Research Article

Efficiency of secondary metabolites produced by *Trichoderma* spp. in the biological control of *Fusarium* wilt in chickpea

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Abstract: The present study was aimed to investigate the effects of secondary metabolites produced by five Trichoderma spp. on the control of Fusarium wilt caused by Fusarium oxysporum f. sp. ciceris (FOC) in chickpea. In vitro biocontrol potentialities of Trichoderma spp. against FOC was tested. Trichoderma secondary metabolites were extracted by solvent extraction methods and evaluated against FOC. In vitro tests showed very good inhibitory effects by all Trichoderma spp. against FOC along with an inhibitory rate up to 73.8% and 27.8%, for direct and indirect contacts, respectively. Additionally, Trichoderma spp. caused a significant decrease in Fusarium wilt disease severity, in particular, T. polysporum showing 64.2% of disease severity reduction. The tested secondary metabolites were also effective against FOC with a significant decrease of mycelial growth from 6% to 76.9%. Similarly, in vivo tests revealed that secondary metabolites were very active in reducing disease severity. It was found that T. polysporum was the most active with 56.9% of disease severity reduction. Chickpea resistance is mostly attributed to polyphenolic compounds. The studied Trichoderma spp. and their secondary metabolites could be used as potential and promising antifungal agents in preventing the occurrence of Fusarium wilt in chickpea.

Keywords: Biological control, *Trichoderma*, *Fusarium oxysporum*, secondary metabolites, Chickpea

Introduction

Chickpea, *Cicer arietinum* L., is one of the most important food legumes worldwide (Sharma *et al.*, 1994). The chickpea cultivation around the world is economically important because of its high nutritional value richness in essential proteins (Al-Snafi, 2016), therapeutic effect (Jukanti *et al.*, 2012) and agricultural interest in biological nitrogen fixation (Gaur and Sen, 1979). Unfortunately, this crop faces

several biotic constraints, especially pathogenic fungi, affecting its yield and quality. Vascular wilt caused by Fusarium oxysporum f. sp ciceris (FOC) has been reported as the most destructive disease of chickpea cultivation in the world. The annual yield losses, caused by the disease were estimated at 10 to 15% (Jiménez-Díaz et al., 2015). However, this pathogen can completely destroy the crop under the favorable conditions for disease development (Trapero-Casas and Jim'enez-D'1az, 1985).

In general, the disease can be reduced by means of chemical, biological or physical disease control methods (Jiménez-Díaz *et al.*, 2015). However, the intensive pesticide



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applications in the conventional production, mainly with fungicides can initiate many emerging problems, including environmental pollution, development of fungicide-resistant pathogens in addition to inducing variety of human and animal health problems (You et al., 2016). Recently, a great research interest has been paid to the alternative management measures, particularly the fungal biological control agents (BCAs) for the protection of worldwide crops. So far, the antagonistic fungi are mainly known by their potential ability to decrease inoculum density of pathogenic fungi (Dennis and Webster, 1971a), and among which fungi of the genus Trichoderma, as biocontrol agents (BCAs), are successfully used as biopesticides worldwide (Vinale et al., 2008). Furthermore, tested Trichoderma the formulations on biological management have proved satisfactory for control of vascular disease agents, eg, Fusarium oxysporum (Vinale et al., 2009). However, these types of formulations have certain disadvantages, such as limited shelf life, high dose efficacy, and present low stability in an adverse environment (Keswani et al., 2014). Since the secondary metabolites are a promising solution for these problems, the conventional formulations of biopesticides can be either improved or replaced, and thus the next-generation of secondary metabolites based formulations may be developed for management of phytopathogens (Keswani et al., 2014). In addition, many species of Trichoderma are commonly able to produce secondary metabolites with a good antibiotic activity (Vinale et al., 2008, 2009, 2012). The present work, therefore, aimed to provide further understanding on the biological control agents belonging to the genus of Trichoderma and their secondary metabolites, targeting the FOC. This biological control model would be based on analyses of these microorganisms, using i), in vitro and in vivo studies of the antagonistic effect of some Trichoderma spp. against FOC, ii), the extraction of secondary metabolites and in vitro as well as in vivo determinations of their antifungal activity against FOC.

Materials and Methods

Fungal strains

The biocontrol agents *Trichoderma harzianum*, *T. viride*, *T. polysporum*, *T. virens*, and *T. atroviride* were isolated from soil rhizosphere of chickpea. *Trichoderma* spp. were identified based on visual macroscopic and microscopic observations according to Gams and Bissett, (1998). Chickpea pathogen (*Fusarium oxysporum* f. sp. *ciceris*) (FOC) was obtained from the infected plant parts of chickpea collected in the region of Mascara, North-Western Algeria.

