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# On the biocontrol by *Trichoderma afroharzianum* against *Fusarium culmorum* responsible of fusarium head blight and crown rot of wheat in Algeria

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## Abstract

**Background:** Durum wheat (*Triticum durum* Desf.) is one of the most important cereals in the world. Unfortunately, the wheat plant is a target of several species of the genus *Fusarium*. This genus causes two serious diseases: fusarium crown rot (FCR) and fusarium head blight (FHB). The search for new indigenous strains of *Trichoderma* with a high potential for biocontrol against these two diseases was the purpose of this study.

**Results:** Biocontrol potential of 15 isolates of *Trichoderma* (T1 to T15), isolated from different rhizosphere soils and Algerian ecosystems, was evaluated against 4 strains of *Fusarium culmorum* (FC11, FC2, FC4, and FC20); the main causative agent of FCR and FHB. The efficacy of biological control by *Trichoderma* spp., evaluated by in vitro tests (direct and indirect confrontation), was confirmed by in vivo bioassays. The in vitro results showed a significant inhibition of mycelial growth of *F. culmorum* species than the control. The highest percentages of inhibition were obtained by T9, T12, and T14 isolates causing a maximum inhibition percentage of 81.81, 77.27, and 80.68%, respectively. T14 was selected for biocontrol in in vivo testing. A tube and pot experiments for FCR against *F. culmorum* showed that T14 decreased the disease severity with 50 and 63.63% reduction, respectively. FHB infection was significantly reduced by T14 in all durum wheat cultivars tested, where %AUDPC (area under the disease progress curve) reduction was 49.77, 43.43, 48.25, and 74.60% for Simeto, Waha, Bousselem, and Setifis genotypes, respectively. Yields also increased significantly for almost all cultivars. The antagonistic T14 was characterized based on molecular tools, using translation elongation factor1-alpha (TEF1- $\alpha$ ) and internal transcribed spacers rDNA (ITS1). The results identified T14 as *T. afroharzianum* with accession numbers attributed by NCBI GenBank as MW171248 and MW159753.

**Conclusions:** *Trichoderma afroharzianum*, evaluated for the first time in Algeria as biocontrol agent, is a promising biocontrol approach against FCR and FHB.

**Keywords:** *Triticum durum*, *Trichoderma afroharzianum*, *Fusarium*, Biocontrol, Pathogenicity, Algeria

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## Background

Durum wheat (*Triticum durum* Desf.) is the main staple food crop in Algeria, where huge quantities are grown and traditionally consumed in the form of semolina and bread. However, it is affected by several abiotic and biotic stresses. Among the latter, fusarium crown rot (FCR) and fusarium head blight (FHB), which are considered two serious fungal diseases of wheat in the world, leading to significant yield losses and reduced market prices due to the poor quality of infected grains (Moya-Elizondo 2013). FCR in wheat causes the invasion of crown and root tissues by *Fusarium* spp. mycelia, which induces their decay and reduces the efficiency of nutrient and water absorption, while FHB is manifested by the bleaching of spikelets (Xu and Nicholson 2009).

Some *Fusarium* species are among the dangerous cereal pathogens with its secondary metabolites such as deoxynivalenol (DON), zearalenone (ZEN), and fumonisin B1, that are among the five most important mycotoxins (Mielniczuk and Skwaryło-Bednarz 2020). Investigations carried out revealed that strains of *F. culmorum* was the dominant and most aggressive species on wheat seedlings associated with FCR and FHB in Algeria (Abdallah-Nekache et al. 2019). In addition to yield losses, *F. culmorum* causes the accumulation of various mycotoxins such as 3-acetyl deoxynivalenol (3ADON) or nivalenol (NIV) (Laraba et al. 2017).

There are no effective and consistent control measures against *Fusarium*. In addition, resistance in commercial cultivars is only partial. Disease control is based primarily on the use of fungicides, and crop techniques such as rotation to reduce the occurrence of the disease (Wegulo et al. 2015). The chemical treatments pose a potential risk to human and animal health and increase environmental pollution, such as altering the beneficial functions of microorganisms living in the soil and rhizosphere ecosystem, and their effectiveness differs according to the fungal species concerned. Recently, biological control agents have been accepted as an ecological alternative (Akrami and Yousefi 2015). It is one of the most promising tools to maintain the current level of agricultural production and to control *Fusarium* spp. (Wegulo et al. 2015; Tian et al. 2016).

*Trichoderma* species have emerged as the most powerful bioprotectors for the management of a number of plant diseases due to their broad-spectrum action (Mukhopadhyay 2005). *Trichoderma* spp., is a cosmopolitan fungus common in different biotopes. It involves different types of interaction with other microorganisms in the rhizosphere. Some *Trichoderma* spp. are opportunistic and show a parasitic lifestyle such as *T. harzianum*, *T. longibrachiatum*, and *T. atroviride*, which live in the soil, without adverse effects on the environment and non-toxic to human health (Ghazalibiglar et al. 2016). The ability to produce volatile and

non-volatile fungitoxic substances vary from one species to another and even between strains of the same species. The production of antibiotics is the most important antagonistic capacity. However, mycoparasitism has been proposed as the primary antagonistic mechanism exhibited by *Trichoderma* spp. (Zeilinger et al. 2016).

*In vitro* methods are considered among the easiest and fastest screening techniques, by using double culture plates, commonly used to select *Trichoderma* strains with potential antagonistic effects and omitting those with no biological activity (Matarese et al. 2011). *Trichoderma* is easily obtained by the multi-tube dilution technique due to its chlamydospore formation and colonization of organic substrates (Khandelwal et al. 2012). Conidia have been commonly used for biological control and conidial biomass can be grown by the submerged culture method or on solid substrate (Harman and Kubicek 1998).

The objectives of the present investigation were (i) isolation of *Trichoderma* spp. and *in vitro* evaluation of the antagonistic power against four strains of *F. culmorum*, the causal agent of FCR and FHB in Algeria; (ii) evaluation of *Trichoderma* as a biocontrol agent against FCR by two *in vivo* tests in the growth chamber; (iii) evaluation of the biocontrol potential against FHB in the pot test in the greenhouse; and (iv) identification of the most antagonistic species by molecular characterization.

## Methods

### Fungal material

For the isolation of the antagonistic agents, 68 soil samples were collected at a depth of 20 to 30 cm, in different provinces of Algeria (Table 1). They were collected at random from the rhizosphere of different wheat fields and also from soils of different ecosystems. They were placed in sterile plastic bags. Fifteen isolates of *Trichoderma* (Table 1) were isolated using the suspension-dilution method (flat dilutions) (Davet and Rouxel 1997); they were purified by monospore culture, then stored as mycelium discs on agar in eppendorf tubes in 20% glycerol at  $-80^{\circ}\text{C}$  (Siou et al. 2013).

Four *Fusarium culmorum* strains coded (FC11, FC2, FC4, and FC20) were submitted to GenBank (NCBI) under accessions numbers MW151664, MW165423, MW165434, and MW165435, respectively, were used. They are characterized by their aggressiveness among a collection of phytopathogenic wheat *Fusarium* species belonging to the fungal library of our laboratory. *Fusarium* strains were used as a pathogen in *in vitro* test. For *in vivo* tests, only FC2 was used.

