Efects of Cold and Salicylic Acid Priming on Free Proline and Sucrose Accumulation in Winter Wheat Under Freezing Stress

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Abstract

This study aims to investigate the efects of cold and salicylic acid (SA) priming on osmolytes accumulation in wheat leaves under freezing stress and its underlying physiological mechanism. The results showed that cold and SA priming treatment significantly enhanced sucrose and free proline contents as compared with non-priming treatment under freezing, resulting in increased leaf water potential, reduced cell death and boosted freezing tolerance. Cold and SA priming-induced free proline accumulation under freezing not only depended on promoting its synthesis, but also on inhibiting its degradation. Interestingly, the synthesis and hydrolysis of sucrose were both increased by cold and SA priming treatment under freezing. Besides, cold and SA priming up-regulated the catabolism of glucose and the assimilation of ammonia as compared with non-priming treatment under freezing stress. Findings of the present study suggested that cold and SA priming could simultaneously promote-free proline and sucrose accumulation in wheat leaves by coordinating carbon and nitrogen metabolism under freezing conditions, and then conferring tolerance to freezing stress. These fndings could provide a new insight into the mechanisms by which cold and SA priming enhanced freezing tolerance in wheat.

Keywords Cold priming · Freezing stress · Free proline · Salicylic acid · Sucrose · Wheat

Introduction

Late spring frost is an important limiting factor in winter wheat production worldwide, including the USA (Gu et al. [2008](#page-11-0)), Australia (Zheng et al. [2015](#page-13-0)), Italy (Bascietto et al. [2019](#page-11-1)) and China (Zhong et al. [2008](#page-13-1)). However, while slightly counterintuitive, the probability and severity of spring frost are predicted to increase due to global warming

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(Gu et al. [2008;](#page-11-0) Ji et al. [2017\)](#page-11-2). In addition to thermodynamical inhibition of metabolic reactions, sub-zero temperatures can also cause other adverse efects such as reactive oxygen species (ROS) over-accumulation, cell dehydration and contraction (Ruellan et al. [2009;](#page-12-0) Shin et al. [2018\)](#page-12-1). Among them, dehydration stress is the main cause of deterioration of the intracellular structures and death of tissues induced by subzero temperatures (Ruellan et al. [2009\)](#page-12-0).

To cope with dehydration damage, wheat plants can induce the accumulation of compatible solutes (such as free proline, sucrose, glycine betaine and polyols) to support osmotic regulation and sustain cell turgor, which play an important role in maintaining water potential in the plant (Wani et al. [2018;](#page-13-2) Wang et al. [2019](#page-12-2)). In addition to their role in osmoregulation, free proline and sucrose can act as signaling molecules to regulate expression of stress-related genes, as ROS scavenger, and as nutrients to help plants recover from stresses (Couee et al. [2006](#page-11-3); Islam et al. [2021](#page-11-4); Szabados and Savourcb [2010\)](#page-12-3). It is well-documented that the accumulation of free proline and sucrose plays a critical role in winter wheat response to freezing stress (Dörfing et al. [1990](#page-11-5); Kamata and Uemura [2004](#page-11-6); Kovács et al. [2011](#page-11-7)).

It is interesting to note that the accumulation of sucrose and free proline often occur simultaneously in plant response to environmental stresses including cold, drought and salt (Cai et al. [2004;](#page-11-8) Ehsan et al. [2012;](#page-11-9) Perezalfocea and Larher [1995;](#page-12-4) Wang et al. [2019](#page-12-2)). Sucrose and proline metabolism are important components of carbon and nitrogen metabolism, respectively (Martins et al. [2020;](#page-12-5) Vágújfalvi et al. [1999](#page-12-6)). Several studies have demonstrated that there was a close relationship between carbon and nitrogen metabolism in plants, which was critical for plant growth and development (Bao et al. [2015;](#page-11-10) Gálvez et al. [1999\)](#page-11-11). It is reasonable to speculate that the coordination between carbon and nitrogen metabolism plays an important role in the simultaneous accumulation of sucrose and free proline in plant response to environmental stimulus.

Priming, defned as a temporal experience of an environmental stimulus (including exposure to the stress cues themselves as well as the exogenous application of biologically active chemicals), can efectively improve the plant's level of resistance to future stress events (Baier et al. [2019](#page-11-12); Zuther et al. [2019](#page-13-3)). Salicylic acid (SA) is an important phenolic derivative widely distributed in the plant kingdom and is known to be a crucial signalling molecule in response to abiotic and biotic stresses (Saleem et al. [2020;](#page-12-7) Sadiq et al. [2020;](#page-12-8) Zaid et al. [2019;](#page-13-4) Ahmad et al. [2019;](#page-11-13) Dong et al. [2014](#page-11-14)). Previous studies have shown that SA and cold temperature priming could efectively enhance wheat freezing tolerance (Sun et al. [2018](#page-12-9); Wang et al. [2020a\)](#page-12-10). Zuther et al. [\(2019\)](#page-13-3) indicted that the accumulation of sucrose and free proline was involved in the cold priming-induced tolerance to freezing in *Arabidopsis*. A series of studies showed that exogenous SA could promote accumulation of free proline in plant leaves under both normal and cold conditions (Min et al. [2018](#page-12-11); Ignatenko et al. [2019\)](#page-11-15). Exogenous SA decreased sucrose content in tomato leaves under normal condition (Poór et al. [2011](#page-12-12)), but increased sucrose content in spinach leaves under cold condition (Min et al. [2018\)](#page-12-11). However, the regulatory mechanisms of cold and SA priming on sucrose and free proline accumulation under freezing stress in wheat are poorly understood.

