

## Plant growth-promoting activity in newly isolated *Bacillus thioparus* (NII-0902) from Western *ghat* forest, India

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**Abstract** A Gram-positive rod-shaped bacterium isolated on nutrient agar plates incubated at  $28 \pm 2^\circ\text{C}$ . The identity of the bacterium was confirmed by sequencing of the 16S rRNA gene and it reveals that it shares highest similarity with *Bacillus thioparus* CECT 7196<sup>T</sup> (99.08%). It was capable of growing at temperatures ranging from 4 to  $40^\circ\text{C}$ , but optimum growth was observed at  $28 \pm 2^\circ\text{C}$ . Strain NII-0902 is endowed with multiple plant growth promotion attributes such as phosphate solubilization, Indole acetic acid (IAA), siderophore and HCN production, which were expressed differentially at sub-optimal temperatures ( $5\text{--}40^\circ\text{C}$ ). It was able to solubilize phosphate ( $17.7 \mu\text{g ml}^{-1}$ ), and produce IAA ( $139.7 \mu\text{g ml}^{-1}$ ) at  $28 \pm 2^\circ\text{C}$ . Qualitative detection of siderophore production and HCN were also observed. At  $5^\circ\text{C}$  it was found to express all the plant growth promotion attributes except HCN production. The ability to colonize roots is a sine qua non condition for a rhizobacteria to be considered a true plant growth-promoting rhizobacteria (PGPR). *Bacillus* sp. NII-0902 has a potential ability to colonize roots visualized by transparency, bacterial growth (turbid, milky and narrow zone) along and around roots and truly supported by scanning electron micrograph. Hence, it is proposed that, *Bacillus thioparus* sp. NII-0902 could be deployed as an inoculant to attain the desired results of bacterization.

**Keywords** PGPR · *Bacillus* · 16S rRNA sequencing · Western *ghat*

### Introduction

Plants are constantly involved in interactions with a wide range of bacteria. These plant-associated bacteria colonize the rhizosphere (rhizobacteria), the phyllosphere (epiphytes), and the inside of plant tissues (endophytes). Bacterial strains that have beneficial effects on plant health are referred to as beneficial plant-associated bacteria, plant-growth-promoting bacteria (PGPB), or plant-growth promoting rhizobacteria (PGPR; Andrews and Harris 2000). PGPB can promote plant growth directly or indirectly, via biocontrol of host plant diseases, production of phytohormones, or improvement of plant nutritional status (Glick 1995). Rhizobia are perhaps the best known beneficial plant associated bacteria because of the importance of the nitrogen fixation that occurs during the *Rhizobium*–legume symbiosis. The co-inoculation of other PGPB with rhizobia is becoming a practical method in the development of sustainable agriculture, because of yield increases seen compared with inoculation with rhizobia alone. PGPB that have been tested as co-inoculants with rhizobia include strains of the following well-known rhizobacteria: *Azospirillum*, *Azotobacter*, *Bacillus*, *Pseudomonas*, *Serratia* and *Streptomyces* (Baudoin et al. 2010; Biari et al. 2008; Ahmad et al. 2008). PGPB have been isolated by screening the rhizosphere, phyllosphere, and the tissues of plants showing particularly vigorous growth in the field. In this study we isolated the effective plant-growth-promoting *Bacillus thioparus* strain NII-0902 from a soil sample collected from dense forest in Western *ghat* India.

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## Materials and methods

### Isolation

The soil used for bacterial isolation was collected from a root-free soil, rhizosphere, and rhizoplane of Western Ghat in west coast of India, located at an altitude of 900 m above mean sea level. The processed soil sample was serially diluted, spread plated on full strength nutrient agar and incubated at 28°C for 48 h. A cream whitish colored bacterial colony was purified by repeated culturing on nutrient agar (NA) and maintained in 20% glycerol at –80°C. All the subsequent experiments were conducted after raising fresh culture.

