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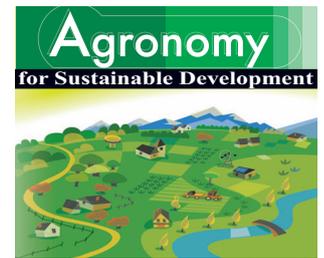
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Research article

Peroxidase changes in *Phoenix dactylifera* palms inoculated with mycorrhizal and biocontrol fungi

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Abstract – In Morocco, yields of date palms are highly decreased by the pathogen fungi *Fusarium oxysporum*. To solve this issue, mycorrhizal fungi and biocontrol agents could enhance plant resistance to pathogens. Here, we studied peroxidases in roots and leaves of *Phoenix dactylifera* in response to inoculation with the fungi *Glomus mosseae* and *Trichoderma harzianum*. Date palm plants were harvested 57 weeks after inoculation with mycorrhizal fungi and biocontrol fungi. We measured the dry biomass, arbuscular colonization, cytochemistry of peroxidase, and peroxidase forms and activities in roots and leaves. Our results show that mycorrhization increased the plant dry biomass by about 57%. The rate of mycorrhizal colonization ranged from 25% to 30%. Peroxidase activity in roots colonized by *T. harzianum* alone was 1.6 times higher than in control plants. Peroxidase activity in roots colonized by *G. mosseae* and *T. harzianum* was about 2 times higher than in control plants. Peroxidase activity in leaves increased by +419% when plants were inoculated by both fungi. The cytochemical results show an accumulation of structural substances in root cell walls after inoculation with *T. harzianum*. These structural substances may increase the mechanical strength of the host cell walls in order to inhibit pathogen invasion. Peroxidase activities were found in plant cell walls; the tonoplast and host plasmalemma in the chloroplast; mitochondrial membranes; and intercellular spaces of plants inoculated with *G. mosseae* and *T. harzianum*. SDS-PAGE analyses of leaf extracts gave a main band at 54 kDa for all the treatments. The stimulatory effect of *Trichoderma* on the peroxidase activity is a resistance mechanism of date plants to pathogens. The use of *Trichoderma* could thus be an alternative to chemicals in crop protection.

Peroxidase / *Trichoderma* / date palm / arbuscular mycorrhizal

1. INTRODUCTION

Phoenix dactylifera is considered the most important plant grown in arid regions. This palm species suffers from a great disease problem caused by *Fusarium oxysporum*, which produces high economic losses in palm groves (Oihabi, 1991). The development of biological control against plant diseases is accepted nowadays as a durable and environmentally-friendly alternative for agrochemicals. Therefore, considerable attention is being paid to the isolation of fungal antagonists that could be as effective as pesticides in the repression of fungal pathogens (Benhamou and Chet, 1996). The mechanism responsible for biocontrol is unknown, although both hydrolytic enzymes (chitinases, glucanases and proteases) and naturally produced antibiotics play an important role.

Association of biocontrol agents such as *T. harzianum* and promoting microorganisms, such as arbuscular mycorrhiza fungi, may contribute to a better protection of the host plant.

Arbuscular mycorrhizal fungi are obligate symbionts that live in association with the roots of most land plants. These fungi grow from the roots into the surrounding soil, forming an external hyphal network, which increases uptake of mineral immobile nutrients (Pedreno et al., 1997; Smith and Read, 1997). In addition, arbuscular mycorrhizal fungi promote plant growth and may contribute to the protection of the host plant against soilborne plant pathogens (Pozo et al., 2002). On the other hand, several *Trichoderma* strains have been reported to be effective in controlling plant diseases (Viterbo et al., 2002). Thus, combinations of arbuscular mycorrhizal fungi and biocontrol agents such as *Trichoderma* could provide levels of disease control which are superior to the effects of the organism when they are used alone (Lindermann, 1988; Datnoff et al., 1995). However, different authors have reported the suppression of arbuscular mycorrhizal root colonization by biocontrol *Trichoderma* species (Calvet, 1989; McAllister et al., 1994), although this effect may depend on the host plant species (Dhillon, 1994) and the timing of inoculation (McAllister et al., 1994). In addition, adverse effects

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of arbuscular mycorrhizal fungi on the population density of *Trichoderma* sp. have also been observed (Green et al., 1999). This type of study has never been carried out on date palms.

