

Article

Insights into Nitric Oxide-Mediated Water Balance, Antioxidant Defence and Mineral Homeostasis in Rice (*Oryza sativa* L.) under Chilling Stress

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Running title: Nitric Oxide-Mediated Chilling Stress Tolerance in Rice

Abstract: Being a chilling-sensitive staple crop, rice (*Oryza sativa* L.) is vulnerable to climate change. The competence of rice to withstand chilling stress should, therefore, be enhanced through technological tools. The present study employed chemical intervention like application of sodium nitroprusside (SNP) as nitric oxide (NO) donor and elucidated the underlying molecular mechanisms of NO-mediated chilling tolerance in rice. At germination stage, germination indicators were interrupted by chilling stress ($5.0 \pm 1.0^\circ\text{C}$ for 8 h day⁻¹), while pretreatment with 100 μM SNP markedly improved the indicators. At seedling stage (14-day-old), chilling stress caused stunted growth with visible toxicity along with alteration of biochemical markers, for example, increase in oxidative stress markers (superoxide, hydrogen peroxide, and malondialdehyde) and osmolytes (total soluble sugar; proline and soluble protein content, SPC), and decrease in chlorophyll (Chl), relative water content (RWC), and antioxidants. However, NO application attenuated toxicity symptoms with improving growth performance which might be attributed to enhanced activities of antioxidants, mineral contents, Chl, RWC and SPC. Furthermore, principal component analysis indicated that water imbalance and increased oxidative damage were the main contributors to chilling injury, whereas NO-mediated mineral homeostasis and antioxidant defense were the critical determinants for chilling tolerance in rice. Collectively, our findings revealed that NO protects against chilling stress through valorizing cellular defense mechanisms, suggesting that exogenous application of NO could be a potential tool to evolve cold tolerance as well as climate resilience in rice.

Keywords: antioxidant system; chilling stress; mineral homeostasis; nitric oxide; oxidative stress; rice

1. Introduction

Climate change noticeably alters the distribution of temperature variability [1], which results in more frequent cold stress events during the crop growing seasons [2]. Prolonged exposure (i.e. several days to weeks) of plants to chilling temperature exerts a severe threat to physiological, biochemical, and molecular processes of plants which results in impairment on seed germination, plant growth and development [2,3]. In semi-arid climates, plants are often exposed to extreme temperatures including hot day and cold night — this fluctuation in temperature results in an additional threat to the plants [4].

In plants, the preliminary symptoms of chilling stress include disruption of plant-water relationships, reduced nutrient uptake, destruction of photosynthetic pigments, change in protein structures, and enzyme activities, which, in turn, exert a serious threat to cell survival resulting in programmed cell death [5,6]. Furthermore, the perturbation in photosynthesis increases the production of excessive reactive oxygen species (ROS) [singlet oxygen ($^1\text{O}_2$), superoxide anion ($\text{O}_2^{\cdot-}$), hydrogen peroxide (H_2O_2), and hydroxyl radical ($\text{OH}^{\cdot-}$)] through Fenton Haber–Weiss reactions [7,8]. Excessive ROS accumulation is responsible for the oxidation of vital biomolecules [9]. Thus, scavenging of these harmful ROS is crucial for the survival of plants, which is mainly accomplished by the antioxidants. The increased antioxidants, therefore, confer tolerance in the plants during chilling stress [6,10,11]. Plants acquired various kinds of defense strategies to cope with the stress-induced adversities, which includes enzymatic antioxidants (catalase, CAT; superoxide dismutase, SOD; peroxidase, POX etc.), non-enzymatic antioxidants (ascorbate, AsA; carotenoids, and phenolic compounds etc.), osmolytes (proline, soluble sugars and proteins) [12,13,14]. On the contrary, any imbalance of antioxidants, osmoregulation, and essential minerals homeostasis lead to the impairment of plant defence mechanisms which results in weakening of plant physiological processes [7]. In this context, chemical priming has been considered as an alternative strategy for improving abiotic stress resistance of plants [15].

Augmenting the defensive responses of plants can minimize the chilling-induced damage, which can be attained by applying exogenous signalling molecules [16]. Nitric oxide (NO), an enormously diffusible and a versatile bioactive inter- and intra-cellular signalling compound, found to be involved in seed germination, water balance, mineral adjustment, gene regulation, and upregulation of antioxidant enzyme activities in plants [17,18,19]. NO networks with other hormones (abscisic acid, auxins, brassinosteroids, cytokinins, ethylene, gibberellins, jasmonates, and salicylic acid, etc.) and signalling molecules (Ca^{2+} , glutathione, and ROS), that alter the developmental activities and stress tolerance of plants [19]. Moreover, NO can possess an energetically more advantageous electron structure, by gaining or losing an electron, to neutralize toxic ROS in the plant cell [19,20]. Furthermore, NO can counteract oxidative stress by directly acting as an antioxidant, regulating the osmotic balance, protecting membrane lipid and stimulating the gene expression of the antioxidant enzymes under various abiotic stress conditions [18,19,20]. In recent years, the exogenous applications of NO showed a protective effect against chilling stress in several plants such as cucumber (*Cucumis sativus*) [21], wheat (*Triticum aestivum*) [22] and Chinese cabbage (*Brassica rapa* subsp. *Pekinensis*) [23]. Additionally, the exogenous application of NO displayed protective effect in salt-stressed chickpea (*Cicer arietinum* L.) [24] and drought-stressed rapeseed (*Brassica napus*) [25]. However, to the best of our knowledge, whether exogenous applications of NO can alleviate chilling-induced oxidative stress in rice plants, has not been reported yet.

Rice is a staple food for more than 60 per cent of the world population [26] and is widely cultivated in tropical, subtropical and temperate climate zones [27]. The optimum temperature for rice cultivation fluctuates between 25°C – 35°C which indicates the reason behind the susceptibility of rice to chilling temperature compared to other cereal crops [28,29]. About 10% of rice cultivable lands are affected by chilling stress [3,27,29]. Particularly during the seedling stage, chilling causes necrotic lesions on leaves, yellowing symptoms, delay leaf development and induce wilting of leaves [3,27] which results in mass

death of rice seedlings in the seedbed. For instance, in the northern region of Bangladesh, cold is usually more substantial (around 2.6°C in January) than other parts of the country and damages rice seedbeds during the winter season [29,30,31]. Protecting rice seedlings from this harsh cold spell is, therefore, necessary for higher rice production.

In this study, firstly, we investigated whether NO mitigates chilling stress in rice both at the germination and seedling stages. Secondly, we also investigated the protective functions of NO in enhancing chilling tolerance of rice seedlings by examining the morphological, physiological and biochemical mechanisms through the evaluation of the following features: (i) germination indices, (ii) plant growth attributes and biomass production, (iii) mineral homeostasis, (iv) photosynthetic pigment, (v) various types of osmolyte status, (vi) chilling-induced oxidative injury in terms of elevated ROS levels and lipid peroxidation, and (vii) enhancement of antioxidant defense system.