### Bacterial identification and characterization

Isolated strain was subsequently differentiated by Gram reaction, salt tolerance and microscopic observation. Sub-sample isolates were initially selected on the basis morphology, physiology and were assessed for Gram reaction. The ability of the isolates to grow in diverse temperature range was carried out by growing each bacterial isolate on nutrient broth and incubated separately at four different temperatures i.e., 5–40°C (5, 15, 30 and 40°C). Results were recorded after every 4 h at 600 nm. The ability of the isolates to grow in different salt concentrations was carried out by inoculating bacterial culture on NA plates supplemented with 0–25% (w/v) NaCl and the plates were incubated at 28 ± 1°C for 3 days. The ability of the isolates to grow in alkaline or acid media was tested in nutrient broth in which the pH was adjusted from 4.0 to 12.0 (at a pH 1.0 interval) and incubated at 28 ± 1°C for 3 days. The ability to use different carbon sources was tested using the Biolog GN Microplate method (Biolog, Hayward, CA). Strain NII-0902 was then screened for traits that might be associated with ability to functions as PGPR, each test performed in triplicate.

### 16S rRNA gene sequencing and phylogenetic analysis

Extraction and amplification of genomic DNA for 16S rRNA gene sequence analysis was carried out as described by Cui et al. (2001). The 16S rRNA gene fragment was amplified by using universal primers. Based on 16S rRNA gene sequences, phylogenetically related bacteria were aligned by using a BLAST search (Altschul et al. 1990) against the GenBank database. Multiple alignments with sequences of related taxa of the genus *Bacillus* were implemented by using CLUSTAL\_X (Thompson et al. 1997). The 16S rRNA gene sequence similarity values were calculated by pairwise comparison (Kimura 1980). A neighbour-joining phylogenetic tree was constructed (Saitou and Nei 1987)

from evolutionary distances calculated using the Jukes–Cantor coefficient (Jukes and Cantor 1969). The topology of the phylogenetic tree was evaluated by the bootstrap resampling method of Felsenstein (1985) with 1000 replicates. The GenBank/EMBL/DDBJ accession number for the isolate is FJ897475.

### Quantitative estimation of phosphate solubilization and IAA production

Initial qualitative estimation of the P-solubilizing activity of the isolate was carried out on Pikovskaya agar (Pikovskaya 1948). Quantitative estimation of tricalcium phosphate (TCP) solubilization and IAA production were carried out at 28°C temperatures. Quantitative estimation of P solubilization was carried out as per standard methodology (Nautiyal 1999), by inoculating 1 ml of bacterial suspension ( $3 \times 10^7$  cells ml<sup>-1</sup>) in 50 ml of NBRIP broth in Erlenmeyer flasks (150 ml), and incubating the flasks for 7 days. At the end of the incubation period the cell suspension was centrifuged at 10,000 rev min<sup>-1</sup> for 10 min and the P content in the supernatant was spectrophotometrically estimated by the ascorbic acid method (Murphy and Riley 1962). Estimation of indole acetic acid (IAA) was done by inoculation of 200 µl of bacterial suspension ( $3 \times 10^8$  cells ml<sup>-1</sup>) in 10 ml Luria–Bertani (LB) broth amended with L-tryptophan (100 µg ml<sup>-1</sup>) and incubating it for a period of 48 h. The IAA content in the culture suspension was estimated by the standard procedure (Gordon and Weber 1951). All the studies were repeated on three independent dates to confirm the results.

### Qualitative measurement of siderophore and hydrocyanic acid (HCN) production

Siderophore and HCN production by the isolate were estimated qualitatively at 28°C. Siderophore production was detected by the Chrome Azurol-S (CAS) assay (Schwyn and Neilands 1987) in 110 mm Petri dishes, and the diameter of the clearing zone was measured. HCN production was inferred by the qualitative method of Bakker and Schipper (1987). The change in the color of the filter paper previously dipped in 2% sodium carbonate prepared in 0.05% picric acid, from yellow to dark brown was rated visually depending on the intensity of the colour change.

