

## Biological Control of Tomato Damping-off Disease Caused by *Rhizoctonia solani* Kuhn

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**ABSTRACT:** Tomato Damping-off disease caused by *Rhizoctonia solani* is considered as one of the most destructive diseases in Egypt. Effect of some biological control agents such as *Trichoderma viride*, arbuscular mycorrhiza (AM) and *Bacillus subtilis*, at seven different mixed treatments, was tested for their ability to control in tomato plants against the disease. According to varietal responses test, all the tested isolates of *R. solani* showed Damping-off symptoms on six tested cultivars; however, they considerably varied in percentages of pre-emergency and post-emergency symptoms. Freda was more susceptible than Alesa cultivar, which was relatively resistant. The combination of AM, with *T. viride*, and *B. subtilis* was the most effective in decrease the infection rate of the tested biological control, compared with the other treatments.

**Kew words:** *Rhizoctonia solani*, *Trichoderma viride*, *Bacillus subtilis*, Arbuscular mycorrhiza.

## INTRODUCTION

Tomato (*Lycopersicon esculentum* L.) is considered one of the most important vegetable crops in Egypt that attack by many soil-borne fungal pathogens (Morsy *et al.*, 2009). *Fusarium solani* and *R. solani* are the most important soil-borne fungal pathogens, which develop in cultured and non-cultured soils, causing the symptoms of damping-off and root rot diseases to wide range of vegetable and crop plants including tomato (Abu-Taleb *et al.*, 2011). The incidence of damping-off was increased from 19 to 90% with increasing the inoculums level of *R. solani*, while the incidence of root rots caused of 10 to 80% losses in different vegetables (Hadwan and Khara, 1992). The tomato cultivars were classified into three groups of resistant, tolerant and susceptible according to their reaction to *Fusarium* and *Rhizoctonia* infection (Moustafa and Khafagi, 1992). Limited information is available on its sustainable management and is generally treated by chemical applications. Overuse of the chemical may result in environmental, human health and pest resistance problem. The increasing awareness of fungicide-related hazards has emphasized the need for adopting biological methods as an alternative disease control method, which is also ecofriendly (Khare *et al.*, 2010).

Biological control is an efficient and environmentally friendly way to prevent damping-off disease. Many microbial species such as *Trichoderma* spp. used to control *R. solani* (Hafez *et al.*, 2013). Mycorrhizal associations are influenced by abiotic and biotic factors, including climate, soil conditions and the identity of host plants. However, the effect of environmental conditions on orchid mycorrhizal associations remains poorly understood (Mujica *et al.*, 2016). Inoculation of carob plants, *Ceratonia siliqua*, with Endomycorrhizae has a significant effect on the growth of these plants. Indeed, the mean values of the

length (61cm) and weight (57.85g) of the aerial part; the length (53.87cm) and weight (53.27g) of the root system, stem diameter (1.15cm) and the number of sheets (139.12) inoculated plants with Mycorrhizal AM are higher than those inoculated simultaneously with the Endomycorrhiza and *Trichoderma harzianum* (Tcomp) respectively, 56.12 cm, 44.47 g, 42.25 cm, 39.82g, 1.03cm, 130.37 and also those only inoculated with *T. harzianum*. Moreover, the frequency (98%) and the intensity (73%) of Mycorrhization being higher in the level of the root seedlings inoculated only with Endomycorrhiza than those co-inoculated with Mycorrhizae AM and *T. harzianum*, respectively 75%; 56% (Zouheir *et al.*, 2016).

Thus, the aim of this work was collecting *R. solani* isolates from different Plant Pathology laboratories, then check their identification by molecular technique, testing their pathogenicity, studying the varietal responses of different tomato cultivars to the disease incidence. Study the efficacy of some biological control agents either single or in combination applications on tomato plants to provide some major resistance subsequently reducing pre- and post-emergence damping-off disease and infection caused by *R. solani*, and hence provide new strategies to control the damping-off disease of tomatoes.

## **MATERIAL AND METHODS**

### **Sources of fungal isolates**

One isolate of tomato damping-off pathogen *R. solani* was isolated from diseased tomato seedlings from Alexandria governorate and 5 isolates were brought from Plant Pathology Laboratories i.e; Agricultural Research Center, Beni Suef, Al Beheira, Al Monufia, Kafr Elsheikh and Faiyum.

