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Article

Biological Control of *Pythium aphanidermatum*, the Causal Agent of Tomato Root Rot by Two *Streptomyces* Root Symbionts

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Abstract: Damping-off caused by *Pythium aphanidermatum*, as one of the most infamous plant diseases, is considered as a significant disease that causes severe damages in greenhouse and field crops in a vast range of hosts especially vegetables. The application of chemical fungicides as a common method to control this disease poses negative side effects on humans and the environmental components. Actinobacteria, especially members of the genus *Streptomyces*, are fascinating biocontrol agents and plant growth-promoting rhizobacteria, which exhibit safer alternative managements to control this disease. The present study aims to explore for bioactives soil *Streptomyces* isolates that are able to control *P. aphanidermatum*, which is the causal agent of damping-off in tomato. Out of a total of 116 actinobacteria isolates collected from the soil, 53 have showed an antagonistic activity against *P. aphanidermatum*, as deduced through in vitro dual cultures. Based on in vitro Petri plate seedling-bioassays (IPSB), from the 53 tested isolates in dual cultures, two isolates coded as H2 and H3, considered as the most bioactive agents, were selected to assess their biocontrol performances against *P. aphanidermatum* in the Sandwich bed-mix technique in greenhouse experiments. In vivo greenhouse statistical studies were performed to compare seven treatments using completely randomized design experiments. Metalaxyl fungicide was applied as the chemical-control treatment. To evaluate biocontrol efficiencies of the two *Streptomyces* symbionts, disease incidence recorded throughout the course of experiment and criteria of number of leaves, length of the plants, plant fresh and dry weights, were measured at the end of experiment and analyzed statistically. The resulted disease incidences for all treatments indicated that the two *Streptomyces* strains had PGPR activity, and they were effective in decreasing disease incidence and improving plant performances regarding number of leaves, height, and plant fresh and dry weights. Based on the phylogenetic analysis of the partial sequences of the 16S rRNA gene, the strain H2 revealed a close relation to six *Streptomyces* species, namely *S. badius*, *S. caviscabies*, *S. globisporus*, *S. parvus*, *S. sindenensis*, and *S. griseoplanus*, with 99.9% similarity. The strain H3 also indicated a close relation of the same similarity to two species, namely *S. flavogriseus* and *S. pratensis*. In overall, collected data dictated that the two bioactive root symbiont *Streptomyces* strains effectively controlled the damping-off disease caused by *P. aphanidermatum*.

Keywords: *Streptomyces*; *Pythium aphanidermatum*; biocontrol; root symbionts; PGPR; 16S rRNA; tomato; damping-off; antagonist

1. Introduction

Tomato (*Lycopersicon esculentum* Mill, Solanaceae) is one of the most important vegetables in many countries with China, India, USA, Turkey, Egypt, Iran, Italy, Spain, Mexico,

and Brazil as the top ten producers of tomato in the world. According to FAO recent reports, in 2018, Iran ranks sixth in the world with the production of over six million tons of tomatoes per year [1]. Many of the soil-borne pathogens threaten this crop and reduce its production efficiency. Damping-off is as one of the devastating soil-borne diseases that causes serious problems in tomato seedlings [2]. *Pythium* spp., *Fusarium* spp., *Rhizoctonia* spp., and *Phytophthora* spp. are as the most frequently plant pathogens associated with damping-off [3]. The *Pythium* spp. that are ubiquitous parasites in soil and water are considered as the most notorious plant pathogens in vegetables especially in the nursery, greenhouse, and field [4,5]. The most common species of *Pythium* in greenhouses include *P. aphanidermatum* (Edson) Fitzpatrick, *P. ultimum*, *P. irregulare*, *P. myriotylum*, *P. spinosum*, and *P. splendens* [6]. *P. aphanidermatum*, the most destructive species in the genus, is a frequent agent of damping-off and root rot of tomato in nurseries and greenhouses [7]. It causes rotting and imposes economic damages on many monocotyledon and dicotyledon plants such as members of the *Solanaceae* family [4]. *Pythium aphanidermatum* is also the causal agent of seed rot, seedling rot, root rot, cottony leak, stem, and rhizome rots as well as pre- and post-emergence disorders in many other economic crops [8,9]. The pre-emergence damping-off symptoms have been associated with damage to seeds or seedlings before their emergence from the soil. Throughout the onset of pre-emergent damping-off rots, infection of the germinating seeds may result in seeds rot or produce water-soaked lesions on the roots and stems of emerging seedlings, ultimately leading to seedling death. In post-emergence, the damping-off occurs after young seedling emergence from the soil. At this stage, the stem becomes thin with brown lesions, the infected crown rots, the hypocotyl becomes water-soaked, the cotyledon leaves may roll downward, and the roots of infected plant rot; then, consequently, the plant dies [10,11]. Oospores, sporangia, and mycelia are surviving forms of *P. aphanidermatum* in the organic debris and soil [5,12,13]. To control this pathogen, several agricultural practices have been applied such as crop rotation, soil-solarization, and the use of chemical fungicides; however, none of them exhibit practicable and sustainable results in highly infected soils. Given the wide host range of this devastating pathogen and its prolonged survival in the soil, it is quite difficult to control the disease [7]. The fungicide metalaxyl has been shown to be very effective to manage the damping-off caused by Oomycetes and to lower the disease incidence. Nevertheless, several reports demonstrated metalaxyl, metalaxyl-M, and Mefenoxam resistance in *Pythium* spp. [14–16]. However, the use of chemical fungicides poses harmful side effects on humans and the environment [17,18]. Awareness of the short and long- term adverse effects of chemical pesticides has led to use of less calamitous approaches and alternative applications such as biological methods. Looking to forefront horizons, the biological methods are being seen as promising and sustainable methods to manage plants diseases. Under the light of such a view and the demand of the general public in recent decades, the biological control agents (BCAs) have been vastly employed to replace chemical agents for the control of plant pathogens. In this regard, several fungal and bacterial agents including *Trichoderma* [19,20], non-pathogenic *Fusarium* [21], *Xanthomonas*, *Serratia* [22], *Bacillus* [23,24], *Pseudomonas* [25], and *Streptomyces* [26–29] have been reported as effective biological control agents. Furthermore, there are several reports on biological control agents of *P. aphanidermatum* in tomato specifically include *Bacillus subtilis* and *Trichoderma* spp. [20,30], *Aspergillus terreus* [31], *Klebsiella oxytoca*, *Exiguobacterium indicum* and *Bacillus cereus* [32], *Talaromyces variabilis* [33], and *Streptomyces* spp. isolate KUMB1.1 [34]. Some of these microorganisms also act as plant growth-promoting fungi (PGPF) or plant growth-promoting rhizobacteria (PGPR). For the first time, Kloepper and Schroth defined PGPRs as plant bacteria that are able to colonize plant roots and improve plant growth [35]. Among PGPRs, *Streptomyces* is the largest genus of phylum Actinobacteria and considered as the type genus of the family Streptomycetaceae with more than 500 species [36,37]. It has attracted a lot of attention regarding their ability to produce antibiotics and biologically active secondary metabolites, which make them the most bioactive microorganisms [38–40]. PGPRs can be applied effectively in the management of plant pathogenic fungi and bacteria [27]. *Streptomyces*