### Bioassay-based plant growth promotion ability on wheat

#### *Seed microbiolization*

Rhizobacterial cell suspensions were prepared by adding sterile saline (0.85% NaCl) to slants containing bacterial

cultures in tryptone soy broth in the exponential phase followed by shaking in order to render a turbid suspension. The concentration was adjusted, in a spectrophotometer, to  $OD_{540} = 0.5$ , what corresponds approximately to  $10^8$  c.f.u./ml. Wheat seeds were dipped into the suspension and allowed to imbed overnight at room temperature (Silva et al. 2003).

#### Root colonization bioassay

Wheat seeds were surface sterilized in 50% ethanol (30 s) followed by 2% NaClO (3 min), washed three times in sterile water, soaked in 5 ml of a suspension of rhizobacteria for 24 h and then transferred to sterile 0.6% water-agar tubes. Control seeds were treated with sterile uninoculated medium. Periodic visual inspections were performed daily in order to detect bacterial growth around arising roots. In case of doubt or difficulty of observation, it proved useful to remove the whole seedling from the agar gel for visual inspection. Assays were carried twice with three replicates per culture.

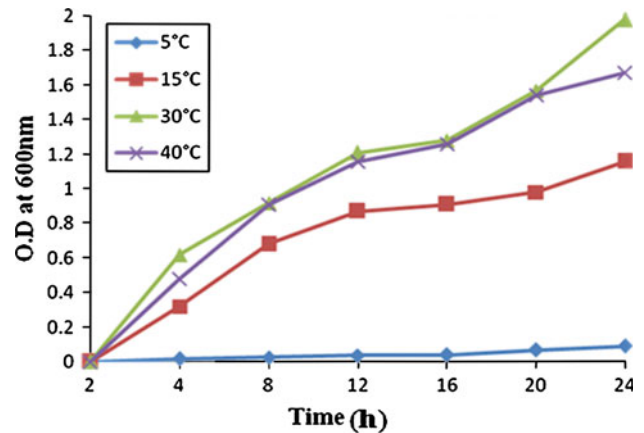
#### Scanning electron microscopic studies

Wheat seedling of 21 day old was randomly selected from growth pots of each treatment for scanning electron microscopic examination. Tissue samples from inoculated and non-inoculated seedling roots of cowpea were thoroughly washed in water to remove soil particles and were fixed in 2% glutaraldehyde (made up in 0.1 M cacodylate buffer) in the refrigerator (8°C) for 1.5 h. Samples were washed two times in the same buffer for 10 min, post fixed in 1% OsO<sub>4</sub> for 4 h, and dehydrated as follows: 30, 50, 70, 85, and 95% ethanol for 15 min; 100% ethanol, two times for 15 min each. For SEM, sputter coating, and a JEOL–JSM 5600LV model operating at 20 kV were used. Root vascular systems and rhizobacteria colonization patterns were observed by SEM.

## Results

#### Characterization of strain NII-0902

The strain NII-0902 was a mesophilic, aerobic, Gram-positive motile bacterium that formed groups of 2–4 cell chains. Cells were rod-shaped measuring 0.5–0.7 μm wide and 1.0–1.7 μm long in 48 h in NA at  $28 \pm 2^\circ\text{C}$  by SEM. Furthermore, colonies grown overnight on NA at 28°C were circular, with smooth margins, cream-white in color and 0.8–1 mm in diameter. Strain NII-0902 grew in 5% (w/v) NaCl but did not grow in the presence of 7, 10 and 15% (w/v) NaCl. Growth was observed at 5, 15, 30 and