In vitro effect of *Trichoderma* spp. on FOC Dual and opposite culture test

The dual culture test was performed in Petri dishes (9 mm in diameter) containing PDA medium as described by Dennis and Webster (1971b). Mycelial plugs (5 mm in diameter) of *Trichoderma* spp. and FOC taken from advancing edge of 7-days PDA culture were placed at equal distance from the periphery (2 cm). Inoculated plates were incubated at 25 ± 3 °C. Control without *Trichoderma* was used. The radial growth inhibition of FOC by *Trichoderma* spp. were determined and compared with the control. Four replications were achieved for each species.

Microscopic examination of the hyphal interactions in dual culture plates was performed. The method was done by taking small portion from the contact hyphal region between *Trichoderma* spp. and FOC on a glass slide and mounted in methylene blue (El-Debaiky, 2017). The contact regions between the antagonistic *Trichoderma* spp. and FOC were investigated using an optical microscope.

In another experiment, the five *Trichoderma* spp. were evaluated for their volatile inhibitory effect according to the techniques described by Dennis and Webster (1971a). In brief, 5mm discs of *Trichoderma* spp. from 7-days old cultures were transferred to the center of PDA Petri dishes, the top of each Petri dish was replaced with bottom of a PDA plate inoculated centrally with FOC, and then sealed with Parafilm for incubation at 25 ± 3 °C. The same setup without *Trichoderma* spp. was used as

control. Four replications were maintained for each species. The diameter of pathogen colony was measured 4, 6 and 8 days after incubation and the inhibition of mycelial growth were determined.

Extraction and evaluation of secondary metabolites against FOC

Isolation of secondary metabolites produced by Trichoderma spp. were carried out as previously described (Vinale et al., 2008). Briefly, two 7 mm diameter plugs of each Trichoderma strain were obtained from actively growing margins of PDA cultures and inoculated to 2 L conical flasks containing 250 mL of sterile potato dextrose broth (PDB). Then the stationary cultures were incubated for 31 days at 25 ± 3 °C, followed by vacuum filtration onto filter paper. The filtered culture broth (2 L) of Trichoderma spp. was after wards extracted exhaustively with ethyl acetate. The antifungal activity of the Trichoderma secondary metabolites was tested against FOC following the direct solubilization in PDA medium (Dubey et al., 2011). Secondary metabolites derived from five species of Trichoderma various were tested at concentrations (50, 100, 250, 500 and 1000ppm) prepared by DMSO (3%).The medium was poured into Petri dishes after flask shaking, and then pathogen plugs were centrally inoculated by making 5mm discs taken from 7-days old culture on the PDA plates. Control received the same quantity of DMSO (3%) mixed with PDA. Also, four replicate plates were tested. The pathogen growth was determined daily by measuring the colony diameter.

The percentage of mycelial growth inhibition in all above experiments was calculated by the formula:, $MGI(\%) = \frac{(C-T)}{C} \times 100$, where MGI:

percent growth inhibition, C = growth in control and T = growth in treatment.

In vivo effect of *Trichoderma* spp. on *Fusarium* wilt severity

The preparation of FOC inoculum was performed according to the technique of Nene and Haware (1980). A mixture containing sieved sand (90 g) and corn flour (10 g) moistened with distilled water (20 ml), according to the proportion of 9/1/2 (w/w/v), respectively, was prepared in plastic bags. The mixture was sterilized three times at 121 °C for 30 min and then inoculated with 25 discs of 5 mm diameter taken from an 8-days FOC culture. The incubation was carried out for 21 days at 25 ± 3 °C. Agitation of the bags every day was carried out in order to allow a homogeneous colonization of the medium by FOC. The obtained inoculum was then incorporated into pots containing the sterilized culture substrate at the rate of 100 g of inoculum per 1 kg of substrate, which is composed of a mixture of sand, soil, and organic matter in the proportions of 1/1/1(v/v/v) sterilized at 121 °C for 24 h.

Chickpea line (Guab 5) seeds were surface disinfected using sodium hypochlorite (1%) for 3 min and rinsed in sterile water. Coating of chickpea seeds with the five species of Trichoderma was carried out according to the method of Mohammad et al. (2011). The chickpea seeds were immersed in 10 ml of the Trichoderma spore suspension at the concentration of 5×10^8 spore/ml for 30 min and placed (10 seed/ dishes) in sterile Petri plates containing two sterile filter papers. Then Petri dishes containing seeds were incubated at 25 ± 3 °C, and the 8-day old plants coated with spore suspension of Trichoderma were carefully transferred into the soil inoculated with FOC.

