Modulation in Plant Growth and Drought Tolerance of Wheat Crop upon Inoculation of Drought-tolerant- *Bacillus* Species Isolated from Hot Arid Soil of India

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Abstract

Drought is one of the most detrimental environmental stressors to plants with the potential to decrease crop yields and affect agricultural sustainability. Native bacteria with beneficial traits enhance plant growth and help avoid and reverse the effects of drought in plants to a greater extent. In the present study, we aimed to (i) isolate drought tolerant *Bacillus* isolates from the rhizosphere soil of wheat crop grown at different locations in Jaisalmer district, Rajasthan state and (ii) further evaluate their ability to enhance plant growth and induce drought tolerance in wheat (Var. HD-2967) grown under drought stress conditions. Of more than 100 isolates, two putative *Bacillus* isolates capable of tolerating 30 % polyethylene glycol-6000 (PEG-6000) [equivalent to -9.80 MPa (Megapascal)] were identified as *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113. These isolates exhibited different plant growth promoting (PGP) attributes such as phosphate solubilization, and production of siderophore, exopolysaccharide, ammonia, indole acetic acid and cytokinin at low osmotic stress of 10% PEG-6000 but shown variable response at higher osmotic stress particularly at 30% PEG-6000. However, they did not show any antifungal activity and one isolate was negative for phosphate solubilization. Of two strains, *B. altitudinis* DT-89 function more prominently with respect to plant growth promotion and drought tolerance to plant in the early stage but protective traits of *B. paramycoides* DT-113 was more prominent after 75 days as evident by increased EPS (164%), root dry weight (144.44%), chlorophyll content (90.26%), SOD (389%) and proline (99.3%). The results support both the strains as a potential candidate to alleviate drought stress and enhance plant growth in the drought regions.

Keywords: Plant growth promoting bacteria, *Bacillus*, EPS, Siderophore, IAA, GA
INTRODUCTION
Drought, as a consequence of climate change, is one of the major threats to agricultural sustainability and has the potential to tremendously reduce crop yields all over the world. This is especially true for tropical countries where drought is the single most important contributor to crop yield reduction.\(^1\) Growth and yield reduction in crops such as maize, barley, wheat and rice are a matter of concern as it can potentially lead to global food scarcity.\(^2\) Water deficit conditions negatively affect seed germination and seed development, fresh weight of the plants, membrane stability index.\(^3\) Furthermore, drought causes decreased diffusion of water-soluble nutrients such as calcium, magnesium, silica, nitrate, and sulphate from the soil.\(^4\) This, in turn, affects metabolic and physiological activities in the plants like carbon dioxide assimilation, stomatal closure, photosynthesis, proline metabolism, nitrogen metabolism, antioxidant defense system, and plant-water relations.\(^5\)

Millions of microorganisms are dwelling in the soil and also form mutualistic associations with plant roots forming complex ecological communities that assist plant growth and development. In the presence of biotic and abiotic stressors, changes in these ecological communities take place that helps enhance tolerance of the plants to environmental stressors.\(^6\) In order to mitigate drought stress, one promising mechanism is the use of plant growth promoting bacteria (PGPB) that imparts drought-tolerance properties to plants.\(^7\) Under drought stress conditions, PGPB encourage root growth by bringing about changes to root architecture which enables enhanced water uptake by the plant. It has also been seen that inoculation of PGPB can enhance shoot growth under the influence of several plant growth mechanisms of microbes.\(^8\) A study has shown that the treatment of maize plants with PGPB induce closure of stomata to conserve water during drought conditions.\(^9\) Other study has shown increased cellular osmolytes of plants for stress tolerance under drought upon PGPB inoculation.\(^10\) One of the characteristic features of drought-stressed plants is the accumulation of free radicals that cause damage to the cellular machinery of plants. PGPB inoculation leads to the production of antioxidants such as peroxidase and catalase that eliminates free radicals and protects plant cells from damage.\(^11\) This phenomenon has been specifically demonstrated in lettuce plants upon inoculation with *Pseudomonas* sp. and *Glomus* sp.\(^10,12\) Other mechanisms that lead to the improvement of crop yield and root biomass include phytohormone production, ACC deaminase production, production of exopolysaccharides, and solubilization of phosphate and EPS production that helps the plant in water and nutrient acquisition from the soil.\(^12,13\)

Apart from the above, PGPB has been instrumental in plants growth as best studied in desert regions where the climate is mainly hot and dry throughout the year. Plants that grow in these regions acquire drought resistance from phytohormones such as IAA, cytokinin, and abscisic acid (ABA) produced by PGPB in the soil.\(^14\) As a result, PGPB are often used for desert farming specifically for phytostimulation, biofertilization, and biocontrol processes.\(^14\) The PGP activities of these bacteria have been shown to be more pronounced in the areas that have long dry periods as compared to areas where drought occurs less frequently.\(^15\) Comparative studies have shown that plants inoculated with PGPB had higher chlorophyll content, larger number of flowers, and more fecund as compared to uninoculated plants.\(^16\) Inoculation of wheat and sorghum seedlings with PGPB demonstrated higher levels of cellular metabolites that increased plant tolerance to high temperatures. In general, PGPB inoculation not only alters the microbiome of plants, but also their metabolome.\(^17\) The direct effects of PGPB inoculation include enhancing water and nutrient uptake, and promoting homeostasis of phytohormones, whereas its indirect effect includes enhancing plant immunity against pathogens.\(^4\) As the agricultural industry is most vulnerable to global climate change, the use of PGPB in promoting drought tolerance in susceptible plants is a promising intervention. The current study advocates the need to isolate the most compatible and drought tolerant *Bacillus* species from hot arid region of India that promote drought tolerance and growth in wheat crop under drought stress conditions in climatic situation of Eastern Uttar Pradesh.
**MATERIALS AND METHODOLOGY**

