

Nitrogen source influences the antioxidative system of soybean plants under hypoxia and re-oxygenation

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ABSTRACT: In this work, we compared nitrate-supplied plants (non-nodulated) with non-nitrate-supplied plants (nodulated) under oxygen privation of root system (hypoxia) and re-oxygenation (post-hypoxia; recovery) in order to verify whether N sources affect the antioxidant system during oxidative stress caused by hypoxia and post-hypoxia conditions. Antioxidant enzymatic activities, ascorbate redox state, and reactive oxygen species (ROS) levels were analyzed in roots and leaves of two soybean genotypes, Fundacep 53 RR and BRS Macota at reproductive stage R2, during hypoxia and post-hypoxia in an experiment carried out in a hydroponic system. The antioxidant system was strongly induced in roots of nitrate-supplied plants of both genotypes, with high activity of superoxide dismutase, ascorbate peroxidase, catalase, glutathione reductase and guaiacol peroxidase. It also increased reduced ascorbate and ascorbate redox state and decreased ROS production under hypoxia and recovery, while in leaves of nodulated and non-nodulated plants, a slight increase on antioxidant system was observed. Nitrate may benefit soybean plants under hypoxic conditions and subsequent re-oxygenation by inducing the antioxidant system mainly in roots to cope with ROS production and reduce oxidative damage.

Keywords: *Glycine max*, waterlogging, oxidative stress, antioxidative enzymes

Introduction

Waterlogging, or flooding, causes oxygen deprivation to plant roots (Limami et al., 2014) and leads to inhibition of mitochondrial oxidative phosphorylation and decrease in ATP production (van Dongen and Licausi, 2015). Besides, hypoxic conditions, mainly re-oxygenation of cells, promote a redox imbalance of mitochondria and chloroplast components leading to overreduction of electron carriers and electron leaking, producing reactive oxygen species (ROS) (Halliwell, 2006).

Several studies have reported that nitrate as N source exerts an important role in many plant species under hypoxia, such as soybean (Oliveira et al., 2013a, b), tobacco (Stoimenova et al., 2003) and tomato (Allègre et al., 2004) by reducing fermentation effects (lactate and ethanol accumulation) (Oliveira et al., 2013a, b) when compared to N₂-fixing or NH₄ supplied plants. The beneficial effects of nitrate have been attributed to nitric oxide (NO) production via the nitrate reductase activity in cytosol (van Dongen and Licausi, 2015) and within mitochondria via cytochrome *c* oxidase (COX) (Gupta and Igamberdiev, 2011).

Most studies have shown beneficial effects of nitrate on N and C metabolism in waterlogged plants and that hypoxia followed by re-oxygenation promotes ROS production. Therefore, nitrate is also supposed to be beneficial by alleviating the effects of ROS production through the induction of the antioxidative activity of enzymes to reduce oxidative damage. Among the antioxidative enzymes, superoxide dismutase (SOD) plays a key role in scavenging superoxide radical (O₂^{•-}) anion into hydrogen peroxide (H₂O₂) (Simova-Stoilova et al., 2012).

Further, H₂O₂ is broken into water and dioxygen by catalase (CAT), guaiacol peroxidase (GPOD) or ascorbate peroxidase (APX) (Gill and Tuteja, 2010). Non-enzymatic antioxidants, which are generally small molecules such as ascorbate (AsA), also contribute to the destruction of H₂O₂, along with glutathione and glutathione reductase (GR) enzyme via the ascorbate-glutathione cycle (Noctor et al., 1998).

Our previous work showed differences between Fundacep (tolerant) and Macota (susceptible) regarding N and C metabolism (Borella et al., 2017). Therefore, due to the lack of information regarding nitrate effects on the antioxidant status of waterlogged root system of soybean, this work investigated whether different N sources (nodulated and nitrate-supplied plants - non-nodulated) influence the antioxidant system of soybean plants by alleviating the oxidative effects caused by hypoxia.

Materials and Methods

Plant material and growth conditions

The study was carried out at Federal University of Pelotas – *Campus* Capão do Leão (31° 52' 32" S and 52° 21' 24" W; altitude: 16 m), with two soybean [*Glycine max* (L.) Merrill] genotypes, Fundacep 53 RR and BRS Macota, identified previously to respond distinctly to hypoxia (Borella et al., 2014, 2017). The experiment comprised two groups, nodulated (N₂-fixing) and non-nodulated (nitrate supplied) plants, grown in a greenhouse under natural light and temperature conditions. Plants were cultivated in 3 L pots (3 plants per pot) in vermiculite and supplied twice a week with 250 mL

of nutrient solution according to the group: NO_3^- as N source for non-nodulated plants and N-free nutrient solution for nodulated (N_2 -fixing), as described previously by Hoagland and Arnon (1950). Nodulated plants were inoculated when cotyledons were fully open by applying 2.5 mL of liquid medium containing 10^9 cells mL^{-1} of *Bradyrhizobium elkanii* strain SEMIA 587 (FEPAGRO) around the stem of each plant on two occasions at 3-d intervals. Hypoxic treatments were initiated with plants at reproductive stage-R2 (Fehr et al., 1971).

For the hydroponic treatment, plants were removed from pots and the root system was carefully washed in tap water to remove vermiculite before transferring to 3 L pots (3 plants per pot) containing N-free nutrient solution (for nodulated plants) or N (NO_3^- 5 mM) nutrient solution (for non-nodulated), both at one-third of normal strength of the Hoagland's solution. The whole root system was kept submerged in the nutrient solution and subjected to hypoxia by flushing N_2 gas for 24 and 72 h. Oxygen concentration in the solution was monitored with an oxygen meter (Handylab OX1). For recovery, after 72 h of hypoxia, plants were returned to 3 L pots containing vermiculite as substrate under normoxic conditions for 24 and 72 h. At harvest, four biological replicates of roots and leaves were removed from each treatment and kept frozen (-80°C) until analysis.

Enzymatic activity assays

To measure the enzymatic activities, leaves and roots (± 0.2 g) were ground using liquid N_2 in porcelain mortars, containing 5 % (w:v) polyvinylpyrrolidone (PVPP) and homogenized in 100 mM potassium phosphate buffer, pH 7.8, containing 0.1 mM ethylenediaminetetraacetic acid (EDTA) and 20 mM sodium ascorbate. The homogenate was centrifuged at 12,000 g for 20 min and the supernatant obtained was used as crude enzyme extract. All steps in the preparation of the enzyme extract were carried out at 4°C . An aliquot of the extract was used to determine protein content as described by Bradford (1976) using bovine serum albumin as standard.

