# Peer

### Aspergillus terreus obtained from mangrove exhibits antagonistic activities against Pythium aphanidermatum-induced damping-off of cucumber

Hanaa Al-Shibli<sup>1</sup>, Sergey Dobretsov<sup>2,3</sup>, Abdulrahman Al-Nabhani<sup>4</sup>, Sajeewa S.N. Maharachchikumbura<sup>1</sup>, Velazhahan Rethinasamy<sup>1</sup> and Abdullah M. Al-Sadi<sup>1</sup>

<sup>1</sup> Department of Crop Sciences, College of Agricultural and Marine Sciences, Sultan Qaboos University, Al-Khoudh, Muscat, Oman

<sup>2</sup> Department of Marine Science and Fisheries, College of Agricultural and Marine Sciences, Sultan Qaboos University, Al-Khoudh, Oman

<sup>3</sup> Center of Excellence in Marine Biotechnology, Sultan Qaboos University, Al-Khoudh, Oman

<sup>4</sup> Department of Pathology, College of Medicine and Health Sciences, Sultan Qaboos University, Al-Khoudh, Oman

### ABSTRACT

A study was conducted to investigate the potential of Aspergillus terreus obtained from Avicennia marina mangrove roots in inhibiting Pythium aphanidermatum and damping-off disease of cucumber. Aspergillus terreus exhibited in vitro inhibition of Pythium aphanidermatum growth. Electron microscope examination revealed that the antagonistic fungal isolate resulted in shrinking and groves in *Pythium* hypha. When Aspergillus terreus culture filtrate was added to Pythium aphanidermatum, it resulted in a significant increase (by 73%) in electrolyte leakage from Pythium hypha compared to the control, as well as significant reduction (by 71%) in oospore production. The Aspergillus terreus culture was also found to produce a cellulase enzyme, which is suggested to be involved in the antagonism against Pythium aphanidermatum. Adding Aspergillus terreus to soil infested with Pythium aphanidermatum significantly reduced percent mortality in cucumber seedlings by 70%. Aspergillus terreus, when applied alone on cucumber seedlings, did not show any suppressive effects on cucumber growth (length and fresh and dry weight). This appears to be the first report of isolation from mangrove of Aspergillus terreus with antagonistic activity against Pythium aphanidermatum-induced damping-off of cucumber. The study indicates that fungal isolates obtained from marine environments may serve as potential biocontrol agents against some plant pathogens.

Subjects Agricultural Science, Microbiology, Plant Science Keywords Biocontrol, Oomycete, Marine fungi, Mangrove, *Avicennia marina* 

### **INTRODUCTION**

*Pythium* species are a major cause of root rot and damping-off diseases in several vegetables, ornamentals, cereals, and other crops (*Al-Sadi et al., 2011; Hatami et al., 2013; Ishiguro et al., 2014; Al-Mahmooli et al., 2015; Gossen et al., 2016). Pythium aphanidermatum* is one of the most commonly distributed and aggressive species of

Submitted 27 April 2019 Accepted 12 September 2019 Published 21 October 2019

Corresponding author Abdullah M. Al-Sadi, alsadi@squ.edu.om

Academic editor Jana U'Ren

Additional Information and Declarations can be found on page 11

DOI 10.7717/peerj.7884

Copyright 2019 Al-Shibli et al.

Distributed under Creative Commons CC-BY 4.0

#### **OPEN ACCESS**

Pythium in the world (*Al-Mawaali et al., 2013; Ishiguro et al., 2014; Le, Smith & Aitken, 2016*). It is the most common cause of damping-off and vine decline diseases of cucumber (*Al-Sadi et al., 2011, 2012; Al-Mawaali et al., 2012, 2013*). It has been reported to result in up to 50% mortality of cucumber seedlings and plants under favorable conditions (*Stanghellini & Phillips, 1975; Al-Sadi et al., 2011*).

Several methods have been developed for the management of *Pythium*-induced diseases. In arid and hot areas of the world, covering soil with transparent polyethylene sheets for a period of 2–8 weeks (solarization) has been effective in reducing populations of *Pythium* and some other fungal species in soil (*Farraq & Fotouh, 2010; Kokalis-Burelle et al., 2017*). Biofumigation has also shown to be effective, but it is less frequently used (*Deadman, Al-Hasani & Al-Sa'di, 2006*). Chemical control through the use of several fungicides such as mefenoxam and hymexazol is widely practiced (*Al-Sadi, 2012; Matić et al., 2019*). However, several studies have shown failure of chemical control due to several reasons, including the development of fungicide resistance, rapid biodegradation of fungicides in soil and delayed activation of the pathogen following application of the fungicide (*Eckert, 1988; Al-Sa'di et al., 2008; Al-Sadi et al., 2012; Al-Balushi et al., 2018*).

The use of biocontrol agents in managing soil-borne diseases is common in several areas of the world, especially under organic agriculture. Several biocontrol agents have been tested and used against *Pythium, Rhizoctonia*, and *Fusarium* species, including *Trichoderma harzianum* and *Pseudomonas aeruginosa* (*Al-Hinai et al., 2010; Blaya et al., 2013; Halo, Al-Yahyai & Al-Sadi, 2018; Kazerooni, Rethinasamy & Al-Sadi, 2018)*. The biocontrol agents interfere with plant pathogens through the production of toxic metabolites and enzymes (e.g., cellulase), competition for food or space and/or direct parasitism (*Bibi et al., 2017; Sharma, Salwan & Sharma, 2017; Vinale et al., 2017; Halo, Al-Yahyai & Al-Sadi, 2018; Zhang et al., 2018*). Biocontrol agents are usually obtained from soils with heavy disease pressures, from plants grown in the same field or from wild plants (*Lyu et al., 2015; Mayo et al., 2017; Raza et al., 2017*). However, no previous attempts have been made to isolate a biocontrol agent against *Pythium aphanidermatum* from mangrove.

Marine fungi are considered critical and important component of the marine ecosystem (*Tisthammer, Cobian & Amend, 2016*). Marine fungi are found on a wide range of substrates, including sediments, algae, sand grains, submerge wood, animals, and plants (*Overy et al., 2019*).

