



Biocontrol Mechanisms of *Bacillus* sp. and its Application as a Bionanocomposite against *Rhizoctonia solani*

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ABSTRACT

Black scurf disease caused by *Rhizoctonia solani* is a main yield limiting factor for potato tuber production as it leads to plant death. Using chemical treatment is not economic, therefore, the present study aimed to isolate naturally occurring antagonistic bacteria that could control *R. solani*. *Bacillus* sp. was selected among of 84 isolates secured from rhizosphere of healthy potato plant based on its ability to suppress the growth of the pathogen *R. solani*. Phylogenetic analysis of this strain based on 16S rRNA gene sequences showed highest similarity (99%) with *Bacillus* sp.; it was deposited in the GenBank under the accession number of MK030136. The strain culture filtrate containing protease, diffusible antibiotic, hydrogen cyanide and siderophore was capable of inhibiting growth of the pathogen up to 15 days compared to 7 and 10 days for other *Bacillus* isolates. It also produces indole acetic acid which promoted plant growth. Morphological and structural changes that took place as a result of *Bacillus* sp. and *R. solani* interaction were evaluated using light, scanning and transmission electron microscopies. The results showed that *Bacillus* sp. caused loss of structural integrity, abnormal coiling, shriveling and lysis of the *R. solani* hyphae, in addition to complete cytoplasm and internal organelles depletion. The *Bacillus* sp. under study was immobilized on nanoclay to form a bionanocomposite, which was stable and exhibited the biocontrol efficiency along 8 months storage. Both in vitro and greenhouse experiments showed high inhibition

KEYWORDS

Antifungal activity; Bacillus sp.; Rhizoctonia solani; biocontrol; mechanism of action, bionanocomposite.

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of *R. solani* radial growth. Results indicate that the prepared bionanocomposite is a promising alternative to the commercial products.

INTRODUCTION

Potato is the third most important food crop in the world after rice and wheat in terms of human consumption (FAO, 2014). It is the world's fourth produced food product, it has the ability to grow in a wide range of climatic conditions, it is highly nutritive and contains high number of vitamins, minerals, proteins, antioxidants, essential amino acids and carbohydrates (FAO, 2009). The Food and Agriculture Organization (FAO) emphasizes on the significant role potatoes play in the world global food system. It is important to maintain a healthy produce in terms of quality and quantity since potatoes provide food, feed and fibre. All of which are needed to sustain the health requirements of a rapidly growing population (Shafi et al., 2017). One of the many problems surrounding potato cultivation that affect the potato economic yield on a global, national and individual farmer level is fungal pathogenesis. Black scurf is one of the critical diseases caused by the fungus *Rhizoctonia solani*. Once this fungus is present in the soil, it is difficult to control due to its broad host range, saprophytic life cycle and persistence of the sclerotia and mycelium in soil and plant material (Ajayi-Oyetundea and Bradley, 2018). Within a few years, this plant disease causes the loss of about 34% of the produce in addition to changes in size and distribution of tubers (Kumar et al., 2017). This problem is a worldwide and is usually addressed by chemical –based-pesticides and fertilizers (Shafi et al., 2017). However, the use of chemical pesticides leads to an increase in production cost, health hazards, environmental pollution and pesticide residual effects on human health and soil ecosystem. In addition to the development of resistant pathogens due

to prolonged exposure to antifungal chemicals. An alternative to chemical pesticides is the use of biological control that entails applying microbial cells or their metabolites to attack fungal pathogens (Tortora et al., 2011; Nandi et al., 2017). Several mechanisms of action are thought to be involved in the process of biological control, all of which are based on ecological interactions, such as competition for space and nutrients, parasitism (production of hydrolytic enzyme such as proteases, cellulases and chitinases), antibiosis and induction of plant defenses (Jamalizadeh et al., 2011). Many microorganisms were reported to have fungal antagonistic activity and can be used as biocontrol agents. The use of *Bacillus* sp. has increased due to its ability to replicate rapidly, form endospores, tolerate extreme pH, temperature and osmotic conditions as well as produce several antibiotics, siderophores and hydrolytic enzymes (Chaurasia et al., 2005). In order to maintain viability and efficiency of a biocontrol agent, the delivery system is prepared as a formulation where the cells are immobilized on a support (Ma et al., 2015). Immobilization of cells on nanoparticles provides advantages over traditional support materials, it increases mass transfer and minimizes diffusion limitation. Therefore, the aim of this study is to isolate bacteria with efficient biocontrol activity against *Rhizoctonia solani*, depict its mechanism of action and test its efficacy as a bionanocomposite in vitro and in green house.

MATERIALS AND METHODS

Collection of soil samples

Four cultivated soil samples were collected in sterile plastic bags from the rhizosphere of healthy plants of potato from four Egyptian governorates: Sharkia, Giza, Beheira and Menofia. The rhizosphere was ditch with intact root system. Samples were air dried for 24 h at ambient temperature before use.

Isolation of bioantagonistic bacteria and Rhizoctonia solani

Ten grams of each soil sample were transferred into 250 ml conical flask containing 90 ml of sterilized saline solution, and shaken for 1 h at 150 rpm. The resulting soil suspension was serially diluted up to 10^{-7} , 0.1 ml of each dilution was plated onto nutrient agar medium, plates were incubated at 30°C for 48 h. Developed colonies were picked up, purified through streaking plate method and sub-cultured on Luria Bertani agar medium slants. A total of 84 purified isolates was screened for antifungal activity against the isolated pathogenic *Rhizoctonia solani* from potato plant (*Lady rossetta*) that showed symptoms of *R. solani* disease. Pieces of potato tubers were submerged in 5% sodium hypochloride for five minutes then washed with sterile distilled water and placed on Petri dishes containing potato dextrose agar (PDA) and incubated at 22°C for five days. Fifty µl of 24 h old bacterial suspensions obtained from different samples were placed on sterile paper disks. Disks were placed on Petri dishes containing PDA surrounding 10 mm diameter disk containing the mycelium of *R. solani* placed in the center of plates which were incubated at 30 °C for 7 days. The inhibition of mycelium growth was checked and was repeated three times. Results are expressed as percentage of growth inhibition of *R. solani* in the presence of the bacterial isolates. The percent growth inhibition (PGI) was calculated using the following formula:

$$PGI (\%) = \frac{KR - RI}{KR} \times 100$$

Where: KR represents the distance (measured in mm) from the point of inoculation to the colony margin on the control Petri plate, and R1 is the distance of fungal growth from the point of inoculation to the colony margin on the treated Petri plate in the direction of the antagonist (**Raupach and Kloepper,**

1998). The zone of inhibition was recorded as the distance between the fungal pathogen and the area of antagonist growth after 7 days. Based on the results of antagonistic activity, 3 isolates were selected for further studies.