Therefore, we hypothesized that cold and SA priming can promote freeze-induced sucrose and free proline accumulation by intensifying the coordination between carbon and nitrogen metabolism, which is contributing to improving wheat freezing tolerance. To test this hypothesis, we investigated the effects of cold and SA priming on cell dehydration, free proline and sucrose accumulation, and key enzymes and metabolites involved in carbon and nitrogen metabolism under freezing stress in wheat leaves.

Materials and Methods

Experiment Design

The locally and widely planted winter wheat (*Triticum aestivum* L. cv Yangmai 16) was planted in plastic pots (22 cm in height and 25 cm in diameter), with a planting density of 7 plants per pot. The fertilizer and water management was carried out as described by Wang et al. ([2020b](#page-13-5)). Wheat plants were grown outdoor until the application of priming treatments.

At the beginning of the jointing stage (when the 1st node was detectable, Zadoks [2010\)](#page-13-6), one-third of the wheat plants were transferred into a growth chamber with a temperature of 16 °C/10 °C (day/night, set according to the ambient temperature) as control treatment (C); one-third of the wheat plants were primed with cold temperature (6 \degree C/2 \degree C, day/night) for 2 days as cold priming treatment (P); one-third of the wheat plants were primed with 100 μM SA (Sigma-Aldrich, St. Louis, MO) by foliage spraying for 2 days (twice a day with an interval of 12 h) under control temperature as SA priming treatment (S). Then, all the plants were grown under control temperature for recovery. After 8 days of recovery, half of each group of the plants were subjected to freezing stress for 2 days (the first day/night temperature was $2 \degree C/0 \degree C$, and the second day/night temperature was−2 °C/−4 °C), while the other half were kept at control temperature (16 °C/10 °C, day/night). Finally, six treatments composed: control (CC), SA priming without freezing stress (SC), cold priming without freezing stress (PC), freezing stress without priming (CF), SA priming plus freezing stress (SF) and cold priming plus freezing stress (PF). The light intensity of the chambers was set at 500 µmol m⁻² s⁻¹ with a photoperiod of 10 h. The latest fully expanded leaves were sampled at the end of freezing treatment for further biochemical and physiological analyses.

Freezing Tolerance Determination

The freezing tolerance of wheat leaves was determined by the LT_{50} (temperature at which 50% of ions in plant tissues leak out) according to Cai et al. ([2004](#page-11-8)) with slightly modifcations. Fresh leaves of C, S and P treatments at 8 days of recovery were cut into 1.5 cm lengths and wrapped in wet gauze, then placed in a glass tube and incubated at 4 °C for 2 h. After incubation, the leaves were treated at diferent freezing temperatures $(0, -3, -6, -9, -12, -12, -15)$ °C) for 2 h in a freezing bath (CDN-1007020F, Southeast Co. Ltd., Ningbo, China). Then, the leaves were thawed at 4 °C overnight and used to measure electrolyte leakage using a electrical conductivity bridge (DDS-307A, LEX Instruments Co. Ltd., Shanghai, China). The mean values of electrolyte leakage of each treatment were utilized to generate a sigmoid curve ftting the Logistic function, and the temperature corresponding to the infection point of the curve is the LT_{50} (Sigmaplot 10.0, Systat Software Inc., CA, USA).

Leaf Water Potential and Histochemical Staining

The WP4C Dewpoint PotentiaMeter (Decagon Devices Inc., WA, USA) was used to detect water potential of the latest fully expanded leaves following the manufacturer's instruction. Trypan blue and Evans blue staining were used to detect dead cells of the latest fully expanded leaves according to Koch and Slusarenko [\(1990](#page-11-16)) and Baker and Mock [\(1994\)](#page-11-17), respectively.

Metabolites Contents

The contents of free proline and sucrose were determined by Lu et al. ([2005](#page-12-13)) and Huber [\(1983\)](#page-11-18) method, respectively. Ammonium measurements were carried out as described by Husted et al. ([2000](#page-11-19)). Glutamate contents were analyzed using an L-8900 High Speed Amino Acid Analyzer (Hitachi Corp., Tokyo, Japan) as described by Zhong et al. ([2018](#page-13-7)). 2-oxoglutarate (2-OG) contents were determined using an ACQUITY UPLC H-Class (Waters Corp., MA, USA) according to Lee and Foy [\(1986\)](#page-12-14).

