

Full Length Research Paper

The effect of aluminium on enzyme activities in two wheat cultivars

N. Nasr^{1*}, J. Carapetian¹, R. Heidari¹, S. Asri Rezaei², N. Abbaspour¹, R. Darvishzadeh^{3,4} and F. Ghezelbash¹

¹Department of Biology, Faculty of Science, Urmia University, Urmia, Iran.

²Faculty of Veterinary Medicine, Urmia University, Urmia, Iran.

³Institute of Biotechnology, Urmia University, Urmia, Iran.

⁴Department of Agronomy and Plant Breeding, Urmia University, Urmia, Iran.

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In order to study the effect of different aluminum (Al) concentrations on the enzyme activities of wheat seedlings and the effect of malate and citrate treatments as chelates for reducing the noxious effect of Al in medium culture, the seedlings of two wheat cultivars, Darab (Al-sensitive) and Maroon (Al-tolerant) were grown on hydroponic solution (non modified Hoagland solution) containing AlCl_3 (0-100-200-300 μM). Factorial experiment was realized in a complete randomized design with three replications. The activity of different enzymes such as ascorbate peroxidase (APX), catalase (CAT), glutathione reductase (GR) and superoxide dismutase (SOD) in root and shoot were measured. Analysis of variance revealed that, the activity of studied enzymes (APX, CAT, GR and SOD) in root and shoot were affected by the main effects of Al concentration. However, in the case of catalase activity in root, the main effect of genotypes as well as genotype \times Al concentration was also significant. APX activity in root was not significantly differed between Maroon and Darab in all Al concentrations. But by increasing Al concentration in root medium APX activity was significantly decreased. In the case of SOD activity, we did not find any difference between the studied genotypes in all Al concentrations but its content in roots was affected by the amount of Al applied in medium. So that by increasing the amount of Al, SOD content increased in the genotypes similarly. The same trend was observed for catalase activity in root. In the case of GR activity, we did not find any difference between the genotypes in all Al concentrations but its content in root was affected by applying Al in medium compared with control medium, so that GR content increased in both genotypes similarly. The activity of investigated enzymes showed the same trend in the shoot. The effect of malate and citrate was also studied on reducing the noxious effect of Al in root. Analysis of variance revealed that, there were significant differences within the treatments on the enzymes activity in root (not in the shoot) except for catalase. However, some interaction effects were significant. This means that malate or citrate application was effective in some Al concentrations.

Key words: Maroon (Al tolerant), Darab (Al sensitive), ascorbate peroxidase (APX), catalase (CAT), glutathione reductase (GR), superoxide dismutase (SOD).

INTRODUCTION

Aluminum (Al) toxicity is a major factor limiting crop

productivity in many acid soils throughout the tropics and subtropics. At mildly acidic or neutral soil pH values, it occurs primarily as insoluble deposits and essentially is biologically inactive. The acidity of the soil gradually increases as a result of the effects of environmental factors, especially acid rain. Acid soils occupy 30 to 40%

*Corresponding author. E-mail: nasr_na2002@yahoo.com. Tel: + 98 9143885342. Fax: +4412776707.

of the arable lands (von Uexküll and Mutert, 1995) and as a consequence of a rapid industrial development and environmental pollution, this area increases from year to year especially in developing countries.

Al stress induces the expression of a number of genes; over 20 of them have already been isolated and characterized in wheat (Snowden et al., 1995; Cruz-Ortega et al., 1997; Hamilton et al., 2001), tobacco (Ezaki et al., 1995; 1996) and Arabidopsis (Richards et al., 1998) and their hypothetical functions in Al toxicity or Al resistance mechanisms have been proposed. Since some of these Al-induced genes are coding for antioxidant enzymes (glutathione-S-transferase, peroxidase, superoxide dismutase, ascorbate peroxidase and catalase), it has been suggested that, there is a strong correlation between Al stress and oxidative stress in plants (Cakmak and Horst, 1991; Richards et al., 1998; Simonovicova et al., 2004). A common feature of several stresses (including Al toxicity) is enhanced production of active oxygen species (AOS), which are generally considered harmful to plant cells (Richards et al., 1998; Tamas et al., 2004). However, it has been recently recognized that hydrogen peroxide plays a central role in several physiological processes, such as defense reaction, stomatal closure, programmed cell death, peroxisome biogenesis, cell wall cross-linking and lignin synthesis (Van Breusegem et al., 2001; Neill et al., 2002). More recently, hydrogen peroxide was also identified as an expression of the signal molecule of several genes in plants (Desikan et al., 2000). The key role of antioxidant enzymes is to reduce or scavenge reactive oxygen species, such as superoxide radicals, singlet oxygen, hydrogen peroxide and hydroxyl radicals. The resulting hydrogen peroxide is removed through the activity of the Asada-Halliwell scavenging cycle in chloroplasts and in cytoplasm by ascorbate peroxidase (APX) (Asada, 1992).