**Sample collection and isolation of *Bacillus* from rhizosphere soil**

Rhizosphere soils were collected from wheat crop during the months of February–March 2017 from the four different sites of agricultural field in Roopsi and Sam villages of Jaisalmer district, Rajasthan India (26.9107° N & 70.9144° E and 26.9798° N, 70.7525° E), respectively. Crop plants of wheat were selected and uprooted, with the soil adhered to the roots, from each site and packed in zip locked polybags and transported to the laboratory. The rhizosphere soil was collected by removing the adhered soil from each plant. The putative *Bacillus* isolates were recovered from rhizosphere soil on nutrient agar (NA) medium supplemented with 5% PEG-6000 using serial dilution method. Briefly, 0.1 ml aliquot of each dilution was placed on Petri plates containing nutrient agar followed by incubation at 37°C for 48 h. The visually distinct colonies were picked up and grew in nutrient broth in order to get pure colony of bacteria. Glycerol stocks (20%) of each bacterium was prepared and stored at -80 °C in a deep freezer. All the subsequent experiments were carried out using fresh cultures.

**In vitro evaluation of drought tolerance potential of *Bacillus* isolates**

**On solid medium using PEG**

Bacterial isolates were grown on NA medium supplemented with varied concentration of PEG-6000 (5–30%) in order to select the most tolerant stain to drought conditions, simulated by osmotic pressure created by addition of PEG-6000 followed by incubation of plates at 28 ± 2 °C for 48 h. The growth of bacteria on PEG-6000 supplemented media was recorded. Isolates that survived at highest concentration of PEG-6000 concentration were selected for further studies.21

**In broth medium using PEG**

Bacterial isolates were grown on nutrient broth (NB) supplemented with 10%, 20%, 30% PEG 6000 equivalents to -1.37 MPa (Megapascal), -4.64 MPa, -9.80 MPa osmotic potential, respectively. The nutrient medium supplemented with PEG-6000 was inoculated with 1 ml of 10^9 cfu ml⁻¹ freshly grown culture followed by incubation at 28 ± 2 °C up to 72 h. Optical density (O.D.) of the culture was taken at 600nm. Lowry et al.22 method was used to determine the protein concentration. Uninoculated medium was used as negative control.

**Identification of putative *Bacillus* isolates**

Two putative bacilli were identified at species level by morphological, biochemical,19 and 16S rRNA gene sequencing methods. The sequencing of 16S rRNA gene of two bacilli was performed using universal primers 27F1 (5'- AGAGTTTGATCMTGGCTCAG-3) and 1494R (5'-CTACGGCTACCTTGTTACGAC-3) at National Center for Microbial Resources (NCMR), Pune, India. The 16S rRNA gene sequences of isolates DT-89 and DT-113 were matched with type strains employing EzTaxon server22 and aligned by CLUSTAL W in MEGA version 7.23 software. The topology of evolutionary tree was evaluated by a bootstrap analysis24 of the N-J method based on 1000 replicates by MEGA 7 software. The processed nucleotide sequences data were submitted in the NCBI GenBank to acquire accession number. Finally, the cultures of the *Bacillus altitudinis* DT-89 (NAIMCC-B-02229) and *Bacillus paramycoides* DT-113 (NAIMCC-B-02943) were deposited in NAIMCC (National Agriculturally Important Microbial Culture Collection), ICAR-NBAIM, Mau, Uttar Pradesh, India.