Bioproducts from Bacillus isolates that affecting the growth of R. solani

Hydrolytic enzymes

The three isolates were screened for their abilities to produce hydrolytic enzymes that can degrade polymeric compounds including chitin, proteins, amylase, lipase and cellulase by adopting the agar plate screening. The isolates were grown on Luria Bertani agar supplemented with substrates for each enzyme as 1% w/v casein, 0.2% carboxymethyl cellulose, 2% colloidal chitin, 1% Tween 20 and 2% w/v soluble starch as substrates for protease, cellulase, chitinase, lipase and amylase, respectively. The five mm plug of each isolate was placed at the center of each enzyme screening agar plate and incubated at 30°C for 24 h. Colonies which exhibited surrounding clear zone were considered as positive results for enzyme production (**Thakaew and Niamsup, 2013**). Each experiment was performed in three replicates.

Volatile antifungal compounds

To verify the production of volatile compounds by the isolates, a culture disk (5 mm in diameter) of the isolates was placed at the center of one half Petri plate containing nutrient agar and 5 mm disk of a 4- day old culture of *R. solani* was placed at the center of another Petri plate containing PDA. Both half plates were placed face to face preventing any physical contact between the bacteria and the pathogen, and were sealed with paraffin film to prevent loss of volatiles formed (**Montealegre et al., 2003**)

Diffusible antifungal compounds

Antagonistic activities of the extracts were tested by agar well diffusion method (**Shanmuga et al.,**

2002). PDA medium (20 mL) was poured into each sterile Petri dish, followed by placement of mycelial disk (5 mm in diameter) of the *R. solani* at the center of the plates. A well (5 mm in diameter) was made by punching the agar with a sterile steel borer at a distance of 3 cm from the central 5mm fungal inoculum. Each *Bacillus* isolate was cultivated in 50 ml nutrient broth and incubated without shaking at 30°C for 72 h. Ten ml aliquots were withdrawn from each flask, centrifuged at 6000 rpm for 20 min. The supernatant was collected and filtered through millipore membrane (0.22µm) and 50 µl of each extract were poured into the well and one well was used as a control. The plates were incubated at 30°C for 7 days.

Siderophore production

Bacillus isolates were cultivated on blue agar medium at 30°C for 24h (Adhikari et al., 2013). An orange halo surrounding bacterial colonies indicated siderophore production.

HCN determination

HCN production test was performed as described by Kumar et al. (2012). Bacteria were heavily inoculated in nutrient agar plates and incubated in an inverted position at 30°C for 24h with filter paper strips dipped in picric acid placed inside the lids. Change of color of the indicator strip from yellow to brown was considered as a positive result.

Production of the plant growth promoting phytohormone IAA.

Production of IAA (indole -3-acetic acid) was determined according to Patten and Glick (2002). 24h old bacterial cultures grown on Luria-Bertani (LB) broth supplemented with L-tryptophane were centrifuged at 6000 rpm for 20 min to obtain culture supernatants. The supernatants (1ml) were mixed vigorously with 4 ml Salkowskis reagent (Gordon and Weber., 1951). Development of pink color indicated a positive result.

Dual culture of isolates and pathogen for long incubation period

To confirm the ability of the isolates for fungicidal action, dual cultures were prepared with *R. solani* and each of the three isolates. A 5 mm disk of pathogen mycelium and 5-mm round filter papers were wet with 50µl of bacterial suspension. The filter paper and pathogen disk were placed opposite to each other on a potato dextrose agar (PDA) plate and incubated at 30°C in the dark for 3 days, during which the growth margins were inspected regularly. The diameters of inhibition areas on the PDA plates were recorded up to 15 days (Yang and Sung, 2011).

Identification of biocontrol active isolates

The isolates were characterized by morphological and biochemical tests according to Bergey's Manual of Systematic Bacteriology (Sneath et al., 1986). The isolate that displayed 15 days of inhibition was subjected to 16S rRNA phylogenetic identification. DNA of a 24 h culture was extracted in 1ml TE buffer (pH 8). The cell suspension was boiled for 10 min to release DNA then chilled on ice for 10 min. The suspension was centrifuged at 10,000 rpm for 5 min. About 50 ng of DNA template was added to 45 µl of PCR reaction solution (Macro Gen) using the following primer sets 27-8 GAGTTTGATCCTGGCTCAG and 1492 GGTTACCTT GTTACGA (Edwards et al., 1989). The amplification was performed as follows: 35 amplification cycles at 94°C for 45 sec, 55°C for 60 sec and 72°C for 60 sec. An aliquot of 5-15 µl of PCR reaction products was electrophoresed on a 1% agarose gel containing ethidium bromide (10 mg/ml in dH₂O) and the DNA bands were visualized under the UV light. The amplified PCR products were submitted to Solgent Co Ltd (South Korea) for purification and sequencing. The resulted sequences were trimmed and assembled in Geneious software (Biomatters). The sequence was compared to the NCBI nucleotide database (www.ncbi.nih.gov/blast)

and phylogenetic tree of the strain given an accession number MK030136 was constructed based on the 16s-rRNA sequence comparisons length polymorphism of the PCR-amplified and sequences from database using blast tree construct in (<https://blast.ncbi.nlm.nih.gov/blast/treeview/treeView.cgi>).