Peroxidases (POD) participate in lignin biosynthesis, cell wall cross-linkage, IAA degradation and disease resistance and convert hydrogen peroxide to water (Asada, 1992; Siegel, 1993). The objective of the present study was to assess the role of antioxidant enzymes, including ascorbate peroxidase (APX), superoxide dismutase (SOD), and catalase (CAT) and glutathione reductase (GR) enzymes as possible mechanisms for Al stress adaptation in wheat.

MATERIALS AND METHODS

Plant materials and experimental design

The seeds of two wheat cultivars, Darab (Al sensitive) and Maroon (Al tolerant) were provided from Agricultural Research Center of Karaj. The seeds of two cultivars were sterilized with 5% (v/v) sodium hypochlorite for 15 min then, were rinsed with distilled H₂O for 15 min and were kept in the dark for 24 h at 25°C. Germinated seeds were placed on a plastic net, which was floated on a continuously aerated solution containing 0.5 mM CaCl₂. The seedlings were kept in the dark for 1 day at 25°C and then were

moved to natural light conditions. Solution was renewed daily and seedlings were selected for treatment by measuring uniform root length. Pre-culture solution were replaced by hydroponic solution (non modified Hoagland) containing AlCl₃ (0-100-200-300 µM) and pH was kept constant at 4. Factorial experiment was realized in a complete randomized design with three replications. Each replication consisted of one Petri dish of ten seedlings per cultivar and AlCl₃ combinations. Treatment solutions were renewed every 3 days with fresh solution (Zakir et al., 2005). The plants were grown for 15 days under a 16 h photoperiod. Then, 15 days old plants were used for the experiments which in citrate and malate were used as phytochelator for decreasing the effect of aluminum toxicity.

Chlorophyll measurement

Chlorophyll content of leaves was determined according to Wintermans and De Mots (1965). Briefly, after extraction in 96% (v/v) ethanol, the absorption was measured in different wavelengths depend on the type of chlorophyll (Kuo and Kao, 2003).

Determination of enzymes activities

For extraction of enzymes, 0.5 g of fresh weight was homogenized with 0.1 M phosphate buffer (pH 6.8) in a chilled pestle. The homogenate was centrifuged at 8000×g for 20 min and the supernatant was used for determining of enzymes activities. The whole extraction procedure was carried out at 4°C. CAT activity was determined by measuring the initial rate of disappearance of H₂O₂ (Kato and Shimidzu, 1987). The decrease in H₂O₂ followed as the decline in absorbance at 240 nm. One unite of CAT was defined as the amount of enzyme which breaks down 1 nmol H₂O₂ per min. Superoxide dismutase (SOD) was determined according to Paoliet et al. (1986). One unit of SOD was defined as the amount of enzyme which inhibits the rate of NADH oxidation by 50% compared with blank. Ascorbate peroxidase (APX) activity was determined according to Nakano and Asada (1981). The decreased in ascorbate concentration was followed at 290 nm. One unit of APX was defined as the amount of enzyme which breaks down 1 µmol of ascorbate per min. Finally, glutathione reductase (GR) was defined as the amount of enzyme which decreases A₃₄₀ (1 u per min). All of the earlier mentioned experiments were repeated three times (Kuo and Kao, 2003).

Data analysis

Analysis of variance was performed using the general linear model (GLM) procedure in the SAS software (SAS Institute Inc., Cary, NC, USA). The main effects of genotype, Al concentration as well as their interaction were determined. To generate a trend analysis, the Proc REG procedure of PC-SAS is specified (SAS Institute Inc., Cary, NC, USA). Commands for each model are placed after the Proc Reg statement. A separate model statement was required for linear, quadratic and cubic trends.

RESULTS

Analysis of variance revealed that, the activity of enzymes such as ascorbate peroxidase (APX), catalase, glutathione reductase (GR) and superoxide dismutase (SOD) in root and shoot were affected by the main effect

Table 1. Analysis of variance summary for enzymes data under different Al concentrations. Data were analyzed using procedures for a completely randomized design.

Source	DF	Mean of square						
		APX-S	Catalase-S	GR-S	SOD-S	Chlorophyll A	Chlorophyll B	Caretenoid
Line	1	0.003 ^{ns}	21.08 ^{ns}	0.00003 ^{ns}	0.05 ^{ns}	4.78 ^{ns}	0.036 ^{ns}	2554.41 ^{ns}
Al concentration	3	175.53 ^{**}	2102.81 ^{**}	0.02 ^{**}	99.60 ^{**}	4.76 [*]	0.023 ^{ns}	58507.74 [*]
Line × Al concentration	3	6.70 ^{ns}	172.31 ^{ns}	0.0008 ^{ns}	4.12 ^{ns}	3.79 ^{ns}	0.035 [*]	48182.51 [*]
Error	16	4.88	75.07	0.0004	2.72	1.22	0.009	13328.22
C.V.		8.17	25.41	7.61	5.80	27.43	23.86	32.50

Table 1. Continued.