2. Materials and Methods

2.1. Germination Stage Experiment

BRRI dhan29, a cold-sensitive, high-yielding and widely cultivated rice in Bangladesh, was used in this experiment. For surface sterilization, seeds were soaked in 2.5% sodium hypochlorite + 2.0% Tween-20 solution for 15 minutes, followed by washed with double distilled water (ddH₂O) four times. After that, 200 seeds were soaked in 100 mL ddH₂O or 100 mL of 100 µM SNP (sodium nitroprusside as the source of NO) solution and kept in dark at 25°C. After 24 h of imbibition, ddH₂O or SNP soaked seeds transferred in two Petri dishes (two pieces of filter papers were placed in each Petri dish with a diameter of 9 cm, and an equal volume of ddH₂O was added to soak the paper) each with 100 seeds. Then one batch of Petri dishes was kept in normal condition and others in chilling stress. Therefore, the study comprised of four treatments as follows “C”, 0 h chilling day⁻¹ + 0 µM SNP (control); “CH”, 8 h chilling day⁻¹ + 0 µM SNP; “SNP”, 0 h chilling day⁻¹ + 100 µM SNP; and “CH + SNP”, 8 h chilling day⁻¹ + 100 µM SNP. The “C” and “SNP” levelled Petri dishes were kept in the dark at 29°C, whereas “CH” and “CH+SNP” levelled Petri dishes were kept at the cold chamber for 8 h day⁻¹ (cold chamber at night, 10.00 PM – 6.00 AM, 5.0±1.0°C). After 8 h of chilling treatment, “CH” and “CH+SNP” levelled Petri dishes were transferred to dark daily at 29°C. However, seeds were considered as germinated when the radicle reached 2 mm in length. From 2nd day after incubation (DAI), number of germinated seeds was recorded up to 5th DAI and by using these germination counts, several germination indices were determined, such as final germination percentage (FGP), germination rate index (GRI), coefficient of velocity of germination (CVG), mean germination time (MGT) [32] and vigor index (VI) [33]. Moreover, the radicle length (RaL), plumule length (PL), radicle fresh weight (RaFW), and plumule fresh weight (PFW) were recorded at 6th DAI.

2.2. Seedling Stage Experiment

Sterilized seeds were imbibed in dH₂O for 24 h followed by incubated in the dark at 29°C for germination. One hundred and fifty germinated rice seeds were sown in 350 mL plastic pot containing ddH₂O. From the 3rd day of sowing, the seedlings were grown in modified Cooper's nutrient solution [34]. The nutrient solution was prepared by adding the following to 10 L of distilled water; KH₂PO₄ (2.63 g), KNO₃ (5.83 g), Ca(NO₃)₂·4H₂O (10.03 g), FeSO₄·7H₂O (2.00 g), MgSO₄·7H₂O (0.79 g), MnSO₄ (0.061 g), H₃BO₃ (0.017 g), CuSO₄·5H₂O (0.004 g), Na₂MoO₄ (0.003 g), ZnSO₄·7H₂O (0.0044 g). The experiment was conducted in a growth chamber set to a constant temperature of 25 ± 2°C and relative humidity of 65-70%, at the Department of Biochemistry and Molecular Biology, Bangladesh Agricultural University, Mymensingh. The nutrient solution was replaced at 3-day intervals throughout the experiment.

The 14-day old rice seedlings were treated with 100 μM SNP in nutrient solution for three days. Moreover, seedlings were treated with freshly prepared 100 μM SNP by the foliar application (10 mL spray⁻¹ pot⁻¹) at 12:00 PM until the end of the experiment. The control plants were sprayed with ddH₂O. The Tween-20 (0.1%, v/v) was added both with SNP and water to ensure the maximum adherence of SNP and water to the leaves of the rice plant. After three days pretreatment, 17-day old SNP-pretreated and non-treated rice seedlings were subjected to chilling stress every day (induced by keeping in a cold chamber at night for 8 h day⁻¹, 10.00 PM – 6.00 AM, 5.0 \pm 1.0°C) followed by recovery at 25°C. Control and SNP treated non-stressed seedlings were kept at normal growth conditions throughout the experimental period. Therefore, the treatment combinations were the same as the germination stage experiments. The experiment was conducted using a randomized complete block design with three independent replicates for each treatment. After 8 days of growth under the above conditions, 25-day-old rice plants were harvested to measure different morpho-physiological and biochemical parameters.

2.3. Analysis of Plant Growth Parameters and Relative Water Content

The length from the shoot base to the leaf tip was measured for shoot length (SL) determination. Likewise, the root length (RL) was determined by measuring the length from the root base to the root tip. From each experiment, 25 seedlings were collected and weighed to determine the fresh weight (FW). The dry weight (DW) of shoot and root was determined after oven drying at 60°C for four days. The FWs and DWs of shoots and roots were expressed as mg seedling⁻¹. The relative water content (RWC) of the rice leaves was measured by the method previously described [35].

2.4. Determination of Photosynthetic Pigment, Carotenoids, and Total Soluble Protein

Chlorophylls (Chl) and carotenoids extraction and determined were done according to the method of Lichtenthaler [36]. Soluble protein content (SPC) was determined by the method developed by Bradford [37] with some modifications. Leaf tissues of rice plants were homogenized with 0.1 M NaOH and vortexed for 30 s followed by centrifuged at 5000 \times g. Then 100 μL of supernatant from the extract was added in 5 mL of Bradford reagent and was incubated for 25 min. Finally, absorbance was taken at 595 nm. The SPC of the samples were determined from the standard curve prepared using casein.

2.5. Determination of H₂O₂, Malondialdehyde Content, and In Situ O₂^{•-}

The levels of H₂O₂ in leaves of rice seedlings were determined by following the method of Velikova et al. [38], using an extinction coefficient of 0.28 $\mu\text{M}^{-1} \text{cm}^{-1}$. Malondialdehyde (MDA) content was measured in rice leaves according to the method of Heath and Packer [39], using an extinction coefficient of 155 $\text{mM}^{-1} \text{cm}^{-1}$. Superoxide (O₂^{•-}) was detected histochemically using nitroblue tetrazolium (NBT) according to the method of Tahjib-Ul-Arif et al. [40].

2.6. Measurement of Proline Content, Total Soluble Sugars, Total Phenolic Compounds, and Ascorbate Content

The method of Zhang and Huang [41] with minor modifications was used to measure the proline content of rice leaves. Total soluble sugars (TSS) was determined using the modified anthrone method as previously reported by Cihra and Brun [42]. Dried shoot samples (0.3 g) were extracted with 80% ethanol, and the resultant extract was filtered. Then 1.0 mL of diluted extract was mixed with 4 mL of 2% anthrone solution and the mixture was then heated for 10 min and then cooled on ice. The absorbance of the solution was then measured at 620 nm using a UV-VIS spectrophotometer (Shimadzu, UV-1201, Kyoto, Japan). The TSS contents of the samples were determined from the standard curve prepared using glucose.

The total phenolic compounds (TPC) was determined according to the method of Singleton and Rossi [43], using the Folin-Ciocalteu reagent. The AsA content was determined according to the method of Tahjib-Ul-Arif et al. [44].

2.7. Assay of Antioxidant Enzyme Activities

The activities of antioxidant enzymes were determined from the 3rd leaves of rice seedlings. Fresh leaf samples (0.05 g) were homogenized using pre-chilled mortar and pestle with 1.0 mL of 50 mM potassium phosphate buffer (pH 8.0) to extract leaf enzymes. The homogenates were centrifuged at 11,500×g for 10 min and the resultant supernatants were collected to analyze the activity of CAT (EC 1.11.1.6) [45], APX (EC 1.11.1.11) and POX (EC: 1.11.1.7) [46]. All procedures were carried out at 4°C and all of the spectrophotometric assays were performed using a UV-VIS spectrophotometer (Shimadzu, UV-1201, Kyoto, Japan).

2.8. Determination of Na⁺, K⁺, Ca²⁺ and Mg²⁺ Content

Twenty five-day-old rice shoots were harvested and washed rigorously with deionized water to eliminate surface contaminants. Seedlings were then oven-dried at 70°C for four days, and the finely powdered plant material was digested with HNO₃: HClO₄ (2:1v/v) mixture at 220°C for 2 h according to the method of Tahjib-Ul-Arif et al. [47]. Na⁺ and K⁺ contents were quantified by flame photometry (Jencon PFP 7, JENCONS-PLS, UK) and Ca²⁺ and Mg²⁺ contents were measured by titration, using disodium ethylene diamine tetra acetate, according to the methods of Schmid and Reilley [48].