Although little is known about the biochemical and physiological processes occurring during the symbiosis establishment, some studies suggest that increased resistance of arbuscular mycorrhizal roots to pathogens may be associated in part with marked metabolic changes in the host, including an enhanced production of peroxidases and phenolic compounds (Spanu and Bonafante-Fasolo, 1988). Peroxidases (EC 1.11.1.7) oxidize a variety of phenolic compounds and have been implicated in many different physiological processes such as incorporation of phenolics into the cell wall, lignification and wound healing, and pathogen defense (Jaizme-Vega and Díaz-Pérez, 1991). An increase in peroxidase activity after inoculation of different plants with pathogenic fungi has been reported by different groups (Münzenberger et al., 1997; Salzer and Hager, 1993). It has been proposed that one role of plant peroxidases is to prevent spreading of pathogens by increasing cell wall rigidity through the incorporation of phenolics (Yedidia et al., 1999). The question arises whether fungal symbionts and saprophytic fungi influence peroxidase activities and isoenzyme patterns in plants in a manner similar to that induced by pathogens (Spanu and Bonafante-Fasolo, 1988). In the present study peroxidase activity was analyzed both in roots and leaves of *P. dactylifera* plantlets inoculated with *G. mosseae* and *T. harzianum* T22.

2. MATERIALS AND METHODS

2.1. Fungi, plants and soil

Palm seeds (*Phoenix dactylifera*) were obtained from Huerto del Cura station (Elche, Spain). The *Glomus mosseae* used as the arbuscular mycorrhiza fungi was multiplied with *Sorghum bicolor* × *sudenense* in pot cultures after 4 months of plant growth. Soil, spores and roots of the hybrid sorghum cultures were used as a crude inoculum for the Palm seedlings. *T. harzianum* T22 was supplied by "Koppert Biological Systems" (Murcia, Spain), and it was used in suspension at 1 g/L/plant concentration. The soil growth medium was a peat; this substrate was sterilized to eliminate indigenous pathogens.

2.2. Germination and growth conditions

Palm seeds were germinated in compartmented growth units of polyvinyl chloride tubes. The substrate was enriched with (15/9/15) NPK and 2 mg of Floranid Permanent. The seedlings were transplanted into pots containing the same substrate, and grown in field conditions. After 2 months, the inoculum (5 g of root and 20 g of spore and mycelium inoculum *G. mosseae*) was added to the seedlings close to the root system. After 16 months, a group of plants was irrigated three times in 1 week with *T. harzianum* T22. Every week, 20 mL of the nutritive solution of Long Ashton was added to each seedling's pot.

Four treatments were used in this experiment: noninoculated controls, plants inoculated with *G. mosseae*, plants inoculated with *T. harzianum* and plants inoculated with both fungi. For these latter, plants were first inoculated with *G. mosseae* for 16 months, then with *Trichoderma*, in order to avoid the effect of *Trichoderma* on arbuscular mycorrhizal root colonization (Siddiqui and Mohmood, 1996).

2.3. Protein extraction

Leaves and young roots were separated, washed under running tap water three times and finally with distilled water, and dried gently. Plant material was homogenized for 2 min at 4 °C in a ratio of 1:12 (w/v) in 0.1 M sodium phosphate buffer (pH 7.0) containing 20% (w/v) polyvinylpyrrolidone. Palm leaves were homogenized with the same buffer without polyvinylpyrrolidone. The homogenate was centrifuged at 37000 g for 20 min at +4 °C and the supernatant was used for enzyme assay. Protein contents were determined according to the method of Bradford (1976) using bovine serum albumin as the standard.

2.4. Determination of soluble peroxidase activities

Soluble peroxidase activity was determined spectrophotometrically in 1.7 mL reaction mixture containing 660 µL of guaiacol 12 mM, 50 µL of H₂O₂ 8 mM and 330 µL of sodium acetate buffer 0.2 M (pH 5.0) by following the increase in the absorbance at 470 nm and using $\epsilon_{470} = 5200 \text{ M}^{-1}\text{cm}^{-1}$. One unit of activity (U) was defined as the amount of peroxidase oxidating 1 µ mole of substrate/min at a temperature and pH specified for each reaction (Goldberg et al., 1983). Five enzymatic assays were carried out for each treatment.