**In vitro assessment of plant growth promoting traits of *Bacillus* spp. under stress conditions**

Plant growth promoting attributes of selected strains were studied at three levels of drought stress (10%, 20%, and 30% PEG-6000) and amended in all the media of the following tests. The bacterial strains were functionally characterized based on production of IAA, GA3, exopolysacharide, siderophore, hydrogen cyanide, ammonium production and ACC deaminase; solubilization of zinc, phosphorus, and antagonisms against phytopathogens by using standard procedures. Bacterial strains were tested for IAA production by procedure given by Ahmad et al.25 Bacterial stains were inoculated separately in 10 ml Luria Bertani (LB) broth in 30 ml tubes. Ten (10) ml LB broth containing supplemented with 200 µg ml⁻¹ L-tryptophan and each tube was inoculated with 48 h grown 100 µl (10^8 ml⁻¹) of culture followed by incubation at 28±2 °C for 4 days with shaking at 125 revolutions per minute (rpm) under normal and osmotic stress conditions. After incubation, a 5 ml culture of each bacterial strain was centrifuged at 9,000 rpm for 15 min. Four (4) ml of Salkowski's
reagent (2% 0.5M FeCl₃ in 35% perchloric acid) along with 100 μl of orthophosphoric acid were added to 2ml of supernatant followed by incubation at 28±2°C in darkness for 1h. The development of pinkish to red colour indicated indole-3-acetic acid (IAA) production. Gibberellic acid (GA3) production by two strains was carried out under normal and osmotic stress conditions as per method of Burrows et al. Briefly, strains DT-89 and DT-113 were grown in LB broth and GA3 was estimated by extracting it with ethyl acetate and measuring absorbance at 680 nm spectrophotometrically. Exopolysaccharide production by two strains was estimated by gravimetric method. For siderophore production assay, bacterial culture was spot inoculated on Chrome Azurol S (CAS) agar and incubated at 28±2°C for 3 days. The appearance of yellow-orange halo around bacterial spot indicates of siderophore production. Quantitative estimation of siderophore was done according to the method of Payne et al. Zinc solubilization was assayed by streaking bacteria on Tris minimal agar medium supplemented separately with 0.1% Zn as zinc oxide, zinc phosphate and zinc carbonate sources. Clear zone around bacterial colony was indicative of solubilization. For quantification of soluble Zn, quantification of soluble Zn from three different insoluble sources solubilized by bacteria in Tri-minimal broth was assayed by using AAS using method described by Fasim et al. The bacterial strains were evaluated for phosphate solubilization on Pikovskaya agar using tricalcium phosphate as an insoluble source of phosphorus according to the standard method of Pikovskaya. The qualitative method of Kremer and Souissi was used to determine HCN production. Ammonia production was assayed by addition of 1 ml Nesseler’s reagent to 72h grown cultures in peptone broth. The positive samples showed yellowish brown colour is an indicative for ammonium production. The ACC deaminase activity was studied as per the method described by Govindaswamy et al. The DT-89 and DT-113 strains were cultured in 5 ml tryptone soya broth separately and incubated at 28±2 °C for 24h at 120 rpm. The cell pellet was harvested by centrifugation at 6000 rpm for 6 min followed by washing two times with sterile distilled water to remove traces of the medium. The washed pellet was suspended in 1 ml sterile distilled water. The bacterium was spotted on Dworkin Foster (DF) minimal medium plates amended with 3mM of ACC, DF medium alone, a negative control, and DF medium containing ammonium sulphate, as a positive control. All the plates were observed after incubation at 28±2 °C for 96 hrs. The bacterium with prominent growth on ACC-DF was indicative of ACC deaminase production. Antagonistic potential of two strains was performed on PDA against Rhizoctonia solani, Macrophomina phaseolina, Fusarium oxysporum f.sp. cumini, F. oxysporum f. sp. carthami and Sclerotium rolfsii employing dual confrontation plate technique. **Plant growth promotion and drought tolerance potential of Bacillus spp. in pot culture**

Two potential drought-adaptive bacteria strains namely DT-89 and DT-113 were taken for evaluation for their growth promoting and drought tolerance inducing ability on wheat crop (Var. HD-2967) in non-sterile soil under net house conditions. Inoculum of strains DT-89 and DT-113 was added in half strength NB and incubated at 28°C for 36 h and pellet of bacteria was obtained after centrifugation at 6000 rpm for 12 min. Bacterial suspension was prepared in 0.85% aqueous saline solution of 0.5 OD equivalent to population of 2.3 x 10⁸ cfu ml⁻¹. Subsequently, seeds of wheat were surface sterilized using 1.0% sodium hypochlorite by exposing it for 30 sec and kept for some time to evaporate trace of sodium hypochlorite. Approximately, 50 g surface sterilized seeds were soaked in 50 ml of 0.5 OD Bacillus species strains suspensions for one hour and then treated seeds were inoculated in the pots and kept in nethouse conditions in the winter of 2019 in Farm of ICAR-National Bureau of Agriculturally Important Microorganisms, Mau. Each pot was containing 10 Kg of sieved (2 mm sieve) alluvial soil and was not fertilized with chemical fertilizers. In total, there were three treatments, (i) Absolute control without any bacterization, (ii) bacterization with DT-89 strain and (iii) bacterization with DT-113 strain with two conditions watered and stressed (50% - that is half dose of water). All the treatments were replicated three times and total 18 pots were used for two stages of sampling. Initially, each pot was sown with 6 seeds for germination and finally 3 seedlings were maintained for each pot throughout experimentation. After germination, plants were maintained under water.
and water stress conditions (50%) throughout experimentation. Plants were harvested after 45 and 75 days, and the soil particles adhered on the roots was removed by washing with tap water. Phenotypic (root and shoot length fresh and dry weight) parameters of the harvested plants were recorded. The leaf samples were preserved at 4 °C for various biochemical estimations like ROS scavenging enzyme activities, proline and chlorophyll content. Chlorophyll content was determined in 80% acetone extract. The absorbance was read spectrophotometrically at 663 and 645 nm. Total chlorophyll concentrations were calculated as per method of Arnon.35 Other parameters were also analysed as mentioned below.

**Estimation of ROS scavenging stress marker enzymes and proline content**

Assays of superoxide dismutase (SOD)37 and peroxidase (POD)38,39 activity were performed utilizing supernatant obtained by centrifuging 500mg of leaves in 1.5 ml of 0.1 M phosphate buffer (pH 7) at 18,000 g, 5°C, 15 minutes.