### Plant material

Durum wheat cultivar Simeto, known for its susceptibility to *Fusarium*, was used for both *in vivo* tests in

**Table 1** Code of *Trichoderma* isolates, with the soil type, region, and province of their isolation

<i>Trichoderma</i> isolate code	Type of soil	Town and region	Provinces
T1	Sand	Tahir plage bazoul	Jijel
T2	Mountainous soil	El akhdaria	Bouira
T3	Mountainous soil	El akhdaria	Bouira
T4	Agricultural sand (potato)	El meghaier	Oued Souf
T5	Forest soil	Oued athmania Barrage grouz	Mila
T6	Agricultural soil	El harrouch	Skikda
T7	Agricultural soil	El harrouch	Skikda
T8	Mountainous soil	Ain zouit	Skikda
T9	Mountainous soil	Ain bouziane	Skikda
T10	Mountainous soil	Ain bouziane	Skikda
T11	Mountainous soil	Ain bouziane	Skikda
T12	Mountainous soil	Khmis meliana	Ain Defla
T13	Agricultural soil (wheat)	Oum tboul	Skikda
T14	Agricultural soil (wheat)	Ouled Rahmoune	Constantine
T15	Forest soil	Lazharia	Tissemsilt

growth chamber (tubes and pots tests) to evaluate the antagonistic effect of *Trichoderma* isolates on the severity of FCR. Three other local durum wheat cultivars (Setifis, Waha, and Bousselem), widely grown in Algeria, were also used for the last test in greenhouse to evaluate the biocontrol effect of *Trichoderma* isolate on FHB. All cultivars used in this study were kindly provided by the National Institute of Agronomic Research of Algeria (INRAA).

#### In vitro antagonistic activity of *Trichoderma* isolates

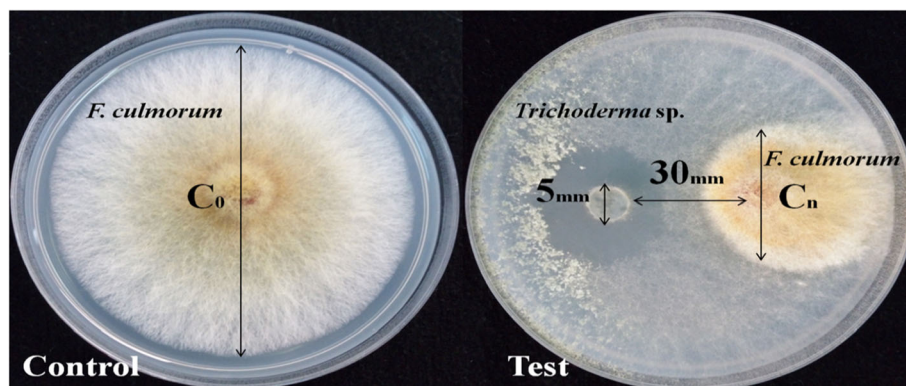
##### Antagonism by direct confrontation (DC)

Confrontations between the antagonist *Trichoderma* sp. and the pathogen *F. culmorum* were carried out according to Hibar et al. (2004). A mycelial disc (5 mm) of each fungus was placed at opposite poles on boxes containing

the potato sucrose agar (PSA) medium, keeping a distance of 30 mm between the two fungi and incubated at 28 °C for 5 days. Three replicates were done for each *Trichoderma* sp./*F. culmorum* combination. Boxes containing only *F. culmorum* strains were used as controls (Fig. 1).

##### Antagonism by indirect confrontation (IC)

It was realized according to Daami-Remadi and El Mahgoub (2001). Discs of the antagonistic isolate *Trichoderma* sp. and the pathogenic strain *F. culmorum* (5 mm of diameter) were deposited in two separate dishes containing the PSA medium. An assembly was carried out by superimposing the two dishes, *Trichoderma* sp. at the bottom and *F. culmorum* at the top. The junction between the two dishes was ensured by parafilm in order



**Fig. 1** Demonstrative schema of the different measurements used to calculate the percentage of mycelial growth inhibition for in vitro confrontation *Trichoderma*/*F. culmorum*

to avoid any loss of volatile substances. The dishes were incubated in the dark at 28 °C for 5 days. The control was formed by superimposing two dishes, the upper one containing a *Fusarium* disc, while the lower one contained only the PSA medium.

#### **Measurement of the inhibition exerted by *Trichoderma* spp. I (%)**

It was estimated according to Hmouni et al. (1996) by calculating the percentage inhibition of mycelial growth of *F. culmorum* strains as follows:  $I(\%) = (1 - (C_n/C_0)) \times 100$

Where  $C_n$ : diameter of the colony of the pathogen (*F. culmorum*) in the presence of the antagonist (*Trichoderma* sp.) (mm) and  $C_0$ : diameter of the control colony (mm) (Fig. 1).

#### **In vivo antagonistic activity of *Trichoderma* isolates**

The highly antagonistic T14 isolate, on the basis of its in vitro efficacy, was chosen to evaluate its biocontrol effect in vivo.

#### **Inocula preparation**

Fungal discs (13 mm of diameter) of the 2 fungi (*Fusarium* and *Trichoderma*), used in the tests, were obtained from 7-day-old young cultures on PSA medium. The inoculation suspension was prepared according to the protocol of Stein et al. (2009). The FC2 strain was grown on 6 Petri dishes on PSA medium for 45 days. Ten milliliter sterile of distilled water with 0.05% (v/v) Tween 20 was added to the mycelial aerial part on the surface of each plate, and then carefully scraped off until the surface part of the fungus was recovered. The mixture of macroconidia, mycelium, and PSA medium was recovered in a beaker and then filtered through a double layer of cheesecloth. The concentration of the macroconidial suspension  $8 \times 10^6$  macroconidia/ml was adjusted, using Malassez cell, for the inoculation test of the spike in greenhouse. The same protocol was followed for the preparation of the spore suspension of the T14 isolate, except that the culture on PSA medium lasted only 1 week (rapid sporulation), with a concentration of  $8 \times 10^8$  spores/ml, for the spike inoculation test in greenhouse. All inocula were stored at 4 °C until use.

#### **Crown inoculation test in assay tubes**

This test was inspired by the method of Asad et al. (2009) with major modifications. Seventy-five assay tubes (14 cm × 3 cm) were filled with 4 cm of cotton, and then 20 ml of distilled water was added to each tube and covered with aluminum foil, then sterilized by autoclaving at 180 °C for 2 h. Three hundred seventy-five seeds of Simeto durum wheat, surface disinfected with 2% sodium hypochlorite (NaClO) for 5 min and rinsed 3

times with sterile distilled water, were placed on the wet cotton swab in each test tube (5 seeds/tube). The tubes were closed with transparent caps and placed in the oven at  $28 \pm 1$  °C for 4 days, where the seedlings reached the one-leaf stage (Zadoks' GS 11), the inoculation took place as follows: 25 control tubes; 25 tubes inoculated by FC2 (13 mm disc); 25 tubes inoculated by FC2 + T14 (13 mm disc for both).

After inoculation, the tubes were transferred to a growth chamber (25/19 °C day/night temperature, 16/8-h light/dark cycle) for 10 days; they were arranged according to a complete randomization plan. When the seedlings reached the two-leaf stage (Zadoks' GS 12), the seedlings were carefully removed from the cotton with water. The FCR severity classes for crown discoloration were assigned to the laboratory on a scale of 0 to 4 where class 0 = healthy crown; 1 = light browning of crown; 2 = half browning of crown; 3 = full browning of crown; and 4 = death of seedlings.

The disease severity (DS) of each treatment was calculated using McKinney's (1923) index, which expresses the percentage of disease severity (i.e., 100) according to the formula:

$$DS = \left\{ \frac{\sum (c \times f)}{n \times N} \right\} \times 100$$

Where  $c$  = disease class,  $f$  = frequency,  $n$  = number of observations, and  $N$  = the highest value of the empirical scale adopted (class 4). The whole test was repeated twice in succession.

#### **Crown inoculation test in pots**

Thirty pots of (8 × 12 cm) were used. For each pot, 5 durum wheat (Simeto) seeds were sterilized and sown as indicated in the previous test. All the pots were placed in the growth chamber (25/19 °C day/night temperature, 16/8-h light/dark cycle). After 3 weeks, the seedlings reached the two-leaf stage (Zadoks' GS 12). The soil around the seedlings was removed, and the inocula discs prepared as mentioned above were placed around the stems, one disc for each stem 2 cm below the soil; the pots were inoculated as follows:

Ten control pots (uninoculated); 10 inoculated pots with FC2 with 13 mm disc; 10 inoculated pots with FC2 + T14 (13 mm disc versus 13 mm disc).

