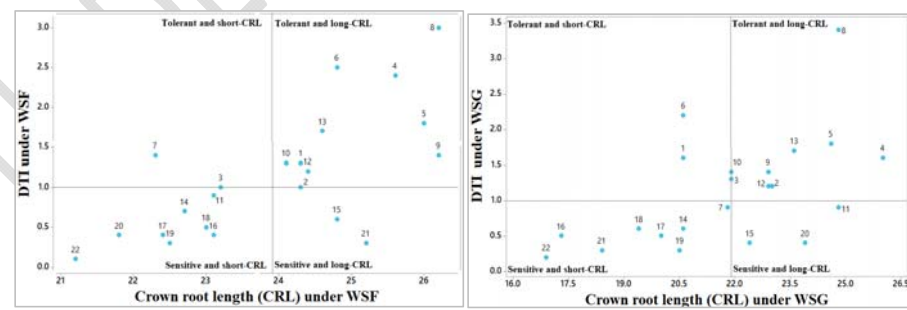
492



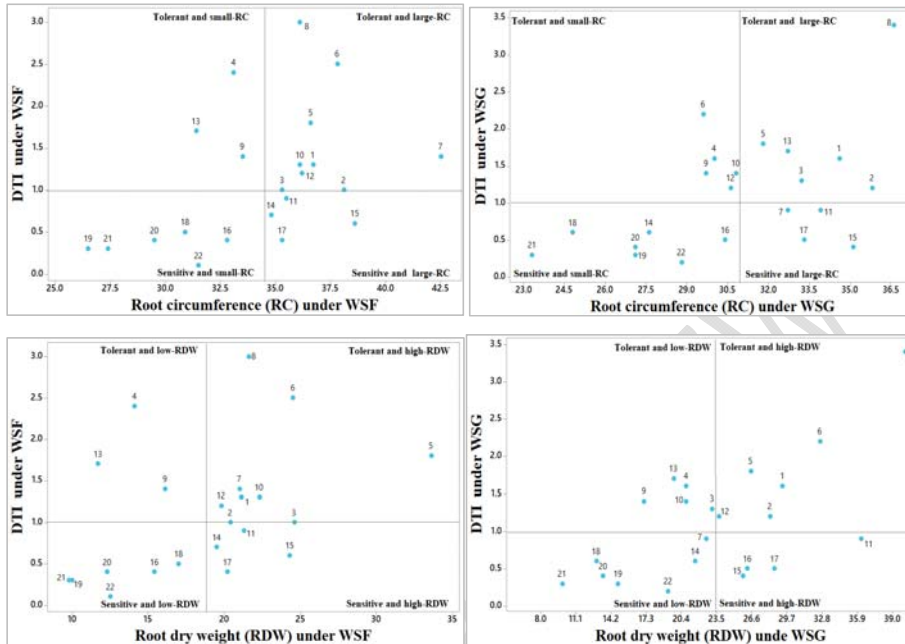
493



494



495



496

497

498

499

500

501

502

503

504

505

506

507

508

509

510

511

512

513

514

**Fig-ure 3.** Relationships between drought tolerance index (DTI) and means of number of whorls carrying brace roots, brace root branching, crown root number, crown root branching, root circumference, crown root length, and root dry weight, of genotypes (from No. 1 to No.22) under water stress at flowering (WSF) and grain filling (WSG) combined across seasons. Broken lines represent mean DTI and root trait.

