

# Physiological and Biochemical Evaluation of Fe-Efficiency in Fe-Deficient Maize Genotypes

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**Abstract:** Iron (Fe) deficiency is prevalent particularly in calcareous soils of arid and semiarid regions. The present study aimed to investigate the response of eight local maize genotypes to Fe deficiency. In addition, a Fe-efficient (WF9) and a Fe-inefficient (ys1) indicator genotypes were used in this study. All genotypes were grown in growth chamber for 21 days in two levels of Fe, sufficient [+Fe (FNS) = 20  $\mu$ M Fe EDTA] and deficient [-Fe (FNS) = 2  $\mu$ M Fe EDTA] Fe supply. Shoots dry weight, Fe concentration and uptake in shoots, active iron concentration, peroxidase activity and chlorophyll content were determined and their validities as screening parameters were discussed. Generally, genotype (WF9) as the Fe efficient indicator and genotype (34) were the less affected by Fe-deficiency. Genotype (ys1) as the Fe inefficient indicator and genotype (62) were highly affected when grown in the nutrient solution with the deficient Fe supply. The present study emphasize that shoot dry weight, Fe uptake, active Fe content, chlorophyll content and peroxidase activity could be used for evaluating the present maize genotypes for Fe efficiency.

**Keywords:** Maize Genotypes, Fe Efficiency

## Introduction

Fe deficiency can reduce maize grain yield by as much as 20% (Godsey *et al.*, 2003). Correcting Fe deficiency through soil amendments and foliar sprays of Fe has been partially successful considering the fact that low availability rather than low total amount usually limits Fe uptake by plant roots. Furthermore, Mengel (1994) suggested that Fe availability in calcareous soils is not the critical process leading to chlorosis but rather the Fe uptake from the root apoplast into the cytosol of root cells turned to be the limiting factor. On the other hand, many food systems in developing countries cannot provide sufficient micronutrient content to meet the demands of their citizens, especially low-income families, because of low content of micronutrients in food is mostly related with low uptake.

There are several solutions such as soil and foliar fertilization, crop systems, application of organic amendments to correct micronutrients deficiency and to increase their density in edible parts of plants. Considering ecological concerns, cultivation and breeding of micronutrient-efficient genotypes in

combination with proper agronomic management practices appear as the most sustainable and cost-effective solution for alleviating food-chain micronutrient deficiency. Micronutrient-efficient genotypes could provide a number of benefits such as reductions in the use of fertilizers, improvements in seedling vigor and resistance to biotic and abiotic stresses. Using bioavailable micronutrient-dense staple crop cultivars can also be used to improve the micronutrient nutritional status of human (Khoshgoftarmanesh *et al.*, 2010).

Genotypic variation for Fe efficiency and resistance to Fe Deficiency Chlorosis (IDC) has been widely reported in maize and many plant species. Thus, potential exists for improving maize plants for Fe efficiency by breeding. The first step in a breeding program is screening and selection of the genetic material. Selection for Fe efficiency and IDC-resistance is made difficult by soil heterogeneity and highly variable environmental conditions that affect expression of the trait in the field. A simple, low cost and reliable screening procedure is required for breeding genotypes for Fe efficiency and IDC-resistance. The existence of

indicator genotypes for Fe efficiency are of significant importance for screening purposes, as they provide a scale of measurement (the hypothetical distance between the efficient and inefficient genotypes) and help to predict the relative position of a certain genotype on the scale (El-Bendary and Roemheld, 1998).

In this study, changes in Fe concentration, Fe uptake, active Fe content, chlorophyll content and peroxidase activity were examined in order to better understand the response triggered by iron starvation.

## Materials and Methods

### Plant Materials

The seed samples of the Fe-efficient (+/+) maize cv. WF9 and the Fe-inefficient (*ys1/ys1*) mutant were kindly provided by Prof. Marty Sachs, Maize Genetics Cooperation Stock Center, University of Illinois, USA. In addition, eight inbred lines used in maize breeding programs in Egypt, namely: 34, 628, 602, 62, 104, 653, 639 and 7 were kindly provided by Prof. Mohamad Abd El Satar, Faculty of Agriculture, Alexandria University, Egypt.