Enzyme Assays

The activities of sucrose-phosphate synthase (SPS) and sucrose synthase (SS, synthetic direction) were detected according to Wardlaw and Willenbrink ([1994](#page-13-8)). The activities of acid-invertase (Ac-INV) and alkaline/neutral-invertase (A/N-INV) were measured according to Miron and Schaffer ([1991\)](#page-12-15) and Vargas et al. ([2007\)](#page-12-16), respectively. The activities of Δ1-pyrroline-5-carboxylate synthetase (P5CS) and proline dehydrogenase (PDH) were determined according to Mario et al. [\(1997\)](#page-12-17) and Rena and Splittstoesser ([1975](#page-12-18)), respectively. The activities of glutamine synthetase (GS), NADH-dependent glutamate dehydrogenase (NADH-GDH) and NADP-dependent isocitrate dehydrogenase (NADP-ICDH) were measured by the method of Lu et al. [\(2005](#page-12-13)).

Genes Expression

Total RNA extraction, cDNA synthesis and quantitative real time PCR (qRT-PCR) were performed as described by Wang et al. [\(2018\)](#page-12-19). The specific primers of genes encoding ornithine- δ-aminotransferase (OAT), pyrroline-5-carboxylate dehydrogenase (P5CDH), hexokinase (HXK), citrate

synthetase (CS), pyruvate kinase (PK) and malate dehydrogenase (MDH) are listed in Table S1. The relative expression levels of genes were calculated according to the $2 - \Delta \Delta C$ t method (Wang et al. [2018\)](#page-12-19), using *Actin* gene as a reference gene.

Statistical Analysis

Statistical analysis was performed using analysis of variance (ANOVA) followed by Duncan's multiple range test (SPSS18.0, SPSS Inc., IL, USA), and $P < 0.05$ was statistically signifcant.

Results

LT50, Leaf Water Potential and Cell Death

As shown in Fig. [1](#page-3-0)A, cold and SA priming treatment (S and P) decreased the LT₅₀ from $-$ 4.4 °C (non-priming treatment, C) to -5.3 °C and -5.8 °C, respectively, demonstrating the beneficial effect of cold and SA priming treatment on freezing tolerance enhancement. The LT_{50} was calculated by electrolyte leakage of leaves. Therefore, we hypothesized that priming with SA and cold temperature could alleviate freeze-induced tissue dehydration in wheat leaves.

As shown in Fig. [1B](#page-3-0), the freezing treatment caused a signifcant decrease in leaf water potential. However, the SF and the PF treatments showed lower reduction of leaf water potential than the CF treatment (Fig. [1](#page-3-0)B). As compared to the CC treatment, the leaf water potential was decreased by 45.6%, 48.5% and 89.5% in the SF, the PF and the CF treatments, respectively. Evans blue and Trypan blue staining showed that more dead cells (as indicated by the blue spots) were present in the CF treatment as compared with the SF and the PF treatments (Fig. [1C](#page-3-0), D). There was no signifcant diference in the leaf water potential and the number of dead cells between the primed and non-primed plants under non-freezing conditions (Fig. [1B](#page-3-0)–D). These results suggested that cold and SA priming could alleviate dehydration-induced cell death under freezing stress.

Activities and Expression Levels of Key Enzymes Involved in Proline and Sucrose Biosynthesis

The activity of SPS plays a crucial role in controlling sucrose biosynthesis in wheat (Vargas et al. [2007\)](#page-12-16), and SS is an important enzyme considered to have reversible sucrose synthesis and cleavage functions (Kumutha et al. [2008](#page-11-20)). As shown in Fig. [2A](#page-4-0) and B, the SC and the PC treatments significantly increased the activities of SPS and SS (synthetic direction) as compared with the CC treatment. Freezing stress obviously enhanced the

Fig. 1 Effects of SA and cold priming treatment on freeze-induced dehydration injury in wheat leaves. **A** LT_{50} . The measurement was taken at the end of the lag phase (8 days of recovery from priming treatment). *C* no priming treatment, *S* SA priming treatment, *P* cold priming treatment. Each value of electrolyte leakage is the mean \pm SE of three biological replicates; **B** Leaf water potential. Each value is the mean \pm SE of three biological replicates. The different lowercase

letters indicate statistically signifcant diferences at *P*<*0.05* level; **C** Trypan blue staining; **D** Evans blue staining. **B**–**D** were determined at the end of the freezing treatment. **C**–**D** blue spots indicate dead cells. *CC* no SA or cold priming+no freezing stress, *SC* SA priming+no freezing stress, *PC* cold priming+no freezing stress, *CF* no SA or cold priming+freezing stress, *SF* SA priming+freezing stress, *PF* cold priming+freezing stress

activities of SPS and SS (Fig. [2](#page-4-0)A, B). However, there were increases in SPS (67.3% and 72.3%, respectively) and SS (44.9% and 47.7%, respectively) activities in the SF and the PF treatments as compared with the CF treatment (Fig. [2A](#page-4-0), B).