Light, scanning electron (SEM) and transmission electron microscopy (TEM) studies for antagonistic effects of Bacillus sp. on mycelia morphology of R. solani.

For light microscopy, *R. solani* cultures before and after treatment with *Bacillus* sp. were mounted on clean glass slides and stained using Lactophenol cotton blue dye. Slides were examined using 40X lens. The antagonistic properties of *Bacillus* sp. were studied using SEM (Kumar *et al.*, 2013). Plugs of mycelium (1mm in diameter) cut from the fungal mycelia growing towards the inhibition zone were managed for SEM by the following procedure: agar discs placed on cover glasses were treated with 2 % osmium tetra oxide vapors for 24h at 20° C. The treated samples were attached to aluminum stubs with double adhesive tape, coated with gold, and then captured using JOEL JMS 5600 scanning electron microscope. Mycelial growth in control plates was observed and compared with the hyphal deformities near the inhibition zone. For TEM assay, cells were infiltrated with acetone for 24h and polymerized at 60 °C for 48 h. Fifty nanometer sections were sectioned, then the sections were placed on copper sieves and contrasted with uranyl acetate and lead citrate for 30 min each (Zhang *et al.*, 2013) , the sections were viewed on JEOL-JEM 1010 TEM.

Bionanocomposite experiments:

Immobilization of Bacillus sp. on nanoclay particles

Bacillus sp. cells were initially cultured for 24 h at 30°C. Ten ml of the preculture (ca. 6×10^7 CFU/

mL) were transferred into 10 mg of clay nanoparticles (bentonite) obtained from Medical Export Company (UK) and cultivated for 72 h at 30°C under static conditions. Immobilization of bacterial cells was characterized by scanning electron microscopy (JOEL JMS 5600), X-Ray diffraction (6000 SHIMADZU), the dried powder of bionanocomposite homogenized was pressed in a stainless-steel sample holder for collecting the XRD patterns using CuK α radiation operating at 40 kV and 40 mA. Fourier Transform Infrared Spectroscopy (FT-IR) scanning was performed from 400 to 4000 nm using FTIR, BRUKER VERTEX 70 device

Shelf life of bionanocomposite and its antagonistic activity during storage time

The viability of *Bacillus* sp. on clay nanoparticles was examined monthly up to 8 months using the dilution plate method. Plates were incubated at 30°C for 24h and the bacterial population was determined. Ten μ l of bionanocomposite solution (2.7.1) stored at room temperature (8 months) were inoculated at the opposite side of PDA plates and incubated with 6mm diameter of *R. solani* as previously mentioned (Isolation of bioantagonistic bacteria and *Rhizoctonia solani*)

Evaluation of bionanocomposite antagonistic activity against R. solani.

The experiments were carried out during the growing season 2019. Sterilized plastic pots (20 cm in diameter), each contains 3 Kg of sterilized soil were infested with *R. solani* grown on sand corn meal medium at the rate of 5% soil weight. Pots were watered regularly for 10 days before planting to insure distribution of inoculum. Since the Lady Roseta cultivar is one of the most sensitive cultivars, it was selected for greenhouse experiment. Healthy potato tubers seeds (cv. Lady Rosetta- susceptible) having about four eyes (buds) and uniform in size and weight possible were selected and washed to

remove existing soil, surface sterilized in 2.0 % sodium hypochlorite solution for 3 min; rinsed 3 times in sterile distilled water then air dried (Tariq *et al.*, 2010). Sterilized tubers were sown at the rate of 3 tubers / pot. Tubers were dipped in either 5L of bio-nanocomposite suspension or Bio Arc commercial formulation (6%) purchased from Central Agriculture Pesticide Laboratory of A.R.C. The immersion time for both treatments was 0, 10, 20, 30 and 60 min. Data were recorded after 60 days of planting using non-soaked tuber (negative control) and Bio Arc soaked tubers (positive control) in infested soil as controls. The reduction of disease incidence was calculated according to the following formula: disease incidence % = number of infected plant /total number of plants assessed \times 100. Plant heights as well as fresh and dry weights were also recorded at the end of the experiment.

Statistical analysis

Statistical analysis was performed using one-way analysis of variance (ANOVA) by Minitab software (version 17). P-Value < 0.05 considered as significant.

RESULTS

Characteristics of *R. solani*

R. solani used in the antagonism was isolated from the infected parts of Lady Rossetta potato plant. Pure culture of isolated fungus was identified according to the cultural properties, morphological and microscopical characteristics (Fig 1). Results showed that *R. solani* slightly melanized hyphae and irregularly shaped. Microscopic observation revealed that the hyphae branched at a 90° angle, constriction of hyphae, formation of septa at a short distance from the point of the hyphal branches' origins and absence of clamp connection and conidia (Ajayl-Oyetunde and Bradley, 2018).

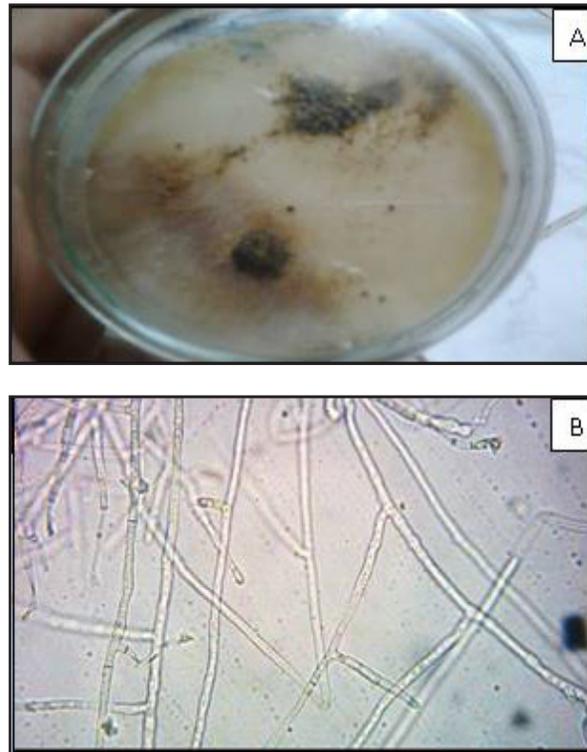


Fig. (1): *R. solani* grown on PDA (A) and under light microscope (B).