### Plant Culture

Seeds were surfaced-sterilized in 10% Clorox for 10 min, rinsed thoroughly in tap water and soaked for 3 to 4h (Mansfield and Key, 1987). The germination was carried out in sand culture at 28°C in dark. After five-days germination seedlings of each genotype were transferred to 1 L pots (5 seedlings/pot), containing a continuously aerated nutrient solution (Hoagland and Arnon, 1950) of the following composition in (mM): Ca(NO<sub>3</sub>)<sub>2</sub>.4H<sub>2</sub>O, (5); KNO<sub>3</sub>, (5); KH<sub>2</sub>PO<sub>4</sub>, (1); MgSO<sub>4</sub>.7H<sub>2</sub>O, (2); micronutrients (in µM): H<sub>3</sub>BO<sub>3</sub>, (10); MnCl<sub>2</sub>, (0.5); ZnSO<sub>4</sub>, (0.5); CuSO<sub>4</sub>, (0.2); Na<sub>2</sub>MoO<sub>4</sub>, (0.1). Fe was supplied as Fe(III)-EDTA (20 µM). After four days of growing in the nutrient solution with the sufficient level of Fe (20 µM), treatment was started by growing genotypes in two levels of Fe: Deficient-Fe supply (-Fe = 2 µM Fe EDTA) or sufficient-Fe supply (+Fe = 20 µM Fe EDTA). All plants were grown in a growth chamber for 21 days 60% relative humidity, light intensity of 120 µmol m<sup>-2</sup> sec<sup>-1</sup> at plant height and (25/20°C) day night regime. The nutrient solution was changed at two days interval.

### Determinations

Shoot samples of 21 days old seedlings were immersed several times in distilled water then oven dried at 105°C, weighed, ground and digested with acids mixture. Fe concentration in shoots was determined using Perkin-Elmer atomic absorption spectrophotometer (Chapman and Pratt, 1978). Active iron (Fe<sup>+2</sup>) concentration in fresh leaves was determined according to Takkar and Kaur (1984). Chlorophyll (chlorophyll a and b) was extracted from fresh leaf tissue with 80%

acetone, then the absorbency was determined at 645 and 663 nm with LKB spectrophotometer and chlorophyll concentration (mg/g f.wt.) were calculated according to the equations of Lichtenthaler (1987). Peroxidase enzyme activity was assayed spectrophotometrically according to Amako *et al.* (1994).

### Statistical Analysis

The experiment was conducted in a completely randomized design with three replicates. Data obtained from samples with the same experimental conditions were statistically analyzed using MSTAT-C Package program (Russell, 1986). When variance generated significant F-value ( $p < 0.05$ ), the mean values of the maize genotypes between and within the two levels of Fe were compared by the Least Significant Difference test (LSD) at the 0.05 level of probability as illustrated by (Little and Hills, 1978).

## Results and Discussion

### Shoots Dry Weight

Shoots (stems and leaves) dry weight of the present maize genotypes was used as a preliminary evaluation for the response to Fe deficiency (Table 1). The results revealed that shoots dry weight of the ten maize genotypes responded differently to iron treatment. Shoot dry weight of the Fe-efficient standard genotype (WF9), in addition to the genotypes (34 and 628) were less affected by Fe-deficiency in the culture media. In contrast, shoots dry weight of the Fe-inefficient yellow stripe (*ys1*) standard genotype in addition to the genotypes (104, 602, 62, 653, 639 and 7) were negatively affected when grown in the Fe-deficient nutrient culture. The percentage relative reduction values of shoot dry weight as an indicator of Fe-efficiency of the present maize genotypes revealed, as expected, that the standard Fe-efficient maize genotype (WF9) was the less affected by the absence of Fe in the nutrient culture. On the other hand, the standard Fe-inefficient yellow stripe (*ys1*) maize genotype was the most highly affected among the present ten maize genotypes. Concerning the response of shoots dry weight to Fe treatment; it seems that genotypes (628, 639, 34 and 602) had similar response to Fe-deficiency as the Fe-efficient standard genotype (WF9). On the other hand, it could be observed that the genotype (62) had similar response to Fe-deficiency as the standard Fe-inefficient (*ys1*) genotype. Between these categories, the genotypes (7 and 653) could be placed as intermediate concerning the response to Fe-deficiency.

El-Bendary and Salama (1998) reported that the increase of Fe-concentration in Hoagland solution resulted in a significant increase in shoot dry weights in varieties of faba bean. Similar results were reported by Krishnasamy *et al.* (2005) on sorghum genotypes and Salama *et al.* (2009) on maize.

