

Research Article

# Physiological and biochemical responses of seedlings of six contrasting barley (*Hordeum vulgare* L.) cultivars grown under salt-stressed conditions

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#### Abstract

Salinity stress affects plant growth and development and underlying metabolisms. To mitigate the effects of the stress, plants responded by changing their physiological and biochemical activities and withstand the stress. The present study aimed to determine barley's (Hordeum vulgare L.) physiological and biochemical response to salinity stress conditions for 7 days and 14 days. Six barley cultivars (Alfa93, DWRB73, DL88, NB1, NB3, NDB1173) were grown under controlled conditions, and different level of salinity stress was applied. In addition, seedling growth, physiological and biochemical parameters, plant leaves RWC, and electrolyte leakage were analyzed. The overall seedling growth, RWC, and electrolyte leakage in salt susceptible lines Alfa93 and DWRB73 were low than the salt-tolerant barley lines (DL88, NB1, NB3, and NDB1173). Electrolyte leakage was 26.0 and 20.6% in Alfa93 and DWRB73, whereas it was 17.6, 14.6, 15.3, and 10.4% in DL88, NB1, NB3, and NDB1173, respectively at 300 mM salinity stress. The loss of photosynthetic pigments under salt stress was high in susceptible lines, salinity treated (300 mM NaCl) Alfa93 plants exhibit 49.5% and 59.5% of Chl-a than control plants after 7 and 14 days of treatment, respectively. However, at 300 mM stress level, NB1 (ST) showed less Ch⊦a loss after 7 days, whereas NDB1173 showed less reduction in Chl-a after 14 days. Antioxidant enzymes such as SOD, POX, CAT, and APX activities in susceptible line Alfa93 and DWRB73 were lower than tolerant lines. PCA analysis demonstrated a positive correlation between antioxidant enzyme activities and genotypes under salinity stress. PCA analysis described DL88 as the most tolerant, and DWRB73 was the most salt susceptible genotype among the studied barley genotypes. The present findings suggest that barley cultivars' physiological and biochemical activities under salinity stress conditions may be used to screen salt-tolerant crops.

Keywords: Antioxidant enzymes, Barley, Chlorophyll, Eelectrolyte leakage, RWC, Salt stress.

# INTRODUCTION

Barley is grown worldwide for various purposes, such as human consumption, feed for livestock, and brewing and malting (Noreen *et al.*, 2021). Barley ranked fourth among cereal crops in total world production after wheat, rice, and corn (Naeem *et al.*, 2021). It is cultivated in various environmental conditions, such as subarctic to sub-tropical (Bera *et al.* 2018).

Salt stress is a major ecological as well as an agro-

nomic problem all over the world.

Around 6.74 million ha of land in India is affected by salinity, and an estimation of 10% rise every year, around 50% of the total arable land by 2050 (Kumar and Sharma 2020).

Not only has the natural salinity, but the salinization by human activity also becomes a serious threat to agricultural production (Ortiz and Jin 2021). Under high salt stress condition, plants uptake high concentration of soluble salt which resists the water movement inside

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the root and cause osmotic stress. The osmotic stress alters the membrane stability and influences the absorption of high concentrations of salt inside cells. As a result, high ions compete with the uptake of essential nutrients and cause nutrition deficiency (Arif *et al.* 2020; Moradi *et al.* 2021).

Due to the excessive salts in the soil decrease its osmotic potentials and the water availability to the roots. It also results in considerable ROS accumulation in roots and leaves (Tanou *et al.*, 2009; Xie *et al.*, 2011). Antioxidant enzymes provide tolerance to the plants against various stresses. Superoxide dismutase initiates the first vital step against oxidative stress in plants by converting superoxide (O2<sup>--</sup>) to hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), and further catalase (CAT), guaiacol peroxidase (POD), and ascorbate peroxidase (APX) catalyze the hydrogen peroxide into water and oxygen (Zhu *et al.*, 2020).

Plants harness the light from the photosynthetically active radiation (PAR) region with the help of pigments in the green part. Three pigments, specifically chlorophylla (Chl-a), chlorophyll-b (Chl-b), and the carotenoids (CRT), are actively involved in photosynthesis. The amount of these pigments in the plants rules the photosynthetic potential and determines its efficiency in utilizing PAR for the biosynthesis process (Kume *et al.* 2018). Therefore, the physiological status of plants is directly related to these pigment concentrations, and changes in these affect the plants' growth and development.

Barley can be used as a model crop to study the mechanism of salinity tolerance because it is the most salinity tolerant crop among other cereal crops. Physiological changes in the plants under salinity stress give insight into the salinity response of plants. Therefore, the present study was carried out to explain physiological and biochemical characteristics such as chlorophyll content, carotenoid content, electrolyte leakage, antioxidative enzymes in providing salinity tolerance to the plants.

