RESPONSES OF BARLEY SEEDLINGS TO SALINITY AND DROUGHT UNDER FREEZE-THAW CONDITIONS

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Abstract. The Qinghai-Tibet Plateau is known for its high altitude, low rainfall and varying temperature, and the crops in this area are susceptible to abiotic stresses induced by drought, salinity and freeze-thaw conditions that cause damages to different properties such as the permeability of biological membrane, osmotic adjustment, and antioxidant enzyme system. Barley (Hordeum vulgare L.) is an indispensable crop on the plateau and plays an important role in agricultural ecosystem as well. In this study, Beiqing 3 was used as experimental material and physiological characteristics, including soluble protein (SP) content, malondialdehyde (MDA) content, antioxidant enzyme activity and relative water content (RWC) of seedlings were examined under freeze-thaw conditions combined with drought and alkali stress. Research results indicated that under the combined stresses of salinity and drought, barley seedlings were damaged by lipid peroxidation, weakened superoxide dismutase (SOD), catalase (CAT) and peroxidase (POD) activities, while osmotic adjustment ability in plants cell was enhanced. We suggested that, in agricultural management, the simultaneous occurrence of two stresses, salinity and drought, should be avoided in the early stage of barley planting to reduce the physiological stress on plants.

Keywords: abiotic stresses, antioxidant enzyme activity, soluble protein, physiological effects, malondialdehyde

Introduction

Lying in the Southern Qinghai-Tibet Plateau, the Brahmaputra Valley, where altitude varies from 2600 m to 3500 m, was known for the complex and changeable terrain, forming a unique plateau climate characteristic (Zhang et al., 2019). The fact that, during early spring, the freeze-thaw often occurs due to the long sunshine hours and the changeable temperature between day and night has various effects on the external morphology and internal physiological metabolism of plants (Arfan et al., 2019; Wang et al., 2019; Xu et al., 2019). It has been reported that the average precipitation of barley planting region is less than 400 mm, besides, 90% of the annual precipitation occurs in June to September, which tends to cause spring drought and affect spring sowing and early crop growth (Hou et al., 2018). The seasonal uneven precipitation and the exceeding evaporation on the Brahmaputra Valley make drought stress prone to occur during crop growth (Bibi et al., 2019), which often causes oxidative stress by the accumulation of reactive oxygen species in plants (Souza et al., 2004), affecting the structure and growth of plant (Duan et al., 2007). Attributing to current climate, the content of sodium bicarbonate (NaHCO₃) and sodium chloride (NaCl) in soil has increased on the
Brahmaputra Valley, where salinization becomes increasingly serious. It has been confirmed that alkali stress imposed more harm to crops than salt stress does (Alvarez-Acosta et al., 2019). Besides, one paper has previously reported that high pH of alkali stress can short root length and seedlings height of rye, reduce the content of water and chlorophyll and decrease the relative transpiration rate (Guo et al., 2012).

Barley, a cereal crop of the genus Gramineae, can grow at an altitude of 3000 ~ 3400 m. Among them, Beiqing 3 has good resistance to cold and drought (Ahmed et al., 2015; He et al., 2015), is the main crop in Tibet and Qinghai. In this experiment, as materials, the Beiqing 3 seedlings were treated with salinity, drought and freeze-thaw stress to artificially simulate the growing environment of plants. The relative water content (RWC), antioxidant enzyme activity, contents of malondialdehyde (MDA) and soluble protein (SP) were examined in order to study the response characteristics of plants to drought, salinity and freeze-thaw.

Materials and methods

Seeds cultivation and salinity treatment

The study was taken out in Northeast China. The full-grained seeds were selected and soaked with 0.1% KMnO₄ solution for 2 h for disinfection, after which the seeds were rinsed with deionized water until the water becoming clear, then we spread 120 seeds evenly on each of 8 culture dishes randomly named FSD, FS, FD, F, SD, S, D and C (Table 1). 1/2 of Hoagland nutrient solution was used to prepare 60 mM NaHCO₃ mixed solution (pH = 8.06), 500 ml of which was added to the cultivated dishes of FSD, FS, SD and S at the same time, 500 ml 1/2 Hoagland nutrient solution was added to the others (FD, F, D and C). 8 dishes of seeds were placed in MGC-450BP light incubator (Shanghai Yiheng Scientific Instruments Co., Ltd) for germination (Fig. 1a), of which the cultivated conditions were set as 12 h light (25 ℃) and 12 h non-light (15 ℃). Daily watering (50 ml) was necessary during the cultivation.

