

The response of the *Rhizobium leguminosarum* bv. *trifolii* wild-type and exopolysaccharide-deficient mutants to oxidative stress

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Abstract

Aims The aim of this study was investigation of the response of *R. leguminosarum* bv. *trifolii* wild-type and its two *rosR* and *pssA* mutant strains impaired in exopolysaccharide (EPS) synthesis to oxidative stress conditions caused by two prooxidants: a superoxide anion generator- menadione (MQ) and hydrogen peroxide (H_2O_2).

Methods The levels of enzymatic (catalase, superoxide dismutase, pectinase and β -glucosidase) and non-enzymatic (superoxide anion generator, formaldehyde, phenolic compounds) biomarkers were monitored using biochemical methods in both the supernatants and rhizobial cells after treatment with 0.3mM MQ and 1.5mM

H_2O_2 . The viability of bacterial cells was estimated using fluorescent dyes and confocal laser scanning microscopy. In addition, the effect of prooxidants on symbiosis of the *R. leguminosarum* bv. *trifolii* strains with clover was established.

Results The tested stress factors significantly changed enzymatic patterns of the rhizobial strains, and the wild-type strain proved to be more resistant to these prooxidants than both *pssA* and *rosR* mutants. Significantly higher activities of both catalase and superoxide dismutase have been detected in these mutants in comparison to the wild-type strain. H_2O_2 and MQ also increased the levels of pectinase and β -glucosidase activities in the tested strains. Moreover, pre-incubation of *R. leguminosarum* bv. *trifolii* strains with the prooxidants negatively affected the viability of bacterial cells and the number of nodules elicited on clover plants.

Conclusions EPS produced in large amounts by *R. leguminosarum* bv. *trifolii* plays a significant protective role as a barrier against oxidative stress factors and during symbiotic interactions with clover plants.

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Introduction

Rhizobium leguminosarum bv. *trifolii* is an aerobic gram-negative bacterium that can live free in the soil

or establish a symbiosis with roots of clover plants (*Trifolium pratense*) (McIntyre et al. 2007). During this process, a sophisticated exchange of recognition signals occurs between both symbiotic partners (Gibson et al. 2008). In response to flavonoids secreted by legume roots, rhizobia synthesize nodulation factors, which are essential for formation of specialized new organs on roots called nodules. Inside nodule cells, bacteria differentiate into their symbiotic forms—bacteroids, which reduce atmospheric nitrogen to ammonia (Gibson et al. 2008). Nitrogenase, a key enzyme of nitrogen fixation, is inactivated rapidly and irreversibly by oxygen. Leghemoglobin occurring in the cortex of nodules constitutes a diffusion barrier that limits permeability to oxygen and prevents the activity of this enzyme. Although this system efficiently reduces the excess of oxygen diffused to nodule cells, a high rate of respiration providing energy for the nitrogen reduction process generates reactive oxygen species (ROS) such as superoxide anion radicals (SOR) and hydrogen peroxide (H_2O_2) (Santos et al. 2001). Additionally, rhizobia living in soil and in the rhizosphere of other plants are exposed to oxidative stress factors. The response of both free-living bacteria and bacteroids to the toxic form of oxygen is an important factor for nodulation and nitrogen fixation. Reactive oxygen species also play a key role as central signal molecules acting in plant adaptation to both biotic and abiotic stresses (Hérouart et al. 2002). Increased levels of ROS enable plants to control and block pathogen penetration (Orikasa et al. 2010). Because pathogenesis and symbiosis are a variation on a common theme, rhizobia are at first recognized as intruders and generation of oxidative burst in host cells is observed (Baron and Zambryski 1995). In order to establish legume-rhizobium symbiosis, there is a special mechanism whereby the typical host defence elements are suppressed. There are many examples showing accumulation of ROS during infection of leguminous plants by rhizobia (Ramu et al. 2002; Pauly et al. 2006). It has also been proposed that reactive oxygen species can stimulate expression of plant and/or bacterial genes that are significant for nodule formation (Hérouart et al. 2002).

Thus, both during progression of symbioses and during life inside formed nodules, rhizobia are exposed to the action of various stress factors including prooxidants. In response to changing environmental conditions, rhizobia have evolved an efficient defence system to inactivate different free radical derivatives

including superoxide anion radicals, hydrogen peroxide, or extremely reactive hydroxyl radicals. Using enzymatic and non-enzymatic antioxidants, these microorganisms can keep a physiologically safe level of ROS throughout the exponential and stationary phases of growth and during the interaction with plants (Matamoros et al. 2003). An investigation of the role of superoxide radicals and H_2O_2 in alfalfa—*Sinorhizobium meliloti* symbiosis showed the presence of such ROS at early stages of the infection process and during the development of infection threads (Santos et al. 2000). Absence of H_2O_2 inside bacterial cells and bacteroids suggests the action of effective antioxidative mechanisms in rhizobia (Hérouart et al. 2002). One of the enzymes which plays a major role in protection of oxidative cell homeostasis in bacteria is superoxide dismutase (SOD), a metalloenzyme catalysing dismutation of O_2^- to H_2O_2 and O_2 (Santos et al. 2000; Saenkham et al. 2007). Hydrogen peroxide, the product of dismutation, is degraded by catalases (CAT), which play a crucial role in the symbiosis during both host plant infection and nitrogen fixation processes (Orikasa et al. 2010).

In addition, cell surface polysaccharides of rhizobia, including exopolysaccharides (EPS) and lipopolysaccharides (LPS), play a key role in both adaptation to environmental conditions and establishment of effective symbioses with leguminous plants (Downie 2010; Mathis et al. 2005). EPS performs several functions, such as nutrient gathering, protection against stress factors and antimicrobial compounds, biofilm formation, and attachment to abiotic surfaces and host plant roots. Moreover, this polysaccharide is a suppressor of the host plant defence reaction, and its low-molecular-weight fraction plays a specific role as a signal molecule in the symbiotic dialogue (Janczarek 2011). EPS-deficient mutants of *Rhizobium leguminosarum* and *Sinorhizobium meliloti* are impaired in invasion of nodule cells and nitrogen fixation (Rolfe et al. 1996; Cheng, and Walker 1998). EPS of *R. leguminosarum* bv. *trifolii* is a heteropolymer composed of repeating units containing five glucose, one galactose and two glucuronic acid residues, modified by acetyl, pyruvyl, and 3-hydroxybutanoyl groups (Janczarek 2011).

Two *pssA* and *rosR* genes play a key role in EPS production in *R. leguminosarum* (Janczarek et al. 2009b). *pssA* encodes a glucosyl-IP-transferase involved in the first step of EPS synthesis and a mutation in this gene results in a total lack of production of this polymer. *rosR* encodes a transcriptional regulator

involved in positive regulation of this process and its mutation substantially (3-fold) decreases the amount of produced EPS (Janczarek and Skorupska 2007; Janczarek et al. 2009a). Both mutants induce formation of small nodules on clover plants that are unable to fix nitrogen. Moreover, the *rosR* mutant shows several other pleiotropic effects such as increased sensitivity to surface-active detergents and some osmolytes, changes in extracellular and membrane protein profiles, and decreased ability to infect roots of the host plant, indicating that the product of this gene is involved in the adaptation of *R. leguminosarum* bv. *trifolii* to stress conditions (Janczarek et al. 2010). On the other hand, multiple copies of *rosR* and *pssA* genes significantly enhance EPS production and the number of nodules induced on clover roots (Janczarek et al. 2009a). The expression of these genes is affected by phosphate and clover root exudates (Janczarek and Skorupska 2011).

Up to now, there are no data concerning mechanisms of oxidative stress response in *R. leguminosarum* bv. *trifolii*, therefore the aim of this study was to investigate the effects of two prooxidants, menadione (2-methyl-1,4-naphthoquinone; MQ) and H_2O_2 , on chosen enzymatic (catalase and superoxide dismutase activities) and non-enzymatic (superoxide anion radicals, formaldehyde, and phenolic compounds) biomarkers related to antioxidative defence. MQ is a redox-cycling compound commonly used as a SOR-generator (Saenkham et al. 2007). These prooxidants were chosen because large amounts of SOR and H_2O_2 are present in infection threads and these ROS species are related to active oxygen derivatives produced by plants during the oxidative burst phenomenon. We investigated how bacteria of *R. leguminosarum* bv. *trifolii* species respond to H_2O_2 - and MQ-mediated oxidative stress when exposed to a low concentration of these compounds for a relatively long incubation period.

Moreover, because hydrolytic enzymes were found to be important for proper infection of host plants by rhizobia, the activities of pectinase (PEC) and β -glucosidase (β -GLU) were also estimated (Lum and Hirsch 2003; Mateos et al. 1992). To investigate the possible role of EPS in *R. leguminosarum* bv. *trifolii* antioxidative defence mechanisms, both *pssA* and *rosR* mutants have been included in these experiments. The present study also evaluates the efficiency of the investigated strains in development of symbiotic interactions with clover using bacteria after pre-treatment with the prooxidants.

Materials and methods

Strains, media, and growth conditions

R. leguminosarum bv. *trifolii* wild-type strain Rt24.2 and its two *rosR* (Rt2472) and *pssA* (Rt5819) mutants were tested for their sensitivity to chosen prooxidants (H_2O_2 and MQ). The bacterial strains used in this study have been described in details in a previous paper (Janczarek et al. 2009a). Rt24.2 and its derivatives were grown in tryptone-yeast (TY) medium at 28 °C (Sambrook et al. 1989). When required, antibiotics were used at the following final concentrations: kanamycin 40 $\mu\text{g mL}^{-1}$ and nalidixic acid 40 $\mu\text{g mL}^{-1}$. To study the influence of the prooxidants on the growth of rhizobial strains, different concentrations of H_2O_2 (0–3 mM) and menadione (0–3 mM) were tested. The viability of cells from these cultures was established by spotting 15- μl aliquots onto TY agar plates and incubation at 28 °C for 3 days. To study the sensitivity of these strains to the prooxidants, different concentrations of H_2O_2 and MQ (up to 3 mM) were added to 48-h cultures and the incubation was continued up to 6 h. After each 30 min, 15- μl aliquots of the cultures were spotted on the plates and bacterial growth was monitored during 3 days.