In higher plants, proline can be synthesized from glutamate and ornithine, which are, respectively, catalyzed by P5CS and OAT (Szabados and Savourcb [2010](#page-12-3)). As shown in Fig. [2](#page-4-0)C and D, the SC treatment clearly increased P5CS activity and the expression level of *OAT* as compared with the CC treatment. The PC treatment caused an obvious increase in the expression level of *OAT*, while it had no significant effect on P5CS activity as compared with the CC treatment (Fig. [2C](#page-4-0), D). Freezing stress significantly up-regulated the expression level of *OAT* (2.31-fold), but it had no significant effect on P5CS activity as compared with the CC treatment (Fig. [2](#page-4-0)C, D). Here, the SF and the PF treatments showed relatively higher P5CS activity (73.1% and 87.3%, respectively) and expression level of *OAT* (1.52-fold and 1.45-fold, respectively) as compared with the CF treatment (Fig. [2](#page-4-0)C, D).

Free Proline and Sucrose Contents

Both free proline and sucrose contents were clearly enhanced by freezing stress (Fig. [3\)](#page-4-1). Moreover, the SF and the PF treatments showed strikingly higher free proline and sucrose contents as compared with the CF treatment (Fig. [3\)](#page-4-1). As compared to the CF treatment, the free proline and sucrose contents were increased by 64.9% and 13.1% under the SF treatment, while increased 60.5% and 20.9% under the PF treatment, respectively. The SC and the PC treatments signifcantly increased the contents of free proline (70.4% and 142%, respectively) and sucrose (28.0% and 32.8%, respectively) as compared with the CC treatment (Fig. [3](#page-4-1)). These results suggested that cold and SA priming could promote-free proline and sucrose accumulation under freezing conditions contributing to alleviating freeze-induced dehydration damage.

Fig. 2 Efects of SA and cold priming on the activities and the expression levels of sucrose and free proline biosynthesis-related enzymes in wheat leaves. **A** Sucrose-phosphate synthase (SPS) activity; **B** Sucrose synthase (SS, synthetic direction) activity; **C** Δ1-pyrroline-5-carboxylate synthetase (P5CS) activity; **D** Relative expression level of *OAT* (encoding ornithine- δ-aminotransferase). **A**–**D** were determined at the end of the freezing treatment. *CC* no

SA or cold priming+no freezing stress, *SC* SA priming+no freezing stress, *PC* cold priming+no freezing stress, *CF* no SA or cold priming+freezing stress, *SF* SA priming+freezing stress, *PF* cold priming + freezing stress. Each value is the mean \pm SE of three biological replicates. For gene expression data, each biological replicate has three technical replicates. The diferent lowercase letters indicate statistically signifcant diferences at *P*<*0.05* level

Fig. 3 Efects of SA and cold priming on sucrose and free proline contents in wheat leaves. **A** Sucrose content; **B** Free proline content. **A**–**B** were determined at the end of the freezing treatment. *CC* no SA or cold priming+no freezing stress, *SC* SA priming+no freezing stress, *PC* cold priming+no freezing stress, *CF* no SA or cold prim-

Activities and Expression Levels of Key Enzymes Involved in Proline and Sucrose Catabolism

Sucrose can be hydrolyzed into glucose and fructose by INV in wheat (Vargas et al. [2007\)](#page-12-16). As shown in Fig. [4A](#page-5-0) and B, the SC treatment had no signifcant efects on the activities of A/N-INV and Ac-INV as compared with the CC treatment. The PC treatment signifcantly increased

ing+freezing stress, *SF* SA priming+freezing stress, *PF* cold priming + freezing stress. Each value is the mean \pm SE of three biological replicates. The diferent lowercase letters indicate statistically signifcant diferences at *P*<*0.05* level

the activities of A/N-INV and Ac-INV as compared with the CC treatment (Fig. [4](#page-5-0)A, B). Freezing stress obviously enhanced the activities of A/N-INV and Ac-INV (Fig. [4](#page-5-0)A, B). The SF and the PF treatments showed relatively higher A/N-INV (20.4% and 8.89%, respectively) and Ac-INV (12.4% and 10.3%, respectively) activities as compared with the CF treatment (Fig. [4A](#page-5-0), B).

Fig. 4 Efects of SA and cold priming on the activities and the expression levels of sucrose and free proline catabolism-related enzymes in wheat leaves. **A** Alkaline/neutral-invertase (A/N-INV) activity; **B** Acid-invertase (Ac-INV) activity; **C** Proline dehydrogenase (PDH) activity; **D** Relative expression level of *P5CDH* (encoding pyrroline-5-carboxylate dehydrogenase). **A**–**D** were determined at the end of the freezing treatment. *CC* no SA or cold priming+no

freezing stress, *SC* SA priming+no freezing stress, *PC* cold priming + no freezing stress, CF no SA or cold priming + freezing stress, *SF* SA priming+freezing stress, *PF* cold priming+freezing stress. Each value is the mean \pm SE of three biological replicates. For gene expression data, each biological replicate has three technical replicates. The diferent lowercase letters indicate statistically signifcant diferences at *P*<*0.05* level

