



# Significance of inoculation with *Bacillus subtilis* to alleviate drought stress in wheat (*Triticum aestivum* L.)

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## Abstract

In the present study, ten morphologically distinct indigenous plant growth-promoting rhizobacteria (PGPR) from wheat roots and rhizosphere were screened for multiple plant growth promoting (PGP) traits and out of ten, two strains (SIR1 and KUR2) possessing maximum PGP traits along with ACC (1-aminocyclopropane-1-carboxylic acid) deaminase activity were evaluated at three water stress regimes in wheat: (i) 80% field capacity (FC), (ii) 60% FC, (iii) 40% FC for 45 days, starting from 15 days after sowing to the maturity. Inoculation of SIR1 strain with wheat at 80% FC, induced a significant increase in plant biomass (root biomass, 66.67%; shoot biomass, 39.09%) together with decreased reactive oxygen species and increased activity of antioxidant enzymes (superoxide dismutase, peroxidase and catalase) over uninoculated control at 80% FC. 16s rDNA analysis of SIR1 strain revealed its lineage to *Bacillus subtilis*. Present investigations demonstrated the potential of bacterial partner (*B. subtilis*) in alleviating drought stress in wheat.

**Keywords** Abiotic stress · Biofertilizers · Drought · Wheat

## Introduction

The main problem hampering the crop growth and productivity is drought, which is the result of global climate change events and is estimated to reduce cereal productivity by 9–10% (Lesk et al. 2016). More than 50% of the cultivable lands by 2050 are going to face devastating drought consequences on plant growth (Vinocur and Altman 2005). Plant and water relationships are affected by drought stress at both cellular and whole plant levels, resulting in various physiological complex processes and phenotypical responses in plant. Oxidative stress is generated within sub cellular compartments due to alleviated levels of reactive oxygen species (ROS). ROS consist of superoxide radical, hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), and the hydroxyl radical, all of these affect building components of the cell (lipids, proteins, carbohydrates,

nucleic acids etc.) and cause cell demise (Naseem and Bano 2014; Mittler 2002). Therefore, there is an increased interest among the scientists in finding solutions to water associated problems such as drought and its impacts on food security. Particularly, there is an utmost need to redress sustainable solutions, which will improve drought tolerance in crop plants, so as to satisfy the food requirement with the limited water resources in today's world (Mancosu et al. 2015).

Crop productivity can be increased by inoculating plants with PGPR (Plant Growth Promoting Rhizobacteria) facing drought stress (Ngumbi and Kloepper 2016). Alleviation of drought stress by PGPR is mediated by alteration of some physiological and biochemical processes (phytohormone levels, antioxidant enzyme activities and organic solutes content). A key trait of PGPR mediated drought stress alleviation is the ability to regulate ethylene formation using the ACC deaminase enzyme, thus acting as a sink of ACC. PGPR hydrolyze the ACC exuded from the roots in the rhizosphere into ammonia and  $\alpha$ -ketobutyrate, and stimulate the extrusion of ACC from the roots to the soil. The lowering of ACC concentration in root tissues reduces the formation of endogenous ethylene, thus promoting plant growth. Plant resistance to drought is enhanced by reducing ethylene-mediated inhibitory effects on plant growth (Saleem et al. 2007; Chandra et al. 2018; Kaushal and Wani 2016;

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Mishra et al. 2017). Biofilms formation is another important mechanism of bacterial drought tolerance and bacteria mediated plant drought tolerance, especially in *Bacillus subtilis*. Variety of macromolecules (oligo- and polysaccharides) in the form of extracellular matrix, released by bacteria helps in plant growth and development by retaining the moisture, thereby improving water availability in root medium (Timusk et al. 2014; Yang et al. 2009; Dimkpa et al. 2009).

The current study is focused on isolation, screening and characterization (16s rDNA) of potent ACC deaminase producers from High hills (dry and wet temperate zones) of North Western Himalayas and their effect on drought stress alleviation in wheat (variety HPW- 42).

## Materials and methods

### Collection and isolation of potential strains

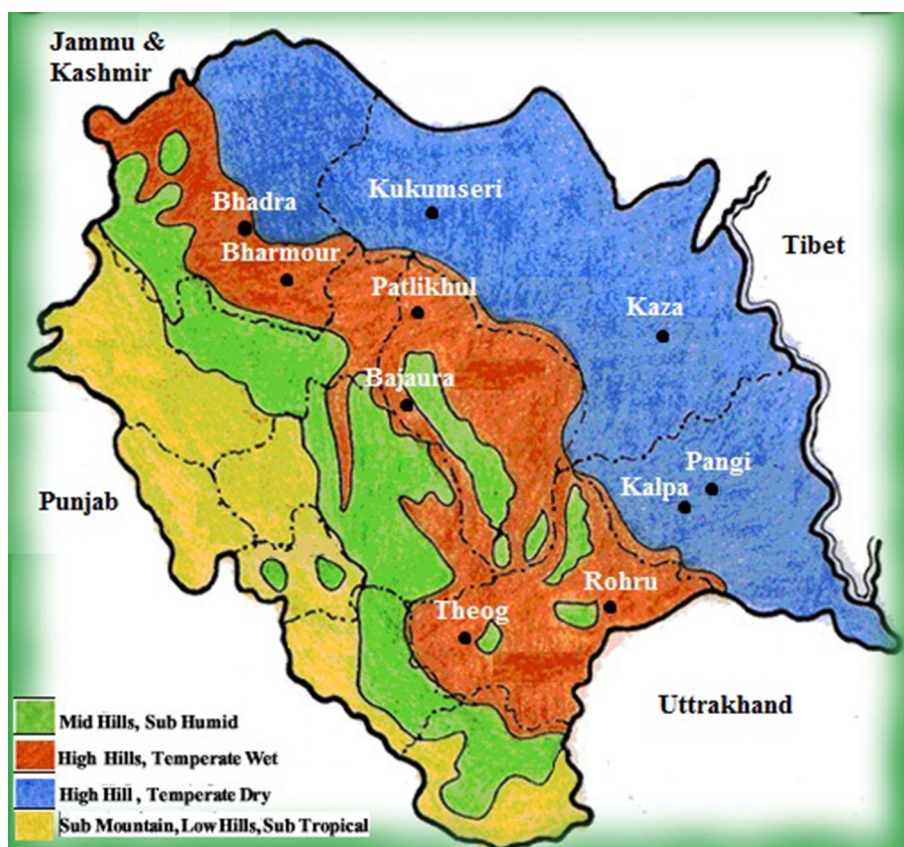
The rhizospheric soil and roots samples of wheat (*Triticum aestivum* L.) plants were collected from high hills (dry and wet temperate zones; Fig. 1) of Himachal Pradesh falling in North Western Himalayan region. Standard serial dilution and pour plate techniques were used for isolation of rhizospheric microbes on nutrient agar, Pikovskaya's and Jensen's medium. Endophytic microorganisms were isolated by