#### MATERIALS AND METHODS

#### Plant materials and experimental conditions

Barley seeds were procured from the Indian Institute of Wheat and Barley Research (IIWBR), Karnal, India. Seeds were screened for salt tolerance, and six cultivars were chosen for study, salt-tolerant (ST) lines DL 88, NB 1, NB 3, NDB 1173, and salt-sensitive (SS) lines Alfa 93 and DWRB 73. Seeds were surface sterilized using 0.1% HgCl<sub>2</sub> solution and then placed on wet filter paper for germination at 25° in the growth chamber at Centre for Biotechnology, M D University, Rohtak, India). After germination, seeds were transferred to plastic pots filled with sand (thoroughly washed with distilled water and autoclaved). Seedlings were supplied with half-strength Hoagland solution (Jones, 1982), and the growth conditions provided were 16hr light and 8 hr dark cycle, day temperature 25°C with 60% relative humidity and night temperature was 18 °C. When seedlings were attained two-leaf stage, salt stresses were introduced gradually (25 mM morning and 25 mM evening up to the desirable levels) to avoid osmotic shock. Plant growth conditions and salinity stress treatment followed Elsawy et al.'s methods (2018) with few modifications. Salinity levels include control (0 mM), 100, 200, and 300 mM NaCl solution in half-strength Hoagland solution. The plant samples were collected after 7 days of salt treatment and 14 days of salt treatment.

#### **Relative water Content**

The leaf's relative water content (RWC) was assessed just after collecting plants, leaf fresh weight (FW) was taken without delay, and then leaves were rehydrated by floating them in distilled water at room temperature for 4 hours. After rehydration, leaf turgid weight (TW) was measured, and then leaves were kept in an oven at 60°C for 48 hours. After 48 hours, the dry weight (DW) of leaves was measured, leaf RWC was calculated using the formula (Barrs and Weatherley, 1962).

RWC (%) =  $\begin{bmatrix} \frac{FW-DW}{TW-DW} \end{bmatrix} x \ 100$ 

.....Eq.1

# Study of relative electrolyte leakage (REL)

Electrolyte leakage of barley leaves was analyzed according to Lakra *et al.* (2015). Leaf samples were collected from the control, and salinity-stressed plants, washed gently with distilled water to eliminate any surface adhering ions. About 100 mg leaf tissue was weighed and immediately dipped into 20 ml milli-Q water, incubating the sample at 60° for 2 hours. After that, samples were cooled to room temperature, and the electrical conductivity (E1) of the solution was measured using a conductivity meter (Okaton, USA). Total conductivity (E2) was determined by autoclaving the solution with the sample at 121° for 15 minutes. Samples were cooled to room temperature, and the conductivity of the solution was measured. Relative electrical conductivity was measured using the formula:

Relative electrolyte leakage % (REL%) = (E1 /E2) ×100 ......Eq.2

# **Determination of photosynthetic pigments**

Chlorophyll content was extracted using the method described by Minocha *et al.* (2009) with few modifications. Approx 50 mg leaves from all varieties were taken and placed in 8 ml DMSO and incubated at 60°C for 4 hours in the dark. The pigment concentration was measured by taking absorbance at 480, 649, and 665 nm using a UV/ vis spectrophotometer (Genetix, Spectro-8). The content of chlorophyll-a (Chl-a), Chlorophyll-b (Chl-b), and carotenoids (CRT) were calculated according to the formula described by Wellburn (1994).

### Antioxidant enzyme activity assay

Antioxidant enzyme activity assay was done according to the method described by Lakra et al. (2015). About 100 mg fresh leaves were collected from the control and salt-stressed plants and freeze in liquid nitrogen. Leaves were homogenized in ice-cold 50 mM potassium phosphate buffer (pH 7.5) consists of 2 mM EDTA and 0.1 mM PMSF. Ascorbate (2 mM) was additionally added to the homogenizing buffer for ascorbate peroxidase (APX). The homogenates were centrifuged at 12000 x g for 10 minutes at 4 °C. The supernatant was collected in fresh microcentrifuge tubes and used for enzyme assay. The activity of superoxide dismutase (SOD) was measured according to the method described by Dhindsa et al. (1981), the ability of the enzyme to inhibit the photochemical reduction of nitroblue tetrazolium (NBT). The guaiacol peroxidase (POD) activity was assayed according to the method of Chance and Maehly (1955), based on the ability of the enzyme to convert guaiacol to tetraguaiacol (ɛ= 26.6 mM<sup>-1</sup>cm<sup>-1</sup>). The catalase (CAT) activity was determined according to the method of Chance and Maehly (1955) by measuring the decomposition of hydrogen peroxide  $(H_2O_2)$  at 240 nm ( $\epsilon$ = 40 M<sup>-1</sup>cm<sup>-1</sup>). The ascorbate peroxidase (APX) activity was determined according to the method described by Nakano and Asada (1981), based on the oxidation of ascorbate by H<sub>2</sub>O<sub>2</sub> and decrease in absorption at 290 nm ( $\varepsilon$ = 2.8 mM<sup>-1</sup>cm<sup>-1</sup>).

