

Antagonism of *Pseudomonas putida* Against *Dematophora nectarix* A Major Apple Plant Pathogen and Its Potential Use as a Biostimulent

Pankaj Prakash Verma*, Sanjana Thakur and Mohinder Kaur

Department of Basic Science (Microbiology section), Dr. Yashwant Singh Parmar University of Horticulture and Forestry, Nauni, Solan - 173 230, India.

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Plant growth promoting rhizobacteria prove a key to sustainable agriculture provided they effectively colonize roots, survive, proliferate in the rhizosphere and enhance plant growth by a variety of mechanisms. In the present study, several samplings were conducted in different apple growing regions of Himachal Pradesh in order to isolate a strain capable of showing multifarious plant growth promoting activities and antagonism against *Dematophora nectarix* a major apple plant pathogen. In this research five isolates of fluorescent *Pseudomonas* sp. were isolated from apple rhizosphere. Out of five isolates, one isolate DE-18 was selected on the bases of high production of PGP activities and was identified by 16S rRNA gene sequencing. The isolate DE-18 showed maximum phosphate solubilisation and siderophore production of 425 µg/ml and 42.18 %SU respectively. Among the five isolate, DE-18 showed production of both protease (40 mm) and chitinase (28 mm). The isolate DE-18 also showed antagonism against *Dematophora nectarix* (38.46%) and *Phytophthora cactorum* (36.18%) and can act as Bioprotectant. The 16S rDNA based phylogenetic analysis demonstrated that the isolate DE-18 belonged to the *Pseudomonas putida* and sequence was deposited in the GenBank nucleotide sequence databases under accession number KU139388.

Keywords: PGPR, *Pseudomonas putida*, *Dematophora nectarix*, Biocontrol efficacy.

Plant growth-promoting rhizobacteria (PGPR) are naturally occurring soil bacteria that aggressively colonize plant roots and benefit plants by providing growth promotion¹. In the last few years, the number of PGPR that have been identified has seen a great increase, mainly because the role of the rhizosphere as an ecosystem has gained importance in the functioning of the biosphere. Various species of bacteria like *Pseudomonas*, *Azospirillum*, *Azotobacter*, *Klebsiella*, *Enterobacter*, *Alcaligenes*, *Arthrobacter*,

Burkholderia, *Bacillus* and *Serratia* have been reported to enhance the plant growth²⁻⁴. There are several PGPR inoculants currently commercialized that seem to promote growth through at least one mechanism; suppression of plant disease (termed Bioprotectants), improved nutrient acquisition (Biofertilizers), or phytohormone production (Biostimulants). Inoculant development has been most successful to deliver biological control agents of plant disease i.e. organisms capable of killing other organisms pathogenic or disease causing to crops.

Pseudomonas sp. is ubiquitous bacteria in agricultural soils and has many traits that make them well suited as PGPR. The most effective

* To whom all correspondence should be addressed.
Mob: +91 9816922958; Fax: +91 1792252844;
E-mail: pankajmicrobio03@gmail.com

strains of *Pseudomonas* have been *Fluorescent Pseudomonas* sp. and have been studied for decades for their plant growth-promoting effects through effective suppression of soil borne plant diseases. Among various biocontrol agents, *Fluorescent pseudomonads*, equipped with multiple mechanisms for biocontrol of phytopathogens and plant growth promotion, are being used widely as they produce a wide variety of antibiotics, chitinolytic enzymes, growth promoting hormones, siderophores, HCN and catalase, and can solubilize phosphorous⁵⁻⁸. Considerable research is underway globally to exploit the potential of fluorescent *Pseudomonads*. Specific strains of the *Pseudomonas fluorescens-putida* group have recently been used as seed inoculants on crop plants to promote growth and increase yields⁹.

Biological control has gained considerable attention and appears to be promising alternative to chemical control. Biological control has significant potential in terms of both environmental and economic issues for incorporation into organic temperate fruit production. A number of studies have demonstrated benefits resulting from application of plant growth promoting and disease suppressive rhizobacteria to subsequent growth of apple in replant soil. Replant problem is caused by abiotic and biotic factors¹⁰. Various species of fungi like *Fusarium equiseti*, *F. oxysporum*, *F. solani*, *Rhizoctonia* sp., *Cylindrocladium* sp., *Rosellinia necatrix*, *Penicillium claviforme*, *P. janthinellum*, *Phytophthora* sp., *Pythium* sp., *Cylindrocarpon* sp., and of nematodes like *Pratylenchus penetrans* and *Xiphinema* sp. have been found associated with replant disease by various workers¹¹. Root rot is a very serious soil-borne disease infecting temperate fruits especially apple. It is caused by *Rosellinia necatrix* Berl. ex Prill. (Anam. *Dematophora necatrix* Hartig). Agarwala¹² (1961) observed this disease on apple trees in Himachal Pradesh. The annual losses estimated due to this disease are about Rs. 1.3 million¹³ which are expected to be much more as the disease is reported to occur in all apple growing regions of the country.