species are saprophytic bacteria, with filamentous nature, Gram-positives, and high GC content in the genome (about 70%) that can also play a role as symbiont bacteria with plants [41,42]. *Streptomyces* spp. are known as biological potential resources for producing more than 75% of bioactive substances and secondary metabolites including more than 80% of the world antibiotics [43–46]. A range of beneficial microorganisms, especially *Streptomyces*, can colonize plant roots in symbiotic interaction. Indeed, colonization is an important behavior in biocontrol interactions, and it can enable symbiont inoculants to compete with indigenous and deleterious microorganisms by competitive exclusion, because they can occupy ecological niches and use nutrients and space resources [47,48]. Loliama et al. [49] proved that *Streptomyces* species could be considered as effective candidates for the biocontrol of damping-off caused by *P. aphanidermatum*. In this regard, *S. rubrolavendulae* S4 was able to control *P. aphanidermatum* in *Amaranthus tricolor* seedlings as effectively as the fungicide metalaxyl. In another study, two endophytic Actinobacteria isolated from healthy maize plants, *Streptomyces* sp. 16R3B and *Streptomyces* sp. 14F1D/2, were able to reduce damping-off in cucumber under greenhouse conditions caused by *P. aphanidermatum* up to 71% and 36%, respectively [50,51]. Consequently, in this research, we focused on in vitro screening and in vivo biocontrol of tomato root rot caused by *P. aphanidermatum* using potent *Streptomyces* spp. Objectives of this research were (1) to isolate biologically efficient *Streptomyces* species from soil and evaluation of their biological control performance against *P. aphanidermatum* in tomato plants; (2) to evaluate the tomato plant growth-promotion properties resulting from bacterial interaction; and (3) characterize and identify molecularly the bioactive *Streptomyces* isolates based on 16S rRNA sequence analyses.

2. Materials and Methods

2.1. Culture Media and Preparation of the Pathogen

Pure culture of *P. aphanidermatum* (accession no. MV599252) obtained from Mycology Laboratory, Department of Plant Protection, Faculty of Agriculture, University of Jiroft, Iran. The pathogen was cultured on potato dextrose agar (PDA, Merck, Darmstadt, Germany) and kept at ambient temperature before use.

2.2. Pathogenicity Assessment

To be assured of the pathogenicity of *P. aphanidermatum* on tomato seedlings (*Lycopersicon esculentum* cv. Moneymaker), damping-off incidence was assessed on tomato seedlings as described by Abreo et al. [52]. In this regard, tomato seeds treated with 2% sodium hypochlorite solution for one min, washed in three series of sterilized water for two min, then placed on 2% water agar (WA) Petri plates and kept in dark to germinate at ambient temperature. Four days later, a mycelial disc of six-millimeter diameter was cut with a sterile cork-borer and placed at the center of each Petri plate at equal distances from the seeds and incubated for 2 to 4 days at 25 °C ($\pm$2 °C). Control Petri plates received blank agar disks in the same manner.

2.3. Soil Samples and Isolating Actinobacteria

Twenty samples of rhizospheric soil were collected from different healthy-looking tomato fields located in different regions of Kerman province, Iran. The soil samples were dried in the shade for 24 h, and bulk soil was removed by sieve shaking. One g of each soil sample was mixed with sterile distilled water at 1:10 ratio (w/v), shaken vigorously for 10 min, settled for 10 min, and 1 mL aliquot of the suspension diluted by sterilized distilled water by 10-fold series up to 10^{-6} dilution. To culture, aliquots of 1 mL of each of 10^{-3}, 10^{-4}, 10^{-5}, and 10^{-6} dilutions mixed with aliquots of 20 mL of relatively cooled (48–49 °C) autoclaved Casein Glycerol Agar (CGA) medium, which was prepared from basic ingredients as described by Küster and Williams [53], in separate 12 cm Petri plates and let to solidify [54]. Two replicates were considered for each dilution. Plates were incubated at 28 °C for a week. When *Streptomyces* colonies became visible, the aseptically

streaking method [55] was used to get pure cultures of *Streptomyces* isolates on CGA slants. The pure Actinobacteria isolates were incubated at 28 °C for 7 days and then refrigerated at 4 °C before use [56].

2.4. In Vitro Screening of Actinobacteria

To screen for anti-*Pythium* Actinobacteria, dual culture bioassays conducted as described by El-Tarabily et al. [57]. Briefly, plugs of 6-day-old actinobacterial isolates and one uninoculated medium plug (as control) cut with a sterile cork borer (diameter of 6 mm) were placed at 2 cm distances from each other on each PDA Petri plate using a sterile forceps. After 48 h at 28 °C, a six mm plug from 3-day-old *P. aphanidermatum* was placed in the center of each PDA Petri plate and incubated at 28 °C for 3 days until the *P. aphanidermatum* colony margin of control reached close to the edge of the plate. The anti-*Pythium* activity of Actinobacteria isolates was determined as follows. Insertion of mean of growth radius of *P. aphanidermatum* in control (r_1) and mean of growth radius of *P. aphanidermatum* in dual culture bioassay (r_2) in the following formula: $\Delta r = r_1 - r_2$. The modified ratings were applied from El-Tarabily et al. [57]: where $\Delta r > 2$ and <4 mm, it was represented as +; where $\Delta r > 4$ and <6 mm, it was represented as ++ and where $\Delta r > 6$ mm, it was represented as +++.

2.5. In Vitro Petri Plate Seedling-Bioassay

In vitro Petri plate seedling-bioassay (IPSB) is a bioassay technique we developed in our lab. To evaluate the potential ability of isolated actinobacteria to inhibit or suppress the pathogenicity of *P. aphanidermatum* in tomato seedlings, the isolated actinobacteria, showing promising results from in vitro dual culture bioassays, were meticulously evaluated in this experiment. In this regard, seeds surface were sterilized (two min in 2% sodium hypochlorite and subsequently 5 min rinse with sterilized distilled water) and placed on well-grown 6-day-old cultures of actinobacteria solid media and shaken until seeds were fully coated with spores. Coated seeds were placed near the margin of 2% WA Petri plates and then received a 6 mm plug of 3-day-old *P. aphanidermatum* culture at the center. To set controls, uncoated seeds and the pathogen were considered as positive controls and uncoated seeds with no pathogen, but a blank agar plug, were considered as negative controls. After a week on a lab bench at ambient temperature, seed rots and seedlings viability in the three treatments were recorded for the tested actinobacteria. All experiments were performed in triplicate, and mean values were recorded.

2.6. In Vitro Root Colonization Bioassay

To evaluate the ability of the active actinobacteria isolates for the colonization of tomato roots, the following steps were performed consecutively. The tomato seeds surface was sterilized (2 min 2% sodium hypochlorite, 5 min successive rinses with sterilized distilled water). The spore suspension of each actinobacteria isolate was prepared at concentrations of 10^8 CFU mL^{-1} by gentle scraping off the spores of the surfaces of 7-day-old cultures using sterile scalpels. Using a NanoDrop™ spectrophotometer (Thermo Scientific™ 840274200), the final concentrations of spore suspensions were adjusted to OD: 1 at 660 nm. For each isolate, the sterilized tomato seeds were immersed in the spore suspension and shaken for 30 min. Subsequently, the inoculated seeds were placed on 2% WA Petri plates and incubated at ambient temperature (25 ± 2 °C) until germinated. When primary roots length reached about 4 cm, approximately 4-6 mm of elongation zones of roots of each treatment was cut with a sterile scalpel, aseptically placed on CGA medium, and then incubated at 28 °C for 3–4 days in the dark at ambient temperature. The growth of actinobacteria at this experiment was considered as positive, and the lack of growth was negative root colonization. Positively root-colonized actinobacteria were considered as tomato root symbionts. All experiments were performed in triplicates and plates were inspected for the growth of actinobacteria daily.