#### Statistical analysis

Statistical analyses of the data weredone by the analysis of variance (ANOVA). The significant differences between the means of stress treatments were determined by the LSD (least significant difference) test at p < 0.01 by SPSS-20.0 (USA). Graphs were prepared using the Microsoft Excel program. Principal component analysis (PCA) was performed using XLSTAT software. The first two principal components were used to derive PCA-biplot, and the possible associations among the genotypes and measured physiological and biochemical traits were determined.

## **RESULTS AND DISCUSSION**

Plant response to salinity stress is a complex phenomenon as it involves changes in morphology to change in metabolism. The changes depend on various factors such as stress level, tolerance potential of the plants, and developmental stage of plants.

# Physiological response of barley cultivars grown under different salt stress

The responses of Indian barley cultivars Alfa93, DWRB73, DL88, NB1, NB3, and NDB1173 to salt stress-induced conditions assessed by comparing the shoot height and fresh weight of the plants are given in Fig. 1; Table 1 and 2.

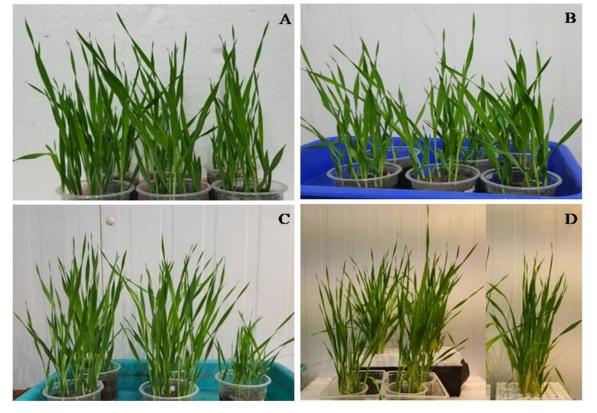
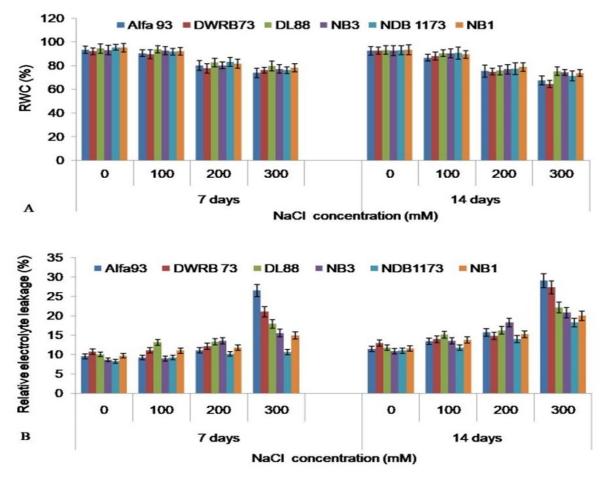


Fig. 1. Effect of salinity treatment on the growth of barley cultivars under (A) control (0 mM NaCl) and saline (B) 100 mM NaCl, (C)200 mM NaCl, and (D) 300 mM NaCl

**Table 1**. Effects of salinity treatment on growth parameter shoot heigh (cm) of *Hordeum vulgare* L. cultivars grown under control (0 mM NaCl) and saline (100, 200, and 300 mM NaCl) conditions. Values are means  $\pm$  SD (n = 3). Asterisks (\*) denote the mean differences are significant from controls at 0.01 level.

	Shoot Height (cm) 7 days					Shoot Height (cm) 14 days					
	0 mM NaCl	100 mM NaCl	200 mM NaCl	300 mM NaCl	0 mM NaCl	100 mM NaCl	200 mM NaCl	300 mM NaCl			
Alfa 93	27.2 ± 2.3	26.8 ± 1.6	19.0 ± 1.3*	17.5 ± 2.1*	30.3 ± 2.2	28.9 ± 3.8	23.8 ± 2.0*	21.3 ± 1.4*			
DWRB73	35.0 ± 1.8	35.2 ± 1.9	31.8 ± 2.7	29.8 ± 1.3*	39.9 ± 1.8	37.9 ± 1.4	36.7 ± 1.5	31.1 ± 1.5*			
DL88	28.9 ± 3.1	27.9 ± 3.2	26.6 ± 2.8	22.4 ± 1.6	31.8 ± 2.2	30.0 ± 1.8	28.9 ± 1.4	25.1 ± 1.5*			
NB3	33.7 ± 1.7	32.3 ± 2.2	29.8 ± 2.3	24.7 ± 1.3*	38.1 ± 2.2	36.7 ± 1.5	31.6 ± 1.5*	25.8 ± 1.3*			
NDB 1173	27.2 ± 3.1	28.6 ± 1.3	26.5 ± 2.1	24.2 ± 1.5	31.9 ± 1.4	31.7 ± 1.6	28.9 ± 1.2	25.8 ± 2.2*			
NB1	29.5 ± 2.6	28.6 ± 1.6	27.0 ± 2.1	26.3 ± 2.1	31.4 ± 1.6	28.1 ± 1.3	24.4 ± 1.8*	23.7 ± 2.0*			