Mangrove is a shrub/small tree that grows in saline environments. The mangrove Avicennia marina is present in coastal areas of several countries, including Oman, China, Iran, and others (*Ghazanfar, 1999; Al-Nadabi & Sulaiman, 2018; Nasr et al., 2018; Tong et al., 2019*). Mangrove has been reported to be rich in biological diversity (*Dias et al., 2012*). Previous studies showed that some fungal species are common in the rhizosphere and different parts of the mangrove tree (*Sanka Loganathachetti, Poosakkannu & Muthuraman, 2017; Prihanto, Caisariyo & Pradarameswari, 2019*). The saline environment, the special life cycle of marine fungi and the unique marine fungal-host interactions make them a valuable source of secondary metabolites (*Balabanova et al., 2018; Deshmukh, Prakash & Ranjan, 2018*).

This study was conducted to investigate whether fungal species associated with mangrove *Avicennia marina* could show antagonistic activities against *Pythium aphanidermatum*. Specific objectives were: (1) isolate and examine the efficacy of *Aspergillus terreus*, obtained from mangrove, on the inhibition of mycelial growth and morphology of *Pythium aphanidermatum*, (2) to its their efficacy in suppressing Pythium damping-off disease, and (3) to examine the production of cellulase enzyme by the fungal isolates. This study will lay the ground for future research on the utilization of marine fungi in managing crop diseases.

### **MATERIALS AND METHODS**

### Sample collection

Breathing roots of mangroves were collected from three trees at Al-Qurum Area in Muscat Governorate, Oman. The collection of samples was done from three plants during October and November 2017. Three leaf/root/stem samples were collected from each plant. Samples (size ~5 cm) were washed with tap water and then cut into small pieces followed by surface sterilization with 1% sodium hypochlorite, and washing three times with sterile distilled water. The samples were then dried on sterilized filter papers. Next, four pieces (~1 cm<sup>3</sup>) from each sample were placed on Petri dishes containing 3% malt extract media (MEA) and 2.5% potato dextrose agar (PDA). The Petri dishes were incubated at room temperature (25 °C) for at least 10 days. Fungi were transferred to new plates. Pure fungal colonies were preserved on 2.5% PDA slants.

# Antagonistic activity of fungal isolates against *Pythium aphanidermatum*

The efficacy of fungal isolates obtained from mangrove roots in inhibiting mycelial growth of *Pythium aphanidermatum* (obtained and identified during a previous study, *Al-Sadi et al.*, *2012*) was investigated. A four mm diameter disc of 10 day-old culture of each fungal isolate was placed at the end of Petri dishes containing 2.5% PDA. A four mm disc of 3-day old *Pythium aphanidermatum* was placed on the other end of the Petri dish. Control plates had *Pythium aphanidermatum* discs on one end, without fungal isolates on the other end. Three replicates were used for the treatments and the control. The plates were incubated at 24 °C up to 10 days, and the experiment was repeated to confirm the results obtained. The size of the inhibition zone was measured after *Pythium aphanidermatum* reached the end of the plate in the control plate.

### Identification of the antagonistic Aspergillus terreus isolate

Since only one fungal isolate (*Aspergillus terreus*) from the previous experiment showed antagonistic activity against *Pythium aphanidermatum*, its identification was confirmed using sequences of the internal transcribed spacer region of the ribosomal RNA gene (ITS rRNA) (*Al-Sadi, Al-Mazroui & Phillips, 2015*). In order to do this, DNA extraction was done from 80 mg of freeze-dried fungal mycelium as described by *Al-Sadi et al. (2012)*. Briefly, the fungal mycelium was ground with 30 mg sterile sand in an Eppendorf tube. Then 600 µl of lysis buffer was added and mixed with the sample. The sample was

incubated at 65 °C for 30 min. After that, 600  $\mu$ l of phenol chloroform isoamyl alcohol (25:24:1) was added followed by centrifugation at 10,000 rpm for 15 min. The last two steps were repeated twice. The supernatant was transferred into new 1.5 ml Eppendorf tubes and 300  $\mu$ l of isopropanol and 10  $\mu$ l of 3M NaOAc were added to the supernatant. The tube was inverted gently and then incubated at -20 °C for 20 min. After that, centrifugation was done for the samples at 10,000 rpm for 2 min by using a refrigerated micro centrifuge. The DNA was washed using 70% ethanol. Then, centrifugation was done for the mixture at 10,000 rpm for 2 min. Ethanol was discarded and the pellet was allowed to dry. The DNA was dissolved in 100  $\mu$ l Milli-Q water and stored at -20 °C.

Amplification of the ITS rRNA region of the isolated fungi was done using the ITS4 (reverse: TCCTCCGCTTATTGATATGC) and ITS5 primers (forward: GGAAGTAAAAGT CGTAACAAGG) (*White et al., 1990*). The PCR reaction mixture and conditions were as described by *Al-Sadi et al. (2012)*.

Sequencing was conducted at Macrogen Inc. (Seoul, Korea). Maximum likelihood analyses including 1,000 bootstrap replicates were run using RaxmlGUI v. 1.3 (*Silvestro & Michalak, 2012*). The *Aspergillus terreus* sequence was deposed at National Center for Biotechnology Information (NCBI) under the accession number MK680140.

# Effect of *Aspergillus terreus* isolate on *Pythium aphanidermatum* morphology

Further work was carried out on the *Aspergillus terreus* isolate by examining the effect of the fungus on *Pythium aphanidermatum* hyphal morphology, which was screened and examined using a Scanning Electron Microscope (SEM, JEOL, Japan JSE- 5600) (*Goldstein et al., 2017*). A piece of mycelium of about 0.5 cm at the inhibition zone was cut. Samples were put in fixative solution to fix the sample tissue. After 1 h of incubation samples were washed two times with buffer (Osmium Tetroxide), 10 min each time. Osmium Tetroxide was added again to the samples and incubated for 1 h to stain the sample lipids and then washed with distilled water. To remove extra water from the sample, ethanol was added to the samples in sequence at concentrations of 25%, 75%, 95% and absolute ethanol, 10 min for each concentration. An automatic drying machine was used to remove the remaining ethanol to reach critical drying point. Finally, the samples were coated with gold and then the samples were examined using the scanning electron microscope.

Differences in morphology between *Pythium aphanidermatum* hypha near the inhibition zone and the control *Pythium aphanidermatum* were examined. The focus was on hypha shape, cytoplasm content, and the end of hypha tips.