SOD activity was assayed by measuring the inhibition of nitroblue tetrazolium (NBT) by SOD enzyme. Three (3ml) reaction mixture contained 100 µL of crude extract, 50 mM sodium phosphate buffer (pH 7.6), 0.1 mM EDTA, 50 mM sodium carbonate, 12 mM L-methionine, 50 µM NBT and 10 µM riboflavin. The reaction mixture without crude extract and irradiation served as blank, whereas SOD reaction was carried out by exposing the reaction mixture to white light for 15 min at room temperature. After 15 min incubation, absorbance was recorded at 560 nm using a spectrophotometer. The time required for the mixture optical density to be increased by 0.1 (Δt) at 436 nm was recorded. In this contribution, the resulting end product is called GDHP (guaiacol dehydrogenation product). Free proline was determined as per Bates et al.40 Briefly, 200 mg of leaf sample was homogenized with 3% (w/v) aqueous sulfoalicylic acid and the supernatant obtained was treated with glacial acetic and ninhydrin at 100°C followed by cooling. Now, the chromophore was extracted with toluene by addition in the reaction mixture and finally reading was taken at 520 nm. Proline content was expressed as mg g⁻¹ fresh weight basis.

**Statistical analysis**

All the experiments were performed in triplicates. Statistical analysis was done using Prism GraphPad (Version 9, CA USA) for obtaining P values. Analysis of variance (ANOVA) was performed and the means were compared with Tukey’s test at P<0.05, P<0.001, P<0.0001. Microsoft Office Excel 2007 was used for graphics and determination of standard deviation.

**RESULTS**

**Screening of drought tolerant isolates**

A total of 113 isolates were recovered from wheat rhizosphere soils of 4 different sites of Jaisalmer district, Rajasthan state to screen out potential drought tolerant isolates. Out of 113 isolates screened for drought tolerance ability, 65 (57%), 29 (25%) and 9 (8%) bacteria grew at 10%, 20% and 30% PEG-6000, respectively. Out of 9 isolates, only two potential bacterial isolates DT-89 and DT-113 were selected for further study. These two isolates had also shown tolerance to temperature up to 65°C. Although, growth of two bacteria at 30% PEG-6000 were low but they grew luxuriantly at 10% and 20% PEG concentration as compared to other bacteria tested in this study (Table 1). Isolate DT-113 was found more tolerant in comparison with isolate DT-89 as indicated by more growth by DT-113 under the stress of PEG-6000.

**Plant growth promoting traits under stress conditions in Bacillus spp.**

The two isolates DT-89 and DT-113 had shown differential behavior with respect to display of functional properties. In general, both the isolates were found positive for production of EPS, IAA, GA 3, siderophore, ammonium and...
Table 1. Plant growth promoting traits and drought tolerance potential of isolates DT-89 and DT-113 of *Bacillus* species under non-stressed and drought stressed conditions. Comparison is made with non-stressed (0% PEG-6000) throughout the experiment. One-way ANOVA was performed followed by Tukey’s test.

| Plant growth promoting traits | Bacillus altitudinis DT-89 | Bacillus paramykoides DT-113 |
|------------------------------|-----------------------------|-----------------------------|
|                               | 0% PEG (0 MPa)       | 10% PEG (-1.37 MPa) | 20% PEG (-4.64 MPa) | 0% PEG / (0 MPa) | 10% PEG (-1.37 MPa) | 20% PEG (-4.64 MPa) | 30% PEG (-9.80 MPa) |
| Growth at different PEG concentration | +++ | ++ | + | + | +++ | +++ | ++ | + |
| EPS production (µg ml⁻¹) Control | 2.59±0.17 | 3.05±0.39 | 3.5±0.32**** | 4.73±0.512**** | P<0.0001 | 1.21±0.025 | 2.876±0.39**** | P<0.0001 | 2.88±0.21**** | P<0.0001 | 3.2±0.32**** | P<0.0001 |
| IAA-equivalents (µg ml⁻¹) Control | 21.9±0.24 | 28.02±0.25 | 34.1±0.23** | 23.88±0.22 | 21.4±0.08 | 32.5±0.21**** | 53.65±0.26**** | 22.76±0.09 |
| GA producing abilities (µg ml⁻¹) Control | 4.5±0.20 | 4.33±0.25 ns | 3.85±0.14* | 3.65±0.90* | 4.16±0.57 | 3.91±0.55 | 3.875±0.14 | 5.375±0.13** |
| Siderophore production (SU%) Control | 22.185±3.75 | 23±0.54 [ns] (P>0.05) | 24.38±0.13 [ns] (P>0.05) | 23.84±0.84 | 13.45±0.03 | 46±0.54**** | 50.54±1.13**** | 102.04±0.41**** |
| ACC-deaminase activity | -ve | -ve | -ve | -ve | -ve | -ve | -ve | -ve |
| Ammonia production | +++ve | ++ve | +ve | +ve | +++ve | +ve | +ve | +ve |
| HCN production | -ve | -ve | -ve | -ve | -ve | -ve | -ve | -ve |
| Ca₃(PO₄)₂ solubilization (µg Pi ml⁻¹) | 157.45 | 105.56 | 75.32 | 36.18 | 257.65 | 205.35 | 105.12 | 65.40 |
| Potassium solubilization | -ve | -ve | -ve | -ve | +++ve | +ve | +ve | +ve |
| Zn₃(PO₄)₂ solubilization (µg Zn ml⁻¹) | 136.23 | 98.45 | 65.90 | 35.96 | 162.45 | 145.90 | 130.90 | 87.80 |
| ZnCO₃ solubilization (µg Zn ml⁻¹) | 32.56 | 21.78 | 10.08 | 05.34 | 26.89 | 18.09 | 07.45 | 06.90 |
| ZnO solubilization (µg Zn ml⁻¹) | 67.78 | 66.78 | 40.89 | 15.99 | 87.05 | 56.78 | 41.89 | 32.23 |

Data show the means ± standard deviation of three replicates.

