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Mycorrhizal fungi and microalgae modulate antioxidant capacity of basil plants

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Abstract

Mycorrhizal fungi, algae and cyanobacteria are some of the most important soil microorganisms and major components of a sustainable soil-plant system. This study presents for the first time evidence of the impact of green alga and cyanobacterium solely and in combination with arbuscular mycorrhizal fungi (AMF) on plant-antioxidant capacity. In order to provide a better understanding of the impact of AMF and soil microalgae on *Ocimum basilicum* L. performance, changes in the pattern and activity of the main antioxidant enzymes (AOEs), esterases and non-enzymatic antioxidants including phenols, flavonoids, ascorbate, and α -tocopherols were evaluated. The targeted inoculation of *O. basilicum* with AMF or algae (alone and in combination) enhanced the antioxidant capacity of the plants and the degree of stimulation varied depending on the treatment. Plants in symbiosis with AMF exhibited the highest antioxidant potential as was indicated by the enhanced functions of all studied leaf AOEs: 1.5-, 2- and more than 10-fold rises of superoxide dismutase (SOD), glutathione-S-transferase (GST) and glutathione reductase (GR), respectively. The greatest increase in the total esterase activity and concentration of phenols, flavonoids and ascorbate was marked in the plants with simultaneous inoculation of mycorrhizal fungi and the green algae. 2,2-diphenyl-1-picryl-hydrazyl (DPPH) free radical scavenging method and ferric reducing antioxidant power (FRAP) assay proved the increased plant antioxidant capacity after co-colonization of green algae and mycorrhizae.

Key words: antioxidants, arbuscular mycorrhizal fungi, *Ocimum basilicum* L., soil microalgae

Introduction

Medicinal plants from the Lamiaceae family have been used since ancient times for their pharmaceutical properties (Bais *et al.* 2002) due to the secondary metabolites such as phenolic compounds (including flavonoids and phenylpropanoids) as well as anthocyanins (Phippen and Simon 2000). Sweet basil (*Ocimum basilicum* L.) belonging to the Lamiaceae family is a rich source of phenolic antioxidant compounds and flavonoids (Juliani and Simon 2002; Jayasinghe *et al.* 2003).

Soil-microbe-plant interactions are complex and there are many ways in which they can affect plant health and productivity. Many species belonging to the Lamiaceae family, including sweet basil, form a symbiotic association with arbuscular mycorrhizal fungi (AMF) (Wang and Qiu 2006). In addition to increasing the uptake of poorly available nutrients such as phosphorus and nitrogen (Toussaint *et al.* 2004) or providing protection against pathogens (Odeyemi *et al.* 2010;

Ziedan *et al.* 2011), AMF can also cause changes in the accumulation of secondary metabolites, including phenolics, in host plant roots (Devi and Reddy 2002; Yao *et al.* 2003). Induction of specific manganese-dependent superoxide dismutase (Mn-SOD) isoforms observed in colonized parts of mycorrhizal tomato roots suggests that localized oxidative bursts might occur in the fungi (Pozo *et al.* 2002). However, it was reported that *Glomus mosseae* directly increases the essential oil content in shoots of *Origanum* sp. (Khaosaad *et al.* 2006) as well as sweet basil (Copetta *et al.* 2006). Reactive oxygen species (ROS) were initially thought as toxic by-products of aerobic metabolism. In recent years, it has been established that ROS play an important signaling role in the plants, controlling processes such as growth, development, response to biotic and abiotic environmental factors and programmed cell death. When ROS are at high concentrations they may initiate destructive oxidative processes such as chlorophyll bleaching, lipid peroxidation, protein oxidation, and damage to nucleic acids (Herbinger *et al.* 2002). To protect themselves from oxidative injury, higher plants induced effective antioxidant systems (Asada 1999). The antioxidant systems consist of antioxidant enzymes and non-enzymatic antioxidants including phenols, flavonoids, ascorbate, glutathione, and α -tocopherols. Plant phenols show marked qualitative and quantitative variations not only at different genetic levels (between and within species and cultivars) but also between physiological and developmental stages (Bunning *et al.* 2010). They also vary in response to environmental factors, such as light intensity and nutrient availability (Yang *et al.* 2004; Mogren *et al.* 2007).

The beneficial effect of arbuscular mycorrhizal inoculation on plants is well known. Relatively little is known about the effects of AMF colonization and the addition of algae on the antioxidant defense system in shoots of medicinal plants, which are often the harvest products.

