

Synergistic Interactions Between Salt-tolerant Rhizobia and Arbuscular Mycorrhizal Fungi on Salinity Tolerance of *Sesbania cannabina* Plants

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Received: 17 February 2016 / Accepted: 8 April 2016 / Published online: 28 May 2016
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Abstract Legumes can host rhizobia and mycorrhizal fungi, and this triple symbiosis might be exploited to improve saline soil fertility. Therefore, a greater understanding of the interaction of rhizobia and arbuscular mycorrhizal fungus during legume growth in saline soil is required. We investigated the efficiency of salt tolerance conferred by rhizobia in mycorrhizal *Sesbania cannabina*. Greenhouse experiments were conducted in which *S. cannabina* plants inoculated with *Glomus mosseae* BGC NM03D (GM), and two rhizobia strains *Agrobacterium pusense* YIC4105 (4105) and *Neorhizobium huautlense* YIC4083 (4083), were exposed to 100 and 200 mM NaCl. Under 200 mM NaCl stress, plants inoculated with 4105, rather than 4083, showed significant increases in shoot and root dry mass compared with non-inoculated plants. Simultaneously, a significant increase over GM-inoculated plants in mycorrhizal colonization and dependency was recorded for 4105 + GM-inoculated plants compared with 4083 + GM-inoculated plants. In addition, under NaCl stress, significant increases in the number and mass of nodules, nitrogenase activity, and leghemoglobin content of nodules occurred in 4105 + GM-inoculated plants compared with 4083 + GM-inoculated plants. Furthermore, the activities of antioxidant enzymes in rhizobia-

inoculated plants were significantly higher in the GM + 4105 group than the 4083 + GM group. The malondialdehyde content of plants from the 4105 + GM group was significantly lower than in the 4083 + GM group. Thus, the results revealed a synergistic relationship among the 4105 and GM in alleviating salt stress in *S. cannabina*. Salt-tolerant rhizobia might improve the salinity tolerance of *S. cannabina* by enhancing the antioxidant system.

Keywords Rhizobia · Arbuscular mycorrhizal fungi · *Sesbania cannabina* · Salt tolerance · Antioxidant enzyme

Introduction

Soil salinity is one of the most widespread threats to agricultural productivity in arid and semiarid areas (Ghazi and Al-Karaki 2006). Currently used techniques, such as improved irrigation practices, changing the soil, and using salt-tolerant plants for farming in salt-affected soils, are either expensive or time consuming (Munns 2002). However, using bioprocesses such as mycorrhizae and root nodules may represent a relatively cost-effective and ecologically friendly method to improve saline soils.

Arbuscular mycorrhizal (AM) fungi exist widely in saline soils, and inoculation with AM fungi improves growth of almost every terrestrial plant under saline stress conditions (Al-Karaki and others 2001; Heikham and others 2009). These fungi have been considered bio-ameliorators in a variety of saline soils (Singh and others 1997; Rao 1998). Rhizobial bacteria can form an exclusive symbiotic relationship with legumes by fixing atmospheric nitrogen to supply the plant with ammonium. They contribute not only to the N sources of leguminous crops, but

Electronic supplementary material The online version of this article (doi:10.1007/s00344-016-9607-0) contains supplementary material, which is available to authorized users.

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also to the N content of the soil, and therefore have a key role in environmentally friendly agricultural practices (Puppo and others 2005). Legumes can grow under nitrogen-limiting conditions because of their ability to establish endosymbiosis with rhizobia. *Sesbania cannabina* is a recognized soil-improving legume, used as green manure to increase the yield of many crops. It is widely adaptable to different adverse climatic conditions, such as soil salinity, drought, and waterlogging. Therefore, rhizobia–*S. cannabina* plant–AM fungi tripartite symbiosis might represent a good strategy for improving saline soil fertility and helping to reintroduce agriculture to these lands.

Many pieces of evidence indicate that rhizobia and AM fungi improve the salinity tolerance of legumes (Rosa and others 2012; Carmen and Roberto 2009). However, studies on how the salt tolerance of these microbe symbionts affects the host plant are limited. Under long-term saline stress, some AM fungi species survive by ecophysiological adaptation (Copeman and others 1996; Weissenhorn and others 1993; Camprubi and Calvet 1996; del Val and others 1999). It is plausible that the AM fungi that are able to survive in saline soils should be considered tolerant species and might be better able to improve the survival and growth of host plants than species from non-saline conditions. However, Tian and others (2004) reported that AM fungi from saline soils do not have a higher capacity to alleviate saline stress in plants than fungi from non-saline soils.

To the best of our knowledge, there have been no reports demonstrating whether the salt tolerance of rhizobia contributes to the survival of plants under salt stress with AM fungi. Any relationship between salt-tolerant rhizobia and AM fungi also remains to be determined. In this study, we inoculated *S. cannabina* plants with a combination of *Glomus mosseae* and two species of rhizobia with significant differences in their salt tolerance to explore this issue.