The soil was then placed back around the stems, and the pots were organized according to a completely random plan. Three weeks after inoculation, the plants reached the early tillering stage (Zadoks' GS 20), each plant was carefully removed from the soil and washed with tap water. Severity classes were assigned on the same scale and the DS was calculated using McKinney's (1923) index, as shown above. Koch's postulates were satisfied by re-isolating FC2 from brown spots on the



coleoptiles for both tests. The whole test was repeated twice in succession. To measure the efficacy of T14 biocontrol against FC2, the reduction percentage in DS is measured for both tests as follows:

$$\%DS \text{ reduction} = 100 - \left\{ \frac{DS_{FC2+T14} \times 100}{DS_{FC2}} \right\}$$

#### **Test of spike inoculation in greenhouse**

A total of 48 pots (24 × 20 cm) filled with a mixture of soil/compost (1/2) previously sterilized at 180° for 2 h (2 repetitions 24 h apart) were used. One hundred twenty seeds of each of the 4 cultivars (Simeto, Waha, Bousselem, Setifis) were surface-sterilized with 2% NaClO for 5 min, and rinsed 3 times with sterile distilled water, were sown (10 seeds/pot) under the soil surface at about 2 cm. The plants were grown in the greenhouse. The soil was watered every 3 days. The inoculation took place at the flowering stage (Zadoks' GS 60) and each spike between the first and the third day after the first anthers emergence was labeled.

The inoculation suspension for each cultivar was done as follows: 10 seeds in a control pot (no inoculation); 10 seeds in a pot for inoculation with FC2 (8 × 10<sup>6</sup> macroconidia/ml); 10 seeds in a pot for inoculation with FC2 + T14 (mixture: 8 × 10<sup>6</sup> macroconidia/ml + 8 × 10<sup>8</sup> spores/ml); and 10 seeds in a pot for inoculation with T14 only, with 2 repetitions of each case for each cultivar (12 pots/cultivar).

The inoculation with the FC2 macroconidial suspension prepared previously as indicated above was carried out by spraying about 2 ml of suspension on both sides of the spike, in a Plexiglas cage. The spikes were then covered with a transparent polyethylene bag for 72 h in order to maintain maximum relative humidity, necessary for the early stages of fungal development. After 15 days of inoculation, a visual disease assessment was performed for each spike by counting the percentage of symptomatic spikelets (PSS) of each inoculated spike. Approximately 8 to 10 heads were noted in each pot.

Because of the variability and specificity of each cultivar (inter and intra-genotype), flowering did not take place on the same day, even within the same genotype. Spike PSS scoring was done separately. For each spike, 5 readings with 4 days interval were done, thus illustrating the kinetics of the disease over approximately 20 days. The area under the disease progress curve (AUDPC) standard illustrated the kinetics of disease progression in PSS between the first and last scoring. AUDPC for FHB incidence was measured by the number of days of observation and calculated as described by Siou et al. (2015):

$$AUDPC = \left\{ \sum_0^n \left[ \left( \frac{Y_i + Y_{i+1}}{2} \right) \times (t_{i+1} - t_i) \right] \right\} / \{t_n - t_0\}$$

Where  $Y_i$  the PSS at the  $i$ th scoring,  $t_i$  the number of days elapsed between inoculation and the last scoring,  $t_0$  the number of days elapsed between inoculation and the 1st scoring and  $n$  the total number of scoring.

At maturity, each spike was harvested. The spikes were deseeded by hand in order to recover all the grains. All the grains were counted and weighed in order to obtain thousand kernel weight (TKW) of inoculated and uninoculated spikes (control) of each cultivar, the reduction percentage of AUDPC is measured in order to estimate the loss percentage of FHB severity resulting from biocontrol by T14.

$$\%AUDPC \text{ reduction} = 100 - \left\{ \frac{AUDPC_{FC2+T14} \times 100}{AUDPC_{FC2}} \right\}$$

Koch's postulates were satisfied by re-isolating the FC2 pathogen from the infected grains.

#### **Physical properties of the grain**

In order to fully understand the effect of T14 treatment on the physical properties of the grains, the different grains from the greenhouse spike inoculation test (control, FC2 infected, FC2 infected and treated with T14, and treated with T14 only) were compared. The four types of grain for each cultivar tested were cleaned manually. Broken and immature kernels were removed. Ten grains were taken at random and their three linear dimensions: length  $L$ , width  $W$ , and thickness  $T$ ; were measured, using a digital caliper (0–150 mm) with an accuracy of 0.01 mm.

#### **Molecular characterization of T14 isolate**

To identify the isolate T14, which presented the best biocontrol efficiency, molecular identification was performed by the BIOfidal laboratory (CEDEX-France).

#### **DNA extraction**

DNA was extracted from mycelium collected by scraping the surface of Petri dishes cultures of purified isolate. One hundred microliters of lysis buffer (50 mM Tris-HCl pH 7.5, 50 mM EDTA, 3% SDS and 1% 2-mercaptoethanol) was added and the nucleic acids were isolated according to the microwave mini-prep procedure described by Goodwin and Lee (1993). The final DNA pellet was supplemented into 100  $\mu$ l TE buffer (10 mM Tris-HCl pH 8.0, 0.1 mM EDTA) and stored at –20 °C until used.

### PCR amplifications and DNA sequencing

The internal transcribed spacer (ITS1) and the translation elongation factor (TEF1- $\alpha$ ) were amplified using primer pairs ITS1 (5' TCC GTA GGT GAA CCT GCG G '3) (White et al. 1990) and EF1-728F (5' CAT CGA GAA GTT CGA GAA GG 3') (Carbone and Kohn 1999). All amplification reactions were performed in a 50- $\mu$ l reaction volume containing 75 mM Tris-HCl pH 9.0, 20 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, 0.01% (w/v) Tween 20, 1.5 mM MgCl<sub>2</sub>, 200 IM each dNTP, 1 unit of thermostable DNA polymerase (GoTaq, Promega), and 400 nM of each relevant oligonucleotide primer. After electrophoresis in 1.2% agarose gels in 0.5 $\times$  TAE buffer (20 mM Tris-acetate pH 8, 0.5 mM EDTA), DNA was visualized by Ethidium bromide staining and UV illumination.

PCR products (amplified DNA) were purified by mini-column centrifugation (NucleoSpinVR Extract II) and the DNA binds in the presence of a chaotropic salt to a

silica membrane. The binding mixture was loaded directly onto NucleoSpinVR Extract II columns. Contaminations were removed by a washing step with ethanolic NT3 buffer. Finally, the pure DNA was eluted under low ionic strength conditions with a slightly alkaline NE buffer (5 mM Tris-Cl, pH 8.5). The sequences of EF1 and ITS region thus obtained were submitted to GenBank database through Submission Portal (a World Wide Web sequence submission server available at NCBI home page: <http://www.ncbi.nlm.nih.gov>).