2.7. DNA Extraction, PCR Amplification, and Electrophoresis

The *Streptomyces* strains (H2 and H3) were cultured on the CGA medium and incubated at 28 °C for a week. Then, aerial mycelia were aseptically scraped off by sterile scalpels, and DNA was extracted by using the CTAB method [58]. For 16S rRNA gene amplification, the universal primers F1 (5′-AGAGTTTGATCITGGCTCAG-3′) and R5 (5′-ACGGITACCTTGTTACGACTT-3′; I = inosine) were used [59]. PCR was carried out in a thermal cycler (Biometra, T-Personal Thermal Cycler) with final volumes of 15 µL containing 7.5 µL 2X PCR master mix (Parstous, Iran), 0.5 µM of each of the primers, and 50 ng DNA template of each strain. The thermal conditions for amplification were: initial denaturation at 96 °C for 2 min, followed by 30 cycles of denaturation at 96 °C for 45 s, annealing at 56 °C for 30 s, and extension at 72 °C for 2 min, and a final extension at 72 °C for 10 min. Lastly, PCR products were electrophoresed on a 1% agarose gel containing 10 µg/mL ethidium bromide to validate the size of amplified fragments compared with the DNA size marker (ExcelBand™ 100 bp+3K DNA Ladder, Smobio, Taiwan).

2.8. DNA Sequencing and Phylogenetic Analysis

PCR products were directly sequenced by Microsynth (Balgach, Switzerland) using the primers F1 and R5. The resulting forward and reverse sequences of the 16S rRNA gene for each strain were assembled into a final consensus sequence using the CAP3 program [60]. The sequences were submitted to GenBank with the accession numbers of MW555322 and MW555323, with lengths of 1346 and 1393 bp, respectively. Then, the sequences were subjected to BLAST (Basic Local Alignment Search Tool) analysis at NCBI (https://blast.ncbi.nlm.nih.gov/Blast.cgi), to compare with those from known *Streptomyces* strains. Then, closely related sequences were retrieved from GenBank and aligned using the ClustalW program included in the MEGA software version 7.0.26. Finally, a phylogenetic tree was constructed using the neighbor-joining statistical method [61] with 1000 bootstrap replicates to confirm inferred clades. The tree was rooted using the 16S rRNA gene sequence from *Mycobacterium helveticum* (GenBank no. MT133249) as an out-group. Moreover, matrix similarity of the sequences (data not shown) was generated by the MatGat software, version 2.0 [62].

2.9. Physiological Characterization

For two active symbiont actinobacteria, evaluation of several enzymatic activities and tolerance to different concentrations of NaCl (0%, 1%, 2.5%, 5%, 10%) performed as described by Palaniyandi et al. [63]. Enzymatic bioassays included productions of gelatinase, catalase [64], citrate [65], and amylase [66]. The production of hydrogen cyanide (HCN) and volatile organic compounds (VOC) [67,68] were also assessed. The vulnerability of their activity to chloroform was performed as described by Jorjandi et al. [69].

2.10. Greenhouse Evaluations

2.10.1. Preparation of Tomato Seed

To conduct greenhouse trials, tomato seeds of Moneymaker cultivar were sterilized (2 min 2% sodium hypochlorite, 5 min successive rinses with sterilized distilled water). Next, the seeds were placed in capped Petri plates bearing Whatman filter paper No. 1 wetted with sterilized water, incubated at ambient temperature until seeds germinated, and then used for greenhouse experiments.

2.10.2. Preparation of *P. aphanidermatum* Inocula

To prepare *P. aphanidermatum* inocula, 100 g of millet seeds were added to 50 mL of distilled water in 500 mL Erlenmeyer flasks. The flasks autoclaved at 121 °C for 40 min on three successive days to sterilize the millet seeds. Afterwards, the inoculation of seeds performed by adding eight plugs from a 3-day-old culture of *P. aphanidermatum* to each flask and incubated at 25 °C (±2 °C) for 4 days. In this period, to achieve uniform colonization, the flasks were shaken at least once a day [70].

2.10.3. Soil Inoculation

Two symbiont actinobacteria isolates (H2 and H3) that showed the highest bioactivity against *P. aphanidermatum* in IPSB were cultured on CGA solid medium in 12 cm Petri plates and incubated at 28 °C. A week later, spore suspension of actinobacteria was prepared aseptically by scarping off the surface of cultured plates in small aliquots of sterilized distilled water using a sterile scalpel. The concentration was adjusted at 10^8 CFU mL^{-1} using a spectrophotometer (NanoDrop™, Thermo Scientific™ 840274200). Then, the suspension of each actinobacteria isolate was mixed with sterilized soil at the ratio of 10 mL kg^{-1} soil. Inoculated soil was incubated in sterile dark plastic bags at 28 °C for three days and shaken twice daily to ensure thorough distribution and homogeneity of actinobacteria in the soil. For control treatment, only sterile distilled water was added to sterilized soil at a ratio of 10 mL kg^{-1} [32].

2.10.4. Design of Sandwich Bed-Mixes

The greenhouse trials were conducted from early June to early Sept 2020. Plastic pots with 20 cm diameter and 15 cm height were used in greenhouse experiments. We designed a Sandwich bed-mix used for plant beds. As schematically indicated in Figure 1, in this method, height wise, the pots were divided into 4 sections (A, B, C, and D, each section was about 3.5 cm). From bottom-up, section D was filled with sterilized soil (no pathogen and no symbiont Streptomyces), section C was filled with soil mixed with inoculum millet seeds (0.5% *w/w*), section B was filled with soil inoculated with one of symbiont *Streptomyces* (H2 or H3), and section A, the top section, was filled with sterilized soil. Tomato seeds (cv. Moneymaker) were evenly sowed in section A. Statistically, greenhouse experiments were performed as a complete randomized design with seven treatments as follows: (1) negative control: all pot sections contained sterilized soil, (2, 3, 4) symbiont *Streptomyces* isolates: layer B contained either one of H2 or H3 and layer C contained only sterilized soil, (5) positive control: layer C contained inoculum millet seeds and layer B contained only sterilized soil, (6, 7, 8) symbiont *Streptomyces* isolates + pathogen: layer B contained either one of H2 or H3 and layer C contained inoculum millet seeds, (9) fungicide Metalaxyl + pathogen: layer C contained inoculum millet seeds, and layer B contained fungicide Metalaxyl (0.1%) [32]. Each treatment was replicated thrice and mean values recorded for the triplicates. The pots kept in a greenhouse with a 16/8 h (light/dark) photoperiod and approximately 60% humidity (Table 1). No fertilizer was used throughout the experiment.
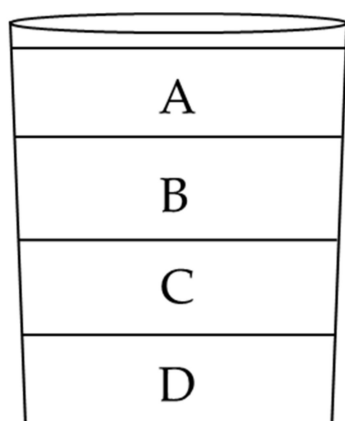
Figure 1. Schematic representation of Sandwich bed-mix used in greenhouse experiments. The pot bears four striated layers of sterilized soils which each receive specific treatment, e.g., in a challenge treatment having both the antagonist and the pathogen, the layers contain the following: (**A**) top sterilized soil, (**B**) sterilized soil plus the antagonist, (**C**) sterilized soil plus the pathogen, and (**D**) bottom sterilized soil.