removing the soil particles from wheat roots with tap water, then roots surface was sterilized using sodium hypochlorite (2%) solution. After surface sterilization is complete, wheat roots were crushed in sterile water. The extracts were centrifuged and supernatant was collected. Standard serial dilution and pour plating techniques were used for isolation of endophytic bacteria on nutrient agar, Pikovskaya's and Jensen's medium.

### Molecular (16s rDNA) Identification of best bacterial strain

Genomic DNA was extracted by the phenol/chloroform extraction method (Sambrook et al. 1989). The bacterial strain SIR1 was grown in nutrient broth at 32 °C and cells were harvested after 72 h of incubation and processed for DNA isolation by Real Genome DNA isolation kit. The PCR amplification was carried out in a final volume of 20 µl. The amplification reaction containing 50 ng of DNA template, 20 p mole each of Universal primers, 0.2 mM dNTPs and 1 U Taq Polymerase in 1X PCR buffer. Reaction was cycled 35 times at 94 °C for 30 s, 58 °C for 30 s, 72 °C for 1 min 30 s followed by final extension at 72 °C for 10 min. Amplified PCR product was separated by gel electrophoresis on 1.2% (w/v) agarose gel. Excision of 1375 bp band from the gel after electrophoresis was done by using gel extraction kit

**Fig. 1** Sampling sites for isolations of bacterial strains falling under High hills (Dry and wet temperate zones) of Himachal Pradesh



(Real genomics). Segments of purified DNA were sequenced from commercial sequencing facility (Xleris lab). 16s rDNA gene sequence was compared via NCBI databases by using BLAST algorithm. On the Basis of maximum similarity first 10 sequences were selected and aligned using multiple alignment software program CLUSTALW. Distance matrix was generated using RDP database and the phylogenetic tree was constructed using MEGA 6.0.

### Plant growth promoting traits

Phosphate solubilization activity of the selected strains was estimated using Pikovskaya (1948) medium. The production of indole-3-acetic acid (IAA) and siderophores were also assessed (Aneja 2010). Ammonia production was also assessed (Aneja 2010). Ability of strains to produce ACC deaminase was assessed by spotting bacterial strains on DF minimal salt medium as amended with 3 mM ACC as sole nitrogen source and micronutrient solution (10 ml) (Dworkin and Foster 1958). The biocontrol potential of the bacterial strains against test fungal pathogens (*Fusarium graminearum*, *Claviceps purpurea* and *Alternaria triticina*) was ascertained by agar streak plate method on potato dextrose agar medium and per cent growth inhibition was determined as described (Vincent 1947). Out of total ten strains only two endophytic strains (SIR1 and KUR2) possessing maximum quantitative and qualitative PGP traits were further selected for net house studies to test the physiological efficacy of wheat under drought stress. All the PGP activities were done in replications (each treatment replicated thrice) and the data recorded was analyzed using completely randomized design (CRD).

### Efficacy of bacterial strains on physiological growth parameters, nutrient uptake and antioxidant enzyme activities

Effect of isolated PGPR for physiological efficacy under water stress conditions on growth of wheat variety HPW-42, was studied. The potting mixture was prepared by mixing sand, soil and farm yard manure (FYM) in the ratio of 1:2:1, sterilized by autoclaving the mixture at 80–100 °C for 30 min and then incubated for 24 h. The procedure was repeated for three successive days (also known as Tyndallization). The pH level, electrical conductivity and organic carbon content of the potting mixture were found to be 6.1, 0.39 dS/m and 0.40%, respectively. Available nutrient (N, P and K) contents of experimental soil before the experiment were recorded to be 291.5, 21.4 and 245.9 kg/ha, respectively. The soil used for pot (15 cm diameter and 15 cm deep) experiment belongs to Entisols order as per USDA (United States Department of Agriculture)

soil taxonomy. The soil texture is sandy loam (28% silt, 17% clay and 55% sand). The maximum water holding capacity (MWHC) and field capacity (1/3 bar tension) of the soil were determined using Keen–Raczowski box and pressure plate apparatus, respectively. The MWHC and the field capacity of the soil were recorded as 42 and 23%, respectively. The pots were filled with oven dried (to ensure that initial moisture content is 0%) and sterilized soil mixture weighing about 1500 g/pot. 80%, 60% and 40% of FC were initially maintained by adding 276, 207 and 138 ml of sterilized water on the basis of calculations made for determination of FC. Subsequently, the soil moisture levels were maintained by gravimetric method. The pot surface was covered by sterile glass beads to check evaporation. The study was set up as a Completely Randomized Block Design (CRD) with three replicates for each treatment at three water stress regimes; 80%, 60% and 40% (water depletion of field capacity) and two PGPR strains (SIR1 and KUR2) with controls (without PGPR inoculation). All the pots were kept to the field capacity up to 45 days after planting, starting 15 days after sowing to maturity. Surface sterilized wheat seeds were dipped in individual culture broth (Nutrient broth medium) of selected strains (SIR1 and KUR2) with cell density about  $10^8$  cells/ml for 1 h. The control seeds were treated with sterilized nutrient broth. Five plants per pot were maintained under net house conditions by taking the following nine treatments: T1: 80% of field capacity; T2: 80% of field capacity + SIR1; T3: 80% of field capacity + KUR2; T4: 60% of field capacity; T5: 60% of field capacity + SIR1; T6: 60% of field capacity + KUR2; T7: 40% of field capacity; T8: 40% of field capacity + SIR1; T9: 40% of field capacity + KUR2 in Completely Randomized Block Design (CRD) with 3 replications.