### Effect of *Aspergillus terreus* isolate on extracellular conductivity and oospore production by *Pythium aphanidermatum*

The effect of *Aspergillus terreus* isolate culture filtrate (CF) on electrolyte leakage of *Pythium aphanidermatum* was investigated using a conductivity meter (*Manhas & Kaur*, 2016; *Halo, Al-Yahyai & Al-Sadi, 2018*). Six conical flasks containing 200 ml of potato dextrose broth (PDB) were incubated with a four mm disk of *Aspergillus terreus* isolate mycelia at 28 °C for 14 days in an incubator shaker. At the end of the incubation period,

*Aspergillus terreus* mycelia were separated from the CF using sterile filter paper followed by centrifugation at 10,000×g. Moreover, the CF was filtered using Ministar filter with 0.2 μm pore size. Further work was conducted as explained by *Halo, Al-Yahyai & Al-Sadi (2018)*. The experiment was repeated twice.

The antifungal activity of *Aspergillus terreus* isolate was tested on oospore production. Petri dishes containing V8 agar (0.2 l of purified vegetable juice and 10 g of agar in 0.8 l of distilled water) were used as a control and V8 agar was mixed with *Aspergillus terreus* isolate CF at a concentration of 20%. After that, a three mm disk of *Pythium aphanidermatum* was placed onto Petri dishes and incubated at 28 °C for 3 weeks. Oospore number was calculated in 30 consecutive squares of a microscope at 40× magnification. The experiment was conducted using three replicates for the treatments and the control, according to *Halo, Al-Yahyai & Al-Sadi (2018)*.

### Analysis of cellulase production by Aspergillus terreus

Measurement of total cellulase activity was done using the filter paper assay. The *Aspergillus terreus* isolate was grown in Basel medium of Mandels (*Oberoi et al., 2010*). The medium was incubated on a rotary shaker at 200 rpm and 28 °C for 10 days. 0.5 ml of fungal the CF was placed in test tubes containing one ml of sodium acetate buffer (50 mM, pH 4.8). Further assays were according to *Halo, Al-Yahyai & Al-Sadi (2018)*. The absorbance was measured by a spectrophotometer at 540 nm (*Ghose, 1987*). Comparing the absorbance at 540 with glucose standard (0.2–5.2) mg/ml was done to calculate the reducing sugar produced in 1 h.

# Effect of *Aspergillus terreus* isolate on damping-off suppression and on growth of cucumber

The effect of *Aspergillus terreus* on Pythium damping-off disease of cucumber was studied using the following protocol. Three different treatments and a control were prepared. Each treatment and the control consisted of five pots (12-cm diameter) and six cucumber seeds (Jabber, F1) in each pot. Potting mix was added to each pot. A nine cm disc of *Pythium aphanidermatum* was placed on the surface of the potting mix of treatments 1 and 2, followed by covering it with two cm potting mix. A total of 25 ml of *Aspergillus terreus*, grown in 2.5% PDB for 10 days on a rotary shaker at 28 °C, was added to the second treatment. The third treatment was irrigated with *Aspergillus terreus* liquid culture (without *Pythium aphanidermatum*). Control pots received neither *Pythium aphanidermatum*, nor *Aspergillus terreus* treatments. Pots were incubated in a growth chamber for ten days under 80% relative humidity at 28 °C. The surviving seedlings of cucumber, seedling height, and fresh and dry weights were determined for each treatment/control. The experiment was repeated two times. Comparison of means was done using Tukey's Studentized range test (SAS software, v.8).

### RESULTS

### Antagonistic activity of fungal isolates against Pythium aphanidermatum

Isolate H2 (*Aspergillus terreus*) produced an inhibition zone of 2.3 mm (Fig. 1). No further growth of *Pythium aphanidermatum* was detected, even after a month.

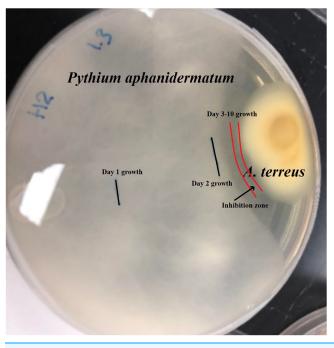


Figure 1 Inhibition zone produced between Aspergillus terreus and Pythium aphanidermatum.Full-size 🖾 DOI: 10.7717/peerj.7884/fig-1

#### Identification of Aspergillus terreus isolate

Growth of culture of H2 isolate in PDA revealed that it belongs to the *Aspergillus* genus. Colonies of *Aspergillus* on PDA attained a diameter of up to 45 mm after 7 days at 25 °C. Colony texture is mostly fluffy and colonies are cinnamon brown and reverse cream-colored to light brown. Conidiophores are long, columnar hyaline and smooth; vesicles nearly globose, biseriate; phialides flask-shaped; conidia  $1.5-2.5 \mu m$  in diameter. The obtained ITS rRNA sequence of the isolate was compared with 10 reference isolates from NCBI. The manually adjusted dataset comprised 506 characters, including gaps. The *Aspergillus* isolate recovered here (H2), clusters with other *Aspergillus terreus* isolates from GenBank with high bootstrap support (100%) (Fig. 2). Morphological and molecular analyses confirm that our isolate is *Aspergillus terreus*.

# Effect of *Aspergillus terreus* isolate on *Pythium aphanidermatum* hyphal morphology

The scanning electron microscope clearly showed that the *Aspergillus terreus* isolate caused significant change in hyphal morphology of *Pythium aphanidermatum* compared with hypha of the control plate. Hypha from the inhibition zone area was shrunk, with deformations and groves, and it showed backward bendings (Fig. 3). The shrinking of hyphae indicated loss of material from the hypha. Bending in hyphae was at the hyphal tips, which was observed on hyphae facing the inhibition zone.