The effect of EPS on survival of *rosR* and *pssA* mutant cells in the presence of prooxidants

To study the influence of EPS on survival of the *rosR* and *pssA* mutants in the presence of prooxidants, bacteria of these mutants and the wild-type strain were grown in 5 ml TY medium at 28 °C for 24 h. Then, the optical density OD_{600} of the cultures was measured and equalized with TY to 0.4. For this experiment, two Rt5819/Rt24.2 and Rt2472/Rt24.2 mixtures containing equal ratios of the mutant culture and the wild-type strain culture were used. In addition, cultures of the Rt24.2, Rt2472 and Rt5819 strains of $OD_{600}=0.2$ were used as controls. 10- μl aliquots of these bacterial cultures and the mixtures were spotted on TY plates containing different concentrations of MQ (0–1 mM) and H_2O_2 (0–3 mM). In order to eliminate the growth of the Rt24.2 cells present in the Rt5819/Rt24.2 and Rt2472/Rt24.2 mixtures, kanamycin was added into agar medium (the Rt5819 and Rt2472 mutant strains are kanamycin-resistant). The plates were incubated at

28 °C and after 3 days bacterial growth was estimated. Three independent experiments were performed in triplicate for each bacterial mixture and culture tested.

Preparation of biological samples

To establish the influence of the oxidants on enzymatic and non-enzymatic parameters of the tested *R. leguminosarum* bv. *trifolii* strains, the bacteria were grown in 450 ml TY medium at 28 °C for 2 days with rotary shaking (120 rpm) to OD₆₀₀ of 0.8. Then, MQ and hydrogen peroxide were added to the final concentrations of 0.3 mM and 1.5 mM, respectively, and the incubation was continued during 3 h. Subsequently, the cultures were centrifuged at 6,000×g for 20 min at 4 °C, and the supernatants and bacterial pellets obtained were used for further experiments.

In order to obtain bacterial extracts, the cell pellets were resuspended in 1 ml of ice-cold MilliQ water and sonicated (six times during 30 s, one pulse per s) on ice following the conditions described by Hérouart et al. (1996). After sonication, the extracts were centrifuged at 10,000×g for 15 min at 4 °C. The clarified extracts were divided into portions and frozen at –70 °C.

Enzyme assays

Superoxide dismutase (SOD)

The activity of SOD was assayed in both culture supernatants and intracellular fractions using the complete set for SOD determination according to manufacturer's instruction (kit for micro-method, Sigma, St. Louis, USA). Superoxide anion radicals were produced by the xanthine oxidase reaction and oxidized with water-soluble tetrazolium salt, WST-1 (2-(4-iodophenyl)-3-(4-nitrophenyl)-5-(2,4-disulfophenyl)-2H-tetrazolium, monosodium salt, which produces a water-soluble formazan dye upon reduction with SOR. The rates of the reduction with SOR are linearly related to the xanthine oxidase activity, and are inhibited by SOD (Beauchamp and Fridovich 1970). Absorbance was measured at 440 nm using the Bio-Tek Microplates Reader and the specific activity of SOD was expressed in U per mg of protein contained in the tested sample.

Catalase (CAT)

Catalase activity was determined according to the method provided by Aebi (1984). Calculation of CAT activity was based on the amount of hydrogen peroxide decomposed by the enzyme during the incubation time (30 s). The changes in absorbance were recorded at 240 nm. The specific activity of catalase was calculated in μ katals per mg of protein contained in the sample.

Pectinase (PEC)

Pectinolytic activity in cell-free culture supernatants was determined using 1 % solution of pectin in 50 mM citrate buffer (pH 4.8) as a reaction substrate. A 0.5-ml sample was mixed with 0.5 ml of 1 % pectin and incubated for 30 min at 40 °C and then, the amounts of liberated reducing sugars were quantified. The concentration of sugars was evaluated using the Lloyd and Whelan method (Lloyd and Whelan 1969). Absorbance was recorded at 520 nm and compared with the calibration curve. One unit of pectinase activity (U) was described as the amount of the enzyme that released 1 μ M galactouronic acid per min under the experimental conditions. The specific activity of pectinase was calculated in U per mg of protein contained in the sample.

β -Glucosidase (β -GLU)

The synthetic substrate p-nitrophenyl- β -D-glucopyranoside (p-NPG) was used to detect β -glucosidase activity, according to the procedure described by Mitchell et al. (1986). Samples of culture supernatants were analysed for this enzymatic activity. At first, 0.75 ml of 10 mM p-NPG in 50 mM citrate buffer (pH 4.8) was added to test tubes. Afterwards, 0.05 ml of the supernatant and 0.425 ml of deionized water was added, and the reaction mixtures were incubated at 30 °C for 30 min. Then, the reaction was stopped by addition of 3 ml of 0.5 M sodium carbonate. The amount of the released product (p-nitrophenol) was determined by spectrophotometric measurement of absorbance at 420 nm. A unit of enzyme activity (U) was expressed as the amount of β -glucosidases, which liberates 1 μ M p-nitrophenol per min under the experimental conditions. The specific activity of β -GLU was calculated in U per mg of protein contained in the analysed sample.

Determination of non-enzymatic components

Relative level of superoxide anion radicals (SOR)

The measurement of the level of SOR was done according to the method described previously (Paździoch-Czochra et al. 2003). The relative level of SOR was measured spectrophotometrically by detection of superoxide-dependent formation of formazone from nitro tetrazolium blue (NBT) in the alkaline conditions. The reaction mixture contained 3 ml of distilled water, 0.05 ml of 1 M NaOH, 0.1 ml of 5 mM NBT, and 0.1 ml of the sample. After incubation for 30 min at room temperature, absorbance at 560 nm was measured. It was observed that the stabilization of SOR in the alkaline conditions prevented precipitation of dark-blue formazone within about 40 min.

Formaldehyde (FA)

The formaldehyde concentration was determined spectrofluorimetrically with Nash reagent (excitation=410 nm, emission=510 nm) using the FluoroMax-2 equipment (Rapoport et al. 1994). The changes in the emission were measured after 10-min incubation at 60 °C and compared with the calibration curve.

Phenolic compounds (PHC)

The concentration of phenolic compounds (hydroxyl-, methoxyl-phenolic acids) was determined with diazosulfanilamide (SA) using the DASA test (Malarczyk 1989). The reaction mixture contained 0.1 ml of SA (1 % SA in 10 % HCl), 0.1 ml of 5 % NaNO₂ solution, and 0.1 ml of the sample. Each sample was stirred thoroughly and neutralized by addition of 1 ml of 20 % Na₂CO₃. Absorbance was recorded at 500 nm and compared with the calibration curve ($y=6.85x-0.0218$, $R^2=0.999$).

Chemiluminometric detection of hydrogen peroxide

The concentration of H₂O₂ in bacterial extracts was determined using the chemiluminescent method developed by Perez and Rubio (2006) based on the luminescence of luminol. In this method, ferricyanide used in the earlier methodology (Warm and Laties 1982) was replaced by Co (II), which is a catalyst for the reaction of luminol with H₂O₂, what significantly increased the

sensitivity of detection. The stock solution was prepared by mixing 10 ml of a 6.5 mM solution of luminol in carbonate buffer pH 10.2 with 2 ml of 0.55 μM CoCl₂ in the same buffer. The prepared reagent was diluted 10-fold after 1 h and was ready to use after 12 h. The signals from the Co(II)–H₂O₂–luminol reaction were detected by a Lumat LB 9506 luminometer (Berthold, Germany). The peak values were recorded and compared with the calibration curve prepared for the different concentrations of H₂O₂. The amount of H₂O₂ in the bacterial extracts was expressed as μg per mg of proteins contained in the analysed sample.

Protein concentration assay

The protein concentration was determined with the Coomassie brilliant blue (G-250) dye-binding method (Bradford 1976) using a Bio-Rad dye stock solution with bovine serum albumin as the standard. All parameters presented in the paper refer to the protein concentration in the sample.

Visualization of intracellular SOD and CAT activity using native PAGE electrophoresis

Bacterial clarified extracts obtained after sonication and centrifugation were concentrated and separated by ultrafiltration using the Microcon Centrifugal Filter Units, 3000 NMWL designed by Millipore. Subsequently, 15 μg of proteins from the samples were introduced into each well of 12.5 % native polyacrylamide gel. The gels were run at 4 °C and 145 V. After protein separation, SOD activities were visualized according to the method of Bayer and Fridovich (1987). The bands representing CAT activity were located using the ferricyanide negative stain as described by Wayne and Diaz (1986).