A diversity of bacterial species has been identified that suppress individual causal elements and enhance growth of plants in replant soil. Biological control of *Phytophthora cactorum*

which contributes to replant disease has been reported in response to application of *Enterobacter aerogenes*¹⁴. Biological control of soil borne pathogens with antagonistic bacteria has gained considerable attention and appears to be promising alternative to chemical control. So the present research was undertaken to isolate and identify a PGPR inoculant that will promote growth and can act as bioprotectant, biofertilizer and a biostimulant.

MATERIALS AND METHODS

Collection of soil samples

Soil samples were collected from rhizosphere of apple plants from Shimla district which is the main apple belt of Himachal Pradesh. The randomly selected apple plants were used for collection of soil samples along with feeder roots from different replant sites of Shimla district viz., Magawta, Shrontha, Siao and Deola. The roots and soil samples from each tree basin were drawn from 15 cm soil depth with the help of soil auger. The samples were carried to the laboratory in polythene bags and were stored in refrigerator at 4° C till further analysis.

Isolation and purification of fungal pathogen

The rhizospheric soil samples were used for isolation of fungal pathogens. The media employed for the isolation of fungus was Potato Dextrose agar (PDA). The plates were incubated at 28 ± 2°C for 3-4 days and the well isolated fungal colonies appeared on plates were purified by subsequent sub culturing. The purified colonies were grown on PDA slants and stored thereafter at 4°C and were subcultured periodically on the same media at 28 ± 2°C. The predominant isolates were identified on the basis of their spore arrangement and fungal isolates were finally identified from National Centre for Fungal Taxonomy, New Delhi.

Isolation of fluorescent *Pseudomonas*, media and growth conditions

Isolation of fluorescent *Pseudomonas* sp. was made from apple rhizosphere in Shimla district of Himachal Pradesh (India). Ten grams of each rhizospheric soil sample were mixed and shaken in 90 ml sterile distilled water blank in 250 ml Erlenmeyer flask for 20-30 minutes to obtain

standard soil suspension. Isolation of fluorescent *Pseudomonas* sp. was made by following the serial dilutions and pour plate method using the specific King's B medium¹⁵. Plates were incubated at $28 \pm 2^\circ\text{C}$ and enumerations of fluorescent *Pseudomonas* colonies were done after 48 h of incubation.

These were identified on the bases of morphological, biochemical and physiological tests viz., Gram's staining, pigment production, oxidase test, catalase test, gelatine liquification, denitrification test and growth at optimum temperature i.e 4°C and 41°C .

Evaluation of PGPR traits

All the *Pseudomonas* isolates from apple rhizosphere were qualitatively and quantitatively characterized by standard protocols for the presence of PGPR traits viz., phosphate solubilisation, siderophore production, HCN production, ammonia production, antifungal activity and for production of hydrolytic enzymes which are known to play an essential role in growth promotion of plants.

Quantification of phosphate solubilisation

Solubilization of P by all the five *Pseudomonas* isolates were estimated using a known amount of inert phosphorus source (tricalcium phosphate) in Pikovskaya's agar medium¹⁶. The composition of the medium was (g/l⁻¹): Glucose, 10.0; $\text{Ca}_3(\text{PO}_4)_2$, 5.0; $(\text{NH}_4)_2\text{SO}_4$, 0.5; $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$, 0.2; MnSO_4 , 0.1; FeSO_4 , 0.0001; Yeast extract, 0.5; Bromocresol purple, 0.1. Quantification of phosphate solubilization was done by spectrophotometric method¹⁷⁻¹⁹. Phosphate solubilizing activity was expressed in terms of tricalcium phosphate solubilization, which in turn represents $\mu\text{g/ml}$ of available orthophosphate as calibrated from the standard curve of KH_2PO_4 (10-100 $\mu\text{g/ml}$).