Antifungal activity of bacterial isolates

Table (1) represents 84 isolates screened for their ability to suppress the growth of *R. solani*. Results show that the antifungal activity was variable and ranged from 11 to 83%. Three isolates exhibited the highest inhibition properties (78, 82 and 83 %) against *R. solani* were isolated from Beheira and Giza.

Bioproducts from bacterial isolates

As shown in Table (2), all bacterial isolates were negative for chitinase, lipase, amylase, cellulase or volatile antibiotics. However, the extracellular enzyme protease was detected only with isolate B1. Other secondary antifungal compounds e.g. diffusible and volatile antibiotics, siderophores, HCN and IAA were detected by both B1 and B3 isolates.

Table (1) : Screening of antagonistic properties of bacteria isolated from rhizosphere of different Governorates on *R. Solani*.

Governorate (number of isolates)		Fungal radical growth (mm)	%Average of inhibition
Giza (23)	1	15-20	83
	5	<40-55	50-67
	2	60-80	33-44
	14	-	0
Beheira (22)	2	15-20	78-82
	5	.40-55	39-50
	7	60-80	11-33
	8	-	0
Sharkia (21)	-	15-20	-
	5	.40-55	39-55
	12	60-80	11-33
	4	-	0
Menofia (18)	-	15-20	-
	6	<40-55	39-50
	4	60-80	22-28
	8	-	0

Table (2) : Characterization of antifungal compounds produced by bacterial isolates B1, B2 and B3.

Isolates	B1	B2	B3
Chitinase	-	-	-
Protease	+	-	-
Lipase	-	-	-
Amylase	-	-	-
Cellulase	-	-	-
Diffusible antibiotic	+	+	+
Volatile antibiotic	-	-	-
Siderophores	+	-	+
Hydrogen Cyanide	+	-	-
Indole Acetic Acid	+	-	-

+, positive result: -, negative result.

Identification of antagonistic bacterial isolates

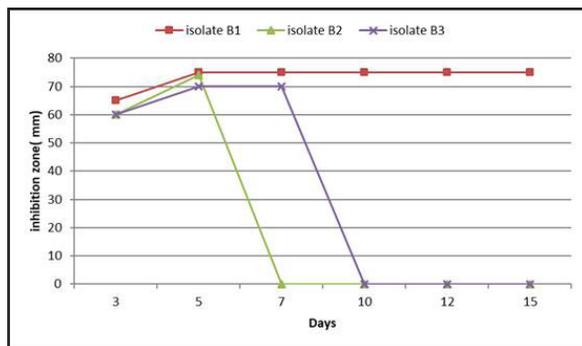
According to Bergey's Manual of Systematic Bacteriology (Sneath *et al*, 1983), the three isolates with the highest inhibition zones belonged to the genus *Bacillus* (Table, 3).

Dual culture of bacterial isolates B1, B2 and B3 and the pathogen

As shown in Fig. (2), the isolate B1 did inhibit the mycelium growth up to 15 days with an inhibition zone of 75mm. On the other hand, the isolate B3 inhibited the fungus growth only until the day 10. No inhibition could be observed for isolate B2

Table (3) : Biochemical and cultural characterization of isolates B1, B2 and B3.

Isolates	B1	B2	B3
Gram staining	Gram positive short rod	Gram positive short rod	Gram positive short rod
Motility	+	+	+
Catalase	+	+	+
Aerobic growth	+	+	+
Anaerobic growth	-	-	-
Acid from:			
Glucose	+	+	+
Sucrose	+	+	+
Fructose	+	+	+
Mannitol	+	+	+
Galactose	+	+	+
Gas production	-	-	-
Hydrolysis of:			
Gelatin	+	+	+
Starch	+	+	-
Indole production	-	-	-
Citrate utilization	+	-	+
Growth in NaCl:			
5%	+	+	+
7%	+	+	+
10%	+	+	+
Oxidase	+	+	+
Nitrate reduction	+	+	-

Fig. (2): Mycelium growth of *R. solani* in dual culture with the three bacterial B1, B2 and B3.

Phylogenetic identification of isolate B1

The 16S rRNA sequence was analyzed by BLAST and shown its identity with 100% sequence

similarity to *Bacillus* sp. Furthermore, the sequence was used to build a phylogenetic tree to form its relationship with its closest identical species using blast tree view and was found to be closely related to *Bacillus* sp. (Fig.3) (Genbank Accession no. MK030136).

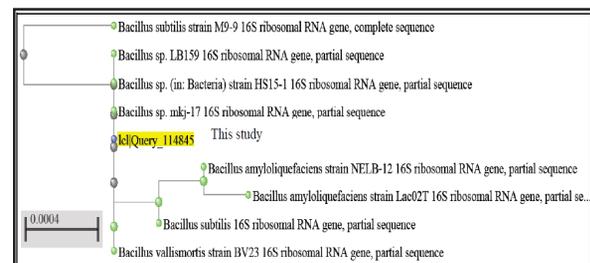


Fig. (3): Phylogenetic identification of the 16S rRNA subunit of the highly effective isolate B1 and its relatedness to other bacilli.

Antagonism of *Bacillus* sp. on hyphae of *R. solani*:

Microscopic examination

The images show that the blue dye did not bind to the fungus after treatment with *Bacillus* sp. (Fig. 4A) as compared to a very deep blue color seen in untreated control samples (Fig. 4B).

Ultrastructural examination

Control sample (Fig. 5 A) shows that the hyphae maintained their normal and intact shape while

in sample treated with bacterial isolate severely changed in the appearance of hypha. The bacterial growth was adhering and colonizing the hyphae at the third day, thus leading to infusion of hyphal tissues. As a result, abnormality of fungal structures was obvious, leading to shrinking and shriveling of hyphae at the 5th day (Fig. 5 B, C). The bacterial colonization over the hyphae resembled a slimy growth and was seen engulfing the hyphae (Fig. 5 D taken on the 7th day).