**Fig. 1** Growth curve of *Bacillus thioparus* NII-0902, at four different incubation temperatures

40°C but not at 45°C (Fig. 1). Optimal growth temperature on NA was  $28 \pm 2^\circ\text{C}$  at pH 7.0. Strain NII-0902 was positive for catalase, nitrate reduction, hydrogen sulphide, gelatin hydrolysis, citrate utilization, hydrolysis of aesculine, lysine decarboxylase and ornithine decarboxylase. Negative to indole, voges Proskauer's, methyl red, urease, phenyl deamination and β-galactosidase. Positive for assimilation of arabinose, adonitol, cellobiose, malonate, arabinose, glucose, lactose, malonate, melibiose, raffinose, rhamnose, saccharose, trehalose and xylose but negative for oxidase reaction, Table 1 shows other features of strain NII-0902 compared with closely related species of genus *Bacillus*.

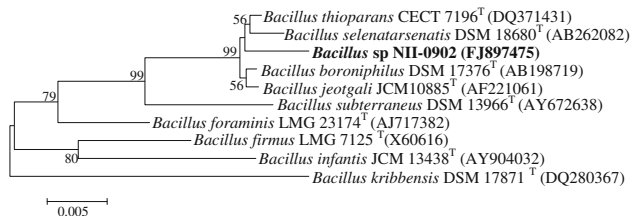
#### Phylogenetic position of the isolate

Strain NII-0902 was assigned to the genus *Bacillus* based on biochemical and morphological criteria. To confirm these results, the 16S rRNA gene sequence was obtained and phylogenetic analysis was carried out. Neighbour-joining tree placed *Bacillus* NII-0902 nested in a clade with some species of the genus *Bacillus*, with a 56% bootstrap value, the clade that contained both strains of *Bacillus thioparus* CECT 7196<sup>T</sup> and *Bacillus selenatarsenatis* DSM 18680<sup>T</sup> (Fig. 2). Strain NII-0902 showed above 99% 16S rRNA gene similarity with both the strains. High degree of similarity among strains of the 16S rRNA gene suggests that all of them may belong to the same species. Nevertheless, there are examples of strains and species with very high 16S rRNA gene sequences identity that differ markedly in their DNA–DNA hybridization values as well as in their morphological and biochemical characteristic that are frequently used to separate one species from the other (Martínez-Murcia et al. 1992; Jaspers and Overmann 2004). High sequence identity could be expected in

**Table 1** Comparison of characteristics of the isolate NII-0902 with its nearest neighbours based on its 16S rRNA gene sequencing

| Characteristic              | NII-0902 | <i>Bacillus thio-parans</i> CECT 7196 <sup>T</sup> | <i>Bacillus selenatarsenatis</i> DSM 18680 <sup>T</sup> |
|-----------------------------|----------|--|---|
| Oxidase                     | –        | –  | –   |
| Catalase                    | +        | +  | +   |
| H <sub>2</sub> S production | +        | –  | +   |
| Gelatin hydrolysis          | +        | W  | +   |
| NaCl tolerance (% w/v)      | 1–7      | 1–5  | 1–5   |
| Optimum pH                  | 7–9      | 7.0  | 7.5–9.0   |
| Maximum growth temp (°C)    | 40       | 45   | 40  |
| Optimum temp (°C)           | 30       | 30–35  | 25–35   |
| ONPG                        | –        | –  | +   |
| Lysine decarboxylase        | +        | –  | ND  |
| Ornithine decarboxylase     | +        | –  | ND  |
| Indole                      | –        | –  | –   |
| Urease                      | –        | –  | ND  |
| Nitrate reduction           | +        | +  | +   |
| Citrate utilization         | +        | +  | ND  |
| Carbon sources utilization  |          |  |   |
| Arabinose                   | +        | –  | –   |
| Rhamnose                    | +        | –  | –   |
| Melibiose                   | +        | –  | –   |
| Glucose                     | +        | +  | +   |

+, positive; –, negative; W, weakly positive; ND, not determined. Comparative data has been collected from Mónica Pérez-Ibarra et al. 2007 and Yamamura et al. 2007