### **Isolation, purification and identification of *R. solani***

Isolation was carried out from fresh tomato seedlings, showing Damping-off symptoms. Seedlings were carefully washed in running tap water to remove the adhering particles, then surface sterilized with 95% ethyl alcohol. The infected parts were cut into small pieces by using sterilized scalpel at the zone of infection. The stem pieces were washed several times in sterile distilled water and dried between two sterilized filter papers, then transferred to the surface of plain agar medium in Petri dishes and incubated at 25 °C for 4-7 days. The developed mycelium was carefully transferred to slant potato dextrose agar (PDA) medium. The growing fungus was kept for subsequent purification and identification Dhingra and Sinclair (1985).

Purification of *R. solani* was carried out, using both single hyphal tip techniques (Hansen, 1926). Pure cultures of the obtained isolates were identified in laboratory on the basis of cultural and microscopic characteristics according to Booth (1971), Alexopoulos and Mims (1979), Ellis (1971) and Barnett and Hunter (1987). Pure cultures of the obtained fungal isolates were then transferred to PDA medium slants and kept at 5°C for further studies.

## DNA isolation

Mycelial cultures of 15 *R. solani* isolates were grown in potato dextrose broth media for 5 days at 28°C. Mycelia were harvested and DNA was extracted using CTAB method used by Murray and Thompson (1980).

## Molecular characterization using ITS

Molecular identification of *R. solani* cultures were carried out based on conserved ribosomal internal transcribed spacer (ITS) region. We amplified the ITS regions between the small nuclear 18S rDNA and large nuclear 28S rDNA, including 5.8S rDNA using universal primer pairs ITS1 (5' -TCCGTA GGTGAACCTGCGG-3') and ITS4 (5' -TCCTCCGCTTATT GATATGC-3') (White *et al.* 1990). Amplification was performed on a Thermal Cycler with 25 reaction mixtures containing 2.5 µl of 10X buffer (10 mM Tris-HCl, pH 8.8); 2.5 mM MgCl<sub>2</sub>; 2 mM each of dNTP; 25 pmol ml<sup>-1</sup> primer (each of ITS-1 and ITS-4); 1U of Taq DNA Polymerase; 60–100 ng genomic DNA. The amplification cycle consists of an initial denaturation at 95 C for 2 min followed by 35 cycles at 94 C for 30 s, 56 C for 1 min, and 72 C for 2 min and a final extension at 72 C for 10 min. Amplified PCR products were separated on an agarose gel (1.5% w/v) in 1X TAE buffer. They were then eluted, and further sequencing was carried out.

## ITS data analysis

The ITS nucleotide sequences for each isolate were then compared to those in the public domain databases NCBI (National Center for Biotechnology information; [www.ncbi.nih.gov](http://www.ncbi.nih.gov)) using Basic Local Alignment Search Tool for Nucleotide Sequences (BLASTN). Alignment of ITS DNA sequences was done using Clustal W program (Vincelli and Tisserat, 2008).

## Varietal responses

In order to check varietal response throughout this study, six isolates of *R. solani* that varied in aggressiveness obtained from diseased tomato plants were used. *R. solani* isolates were selected according to their mycelia and sclerotia characteristics. Six isolates were individually tested for their pathogenicity on Alesa, Alia, Princesa, Jana, Alex. and Frida tomato cultivars under greenhouse conditions. Pots capacity 1 kg were sterilized by immersing them in 5% formalin solution, then left 3 weeks to allow formaldehyde evaporation and filled with autoclaved aerated clay soil. Fungal inocula were grown on sterilized PD medium in 250 ml flasks for 2 weeks at 18°C. Soil infestation with the tested pathogen was carried out using the inoculum which was mixed thoroughly with the soil in each, watered and left for one week to secure establishment of the inoculated fungi. Control pots were filled with the same soil mixed and watered. Seeds of tomato cultivars were surface sterilized using 1% sodium hypochlorate solution for 2 minutes, washed with sterilized water, dried and sown at a depth of 2 cm and watered regularly 7 days under greenhouse conditions. The number of pre-emergence and post-emergence damping-off seedlings were recorded 10-30 days after planting.

