RESEARCH ARTICLE



The protective effects of polyamines on salinity stress tolerance in foxtail millet (*Setaria italica* L.), an important C4 model crop

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Abstract Soil salinity is a major abiotic stress that adversely affects crop growth, development and productivity worldwide. In this study, the individual and synergistic roles of putrescine (Put) and spermidine (Spd) in salinity stress tolerance of foxtail millet (*Setaria italica* L.) was assessed. In the present study, plants treated with combined biogenic amines Put + Spd possess very efficient antioxidant enzyme systems which help to control the uninhibited oxidation and protect the plants from oxidative damage by ROS scavenging. Additionally, lower concentration of Put + Spd under NaCl stress showed reduced hydrogen peroxide, electrolyte leakage and caspase-like activity than control. FTIR analysis underlying the ability of PAs induced tolerance and the chemical bonds of Put + Spd treated plants were reminiscent of control plants. Moreover, histochemical analysis with 2',7'-3,3'-Didichlorofluorescein diacetate (DCF-DA), aminobenzidine (DAB) and nitrotetrazolium blue chloride (NBT) revealed that ROS accumulation was inhibited by combined PAs under salt stress condition. These results showed that Put + Spd significantly improve the endogenous PAs, which enhance high-salinity stress tolerance by detoxifying ROS. For the first time, the synergistic ROS scavenging ability of Put along with Spd was investigated upon salinity tolerance in C4 model foxtail millet crop. Overall, our findings illustrated the implication for improving salinity tolerance of agronomically important crop species.

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Graphic abstract



Keywords Confocal laser scanning microscope · Fourier transform-infrared spectroscopy · Histochemical analysis · Polyamines · Reactive oxygen species · Salinity stress

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Abbrevia	lions
APX	Ascorbate peroxidae
CAT	Catalase
CLSM	Confocal laser scanning microscope
DAB	3,3-diaminobenzidine
DAO	Diamine oxidase
EL	Electrolyte leakage
FTIR	Fourier transform-infrared spectroscopy
GPX	Guaiacol peroxidise
GR	Glutathione reductase
GSH	Glutathione
GSSG	Glutathione disulfide
H_2O_2	Hydrogen peroxide
MDA	Malondialdehyde
NaCl	Sodium chloride
$NADH^+$	Nicotinamide adenine dinucleotide
NADPH	Nicotinamide adenine dinucleotide phosphate
NBT	Nitrotetrazolium blue chloride
PAO	Polyamine oxidase
PAs	Polyamines
POD	Peroxidase
Put	Putrescine
ROS	Reactive oxygen species

Introduction

RWC

SOD

Spd

Salinity stress is detrimental to overall crop plant growth, production and yield, which cause economic losses, threatening global agriculture, particularly in steppes and arid zones (Panuccio et al. 2018). Salinization affects \sim 20% of world agricultural lands (Zhao et al. 2017). In general, under high salt stress, plants undergo several morphological, biochemical, physiological, and molecular level alterations which eventually lead to cell damage and growth reduction in plants (Satish et al. 2016; Sun et al. 2020). Salinity stress could induce generation of reactive oxygen species (ROS) in plants. ROS are harmful to macromolecules, cell signalling in proliferation which includes singlet oxygen ($^{1}O_{2}$), hydrogen peroxide ($H_{2}O_{2}$) and superoxide anion radical (\overline{O}_2) , that leads to membrane damage, degradation of protein, nucleic acid and eventually cell death (Huang et al. 2013; Kissoudis et al. 2014; Singh et al. 2019). The level of ROS in the stressed plants was regulated by the antagonism between the ROS producers and the scavengers (Asada 2006). Plants possess several stress-defence mechanisms which include the

Relative water content

Superoxide dismutase

Spermidine

accumulation of compatible soluble solutes and the production of antioxidant enzymes such as superoxide dismutase (SOD), catalase (CAT), ascorbate peroxidase (APX), and other peroxidases (POXs) to remove the excess ROS (Miller et al. 2010). These defence mechanisms can also be induced and/or enhanced by certain chemical compounds such as methyl jasmonate, sodium nitroprusside, melatonin, salicylic acid, strigolactone and some polyamines (PAs) (Yoon et al. 2009; Van Ha et al. 2014; Khan et al. 2015; Savvides et al. 2016). Hence, it is imperative to establish a new approach to enrich high salinity tolerance in plants.

The exogenous application of plant growth regulators is found to be effective in improving the resistance against various abiotic stresses viz., water deficit, water submergence and salinity (Liu et al. 2015; Satish et al. 2018). Nowadays, PAs have been considered as new plant growth regulators which are used to regulate plant growth and to enhance the competence of plants for abiotic stress tolerance (Liu et al. 2015; Sang et al. 2016). PAs are lowmolecular weight aliphatic amines ubiquitously present in all the organisms with key functions such as the regulators of protein synthesis, DNA replication, cell division, growth and several biological activities (Roychoudhury et al. 2011). Spermidine (Spd), spermine (Spm) and putrescine (Put) are the most common natural PAs present in plants acting as signaling molecules in plant-environmental stresses (Tiburcio et al. 2014). Among these, Spd could be significantly active in plant stress signal transductions thus activating stress tolerance mechanism (Hu et al. 2012; Liu et al. 2015). Similarly, several researchers reported that Put can effectively reduce ROS accumulation and improve plant tolerance against stress induced toxicity (Mandal et al. 2013; Wang et al. 2013). Foliar spraying of Spd in plants has resulted in increased accumulation of PAs, proline, glutathione (GSH) and the antioxidant enzyme activities in stressed plants (Duan et al. 2008; Sun et al. 2020). Furthermore, the exogenous Spd found to improve stress tolerance to aging (Wang et al. 2000), drought (Nemeth et al. 2002; Zhou et al. 2015), salinity (Fariduddin et al. 2018), heat (Tang et al. 2018), chilling (Fu et al. 2019), heavy metal stress (Gong et al. 2016) and submergence (Liu et al. 2015) stresses in different plant species. However, the combination of PAs decreased the dosage of application and modulates stress tolerance. The efforts were made to study the antioxidant enzyme activities and ROS scavenging mechanism of PAs and synergistic effects of Put and Spd and their roles in foxtail millet salinity tolerance.