Determination of cell viability of rhizobial strains using confocal laser scanning microscopy

In order to determine the viability of rhizobial cells after the treatment with the prooxidants, bacteria of the wild-type and the *pssA* and *rosR* mutant strains were grown in TY medium at 28 °C for 48 h (3 6-ml cultures for each strain were prepared). OD₆₀₀ of each culture was measured and equalized with TY medium to 0.4. Next, the cultures were divided into three 2-ml portions. The first ones were used as control cultures, whereas 1.5 mM H₂O₂ and 0.3 mM MQ were added to the second and

to the third portions, respectively, and the incubation was continued during 3 h. The two-component Bacterial Viability kit (LIVE\DEAD *BacLight* kit, Invitrogen) was used to investigate the influence of the prooxidants on viability of rhizobial cells. Live cells were stained with Syto-9 dye, whereas dead cells with propidium iodide. 1-ml portions of cultures were centrifuged at $9,000\times g$ for 10 min. Obtained pellets were resuspended in 1 ml 0.85 % NaCl and centrifugation was repeated. Then, the bacterial pellets were resuspended in 1 ml 0.85 % NaCl and 50- μ l portions thereof were supplemented with Syto-9 and propidium iodide to the final concentrations of 5 μ M and 30 μ M, respectively, and left in darkness for 20 min at room temperature. Subsequently, bacterial suspensions (50- μ l) were introduced into wells of 96-well polystyrene plates and analysed using a microscope. In order to determine the number of cells used for this analysis, a set of dilutions of the culture of $OD_{600}=0.4$ was prepared for each strain, and 100- μ l portions thereof were loaded onto agar plates, incubated at 28 °C for 3 days, and the number of colonies was counted. 1 ml of these cultures of the Rt24.2, Rt5819, and Rt2472 strains contained 8.16×10^8 , 8.82×10^8 , and 8.23×10^8 cells, respectively. The Olympus SV1000 microscope was used for visualization of the prooxidant-treated bacterial cells. For each strain and each condition, the experiment described was repeated three times. The ratio of live to dead cells was calculated using the ImageJ 1.43e software (Wayne Rasband, NIH, USA).

Plant tests

Red clover (*Trifolium pratense* cv. Diana) seeds were surface sterilized using 0.1 % $HgCl_2$ (3 min), three washes with sterile water, 70 % ethanol (3 min), and next three washes with water. Subsequently, the seeds were placed on Fåhræus agar plates where they germinated. Then, seedlings were transferred onto Fåhræus slants for further growth (Vincent 1970). Four-day-old seedlings were inoculated with bacterial suspensions of Rt24.2 and its derivatives (Rt5819 and Rt2472) of $OD_{600}=0.2$ (100 μ l per plant) and grown 28 days under natural light supplemented with artificial light (14-h day at 24 °C and 10-h night at 18 °C) in a greenhouse. To study the effect of menadione and H_2O_2 on nodulation ability of these strains and their effectiveness in symbiosis with clover, 10-ml bacterial suspensions of OD_{600} of 0.2 were incubated with 1.5 mM H_2O_2 or 0.3 mM MQ

for 1 h at 28 °C. To prevent the deleterious effect of these compounds to plants, the bacteria after incubation were centrifuged 10 min at $5,000\times g$, and the pellets obtained were washed twice in 5 ml sterile water. Finally, 5-ml bacterial suspensions in water of $OD_{600}=0.2$ were prepared and 100 μ l portions were used to infect individual clover plants. The plants were inspected for root nodule formation every week, and after 4 weeks they were harvested and wet shoot and root masses were estimated.

Statistical analysis

The results are mean \pm SD from three experiments ($n=3$) performed in triplicate. Comparison of values between untreated and prooxidant-treated bacteria of the particular strain and between different strains in the same conditions were performed using a one-way ANOVA to find statistically significant differences. In cases where the null hypothesis (all population means are equal) was rejected at the $\alpha=0.05$ level, the Tukey's HSD (Honestly Significant Difference) test was applied. Significant differences among treatments and the tested strains were considered at the level of p values <0.05 .

Chemicals

Most of the reagents and substrates were obtained from Sigma (St. Louis, USA), POCH (Poland) and Invitrogen (USA).

Results

Growth and survivability of *R. leguminosarum* bv. *trifolii* in response to H_2O_2 and menadione

In order to establish the sensitivity and bacterial response to H_2O_2 - and menadione-mediated oxidative stress, the growth and survival of the *R. leguminosarum* bv. *trifolii* wild-type and two *pssA* and *rosR* mutant strains were monitored in the presence of various concentrations of these compounds. Two different treatments were used. In the first one, cell cultures of low density were exposed for a long period (96 h) to different concentrations of the stressors and the growth was determined by measurement of cell turbidity. In general, hydrogen peroxide affected the growth of these bacteria

less drastically than MQ (Fig. 1). It was observed that 1 mM H₂O₂ did not affect the bacterial growth (with the exception of the *rosR* mutant). 1.5 mM H₂O₂ moderately

decreased the growth of these strains (Fig. 1a–c), whereas the 3 mM concentration of this prooxidant resulted in growth inhibition for all tested strains. Among these

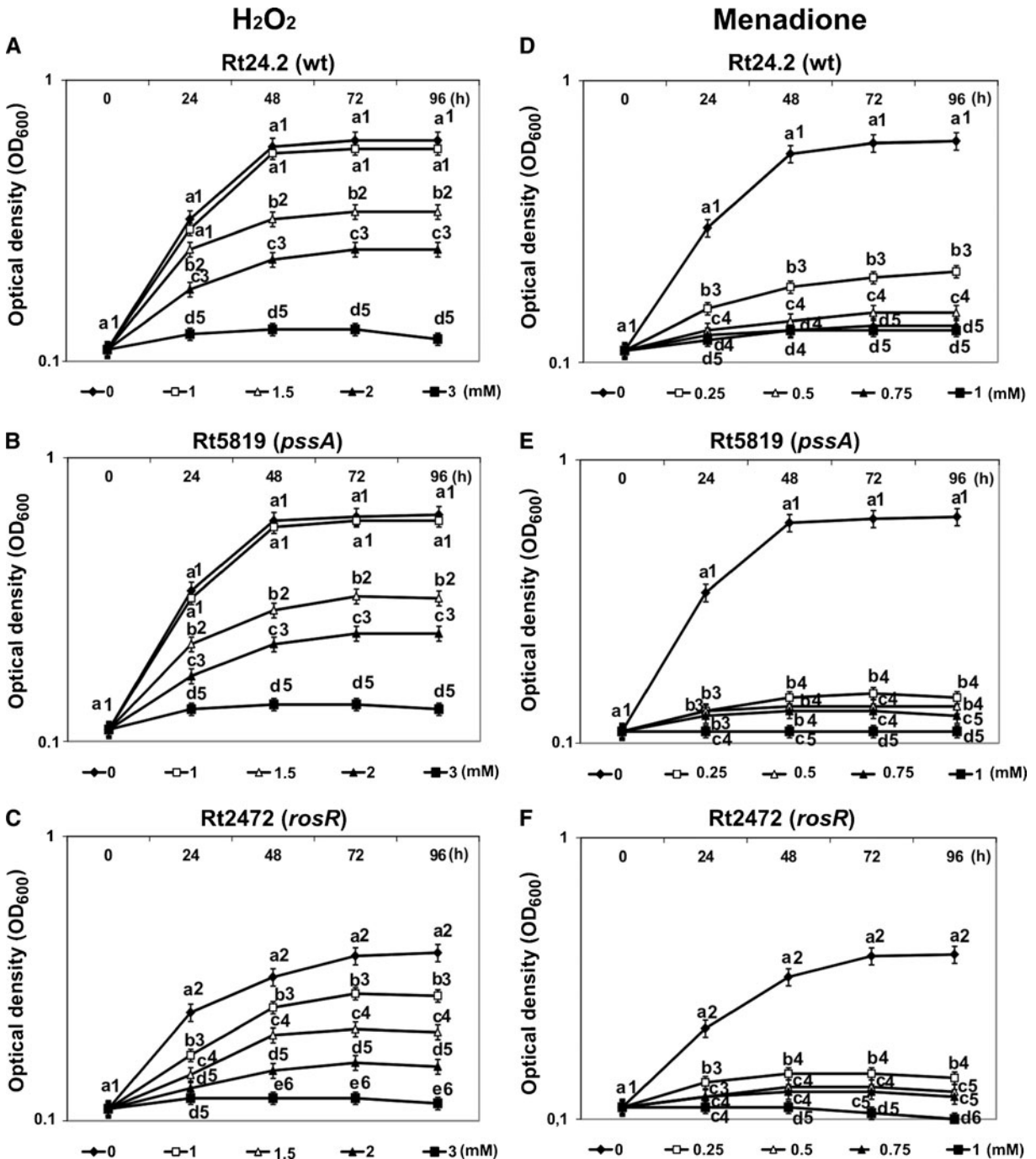


Fig. 1 The growth rate of the *R. leguminosarum* bv. *trifolii* wild-type and the *pssA* and *rosR* mutants cultivated in TY medium supplemented with different concentrations of MQ and H₂O₂. Data are mean ± SD for three experiments. Differences statistically

significant ($p < 0.05$; ANOVA, post hoc Tukey’s test) are marked by different letters (for particular strain) and different numbers (for particular treatment)

strains, the *rosR* mutant showed the highest sensitivity to hydrogen peroxide.

In contrast to H₂O₂, a strong negative effect of MQ on the growth rates of all the tested rhizobial strains was found (Fig. 1d–f). Even such a low concentration of MQ as 0.25 mM negatively affected the bacterial growth, and 0.75 mM of this compound resulted in total inhibition of the growth of these strains. Moreover, both *pssA* and *rosR* mutants deficient in EPS production were significantly more sensitive to this prooxidant than the wild-type strain, indicating a significant role of this polymer in protection against oxidative stress factors.

The next experiments were carried out in order to establish the highest concentration and time of treatment with the prooxidants which did not drastically affect the cell viability. To this end, different concentrations of H₂O₂ and MQ (up to 3 mM) were added to 72-h cultures of the tested strains and the incubation was continued up to 6 h. After each 30 min, aliquots of the cultures were taken and loaded onto agar plates, and the growth of colonies was monitored. We found that bacteria at the stationary-phase of growth were less sensitive to the oxidants in comparison to those growing in the presence of these factors. It was observed that the wild-type and both mutants survived the treatment with 0.3 mM MQ and 0.45 mM up to 5 h and 3 h, respectively.

Moreover, the bacteria retained viability after the 6-h incubation in the presence of 2 mM H₂O₂. Based on these results, we chose 0.3 mM MQ and 1.5 mM H₂O₂ as the most appropriate for all of the tested strains for further experiments.