Cyanobacteria and green algae release into the soil a large number of substances that play an important role in building up soil fertility, as well as acting as plant growth promoting agents, which results in increased productivity. These microorganisms contain growth-promoting regulators (auxin, gibberellic acid, and cytokinins), nutrients, sugar, amino acids, vitamins, and other secondary metabolites (Karthikeyan *et al.* 2007). Cyanobacteria, more commonly known as blue-green algae, are able to convert insoluble phosphorus in the soil into forms accessible to plants and some of them fix the atmospheric dinitrogen and convert it into a bioavailable form of ammonium (Sahu *et al.* 2012). *Synechocystis* sp. strain R10 was isolated from the shallow lake in Rupite, Bulgaria. The effect of light and temperature on growth, biochemical composition, enzymatic antioxidant defense and biological

activity of *Synechocystis* sp. R10 (Gigova *et al.* 2012) was investigated. However, the role of microalgae as a plant antioxidant, especially in relation to their functioning in the rhizosphere, has not been explored in depth. Most research has focused on their growth promotion (influence on plant photosynthesis and yield). Due to the lack of information, the aim of the presented research was to evaluate the influence of AMF (*Claroideogloium claroideum* EEZ 54) and mono-algal cultures (*Scenedesmus incrassatulus* R83 and *Synechocystis* sp. R10) in the soil on the antioxidant capacity of basil (*O. basilicum*) plants.

Materials and Methods

Biological materials and growth conditions

Basil plants (*O. basilicum* var. *purpurascens* Benth.) were grown from seeds in 2 kg transparent plastic boxes (five plants per box for each treatment) of polyethylene glycol terephthalate (PET) with 92% transmittance in the range of 380–710 nm in a growth chamber from May to September under a light intensity of $320 \mu\text{mol} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ at 21–25°C (night-day) and a 15 h photoperiod. The relative humidity ranged from 40 to 65%. The plants developed on a soil : perlite substrate (3 : 1, v/v). All pots were adjusted daily to 60% water holding capacity. The soil (leached cinnamonic forest soil – Chromic Luvisols, FAO – 30–40 cm depth) had the following agrochemical characteristics: pH (H_2O) – 6.2; $8 \text{ mg} \cdot \text{kg}^{-1}$ soil total mobile nitrogen ($\text{N-NO}_3^- + \text{N-NH}_4^+$); $30 \text{ mg} \cdot \text{kg}^{-1}$ soil P_2O_5 ; $120 \text{ mg} \cdot \text{kg}^{-1}$ soil K_2O .

The mycorrhizal strain (*C. claroideum*, ref. EEZ 54) was kindly provided by the AMF collection of Estación Experimental del Zaidín (CSIC Granada, Spain). Mycorrhizal inoculation was done by placing the seeds over a thin layer of the AMF inoculum ($2 \text{ g} \cdot \text{kg}^{-1}$ soil substrate) following the layering method of Jackson *et al.* (1972). The inoculum consisted of colonized roots and soil from 4-month-old oat pot cultures.

Algae strains came from the culture collection of the Experimental Algology Department (Institute of Plant Physiology and Genetics, Bulgarian Academy of Sciences). Monoalgal cultures of *Scenedesmus incrassatulus* R83 and *Synechocystis* sp. R10 were grown autotrophically in a medium according to Šetlik (1967), modified by Georgiev *et al.* (1978) with a 1/2 concentration of nutrients and mineral medium after Aiba and Ogawa (1977), respectively. The final concentration of algal suspensions used for watering was $0.5 \text{ g} \cdot \text{l}^{-1}$. The algae were maintained and prepared for watering of basil plants with uninterrupted illumination from luminescent lamps ($75 \mu\text{mol} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$ light intensity), and bubbling with $3 \text{ cm}^3 \cdot \text{s}^{-1}$ air, enriched with 0.5% CO_2 (Petkov 1995). Algal suspension for

inoculation was added twice after seed germination instead of watering with sterilized water (Petkov 1995).

Six treatments were compared: 1 – control non-inoculated plants (C); 2 – plants, inoculated with *C. claroideum* EEZ 54 (AM); 3 – plants, inoculated with *S. incrassatulus* R83 (AL1); 4 – plants, inoculated with *Synechocystis* sp. R10 (AL2); 5 – plants, inoculated with *C. claroideum* EEZ 54 and *S. incrassatulus* R83 (AM + AL1); 6 – plants, inoculated with *C. claroideum* EEZ 54 and *Synechocystis* sp. R10 (AM + AL2).

Native polyacrylamide gel electrophoresis (PAGE) and enzyme activity staining

Fresh leaves (0.50 g FW) from plants exposed to different treatments were homogenized in 0.1 mM K-phosphate buffer (pH 7.8), containing 2.0 mM disodium ethylenediaminetetraacetate dihydrate ($\text{Na}_2\text{-EDTA}$), 1 mM phenyl methyl sulfonyl fluoride (PMSF), 2% polyvinylpyrrolidone K-40 (w/v) and 10% glycerol. The homogenates were centrifuged at $12\,000 \times g$ for 30 min. Both steps were carried out at 0–4°C. The concentration of soluble protein in the supernatants was measured by the dye-binding assay (Bradford 1976) with bovine serum albumin grade V (Sigma) as a standard. For determination of changes in the isoenzyme profile and activities of superoxide dismutase (SOD, EC 1.15.1.1), glutathione-S-transferase (GST, EC 2.5.1.18), glutathione-reductase (GR, EC 1.6.4.2) and esterase (EST, EC 3.1.1.x), equal amounts (35 µg) of leaf proteins were subjected to discontinuous PAGE under non-denaturing, non-reducing conditions essentially as described by Laemmli (1970), but without sodium dodecyl sulfate (SDS). Electrophoretic separation was performed by using 4% and 10% polyacrylamide in the stacking and resolving gels, respectively, for 4–5 h at a constant current of 35 mA per gel. After completion of electrophoresis, separate gels were stained for the activities of the respective enzymes.