*, **, ****: Significant at the P < 0.05 and P < 0.01, P < 0.001, P < 0.0001 probability levels, respectively. ns, Non-significant. MPa = Megapascal.
solubilization of P and Zn but no ACC deaminase and HCN production were obtained (Table 1).

Isolate DT-89 showed 28% (P>0.05), 55.7% (P<0.001), and 9% (P>0.05) increase in IAA production at 10%, 20% and 30% PEG 6000, respectively, in comparison to 0% PEG supplemented group. Similarly, isolate DT-89 showed increase in EPS by 17.765%, 35.13% and 19.30% (P<0.0001) in comparison to normal conditions. In case of isolate DT-113, a significant high increase of 137%, 137% and 164% (P<0.0001) was found in EPS content at 10%, 20% and 30% PEG 6000 concentration respectively. In comparison to no PEG condition, GA3 activity decreased insignificantly by 3.7%, 14.4% (P<0.05) and 18.8% (P<0.05), at 10%, 20% and 30% PEG 6000 supplementation respectively in DT-89. Further, insignificant 3.67%, 9.89%, and significant 28.19% (P<0.05) increase was observed at 10%, 20% and 30% PEG 6000 conditions, respectively, for siderophore production by isolate DT-89 in comparison to normal control. On the contrary, DT-113 showed significant increase (240%-658%) in siderophore (P<0.0001) at all levels of stress conditions.

In case of isolate DT-113, significant increase of 51.86% (P<0.001) and 150.7% (P<0.0001) in IAA production were observed at 10% and 20% PEG, respectively, in comparison to normal conditions. On the contrary, the GA3 production showed a marked non-significant decrease of 6%, 6.8% and a significant increase of 29.2% (P<0.01) at 10%, 20% and 30% PEG 6000 by strain DT-113 in comparison to the normal condition.

EPS and siderophore production by both the isolates under stress of 30% PEG was highest whereas other traits like IAA, GA3, phosphate- and zinc- solubilization reduced gradually up to 30% PEG. IAA production was also increased up to 20% stress level. Increase production of EPS, siderophore and IAA on increasing stress by increased percent of PEG 6000 might contribute to induction of stress tolerance in Bacillus. Both the isolates were able to solubilize calcium phosphate, zinc phosphate, zinc oxide and zinc carbonate but maximum solubilization was achieved in calcium phosphate (257.65 μg Pi ml⁻¹) and zinc phosphate (162.45 μg Zn ml⁻¹) and least was noted with zinc carbonate (5.34 μg Zn ml⁻¹) (Table 1). Further,
the extent of solubilization was reduced upon increasing level of stress. It is interesting to note that although ACC deaminase production is associated with induction of stress tolerance but none of the isolates showed significant production of ACC deaminase in this study. Both the isolates did not show any antagonism toward most of the phytopathogens except isolate DT-113 which showed weak antagonism to *Fusarium oxysporum* f.sp. *carthami* under no stress and stress of 10% PEG in this study (Table 2). Overall, isolate DT-113 displayed better stress tolerance and plant growth promoting attributes as compared to isolate DT-89.

**Identification of *Bacillus* isolates**

The putative *Bacillus* isolates designated as isolates DT-89 and DT-113 were characterized by morphological, physiological, biochemical and 16S rRNA gene sequencing methods (Table 3). The colony morphology of the isolate DT-89 was undulated, raised, rough with creamy white color pigmentation whereas DT-113 was entire, raised, smooth and creamy white. Under in

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**Table 3. Morphological, physiological, biochemical and molecular characterization of strains DT-89 and DT-113 of *Bacillus* species**

| Characteristics                              | Bacillus altitudinis DT-89 | Bacillus paramycoides DT-113 |
|----------------------------------------------|----------------------------|----------------------------|
| **Isolation source**                         | Rhizosphere of wheat crop | Rhizosphere of wheat crop |
| **Morphological characteristics**           |                            |                            |
| Colony configuration                         |                            |                            |
| Colony margin                                | Undulate                   | Entire                     |
| Colony elevation                             | Raised                     | Raised                     |
| Colony surface                               | Rough                      | Smooth                     |
| Colony pigmentation                          | Creamy white               | Creamy white               |
| Gram reaction                                | + ve                       | + ve                       |
| Endospor                                     | + ve                       | + ve                       |
| **Physiological characteristics**            |                            |                            |
| Growth at/in                                 | Aerobic                    | Aerobic                    |
| Temperature, 15-37°C                          | +++ ve                     | +++ ve                     |
| pH 6.0                                       | + ve                       | + ve                       |
| pH 7.0-10.0                                  | +++ ve                     | +++ ve                     |
| NaCl (%) 5%                                  | +++-ve                     | +++-ve                     |
| NaCl 4.0-15.0                                | 10 % = -ve                 | 15% = -ve                  |
| **Biochemical characteristics**              |                            |                            |
| Indole production                            | - ve                       | - ve                       |
| Methyl red test                              | ++ ve                      | ++ ve                      |
| Voges-Proskauer test                         | - ve                       | - ve                       |
| Citrate utilization                          | - ve                       | - ve                       |
| H₂S production                              | - ve                       | - ve                       |
| Oxidase                                      | + ve                       | + ve                       |
| Catalase                                     | ++ ve                      | ++ ve                      |
| Starch hydrolysis                            | + ve                       | + ve                       |
| Nitrate reduction                            | - ve                       | + ve                       |
| Casein/protease hydrolysis                   | ++ ve                      | ++ ve                      |
| Acid production from                         |                            |                            |
| Dextrose                                     | +++ ve                     | +++ ve                     |
| Lactose                                      | - ve                       | ++ ve                      |
| **16S rRNA gene sequence**                   |                            |                            |
| NCBI Gene bank accession no.                 | MK547281                   | MW959783                   |
| Accession of cultures deposited in MRC       | NAIMCC-B-02229             | NAIMCC-B-02943             |