Siderophore detection and quantification

Siderophore production by *Pseudomonas* isolates was detected by observing orange halos production around the bacterial colony on CAS agar plates²⁰ after 72 h of growth. For quantification of siderophores, to 0.5 ml of cell free culture supernatant of each *Pseudomonas* isolate grown in liquid CAS medium, 0.5 ml of CAS reagent and 10 μl of shuttle solution was added and absorbance was measured at 630 nm against a blank. Siderophores content was expressed as percentage siderophore units using the formula:

$$\% \text{ Siderophore units (\%SU)} = (\text{Ar}-\text{As}) / \text{Ar} \times 100$$

Where Ar = absorbance of reference at 630 nm (CAS reagent) and As = absorbance of sample at 630 nm

HCN and ammonia detection

Bacterial cultures were tested for the production of ammonia in peptone water. Freshly grown cultures were inoculated in 10 ml peptone water and incubated for 48-72 h at $28 \pm 2^\circ\text{C}$. After incubation, Nessler's reagent was added in each tube. Development of brown to yellow colour will be a positive test for ammonia production²¹ and King's B medium amended with 0.44% of Glycine was used for detection of hydrogen cyanide following the method of Bakker and Schippers²² (1987)

Antifungal activity

Isolated bacterial cultures were tested for growth inhibitory effect on the mycelium growth of *Dematophora* sp., *Fusarium* sp., *Phytophthora* sp., *Pythium* sp. and *Rhizocotina* sp, the major fungal pathogens of apple plants by well plate assay method²³.

Chitinase and Protease activity

Chitinase activity was measured according to Chernin²⁴ et al. (1995) and protease activity according to Kaur²⁵ et al. (1988).

DNA Extraction and amplification of 16S rRNA gene fragments

The fluorescent *Pseudomonas* isolate DE-18 was selected on the basis of overall plant growth promoting and disease suppressing activities for their genotypic characterization by 16S rRNA gene sequencing. Genomic DNA was extracted using conventional method (Phenol: Chloroform method) and the DNA was quantified via agarose gel electrophoresis (using 1.0% agarose). 16S rRNA gene analysis was done using *Pseudomonas* specific oligonucleotide primer sequences viz., FP-1 (20-mer): GGTCTGAGAGGATGATCAGT and RP-1 (18-mer): TTAGCTCCACCTCGCGGC in MJ Mini BIO-RAD personal thermal cycler-100 (PTC-100) with a total of 35 cycles. These primers were designed and were used to find their taxonomic affiliation to fluorescent *Pseudomonas* species of group I which mainly comprise *Pseudomonas aeruginosa*, *Pseudomonas putida* and *Pseudomonas fluorescence*.

The PCR amplification was carried out in 0.2 ml PCR tubes with 50 µl reaction volume consisting of following components: 5.0 µl Taq buffer A, 3.0 µl dNTP Mix, 1.0 µl FP-1, 1.0 µl RP-1, 0.40 µl Taq DNA polymerase, 2.0 µl genomic DNA, 37.6 µl double distilled water. A PCR program was implemented as follows: denaturation at 94°C for 1min, two minutes annealing at 55°C and extension for 2 min at 72°C. The final extension was for 10 min at 72°C. For DNA sequencing, eluted amplified

DNA product was first purified followed by sequencing in *B. Genei*, India pvt. Ltd. (Bangalore)

Phylogenetic analysis

The sequencing results were aligned to 16S rRNA gene sequences from GenBank database at NCBI (National Centre of Biotechnological information) through web site <http://www.ncbi.nlm.nih.gov> using BLAST program for screening of sequence similarity. Phylogenetic reconstruction was accomplished with the

Table 1. Details of soil samples and location used for isolation of fluorescent *Pseudomonas* sp

fluorescent <i>Pseudomonas</i> isolates	Location	District	State
M1	Magawta	Shimla	Himachal Pradesh
M2	Magawta	Shimla	Himachal Pradesh
SH1	Shrontha	Shimla	Himachal Pradesh
SO1	Siao	Shimla	Himachal Pradesh
DE-18	Deola	Shimla	Himachal Pradesh

Table 2. Morphological characterization of fluorescent *Pseudomonas* sp. isolated from apple rhizosphere

fluorescent <i>Pseudomonas</i> isolates	Form	Colony morphology			Gram reaction	Cell shape
		Elevation	Margin	Opacity		
M1	Circular	Raised	Entire	Translucent	-	Rod
M2	Circular	Raised	Entire	Translucent	-	Rod
SH1	Circular	Flat	Entire	Transparent	-	Rod
SO1	Circular	Flat	Entire	Transparent	-	Rod
DE-18	Circular	Raised	Entire	Transparent	-	Rod