For more precise determination of cell viability of the *R. leguminosarum* bv. *trifolii* wild-type and the *pssA* and *rosR* mutants treated with these prooxidants, we calculated the ratio of live to dead cells using dyeing specific for these two types of cells (Fig. 2). The results obtained demonstrated that the mutation in *rosR* gene negatively affected cell viability of this strain (Table 1). In the 48-h culture of the wild-type strain, 95.1 % of the bacteria were alive, whereas 85.3 % of the cells were live in the case of *rosR* mutant. However, in the *pssA* mutant, a similar level of live cells to that of the wild-type strain was observed. The treatment with 1.5 mM H₂O₂ during 3 h did not significantly affect the viability of these strains. However, 0.3 mM MQ exerted a strong negative effect on cell viability, which was especially visible in the case of the non-mucoid *pssA* mutant (40.38 % of live cells) (Table 1).

In summary, all these growth experiments indicate that the proper level of EPS is important for bacterial survival and adaptation to stress conditions.

The effect of EPS on survival of the *rosR* and *pssA* mutant cells in the presence of prooxidants

To study the influence of EPS on survival of the *rosR* and *pssA* mutants in the presence of H₂O₂ and MQ, 10- μ l aliquots of culture mixtures of the mutant and the wild-type strain (Rt2472/Rt24.2 and Rt5819/Rt24.2) were tested on agar plates containing different concentrations of these prooxidants (0–3 mM). In addition, cultures of the wild-type and the *rosR* and *pssA* mutant strains were used as controls. Since both Rt5819 and Rt2472 strains are kanamycin-resistant, this antibiotic was added to the medium in order to eliminate the growth of the wild-type bacteria. After 3-day incubation, we found Rt24.2 strain colonies on plates containing 3 mM H₂O₂ and 0.45 mM MQ, respectively. In contrast, both *rosR* and *pssA* mutants showed significantly higher sensitivity to these factors and only few colonies of these bacteria were visible in the presence of 2 mM H₂O₂ and 0.25 mM MQ. In the case of Rt2472/Rt24.2 and Rt5819/Rt24.2 mixtures, we observed that the mutant bacteria tolerated significantly higher concentrations of the tested prooxidants (3 mM H₂O₂ and 0.45 mM MQ, respectively) than the mutant cells growing in the absence of EPS derived from the Rt24.2 strain (Table 2). These data confirm that EPS produced by the wild-type cells improved survival of mutant cells which are impaired in the synthesis of this extracellular polysaccharide, and the presence of EPS surrounding rhizobial cells ensures their better survival under stress conditions.

The effect of prooxidant (MQ and H₂O₂) treatment on CAT activity of *R. leguminosarum* bv. *trifolii* strains

In order to determine the response of *R. leguminosarum* bv. *trifolii* to H₂O₂- and MQ-mediated oxidative stress, the levels of CAT activity and the concentrations of intracellular H₂O₂ were determined in the strains untreated and treated with these prooxidants. It was observed for the prooxidant-untreated bacteria that both *pssA* and *rosR* mutants had more than two times higher specific CAT activity in comparison to that of the wild-type strain (Fig. 3a). Moreover, the addition of the prooxidants had a high stimulation effect in Rt24.2

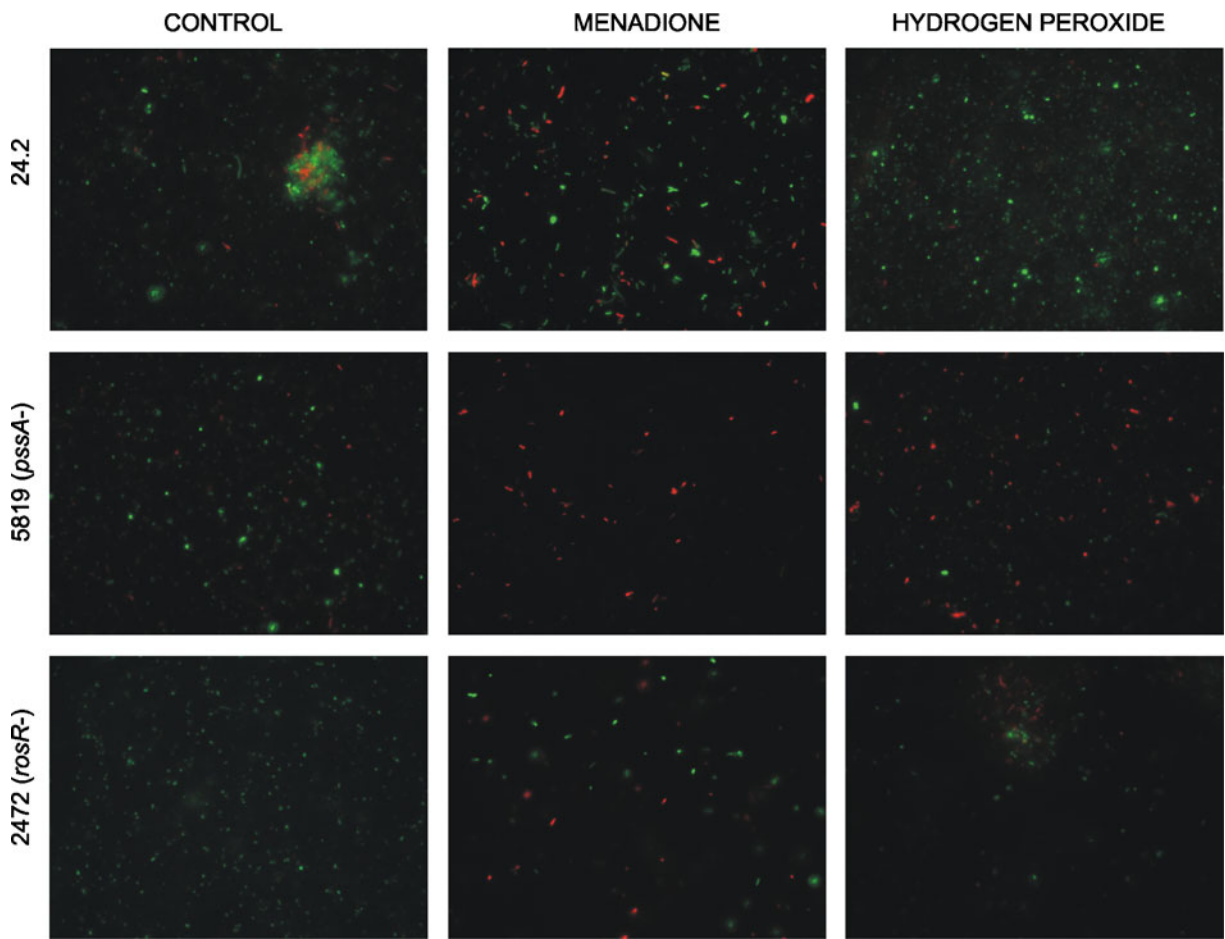


Fig. 2 Cell viability of the *R. leguminosarum* bv. *trifolii* wild-type and the *pssA* and *rosR* mutant strains treated with prooxidants (MQ and H₂O₂) (200× magnification)

strain. The value noted for the H₂O₂-treated bacteria was 2.5-fold higher in relation to the control and even 4.5-fold higher in the MQ-treated cells. In the case of the *pssA* mutant, the presence of the prooxidants slightly decreased the level of CAT activity. For the *rosR* mutant, the level of CAT

activity in the presence of MQ was slightly higher in relation to the strain untreated with this factor. H₂O₂ caused even higher increase in CAT activity in the *rosR* mutant cells in comparison to MQ.

In addition, the levels of intracellular H₂O₂ in the strains untreated and treated with these stress factors

Table 1 The influence of prooxidants on cell viability of the *R. leguminosarum* bv. *trifolii* wild-type and the *pssA* and *rosR* mutant strains

Prooxidant treatment	Viability of <i>R. leguminosarum</i> bv. <i>trifolii</i> strains (%)		
	Rt24.2 (wild-type)	Rt5819 (<i>pssA</i> mutant)	Rt2472 (<i>rosR</i> mutant)
Non-treated	95.1±3.4 ^{a1}	92.9±3.6 ^{a1}	85.1±3.5 ^{a2}
1.5mM H ₂ O ₂	93.2±3.8 ^{a1}	92.3±4.3 ^{a1}	84.0±3.3 ^{a2}
0.3mM MQ	71.3±3.5 ^{b1}	40.4±2.1 ^{b3}	61.8±2.9 ^{b2}

The given values (± standard deviation) are averages of three biological repeats performed in triplicate. The data within the column followed by different letters (for a particular strain) and within the rows followed by different numbers (for a particular treatment) significantly different at $p < 0.05$ (ANOVA, post hoc Tukey's test)

Table 2 The influence of exopolysaccharide on survival of the *R. leguminosarum* bv. *trifolii* *pssA* and *rosR* mutant strains

Bacterial culture or culture mixture	The number of bacterial colonies growing in the presence of different concentrations of prooxidants								
	Hydrogen peroxide (mM)					Menadione (mM)			
	1.0	1.5	2.0	2.5	3.0	0.25	0.45	0.75	1.0
Rt24.2 (wild-type)	un.	un.	25±4 ^{A,a}	11±3 ^{B,b}	4±1 ^{B,c}	26±5 ^{A,a}	14±3 ^{B,b}	0±0 ^{A,d}	0±0 ^{A,d}
Rt5819 (<i>pssA</i>)	un.	un.	2±1 ^{C,a}	0±0 ^{C,b}	0±0 ^{C,b}	3±1 ^{C, a}	0±0 ^{C,b}	0±0 ^{A,b}	0±0 ^{A,b}
Rt2472 (<i>rosR</i>)	un.	un.	9±2 ^{B,a}	0±0 ^{C,b}	0±0 ^{C,b}	8±2 ^{B,a}	0±0 ^{C,b}	0±0 ^{A,b}	0±0 ^{A,b}
Rt5819/Rt24.2	un.	un.	32±4 ^{A,a}	20±3 ^{A,a}	8±2 ^{A,b}	31±5 ^{A,a}	24±5 ^{A,a}	0±0 ^{A,c}	0±0 ^{A,c}
Rt2472/Rt24.2	un.	un.	38±6 ^{A,a}	26±6 ^{A,a}	11±3 ^{A,b}	36±6 ^{A,a}	28±6 ^{A,a}	0±0 ^{A,c}	0±0 ^{A,c}

un. uncountable number of bacterial colonies visible as a monolayer spot

^{A, B, C} Within the same column, means followed by different letters are significantly different at $p < 0.05$, ANOVA, post hoc Tukey's test
^{a,b,c,d} Within the same row, means for the individual strain followed by different letters are significantly different at $p < 0.05$, ANOVA, post hoc Tukey's test

were determined (Fig. 3b). In general, this analysis revealed that the concentration of H₂O₂ in cells is