The superoxide dismutase activity (SOD; EC 1.15.1.1) was tested as described by Giannopolitis and Ries (1977) by monitoring the inhibition of the nitroblue-tetrazolium (NBT) coloration at 560 nm. The catalase activity (CAT; EC 1.11.1.6) was determined according to the method described in Azevedo Neto et al. (2006) by monitoring hydrogen peroxide consumption measuring decline in absorbance at 240 nm. The ascorbate peroxidase activity (APX; EC 1.11.1.11) was determined according to the method described by Nakano and Asada (1981) measuring the rate of ascorbate oxidation at 290 nm. The glutathione reductase activity (GR; EC 1.6.4.2) was measured according to Cakmak et al. (1993) by following the decrease in absorbance at 340 nm due to NADPH oxidation. The guaiacol peroxidase activity (GPOD; EC 1.11.1.7) was analyzed following the method described by Urbanek et al. (1991) by monitor-

ing the tetraguaiacol production by reducing hydrogen peroxide at 470 nm. The glutathione S-transferase activity (GST; EC 2.5.1.13) was performed as described by Dalton et al. (2009) by measuring absorbance at 340 nm using 2,4-dinitrochlorobenzene as substrate.

Ascorbate content

The contents of reduced ascorbate (AsA) and total ascorbate [(AsA + oxidized ascorbate (DHA))] were quantified as described by Arakawa et al. (1981). Tissues (0.2 g) were ground in 5 % trichloroacetic acid (TCA), homogenized and centrifuged at 10,000 g for 15 min at 4°C . Total ascorbate from supernatant was determined after reduction of DHA by dithiothreitol (DTT). The reaction medium consisted of 5 % TCA, 0.06 % DTT and 0.2 M sodium phosphate buffer, pH 7.0. After incubation at room temperature for 10 min, 0.24 % N-ethylmaleimide was added and the pH of each tube adjusted to between 1 and 2 with 20 % TCA. Afterwards, phosphoric acid 4 % (H_3PO_4), bathophenanthroline 0.5 % and ferric chloride 0.03 % (FeCl_3) were added and incubated for 90 min at 30°C . The absorbance was read at 534 nm. The ascorbate was determined as described above, but replacing the DTT for absolute ethanol in equal volume. The DHA values were obtained by the difference between the values of total ascorbate and reduced ascorbate. The ascorbate redox state was calculated as [(AsA)/(AsA + DHA)] $\times 100$ and expressed as percentage of reduced ascorbate from the total ascorbate (Bonifacio et al., 2011).

$\text{O}_2^{\cdot-}$ content

The assay of $\text{O}_2^{\cdot-}$ content was determined according to Li et al. (2010). The tissues (0.2 g) were ground in 65 mM phosphate buffer, pH 7.8, and centrifuged at 5,000 g for 10 min. The supernatant was mixed with 65 mM phosphate buffer, pH 7.8, and 10 mM hydroxylamine hydrochloride, and incubated at 25°C for 20 min. Then, sulphanilamide and α -naphthylamine were added to the mixture, resulting in a final concentration of 17 mM and 7 mM, respectively. The solution absorbance at 530 nm was measured after incubation for 20 min at 25°C . A standard curve with nitrite (NO_2^-) was used to calculate $\text{O}_2^{\cdot-}$ generation rate.

H_2O_2 content and lipid peroxidation measurement

Hydrogen peroxide levels and lipid peroxidation were determined according to Velikova et al. (2000). The tissues (0.2 g) were ground in 0.1 % (w:v) trichloroacetic acid (TCA). The homogenate was centrifuged (12,000 g, 4°C , 20 min) and the supernatant was used for the analyses. An aliquot of the supernatant was added to 10 mM potassium phosphate buffer, pH 7.0 and 1 M potassium iodide to determine H_2O_2 . Reaction absorbance was measured at 390 nm. The H_2O_2 content was given on a standard curve prepared with known H_2O_2 concentrations. Lipid peroxidation was determined by using thiobarbituric acid (TBA), which determines malondial-

dehyde (MDA) as a product of lipid peroxidation. The supernatant was added to 0.5 % (w:v) TBA in 10 % TCA solution. The mixture was incubated in boiling water (90 °C) for 20 min; afterwards, the reaction was stopped in an ice bath for 10 min. Then, the samples were centrifuged at 10,000 g for 5 min and absorbance was read at 535 nm. The value for non-specific absorption at 600 nm was subtracted. The amount of MDA-TBA complex was calculated from the extinction coefficient ($\epsilon = 155 \times 10^3 \text{ M}^{-1} \text{ cm}^{-1}$).

Statistical analysis

Each treatment consisted of four replicates and each replicate consisted of one pot containing three plants (material pooled), in a completely randomized design. The data were analyzed by one-way analysis of variance (ANOVA). When *F* was significant, the means of O₂ availability treatments (normoxia, hypoxia, and recovery) were compared for each N assimilation system (nodulated or non-nodulated plants) or the N assimilation systems were compared for each O₂ availability treatment by the Tukey's test ($p \leq 0.05$). The statistical

analyses were performed using the SAS 8.0 statistical software program (Statistical Analysis System, v. 8.0).

Results

Oxygen concentration in the solution of both nodulated and non-nodulated plants was about 6.5 mg L⁻¹ under normoxia. Under hypoxia, oxygen concentration decreased rapidly to 0.5 mg L⁻¹ within 5 h, reaching 0.25 mg L⁻¹ in 24 h until the end of the experiment (72 h) (data not shown).