Table 1. Sandwich bed-mix treatments applied in greenhouse for plant beds to evaluate anti-*Pythium aphanidermatum* bioactivity of two *Streptomyces* symbionts on tomato seedlings (cv. Moneymaker).

Treatment No.	Treatment	Treatment Description
1	Positive control	Millet seed *Pythium aphanidermatum* inoculum (0.5% w/w)
2	*P. aphanidermatum* + H2	10 mL 10^8 CFU spore suspension of H2 + Millet seed *Pythium aphanidermatum* inoculum (0.5% w/w)
3	*P. aphanidermatum* + H3	10 mL 10^8 CFU spore suspension of H3 + Millet seed *Pythium aphanidermatum* inoculum (0.5% w/w)
4	*P. aphanidermatum* + Metalaxyl (chemical fungicides)	Millet seed *Pythium aphanidermatum* inoculum (0.5% w/w) + Metalaxyl (1%)
5	Symbiont *Streptomyces* isolate H2	10 mL 10^8 CFU spore suspension of H2 kg^{-1} sterilized soil
6	Symbiont *Streptomyces* isolate H3	10 mL 10^8 CFU spore suspension of H3 kg^{-1} sterilized soil
7	Negative control	10 mL sterile water kg^{-1} sterilized soil

2.10.5. Evaluation of Plants Responses

After three months of greenhouse trials, the plant were uprooted by gentle washes with tap water. To assess the performance of the applied symbiont *Streptomyces* isolates, several parameters were measured and mean values of triplicates were calculated. Evaluated parameters are indicated as follows.

Disease Incidence

In the Sandwich bed-mix experiments, from the beginning and throughout the experiment, the number of germinated and perished seeds and seedlings was recorded. Compared to the number of plants in negative control, the percentage of disease incidences was calculated in triplicate treatments and used in statistical assessments.

The disease incidences was calculated using the following formula [71].

$$Disease\ incidence\ (\%) = 100 \times \sum\nolimits_{i=0}^{n} X / \sum\nolimits_{i=0}^{n} Y \tag{1}$$

where X = number of dead tomato plants, Y = total number of tomato plants in each treatment, n = rating scale (0–4).

Number of Leaves

At the end of the experiment, before uprooting, the number of leaves per plant was counted in triplicate, and the mean values were recorded.

Length of the Plants

Plant length including the length of stem from apex plus that of tap root to the tip. The plant length was monitored in triplicates in cm and mean values recorded.

Plant Fresh Weights

Plant weights comprised the total weight of plants, including both above and underground parts. After digging up the plants and gently washing the root, excess moisture was removed thoroughly by tissue papers. The total fresh weight of plants was recorded in triplicate treatments.

Plant Dry Weights

Following recording the total fresh weights of plants, specimens were placed in labeled pre-weighed paper bags and dried in 60 °C oven for 48 h. After drying, bags were weighed

accurately, and the subtraction of first and second weights was considered as plant dry weights for the triplicate treatments.

2.11. Statistical Analyses

Statistical analyses of mean values of greenhouse triplicate treatments were performed for analysis of variances (ANOVA) of completely randomized design experiment. The means were compared by Duncan's Multiple Range Test. SAS software v9.1 (SAS Institute Inc., Cary, NC 27513–2414, CA, USA) was applied to analyze data in the significant differences at $p \leq 0.05$.

3. Results

3.1. Pathogenicity Test and Damping-Off Incidence

In pathogenicity assessment experiment, infected seedling roots became necrotic, which was indicative of a pathological reaction. The pathogen–seedling interaction revealed that *P. aphanidermatum* was pathogenic on *Lycopersicon esculentum* cv. Moneymaker. The symptoms included firstly a discoloration of root and hypocotyl; then, the root and hypocotyl became water soaked and rotten (Figure S1, Supplementary Materials). In this bioassay evaluation, the disease incidence was observed in eight of nine tested seedlings.

3.2. Isolated Soil Actinobacteria and In Vitro Screening

By macroscopic and microscopic evaluations of culture plates, 160 actinobacteria pure isolates collected from soil cultures and stored at 4 °C as slants for further studies. The anti-*Pythium* activity of the 160 actinobacteria was isolated from soil conducted by dual culture bioassays and evaluated for inhibition bioactivity. From 160 actinobacteria isolates, 53 isolates exhibited moderate or strong inhibition activity against *P. aphanidermatum*. These active isolates selected as described by El-Tarabily et al. [57]. Both H2 and H3 isolates that showed a high antagonistic effect against *P. aphanidermatum* when tested on tomato seedlings (Figure 2).
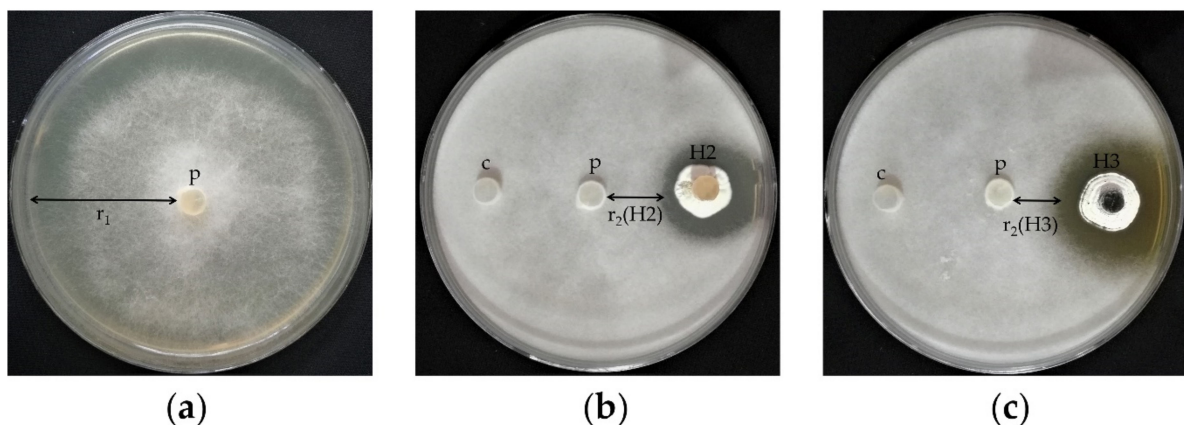

(a)   **(b)**   **(c)**

Figure 2. In vitro dual culture bioassay to select the bioactive actinomycetes isolates against *Pythium aphanidermatum*, quantitatively. (**a**) Arrow (r1) indicates the pathogen colony radius in control plate (7 cm diameter) in comparison with plate; (**b**) and (**c**) indicate the inhibitory effect on radii of *P. aphanidermatum* against actinomycetes isolates H2 and H3 (r2 (H2) and r2 (H3)). The blank agar plug (**c**) in each bioassay plate considered as control. Differential anti-*Pythium* activity was calculated from formula given in Section 2.4.