Severity of *Fusarium* wilt disease was assessed at 2 day intervals, on a scale ranging from 0 to 4 according to the percentage of foliage with yellowing or necrosis, where 0 =0%, 1 = 1 to 33%, 2 = 34 to 66%, 3 = 67 to 100%, and 4 = dead plant). Also, the evaluation was carried out according to the method of Traperos Cases and Jim'enez-D'1az (1985), since disease incidence (DI) was assessed by counting the number of plants showing symptoms. Hence, incidence symptoms (0 or 1) and severity data (ranged from 0 to 4) were used to calculate disease index intensity (*Dis*) using the equation *Dis* = (I × S)/4. In addition, the AUDPC value was mainly used to estimate the area under disease progress curve of each treatment:

$$AUDPC = \sum_{i+1}^{n} \left[(x_i + x_{i+1}) / 2 \right] (t_{i+1} - t_i)$$

In vivo effect of secondary metabolites produced by *Trichoderma* spp.

To test the in vivo effects of Trichoderma extracts, 1000 ppm of secondary metabolites was prepared in 1 L of sterile distilled water. Chickpea line (Guab 5) seeds were surface disinfected for 3 min using sodium hypochlorite (1%), rinsed in sterile water for 3 min and then germinated (10 seed/ disc) for 8 days in Petri plates containing 2 sterilized filter paper. Petri plates containing seeds were incubated under laboratory conditions (25 \pm 3 °C). Germinated seed were transferred to peat substrate, sterilised at 121 °C/ 30 min. Two-weekold roots of chickpea plants grown on peat were immersed in each extract to ensure better application and carefully transferred into plastic pots of 20 cm diameter and 30 cm height) filled with the substrate composed of the mixture soilsand-compost (1v: 1v: 1v), sterilized in autoclave for 40 min at 120 °C and then sprayed again by the different Trichoderma extracts. Five seedlings were planted per pot. Two control treatments were used: (i) negative control pots uninoculated with FOC and treated with sterile distilled water; (ii) positive control pots inoculated with FOC and treated with aqueous DMSO (3%).

Determination of total polyphenol and flavonoid contents

Total polyphenol content was evaluated according to the Folin–Ciocalteu procedure as described by Ardestani and Yazdanparast (2007). Briefly, a host plant extract was prepared by adding10 ml of 80% methanol to 250 mg of the dried-milled chickpea under a slow shaking. The obtained solution was filtered and 0.5 ml of the methanolic extract was mixed with 2.5 ml of Folin-Ciocalteu's reagent (1:10 diluted with distilled water) and 2 mL of 7.5% Na₂CO₃ solution in a test tube under shaking. Thereafter, the mixture was incubated at 30 °C in a hot water bath for 90

min, and the absorbance of the mixture was measured 765 nm using at а spectrophotometer. Total polyphenol content was given as milligram gallic acid equivalents/g dried extract, using the blank sample composed of water and reagents. All the measurements were replicated four times.

Total flavonoid content was evaluated according to the aluminum chloride colorimetric method (Chua *et al.*, 2011). A volume of 1 ml of the methanolic extract was added to 2 ml of the methanolic solution containing 2% AlCl₃. The absorption of pink color of mixture was measured after 15 min at 430 nm. Total flavonoid content was given in milligram of quercetin equivalents (QE) per gram of extract.

Statistical Analysis

In the present study each experiment was performed four times and displayed as mean \pm SE. The statistical analysis was performed by one and two-way analysis of variance (ANOVA). Comparisons between results of each treatment were compared with Tukey post hoc test (p < 0.05). Statistical tests were performed using software package STATISTICA 8.

Results

In vitro antagonistic activity of *Trichoderma* spp. against FOC

Results of *in vitro* tests showed that *Trichoderma* spp. exhibited strong inhibitory effects on mycelial growth of FOC (Fig. 1). We noticed significant differences in terms of inhibitory activity among the tested plates (Fig. 2). After eight days, the highest inhibitory effect was obtained with *T. harzianum* and *T. polysporum* (83.47% and 83.05%, respectively).Whereas, *T. viride*, *T. atroviride* and *T. virens* showed an inhibition activity which ranged from 73.81% to 77.68%.

Microscopic observations of the contact zone between the two protagonists showed profound changes in the mycelium of FOC manifested by massive winding of *Trichoderma* mycelium over that of FOC, transformation in cords of the mycelial filaments, lysis of pathogen mycelia, vacuolation and early aging by wall thickening and chlamydospore formation (Fig. 3).