### **Effect of antagonistic fungi on the disease incidence caused by *R. solani***

Plastic pots were filled with autoclaved sandy loam soil as previously mentioned. Sowing was carried out 8 days after inoculation. Five seeds of vicar genotype were planted (3 replicates). Before planting, seeds were surface sterilized by dipping in 1% sodium hypochlorite for 2 min., washed with distilled water, then dried under laminar flow. Seeds coated were performed by wetting them with sterile water containing molasses, air dried and then placed on the surface of 14 days-old culture of

antagonistic fungi in Petri plates in that conidia were abundant. Control treatment was done by soaking seeds in distilled water. Data of the disease incidence was recorded after 15 and 45 days.

### **Biocontrol agents**

Certain antagonistic microorganisms, *T. viride*, AM and *B. subtilis* were used in greenhouse experiments to study their effectiveness as biological control against the highly pathogenic isolate of *R. solani*.

### **Source and preparation of inoculate**

#### **Preparation of pathogen isolate**

Virulent Isolates of *R. solani* were selected from pathogenicity tests. Mycelium suspension was prepared by growing isolate in wheat grain medium and incubated at 25-28 °C for 15 days.

### **Source Mycorrhiza**

Arbuscular mycorrhiza (*Glomus intraradices*) was used in this experiment. The soil was mixed with 20 ml mycorrhiza one week before planting as described by Malibari *et al.* (1990). Also, 10 ml inoculums were added with the seeds at sowing time. The arbuscular mycorrhiza (AM) was obtained from Germany and activated in the Soil Microbiology Lab - Soil and Agricultural Chemistry Department, Faculty of Agriculture, Saba Basha, Alexandria University, Alexandria, Egypt.

### **Soil preparation**

The clay soil was used in the present experiment. Soil samples each weighing 1000 g were separately filled into plastic pots (15cm). Five seeds of tomato were sown in each pot. The NPK, 20:20:20 fertilizer was added at two equal doses (20 days and 34 days after sowing) at rate 92 mg N/ 50 ml water for each pot. The biocontrol AM were applied throughout this experiment. One week before planting, seeds were mixed with mycorrhiza and in each hill as 10g each treatment contains AM alone or mixed with other treatment.

### **Statistical analysis**

The collected data were arranged in a completely randomized design and replicated four times. Data were statistically analyzed for ANOVA and means compares to fulfill the significance according to Steel and Terrie (1982). Significance level of  $\alpha=0.05$  was used in all analysis.

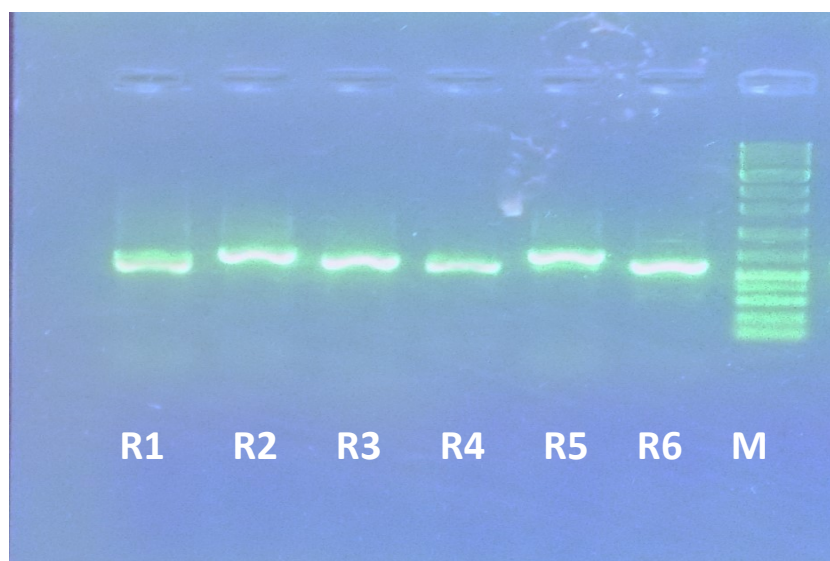
## RESULTS AND DISCUSSION

### Molecular identification of *R. solani* isolates based on ITS

The identification of *R. solani* isolates using the ITS1 and ITS4 regions, including the 5.8S rDNA gene of the isolates studied (Fig. 1). *Rhizoctonia* sequences obtained from amplification of conserved ribosomal ITS region (Fig. 2) was compared with sequences from National Center for Biotechnology Information (NCBI) database using BLAST 2.0 (<http://www.ncbi.nlm.nih.gov/BLAST>). These sequences were identified and deposited in NCBI GenBank (Table 1). All the isolates identified as *R. solani*.