Table 3. Physiological and biochemical characterization of fluorescent *Pseudomonas* sp. isolated from apple rhizosphere

Parameters	Fluorescent <i>Pseudomonas</i> isolates				
	M1	M2	SH1	SO1	DE-18
Catalase test	+	+	+	+	+
Gelatin liquification	+	+	+	+	-
Lecithinase activity	+	+	-	-	-
Oxidase test	+	+	+	+	+
Spore staining	-	-	-	-	-
Starh hydrolysis	-	-	-	-	-
Tween 80 hydrolysis	-	-	-	-	-
Denitrification	+	+	+	+	+
Growth at 4°C	+	+	+	-	+
Growth at 41°C	-	+	-	+	-

(-) Indicates negativity of test (+) indicates positivity of test

phylogeny MEGA 6.06 inference package. Phylogenetic trees were constructed from distance matrices by the neighbour-joining method and tree topology was estimated by bootstrap analysis, which includes 1000 replicate data sets.

RESULTS AND DISCUSSION

In the present research work, five isolates of fluorescent *Pseudomonas* sp. were isolated from Apple rhizosphere of Magawta, Shrontha, Siao and Deola (Shimla distt., Himachal Pradesh) Table 1. These isolates were further subjected to morphological, physiological and biochemical characterization. All the isolates of fluorescent *Pseudomonas* were studied in detail for colony, colour, growth type, fluorescence, and cell shape. It was evident from the observations that all the five isolates viz., M1, M2, SH1, SO1 and DE-18 produced round shaped colonies with an entire margin and rod shaped cells Table 2. Through morphological and biochemical identification, all the isolates were found to be gram-negative, aerobic, oxidase positive, catalase positive with a fluorescent pigmentation. However, negative

responses were also identified for some *Pseudomonas* isolates such as for gelatin hydrolysis and their ability to grow at 4°C and 41°C. The isolate DE-18 could not hydrolyse gelatine and isolates M1, SH1 and DE-18 did not showed growth at 41°C Table 3. Reynolds²⁶ (2004) also characterized isolates on the bases of biochemical tests including oxidase, catalase, gelatin hydrolysis, nitrate reduction and performing growth at 4 and 41°C and identified them as *P. fluorescens*.

The fungal cultures from rhizospheric soil were purified by single spore and hyphal tip isolation technique. The identification of the isolated fungus was done by studying the morphological characters and by microscopic examination of each isolate. The cultures were sent to National Centre for Fungal Taxonomy, Inderpuri, New Delhi for identification to species level and the fungi were identified as *Fusarium solani* and *Nigrospora oryzae* Table 4 Figure 1.

All the fluorescent *Pseudomonas* isolates were identified as potential phosphate solubilizers based on their capacity to solubilize tricalcium phosphate [$\text{Ca}_3(\text{PO}_4)_2$] by the formation of halo

Table 4. Identification of fungal pathogens isolated from apple rhizosphere

Sr. no	Host	Identification number	Final identification	Replant site
1	Apple rhizosphere	6219.15	<i>Fusarium solani</i>	Magawta (Shimla distt.)
2	Apple rhizosphere	6220.15	<i>Nigrospora oryzae</i>	Deola (Shimla distt.)

Table 5. Characterization of indigenous fluorescent *Pseudomonas* sp. for plant growth promoting traits

fluorescent <i>Pseudomonas</i> isolates	Phosphate solubilisation* (Pi)($\mu\text{g/ml}$)	siderophore production ***% SU	Plant growth promoting traits				
			HCN production	Ammonia production	Plant growth regulators ($\mu\text{g/ml}$)		
					Auxins	Gibberellins	Cytokinins
M1	362.5	20.31	+	++++	8.0	250	75
M2	350	6.25	++	+++	10.0	200	600
SH1	162.5	14.06	+	++	3.0	100	300
SO1	225	4.68	+	+++	2.0	120	30
DE-18	425	42.18	++	++	14.0	160	60

*Phosphate solubilizing activity expressed in terms of tricalcium phosphate solubilization which in turn represents $\mu\text{g/ml}$ of soluble inorganic phosphate (Pi) in supernatant as calibrated from the standard curve of KH_2PO_4 (10-100 $\mu\text{g/ml}$).

**The siderophore unit (% SU) expressed as percent reduction in blue color of chrome azurol-S as compared to reference i.e.