The observations on root/shoot length and biomass were recorded following standard methods. The total N contents in plant samples were analyzed by the microkjeldhal's method (Helrich 1990). For P and K contents, leaf samples were digested in a diacid mixture of HNO<sub>3</sub>: HClO<sub>4</sub> (4:1) and final volume was made to 100 ml (Jackson 1973). Phosphorous percentage in the digested sample was determined by vanado-molybdate yellow colour method (Jackson 1973). Potassium contents in the digest were determined using flame photometer (Jackson 1967). Nutrient uptake (mg/plant) was calculated by multiplying NPK concentrations of the plant with total dry matter content. Total chlorophyll contents and relative leaf water contents were determined (Witham et al. 1971). Total amino acid content was estimated using Folin & ciocalteu's phenol reagent (Folin and Ciocalteu's phenol reagent, Hi-LR™, HiMedia Laboratories Pvt. Ltd., 23, Vadhani Ind. Est., LBS Marg, Mumbai, India, Cat. No. RM10822-100 ml) (Lowry et al. 1951).

## Antioxidant enzyme assay

Crude enzyme extract was prepared by homogenizing 0.5 g of frozen leaf and roots tissues in extraction buffer consisting of 0.5% Triton X-100 and 1% PVP (polyvinyl pyrrolidone) in 100 mM potassium phosphate buffer (pH 7.0) in a chilled pestle mortar. After homogenization, centrifugation was done at 10,000 rpm for 20 min and supernatant was used for antioxidant enzymes assays as given below:

### Superoxide dismutase

Superoxide dismutase activity was assayed by measuring its ability to inhibit the photochemical reduction of nitroblue tetrazolium (NBT) at 560 nm (Beauchamp and Fridovich 1971). The enzyme activity was expressed in U/g fresh weight.

### Catalase activity

Total CAT activity was assayed spectrophotometrically by monitoring decrease in absorbance at 240 nm per minute as a result of decomposition of  $H_2O_2$  (Chandlee and Scandalios 1984). The enzyme activity was expressed in U/g fresh weight.

### Peroxidase activity

Peroxidase activity was determined by measuring the quantity of purpurogallin formed spectrophotometrically at 420 nm (Kumar and Khan 1982). The enzyme activity was expressed in U/g fresh weight.

## Statistical analysis

Data obtained from the net house and laboratory experiments was analyzed using completely randomized design (CRD) with three replications. Statistical analyses of data were done by using SPSS version 16 (SPSS Inc., Chicago, IL, USA) and Microsoft Excel (Microsoft, Redmond, WA, USA) at  $P=0.05$  level of significance.

## Results and discussion

### Collection and isolation of potential strains

Marked variations were observed in both endophytic and rhizospheric bacterial count associated with wheat falling under high hills (Fig. 1) of Himachal Pradesh. On purification, out of total 71 bacterial strains only 10 strains (KIR2, SIR1, KIS2, LSR1, KUS3, KUR2, CHR1, CHS2, SHR1 and SHS1) were able to form clear halo zone on the Pikovskaya's

(PVK) media and showed growth on Jensen's medium. Out of ten strains only three (KIS2, KUS3 and CHS2) were rhizospheric in origin while the rest were endophytic in origin, all of which were further screened for other quantitative PGP traits and ACC deaminase activity. The variation in microbial population in the rhizosphere and roots may be attributed to the age of plant, cultivar type, sampling time, physico-chemical properties of soil, positive influence of root exudates that act as regulators of microbial growth and environmental conditions of study area (Sood et al. 2018a, b, 2019).

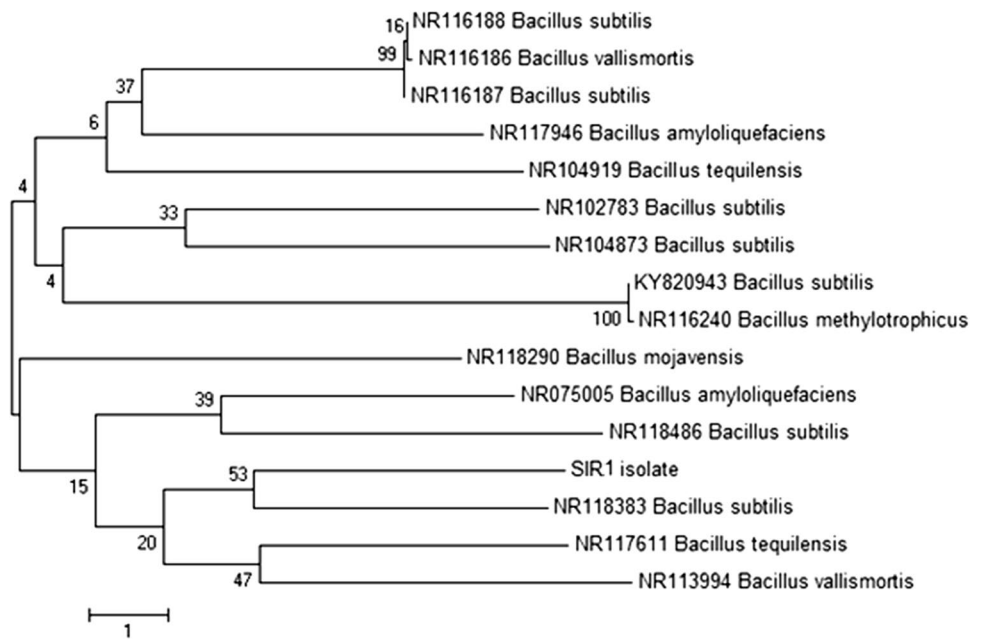
### Molecular (16s rDNA) Identification

Identification of the best performing strain (SIR1) under net house conditions was confirmed by 16s rDNA gene sequencing. Phylogenetic tree generated by the neighbour-joining method for the SIR1 is shown in Fig. 2. Based on the nucleotide homology and phylogenetic tree generated, SIR1 strain was found to be *Bacillus subtilis* (GenBank Accession Number: KX379531) by securing 99% homology with *Bacillus subtilis* strain SBMP4 (GenBank Accession Number: NR118383). Similar molecular characteristics for the member of genus *Bacillus* were reported (Kumar et al. 2015; Wahyudi et al. 2011; Majeed et al. 2015). Further, it has also been demonstrated that bacteria belonging to genus *Bacillus*, not only proliferate rhizosphere, but also reside inside the root tissue of crop plants (Hallmann et al. 1997).