2.5. Electrophoresis

The protein extracts were brought to 60% saturation with solid ammonium sulfate (NH₄)₂SO₄ under continuous stirring at 4 °C. After 1 h, the solution was centrifuged at 4 °C for 20 min at 120 000 g, then the pellets were resuspended in a small volume of 0.1 M sodium phosphate buffer (pH 5.0). The crude extracts were dialyzed overnight against the same buffer (sodium acetate pH 5.0) at 4 °C.

20 µL of each extract (0.02 mg of protein) were subjected to Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) with the presence of 2-mercaptoethanol of the ammonium sulfate. The pellet was on 12% acrylamide gel, according to the method of Laemmli (1970). Gels were internally calibrated using a high-molecular-mass marker kit (Sigma) containing myosine (205 KDa), β galactosidase (116 KDa), phospholipase b (97.4 KDa), bovine albumin (66 KDa), egg albumin (45 KDa) and carbonic anhydrase (29 KDa). The gels were stained for protein visualization with Coomassie Brilliant Blue.

Non-denaturing polyacrylamide gel electrophoresis (PAGE) was carried out at 4 °C on 10% acrylamide gels: both

the resolving and the stacking gels were prepared with Tris-HCl buffer pH 6.8. After the electrophoresis run, the gels were incubated for 60 min in 15 mM sodium acetate buffer containing 8 mM H₂O₂ and 0.4 M guaiacol. All electrophoresis was carried out in a mini-Protein cell (Bio-Rad).

Cationic electrophoresis

The extracts were mixed (v/v) with a buffer solution (3.1 mL stacking buffer, 5 mL glycerol, 0.5 mL methyl green (0.04%) and 1.4 mL distilled water). 20 µL of each was separated in 35 M β-alanine buffer pH 4.5. Discontinuous, non-denaturing acrylamide gel electrophoresis was prepared with 4% resolving gel (stacking buffer pH 6.8, ammonium persulfate 10%, acrylamide 30%, TEMED and water) and 10% separating gel (separating buffer pH 4.3, ammonium persulfate 10%, acrylamide 30%, TEMED and water). Gels were soaked for 30 to 60 min in 15 mM sodium acetate buffer pH 5, containing 8 mM H₂O₂ and 0.4 M guaiacol, for staining.

2.6. Visual quantification of root colonization

The root samples from the treatments with *G. mosseae* and *G. mosseae* + *T. harzianum* were cleared in a KOH (10%) solution and stained with Trypan Blue according to Phillips and Hayman (1970) using lactoglycerol. Mycorrhizal colonization was determined by the method of Trouvelot et al. (1986). An average of 150 root pieces per plant and 5 plants per treatment were examined. The results are given as percents of the root lengths infected. The presence of the mycelium of *Trichoderma* attached to the surface of the roots was checked visually under a magnifying glass.

2.7. Transmission electron microscopy and cytochemical peroxidase localization

Root and leaf samples were collected 16 months after inoculation with *G. mosseae* and 21 days after inoculation with *Trichoderma*, rinsed with distilled water, then cut into small pieces (1–2 cm) and immersed in 1.25% (v/v) glutaraldehyde in 0.1 M phosphate buffer (pH:7.3) for 1 h at 4 °C. Samples were incubated in 3,3-diaminobenzidine, H₂O₂ as substrates and 3-Amino-1,2,4-triazole inhibitor of catalase in Tris-HCl buffer (pH:7.6) for 30 min in the dark. After that, samples were washed twice for 5 min in Tris-HCl buffer (pH: 7.6) and postfixed in 1% (w/v) osmium tetroxide in the same buffer for 30 min, dehydrated in a graded ethanol series, and embedded in Epon-araldite resin and polymerized at 60 °C for 48 h.

Ultrathin sections (90 µm) were cut with a diamond knife and collected on formvar-coated nickel grids. The sections were treated with uranyl acetate and lead citrate before observations in the transmission electron microscope. For each treatment, an average of five samples from four different leaves and roots were investigated. For each sample, 5 to 10 ultrathin sections were examined.