Bands of SOD activity were localized on gels as described by Azevedo *et al.* (1998). The gels were soaked in 0.1 mM nitroblue tetrazolium, 0.05 mM riboflavin, and 0.3% (v/v) tetramethyl ethylene diamine (TEMED) in 50 mM potassium phosphate buffer (pH 7.8) for 20 min in the dark. The gels were then placed in distilled water and exposed to a light box until they became uniformly blue, while bands where SOD was present were achromatic (10–15 min). The metalloforms of SOD were identified by staining parallel gels preincubated for 30 min with selective inhibitors (2 mM KCN for inhibiting Cu/ZnSOD and 5 mM H_2O_2 for inhibiting both Cu/ZnSOD and FeSOD (Azevedo *et al.* 1998).

Glutathione reductase activity was detected by incubating the gels in 250 mM Tris-HCl buffer (pH 7.8), containing 0.24 mM 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), 0.34 mM

2,6-dichlorophenolindophenol (DPIP), 3.6 mM oxidized glutathione (GSSG), and 0.4 mM reduced form of nicotinamide adenine dinucleotide phosphate (NADPH) in darkness for 1 h (Anderson *et al.* 1995).

For GST activity staining, the method described by Ricci *et al.* (1984) was used. Briefly, after the electrophoretic run, the gel was equilibrated in 100 mM potassium-phosphate buffer, pH 6.5, for 10 min, and transferred to a reaction mixture containing 4.5 mM glutathione (GSH), 1 mM 1-chloro-2,4-dinitrobenzene (CDNB), and 1 mM nitroblue tetrazolium (NBT) in 0.1 M potassium-phosphate buffer, pH 6.5, at 37°C for 10 min. Further, the gel was incubated at room temperature in 100 mM Tris/HCl, pH 9.6, containing 3 mM phenazine methosulphate (PMS). The activity band appeared as an achromatic zone against a blue background.

Esterase isoforms and activity were visualized by staining the gels with α -naphthyl acetate (0.06%, dissolved in a few drops of acetone) and Fast blue BB (0.1%) in 200 mM Tris (pH 7.0) with gentle shaking at room temperature until black bands appeared (Murphy *et al.* 1996).

Determination of total phenolic compounds and flavonoids

Dry leaf samples (1 g) were ground and thoroughly extracted with 96% (v/v) methanol. Concentrations of phenolic compounds were determined spectrophotometrically using the Folin-Ciocalteu reagent and calculated as caffeic acid equivalents (Pfeffer *et al.* 1998). Flavonoids in plant tissues were measured spectrophotometrically according to Zhishen *et al.* (1999), using the standard curve of catechin.

Antioxidant potential assay

Spectrophotometric quantification of water-soluble (WS-AOC) and lipid-soluble (LS-AOC) antioxidant capacity, expressed as equivalents of ascorbate and α -tocopherol, were performed through the formation of the phospho-molybdenum complex (Prieto *et al.* 1999). The assay was based on the reduction of Mo (VI) to Mo (V) by the sample analysis and the subsequent formation of a green phosphate/Mo (V) complex at an acidic pH. Dry leaf material (0.5 g) was ground with pestle and mortar to a fine powder and 3 ml dH_2O was added and the suspension was homogenized, transferred to tubes and shaken for 1 h at room temperature in the dark. The suspension was filtered and the extraction repeated with 3 ml dH_2O . The pellet was washed again with 2 ml dH_2O . For lipid soluble antioxidant capacity (expressed as α -tocopherol), the procedure was the same except that the extraction was carried out with hexane as a solvent. The method was optimized and characterized on linearity interval, repeatability and reproducibility and molar absorption

coefficients for the quantitation of water-soluble and lipid-soluble antioxidant capacities expressed as equivalents of ascorbate and α -tocopherol (Prieto *et al.* 1999). Absorption coefficients were: $(3.4 \pm 0.1) \times 10^3 \text{ M}^{-1} \cdot \text{cm}^{-1}$ for ascorbic acid and $(4.0 \pm 0.1) \times 10^3 \text{ M}^{-1} \cdot \text{cm}^{-1}$ for α -tocopherol.

Ferric reducing antioxidant power (FRAP) assay

The FRAP reagent was freshly prepared by mixing acetate buffer (300 mM, pH 3.6), TPTZ (tripyridyltriazine) solution (10 mM TPTZ in 40 mM HCl), and $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ (20 mM) at a ratio of 10 : 1 : 1 (Benzie and Strain 1996). To perform the assay, 900 μl of FRAP reagent, 90 μl of distilled water and 30 μl of the leaf extract were mixed and incubated at 37°C for 15 min. The absorbance was measured at 595 nm using FRAP working solution as a blank.