Foxtail millet (*Setaria italica* (L.) is an important small millet, serving as a staple food for many people in temperate, tropical and subtropical Asian and African countries (Sreenivasulu et al. 1999). Generally, foxtail millet plants

are nutritionally high value crop possessing effective stress tolerance mechanisms to various abiotic stresses such as drought and salinity (Sreenivasulu et al. 2000; Zhang et al. 2005). Previously, Sudhakar et al. (2015) reported that PAs may be associated with the salinity stress tolerance in foxtail millet.

However, it is still unclear how the exogenous PAs improve the response in foxtail millet plants. Additionally, the synergistic effect of Put with Spd upon high salinity stress has not been studied so far. In the present study, we have investigated the effects of exogenous PAs on protection against salinity stress in foxtail millet by analyzing plant growth, physiological, biochemical and antioxidant enzymes activity. To the best of our knowledge, this is the first study to explore the synergistic effect of biogenic amines (Put and Spd) on salinity stress tolerance in foxtail millet. Perhaps, this study will aid in future for computational transcriptomic analyses of this important C4 model species.

Materials and Methods

Plant material and experimental conditions

The healthy foxtail millet seeds procured from Tamil Nadu Agricultural University, Coimbatore, Tamil Nadu, India were surface sterilized according to Rathinapriva et al. (2018). Seeds were sown on MS medium (Murashige and Skoog 1962) buffered at pH 5.75 \pm 0.5 with 3% w/v sucrose, 0.8% agar-agar type I (Himedia, Mumbai, India) and sodium chloride (NaCl; 200 mM) (Himedia, Mumbai, India). The rationale for 200 mM NaCl was previously described by Sudhakar et al. (2015). The autoclaved MS medium was supplemented with various concentrations of putresine (Put; 0.5, 1, 1.5 and 2 mM), spermidine (Spd; 0.5, 1, 1.5 and 2 mM) (Sisco Research Laboratories, Mumbai, India) individually or in combination Put + Spd (0.25 + 0.25, 0.5 + 0.5, 1 + 1 and 1.5 + 1.5 mM)respectively. To determine more suitable PAs (Put, Spd and Put + Spd) concentrations for modulating the growth of shoot and root, the preliminary assays were performed. Four-days-old germinated seedlings were placed in a growth chamber (Sanyo Versatile Environmental Test Chamber, Japan, Model No MLR351H) programmed with temperature 25 °C (\pm 1 °C), 16 h light (200 µM photons $m^{-2} s^{-1}$), 8 h dark photoperiod and 70% relative humidity for 15 days. Assays were performed using 15 bottles (5 seeds per bottle) for each of the three independent biological repeats. Foxtail millet plants grown for 15 days on MS medium without NaCl and PAs are used as a control (unstressed) and in treated control, plants were exposed to 200 mM NaCl without PAs.

Effects of PAs on germination and growth parameters under salinity stress

The survival and growth of foxtail millet plants were assessed through germination percentage (GP) and the morphological factors such as fresh weight (FW), dry weight (DW), shoot and root length after 15 d of treatment. The germination rate was based on the percentage of emergence of seedling after 4 d of darkness incubation.

For DW measurement, plant samples were dried in oven at 70 °C for 48 h, then allowed to cool in room temperature (RT) and weighed.

The salinity induced physiological changes ameliorated by Put and Spd treated and untreated plants were assessed with leaf relative water content (RWC), leaf electrolyte leakage (EL) and caspase-like activity. RWC (%) was calculated by the following formula,

RWC (%) =
$$[(FW - DW)/(TW - DW)] \times 100$$

Leaf samples were allowed to dip overnight in deionized water for swelling, shortly air dried and weighed to determine the turgid weight (TW) (Sapeta et al. 2013). Electrolyte leakage (EL) was determined by the electrical conductivity method (Tabot and Adams 2013). The EL was calculated as:

 $EL(\%) = (E1/ET) \times 100$

where E1 is the initial electrical conductivity of the samples measured using a conductivity/TDS/ °C gauge (Cyberscan 200, Eutech Instruments, Singapore) and ET is the final electrical conductance of the solution. The caspaselike activity was assessed based on the protocol of Poborilova et al. (2013).

Photosynthetic pigment analysis

Based on the preliminary analyses, 1 mM of Spd, 1 mM of Put and 0.5 + 0.5 mM of Spd + Put were found to be effective and selected for further experiments. The total chlorophyll (a + b) and carotenoid content of the 200 mM NaCl with 1 mM Spd, 1 mM Put and Put + Spd (0.5 + 0.5 mM) treated plant leaves were measured according to Lichtenthaler and Buschmann (2001) method. Each sample of 0.1 g fresh leaves was weighed from the treated plants, finely ground and immersed in 98% methanol (Himedia, Mumbai, India). Samples were extracted by incubating at 4 °C for 24 h in complete darkness. The chlorophyll content of methanol extract was measured at 652.4 and 665.2 nm and carotenoid content was measured at 480 nm, respectively using UV-vis spectrophotometer (UV-2450, Shimadzu Analyticals, Japan). 98% methanol was used as a blank.