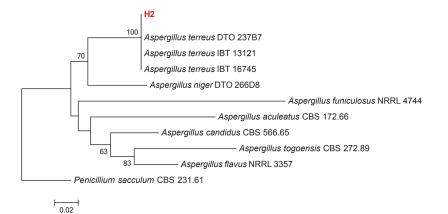
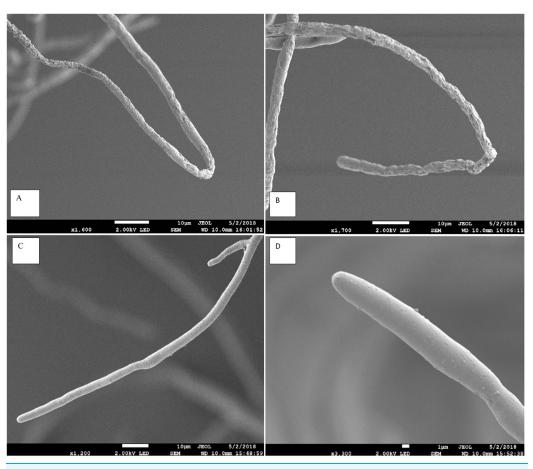


Figure 2 Maximum-likelihood phylogenetic tree inferred from ITS sequences of 11 taxa. Isolates of the present study are depicted in red bold lettering and bootstrap values above 50% are shown in the nodes. The tree is rooted to *Penicillium sacculum* (CBS 231.61).





**Figure 3 Effect of** *Aspergillus terreus* **on morphology of** *Pythium aphanidermatum* **hyphae using scanning electron microscope (SEM).** (A & B) abnormal *Pythium aphanidermatum* hypha: shrunken and wrinkled under effect of *Aspergillus terreus*. (C & D) Normal hypha of Pythium aphanidermatum (control). Full-size DOI: 10.7717/peerj.7884/fig-3

# Effect of *Aspergillus terreus* isolate on electrolyte leakage and production of oospores

Applying the CF of *Aspergillus terreus* on *Pythium aphanidermatum* resulted in significantly increased extracellular conductivity in comparison with the control after 24 h, with 0.22250 mS/cm for the CF of *Aspergillus terreus* and 0.06 mS/cm for control (P < 0.05). This result suggests that treating *Pythium aphanidermatum* mycelium with the CF of *Aspergillus terreus* results in electrolyte leakage. The production of oospores by *Pythium aphanidermatum* in CF of *Aspergillus terreus* (47 oospores) was significantly reduced in comparison with the control (161 oospores, P < 0.05).

### Analysis of cellulase activity by the Aspergillus terreus isolate

The standard graph was plotted with concentration of glucose on the *x*-axis and absorbance value at 540 nm on *y*-axis. Linear equation of glucose stander curve is y = 0.0297x + 0.2333,  $R^2 = 0.9542$ . The cellulase enzyme activity on filter paper was found to be 2.25 FPU/ml in the CF of *Aspergillus terreus* after 60 min of incubation. One filter paper unit (FPU) is defined as the amount of cellulase enzyme that liberates one µmol glucose per ml per minute.

# Effect of *Aspergillus terreus* on damping-off suppression and on growth of cucumber

When cucumber seedlings inoculated with *Pythium aphanidermatum* were treated with *Aspergillus terreus*, the survival rate significantly increased by 70%, to levels similar to when *Aspergillus terreus* was used alone (Fig. 4). In addition, no significant effect was found of *Aspergillus terreus* on shoot length or dry and fresh weight of cucumber seedlings (Fig. 5).

### **DISCUSSION**

The isolation of an antagonistic fungal isolate to *Pythium aphanidermatum* from *Avicennia marina* indicates that mangroves can serve as a source of microbes with potential biological applications. Previous studies have shown that fungal isolates from mangroves showed potent bioactivities. *Aspergillus* sp. D1 isolated from mangrove proved to produce L-asparaginase, which inhibits the formation of acrylamide compounds (*Prihanto, Caisariyo & Pradarameswari, 2019*). *Alternaria longipes* extract from mangrove exhibited antidiabetic activity for type 2 diabetes (*Ranganathan & Mahalingam, 2019*). *Liu et al. (2017b)* reported that some extracts of endophytic fungi from mangrove can inhibit lung cancer cell growth.

There is a recent move toward the search for antagonistic fungal isolates from mangrove against soil-borne fungal pathogens of crops. A study by *Hamzah et al. (2018)* revealed that some fungal isolates obtained from mangrove can inhibit the growth of the soil-borne fungus *Fusarium solani*. *Cladosporium oxysporum* isolated from mangrove was found to have biocontrol potential against *Rhizoctonia solani*, the cause of cucumber damping-off (*Sureshkumar & Kavitha, 2019*).

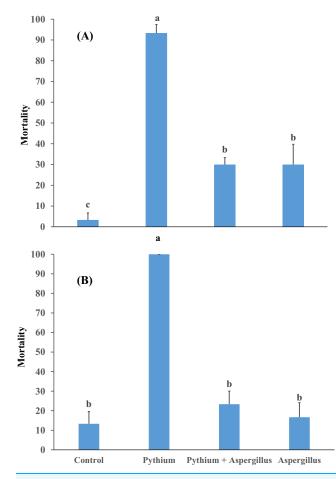


Figure 4 Effect of *A. terreus* on percent seedling mortality in cucumber inoculated with *P. aphanidermatum* ((A) trial 1, (B) trial 2). Columns with different letters indicate that they are significantly different from each other at P < 0.05 (Tukey's Studentized range test, SAS v.8). Error bars are displayed for the different treatments. Full-size  $\square$  DOI: 10.7717/peerj.7884/fig-4

Our study showed that our *Aspergillus* isolate inhibited the growth of *Pythium aphanidermatum*. Identification of the isolate showed that it is *Aspergillus terreus*. *Aspergillus terreus* also resulted in changes in *Pythium aphanidermatum* hyphal morphology, where the affected hypha were shrunk, twisted, and with groves. This gave indication that the isolate not only inhibits growth but also causes abnormalities in the pathogen hypha. A previous study showed that *Aspergillus terreus* isolated from the rhizosphere of cucumber showed antagonistic activities against *Fusarium*, *Rhizoctonia*, and *Sclerotinia*-induced root diseases in cucumber (*Lyu et al., 2015*). Another study showed that some antagonistic fungal isolates inhibiting the growth of *Pythium aphanidermatum* can also result in morphological abnormalities in their hypha (*Halo, Al-Yahyai & Al-Sadi, 2018*).