% SU = $(\text{Ar}-\text{As})/\text{Ar} \times 100$

where, Ar = Absorbance of reference solution at 630 nm; As = Absorbance of test solution at 630 nm

zone on Pikovskaya's agar medium. The maximum P solubilisation was recorded by DE-18 (425 µgml⁻¹) compared to other isolates M1, M2, SH1 and SO1 (362.5, 350, 162.5 and 225 respectively) Table 5. Verma²⁷ et al. (2014) also recorded phosphate solubilizing activity of *Pseudomonas* isolates in the range of 155µg/ml to 410 µg/ml. Most of phosphorus in soil is present in the form of insoluble phosphates and cannot be utilized by

the plants. PGPR have been shown to solubilize precipitated phosphates and enhance phosphate availability to plant that represent a possible mechanism of plant growth promotion under field conditions²⁸.

All the tested isolates of fluorescent *Pseudomonads* were positive for the production of siderophores, HCN, ammonia, lytic enzymes and plant growth-promoting hormones viz., auxins, gibberellins and cytokinins. The maximum siderophore production was showed by isolate DE-18 (42.18 %SU) and minimum by SO1 (4.68 %SU) (Table 5). In the development of the antagonism the siderophore production of the bacteria has an important role and such bacteria may function as stress factors including local and systematic host resistance²⁹. Iron competition in *Pseudomonads* has been intensively studied and the role of the pyoverdine siderophore produced by many *Pseudomonas* sp. has been clearly demonstrated in the control of *Pythium* and *Fusarium* species³⁰. Different types of siderophores in *Pseudomonas* species have antagonistic effect against *F. graminearum*³¹. Microbial production of HCN has been suggested as an important antifungal feature to control root fungi pathogen. Cyanide acts as a general metabolic inhibitor to avoid predation or competition. The host plants are generally not harmfully affected by inoculation with HCN

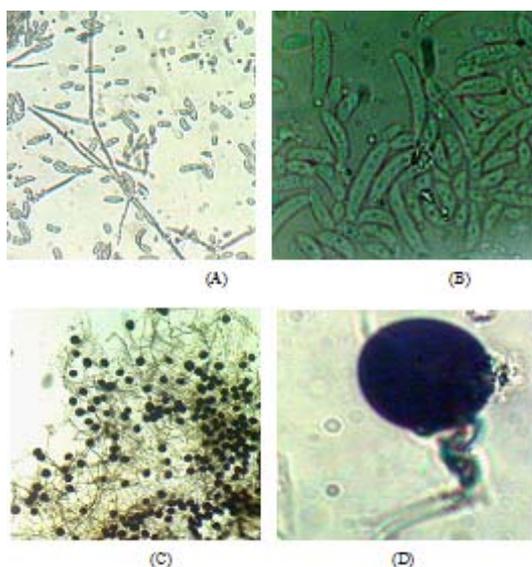


Fig. 1. Microphotographs of *Fusarium solani* (A, B), *Nigrospora oryzae* (C, D)

Table 6. Characterization of fluorescent *Pseudomonas* sp. for antifungal activity against different apple plant pathogens

fluorescent <i>Pseudomonas</i> isolates	<i>Dematophora nectarix</i> (Control=65 mm)		<i>Phytopathora cactoram</i> (Control=47 mm)		Lytic enzymes	
	mm dia	%I*	mm dia	%I*	**Protease (mm dia)	**Chitinase (mm dia)
M1	46	29.23	-	-	12	-
M2	-	-	37	21.27	27	-
SH1	-	-	-	-	30	-
SO1	-	-	-	-	-	-
DE-18	40	38.46	30	36.18	40	28

* % Inhibition of mycelial growth of indicator test fungi by 100µl of 72 h old supernatant grown in nutrient broth at 28+2°C under shaking condition by well plate. All other isolates did not showed antifungal activity against *Fusarium oxysporium*, *Pythium ultimum* and *Rhizocotina* sp

$$\text{Per cent inhibition (\%Inhibition)} = \frac{C-T}{C} \times 100$$

Where, C: Growth of mycelia in control; T: Growth of mycelia in test

**Protease and chitinase activities were expressed in terms of mm diameter of clear zone produced around the well by 100µl of 72 h old supernatant at 28+2°C on skim milk agar after 48 h and minimal agar amended with 1% chitin after 4 days flooded with iodine solution respectively.

producing bacteria and rhizobacteria can operate as biological control agents³².