3.3. In Vitro Petri Plate Seedling Bioassay

The results of in vitro Petri plate seedling bioassay (IPSB) are presented in Figures 3 and 4. To shorten this section, the results in Figure 3 are indicted for the actinobacterium H2 isolate only. In less than 10 days, in Petri plates, the test well indicated the potential ability of active actinobacteria for root symbiosis and suppression of the *P. aphanidermatum* in tomato seedlings. Statistically, the protection (%viability) of germinating tomato seeds

coated with each of the actinobacteria was represented as follows: 9% for positive control (treated with *P. aphanidermatum* but no symbionts); 100% for negative control (untreated with neither pathogen nor symbiont); 73% and 64% for seeds coated with *P. aphanidermatum* and either of symbiont H2 or H3, respectively. As indicated in Figure 3, IPSB is relatively a fast bioassay and may be considered as a semi in vivo evaluation experiment. The symbiont actinobacteria exert their bioactivity via antibiosis and competitive exclusion [72]. In this bioassay, all 53 actinobacteria isolates, which were concluded as active in dual culture bioassays, were screened through IPSB among which the two isolates of H2 and H3 were selected for further evaluations including biochemical bioassays, molecular identifications, and in vivo greenhouse trials. According to El-Tarabily et al. [57], Δr values included 2.0 for H2 and 2.1 for H3, which coincide with IPSB results.
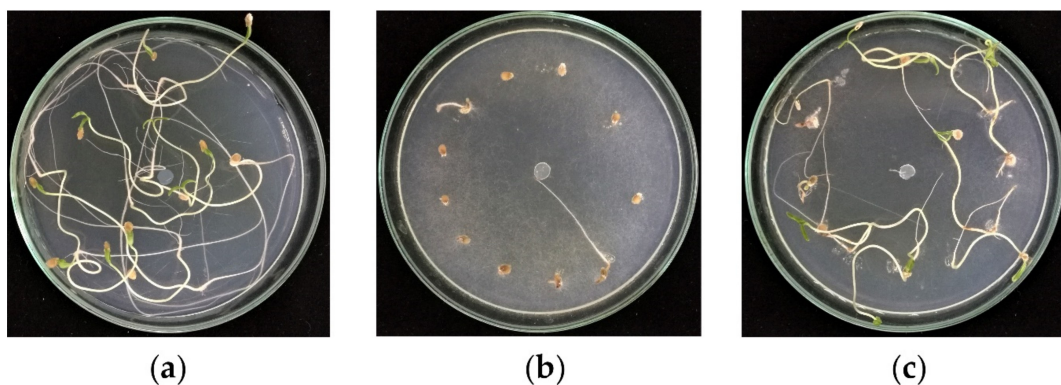

(**a**)　　　　(**b**)　　　　(**c**)

Figure 3. Evaluation of biocontrol activity of symbiont H2 actinobacterium isolate against *Pythium aphanidermatum* performed by tomato seed coating via in vitro Petri plate seedling bioassay (IPSB): (**a**) germinating untreated seeds (negative control); (**b**) germinating uncoated seeds placed from a 6 mm plug of *P. aphanidermatum* (positive control); (**c**) germinating coated seeds with spores of active actinobacteria isolates placed at equal distances of 2 cm vicinity of a 6 mm plug of *P. aphanidermatum* (challenge plate).

Figure 4. In vitro evaluation result of antagonistic activity of two symbiont actinobacteria isolates of H2 and H3 against pathogen *Pythium aphanidermatum* in tomato seedlings performed by tomato seed-coating via in vitro Petri plate seedling-bioassay (IPSB). Each picture is representative of its related treatment. −S+P: no symbiont actinobacteria + pathogen, the positive control (PC); +S+P: symbiont actinobacteria (H2 or H3) + pathogen and −S−P: no symbiont actinobacteria + no pathogen, the negative control (NC).

3.4. In Vitro Root Colonization Bioassay

The formation and growth of colonies on CGA medium indicated that both actinobacteria isolates, H2 and H3, colonized the tomato seedling roots (Figure 5). Apparently, spores of seed-coated actinobacteria germinate and utilize exudates of spermosphere and subsequently propagate on exudates of root surface; hence, they would act as symbionts in a symbiotic relationship where both plant and participating actinobacteria benefit from each other, which is a relationship referred to as mutualism. This process often leads to fully colonized root system. Root colonization is an advantage of such symbionts in their biocontrol behavior [73].
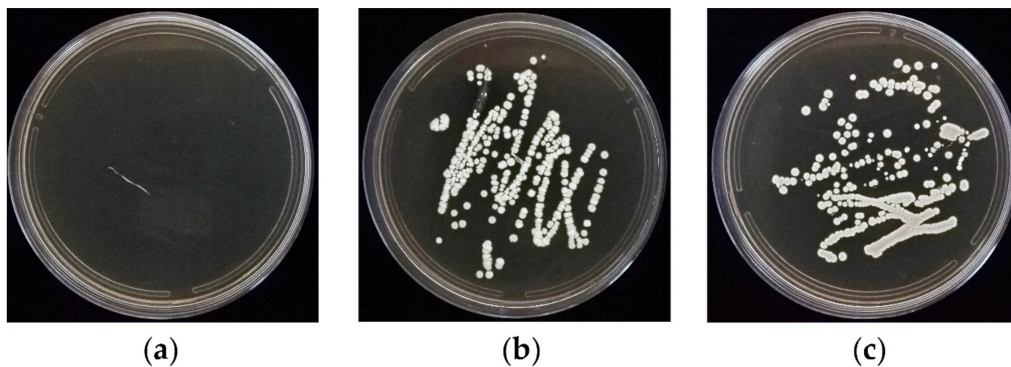

(**a**) (**b**) (**c**)

Figure 5. Result of root colonization of tomato seedling roots by two actinobacteria symbionts of H2 and H3. The growth of symbionts colonies on CGA medium indicated that both actinobacteria isolates, H2 and H3, colonized the tomato seedling roots. (**a**) Control; includes seeds treated only with sterilized distilled water (no symbiont isolates used); (**b**,**c**) colonies formed from aseptic smearing of cut 4–6 mm of elongation zones of roots of each treatment on CGA medium.

3.5. PCR Amplification of 16S rRNA Gene and Phylogeny Analysis

PCR with the 16Sr RNA gene universal primers F1 and R5 amplified a fragment with the expected size of 1490 bp from the *Streptomyces* strains H2 and H3 (Figure 6). Based on the 16S rRNA gene sequences phylogenetic tree (Figure 7) and matrix similarity, the H2 strain was clustered together with the species *S. badius*, *S. caviscabies*, *S. globisporus*, *S. parvus*, *S. sindenensis*, and *S. griseoplanus*, and the strain H3 was clustered with the species *S. flavogriseus*, and *S. pratensis* with 99.9% similarity.
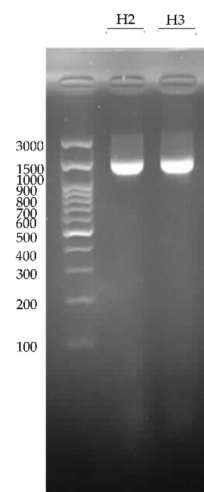