**Table (1). Accession number of *R. solani* isolates on Gene Bank**

Isolate code	Gene Bank Accession no.
Rs01	MH687911
Rs02	MH687912
Rs03	MH687913
Rs04	MH687914
Rs05	MH687915
Rs06	MH687916



**Fig. (1). DNA profile generated by ITS primers on *Rhizoctonia solani* isolates; R1=Alexandria, R2= Beheira, R3= Beni Suef, R4= Faiyum, R5= Monufia, R6= Kafar Elsheikh, M=100bp marker.**

**>Rs01**

tcattattga attnatgtag agtttggtg tagctggctc ctaattaaac ttgggggcat  
gtgcacacct ttctcttca tcccatacac acctgtgcac ctgtgagaca gatgtttgt gggggggaag  
gaactttatt ggaccttcta ctccccctng acttctgtct acttaattca tataaactca atttatttta  
aaacggatgt aatggatgta acacatctaa tactaagttt caacaacgga tctcttggct ctgcatcga  
tgaagaacgc agcgaatgc gataagtaat gtgaattgca gaattcagtg aatcatcgaa tcttgaacg  
caccttgccg tccttggat tccttggagc atgcctgttt gagtatcatg aaatctcaa agtcaaacct  
ttgttaatt caattggtc tgcttggta ttggaggatt attgcagctt cacacctgct cctcttgtg  
cattagctgg atctcagtgat tatgcttgg tccactcagc gtgataagtt atctatcgct gaggacaccc  
tgtaaaaag gggtgccaa ggtaaatgca gatgaaccgc ttctaagatg ccattgactt ggacaatatc  
taatttatga tctgatctca aatc

**>Rs02**

aaggatcatt attgaattaa tgtagagttt ggtttagctt ggctcctaat taaacttggg  
ggcatgtgca cacctttctc ttcatocca tacacacctg tgcacctgtg agacagatgt tttgtggggg  
ggaaggaact ttattggacc ttctactccc ccttgacttc tgttactta attcataaa actcaattta  
ttttaaaccg gatgtaatgg atgtaacaca tctaacta agttcaaca acggatctct tggctctcgc  
atcgatgaag aacgcagcga aatgcgataa gtaatgtaa ttgcagaatt cagtgaatca tcaaatctt  
gaacgcacct tgcctcctt ggtattcctt ggagcatgcc tgtttgagta tcatgaaatc tcaaagtca  
aacctttgt taattcaatt ggttctgctt tggattgga ggattattgc agcttcacac ctgctcctct  
ttgtcatta gctggatctc agtgttatgc ttggttccac tcagcgtgat aagttatcta tgcctgagga  
caccctgtta aaaaggggtg gccaaagtaa atgcagatga accgcttcta atagtccatt gacttgaca  
atatctaatt tatgatctga tctcaa

**>Rs03**

tcattattga attnatgtag agtttggtg tagctggctc ctaattaaac ttgggggcat  
gtgcacacct ttctcttca tcccatacac acctgtgcac ctgtgagaca gatgtttgt gggggggaag  
gaactttatt ggaccttcta ctccccctng acttctgtct acttaattca tataaactca atttatttta  
aaacggatgt aatggatgta acacatctaa tactaagttt caacaacgga tctcttggct ctgcatcga  
tgaagaacgc agcgaatgc gataagtaat gtgaattgca gaattcagtg aatcatcgaa tcttgaacg  
caccttgccg tccttggat tccttggagc atgcctgttt gagtatcatg aaatctcaa agtcaaacct  
ttgttaatt caattggtc tgcttggta ttggaggatt attgcagctt cacacctgct cctcttgtg  
cattagctgg atctcagtgat tatgcttgg tccactcagc gtgataagtt atctatcgct gaggacaccc  
tgtaaaaag gggtgccaa ggtaaatgca gatgaaccgc ttctaagatg ccattgactt ggacaatatc  
taatttatga tctgatctca aatc