### Statistical analysis

Data from in vitro and in vivo assays were subjected to analysis of variance (ANOVA). The histograms were made, using Software SPSS (IBM SPSS Statistics version 25), and means compared by Tukey's test ( $P < 0.05$ ). The correlation between AUDPC<sub>FC2</sub> and AUDPC<sub>FC2 + T14</sub> was determined by Pearson correlation. The

**Table 2** The percentage of inhibition of *Trichoderma* isolates on *Fusarium culmorum* strains in both cases of confrontation (direct and indirect) after 5 days incubation

Mean inhibition I%	FC11		FC2		FC4		FC20	
	Direct	Indirect	Direct	Indirect	Direct	Indirect	Direct	Indirect
T1	30.46 $\pm$ 2.51 <sup>a</sup>	10.00 $\pm$ 1.08 <sup>a</sup>	14.77 $\pm$ 1.37 <sup>a</sup>	0.00 $\pm$ 0.00 <sup>a</sup>	18.18 $\pm$ 1.74 <sup>a</sup>	6.97 $\pm$ 2.40 <sup>a</sup>	11.49 $\pm$ 1.51 <sup>a</sup>	9.52 $\pm$ 1.50 <sup>abc</sup>
T2	63.57 $\pm$ 4.51 <sup>de</sup>	17.50 $\pm$ 2.40 <sup>ab</sup>	31.81 $\pm$ 2.97 <sup>b</sup>	0.00 $\pm$ 0.00 <sup>a</sup>	67.04 $\pm$ 3.05 <sup>c</sup>	30.23 $\pm$ 1.29 <sup>e</sup>	41.37 $\pm$ 1.62 <sup>c</sup>	13.09 $\pm$ 2.34 <sup>cd</sup>
T3	63.57 $\pm$ 6.64 <sup>de</sup>	36.25 $\pm$ 4.21 <sup>cde</sup>	29.54 $\pm$ 2.8 <sup>b</sup>	0.00 $\pm$ 0.00 <sup>a</sup>	76.13 $\pm$ 3.66 <sup>def</sup>	17.44 $\pm$ 0.93 <sup>c</sup>	49.42 $\pm$ 1.98 <sup>cd</sup>	20.23 $\pm$ 1.95 <sup>ef</sup>
T4	51.65 $\pm$ 4.19 <sup>bcd</sup>	28.75 $\pm$ 4.54 <sup>c</sup>	57.95 $\pm$ 6.54 <sup>c</sup>	0.00 $\pm$ 0.00 <sup>a</sup>	73.86 $\pm$ 2.84 <sup>cdef</sup>	23.25 $\pm$ 1.58 <sup>d</sup>	49.42 $\pm$ 7.02 <sup>cd</sup>	5.95 $\pm$ .95 <sup>ab</sup>
T5	47.68 $\pm$ 5.13 <sup>b</sup>	27.50 $\pm$ 4.51 <sup>bc</sup>	48.86 $\pm$ 1.83 <sup>c</sup>	17.07 $\pm$ 2.25 <sup>b</sup>	70.45 $\pm$ 1.63 <sup>cd</sup>	16.27 $\pm$ 1.26 <sup>c</sup>	49.42 $\pm$ 5.28 <sup>cd</sup>	14.28 $\pm$ 2.38 <sup>cd</sup>
T6	48.34 $\pm$ 3.68 <sup>bc</sup>	32.50 $\pm$ 2.38 <sup>cde</sup>	47.72 $\pm$ 2.97 <sup>c</sup>	18.29 $\pm$ 2.96 <sup>b</sup>	65.90 $\pm$ 5.87 <sup>c</sup>	38.37 $\pm$ 2.03 <sup>f</sup>	43.67 $\pm$ 1.32 <sup>c</sup>	15.47 $\pm$ 1.80 <sup>de</sup>
T7	39.07 $\pm$ 3.31 <sup>ab</sup>	30.00 $\pm$ 2.32 <sup>c</sup>	54.54 $\pm$ 2.74 <sup>c</sup>	0.00 $\pm$ 0.00 <sup>a</sup>	85.22 $\pm$ 2.29 <sup>g</sup>	46.51 $\pm$ 0.78 <sup>g</sup>	48.27 $\pm$ 2.58 <sup>cd</sup>	4.76 $\pm$ 1.22 <sup>a</sup>
T8	48.34 $\pm$ 4.62 <sup>bc</sup>	42.50 $\pm$ 4.84 <sup>ef</sup>	14.77 $\pm$ 2.74 <sup>a</sup>	19.51 $\pm$ 2.61 <sup>bc</sup>	72.72 $\pm$ 2.32 <sup>cde</sup>	30.23 $\pm$ 1.07 <sup>e</sup>	52.87 $\pm$ 2.06 <sup>de</sup>	13.09 $\pm$ 1.64 <sup>cd</sup>
T9	67.54 $\pm$ 1.82 <sup>e</sup>	48.75 $\pm$ 4.30 <sup>fg</sup>	70.45 $\pm$ 2.71 <sup>d</sup>	29.26 $\pm$ 1.96 <sup>de</sup>	81.81 $\pm$ 2.43 <sup>fg</sup>	27.90 $\pm$ 3.14 <sup>de</sup>	67.81 $\pm$ 1.71 <sup>h</sup>	25.00 $\pm$ 1.42 <sup>f</sup>
T10	50.99 $\pm$ 7.87 <sup>bcd</sup>	30.00 $\pm$ 2.61 <sup>c</sup>	54.54 $\pm$ 5.10 <sup>c</sup>	0.00 $\pm$ 0.00 <sup>a</sup>	73.86 $\pm$ 3.18 <sup>cdef</sup>	32.55 $\pm$ 2.20 <sup>e</sup>	56.32 $\pm$ 1.94 <sup>def</sup>	10.71 $\pm$ 1.87 <sup>cde</sup>
T11	44.37 $\pm$ 7.26 <sup>ab</sup>	35.00 $\pm$ 2.95 <sup>cde</sup>	13.63 $\pm$ 3.01 <sup>a</sup>	24.39 $\pm$ 2.32 <sup>cd</sup>	39.77 $\pm$ 2.72 <sup>b</sup>	48.83 $\pm$ 1.64 <sup>g</sup>	25.28 $\pm$ 1.82 <sup>b</sup>	47.61 $\pm$ 3.86 <sup>h</sup>
T12	67.54 $\pm$ 7.51 <sup>e</sup>	50.00 $\pm$ 3.73 <sup>fg</sup>	77.27 $\pm$ 6.53 <sup>d</sup>	32.92 $\pm$ 1.23 <sup>e</sup>	71.59 $\pm$ 1.76 <sup>cd</sup>	55.81 $\pm$ 2.85 <sup>h</sup>	62.06 $\pm$ 2.35 <sup>fgh</sup>	48.80 $\pm$ 1.95 <sup>h</sup>
T13	45.69 $\pm$ 2.31 <sup>b</sup>	31.25 $\pm$ 4.39 <sup>cd</sup>	34.09 $\pm$ 2.19 <sup>b</sup>	2.43 $\pm$ 1.00 <sup>a</sup>	73.86 $\pm$ 1.69 <sup>cdef</sup>	12.79 $\pm$ 1.02 <sup>bc</sup>	42.52 $\pm$ 1.26 <sup>c</sup>	30.95 $\pm$ 1.04 <sup>g</sup>
T14	62.91 $\pm$ 2.52 <sup>cde</sup>	56.25 $\pm$ 4.18 <sup>g</sup>	79.54 $\pm$ 4.65 <sup>d</sup>	21.95 $\pm$ 2.91 <sup>bc</sup>	80.68 $\pm$ 2.12 <sup>efg</sup>	44.18 $\pm$ 1.45 <sup>g</sup>	65.51 $\pm$ 1.61 <sup>gh</sup>	13.09 $\pm$ 2.31 <sup>cd</sup>
T15	46.35 $\pm$ 4.13 <sup>b</sup>	41.25 $\pm$ 3.81 <sup>def</sup>	57.95 $\pm$ 1.86 <sup>c</sup>	17.07 $\pm$ 2.11 <sup>b</sup>	72.72 $\pm$ 2.20 <sup>cde</sup>	10.46 $\pm$ .79 <sup>ab</sup>	58.62 $\pm$ 1.58 <sup>efg</sup>	5.95 $\pm$ 0.86 <sup>ab</sup>

The values (mean  $\pm$  Std. deviation) with different letters denote the statistical significance determined by ANOVA followed by Tukey test ( $P < 0.05$ )

coefficient of correlation was found to be significant at the 5% level ( $P \leq 5\%$ ).