**Fig. 2.** Different physiological responses of plants under salinity stress treatment (A) Relative Water Content (RWC%), and (B) Relative Electrolyte leakage in leaves of barley cultivars under control and saline (0 mM, 100 mM, 200 mM, and 300 mM NaCl) conditions after 7 days and 14 days.

#### Relative water content (RWC %)

Leaf RWC of barley cultivars under control and salinitystressed conditions for 7 days and 14 days ranged from 95.3% to 73.8% and 93.2% to 64.4%, respectively (Fig. 2A). When salinity stress increased, the RWC of the plants decreased. The most reduction in RWC of 7 days was observed in susceptible lines than tolerant varieties. In control plants (both SS and ST), there were no significant differences in RWC, whereas, at 300 mM salinity stress level, a steep decline was observed in SS plants. In the susceptible barley lines, Alfa93, and DWRB73 lines at 300 mM, RWC was 78.8% and 76.1% after 7 days, and after 14 days, it was decreased to 67.4% and 64.4%, respectively. In the tolerant barley lines at 300 mM stress level, DL88 exhibits 79.8% after 7 days and 75.2 % after 14 days, and a similar pattern

Table 2. Effects of salinity treatment on growth parameter shoot fresh weight (mg) of Hordeum vulgare L. cultivars grown under control (0 mM NaCl) and saline (100, 200 and 300 mM NaCI) conditions. Values are means ± SD (n = 3). Asterisks (\*) denote the mean differences are significant from controls at 0.01 level

	Shoot weight (mg) 7 days	days			Shoot weight (mg) 14 days	g) 14 days		
	0 mM NaCl	100 mM NaCI	200 mM NaCI	300 mM NaCl	0 mM NaCI	100 mM NaCI	200 mM NaCI	300 mM NaCI
Alfa 93	1151.3 ± 71.4	1100.0 ± 94.9	756.3 ± 20.5	591.1 ± 39.4*	1118.3 ± 50.0	1058.8 ± 65.7	924.8 ± 72.3*	665.4 ± 40.3*
DWRB73	1298.7 ± 335.1	1040.7 ± 40.6	967.2 ± 173.8	1002.8 ± 114.9*	1390.3 ± 190.3	1269.3 ± 188.0	979.0 ± 52.1	785.7 ± 71.5
DL88	1037.1 ± 72.9	918.5 ± 125.1	869.5 ± 51.3	620.2 ± 98.8	1106.2 ± 183.8	1042.3 ± 179.4	926.5 ± 125.2	750.0 ± 23.8*
NB3	1519.2 ± 83.3	1294.0 ± 40.9	831.5 ± 41.4	788.0 ± 100.5*	1379.0 ± 232.7	1307.3 ± 134.8	1092.7 ± 97.4*	809.7 ± 80.7*
NDB 1173	930.7 ± 94.3	983.3 ± 173.3	769.5 ± 51.2	715.3 ± 45.4*	1189.0 ± 110.5	$1090.8 \pm 131.3$	1011.7 ± 33.0	818.8 ± 61.9*
NB1	1081.8 ± 199.2	967.7 ± 39.5	981.2 ± 105.8	932.0 ± 147.2	1330.2 ± 189.4	1027.8 ± 149.6	907.8 ± 20.1	897.2 ± 110.7

of decrease in RWC was observed in other tolerant lines.

The RWC is a valuable tool for indicating the water relation in plants. The relationship between plant and water depends significantly on the age and nature of plant materials (González and González-Vilar, 2001). The present study on RWC in barley plants demonstrated the relationship between salt stress and water content in plants. Although the water content of plants decreased with the increase of salinity level, the RWC of salt-tolerant was greater than those of salt susceptible plants. He et al. (2019) reported a decrease in the RWC of two barley cultivars (Kunlun 14 and Ganpi6) when salinity stress was imposed for 48 hours. In the present study, a lower level of in RWC of the susceptible line Alfa93 and DWRB73 and a higher level of RWC of tolerant lines DL88, NB1, NB3, and NDB1173 under salinity stress-induced conditions are in agreement with the result of Mahlooji et al. (2018) who reported a higher RWC of tolerant barley genotype (Khatam) than the sensitive genotype (Morocco) under salinity.