Figure 6. Agarose gel electrophoresis (1%) image of PCR amplification of the 16S rRNA gene of two actinomycetes isolates. M: molecular marker (ExcelBand™ 100 bp+3K DNA Ladder, Smobio, Taiwan); lines H2 and H3 represent PCR product of the related actinomycetes.
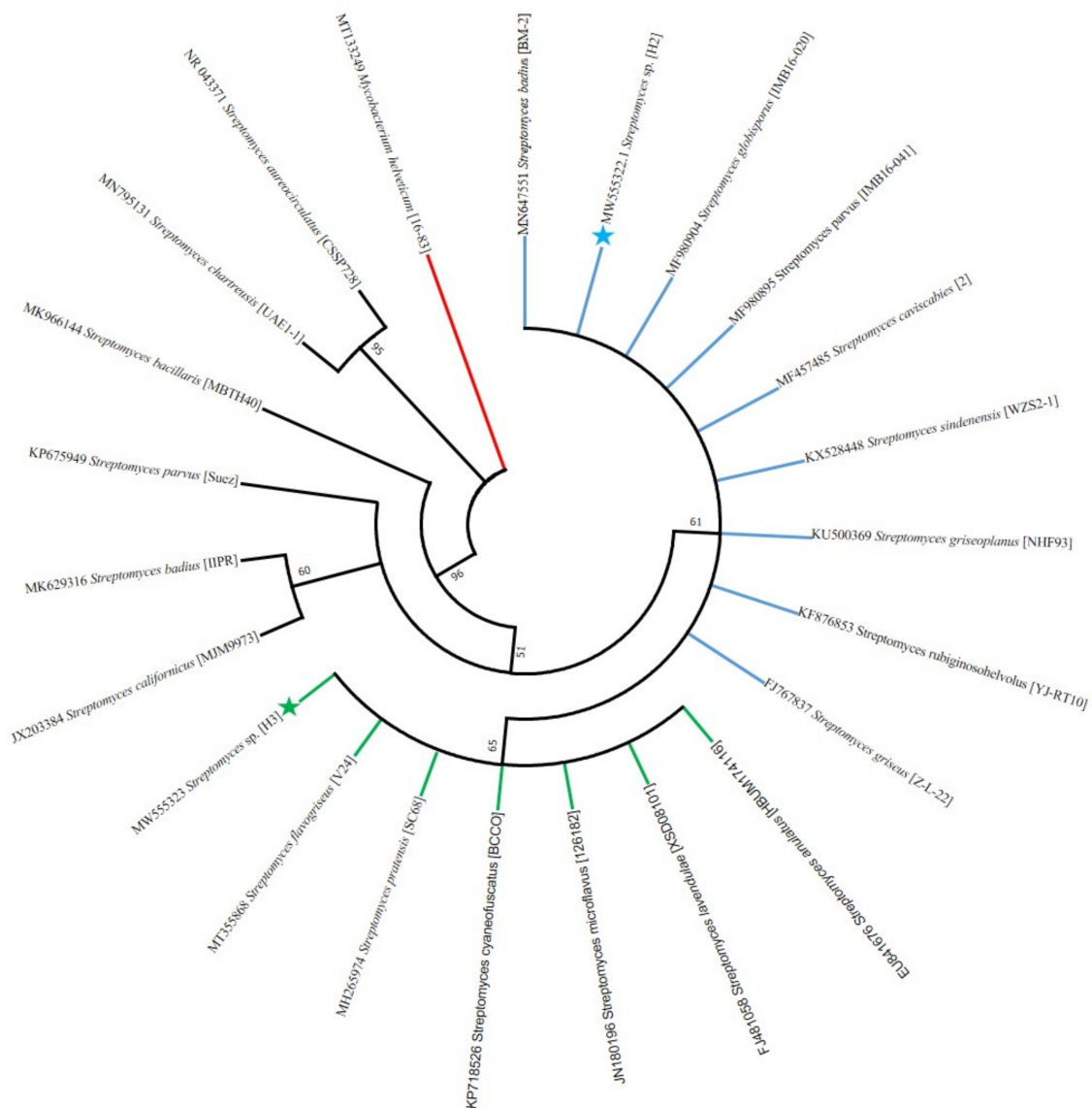
Figure 7. Phylogenetic relations of the *Streptomyces* strains H2 and H3 isolates with 20 known *Streptomyces* strains, based on 16S rRNA gene sequences obtained from GenBank. The tree was constructed using the neighbor-joining method in the MEGA program, version 7.0. Bootstrap values more than 50% from 1000 resampling are shown at the nodes. The 16S rRNA gene sequence from *Mycobacterium helveticum* 16–83 (accession no. MT133249) was used as the out-group.

3.6. Physiological Characterization

The results of physiological characterization of two active symbiont *Streptomyces* strains are indicated in Table 2. Results revealed that both isolates produced gelatinase, catalase, and amylase and could utilize the citrate as a carbon source. Chloroform bioassay showed that bioactive compounds of H2 strain are sensitive to chloroform and became inactive, while those of the H3 strain were resistant to chloroform. Evaluation of the production of volatile organic compounds and HCN revealed that neither strain was able to produce either compound. The maximum tolerable concentrations of NaCl for H2 and H3 strains were 5% for each.

Table 2. Results of physiological characterization of two active symbiont *Streptomyces* strains of H2 and H3.

Bioassay	H2	H3
Catalase production	+	+
Gelatinase production	+	+
Amylase production	+	+
Citrate utilization	+	+
HCN [1] production	−	−
VOC [2] production	−	−
Deactivated by chloroform	+	−

[1] HCN: Hydrogen Cyanide, [2] VOC: Volatile Organic Compound.

3.7. Greenhouse Evaluations

Greenhouse results indicated that the two symbiont *Streptomyces* strains acted as valuable means to subside the disease onset and helped enhance the plants' performance (Figure 8). Results of evaluated criteria are described as follows.
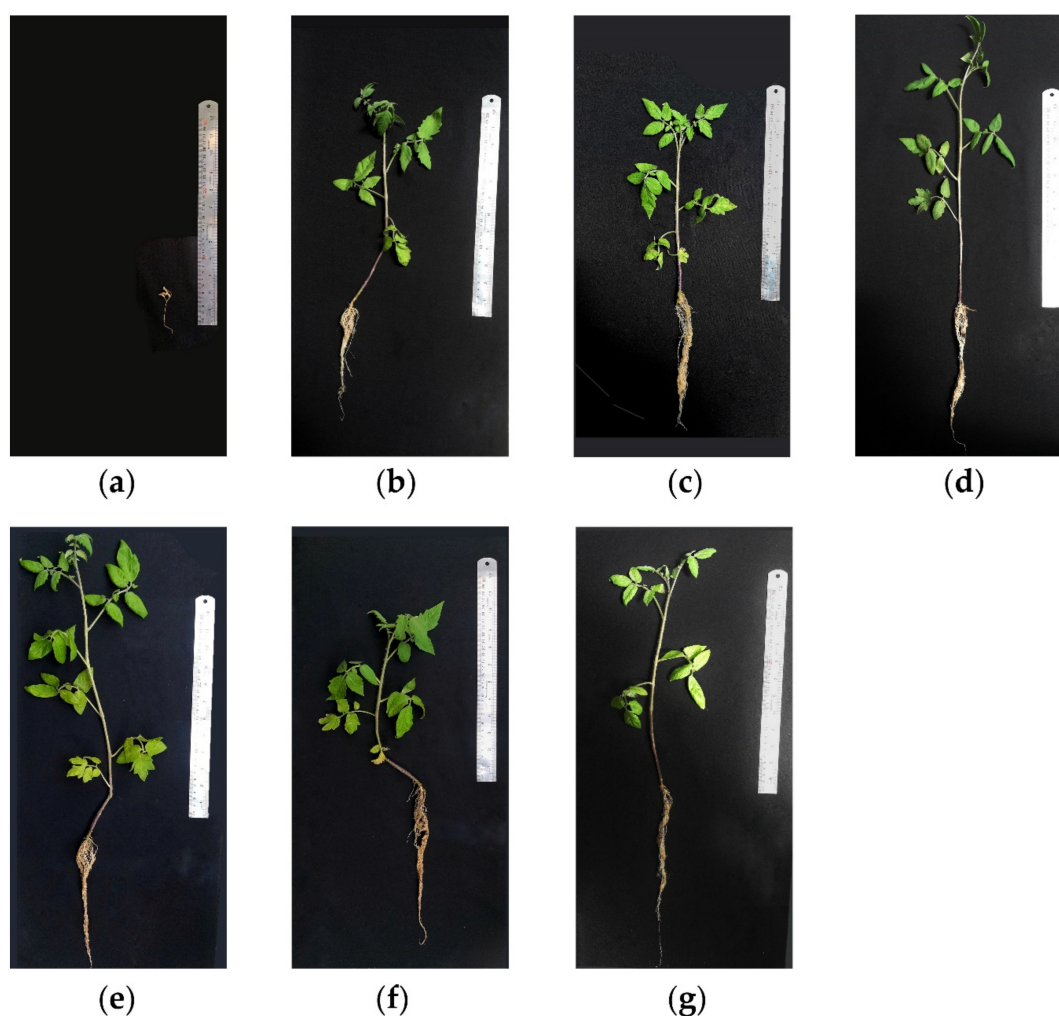