2.9. Statistical Analysis

A one-way analysis of variance followed by Tukey's test was performed using Minitab 17.0. MetaboAnalyst 4.0 was used for cluster analysis and the generation of heatmap from normalized mean values [49]. The PCA analysis was done in R version 3.5.2 using the packages ggplot2, factoextra, and FactoMineR [50,51]

3. Results

3.1. Exogenous NO improved germination parameters of rice seeds under chilling stress

The protective role of NO on the germination capacity of rice seeds under chilling stress was evaluated by determining the PL, PFW, FGP, GRI, CVG and VI. Under chilling stress condition, PL, PFW, FGP, GRI, CVG and VI were markedly decreased by 29.09%, 47.06%, 24.38%, 55.17%, 32.32%, and 68.25%, respectively, compared with that of the control treatment (Table 1). On the other hand, exogenous application of SNP on chilling-stressed seedlings elevated the PL, PFW, FGP, GRI, CVG and VI by 36.75%, 55.55%, 19.67%, 48.25%, 18.65% and 103.08%, respectively, whereas declined MGT (by 12.8%) relative to only chilling-stressed rice seedlings (Table 1). Moreover, compared with control, RaL, RaFW, and MGT were increased significantly by 46.15%, 38.89% and 42.26% in response to only chilling-stressed treatment (Table 1).

Table 1 Effects of exogenous sodium nitroprusside (SNP) on germination indices and growth parameters of rice at germination stage grown under chilling stress

Treatment	Germination indices					Growth parameters			
	FGP (%)	MGT (days)	GRI (% day ⁻¹)	CVG (% day ⁻¹)	VI	PL (cm)	RaL (cm)	PFW (mg)	RaFW (mg)
C	80.67±1.09 ^b	2.07±0.0 ^c	52.79±0.6 ^b	48.47±0.5 ^b	290.27±1.0 ^b	5.50±0.05 ^a	3.90±0.3 ^b	51.00±1.69 ^a	6.00±0.05 ^b
CH	61.00±1.24 ^d	2.94±0.0 ^a	23.67±1.0 ^d	32.81±0.6 ^d	92.16±3.48 ^d	3.90±0.09 ^b	5.70±0.1 ^a	27.00±1.24 ^b	8.33±0.27 ^a
SNP	86.67±1.19 ^a	1.85±0.0 ^d	59.20±1.09 ^a	51.60±0.6 ^a	325.59±6.6 ^a	5.50±0.05 ^a	3.20±0.1 ^b	47.00±2.05 ^a	6.33±0.27 ^b
CH+SNP	73.00±0.47 ^c	2.56±0.0 ^b	35.09±0.19 ^c	38.92±0.3 ^c	187.16±2.9 ^c	5.33±0.05 ^a	5.33±0.0 ^a	42.00±1.41 ^a	7.33±0.27 ^{ab}

Data presented are means of three independent replicates ± standard errors ($n = 3$). Different letters in each column represent significant differences at $P < 0.05$ (Tukey's honest significant differences test). "C", 0 h chilling day⁻¹ + 0 µM SNP (control); "CH", 8 h chilling day⁻¹ + 0 µM SNP; "SNP", 0 h chilling day⁻¹ + 100 µM SNP; and "CH+SNP", 8 h chilling day⁻¹ + 100 µM SNP. Final germination percentage (FGP); mean germination time (MGT); germination rate index (GRI); coefficient of velocity of germination (CVG); vigor index (VI); plumule length (PL); radicle length (RaL); plumule fresh weight (PFW) and radicle fresh weight (RaFW).

3.2. Exogenous NO enhanced growth, biomass and photosynthetic pigments of rice plants under chilling stress

To determine whether SNP was involved in improving the toxic effects of chilling, we monitored the growth parameters of rice seedlings. Rice seedlings treated with only chilling displayed a significant reduction in SL, SFW, RFW, SDW and RDW (by 17.92%, 41.04%, 49.22%, 51.55%, and 44.32%, respectively) as compared with that of the non-stressed seedlings (Table 2). However, exogenous application of SNP to chilling-stressed plants relieved the lethal effects of chilling stress by significantly enhancing the SL, SFW, RFW, SDW and RDW (by 22.99%, 57.09%, 50.78%, 29.075%, and 86.92%, respectively) in comparison with that of the only chilling-stressed plants. Exogenous application SNP on chilling free rice seedlings showed no significant effect on the growth of rice seedlings except the SFW compared with the control (Table 2).

Table 2 Effects of exogenous sodium nitroprusside (SNP) on growth and photosynthetic pigments of rice seedlings grown under chilling stress

Treatment	Shoot Length (cm seedling ⁻¹)	Root Length (cm seedling ⁻¹)	Fresh weight (mg seedling ⁻¹)		Dry weight (mg seedling ⁻¹)		Chlorophyll content (mg g ⁻¹ FW)		
			Shoot	Root	Shoot	Root	Chl a	Chl b	Chl a+b
C	17.67±0.36 ^a	8.17±0.49 ^{ab}	203.67±0.72 ^a	27.00±0.47 ^a	12.83±0.36 ^a	8.53±0.53 ^a	0.27±0.03 ^a	0.09±0.01 ^{ab}	0.37±0.03 ^a
CH	14.50±0.24 ^b	10.00±0.47 ^{ab}	98.67±1.44 ^d	15.03±0.65 ^c	7.57±0.30 ^c	4.33±0.35 ^b	0.06±0.01 ^b	0.02±0.01 ^c	0.08±0.02 ^c
SNP	18.47±0.42 ^a	6.77±0.99 ^b	190.67±2.41 ^b	25.67±0.98 ^{ab}	12.97±0.16 ^a	7.03±0.26 ^a	0.35±0.02 ^a	0.13±0.02 ^a	0.48±0.01 ^a
CH+SNP	17.83±0.36 ^a	11.17±0.59 ^a	155.00±0.94 ^c	22.67±0.27 ^b	9.77±0.32 ^b	8.10±0.20 ^a	0.16±0.01 ^b	0.06±0.01 ^{bc}	0.22±0.01 ^b

Data presented are means of three replicates ± standard errors ($n = 3$). Different letters in each column represent significant differences at $P < 0.05$ (Tukey's honest significant differences test). "C", 0 h chilling day⁻¹ + 0 µM SNP (control); "CH", 8 h chilling day⁻¹ + 0 µM SNP; "SNP", 0 h chilling day⁻¹ + 100 µM SNP; and "CH+SNP", 8 h chilling day⁻¹ + 100 µM SNP.

We determined the Chl content of rice leaves to know the role of SNP in protecting photosynthetic machinery under chilling stress. Chilling treatment caused a significant decrease in Chl *a*, Chl *b* and Chl *a+b* contents (by 76.99%, 82.89% and 78.51%, respectively) compared with that of control (Table 2). On the contrary, exogenously applied SNP significantly alleviated the inhibitory effects of chilling (by 157.54%, 286.21%, and 183.85%) relative to only chilling-stressed treatment. Also, a significant elevation of Chl *a*, Chl *b* and Chl *a+b* content, with relative to the control, was observed in SNP-treated rice seedlings without chilling stress (Table 2).