Materials and Methods

Preparation of Inocula

Rhizobia strains were isolated from *Sesbania cannabina* plants collected from DongYing, ShanDong province (37.76°N, 118.98°E). The isolates were grown on yeast extract mannitol agar (YEMA) medium and incubated at 28°C (Vincent 1970). DNA was extracted from bacterial cultures using the sodium dodecyl sulfate/cetyl trimethylammonium bromide lysis and phenol/chloroform extraction method. The housekeeping gene *recA* was amplified using the following primers:

recA-41F:TTC GGC AAG GGM TCG RTS ATG
recA-640R:ACA TSA CRC CGA TCT TCA TGC

After the approximately 600-bp *recA* amplicons were clustered and compared using MEGA5.05 program, 18 representative genotypes were selected. These isolates were designated as YIC4009, YIC4027, YIC4031, YIC4032, YIC4056, YIC4071, YIC4072, YIC4083, YIC4103, YIC4104, YIC4105, YIC4108, YIC4121, YIC4260, YIC4261, YIC5077, YIC5079, and YIC5082 (Fig. S1).

Then, the approximately 1500-bp 16S rDNA genes were amplified from the 18 representative strains using the following primers:

16S-27F: AGA GTT TGA TCC TGG CTC AG
 16S-1492R: AAG GAG GTG ATT CCA GCC

The sequences were edited and assembled using BioEdit 7.0 (<http://www.mbio.ncsu.edu/BioEdit/bioedit.html>). BLAST searches were performed using the NCBI server (www.ncbi.nlm.nih.gov/blast/Blast.cgi). A phylogenetic tree was constructed from the 16S rRNA gene sequences in comparison with 16SrRNA gene sequences from several standard bacterial strains (Figure S1). Two genospecies, characterized as *Agrobacterium pusense* YIC4105 (4105) and *Neorhizobium huautlense* YIC4083 (4083), were selected based on their salt-tolerance traits and capacity to promote *S. cannabina* growth.

The original inoculum of the AM fungi *Glomus mosseae* (GM, BGC NM03D) provided by the Institute of Plant Nutrition and Resources (Beijing, China) was propagated in pot culture on *Trifolium repens* for 8 weeks. The spore intensity was 116 spores per 10 ml inoculum.

Experimental Design and Biological Treatments

The experiment had a 6 × 3 factorial design which was composed of six inoculation treatments and three salinity levels. Six inoculation treatments were non-inoculation control (control), inoculations with AM fungi, and rhizobia individually and in combination (GM, 4083, 4105, GM + 4083, and GM + 4105). Three salinity levels were 0, 100, and 200 mM NaCl. Treatments were completely randomized and replicated three times. The NaCl solution was added to the medium at a rate of 100 ml/7 days, five times in total. Seeds of *Sesbania cannabina* (Retz.) Pers. (Shandong Academy of Agricultural Sciences, Shandong, China) were sterilized in 10 % hydrogen peroxide for 10 min and rinsed several times with distilled water before use. Ten seeds were germinated at 28 °C for 48 h and sown (thinned to three uniform seedlings after germination) in a single 2-L pot containing autoclaved zonalite in a greenhouse with day/night temperatures of 30/22 °C, 60 ± 2 % relative humidity, and a photoperiod of 14/10 h light/dark. AM inoculums and rhizobia were applied as the seeds were sown. 1/4 Hoagland solution (Hoagland 1950) was supplied regularly and the pots were weighed every week to

adjust the water content. After five weeks, plants were harvested for analysis.

Mycorrhizal Colonization and Dependency

The percentage of mycorrhizal colonization in the roots was calculated by the gridline intersection method (Giovannetti and Mosse 1980), after staining with trypan blue (Phillips and Hayman 1970).

Mycorrhizal colonization (%)

$$= \frac{\text{(Number of root segments colonized)}}{\text{(Number of root segments studied)}} \times 100$$

According to Plenchette and others (1983), the mycorrhizal dependency was calculated as follows:

$$\text{Dependency of growth} = \frac{\text{(Dry weight}_M - \text{Dry weight}_{NM})}{\text{Dry weight}_M} \times 100$$

where 'M' represents the mycorrhizal plants and 'NM' the non-mycorrhizal plants.

Nitrogenase Activity and Leghemoglobin Content of Nodules

The excised nodulated roots were placed in 100-ml bottles sealed with rubber plugs. Ten ml of air was taken out and the equivalent amount of acetylene gas was injected into the bottle, which was incubated at 28 °C for 4 h. Gas samples (100 µl) from the bottles were then injected into a gas chromatograph (7890A series GC, Hewlett-Packard, Palo Alto, CA, USA) equipped with a flame ionization detector and an HP-PLOT AL2O3/KCL column (30 m, 0.25 mm, 5 µm; Agilent, Santa Clara, CA, USA) was used. A calibration curve was constructed using pure ethylene. After the nitrogenase activity was determined, the nodules of each root were counted and measured for fresh and dry mass.

One gram of fresh nodules was ground with 5 ml of distilled water. The homogenate was centrifuged at 12,000×g for 15 min. Leghemoglobin in the supernatant was determined by the method of Hartree (1957), which is based upon the conversion of hematin to pyridine hemochromogen. A standard curve was prepared using graded concentrations of hemin. Unless stated otherwise, all reagents used in present study were purchased from Sigma-Aldrich (St. Louis, MO, USA).