### Plant growth promoting traits

Microorganisms possessing multifarious PGP traits are beneficial for enhancing plant growth under abiotic stress. Significant variations were observed in the plant growth promoting activities of the isolated PGPR (Table 1). The phosphate solubilization ranged from 93.00 to 190.00  $\mu\text{g/ml}$  in liquid PVK medium. The maximum (190.00  $\mu\text{g/ml}$ ) P-solubilisation was noted for SIR1 strain which was at par with KUR2 (181.00  $\mu\text{g/ml}$ ) whereas the minimum (93.00  $\mu\text{g/ml}$ ) was recorded for SHS1. Possible mechanisms adopted by the strains for phosphate solubilisation includes different array of solubilization reactions, which include acidification, chelation, exchange reactions, and production of gluconic acid, to release monobasic and dibasic forms of phosphorus from insoluble phosphorus (Ngumbi and Klopper 2016; Sood et al. 2018a, b, 2019), which might have resulted in the increased available P content, thus enhancing the growth of plants under drought stress. The indole acetic acid (IAA) and siderophore production ranged from 17.33 to 29.67  $\mu\text{g/ml}$  and 25.00 to 57.00%, respectively. The maximum IAA (29.67  $\mu\text{g/ml}$ ) and siderophore production (57.00%) was shown by SIR1 strain, followed by KUR2. Auxin, also referred to as indole-3-acetic acid (IAA), is an

**Fig. 2** Neighbour-joining tree based on 16s rDNA gene sequence data of *Bacillus subtilis* (SIR1) and various species of genus *Bacillus*



important regulator of plant growth and development (Glick et al. 2007; Glick 1995). Recently, Chandra et al. (2018) have reported the effect of bacterial IAA on root growth for enhanced uptake of nutrients, thereby, anchoring tall sugarcane plants with soil under drought stress. Similar studies on the effect of bacterial IAA on plant growth under abiotic stress have also been conducted in wheat, maize, pepper and *Arabidopsis thaliana* etc. (Sorty et al. 2016; Vardharajula et al. 2011; Lim and Kim 2013; Cohen et al. 2015). The plants facing stressed conditions are more susceptible to phytopathogens. The possible solution to combat this susceptibility is to provide defense system against these phytopathogens by using the strains that were efficient producers of bacterial siderophores. This can be achieved either directly by supplying iron to plants or indirectly by depriving the fungal pathogens of iron, thereby limiting their growth (Glick 2012; Bagyaraj 2011). All the strains showed ammonia production and antagonism against *Fusarium graminearum*, *Alternaria trititica* and *Claviceps purpurea*. Ammonia produced by bacterial strains, suppresses the growth of certain fungi due to its potent antagonistic effect, which complemented the biocontrol activity of the tested strains. Only three strains namely KIR2, LSR1 and SHR1 were not found to be ACC deaminase producers. The active ACC deaminase enzyme of bacterial origin might have played a significant role in the reduction of ACC levels in roots, as reported in previous research (Glick 2014). Onset of stress signals triggers the enhanced production of ethylene (ET) and 1-Aminocyclopropane-1-carboxylate (ACC) during the initial phase of stress induction. Bacteria exert beneficial effects on plants under abiotic stress by regulating the ET level (Saleem et al. 2018). PGPR possessing ACC

deaminase activity results in hydrolysis of ACC into ammonia and alpha-ketobutyrate. As a result, the levels of ethylene get reduced and plants are able to maintain normal growth (Glick and Bashan 1998). The results are in line with the previous findings (Khan et al. 2016). Similarly, Saikia et al. (2018) reported consortium effect of three ACC-deaminase producing rhizobacteria (*Ochrobactrum pseudogrignonense* RJ12, *Pseudomonas* sp. RJ15 and *Bacillus subtilis* RJ46) in enhancing the crop physiological parameters of *Vigna mungo* L. and *Pisum sativum* L. under drought stress. On the basis of PGPR traits and antifungal activity, SIR1 and KUR2 were found to possess the maximum PGP traits and were thus selected for net house studies.

### Efficacy of bacterial strains on physiological growth parameters, antioxidant enzyme activities and nutrient uptake

The adverse effect of drought stress was noticed in uninoculated water stress (drought) imposed wheat plants with reduced growth, relative water content and less chlorophyll content (Table 2). However, significant ( $p=0.05$ ) plant growth and development was noticed in bacteria-inoculated plants under drought stress (Table 2). The action of SIR1 strain was found to be more promising than the action of KUR2. At 80% FC soil moisture level, plants receiving SIR1 inoculation increased the shoot and root length of wheat significantly ( $p=0.05$ ) by 14.22% and 23.51% over uninoculated control maintained at same stress level (80% FC). However, these values in KUR2 inoculated plants were 8.10% and 4.60% for shoot and root length, respectively. Similar trend was observed in SIR1 and KUR2 inoculated