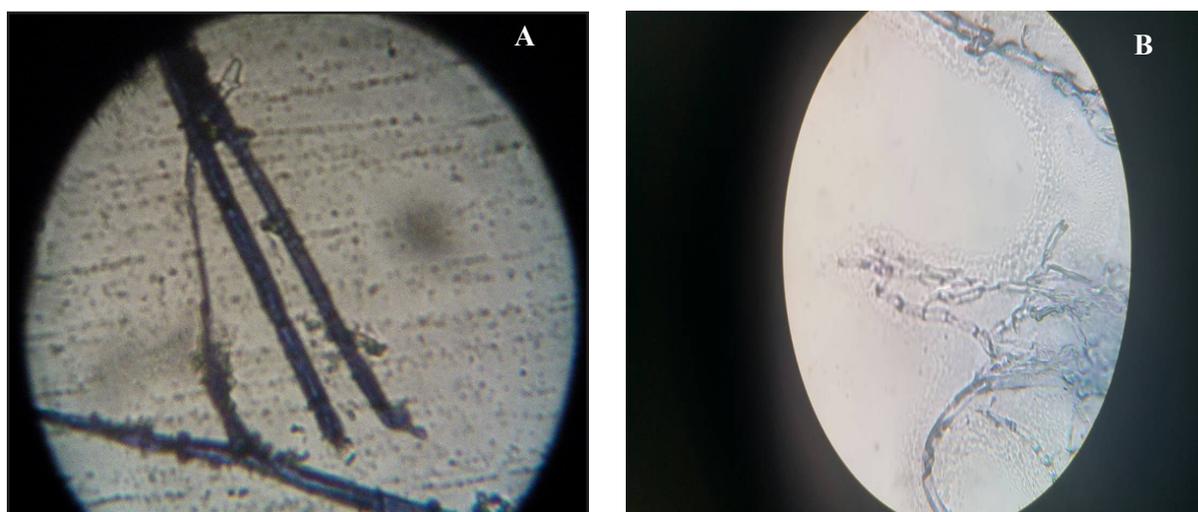


Fig. (4): Light microscope images of *Rhizoctonia solani* treated with *Bacillus* sp. (B) as compared to control samples (A). Both samples were stained with lactophenol cotton blue. Magnification using 40X lens.

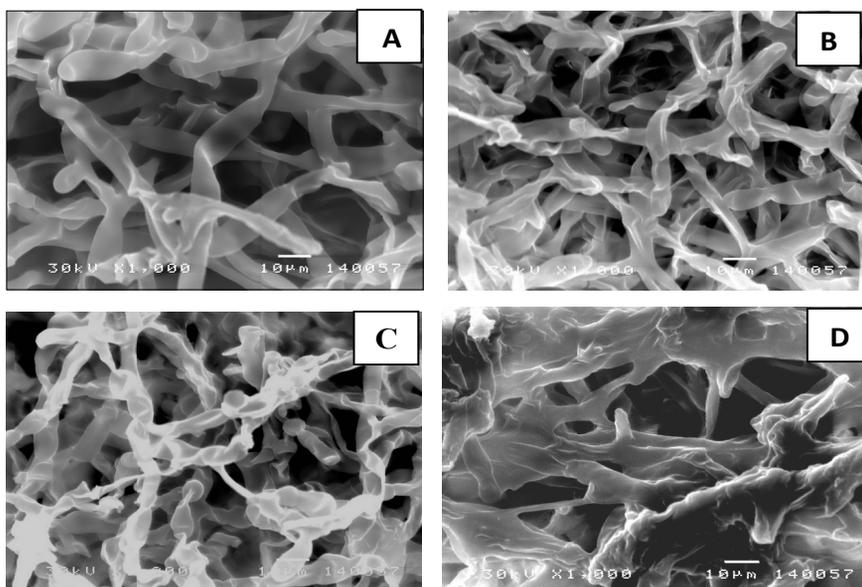


Fig. (5): Scanning electron micrographs of untreated (A) and treated hyphae (B, C and D)..

The TEM analysis was used to estimate the structural characteristics of young hyphae treated with the biocontrol agent (*Bacillus* sp.). Control *R. solani* hyphae produced regular cell walls, with equal widths and distinctive layers, septa with uniform composition as well as dense cytoplasm with good distribution and clearly distinct organelles, such as nucleus and lipid body (Fig. 6 A, B). In contrast, treatment with *Bacillus* sp. caused significant

structural destruction of hyphae, especially in cell walls and plasma membranes, treated cell wall had no distinct layers. Furthermore, plasma membranes were detached from cell walls and septa. Septa had no distinct layers or structures. The damaged cell walls and membranes allowed the cell contents to leak out (Fig. 6 C, D) as compared to the control cells.



Fig. (6): Transmission electron micrographs showing transverse section of untreated hyphae cells (A) and septum (B) as compared to hyphae treated with *Bacillus* sp. (C) and that lacking septum (D).

Physical characteristics of adhesion of *Bacillus* sp. on clay nanoparticles

Bacillus sp. attached on clay nanoparticles were characterized using either SEM, XRD or FT-IR. Fig. (7 A3) shows rod - shaped *Bacillus* sp. aggregates adhered on surface of clay nanoparticles. The control without the bacterium (Fig. 7A1) and bacterial cells

only (Fig. 7 A2) did not show any aggregates. As shown in Fig. (7B), the patterns show a mean peak of the clay nanoparticles at 7.31° , 20° and 27° which are the characteristic peaks of the crystalline bentonite composing the nanoclay under study. While in case of bionanocomposite, the peaks representing clay nanoparticles exhibited change in their intensities.

The FT-IR spectra for the clay nanoparticles, *Bacillus* sp. and bionanocomposite are shown in Fig. (7C). The spectrum of clay nanoparticles represents stretching vibration bands of water molecules adsorbed on clay shifted from 3450 to 3354 cm^{-1} and

from 1633 to 1664 cm^{-1} after bacterial adhesion. The band located at 1658 cm^{-1} , which is assigned to the vibration of amide I, was observed in pure *Bacillus* sp. spectrum. This band was shifted to 1664 cm^{-1} for *Bacillus* sp. nanocomposite.