The degradation of proline is catalyzed by PDH and P5CDH in higher plants (Yang et al. [2016\)](#page-13-9). As shown in Fig. [4](#page-5-0)C and D, the SC treatment signifcantly decreased PDH activity, and up-regulated the expression level of *P5CDH* as compared with the CC treatment. The PC treatment obviously up-regulated PDH activity and the expression level of *P5CDH* as compared with the CC treatment (Fig. [4C](#page-5-0), D). The CF treatment signifcantly increased PDH activity (18.7%) and the expression level of *P5CDH* (4.47 fold) as compared with the CC treatment (Fig. [4C](#page-5-0), D). Here, there were decreases in PDH activity (18.4% and 11.4%, respectively) and expression level of *P5CDH* (0.76-fold and 0.75-fold, respectively) in the SF and the PF treatments as compared with the CF treatment (Fig. [4C](#page-5-0), D).

Ammonium Assimilation

The activity of GS and NADH-GDH play critical role in regulating the assimilation of ammonia for glutamate synthesis in plants (Mifin and Habash [2002\)](#page-12-20). There was no substantial diference in the contents of ammonium and activities of GS and NADH-GDH between the primed and non-primed plants under non-freezing conditions, except for a signifcant increase of NADH-GDH activity in the SC treatment as compared with the CC treatment (Table [1](#page-6-0)). The CF treatment caused a signifcant increase in ammonium content (115%) and GS activity (12.2%), and a slight increase in NADH-GDH activity (8.0%) as compared with the CC treatment (Table [1](#page-6-0)). The SF and the PF treatments obviously increased the activities of GS (13.0% and 17.4%, respectively) and NADH-GDH (20.1% and 16.3%, respectively), and dramatically decreased ammonium content (46.8% and 39.2%, respectively) as compared with the CF treatment (Table [1](#page-6-0)). These results suggested that cold and SA priming promoted the assimilation of ammonium under freezing. Here, we investigated glutamate content in each treatment to further confrm this inference. Interestingly, the SF and the PF treatments induced a signifcant reduction by 28.7% and 26.5% in glutamate content in wheat leaves as compared with the CF treatment (Table [1](#page-6-0)).

Table 1 Efects of SA and cold priming on the levels of ammonium and glutamate, and the activities of GS and NADH-GDH in wheat leaves

The measurement was taken at the end of the freezing treatment. Each value is the mean \pm SE of three biological replicates. The diferent lowercase letters indicate statistically signifcant diferences at *P*<*0.05* level

GS glutamine synthetase, *NADH-GDH* NADH-dependent glutamate dehydrogenase, *CC* no SA or cold priming+no freezing stress, *SC* SA priming+no freezing stress, *PC* cold priming+no freezing stress, *CF* no SA or cold priming+freezing stress, *SF* SA priming+freezing stress, *PF* cold priming+freezing stress

NADP‑ICDH Activity and 2‑OG Content

In plants, 2-OG is an important carbon skeleton for glutamate synthesis (Mifin and Habash [2002\)](#page-12-20), and cytosolic NADP-ICDH plays a key role in 2-OG synthesis (Wang et al. [2007\)](#page-12-21). As shown in Fig. [5](#page-6-1), the SC and the PC treatments insignifcantly increased NADP-ICDH activity, and remarkably increased the content of 2-OG as compared with the CC treatment. The CF treatment induced a slight increase in NADP-ICDH activity (13.0%), but obviously enhanced the accumulation of 2-OG (61.3%) as compared with the CC treatment (Fig. [5\)](#page-6-1). Interestingly, the SF and the PF treatments clearly increased NADP-ICDH activity by 23.0% and 35.1%, but signifcantly decreased 2-OG content by 14.9% and 11.9% as compared with the CF treatment, repectively (Fig. [5\)](#page-6-1).

Expression Levels of Key Enzymes Involved in Carbon Metabolism

Expression levels of four key genes (*HXK*, *CS*, *PK* and *MDH*) involved in carbon metabolism were investigated here. There was no substantial diference in expression levels of these genes between the primed and non-primed plants under non-freezing conditions, except for a signifcant up-regulation of the expression level of *HXK* in the

Fig. 5 Efects of SA and cold priming on the activities of NADP-ICDH and the contents of 2-OG in wheat leaves. **A** NADP-dependent isocitrate dehydrogenase (NADP-ICDH) activity; **B** 2-oxoglutarate (2-OG) content. **A**–**B** were determined at the end of the freezing treatment. *CC* no SA or cold priming+no freezing stress, *SC* SA

priming+no freezing stress, *PC* cold priming+no freezing stress, *CF* no SA or cold priming+freezing stress, *SF* SA priming+freezing stress, PF cold priming + freezing stress. Each value is the mean \pm SE of three biological replicates. The diferent lowercase letters indicate statistically signifcant diferences at *P*<*0.05* level

Fig. 6 Efects of SA and cold priming on the expression levels of genes involved in glycolysis and the tricarboxylic acid (TCA) cycle in wheat leaves. **A** Relative expression level of *HXK* (encoding hexokinase); **B** Relative expression level of *CS* (encoding citrate synthetase); **C** Relative expression level of *PK* (encoding pyruvate kinase); **D** Relative expression level of *MDH* (encoding malate dehydrogenase). **A**–**D** were determined at the end of the freezing treat-

SC treatment as compared with the CC treatment (Fig. [6](#page-7-0)). The CF treatment signifcantly up-regulated the expression levels of *HXK* (1.81-fold) and *CS* (1.82-fold), while it had no signifcant efects on the expression level of *PK* and *MDH* as compared with the CC treatment (Fig. [6](#page-7-0)). The SF and the PF treatment evidently up-regulated the expression level of *MDH* (2.22-fold and 1.75-fold, respectively), and gently up-regulated the expression level of *HXK*, *PK* and *CS* as compared with the CF treatment (Fig. [6\)](#page-7-0).