Total antioxidant capacity

Free radical-scavenging activity in leaf tissues was determined from the bleaching of the purple methanol solution (2, 2-diphenyl-1-picryl-hydrazyl, DPPH \cdot), according to Tepe *et al.* (2006). DPPH \cdot is a stable radical with a maximum absorption at 517 nm that can readily undergo reduction by an antioxidant. The percent inhibition of the DPPH \cdot radical (I%) was calculated by the following equation:

$$I\% = [(A_{\text{blank}} - A_{\text{sample}}) / A_{\text{blank}}] \times 100,$$

where: A_{blank} – the absorbance of the control reaction (containing all reagents except the test compound), A_{sample} – the absorbance of the test compound.

Statistical analysis

The experiment had a completely randomized block design with four replications. The data were subjected to one-way analysis of variance (ANOVA) for comparison of means, and significant differences were calculated according to Fisher's least significance difference (LSD) test at the 5% significance level using a statistical software package (Statgraphics Plus, version 5.1 for Windows). Data are presented as means \pm standard error.

Results

This study focused on the impact of AMF and microalgae on *O. basilicum* performance, changes in the pattern and activity of the main antioxidant enzymes, esterases and non-enzymatic antioxidants including phenols, flavonoids, ascorbate, and α -tocopherols. Seven common bands of SOD activity were detected in leaf protein samples from plants exposed to treatments (Fig. 1A, a). Band numbers 1 and 2 were identified as Mn-SOD enzymes by their insensitivity to inhibition by both H_2O_2 (Fig. 1A, b) and cyanide (Fig. 1A, c), while band numbers 3 to 7 were sensitive to both inhibitors, suggesting that they represented a CuZnSOD activity. FeSOD activity was not detected in *O. basilicum* L.

Compared to the control untreated plants (C), the relative total activity of SOD, the first enzyme which scavenges reactive oxygen species, did not change after combined treatment with *C. claroideum* EEZ 54 and *S. incrassatulus* R83 (AM + AL1), while the activity was higher in plants treated only with AM, and lower

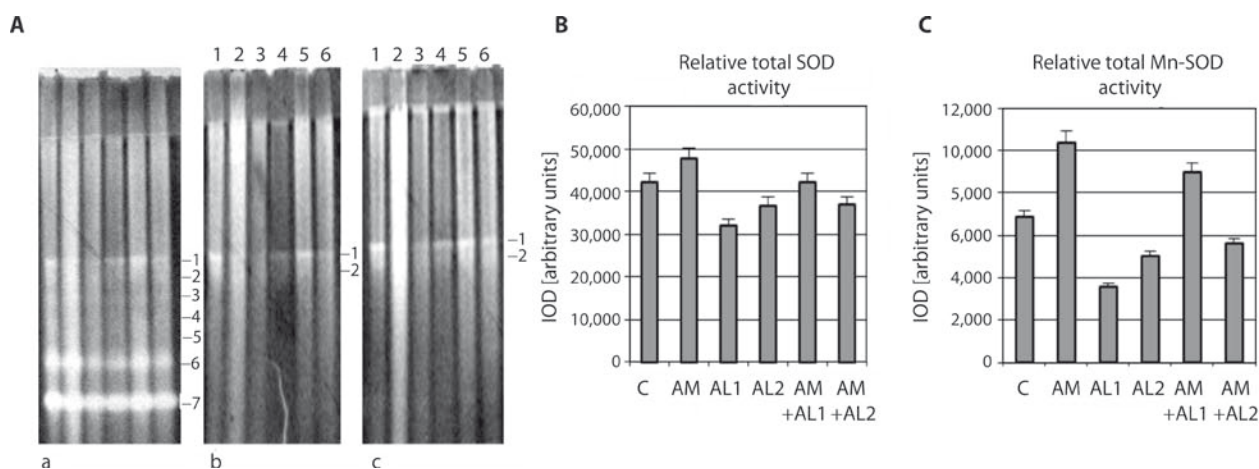


Fig. 1. Activity of superoxide dismutase (SOD) of non-inoculated (control) and inoculated (AMF and/or algae) *Ocimum basilicum* plants. In a gel SOD isoforms (A) pattern and intensity (a), identification of Mn-SOD (b), and identification of CuZnSOD (c) in leaves after native PAGE of the samples: lane 1– control untreated plant (C); 2 – AM; 3 – AL1; 4 – AL2; 5 – AM+AL1; 6 – AM+AL2. Equal amounts of protein (35 μg) were loaded per well. The isoforms of SOD are numbered from cathode to anode. SOD1 and SOD2 represent Mn-SOD; and SOD3 to 7 – CuZnSOD activity. Relative total SOD activity (B) and total activity of MnSOD (C) of the same samples expressed as a sum of the values (in arbitrary units) for integrated optical density (IOD) of the respective bands. Results shown are representative of three independent trials

in plants treated with AL1 (Fig. 1B). Application of *Synechocystis* sp. R10 alone and in combination with *C. claroideum* EEZ 54 (AL2; AM + AL2) slightly lowered the total SOD activity. The same trend of changes was observed in total Mn-SOD activity (Fig. 1C), but the increase and decrease were more pronounced, reaching 50% in both cases. Mn-SOD was also more responsive to the AM + AL1 treatment (about 30% rise).