Determination of compatible soluble solute

Leaf protein content was quantified according to Bradford (1976) and bovine serum albumin (BSA) was used as a standard. Free proline content was extracted and determined by Bates et al. (1973) protocol on the basis of ninhydrin reaction and utilizing the L-Proline (Hi-Media, Mumbai, India) as a standard.

Determination of cell membrane damage

Hydrogen peroxide content in the treated plants was individually determined by the method described by Del Pozo and Lam (1998). Absorbance of H_2O_2 content was recorded with a UV–vis spectrophotometer (UV-2450, Shimadzu Scientific Instruments, Japan) at 390 nm. The H_2O_2 content was calculated through comparing its optical density value in the standard graph. Lipid peroxidation was detected by quantifying the total amount of malondialdehyde (MDA) contents measured by thiobarbituric acid (TBA) reaction method according to Heath and Packer (1968).

Assays of antioxidant enzyme activity and free radical production

In order to determine the reaction products of PAs oxidation, SOD, GR, APX, GPX and GSH were analyzed.

SOD activity was assayed according to Paoletti and Mocali (1990) by evaluating its ability to inhibit the oxidation of NADPH into NADP⁺ and it was measured by inhibition (50%) in the reduction of nitrobluetetrazolium chloride (NBT) at 340 nm for 5 min. CAT activity was determined by measuring the decomposition of H_2O_2 content at 240 nm for 3 min, as described by Aebi (1984). GR activity was determined as defined by Carlberg and Mannervik (1975). Enzyme assay was quantified in basic of oxidized glutathione decomposition of NADPH by monitoring the change in absorbance at 340 nm. APX assay was conducted according to Nakano and Asada (1981) and measured by estimating the rate of ascorbate oxidation at 290 nm for 3 min. Guaiacol peroxidase (GPX) activity was determined as described by Lin and Kao (1999). The enzyme activity was quantified in terms of tetraguaiacol formation in 1 min of increase in absorbance at 470 nm.

GSH level was measured by DTNB (5,5'-dithio-bis-[2nitrobenzoic acid]) to form yellow coloured 5-thio-2-nitrobenzoic acid reaction method and the glutathione was used as a standard according to Sedlak and Lindsay (1968). Reaction mixture was then measured at 412 nm.

Assay of polyamine and diamine oxidase activity

Diamine oxidase (DAO) and polyamine oxidase (PAO) activity were assessed as described by Su et al. (2005) by quantifying the generation of H_2O_2 as a product of the oxidation of PAs. The enzyme reaction was initiated by the addition of Spd for PAO and Put for DAO determination and change in the optical density at 555 nm for 1 min was considered as one enzyme activity unit.

Histochemical detection of H_2O_2 and \bar{O}_2

In situ H_2O_2 and \overline{O}_2 production in foxtail leaves exposed to 200 mM NaCl, Put, Spd (1 mM) and Put + Spd (0.5 + 0.5 mM) were detected by the chromogenic substrate 3,3-diaminobenzidine (DAB) and nitrotetrazolium blue chloride (NBT) (Sisco Research Laboratories, Mumbai, India). Briefly, youngest leaves were excised from treated and control foxtail millet plants and immediately immersed into 0.1% DAB in 10 mM (pH 7.0) potassium phosphate solution for 5 h at RT (Mostofa et al. 2015). The DAB staining solution was decanted, replaced with an acetic acid: ethanol (1:3; v/v) for destaining of chlorophyll and photographed. The experiment was conducted using triplicates with the total of 20 seedlings for each treatment.

Analysis of H₂O₂ in confocal laser scanning microscopy

The production of ROS in the foxtail millet leaves treated with water control, 200 mM NaCl and Put + Spd were observed with 2',7'–dichlorofluorescein diacetate (DCF-DA) staining as described by Rossi et al. (2017). Lower epidermis of young leaves were peeled and immersed in 10 mM Tris–HCl (pH 6.1) in dark for 2 h then the samples were transferred to DCF-DA (25 μ M) solution for 20 min. The stained leaf samples were washed twice with Tris–HCl buffer and then images were visualized under confocal laser scanning microscopy (CLSM) (LSM, Jena, Carl Zeiss, Germany). The DCF-DA fluorescence intensities were analyzed using COMSTAT2 software procured from Dr. Claus Sternberg, DTU Systems Biology, Technical University of Denmark, Denmark.

FTIR analysis

Fourier transform-infrared spectroscopic (FTIR) analysis was performed to understand the metabolic changes of NaCl (200 mM), control (unstressed) and PAs (Put, Spd and Put + Spd) exposed foxtail millet plants. In the experiment, 100 mg of dried leaves were ground to a fine powder and compressed into pellets with KBr using Pelletization method. FTIR spectra was observed in the range

of wavelengths 400 to 4000 cm^{-1} using Nicolet iS 5 FTIR instrument (Thermo Scientific, USA) and the peaks were identified using OMNIC software (Kannappan et al. 2017).