Table 1. Shoots (stems + leaves) dry weight (g/plant) and calculated relative reduction (RD%) of shoots dry weight of the ten maize genotypes grown in nutrient solution for three weeks in sufficient [+Fe (FNS) = 20  $\mu$ M Fe EDTA] and deficient [-Fe (FNS) = 2  $\mu$ M Fe EDTA] Fe supply

Genotypes	Shoot dry weight (g/plant)		RD%**
	+Fe(FNS)*	-Fe(FNS)*	
34	0.07	0.06	-14
628	0.08	0.07	-13
602	0.11	0.09	-18
62	0.07	0.04	-43
104	0.09	0.06	-33
653	0.09	0.06	-33
639	0.15	0.13	-13
7	0.07	0.05	-29
WF9	0.11	0.10	-9
ys1	0.07	0.04	-43
L.S.D (0.05)1	0.02	0.02	
L.S.D (0.05)2	0.02		

1. L.S.D between genotype means at the same level of Fe.

2. L.S.D for the same genotype at different levels of Fe.

\*FNS: Full Nutrient Solution

$$**\text{Relative Reduction (RD) \%} = (-100) \frac{(+Fe) - (-Fe)}{(+Fe)}$$

Table 2. Fe-concentration ( $\mu$ g/g DW) and calculated relative reduction (RD%) in shoots (stems + leaves) of the ten maize genotypes grown in nutrient solution for three weeks in sufficient [+Fe (FNS) = 20  $\mu$ M Fe EDTA] and deficient [-Fe (FNS) = 2  $\mu$ M Fe EDTA] Fe supply

Genotypes	Iron concentration ( $\mu$ g /g DW)		RD%
	+Fe(FNS)*	-Fe(FNS)*	
34	180	172	-4.4
628	146	127	-13.0
602	163	154	-5.5
62	93	54	-42.0
104	101	97	-4.0
653	160	142	-11.3
639	99	43	-56.6
7	156	149	-4.5
WF9	162	151	-6.8
ys <sub>1</sub>	68	36	-47.0
L.S.D (0.05)1	17	17	
L.S.D (0.05)2	16		

1. L.S.D between genotypes means at the same level of Fe

2. L.S.D for the same genotypes at different levels of Fe

\*FNS: Full Nutrient Solution

$$**\text{Relative Reduction (RD) \%} = (-100) \frac{(+Fe) - (-Fe)}{(+Fe)}$$

Krishnasamy *et al.* (2005) screened sorghum genotypes and they stated that Fe-efficient genotype should not only be able to absorb more Fe from deficient soils but should also produce more dry matter and grain yield. Celik and Katkat (2008) found the root and shoot dry weight values of maize varieties grown in iron free nutrient solution were severely

affected and gave the lowest dry weight values. Similar results were observed by Jelali *et al.* (2012).

### Fe-Concentration

Significant differences were observed among the present maize genotypes for Fe concentration in shoots either at the same level of Fe or at the different levels of Fe-treatment (Table 2). At Fe supply (+Fe), the highest Fe-concentrations in shoots were detected in genotypes (34, 602 and WF9), while the lowest Fe-concentration in shoots were detected in ys1 genotype. At deficient Fe supply (-Fe), the highest Fe-concentrations in shoots were observed in genotypes (34, 602, WF9), while the lowest Fe-concentrations in shoots were observed in genotypes (ys1 and 639). On the other hand, different response to deficient Fe supply was observed in Fe-concentrations when compared with the same genotypes in Fe treatment. Fe-concentrations in shoots of (ys1, 639, 653, 62 and 628) genotypes grown in deficient Fe-supply were negatively affected when compared with the same genotypes grown in sufficient Fe-supply. No significant differences were observed for Fe-concentration in shoots of (WF9, 7, 34, 602 and 104) genotypes for both treatments.