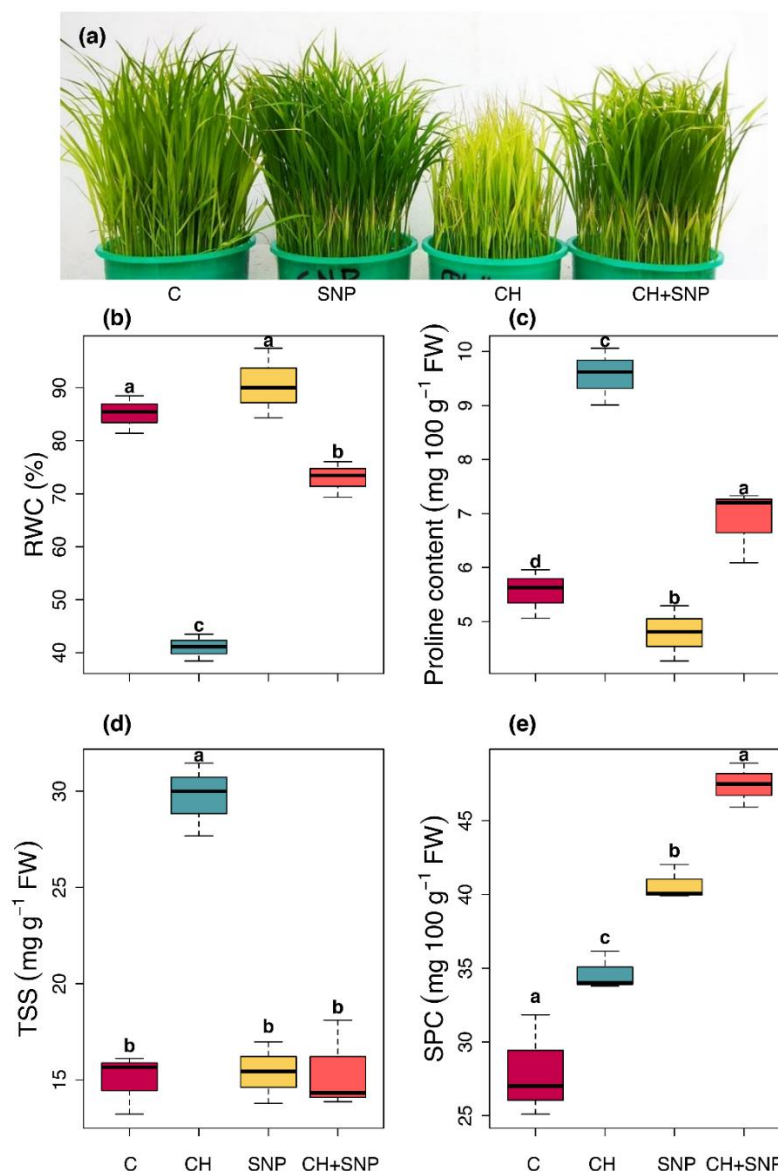


Figure 1. Effects of exogenous sodium nitroprusside (SNP) on (a) phenotype, (b) relative water content (RWC), (c) proline content, (d) total soluble sugar (TSS), and (e) soluble protein content (SPC) of rice seedlings grown under normal or chilling stress condition for 8 days. Data represented as the means of three independent replicates for each treatment ($n = 3$). The vertical bars indicate standard errors. Different letters

on the top of each bar denote statistically significant differences at $P < 0.05$, based on Tukey's test. "C", 0 h chilling day⁻¹ + 0 μ M SNP (control); "CH", 8 h chilling day⁻¹ + 0 μ M SNP; "SNP", 0 h chilling day⁻¹ + 100 μ M SNP; and "CH+SNP", 8 h chilling day⁻¹ + 100 μ M SNP.

3.3. Exogenous NO improved phenotype, water relation and regulated osmolyte status of rice plants under chilling stress

Application of SNP on chilling-stressed rice seedling improved the phenotype (Figure 1a; the photograph was taken on day 25th of the experiment). Chilling stress-induced yellowing symptoms on the leaves and deteriorated the plant growth in terms of shoot length of rice seedlings (Figure 1a). On the other hand, exogenous application of SNP mitigated the lethal effects of chilling stress as displayed improved phenotypes of rice seedlings (Figure 1a). Chilling stress declined the RWC of rice plants by 44.32% but increased proline, TSS and SPC by 72.31%, 98.07% and 23.82% compared with that of the non-stressed treatment (Figure 1b-e). However, a notable increase in RWC and SPC (77.75% and 36.92%, respectively), but a marked decline in proline and TSS (28.12% and 48.05%, respectively), were recorded in SNP-treated chilling-stressed seedlings compared with only chilling-stressed rice seedlings (Figure 1b-e). Moreover, the application of SNP to non-stressed rice seedlings resulted in enhanced SPC (by 45.30%) but in declined proline content (by 13.69%) as compared with that of control rice seedlings (Figure 1b, e).

3.4. Exogenous NO decreased lipid peroxidation, ROS accumulation and enhanced activities of antioxidant enzyme of rice plants under chilling stress

To examine whether exogenous SNP mitigated chilling-induced excess ROS accumulation and membrane damage in rice seedlings, we examined O₂^{•-}, H₂O₂ and MDA contents. Superoxide (O₂^{•-}) was visualized histochemically in leaves by NBT staining. Chilling-stressed rice seedlings showed the highest O₂^{•-} accumulation (observed as dark blue spots within the leaf blade) (Figure 2a). In contrast, SNP-treated chilling-stressed seedlings showed less O₂^{•-} accumulation as compared with seedlings treated with chilling stress only (Figure 2a).

The production of MDA and H₂O₂ was markedly increased by 85.74% and 197.63 % at only chilling treatment as compared with that of the control treatment (Figure 2b, c). Supplying SNP to the chilling-stressed rice seedlings resulted in reductions in MDA and H₂O₂ production by 25.55% and 36.91% in comparison with only chilling-stressed rice seedlings (Figure 2b, c). Furthermore, under non-stressed conditions, exogenous SNP exhibited no significant increase in the contents of MDA and H₂O₂ (Figure 2b, c).

The rice seedlings exposed to chilling stress had increased APX and POX activity by 29.75% and 94.19% at chilling treatment, respectively, as compared with that of the control (Figure 3b, c). In contrast, SNP supplementation to chilling-stressed seedlings further showed enhancement of APX and POX activity by 28.40% and 31.80%, respectively, as compared with only chilling-stressed seedlings (Figure 3b, c). The significant declination in CAT activity by 40.60% was recorded at chilling treatment relative to that of the control treatment. The SNP application to the chilling-stressed rice seedling elevated the CAT activity by 62.68% compared with only chilling-stressed seedlings (Figure 3a). Moreover, in non-stressed plants, the application of SNP enhanced CAT, APX, and POX activity significantly by 13.60%, 94.71%, and 55.01% in comparison with that of the control seedlings (Figure 3a-c).