Antioxidant enzymatic activity

The antioxidant enzymatic system of plants subjected to hypoxia and subsequent re-oxygenation are shown in roots (Figures 1A, B, C, D, E and F and 3A, B, C, D, E and F) and leaves (Figures 2A, B, C, D, E and F and 4A, B, C and D) of nodulated (N₂-fixing) and non-nodulated (nitrate-supplied) soybean plants of two genotypes, Fundacep 53 RR and BRS Macota. The enzymatic activity increased significantly in roots during hypoxia and recovery in non-nodulated plants. SOD and

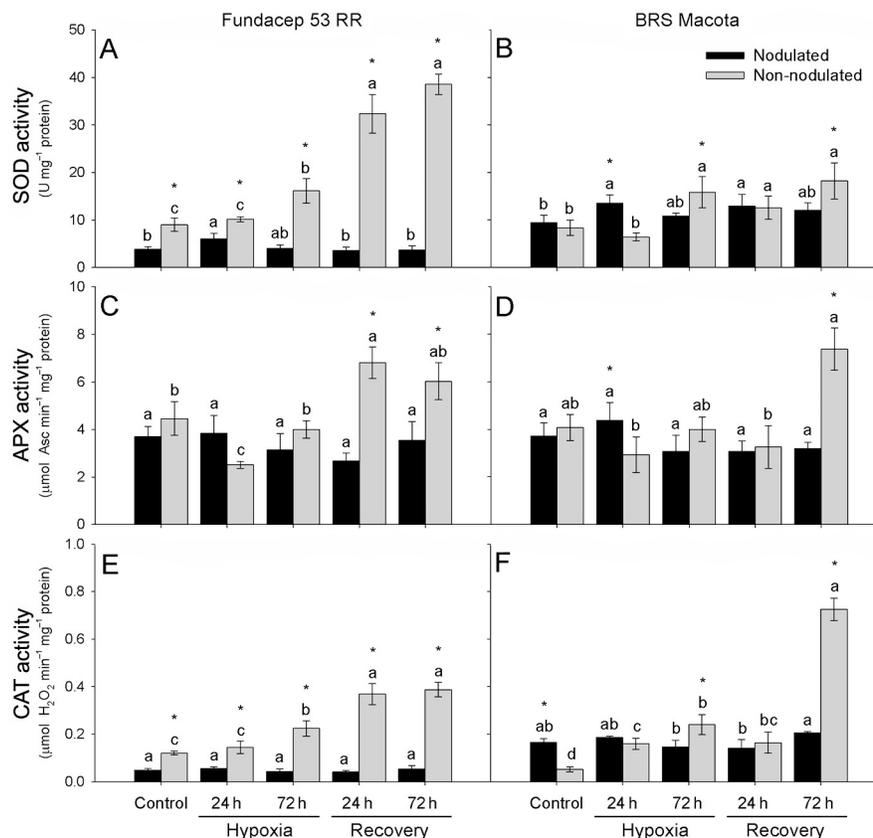


Figure 1 – Superoxide dismutase (SOD – A and B), ascorbate peroxidase (APX – C and D) and catalase (CAT – E and F) activity in roots of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey's test ($p \leq 0.05$) among O₂ availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey's test ($p \leq 0.05$) between nodulated and non-nodulated for each O₂ availability treatment. Values represent the mean \pm SD ($n = 4$).

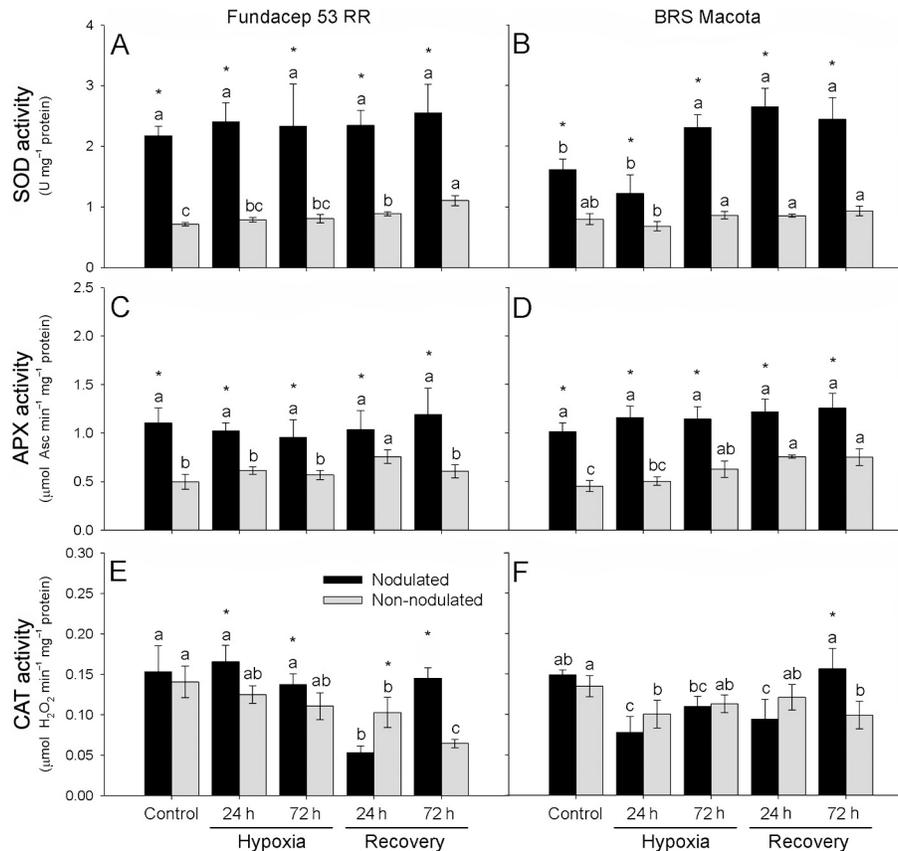


Figure 2 – Superoxide dismutase (SOD – A and B), ascorbate peroxidase (APX – C and D) and catalase (CAT – E and F) activity in leaves of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey's test ($p \leq 0.05$) among O_2 availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey's test ($p \leq 0.05$) between nodulated and non-nodulated for each O_2 availability treatment. Values represent the mean \pm SD ($n = 4$).

CAT increased in roots of both waterlogged non-nodulated soybean genotypes at 72 h and were higher than in the control during recovery with strong effect on Fundacep 53 RR. In nodulated soybean plants, an increase in SOD activity was observed at 24 h of hypoxia in both genotypes and remained until the end of the experiment in BRS Macota. In Fundacep 53 RR, the activity decreased with recovery. The CAT activity did not differ from that in the control during the entire experiment in both genotypes. Plants supplied with nitrate increased SOD and CAT activities by about 3-fold compared to nodulated plants in both genotypes (Figures 1A and B).

APX activity only increased upon return to normoxia in nitrate-supplied plants while did not differ in nodulated plants in both genotypes (Figures 1C and D). Interestingly, the APX activity was responsive during recovery with higher increase in the activity in nitrate-supplied plants and a faster increase in Fundacep 53 RR than BRS Macota (Figures 1C and D).