**Fig. 2** Neighbour-joining tree derived from 16S rRNA gene sequences showing the relationships between strain NII-0902 and related *Bacillus* species. Bootstrap percentages (based on 1,000 replications) greater than 50% are given at branching points. Bar 0.005 substitutions per nucleotide position. *Bacillus kribbensis* DSM 17871<sup>T</sup> was used as the outgroup to root the tree. Accession numbers are given in parentheses

*Bacillus* as the rate of evolution of the 16S rRNA gene is exceptionally low due a bottle neck in its evolution (Palys et al. 2000).

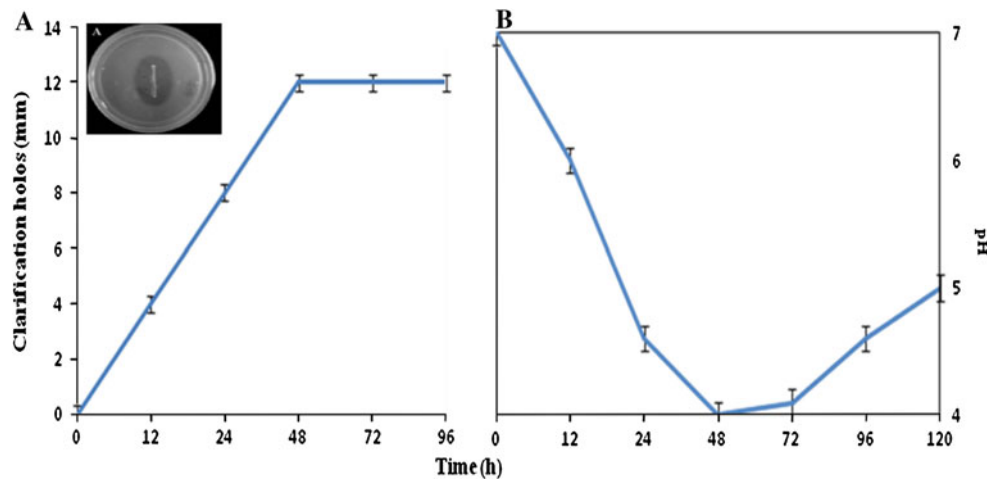
#### Plant growth attributes

The different plant growth promotion traits of the isolate were determined at different incubation temperatures from 5 to 40°C (Table 2). At 5°C the isolate was able to solubilize phosphate (8.0 µg of ml<sup>-1</sup> day<sup>-1</sup>), and produce IAA (38.68.0 µg of ml<sup>-1</sup> day<sup>-1</sup>). The P-solubilizing ability of the isolate was evidently visible on plates of Pikovskaya agar, where it produced a clear halo zone (Fig. 3a). Although the zone of solubilization around the bacterial colony on Pikovskaya agar after varies from 0 to 72 h. The pH of the P-solubilization in broth was found to decline, in each case, due to bacterial activity; lowering of pH coincided with increase in the efficiency of phosphate-solubilizing activity. The pH was found to decline from 7.0 to 4.0–3.0 (Fig. 3b). Qualitative detection of siderophore production and HCN were also observed in all tested temperature. It was interesting to observe that the isolate

**Table 2** Plant growth promoting attributes of *Bacillus thio-parus* NII-0902 at different incubation temperature

| Incubation temperature (°C) | P-solubilization Ca <sub>3</sub> (PO <sub>4</sub> ) <sub>2</sub> µgml <sup>-1</sup> day <sup>-1</sup> | IAA         | HCN production | Siderophore |
|-----------------------------|---|-------------|----------------|-------------|
| 5                           | 8.07 ± 1.2  | 12.3 ± 0.9  | –              | +           |
| 25                          | 11.75 ± 0.3   | 120.6 ± 1.8 | ++             | ++          |
| 30                          | 17.70 ± 0.8   | 139.7 ± 1.2 | ++             | ++          |
| 40                          | 9.61 ± 0.5  | 56.0 ± 1.0  | ++             | +           |