Formatted: Indent: Left: 0", First line: 0",  
Space After: 0 pt

Formatted: Font: Not Bold

Figure (3) indicates that the 1<sup>st</sup> group "tolerant and high value of root trait" included the genotypes No. 10 and 12 under WSF, No. 10, 13, 6, 5 and 1 under WSG for number of whorls carrying brace roots, No. 10, 12, 1 and 5 under WSF, No. 10, 13, 1, 5 and 6 under WSG for number of brace roots, No. 4, 13, 10, 12, 3 and 5 under WSF, No. 5, 6, 10, 12, 4 and 3 under WSG for brace root angle, No. 6, 9, 10, 1 and 7 under WSF, No. 9, 6, 5, 1 and 10 under WSG for brace root branching, No. 6, 8, 7, 1, 5, 3 and 2 under WSF, No. 3, 1, 6, 8, 12 and 2 under WSG for number of crown roots, No. 10, 13, 12 and 5 under WSF, No. 10, 6, 12, 5 and 2 under WSG for crown root angle, No. 6, 8, 9, 1, 7 and 5 under WSF, No. 8, 6, 1, 9, 4, 3 and 12 under WSG for crown root branching, No. 8, 4, 5, 6, 9, 12, 1, 10 and 2 under WSF, No. 8, 4, 5, 13, 9, 2, 3, 10 and 12 under WSG for crown root length, No. 7, 6, 8, 5, 1, 10, 12, 2 and 3 under WSF, No. 8, 2, 1, 3, 13 and 5 under WSG for root circumference and No. 5, 6, 8, 10, 7, 1, 12, 3 and 2 under WSF, No. 8, 6, 1, 5, 2 and 12 under WSG for root dry weight.

**Mechanisms of drought tolerance of the most tolerant and high-yielding genotypes:**



515 The above-mentioned results (Figs. 2 and 3) helped us to identify the root traits that  
516 characterize the most drought tolerant and high-yielding genotypes, in descending order, as  
517 follows:

5181. **Genotype No. 8 (SC-P-3444):** Five traits (high CN, CB, large RC, long CRL and heavy RDW) under both WSF and WSG.

5202. **Genotype No. 6 (SC-128):** Four traits (high CN, CB, BB, large RC and heavy RDW) under both  
521 WSF and WSG.

5223. **Genotype No. 4 (SC-Egaseed-77):** Two traits (steep brace root; i.e. large BA and long CRL)  
523 under both WSF and WSG.

5244. **Genotype No. 5 (SC-10):** Six traits (high CN, CB, BA, RC, long CRL and heavy RDW) under  
525 WSF and five traits (high BA, CA, large RC, long CRL and heavy RDW) under WSG.

5265. **Genotype No. 13 (TWC-324):** Two traits (steep brace root; i.e. large BA and long crown root  
527 (CRL) under WSF and two traits (large RC and long CRL) under WSG.

5286. **Genotype No. 9 (SC-166):** Two traits (high CB and long crown root CRL) under both WSF and  
529 WSG.

5307. **Genotype No. 10 (SC-P-32D99):** Four traits (steep crown root; CA steep brace root; BA, long  
531 crown root; CRL and heavy root dry weight; RDW) under both WSF and WSG and one trait  
532 (heavy RDW) under WSF.

5338. **Genotype No. 12 (Watania TWC-11):** Seven traits (BW, BN, BA, CA, CRL, RC and RDW)  
534 under WSF and six traits (BA, CN, CA, CB, CRL and RDW) under WSG.

535 The present study suggested that further investigation should be conducted to determine the  
536 underlying root mechanisms contributing to the selection of water-efficient hybrids of maize.

537 In a recent study [31] (Shao et al., 2019), maize genotypes with less variation in root  
538 size, medium root size, medium broad root system and more inter-row root distribution help to  
539 reduce root-to-root competition and tend to have higher yield at high planting density.

## 540 CONCLUSIONS

541 Correlation analysis of the present study concluded that drought tolerance in maize had a  
542 strong and positive association with crown root length, root circumference and root dry weight  
543 under both WSF and WSG, a negative correlation with brace root whorls, and a positive  
544 correlation with crown root number under WSF and brace root branching and crown root  
545 branching under WSG. These root traits could be considered as putative mechanisms of drought

Formatted: Indent: Left: -0.25"

Formatted: Space Before: 0 pt, After: 0 pt

546 tolerance. The present study suggested that further investigation should be conducted to  
 547 determine the underlying plant mechanisms contributing to the selection of water-efficient  
 548 hybrids of maize. The cultivars Pioneer-3444, SC-128, Egaseed-77, SC-10 and TWC-324  
 549 showed the most drought tolerance and the highest yielding in a descending order; each had a  
 550 number of such drought tolerance mechanisms. These cultivars should be retested for drought  
 551 tolerance and grain productivity under drought stress and could be offered to plant breeding  
 552 programs for improving tolerance to drought and high grain yield.