Superoxide Dismutase Activity

The assay is based on the formation of formazone, which absorbs at 560 nm, by nitroblue tetrazolium and O₂⁻ radical. Superoxide dismutase (SOD; E.C.1.15.1.1) decreases this absorbance by reducing the formation of the O₂⁻

radical (Dhindsa and others 1981). Leaves from *S. canabina* frozen in liquid nitrogen were ground with phosphate buffer, centrifuged at 4 °C, and the supernatant was used as the enzyme sample. The reaction mixture contained 14.5 mM methionine, 100 µM nitroblue tetrazolium chloride, 0.1 mM EDTA-Na₂, 50 mM phosphate buffer (pH 7.8), 50 mM sodium carbonate, and 50 µl of enzyme. The reaction was started by adding 2.25 µM riboflavin and illuminating. A complete reaction mixture without enzyme served as the control. Switching off the light stopped the reaction. A non-irradiated complete reaction mixture served as the blank. One unit of enzyme activity was taken as the amount of enzyme that reduced the absorbance reading by 50 % in comparison with the control.

Catalase Activity

The catalase activity (CAT; E.C. 1.11.1.6) assay is based on a decrease in the absorbance of H₂O₂ at 240 nm over a time period, according to Aebi (1984). The reaction mixture consisted of 50 mM (pH 7.0) phosphate buffer, 12.5 mM hydrogen peroxide, and 50 µl of enzyme. Adding H₂O₂ started the reaction. Enzyme activity was computed by calculating the amount of H₂O₂ decomposed in 1 min. A standard curve was prepared using graded concentrations of H₂O₂. The enzyme activity was represented as the H₂O₂ reduced per min per mg protein.

Peroxidase Activity

The peroxidase activity (POX; E.C. 1.11.1.7) assay is based on the increase in optical density caused by the oxidation of guaiacol to tetraguaiacol (Castillo and others 1984). The reaction mixture contained 8 mM guaiacol, 5 mM H₂O₂, 100 mM phosphate buffer (pH 6.4), and 150 µl of enzyme extract. Absorbance from the formation of tetraguaiacol was recorded at 470 nm; the extinction coefficient ε was 26.6 mM⁻¹ cm⁻¹. The enzyme activity was represented as µmol tetraguaiacol formed per min per g fresh weight (FW).

Ascorbate Peroxidase Activity

The ascorbate peroxidase (APOX; E.C. 1.11.1.11) activity assay is based on the decrease in absorbance of ascorbic acid at 290 nm resulting from its oxidation (Nakano and Asada 1981). The enzyme sample was prepared with 1 mM ascorbic acid in addition to the extraction buffer. The reaction mixture contained 50 mM phosphate buffer (pH 7.0), 0.5 mM ascorbic acid, 0.1 mM EDTA-Na₂, 0.2 mM H₂O₂, and 100 µl of enzyme. The reaction was started with the addition of 0.2 ml H₂O₂. The decrease in absorbance at 290 nm for 30 s was measured. A standard curve was

prepared using graded concentrations of ascorbic acid. The enzyme activity was represented as the concentration of ascorbic acid oxidized per min per mg protein.

Glutathione Reductase Activity

The glutathione reductase (GR; E.C. 1.11.1.9) activity assay is based on the formation of a colored complex that absorbs at 412 nm by the reduction of glutathione with 5, 5-dithiobis-2-nitrobenzoic acid (DTNB) (Smith and others 1988). The reaction mixture contained 50 mM phosphate buffer (pH 7.5) and 0.1 mM EDTA, 0.5 mM DTNB, 0.5 mM NADPH, 5 mM GSSG (oxidized glutathione), and 100 μ l of enzyme. The reaction was started by adding 2 mmol GSSG. The activity was expressed as total absorbance at 412 nm (Abs. 412) per mg protein per min.

Determination of Malondialdehyde Content

Lipid peroxidation was determined by measuring malondialdehyde (MDA) formation, using the thiobarbituric acid reaction described by Sudhakar and others (2001). The colored complex absorbs at 532 nm. Leaves from *Sesbania cannabina* were homogenized in 0.1 % trichloroacetic acid. The homogenate was centrifuged at 13,000 \times g for 20 min. The supernatant was used to estimate the MDA content. Thiobarbituric acid (TBA) in 20 % TCA was added to the supernatant. The mixture was heated at 95 $^{\circ}$ C for 20 min and then cooled in an ice bath. After centrifugation at 10,000 \times g for 10 min, the absorbance of the supernatant was recorded. Non-specific absorption values recorded at 600 nm were subtracted from the values recorded at 532 nm. The MDA content was calculated according to its extinction coefficient, ϵ (155 mM $^{-1}$ cm $^{-1}$).

Statistical Analyses

Data were compiled using Microsoft Excel (Redmond, WA, USA). Values are represented as means \pm SDs of three replicates for each treatment. Student's *t* test, one-way ANOVA, and Duncan's multiple range test were used to identify significant differences, using SPSS ver. 13.0 (SPSS Inc., Chicago, IL, USA).

Results

Effects on Plant Growth

Two rhizobia, 4105 and 4083, with significant differences in salt tolerance, were isolated and selected for inoculation (Fig. 1). These two strains had similar capacities to promote *S. cannabina* growth: the increases in shoot and root

dry mass of 4105- and 4083-inoculated plants were not significantly in 0 mM NaCl (Table 1).