Table 1. Experimental design of groups under salinity (S), drought (D) and freeze-thaw (F) stress

|                | FSD | FS | FD | F  | SD | S  | D  | C  |
|----------------|-----|----|----|----|----|----|----|----|
| Salinity       | +   | +  | -  | -  | +  | +  | -  | -  |
| Drought        | +   | -  | +  | -  | +  | -  | +  | -  |
| Freeze-thaw    | +   | +  | +  | +  | -  | -  | -  | -  |

+ add stress, - no stress

Drought treatment

After seedlings were cultivated to 15 cm high with 2 or 3 leaves (around 1 week), they were treated with drought stress. NaHCO₃ mixed solution was used to prepare 20% PEG-6000 mixed solution for combined treatment of salinity and drought stress, and 1/2 Hoagland nutrient solution was used to prepare 20% PEG-6000 solution. The solution in the cultivated dishes of Group FSD and Group SD was replaced with 500 ml PEG-NaHCO₃ mixed solution, in the cultivated dishes of Group FD and Group D replaced with 500 ml PEG solution, in the cultivated dishes of Group FS and Group S
replaced with 500 ml NaHCO$_3$ solution, in the cultivated dishes of Group F and Group C replaced with 500 ml 1/2 Hoagland nutrient solution. The drought treatment lasted for 48 h without watering.

**Figure 1.** Photos of experimental equipment. (a): MGC-450BP light incubator (Shanghai Yiheng Scientific Instruments Co., Ltd). (b): BPHJ-120A high-low-temperature test chamber (Shanghai Yiheng Scientific Instruments Co., Ltd)

**Freezing and thawing stress treatment and sampling**

After drought treatment, the cultivated dishes of Groups FSD, FS, FD and F were put into BPHJ-120A high-low-temperature test chamber (Shanghai Yiheng Scientific Instruments Co., Ltd) to carry out a freeze-thaw cycle for a period of 14 h (Fig. 1b), with the temperature curve being set as 15, 10, 5, 0, −5, 0, 5 and 10 ℃, while other cultivated dishes of Groups SD, S, D and C were maintained in light incubator under previous culture conditions (Fig. 2a). Initially, the cultivated dishes were placed in the chamber at 15 ℃ that closed to room temperature at night. Controlled precisely by program, the temperature decreased to −5 ℃ steadily at a speed around 0.04 ℃/min, and then the temperature increased from −5 to 10 ℃ at a speed around 0.04 ℃/min. After the freeze-thaw cycle being started, five parallel samples were taken every 2 hours from 8 cultivated dishes at random according to the required amount of the measurement, the corresponding sampling temperature was 10, 5, 0, −5, 0, 5, 10 ℃ respectively (Gong et al., 2020). All the samples were firstly wrapped up with tin foil paper, secondly fixed in liquid nitrogen immediately for 50 s and finally put into the ultra-low-temperature freezer at −80 ℃ for storage in order to measure the content of MDA and soluble protein, SOD, POD and CAT activity. At the same time, fresh leaves were taken to determinate RWC.
Figure 2. Photos of experimental culture and equipment. (a): 9-day barley seedlings in light incubator. (b): UV-6100 UV–visible spectrophotometer (Metash Co. Ltd)

Analysis

Relative water content (RWC)

The relative water content of seedlings was determined by the oven drying method (Colom and Vazzana, 2001). For each sample (around 0.1 g), fresh weight supposed to be measured and recorded as $F_W$ after drying the surface of leaves with filter paper. Completely being immersed in distilled water until the weight of leaves being constant, the leaves were taken out and wiped up with filter paper. The saturated fresh weight of the leaves at this time was measured and recorded as $T_W$. Finally, the leaves were de-enzymed for 15 min in oven that was heated up to 105 °C, and then dried to a constant weight in 80 °C. The dry weight was measured and recorded as $D_W$. The RWC of leaves is calculated by formula $Eq. 1$:

$$RWC = \frac{(T_W - D_W)}{(F_W - D_W)} \times 100\% \quad (Eq.1)$$

Soluble protein (SP) content

The soluble protein content in seedlings was determined by the Coomassie brilliant blue method (Kong and Yi, 2008). 0.1 g leaves were selected randomly and shredded into a mortar, and then ground until homogenized with 5 ml distilled water, which next was
centrifuged with a TDL-40B centrifuge (Anting Scientific Instrument Factory, Shanghai) at a speed of 3000 r/min for 10 min. 1 ml of the supernatant was diluted in 5 times with 4 ml distilled water, of which 1 ml diluted supernatant was taken into a test tube with 5 ml of Coomassie brilliant blue solution being added. After the mixed solution being shaken and placed for 2 min, the absorbance of the solution was measured at 595 nm with a UV-6100 UV–visible spectrophotometer (Metash Co. Ltd) (Fig. 2b). The soluble protein content was calculated by standard curves.