**>Rs04**

tcattattga attnatgtag agtttggtg tagctggctc ctaattaaac ttgggggcat  
gtgcacacct ttctcttca tcccatacac acctgtgcac ctgtgagaca gatgtttgt gggggggaag  
gaactttatt ggaccttcta ctccccctng acttctgtct acttaattca tataaactca atttatttta  
aaacggatgt aatggatgta acacatctaa tactaagttt caacaacgga tctcttggct ctgcatcga  
tgaagaacgc agcgaatgc gataagtaat gtgaattgca gaattcagtg aatcatcgaa tcttgaacg  
caccttgccg tccttggat tccttggagc atgcctgttt gagtatcatg aaatctcaa agtcaaacct  
ttgttaatt caattggtc tgcttggta ttggaggatt attgcagctt cacacctgct cctcttgtg  
cattagctgg atctcagtgat tatgcttgg tccactcagc gtgataagtt atctatcgct gaggacaccc  
tgtaaaaag gggtgccaa ggtaaatgca gatgaaccgc ttctaagatg ccattgactt ggacaatatc  
taatttatga tctgatctca aatc

**>Rs05**

aattaaactt gggggcatgt gcacaccttt ctcttcatc ccatacacac ctgtgcacct  
gtgagacaga tgtttgtgg gggggaagga actttattgg accttctact ccccttgac ttctgtctac  
ttaattcata taaactcaat ttattttaa acggatgtaa tggatgtaac acatctaata ctaagttca  
acaacggatc tcttggtct cgcacgatg aaraacgcag cgaaatgcga taagtaatgt gaattgcaga  
attcagtgaa tcatcgaatc ttgaacgca ccttgcgctc ctgggtattc ctggagcat gcctgttga  
gtatcatgaa atctcaaag tcaaacttt tgtaattca attggtctg ctttggatt ggaggattat  
tgcagctca cacctgctcc tcttgtgca ttagctggat ctacagtta tgcttggtc cactcagcgt  
gataagttat ctatcgtga ggacaccctg ttaaaaagg gtggccaagg taaatgcaga tgaaccgctt  
ctaatagtcc attgacttg acaatatcta attatgatc tgatcctaaa tcaggttaga ctaccgctg  
aactaagca tatcaataag c

**>Rs06**

tagctggctc ctaattaaac ttgggggcat gtgcacacct ttcttttca tccatacac  
acctgtcac ctgtgagaca gatgtttgt gggggggaag gaactttatt ggaccttcta cccccctg  
acttctgtct actaattca tataaactca atttattta aacggatgt aatggatga acacatctaa  
tactaagttt caacaacgga tctctggct ctgcacgca tgaagaacgc agcgaaatgc gataagtaat  
gtgaattgca gaattcagtg aatcatcgaa tcttgaacg caccttgcgc tccttggat tcctggagc  
atgcctggtt gactatcatg aatcttcaa agtcaaact tttgtaatt caattggctc tgcttggta  
ttggaggatt attgcagctt cacacctgct cctcttctg cattagctgg atctcagtg tatgcttgg  
tccactcagc gtgataagtt atctatcgt gaggacccc ttttaaaaag gggggcca ggtaatgca  
gatgaaccgc ttctaagct ccattgact ggacaatc taattatga tctgatcctaatcaggtag  
gactaccgc tgaactaag c

**Fig. (2). Sequence of different isolates of *R. solani* submitted to Gene Bank**

**Pathogenicity experiments**

Inoculation of the six tested tomato cultivars; Alia, Freda, Jana, Alesa, Princesa and Alex., was carried out using the tested *R. solani* isolates in greenhouse. Data presented in Table (2) showed disease damping off in all the investigated locations. Frequency of the total isolates from root and stem, were estimated for both isolation techniques in all regions of the six investigated governorates. From the obtained results, all the tested *R. solani* isolates were significantly virulent on susceptible Freda, Alia, Jana and Princesa cultivars (Infection rates were 71.42, 48.57, 33.33 and 27.61%, respectively). However, in resistant Alesa and Alex cultivars, infection rates from 24.76 and 19.04%. However, they considerably varied in disease damping off. The highest infection values were obtained by isolate of Beni Suef and the lesser infection values was obtained by isolate from Alexandria, compared with the other tested isolates.