**Table 1** Screening of bacterial isolates for plant growth promoting activities

Isolates	Agro climatic zones	P solubi- lization in liquid medium (µg/ ml)	IAA (µg/ml)	SPE (%)	ACC- deaminase activity	Ammonia produc- tion	% Antagonism against:		
							<i>F. grame- narum</i>	<i>C. purpurea</i>	<i>A. triticini</i>
KIR2*	Dry Temper- ate	177.67 <sup>bc*</sup>	21.67 <sup>bcde</sup>	40.00 (39.21) <sup>bcd</sup>	–	+++	38.00 (38.03) <sup>abc</sup>	41.00 (39.8) <sup>abc</sup>	38.33 (38.23) <sup>abc</sup>
SIR1*	High Hills	190.00 <sup>a</sup>	29.67 <sup>a</sup>	57.00 (51.00) <sup>a</sup>	+	+++	43.11 (39.00) <sup>a</sup>	55.50 (42.10) <sup>a</sup>	42.87 (40.88) <sup>a</sup>
KIS2	Dry Temper- ate	161.00 <sup>d</sup>	20.00 <sup>bcdefg</sup>	25.00 (29.94) <sup>ghij</sup>	+	+	25.33 (30.15) <sup>fgh</sup>	30.33 (33.39) <sup>e</sup>	23.67 (29.07) <sup>ghij</sup>
LSR1*	Dry Temper- ate	96.67 <sup>hi</sup>	21.00 <sup>bcdef</sup>	42.67 (40.76) <sup>bc</sup>	–	+	30 (33.20) <sup>def</sup>	22.67 (28.40) <sup>g</sup>	24.33 (29.38) <sup>fghi</sup>
KUS3	High Hills	130.67 <sup>g</sup>	23.00 <sup>bcd</sup>	33.00 (35.04) <sup>f</sup>	+	+++	27.33 (31.49) <sup>defg</sup>	29.67 (32.98) <sup>ef</sup>	29.33 (32.77) <sup>def</sup>
KUR2*	High Hills	181.00 <sup>ab</sup>	23.67 <sup>b</sup>	44.67 (41.92) <sup>b</sup>	+	+++	39.00 (38.61) <sup>ab</sup>	43.33 (41.15) <sup>ab</sup>	40.00 (39.21) <sup>ab</sup>
CHR1*	High Hills	149.67 <sup>ef</sup>	18.67 <sup>efghi</sup>	30.67 (33.60) <sup>fg</sup>	+	+	22.67 (28.34) <sup>ghi</sup>	19.33 (26.05) <sup>ghi</sup>	34.67 (36.04) <sup>bcd</sup>
CHS2	High Hills	154.00 <sup>de</sup>	17.33 <sup>fghij</sup>	39.33 (38.81) <sup>bcde</sup>	+	+	31.00 (33.81) <sup>de</sup>	37.00 (37.43) <sup>cd</sup>	34.67 (35.97) <sup>bcde</sup>
SHR1*	High Hills	97.67 <sup>h</sup>	19.33 <sup>cdefgh</sup>	26.00 (30.59) <sup>gh</sup>	–	++	31.67 (34.22) <sup>d</sup>	19.00 (25.83) <sup>ghij</sup>	26.67 (31.03) <sup>fgh</sup>
SHS1*	High Hills	93.00 <sup>hij</sup>	23.33 <sup>bc</sup>	25.67 (30.38) <sup>ghi</sup>	+	+	31.00 (33.81) <sup>de</sup>	21.33 (27.43) <sup>gh</sup>	29.00 (32.56) <sup>defg</sup>
C. D. (p=0.05)		10.32	4.00	3.73			3.53	3.04	4.80

Values in parentheses are arc sine-transformed

IAA indole-3-acetic acid, SPE siderophore production efficiency. Values in parentheses are arc sine-transformed. Within a column, means followed by the same letter are not significantly different (critical difference (CD) at  $P=0.05$ ). For ACC deaminase activity: +, activity present; –, no activity. For ammonium production: +, light brown colour; ++, dark brown colour; +++, orange brown colour; –, no activity. \*: isolates endophytic in origin

For ACC deaminase activity: + activity present; – no activity. For ammonium production: + light brown colour; ++ dark brown colour; +++ orange brown colour; – no activity. \*: isolates endophytic in origin

Within a column, means followed by the same letter are not significantly different (critical difference (CD) at  $P=0.05$ )

treatments maintained at 60 and 40% FC soil moisture levels in contrast to uninoculated controls at respective stress levels (60 and 40% FC). This is also confirmed by significant increase in shoot and root biomass (dry weight) compared to non-inoculated plants. Compared to uninoculated control plants at 80% FC soil moisture, 39.09% and 66.00% increase in shoot and root biomass was observed in SIR1-treated stressed plants at 80% FC. However, this rate for KUR2 inoculated stressed plants was 3.64% and 11.00%, respectively (Table 2). 33.33%, 15.56% increase in shoot biomass and 66.67%, 50.00% in root biomass was observed in the treatments receiving SIR1 and KUR2 inoculation at 60% FC compared to their uninoculated control stressed plants. This increase for plants inoculated with SIR1 and KUR2 strains at 40% FC was noted as 26.67%, 16.67% (shoot biomass) and 40.00%, 20.00% (root biomass), respectively. The possible reason for enhanced shoot/root length and their biomass in the inoculated treatments is the release

of variety of plant growth regulators i.e. phytohormones in presence of different plant flavonoids in the rhizospheric region, which results in alteration of root architecture that may lead to an increase in the total root surface area, and consequently improved water and nutrient uptake, particularly N (asymbiotic N-fixation) and P (P-solubilization), with positive effects on plant growth as a whole (Montano et al. 2014). For example, Timmusk et al. (2014) reported that wheat plants treated with PGPR had 78% higher biomass than non-treated plants under drought stress, exhibiting the potential of PGPR to enhance plant performance under stress conditions. Significant ( $p=0.05$ ) increase of 2.22%, 6.17%, 4.23% and 1.11%, 2.06%, 2.82% in RWC (relative water content) was observed in SIR1 and KUR2 inoculated stress (80, 60 and 40% FC) imposed wheat plants, as compared to the uninoculated stressed plants at respective stress levels. Bacterial inoculation had a direct effect on plant osmolytes (Table 2). Treatments receiving SIR1