### 553 REFERENCES

- 554 1] **Bai LP, Sui FG, Ge TD, Sun ZH, Lu YY, Zhou GS (2006)**. Effect of soil drought stress on leaf water  
 555 status, membrane permeability and enzymatic antioxidant system of maize. *Pedosphere* 16:326–332.
- 556 2] **Witt S, Galicia L, Lisee J, Cairns J, Tiessen A, Araus JL, Palacios-Rojas and N, Fernie ARR (2012)**  
 557 Metabolic and phenotypic responses of greenhouse-grown maize hybrids to experimentally controlled  
 558 drought stress. *Mol Plant* 5:401–417.
- 559 3] **Edmeades, G.O.; Bolanos, J.; Hernandez, M. and Ballo, S. (1993)**. Causes for silk delay in a low land  
 560 tropical maize population. *Crop Sci.*, 33: 1029-1035.
- 561 4] **Lynch, J.P. (1995)**. Root architecture and plant productivity. *Plant Physiol.*, 109: 7-13.
- 562 5] **Lynch, J.P. (2007)**. Roots of the second green revolution. *Aust. J. Bot.*, 55: 493-512.
- 563 6] **Hund, A.; Reimer, R. and Messmer, R. (2011)**. A consensus map of QTLs controlling the root length of  
 564 maize. *Plant and Soil*, 344: 143-158.
- 565 7] **Rauf, S. and Sadaqat, H.A. (2008)**. Effect of osmotic adjustment on root length and dry matter  
 566 partitioning in sunflower (*Helianthus annuus* L.) under drought stress. *Acta Agric Scand, Section B, Soil*  
 567 *& Plant Sci.*, 58(3): 252-260.
- 568 8] **Rauf, S.; Sadaqat, H.A.; Ahmed, R. and Khan, I.A. (2009)**. Genetics of root characteristics in  
 569 sunflower (*Helianthus annuus* L.) under contrasting water regimes. *Ind. J. Plant Physiol.*, 14: 319-327.
- 570 9] **King CA, Purcell LC, Brye KR, (2009)**. Differential wilting among soybean genotypes in response to  
 571 water deficit. *Crop Sci.* 49: 290–298.
- 572 10] **Trachsel, S.; Kaepller, S.M.; Brown, K.M. and Lynch, J.P. (2011)**. Shovelomics: high throughput  
 573 phenotyping of maize (*Zea mays* L.) root architecture in the field. *Plant Soil*, 341: 75-87.
- 574 11] **Fageria, N.K. (1992)**. Maximizing Crop Yields. Dekker. New York; 423.
- 575 12] **Steel, R.G.D.; Torrie, G.H. and Dickey, D.A. (1997)**. Principles and Procedures of Statistics: A  
 576 Biometrical Approach. 3rd ed. McGraw-Hill, New York, USA.
- 577 13] **Singh, P. and Narayanan, S.S. (2000)**. Biometrical Techniques in Plant Breeding. Kalayani Publishers,  
 578 New Delhi, India.
- 579 14] **Al-Naggar, A.M.M.; Abdalla, A.M.A.; Gohar, A.M.A. and Hafez, E.H.M. (2016a)**. Tolerance of 254  
 580 maize doubled haploid lines × tester crosses to drought at flowering and grain filling. *J. Appl. Life Sci.*  
 581 *Inter.*, 9(4): 1-18.
- 582 15] **Al-Naggar A. M. M., M. M. Shafik and M. O. Elsheikh (2018)**. Correlations and heritability for root  
 583 architecture traits of maize under water stress at flowering and grain filling. *Bioscience Research*, 15(4):  
 584 4571-4583.

Formatted: Indent: Left: -0.3"

Formatted: Portuguese (Brazil)

Formatted: Indent: Left: -0.3", Space After:  
0 pt

Formatted: Indent: Left: -0.3"

585 16] **Al-Naggar, A.M.M.; Shabana, R. Mahmoud, A.A.; Abdel El-Azeem, M.E.M. and Shaboon S.A.M.**  
586 (2009). Recurrent selection for drought tolerance improves maize productivity under low-N conditions.  
587 Egypt. J. Plant Breed., 13: 53-70.