Statistical analysis

All the experiments were done in a completely randomized design with minimum of three replicates. Values in the text and tables were indicated Mean values \pm SD of three independent experiments. Statistical analysis was done by one-way analysis of variance (ANOVA) using Duncan's multiple range test, with Statistical Package for the Social Sciences software version 17.0 (SPSS, IBM Statistics) and taking $p \leq 0.05$ as significant.

Results

PAs enhances salinity stress tolerance in foxtail millet

The role of PAs in the enhancement of salinity stress tolerance was investigated by measuring the GP and growth



Fig. 1 Polyamines ameliorates salinity tolerance in foxtail millet. Control (Unstressed), NaCl (Treated control; 200 mM), NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM), NaCl + Spd (NaCl; 200 mM, Spermidine; 1 mM) and NaCl + Put + Spd (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM) (left to right)

parameters of foxtail millet plants (Fig. 1) (Fig. S1). After 14 d incubation with Put, Spd (0.5, 1, 1.5 and 2 mM) and (0.25 + 0.25,0.5 + 0.5, Put + Spd1 + 1and 1.5 + 1.5 mM), results revealed that the (Put + Spd; 0.5 + 0.5 mM) treated plants showed a 98.2% increased germination rate than single Put and Spd plants upon NaCl stress condition. PAs treatments upon salinity stress revealed significant variations in the foxtail millet seedlings growth and development, such as fresh weight, dry weight of shoots and roots and length of shoot and roots (Table 1). In 200 mM NaCl stress, a severe reduction of the shoot and root biomass was observed, wherein NaCl (200 mM) stress condition along with Put, Spd and Put + Spd, treatments alleviated the salinity stress mediated growth reduction in both shoots and roots. Fresh and dry weight of shoots and roots showed that PAs treatment increased the biomass content in foxtail millet plants (Table 1).

Similarly, the height of the shoot and root length measurement signifies that Put (1 mM) and Spd (1 mM) and Put + Spd (0.5 + 0.5 mM) application enrich the growth of salinity exposed foxtail millet than the other concentrations of Put and Spd exposed plants. These results evidenced that Put + Spd treatment significantly enhance the plant biomass and also influence plant growth recovery that leads tolerance to high salinity stress in treated plants. The 200 mM NaCl along with Put (1 mM) and Spd (1 mM) and Put + Spd (0.5 + 0.5 mM) concentrations were chosen for further experiments.

Effects PAs on physiological parameters

The physiological changes in foxtail millet leaves were analyzed to understand the mechanism underlying tolerance to NaCl stress condition under PAs treatments. Foxtail millet plants exposed to high salinity stressed condition showed increased EL and caspase-like activity and decreased in RWC in shoots compared to NaCl with exogenous PAs and water control treatments (Fig. 2). The RWC in salinity stressed leaf samples was progressively increased, whereas the reduction in EL and caspase-like activity was noticed by the application of Put, Spd and Put + Spd. However, the level of EL was significantly diminished in the control plants which were exposed to high salinity stress alone without any PAs applications. In comparison with the control (unstressed) plants, the EL (63%) and caspase-like activity (79%) increased, but decreased RWC (24%) upon NaCl exposure. Furthermore, in PAs treated plants, the EL and caspase activity was reduced by 48% and 57% in Put + Spd, 39% and 43% in Spd whereas 30% and 36% in Put respectively, as compared with NaCl exposed plants (Fig. 2a-c). These results