Although these data may indicate that Fe-concentration in shoots dry weight of the present maize genotypes may be an indicator for Fe-efficiency, many previous results were in disagreement with this conclusion. Confirmed results from maize and other cereal species indicated that the concentration of Fe, among other micronutrients, is not a reliable parameter for distinguishing sensitivity to Fe-deficiency among plant genotypes. Wallace *et al.* (1976a) and Katyal and Sharma (1980) concluded that Fe concentration techniques for diagnosing Fe deficiency in plant are generally considered unsatisfactory, because total Fe concentration in plant does not correlate with plant growth response to Fe. El-Bendary and Roemheld (1998) reported that Fe-concentration in shoot was not a suitable parameter for evaluating Fe-efficiency and it was not related to Fe-efficiency. El-Bendary *et al.* (1999) found that total Fe concentration in leaves is not a suitable indicator for assessing chlorosis expression in plants. Pestana *et al.* (2003) found that in the case of the Fe-deficiency, sometimes, the total iron content might not reflect the iron nutritional status of the plant. It has been established that when plants are grown under Fe-deficiency in field conditions, the total leaf Fe concentration is generally the same or even higher than in Fe-sufficient plants. Nenova and Stoyanov (1999) found that Fe-deficiency resulted in rapid and strong decrease of root Fe concentration. Despite the low total Fe in the leaves of chlorotic plants, the Fe concentration in leaves did not reflect the Fe supply to plants.

### Iron Uptake

Shoots content of the present genotypes showed that the Fe efficient standard genotype (WF9) and genotype (34) were less affected by Fe deficiency in the nutrient solution (Table 3). In contrast, shoots uptake of the Fe inefficient yellow stripe (ys1) standard genotype, in addition to genotypes (62 and 639) were highly affected when grown in the Fe deficient nutrient culture. These results are confirmed by the Fe efficiency scale expressed as the percentage relative reduction values of shoots uptake of the present genotypes. Moreover, a clear correlation was observed between Fe-deficiency symptoms and Fe-uptake pattern of the present genotypes. WF9, 34, 602 and 628 with slight deficiency symptoms contained more iron in shoots than other genotypes with severe deficiency symptoms. In addition, the present results revealed that among the Fe deficient maize genotypes, ys1 and 62 exhibited the highest deficiency symptoms correlated with lowest iron content in shoots.

The different uptake pattern of iron for the present genotypes is in agreement with the findings of (Clark and Brown, 1974; EL-Bendary *et al.*, 1999) and others; they found that maize genotypes grown in calcareous soil showed wide differences in Fe uptake and utilization. Moreover, the different uptake pattern of iron of the present maize genotypes is in agreement with the fact that some plant species and cultivars of the same species have evolved a mechanism for more efficient uptake under deficiency of some elements (El-Bendary *et al.*, 1998; Cakmak *et al.*, 1999). El-Bendary and Roemheld (1998) suggested that shoot uptake is suitable for selecting the most inefficient lines of maize.

### Active Iron Concentration

Data presented in Table 4 showed that active iron content in the leaves of (34 and WF9) genotypes was relatively higher under both Fe treatments. When plants grown with adequate Fe supply in the growth media, the active iron content in leaves was (41.3 and 36.3 µg/g F.W) respectively, while under Fe deficiency the active iron content reached (30.11 and 27.60 µg/g F.W), at 21 days old. For ys1, 62 and 639 genotypes, the active iron content in the Fe deficient leaves of ys1, 62 and 639 was (17.4 µg/g F.W, 16.1 µg/g F.W and 13.5 µg/g F.W) respectively. Under Fe deficiency treatment the active iron decreased in ys1, 62 and 639 genotypes by 47.4, 38.6 and 37.6%, respectively. For 7, 653, 602, 34, 628, WF9 and 104 genotypes, the active iron content in the leaves was decreased by (29.6, 28.9, 27.9, 27.1, 26.3, 24 and 22%).

These results are in a good agreement with the results of Mehrotra and Gupta (1990), they concluded that the high level of active iron detected in resistant chickpea cultivar might be the source of the ability to tolerate iron deficiency. Similar results were obtained by Ohwaki and Sugahara (1993).