+ve: positive; -ve: negative; MRC: Microbial Resource Centre
vitro conditions, the isolates DT-89 and DT-113 can tolerate up to 5% and 10% NaCl, respectively and can withstand pH 7.0–10.0 and temperature up to 65°C. Both isolates DT-89 and DT-113 isolates were positive for methyl red, catalase, starch hydrolysis, casein hydrolysis, dextrose utilization, and negative for indole production, Voges-Proskauer test, citrate utilization, oxidase, nitrate reduction and lactose utilization. 16S rRNA gene sequence based phylogenetic analysis indicated that isolates DT-89 was found to be closely related to *Bacillus altitudinis* 41KF2b (99.01%), and DT-113 was similar to *Bacillus paramycoides* MCCC 1A04098 (99.01%) (Fig. 1).

**Plant growth promotion and drought tolerance induction in wheat by Bacillus spp.**

Two drought-tolerant *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 were assessed for their effect on plant growth promoting attributes, chlorophyll content and drought tolerance inducing ability on wheat crop under normal and water stress (drought stress) in net house conditions. In general, the growth associated traits such as shoot and root dry weight, and chlorophyll content were increased in inoculated plants at both normal and 50% water stress conditions over the uninoculated plants (Fig. 2). At 45 days, induction of drought stress

![Phylogenetic tree](image-url)
reduced shoot dry weight by 46.75% (P<0.0001), 9.92% and 9.31% in uninoculated control, DT-89 and DT-113 strains, respectively, as compared to non-stressed condition (watered). Further, on comparison among treatments under drought stressed conditions, strains DT-89 and DT-113 enhanced 54.87% (P<0.0001) and 78.04% (P<0.0001) shoot dry weight, respectively at 45 days of sowing. Similarly, after 75 days of sowing, induction of drought stress reduced shoot dry weight by 45.47% (P<0.0001), 36.40% and 41.70% (P<0.0001) in uninoculated control, DT-89 and DT-113 strains, respectively as compared to watered condition. In case of root dry weight, induction of drought stress reduced root dry weight by significantly by 41.17% (P<0.0001), 40.00% (P<0.0001) and 25.00% in uninoculated control, DT-89 and DT-113 strains, respectively, as compared to respective non-stressed condition (watered) (Fig. 3) and 75.55% (P<0.0001) and 144.44% (P<0.0001) respectively after 75 days of sowing. Similarly, at 75 days of sowing induction of drought stress reduced shoot dry weight by 45%, 35.77% and 44.55% (P<0.0001) in uninoculated control, DT-89, and DT-113 strains respectively. In general, strains DT-89 and DT-113 inoculation induced drought tolerance as indicated by enhanced shoot and root dry weight noted both at 45 and 75 days and maximum being by strain DT-113. In case of chlorophyll content, at 45 days, drought reduced total chlorophyll content by 48.77% (P<0.0001) in uninoculated control but inoculation by DT-89 and DT-113 strains reduced 29.98% and 18.6%, respectively in comparison to their respective normally watered plants groups. In comparison to uninoculated drought stress an increase of 175% (P<0.0001) and 150% (P<0.0001) by DT-89 and DT-113 respectively was observed for 45 days growth treatment. Similarly, at 75 days, drought reduced total chlorophyll content by 22.24% in comparison to uninoculated control (Fig. 4). Further under drought stress 39% (P<0.001) increase was recorded in DT-113 strain inoculated plants in comparison to drought control at 75 days treatment group.

**Effect of PGPB inoculation on ROS scavenging enzymes and proline under drought stress**

In general, drought stress increased superoxide dismutase (SOD), peroxidase (POD) and proline content in wheat plants under uninoculated control condition but inoculation of bacilli further enhanced the activity of enzymes and proline content (Fig. 5, Fig. 6, Fig. 7). *Bacillus altitudinis* DT-89 showed a significant variation of 137% (P<0.0001) in SOD activity at 45 days but
214% increase (P<0.0001) in SOD activity in 75 days treatment in comparison to untreated plants. Similarly, *Bacillus paramycoides* DT-113 showed 151% (P<0.0001), and 389% increase (P<0.0001) in SOD activity at 45 days and 75 days treatment groups respectively in comparison to uninoculated drought control. The SOD content was higher at 75 days as compared to 45 days with maximum SOD as noted upon inoculation of strain DT-113. However, POD activity was increased by 90% (P<0.0001) and 19.63% (P>0.05) by DT-89 treatment at 45 days and 75 days period respectively under drought stress.

Fig. 3. Influence of *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 on shoot dry weight (g) in wheat crop (HD-2967) after 45 and 75 days of sowing under non-stressed and drought (water) stressed conditions. Data represent means ± SD, n = 3. Significance levels are represented as * (P<0.05), ** (P<0.01), *** (P<0.001), **** (P<0.0001).