As shown in Table 1, the inhibitory effect of salinity on shoot and root dry mass accumulation of *S. cannabina* increased as the NaCl concentration increased. NaCl at 100 and 200 mM induced a 25.3 and 51.8 % decline in dry matter accumulation of *S. cannabina* plants, respectively, compared with plants grown in 0 mM NaCl. In the single inoculation group, 4105 and GM, but not 4083, induced a clear improvement in dry mass accumulation of *S. cannabina* plants under 200 mM NaCl treatment, compared with non-inoculated plants. In the dual inoculation group, 4105 + GM was more efficient than 4083 + GM in increasing plant growth under salinity stress. Under 100 and 200 mM NaCl, the increase of 4105 + GM plants over non-inoculated plants was 2.54 and 2.38 times, whereas the increases of 4083 + GM plants were 1.93 and 1.71 times, respectively, (Table 1).

Notably, GM promoted stronger increases in root dry mass compared with shoot dry mass. This effect also seemed to occur in dual-inoculated plants (Table 1).

Mycorrhizal Colonization and Dependency

The results of mycorrhizal colonization showed that different levels of NaCl did not affect the colonization ability of GM significantly. The mycorrhizal infection of *S. cannabina* roots observed under different saline concentrations was not significantly different (Fig. 2a). Moreover, the colonization rate was even higher when the mycorrhizal plants were inoculated with 4105 compared with 4083 under 100 and 200 mM NaCl (Fig. 2a).

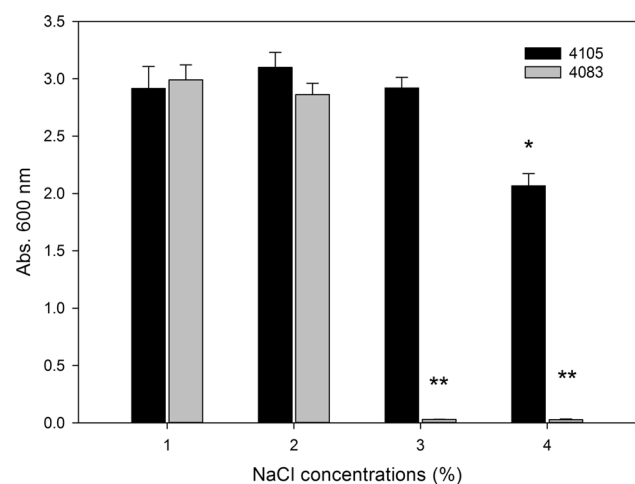


Fig. 1 Optical density at 600 nm of *Rhizobium pusense* YIC4105 and *Rhizobium huaatlense* YIC4083 in tryptone-yeast media with different NaCl levels after 48 h. Bars with different letters indicate significant differences (Student's *t* test, *, $P < 0.05$; **, $P < 0.01$)

Table 1 Dry weight of shoots and roots of *Sesbania cannabina* plants inoculated with rhizobium and *Glomus mosseae* grown under different levels of NaCl

NaCl (mM)	Dry weight (g)					
	Control	4083	4105	GM	GM + 4083	GM + 4105
0						
Shoot	5.83 ± 0.49a	13.02 ± 2.11b	12.73 ± 0.82b	7.09 ± 1.13a	13.86 ± 0.58b	13.36 ± 0.61b
Root	7.14 ± 1.14a	11.89 ± 0.57c	11.18 ± 1.05c	8.83 ± 0.49b	12.18 ± 0.54c	12.2 ± 1.06c
100						
Shoot	5.13 ± 0.39a	7.75 ± 1.03bc	8.68 ± 0.60c	6.87 ± 1.06b	9.06 ± 0.72c	12.42 ± 0.73d
Root	4.56 ± 0.37a	7.39 ± 0.62b	8.35 ± 0.51b	8.4 ± 1.02b	9.68 ± 0.90c	12.18 ± 0.65d
200						
Shoot	2.9 ± 0.87a	3.81 ± 0.14ab	4.58 ± 0.37bc	5.45 ± 0.64 cd	5.8 ± 0.74d	7.56 ± 0.27e
Root	3.62 ± 0.09a	3.44 ± 0.03a	4.84 ± 0.45b	6.17 ± 0.85c	5.35 ± 0.61bc	7.97 ± 0.39d

Means with different letters among inoculations within an NaCl level are significantly different (one-way ANOVA, Duncan's multiple range test, $P < 0.05$)

The mycorrhizal dependency was calculated in mycorrhizal plants with or without rhizobial inoculation. It is clear that the maximum mycorrhizal dependency of dry mass was obtained at the higher NaCl level of 200 mM. However, the mycorrhizal dependency of 4083-inoculated plants was not significantly different from GM-inoculated plants, while the mycorrhizal dependency of 4105-inoculated plants was 39 % higher than mycorrhizal plants under NaCl stress of 200 mM (Fig. 2b). The results above indicated that 4105 had greater capacity than 4083 to promote *S. cannabina* growth under salt stress in a single inoculation and in a double inoculation with GM.

Effect of Arbuscular Mycorrhizal Fungi on Nodulation and Nitrogen Fixation

To study the effect of AM fungi on rhizobia in dual-inoculated *S. cannabina*, nodulation and nitrogen fixation were investigated.

Data presented in Table 2 revealed that the number and mass of nodules significantly decreased under treatment with 200 mM NaCl in rhizobia-inoculated and dual-inoculated plants. Interestingly, dual inoculation of 4105 and GM increased the average number of nodules by 11.33 compared with the 4105 plants, whereas dual inoculation with 4083 and GM increased the average number of nodules by 5 compared with the 4083 plants. Total nodule weight per plant followed a similar pattern to the number of nodules.