Table 3. Iron uptake (µg/plant) in shoots (stems + leaves) and calculated relative uptake reduction (RD %) of the ten maize genotypes grown in nutrient solution for three weeks in sufficient [+Fe (FNS) = 20 µM Fe EDTA] and deficient [-Fe (FNS) = 2 µM Fe EDTA] Fe supply

Genotypes	Iron uptake (µg/plant)		
	+Fe(FNS)*	-Fe(FNS)*	RD%**
34	12.6	10.3	-18
628	11.7	8.9	-24
602	18.0	13.8	-23
62	6.5	2.2	-66
104	9.1	5.8	-36
653	14.4	8.5	-41
639	14.8	5.6	-62
7	10.9	7.4	-32
WF9	17.8	15.0	-16
ys1	4.7	1.4	-70
L.S.D (0.05)1	2.7	2.7	
L.S.D (0.05)2	2.9		

1. L.S.D between genotype means at the same level of Fe

2. L.S.D for the same genotype at different levels of Fe

\* FNS: Full Nutrient Solution

$$** \text{Relative Reduction (RD) \%} = (-100) \frac{(+Fe) - (-Fe)}{(+Fe)}$$

Table 4. Active iron (Fe<sup>2+</sup>) concentration (µg/g F.W) in seedling leaves of the ten maize genotypes grown in nutrient solution for three weeks in sufficient [+Fe (FNS) = 20 µM Fe EDTA] and deficient [-Fe (FNS) = 2 µM Fe EDTA] Fe supply

Genotypes	Active iron (Fe <sup>2+</sup> ) µg/g F.W		
	+Fe(FNS)*	-Fe(FNS)*	RD%**
34	41.3	30.1	-27.1
628	28.9	21.3	-26.3
602	32.3	23.3	-27.9
62	25.8	16.1	-37.6
104	28.8	22.2	-22.9
653	30.4	21.6	-28.9
639	22.0	13.5	-38.6
7	30.7	21.6	-29.6
WF9	36.3	27.6	-24.0
ys1	33.1	17.4	-47.4
L.S.D (0.05)1	3.9	3.9	
L.S.D (0.05)2	3.9		

1. L.S.D between genotype means at the same level of Fe

2. L.S.D for the same genotype at different levels of Fe

\*FNS: Full Nutrient Solution

$$** \text{Relative Reduction (RD) \%} = (-100) \frac{(+Fe) - (-Fe)}{(+Fe)}$$

They reported that the genotypic differences of sensitive and resistant cultivars of chickpea to Fe-deficiency were attributed to the active iron in the leaves when grown under Fe-stress. Krishnasamy *et al.* (2005) indicated that the active Fe content of sorghum leaf blades was decreased with increasing level of chlorosis. Moreover, Ramirez *et al.* (2002) stated that active iron content may

be used as a reliable tool for diagnosis of Fe efficiency for rice. Mohamed *et al.* (2003) concluded that total Fe cannot be used as a criteria to differentiate between the Fe-deficient and non-deficient plants.

### Chlorophyll Content

The mean values of total chlorophyll content of the ten maize genotypes in the presence and absence of iron supply are shown in Table 5. After 21 days, the data showed that the total chlorophyll content differed in all genotypes for both treatments. Under Fe deficiency treatment the total chlorophyll decreased in ys1, 62 and 639 genotypes by 72.1, 65.4 and 64.5%, respectively. The decrease was markedly observed in ys1 genotype, more than (72%) reduction compared with WF9 (42.9% reduction).

Chlorosis of young leaves is often the first visual sign of iron deficiency. It is associated not only with loss of chlorophyll, as several steps of its biosynthesis depend on Fe, but also with changes in the expression and assembly of other components of the photosynthetic apparatus (Terry and Abadía, 1986). Early chlorosis caused decreased photosynthetic rate, less photosynthates production, nitrogen fixation and consequently yield losses (Singh and Sahu, 1993). Iron deficient plants showed visible symptoms on their youngest leaves, which became yellow (chlorotic) due to a decrease in chlorophyll content and had lower net photosynthetic rate (Price, 1968; Terry, 1980; Misra and Srivastava, 1994; Abadia *et al.*, 2000). Gogorcena *et al.* (2001) found that iron deficiency caused moderate decreases (about 8%) in the dark-adapted efficiency of PSII. Zhang *et al.* (2012) found that Fe-deficiency induced chlorosis of plants growing on calcareous soil and Fe-deficiency slightly reduced the chlorophyll ratio. Earlier, it was suggested that the determination of leaf chlorophyll content can be a diagnostic tool to quantify Fe chlorosis (Abadia, 1992). On the other hand, El-Baz *et al.* (1998) found that chl. b showed no response to different Fe supply in snap and faba beans.