The production of IAA by PGPR generally affects the root system, increasing the size and number of adventitious roots and also the root subdivision, enabling a bigger soil amount to be exploited by the roots, thus providing large amounts of nutrients accessible to the plant³³. The isolates DE-18 and M2 produced 14.0 and 10.0 μgml^{-1} of auxins respectively, whereas all the other isolates produced auxins between 2.0 and 8.0 μgml^{-1} (Table 5). Similar results were also observed by Sharma³⁴ et al. (2014) in which all the screened fluorescent *Pseudomonas* sp. produced auxin in concentration of 5.1-16.9 μgml^{-1} .

The production of indole acetic acid by the strains of *Pseudomonas* sp. responsible for increasing root elongation was also reported by

O' Sullivan and O' Gara³⁵, (1992). However, IAA production by PGPR can vary among different species and strains, and it is also influenced by culture condition, growth stage and substrate availability³⁶. All the isolates also produced gibberellins and cytokinins in the range of 100-250 μgml^{-1} and 30-600 μgml^{-1} respectively (Table 5). Thakur³⁷ et al. (2013) also observed the production of gibberellins and cytokinins by fluorescent *Pseudomonas* isolates in the range 15.20 to 179.48 $\mu\text{g/ml}$ and 51.20 to 179.48 $\mu\text{g/ml}$ respectively.

The results of inhibition of mycelium growth of the assayed phytopathogenic fungi showed inhibition against only two fungal pathogens among the *Fusarium oxysporium*, *Pythium ultimum*, *Dematophora nectarix*, *Phytophthora cactorum* and *Rhizocotina* sp. From



Fig. 2. Antifungal activity of *Pseudomonas putida* DE-18 against (A) *Phytophthora cactorum* and (B) *Dematophora nectarix*

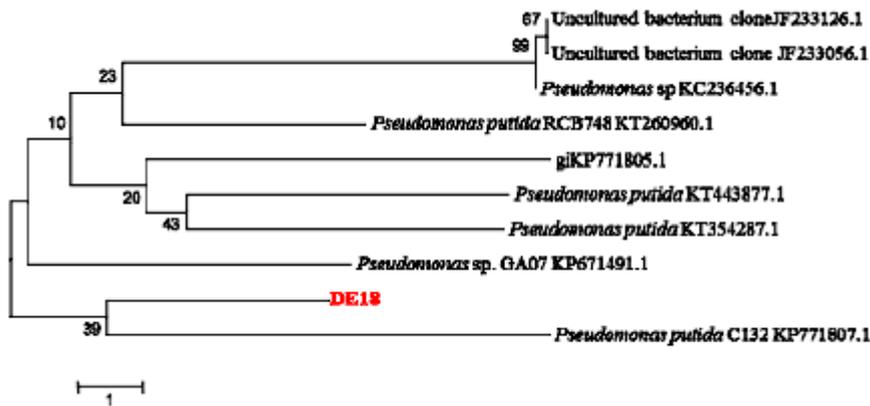


Fig. 3. Phylogenetic tree of *Pseudomonas* sp. strain DE-18 and related species constructed on the basis of 16S rDNA gene sequence using the neighbor-joining method

the assayed five bacterial cultures, the maximum inhibition of mycelium growth of 38.46 % and 36.18 % was found by isolate DE-18 against *Dematophora nectarix* and *Phytopathora cactorum* Table 6 and Figure 2. *Pseudomonas putida* and *Pseudomonas fluorescens*, are frequently isolated from soil and plant tissue and reported as potential biocontrol agents of phytopathogens³⁸⁻⁴⁰. Based on the antagonistic potential and other multifarious plant growth promoting characteristics isolate DE-18 was genotypically characterized by 16S rDNA gene sequencing.

The potential isolate DE-18 was identified based on their 16S rRNA gene using the universal primer set and BLAST analysis. The 16S rDNA based phylogenetic analysis demonstrated that the isolate DE-18 belonged to the genus *Pseudomonas* sp. Figure 3 described the relationship between the isolated strain and the nearest phylogenetic relatives. Similarity calculations after neighbor-joining analysis indicated that the closest relative of isolate DE-18 was *Pseudomonas putida* (90 %). The determined 16S rDNA sequence of strain *Pseudomonas putida* DE-18 was deposited in the GenBank nucleotide sequence databases under accession number KU139388.

This study is assumed to be important as the agriculturally beneficial antibiotic-producing *P. putida* could be one of the potential candidates in the development of microbial pesticides for sustained crop productivity. Hence this *P. putida* DE-18 strain have a potential of being developed as a bio-inoculant for application in a agriculture and horticulture crops.

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