## Results

### In vitro effect of *Trichoderma* isolates on *F. culmorum* strains

The antagonistic effect in direct confrontation (DC) of *Trichoderma* isolates against *F. culmorum* varies widely from isolate to isolate. Three isolates T9, T12, and T14 had the highest and most stable levels of inhibition with all strains of *F. culmorum*. Inhibition values ranged from 62.06 to 81.81% (Table 2, Fig. 2). The antagonistic effect in indirect confrontation (IC) is less important for all *Trichoderma* spp., except T11 against FC2, FC4, and FC20 where IC is greater than DC (24.39% and 48.83%, 47.61%), respectively (Table 2, Fig. 3b–d). T9, T12, and T14 were also the most reliable in the IC with maximum values of 48.75%, 55.81%, and 56.25%, respectively.

The T1 isolate recorded the lowest rates in both confrontations (DC and IC), and with all strains of *F. culmorum*. It is therefore far from being chosen for a biocontrol role against FCR and FHB. On the other hand, maximum growth inhibition (85.22%) was obtained by T7 against FC4 in DC. Some *Trichoderma* isolates had a very low inhibition percentage with null

values for T1, T2, T3, T4, T7, and T10 for IC with FC2 (Table 2). A graphical presentation summarizes the DC and IC results of the in vitro test, to better compare the effect of *Trichoderma* isolates on each strain of *F. culmorum* separately was shown in Fig. 3).

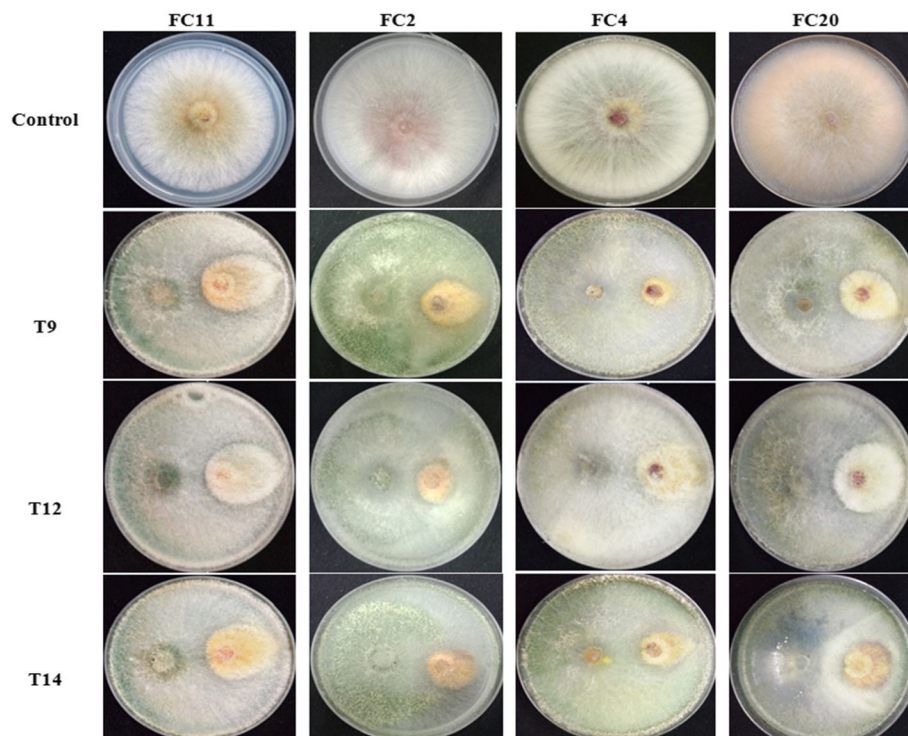
### In vivo effect of T14 in reducing FCR in both tube and pot tests

FC2 induced FCR for the cultivar Simeto with 68 and 60.5%, respectively. A significant decrease in the aggressiveness of DS disease per T14 with considerable %DS reduction rates of 50 and 63.63% was observed (Table 3).

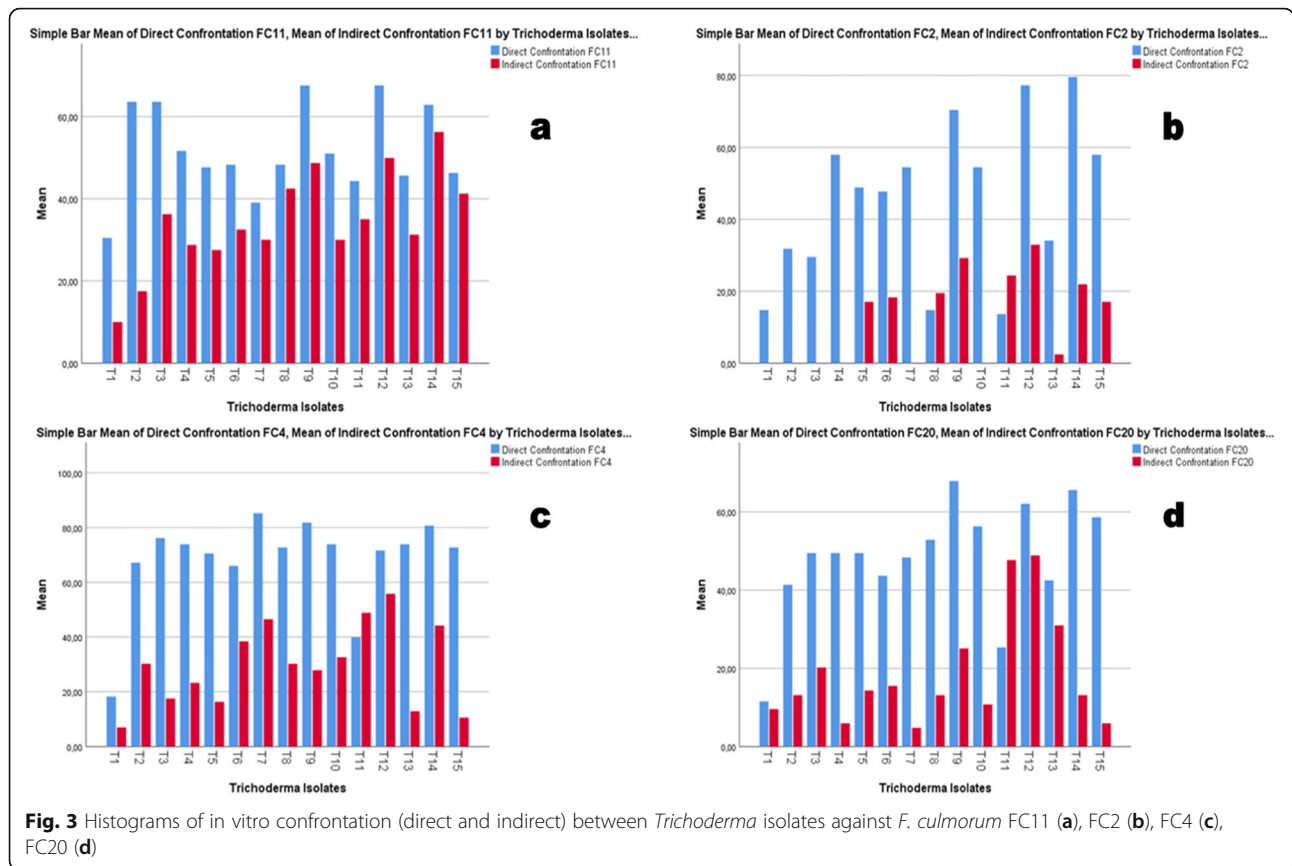
### In vivo effect of T14 in the reduction of FHB

All cultivars were affected by FHB due to FC2 inoculation (Table 4 and Fig. 4). The highest disease severity recorded by the AUDPC standard was in Simeto and Waha cultivars with 63.37 and 56.27%, respectively, while cultivars Bousselem and Setifis showed some resistance to FHB with AUDPC = 18.94% and 16.38%, respectively (Table 4 and Fig. 5a).