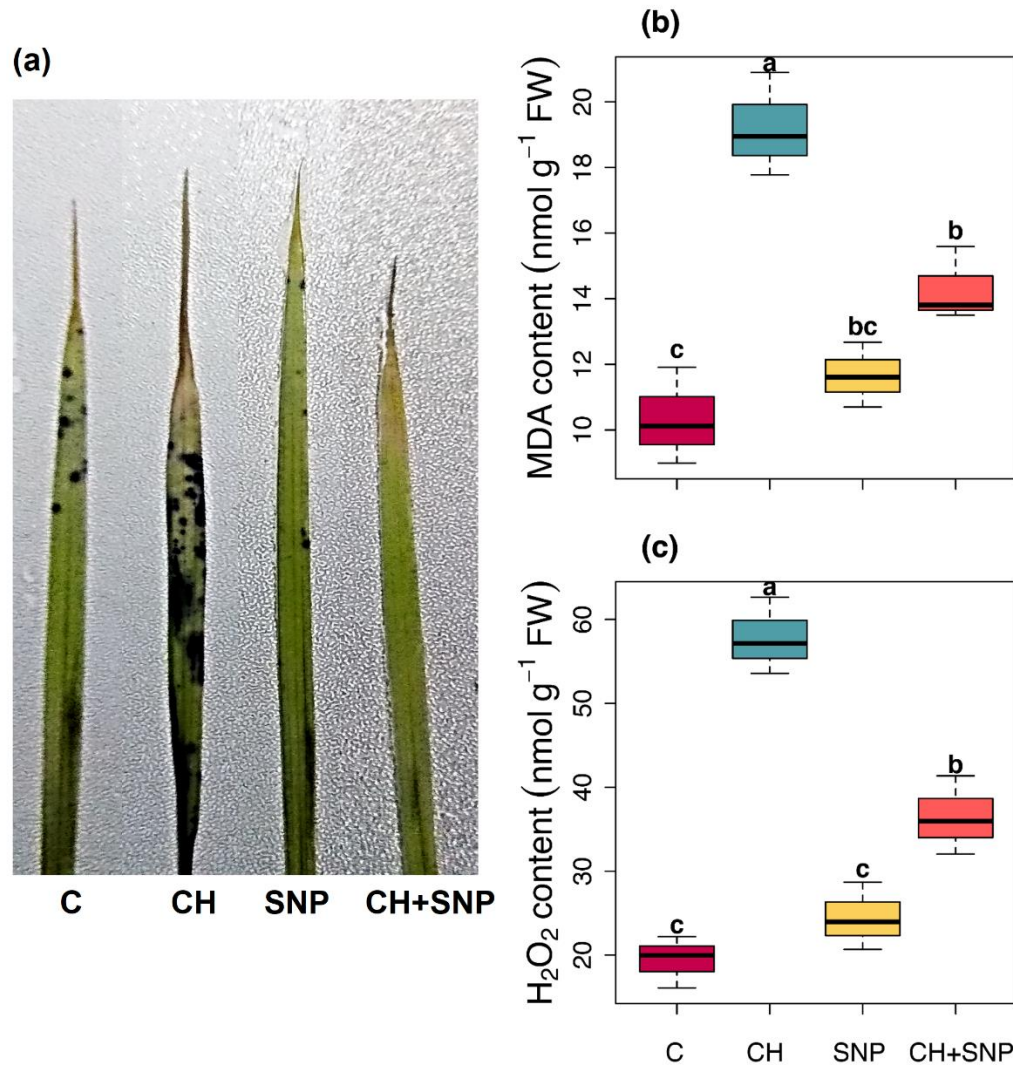


Figure 2. Effects of exogenous sodium nitroprusside (SNP) on (a) superoxide accumulation, (b) malondialdehyde (MDA) content, and (c) H₂O₂ content of rice seedlings grown under normal or chilling stress condition for 8 days. Plotted data represent the average (\pm) of three independent replicates for each treatment ($n = 3$, three leaves per replicate). The vertical bar indicates the standard error. The letter on top of each bar denotes a statistically significant difference at $P < 0.05$ (Tukey's honest significant differences test). "C," 0 h chilling day⁻¹ + 0 μ M SNP (control); "CH," 8 h chilling day⁻¹ + 0 μ M SNP; "SNP," 0 h chilling day⁻¹ + 100 μ M SNP; and "CH + SNP," 8 h chilling day⁻¹ + 100 μ M SNP.

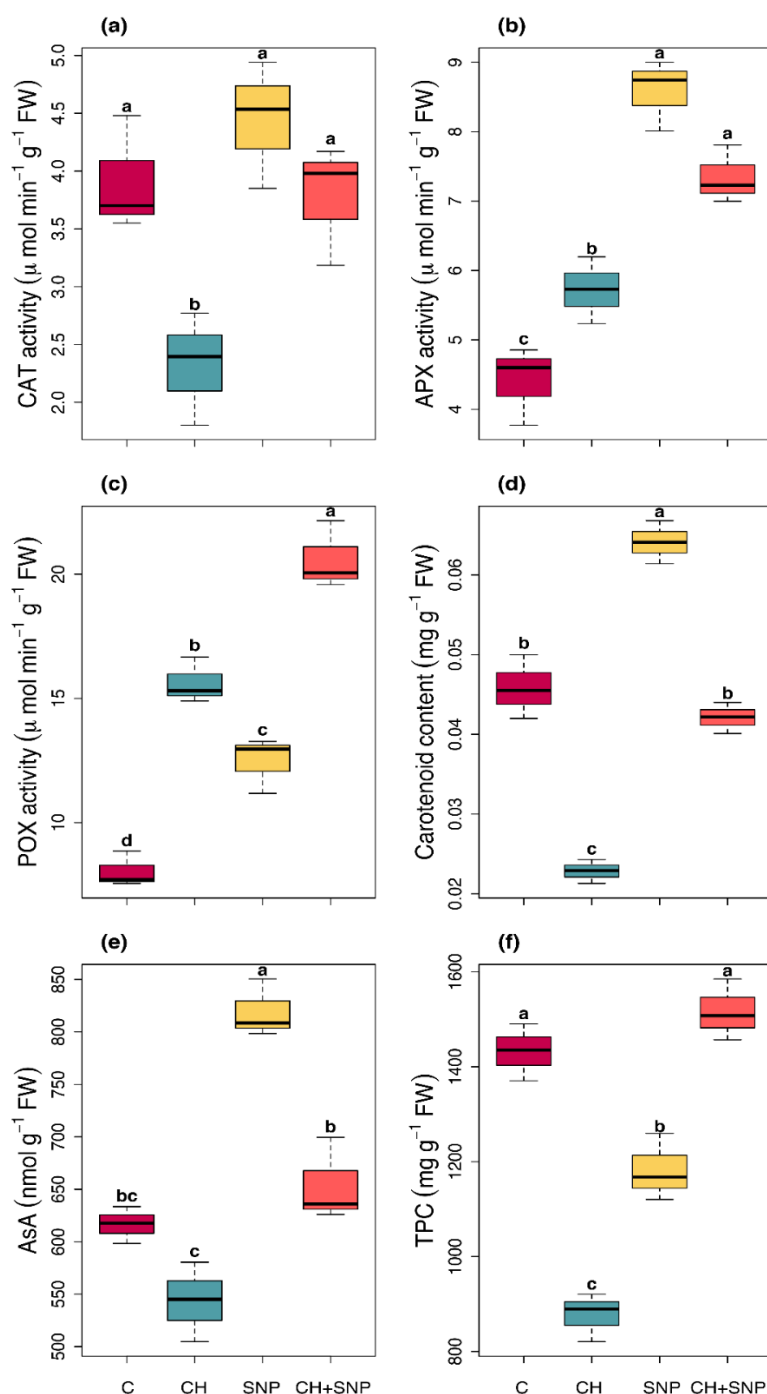
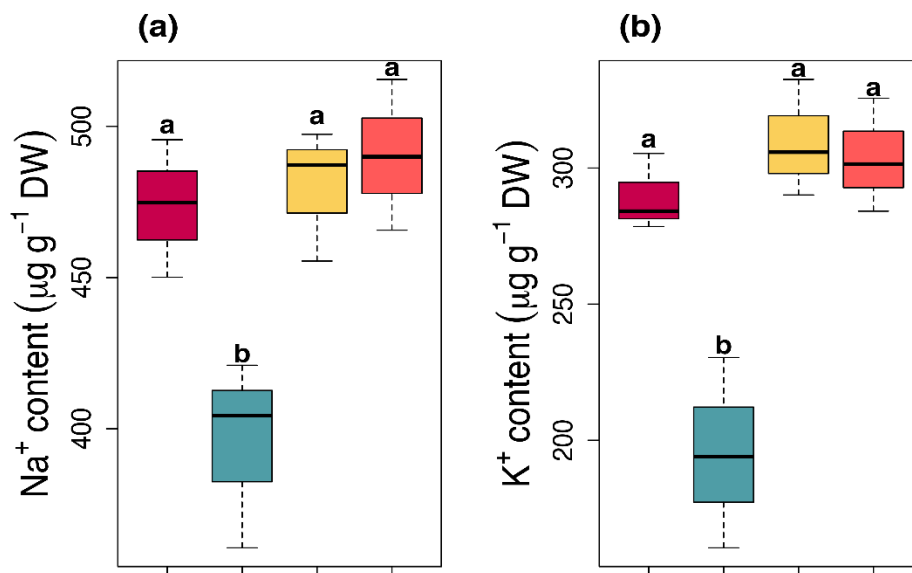


Figure 3. Effects of exogenous sodium nitroprusside (SNP) on (a) catalase (CAT) activity (b), ascorbate peroxidase (APX) activity, (c) peroxidase (POX) activity, (d) carotenoids content, (e) ascorbic acid (AsA) content, and (f) total phenolic compounds (TPC) content of rice seedlings grown under normal or chilling stress condition for 8 days. Data represented in figures are the mean (\pm) of three independent replicates for each treatment ($n = 3$, three leaves per replicate). The vertical bar indicates standard error. Different letters represent significant differences at $P < 0.05$ (Tukey's honest significant differences test). "C," 0 h chilling day $^{-1}$ + 0 μM SNP (control); "CH", 8 h chilling day $^{-1}$ + 0 μM SNP; "SNP", 0 h chilling day $^{-1}$ + 100 μM SNP; and "CH + SNP", 8 h chilling day $^{-1}$ + 100 μM SNP.