The CF of the *Aspergillus terreus* isolate affected oospore production and resulted in electrolyte leakage from *Pythium* hyphae. This indicates the *Aspergillus terreus* utilizes several strategies to affect the growth and development of *Pythium aphanidermatum*. *Aspergillus terreus* was found to produce cellulase enzyme. Electrolyte leakage from

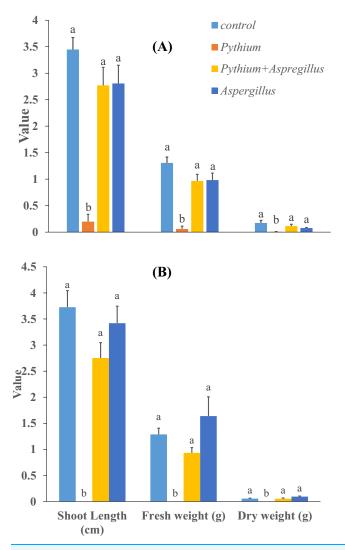


Figure 5 Effect of Aspergillus terreus on shoot length, fresh and dry weight of cucumber seedlings ((A) trial 1, (B) trial 2). Bars with different letters are significantly different from each other (Tukey's Studentized range test, SAS). Error bars are displayed for the different treatments. Full-size DOI: 10.7717/peerj.7884/fig-5

*Pythium* hypha as well as deformations in the *Pythium* hypha might be related in part to the activity of the cellulase enzyme of *Aspergillus terreus*. Several studies indicated the effects of cellulase enzyme of the biocontrol agent, including *Aspergillus* spp., on pathogen hyphae (*Madi et al., 1997; Ayyadurai et al., 2006; Gao et al., 2008; Lee, Tan & Ting, 2014; Sae-Lee & Boonmee, 2014; Halo, Al-Yahyai & Al-Sadi, 2018).* 

The use of *Aspergillus terreus* in *Pythium aphanidermatum*-infested soil significantly reduced damping-off disease of cucumber. This may be related to the effects of *Aspergillus terreus* on *Pythium aphanidermatum* growth, oospore production, and hypha development. Additionally, no negative effects were detected for *Aspergillus terreus* on cucumber growth. Similarly, previous studies showed that antagonistic microorganisms, including *Aspergillus terreus*, either have positive or no negative effects on the growth of

cucumber and other crops (*Elgorban, Bahkali & Al-Sum, 2013*). However, in one trial, the survival rate of cucumber seedlings was higher in the control compared to soil infested with *Aspergillus terreus*. Although no disease was caused by *Aspergillus terreus* in cucumber, and since it was not consistent in the other trial, it may be related to some experimental factors other than pathogenicity of *Aspergillus terreus* on cucumber.

### CONCLUSIONS

Marine fungi received less attention than their terrestrial counterparts. Most of the previous work focused on fungi from soils or plants, with little attention been given to fungi from the marine environment (Al-Balushi et al., 2017; Chen et al., 2017; Dertli & Con, 2017; Liu et al., 2017a; Yao et al., 2017; Huang et al., 2018; Perera et al., 2018; Yang et al., 2018). Although a previous study showed that Aspergillus terreus from desert plants can have antagonistic activities against Pythium aphanidermatum (Halo, Al-Yahyai & Al-Sadi, 2018), this study is the first to characterize a fungal isolate from mangrove (Avicennia marina) with antagonistic activity against Pythium aphanidermatum-induced damping-off of cucumber. This suggests that fungal antagonism is not affected by the origin of isolates, that is, terrestrial isolates and marine isolates of the same species can both show antagonism against terrestrial pathogens. Isolation of a fungal strain from mangrove with potent activity against a crop pathogen necessitate more future studies focusing on the potential activities of mangrove-associated fungal species. In addition, these studies should be coupled with conservation programs of the biologically-rich mangrove "forests." Also, the ability of this isolate to interfere with germination of *Pythium aphanidermatum* spores should be investigated. Several Aspergillus species are known to have negative effects on humans through the production of different types of mycotoxins. Therefore, future studies should investigate the potential presence of negative effects of this isolate on humans or human food.

### ADDITIONAL INFORMATION AND DECLARATIONS

#### Funding

This work was supported by the TRC project (RC/AGR/FISH/16/01), the EG/AGR/CROP/ 16/01 project, Sultan Qaboos University, and the Oman Animal and Plant Genetic Resources Center. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

#### **Grant Disclosures**

The following grant information was disclosed by the authors: TRC project (RC/AGR/FISH/16/01), the EG/AGR/CROP/16/01 project, Sultan Qaboos University, and the Oman Animal and Plant Genetic Resources Center.

#### **Competing Interests**

The authors declare that they have no competing interests.

#### **Author Contributions**

- Hanaa Al-Shibli conceived and designed the experiments, performed the experiments, analyzed the data, prepared figures and/or tables, authored or reviewed drafts of the paper, approved the final draft.
- Sergey Dobretsov conceived and designed the experiments, analyzed the data, contributed reagents/materials/analysis tools, authored or reviewed drafts of the paper, approved the final draft.
- Abdulrahman Al-Nabhani performed the experiments, analyzed the data, contributed reagents/materials/analysis tools, prepared figures and/or tables, authored or reviewed drafts of the paper, approved the final draft.
- Sajeewa S.N. Maharachchikumbura analyzed the data, contributed reagents/materials/ analysis tools, prepared figures and/or tables, authored or reviewed drafts of the paper, approved the final draft.
- Velazhahan Rethinasamy conceived and designed the experiments, contributed reagents/materials/analysis tools, authored or reviewed drafts of the paper, approved the final draft.
- Abdullah M. Al-Sadi conceived and designed the experiments, analyzed the data, prepared figures and/or tables, authored or reviewed drafts of the paper, approved the final draft.

#### **Data Availability**

The following information was supplied regarding data availability:

The sequence data of the fungal isolate is available at GenBank: MK680140.

#### **Supplemental Information**

Supplemental information for this article can be found online at http://dx.doi.org/10.7717/ peerj.7884#supplemental-information.