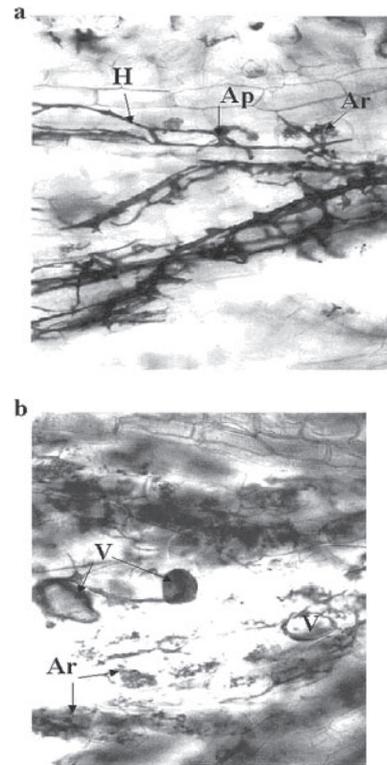


Figure 1. Root of date palm stained with trypan blue that is extensively colonised by *Glomus mosseae*. **a** and **b**: infection began by formation the appressorium (Ap), a dense group of hyphal within the cortex, intracellular fungal hyphae penetrate a root and differentiate to form arbuscules (Ar) and vesicles in cortex cells (V). The magnification of the panels (**a** and **b**) is × 200.

2.8. Statistical analysis

Significances between means were tested by two tests: Duncan's multiple range test ($P = 0.05$) and the Newman-Keuls test ($P = 0.05$). Five replications were pooled.

3. RESULTS AND DISCUSSION

3.1. Arbuscular mycorrhizal colonization

At harvest, 57 weeks after mycorrhization, *G. mosseae* allowed an improvement of the dry biomass of about 57%. The typical arbuscular mycorrhizal structures such as vesicles, arbuscules and intraradical mycelium were observed in the mycorrhizal plants (Fig. 1). The percentage of mycorrhizal colonization at harvest time varied between 25 and 30% and no mycorrhizal infection was found in control plants. This percentage of colonization was very similar to those obtained for other palm species (Jaizme-Vega and Díaz-Pérez, 1991; Morte and Honrubia, 2002). The mycorrhizae infection may introduce metabolic modifications. As far as the enzymes are proteins, any metabolic changes may affect the total protein contents.

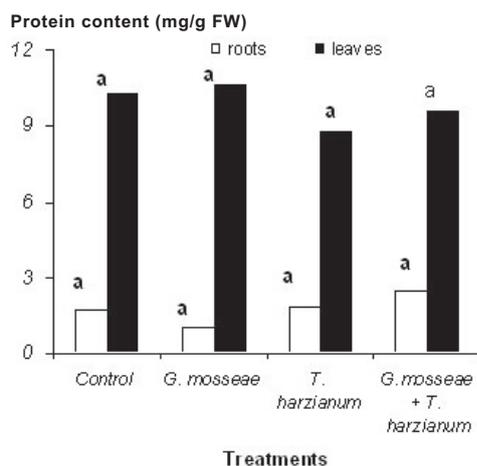


Figure 2. Protein content in roots (□) and leaves (■) of noninoculated (control), inoculated with *G. mosseae* (Gm), *T. harzianum* (Th), or with both *G. mosseae* and *T. harzianum* (Gm + Th) *P. dactylifera* plants. Plants were inoculated and grown under greenhouse conditions for 16 months with *G. mosseae* and then irrigated three times with a *Trichoderma harzianum* T22 (Koppert Biology systems) solution (1 g L⁻¹ per plant). The substrate used was a mixture of peat and vermiculite. Significances between means were tested by two tests: Duncan's multiple range test ($P = 0.05$). Five replications were pooled. Results with a similar letter are considered non significant.

3.2. Protein content

The protein content in leaves was not affected by the inoculation with *T. harzianum* and/or *G. mosseae*, but roots of plants inoculated with *G. mosseae* presented a protein content significantly lower than the leaves. In general, the protein content in leaves was 4 to 10 times higher than in roots (Fig. 2).