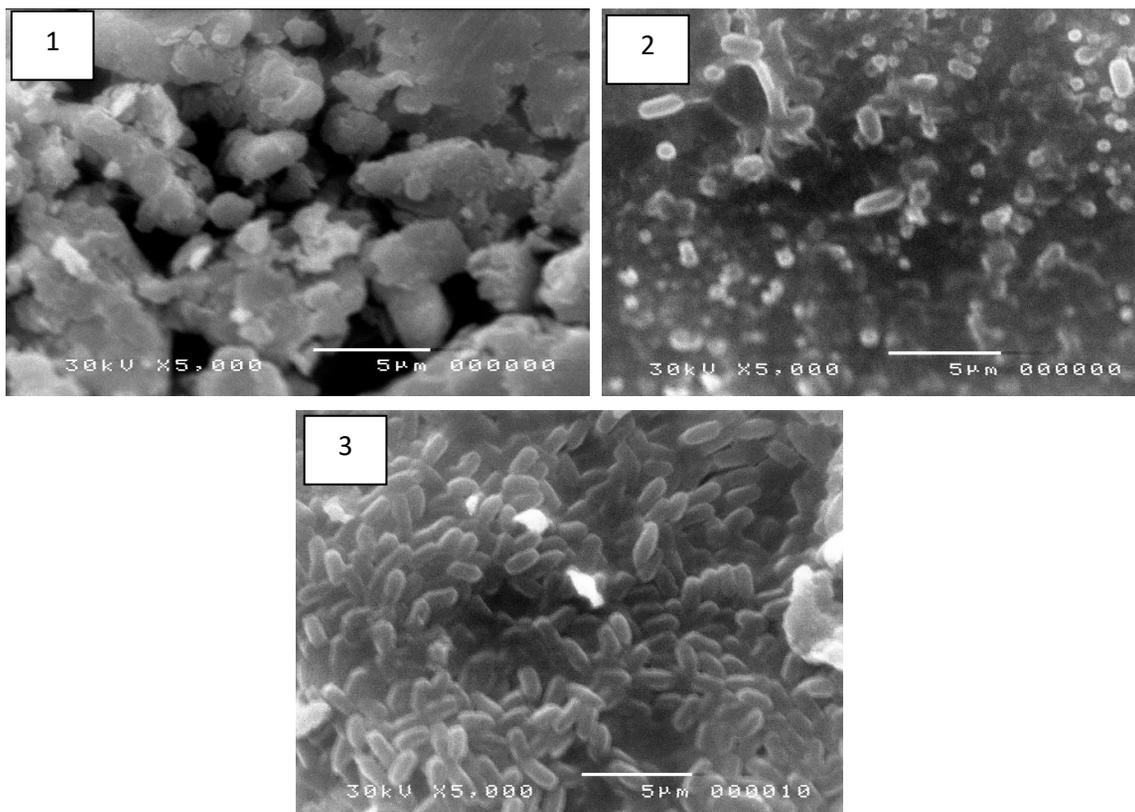


Fig. (7A): SEM images of clay nanoparticles only (1), *Bacillus* sp. cells (2) and *Bacillus* sp. adhered on clay nanoparticles (3).

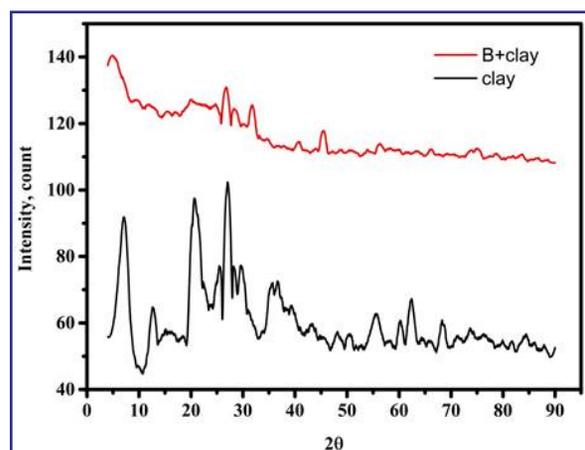


Fig. (7B): X-ray diffraction (XRD) for nanoclay and bionanocomposite.

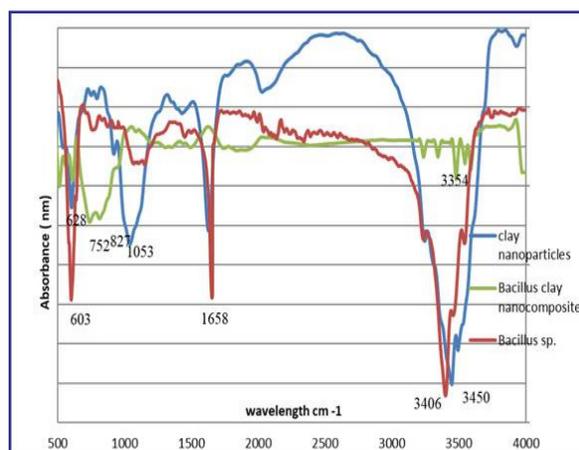


Fig. (7C): FTIR spectroscopy representing nanoclay, *Bacillus* and bionanocomposite of *Bacillus* immobilized on nanoclay.

Shelf life of bionanocomposite and its antagonistic activity against *R. solani*

The viability of *Bacillus* sp. in clay nanoparticles powder (Fig. 8) showed slight decrease in counts among 7 months with significant difference $p < 0.05$ and the lowest viability rate was observed at 8th month (ca. 3×10^5 CFU/ml⁻¹). Bioactivity results of *Bacillus* nanocomposite exhibited high inhibition against *R. solani* (88%) in vitro during 2-month storage period.