Exposing the plants to NaCl stress of 100 and 200 mM resulted in a sharp reduction in the leghemoglobin content in rhizobia-inoculated plants. However, significantly less damage to the leghemoglobin protein was recorded in 4105 + GM-inoculated plants under stress compared with

4105 + GM- and 4105-inoculated plants. Nodule activity, according to the ethylene transformation from acetylene, corresponded to the reduction in nodule dry mass and leghemoglobin content under salt stress (Table 2). The efficiency of *S. cannabina*–4105 symbiosis increased significantly and the symbiotic performance was better in the 4105-inoculated than the 4083-inoculated plants under salt stress (Table 2).

Antioxidant Enzyme Activities

Salt stress could lead to the excessive production of reactive oxygen species (ROS), such as superoxide O_2^- radicals and hydrogen peroxide H_2O_2 , resulting in nodule senescence. Antioxidant enzymes, such as SOD, CAT, and POX, play key roles in the detoxification of ROS (Moran and others 2003). ROS-generating and activated antioxidant systems have been found not only at biotic/abiotic stress sites but also in distal leaf, acting as systemic signal (Freeman 2003; Martha and Clarence 1999). To exclude the ROS interference in AM fungal tissues, shoots instead of roots were used for measurements of antioxidant enzymes activities. The results showed that SOD activity (Fig. 3a) was elevated in response to moderate (100 mM) and heavy (200 mM) salt stress in 4105 + GM-inoculated plants. NaCl levels of 100 and 200 mM increased the SOD activity by 1.52 and 3.99 times, respectively, in 4105 + GM-inoculated plants compared with the corresponding plants at 0 mM. Higher SOD activity (1.93 times) was also observed in 4105 plants at 100 mM, but not at 200 mM NaCl, compared with non-stressed plants (Fig. 3a). The SOD activity of 4105 + GM plants was significantly higher than that in 4083 + GM plants. Increased levels of CAT and POX (Fig. 3b, c) activities

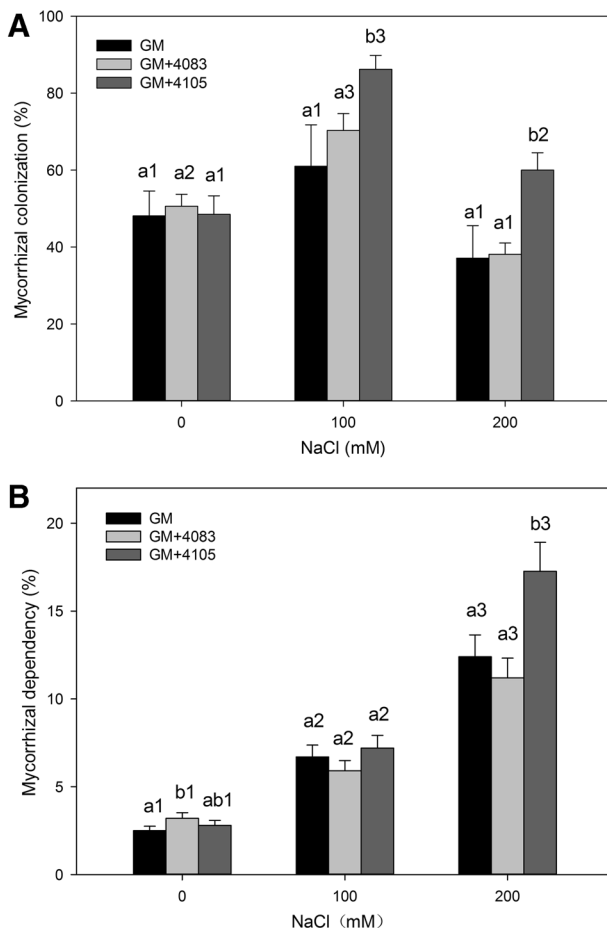


Fig. 2 *Glomus mosseae* mycorrhizal colonization (%) and mycorrhizal dependency (%) of *Sesbania cannabina* plants inoculated with rhizobium under NaCl stress. **a** Percentage of *Glomus mosseae* colonization of the roots of *Sesbania cannabina* plants inoculated with *Rhizobium pusense* YIC4105 and *Rhizobium huautlense* YIC4083. **b** Mycorrhizal dependency (%) of *Sesbania cannabina* plants inoculated with *Rhizobium pusense* YIC4105 and *Rhizobium huautlense* YIC4083. Bars with different letters among the inoculations within an NaCl level are significantly different, and with those different numbers among salinity levels within the same inoculation are significantly different (one-way ANOVA, Duncan’s multiple range test, $P < 0.05$)

were also observed in 4105 + GM-inoculated plants compared with non-inoculated plants subjected to moderate to heavy salt stress. The presence of 4105 and GM in the plant root increased CAT activity by 1.35 and 3.24 times at 100 and 200 mM, respectively, relative to non-stressed controls. The POX activity of stressed 4105 + GM-inoculated plants was 1.23 and 2.07 times higher than the corresponding non-stressed controls, at 100 and 200 mM, respectively. The dual inoculations of 4083 and GM also increased CAT and POX activity under saline stress, but not to as high a level as 4105 + GM-inoculation. SOD and POX activities of 4105 + GM plants were significantly higher than those in 4083 + GM plants (Fig. 3b, c).