**Malondialdehyde (MDA) content**

Malondialdehyde (MDA) content in seedlings was determined by the thiobarbituric acid method (Kong and Yi, 2008). 0.5 g leaves were selected randomly and shredded into a mortar, and then ground into a homogenate with 5 ml 10% trichloroacetic acid (TCA) solution, which next was centrifuged at a speed of 4000 r/min for 10 min. Then 2 ml of the supernatant into was taken and fixed with 2 ml 0.6% thiobarbituric acid (TBA) solution. Mixtures was bathed in 99 °C water for 15 min, then cooled quickly in 5 min and centrifuged again at a speed of 4000 r/min for 10 min with a TDL-40B centrifuge (Anting Scientific Instrument Factory, Shanghai). The absorbance of supernatant was measured at 532 nm, 600 nm, and 450 nm with a UV-6100 UV–visible spectrophotometer (Metash Co. Ltd). The MDA concentration and MDA content were calculated according to formulas Eq.2 and Eq.3.

\[
MDA \text{ concentration (μmol/L)} = 6.45 \times (D_{532} - D_{600}) - 0.56 \times D_{450} \quad (\text{Eq.2})
\]

\[
MDA \text{ content (μmol/g)} = c_{MDA} \times V_T / F_W \quad (\text{Eq.3})
\]

where:
- \(D_{450}, D_{532}, D_{600}\) are the absorbance at 450nm, 532nm and 600nm, respectively.
- \(c_{MDA}\) is MDA concentration (μmol/L);
- \(V_T\) is the volume of TCA solution (ml);
- \(F_W\) is the fresh weight of seedlings (g).

**Catalase (CAT), superoxide dismutase (SOD) and peroxidase (POD) activities**

The activities of CAT, SOD and POD were determined with the CAT, SOD and POD kits provided by Nanjing Jiancheng Biological Institute (Bao et al., 2017). A parallel sample (around 0.25 g) were randomly taken and ground to a homogenate with 5 ml phosphate buffer on ice. After centrifugations at a speed of 2500 r/min, 3500 r/min and 3500 r/min with a TDL-40B centrifuge (Anting Scientific Instrument Factory, Shanghai), respectively for 10 min, the supernatant was used for following measurements according to instructions of kits.

**Data processing**

The experiments were repeated five times, and the data were expressed as mean ± standard error (SE) (n=5), which statistically performed with R 3.3.1 statistical software (R Foundation for Statistical Computing, Vienna, Austria) for one-way analysis of variance (ANOVA). When the variables were uniform, the significance analysis of data was analyzed using Duncan model, otherwise using Games-Howell model (Warner 2007). Pearson correlation coefficient was used to describe the correlation between
variables. All results were shown in bars in figures plotted by Origin 8.0 software. Different letters presented in figures indicated significant differences between different treatment groups at the same time.

**Results**

**Effect on relative water content of seedlings**

In this experiment, the relative water content (RWC) in of 8 treatment groups decreased during freeze-thaw cycle. As shown from figure Fig. 3, RWC in barley seedlings had a maximum decrease under combined stresses of salinity and drought. The RWC in seedlings of Groups FS, FD and F had no significant differences compared to that in Groups S, D and C, respectively. However, in thawing stage, RWC in seedlings of Group FSD showed significant differences compared with that of Group SD. Notably, the RWC in barley seedlings of 4 freeze-thaw groups showed a sequence as F > FS > FD > FSD. Consistently, a similar order of RWC in seedlings can be observed among 4 non-freeze-thaw groups as well, that is, C > S > D > SD.