The obtained results revealed an inhibitory effect of volatile substances on the mycelial growth of FOC as compared to control (Fig. 2). Here, FOC was found to be more sensitive to the volatile substances of *T. harzianum*, at 8 days of incubation, leading subsequently to maximum mycelial growth inhibition of FOC (54.79%). However, the lowest rate of mycelial growth inhibition was observed with *T. atroviride* (27.86%).

In vitro effects of *Trichoderma* spp. secondary metabolites

The *in vitro* antifungal activity of *Trichoderma* spp. secondary metabolite was investigated

against FOC (Fig. 4). We noticed significant differences in antifungal activity for the tested concentrations of secondary metabolites ($p \leq$ 0.05), in addition to the different degrees of antifungal activity of each species (Table 1). In this study, the incorporation method showed a significant reduction of mycelial growth of FOC and the rate of reduction was gradually increased by increasing the concentration of extract. T. polysporum and T. harzianum were the most effective in inhibiting FOC mycelial growth for all tested concentrations. The level of mycelial growth inhibition by T. polysporum ranged from 45.8% to 76.94%. Trichoderma virens showed the lowest antifungal activity with percentages ranging from 6.08% to 33.94%.

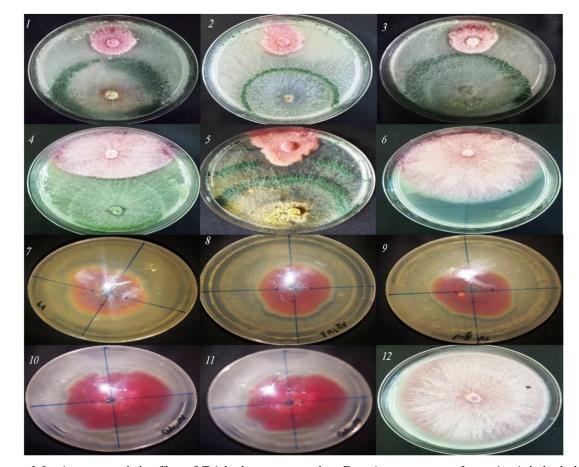


Figure 1 *In vitro* antagonistic effect of *Trichoderma* spp. against *Fusarium oxysporum* f. sp. *ciceris* in both dual culture (1 to 6) (1- *Trichoderma viride*, 2-*T. harzianum*, 3- *T. polysporum*, 4- *T. virens*, 5- *T. atroviride*, 6-Control) and opposite cultures (7 to 12) (7- *Tricoderma viride*, 8-*T. harzianum*, 9- *T. polysporum*, 10- *T. virens*, 11- *T. atroviride*, 12- Control).

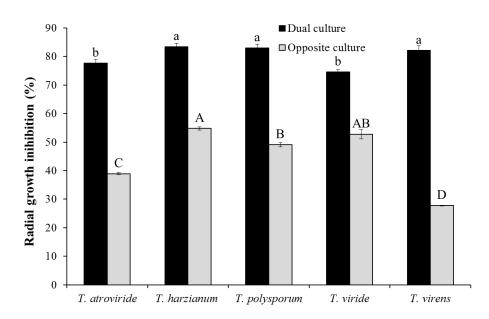


Figure 2 Inhibition of *Fusarium oxysporum* f. sp. *ciceris* mycelial growth in dual and opposite culture. Values represent the mean of four replicates \pm SE. Data marked by different letters in a bar indicate significant difference (Tukey's test, P < 0.05).

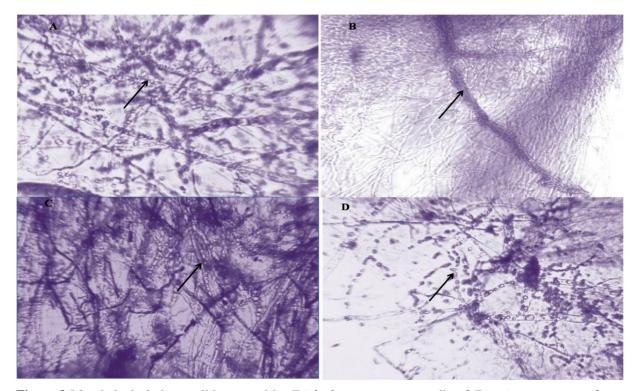


Figure 3 Morphological abnormalities caused by *Trichoderma* spp. on mycelia of *Fusarium oxysporum* f. sp. *ciceris* (8 days treatments). A) Massive winding of *Trichoderma* mycelium over that of *Fusarium oxysporum* f. sp. *ciceris*, B) transformation in cords of the mycelial filaments, C) vacuolation, D) lysis of pathogen mycelia.