Green house studies

The bionanocomposite positively increased the fresh, dry weight and length of both shoots and roots over all dipping periods compared with the Bioark

(positive control). At zero-time dipping, only 22 % disease incidence could be observed using the bionano composite, compared with 66 % with the Bioark (Table 4).

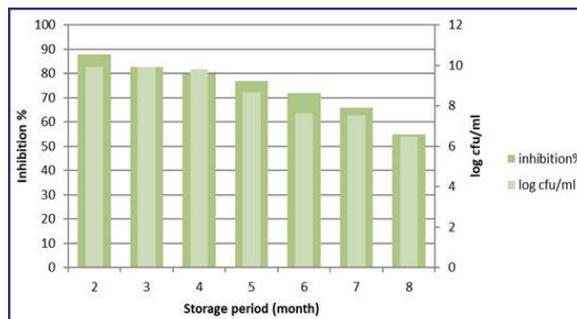


Fig. (8): Viability and bioactivity of bionanocomposite on radial growth rate of *R.solani* during different storage periods.

Table (4) : Effects of various treatments on fresh and dry weights (g), length (cm) of roots and shoots of potato infected by *R. solani*.

Treatments	Immersion time	Root			Shoot			% Disease incidence
		Fresh weight	Dry weight	Length	Fresh weight	Dry weight	Length	
Infested (negative control)		0.4	0.07	3.5	16	1.5	10.3	100
BIOARK (positive control)	zero time	1.23 ^c	0.32 ^g	6.2 ^h	14.87 ^j	2.23 ^g	16.3 ^g	66
	10 min	1.23 ^c	0.41 ^f	7.67 ^g	18.67 ⁱ	2.53 ^g	17.3 ^g	44
	20 min	1.37 ^{de}	0.47 ^f	8.60 ^{fg}	20.03 ^h	3.13 ^f	18.53 ^f	33
	30 min	1.57 ^{bcd}	0.59 ^e	9.03 ^{cf}	20.93 ^g	3.46 ^{ef}	19.13 ^{ef}	11
	60 min	1.77 ^{ab}	0.69 ^d	9.93 ^c	24.43 ^f	4.07 ^d	19.83 ^c	0
Bionano composite	zero time	1.47 ^{cd}	0.59 ^e	12 ^d	28.9 ^c	3.90 ^{de}	29.50 ^d	22
	10 min	1.57 ^{bcd}	0.69 ^d	17.83 ^c	34.07 ^d	5.47 ^c	33.83 ^c	11
	20 min	1.63 ^{bc}	0.9 ^c	19.50 ^b	36.13 ^c	6.23 ^b	36.5 ^b	0
	30 min	1.87 ^a	1.17 ^b	19.83 ^b	37.13 ^b	6.73 ^a	40 ^a	0
	60 min	1.97 ^a	1.39 ^a	21.17 ^a	40.83 ^a	6.90 ^a	39.9 ^a	0

Means with the same letters of the same assessment are not significantly different ($P=0.05$)

DISCUSSION

A total of eighty –four bacterial isolates was isolated from healthy potato plants collected from different areas of Egypt. Antagonism of all isolates was first evaluated against *R. solani* by adopting dual culture assay. On the basis of these tests, three efficient

antagonistic bacteria i.e., B1, B2, B3 were selected and belonged to the genus *Bacillus* and significantly controlled *R. solani*. The inhibition efficiencies recorded were 83, 78 and 82 %, respectively. These inhibition capabilities were apparently higher than those observed by other researches such as **El Ben-**

dary *et al.* (2016) using *B. subtilis* (40%) or *Bacillus* 29 Es (70%). It was reported that the production of protease enzyme and antifungal compounds by *Bacillus* are directly related to fungal growth inhibition because of the ability of *Bacillus* to degrade fungal cell wall and disrupt its internal organelles (Ben Khedher *et al.*, 2015). Only the isolate B1 produced hydrogen cyanide, which was reported to block the cytochrome oxidase pathway and is highly toxic to aerobic microorganisms at low concentrations (Kumar *et al.*, 2012). The Isolates B1 and B3 produced siderophores which are known to chelate most of the available iron thus preventing the proliferation of the pathogen due to lack of the iron (Bharucha *et al.*, 2013). In addition to being the only isolate producing hydrogen cyanide, isolate B1 was the only one that produced indole acetic acid (IAA). IAA is considered important for plant growth promotion, enhancing nutrient uptake and effectively colonizing the rhizosphere (Jadhav *et al.*, 2017). The isolate B1 generated a fungicidal effect in solid culture medium. In agreement with these results, Torres *et al.* (2017) observed that the metabolites produced by *Bacillus amyloliquefaciens* PGPBacCA1 generated a fungicidal effect in solid culture medium. On the other hand, the isolates B2 and B3 only caused a fungistatic effect on this pathogen. A possible explanation can be associated to the low concentration or even the absence of the active antifungal compound in the narrow interface between the fungal mycelium and the bacterial colony. Moreover, the antagonistic effect produced by these strains may also be related to a competition for space or nutrients, instead of a fungicidal effect due to antifungal bacterial metabolites. Phylogenetic analysis revealed that the isolate B1 is closely related to *Bacillus* sp. Light microscope images of cotton blue stained fungus before and after treatment with *Bacillus* sp. showed that the blue dye did not bind to the fungus after its treatment as compared to a very deep blue color seen in untreated control samples. Lactophenol cotton blue is a dye used for staining as well as for wet mounting of fun-