Discussion

Despite the important role of sucrose and proline in winter wheat resisting freezing stress has been well recorded, the underlying mechanisms of freeze-induced sucrose and

ment. *CC* no SA or cold priming+no freezing stress, *SC* SA priming+no freezing stress, *PC* cold priming+no freezing stress, *CF* no SA or cold priming+freezing stress, *SF* SA priming+freezing stress, PF cold priming + freezing stress. Each value is the mean \pm SE of three biological replicates, and each biological replicate has three technical replicates. The diferent lowercase letters indicate statistically signifcant diferences at *P*<*0.05* level

free proline accumulation are not completely understood. In the present study, the synthesis and degradation of sucrose were both clearly increased by freezing, as revealed by a signifcant higher SPS, SS (synthetic direction), A/N-INV and Ac-INV activities under freezing conditions than nonfreezing conditions (Figs. [2](#page-4-0)A, B, and [4A](#page-5-0), B). These results were similar to those of Savitch et al. ([2000\)](#page-12-22), who found that chilling stress prominently increased the activities of SPS, SS (synthetic direction), A/N-INV and Ac-INV in wheat leaves. Kaplan et al. [\(2007](#page-11-21)) indicated that starch degradation played an important role in chilling-induced sugar accumulation in *Arabidopsis*. The degradation of starch leaded to the accumulation of glucose-6-phosphate and fructose-phosphate (immediate precursors for sucrose biosynthesis), then to sucrose and then to fructose and glucose (Kaplan et al. [2007](#page-11-21)). In the light of the above studies, it appears that the accumulation of sucrose in wheat leaves under cold temperature stress might be due to a faster rate or an earlier start of its synthesis than degradation. Zhang et al. [\(2019](#page-13-10)) reported that freezing stress signifcantly increased sucrose content and SPS activity, but decreased INV activity in wheat ears at the booting stage. These studies suggested that the diferential mechanisms of freeze-induced sucrose accumulation found in diferent organs of wheat.

Previous studies reported that the increased INV activity could contribute to producing more hexoses (fructose and glucose) that would be metabolized into energy and intermediate metabolites for plant resistance to stresses (Sehar et al. [2019;](#page-12-23) Savitch et al. [2000;](#page-12-22) Vargas et al. [2007\)](#page-12-16). This was further confrmed by our results that the up-regulation of INV activity and the expression levels of *HXK* (encoding hexokinase, the crucial rate-limiting enzyme relating to the glycolytic pathway) and *CS* (encoding citrate synthetase, the crucial rate-limiting enzyme relating to the tricarboxylic acid cycle) induced by freezing treatment (Figs. [4A](#page-5-0), B, and [6](#page-7-0)A, B). In addition, increased INV activity was associated with the transport of sucrose to the vacuole where it is used for fructan synthesis, which also played an important role in the regulation of freezing tolerance in wheat (Amani [2008](#page-11-22); Vágújfalvi et al. [1999\)](#page-12-6).

Previous studies indicated that both the glutamate pathway and the ornithine pathway were implicated in chillinginduced proline synthesis (Cao et al. [2012;](#page-11-23) Yang et al. [2016](#page-13-9)). However, in this study, freezing treatment caused a signifcant up-regulation of the expression level of gene encoding OAT, but it had no signifcant efect on P5CS activity (Fig. [2C](#page-4-0), D). Yang et al. ([2009\)](#page-13-11) and (2016) found that the glutamate pathway was activated prior to the ornithine pathway when maize and *Jatropha curcas* seedlings responded to stress factors such as chilling temperature and hydrogen peroxide. Due to the activity of P5CS determined at the end of freezing treatment here, the role of the glutamate pathway in proline synthesis at the early time points of wheat plants response to freezing cannot be excluded. However, our results suggested that the ornithine pathway played an important role in proline synthesis at the later stage of freezing stress in wheat.

The accumulation of proline depends not only on its synthesis, but also on its degradation (Liu et al. [2020](#page-12-24); Szabados and Savourcb [2010](#page-12-3)). Cao et al. [\(2012\)](#page-11-23) and Zeng et al. ([2015\)](#page-13-12) indicated that the increase in proline level in loquat fruit and bamboo shoots under chilling stress were related with lower PDH activity. Intriguingly, freezing treatment significantly up-regulated the activity of PDH and the expression level of gene encoding P5CDH in this study (Fig. [4](#page-5-0)C, D). Szabados and Savourcb [\(2010\)](#page-12-3) indicated that proline had a feedback efect on transcriptional regulation of proline metabolism. Thus, the up-regulation of activity of PDH and expression level of *P5CDH* at the end of freezing treatment might be induced by the accumulated free proline.