3.3. Peroxidase activity

In control plants, peroxidase activity (expressed as U/g FW) was 15 U/g FW, and this value was approximately 2.5 times higher in roots than in leaves (Fig. 3) and significantly lower than the activity measured in date palm and other palm species by Sakharov et al. (2001). The peroxidase activity is completely absent in the leaves of the Madagascar palm (*Chrysalidocarpus lutescens*), and it was only 50 U/g of leaf in leaves of the coconut palm (*Cocos nucifera*) (Sakharov et al., 2001). This author added that the highest peroxidase activity was detected in leaves of ruffle palm (*Aiphanes cariotifolia*), royal palm (*R. regia*), date palm (*Phoenix dactylifera*) and African oil palm (*Elaeis guineensis*) with activities of 1150, 700, 580 and 570 U per g leaves, respectively.

Peroxidase activity was higher in roots colonized by *T. harzianum* alone or in combination with *G. mosseae* than in noninoculated control plants (Fig. 3). No significant difference in peroxidase activity was observed between noninoculated control plants and plants colonized by *G. mosseae*, indicating that most of the activity was derived from the *T. harzianum*. A similar result was reported by Yedidia et al. (1999), who observed that the inoculation of *Cucumber* plants with *Trichoderma* increases peroxidase and chitinase activities

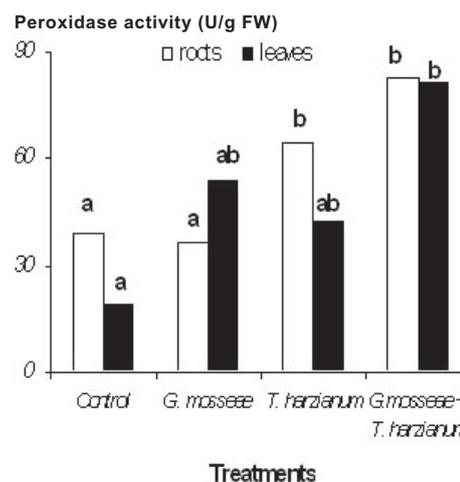


Figure 3. Peroxidase activity in roots (□) and leaves (■) of noninoculated (control), inoculated with *G. mosseae* (Gm), *T. harzianum* (Th), or with both *G. mosseae* and *T. harzianum* (Gm + Th) *P. dactylifera* inoculated. Soluble peroxidase activity was determined spectrophotometrically by following the increase in absorbance at 470 nm and using $\epsilon_{470} = 5.200 \text{ M}^{-1}\text{cm}^{-1}$. Significances between means were tested by two tests: Duncan's multiple range test ($P = 0.05$). Five replications were pooled. Results with a similar letter are considered non significant.

both in roots and leaves. The delay between plant inoculation with *G. mosseae* and the determination of peroxidase activities (16 months and three weeks) could explain the fact that the measured peroxidase activities showed similar values compared with the control plants. Spanu and Bonfante-Fasolo (1988) observed that higher cell wall peroxidase activities occur at the beginning of symbiotic establishment of *Glomus versiforme* in garlic plants; at more advanced stages these activities decreased to the same level as in nonmycorrhizal plants. Transient increases in peroxidase activity have also been detected in root cell walls of *Allium porrum* during early stages (16 days) of colonization with the vesicular-arbuscular mycorrhizal fungus *G. versiforme*; after that (50 days), there was a decrease in activity which reached the same level as that of noncolonized plants (Spanu and Bonafante-Fasolo, 1988). Mycorrhizal fungi increased peroxidase and polyphenoloxidase activities in *Ziziphus xylopyrus* roots (Mathur and Vyas, 1996). The increased peroxidase activity by mycorrhizal fungi may be due to an increased P uptake resultant from the symbiosis. McArthur and Knowles (1992) reported a positive correlation between root phosphorus uptake and peroxidase activity.