PAs concentration (mM)		Growth parameters (Mean \pm SD)						
		FW (mg)		DW (mg)		Length (cm)		
		Shoot	Root	Shoot	Root	Shoot	Root	
Control		$94.3\pm0.7^{\rm i}$	$23.2\pm0.9^{\rm i}$	$9.4\pm0.6^{\mathrm{g}}$	$3.1\pm0.4^{\rm e}$	$12.5\pm0.8^{\rm h}$	5.4 ± 0.6^{ef}	
NaCl	200 mM	38.6 ± 0.6^a	$7.9 \pm 0.3^{\rm cd}$	1.6 ± 0.7^{a}	0.3 ± 0.1^{a}	$6.7\pm0.2^{\rm bc}$	$2.8 \pm 0.8^{\mathrm{abcd}}$	
NaCl + Put	0.5	$76.8 \pm 0.9^{\circ}$	$5.5\pm0.6^{\mathrm{b}}$	$2.4 \pm 0.2^{\mathrm{abc}}$	0.6 ± 0.3^{ab}	$7.9\pm0.4^{\rm cd}$	3.3 ± 0.7^{abcd}	
	1	$81.2 \pm 0.4^{\rm f}$	6.0 ± 0.5^{b}	$2.8 \pm 0.8^{\mathrm{abcd}}$	$0.9 \pm 0.6^{\mathrm{abc}}$	$8.4 \pm 1.0^{\text{def}}$	3.7 ± 0.5^{cd}	
	1.5	$79.3 \pm 1.0 d^{e}$	4.1 ± 0.2^{a}	2.2 ± 0.9^{ab}	$0.7 \pm 0.1^{\mathrm{abc}}$	6.8 ± 0.3^{bc}	$2.6 \pm 0.4^{\mathrm{abc}}$	
	2	$73.9 \pm 1.1^{\rm b}$	$3.8\pm0.8^{\rm a}$	1.7 ± 0.7^{ab}	0.5 ± 0.2^{ab}	6.2 ± 0.6^{ab}	$2.1\pm0.9^{\rm a}$	
NaCl + Spd	0.5	80.4 ± 0.5^{ef}	9.4 ± 0.7^{e}	3.1 ± 0.4^{bcde}	$1.1 \pm 0.8^{\mathrm{abc}}$	8.2 ± 0.7^{de}	3.5 ± 0.3^{bcd}	
	1	85.3 ± 0.6^{g}	$10.6 \pm 0.3^{\rm f}$	3.7 ± 0.8^{cdef}	1.4 ± 0.9^{abcd}	$9.3 \pm 0.5^{\rm ef}$	4.0 ± 0.2^{de}	
	1.5	$78.9\pm0.8^{\rm de}$	8.3 ± 0.5^{cde}	2.9 ± 0.7^{abcd}	1.3 ± 0.3^{abcd}	$6.1 \pm 0.9^{\mathrm{ab}}$	$2.8 \pm 0.4^{\mathrm{abcd}}$	
	2	$75.1\pm0.9^{\rm b}$	$7.2\pm0.8^{\rm c}$	2.6 ± 0.3^{abcd}	$0.9\pm0.2^{ m abc}$	$5.3\pm0.4^{\rm a}$	2.4 ± 0.7^{ab}	
NaCl + Put + Spd	0.25 + 0.25	$84.8 \pm 1.0^{\rm g}$	$11.4\pm0.6^{\rm f}$	$4.3\pm0.9^{\text{ef}}$	1.8 ± 0.5^{cd}	$9.5\pm0.8^{\rm f}$	$6.3\pm0.4^{\mathrm{g}}$	
	0.5 + 0.5	88.5 ± 1.1^{h}	14.9 ± 0.9^{h}	$5.1 \pm 1.0^{\rm f}$	2.2 ± 0.8^{de}	11.0 ± 0.6^{g}	6.6 ± 0.3^{g}	
	1 + 1	$81.6 \pm 1.2^{\rm f}$	$12.9\pm0.4^{\rm g}$	$4.8\pm0.9^{\rm f}$	1.5 ± 0.6^{bcd}	$10.9\pm0.7^{\rm g}$	$5.9\pm0.8^{\rm fg}$	
	1.5 + 1.5	$78.6\pm0.4^{\rm d}$	8.8 ± 0.2^{de}	$4.0 \pm 1.0^{\text{def}}$	$1.0 \pm 0.1^{\rm bc}$	$8.6 \pm 0.9^{\text{def}}$	$5.1 \pm 0.6^{\mathrm{ef}}$	

Table 1 Effect of exogenous Put, Spd and Put + Spd treatment on the growth parameters of 15 d old foxtail millet exposed to NaCl stress

Here, FW and DW represent fresh weight and dry weight of foxtail millet seedlings. Similarly, NaCl + Spd, NaCl + Put and NaCl + Put + Spd indicate spermidine (NaCl; 200 mM, Spd; 0.50 - 2.0 mM), putrescine (NaCl; 200 mM, Put; 0.50 - 2.0 mM) and putrescine + spermidine (NaCl; 200 mM, Put + Spd; 0.25 + 0.25, 0.50 + 0.50, 1.00 + 1.00 and 1.50 + 1.50 mM), respectively. Values are Mean \pm standard deviation (SD) of independent triplicates (n = 3). Different superscripted letters (a-i) within the column indicates statistically significant differences among the treatments according to Duncan's multiple range test ($p \le 0.05$). The optimum values for Put, Spd, Put+Spd are shown in bold for all the tested growth parameters



Fig. 2 Effect of PAs on the physiological parameters in foxtail millet with or without NaCl stress. A. EL (Electrolyte leakage; %) and B. Caspase-like activity (µmol H₂O₂ min⁻¹ g FW) C. RWC (Relative water content, %). Control (Unstressed), NaCl (200 mM), NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM), NaCl + Spd (NaCl; 200 mM, Spermidine; 1 mM) and NaCl + Put + Spd (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM). The data correspond to the average of three replicates. Values represent the Mean ± SD (analyzed by SPSS version 17.0), bars with different letters are significantly different at $p \le 0.05$ based on Duncan's multiple range test

firmly showed the regulatory roles of combined Put + Spd in high salinity stress.

PAs rescue the losses of chlorophyll, carotenoids and soluble protein

The synergistic effect of PAs on the photosynthetic pigments of foxtail millet under NaCl stress was determined in terms of total chlorophyll (a + b), carotenoids and soluble protein content. Salinity stress sharply increased the total chlorophyll (a + b) content, (54, 47 and 60%) (Fig. 3a, b) carotenoid content (50, 35 and 62%) and total soluble protein (15, 18 and 23%) in Spd, Put and Put + Spd supplements respectively, compared with the NaCl treated plants (Fig. 3c) (Fig. S2). The total chlorophyll (a + b), carotenoids and soluble protein contents were significantly reduced when the plants were exposed to NaCl alone (Fig. 3) (Fig. S2). The damaging effects of NaCl on the total chlorophyll (a + b), carotenoids and soluble protein contents were considerably declined by the exogenous application of combined PAs.