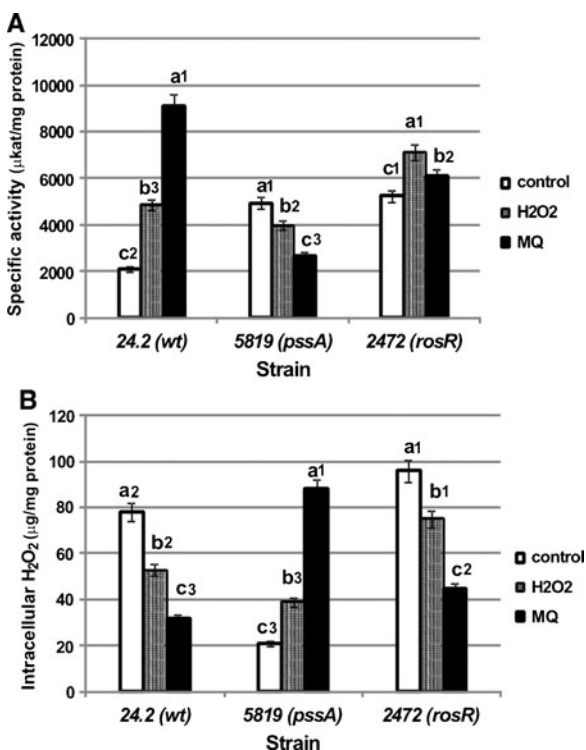


Fig. 3 Specific CAT activity (a) and intracellular level of H₂O₂ (b) assayed in bacteria of *R. leguminosarum* bv. *trifolii* wild-type strain and its derivatives after treatment with prooxidants. Data are mean ± SD for three measurements. Differences statistically significant ($p < 0.05$; ANOVA, post hoc Tukey's test) are marked by different letters (for particular strain) and different numbers (for particular treatment)

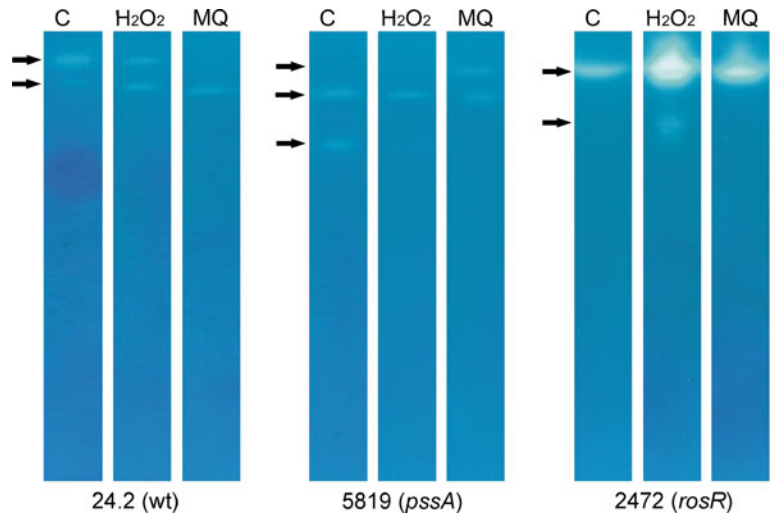
opposite to the level of specific CAT activity (when the individual strain and condition was compared). In the wild-type strain, the lowest CAT activity and the highest concentration of H₂O₂ were observed in the absence of the stressors, whereas the treatment of bacteria with these factors increased the levels of CAT activity and decreased the amount of H₂O₂. The *rosR* mutant exhibited a profile of H₂O₂ levels very similar to that of the wild-type strain, although the values were significantly higher. In contrast to these strains, the untreated bacteria of the *pssA* mutant had the lowest concentration of H₂O₂, and higher levels of H₂O₂ in the presence of the prooxidants.

Moreover, the detection of CAT activity on native PAGE gels was performed, which revealed the presence of two or three bands exhibiting enzyme activity (Fig. 4). In the case of the *rosR* mutant, bands with stronger intensity of CAT activity in the control as well as in the prooxidant-treated samples were observed. Moreover, an additional band of CAT activity was detected in this mutant treated with H₂O₂.

The effect of MQ and H₂O₂ treatment on SOD activity

Our previous studies indicated that mutations in *rosR* and *pssA* genes resulted in several pleiotropic effects, among them disturbances in membrane integrity causing unspecific protein leakage (Janczarek et al. 2009a, 2010). Therefore, the levels of both extracellular and intracellular SOD activity have been determined. We have found that the specific activity of SOD in the supernatant obtained from the Rt24.2 culture reached a

Fig. 4 Native PAGE electrophoresis of intracellular CAT activity in the *R. leguminosarum* bv. *trifolii* wild-type and *pssA* and *rosR* mutants treated with prooxidants (MQ and H₂O₂). 15 µg of proteins were introduced into each well of 12.5 % native polyacrylamide gel



very low level (below 1 U per mg of extracellular protein) (Fig. 5a). The presence of the stress factors (especially MQ) in the cultures of this strain enhanced extracellular SOD activity in comparison to the control.

The supernatants of both *pssA* and *rosR* mutants untreated with prooxidants demonstrated significantly higher SOD activity in comparison to the Rt24.2. In addition, the presence of the stressors in the cultures of these mutants resulted in a significant increase in extracellular SOD activity, with the exception of the *pssA* mutant treated with H₂O₂. The most important changes appeared in the case of this strain treated with MQ (~13.5 U per mg of protein). Furthermore, the addition of H₂O₂ stimulated the SOD activity in the Rt24.2 and even higher in the *rosR* mutant. These results show that the mutations in both *pssA* and *rosR* genes result in remarkably higher activities of SOD in control cultures in relation to the wild-type and in most cases the stress factors additionally increase the level of this activity (Fig. 5a).

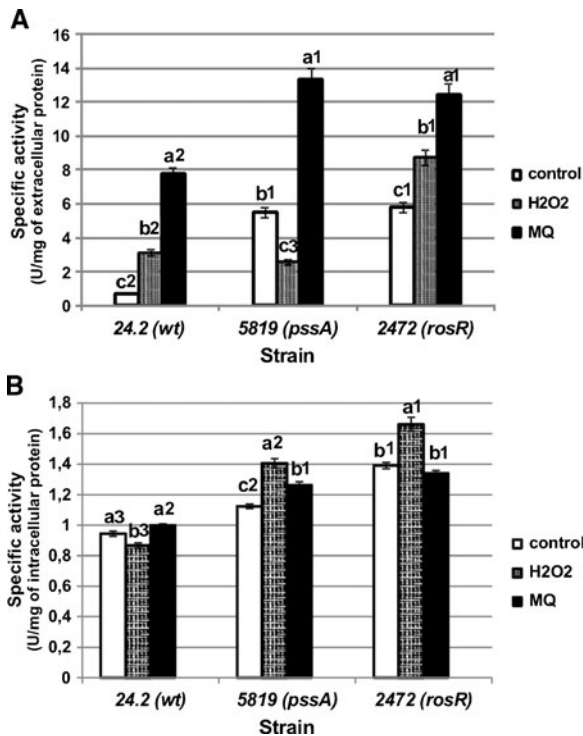
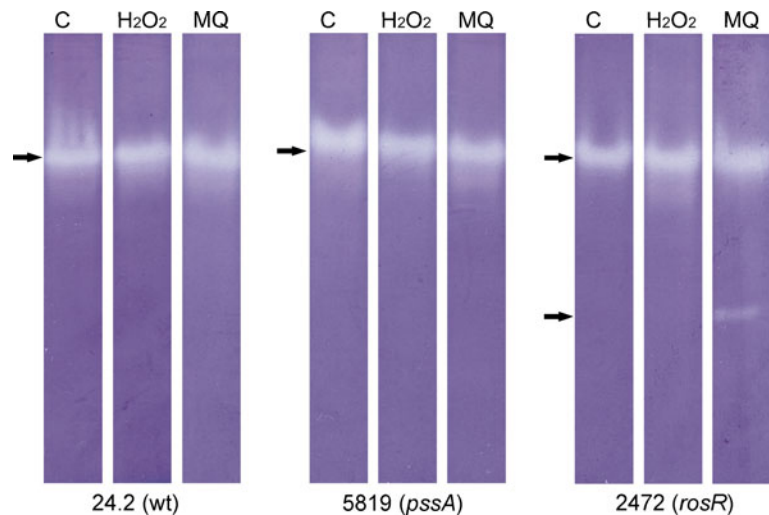


Fig. 5 The effect of prooxidants on extracellular (a) and intracellular (b) specific SOD activity in the wild-type *R. leguminosarum* bv. *trifolii* and the *pssA* and *rosR* mutants. Data are mean ± SD for three measurements. Differences statistically significant ($p < 0.05$; ANOVA, post hoc Tukey’s test) are marked by different letters (for particular strain) and different numbers (for particular treatment)

The intracellular specific activity of SOD in the Rt24.2 was similar to that established in the supernatant of this strain. However, the addition of the prooxidants did not significantly affect the level of this activity within the wild-type bacteria (Fig. 5b). Both the Rt5819 and Rt2472 mutants displayed a moderate increase in the intracellular SOD activity in comparison to the wild-type strain. Moreover, the presence of 1.5 mM of H₂O₂ resulted in a significant increase in the specific SOD activity in these mutants. The highest activity was detected in the case of the H₂O₂-treated *rosR* mutant (~1.65 U per mg of protein). The SOD activities analysed in native electrophoresis showed the presence of one or two bands exhibiting the activity of this enzyme in all of the tested strains both in the control samples and after induction by H₂O₂ and MQ (Fig. 6). The intensity of SOD bands in the strains treated with

Fig. 6 Native PAGE electrophoresis of intracellular SOD activity in the *pssA* and *rosR* mutants and the wild-type *R. leguminosarum* bv. *trifolii* bacteria treated with hydrogen peroxide and menadione (15 μ g of proteins were introduced into each well of 12.5 % native polyacrylamide gel)



the prooxidants seemed stronger than in the control. Interestingly, an additional band of SOD activity was detected in the MQ-treated cells of the *rosR* mutant.