Since the "infection development stage" may influence the level of peroxidase, the observed increase in peroxidase activity may likely be attributed to the *Trichoderma* infection rather than the mycorrhizal establishment. On the other hand, Salzer and Hager (1993) described the suppression of the elicitor induction of peroxidase by auxins. As many mycorrhizal fungi produce auxin (Münzenberger et al., 1997), it is conceivable that peroxidase activity is suppressed by fungal auxin in the intact system. The long time needed (at least three months) to get a good development of mycorrhizal infection

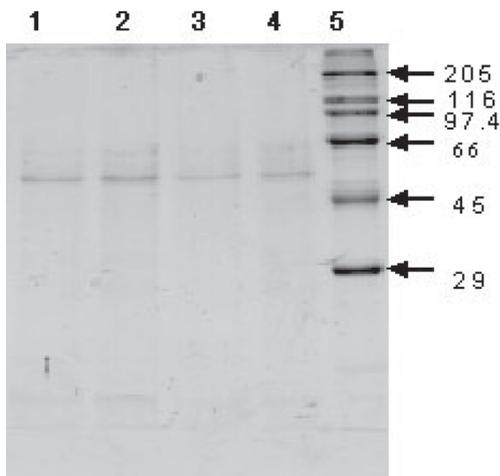


Figure 4. SDS-PAGE of palm leaf extracts: (1) 20 μ L of plants inoculated with *G. mosseae*; (2) 20 μ L of plants inoculated with *T. harzianum*; (3) 20 μ L of control noninoculated plants; (4) 15 μ L of plants inoculated with *G. mosseae* and *T. harzianum* T22; (5) 10 μ L of molecular markers. Gels were internally calibrated using a high-molecular-mass marker kit (Sigma SDS-6H) containing: myosine (205 kDa); β -galactosidase (116 kDa); phospholipase b (97.4 kDa); bovine albumine (66 kDa); albumine egg (45 kDa) and carbonic anhydrase (29 kDa). The gels were stained for protein visualization with Coomassie Brilliant Blue.

in *Phoenix* species (Morte and Honrubia, 2002; Yedidia et al., 1999; Dreyer et al., 2001) makes the measurement of the levels of peroxidase activity difficult at the early stages of mycorrhizal colonization.

In leaves, significant differences were observed only between noninoculated control plants and plants colonized by both fungi (Fig. 3). This increase suggests the existence of a signal that is transported from the infection sites throughout the whole plant, inducing the synthesis of defensive enzymes such as peroxidase. A similar response has been described in tomato (Bergey et al., 1996). To approach the overall metabolic modifications a separation of the native total protein is fulfilled.

3.4. Electrophoresis method

The SDS-PAGE analyses of palm leaf extracts gave a main band at 54 kDa for all the treatments (Fig. 4). This band could correspond to the peroxidase, as the estimated molecular mass is comparable with that found by Sakharov et al. (2000) for *P. dactylifera* peroxidase. However, further purification is necessary to ascertain that this band corresponds to peroxidase. Using native cationic electrophoresis, several basic peroxidase isoenzymes were detected in roots for all fungal treatments and control (Fig. 5). The electrophoretic pattern was different in control plants with respect to fungal inoculated plants. When native anionic electrophoresis was carried out at pH 6.6, two bands corresponding to acid peroxidase isoenzymes were observed in roots of *T. harzianum*-infected plants and *G. mosseae*-colonized plants. One of these bands was not present in plants inoculated with both *T. harzianum*

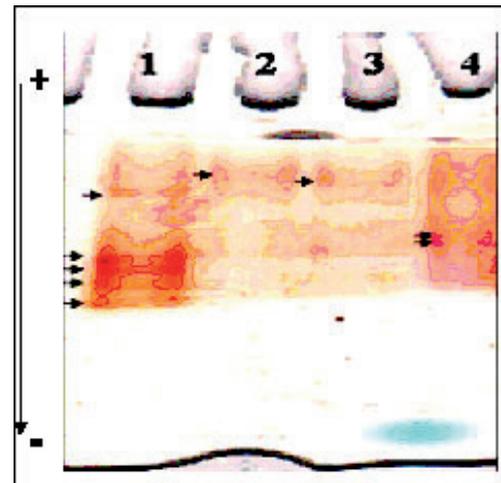


Figure 5. Native cationic electrophoresis in roots of palm extracts (1) control (2) plants inoculated with *G. mosseae*; (3) plants inoculated with *T. harzianum* T22; (4) plants inoculated with *G. mosseae* and *T. harzianum* T22. Peroxidase was visualized following the method described by Reisfeld and co-workers and modified by Escribano et al.