Four well-resolved bands of GR were detected in *O. basilicum* leaf extracts, following all treatments. Despite the similarity in GR patterns (Fig. 2A), the activity of this enzyme varied between the treatments. In control (C), AL2 and AM + AL1 treated plants, a relatively low total GR activity was maintained

(Fig. 2B). In AL1, AM + AL2 and AMF treatments, a gradual increase in enzyme activity was observed (by 200, 346 and 1150%, respectively).

GST was presented by seven isoforms (Fig. 3A). All treatments, except AL1, led to an increase in total activity from 148% for AM + AL2 to 288% for AL2, mainly due to stimulation of isoforms 3, 4, 5 and 6 (Fig. 3B).

Five esterase enzymes with a capacity to hydrolyze non-specific ester substrates such as α -naphthyl acetate were visualized on native polyacrylamide gels. The EST2 and EST4 were the most intense in all samples (Fig. 4A). Although the intensity of each band varied between treatments, relative total EST activity (Fig. 4B) increased or decreased slightly (in the range of 10 to

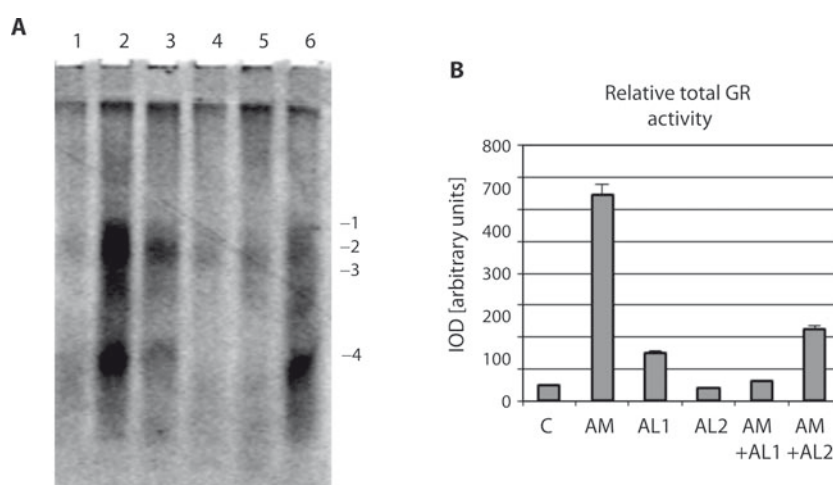


Fig. 2. Glutathione reductase (GR) activity of non-inoculated (control) and inoculated (AMF and/or algae) *Ocimum basilicum* plants. In a gel GR isoform pattern (A) and intensity in leaves after native PAGE of the samples (B): lane 1 – control untreated plant (C); 2 – AM; 3 – AL1; 4 – AL2; 5 – AM+AL1; 6 – AM+AL2. Relative total GR activity of the same samples expressed in arbitrary units. Results shown are representative of three independent trials

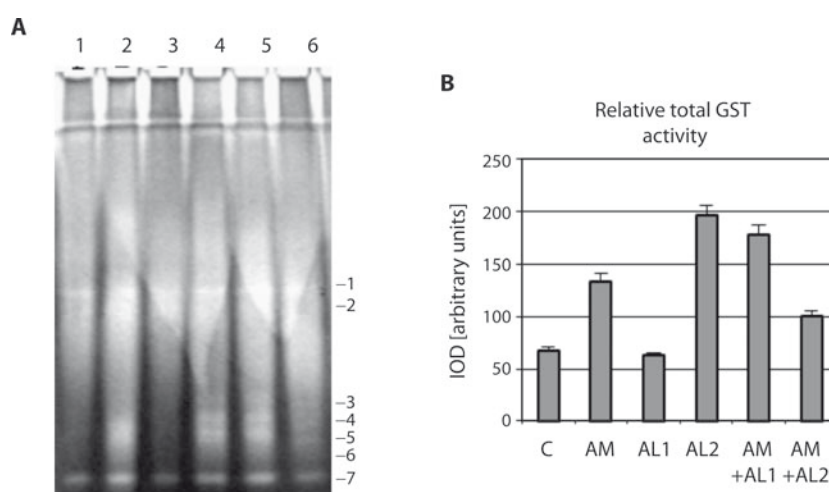


Fig. 3. Glutathione-S-transferase (GST) activity of non-inoculated (control) and inoculated (AMF and/or algae) *Ocimum basilicum* plants. In a gel GST isoform pattern (A) and intensity in leaves after native PAGE of the samples (B): lane 1 – control untreated plant (C); 2 – AM; 3 – AL1; 4 – AL2; 5 – AM+AL1; 6 – AM+AL2. Relative total GST activity of the same samples expressed in arbitrary units. Results shown are representative of three independent trials