A remarkable improvement for the thousand kernel weight (TKW) inoculated and treated with FC2 + T14 was observed, compared to the one inoculated with FC2



**Fig. 2** Direct in vitro confrontation between *Trichoderma* isolates (T9, T12, and T14) against strains of *F. culmorum* (FC11, FC2, FC4, FC20)



in two cultivars, but a slight decrease was observed for Setifis and Waha, where  $TKW_{FC2 + T14} < TKW_{FC2}$ . The antagonistic effect of T14 in the FC2 + T14 mixed inoculum significantly reduced the percent severity of FHB disease for all cultivars with very high reduction rates (74.60% for Setifis) (Table 4 and Fig. 5b). A slight decrease of  $TKW_{T14}$  than the control ( $TKW_{T14} < TKW_{control}$ ), was recorded for all tested cultivars. This decrease remained negligible and significantly better than the *Fusarium*-treated  $TKW_{T14} > TKW_{FC2 + T14} > TKW_{FC2}$ , only the cv. Setifis was the exception (Table 4).

**Effect of T14 on physical properties of grain**

Results showed that for the length parameter L, the difference among the four grain types was not statistically significant, even for the four cultivars. On the other hand, the parameters width W and thickness T

show a very significant variation, the measurements of grains infected with FC2 are the smallest, followed by grains infected with FC2 and treated with T14, due to the antagonistic effect of T14 used as biocontrol agent, the values of the control grains and the one treated only with T14 are almost similar (Table 5).

**Molecular identification of T14**

The amplification of DNA regions with primers TEF1- $\alpha$  and ITS1 was successfully used to identify T14. The final sequences size used for identification analysis were 565 and 533 bp, respectively (Fig. 6). TEF1- $\alpha$  and ITS 1 sequences were submitted to NCBI GenBank and accession numbers were given as MW171248 and MW159753, respectively. The isolate T14 was identified as *Trichoderma afroharzianum*.

**Table 3** Severity of FCR inoculated by FC2 and reduction percentage induced by T14, in both in vivo tube and pot tests

Inoculum	Tubes test		Pots test	
	DS (mean $\pm$ Std. Dev)	%DS reduction	DS (mean $\pm$ Std. Dev)	%DS reduction
FC2 (disc 13 mm)	68 $\pm$ 2.35	0	60.5 $\pm$ 7.08	0
FC2 (disc) + T14 (disc)	34 $\pm$ 6.62	50	22 $\pm$ 3.08	63.63
Control	0	/	0	/



**Table 4** Biocontrol of T14 on FHB tested on 4 durum wheat cultivars, by the AUDPC standard (FC2, FC2 + T14, %reduction), and TKW (control, FC2, FC2 + T14, T14)

Cutivars	AUDPC			TKW (gram)			
	FC2	FC2 + T14	Reduction%	Control	FC2	FC2 + T14	T14
Simeto	63.37 ± 4.52 <sup>b</sup>	31.83 ± 3.16 <sup>b</sup>	49.77	60.38 ± 3.26 <sup>b</sup>	21.93 ± 2.31 <sup>a</sup>	34.15 ± 1.33 <sup>a</sup>	57.60 ± 3.52 <sup>b</sup>
Waha	56.27 ± 3.43 <sup>b</sup>	31.83 ± 3.84 <sup>b</sup>	43.43	58.65 ± 3.59 <sup>b</sup>	39.25 ± 1.73 <sup>b</sup>	32.77 ± 3.15 <sup>a</sup>	40.64 ± 2.22 <sup>a</sup>
Bousselem	18.94 ± 2.10 <sup>a</sup>	9.80 ± 1.34 <sup>a</sup>	48.25	56.91 ± 3.77 <sup>b</sup>	37.38 ± 1.60 <sup>b</sup>	44.18 ± 2.35 <sup>b</sup>	50.99 ± 2.35 <sup>b</sup>
Setifis	16.38 ± 1.37 <sup>a</sup>	4.16 ± 1.28 <sup>a</sup>	74.60	41.92 ± 2.37 <sup>a</sup>	39.62 ± 3.23 <sup>b</sup>	37.81 ± 5.08 <sup>ab</sup>	35.87 ± 4.93 <sup>a</sup>

The values (mean ± Std. deviation) with different letters denote the statistical significance determined by ANOVA followed by Tukey test ( $P < 0.05$ )

## Discussion

In this study, the antagonistic activity of 15 unidentified *Trichoderma* isolates was evaluated in vitro and in vivo against 4 strains of *F. culmorum*, common causative agents of FCR and FHB in wheat. *Trichoderma* is a non-pathogenic fungus that provides protection to many crops against fungal diseases caused by the genus *Fusarium* (Tsegaye Redda et al. 2018). It is used as a biocontrol agent, avoiding the adverse effects that accompany chemical control.

Double culture techniques, as described by many previous studies, have been widely used in tests of antagonistic activities (Srivastava et al. 2010). The direct confrontation experiment showed that all *Trichoderma* isolates were capable of inhibiting mycelial growth of all four pathogenic strains, with rates differing from isolate to isolate and strain to strain as found by Khan et al. (2006). Differences between strains can reach an interval of more than 50%, either in vitro or in vivo.

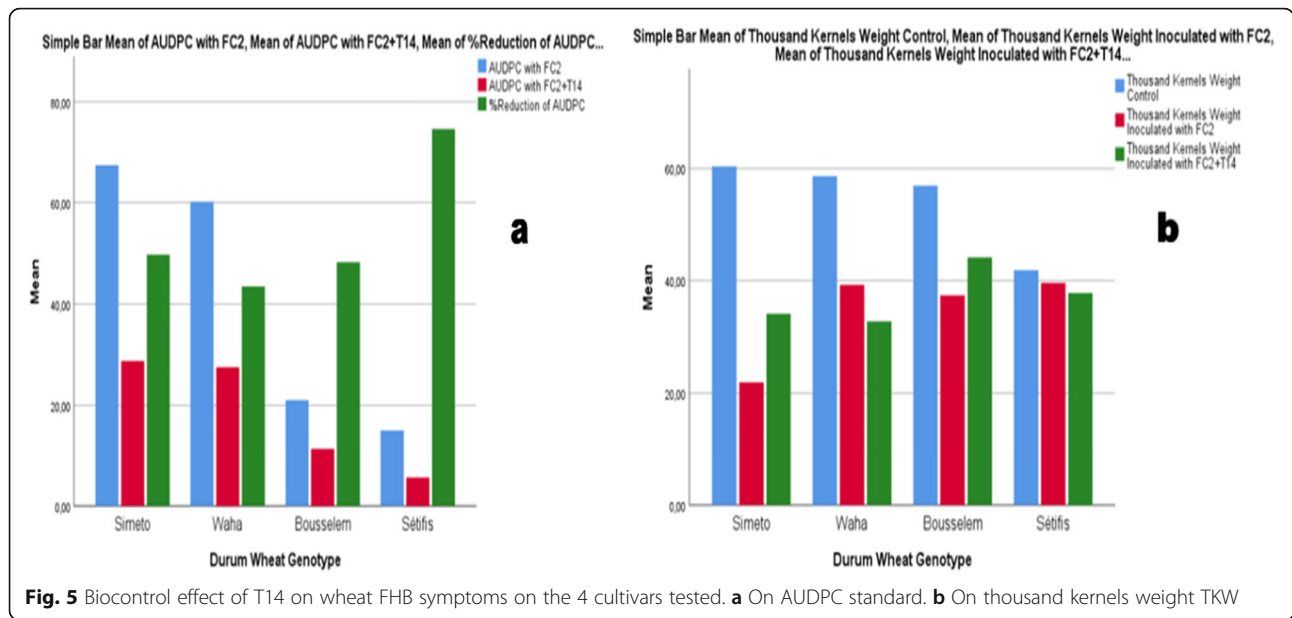
Isolates T9 (67.54, 70.45, 81.81, and 67.81%), T12 (67.54, 77.27, 71.59, and 62.06%), and T14 (62.91, 79.54,

80.68, and 65.51%) showed the most significant rates of reduction than the controls against FC11, FC2, FC4, and FC20, respectively (Table 2 and Figs. 2 and 3). These results are significantly better than those obtained by Ann et al. (2017), where inhibition of mycelial growth of *F. verticillioides* by *Trichoderma* sp. was only 32% compared to control in an in vitro bi-culture test. They were similar to the result obtained by Dendouga et al. (2016), where the rate of growth reduction was 70.68%, representing the efficacy of biological control of *T. harzianum* against *F. culmorum* by in vitro assays. We suppose therefore that T9, T12, and T14 are part of the same species, according to some reports in the literature showing that there are no or slight differences among strains of the same species, for the control of *Fusarium* spp. (Khan et al. 2006).