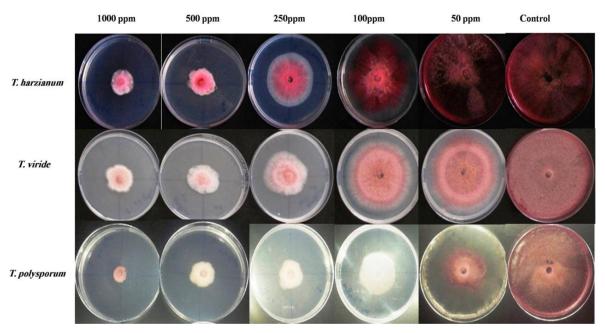


Figure 4 Effects of different concentrations of *Trichoderma* spp. secondary metabolites on mycelial growth of *Fusarium oxysporum* f. sp. *ciceris* after 8 days of incubation at 25 ± 3 °C. Control was treated with DMSO (3%).

Table 1 Effect of secondary metabolites isolated from culture filtrates of various species of *Trichoderma* on mycelial growth of *Fusarium oxysporum* f. sp. ciceris.

Species	Mycelial growth inhibition (Mean \pm SE) (%)				
	1000 ppm	500 ppm	250 ppm	100 ppm	50 ppm
T. harzianum	71.69 ± 0.35^{ab}	60.17 ± 2.07^{bcde}	52.99 ± 1.06^{def}	17.94 ± 2.79^{klmn}	8.98 ± 1.72^{mno}
T. polysporum	$76.94 \pm 1.09^{\rm a}$	68.54 ± 2.08^{abc}	61.33 ± 4.46^{bcd}	$53.29 \pm 1.79^{\text{def}}$	45.8 ± 2.47^{efgh}
T. viride	53.88 ± 0.92^{cdef}	$48.20 \pm 1.22^{\text{defg}}$	40.75 ± 6.16^{fghi}	21.53 ± 2.90^{jklm}	8.38 ± 2.95^{mno}
T. atroviride	52.83 ± 3.21^{def}	$32.29 \pm 1.87^{\text{hijk}}$	$13.39\pm2.13l^{mno}$	4.70 ± 0.34^{no}	$00\pm00^{\circ}$
T. virens	33.94 ± 7.59^{ghij}	26.79 ± 2.53^{ijkl}	$14.75\pm1.6l^{mno}$	6.08 ± 0.61^{no}	$00\pm00^{\rm o}$

Values represent the mean of four replicates \pm SE (standard errors). Data marked by different letters in a column indicate significant difference (Tukey's test, P < 0.05).

In vivo antagonistic activity of *Trichoderma* spp. and their secondary metabolites on *Fusarium* wilt severity

The effect of the different *Trichoderma* spp. on severity of *Fusarium* wilt, achieved after 40 days of treatment, are displayed in Figures 5-6. Chickpea plants treated with different species of *Trichoderma* exhibited AUDPC values ranging from 730.63 to 1468.88, whereas the untreated control showed an AUDPC value of 2045, which is significantly higher than all treated plants. Importantly, the *in vivo* study showed that *T*.

polysporum, T. harzianum and T. viride were the most effective against wilt disease in which the AUDPC values were significantly reduced (P \leq 0.05). The mean values of the AUDPC were 730.63, 835.25 and 960 representing 64.27%, 59.15% and 53.05% of disease severity reduction, respectively. Treatment of chickpea plants with T. virens and T. atroviride showed the lowest effect against Fusarium wilt severity.

Plants treated with secondary metabolites of different *Trichoderma* spp. showed AUDPC values ranging from 880.5 to 1329, which related

to 56.94% and 35.01% of reduction in disease severity, respectively, compared to the AUDPC of positive control (2045) (Fig. 6). *Trichoderma polysporum* and *T. viride* secondary metabolites showed AUDPC values of 880.5 and 942.75 which correspond to 35.01% and 47.33% of disease reduction, respectively.

Determination of total polyphenolic and flavonoid content in plants treated with *Trichoderma* spp.