### Peroxidase Activity (POD)

Data presented in Table 6 showed the changes in peroxidase activity in leaves of the ten maize genotypes grown in nutrient solution for three weeks in sufficient and deficient Fe supply. The results revealed that the activity of peroxidase in leaves of the present genotypes was depressed markedly in Fe deficient treatment when compared with sufficient Fe treatment. Supplying plants with sufficient iron +Fe (FNS) induced significant increase in POD activity for all maize genotypes as compared with deficient iron -Fe (FNS) treatment.

Table 5. Total chlorophyll (mg/g F.W) content and calculated relative Reduction (RD%) of the ten maize genotypes grown in nutrient solution for three weeks sufficient [+Fe (FNS) = 20 µM Fe EDTA] and deficient [-Fe (FNS) = 2 µM Fe EDTA] Fe supply

Genotypes	Total Chlorophyll (mg/g F.W)		
	+ Fe(FNS)*	- Fe(FNS)*	RD%**
34	1.6	0.87	-44.6
628	1.4	0.68	-51.1
602	2.2	1.23	-44.8
62	0.8	0.29	-65.4
104	1.7	0.76	-55.2
653	1.2	0.54	-54.8
639	0.7	0.26	-64.5
7	1.5	0.74	-49.2
WF9	1.7	0.99	-42.9
ys1	1.4	0.39	-72.1
L.S.D (0.05)1	0.1	0.10	
L.S.D (0.05)2	0.2		

1. L.S.D between genotype means at the same level of Fe  
 2. L.S.D for the same genotype at different levels of Fe  
 \*FNS: Full Nutrient Solution

$$**Relative\ Reduction\ (RD)\ \% = (-100) \frac{(+Fe) - (-Fe)}{(+Fe)}$$

Table 6. Peroxidase (POD) activity (U/g F.W/min) in the leaves of the ten maize genotypes grown in nutrient solution for three weeks in sufficient [+Fe (FNS) = 20 µM Fe EDTA] and deficient [-Fe (FNS) = 2 µM Fe EDTA] Fe supply

Genotypes	POD activity (U/g F.W/min)		
	+ Fe(FNS)*	- Fe(FNS)*	RD%**
34	12.8	10.3	-19.5
628	12.6	9.0	-28.6
602	15.6	10.4	-33.3
62	10.6	5.3	-50.0
104	14.5	10.9	-24.8
653	11.4	8.4	-26.3
639	5.7	3.3	-42.1
7	12.3	7.7	-37.4
WF9	15.5	14.2	-8.4
ys1	9.2	4.3	-53.3
L.S.D (0.05)1	1.9	1.9	
L.S.D (0.05)2	1.9		

1. L.S.D between genotype means at the same level of Fe  
 2. L.S.D for the same genotype at different levels of Fe  
 \*FNS: Full Nutrient Solution

$$**Relative\ Reduction\ (RD)\ \% = (-100) \frac{(+Fe) - (-Fe)}{(+Fe)}$$

WF9 genotype showed the highest value for (POD) activity under both treatments of iron. In plants grown with adequate Fe supply in the growth media, the POD activity in leaves of WF9 genotype was (15.5 U/g F.W/min) and under iron deficient treatment was (14.20 U/g F.W/min) and the relative reduction was (8.40%). In contrast, POD activity in leaves of ys1 plants grown in adequate Fe supply was (9.2 U/g F.W/min), while under Fe deficient treatment the POD activity was (4.3 U/g F.W/min).

Table 7. Maize genotypes (G) arranged in ascending order according to percentage Reduction (RD%) due to Fe deficiency for the different plant parameters used to evaluate Fe efficient and inefficient genotypes

Shoot dry weight (g/plant)		Fe-concentration ( $\mu\text{g/g DW}$ )		Iron uptake ( $\mu\text{g/plant}$ )		Active iron ( $\text{Fe}^{2+}$ ) $\mu\text{g/g F.W}$		POD activity (U/g. F.W/min)		Total Chlorophyll (mg/g F.W)	
G	RD%	G	RD%	G	RD%	G	RD%	G	RD%	G	RD%
WF9	-9	104	-4	WF9	-16	104	-22.9	WF9	-8.4	WF9	-42.9
628	-13	34	-4.4	34	-18	WF9	-24.0	34	-19.5	34	-44.6
639	-13	7	-4.5	602	-23	628	-26.3	104	-24.8	602	-44.8
34	-14	602	-5.5	628	-24	34	-27.1	653	-26.3	7	-49.2
602	-18	WF9	-6.8	7	-32	602	-27.9	628	-28.6	628	-51.1
7	-29	653	-11.3	104	-36	653	-28.9	602	-33.3	653	-54.8
104	-33	628	-13	653	-41	7	-29.6	7	-37.4	104	-55.2
653	-33	62	-42	639	-62	62	-37.6	639	-42.1	639	-64.5
62	-43	ys <sub>1</sub>	-47	62	-66	639	-38.6	62	-50	62	-65.4
ys <sub>1</sub>	-43	639	-56.6	ys <sub>1</sub>	-70	ys <sub>1</sub>	-47.4	ys <sub>1</sub>	-53.3	ys <sub>1</sub>	-72.1