gi, cotton blue imparts blue colouration to the fungal spores and hyphae, its lack of binding indicates the disruption of fungal hyphae, lysis, and abnormalities such as abnormal enlargement of hyphae, terminal swelling, wrinkling and shrinking of hyphae (Parija and Prabhakar, 1995). The effect of *Bacillus* sp. on the morphological structure of fungal hyphae observed by SEM indicate that the mode of action of *Bacillus* sp. against *R. solani* includes two forms of antagonism, e.g. antibiosis and parasitism. Loss of structural integrity of pathogen hyphae due to antibiosis of *Bacillus* sp. was observed. Hyper-parasitism on fungal hyphae was evident through maceration, shrinking, shriveling and abnormal coiling (Kumar *et al.*, 2012). TEM analysis showed that *Bacillus* sp. caused significant structural destruction of hyphae, especially in cell walls and plasma membranes, treated cell wall had no distinct layers. Furthermore, plasma membranes were detached from cell walls and septa. Septa had no distinct layers or structures. The damaged cell walls and membranes allowed the cell contents to leak out. These results agree with the findings of Gong *et al.* (2015) as the clay particles possess a high specific surface area and cationic exchange capacity which combine to create a high adsorption capacity, the attachment of microbial cells on clay particles includes both electrostatic and non-electrostatic mechanisms. In the electrostatic interaction, the cationic surface of clay particles attracts the negatively charged cell walls of Gram-positive bacteria due to presence of proton active functional groups as hydroxyl and amide group (Tourney and Ngwenya, 2014). The non-electrostatic mechanisms mainly include hydrogen bonding, Van der Waals forces and hydrophobic interaction. The attachment of cells onto clay minerals can be influenced by the production of extracellular polymeric substance that increases the attachment of bacteria on clay due to their polymeric interaction. In this regard, clay nanoparticles are efficient support materials and provide a protective habitat for microorganisms by forming biofilm. These results agree

with the findings of **Fathima et al. (2012)**. The peaks in XRD patterns representing clay nanoparticles exhibited change in their intensities. This can be attributed to the coverage of the clay nanoparticles with *Bacillus* sp. These findings are in agreement with **Biswas et al. (2017)** who showed that the insertion of other organic molecules (EPS and other macromolecules produced by the bacterial activity), leads to slight increase of basal space of montmorillonite and a higher production of these compounds might have a significant impact in the interlayer structure. The FT-IR spectra for the clay nanoparticles in the present study represent stretching vibration bands of water molecules adsorbed on clay shifted from 3450 to 3354 cm^{-1} and from 1633 to 1664 cm^{-1} after bacterial adhesion. These results suggest that the water molecules adsorbed on the clay minerals are involved in bacterial adhesion. The shifted frequencies have been assigned to hydrogen bonding interactions between the bacterial cell components and water molecules at clay surfaces (**Tavanaei et al., 2017**). In *Bacillus* sp. nanocomposite spectrum represents conformational or orientation changes of cell surface proteins upon interaction with the fibrous clays. **Tavanaei et al. (2017)** showed the same band of pure *P. putida* spectrums (1658 cm^{-1}) and was shifted to 1657 cm^{-1} and 1648 cm^{-1} for *P. putida*-palygorskite and *P. putida*-sepiolite complexes, respectively. Nano clay as a carrier plays an important role in *Bacillus* formulation to make it applicable under field conditions, elongate the shelf life and commercially appropriate. Also, they have a high aspect ratio that affords more interactive surfaces when exfoliated and dispersed well. Clays have been exploited for immobilization due to their mechanical and thermal stabilities, chemical inertness, well defined layered structures and ion exchange properties (**Ghormade et al., 2011**). Most formulations maintained bacterial viability until 6 months of storage at 22°C when they used talc as the carrier material (**Klein et al 2016**). According to **Ya et al. (2012)**, commercial products should have at least 6 months of shelf life and a pref-

erence to be stored at room temperature. The greenhouse experiment in the present study proved the biocontrol ability of the bionanocomposite against *R. solani*. Indeed, potato tuber (seeds) treatment with bionanocomposite formulation significantly suppressed *R. solani* growth and enhanced plant growth, when compared to the untreated plants. Likewise, **Idris et al. (2009)** reported that *Bacillus* strains significantly enhanced plant growth and inhibited soil-borne pathogen when using an antagonist-treated tuber. On the other hand, coating of tomato seeds with *B. subtilis* RB14-C did not protect tomato plants against *R. solani*. (**Szczzech et al., 2006**). Consequently, the efficacy of seed inoculation with *B. subtilis* was not established for all model plants. Bionanocomposite was able to control stem canker (100% biocontrol efficacy) more efficiently than the commercial biofungicides BioARK at different immersion times. **Tariq et al. (2010)** reported that *Pseudomonas* spp. StT2 and StS3 reduced the proportion of infected tubers by *R. solani* to 40% for cv. Spunta and to 74% for cv. Nicola. In addition to its ability to protect potato plants against *R. solani* more efficaciously than *Pseudomonas* spp., *Bacillus* sp. has the advantage to form endospores that facilitate its production and recovery at a large scale, its formulation and its use in pots and in field treatments. Moreover, *Bacillus* sp. significantly promoted the growth of potato plants. Indeed, bionanocomposite inoculation into potato tuber in the presence of the pathogen increased the height and the weight of stems, and the weight of roots significantly, in comparison to the untreated control. The advancement of plant growth can involve direct and indirect mechanisms. Direct growth promotion is due to bacterial secretion of phytohormones and metabolites that can influence root by overproduction of root hairs and lateral roots and then increase nutrient and water uptake, thus participating to growth. The indirect promotion of plant growth can be due to antibiosis, competition for space and nutrients, parasitism or lysis of pathogen hyphae, inhibition of pathogen-produced en-

zymes or toxins, and through induced systemic resistance. (Ben khedher *et al.*, 2015). Taking into account the harmful effect of *Bacillus* sp. on fungus hyphae. This suggests that the promotion of plant growth by bionanocomposite may be mainly associated with direct and indirect effect, also clay nanoparticles are ideal for maintaining survival and viability of *Bacillus* sp. and rapid adhesion with colonization of clay nanoparticles.

CONCLUSION

An important perspective of the present study is the isolation of *Bacillus* sp. that has a strong antifungal activity against *Rhizoctonia solani*. This bacterium produces an array of metabolites, each playing an important role in the antagonism process ensuring a highly effective and prolonged control as compared to other available bacteria. This bacterium and its metabolites also provide an attractive option for sustainable agriculture due to their ability to stimulate plant growth and production. Preparing the biocontrol agent as a nanocomposite formulation increases its efficiency and longevity, thus providing a safe and effective alternative to traditional commercial formulations.

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