Plants treated with all Trichoderma spp. induced production of polyphenol compounds in chickpea plants with rate ranging from 25.57 to 34.77 mg/g, while positive and exhibited negative controls average polyphenol levels of 9.25 and 23.29 mg/g, respectively. The highest level of polyphenols was recorded in the plants treated with T. harzianum (34.77 mg/g) (Fig. 7A). In terms of flavonoids, plants treated with various Trichoderma spp. also showed high values of flavonoid contents ranging from 7.93 to 13.22 mg/g, which are significantly higher than those of the positive and negative controls

(1.57 and 5.45 mg/g, respectively). Here too, the highest levels of flavonoids were recorded in plants treated with *T. harzianum* (13.22 mg/g) (Fig. 7B).

To explore the potential effect of secondary metabolites of Trichoderma spp. in inducing plant defense system, polyphenolic and flavonoid contents were measured in chickpea seedling treated with different secondary metabolites of Trichoderma spp. (Fig. 7). Marked decreases in polyphenol and flavonoid contents of seedling treated with secondary metabolites were noticed. As shown in figure 7A, plants treated with secondary metabolites of T. polysporum and T. harzianum showed high polyphenol levels (36.2 and 27.8 mg/g, respectively). Lower values were recorded for T. atroviride 18.7 mg/g. Flavonoid contents were found to vary within plants treated with secondary metabolites of Trichoderma spp. (Fig. 7B). The high values were recorded in plants treated with secondary metabolites of T. viride (9.09 mg/g), whereas, the low values were found in plants treated with T. harzianum (7.48 mg/g).



Figure 5 Effects of *Trichoderma* spp. (1- *Trichoderma harzianum*, 2- *T. polysporum*, 3- *T. atroviride*, 4- *T. viride*, 5- *T. virens*, 6- Control) on the reduction of disease severity caused by FOC.

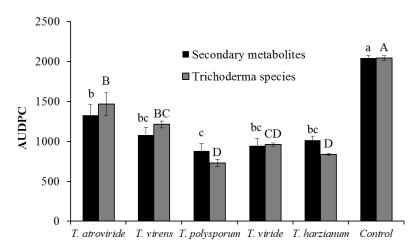


Figure 6 Effects of *Trichoderma* spp. and their secondary metabolites on the reduction of *Fusarium* wilt disease severity. Values represent the mean of four replicates \pm SE. Data marked by different letters in a bar indicate significant difference (Tukey's test, P < 0.05).

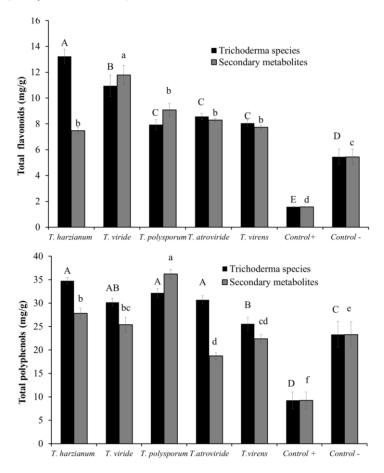


Figure 7 Total polyphenol and flavonoid contents in chickpea seedlings treated with different species of *Trichoderma* spp. and their secondary metabolites. Values represent the mean of four replicates \pm SE. Data marked by different letters in a bar indicate significant difference (Tukey's test, P < 0.05).

Discussion

The main objective of this study was to investigate biological control of chickpea against FOC through application of different species of Trichoderma spp. Our results demonstrated effect the potential of secondary Trichoderma spp. and their metabolites in decreasing the severity of wilt disease of chickpea. Several studies have demonstrated the successful use of Trichoderma spp. as biological control agents against a wide range of economically important soil phytopathogens, particularly borne Sclerotinia sclerotiorum (Zhang et al., 2016), Verticillium dahliae, Penicillium verrucosum, Aspergillus carbonarius (Kumar et al., 2014) and Fusarium oxysporum f. sp ciceris (Dubey et al., 2007).

In this work, five species of Trichoderma showed an important inhibitory zone of mycelial growth of FOC at about 73.81% to 83.47% in dual culture experiments. Microscopic observations of the contact zone between Trichoderma spp. and FOC showed a marked alteration in the mycelium of FOC, evidenced by an important lysis, transformation in cords of the mycelial filaments and a coil of the Trichoderma mycelium on FOC. These results are in agreement with previous studies (Saravanakumar et al., 2016; Toghueo et al., 2017). El-Debaiky, Our results 2016; corroborate those of Dubey et al. (2007) who reported that T. viride, T. harzianum and T. virens inhibited FOC mycelial growth from 50 to 60% in dual culture. Moreover, Zhang et al. (2016) reported that T. harzianum inhibited mycelial growth of S. sclerotiorum with an effectiveness of 56.3% in dual culture tests. Nonetheless, T. atroviride and T. harzianum caused over 28.8% inhibition of mycelial growth of Fusarium oxysporum f. sp. Phaseoli in dual culture (Otadoh et al., 2011).