![Figure 3. The relative water content in barley seedlings under different treatment. The letter F, S, D and C represent freeze-thaw stress, salinity stress, drought stress and control, respectively. The temperature 10, 5, 0, -5, 0, 5 and 10 °C means the corresponding sampling temperature. The different low-case letters mean the significant difference at the same temperature (P < 0.05)](image-url)

**Effect on soluble protein content of seedlings**

It can be observed that the soluble protein (SP) content in barley seedlings of 4 treatment groups under freeze-thaw stress (FSD, FS, FD and F) was higher than that of 4 treatment groups without freeze-thaw stress (SD, S, D and C), respectively (Fig. 4). The SP content in seedlings of Group SD and Group S was significantly higher than that of Group D and Group C (P < 0.05), which indicated that under non-freeze-thaw conditions, the SP content in seedlings increased due to the occurrence of salinity stress.
Nevertheless, the SP content in seedlings of groups under either single freeze-thaw stress or single drought stress had no significant difference compared with that of control group ($P > 0.05$). In the case of freeze-thaw stress, the SP content in seedlings of Groups FSD, FS, FD and F increased during the period of freeze-thaw stage. Among them, Group FSD reached the maximum at 0°C (thawing stage), which exhibited a further 41.2% increase than the minimum 0°C (freezing stage). Somewhat differently, the other 3 groups (FS, FD and F) all reached the maximum value at 5°C (thawing stage), and were 46.4%, 86.6% and 72.7% higher than the minimum value, respectively.

![Figure 4](image_url)  
*Figure 4. The soluble protein content in barley seedlings under different treatment. The letter F, S, D and C represent freeze-thaw stress, salinity stress, drought stress and control, respectively. The temperature 10, 5, 0, -5, 0, 5 and 10 °C means the corresponding sampling temperature. The different low-case letters mean the significant difference at the same temperature ($P < 0.05$)*

**Effect on MDA content of seedlings**

*Figure 5* shows that the malondialdehyde (MDA) content in barley seedlings of all experimental groups was higher than that of the control group. The non-freeze-thaw groups (SD, S, D and C) fluctuated little during a 14-hour freeze-thaw period, however the MDA content in barley seedlings of Group SD and Group S was significantly higher than that of Group D and C ($P < 0.05$). Besides, we have noticed that MDA content in barley seedlings in response to Group D was significantly higher than that in response to blank treatment ($P < 0.05$). Under the freeze-thaw stress, the MDA content in barley seedlings of single freeze-thaw group (F) was significantly lower than that of Groups FSD and FD ($P < 0.05$).

**Effect on SOD activity**

The SOD activity in barley seedlings of Groups FSD and SD had no significant difference compared with control group (C) (*Fig. 6*). The SOD activity significantly enhanced owing to the occurrence of drought stress in barley seedlings ($P < 0.05$), while significantly weakened due to the salinity stress in seedlings ($P < 0.05$). Under freeze-
thaw stress, the SOD activity in barley seedlings of Groups FSD, FS, FD and F decreased at first and then increased. During the freeze-thaw cycle, except for 5°C, the SOD activity in barley seedlings of Group FD was significantly higher than that of Groups FSD, FS and F ($P < 0.05$).

**Figure 5.** The malondialdehyde (MDA) content in barley seedlings under different treatment. The letter F, S, D and C represent freeze-thaw stress, salinity stress, drought stress and control, respectively. The temperature 10, 5, 0, -5, 0, 5 and 10 °C means the corresponding sampling temperature. The different low-case letters mean the significant difference at the same temperature ($P < 0.05$).

**Figure 6.** The SOD activity of barley seedlings under different treatment. The letter F, S, D and C represent freeze-thaw stress, salinity stress, drought stress and control, respectively. The temperature 10, 5, 0, -5, 0, 5 and 10 °C means the corresponding sampling temperature. The different low-case letters mean the significant difference at the same temperature ($P < 0.05$).
**Effect on CAT activity**

It can be seen from Figure 7 that, under non-freeze-thaw conditions, the CAT activity in barley seedlings of Groups SD, S and D was significantly lower than that of the control group \((P < 0.05)\). CAT activity in seedlings of either Group F or Group FS showed a trend of initially increasing and then decreasing, while that of Group FD showing a general downward trend, and that of Group FSD showing an upward trend. Take if further, the CAT activity in seedlings of Group FSD was significantly lower than that of Group F \((P < 0.05)\), and that of the Groups FS and FD was significantly lower than that of Group F only in the latter thawing case \((P < 0.05)\).