Source	DF	Mean of square			
		APX-R	Catalase-R	GR-R	SOD-R
Line	1	18.69 ^{ns}	6253.30 ^{**}	0.002 ^{ns}	10.51 ^{ns}
Al concentration	3	176.44 ^{**}	16370.70 ^{**}	0.02 ^{**}	99.25 ^{**}
Line × Al concentration	3	9.02 ^{ns}	1124.86 ^{**}	0.0009 ^{ns}	5.07 ^{ns}
Error	14	7.17	194.52	0.0006	4.04
C.V.		15.07	13.20	6.86	5.67

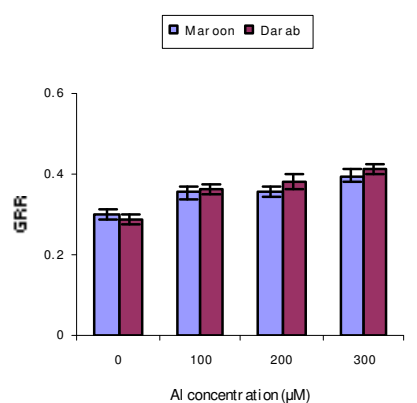
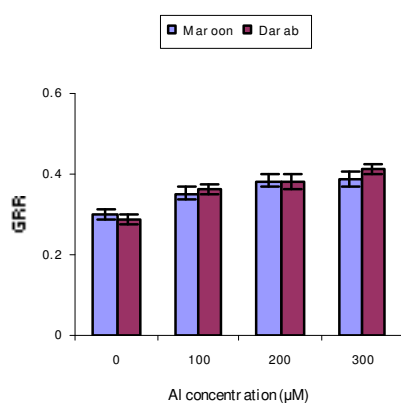
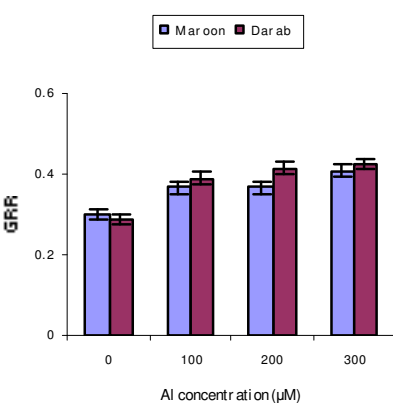
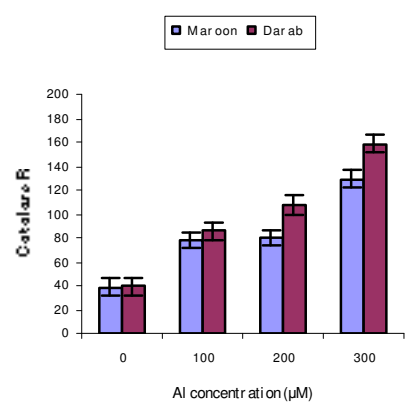
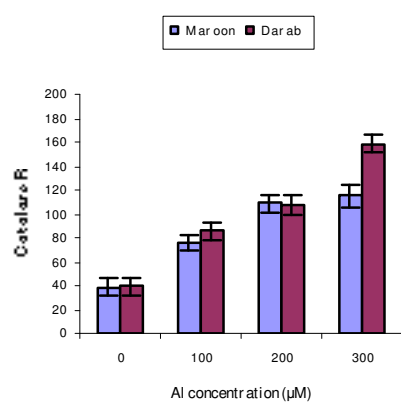
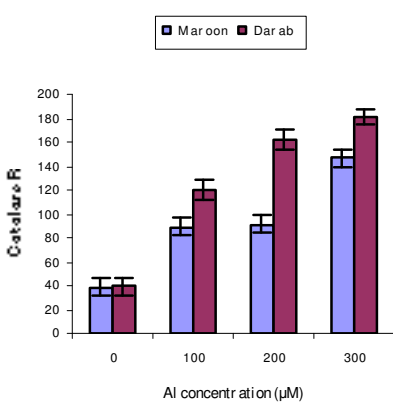
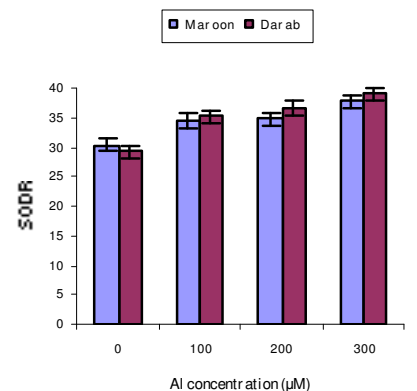
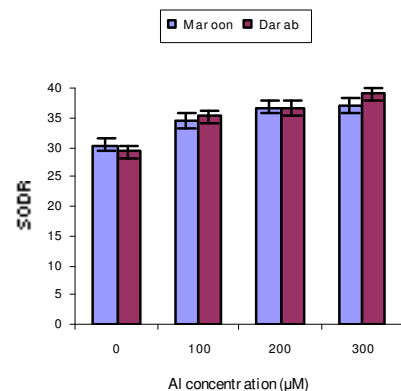
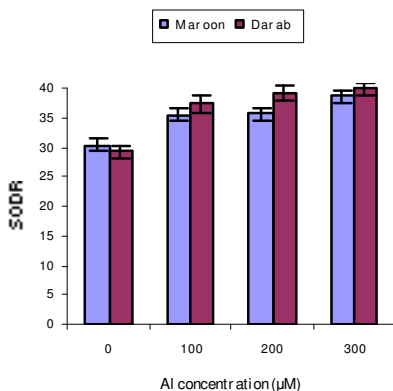
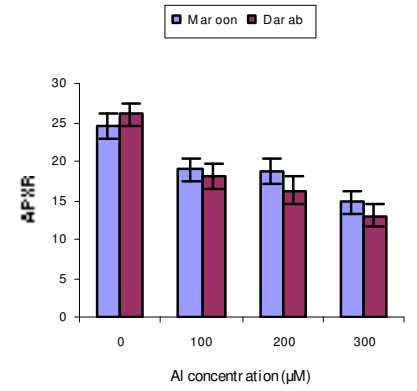
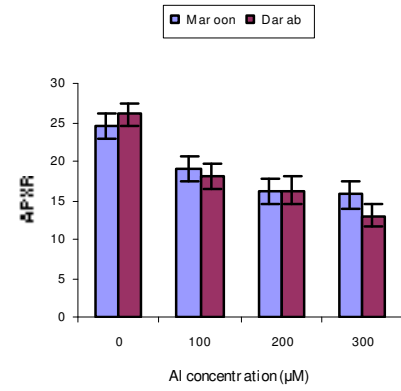
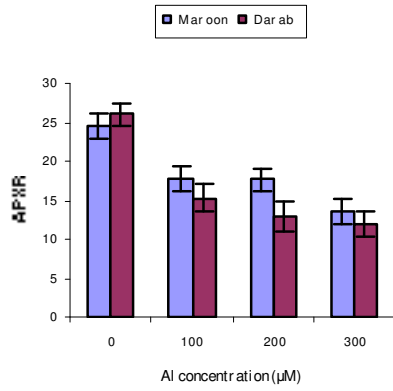
CV: Coefficient of variation. df = Degrees of freedom; **, *: Significant at 0.01 and 0.05 probability level; ns: non significant.