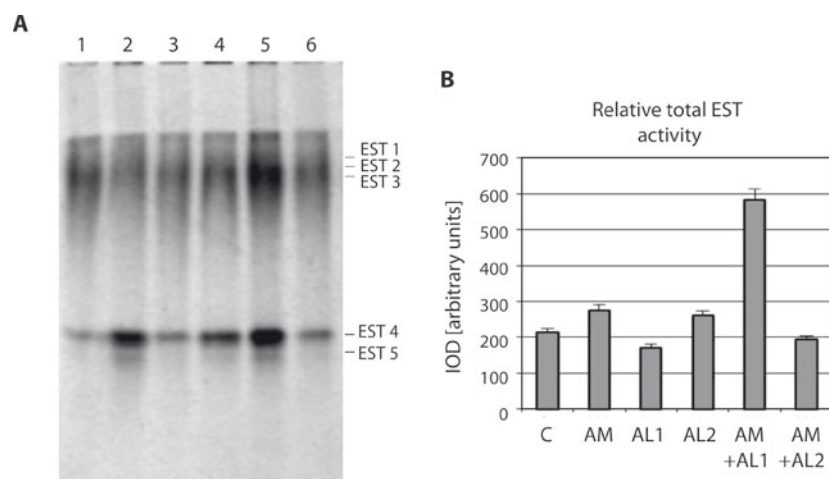


Fig. 4. Esterase (EST) activity of non-inoculated (control) and inoculated (AMF and/or algae) *Ocimum basilicum* plants. In a gel EST isoform pattern (A) and intensity in leaves after native PAGE of the samples (B): lane 1 – control untreated plant (C); 2 – AM; 3 – AL1; 4 – AL2; 5 – AM+AL1; 6 – AM+AL2. Relative total EST activity of the same samples expressed in arbitrary units. Results shown are representative of three independent trials

30%) compared to that of control plants (accepted for 100%). Only AM + AL1 treated plants showed significantly higher total enzyme activity (274%). It should be noted, however, that slow (EST1 to 3) and faster (EST4 and 5) moving enzyme forms behaved differently. The slow moving ESTs showed a drop in activity (except for AM + AL1 treatment) in a narrow range (63–88% from that of the control), while the activity of the faster moving EST4 and EST5 was increased. The relative total EST4 + EST5 activity was lowest in the control (accepted for 100%) and highest in the AM + AL1 plants (1064%). AM and AL2 treatments also markedly elevated the activity (by about 600 and 320%, respectively). Comparatively less stimulation was observed in the AL1 and AM + AL2 treated plants (by 54 and 60%, respectively).

Green algae showed more impact on total phenol content than cyanobacterium (Fig. 5). The content of total phenol compounds in nonmycorrhizal and mycorrhizal plants was less when they were treated with cyanobacterium (AL2; AM + AL2) in comparison with treatment with green algae (AL1; AM + AL1). The greatest flavonoid content rise was marked in the plants with tandem inoculation of mycorrhizal fungi and algae (AM + AL1; AM + AL2) while the lowest concentration was detected in control variant.

The content of lipid soluble metabolites with antioxidant capacity expressed as α -tocopherol in basil plants significantly increased with AMF treatment (Fig. 5) followed by inoculation with green alga solely (AL1) and in combination with AMF (AM + AL1). Water soluble-AOC increased with the addition of AMF together with microalgae. Basil plants treated with cyanobacterium solely (AL2) had less WS-AOC (Fig. 5). Total phenols, WS-AOC and LS-AOC

contents in mycorrhizal plants are higher when they are treated with green algae (AM + AL1), in comparison when they are treated with a cyanobacterium (AM + AL2). The antioxidant capacities measured by DPPH and FRAP methods in basil plants decreased in the control compared with inoculated treatments (Fig. 5). Green microalgae (AL1; AM + AL1) in the soil led to the highest antioxidant capacities in the basil leaves. Targeted treatments with cyanobacterium (AL2; AM + AL2) had lower FRAP and DPPH values (Fig. 5).

Discussion

Soil organisms play a crucial role in the functioning of agricultural ecosystems. Mycorrhizal fungi, algae, and cyanobacteria are some of the most important soil microorganisms and major components of a sustainable soil-plant system. This study presents for the first time evidence of the impact of green alga and cyanobacterium solely and in combination with AMF on plant antioxidant activity. In order to get a better understanding of the impact of AMF and soil microalgae on *O. basilicum* performance, the changes in the pattern and activity of the main antioxidant enzymes, esterases and non-enzymatic antioxidants including phenols, flavonoids, ascorbate, and α -tocopherols were evaluated.

Our results showed that the treatments of *O. basilicum* with AMF or algae (alone and in combination) stimulated the antioxidant capacity of the plants and the degree of stimulation varied depending on the treatment. Moreover, the effect of each treatment was due to activation of specific antioxidant enzymes.