### REFERENCES

- Al-Balushi ZM, Agrama H, Al-Mahmooli IH, Maharachchikumbura SSN, Al-Sadi AM. 2018. Development of resistance to hymexazol among *Pythium* species in cucumber greenhouses in Oman. *Plant Disease* 102(1):202–208 DOI 10.1094/PDIS-11-16-1680-RE.
- Al-Balushi IM, Bani-Uraba MS, Guizani NS, Al-Khusaibi MK, Al-Sadi AM. 2017. Illumina MiSeq sequencing analysis of fungal diversity in stored dates. *BMC Microbiology* 17(1):72 DOI 10.1186/s12866-017-0985-7.
- **Al-Hinai A, Al-Sadi A, Al-Bahry S, Mothershaw A, Al-Said F, Al-Harthi S, Deadman M. 2010.** Isolation and characterization of *Pseudomonas aeruginosa* with antagonistic activity against *Pythium aphanidermatum. Journal of Plant Pathology* **92**:653–660.
- Al-Mahmooli IH, Al-Fahdi AR, Al-Sadi AM, Deadman ML. 2015. First report of root rot and crown necrosis caused by *Pythium aphanidermatum* on *Phaseolus vulgaris* in Oman. *Plant Disease* **99(3)**:419 DOI 10.1094/PDIS-08-14-0795-PDN.
- Al-Mawaali QS, Al-Sadi AM, Al-Said FA, Deadman ML. 2013. Etiology, development and reaction of muskmelon to vine decline under arid conditions of Oman. *Phytopathologia Mediterranea* 52:457–465.

- Al-Mawaali QS, Al-Sadi AM, Khan AJ, Al-Hasani HD, Deadman ML. 2012. Response of cucurbit rootstocks to *Pythium aphanidermatum*. Crop Protection 42:64–68 DOI 10.1016/j.cropro.2012.07.017.
- **Al-Nadabi A, Sulaiman H. 2018.** Carbon sink potential of *Avicennia marina* in the Al-Qurm nature reserve. In: *IOP Conference Series: Earth and Environmental Science*. Vol. 15. Bristol: IOP Publishing.
- **Al-Sadi AM. 2012.** Efficacy of mefenoxam is affected by a lag period between application and inactivation of *Pythium* species. *Phytopathologia Mediterranea* **51**:292–297.
- Al-Sadi AM, Al-Ghaithi AG, Al-Balushi ZM, Al-Jabri AH. 2012. Analysis of diversity in *Pythium aphanidermatum* populations from a single greenhouse reveals phenotypic and genotypic changes over 2006 to 2011. *Plant Disease* **96(6)**:852–858 DOI 10.1094/PDIS-07-11-0624.
- Al-Sadi AM, Al-Mazroui SS, Phillips AJL. 2015. Evaluation of culture-based techniques and 454 pyrosequencing for the analysis of fungal diversity in potting media and organic fertilizers. *Journal of Applied Microbiology* 119(2):500–509 DOI 10.1111/jam.12854.
- Al-Sadi AM, Al-Said FA, Al-Kiyumi KS, Al-Mahrouqi RS, Al-Mahmooli IH, Deadman ML.
  2011. Etiology and characterization of cucumber vine decline in Oman. *Crop Protection* 30(2):192–197 DOI 10.1016/j.cropro.2010.10.013.
- Al-Sa'di AM, Drenth A, Deadman ML, Al-Said FA, Khan I, Aitken EAB. 2008. Association of a second phase of mortality in cucumber seedlings with a rapid rate of metalaxyl biodegradation in greenhouse soils. *Crop Protection* 27(7):1110–1117 DOI 10.1016/j.cropro.2008.01.007.
- Ayyadurai N, Ravindra Naik P, Sreehari Rao M, Sunish Kumar R, Samrat SK, Manohar M, Sakthivel N. 2006. Isolation and characterization of a novel banana rhizosphere bacterium as fungal antagonist and microbial adjuvant in micropropagation of banana. *Journal of Applied Microbiology* 100(5):926–937 DOI 10.1111/j.1365-2672.2006.02863.x.
- Balabanova L, Slepchenko L, Son O, Tekutyeva L. 2018. Biotechnology potential of marine fungi degrading plant and algae polymeric substrates. *Frontiers in Microbiology* 9:1527 DOI 10.3389/fmicb.2018.01527.
- Bibi F, Ullah I, Alvi SA, Bakhsh SA, Yasir M, Al-Ghamdi AAK, Azhar EI. 2017. Isolation, diversity, and biotechnological potential of rhizo- and endophytic bacteria associated with mangrove plants from Saudi Arabia. *Genetics and Molecular Research* 16(2):gmr16029657 DOI 10.4238/gmr16029657.
- Blaya J, López-Mondéjar R, Lloret E, Pascual JA, Ros M. 2013. Changes induced by *Trichoderma harzianum* in suppressive compost controlling Fusarium wilt. *Pesticide Biochemistry and Physiology* **107(1)**:112–119 DOI 10.1016/j.pestbp.2013.06.001.
- Chen Y-L, Xu T-L, Veresoglou SD, Hu H-W, Hao Z-P, Hu Y-J, Liu L, Deng Y, Rillig MC, Chen B-D. 2017. Plant diversity represents the prevalent determinant of soil fungal community structure across temperate grasslands in northern China. *Soil Biology and Biochemistry* 110:12–21 DOI 10.1016/j.soilbio.2017.02.015.
- **Deadman M, Al-Hasani H, Al-Sa'di AM. 2006.** Solarization and biofumigation reduce *Pythium aphanidermatum* induced damping-off and enhance vegetative growth of greenhouse cucumber in Oman. *Journal of Plant Pathology* **88**:333–335.
- **Dertli E, Çon AH. 2017.** Microbial diversity of traditional kefir grains and their role on kefir aroma. *LWT*—*Food Science and Technology* **85**:151–157 DOI 10.1016/j.lwt.2017.07.017.
- Deshmukh SK, Prakash V, Ranjan N. 2018. Marine fungi: a source of potential anticancer compounds. *Frontiers in Microbiology* 8:2536 DOI 10.3389/fmicb.2017.02536.
- Dias ACF, Taketani RG, Andreote FD, Luvizotto DM, da Silva JL, Nascimento RS, De Melo IS. 2012. Interspecific variation of the bacterial community structure in the phyllosphere of the

three major plant components of mangrove forests. *Brazilian Journal of Microbiology* **43(2)**:653–660 DOI 10.1590/S1517-83822012000200030.