The enzymatic activities were different in leaves compared to roots under normoxic conditions. Despite increased SOD and APX activity with recovery in plants

supplied with nitrate (Figures 2A-D), the activity of these enzymes did not change in nodulated plants, except for the SOD activity in BRS Macota at 72 h of hypoxia (Figure 2B), remaining higher even during recovery. In addition, SOD and APX activities were higher in nodulated plants than in nitrate-supplied plants (non-nodulated) in comparison with their activities in roots of both plant groups (Figures 1A-D). On the other hand, the CAT activity appears to be non-responsive in leaves (Figures 2E and F) as it is in roots.

In addition to SOD, APX and CAT (Figures 1A, B, C, D, E, F and 2A, B, C, D, E and F), GR and GPOD activities increased in roots similarly to CAT with 72 h of hypoxia, although they were higher during recovery, in nitrate-supplied plants (Figures 3A and B), with a pronounced increase of GR in Fundacep 53 RR. The GR activity did not increase in nodulated plants. In leaves, GR was more active during hypoxia with increased activity at 72 h in leaves of both genotypes in nitrate-supplied plants, while in nodulated plants, the activities were similar to those in the control (Figures 4A and B). In contrast, GPOD was found as the most active peroxidase

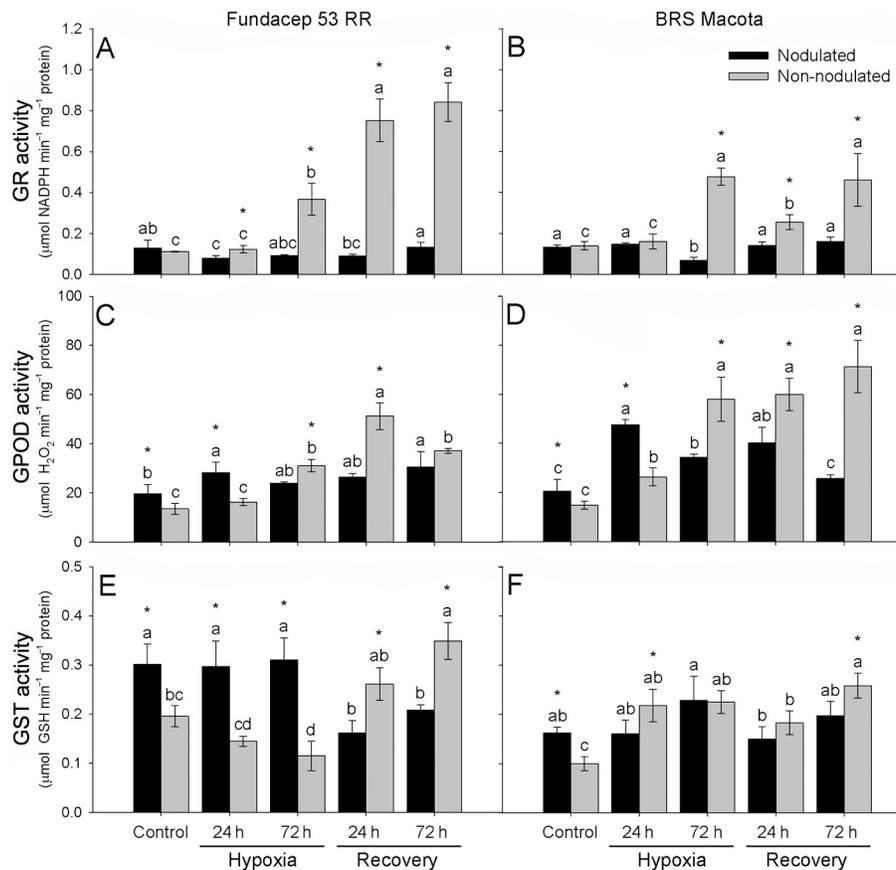


Figure 3 – Glutathione reductase (GR – A and B), guayacol peroxidase (GPOD – C and D) and glutathione S-transferase (GST – E and F) activity in roots of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey's test ($p \leq 0.05$) among O_2 availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey's test ($p \leq 0.05$) between nodulated and non-nodulated for each O_2 availability treatment. Values represent the mean \pm SD ($n = 4$).

in roots under hypoxia (Figures 3C and D), whereas no activity of this enzyme was detected in leaves.

The GST activity changed significantly in roots during hypoxia in non-nodulated plants, decreased in Fundacep 53 RR and increased in re-oxygenation. In BRS Macota, the GST activity increased during hypoxia and was kept at high levels during recovery (Figures 3E and F). In leaves of non-nodulated plants, the GST activity increased markedly during hypoxia in both genotypes and was kept at higher levels during recovery (Figures 4C and D). In nodulated plants, GST was more active during hypoxia in roots and leaves (Figures 4C and D) in both genotypes.

Ascorbate redox state

In the most treatments, reduced ascorbate was higher in nitrate-supplied plants and in both genotypes. The content was higher during recovery than during hypoxia in both, roots (Figures 5A-D) and leaves (Figures 6A-D). On the other hand, increased ascorbate redox state was higher during recovery only in non-nodulated

plants in roots and leaves, and these responses were greater in BRS Macota genotype.

Oxidative damage

Superoxide, hydrogen peroxide and lipid peroxidation in roots and leaves are shown in Figures 7A, B, C, D, E, F and 8A, B, C, D, E and F, respectively. In roots of nitrate-supplied plants with 24 h of hypoxia, the content of superoxide remained low, similar to the control, and later reduced at 72 h of hypoxia in both genotypes. In turn, during recovery in Fundacep 53 RR, the content was below the control levels (Figures 7A and B), reflecting the SOD activity in roots (Figures 1A and B). In nodulated plants, the production of superoxide decreased during hypoxia and increased with the return to normoxic conditions (Figures 7A and B).