![Figure 7. The CAT activity in barley seedlings under different treatment. The letter F, S, D and C represent freeze-thaw stress, salinity stress, drought stress and control, respectively. The temperature 10, 5, 0, -5, 0, 5 and 10 °C means the corresponding sampling temperature. The different low-case letters mean the significant difference at the same temperature \((P < 0.05)\).](image)

**Effect on POD activity**

In this experiment, during freeze-thaw cycle, the POD activity in barley of Groups FD and F showed a trend of initially increasing and then decreasing, while that of Groups FSD and FS showing an increasing trend \((\text{Fig. 8})\). When the temperature dropped to 10 °C (freezing stage), the POD activity in barley seedlings of Groups F, S and D were significantly lower than that of control group \((P < 0.05)\). Under salinity stress, the POD activity in seedlings significantly decreased within a freeze-thaw cycle \((P < 0.05)\). Nevertheless, no significant difference was observed in POD activity between Groups FD and D. Accordingly, under freeze-thaw stress, there was a significant reducing of POD activity of groups subjected to salinity treatment, but no effect on that of groups subjected to drought treatment.

**Correlation analysis**

It can be observed from Table 2 that the content of SP and MDA in seedlings were significantly positively correlated under freeze-thaw conditions \((P < 0.01)\). There was a...
significant negative correlation between SP content and antioxidant enzyme activity ($P < 0.01$). MDA was significantly negatively correlated with antioxidant enzyme activity and RWC, while CAT and POD were positively correlated ($P < 0.01$).

**Figure 8.** The POD activity in barley seedlings under different treatment. The letter F, S, D and C represent freeze-thaw stress, salinity stress, drought stress and control, respectively. The temperature 10, 5, 0, -5, 0, 5 and 10 °C means the corresponding sampling temperature. The different low-case letters mean the significant difference at the same temperature ($P < 0.05$)

**Table 2.** Pearson correlation analysis between relative water content (RWC), soluble protein (SP) content, malondialdehyde (MDA) content, SOD, CAT and SOD activity in barley seedlings of freeze-thaw treatment groups

|         | RWC   | SP     | MDA    | SOD    | CAT    | POD    |
|---------|-------|--------|--------|--------|--------|--------|
| RWC     | 1     | -0.524** | -0.427* | -0.341 | 0.726** | 0.640** |
| SP      |       | 1      | 0.750** | -0.351 | -0.799** | -0.868** |
| MDA     |       |        | 1      | -0.407* | -0.663** | -0.798** |
| SOD     |       |        |        | 1      | 0.052  | 0.213  |
| CAT     |       |        |        |        | 1      | 0.833** |
| POD     |       |        |        |        |        | 1      |

* Significant correlation at 0.05 level (both sides). ** Significant correlation at the 0.01 level (both sides)

**Discussion**

Either drought or alkaline salt can lead to a large amount of water loss in seedlings by reducing the osmotic pressure of plant cells (Shereen et al., 2019). It is one important conclusion from Alexander’s research that the RWC of leaves is positively related to plant’s stress resistance (Alexander et al., 2019). It has been reported that under the combined effects of water loss and high temperature, the reduction of RWC in Australian durum is greater than that under single stress (Liu et al., 2019). Consistently, in this
experiment, the combined effects on RWC in barley seedling of salinity and drought were severer than the additive of salinity and drought alone; thus, interactions between the two stress factors were synergistic for RWC. Under freeze-thaw conditions, a decrease in temperature can not only ice the water in plant cells, but also caused a dehydration, resulting in a decrease in RWC (Iseri et al., 2013). As have shown that at the freezing stage, the decreased RWC contributed to alleviate the damage caused at freezing stage to plants and maintain the osmotic balance of plant cells (Hao et al., 2009).

Most of the soluble proteins in plants are enzymes involved in metabolism (Bao et al., 2019). An increase of soluble proteins can maintain the cell's higher osmotic potential, enhance the capacity of water absorption and holding, maintaining plant growth and improving resistance to stress (Yin et al., 2004). Here we observed from experiments that the SP content of barley seedlings increased under either drought or freeze-thaw stress (Groups F and D), while a higher accumulation of SP was measured in the groups subjected to basic-salt stress (FSD, FS, SD and S). These observations may attribute to the expression of resistant proteins in plant cells stimulated by alkaline stress, increasing the content of SP participating in osmotic adjustment in cells, thus making plants adapt to the external environment (Hazman et al., 2016). Under freeze-thaw stress, the SP content in seedlings of groups subjected to freeze-thaw stress decreased with the dropping temperature, which may be due to the accelerated decomposition of soluble proteins in cells, providing plants with energy to relieve the damage caused by stresses (Bae et al., 2006). At thawing stage, with the temperature rising to 10 ℃, the SP content in seedlings increased. These findings are similar to the results of Lee’s research, in which they examined the proteomic changes of rice roots under low temperature stress and found that the expressions of 27 proteins were up-regulated at 10 ℃ (Lee et al., 2009).