Reduction in the RWC of the plant leaves under salinity -induced conditions may occur due to loss of turgor in the leaves under salinity stress which resulted in limited water availability and caused dehydration at the cellular level (Soni *et al.*, 2021).

# Relative electrolyte leakage

Electrolyte leakage can be used as an indicator for membrane damage caused by abiotic stresses on the membrane. Electrolyte leakage was analyzed under salinity stress-induced conditions. The relative percent electrolyte leakage was higher in treated plants than in control plants (Fig. 2B). No significant change was observed in control plants, whether salinity tolerant or susceptible but observed at 300 mM significant difference between them. Tolerant lines exhibited decreased electrolyte leakage than susceptible lines, in 7 days, susceptible lines Alfa93, DWRB73 showed 26.0 and 20.6 percent relative electrolyte leakage, respectively, whereas tolerant lines DL88, NB1, NB3, and NDB1173 exhibits 17.6, 14.6, 15.3, and 10.4 percent relative electrolyte leakage respectively, a similar pattern of relative electrolyte leakage was also observed after 14 days treatment.

Salinity stress may modify the physical structure of the plasma membrane by a change in chemical composition and organic acids (Bajji *et al.*, 2001). The membrane injury depends on the level of osmotic stress and the duration (Kocheva *et al.*, 2004). Electrolyte leakage gradually increases when salt stress levels increase, and the level of this leakage depends on cultivars, such as salt-tolerant cultivars with lower electrolyte leakage than susceptible cultivars (Mahlooji *et al.*, 2018; Zeeshan *et al.*, 2020). In the present study, electrolyte leakage

age in salinity susceptible lines Alfa93, DWRB73 were higher in comparison to the salinity tolerant lines DL88, NB1, NB3, and NDB1173. A similar result was also reported by Mahlooji et al. (2018), who reported salt susceptible barley line Morocco exhibited higher electrolyte leakage than salt-tolerant line Khatam under salinity stress conditions. When comparing the electrolyte leakage in control and 300 mM NaCl treated barley cultivar CM72, wheat cultivars Suntop (ST) and Sunmat (SS) higher value of electrolyte leakage was observed in 300 mM NaCl treated plants (Zeeshan et al., 2020). Elsawy et al. (2018) studied the effects of salt stress in two Egyptian barley cultivars, Giza 126 (SS) and Giza 128 (ST). They reported that stress exhibited higher electrolyte leakage than control plants; also, tolerant lines had reduced electrolyte leakage than susceptible lines under salt stress-induced conditions (200 mM NaCl). Electrolyte leakage may be an important tool in the screening of salt susceptible and tolerant cultivars.

## Effect of salt stress on photosynthetic pigment

Salinity stress negatively affected the photosynthetic pigments and an increase in salinity stress caused loss of photosynthetic pigments. However, among cereal crops, barley is somewhat salt tolerant. Changes in photosynthetic pigments are shown in Fig. 3. The highest loss of photosynthetic pigments, Chl-a, was recorded in salt susceptible lines and was 49.5% and 59.5% in Alfa93 in 7 and 14 days respectively under 300 mM salt stress level, while another susceptible line, DWRB73, exhibits loss of 52.1% and 64% in 7 and 14 days respectively. In comparison to susceptible lines, tolerant lines showed less loss of photosynthetic pigments. Among tolerant lines under 300 mM salt stress, NB1 showed less loss in Chl-a in 7 days, while in 14 days, NDB1173 was showed less loss in Chl-a. The loss in Chl-b under 300 mM salt stress was 54.7 and 62.2% in 7days in Alfa93 and DWRB73, respectively, while in salt-tolerant lines, loss in Chl-b was higher than susceptible lines 14 days. A similar pattern like Chl-a in CRT was also observed. Susceptible lines lose more carotenoids pigment than tolerant lines.

Chlorophylls are fundamental pigments in plants to absorb light and release electrons. However, various types of chlorophyll exist in plants, but only two types are possessed by the terrestrial plants: Chl-a and Chlb. These two pigments form light-harvesting complexes, which absorb the most light (Kume *et al.*, 2018). Carotenoid is the part of the photosystem and chlorophylls and is located in chromoplasts (Costache *et al.*, 2012).