Fig. 4. Influence of *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 on total chlorophyll (mg g⁻¹ fresh weight) in wheat crop (HD-2967) after 45 and 75 days of sowing under non-stressed and drought (water) stressed conditions. Data represent means ± SD, n = 3. Significance levels are represented as * (P<0.05), ** (P<0.01), *** (P<0.001), **** (P<0.0001).
in comparison to uninoculated drought control. While DT-113 exhibited 36% and 21% increase of drought activity in comparison to drought control. Level of proline content was enhanced by inoculation of bacilli irrespective of unstressed and drought stressed conditions, but drought stressed plants retained more proline content. Strain DT-89 increased proline by 39.28% (P<0.0001) and 155% (P<0.0001) at 45 days and 75 days treatment, respectively, in comparison to uninoculated control drought plants. In case of strain DT-113 treated plants proline value increased by 70.24% (P<0.0001) and 99.3% (P<0.0001) in 45 days and 75 days treatment plants, respectively, in comparison to uninoculated control drought stressed plants. Under drought stress condition, inoculation of strains DT-89 and DT-113 further stimulated production of proline suggesting positive role

**Fig. 5.** Influence of *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 on drought induced enzymes superoxide dismutase (SOD) levels (Unit mg⁻¹ protein) in wheat crop (HD-2967) after 45 and 75 days of sowing under non-stressed and drought (water) stressed conditions. Data represent means ± SD, n = 3. Significance levels are represented as * (P<0.05), ** (P<0.01), ****(P<0.001), ****(P<0.0001)

**Fig. 6.** Influence of *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 on drought induced enzymes Peroxidase (POD) levels (Enzyme unit L⁻¹) in wheat crop (HD-2967) after 45 and 75 days of sowing under non-stressed and drought (water) stressed conditions. Data represent means ± SD, n = 3. Significance levels are represented as * (P<0.05), ** (P<0.01), ****(P<0.001), ****(P<0.0001)
of bacilli in improvement in plant growth under drought stress. Based on above results, it has been inferred that *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 have the ability to promote plant growth and inducing drought tolerance in wheat plant by enhancing enzymes and proline production.

**DISCUSSION**

The bacilli represent an important member of microbial community present in soil. Studies have found that *Bacillus* species occur predominantly in the rhizosphere of wheat and barley.41,42 *Bacillus* species have bestowed with properties like endospore formation that help them to survive in the rhizosphere under abiotic stress and thus can manifest their role as plant growth promoting rhizobacteria by diverse modes of action. They can exhibit important mechanisms of biological control, biostimulation and biofertilisation.43 In the present study, 113 isolates of *Bacillus* were screened out for drought tolerance behavior and out of which *Bacillus altitudinis* DT-89 and *Bacillus paramycoides* DT-113 have been chosen based on their ability to survive under high water stress conditions and possess plant growth promoting ability. Two strains DT-89 and DT-113 were isolated from rhizosphere of wheat crop grown at different agricultural sites of Jaisalmer district of Rajasthan state, India in assumption that they will exert drought stress tolerance in wheat crop cultivated in northern Gangetic plain where drought stress prevails continually. The drought tolerance ability present in many bacteria like *Pseudomonas*, *Burkholderia*, *Pantoea* and other genera but *Bacillus* species are predominantly fall in the category of drought tolerance. Multiple reports supported the use of *Bacillus* as promising bacteria as they exhibit high degree of adaptability to diverse environmental stress conditions owing to endospore formation capability. Various strains of *Bacillus* species like *Bacillus subtilis* GB03, *B. amyloliquefaciens* FZB4244, *Bacillus* spp. produce important metabolites thus are being used in biofertilizer and biopesticide formulations. Vardharajulu et al45 reported strains of *Bacillus* namely *B. licheniformis*, *B. amyloliquefaciens*, *B. thuringiensis*, *B.subtilis* and *Paenibacillus favisporus* can alleviate negative response of drought by increasing root adhering soil/root tissue ratio, soil aggregate stability, plant biomass, relative water content, leaf water potential, preventing leaf water loss. Also, *Bacillus* spp. increased sugars, proline, free amino acids, reduced electrolyte leakage as well as decreased antioxidant enzymes like catalase, glutathione peroxidase, ascorbate peroxidase. For example, Moreno-Galvan et al.46 evaluated five *Bacillus*

![Fig. 7. Effect of *Bacillus altitudinis* DT-89 and *Bacillus paramycoides DT-113 on proline content (mg g⁻¹ tissue) in wheat crop (HD-2967) after 45 and 75 days of sowing under non-stressed and drought (water) stressed conditions. Data represent means ± SD, n = 3. Significance levels are represented as *(P<0.05), **(P<0.01), ****(P<0.001), *****(P<0.0001).](image-url)
spp. strains which ameliorated drought stress by increasing plant height, dry biomass and K⁺ and P⁺ uptake in maize plants. Bacillus species are reported to upgrade ability of plants to resist adverse environmental abiotic stresses including drought. Bacillus synergize with other root harboring microbes to promote plant growth, mineral nutrition, and conferring stress tolerance.47