- **Eckert JW. 1988.** Historical development of fungicide resistance in plant pathogens. In: Delp CJ, ed. *Fungicide Resistance in North America*. St. Paul: APS, 1–3.
- Elgorban AM, Bahkali AH, Al-Sum BA. 2013. Biological control of root rots and stems canker of tomato plants caused by *Rhizoctonia solani* in Saudi Arabia. *Journal of Pure and Applied Microbiology* 7:819–826.
- Farraq ESH, Fotouh YO. 2010. Solarization as a method for producing fungal-free container soil and controlling wilt and root-rot diseases on cucumber plants under greenhouse conditions. *Archives of Phytopathology and Plant Protection* 43(6):519–526 DOI 10.1080/03235400701875679.
- Gao J, Weng H, Zhu D, Yuan M, Guan F, Xi Y. 2008. Production and characterization of cellulolytic enzymes from the thermoacidophilic fungal *Aspergillus terreus* M11 under solid-state cultivation of corn stover. *Bioresource Technology* 99(16):7623–7629 DOI 10.1016/j.biortech.2008.02.005.
- Ghazanfar SA. 1999. Coastal vegetation of Oman. *Estuarine, Coastal and Shelf Science* 49:21–27 DOI 10.1016/S0272-7714(99)80004-3.
- Ghose T. 1987. Measurement of cellulase activities. *Pure and Applied Chemistry* 59(2):257–268 DOI 10.1351/pac198759020257.
- Goldstein JI, Newbury DE, Michael JR, Ritchie NWM, Scott JHJ, Joy DC. 2017. Scanning electron microscopy and x-ray microanalysis. New York: Springer-Verlag
- Gossen BD, Conner RL, Chang K-F, Pasche JS, McLaren DL, Henriquez MA, Chatterton S, Hwang S-F. 2016. Identifying and managing root rot of pulses on the northern great plains. *Plant Disease* 100(10):1965–1978 DOI 10.1094/PDIS-02-16-0184-FE.
- Halo BA, Al-Yahyai RA, Al-Sadi AM. 2018. Aspergillus terreus inhibits growth and induces morphological abnormalities in Pythium aphanidermatum and suppresses Pythium-induced damping-off of cucumber. Frontiers in Microbiology 9:95 DOI 10.3389/fmicb.2018.00095.
- Hamzah TNT, Lee SY, Hidayat A, Terhem R, Faridah-Hanum I, Mohamed R. 2018. Diversity and characterization of endophytic fungi isolated from the tropical mangrove species, *Rhizophora mucronata*, and identification of potential antagonists against the soil-borne fungus, *Fusarium solani*. *Frontiers in Microbiology* **9**:1707 DOI 10.3389/fmicb.2018.01707.
- Hatami N, Aminaee MM, Zohdi H, Tanideh T. 2013. Damping-off disease in greenhouse cucumber in Iran. Archives of Phytopathology and Plant Protection 46(7):796–802 DOI 10.1080/03235408.2012.752145.
- Huang S, Maharachchikumbura SSN, Jeewon R, Bhat DJ, Phookamsak R, Hyde KD,
  Al-Sadi AM, Kang J. 2018. *Lecanicillium subprimulinum* (Cordycipitaceae, hypocreales), a novel species from baoshan, Yunnan. *Phytotaxa* 348(2):99–108
  DOI 10.11646/phytotaxa.348.2.4.
- Ishiguro Y, Otsubo K, Watarai M, Iwashita T, Tomita M, Suematsu M, Fukui H, Suga H, Kageyama K. 2014. Seedling blight of *Glycyrrhiza uralensis* caused by *Pythium myriotylum*, *P. aphanidermatum* and *P. spinosum* and identifying primary inoculum sources using multiplex PCR detection. *Journal of General Plant Pathology* 80(3):230–236 DOI 10.1007/s10327-014-0508-4.
- Kazerooni EA, Rethinasamy V, Al-Sadi AM. 2018. *Talaromyces pinophilus* inhibits *Pythium* and *Rhizoctonia*-induced damping-off of cucumber. *Journal of Plant Pathology* **101(2)**:377–383 DOI 10.1007/s42161-018-0186-8.

- Kokalis-Burelle N, McSorley R, Wang KH, Saha SK, McGovern RJ. 2017. Rhizosphere microorganisms affected by soil solarization and cover cropping in *Capsicum annuum* and *Phaseolus lunatus* agroecosystems. *Applied Soil Ecology* 119:64–71 DOI 10.1016/j.apsoil.2017.06.001.
- Le DP, Smith MK, Aitken EAB. 2016. An assessment of *Pythium* spp. associated with soft rot disease of ginger (*Zingiber officinale*) in Queensland, Australia. *Australasian Plant Pathology* 45(4):377–387 DOI 10.1007/s13313-016-0424-5.
- Lee JM, Tan WS, Ting ASY. 2014. Revealing the antimicrobial and enzymatic potentials of culturable fungal endophytes from tropical pitcher plants (*Nepenthes* spp.). *Mycosphere* 5(2):364–377 DOI 10.5943/mycosphere/5/2/10.
- Liu L, Liu Y, Hui R, Xie M. 2017a. Recovery of microbial community structure of biological soil crusts in successional stages of Shapotou desert revegetation, northwest China. *Soil Biology and Biochemistry* 107:125–128 DOI 10.1016/j.soilbio.2016.12.030.
- Liu X, Wu X, Ma Y, Zhang W, Hu L, Feng X, Li X, Tang X. 2017b. Endophytic fungi from mangrove inhibit lung cancer cell growth and angiogenesis in vitro. *Oncology Reports* 37(3):1793–1803 DOI 10.3892/or.2017.5366.
- Lyu H, Niu YC, Deng H, Lin XM, Jin CL. 2015. Suppression of three soil-borne diseases of cucumber by a rhizosphere fungal strain. *Chinese Journal of Applied Ecology* 26:3759–3765.
- Madi L, Katan T, Katan J, Henis Y. 1997. Biological control of *Sclerotium rolfsii* and *Verticillium dahliae* by *Talaromyces flavus* is mediated by different mechanisms. *Phytopathology* 87(10):1054–1060 DOI 10.1094/PHYTO.1997.87.10.1054.
- Manhas RK, Kaur T. 2016. Biocontrol potential of *Streptomyces hydrogenans* strain DH16 toward *Alternaria brassicicola* to control damping-off and black leaf spot of *Raphanus sativus*. *Frontiers in Plant Science* 7(387230):1869 DOI 10.3389/fpls.2016.01869.
- Matić S, Gilardi G, Gisi U, Gullino ML, Garibaldi A. 2019. Differentiation of *Pythium* spp. from vegetable crops with molecular markers and sensitivity to azoxystrobin and mefenoxam. *Pest Management Science* 75(2):356–365 DOI 10.1002/ps.5119.
- Mayo S, Gutiérrez S, Cardoza RE, Hermosa R, Monte E, Casquero PA. 2017. *Trichoderma* species as biocontrol agents in legumes. In: Clemente A, Jimenez-Lopez JC, eds. *Legumes for Global Food Security*. New York: Nova Science Publishers, 73–100.
- Nasr S, Mohammadimehr M, Geranpayeh Vaghei M, Amoozegar MA, Shahzadeh Fazeli SA. 2018. *Aureobasidium mangrovei* sp. nov., an ascomycetous species recovered from Hara protected forests in the Persian Gulf, Iran. *Antonie Van Leeuwenhoek* 111(9):1697–1705 DOI 10.1007/s10482-018-1059-z.
- **Oberoi HS, Chavan Y, Bansal S, Dhillon GS. 2010.** Production of cellulases through solid state fermentation using kinnow pulp as a major substrate. *Food and Bioprocess Technology* **3(4)**:528–536 DOI 10.1007/s11947-008-0092-8.
- Overy PD, Rämä T, Oosterhuis R, Walker KA, Pang K-L. 2019. The neglected marine fungi, sensu stricto, and their isolation for natural products' discovery. *Marine Drugs* 17(1):42 DOI 10.3390/md17010042.
- Perera RH, Maharachchikumbura SN, Hyde KD, Bhat DJ, Camporesi E, Jones EBG, Senanayake IC, Al-Sadi AM, Saichana N, Liu JK, Liu ZY. 2018. An appendage-bearing coelomycete *Pseudotruncatella arezzoensis* gen. and sp. nov. (amphisphaeriales genera incertae sedis) from Italy, with notes on *Monochaetinula*. *Phytotaxa* 338(2):177–188 DOI 10.11646/phytotaxa.338.2.2.
- Prihanto AA, Caisariyo IO, Pradarameswari KA. 2019. Aspergillus sp. as a potential producer for L-Asparaginase from mangrove (Avicennia germinans). In: IOP Conference Series: Earth and Environmental Science. Vol. 230. Bristol: IOP Publishing, 1–7.