3.5. Exogenous NO enhanced non-enzymatic antioxidants and mineral content of rice plants under chilling stress

Several non-enzymatic antioxidants viz. AsA, carotenoids and TPC were measured to assess the role of SNP on these non-enzymatic antioxidants. In comparison with that of the control seedlings, the level of carotenoid and TPC declined significantly (by 50.18% and 38.77 %, respectively), whereas AsA content decreased non-significantly in chilling-stressed seedlings (Figure 3d-f). However, chilling-stressed plants supplemented with SNP showed enhancement in AsA, carotenoids, and TPC (by 20.31%, 84.37% and 72.94%, respectively) in comparison with only chilling-stressed seedlings (Figure 3d-f). Furthermore, the application of SNP to non-stressed rice plants resulted in 32.87% and 39.85 % higher levels of AsA and carotenoid contents whereas 17.43% lower level of TPC as compared with that of the non-treated control rice seedling (Figure 3 d-f).

The Na^+ and K^+ contents were declined by 16.51% and 32.61 % in chilling-stressed rice seedlings, compared with that of the control seedlings (Figure 4 a,b). However, supplementation of SNP to chilling-stressed rice seedlings showed amplification in Na^+ and K^+ contents by 24.05% and 55.77% as compared with chilling-stressed rice seedlings (Figure 4 a,b). Then the Mg^{2+} and Ca^{2+} displayed no significant change in rice seedlings at chilling treatment relative to that of the control treatment (Figure 4 c,d). On the other hand, supplementation with SNP to chilling-stressed seedlings showed enhanced shoot Na^+ , K^+ , Mg^{2+} and Ca^{2+} content by 24.05%, 55.77%, 7.14% and 95.06 % as compared to only chilling-stressed rice seedlings (Figure 4). Moreover, in non-stressed conditions, SNP treatment showed no significant change in Na^+ and K^+ content but significant change on Mg^{2+} and Ca^{2+} content by 16.75% and 312.08% respectively as compared with that of the control condition (Figure 4).



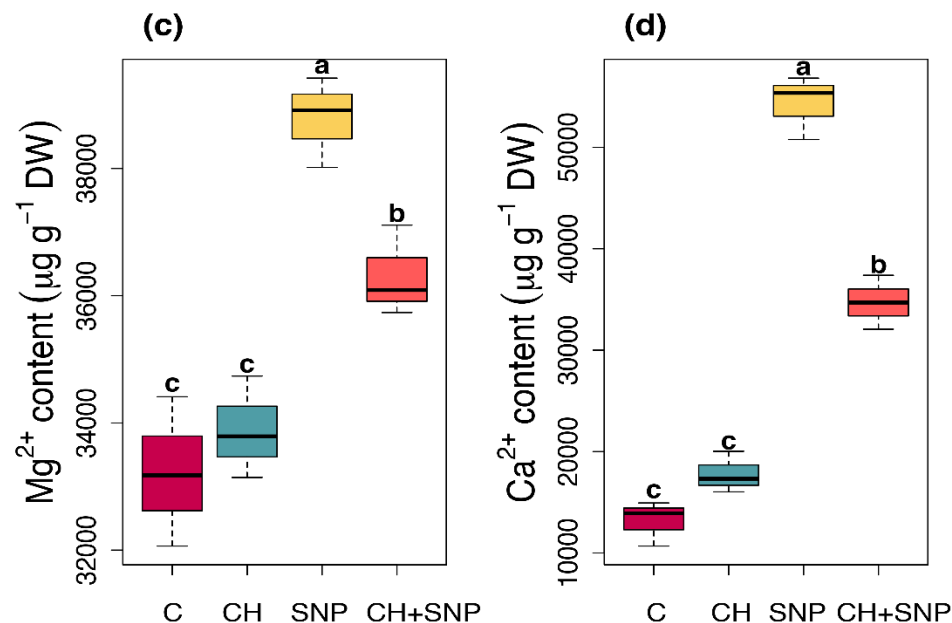


Figure 4. Effects of exogenous sodium nitroprusside (SNP) on: (a) sodium content (Na^+) content (b) potassium content (K^+) (c) magnesium content (Mg^{2+}) content (d) calcium content (Ca^{2+}) of rice seedlings under normal or chilling stress condition for 8 days. Data represented in figures are the mean (\pm) of three replicates for each treatment ($n = 3$, three leaves per replicate). The vertical bar indicates standard error. The letter on the top of bar denotes the statistically significant difference at $P < 0.05$ (Tukey's honest significant differences test). "C", 0 h chilling day⁻¹ + 0 μM SNP (control); "CH", 8 h chilling day⁻¹ + 0 μM SNP; "SNP", 0 h chilling day⁻¹ + 100 μM SNP and "CH + SNP", 8 h chilling day⁻¹ + 100 μM SNP.

3.6. Hierarchical clustering, PCA and Correlation analysis of NO-treated rice seedling under normal and chilling conditions

Average mean values of all morpho-physiological and biochemical data were set to perform heatmap and hierarchical clustering, and PCA. Hierarchical clustering revealed the classification of these parameters into three clusters (cluster-A, cluster-B, and cluster-C) (Figure 5a). Cluster-A encompass germination indexes (PL, FGP, VI, GRI, CVG and PFW), growth-related attributes (SL, RDW, SFW and SDW), photosynthetic pigment (Chl *a*, Chl *b* and Chl *a+b*), antioxidants (Car, AsA, TPC and CAT), mineral contents (Na^+ and K^+) and RWC of leaves of rice plants. In comparison with control, all the parameters of cluster-A displayed a decreasing trend in chilling-stressed rice plants, while an increasing trend was observed in SNP-treated chilling-stressed rice plants compared with chilling-stressed only plants (Figure 5a). Interestingly, only SNP treated non-stressed rice plants showed the highest level of increase of these parameters classified under cluster-A. The TSS, proline, RaFW, MDA, H_2O_2 , RL, RaL and MGT level in leaves were grouped in cluster-B. Compared with non-stressed treatment, Cluster-B parameters showed an increasing pattern in only chilling-treated rice plants, while they displayed a decreasing trend in SNP-treated plants and SNP-treated chilling-stressed rice plants. Cluster-C comprises APX, Mg^{2+} , Ca^{2+} , POX and TSP content, which showed a moderate increase in chilling-stressed rice plants compared with control plants whereas showed the highest increase in SNP-treated chilling-stressed plants compared with only chilling-stressed rice plants.

The PCA was performed to determine the association of the morpho-physiological and biochemical parameters with the treatment groups. The PCA biplot reveals clear segregation of the four

treatment groups (C, CH, SNP and CH+SNP) and their biological replicates. The first two PCA components combinedly explained 81.6% of the data variability (Figure 5b). SPC, POX, APX, Mg^{2+} , and Ca^{2+} were loaded into the PC1, whereas CAT, Chl *a*, Chl *b*, Chl *a+b*, RWC, PL were loaded into the PC2 (Supplementary Table 1). Results displayed that cluster-A variables were moderately and strongly associated with control and SNP treatments, respectively, whereas heatmap cluster-C variables were intensely interlinked to SNP application followed by chilling treatments. Heatmap cluster-B variables were strongly associated with chilling treatment only (Figure 5a, b).