The hydrogen peroxide content was lower than in the control even under hypoxia and recovery in roots of nitrate-supplied plants than in nodulated plants (Figures 7C and D), probably due to the activity of enzymes that are responsive to scavenging. The level of lipid peroxida-

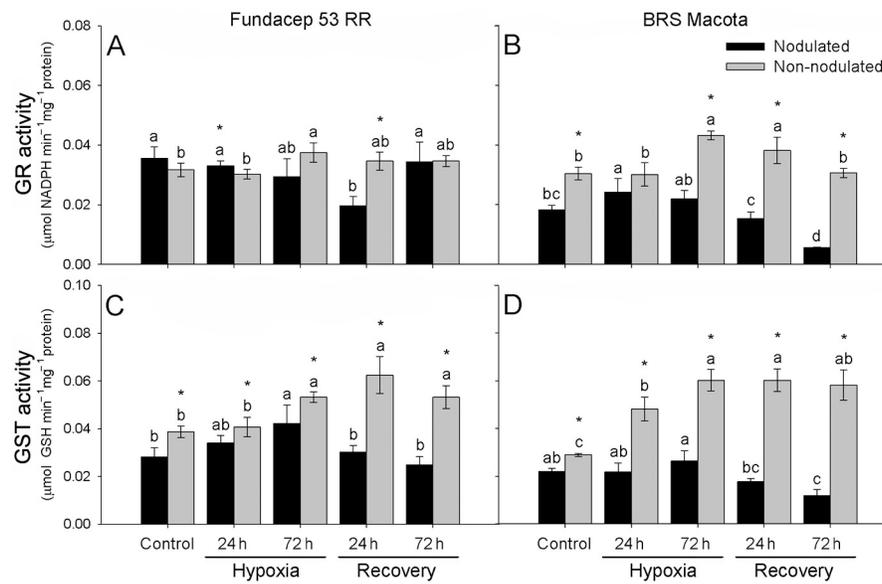


Figure 4 – Glutathione reductase (GR – A and B), glutathione S-transferase (GST – C and D) activity in leaves of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey's test ($p \leq 0.05$) among O_2 availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey's test ($p \leq 0.05$) between nodulated and non-nodulated for each O_2 availability treatment. Values represent the mean \pm SD ($n = 4$).

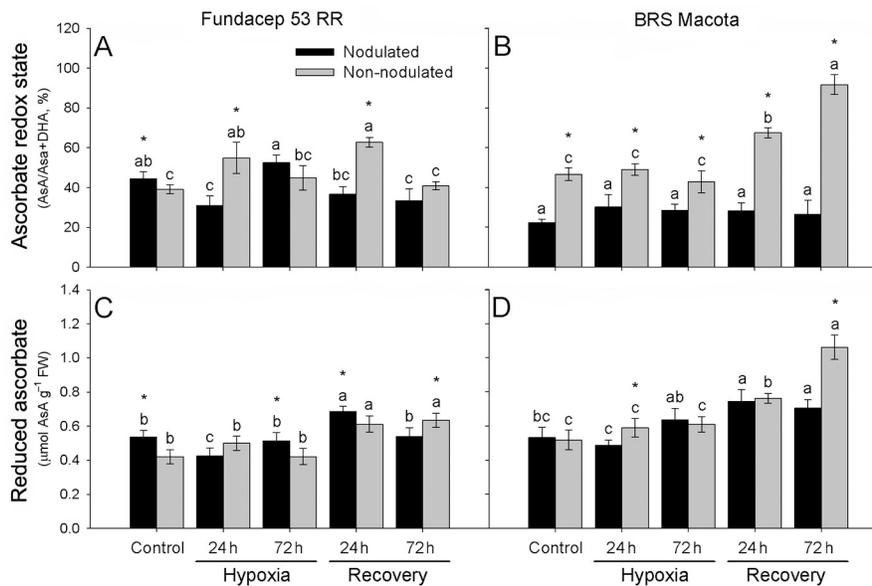


Figure 5 – Ascorbate redox state (A and B) and reduced ascorbate content (AsA – C and D) in roots of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey's test ($p \leq 0.05$) among O_2 availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey's test ($p \leq 0.05$) between nodulated and non-nodulated for each O_2 availability treatment. Values represent the mean \pm SD ($n = 4$).

tion decreased in roots of both genotypes during hypoxia and recovery. However, the level modulation was higher in nitrate-supplied plants in which the levels did not increase as much as in control after hypoxia in roots of nodulated plants in both genotypes (Figures 7E and F).

In leaves, differently from roots, an increase in superoxide production was observed in nitrate-supplied plants during hypoxia and remained at higher levels than in control even at 72 h of recovery. In nodulated plants, the content did not change in Fundacep 53 RR, it

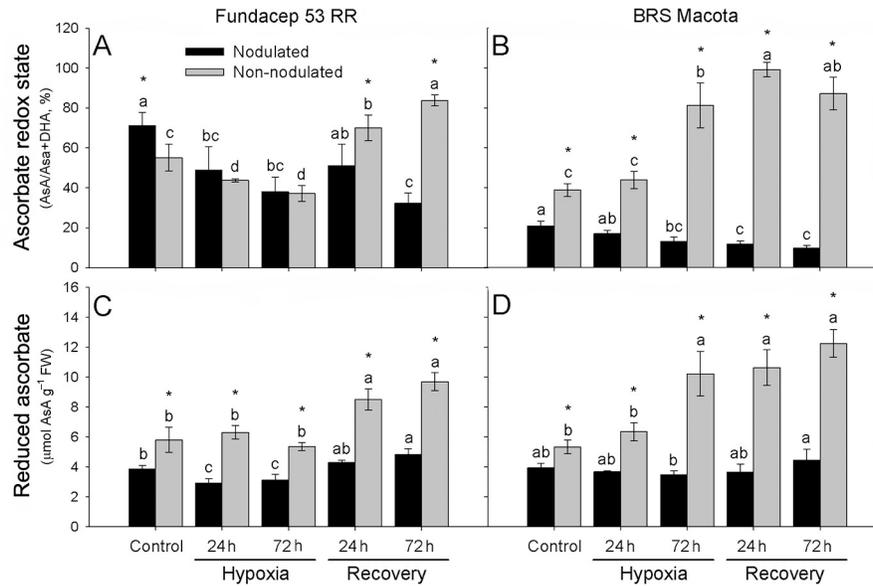


Figure 6 – Ascorbate redox state (A and B) and reduced ascorbate content (AsA – C and D) in leaves of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey’s test ($p \leq 0.05$) among O₂ availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey’s test ($p \leq 0.05$) between nodulated and non-nodulated for each O₂ availability treatment. Values represent the mean ± SD (n = 4).