588 17] **Al-Naggar, A.M.M.; Soliman, S. M and M. N. Hashimi (2011).** Tolerance to drought at flowering  
589 stage of 28 maize hybrids and populations. Egypt. J. Plant Breed., 15(1): 69-87.

590 18] **Al-Naggar, A.M.M.; Atta, M.M.M.; Ahmed, M.A. and Younis, A.S.M. (2016b).** Influence of deficit  
591 irrigation at silking stage and genotype on maize (*Zea mays* L.) agronomic and yield characters. J. Agric.  
592 and Ecol. Res. Inter., 7(4): 1-16.

593 19] **Al-Naggar, A.M.M., Shabana R., Hassanein M.S. and Metwally A.M.A. (2017a).** Effects of genotype,  
594 plant density and their interaction on maize yield and traits related to plant density tolerance. Bioscience  
595 Research, 14 (2): 395- 407.

596 20] **Al-Naggar, A.M.M., Shabana R., Hassanein M.S., Elewa T. A., Younis A.S.M. and Metwally A.M.A.**  
597 (2017b). Secondary Traits and Selection Environment of Plant Density Tolerance in Maize Inbreds and  
598 Testcrosses. Journal of Advances in Biology & Biotechnology 14(3): 1-13.

599 21] **Wright, G.C. and Nageswara Rao, R.C. (1994).** Groundnut Water Relations. In: Smartt J (Ed.) The  
600 groundnut crop: A Scientific Base for Improvement. Chapman and Hall, London, UK., 281-325.

601 22] **Henry, A.; Gowda, V.R.P. and Torres, R.O. (2011).** Variation in root system architecture and drought  
602 response in rice (*Oryza sativa*): phenotyping of the OryzaSNP panel in rainfed lowland fields. Field  
603 Crops Res., 120: 205-214.

604 23] **Hund, A.; Trachsel, S. and Stamp, P. (2009b).** Growth of axile and lateral roots of maize: I  
605 Development of a phenotyping platform. Plant and Soil, 325(1): 335-349.

606 24] **Hund, A. (2010).** Genetic variation in the gravitropic response of maize roots to low temperatures. Plant  
607 Root, 4: 22-30.

608 25] **Sattelmacher, B.; Horst, W.J. and Becker H.C. (1994).** Factors that contribute to genetic variation for  
609 nutrient efficiency of crop plants. Zeitschrift für Pflanzenernährung und Bodenkunde, 157: 215-224.

610 26] **Worku, M.; Banziger, M.; Erley, G.S.A.; Alpha, D.F.; Diallo, O. and Horst, W.J. (2007).** Nitrogen  
611 uptake and utilization in contrasting nitrogen efficient tropical maize hybrids. Crop Sci., 47: 519-528.

612 27] **Al-Naggar, A.M.M., Shabana R.A., Atta M.M.M. and Al-Khalil T.H. (2015).** Maize response to  
613 elevated plant density combined with lowered N-fertilizer rate is genotype-dependent. The Crop Journal 3  
614 (2): 96-109.

615 28] **Fageria, N.K. and Baligar, V.C. (1994).** Screening crop genotypes for mineral stresses. In: Adaptation  
616 of Plants to Soil Stress, (Eds. Maranville, J., W. Baligar, V. C., Duncan, R. R. and Yohe, J. M.),  
617 Nebraska-Lincoln Press, Inc, United states, NE. 152-159.

618 29] **Fageria, N.K. and Baligar, V.C. (1997a).** Phosphorous-use efficiency by corn genotypes. J. Plant Nutr.,  
619 20: 1267-1277.

620 30] **Fageria, N.K. and Baligar, V.C. (1997b).** Integrated plant nutrient management for sustainable crop  
621 production-An over. Inter. J. Trop. Agri., 15: 7-18.

622 31] **Shao Hui, Dongfeng Shi, Wenjun Shi, Xiangben Ban, Yachao Chen, Wei Ren, Fanjun Chen and**  
623 **Guohua Mi (2019).** Genotypic difference in the plasticity of root system architecture of field-grown  
624 maize in response to plant density. Plant Soil pp 1-17. doi.org/10.1007/s11104-019-03964-8

625 |  
626 |  
627 |  
628 |

UNDER PEER REVIEW