Values are mean of three independent observations, +, moderate; ++, strong activity



**Fig. 3 a** *Bacillus thiooparus* NII-0902 was inoculated on Pikovskaya agar media plates, and incubated for 5 days at 28°C. The clarification halos show P-solubilization and the maximum size of the clarification halos was reached after 96 h. **b** In a separate experiment using liquid National Botanical Research Institute's Phosphate (NBRIP) medium

for bacterial growth, P-solubilization was confirmed by checking pH of the media on daily basis. The pH of the medium was observed to fall continuously up to 72 h. Values given are means of three replicates. *Error bars* indicate standard deviation and are shown when larger than the symbol

**Table 3** Effect of inoculation of *Bacillus thiooparus* NII-0902 on the growth of wheat seedlings

| Plant                               | Treatment                    | Number of roots | Shoot length (cm plant <sup>-1</sup> ) | Root length (cm plant <sup>-1</sup> ) | Plant height (cm plant <sup>-1</sup> ) | Dry weight (mg plant <sup>-1</sup> ) |
|-------------------------------------|------------------------------|-----------------|--|---------------------------------------|--|--------------------------------------|
| <i>Triticum aestivum</i><br>(Wheat) | Control (DW)                 | 2.8 ± 0.2       | 13.9 ± 0.8                             | 9.8 ± 0.8                             | 23.7 ± 1.0                             | 18.2 ± 0.8                           |
|                                     | LB + saline (0.8%)           | 3.2 ± 0.1       | 15.9 ± 1.0                             | 10.8 ± 0.6                            | 26.7 ± 1.2                             | 19.0 ± 1.4                           |
|                                     | <i>Bacillus</i> sp. NII-0902 | 5.4 ± 0.9       | 20.5 ± 1.0                             | 16.8 ± 1.0                            | 37.3 ± 2.0                             | 32.4 ± 2.0                           |

In a column, treatment means having a common letter(s) are not significantly different at the 5% level by DMRT (Duncan's New. Multiple Range Test). 10 ml of culture filtrate (10<sup>8</sup> CFU) was given to sprouts and growth promotion was observed in wheat after 3 week. Distilled water (DW) and saline + Luria-Bertani (LB) media were used as control

was able to retain its functional traits even at 5°C, which was the lower temperature extreme for its growth, while higher values for all parameters were recorded at 28 ± 2°C. The plant growth promotion potential of *Bacillus thiooparus* sp. NII-0902 was determined by a root colonization bioassay in wheat seeds. It was observed that the bacterized seedlings recorded 71.4 and 47.4% higher root and shoot lengths compared to uninoculated control (Table 3). A corresponding increase in the root and shoot biomass was also observed in bacterized seedlings. It was observed that seed bacterization resulted in greater enhancement of the root growth, as compared to the shoot growth.

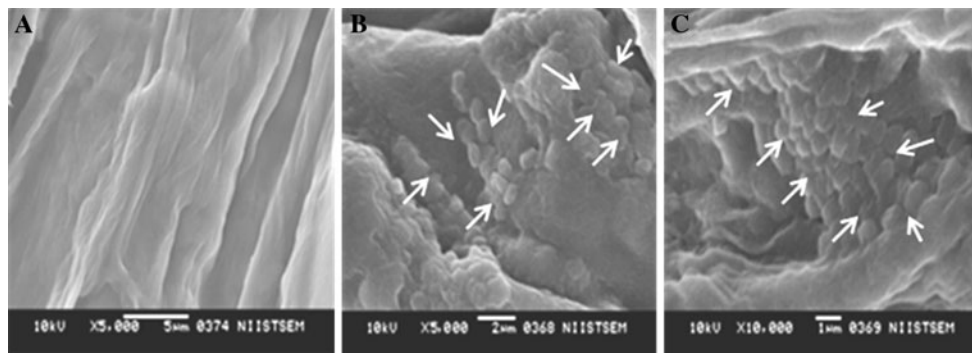
#### Scanning electron microscopic observations