Figure 8. Pictorial results of greenhouse statistical evaluations of interaction between two *Streptomyces* root symbionts, H2 and H3 strains, and the causal agent of damping-off disease, *Pythium aphanidermatum*, on criterion of plant length in tomato seedlings. Each picture is representative of its related treatment: (**a**) tomato seedling treated with *P. aphanidermatum* (positive control); (**b**) seedlings treated neither with pathogen nor symbiont (negative control); (**c** and **d**) seedlings treated with either symbiont *Streptomyces* strains of H2 or H3; (**e,f**) seedlings treated with pathogen and either of symbiont isolates H2 or H3 respectively; (**g**) seedling treated with Metalaxyl (chemical fungicide).

3.7.1. Biocontrol Efficacy of Symbiont Streptomyces Strains

The efficacy of symbiont *Streptomyces* strains evaluated through several parameters are described in following subsections.

Disease Incidence

As indicated in Table 3, the highest disease incidence (47%) belongs to positive control (PC) and the lesser incidences belong to negative control (NC), two symbiont *Streptomyces* strains, and Metalaxyl (Chemical fungicide) treatments. In treatments that received the pathogen and each of two symbiont *Streptomyces* strains, disease incidences (DI) were 31.3% and 21% for symbiont *Streptomyces* strains H2 and H3, respectively. For a better insight into the efficacies of two antagonists, disease incidence adjusted to positive control (%), equalizing disease incidence of positive control to 100% and adjusting the values of other treatments to it, respectively. Accordingly, reduced disease incidence (compared to positive control) was obtained by subtracting disease incidence from 100%. Indeed, the values of reduced disease incidence (%) are indicative of quantitative representation of biocontrol activity of *Streptomyces* strains of H2 and H3.

Number of Leaves

However, as indicated in Table 3, there existed a statistical difference between these treatments and positive control, while there was no statistical difference between negative control, Metalaxyl, and symbiont *Streptomyces* strains at $p \leq 0.05$.

Plants Length

As indicated in Table 3, there existed statistical differences between the plant lengths of the positive control and all other treatments; however, there were no statistical differences among the negative control, H2+ pathogen, and H3+ pathogen at $p \leq 0.05$.

Fresh and Dry Plant Weights

Compared to positive and negative controls, analysis of fresh and dry plant weights showed that *Streptomyces* strain H2 promoted both fresh and dry plant weights while strain H3 enhanced only plant dry weight (Table 3). Considering this criterion, *Streptomyces* strain H2 may be acting as PGPRs.

Table 3. Results of in vivo greenhouse evaluations of two *Streptomyces* root symbionts, H2 and H3 strains, against *Pythium aphanidermatum*, which is the causal agent of damping-off disease in tomato. Evaluated criteria included mean values of triplicates in seven treatments. Recorded criteria in treatments comprised: percentage disease incidence, number of leaves, plant length, fresh and dry stem weights.

Treatment No.	Treatment	%DI [1]	%APC [2]	%RDI [3]	MNL [4]	MPL [5] (cm)	MFSW [6] (g)	MDSW [7] (g)
1	*P. aphanidermatum* [8]	47 [a]	100%	0%	1.9 [c]	8 [c]	0.8 [c]	0.1 [e]
2	*P. aphanidermatum* + H2	31.3 [ab]	66.6%	33.4%	4.4 [b]	42.3 [ab]	3.0 [b]	0.3 [cd]
3	*P. aphanidermatum* + H3	21 [b]	44.7%	55.3%	4.9 [ab]	43.3 [ab]	2.6 [b]	0.2 [de]
4	*P. aphanidermatum* + Metalaxyl	0 [b]	0%	100%	5 [ab]	40.4 [ab]	2.6 [b]	0.3 [cd]
5	*Streptomyces* strains H2	0 [b]	0%	100%	6.8 [a]	43.8 [ab]	5.6 [a]	0.6 [a]
6	*Streptomyces* strains H3	0 [b]	0%	100%	6 [ab]	46 [a]	3.9 [b]	0.5 [a]
7	NC [9]	0 [b]	0%	100%	5.5 [ab]	40 [ab]	3.4 [b]	0.4 [bc]

[1] Disease incidence (DI, %); [2] disease incidence adjusted to positive control (APC, %); [3] reduced disease incidence compared to positive control (RDI, %); [4] mean of number of leaves (MNL); [5] mean of plant length (MPL); [6] mean of fresh stem weights (MFSW); [7] mean of dry stem weights (MDSW); [8] positive controls (*P. aphanidermatum*); [9] negative controls (no pathogen, no symbiont). For a given parameter, different letters indicate significant differences between the different treatments (Duncan's test; $p < 0.05$).

4. Discussion

Damping-off and root diseases of vegetable crops can be controlled using chemical treatments. However, these chemicals negatively impact the environment and humans. Therefore, application of environmentally safe pesticides is a worldwide need in sustainable agricultural pest managements.

Pythium aphanidermatum is a soil-borne phytopathogen and a major causal agent of damping-off disease in many plant species. The pathogen severely damages tomato plants especially in nurseries and greenhouses all over the world [20]. In the present research, 160 pure culture actinobacteria isolated from rhizospheric soil of tomato healthy-appearing fields were screened, in vitro and in vivo, for their anti-*Pythium* activity. This step provided two *Streptomyces* root symbionts, H2 and H3, both with potent biocontrol activities to control tomato damping off disease, which were further characterized in detail at both physiological and molecular levels.