The activity changes of antioxidant enzyme in plants caused by abiotic environmental stresses may have an effect on physiological characteristics to reduce damage (Ahsan et al., 2007). In a previous study, Zeng et al. (2019), using methods of indoor cultivating of soybean seedlings and experiments, disclosed that a large amount of CAT transcription and significant enhancement of enzyme activity were observed under high aluminum stress. Researches have shown that the antioxidant enzyme activities in leaves of *Pyracantha fortuneana* and *Rosa cymosa* are significantly enhanced under severe drought stress, indicating the strong resistance to drought stress of these species, however the antioxidant enzyme activities are greatly weakened in leaves of *Broussonetia papyrifera* and *Cinnamomum bodinieri* under same conditions, indicating the weaker resistance to drought stress (Liu et al., 2011). Here, our experiment showed that CAT and POD activities in barley seedlings significantly weakened under non-freeze-thaw stress. The results suggested a possible reason of lipid peroxidation on the cell membrane affected by stresses duration, leading to the damage to cells and the effect on the synthesis of substances like proteins in cells, at last reducing the antioxidant enzyme activity. This is consistent with the study by Gao et al. (2012). Moreover, an observed decrease in SOD activity in the leaves of *Camptotheca acuminata* seedlings accompanied low temperature stress, which was found by Feng et al. (2002). It was worth noting that CAT and POD activities increased, while SOD activity decreased with a decrease of temperature, which could be explained by the role played by SOD as the first line in defense and eliminating reactive oxygen species (ROS). A large consumption of SOD in the process of eliminating ROS and an inefficient synthesis of enzymes in the case of low temperature were confirmed in the research results of Bao et al. (2019).
Environmental stress can disrupt the homeostasis of cells and the dynamic balance between production and clearance of ROS, leading to excessive accumulation of ROS in cells, causing the oxidative damage to biomolecules such as lipids (Mano, 2012), proteins (Dean et al., 1997) and nucleic acids (Cadet et al., 2003), and the disruption of osmotic balance in plants (Bian et al., 2018), which were discussed in detail in numerous studies. MDA is the end product of lipid peroxidation and can be induced by stress in plants organ, e.g. leaves, shoots or roots (Iseri et al., 2013; Karagoz et al., 2018). In this paper, MDA content in barley seedlings increased under the single or combined stress of salinity and drought, importantly, MDA content accumulated more under single salinity stress than that under combined stress. In addition to osmotic stress on seedlings, salinity stress, compared with drought, is accompanied by high pH stress as well, causing damage to plant cell membranes and eventually leading to the accumulation of MDA, which has been confirmed in a study by Ali et al. (2011). The freeze-thaw manipulation treatments decreased the MDA content of barley seedlings, during which, the activities-enhanced CAT played a key regulatory role (Wu et al., 2018).

Conclusion

In summary, as an important crop on the Qinghai-Tibet Plateau, barley has equipped with great resistance to the freeze-thaw environment within the long-term evolution. Considering current global warming, soil salinization and drought have become increasingly serious, resulting in physiological responses like the accumulation of MDA and the changes in antioxidant enzymes activity. Herein we showed that either single or compound stresses of drought, salinity and freeze-thaw could make MDA accumulated excessively in seedlings because of the imbalance between oxygen free radical reaction and lipid peroxidation reaction, which caused oxidative stress on plants, affecting the stability of plant cells. Moreover, the contents of MDA and SP in barley seedlings increased significantly under the combined stresses of salinity and drought, while the RWC significantly reduced. As a conclusion, to avoid simultaneous occurrence of drought and salinity stress, the intensity of spring irrigation is supposed to increase in areas with severe drought stress. Though the resistance characteristics of plants under one freeze-thaw cycle were studied in this paper, in view of multiple freeze-thaw cycles in nature, one important future direction of physiological responses to freeze-thaw stress is studying the different resistance characteristics of plant between under one freeze-thaw cycle and under multiple freeze-thaw cycles.

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