of Al concentration. Only concerning to catalase activity in root, the main effect of genotype as well as genotype × Al concentration was significant (Table 1). As shown in Figure 1, the APX activity in root was not significantly different between Maroon and Darab in all studied Al concentrations. However, by increasing the Al concentration in medium, APX activity in root was significantly decreased. In the case of SOD activity, there was no difference between both studied genotypes in all applied Al concentrations. However, its content was affected by the amount of Al concentration. Such that by increasing the amount of Al its content increased in the genotypes similarly. The same trend was observed for the catalase activity in root. As for superoxide dismutase, there was no difference between two studied genotypes in all applied Al concentrations from the view point of GR activity. But in comparison to control plants glutathione reductase activity in roots was affected by applying Al in medium and its content increased in two genotypes similarly. The activity of the investigated enzymes showed the same trend in the shoots.

A separate regression analysis considering Al concentration as an independent variable and the enzymes activity as a dependent variable was conducted for each genotype. The activity of APX in Maroon and Darab root best fit the cubic and linear model, respectively, as indicated by a significant T-value (Table 2). This means that, activity of APX in root of these genotypes followed

the different basic trend. R^2 values for Maroon and Darab were 0.99 and 0.61, respectively. This also means that, 99 and 61% of the variation was explained by the model. These values are high because R^2 values for biological data generally range from 0.50 to 0.90, whereas, a low R^2 for non-biological data may be 0.90 (Kleinbaum and Kupper, 1978). In contrast, analysis of APX activity in shoot using polynomial contrasts indicated that, the response of Maroon and Darab seedlings best fit the linear and cubic model. Concerning to other enzymes we also found a difference in fitted model between stem and root (Table 2).

We have also studied the effect of genotype and Al concentration on the chlorophyll-A, chlorophyll-B and caretenoid content. Analysis of variance for chlorophyll-content revealed that, only the main effect of Al concentration was significant. In the case of chlorophyll-B content, the interaction between genotype and Al concentration was significant. For caretenoid both the main effect of Al concentration and the interaction effect between genotype and Al concentration were significant. In the second experiment, the effect of citrate and malate treatments was studied on decreasing the noxious effect of Al in medium. Analysis of variance showed that, there were significant differences within applied treatments on studied enzymes activities in root not in the shoot except for catalase activity (Table 3). However, some interaction effects were significant. This means that, the effect of