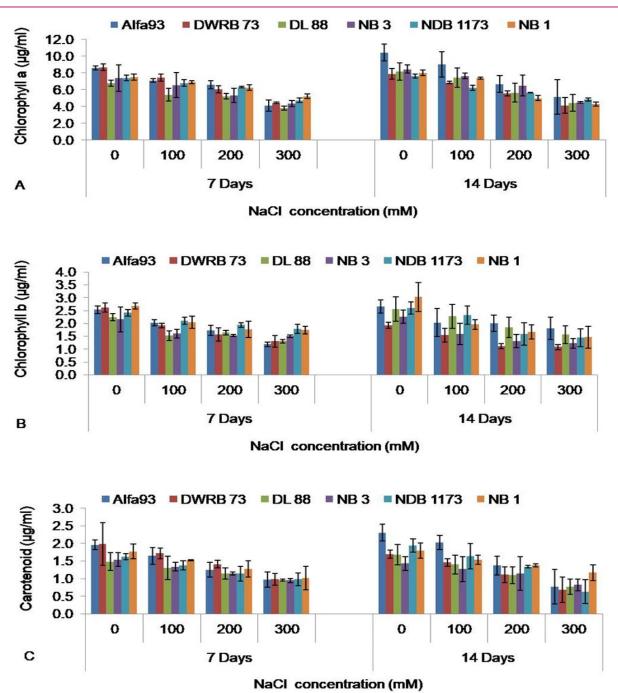
# Antioxidant enzyme activities

The antioxidant enzyme activities of the six barley cultivars under control and salinity-stressed conditions are shown in Fig. 3. The SOD levels under control condi-

tions were lower than stressed levels for both 7 days and 14 days. SOD activity was significantly lower in Alfa93 and DWRB73 than in the DL88, NB1, NB3, and NDB1173 barley lines (Fig. 4). Salinity had variable effects on SOD activity in all tolerant and susceptible cultivars.Salt tolerant lines demonstrate an increasing trend of SOD activity along with susceptible lines but different activity levels. Susceptible lines Alfa93 and DWRB73 under 300 mM salinity stress showed an increase of 85% and 79% in 7 days, and 96% and 108% in 14 days respectively in comparison to control. Among salt-tolerant lines, NB3 had shown the highest activity under 300 mM salt stress than control, 110% activity in 7 days, and 140% activity in 14 days. APX activity increased with an increase in salinity stress. After 7 days of treatment, a 60% increase in activity was observed in tolerant line NDB1173 and then NB1, in which activity was increased to 46.2%. After 14 days,Alfa93 and DWRB73 exhibit 76% and 36.9% increase in APX activity, whereas NDB1173 exhibits 100.5% increased activity. At 300 mM NaCl, CAT activity is 91% and 50.3% higher in DWRB73 and Alfa93 than in control plants. However, NB1 showed the highest 83.4% increase in CAT activity in 300 mM NaCl treated plants in salt-tolerant lines. Salinity stress increases the POX activity also: salt-tolerant lines exhibit more POD activity than susceptible lines. After 14 days, salt-tolerant lines DL 88, NB1, NB3, and NDB173, exhibits POD activity were recorded 39.3%, 65.8%, 42.7%, and 33.7% respectively under 300 mM NaCl treatment than control plants after 14 days.

Salinity stress induces ROS accumulation in plants, affecting membrane integrity and other cellular components that resulted in reduced growth and development (Tuna *et al.*, 2007). However, plant defence systems readily mitigate the salinity-induced ROS by enhanced antioxidants such as SOD, POD, CAT, and APX (Noreen *et al.*, 2021).

APX activity increased by 76% and 36.9% in Alfa93 and DWRB73 after 14 days of salt treatment, whereas in NDB1173 (ST) increase in activity was 100.5%. CAT activity in DWRB73 and Alfa93 was 91% and 50.3% higher than the control plants at 300 mM salinity stress. The results of CAT activity in leaves of salinity susceptible cultivars Alfa93 and DWRB73 of the present study are in disagreement with the result of Elsawy et al. (2018), they reported no significant difference in CAT activity in leaves of control and salt-treated barley cultivar Giza 126. In line with our results, Abdel Latef et al. (2019) reported a gradual increase in POD, CAT, and APX in two Egyptian wheat cultivars, Gemmiza 11 (SS), Misr 1 (ST), when salinity stress increases. CAT activity was high in salt-sensitive cultivar Gemmiza 11 under salinity stress, while POD and APX were higher in tolerant cultivar Misr1. Two contrasting Egyptian wheat cultivars, Sakha 95 and Misr 2, exhibit high activ-