**Table 2** Effect of liquid bacterial inocula on physiological parameters of wheat under water stress

Treatments	Shoot length (cm)	Root length (cm)	Root biomass (g/plant)	Shoot biomass (g/plant)	Relative water content (%)	Total amino acid content (mg/g FW)	Total chlorophyll content (mg/g FW)
T1 (80% of field capacity)	52.53 <sup>c</sup>	9.78 <sup>cd</sup>	0.09 <sup>bcdef</sup>	1.10 <sup>bc</sup>	90.00 (9.54) <sup>c</sup>	0.32 <sup>gh</sup>	0.18 <sup>abc</sup>
T2 (80% of field capacity + SIR1)	60.00 <sup>a</sup>	12.08 <sup>a</sup>	0.15 <sup>a</sup>	1.53 <sup>a</sup>	92.00 (9.64) <sup>a</sup>	0.37 <sup>efg</sup>	0.24 <sup>a</sup>
T3 (80% of field capacity + KUR2)	56.82 <sup>b</sup>	10.23 <sup>b</sup>	0.10 <sup>ab</sup>	1.14 <sup>b</sup>	91.00 (9.59) <sup>b</sup>	0.35 <sup>fgh</sup>	0.21 <sup>ab</sup>
T4 (60% of field capacity)	45.27 <sup>ef</sup>	9.17 <sup>f</sup>	0.06 <sup>bcde</sup>	0.45 <sup>def</sup>	81.00 (9.06) <sup>f</sup>	0.40 <sup>def</sup>	0.16 <sup>abcde</sup>
T5 (60% of field capacity + SIR1)	47.40 <sup>d</sup>	9.84 <sup>c</sup>	0.10 <sup>ab</sup>	0.60 <sup>d</sup>	86.00 (9.33) <sup>d</sup>	0.42 <sup>cde</sup>	0.18 <sup>abc</sup>
T6 (60% of field capacity + KUR2)	45.85 <sup>e</sup>	9.63 <sup>cde</sup>	0.09 <sup>bc</sup>	0.52 <sup>de</sup>	82.67 (9.15) <sup>e</sup>	0.44 <sup>bcd</sup>	0.17 <sup>abcd</sup>
T7 (40% of field capacity)	33.75 <sup>i</sup>	6.77 <sup>i</sup>	0.05 <sup>bcdef</sup>	0.30 <sup>fghi</sup>	71.00 (8.49) <sup>i</sup>	0.47 <sup>abc</sup>	0.10 <sup>cdefgh</sup>
T8 (40% of field capacity + SIR1)	38.97 <sup>g</sup>	7.31 <sup>g</sup>	0.07 <sup>bcd</sup>	0.38 <sup>efg</sup>	74.00 (8.66) <sup>g</sup>	0.51 <sup>a</sup>	0.14 <sup>bcdef</sup>
T9 (40% of field capacity + KUR2)	34.88 <sup>h</sup>	7.26 <sup>gh</sup>	0.06 <sup>bcde</sup>	0.35 <sup>efgh</sup>	73.00 (8.62) <sup>h</sup>	0.49 <sup>ab</sup>	0.11 <sup>cdefg</sup>
CD <sub>0.05</sub>	0.87	0.21	0.05	0.19	0.13	0.06	0.08

Figures in parentheses are square root transformed values

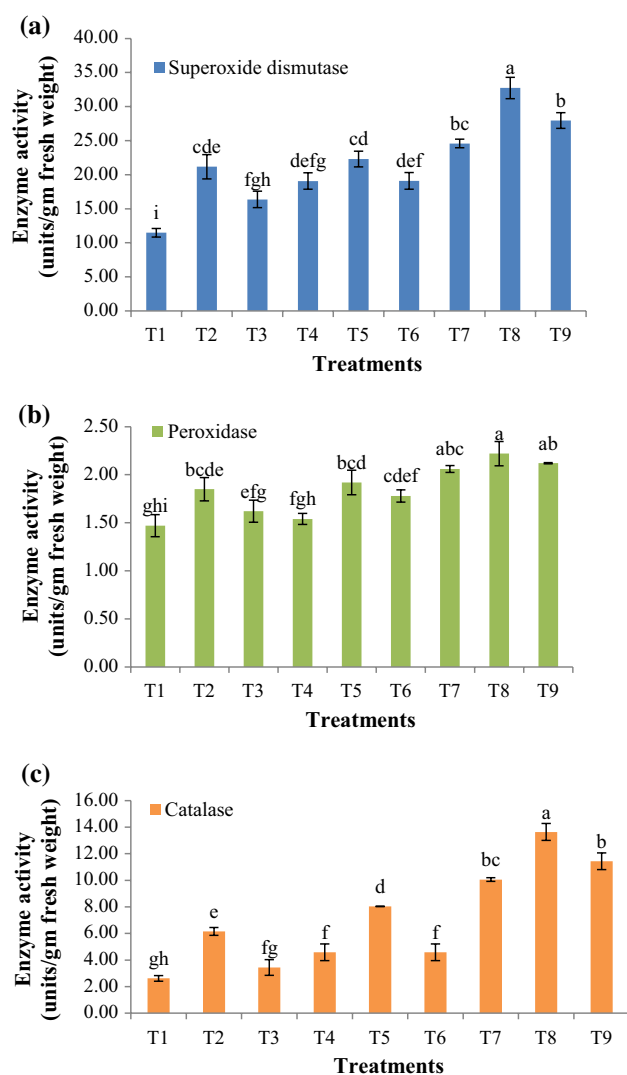
Figures followed by the same superscript within a column are not significantly different but statistically significant over other treatment combinations based on C. D. (critical difference,  $P=0.05$ ); T1: 80% of field capacity; T2: 80% of field capacity + SIR1; T3: 80% of field capacity + KUR2; T4: 60% of field capacity; T5: 60% of field capacity + SIR1; T6: 60% of field capacity + KUR2; T7: 40% of field capacity; T8: 40% of field capacity + SIR1; T9: 40% of field capacity + KUR2; number of replications per treatment ( $n=03$ )