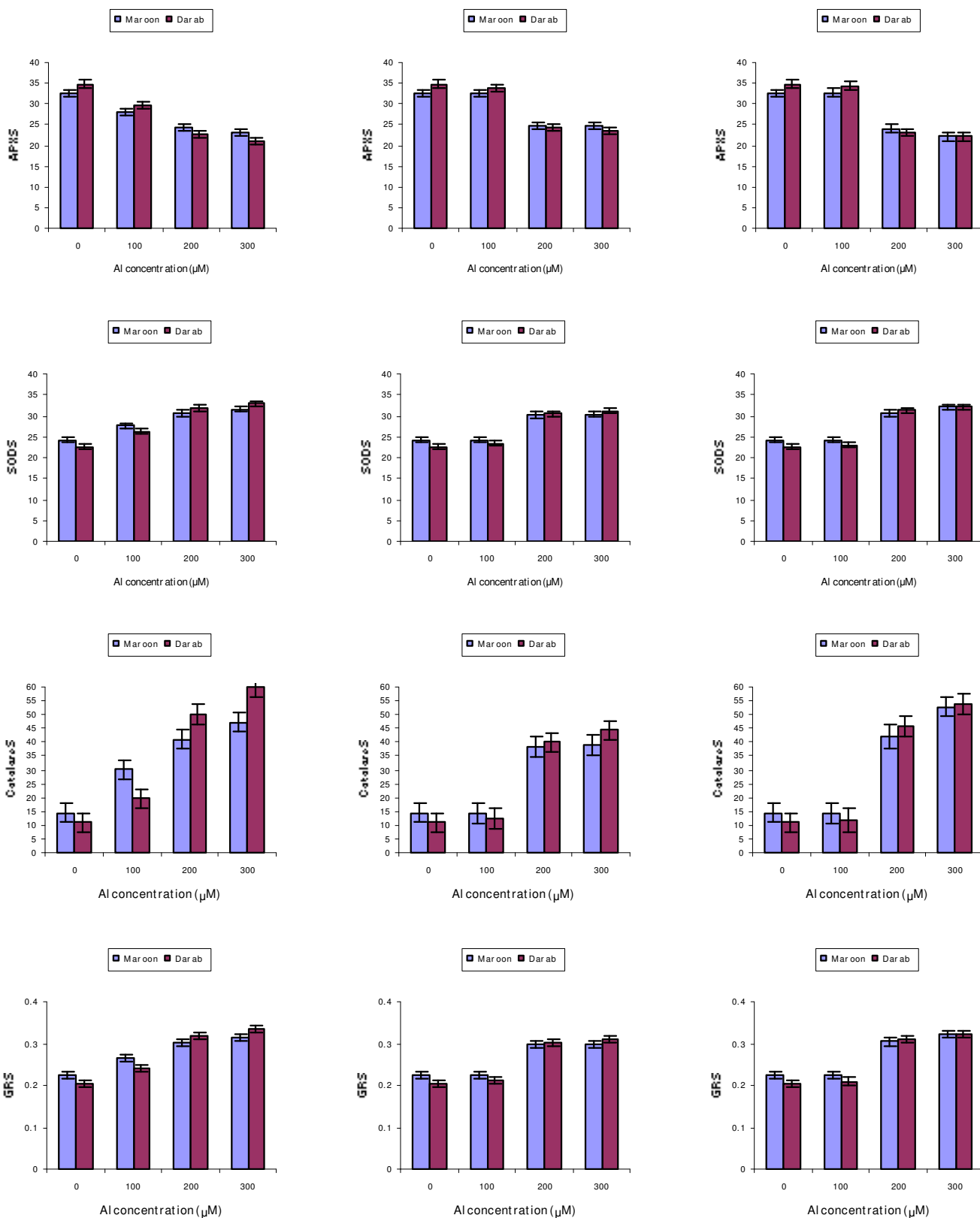


Figure 1. Effect of malate and citrate treatments on reducing the noxious effect of Al in medium culture. The first column from left show the effect of just different Al concentrations in medium culture on the different enzyme activities. The second column show the effect of just Al concentrations in medium culture together with the malate on the different enzyme activities and the third column show the effect of Al concentrations in medium culture together with the citrate on the different enzyme activities.

Table 2. Summary table for wheat enzymes data in different AI concentrations using regression analysis.

Character	Line	Source	Linear				Quadratic				Cubic			
			Pr> T ^a	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE
	R	Intercept	-	-	23.35	0.75	-	-	24.06	0.80	-	-	24.58	0.28
		Concentration	**	0.86	-0.03		**	-	-0.05	0.01	**	-	-0.14	0.01
		Concentration ²	-	-	-	-	ns	0.88	0.00007	0.00004	**	-	0.0009	0.0001
		Concentration ³	-	-	-	-	-	-	-	-	**	0.99	-0.000002	2.107168E-7
APX-R	S	Intercept	-	-	23.85	2.26	-	-	25.81	2.18	-	-	26.04	2.34
		Concentration	**	0.61	-0.05	0.01	*	-	-0.12	0.04	-	-	-0.17	0.11
		Concentration ²	-	-	-	-	ns	0.71	0.0003	0.0001	-	-	0.0007	0.001
		Concentration ³	-	-	-	-	-	-	-	-	-	0.68	-0.000001	0.000002
	R	Intercept	-	-	31.99	1.41	-	-	32.82	1.64	-	-	32.70	1.78
		Concentration	**	0.63	-0.03	0.008	ns	-	-0.06	0.03	ns	-	-0.04	0.07
		Concentration ²	-	-	-	-	ns	0.62	0.00008	0.00008	ns	-	-0.00009	0.0006
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.58	3.915E-7	0.000001
APX-S	S	Intercept	-	-	34.33	0.71	-	-	35.24	0.63	-	-	34.84	0.30
		Concentration	**	0.94	-0.05	0.004	**	-	-0.08	0.01	ns	-	-0.01	0.01
		Concentration ²	-	-	-	-	*	0.96	0.00009	0.00003	**	-	-0.0005	0.0001
		Concentration ³	-	-	-	-	-	-	-	-	**	0.99	0.000001	2.227478E-7
Caretenoid	R	Intercept	-	-	354.95	27.73	-	-	355.86	34.04	-	-	356.91	37.008
		Concentration	ns	0.02	-0.07	0.15	ns	-	-0.09	0.55	ns	-	-0.26	1.42
		Concentration ²	-	-	-	-	ns	0.02	0.00009	0.002	ns	-	0.002	0.01
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.02	-0.000004	0.00003
	S	Intercept	-	-	571.75	71.02	-	-	592.47	86.17	-	-	615.07	86.69
		Concentration	**	0.52	-1.37	0.38	ns	-	-2.00	1.38	ns	-	-5.54	3.33
		Concentration ²	-	-	-	-	ns	0.48	0.002	0.004	ns	-	0.04	0.03
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.50	-0.00008	0.00007