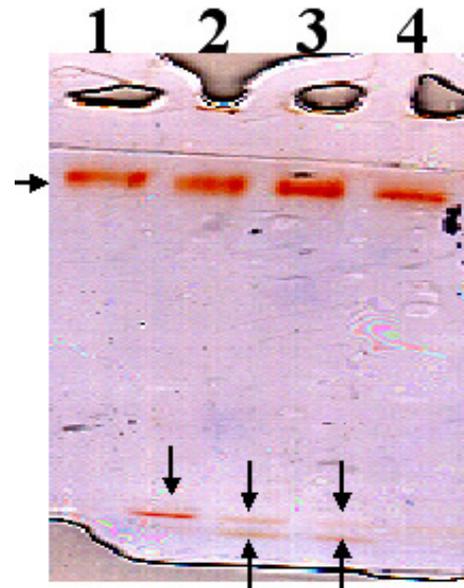


Figure 6. Native anionic electrophoresis in roots of palm extracts (1) Control (2) plants inoculated with *G. mosseae* and *T. harzianum* T22; (3) plants inoculated with *T. harzianum* T22; (4) plants inoculated with *G. mosseae*. The band corresponding to acid peroxidase isoenzymes is absent in plants control.

and *G. mosseae* and both of them were absent in control plants (Fig. 6). More experiments have to be carried out to clarify these results.