**Table (2). Varietal responses of six tomato cultivars to the tested isolates of *R. solani***

Infection rate of Pathogen factor (%)			
Isolates	Pre-emergence	Post-emergence	infection
Beni Suef	74.44 <sup>a</sup>	10.0 <sup>a</sup>	84.44 <sup>a</sup>
Monufia	50.00 <sup>b</sup>	7.77 <sup>ab</sup>	57.77 <sup>b</sup>
Beheira	24.44 <sup>c</sup>	7.77 <sup>ab</sup>	32.22 <sup>c</sup>
Alexandria	18.88 <sup>c</sup>	5.55 <sup>ab</sup>	24.44 <sup>c</sup>
Kafr Elsheikh	23.33 <sup>c</sup>	7.77 <sup>ab</sup>	31.11 <sup>c</sup>
Faiyum	25.55 <sup>c</sup>	6.66 <sup>ab</sup>	32.22 <sup>c</sup>
Control	0.00 <sup>d</sup>	0.00 <sup>b</sup>	0.00 <sup>d</sup>
L.S.D 0.05	10.85	8.15	12.13
Infection rate of Tomato cultivar factor (%)			
Cultivar	Pre-emergence	Post-emergence	infection
Freda	60.00 <sup>a</sup>	11.43 <sup>a</sup>	71.42 <sup>a</sup>
Alia	40.95 <sup>b</sup>	7.61 <sup>a</sup>	48.57 <sup>b</sup>
Jana	27.61 <sup>c</sup>	5.71 <sup>a</sup>	33.33 <sup>c</sup>
Alex	20.95 <sup>cd</sup>	3.80 <sup>a</sup>	24.76 <sup>cd</sup>
Princesa	21.90 <sup>cd</sup>	5.71 <sup>a</sup>	27.62 <sup>cd</sup>
Alesa	14.28 <sup>d</sup>	4.76 <sup>a</sup>	19.04 <sup>d</sup>
L.S.D 0.05	10.04	7.55	11.23

Values followed by the same letter(s) in each column didn't differ significantly according to Fisher's LSD Test ( $P \leq 0.05$ ).

### The effect of biological control agents on disease incidence

In order to check the efficacy of the tested biocontrol treatment on controlling damping off disease of tomato plant of resistant, Alesa and susceptible Freda tomato cultivars were treated with AM, *T. viride* and *B. subtilis* with *R. solani* infection rates was determined in both cultivars at the end of experiments. Results were statistically analyzed and presented in Table (3) obtained in the treatment *T. viride* and *B. subtilis* in agreement with Riad *et al.* (2016). All tested bioagents reduced the linear growth of *F. solani*. Complete inhibition of the linear growth was obtained with *T. viride*, *T. harzianum* and *B. subtilis*.

Also, the results obtained in the treatment *T. viride* and AMF agreed with who obtained with Tanwar *et al.* (2013). The influences of AM fungi (*G. intraradices*) and *Acaulospora laevis* (A) with *T. viride* (T) and *P. fluorescens* (P) alone and in combinations on growth of broccoli plants under glasshouse conditions.

This study was similar with Riad *et al.* (2016) who reported dry root rot and slow decline diseases of citrus caused by *F. solani* and *Tylenchulus semipenetrans*, respectively, are serious diseases attacking many groves in Egypt. The most effective treatment against *F. solani* was compost + mixture of *T. harzianum* + *T. viride* which reduced disease incidence and severity by 87.5%. The highest reduction in total count of *F. solani* was obtained with compost + mixture of *T. harzianum* + *T. viride* which reduced total count by 82.1%.