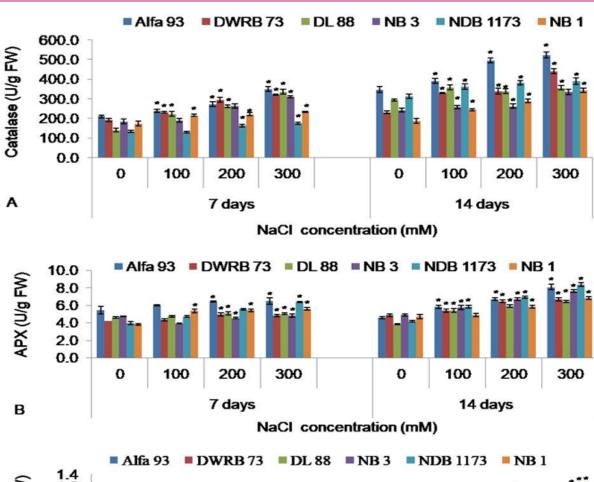
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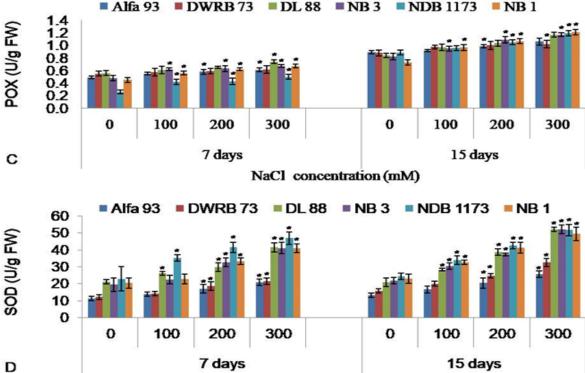
**Fig. 3.** The effect of salinity stress treatment on photosynthetic pigments (A) Chlorophyll a (B) Chlorophyll b and (C) Carotenoid in leaves of barley cultivars under control and saline (0 mM, 100 mM, 200 mM, and 300 mM NaCl) conditions after 7 days and 14 days.

ity of antioxidant enzymes under salinity stress of 150 mM NaCl than control plants (Yassin *et al.*, 2019). In the present study, the enzymatic activities of the salt-tolerant lines DL88, NB1, NB3, and NDB1173 were higher in salinity-treated plants than in control plants. Also, similar pattern was found in susceptible lines Alfa93 and DWRB73. When comparing the enzymatic activities between tolerant and susceptible lines SOD, POD, APX were high in tolerant lines. However, CAT activity was recorded high in susceptible lines Alfa93 and DWRB73.

# Principal component analysis of physiological parameters

Principal component analysis of the studied physiological parameters studied under salt-stressed conditions was done for the barley genotypes. All parameters were loaded into two major principal components (F1 and F2), which described the cumulative variance of 78.6, 78.8, 86.4, and 79.7% in control, 100 mM, 200 mM, and 300 mM, respectively, after 7 days of salt treatment. Similarly, the cumulative variance after 14 days were 69.4, 69.5, 66.5, and 80.0% in control, 100





NaCl concentration (mM)

**Fig. 4.** Effect of salinity stress treatment on (A) CAT, (B) APX, (C) POX, and (D) SOD activities in leaves of barley cultivars under control and saline (0 mM, 100 mM, 200 mM, and 300 mM NaCl) conditions after 7 days and 14 days. Values are mean  $\pm$  SD of three replicates, and asterisks denote significant differences from controls (P < 0.01)

mM, 200 mM, and 300 mM, respectively. After 7 days of treatment, PC1 accounted for 57.7% of the variation in the PCA plot of controls and was positively correlated with RWC and SOD. However, photosynthetic pigments (Chl-a, Chl-b, and CRT), REL, and antioxidant enzymes except SOD were negatively correlated. PC2 accounted for 20.9% of the variation and was positively affected by photosynthetic pigments (Chl-a, Chl-b, and CRT), REL, and RWC but negatively affected by the antioxidant enzymes. Upon comparing the PCs of control and 300 mM, NaCl treated samples of 7 days and 14 days, a cumulative variance of control was 78.6% and 69.4 respectively, whereas, 300 mM NaCl treated sample was 79.8% and 80.0%, respectively (Table 3, Fig. 5 and 6).

The PCA also demonstrated the different responses of barley genotypes under salt stress conditions. The biplot analysis signifies that the antioxidant enzyme SOD and APX were positively associated with NB1 and NDB1173, POX and CAT associated with DL88 and NB3 in 300 mM salt-treated plants for 7 days. When analyzed, the biplot of 14 days salt-treated plants POX and SOD were associated with NDB1173, APX, and CAT was associated with Alfa93. According to Guellim et al. (2020), genotype position from the centroid described the tolerance of plants to the stress, least the distance more the tolerance. The present study showed that the centroid position of Alfa93 and DWRB73 was at a distant position thanNB1, NDB1173, NB3, and DL88 in 14 days salt-stressed plants. Barley genotypes were ranked according to their tolerance level after 14 days of salinity treatment DL88 was the most tolerant than NB3, NB1, and NDB1173, least tolerant was DWRB73.