Fig. 3 Effect of PAs on the photosynthetic pigments in foxtail millet seedlings upon NaCl stress. A. Total chl $(a + b) \pmod{g^{-1}}$ FW), B. Estimation of Chl and C. Carotenoids $(\arg g^{-1}$ FW). Here, Control (Unstressed), NaCl (200 mM), NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM), NaCl + Spd (NaCl; 200 mM, Spermidine; 1 mM) and NaCl + Put + Spd (NaCl; 200 mM, Spermidine; 200 + 0.5 + 0.5 mM), respectively. The data correspond to the average of three replicates. Values represent the Mean \pm SD (analyzed by SPSS version 17.0), bars with different letters are significantly different at $p \le 0.05$ based on Duncan's multiple range test

Application of PAs inhibits the accumulation of proline

The role of PAs on osmoprotectant and ROS quencher was investigated by calculating the level of proline production under salinity stress. In this study, the proline contents were much higher in leaves of foxtail millet subjected to NaCl stress compared with the content in PAs treated and unstressed control plants. The addition of PAs decreased the proline accumulation by 22% in Put, 25% in Spd and 34% Put + Spd in comparison with NaCl treated plants, respectively (Fig. S3). These decreases were more significant in water control and NaCl with Put, Spd and Put + Spd treated plants. PAs applications are the most effective treatment, considerably controlling proline accumulation in salinity stressed foxtail millet plants. However, the combined Put + Spd application significantly regulates proline accumulation, which implies the synergistic effect of PAs on salinity stress.

PAs alleviates NaCl induced cell membrane damage

The damage in cell membrane was assessed by analyzing the MDA and H_2O_2 content which acts as prime indicators of oxidative stress. A drastic increase in the level of H_2O_2 and lipid peroxidation was noticed in NaCl (70 and 73%) compared with the control plants. The supplementation of Spd and Put reduced NaCl induced oxidative stress as evidenced by decreased levels, 44 and 45% in H_2O_2 and 44 and 39% in MDA respectively, when compared to the levels in plants treated only with NaCl stress (Fig. S4A, B). Moreover, co-application of Put + Spd to stressed plants resumed oxidative stress by increasing the level of H_2O_2 (52%) and MDA activity (58%) as compared with NaCl treated control. These results confirmed that Put + Spd increases salinity tolerance in foxtail millet plants by inhibiting the accumulation of toxic ROS contents.

Assays of antioxidant enzyme activity and free radical production

In order to explore the potential regulatory roles of PAs in alleviation of salinity induced oxidative stress, various antioxidant enzymes viz. SOD, CAT, APX, GR, GPX and GSH activities were estimated (Fig. 4). In the present study, NaCl treatment stimulated significant increases in the SOD (31%), CAT (70%), APX (71%), GR (76%), GPX (55%) and GSH (61%) respectively, and resulted in a greater accumulation of H₂O₂, \overline{O}_2 and MDA compared with the levels in unstressed control plants. This tendency is more obviously found in Spd than in Put, excluded for H₂O₂ content. SOD activity was considerably rising by 61, 56 and 69% in Spd, Put and Put + Spd treatments, whereas

CAT activity increase by 25% in Spd, 19% in Put and 35% in co-treatment of Put + Spd as compared with the NaCl treated plants, which reveals the decrease in ROS (H₂O₂ and \bar{O}_2) contents (Fig. 4a, b).

In comparison with NaCl treated control, APX and GPX activities raised (53 and 52%) and (62 and 57%) in NaCl + Spd and NaCl + Put treated plants. Meanwhile, the combined application of Put + Spd efficaciously increased the APX (63%) and GPX (72%) activities (Fig. 4c, d). The supplementation of PAs significantly reduced the activities of GR (39%) and GSH (50%) in NaCl + Put + Spd treatment when compared with the NaCl treatment (Fig. 4e, f). No remarkable changes in the level of antioxidant enzymes were noticed between the Put and Spd treated plants, however the synergistic effect of Put + Spd boost the antioxidant capability by enhancing the ROS scavenging mechanism in salinity stressed foxtail millet plants.

Assay of PAO and DAO activity

The generation of excess level of H_2O_2 through catalytic reaction was estimated by DAO and PAO, which play a key role in PAs homeostasis in plants. In the NaCl treated plants, DAO and PAO were severely increased by 62% and 81% when compared with the unstressed control plants. However, the DAO activity was gradually declined with the augmentation of Put (19%), Spd (22%) and Put + Spd (44%) treatment (Fig. S5A). Similarly, PAO activity was consistently decreased with 1 mM concentration of Put by 18%, Spd by 22% and Put + Spd (0.5 + 0.5 mM) registered a 29% decline as compared to salinity stressed plants (Fig. S5B). These results clearly specify the synergistic effect of Put + Spd enhancing salinity tolerance in foxtail millet.

Histochemical detection of H_2O_2 and \bar{O}_2

To elucidate whether the application of PAs in foxtail millet plants deals with ROS, detoxification induced by salinity stress were assessed by visualization of H_2O_2 and O $_2$ production using DAB and NBT staining. Foxtail millet plants were more extensively stained under NaCl stress condition than the controls (Fig. 5). The intensity of the DAB and NBT staining were considerably diminished in NaCl + Put and NaCl + Spd treated plants, respectively, signifying that PAs inhibited the accumulation of H_2O_2 and O_2 (Fig. 5a, b). Plants treated with NaCl + Put + Spd showed a minimal concentration of H_2O_2 and O_2 , as similar to the control plants (Fig. 5a, b). These results mimic the role of Put + Spd in controlling the ROS homeostasis in salinity treated plants.