Character	Line	Source	Linear				Quadratic				Cubic			
			Pr> T ^a	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE
Catalase-R	R	Intercept	-	-	42.56	6.46	-	-	43.99	7.88	-	-	38.91	3.00

Table 2. Contd.

		Concentration	**	0.88	0.33	0.04	ns	-	0.28	0.13	**	-	1.08	0.12
		Concentration ²	-	-	-	-	ns	0.88	0.0001	0.0004	**	-	-0.008	0.001
		Concentration ³	-	-	-	-	-	-	-	-	**	0.98	0.00002	0.000002
	S	Intercept	-	-	52.19	12.42	-	-	39.78	10.78	-	-	39.31	11.80
		Concentration	**	0.85	0.47	0.07	**	-	0.94	0.19	ns	-	1.04	0.53
		Concentration ²	-	-	-	-	*	0.91	-0.002	0.0006	ns	-	-0.003	0.005
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.90	0.000002	0.00001
	R	Intercept	-	-	16.75	5.39	-	-	14.40	6.44	-	-	14.40	7.01
		Concentration	**	0.55	0.11	0.03	ns	-	0.18	0.10	ns	-	0.18	0.27
		Concentration ²	-	-	-	-	ns	0.53	-0.0002	0.0003	ns	-	-0.0002	0.002
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.47	-2.58889E-8	0.000005
Catalase-S	S	Intercept	-	-	8.50	2.62	-	-	8.79	3.22	-	-	10.91	0.99
		Concentration	**	0.94	0.18	0.01	**	-	0.17	0.05	**	-	-0.16	0.04
		Concentration ²	-	-	-	-	ns	0.93	0.00003	0.0002	**	-	0.003	0.0003
		Concentration ³	-	-	-	-	-	-	-	-	**	0.99	-0.000007	7.361438E-7
	R	Intercept	-	-	4.60	0.39	-	-	4.34	0.45	-	-	4.33	0.49
		Concentration	ns	0.02	-0.0009	0.002	ns	-	0.007	0.007	ns	-	0.009	0.02
		Concentration ²	-	-	-	-	ns	0.13	-0.00003	0.00002	ns	-	-0.00005	0.0002
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.14	5.5E-8	3.681328E-7
Chlorophyll A	S	Intercept	-	-	5.50	0.57	-	-	5.56	0.70	-	-	5.59	0.75
		Concentration	**	0.61	-0.01	0.003	ns	-	-0.02	0.01	ns	-	-0.02	0.03
		Concentration ²	-	-	-	-	ns	0.56	0.000006	0.00004	ns	-	0.00005	0.0003
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.51	-9.46111E-8	5.624392E-7

Character	Line	Source	Linear				Quadratic				Cubic			
			Pr> T ^a	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE
	R	Intercept	-	-	0.41	0.04	-	-	0.41	0.04	-	-	0.42	0.04
Chlorophyll B		Concentration	ns	0.06	0.0001	0.0002	ns	-	0.00008	0.0007	ns	-	-0.001	0.002
		Concentration ²	-	-	-	-	ns	0.06	2.155602E-7	0.000002	ns	-	0.00001	0.00002
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.12	-2.4376E-8	3.331914E-8

Table 2. Contd.