Glutathione ascorbate peroxidase (APOX) and reductase (GR), which are enzymes involved in the ascorbate–glutathione pathway that is responsible for the removal of H_2O_2 , were also investigated. No significant increase in APOX activity was recorded in single inoculation of rhizobia or in dual inoculations of rhizobia and arbuscular mycorrhizal fungi (Fig. 3d). An increase in GR activity was recorded only in 4083 and 4105 + GM-inoculated plants under heavy salt stress (Fig. 3e). However, the difference between the APOX and GR activities in 4105 + GM and 4083 + GM-inoculated plants was not significant.

Lipid Peroxidation

Salt stress-induced accumulation of ROS leads to oxidation damage of cell components, such as peroxidation of polyunsaturated fatty acids in membranes via free radical reactions. The results shown in Fig. 4 revealed that only the MDA content of 4105 + GM-inoculated plants was not significantly increased compared with corresponding plants under 0 mM NaCl treatment. This was consistent with the data from the antioxidant enzyme assays.

Discussion

In stressed edaphic environments, many plants establish a symbiosis with soil microorganisms, including AM fungi or rhizobium in legumes. Both symbionts improve plant growth under severe environmental conditions (Mohamed and others 2014). In fact, it is common that dual symbiosis of rhizobium and AM fungi enhances the growth and yield of many legumes (Zahran 1999; Kazunori and others 2013). In the present study, rhizobial bacteria 4083 promoted the growth of *S. cannabina* under moderate salt stress, whereas 4105 showed a clear improvement under moderate and severe salt stress. The AM fungus GM had a similar growth-promoting effect to 4105 (Table 1). The plant growth response to dual symbiosis is influenced by microbial strains, as well as by the compatibility of the interactions among them (Azcón and others 1991). In AM fungi and rhizobium dual-inoculated plants, 4105 + GM was more efficient than 4083 + GM in increasing plant growth under salinity stress. In addition, the dry mass of dual-inoculated plants was higher than in the corresponding rhizobial-inoculated plants (Table 1).

There is ample evidence of a positive effect of dual inoculation with different rhizobium strains and AM fungi on legume growth (Aryal and others 2003; Mortimer and others 2008). Here, we examined whether this positive effect also occurred under saline stress. Interestingly, the colonization rate was even higher when the mycorrhizal

Table 2 Nodulation, nitrogenase activity, leghemoglobin content of *Sesbania cannabina* plants inoculated with rhizobium and *Glomus mosseae* grown under different levels of NaCl

Inoculation	NaCl(mM)	Nodule/plant			Nitrogenase activity (moles C ₂ H ₄ g ⁻¹ FW nodules h ⁻¹)	Leghemoglobin content (mg g ⁻¹ FW nodules)
		Number	Fresh weight (g)	Dry weight (g)		
4083	0	47.33 ± 2.08a	1.23 ± 0.07ab	0.26 ± 0.02a	2.32 ± 0.15a	1.33 ± 0.14a
	100	52.67 ± 2.52b	1.22 ± 0.05ab	0.27 ± 0.01ab	1.88 ± 0.11ab	0.81 ± 0.17a
	200	2.67 ± 1.53a	0.14 ± 0.06a	0.04 ± 0.02a	0.85 ± 0.13a	0.32 ± 0.12a
4105	0	44.00 ± 4.58a	1.11 ± 0.06a	0.26 ± 0.01a	2.17 ± 0.16a	1.43 ± 0.15a
	100	43.67 ± 2.52a	1.17 ± 0.05a	0.26 ± 0.02a	1.91 ± 0.10ab	0.83 ± 0.11a
	200	9.00 ± 2.00b	0.26 ± 0.07a	0.06 ± 0.02a	0.94 ± 0.11a	0.43 ± 0.11a
GM + 4083	0	43.33 ± 3.51a	1.27 ± 0.04b	0.29 ± 0.02a	2.37 ± 0.19a	1.42 ± 0.15a
	100	55.33 ± 4.04b	1.30 ± 0.06bc	0.29 ± 0.01bc	1.75 ± 0.12a	0.87 ± 0.13a
	200	7.67 ± 1.53b	0.21 ± 0.09a	0.06 ± 0.03a	1.01 ± 0.08a	0.41 ± 0.12a
GM + 4105	0	42.00 ± 7.00a	1.14 ± 0.09a	0.26 ± 0.02a	2.23 ± 0.15a	1.41 ± 0.12a
	100	56.00 ± 2.65b	1.37 ± 0.07c	0.30 ± 0.02c	2.08 ± 0.09b	0.94 ± 0.07a
	200	20.33 ± 3.79c	0.89 ± 0.10b	0.20 ± 0.02b	1.33 ± 0.12b	0.77 ± 0.06b

Means with different letters among inoculations within a NaCl level are significantly different (one-way ANOVA, Duncan's multiple range test, $P < 0.05$)

plants inoculated with 4105 were treated with 200 mM NaCl (Fig. 2a). The mycorrhizal dependency of 4105-inoculated plants was 39 % higher than mycorrhizal plants under 200 mM NaCl. However, the mycorrhizal dependency of 4083-inoculated plants was not significantly different from GM-inoculated plants (Fig. 2b).