The result of the present study revealed that five *Trichoderma* spp. could significantly inhibit mycelial growth of FOC in opposite culture. Thus, we could also anticipate that the antagonistic *Trichoderma* inhibited the mycelial growth of FOC through production of nonvolatile substances. These are in agreement with those obtained by Dubey *et al.* (2007), who reported that *T. viride*, *T. harzianum* and *T. virens* showing a highest ability to inhibit mycelial growth of FOC from 50 to 60% in indirect tests. The volatile metabolites from various *Trichoderma* spp. were effective in inhibiting *S. sclerotiorum*, *F. solani* and *R. solani* growth with inhibitory zone ranging from 33 to 71% (Qualhato *et al.*, 2013).

The antagonistic activity of Trichoderma as biocontrol agent against various plant diseases can be achieved by a number of bioactive compounds including secondary metabolites (Zeilinger et al., 2016; Li et al., 2018). In the Trichoderma present study. secondary metabolites showed a significant reduction of mycelial growth of FOC. It was found that the rate of reduction gradually increased by increasing the concentration of secondary metabolites. The observed variation in fungicidal activity among the five Trichoderma spp. could be attributed to the different types of chemical compounds produced in the species. As reported, Trichoderma spp. secreted various volatile and non-volatile compounds like, alkylpyrones, polyketides, sesquiterpenes, peptaibols, siderophores and terpenoids which are crucial factors for mycoparasitism and antibiosis by Trichoderma spp. (Zeilinger et al., 2016). These substances involve a very large variety of action mainly referred to a function of their chemical composition. In this regard, Vinale et al. (2013) reported that Harzianic acid is a T. harzianum secondary metabolite having antifungal polysporum activities. Also. Т. produces common families of secondary metabolites, such trichopolyns, alamethicins, hypelcins, as trichosporins, trichocellius, trichokindins and aibellin, possessing variety of biological membrane activities derived from their modifying properties, such as the formation of voltage-gated ion channels, hemolysis, uncoupling of oxidative phosphorilation (New et al., 1996; Iida et al., 1999). Obtained results corroborate those of Dubey et al. (2011) who demonstrated that T. viride, T. virens and T.

harzianum secondary metabolites exhibited pronounced antifungal activity against FOC ranging from 47.8% to 78% at 75 ppm. Similar effect was recorded by Toghueo *et al.* (2016) when investigating the antagonistic effects of secondary metabolites secreted by *T. atroviridae* at different concentrations showing a high mycelial growth inhibition of *F. solani*. Moreover, Bae *et al.* (2016) reported significant inhibitory effects of the metabolites of *T. atroviride* and *T. virens* against *Phytophthora* isolates. It is worth noting that in this study, only the concentrations of 3, 50 and 100 µl/ml of bioactive extracts of *Trichoderma* significantly increased fungicidal effect against FOC.

The results from this study demonstrated that Trichoderma spp. exhibited antifungal activity in vivo and reduced severity of Fusarium wilt of chickpea. The efficacy of Trichoderma spp in biocontrol of Fusarium wilt could be attributed to the reduction of soil fungal populations and induction of host resistance, thereby confirming the antifungal activity of Trichoderma spp. in vivo. Indeed, the biocontrol Trichoderma agents present numerous advantages to their linked host plants which are useful in combating biotic stresses. A wide range of biologically active metabolites are deployed by Trichoderma spp. for pathogen exclusion such as antibiosis (Sharma et al., 2017). It is assumed that *Trichoderma* spp. the production secondary induced of metabolites causing direct antimicrobial effect, or indirect defense by stimulating host plant defense (Vinale et al., 2012). Therefore, the results of the experiments indicated Trichoderma secondary metabolites are directly implicated in the establishment of plant defense genes, as previously reported for peptaibols (Viterbo et al., 2007). In particular, peptaibols induced an over-expression of defense related genes implicated in the systemic responses of plant (Vinale et al., 2008).