GR-R	S	Intercept	-	-	0.48	0.06	-	-	0.41	0.06	-	-	0.42	0.06
		Concentration	*	0.36	-0.0009	0.0003	ns	-	0.001	0.0009	ns	-	0.0002	0.002
		Concentration ²	-	-	-	-	*	0.55	-0.000007	0.000003	ns	-	0.000003	0.00002
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.51	-2.17434E-8	4.650606E-8
	R	Intercept	-	-	0.31	0.007	-	-	0.31	0.008	-	-	0.30	0.003
		Concentration	**	0.86	0.0003	0.00004	**	-	0.0005	0.0001	**	-	0.001	0.0001
		Concentration ²	-	-	-	-	ns	0.88	-6.58333E-7	4.115032E-7	**	-	-0.000009	9.691061E-7
		Concentration ³	-	-	-	-	-	-	-	-	**	0.99	1.738889E-8	2.130032E-9
	S	Intercept	-	-	0.31	0.02	-	-	0.29	0.02	-	-	0.29	0.02
		Concentration	**	0.62	0.0005	0.0001	*	-	0.001	0.0004	ns	-	0.002	0.001
		Concentration ²	-	-	-	-	ns	0.73	-0.000002	0.000001	ns	-	-0.000007	0.00001
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.70	1.055556E-8	1.96756E-8
GR-S	R	Intercept	-	-	0.23	0.01	-	-	0.22	0.02	-	-	0.22	0.02
		Concentration	**	0.62	0.0003	0.00007	ns	-	0.0005	0.0003	ns	-	0.0004	0.0007
		Concentration ²	-	-	-	-	ns	0.62	-7.66667E-7	7.969425E-7	ns	-	8.833333E-7	0.000006
		Concentration ³	-	-	-	-	-	-	-	-	ns	0.58	-3.66667E-9	1.25339E-8
S	Intercept	-	-	0.21	0.007	-	-	0.20	0.008	-	-	0.20	0.003	
	Concentration	**	0.93	0.0005	0.00004	**	-	0.0006	0.0001	**	-	-0.0002	0.0001	
	Concentration ²	-	-	-	-	ns	0.93	-5.83333E-7	4.180053E-7	**	-	0.000007	0.000001	
	Concentration ³	-	-	-	-	-	-	-	-	**	0.98	-1.73333E-8	2.475185E-9	

Character	Line	Source	Linear				Quadratic				Cubic			
			Pr> T ^a	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE	Pr> T	R ²	Estimate	SE
SOD-R	R	Intercept	-	-	31.24	0.56	-	-	30.71	0.60	-	-	30.32	0.21
		Concentration	**	0.86	0.03	0.003	**	-	0.04	0.01	**	-	0.10	0.008
		Concentration ²	-	-	-	-	ns	0.88	-0.00005	0.00003	**	-	-0.0006	0.00007
		Concentration ³	-	-	-	-	-	-	-	-	**	0.98	0.000001	1.580746E-7
	S	Intercept	-	-	30.86	1.70	-	-	29.39	1.63	-	-	29.22	1.76
		Concentration	**	0.61	0.03	0.009	*	-	0.09	0.03	ns	-	0.13	0.08
		Concentration ²	-	-	-	-	ns	0.71	-0.0001	0.00009	ns	-	-0.0006	0.0007

Table 2. Contd.

		Concentration ³	-	-	-	-	-	-	-	-	ns	0.68	8.286111E-7	0.000002
SOD-S	R	Intercept	-	-	24.76	1.06	-	-	24.13	-	-	-	24.22	1.33
		Concentration	**	0.63	0.03	0.006	ns	-	0.04	ns	-	-	0.03	0.05
		Concentration ²	-	-	-	-	ns	0.63	-0.00006	ns	-	-	0.00007	0.0005
		Concentration ³	-	-	-	-	-	-	-	ns	0.58	-2.93944E-7	9.940348E-7	
S	S	Intercept	-	-	22.89	0.53	-	-	22.28	0.52	-	-	22.62	0.18
		Concentration	**	0.94	0.04	0.003	**	-	0.06	0.008	ns	-	0.002	0.007
		Concentration ²	-	-	-	-	*	0.96	-0.00006	0.00003	**	-	0.0005	0.00006
		Concentration ³	-	-	-	-	-	-	-	-	**	0.99	-0.000001	SE

malate or citrate application in some Al concentrations is effective compared with other Al concentrations. As shown in Figure 1, the application of citrate and malate approximately improved enzymes activities; however, their applications did not change the observed trends (Figure 1).

DISCUSSION

A uniform increase in superoxide dismutase (SOD), catalase (CAT) and glutathione reductase (GR) activities associated with a gradual decrease in APX activity was detected by increasing Al concentration in medium culture. SOD is an important enzyme that is associated with anti-oxidative stress in plants (Guo et al., 2004). Increased superoxide dismutase activity may be due to an elevated content of superoxide radical. It has been shown that, Al enhanced SOD activity in root tips of soybean (Cakmak and Horst, 1991), roots of Arabidopsis (Richards et al., 1998), and roots of sorghum (Peixoto et al., 1999). Lee et al. (2001) suggested that, enhanced activity of superoxide dismutase may function in signaling of oxidative stress and hence, leads to the induction

of antioxidant enzymes associated with an H₂O₂ scavenging system, particularly an ascorbate-glutathione cycle. Al-induced high activity of SOD that was observed in the present study is consistent with previous results obtained from different plants (Boscolo et al., 2003; Tamas et al., 2003; Meriga et al., 2004; Sharma and Dubey, 2007).