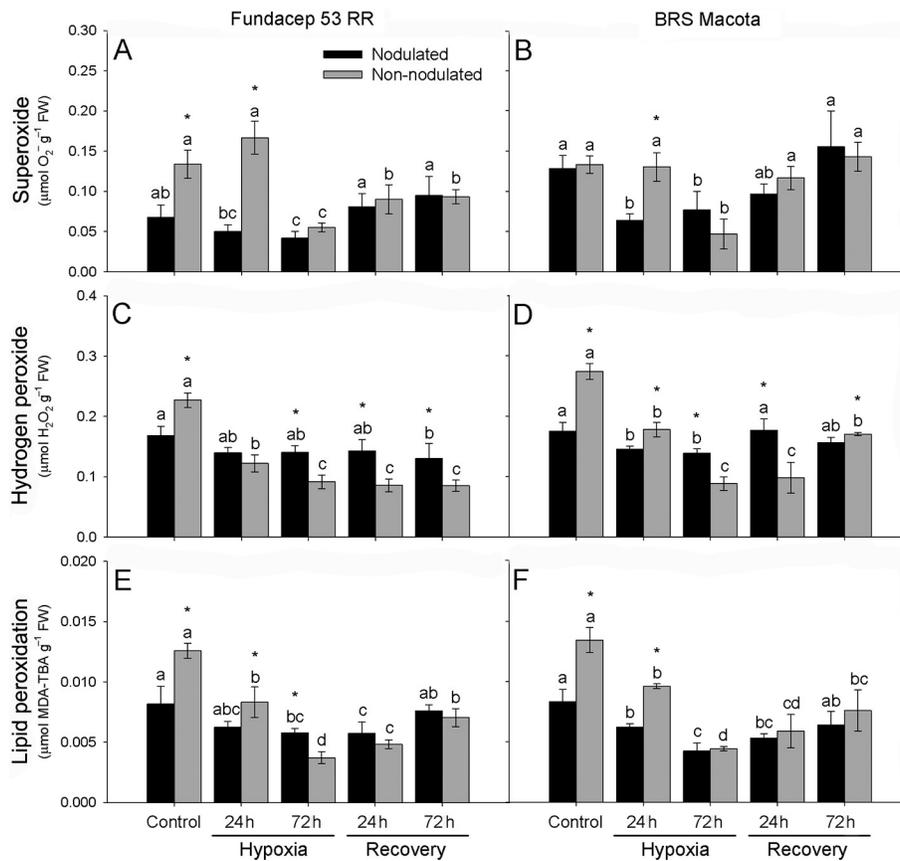


Figure 7 – Superoxide (A and B), hydrogen peroxide (C and D) content and lipid peroxidation (E and F) in roots of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey’s test ($p \leq 0.05$) among O₂ availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey’s test ($p \leq 0.05$) between nodulated and non-nodulated for each O₂ availability treatment. Values represent the mean ± SD (n = 4).

increased in BRS Macota during hypoxia and decreased in control after 72 h of recovery (Figures 8A and B). Hydrogen peroxide production was lower than in control at 24 h of hypoxia and increased later. During recovery, the content increased similarly to control in nodulated plants. In non-nodulated plants, no great changes were observed in hydrogen peroxide levels (Figures 8C and D). Lipid peroxidation did not alter significantly during hypoxia and post-hypoxia treatments in Fundacep 53 RR of nodulated plants. In BRS Macota, lipid peroxidation increased at 72 h of hypoxia and decreased to control levels during recovery. In non-nodulated plants, no great changes were observed in lipid peroxidation during hypoxia in comparison to normoxia in both genotypes. During post-hypoxia treatments, lipid peroxidation decreased in Fundacep 53 RR while it did not alter in BRS Macota (Figures 8E and F).

Discussion

In this work, we described the influence of nitrate on alleviating the effects of oxidative damage caused by

ROS via induction of enzymatic and non-enzymatic antioxidants in leaves and mainly in roots, during and after hypoxia of the root system of non-nodulated plants (nitrate-supplied plants) in comparison to nodulated plants (plants assimilating ammonium, via N_2 fixation) of two soybean genotypes, Fundacep 53 RR and BRS Macota. Nodulated plants were chosen rather than supplying them directly with ammonium, as ammonium is toxic and possibly influences antioxidant metabolism.

Since the study conducted by Arnon (1937), nitrate has been investigated and related to its beneficial effects in plants under oxygen deficiency (Horchani et al., 2010; Horchani et al., 2011). However, these studies have focused primarily on C and N metabolism (Horchani et al., 2010; Oliveira et al., 2013a, b; Oliveira and Sodek, 2013; Lanza et al., 2014). In agreement to experiments on nitrate benefits, a similar pattern was also observed in relation to oxidative metabolism under hypoxia and post-hypoxia conditions. In roots of non-nodulated plants, the induction of enzymes SOD, APX, CAT, GR and GPOD (Figures 1A, B, C, D, E, F and 3A, B, C, D, E and F) and non-enzymatic antioxi-