Due to glutamate is an important substrate for proline synthesis, the accumulation of proline is closely associated with ammonia metabolism (Martins et al. [2020](#page-12-5)). In the present study, freezing stress obviously enhanced ammonia content in wheat leaves (Table [1\)](#page-6-0). This is in accordance with previous studies that the enhancement of photorespiration capacity and ROS-induced proteolysis under stress conditions could lead to an increase in intracellular ammonia (Skopelitis et al. [2006;](#page-12-25) Wang et al. [2007\)](#page-12-21). The synthetic precursor of 2-OG is citrate, an intermediate metabolite of the tricarboxylic acid cycle (Gálvez et al. [1999\)](#page-11-11). Therefore, freeze-enhanced the expression of *HXK* and *CS* and the activity of NADP-ICDH could contribute to the increase in 2-OG level (Figs. [5](#page-6-1) and [6](#page-7-0)A, B). Previous studies indicated that accumulation of ammonia and 2-OG could regulate the activities of GS and NADH-GDH (Gálvez et al. [1999](#page-11-11); Wang et al. [2016\)](#page-12-26). Thus, the increased ammonia and 2-OG might induce the increasement of GS and NADH-GDH activities under freezing, resulting in a signifcant increase in glutamate content that could be used for proline synthesis (Table [1\)](#page-6-0). These results were consistent with Lu et al. ([2005](#page-12-13)), who found that the accumulation of free proline under chilling stress in rice was in good agreement with the induction of GS and NADP-ICDH activities. All these fndings suggested that the coordination of carbon and nitrogen metabolism played an important role in the freeze-induced simultaneous accumulation of sucrose and free proline in wheat leaves.

It has been well-documented that winter wheat can acquire tolerance to sub-zero temperatures by a process known as cold acclimation or cold hardening (Kaplan et al. [2007](#page-11-21); Savitch et al. [2000\)](#page-12-22). Nevertheless, there are some differences between cold acclimation and cold priming, which have been well elucidated in several recent papers (Baier et al. [2019;](#page-11-12) Zuther et al. [2019](#page-13-3)). Previous studies demonstrated that accumulation of compatible solutes plays an important role in the cold acclimation-induced freezing tolerance in wheat (Vágújfalvi et al. [1999;](#page-12-6) Wardlaw and Willenbrink [1994](#page-13-8)). However, the effects of cold priming on compatible solutes accumulation under freezing remain largely unidentifed. In this study, we found that cold priming treatment signifcantly increased free proline and sucrose contents in wheat leaves as compared with non-priming treatment under both non-freezing and freezing conditions (Fig. [3\)](#page-4-1). This is in good agreement with a recent study by Zuther et al. [\(2019](#page-13-3)), who reported that the accumulation of osmoprotective compounds such as galactinol, sucrose and proline were involved in the cold priming-induced freezing tolerance in *Arabidopsis*. In addition, our previous studies reported that drought primed plants showed signifcantly higher levels of the compatible solutes such as free proline, sucrose and glycine betaine as compared with non-primed pants under the subsequent drought stress (Wang et al. [2019,](#page-12-2) [2018](#page-12-19)). Taken together, these results suggested that accumulation of compatible solutes is an important mechanism of the priming events inducing stress tolerance in wheat plants.

Numerous recent reports have shown that exogenous SA could promote accumulation of free proline under stressful conditions (Sedaghat et al. [2020;](#page-12-27) Min et al. [2018;](#page-12-11) Estaji and Niknam [2020](#page-11-24)). The regulatory effect of SA on accumulation of free proline in this study was consistent with the previous studies (Fig. [3](#page-4-1)B). However, previous reports about the regulatory efect of SA on sucrose level was inconsistent. Poór et al. [\(2011\)](#page-12-12) and Dong et al. ([2011](#page-11-25)) found that SA signifcantly decreased sucrose content in tomato and cucumber leaves, but increased its content in roots under salinity stress. La et al. ([2019a](#page-11-26)) and Min et al. ([2018](#page-12-11)) reported that SA markedly enhanced sucrose content in *Brassica rapa* and spinach leaves under drought and freezing stress, respectively. In this study, SA signifcantly increased sucrose level in wheat leaves under both non-freezing and freezing conditions (Fig. [3](#page-4-1)A). These results indicated that the regulatory efect of SA on sucrose level might be dependent on species and type of organs and stresses.