Streptomyces spp. are not only considered for the production of antibiotics but also as a potential biological control of plant pathogenic fungi and bacteria. For instance, *Streptomyces* sp. strain CACIS-1.5CA has been shown to have high bioactivity against several plant pathogenic fungi by inhibition of their spore germination [74]. The antifungal and antibacterial activity of *Streptomyces* sp. strain FJAT-31547 controlled several species of phytopathogenic fungi and bacteria in tomato [71]. Similarly, *Streptomyces* spp. isolate KUBPMB1.1 was efficient to control the *P. aphanidermatum* [34]. The main biocontrol mechanisms of *Streptomyces* species consist of antibiosis by the production of antibiotics [75] and hydrolyzing enzymes. Their activity against phytopathogens consist of producing extra cellular hydrolyzing and degrading enzymes such as cellulases, amylases, chitinases, lipases, etc. [76]. Actinobacteria form an important part of the microbial communities in the soil but also in the rhizosphere and endospheric zones of plant roots [77]. To elucidate the root colonization affinity of *Streptomyces* spp., we developed an "in vitro Petri plate seedling-bioassay" (IPSB), which is relatively a fast bioassay and may be considered as a semi in vivo evaluation bioassay. The IPSB well indicated the potential ability of active actinobacteria for root symbiosis. By implementing this bioassay, we screened 53 actinobacteria isolates, which were demonstrated to be active in dual culture bioassays.

The bioassay test indicated a significant effect of two symbiont *Streptomyces* strains, H2 and H3, on the survival of tomato seedlings inoculated by *P. aphanidermatum*. Both H2 and H3, which revealed their biocontrol potencies, were used to compose main bulk of this paper. *Streptomyces* strain H3 retained its antagonistic activity after exposure to chloroform, a reagent that denatures enzymes, which indicates the involvement of some non-proteinaceous anti-*Pythium* secondary metabolite(s). These results are consistent with other previous results showing that actinobacteria in general, and *Streptomyces* sp. in particular, are of great importance in the field of biotechnology, as producers of a wild range of bioactive secondary metabolites of which many with antibiotic activities that are used in medicine and agriculture [78]. Biochemical bioassay results showed that the two *Streptomyces* strains of H2 and H3 are tolerant to relatively high concentrations of NaCl. Such criteria imply that these strains may be appropriate candidates for biocontrol approaches in saline soils. The biochemical bioassays also showed that two *Streptomyces* strains are able to produce catalase, amylase, and gelatinase. The utilization of citrate showed that these strains can use citrate as a carbon source.

The 16S rRNA-phylogeny confirmed a close relationship of the H2 to the species *S. badius*, *S. caviscabies*, *S. globisporus*, *S. parvus*, *S. sindenensis*, and *S. griseoplanus*, and H3 isolate to *S. flavogriseus*, and *S. pratensis* among 675 validly published *Streptomyces* species (https://lpsn.dsmz.de/genus/streptomyces). The 16S rRNA gene sequencing is not always able to differentiate *Streptomyces* species [79], and it is more proper to discriminate moderately or distantly related species. Hence, to clarify the phylogeny of closely related species in the 16S rRNA gene clades, a multilocus sequence analysis scheme using the other housekeeping genes, *atpD*, *gyr*B, *recA*, *rpo*B, and *trpB* is suggested [80].

Actinomycetes have been reported as efficient biocontrol agents of different soil pathogens through the production of a range of antibiotics with both antibacterial and antifungal action [80] against *Fusarium* spp. [81], *Phytophthora* spp. [82], *Rhizoctonia* spp. [83], *Verticillium* spp. [84], and *Pythium* spp. [85]. The efficacy of bioactivity of H2 and H3 isolates was evidenced in greenhouse trials; however, statistical evaluations showed that similar to the fungicide Metalaxyl, both *Streptomyces* root symbionts significantly suppress the damping-off disease caused by *P. aphanidermatum*. These results are in accordance with reports indicating the antifungal activity of various strains of *Streptomyces* [80]. Furthermore, *Streptomyces* strains were able to reduce the in vitro growth of *Rhizoctonia* and significantly reduce the disease index triggered by this agent when tomato seeds were inoculated with these strains [86]. In addition, *S. anulatus* S37 isolated from the rhizosphere of healthy wild *Vitis vinifera* have been shown as a plant growth-promoting bacteria that enhances the disease resistance toward several pathogens including *B. cinerea* [87]. Hence, when safety procedures passed, these two *Streptomyces* strains can be considered in IPM as appropriate biofungicides. Interestingly, in the criteria of plant weight, the two strains behaved as PGPRs. The increase in plant growth due to antagonist application may be due to the plant growth-promoting characteristics of biocontrol agents [78,88]. Indeed, the greenhouse results indicate that these two root symbionts can be promising candidates to invest on as novel biological control agents in line with the goals of sustainable agriculture. Notably, none of them showed adverse allelopathic effects on plant growth; hence, they are well compatible root symbionts in tomato plant.

In this regard, the study of the effects of both bioactive strains on the rhizosphere microflora (beneficial and/or pathogenic microorganisms) would be useful in future studies, and it would be a great step forward along with goals of sustainable agriculture.

As our results indicate, root symbionts can be considered as promising root-colonizing candidates in integrated management of the damping-off diseases of vegetables in the future. To make such desire come through, further investigations should improve our knowledge of their physiology and environmental accomplishments, including their beneficial or adverse effects on biological stability of the soil; their interaction with other plant root pathogens; their interaction with root mycorrhizae, their interaction with roots of weedy plants, their interaction with other biocontrol agents in soil, their interaction with phyllosphere pathogens, their interaction with endophyte microorganisms, their residual adverse effects on humans through edible plant roots as carrots and even the probable consequences of their genetic transformation such as transduction or conjugation with other prokaryotes, their behavior in other new habitats as forests, rivers, oceans, etc.

This investigation indicated that there were no significant differences between disease control by the chemical fungicide (Metalaxyl®) and either of the two Streptomyces root symbionts, H2 and H3. This perception indicates that there is promise in considering them as candidates for seed treatment and/or soil applications in integrated management of tomato damping-off disease in the future. The current results indicate that these two symbionts are desirable candidates for future investiture in projects with organic agricultural goals.

This is the first report of biocontrol of damping-off in tomato plants caused by *P. aphanidermatum* managed by applying *Streptomyces* in Iran and the second report of employment of *Streptomyces* spp. for biological control of damping-off in tomato caused by *P. aphanidermatum* [34].

Certainly, a wider evaluation of antimicrobial activities of the two bioactive symbionts against other tomato phytopathogens would further add to the knowledge on the bioactivity of these bioactive *Streptomyces*. Furthermore, the assessment of antifungal activity of the two bioactive agents against other causal agents of tomato root disease agents such as *Fusarium* spp., *Verticillium dahlia*, *Rhizoctonia solani*, and *Phytophthora* spp. [3,89] can improve our knowledge on their future applications.

5. Conclusions

Streptomyces spp. are clearly an important part of the soil and/or root microbiome and actively contribute to the control of plant soil pathogens. However, very few products based on *Streptomyces* strains or their bioactive molecules are available on the market. This investigation revealed that (a) the two symbionts *Streptomyces* strain H2 and H3 are appropriate candidates for future investigations as bio-control agents for root fungal pathogens of tomato plants, (b) they behave as plant growth-promoting rhizobacteria able to increase fresh and dry weight of tomato plants; and (c) their biosafety and environmental suitability is yet to be meticulously revealed; however, their immediate field application is not recommended yet.

Supplementary Materials: The following are available online at https://www.mdpi.com/article/10.3390/agronomy11050846/s1, Figure S1: Assessment of *Pythium aphanidermatumpathogenicity* on tomato seedlings (*Lycopersicon esculentum* cv. Moneymaker) in vitro. Healthy non-inoculated seedling (left) and discoloredand *P. aphanidermatum*-killed seedling (right).

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