inoculation at respective stress levels (80, 60 and 40% FC) showed significant increase of 15.63%, 5.00% and 8.51% in total amino acid content over their uninoculated stressed plants maintained at 80, 60 and 40% FC, followed by KUR2 with 9.30%, 10% and 4.26% increase. The increased concentration of solutes and osmolytes are key elements of plant response facing abiotic stress. Osmolyte accumulation and cellular osmotic potential are correlated i.e. as the osmolyte concentration increases, there is a decrease in the cellular osmotic potential. This decrease helps in absorption of water from the drying soil and as a result, turgor pressure of the cell gets increased thereby improving the physiological efficacy of plants under drought stress (Harb et al. 2010). Total chlorophyll content was determined to examine the impact of the rhizobacterial strains on the photosynthetic potential of host plants. Plants receiving SIR1 and KUR2 inoculation at 80%, 60% and 40% FC showed significant ( $p=0.05$ ) increase of 33.33%, 12.50%, 40.00% and 16.67%, 6.25%, 10% in total chlorophyll content in comparison to uninoculated stressed plants (negative controls at respective stress levels). This indicates the competence of select strains at upholding the chlorophyll content under drought conditions as reported by Glick (2014).

The activation of a plant's inherent enzymatic and non-enzymatic systems is always crucial for the detoxification of the ROS under stress conditions. The results of this study clearly indicate that treatments receiving SIR1 and KUR2 inoculation ( $p=0.05$ ) stimulates the SOD activity in wheat plants much significantly over uninoculated stressed controls (Fig. 3). An increase of 84.40%, 42.60% at 80% FC, 16.94%, 0.10% at 60% FC and 33.16%, 13.79% at 40% FC

was recorded in SIR1 and KUR2 inoculated plants in comparison to the uninoculated stressed controls maintained at 80%, 60% and 40% FC soil moisture, respectively. POD and CAT were located in every ROS-producing compartment and functioned as a fine regulator of intracellular ROS level. As with SOD, a similar increasing trend was observed in POD and CAT activities, with 134.73%, 31.3% increase at 80% FC, 74.95%, 0.00% at 60% FC and 35.72%, 13.73% at 40% FC in case of CAT and 25.85%, 10.20% at 80% FC, 24.68%, 15.58% at 60% FC and 7.77%, 2.91% at 40% FC in case of POD. Moreover, an increasing trend was also noticed in stress-induced plants without inoculation, but this trend was not as prominent as it was in the SIR1 and KUR2 inoculated plants (Fig. 3). This increase in enzymatic activity may serve to minimize oxidative injury and contribute to the drought stress tolerance by degradation of reactive oxygen species (ROS), such as hydrogen peroxide, singlet oxygen, superoxide radical, and hydroxyl radical (Hasanuzzaman et al. 2014). Our studies are in accordance with Kasim et al. (2013) who reported that priming of wheat seeds with plant-growth promoting bacteria (*Bacillus amyloliquefaciens* 5113 and *Azospirillum brasilense* NO40) significantly alleviated the deleterious effect of drought stress on wheat.

Induction of drought stress hindered the nutrient uptake from soil and a decrease in nutrient (NPK) content of wheat plants was observed with increase in imposed drought stress (Table 3). A decrease of 32.50%, 23.40% in N, 11.36%, 9.30% in P and 18.31%, 15.33% in K content was observed in the stressed plants grown at 80% FC over SIR1 and KUR2 inoculated plants at same water stress regime. Similar decrease was observed in uninoculated



**Fig. 3** Effect of liquid bacterial inoculum on antioxidant enzymes activities **a** superoxide dismutase, **b** peroxidase and **c** catalase. T1: 80% of field capacity; T2: 80% of field capacity + SIR1; T3: 80% of field capacity + KUR2; T4: 60% of field capacity; T5: 60% of field capacity + SIR1; T6: 60% of field capacity + KUR2; T7: 40% of field capacity; T8: 40% of field capacity + SIR1; T9: 40% of field capacity + KUR2

treatments over SIR1 and KUR2 inoculated ones, maintained at 60 and 40% FC soil moisture levels [18.75%, 17.20% (N), 29.41%, 27.27% (P) and 15.04%, 5.83% (K) at 60% FC; 5.63%, 4.29% (N), 20.51%, 18.42% (P) and 12.21%, 8.00% (K) at 40% FC]. Drought stress also

significantly reduced nutrient uptake in the uninoculated plants over the inoculated ones. This decrease in nutrient uptake was 53.76%, 29.53% for N uptake, 39.54%, 16.33% (P uptake) and 42.27%, 21.97% (K uptake) in the uninoculated stressed plants grown at 80% FC as compared to the ones that received SIR1 and KUR2 inoculation. Similar trend for nutrient uptake was observed in uninoculated treatments maintained at 60% and 40% FC soil moisture levels over plants receiving SIR1 and KUR2 inoculation (40.37%, 30.17% (N uptake), 28.87%, 16.44% (P uptake) and 37.58%, 21.15% (K uptake) at 60% FC; 26.13%, 18.48% (N uptake), 38.07%, 29.68% (P uptake) and 31.62%, 22.33% (K uptake) at 40% FC). The significant increase in nutrient (NPK) content in the PGPR inoculated treatment as compared to uninoculated controls might be attributed to the increase in different fractions of available N, P and K in soil, stimulation of root formation or increase in plant growth stimulating mechanisms (direct mechanisms), which can positively influence plant growth and crop yields. Similarly, the plant growth promoting effect of strains favored the wheat plants accumulating more macronutrients in the biomass. Our results are in accordance with the findings of Kumar et al. (2014) who reported maximum nutrients (macro and micronutrients) acquisition and content in grain of wheat receiving different combinations of microbial strains (*Bacillus megaterium*, *Arthrobacter chlorophenicus* and *Enterobacter* sp.) under pot and field experiments. The results of *in-vivo* experiments revealed tremendous improvements in plant growth promotion of wheat plants under water deficit conditions when the two strains (SIR1 and KUR2) were applied at different levels of water deficit stress as compared to uninoculated treatments (Fig. 4).