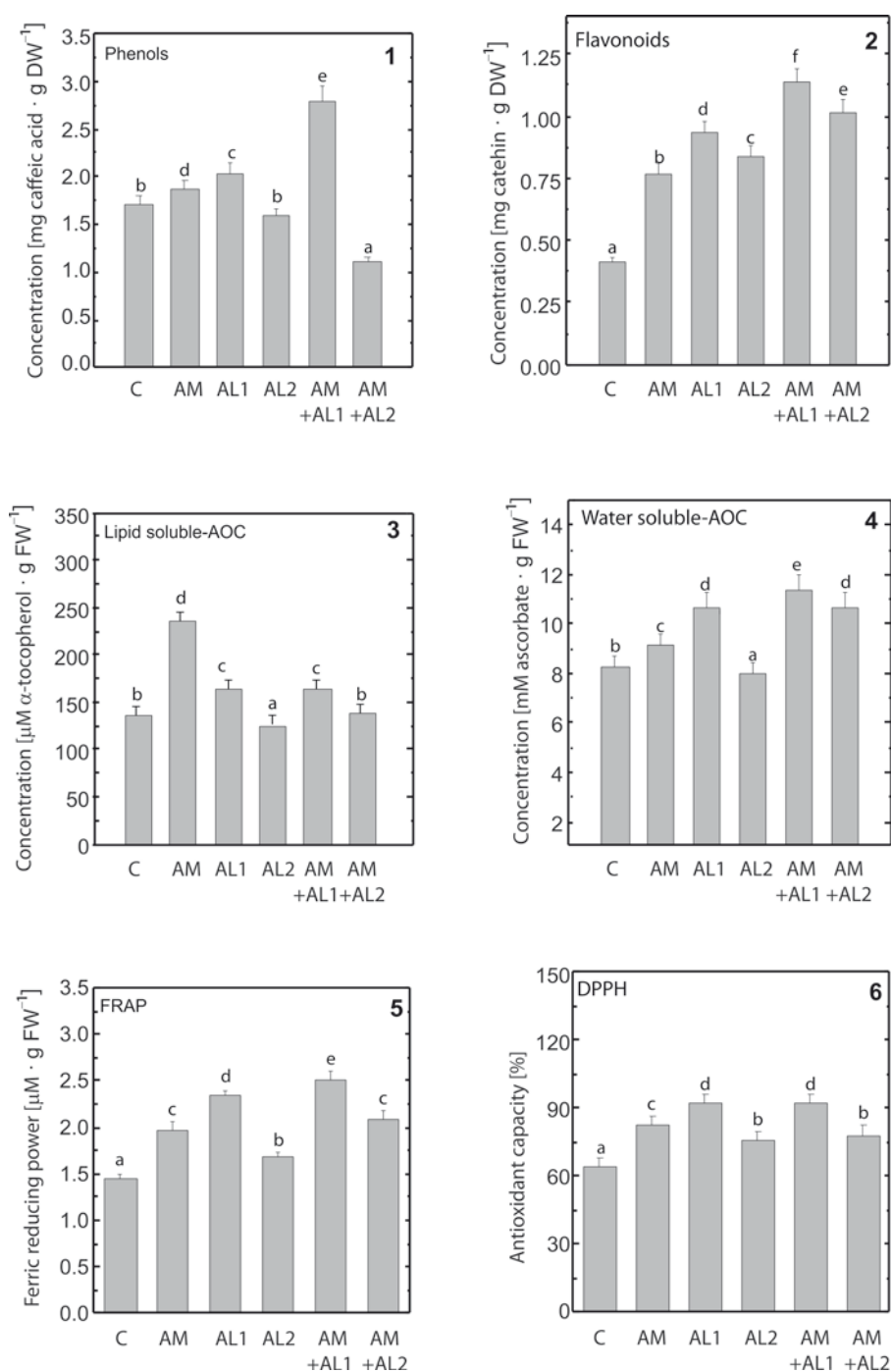


Fig. 5. Total phenolic (1), flavonoid concentrations (2), lipid-soluble (3), water-soluble antioxidant capacities (AOC) (expressed as equivalents of ascorbate and α -tocopherol) (4), ferric reducing antioxidant power (FRAP) (5) and free radical-scavenging activity (DPPH) (6) of non-inoculated (control, C) and inoculated (AMF and/or algae) *Ocimum basilicum* plants. FW – fresh weight, DW – dry weight. Values are means \pm SE. Letters in common within a graph indicate no significant differences assessed by Fisher LSD test ($p \leq 0.05$) after performing ANOVA ($n = 9$)

Plants treated with AMF exhibited the highest antioxidant potential as was indicated by the enhanced functions of all studied leaf AOE (1.5-, 2- and more than 10-fold rises of Mn-SOD, GST and GR, respectively). In the AM + AL1 treated plants the activities of Mn-SOD and especially of GST were stimulated (by about 30 and 160%, respectively), while in the AM + AL2

plants the activities of both GR and GST were higher than the control. AL1 treatment stimulated the activity of GR by about three times. GST was the only positive responder to the AL2 treatment. When applied alone, the algae apparently had less effect and this was probably related to the lack of direct contact between them and plants. When basil plants are double inocu-

lated with *C. claroideum* EEZ 54 and *S. incrassatulus* R 83 (AM + AL1) or *Synechocystis* sp. R 10 (AM + AL2) however, the activity of plant enzymes differed from that after AMF only inoculation, suggesting that algae impact the AMF.