The marked increase of catalase activity with increasing Al levels in Maroon and Darab varieties may indicate enhanced production of reactive oxygen species (ROS) under an excess of aluminium (Al). This enhanced activity seems to be related to increased oxidative stress tolerance (Allen, 1995). Among the enzymatic systems considered to play an important role in the cellular defense strategy against oxidative stress, catalase (CAT) plays an important role as it decomposes H₂O₂ to water and O₂. It has been previously suggested that, accumulation of H₂O₂ caused by various environmental stresses would result in the enhanced activity of catalase in order to protect plant cells (Mizuno et al., 1988).

Superoxide dismutase (SOD) and catalase may cooperatively contribute to suppress the lipid peroxidation caused by H₂O₂ and diamide as well as aluminium stress in *Andropogon* and *Miscanthus*.

Hydrogen peroxide which is cytotoxic and acts both as an oxidant and reductant is detoxified by catalase activity.

Al, in the present study, decreased APX activity in root and shoot of both studied genotypes. Reports available on effect of Al on ascorbate peroxidase activity have been some what contradictory. Aluminium has been shown to enhance activity of APX in *Cucurbita pepo* (Dipierro et al., 2005) and rice (Sharma and Dubey, 2007). Activity of chloroplastic-APX in rice, however, was inhibited by Al (Sharma and Dubey, 2007).

Interestingly, the greater catalase activity in Darab as an Al-sensitive genotype than in Maroon as an Al-resistant may be attributed to higher oxidative stress in Darab. Differences in response of Al-resistant in relation to Al-sensitive plant genotypes with respect to activities of antioxidant enzymes have been recorded. For instance, Al-resistant genotypes of wheat (Darko et al., 2004) or rice (Ma et al., 2007) showed significantly higher catalase and ascorbate peroxidase activities than that of Al-sensitive counterparts under Al- treatments. The opposite, however, was true for barley (Tamas et al., 2003) and maize (Boscolo et al., 2003) that registered a greater

Table 3. Analysis of variance summary for wheat enzymes data under different Al concentrations as well as malate and citrate treatments. Data were analyzed using procedures for a completely randomized design.

Source	DF	Mean of square			
		APXs	SODs	Catalases	GRs
Genotype	1	2.78 ^{ns}	1.82 ^{ns}	8.16 ^{ns}	0.0003 ^{ns}
Al concentration	3	547.71**	309.62**	5911.81**	0.05**
Treatment	2	21.76**	11.59**	330.50**	0.002**
genotype× treatment	2	0.72 ^{ns}	0.30 ^{ns}	6.74 ^{ns}	0.00003 ^{ns}
Al concentration × treatment	6	8.75**	4.54**	90.31*	0.0006*
genotype× Al concentration	3	11.27**	6.54**	141.55*	0.001**
genotype× Al concentration × treatment	6	0.45 ^{ns}	0.32 ^{ns}	32.09 ^{ns}	0.00008 ^{ns}
Error	46	2.20	1.23	37.91	0.0002
CV		5.29	4.00	20.13	5.28

Table 3. Continued.

Source	DF	Mean of square			
		APXr	SODr	Catalaser	GRr
Genotype	1	19.40 ^{ns}	10.91 ^{ns}	7344.42**	0.002 ^{ns}
Al concentration	3	433.56**	243.87**	35592.00**	0.04**
Treatment	2	11.20 ^{ns}	6.30 ^{ns}	2407.24**	0.001 ^{ns}
genotype× treatment	2	2.61 ^{ns}	1.47 ^{ns}	695.41*	0.0003 ^{ns}
Al concentration × treatment	6	2.16 ^{ns}	1.22 ^{ns}	388.22*	0.0002 ^{ns}
genotype× Al concentration	3	12.96 ^{ns}	7.29 ^{ns}	1117.79**	0.001 ^{ns}
genotype× Al concentration × treatment	6	2.23 ^{ns}	1.25 ^{ns}	441.54*	0.0002 ^{ns}
Error	43	7.00	3.94	150.00	0.0006
CV		14.29	5.69	12.96	6.91

CV: Coefficient of variation. df = Degrees of freedom; ***, **, *, significant at 0.001, 0.01 and 0.05 probability level; ns: non significant.

superoxide dismutase activity in Al-sensitive genotypes than that of Al-resistant genotypes under Al treatments. It was thus, indicated that induction or inhibition of specific antioxidant enzymes in plants in response to Al-stress was genotype dependent. We suppose that, induction of antiperoxi-dation enzymes by Al treatment may contribute to their Al tolerance by a suppression of oxidative damages caused by Al stress. Similar enhanced tolerance to oxidative stress and/or heavy metal stress was also observed by expression of antioxidant enzymes [glutathione S-transferase (GST), superoxide dismutase (SOD), peroxidase and dehydroascorbate reductase] in transgenic plants (Ezaki et al., 2000; Lee et al., 2007).

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Abbreviations

APX, Ascorbate peroxidase; **CAT**, catalase; **GR**, glutathione reductase; **SOD**, superoxide dismutase; **ROS**, reactive oxygen species; **Al**, aluminum; **GST**, glutathione S-transferase.

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