The T7 isolate recorded the highest rate (85.22%) against FC4, but this high score was not generalized against other strains of *F. culmorum*. The T7 isolate was excluded from in vivo testing due to its specificity as an FC4 antagonist only (Table 2 and Fig. 3c). In contrast,



**Fig. 4** Photos illustrate the different stages of FHB infection, after inoculation with FC2. **a** Healthy spike (control). **b** Start of FHB. **c** Spread. **d** 50% fusarium spike. **e** 100% fusarium spike. **f** Different stages of *Fusarium* disease in the same cultivar. **g** Healthy kernels. **h** Fusarium kernels



T1 recorded the lowest rates among all isolates with 30.46, 14.77, 18.18, and 11.49% against FC11, FC2, FC4, and FC20, respectively (Table 2 and Fig. 3).

*Trichoderma* species are known to produce many volatile organic compounds (Stoppacher et al. 2010), and they also play a major role in inhibiting the growth of the pathogen. Indirect (remote) confrontation experiments showed a reduction in pathogen growth that differs from strain to strain. The T9, T12, and T14 isolates were again more effective than the other isolates studied, the reduction in growth of all 4 strains of *F. culmorum* was stable and balanced with T9 (48.75, 29.26, 27.90, and 25%), T12 (50, 32.92, 55.81, and 48.80%), and T14 (56.25, 21.95%, 44.18, and 13.09%) versus FC11, FC2, FC4, and FC20, respectively (Table 2 and Fig. 3). Behzad et al. (2008) showed similar results with volatile inhibitory substances produced by

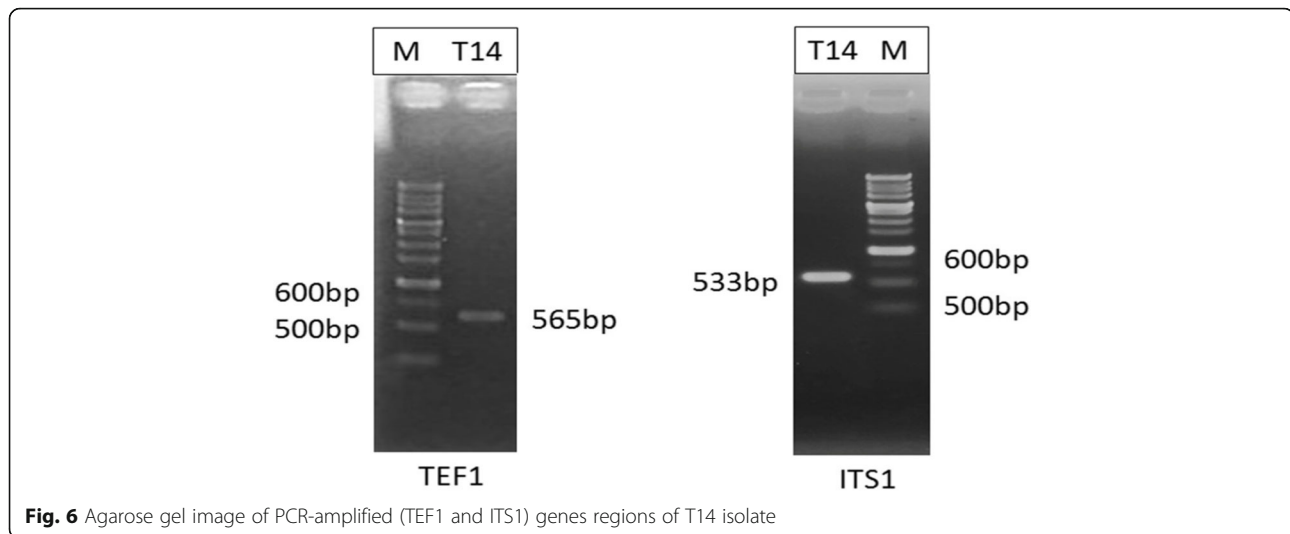
*Trichoderma* isolates against *F. graminearum*. The major advantage of antibiosis via volatile substances is that these substances can remain close to the spikes, thus controlling the inoculum of airborne pathogens without even establishing actual physical contact with them.

In addition, the *I*% recorded for DC is significantly higher than that recorded for IC, with the exception of the T11 isolate with FC2, FC4, and FC20, where the indirect inhibition rates were better than the direct ones, which can be explained by the volatile substances that characterize T11 compared to the others (Table 2 and Fig. 3b–d). The T1 isolate always remained in last place with non-significant and almost negligible rates (10, 0, 6.97%, and 9.52%) versus FC11, FC2, FC4, and FC20, respectively (Table 2 and Fig. 3). These results indicated

**Table 5** Comparison of physical properties (length, width, thickness) of the grains of the 4 cultivars tested (control, FC2 infected, FC2 infected and treated with T14, and treated with T14 only)

Physical properties of grain	Length ± Std deviation(mm)				Width ± Std deviation(mm)				Thickness ± Std deviation(mm)			
	Control	Infected FC2	Treated FC2 + T14	Treated T14	Control	Infected FC2	Treated FC2 + T14	Treated T14	Control	Infected FC2	Treated FC2+T14	Treated T14
Simeto	8.58 ± 0.30 <sup>b</sup>	7.97 ± 0.39 <sup>ab</sup>	8.39 ± 0.42 <sup>b</sup>	8.54 ± 0.22 <sup>b</sup>	3.19 ± 0.17 <sup>b</sup>	1.69 ± 0.42 <sup>a</sup>	2.60 ± 0.26 <sup>ab</sup>	2.97 ± 0.37 <sup>ab</sup>	3.56 ± 0.13 <sup>b</sup>	2.55 ± 0.34 <sup>ab</sup>	3.37 ± 0.12 <sup>b</sup>	3.48 ± 0.16 <sup>b</sup>
Waha	7.99 ± 0.17 <sup>a</sup>	6.90 ± 0.23 <sup>a</sup>	7.27 ± 0.32 <sup>a</sup>	7.63 ± 0.40 <sup>a</sup>	3.23 ± 0.10 <sup>b</sup>	2.14 ± 0.07 <sup>b</sup>	2.39 ± 0.35 <sup>a</sup>	3.15 ± 0.17 <sup>b</sup>	3.39 ± 0.18 <sup>ab</sup>	2.48 ± 0.38 <sup>ab</sup>	2.84 ± 0.31 <sup>a</sup>	3.36 ± 0.22 <sup>ab</sup>
Bousselem	7.88 ± 0.54 <sup>a</sup>	8.13 ± 0.25 <sup>b</sup>	8.23 ± 0.31 <sup>b</sup>	8.31 ± 0.25 <sup>b</sup>	2.56 ± 0.16 <sup>a</sup>	2.08 ± 0.26 <sup>ab</sup>	2.86 ± 0.14 <sup>b</sup>	3.11 ± 0.12 <sup>ab</sup>	3.16 ± 0.23 <sup>a</sup>	2.44 ± 0.24 <sup>a</sup>	3.21 ± 0.12 <sup>b</sup>	3.17 ± 0.06 <sup>a</sup>
Setifs	7.88 ± 0.25 <sup>a</sup>	7.73 ± 0.16 <sup>b</sup>	7.99 ± 0.19 <sup>b</sup>	7.38 ± 0.23 <sup>a</sup>	3.32 ± 0.12 <sup>b</sup>	2.34 ± 0.29 <sup>b</sup>	2.44 ± 0.19 <sup>a</sup>	2.84 ± 0.16 <sup>a</sup>	3.42 ± 0.17 <sup>b</sup>	2.84 ± 0.25 <sup>b</sup>	3.16 ± 0.14 <sup>b</sup>	3.41 ± 0.19 <sup>ab</sup>

The values (mean ± Std. deviation) with different letters denote the statistical significance determined by ANOVA followed by Tukey test (*P* < 0.05)



**Fig. 6** Agarose gel image of PCR-amplified (TEF1 and ITS1) genes regions of T14 isolate

that not all antagonist isolates are equally effective against the pathogen.