Unlike rhizobia, legume-rhizobium symbiosis and nodule formation on legumes are more sensitive to salt (Zahran 1991). Salt stress not only inhibits the initial steps of nodule formation, but also affects nitrogen fixation of legumes (Delgado and others 1994; Nair and others 1993; Salwa and others 2005). The reduction of N₂-fixing activity by salt stress is usually attributed to a reduction in respiration of the nodules (Walsh 1995) and a reduction in leghemoglobin production by nodules (Delgado and others 1994). *Agrobacterium pusense* YIC4105 and *Neorhizobium huautlense* YIC4083 used in this work could tolerate 4 and 2 % NaCl in tryptone-yeast media, respectively. However, the number, mass of nodules, leghemoglobin content, and nitrogenase activity were all significantly suppressed by 200 mM NaCl treatments in rhizobia-inoculated plants. In addition to intrinsic protective systems of plants against saline stress, arbuscular mycorrhizal symbiosis can alleviate saline-induced nodule senescence in legume plants (Porcel and others 2003). The present study showed that dual inoculation of 4105 + GM increased the number and weight of nodules to a greater extent than in 4083 + GM-inoculated plants. In addition, significantly lower damage to the leghemoglobin content and N₂-fixing efficiency was recorded in 4105 + GM-inoculated plants under stress

compared with 4083 + GM as well as 4105-inoculated plants (Table 2). Notably, although the leghemoglobin content and nitrogenase activity of all rhizobia-inoculated plants declined under saline stress, the nodule number and mass did not; the nodule number of 4083-inoculated plants and the fresh weight of 4105-inoculated plants even increased significantly under 100 mM NaCl (Table 2). The data revealed that the rhizobium actively infected the roots and developed nodules; however, saline stress prevented the nodules from enlarging or activating symbiosis. Similar results have been observed in previous studies: Anthraper and DuBios (2003) reported that although salt-stressed plants produced more nodules than the controls, most of the nodules were small or inactive. In addition, an increase in the average nodule weight with increasing salinity level was observed in chickpeas (Garg and Singla 2004) and faba beans (Cordovilla and others 1999).

As a consequence of salinity stress, the accumulation of ROS can cause oxidative damage to membrane lipids, proteins, and nucleic acids, eventually leading to programmed cell death (Gomez and others 1999; Hernandez and Almansa 2002). Increased antioxidative enzyme activities could be involved in the beneficial effects of mycorrhizal colonization on the performance of plants grown under semiarid conditions (Alguacil and others 2003). Our results revealed that SOD activity increased significantly in 4105 and 4105 + GM plants compared with 4083 and 4083 + GM plants under salt stress. This induction of SOD activity was consistent with changes in hydrogen peroxide-