The principal component analysis enabled recognizing the physiological traits associated with the salinity stress and representing the level of salt stress tolerance among the genotypes. In the present study, antioxidant enzymes activity under salt stress and other physiological parameters such as REL, RWC, and photosynthetic pigments positively correlated with the tolerant genotypes. The result of the present study is also in agreement with the results of Ahmadi *et al.* (2020) who evaluated the physiological and biochemical response of wheat genotypes. PC analysis revealed the and tolerant genotypes exhibit enhanced responses to salinity stress.

Pour-Aboughadareh *et al.* (2020) reported that the antioxidant enzyme activity was positively correlated with the stress tolerance of the plant. In the present study, antioxidant enzyme activity was increased when barley plants were subjected to salinity stress. Higher activity was observed in tolerant lines than susceptible lines except for CAT activity. This enhanced enzyme activity may be used as a marker for screening of the salinity tolerance in the plants.

### Conclusion

This study indicated the effects of 7 days and 14 days prolonged salinity stress on barley plants (Alfa93, DWRB73, DL88, NB1, NB3, NDB1173) and their impact on physiology like decreased growth, loss in biomass, changes in photosynthetic pigments, and biochemical activities like antioxidant enzymes activity. The findings illustrated salinity stress, stress level, reduced growth, photosynthetic pigments, and antioxidant enzyme activity. Therefore, the elevated activity of antioxidant enzymes in salinity-stressed barley plants may be the mechanism of plants to tolerate the stress. The antioxidant enzyme (SOD) initiated the antioxidative process, which the CAT, POD followed. Plant's response in the form of changes in physiological as well as biochemical activity suggested that the strategies were adopted by the plants to mitigate stress. Exploiting these characteristics of the plants in the screening of tolerant and susceptible lines incorporates them in selecting lines for cultivation in salt-affected areas.

Table 3. Eigenvalues, variability (%), and cumulative (%) of PC1 (F1) and PC2 (F2) axes of PCA of physiological parameters.

	0mM NaCl		100mM NaCl		200mM NaCl		300mM NaCl		
	F1	F2	F1	F2	F1	F2	F1	F2	
Eigen value	3.829	2.413	4.063	2.192	3.686	2.295	4.821	2.381	-
Variability (%)	42.549	26.810	45.141	24.356	40.960	25.497	53.571	26.453	14 Days
Cumulative %		69.360		69.497		66.456		80.025	
Eigen value	5.192	1.885	4.127	2.962	4.803	2.969	4.565	2.612	
Variability (%)	57.692	20.940	45.858	32.908	53.370	32.984	50.718	29.027	07 Days
Cumulative %		78.632		78.766		86.354		79.745	

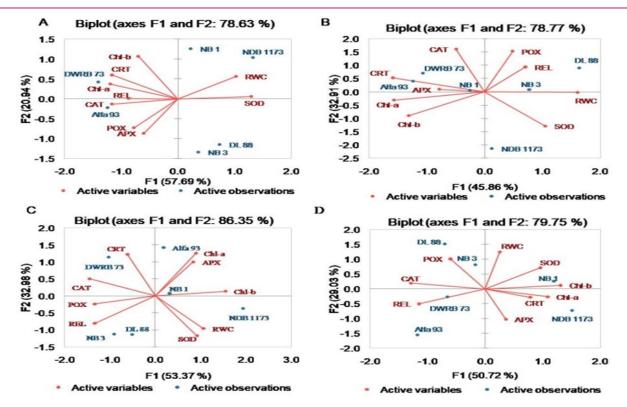
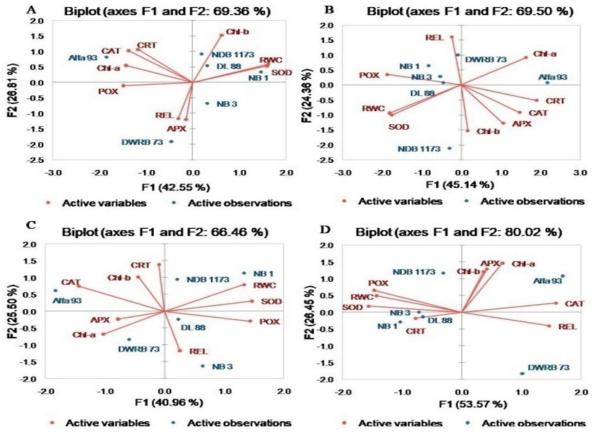


Fig. 5. Principle Component Analysis (PCA) Biplots of 7 days salt stress treated samples (A) Control, (B) 100 mM NaCl, (C) 200 mM NaCl, (D) 300 mM NaCl.



**Fig. 6.** Principle component analysis (PCA) Biplots of 14 days salt stress treated samples (A) Control, (B) 100 mM NaCl, (C) 200 mM NaCl, (D) 300 mM NaCl.

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#### **Conflict of interest**

The authors declare that they have no conflict of interest.

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