Primary root sections of cowpea bacterized with NII-0902 which showing potential plant growth ability was examined by SEM. The results revealed that cells of isolates NII-0902 were consistently distributed on the surface of roots (Fig. 4b, c). Surface furrows appeared to be located at

epidermal cell junctions. Root seedlings free of inoculant bacteria typically revealed a smooth, undamaged epidermal root surface (Fig. 4a). Root surfaces from isolate NII-0902 inoculated seedlings were colonized with many clusters of cells associated with fibrillar material, which contributed to the formation of microcolonies. Micro colony formation by deleterious bacteria on root surfaces frequently occurs with effective colonization.

#### Discussion

Phosphate is abundant in several soils and is one of the major nutrients limiting the plant growth. The overall phosphate use efficiency following phosphate fertilizer application is low because of the formation of insoluble complexes (Vassilev and Vassileva 2003). Hence, frequent application of soluble forms of inorganic phosphate is necessary for crop production and which leaches to the ground water and results in eutrophication of aquatic systems (Del Campillo et al. 1999). In view of environmental



**Fig. 4** Scanning electron micrograph of wheat seedlings. **a** Root surface of untreated seedlings free of bacteria; **b, c** Seedlings treated with *Bacillus thioeparus* sp. NII-0902. Formations of microcolonies or clustered cells are denoted by *arrowheads*

concerns and current developments in sustainability, research efforts are concentrated on elaboration of techniques that involve the use of less expensive, though less bio-available sources of plant nutrients such as rock phosphate and by application of phosphate solubilizing bacteria and the agronomic effectiveness can be enhanced (Whitelaw 2000). The present study deals with the cold tolerance and plant growth promotion potential of *Bacillus thioeparans* sp. NII-0902 isolated from the Western *ghat* forest soil. As the isolate grew at temperatures ranging from 5 to 40°C, it would be appropriate to call it as cold tolerant. Many cold tolerant plant growth promoting rhizobacteria have been reported from different habitats including *Rhodococcus*, *Pantoea*, *Serratia* and *Pseudomonas* species (Pankaj et al. 2007; Selvakumar et al. 2008a, b, 2009). In this study, an increase in the nutrient uptake as a consequence of seed bacterization has been demonstrated. This phenomenon can be attributed to the ability of the isolate to produce IAA, as IAA positively influences root growth and development, thereby enhancing nutrient uptake (Khalid et al. 2004). Another contributing factor could be the increased availability of phosphorous in the rhizospheric region, as a result of P solubilization by the isolate. It is a well-established fact that improved phosphorous nutrition influences overall plant growth and root development (Jones and Darrah 1994). Further Plant growth promotion activity of the isolate were demonstrated through a plant-based bioassay in tomato seeds. Its use as inoculant resulted in statistically significant increment in root and shoot biomass of wheat seedlings after 21 days of growth. These findings will strongly supported by the previous finding. (Selvakumar et al. 2008a, b, 2009).

From the present study, we demonstrate that the P-solubilizing soil bacteria could serve as efficient biofertilizer candidates for improving the P-nutrition of crop plants. The advantage of using natural soil isolates over the genetically manipulated or the one which has been isolated from a different environmental set up is the easier adaptation and

succession when inoculated into the plant rhizosphere. In conclusion the bacterium-induced plant growth promotion observed in this study was attained under different temperature conditions. Hence, it is proposed that strains NII-0902 can be deployed as a bioinoculants to increase the available phosphorous in soil, helps to minimize the P-fertilizer application, reduces environmental pollution and promotes sustainable agriculture. Future studies are required to prove the nature of these isolates and to harness their potential as bio-inoculants in agriculture.

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