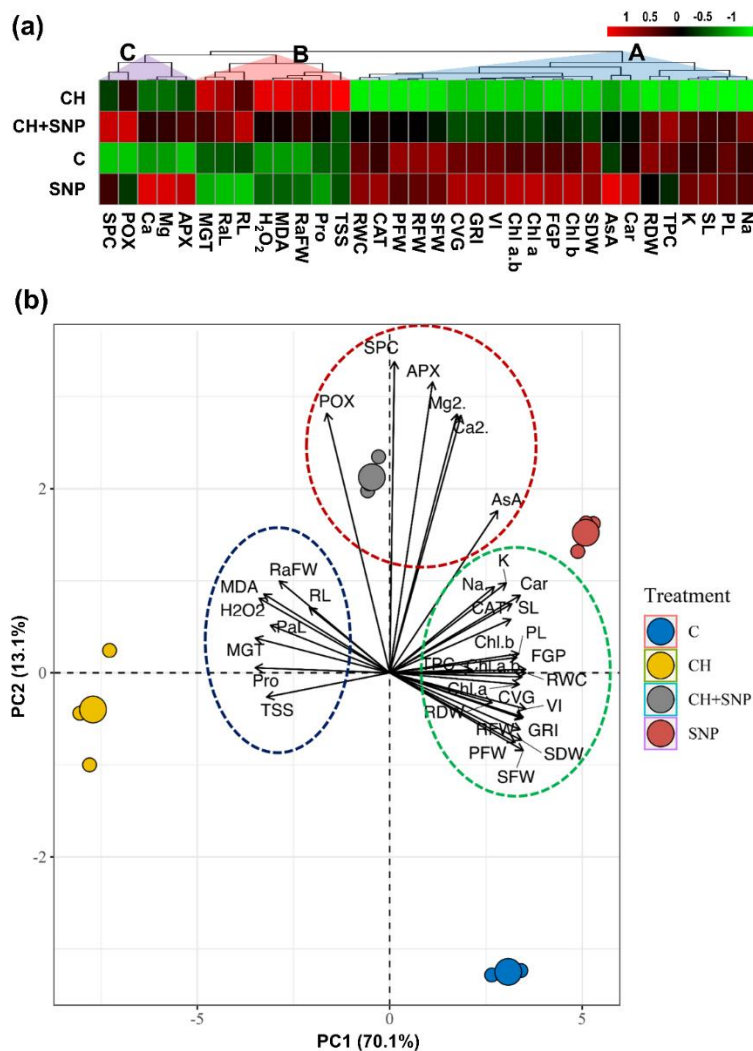


Figure 5. (a) Hierarchical clustering with heatmap and (b) principal component analysis (PCA) show the treatment-variable relationships. In hierarchical clustering and heatmap, the mean values of various parameters obtained in this study were normalized and clustered. Three distinct clusters (cluster-A, -B and -C) were identified at the variable level. The colour scale displays the intensity of normalized mean values of different parameters. The entire dataset was analyzed using PCA. The variables included final germination percentage (FGP), mean germination time (MGT), germination rate index (GRI), coefficient of velocity of germination (CVG), vigor index (VI), shoot length (SL), root length (RL), shoot fresh weight (SFW), root fresh weight (RFW), radicle length (RaL), plumule length (PL), radicle fresh weight (RaFW), plumule fresh weight (PFW), catalase (CAT), ascorbate peroxidase (APX), peroxidase (POX), malondialdehyde (MDA), hydrogen peroxide (H_2O_2), chlorophyll *a* (Chl *a*), chlorophyll *b* (Chl *b*), chlorophyll *a+b* (Chl *a+b*), carotenoid (Car), ascorbate (AsA), proline (Pro), total soluble sugar (TSS), total phenolic contents (TPC), soluble protein content (SPC), shoot dry weight (SDW), root dry weight (RDW), relative water content (RWC), sodium (Na),

potassium (K), magnesium (Mg²⁺), calcium (Ca²⁺). “C”, 0 h chilling day⁻¹ + 0 μ M SNP (control); “CH”, 8 h chilling day⁻¹ + 0 μ M SNP; “SNP”, 0 h chilling day⁻¹ + 100 μ M SNP and “CH + SNP”, 8 h chilling day⁻¹ + 100 μ M SNP.

4. Discussion

Finding an economical and effective technique to reduce damages caused by chilling stress in rice is urgent because it causes significant yield loss. Exogenous application of some signaling molecules have enormous potentiality to reduce the deleterious effects of abiotic stresses such as chilling stresses from both theoretical and practical perspectives [52,53]. Nitric oxide (NO), a signaling molecule and secondary messenger, has been shown that to counteract with excessive ROS-induced toxicity and positively regulate many physiological processes under different stress conditions [18,54,55]. Therefore, NO could potentially be used to reduce the adverse effects of chilling stress. In the present study, we evaluated whether NO would enhance the chilling tolerance of rice seedlings when applied exogenously.

Seed germination is the first step for successful crop establishment, which critically depends on an optimum temperature, for example, 25-35°C for rice [27]. Usually, the temperature below the optimum level severely hampers the seed germination process [3]. In the current experiment, chilling stress (8 h 5.0 \pm 1.0°C day⁻¹) inhibited the germination indices, such as FGP, MGT, GRI, CVG and VI (Table 1), which corroborate with the previous reports in rice under chilling and salt stress [32,56,57,58]. Priming with NO, on the other hand, successfully ameliorated chilling-induced impairment of seed germination in rice (Table 1). Besides, exogenous NO pretreatment enhances β -amylase activity in seeds [59,60], which might contribute to improving the seed germination (Table 1).

The adverse impacts of abiotic stress on plants can be evaluated by measuring the morphological parameters such as growth and biomass of plants [61]. In the present experiment, the SL, SFW, RFW, SDW and RDW of the chilling-stressed rice plants were lower compared with that of control plants (Table 2), which perhaps occurred due to the disruption of water relation (Figure 1b), elevation of ROS content and lipid peroxidation (Figure 2) and an imbalance of ionic homeostasis (Figure 4). Exogenous application of NO, however, recuperated the growth performance of chilling stressed rice seedling as evident by their improved phenotypic appearance (Figure 1a) and growth-related features, including SL, SFW, RFW, SDW and RDW (Table 1), perhaps by decreasing ROS accumulation and lipid peroxidation (Figure 2a, c). Moreover, NO-induced growth improvement might be ascribed to its role in increasing membrane fluidity and accelerating cell expansion processes [62]. Consistent with our results, it has also been reported that exogenous NO improved plant growth of cucumber (*Cucumis sativus*) [21], wheat [22] and Chinese cabbage (*Brassica rapa* subsp. *Pekinensis*) [23] under chilling stress condition. Moreover, our results were verified by PCA, which suggested that chilling-stressed rice plants that treated with NO exhibited a stronger correlation with growth-related attributes in comparison with chilling-stressed only plants (Figure 5b). Surprisingly, in the present study, RL of chilling-stressed plants increased slightly compared to the control plants (Table 2), which differs that of the findings of Fan *et al.* [23]. A similar increase of RL was also observed previously in *Silene vulgaris* under mild drought conditions [63].