The influence of prooxidants on the levels of SOR, FA, and PHC

The levels of some low-molecular-weight compounds (SOR, FA, and PHC) connected with cellular stress response were estimated in the rhizobial strains. The data were compared to those obtained for the control of the wild-type strain Rt24.2 (100 %) and presented in Table 3 as a percentage of their values. In general, the extracellular and intracellular levels of these compounds significantly differed between the Rt24.2 and both Rt5819 (*pssA*) and Rt2472 (*rosR*) mutants. In the wild-type strain, H₂O₂ caused an increase in both the intracellular and extracellular levels of formaldehyde (FA) and extracellular phenolic compounds (PHC), whereas MQ resulted in higher amounts of intra- and extracellular superoxide anion radicals (SOR) (Table 3). The *pssA* mutant displayed a 2.5-fold lower level of extracellular PHC than the wild-type. Moreover, the addition of H₂O₂ to the culture of this strain resulted in an increase in all of the assayed extra- and intracellular compounds, with the exception of the intracellular PHC. Additionally, MQ present in the cultures increased the amounts of extracellular SOR and FA. However, the highest differences in the levels of these compounds were found in the case of the Rt2472 strain (Table 3). Cells of this mutant showed a higher level of intracellular FA (140 % of the value for Rt24.2) and a very low level of PHC (39.36 %). Moreover, significantly increased amounts

of SOR (153 %), FA (225 %), and PHC (171 %) were found in the culture supernatant of this mutant in comparison to the wild-type, and both prooxidants additionally affected the levels of these compounds.

The effect of MQ and H₂O₂ on the activity of pectinase and β -glucosidase

Biochemical studies concerning extracellular pectinase-like activity in cultures of the wild-type *R. leguminosarum* bv. *trifolii* and the *pssA* and *rosR* mutants showed significant differences between these strains. A nearly 2-fold lower pectinolytic activity was found in the supernatant of the *pssA* mutant in relation to the Rt24.2 (Fig. 7a). On the other hand, a 2-fold higher activity was evidenced in the supernatant of the *rosR* mutant compared to that in the parental strain. In addition, it was observed that both H₂O₂ and MQ caused an increase in the pectinolytic activity in the Rt24.2 and the Rt5819 strains. The most visible stimulation effect was observed in the cultures treated with 0.3 mM MQ, reaching an approximate value of ~54U and ~40U per mg of protein for the wild-type and the *pssA* mutant, respectively. In contrast, the Rt2472 mutant displayed a distinct profile of this enzymatic activity in comparison to the remaining strains. The highest pectinolytic activity was detected in the culture of this mutant with no added prooxidants. In the presence of H₂O₂ and MQ, lower levels of this enzymatic activity were established, suggesting no stimulation effect of these factors on the *rosR* mutant.

The studies concerning the activity of extracellular β -glucosidase (β -GLU) revealed a similar enzymatic

Table 3 The influence of prooxidants on the levels of intracellular and extracellular superoxide anion radicals (SOR), formaldehyde (FA) and phenolic compounds (PHC) determined in the *R. leguminosarum* bv. *trifolii* wild-type and the *pssA* and *rosR* mutants

Strains	Intracellular level*			Extracellular level*		
	SOR (%)	FA (%)	PHC (%)	SOR (%)	FA (%)	PHC (%)
Rt24.2	100±6.2 ^{b1}	100±2.7 ^{b3}	100±7.5 ^{a1}	100±11.6 ^{b2}	100±5.6 ^{c2}	100±4.0 ^{b2}
Rt24.2 H ₂ O ₂	105.6±2.4 ^{b2}	122.3±2.5 ^{ac3}	74.8±2.9 ^{b2}	98.9±3.8 ^{b3}	148.5±14.4 ^{a1}	133.59±2.1 ^{a2}
Rt24.2 MQ	138.9±6.0 ^{a1}	111.8±5.9 ^{bc1}	51.6±2.7 ^{c2}	145.5±5.6 ^{a3}	114.1±3.4 ^{b2}	84.45±1.4 ^{c2}
Rt5819	116.7±7.6 ^{b1}	119.3±11.4 ^{b2}	98.4±3.4 ^{a1}	110.3±3.5 ^{c2}	103.9±1.3 ^{c2}	40.60±0.1 ^{c3}
Rt5819 H ₂ O ₂	138.0±4.4 ^{a1}	199.3±0.6 ^{a1}	97.8±8.6 ^{a1}	173.9±9.7 ^{b1}	123.7±12.9 ^{b1,2}	144.49±3.4 ^{a1}
Rt5819 MQ	111.1±2.9 ^{b2}	106.6±10.4 ^{b1}	51.9±4.1 ^{b2}	211.8±5.8 ^{a1}	148.3±5.5 ^{a1}	87.74±1.2 ^{b2}
Rt2472	94.4±4.0 ^{a1}	141.0±13.3 ^{a1}	39.4±2.3 ^{b2}	153.8±0.8 ^{a1}	225.9±14.0 ^{a1}	171.1±1.6 ^{a1}
Rt2472 H ₂ O ₂	72.2±8.4 ^{b3}	156.7±5.1 ^{a2}	73.0±5.4 ^{a2}	149.6±4.2 ^{bc2}	125.8±5.5 ^{b2}	66.0±0.3 ^{c3}
Rt2472 MQ	66.7±4.7 ^{b3}	110.5±3.1 ^{b1}	84.7±7.0 ^{a1}	156.5±1.5 ^{ac2}	114.8±1.1 ^{c2}	104.7±1.1 ^{b1}

* Relative levels of superoxide anion radicals (SOR), formaldehyde (FA) and phenolic compounds are given in % of the control values of the Rt24.2 strain. The given values (± standard deviation) are averages of three independent experiments performed in triplicate

^{a,b,c} Different letters indicate significantly different values for a particular SOR assayed in the individual strain untreated and treated with MQ and H₂O₂ (data within the column compared) ($p \leq 0.05$; ANOVA, post hoc Tukey's test)

^{1,2,3} Different numbers indicate significantly different values for individual SOR assayed in Rt24.2, Rt5819 and Rt2472 strains in particular conditions (untreated strains; the strains treated with MQ; and the strains treated with H₂O₂) ($p \leq 0.05$; ANOVA, post hoc Tukey's test)

profile for both the wild-type and the *pssA* mutant (Fig. 7b). A very low level of the β -GLU activity in the culture supernatants of these strains was observed. The presence of 1.5 mM H₂O₂ and 0.3 mM MQ resulted in a higher enzymatic activity, and a significant increase (4-fold for the Rt24.2 and nearly 6-fold for the Rt5818) was observed after addition of MQ to the bacterial cultures. In contrast, the *rosR* mutant displayed a different enzymatic profile in relation to these strains. In the culture supernatant of Rt2472, a 3.5-fold higher β -GLU activity in comparison to that of the wild-type and the *pssA* mutant was observed. The addition of H₂O₂ did not significantly change the level of this activity in the Rt2472 supernatant. Moreover, MQ only slightly increased the level of extracellular β -GLU activity.

The influence of prooxidants on symbiosis of *R. leguminosarum* bv. *trifolii* strains with clover

The wild-type strain and the *pssA* and *rosR* mutants were used to study the effect of menadione and H₂O₂ on symbiosis of *R. leguminosarum* bv. *trifolii* with clover. In this experiment, before infection of clover plants these strains were incubated for 1 h at 28 °C in the presence of 1.5 mM H₂O₂ and 0.3 mM MQ. Control plants were infected by bacteria that were not treated by these stress factors. We have observed that both tested

stressors affected the nodulation ability and, in consequence, efficiency in nitrogen fixation of the wild-type strain (Table 4). MQ exerted a stronger negative effect than H₂O₂ on the symbiotic properties of this strain. This was especially visible during the first week of the experiment, when MQ caused a significant reduction (to 40 %) of nodules induced by these bacteria on clover roots. The *pssA* and *rosR* mutants are strongly impaired in symbiosis. All nodules induced on clover plants by the *pssA* mutant are small, white, and non-functional. In addition, the *rosR* mutant elicits formation of nodules with significantly less effectiveness than those induced by the wild-type strain. In the case of these mutants, prooxidant-treated bacteria induced a decreased number of nodules than untreated cells. The shoot mass of host plants reflects the efficiency of the nitrogen-fixation process. The shoot weight in the plants infected by H₂O₂-treated Rt24.2 was significantly lower than in those infected by untreated bacteria. Moreover, MQ had even a stronger effect on this symbiotic property of the Rt24.2, resulting in a decrease in the plant shoot mass to 54.3 % of the control. For both the Rt5819 and Rt2472 mutants, the further reduction of the plant shoot mass in the presence of the prooxidants was very low, since the shoot masses of plants infected by these mutants untreated with the stressors were only moderately higher than those of uninfected plants.