In the present study, Bacillus altitudinis DT-89 and Bacillus paramycoides DT-113 had shown tolerance up to 30% of PEG 6000 indicating their high-level tolerance which is in agreement with Vardharajulu et al.45 who reported that B. amyloliquefaciens, B. subtilis and B. licheniformis could tolerate up to 25% of water stress. In this study, our strains DT-89 and DT-113 have displayed a few PGP traits such as production of siderophore, ammonia, exopolysaccharide (EPS), IAA, GA3 and solubilization of zinc, phosphate and potassium under non-stressed and drought stress conditions. These properties have been reported as markers for biofertilization, biostimulation, and biocontrol by Bacillus species isolated from wheat rhizospheres.42 The potential of PGPB for phosphate solubilization and siderophores production is considered an essential trait to promote its growth and suppress the growth of soil pathogens. Additionally, by inducing the production of phytohormones and other metabolites in plants, PGPB enhances the plant’s capability to thrive under environmental stressors.48 Production of biomolecules by DT-89 and DT-113 strains under drought stress conditions is of special significance because in arid areas they extend these benefits for plant growth promotion under harsh conditions. In this study, most of the tested PGP traits were found to be expressed under osmotic stress conditions too. Traits like siderophore production, exopolysaccharide production, and gibberellic acid production were found to be highest at 30% PEG-6000 in the growth medium. On the other hand, the production of IAA was found to be highest in the presence of 20% PEG-6000 and was non-significantly changed in the presence of 30% PEG-6000. These findings are in contrast to those reported by Bandeppa et al.,15 who found the maximum expression of most PGP traits in the presence of 20% PEG-6000 in the growth media. Additionally, Sandhya et al.,49 reported a decrease in expression of PGP traits upon exposure to water deficit stress when compared to normal growth conditions. However, in the present study, we found a beneficial outcome of osmotic stress on the expression of PGP traits in these two strains.

In the presence of environmental stressors, it has been reported that plant phytohormone production is disturbed thereby affecting plant growth and seed germination. External application of these phytohormones to plants exposed to drought has the capability to enhance plant growth.50 Therefore, production of phytohormones is considered to be an important PGP trait. In this study, the strains were found to produce IAA and GA 3 upon exposure to osmotic stress, therefore having the potential to supplement the endogenous plant hormones and contribute to plant growth under stress conditions. It also has potential to enhance seedling vigor and seed germination upon inoculation of drought stressed plants with these strains.15 Production of exopolysaccharide by the bacteria has also been reported to improve stress resistance in plants by enhancing water retention and mitigating drought stress.51 In the present study, both strains showed the capability of exopolysaccharide production under osmotic stress which is an important PGP trait for promoting drought tolerance in plants.

Plant growth promotion and drought stress tolerance induction in wheat and other crops by inoculation of Bacillus species have been documented by many researchers in the past. For example, Devarajan et al.52 reported induction of stress tolerance in rice by Bacillus endophyticus PB3, B. altitudinis PB46, and B. megaterium PB50. Likewise, Sunar et al.53 isolated the B. altitudinis strain from the rhizosphere region of Sechium edule plant in Darjeeling and demonstrated its root and shoot growth promotion properties under drought conditions. A decline in chlorophyll content and photosynthetic rate under drought stress was reported by inoculation of PGPR Bacillus subtilis HAS31.54 Other reports suggested similar results of decreased chlorophyll content in drought stress55 Alteration in chloroplast structure and its photo-oxidation by oxidative stress in drought affects the photosynthetic productivity.4,56 Proteomics studies on molecular and physiological mechanisms of effect of Bacillus velezensis 5113 on
abiotic stress including drought suggests their role in preserving the plant photosynthetic machinery by maintaining higher contents of photosynthetic pigments. Bacillus amyloliquefaciens 5113 and Azospirillum brasilense NO40 improved drought stress tolerance in wheat, by increasing root growth and lateral root formation. Similarly, Rashid et al. who reported that Bacillus megaterium and B. licheniformis induced drought tolerance in wheat by enhancing relative water content (59%), chlorophyll a, b and carotenoid (260, 174 and 70%), protein content (136%), proline content (117%), fresh weight (35–192%), and dry weight (58–226%) of wheat and decreasing in MDA content (57%). Our results suggest that Bacillus altitudinis DT-89 and Bacillus paramycoides DT-113 have ability to increase shoot and root biomass, chlorophyll, SOD, POD, and proline content in the wheat crop under drought stress to a significant level.

CONCLUSION

In this study, two potential drought-tolerant Bacillus altitudinis DT-89 and Bacillus paramycoides DT-113 were isolated from the rhizosphere of wheat crop from Jaisalmer district of Rajasthan state, India. The two DT-89 and DT-113 strains were found to exhibit tolerance to around 30% PEG-6000 stress under in vitro condition and at the same time they displayed multitudes of traits pertaining to plant growth promotion. In addition, their inoculation improved shoot, root, chlorophyll content, SOD, POD and proline content in wheat plant under drought stress conditions. Of two strains, Bacillus paramycoides DT-113 had stimulated wheat growth maximum by conferring drought stress tolerance. Hence, Bacillus paramycoides DT-113 may be advocated as a bioinoculant for enhancing plant growth and imparting drought stress tolerance to wheat crop under water deficit condition prevailing in many parts of India.

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CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

AUTHORS’ CONTRIBUTION

SKS and NB conceived and designed the experiments. VKY conducted experiments, collected data and prepared first draft of the manuscript. NB and SKS revised the manuscript and structured the work in proper shape. All the authors revised manuscript and approved final version for submission.

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DATA AVAILABILITY

All datasets generated or analyzed during this study are included in the manuscript.

ETHICS STATEMENT

Not Applicable.

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