These results were in agreement with Patro *et al.* (2018) who reported that during *Kharif* 2016 and 2017, The disease severity and yield parameters (grain yield and straw yield) were evaluated against banded blight using different combinations of potential biocontrol agents *viz.*, *B. subtilis*, *P. flourescens* and *T. asperellum* in the field during 2016 and 2017. Among all treatments applied treatment T7 (*i.e.* Soil application of value added *P. flourescens* + *T. asperellum* + *B. subtilis* (one kg talc formulation mixed in 25 kg FYM or vermicompost, incubated for 15 days) applied over an acre at the time of sowing) showed maximum reduction in disease intensity (22.81%) and (50.67 %) with higher grain and fodder yield over control. Ainhoa *et al.*, (2010) reported that the AM-inoculated plants were effective in controlling *Fusarium* wilt, *G. mosseae* - inoculated plants showing the greatest capacity for reduction of disease incidence. The *T. harzianum*-inoculated plants were more effective than AMF-inoculated plants with regard to suppressing disease incidence. Co-inoculation of plants with the AMF and *T. harzianum* produced a more effective control of *Fusarium* wilt than each AMF inoculated alone, but with effectiveness similar to that of *T. harzianum*-inoculated plants.

**Table (3). Responses of tomato cultivars treated with biocontrol agents to infection with *R. solani***

Treatment	Infection percent %		
	Pre-emergence	Post-emergence	Infection %
AM+B+T *	8.33 <sup>c</sup>	0.00 <sup>b</sup>	8.33 <sup>c</sup>
B+T	15.00 <sup>bc</sup>	3.33 <sup>b</sup>	18.33 <sup>bc</sup>
AM+T	8.33 <sup>c</sup>	1.66 <sup>b</sup>	10.00 <sup>bc</sup>
AM+B	13.33 <sup>bc</sup>	0.00 <sup>b</sup>	13.33 <sup>bc</sup>
T	15.00 <sup>bc</sup>	3.33 <sup>b</sup>	20.00 <sup>bc</sup>
B	18.33 <sup>bc</sup>	5.00 <sup>ab</sup>	21.66 <sup>b</sup>
AM	21.66 <sup>ab</sup>	0.00 <sup>b</sup>	21.66 <sup>b</sup>
Control	31.66 <sup>a</sup>	10.00 <sup>a</sup>	41.66 <sup>a</sup>
L.S.D 0.05	10.30	5.61	10.90
Tomato cultivar factor			
Cultivar	Pre-emergence	Post-emergence	Infection %
Alesa	6.25 <sup>b</sup>	3.75 <sup>a</sup>	8.33 <sup>b</sup>
Freda	26.66 <sup>a</sup>	1.25 <sup>a</sup>	30.41 <sup>a</sup>
L.S.D 0.05	5.15	2.80	6.69
Pathogen factor			
Isolate	Pre-emergence	Post-emergence	Infection %
Beni Suef	29.58 <sup>a</sup>	3.33 <sup>a</sup>	32.91 <sup>a</sup>
Alexandria	3.33 <sup>b</sup>	1.66 <sup>a</sup>	5.83 <sup>b</sup>
L.S.D 0.05	5.15	2.80	5.45

Values followed by the same letter(s) in each column didn't differ significantly according to Fisher's LSD Test ( $P \leq 0.05$ ).

\* AM = Mycorrhiza, T = *T. viride*, B = *B. subtilis*.

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الملخص العربي  
المكافحة الحيوية لمرض الذبول الطرى فى الطماطم المتسبب عن الفطر  
ريزوكتونيا سولانى

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يعتبر مرض الذبول الطرى فى الطماطم المتسبب عن الفطر ريزوكتونيا سولانى أحد أهم الامراض المدمرة للمحصول فى مصر. تم دراسة تأثير المكافحة الحيوية باستخدام فطر تريكوديرما فيرىدى والميكوريزا الوعائية وبكتيريا باسيلس ستلس لاختبار قدرتهم على مكافحة المرض. طبقا لاختبار إستجابة الأصناف فان جميع عزلات المسبب المرضى ريزوكتونيا سولانى أظهرت أعراض الذبول الطرى فى أصناف الطماطم الستة المختبرة. وقد أظهرت الأختلافات بين الاصناف الستة المختبرة حدوث أعراض ذبول بادرات الطماطم قبل وبعد ظهور البادرات . وقد أظهرت النتائج أن صنف فريدة هو أكثر الاصناف حساسية للإصابة بينما صنف أليسا هو الأكثر مقاومة. وقد تبين أن مخلوط كل من فطر الميكوريزا و التريكوديرما فيرىدى والبكتيريا باسيلس ستلس أكثر فاعلية فى خفض نسبة الإصابة مقارنة بالمعاملات الأخرى.