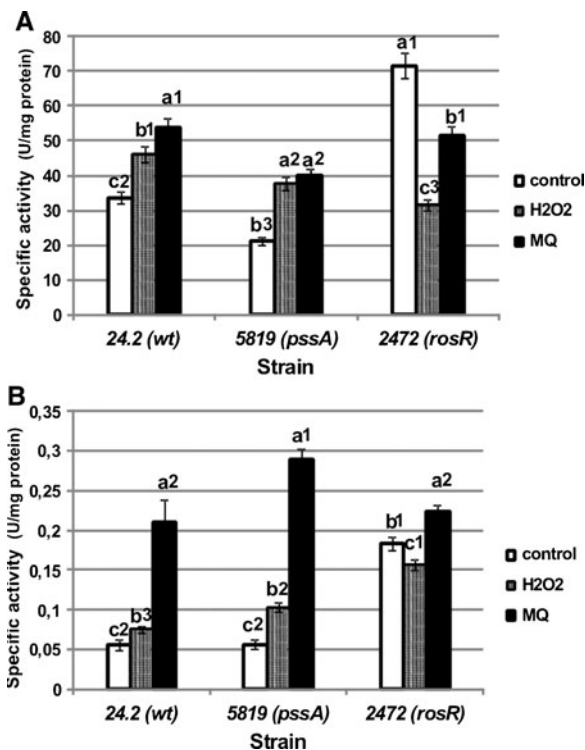


Fig. 7 The effect of prooxidants on extracellular PEC (a) and β -GLU (b) specific activity in the *R. leguminosarum* bv. *trifolii* wild-type and *pssA* and *rosR* mutants. Data are mean \pm SD for three measurements. Differences statistically significant ($p < 0.05$; ANOVA, post hoc Tukey's test) are marked by different letters (for particular strain) and different numbers (for particular treatment)

In summary, our data indicate that both stress factors affect the first step of symbiosis (induction of nodule formation on the host roots), that, in consequence, leads to a decrease in symbiotic efficiency.

Discussion

The oxidative stress phenomenon, i.e. the subject of the present work, very often appears to be a consequence of reactions triggered by prooxidative targets such as menadione, paraquat, metal ions and others, when the levels of produced ROS increase beyond the antioxidant capacity of cells (Jamieson 1998; Corticeiro et al. 2006). Since the physiological life cycle of rhizobia during the legume-rhizobia symbiosis is strictly correlated with the relatively high concentration of reactive oxygen intermediates such as SOR, hydroxyl radicals, and hydrogen peroxide, it is evidenced that

these organisms must evolve efficient anti-stress mechanisms (H rouart et al. 2002; Saenkham et al. 2007).

The classic bacterial ROS defence arsenal comprises a complex of enzymatic and non-enzymatic antioxidants. Bacteria are able to respond directly to the increased level of ROS by enhancing the expression of ROS decaying enzymes (catalases, superoxide dismutases and peroxidases) (Dombrecht et al. 2005).

The generation and accumulation of several ROS is a first plant defence reaction that blocks pathogen attack, but Jamet et al. (2007) confirmed that a suitable amount of H₂O₂ is required for establishment of the effective *Medicago sativa*–*S. meliloti* symbiosis. An increase in the concentration of free oxygen radical derivatives was also indicated during initial stages of *R. leguminosarum* bv. *trifolii* symbiosis with clover (Kopcińska 2009).

We observed several differences in enzymatic activity patterns of the *R. leguminosarum* bv. *trifolii* wild-type strain and two *pssA* and *rosR* mutants in relation to oxidative stress response. Although the antioxidative mechanisms in other groups of organisms have been often described, the profile of responses to oxidative stress is still poorly understood in rhizobia. The study reported here is probably the first attempt to demonstrate the response of *R. leguminosarum* bv. *trifolii* to chemically caused oxidative stress and a role of EPS in this process based on the comparison of defence parameters between the wild-type and the *pssA* and *rosR* mutants. The genetically modified bacterial strains used in this work had been well characterized in an earlier report with respect to EPS synthesis, symbiotic properties, and response to osmotic stress (Janczarek et al. 2009a; 2010).

The implementation of the experimental program presented in this study has provided interesting data that can be analysed in two ways: estimation whether the changes in chosen biochemical parameters depend on the genetic modification of rhizobial strains and investigation of the response of these bacteria to the presence of prooxidants. The comparison of CAT, SOD, PEC, and β -GLU activities noted in the control variants (without addition of prooxidants) of the wild-type and the *pssA* and *rosR* mutants showed that the values of specific CAT and extra- and intracellular SOD activities were significantly higher in the mutants than in the wild-type strain. In the case of PEC and β -GLU activity, only the *rosR* mutant displayed a higher activity in comparison to the wild-type. These data suggest

Table 4 Symbiotic properties of the *R. leguminosarum* bv. *trifolii* wild-type strain 24.2 and its derivatives in the presence of oxidative stress factors

Clover plants infected by strains	Nodule no. per plant* (days after infection)				Fresh shoot weight (mg plant ⁻¹)				Fresh root weight (mg plant ⁻¹)			
	7	14	21	28	7	14	21	28	7	14	21	28
Rt24.2	1.2±0.6 ^{a1} (100 %)	4.5±1.9 ^{a1} (100 %)	8.7±3.0 ^{a1} (100 %)	12.1±3.2 ^{a1} (100 %)	54.1±9.8 ^{a1} (100 %)	42.2±7.8 ^{a1} (100 %)						
Rt24.2 1.5 mM H ₂ O ₂	0.6±0.4 ^{a1} (50 %)	3.1±1.5 ^{a1} (69 %)	6.6±2.3 ^{a1} (75.9 %)	10.1±2.8 ^{a1} (83.5 %)	44.3±7.6 ^{a1} (81.88 %)	48.0±8.9 ^{a1} (113.7 %)						
Rt24.2 0.3 mM MQ	0.5±0.3 ^{a1} (41.7 %)	3.0±1.7 ^{a1} (66.7 %)	6.7±2.1 ^{a1} (77 %)	9.0±1.9 ^{a1} (74.4 %)	31.0±5.2 ^{b1} (57.3 %)	45.1±6.6 ^{a1} (106.8 %)						
Rt5819 [#]	0.3±0.1 ^{a2} (100 %)	3.4±2.2 ^a (100 %)	5.6±2.2 ^{a1} (100 %)	6.2±1.8 ^{a2} (100 %)	33.3±6.1 ^{a2} (100 %)	36.6±5.7 ^{a1} (100 %)						
Rt5819 [#] 1.5 mM H ₂ O ₂	0.1±0.1 ^{a2} (66.7 %)	1.4±1.0 ^{a1} (41.2 %)	3.6±1.7 ^{a1} (64.3 %)	5.6±1.4 ^{a2} (90.3 %)	31.2±6.3 ^{a1} (93.7 %)	39.1±6.3 ^{a1} (106.8 %)						
Rt5819 [#] 0.3 mM MQ	0.0±0.0 ^{a2} (0 %)	1.6±1.2 ^{a1} (47.0 %)	4.3±1.6 ^{a1} (76.8 %)	5.6±1.3 ^{a2} (90.3 %)	30.1±5.9 ^{a1} (90.4 %)	37.8±6.2 ^{a1} (103.3 %)						
Rt2472	0.7±0.4 ^{a2} (100 %)	2.9±1.6 ^{a1} (100 %)	5.0±1.8 ^{a1} (100 %)	6.9±2.3 ^{a2} (100 %)	44.6±6.1 ^{a2} (100 %)	34.6±6.9 ^{a1} (100 %)						
Rt2472 1.5 mM H ₂ O ₂	0.4±0.1 ^{a1} (57.1 %)	2.1±1.3 ^{a1} (72.4 %)	4.2±1.7 ^{a1} (84.0 %)	6.1±2.0 ^{a1} (88.4 %)	42.3±6.8 ^{a1} (94.8 %)	33.8±7.1 ^{a1} (97.7 %)						
Rt2472 0.3 mM MQ	0.2±0.1 ^{a1} (28.6 %)	1.7±1.2 ^{a1} (58.6 %)	3.0±1.4 ^{a1} (60.0 %)	5.2±1.9 ^{a2} (75.3 %)	40.1±8.8 ^{a1} (89.9 %)	37.3±5.9 ^{a1} (107.8 %)						
Uninoculated plants	–	–	–	–	32.7±3 ²	33.9±4 ¹						

* Plants were harvested 28 days after inoculation. Given values (± standard deviation) are averages of three independent experiments with 20 plants for each treatment

All nodules induced by the Rt5819 mutant on clover plants were small, white and ineffective in nitrogen fixation

^{a,b,c} Different letters indicate values within the column for the individual strain untreated and treated with prooxidants, which are significantly different ($p \leq 0.05$; ANOVA, post hoc Tukey's test)

^{1,2,3} Different numbers indicate significantly different values between Rt24.2, Rt5819 and Rt2472 strains in the particular treatment ($p \leq 0.05$; ANOVA, post hoc Tukey's test)

rebuilding of the metabolism of the mutants (e.g. an increase or decrease in some biochemical parameters) and a significant role of EPS in adaptation of rhizobia to stress conditions.