3.5. Histochemical studies

T. harzianum T22 penetrated into the epidermal root cells. The *T. harzianum* hyphae were characterized by a high

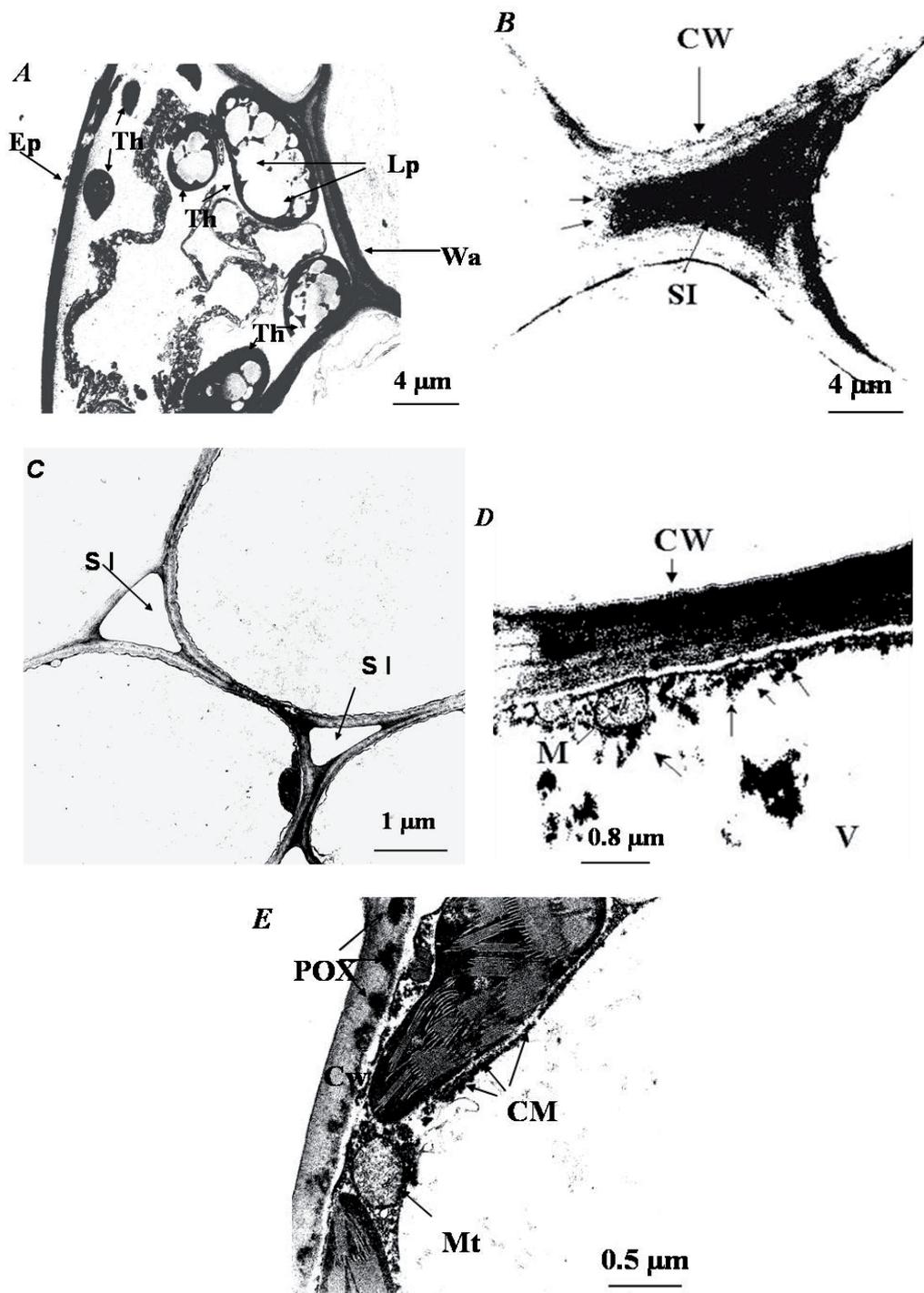


Figure 7. (A) Epidermal root cell of *P. dactylifera* colonized by *T. harzianum* hyphae (Th), lipid vesicles (Lp), wall appositions (Wa); (B, D) Cortical root cell of *P. dactylifera* palms inoculated with both *T. harzianum* and *G. mosseae* fungi: electron-dense deposits indicated a peroxidase activity. These were observed on the internal side of the cell walls (CW), at the intercellular spaces (IS) and vacuole (V); (C) Cortical root cell of *P. dactylifera* control without peroxidase activity at the intercellular spaces (IS); (E) Leaf cell of *P. dactylifera* palms inoculated with both *T. harzianum* and *G. mosseae*: peroxidase activity (POX) was found in the cell walls (CW), mitochondria membrane (Mt) and cytoplasm membrane (T).

electron density of their cytoplasm and numerous lipid vesicles (Fig. 7A). The formation of heterogeneous wall appositions was observed in the noninfected host cell adjacent to infected cells. They were elongated deposits along a large portion of the host cell wall (Fig. 7A). A similar observation was reported by Yedidia et al. (1999) with cucumber roots infected by *Trichoderma harzianum* T22. It was suggested that host cells produce signals to mobilize defense strategies such as the accumulation of heterogeneous wall appositions, and this dense material originated from an aggregation of the host cytoplasm, or was newly synthesized as a response to infection (Yedidia et al., 1999).

Peroxidases are normally present in roots, leaves and stems of plants. These enzymes can be found in vacuoles (Pedreño et al., 1993), the apoplast (Takahama and Oniki, 1992), the plasmalemma, and inside and outside the cell wall (Pedreño et al., 1995; Mäder, 1992). In *Phoenix dactylifera* peroxidase activity was observed on the internal side of the cell walls, and in the intercellular spaces (Fig. 7B) and the vacuole of palm root inoculated with both fungi, *G. mosseae* and *T. harzianum* (Fig. 7C). In the leaves of the same treatment peroxidase was localized in the cell wall, the host plasmalemma, the tonoplast and the chloroplast membrane (Fig. 7D).

4. CONCLUSION

Mycorrhization allowed an improvement of the dry biomass. Quantitative and qualitative peroxidase study showed significant differences between controls and plants inoculated with *Trichoderma harzianum* T22 alone or combined with *Glomus mosseae*. The inoculation with *Trichoderma* on already mycorrhized palm date seedlings seems to induce a high peroxidase activity in leaves and roots. Peroxidase cytochemical localization was established in the intercellular regions, cell walls, cytoplasm and in chloroplastic and mitochondrial membranes. The stimulatory effect of *Trichoderma* on peroxidase activity production could be one of many resistance mechanisms of plants to pathogens. The combination of mycorrhiza and *Trichoderma* showed synergetic effects.

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