B The second se





Fig. 4 Effect of exogenous PAs on antioxidant enzymes in NaCl affected foxtail millet seedlings. A. Superoxide dismutase (SOD; U mg⁻¹ leaf protein min⁻¹), B. Catalase (CAT; U mg⁻¹ leaf protein min⁻¹), C. Ascorbate Peroxidase (APX; U mg⁻¹ leaf protein min⁻¹), D. Guaiacol peroxidase (GPX; μ g⁻¹ leaf protein), E. Glutathione Reductase (GR; U mg⁻¹ protein) and F. Glutathione (GSH; mg g⁻¹ FW). Here, Control (Unstressed), NaCl (200 mM), NaCl + Spd

CLSM analysis

The cell membrane damage was assessed by examining ROS levels in the leaf cells. The fluorescence intensity ratio of NaCl stressed plants is significantly higher than the unstressed control and PAs treated plants. Salinity stress

(NaCl; 200 mM, Spermidine; 1 mM), NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM) and NaCl + Put + Spd (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM), respectively. The data correspond to the average of three replicates. Bars represent SD of the mean (n = 3). Different letters (a–e) indicate significant differences among the treatments at $p \le 0.05$, according to Duncan's multiple range test

generates increased ROS production within the leaf cells. The fluorescence intensity of the DCF-DA staining was considerably diminished in Put and Spd treated plants, respectively. Furthermore, the ROS production was profoundly reduced by the co-application of Put + Spd which was not statistically different from the unstressed control

plants (Fig. 6). These observations precisely show that a combined application of PAs (Put + Spd) considerably ameliorates salinity tolerance in foxtail millet plants.

FTIR analysis

Control

NaCl

The cell wall damage as well as protein damage is the key indicator of stress response during any stress conditions in the plants. In this study, the molecular changes during the NaCl exposure was observed using FTIR analysis by comparing the functional group intensity variations among the control and treated plants. The amide I absorption region acts as key indicator which is particularly sensitive to salinity stress. The absorption regions such as 3050 to 2800, 1750 to 1250, and 1250 to 900 cm^{-1} bands corresponding to lipids, amides and carbohydrates, respectively, were significantly reduced in the NaCl treated plants; whereas these regions gradually enriched in Put, Spd and Put + Spd treatment. The enriched intensities of the mentioned regions suggested that the cell wall and protein damage were successfully renovated during these Put, Spd, Put + Spd treatment upon NaCl exposure. The absorption range around 3537 cm⁻¹ indicates O-H and N-H stretching that predominantly found in chlorophyll protein and carbohydrates (Fig. 7). In FTIR analysis, the enriched (NaCl: (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM)intensities of these above mentioned regions clearly reveal

staining of $O_2^{\bullet-}$ (Left to right) Unstressed control, NaCl (200 mM), NaCl + Spd (NaCl; 200 mM, Spermidine; 1 mM), NaCl + Put 200 mM, Putrescine; 1 mM), NaCl + Put + Spd

that these two regions has been improved in Put, Spd and Put + Spd treatment, which stated that PAs enhances the salinity tolerance in foxtail millet plants.

Discussion

Salinity is responsible for multidimensional stresses such as ionic, osmotic, oxidative stress and hormonal imbalances in plants also it affect the photosynthetic machinery, lipid and protein metabolic process (Rasool et al. 2013). Salinity stress alters the genome, epigenome, transcriptome and proteome of the plants in the form of physiological, morphological, and biochemical dynamism. Plants have in turn evolved many tangled tolerance mechanisms to withstand the salinity stress. However, these tolerant mechanisms vary among diverse plant species.

The salinity stress greatly suppressed the GP, FW, DW and plant height. It also affected photosynthetic pigments, RWC and soluble protein, indicating that NaCl impaired the photosynthetic mechanism and deactivate the enzyme activities (Parida and Das 2005; Hameed and Ashraf 2008; Hussain et al. 2015; Satish et al. 2016). Reduction in plant biomass acted as a reliable indicator for evaluating the

Fig. 5 a Histochemical detection of H_2O_2 in salinity induced foxtail millet seedlings under PAs. DAB staining of H₂O₂ (Left to right) Here, Control (Unstressed), NaCl (200 mM), NaCl + Spd (NaCl; 200 mM, Spermidine; 1 mM), NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM), NaCl + Put + Spd (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM). **b** Effect of PAs on histochemical detection of



A



Fig. 6 Analysis of cell membrane damage through confocal laser scanning microscopy in foxtail millet leaves treated with PAs (Left to right). **a** Unstressed control, **b** NaCl – 200 mM, **c** NaCl + Spd

degree of damage and the tolerance capability (Sun et al. 2020). This study showed that RWC, total chlorophyll (a + b), carotenoid, soluble protein, plant growth and biomass were significantly alleviated with PAs treatments. The photosynthetic pigments content have been significantly reduced upon high salinity stress exposure, this result coincide with the findings of Dąbrowski et al. (2017). The enrichment of chlorophyll biosynthesis was alleviated by exogenous application of PAs. However, the synergistic effect of Put with Spd upon high salinity stress in foxtail millet plants has not been studied so far.