- Ranganathan N, Mahalingam G. 2019. Secondary metabolite as therapeutic agent from endophytic fungi Alternaria longipes strain VITN14G of mangrove plant Avicennia officinalis. Journal of Cellular Biochemistry 120(3):4021–4031 DOI 10.1002/jcb.27686.
- Raza W, Ling N, Zhang R, Huang Q, Xu Y, Shen Q. 2017. Success evaluation of the biological control of *Fusarium* wilts of cucumber, banana, and tomato since 2000 and future research strategies. *Critical Reviews in Biotechnology* 37(2):202–212 DOI 10.3109/07388551.2015.1130683.
- Sae-Lee R, Boonmee A. 2014. Newly derived GH43 gene from compost metagenome showing dual xylanase and cellulase activities. *Folia Microbiologica* 59(5):409–417 DOI 10.1007/s12223-014-0313-7.
- Sanka Loganathachetti D, Poosakkannu A, Muthuraman S. 2017. Fungal community assemblage of different soil compartments in mangrove ecosystem. *Scientific Reports* 7(1):8560 DOI 10.1038/s41598-017-09281-3.
- Sharma V, Salwan R, Sharma PN. 2017. The comparative mechanistic aspects of *Trichoderma* and probiotics: scope for future research. *Physiological and Molecular Plant Pathology* 100:84–96 DOI 10.1016/j.pmpp.2017.07.005.
- Silvestro D, Michalak I. 2012. RaxmlGUI: a graphical front-end for RAxML. Organisms Diversity & Evolution 12(4):335–337 DOI 10.1007/s13127-011-0056-0.
- Stanghellini ME, Phillips JM. 1975. *Pythium aphanidermatum*: its occurrence and control with pyroxychlor in the Arabian desert at Abu Dhabi. *Plant Disease Reporter* **59**:559–563.
- Sureshkumar P, Kavitha S. 2019. Bioprospecting potential of mangrove fungus from vellar estuary, southeast coast of india for biocontrol of damping off on mustard. *Research Journal of Biotechnology* 14:72–78.
- **Tisthammer KH, Cobian GM, Amend AS. 2016.** Global biogeography of marine fungi is shaped by the environment. *Fungal Ecology* **19**:39–46 DOI 10.1016/j.funeco.2015.09.003.
- Tong T, Li R, Wu S, Xie S. 2019. The distribution of sediment bacterial community in mangroves across China was governed by geographic location and eutrophication. *Marine Pollution Bulletin* 140:198–203 DOI 10.1016/j.marpolbul.2019.01.046.
- Vinale F, Nicoletti R, Lacatena F, Marra R, Sacco A, Lombardi N, d'Errico G, Digilio MC, Lorito M, Woo SL. 2017. Secondary metabolites from the endophytic fungus *Talaromyces pinophilus*. *Natural Product Research* 31(15):1778–1785 DOI 10.1080/14786419.2017.1290624.
- White TJ, Bruns TD, Lee S, Taylor J. 1990. Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: Innis MA, Gelfand DH, Sninsky JJ, White TJ, eds. *PCR Protocols: A Guide to Methods and Applications*. New York: Academic Press, 315–322.
- Yang J, Maharachchikumbura SSN, Liu J-K, Hyde KD, Gareth Jones EB, Al-Sadi AM, Liu Z-Y. 2018. Pseudostanjehughesia aquitropica gen. et sp. nov. and Sporidesmium sensu lato species from freshwater habitats. Mycological Progress 17(5):591–616 DOI 10.1007/s11557-017-1339-4.
- Yao Q, Liu J, Yu Z, Li Y, Jin J, Liu X, Wang G. 2017. Three years of biochar amendment alters soil physiochemical properties and fungal community composition in a black soil of northeast China. Soil Biology and Biochemistry 110:56–67 DOI 10.1016/j.soilbio.2017.03.005.
- **Zhang L, Niaz SI, Wang Z, Zhu Y, Lin Y, Li J, Liu L. 2018.** α-Glucosidase inhibitory and cytotoxic botryorhodines from mangrove endophytic fungus *Trichoderma* sp. 307. *Natural Product Research* **32(24)**:2887–2892 DOI 10.1080/14786419.2017.1385023.