One of the main goals of this work was to examine the potential involvement of secondary metabolites in the induction of systemic resistance through *Trichoderma*-plant interaction. Biochemical analysis revealed a strong accumulation of polyphenols and flavonoids in the plant treated with various Trichoderma spp. and their secondary metabolites. This is likely due to the positive effect of Trichoderma spp. in reducing the severity of chickpea wilt disease. Interestingly, polyphenol compounds play a major role owed to their accumulation, basically known as important compounds in resistant plants and mav be involved in the crosslinking, suberification and lignifications aiming in particular to limit the action of compressive forces and that of parasite hydrolases (Clérivet et al., 1996). As previously reported, phenolic and flavonoid compounds were identified to be involved in cell wall lignifications and in the decrease of Fusarium wilt severity by limiting pathogen penetration (Jin et al., 2012). Mona et al. (2017) also reported that T. harzianum increased the rate of phenol and flavonoid contents in tomato seedling. A very recent study showed accumulation of 25 abiotic and biotic stress-responsive metabolites including flavonol and flavonoid compounds in seedling of onion treated with Τ. harzianum (Abdelrahman et al., 2018). Additionally, El-Sharkawy et al. (2018) reported that application of Trichoderma spp. significantly reduced disease severity by inducing peroxidase and polyphenol oxidase enzymes, and increasing the total phenol content in host plant. The use of T. polysporum exhibited the highest effectiveness in biocontrol of melon wilt by about 44.85% (Gava and Pinto, 2016). Sunpapao et al. (2018) studied three species of Trichoderma and recorded more than 66.21% inhibition in disease severity in leaf spot caused by Curvularia oryzae in oil palm seedlings. In terms of disease resistance induction, Pascale et al. (2017) investigated the effect of two Trichoderma strains and their secondary metabolites against powdery mildew caused by Uncinula necator; they found that T. harzianum and T. atroviride increased polyphenols rate in grapes.

Overall, this study provided solid evidence that *Trichoderma* spp. are the most potential biocontrol agents against FOC. The use of some *Trichoderma* secondary metabolites reduced significantly disease severity and enhanced systemic resistance. This could have a significant beneficial impact on the management of diseases in crop plants.

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کارایی متابولیتهای ثانویه تولید شده توسط *.Trichoderma* spp در کنترل بیولوژیکی پژمردگی فوزاریومی نخود

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چكیده: تحقیق حاضر بهمنظور ارزیابی اثر متابولیتهای ثانویه تولید شده توسط پنج گونه Fusarium oxysporum f. sp. ciceris از گونههای Frichoderma sp. Trichoderma sp. در نفرایط درون (FOC) در نخود انجام شد. توانایی بیوکنترلی گونههای Trichoderma sp. علیه FOC در شرایط درون شیشهای آزمایش شد. متابولیتهای ثانویه *تریکودرما* با روش استخراج به کمک حلال استخراج شد و علیه FOC مورد ارزیابی قرار گرفت. آزمایشهای درون شیشهای اثرات بازدارندگی خیلی خوبی را در نمام گونههای FOC مورد ارزیابی قرار گرفت. آزمایشهای درون شیشهای اثرات بازدارندگی خیلی خوبی را در غیرمستقیم به ترتیب تا ۲۸/۸ و ۲۷/۸ درصد بود. بهطوری که نرخ بازدارندگی در تماس مستقیم و قابل توجهی در شدت بیماری پژمردگی فوزاریومی شدند. بهطور ویژه گونه مورد آزمایش موجب کاهش قابل توجهی در شدت بیماری پژمردگی فوزاریومی شدند. بهطور ویژه گونه Trichoderma sp. ترمام گونههای Trichoderma sp. درصد بود. بهطور ویژه گونه موجب کاهش قابل توجهی در شدت بیماری را کاهش داد. همچنین متابولیتهای ثانویه مورد آزمایش موجب کاهش نشان داد که متابولیتهای ثانویه در کاهش شده بیماری بسیار فعال بودند. گونه موجب کاهش نشان داد که متابولیتهای ثانویه در کاهش شدت بیماری بسیار فعال بودند. گونه موجب کاهش ترکیبات پلیفنولی نسبت داده میشود. گونههای ۵۶/۹ درصد کاهش نشان داد. مقاومت نخود غالباً به نشان داد که متابولیتهای ثانویه در کاهش شدت بیماری با ۵۶/۹ درصد کاهش نشان داد. مقاومت نخود غالباً به نشان داد که متابولیتهای ثانویه در کاهش شدت بیماری با ۵۶/۹ درصد کاهش نشان داد. مقاومت نخود غالباً به نشان داد که متابولیتهای ثانویه در کاهش شدت بیماری با ۵۶/۹ درصد کاهش نشان داد. مقاومت نخود غالباً به نورزیوم آنها میتوانند بهعنوان عوامل ضدقارچی بالقوه و امیدبخش در جلوگیری از وقوع پژمردگی فوزاریومی نخود باشد.

واژگان کلیدی: کنترل بیولوژیکی، تریکودرما، Fusarium oxysporum، متابولیتهای ثانویه، نخود