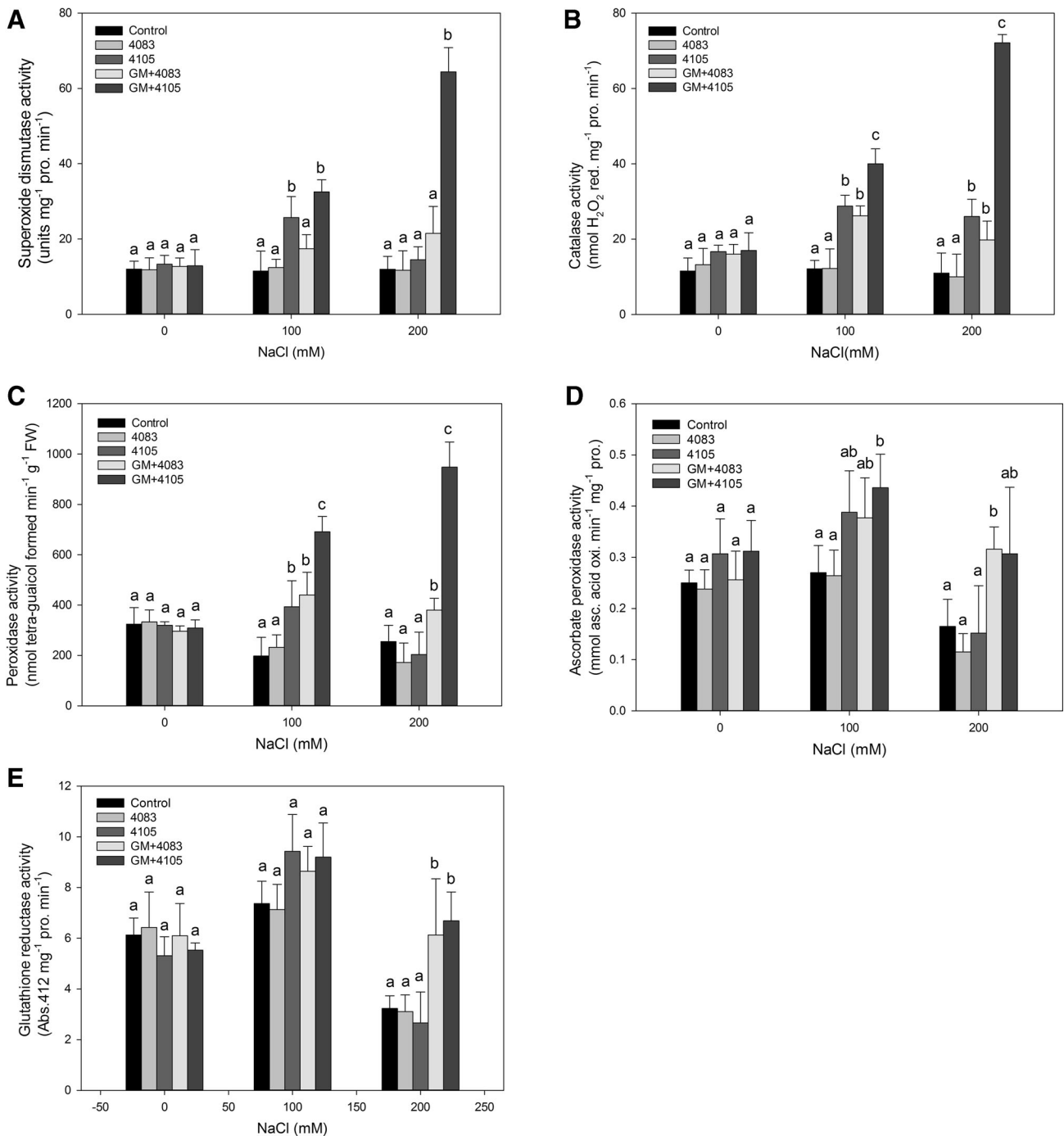


Fig. 3 Effect of AM inoculation on the antioxidant enzymes activity of *Sesbania cannabina* plants inoculated with rhizobium and *Glomus mosseae* under salt stress. **a** Effect of AM inoculation on the superoxide dismutase (SOD) activity (units mg^{-1} protein min^{-1}) of *Sesbania cannabina* plants. **b** Effect of AM inoculation on the catalase (CAT) activity (nmol H_2O_2 reduced mg^{-1} prot. min^{-1}) of *Sesbania cannabina* plants. **c** Effect of AM inoculation on the peroxidase (POX) activity (nmol tetraguaiacol formed min^{-1} g^{-1} FW)

of *Sesbania cannabina* plants. **d** Effect of AM inoculation on the ascorbate peroxidase (APOX) activity (1 mol asc. acid oxidized min^{-1} mg^{-1} protein) of *Sesbania cannabina* plants. **e** Effect of AM inoculation on the glutathione reductase (GR) activity (absorbance at 412 nm mg^{-1} protein min^{-1}) of *Sesbania cannabina* plants. Bars with different letters among inoculations within NaCl levels are significantly different (one-way ANOVA, Duncan’s multiple range test, $P < 0.05$)

scavenging enzymes, CAT and POD (Fig. 3). The increased SOD activity induced by O_2^- radicals results in increased H_2O_2 levels, and this is accompanied by an

increased enzymatic capacity to decompose H_2O_2 (Hernandez and Almansa 2002). Although the difference between the APOX and GR activities in 4105 + GM and

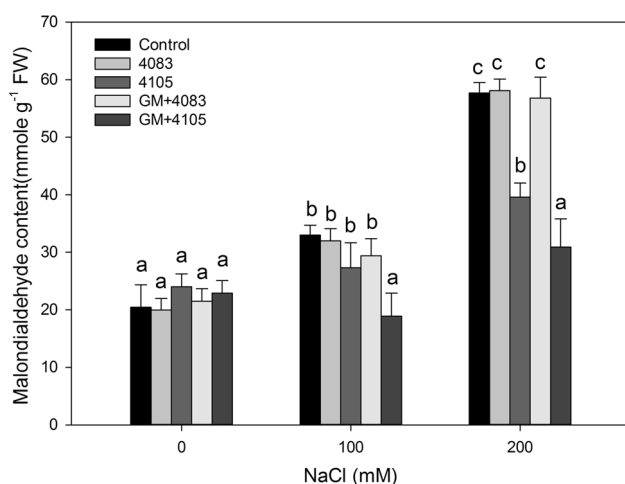


Fig. 4 Effect of AMF and *Rhizobium* inoculation on lipid peroxidation (MDA contents) (mmol g^{-1} FW) in *Sesbania cannabina* plants under salt stress. Bars with different letters among inoculations within NaCl levels are significantly different (one-way ANOVA, Duncan's multiple range test, $P < 0.05$)

4083 + GM-inoculated plants was not significant (Fig. 3d, e), the MDA content of 4105 + GM-inoculated plants was significantly lower than that in 4083 + GM-inoculated plants (Fig. 4).

In summary, the present study revealed that the salt tolerance of rhizobia contributes to the host plant under salt stress, together with AM fungi. In higher salt-tolerant rhizobia-inoculated plants, antioxidative enzyme activities, such as SOD, POX, CAT, and GR, were induced to protect the plants from the oxidative effects of the ROS. The higher salt-tolerant strain obviously had greater protective effects, and exhibited a synergistic interaction with AM fungi to improve the salinity tolerance of *S. cannabina*. The results suggested that screening salt-tolerant rhizobia that have synergistic interactions with AM fungi in triple symbiosis could be a useful strategy to enhance the tolerance of legumes to saline stress and, therefore, improve the fitness of plants. The development of sustainable biofertilizer technology is desirable for maximizing environmentally friendly crop production in saline soils (Singh and others 2011).

Acknowledgments This work was financed by the Strategic Priority Research Program of the Chinese Academy of Sciences (Grant No. XDA11020403), the Key Research Program of the Chinese Academy of Sciences (Grant No. KZZD-EW-14), the National Natural Science Foundation of China (31370108 and 31570063), One Hundred-Talent Plan of Chinese Academy of Sciences (CAS), Yantai Science and Technology Project (2013JH021).

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