While being extensively studied primarily in the roots, a rise of the antioxidant status in the aboveground plant parts has been reported in different AM-inoculated plants compared to non-mycorrhizal plants. For example, higher levels of antioxidant enzyme activities were observed in leaves of tomato plants colonized by *Glomus mosseae* (Abdel Latef and Chaoxing 2011). According to Rozpądek *et al.* (2014), SOD activity measured in *Cichorium intybus* L. significantly increased in AMF inoculated plants, compared to non-inoculated plants. Estrada *et al.* (2013) also found that AMF symbiosis induced SOD and CAT activities of maize shoots and roots, emphasizing that the degree of induction depended on the AMF species. Despite the symbiotic nature of AMF associations, enhancement of AOE functions may be a result of general plant defense response to fungal colonization.

In earlier reports, stimulation of leaf esterase activity was observed in duckweed (*Lemna minor*) exposed to lead, cadmium, chromium, zinc, copper and mercury (Mukherjee *et al.* 2004) and also in *Centaurea ragusina* under salt treatments (Radić and Pevalek-Kozlina 2010). In these studies, the authors imply that the induction of esterases confers tolerance of plants to the applied treatments. Furthermore, plant esterases are known to actively participate in many biological processes, such as activation of signal molecules (Stuhlfelder *et al.* 2004), regulation of cell growth and organ development (Gao *et al.* 2009) as well as regulation of secondary metabolite levels and bioactivity (Dogru *et al.* 2000; Ganjewala and Luthra 2009). Based on this knowledge, it is tempting to speculate that the activated specific esterases (EST4 and EST5) of *O. basilicum* participate in the improvement of plant growth and development observed following all treatments (data not shown).

Some antioxidants are high molecular weight proteins and enzymes, while others are small molecular weight compounds such as phenols. Their antioxidant properties are associated with their ROS scavenging potential (Sakihama and Yamasaki 2002). Phenolic production in plants can be affected by biotic and abiotic factors (Toussaint *et al.* 2007). AMF colonization and microalgae treatment may be a way to alter or enhance phenolic production or composition within the host plant. It is also known that the accumulation of phenols in plants is dependent on the developmental stage of the symbiosis (Abdel-Lateif *et al.* 2012). In the present study, these compounds were analyzed in the advanced stage of mycorrhiza formation. The total

phenols of *O. basilicum* are increased, when plants are inoculated with AMF and green algae (Fig. 5). Similar results for the phenol content have already been reported for several other mycorrhized plants (Jurkiewicz *et al.* 2010; Zubek *et al.* 2015). Phenolic compounds may play a key role in the crosstalk between plants and symbiotic fungi, as they affect spore germination, the growth of hyphae and stimulate root colonization (Abdel-Lateif *et al.* 2012; Pusztahelyi *et al.* 2015). There is a lack of studies on whether the microalgae have some role for phenols and flavonoids synthesis in plants. The contents of total phenols in aerial parts of basil do not change significantly when the plants are mycorrhized (Hazzoumi *et al.* 2015). Our results indicate that basil antioxidant potential mainly defined by its high concentration of phenolic compounds and flavonoids. These results confirm previous reports showing that total phenols and flavonoids contributed significantly to the antioxidant capacity in the plants of *Lamiaceae* (Matkowski *et al.* 2008).

Conclusions

The present study provides evidence that dual inoculation can be a simple and useful method for obtaining higher contents of total phenols, flavonoids and water soluble antioxidants followed by an increase of antioxidant activity in *O. basilicum*. The production of higher yield and quality in medicinal plants, through conventional methods, often requires external inputs such as fertilizers and pesticides. In this context, the use of bioinoculants, a natural alternative to chemical fertilizers, is likely to promote the production of active ingredients.

More detailed studies are needed, however, to clarify the exact role of mycorrhizal fungi in increasing the antioxidant activity and regulation of defense responses of the host plant. On the other hand, AMF has been found to improve resistance to pathogens, heavy metals and salinity, and influence the level of secondary metabolites in plants (Abdel Latef and Chaoxing 2011; Estrada *et al.* 2013; Zubek *et al.* 2015; Mollavali *et al.* 2016). With this in mind, it cannot be ruled out that the increase in antioxidant capacity is a prerequisite for better tolerance of the plants to subsequent abiotic and/or biotic stress.

AM associations bring about significant changes in the host plant and its environment: at the rhizosphere level, they influence soil structure, carbon deposition in the soil, and microbial diversity, in part through changes in root exudation. These shifts in the microbial communities of the rhizosphere may indirectly influence the outcome of plant interactions with other organisms, including soil microalgae.

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