The detailed study on the response of rice plants to chilling stress suggested that chilling-induced growth retardation might be due to the imbalance in ionic homeostasis. However, exogenous NO application reestablished Na⁺ and K⁺ balance and further increased the Ca²⁺ and Mg²⁺ contents in rice plants which might improve the growth of rice plants (Table 1; Figure 4). This finding was strongly supported by PCA results which showed a relatively stronger correlation of mineral contents with NO-treated chilling-stressed plants than that with the chilling-stressed only plants (Figure 5b). Chilling stress decreased the K⁺ content (Figure 4b), which might trigger the partial or complete closure of stomata, thereby decreasing the water flow (Figure 1b) and CO₂ assimilation [64] in rice plants, as K⁺ is crucial for stomatal opening and closing [65]. Under lower K⁺ level, Na⁺ can perform some metabolic functions, due to the structural similarity of two ions in solution, thereby conferring tolerance to plants [66]. However, our results showed

that the chilling stress decreased the Na⁺ content in rice plants as well (Figure 4a). A similar decrease of Na⁺ and K⁺ contents was observed under various abiotic stress conditions [67,68,69,70]. Furthermore, non-stressed plants treated with NO showed higher accumulation of Ca²⁺ and Mg²⁺, which indicated that NO might upregulate the divalent cations uptake capacity of rice plants which led to the further enhancement of plant growth (Table 1; Figure 4c, d).

Along with the growth, chilling stress drastically declined the Chl pigment content (Chl *a* and Chl *b*) as compared with that of control (Table 3). A similar decline of chlorophylls was also reported in Bermuda grass (*Cynodon dactylon*) [71] and *Stevia rebaudiana* [72] plants under chilling stress. Exogenous application of NO increased Chl content in chilling-stressed rice plants (Table 3), as also observed in Chinese cabbage [23] and cucumber [21] under chilling stress. This enhancement of Chl content occurred might be due to the elevation of the uptake and transportation of Mg²⁺ (Figure 4c), which is indispensable for chlorophyll synthesis [73], decreased lethal ROS and NO-mediated stability and integrity of the subcellular structures under stressed condition [74]. This enhanced Chl content might enhance the growth-related attributes of NO-treated chilling-stressed rice plants (Table 2). Moreover, our PCA findings endorsed that chilling-exposed rice plants treated with NO had a stronger positive connection with Chl content than chilling-stressed only rice plants (Figure 5b).

Osmoregulation supports plants to sustain proper growth and development under stressful environments [3]. Thus, plant cell accumulates compatible solutes like proline, soluble carbohydrates, and SPC to maintain the osmotic potential and consequently retains plant water balance [75,76,77]. Our data revealed that only chilling-stressed plants showed a sharp accumulation of proline, TSS and SPC in rice leaves (Figure 1b, c, d), which corroborates with the findings in cold-stressed pepper (*Capsicum annuum* L.) [78], rice [79], *Stevia rebaudiana* [72], maize (*Zea mays* L.) [80] and wheat [81]. However, a sharp drop of RWC (Figure 1b) indicated that the increased level of these osmolytes was insufficient to alleviate the chilling effect on the growth of rice plants (Figure 1a). On the other hand, supplementation of NO allowed plants to maintain an optimum amount of water potential which consequently decreased proline and TSS content (Figure 1b, c). Moreover, we found a negative correlation between proline content and RWC (Supplementary Figure 1). On the contrary, exogenous application of NO further enhanced SPC as also observed in cucumber [82] and chickpea [24] under salinity stress.

Chilling stress alters the coordination between generation and scavenging of ROS in plant cells, thereby triggering cell death [83]. Our results showed that O₂^{•-} and H₂O₂ contents increased in the leaves of rice plants when subjected to chilling stress (Figure 2a, c), which triggered the oxidation of membrane lipids [84], and consequently elevated MDA content (Figure 2b). Furthermore, a positive correlation was found between H₂O₂ and MDA content (Supplementary Figure 1), and our PCA results showed a strong correlation of ROS and MDA with chilling stress only treatment (Figure 5b). The enhanced production of H₂O₂ might be due to the imbalanced antioxidant enzyme activities (Figure 3a-c). These results demonstrated the adverse effects of chilling stress, i.e., stunted growth of rice seedlings through increasing ROS level and lipid peroxidation under chilling stress, that was in line with the previous studies on wheat [22,84], chickpea [85] and maize (*Zea mays* L.) [86]. On the other hand, exogenous application of NO to rice plants relieved adverse effects of chilling-induced oxidative stress, as supported by declined O₂^{•-}, H₂O₂ and MDA contents in rice plants (Figure 2a-c), which was in accordance with previous findings in wheat [22], Chinese cabbage [23] and orange (*Citrus sinensis*) [87] under chilling stress. Similarly, our PCA results also confirmed that NO-treated salt-stressed treatment had a positive association with H₂O₂, MDA. However, the association prolonged to a smaller extent relative to chilling-stressed only treatment (Figure 5b), suggesting that NO treatment could reduce oxidative damages (Figure 2a-c) and facilitated improvements of growth parameters of the rice plants (Table 2).

The increase of antioxidants activities is related to better oxidant management under stressed conditions [88,89]. In this investigation, the activities of APX and POX increased whereas CAT activity decreased in chilling-stressed plants compared with that of control plants (Figure 3a-c). The activities of these studied antioxidant enzymes were insufficient to neutralize the excess ROS, as also evident by the higher accumulation of H₂O₂ and MDA in chilling-stressed rice plants (Figure 2a-c). Consequently, growth and biomass inhibition of rice plants were evident in only chilling-stressed plants (Table 2). A similar decrease of CAT activity under chilling stress was noticed in wheat plants [84] and an increase of APX and POX were found in cucumber and wheat plants under chilling stress [22, 90]. Application of NO in chilling-stressed plants enhanced CAT, APX, and POX activities compared to only chilling stressed plants (Figure 3a-c). Importantly, our result showed that CAT and H₂O₂ negatively correlate with each other (Supplementary Figure 1), and PCA revealed that NO-treated chilling-stressed treatment had a stronger positive association with POX, APX, and CAT activity than chilling-stressed only treatment (Figure 5b). These findings revealed positive effects of exogenous NO in boosting up ROS-scavenging ability, therefore the growth of rice seedlings under chilling stress. A similar findings have been demonstrated in several plant species, such as wheat [22] and orange [87]. The current study showed that chilling stress declined the AsA, carotenoids, and TPC (Figure 3d-f); similar results were observed in rice [91] and *Vitis vinifera* under chilling [92,93]; tomato [94] and *Solanum lycopersicum* [95] under salt stress. On the other hand, the application of NO to chilling-stressed plants significantly increased AsA, TPC, and carotenoids contents (Figure 3d-f), which was also supported by PCA that showed a stronger correlation between these antioxidant content and NO-supplemented chilling-stressed treatment (Figure 5b). Several studies have also reported the NO-mediated increase in the non-enzymatic antioxidants which alleviate a wide range of abiotic stress in various plants, such as *Solanum Lycopersicum* [95] and tomato [94] under salt stress. Enhanced non-enzymatic antioxidants might scavenge chilling-induced ROS (Figure 2a, c), which contributed to the enhanced growth and development of rice plants (Figure 1a; Table 2).

In conclusion, our study provides the first evidence of exogenous NO-mediated chilling stress tolerance mechanisms in rice plants. The beneficial effect of NO treatments might be attributed to the alleviation of chilling stress-induced over-accumulation of ROS, possibly by enhancing the activities of ROS-scavenging enzymatic antioxidants (CAT, APX, and POX) and non-enzymatic antioxidants (AsA, TPC, and carotenoids). Moreover, exogenous application of NO was found to be effective in osmotic adjustment by modulating the contents of proline, SPC, and TSS in the rice plants growing under chilling stress, which might help in promoting water absorption and retention of rice seedlings under chilling stress. Besides, NO treatment effectively upregulate the ionic homeostasis under chilling stress. Finally, our results propose that exogenous NO application could be a practical and effective approach in mitigating the adverse effects of chilling stress, subsequently to ensure sustainable rice production. Further molecular studies should be carried out to gain more in-depth insight into the better understanding of the comprehensive mechanisms of NO-induced chilling stress tolerance in rice plants.

Conflict of interest: The authors declare that they have no conflict of interest.

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