Accumulation of proline, protein and imbalance water contents are the general responses to salinity stress (Zhang et al. 2014). In this study, protein and RWC were relatively lower and proline accumulation was higher in salinity

(NaCl; 200 mM, Spermidine; 1 mM), **d** NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM) and **e** NaCl + Put + Spd (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM)

treated plants when compared with the control foxtail millet which is highly correlated with the severity of the cell damage. These observations support the findings of Abdel-Latef (2005) and Wasti et al. (2012) in wheat and Zhang et al. (2014) in tomato. The application of Put and Spd restored the RWC without much accumulation of proline content which is inversely proportional to the NaCl stressed plants. This may be due to enhancement of the pathway for synthesis of protein and proline from glutamine or transforming the other amino acids into proline (Kong-Ngern et al. 2005; Zhang et al. 2014). In plant cells, the level of ROS was reported to be increased under high salinity stress condition (Borsani et al. 2005; Chawla et al. 2013). Similarly, in our study, the foxtail millet plants treated with NaCl showed a drastic oxidative stress by



Fig. 7 Analyses the metabolic profile of PAs treated foxtail millet under salinity stress through FTIR spectroscopy. The data correspond to the average of three replicates. Here, control (Unstressed), NaCl (200 mM), NaCl + Spd (NaCl; 200 mM, Spermidine; 1 mM), NaCl + Put (NaCl; 200 mM, Putrescine; 1 mM), NaCl + Put + Spd (NaCl + Putrescine + Spermidine; 200 + 0.5 + 0.5 mM), respectively

higher levels of H_2O_2 and \overline{O}_2 . Salinity induced oxidative damage by membrane lipid peroxidation were correlated with a significant increase in MDA level. Additionally, this study revealed encouraging results in plant defence against the toxicity of NaCl upon Put + Spd co-application.

In the present investigation, FTIR has been used to analyse the functional group intensity variations under PAs treatment which is associated with salinity stress tolerance. The FTIR exposes the molecular-vibrational transitions which offer characteristic details on molecular configurations in biological samples (Griffiths 1978). The lipids, amides and carbohydrates absorption regions were significantly reduced upon salinity stress while PAs treatment enriched these functional groups. Previously, studies have proven the deconvolution and analysis of two absorption regions such as amide I (1580-1700 cm⁻¹) and pectin accumulation (1745 cm^{-1}) as key indicators of stress levels (Griffiths 1978; Barth 2000; Yang and Yen 2002). Moreover, the amide I absorption region is particularly sensitive to salt stress (Yang and Yen 2002). The absorption regions such as 3050 to 2800, 1750 to 1250, and 1250 to 900 cm^{-1} bands corresponding to lipids, amides and carbohydrates, respectively, were higher in normal plants (Lahlali et al. 2014). The enriched intensities of these regions revealed that the amide and lipid regions has been enhanced in PAs treatment, which specified the protective action of membrane lipid, cell wall pectin, chlorophyll protein and carbohydrates, than that of NaCl stressed plants.

Several studies have revealed that Put is the main precursor of PAs biosynthesis in plants, which can significantly ameliorate the salinity stress. PAs are important regulators to overcome abiotic stress in plants through influencing the antioxidant activities (Parvin et al. 2014; Tajti et al. 2018). Furthermore, PAs have a pivotal role in complex signaling, cell proliferation, interacting with macromolecules, endogenous phytohormone, metabolic profiles and have potential to influence in various plant defence mechanisms at enzymatic and gene expression levels (Liu et al. 2015; Tajti et al. 2018). The application of PAs significantly increases the H₂O₂-scavenging activity through antioxidant enzymes production in salinity stressed plants, resulting in decline of H₂O₂ level. These results are in agreement with the findings of Mittler et al. (2004). The rise in these ROS-scavengers may accelerate the reduction of toxic ROS by converting H_2O_2 into the water to decrease excess H_2O_2 content (Mittler et al. 2004). In this study, exogenous treatment of Put and Spd strongly induces the activation of antioxidant enzymes and PAs biosynthesis can confer salinity tolerance in foxtail millet plants. Numerous studies illustrated that PAs could ameliorate antioxidant ability in various plant species such as rice (Roychoudhury et al. 2011), ginseng (Parvin et al. 2014), wheat (Rady and Hemida 2015), zoysiagrass (Li et al. 2016), finger millet (Satish et al. 2018), and cucumber (Duan et al. 2008; Wu et al. 2018). Our observations revealed that the synergistic effect of combined Put + Spdtreatment enhances salinity tolerance in foxtail millet with their efficient elimination of ROS by scavenging system. Several reports (Nayyar and Chander 2004; Gong et al. 2016; Nahar et al. 2016; Paul et al. 2018; Chen et al. 2018) including this study, proved the regulatory roles of PAs on antioxidant defence, however PAs based ROS detoxification mechanism has still remained to be detected in plants. Based upon the above information, a schematic representation was portrayed in Fig. S6 which elucidates the biochemical mechanisms involving NaCl induced toxicity and the exogenous PAs (Put + Spd) mediated salinity stress tolerance in foxtail millet plants.

Conclusion

This is the first report demonstrating the salinity tolerance mechanism of combined PAs such as Put along with Spd in foxtail millet. The morphological, physiological and biochemical responses upon Put + Spd (0.5 + 0.5 mM) treatment showed higher adaptive mechanism than individually (1 mM Spd and 1 mM Put) treated plants. Further, the synergistic effect of Put + Spd confers salinity tolerance through inducing antioxidant enzymes and osmoprotectants. Overall, the exposure of combined PAs under NaCl stress coordinates the complex physiological and biochemical processes. Hence, the exogenous

co-application of lower concentration of Put + Spd can be easily adopted to ameliorate the salinity stress in foxtail millet.

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Author Contributions Conceived and designed the experiments: PR & MR. Performed the experiments: PR MB KR RA RR. Analyzed the data: PR LS SP. Contributed reagents/materials/analysis tools: MR. Wrote the paper: PR. All the authors have read the manuscript and approved for publication.

Compliance with ethical standards

Conflict of interest The authors don't have any conflict of interest to declare.

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