The relative reduction in POD activity was (53.3%). The magnitude inhibition in POD activity due to Fe deficiency was clearly observed in 639, 62 and ys<sub>1</sub> genotypes, the values were 3.3, 5.3 and 4.3 (U/g F.W/min) (more than 42, 50 and 53% depression). It is also interesting to note that POD activity in the fresh leaves of WF9 maize genotype, grown in either adequate or deficient Fe treatment, showed more or less the same value (15.5 and 14.2 U/g F.W/min). Genotype WF9 is known to be more efficient in the uptake and translocation of Fe from the roots into the leaves tissues.

Iron deficiency has been generally known to affect plant growth in several plant species (Rombolà *et al.*, 2005; Pestana *et al.*, 2005). As shown above in our results, significant differences in the pattern of plant growth were found to depend on the treatment and genotype. Furthermore, it was noticed that Fe deficiency caused reduced activity of POD enzyme. The reduction in POD activity was higher in the susceptible genotype (ys<sub>1</sub>) when compared with the non susceptible one (WF9). These results suggested that under Fe deficiency, the tolerant genotype tried to keep Fe-dependent PODs functioning, probably to counteract H<sub>2</sub>O<sub>2</sub> accumulation. In general, it can be suggested that Fe deficiency led to a drastic decrease in POD activity in the susceptible genotype (ys<sub>1</sub>) when compared with the tolerant one (WF9), indicating that the H<sub>2</sub>O<sub>2</sub> scavenging mechanism was less effective in the susceptible genotype (ys<sub>1</sub>). It turned out that the active involvement of this antioxidant enzyme was related, at least in part, to the tolerance to Fe-deficiency-induced oxidative stress.

Susceptibilities of the various maize genotypes were expressed as percentage reduction, from Fe sufficient level, due to Fe deficiency. These calculations were made for the different plant parameters used for screening the maize genotypes in the present study and arranged in ascending order according to the percentage reduction (RD%) values (Table 7). WF9, as the Fe efficient indicator and genotype (34) were the less affected genotypes and occupied the top of the table in four parameters out of six. On the other hand, the Fe inefficient indicator genotype (ys<sub>1</sub>) was the most

affected genotype and occupied the bottom of the table in five parameters out of six. Similar response to Fe deficiency was noticed for genotype (62), it occupied the bottom of the table in four parameters out of six.

## Conclusion

Although visual diagnostic symptoms are an extremely valuable tool for the rapid evaluation of the nutrient status of a plant, they are only some of the tools available. Other major tools include plants fresh and dry weights, tissue analysis, chlorophyll content, enzymes activity and others. These methods all vary in their precision, rapidity and their ability to predict future nutrient status. Because of the close interaction between plant growth and the environment, all predictions of future nutrient status must make assumptions about how the environment will change in that time frame. The other tools adopted in this study to evaluate the eight maize genotypes in addition to the Fe efficient (WF9) and the Fe inefficient (ys<sub>1</sub>) genotypes were shoot dry weight, Fe concentration, Fe uptake, active iron content, chlorophyll content and peroxidase activity. In general, the results indicated that the present maize genotypes showed different response when grown in Fe deficient media and could be classified according to the above mentioned parameters to efficient, moderate and inefficient. However, some genotypes showed different behavior concerning Fe efficiency when compared by different tools. For instance, genotype (639) was classified by shoot dry weight parameter as Fe efficient while it was classified as Fe inefficient by adopting Fe concentration as a parameter for evaluating the present maize genotypes for Fe efficiency.

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### Author's Contributions

All authors equally contributed in this work.

### Ethics

This article is original and contains unpublished material. The corresponding author confirms that all of the other authors have read and approved the manuscript and no ethical issues involved.

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