The wild-type bacteria were more resistant to the presence of prooxidants than the *rosR* and *pssA* mutants (Figs. 1 and 2). The addition of prooxidants to the culture of the investigated strains resulted in an increase of the ROS level: SOR (the product of menadione redox cycles) and H_2O_2 . The results show that the presence of MQ increased the relative level of extracellular SOR in comparison to the cultures of the non-treated wild-type strain (Table 3). The overproduction of SOR in the supernatants was strictly connected with high extracellular SOD activities (Fig. 5a). The presented data show that the mutation in both *pssA* and *rosR* genes significantly change the level of extra- and intracellular dismutase activity in the controls as well as in the stressed bacterial cultures. A high activity of SOD was noted in the *pssA* and *rosR* mutants and this observation was correlated with the high SOR level in these bacteria. The suitable level of SOD activities in rhizobial cells is necessary for their symbiotic properties. It was detected that the SOD-deficient mutants of *S. meliloti* did not reach the appropriate differentiation into nitrogen-fixing bacteroids (Santos et al. 2000). SOD catalyses the dismutation of SOR to hydrogen peroxide and oxygen (McCormick et al. 1998). In *S. meliloti*, a single superoxide dismutase was found in free-living growth conditions (Santos et al. 1999). In contrast, *Agrobacterium tumefaciens* possesses three SOD enzymes, among them one periplasmic and two cytoplasmic isozymes (Saenkham et al. 2007). A mutant lacking all three SODs is very sensitive to menadione treatment. In the cells of *R. leguminosarum* bv. *trifolii* wild-type and the *pssA* and *rosR* mutants untreated and treated with the MQ and H_2O_2 , we detected a single band corresponding to the SOD activity (Fig. 6). Moreover, an additional band was observed for the MQ-treated *rosR* mutant. Our data are in agreement with those described recently by Krehenbrink and co-workers (2011), who demonstrated that only one superoxide dismutase SodA was expressed at the stationary phase in *R. leguminosarum* bv. *viciae* 3841. This protein was detected in the periplasm, although it did not have a signal peptide. Bioinformatic analysis of genomic sequences of several *R. leguminosarum* strains revealed that bacteria from this species possess two distinct genes encoding SOD proteins. These two SOD proteins are probably induced in MQ-treated cells of the *rosR* mutant.

For detoxification of H_2O_2 , bacterial cells have evolved a specific mechanism based on catalase activity (Pauly et al. 2006). H_2O_2 is a signal molecule involved in plant response to pathogen attack and other stress conditions as well as nodulation (Rubio et al. 2004). It is also known that the proper level of H_2O_2 is required for optimal establishment of symbiosis. This confirms a phenotype of *S. meliloti* bacteria overexpressing the *katB* gene for catalase. The higher amount of catalase significantly decreased the level of H_2O_2 in comparison to the wild-type strain and induced formation of a lower number of nodules on *M. sativa* roots (Jamet et al. 2007). Our data indicated about a 2.5-fold higher CAT activity in both the *pssA* and *rosR* mutants in relation to the wild-type (Fig. 3a). The addition of the prooxidants increased the activity of CAT in the wild-type strain and in the *rosR* mutant, whereas this dependence was not observed in the *pssA* mutant. Moreover, cells of the wild-type and the *rosR* mutant treated with these prooxidants had significantly lower levels of intracellular H_2O_2 than the non-treated bacteria (Fig. 3b). The above results suggest that one of the consequences of metabolic changes caused by these mutations in *R. leguminosarum* bv. *trifolii* is a higher activity of the antioxidative SOD and CAT enzymes. *In silico* sequence analysis performed by us indicated that *R. leguminosarum* strains usually contain one housekeeping chromosomal gene and one to two genes located on plasmids that encode catalases. Jamet et al. (2003, 2005) have indicated that *S. meliloti* possesses three distinct catalases: two monofunctional KatA and KatC enzymes and one bifunctional catalase-peroxydase KatB. The *katB* gene is constitutively expressed during bacterial growth in the culture and is not expressed under oxidative stress conditions, whereas the expression of the *katA* and *katC* genes was induced in these conditions. A *katB katC* double mutant displays abnormal infection of *M. sativa* plants and elicits essentially fewer nodules than the wild-type (Jamet et al. 2003). We have detected two to three bands indicating the CAT activity in all of the tested *R. leguminosarum* bv. *trifolii* strains, both untreated and treated with the MQ and H_2O_2 (Fig. 4). The highest CAT activity was observed in the *rosR* mutant cells both untreated and treated with the prooxidants, suggesting the possibility of biotechnological application of this strain in several bioremediation processes (Nakayama et al. 2008). Moreover, the activity of an additional CAT in the H_2O_2 -induced cells of this strain has been detected. These data suggest the presence of the three CAT enzymes in *R. leguminosarum* bv. *trifolii*,

similarly as it was evidenced in *S. meliloti*, where three catalases were differently expressed under specific environmental conditions (Jamet et al. 2003). Yuan et al. (2005) have reported that phosphate limitation induced expression of the major H₂O₂-inducible catalase gene (*katA*) in this organism.

Based on the enzymatic assays, we detected significantly higher catalase activity in the *R. leguminosarum* bv. *trifolii* wild-type strain treated with H₂O₂ and especially with MQ in comparison to the untreated bacteria of this strain (Fig. 3a). However, these results did not correlate with the intensity of the CAT activity obtained using native gel detection (Fig. 4). Also for the Rt2472 mutant, differences in the CAT activity for these two techniques were found. This phenomenon is difficult to explain clearly, but similar differences between native gel detection and spectrophotometric measurement of the CAT activity were presented in several reports concerning various organisms such as *Saccharomyces*, *Neurospora crassa*, *Aspergillus nidulans*, and maize (Gamero-Sandemetrio et al. 2013; Schliebs et al. 2006; Wadsworth and Scandalios 1990; Kawasaki and Aguirre 2001). Moreover, Gamero-Sandemetrio et al. (2013) observed similar discrepancy between zymogram and enzymatic assays for SOD activity in several *S. cerevisiae* strains, indicating that native gel detection cannot always be considered as a reliable quantitative technique.

Also, we have observed that pre-incubation of *R. leguminosarum* bv. *trifolii* strains with the prooxidants (especially with the MQ) negatively affected the number of nodules elicited on clover plants. This effect was the most intensively visible during the first 14 days of the experiment (Table 4). In addition, these stress factors influenced the mass of shoot plants. Previously, it was observed that the *pssA* mutant totally defective in EPS synthesis induced almost empty nodules inefficient in nitrogen fixation. Additionally, the *rosR* mutant produced 3-fold less EPS and elicited significantly fewer nodules, which were occupied by ten times fewer bacteria than the wild-type strain (Janczarek et al. 2009a, 2010). The part-after-part estimation of the several enzymatic as well as non-enzymatic biochemical and physiological features of the wild-type and the *pssA* and *rosR* mutants cultured under oxidative stress conditions confirmed the crucial role of these genes involved in EPS synthesis in oxidative stress defence mechanisms. It was observed that the changes in amounts of EPS in rhizobial strains exerted a negative effect on their

oxidative stress response and symbiotic efficiency. The results obtained suggest that the lack of or a significantly decreased level of EPS in the *pssA* and *rosR* mutants, respectively, resulted in overproduction of enzymatic antioxidants in order to compensate for the lack of appropriate amounts of EPS, which has a significant protective role against environmental stress conditions. Unfortunately, the high activity of SOD and CAT enzymes resulting in excessive elimination of SOR and H₂O₂ is most probably the cause of low effectiveness of symbiosis in the wild-type strain in these stress conditions. Recently, Davies and Walker (2007) identified novel *S. meliloti* mutants compromised for oxidative stress protection and symbiosis, which were sensitive to H₂O₂ and symbiotically defective on *M. sativa*. These mutations affected a wide variety of cellular processes, including succinoglycan synthesis. Similarly, D’Haeze and others (2004) indicated that ORS571-X15 mutant of *Azorhizobium caulinodans* producing significantly less EPS than the wild-type strain was more sensitive to H₂O₂ and elicited only pseudo-nodules on its host plant, *Sesbania rostrata*. Their studies demonstrated a negative correlation between the amount of H₂O₂ present within bacteria and the quantity of produced EPS. A protective role of EPS was also shown in *ex planta* experiments. Therefore, the authors proposed that this polymer forms a diffusion barrier to protect rhizobial cells against huge amounts of H₂O₂ encountered during early stages of legume infection and nodule development (D’Haeze et al. 2004). Additionally, the data presented in this work indicate that EPS added to cultures of the *pssA* and *rosR* mutants improves survival of these bacteria in the presence of the prooxidants to levels near to those observed for the wild-type bacteria (Table 2). This suggests that although mutations in both *rosR* and *pssA* genes have pleiotropic effects in *R. leguminosarum* bv. *trifolii*, the lack of EPS surrounding rhizobial cells seems to be a main cause of decreased survival and adaptation ability of these bacteria to stress conditions.

A very important phase in the infection of clover roots by *R. leguminosarum* bv. *trifolii* leading to effective nodule formation is movement of bacteria inside root hairs (Mateos et al. 1992). This step requires degradation of polysaccharides building the root cell walls by hydrolytic enzymes such as PEC or/and β -GLU. Mateos and co-workers (1992) confirmed the presence of cell-associated pectinolytic and cellulolytic enzymes in *R. leguminosarum* bv. *trifolii*. We have observed a stimulation effect of the prooxidants on the extracellular

PEC and β -GLU activities for the wild-type *R. leguminosarum* bv. *trifolii* as well as the *pssA* and *rosR* mutants. The increased amounts of the described enzymes detected under oxidative stress conditions can be interesting from the biotechnological point of view. Special attention is focused on the enzymatic pattern of the *rosR* mutant, which differs significantly from those of the wild-type and *pssA* mutant strains (Fig. 7). Previously, we have established that the transcriptional regulator RosR of *R. leguminosarum* bv. *trifolii* positively regulates the *pssA* expression (Janczarek and Skorupska 2007; Janczarek and Urbanik-Sypniewska 2013). This gene encodes a protein, which besides its enzymatic activity is probably involved additionally in other cellular processes (Janczarek and Skorupska 2003). Bittinger and Handelsman (2000) have reported that RosR of *Rhizobium etli* regulates expression of many genes, including those involved in polysaccharide production, carbohydrate metabolism, and plant infection.

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