In the present work, it was shown that cold and SA priming-induced free proline accumulation by promoting its synthesis and inhibiting its degradation in wheat leaves as compared with non-priming treatment under freezing conditions (Figs. [2](#page-4-0)C, D, and [4](#page-5-0)C, D). The results were consistent with La et al. [\(2019b\)](#page-12-28) who found that exogenous SA increased free proline content by up-regulating the expression level of *P5CS* and down-regulating the expression level of *PDH* in *Brassica rapa* leaves under drought stress. Previous studies indicated that exogenous SA could enhance sucrose level through increasing the activity of SPS and the degradation of starch in *Brassica rapa* leaves under drought stress (La et al. [2019a\)](#page-11-26). Our results showed that cold and SA priming promoted sucrose accumulation by increasing the activities of SPS and SS (synthetic direction) in wheat leaves under freezing stress (Fig. [2](#page-4-0)A, B). However, the role of starch degradation in SA and cold priming-induced the accumulation of sucrose in wheat leaves under freezing need to be investigated further.

In this study, priming treatment signifcantly increased activities of GS and NADH-GDH, and decreased level of ammonia as compared with non-priming treatment under freezing stress (Table [1](#page-6-0)). This indicated that cold and SA priming improved ammonia assimilation under freezing stress, which could provide more synthetic precursor (glutamate) for the synthesis of proline. Meanwhile, cold and SA primed plants showed relatively higher activities of INV and NADP-ICDH, and higher expression level of genes involved in the glycolytic pathway (such as *HXK* and *PK*) and the tricarboxylic acid cycle (such as *CS* and *MDH*) as compared with non-primed plants under freezing stress (Figs. [4–](#page-5-0)[6](#page-7-0)), indicating that cold and SA priming promoted the catabolism of sucrose and glucose under freezing stress. This might contribute to producing more energy and carbon skeleton (2-OG) for ammonia assimilation. Collectively, these results suggested that cold and SA priming could intensify the coordination between carbon and nitrogen metabolism to enhance the accumulation of compatible solutes in wheat leaves under the later freezing stress, and therefore efectively alleviated freeze-induced dehydration damage.

But interestingly, SA and cold primed plants showed signifcantly lower level of 2-OG as compared with non-primed plants under freezing conditions (Fig. [5B](#page-6-1)). This might be due to the accelerated ammonia assimilation by the priming treatment under freezing conditions (Table [1\)](#page-6-0). In addition, the higher level of free proline in primed plants might explain why the primed plants accelerated ammonia assimilation but had lower glutamate contents as compared with the non-primed plants under freezing conditions (Table [1\)](#page-6-0).

Our previous study found that cold priming significantly increased SA level in wheat leaves, and the results of pharmacological experiment indicated that SA signal was involved in the cold priming-induced tolerance to freezing in wheat (Wang et al. [2021\)](#page-13-13). In the present study, the results indicated that SA and cold priming have similar mechanisms in inducing sucrose and free proline accumulation in wheat leaves to alleviate freeze-caused dehydration damage (Fig. [7\)](#page-10-0). This could provide indirectly evidence for the involvement of SA in the cold priming-induced freezing

$SA/Cold$ priming + Freezing temperatures

Improved Freezing Tolerance

Fig. 7 Schematics of SA and cold priming simultaneously enhanced freeze-induced sucrose and proline accumulation by intensifying the coordination of carbon and ammonia metabolism in wheat leaves. *SPS* sucrose-phosphate synthase, *SS* sucrose synthase, *INV* invertase, *NADP-ICDH* NADP-dependent isocitrate dehydrogenase; *TCA cycle* tricarboxylic acid cycle, *2-OG* 2-oxoglutarate, *NADH-GDH* NADHdependent glutamate dehydrogenase, *GS* glutamine synthetase, *GOGAT* glutamate synthase, *P5CS* Δ1-pyrroline-5-carboxylate syn-

tolerance, and should help the introduction of SA into practices of anti-freezing stress cultivation in wheat.

Conclusions

In conclusion, the coordination between carbon and nitrogen metabolism played important role in the simultaneous accumulation of sucrose and free proline in wheat leaves response to freezing stress, and cold and SA priming could intensify this coordination to enhance sucrose and free proline levels, which was contributing to the alleviation of freeze-induced cell dehydration (Fig. [7\)](#page-10-0). It has become clear

thetase, *OAT* ornithine- δ-aminotransferase, *PDH* proline dehydrogenase, *P5CDH* pyrroline-5-carboxylate dehydrogenase, *ROS* reactive oxygen species. Black arrows indicate the positive regulation of freezing temperature in metabolites level, enzymes activity and genes expression. Red arrows indicate the positive or negative regulation of SA and cold priming in metabolites level, enzymes activity and genes expression

that carbon and nitrogen metabolism are regulated to enable the coordination essential for plant growth and development. To the best of our knowledge, this is the frst report owing that the important role of the coordination between carbon and nitrogen metabolism in wheat responded to freezing stress. However, the carbon and nitrogen metabolisms of plants involve many processes, the more detailed mechanisms of plant coordinated carbon and nitrogen metabolism in response to environmental stresses need further study.

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Author Contributions WW, XW and DJ conceived and designed research. WW, ZL and MH conducted experiments. JC, QZ and ZH guided experiments. WW and AK wrote the manuscript. All authors read and approved the manuscript.

Declarations

Conflict of interest The authors declare no confict of interest.

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