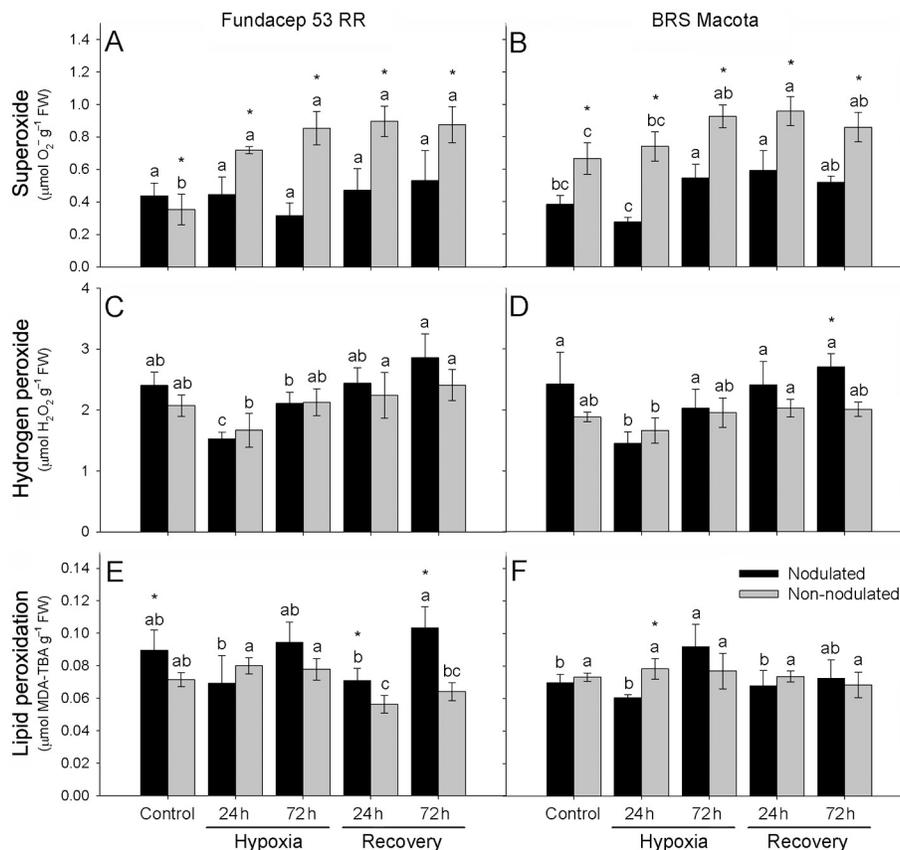


Figure 8 – Superoxide (A and B), hydrogen peroxide (C and D) content and lipid peroxidation (E and F) in leaves of two nodulated and non-nodulated soybean genotypes (Fundacep 53 RR and BRS Macota) under hypoxia and recovery conditions. Means followed by the same letters do not differ by the Tukey's test ($p \leq 0.05$) among O_2 availability treatments for nodulated or non-nodulated, separately. Asterisk (*) indicates significant differences by the Tukey's test ($p \leq 0.05$) between nodulated and non-nodulated for each O_2 availability treatment. Values represent the mean \pm SD ($n = 4$).

dants (Figures 5A, B, C and D), as well as their efficiency in scavenging ROS production (Figures 7A, B, C, D, E and F) was markedly more efficient in comparison to the roots of nodulated plants. The ROS content in control plants of non-nodulated one was much higher than that in nodulated plants. However, the hypoxia and recovery treatments induced decreases ROS content of non-nodulated plants similarly to the ROS content in nodulated plants. Besides that, increases in enzyme activity of non-nodulated plants was above the activity of nodulated ones. These results showed that the two forms of N assimilation, nitrate (non-nodulated) and N_2 -fixing (nodulated) in plants, influence differently the antioxidant defence system under stressful conditions caused by O_2 privation.

In leaves, despite the slight induction of the antioxidant system, an increase of both enzymatic and non-enzymatic antioxidants was observed (Figures 2A-F, 4A-D and 6A-D). These results may be attributed to the short duration of hypoxia (3 days) to which plants were submitted. In addition, leaves were kept under normoxic conditions that may have alleviated the effects of oxygen deprivation in comparison to roots, which were directly affected.

In non-nodulated plants, an increased induction of SOD activity was observed in roots under hypoxia. During recovery, induction of SOD activity was stronger, at least in Fundacep 53 RR, while in nodulated plants, the SOD activity appears to be induced only under hypoxic conditions (Figures 1A and B). These results agree with the reduction of $O_2^{\cdot-}$ anion production by SOD activity (Figures 7A and B). Furthermore, in non-nodulated roots of Fundacep 53 RR, $O_2^{\cdot-}$ production did not reach levels of the control during recovery (Figure 7A) due to high SOD activity (Figure 1A).

As reported, SOD constitutes the first line of defence against ROS, playing an important role in detoxification of $O_2^{\cdot-}$ into H_2O_2 (Gill and Tuteja, 2010). Although ROS production increased in several plant species under hypoxia (Bai et al., 2010; Simova-Stoilova et al., 2012), it also correlated to the period of hypoxic treatment (Blokhina and Fagerstedt, 2010b). On the other hand, ROS production reduces under hypoxia conditions due to oxygen deprivation and limited functioning of the electron transport chain (Sairam et al., 2011). Moreover, distinct SOD responses to oxygen deprivation stress (anoxia and hypoxia) on different plant species have always been contradictorily described, depending on experiment design or re-oxygenation (Blokhina et al., 2003). The decrease of both $O_2^{\cdot-}$ and H_2O_2 production under hypoxia were attributed to a shift from aerobic respiration to fermentation with blockage of the mitochondrial site of ROS production (Sairam et al., 2011).

In nitrate-supplied plants (soybean), fermentation has been reported to be modulated by nitric oxide (NO) production, which leads to a decrease in lactate and ethanol content (Oliveira et al., 2013a, b). NO

acts as an alternative pathway to recycle nicotinamide adenine dinucleotide (NAD^+) from its reduced form (NADH) under low-oxygen conditions via futile NO cycle, where nitrite reduction by nitrate reductase leads to NO production in the cytosol (Limami et al., 2014; van Dongen and Licausi, 2015).

Another pathway is the reduction of nitrite via cytochrome *c* oxidase (COX), linked to membrane proton translocation (Gupta et al., 2005; Gupta et al., 2011; Gupta and Igamberdiev, 2011; Oliveira et al., 2013b). NO is then oxidized to nitrate again by class-1 non-symbiotic haemoglobin (Igamberdiev and Hill, 2004), besides playing other roles in plants, such as in nodules (Sainz et al., 2015). NO has emerged as an important free radical signal in plants (Neill et al., 2008) and may affect the induction of the antioxidant system as observed in roots of nitrate-supplied plants, however, further studies are needed to clarify its effects.

ROS act in oxidative damage to membrane cells (Gill and Tuteja, 2010), with deleterious consequences and signaling roles in biological systems (Blokhina and Fagerstedt, 2010a). Among the consequences are damage to proteins, lipids, carbohydrates, and DNA, which ultimately results in cell death (Gill and Tuteja, 2010). However, further investigations are needed to confirm NO influence on antioxidant modulation.

Although NO production is enhanced in hypoxia (Gupta et al., 2011), the increased activity of antioxidant enzymes (Figures 1A-F and 3A-F) under recovery conditions may be linked to longer exposure to hypoxia and the effects of the re-oxygenation, which are well reported as responsive for oxidative burst in the cells, leading to enzyme induction to counteract possible oxidative damage (Sairam et al., 2009; Kumutha et al., 2009; Simova-Stoilova et al., 2012)

CAT (Figures 1E and F), GR (Figures 3A and B) and GPOD (Figures 3C and D) have important roles in detoxification of H_2O_2 in roots of non-nodulated plants in both genotypes under hypoxia and recovery (Figures 7C and D). APX, another enzyme that acts in the scavenge of H_2O_2 , was responsive to the return to normoxic conditions in roots (Figures 1C and D) of non-nodulated plants, in agreement to gene expression and activity of APX in soybean seedling, which was responsive only after hypoxic stress (Shi et al., 2008). In nodulated plants, GPOD was important in detoxification of H_2O_2 , under hypoxia and recovery (Figures 3C and D). Blokhina et al. (2003) proposed the hypothesis that CAT acts earlier in response to hydrogen peroxide production compared to other enzymes, partly explaining its high activity in roots.