A rapid neutralization of the pathogen was observed. All *Trichoderma* isolates were fast growing, with the mycelial mass invading the Petri dish in only 3 days, while the strains of the pathogen require 6 days. Rapid germination was a decisive element for the competitiveness of *Trichoderma* spp. in antagonism. The same observation was made by El-Komy et al. (2015). In addition to the rapid development of *Trichoderma* strains, they are naturally resistant to many toxic compounds, including herbicides, fungicides and pesticides, and phenolic compounds, and recover very quickly after sub-lethal doses of some of these compounds are added (Harman et al. 2004).

In the in vivo tests, in growth chamber or in greenhouse, T14 isolate was selected to be applied in vitro. This choice was supported by three reasons; firstly, the highest and most stable inhibition rates recorded against the four strains of *F. culmorum*; secondly, T14 recorded the best inhibition score against the pathogen FC2 in in vitro tests (79.54%) (Table 2); and thirdly, the fact that it came from wheat rhizosphere. So, it is already adapted to the environment where it was to be applied. However, the choice of FC2 as a pathogen was random. These experiments were very useful in evaluating the in vivo efficacy of *Trichoderma* biocontrol in reducing FCR and FHB.

According to Erginbas-Orakci et al. (2016), there are several collar inoculation techniques for FCR (seedling dipping, droplet at the base of the stem and colonized grain). The disc technique used in this study was new and used for the first time in this study, taking into consideration the work of Dhingra and Sinclair (1985) who found that the addition of adhesive agents such as gelatin, agar, or methylcellulose is advantageous for inoculation in greenhouse and in field. The inoculation with

FC2 was performed by a mycelial disc around the stems, and the treatment is also applied in the same way for more equality in competitiveness, and more targeting of the crown.

The results obtained in growth chamber showed that FC2 strain induced FCR in all tubes and pots, with a DS% of 68 and 60.5%, respectively. While a very significant decrease in DS% was recorded in tubes and pots treated with T14 strain, with a reduction rate of 50 and 63.63%, respectively (Table 3). Results close to those obtained by Lu et al. (2020) with an in vivo inhibition rate of more than 50% by a strain of *Trichoderma* against *F. graminearum*, and *F. verticillioides* agents responsible for corn stem rot in China. Also, obtained results appeared much more interesting compared to the one obtained by Ghanbarzadeh et al. (2016), where a strain of *T. harzianum* reduced the incidence of basal rot disease in onions by only 25% against *F. proliferatum*. Nevertheless, the fungicidal seed treatment was inferior to that with *Trichoderma*.

In in vivo tests in the greenhouse, the application of the T14 isolate (highly antagonistic in the in vitro test) against the FC2 strain, led to a significant reduction in the AUDPC standard of FHB than the untreated inoculated control. The highest rate of reduction was achieved by the cultivar Setifis (74.60%), followed by Simeto (49.77%) (Table 4 and Fig. 5a). Obtained results are in agreement with those of Saharan et al. (2008), in which, *Trichoderma* spp. had confirmed as the most potent agents against *Fusarium* spp. and induce inhibitory effects on the mycelial growth of *F. graminearum* and *F. semitectum*, the causative agent of FHB in wheat. Other results obtained by Mahmoud (2016) revealed that *T. harzianum* significantly reduced the severity of FHB caused by *F. graminearum* and *F. culmorum*. A highly

significant Pearson correlation between  $AUDPC_{FC2}$  and  $AUDPC_{FC2+T14}$  with  $r = 0.955$ ,  $p < 0.001$ , for the 4 wheat cultivars tested, confirming the positive role induced by the treatment with T14.

The lowest FHB severity was achieved by the cultivar Setifis with  $AUDPC_{FC2} = 16.38\%$ , followed by Bousselem (18.94%), compared to the other cultivars tested (Table 4 and Fig. 5a). This confirms the varietal resistance of these two cultivars to FHB. It is also observed that there was a clear improvement in TKW between the diseased grain and that treated with T14, except Waha and Setifis, where the  $TKW_{FC2 + T14} < TKW_{FC2}$  for unknown reasons (Table 4 and Fig. 5b). The decrease in  $TKW_{T14}$  compared to the control ( $TKW_{T14} < TKW_{control}$ ) was very small and negligible, which encouraged the use of T14 as a preventive treatment without affecting yield.

Study of the physical parameters of the grains from the spike inoculation test in the greenhouse (control, FC2 infected, FC2 infected and treated with T14, and treated with T14 only) revealed that the length parameter L was not at all influenced by FC2 infection; on the other hand, the width parameter W and thickness parameter T showed that the measurements of FC2-infected grains were smaller than those infected by FC2 and treated with T14, which confirmed the biocontrol effect of T14. The difference between control and T14-treated grains were almost similar, which encourage the use of T14 as a preventive treatment without influencing the physical properties of the grains (Table 5).

The identification of microscopic characteristics, as well as the sequences of ITS and TEF1- $\alpha$  (Fig. 6), allowed to identify T14 isolate as *T. afroharzianum*, recently declared as the first report in Algeria (Benttoui et al. 2020; Haouhach et al. 2020). Therefore, the present work was the first to use *T. afroharzianum* as a biocontrol agent in Algeria.

Species identification using molecular characterization tools is very useful in answering the question of whether a particular taxon is present on particular hosts or plants (Abd-Elsalam et al. 2010). This will reduce the severity of FCR and FHB diseases in wheat by using the appropriate biocontrol agent.

## Conclusion

In conclusion, biocontrol agents could play an important role in the protection of wheat. Among the 15 *Trichoderma* spp. isolated in this study, *T. afroharzianum* showed promising results for the control of FCR and FHB in wheat. *Trichoderma afroharzianum* evaluated for the first time in Algeria as a biocontrol agent is recommended as a preventive treatment without affecting yield and without influencing the physical properties of the grains. Additionally, it comes from agricultural soil of wheat, so it is already adapted to the environment where it is to be applied.

## Abbreviations

FCR: Fusarium crown rot; FHB: Fusarium head blight; %GI: Germination inhibition percentage; AUDPC: Area under the disease progress curve; ANOVA: Analysis of variance; TEF1- $\alpha$ : Translation elongation factor1-alpha; ITS1: Internal transcribed spacers; DC: Direct confrontation; IC: Indirect confrontation; PSA: Potato sucrose agar; DS: Disease severity; NaClO: Sodium hypochlorite; PSS: The percentage of symptomatic spikelets; TKW: Thousand kernel weight

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## Authors' contributions

HB conceived the work, designed and performed the experiments, analyzed the data, and wrote the paper. IB was a major contributor in writing the manuscript, reviewing, and editing the paper. WH contributed to the isolation of *Trichoderma* strains. SB contributed to the isolation of *Trichoderma* strains. AB offered the durum wheat cultivars tested. DK was responsible for the supervision and project administration. All authors have read and approved the final manuscript.

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## Availability of data and materials

All data generated or analyzed during this study are included in this published article.

## Declarations

### Ethics approval and consent to participate

Not applicable

### Consent for publication

Not applicable

### Competing interests

The authors declare that they have no competing interests

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