APX and GR are reported as responsible for scavenging H_2O_2 (Blokhina and Fagerstedt, 2010a, b; Gill and Tuteja, 2010), along with non-enzymatic antioxidants via the ascorbate-glutathione cycle (Bonifacio et al., 2011), where GR and GSH are used to reduce AsA, oxidized by APX (Blokhina and Fagerstedt, 2010b). The redox states of GSH and AsA in wheat roots were re-

ported to be directly dependent on oxygen concentration and reflected oxidative burst during re-aeration (Biemelt et al., 1998). Although, ROS production, $O_2^{\cdot-}$ and H_2O_2 , did not increase in roots (Figures 7A-F) during recovery, the increased ascorbate redox state and decreased AsA may be attributed to APX (Figures 1C and D; Figure 2C and D) and GR activity (Figures 3A and B; Figure 4A and B). On the other hand, CAT was not responsive in leaves (Figures 2E and F) as it was in roots (Figures 1E and F), while APX and GR increased mainly in response to recovery, which may be explained by the fluctuation of NAD(P)H/NAD(P)⁺ ratio under hypoxia (Stoimenova et al., 2007) to keep the ascorbate-glutathione cycle operating properly, whereas CAT itself does not need NAD(P)H to break down H_2O_2 (Blokina and Fagerstedt, 2010b).

On the other hand, dehydroascorbate (DHA) played an important role in the mitochondria respiratory electron transport chain in ascorbate regeneration (Blokina and Fagerstedt, 2010b). Interestingly, DHA also participated efficiently in NO scavenge (Kytzia et al., 2006).

In addition to the efficient enzymatic system operating to scavenge $O_2^{\cdot-}$ and H_2O_2 to avoid lipid peroxidation in roots (Figures 7E and F) and leaves (Figures 8E and F), enzyme GST may have an important role by tagging oxidized/degraded products, such as fatty acids and nucleic acids, for removing or by acting as a peroxidase to directly scavenge peroxides and remove lipid peroxidation products (Dalton et al., 2009). ROS is dangerous because of its ability to initiate a chain reaction on polyunsaturated fatty acids that leads to lipid peroxidation (Bai et al., 2010). Free fatty acids (FFAs) are recognized as powerful uncoupling agents activating mitochondrial uncoupling proteins (UCPs), leading to a severe membrane damage and further cell death (Blokina and Fagerstedt, 2010b).

The efficient antioxidant system as observed in roots of both genotypes (Figures 1A-F and 3A-F) may have an important role against ROS (Figures 7A-F). Furthermore, ROS production can also be avoided by uncoupling proteins (UCPs), as ROS acts on decreasing the electrochemical gradient (Δ_{vm}) when the mitochondria electron transport chain is over reduced (Schönfeld and Wojtczak, 2008). In addition, in non-nodulated plants, nitrate exerts an important role in the mitochondrial electron transport chain leading to ATP synthesis under hypoxia (Horchani et al., 2011) via oxidation of NADH and NADPH (Stoimenova et al., 2007).

The effect of waterlogging in roots induces stomatal closure in the shoots, once hypoxia decreases water absorption by roots and consequently hydraulic conductivity through xylem sap (Herrera, 2013). Besides, waterlogging could promote a redox imbalance of the chloroplast electron transport chain, which may have partly influenced high production of $O_2^{\cdot-}$ in leaves (Figures 8A and B). In addition, waterlogging can also influence the reduction of chlorophyll and carotenoids contents, ethylene production and disruption of photosynthate translocation (Blokina and Fagerstedt, 2010a, b).

The redox imbalance generated by an over reduction in the electron transport chain of chloroplasts is due to a decrease in ATP and NADPH consumption via Calvin-Benson cycle promoted by a limited CO_2 entry that slows down the carboxylation reaction of Rubisco enzyme. Thus, the over reduction of ETC also leads to electron escape that react with oxygen to produce ROS (Blokina and Fagerstedt, 2010a). In roots, waterlogging reduces the activity of mitochondrial ETC, which, in turn, reduces the possibility of electron to escape and ROS production, as reported by several authors (Azevedo Neto et al., 2006; Bai et al., 2010). In leaves, however, it is different, as ETC is driven by light absorption.

The increase in SOD activity in leaves (Figures 2A and B) may reflect $O_2^{\cdot-}$ production, once NO was not reported to be produced from nitrite reduction in leaves under hypoxic conditions (Gupta et al., 2005), although nitrate is transported through xylem sap from roots to shoot in soybean plants (Oliveira et al., 2013a; Lanza et al., 2014) and the lack ability of leaf mitochondria to produce NO may somehow be related to photosynthesis (Gupta et al., 2005).

Although genotypes Fundacep 53 RR and BRS Macota respond distinctly to hypoxia, tolerant and sensitive, respectively (Borella et al., 2014, 2017), the antioxidative metabolism here studied was not correlated with tolerance mechanisms that differentiate genotypes and it is probably due to the short duration of flooding of the root system (Wang et al., 2009). However, tolerant species increase the activity of antioxidant enzymes to counteract the oxidative effects (Sairam et al., 2009; Simova-Stoilova et al., 2012). According to our findings and others reported in studies, nitrate exerts beneficial effects on soybean plants by inducing antioxidant enzymatic and non-enzymatic compounds that may lead to a prolonged tolerance in comparison to non-nitrate-supplied plants.

Conclusions

Our data show that nitrate exerts beneficial effects on soybean plants under hypoxic conditions and consequent re-oxygenation by inducing the antioxidant system, mainly in roots, to cope with possible oxidative damage caused by ROS production. The enzymatic antioxidant system of soybean is also much more responsive during recovery from hypoxia stress than during the period of oxygen deprivation.

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Authors' Contributions

Conceptualization: Borella, J., Amarante, L., Oliveira, D.S.C., Braga, E.J.B., Oliveira, A.C.B. Data acquisition: Borella, J., Becker, R., Lima, M.C. Data Analysis: Borella, J., Becker, R., Lima, M.C., Amarante, L. Design of methodology: Borella, J., Amarante, L., Oliveira, D.S.C., Braga, E.J.B., Oliveira, A.C.B. Writing and editing: All authors have writing and approved the final version of the manuscript.

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