## Conclusions

It can be concluded from current study that inoculation of wheat plants with endophyte (*Bacillus subtilis*) significantly alleviated drought stress and consequently improved growth. The application of *Bacillus subtilis* may thus be recommended for field trials in high hills and dry temperate zone of Himachal Pradesh for sustainable crop production. It can further help in improving soil health and conservation of natural resources.



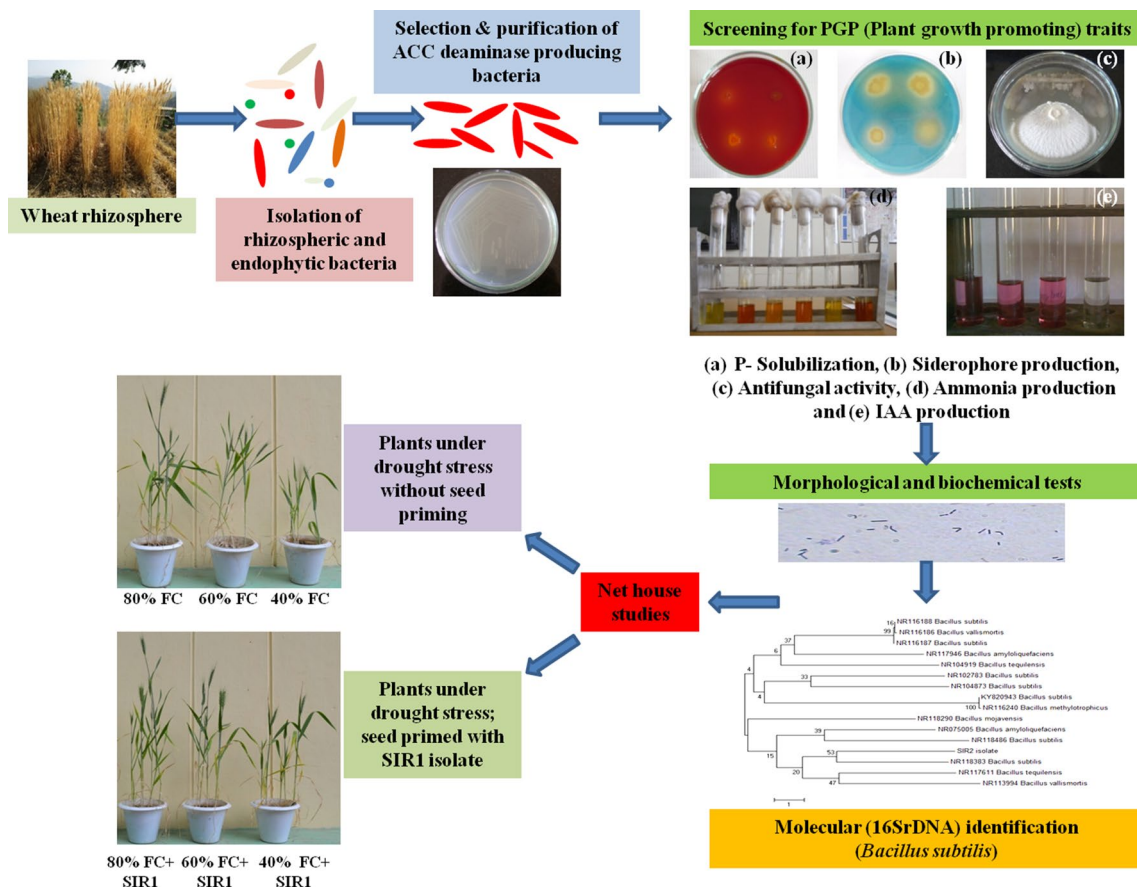


Fig. 4 Graphical representation of the summary

Table 3 Influence of PGP endophytic bacteria isolates on NPK content and uptake of wheat under varied levels of drought stress

Treatments	Nutrient content (%)			Nutrient uptake (mg/plant)		
	N	P	K	N	P	K
T1 (80% of field capacity)	1.08 (1.44) <sup>h</sup>	0.39 (1.18) <sup>c</sup>	1.16 (1.47) <sup>fg</sup>	12.41 <sup>c</sup>	4.51 <sup>bc</sup>	13.32 <sup>c</sup>
T2 (80% of field capacity + SIR1)	1.60 (1.61) <sup>a</sup>	0.44 (1.20) <sup>a</sup>	1.42 (1.56) <sup>a</sup>	26.84 <sup>a</sup>	7.46 <sup>a</sup>	23.90 <sup>a</sup>
T3 (80% of field capacity + KUR2)	1.41 (1.55) <sup>cd</sup>	0.43 (1.20) <sup>ab</sup>	1.37 (1.54) <sup>b</sup>	17.61 <sup>b</sup>	5.39 <sup>b</sup>	17.07 <sup>b</sup>
T4 (60% of field capacity)	1.30 (1.52) <sup>fg</sup>	0.24 (1.11) <sup>h</sup>	1.13 (1.46) <sup>hi</sup>	6.69 <sup>ef</sup>	1.70 <sup>def</sup>	5.78 <sup>efg</sup>
T5 (60% of field capacity + SIR1)	1.60 (1.61) <sup>a</sup>	0.34 (1.16) <sup>e</sup>	1.33 (1.53) <sup>bc</sup>	11.22 <sup>cd</sup>	2.39 <sup>d</sup>	9.26 <sup>d</sup>
T6 (60% of field capacity + KUR2)	1.57 (1.60) <sup>ab</sup>	0.33 (1.15) <sup>ef</sup>	1.20 (1.48) <sup>f</sup>	9.58 <sup>cde</sup>	1.46 <sup>efgh</sup>	7.33 <sup>de</sup>
T7 (40% of field capacity)	1.34 (1.53) <sup>f</sup>	0.31 (1.15) <sup>fg</sup>	1.15 (1.47) <sup>gh</sup>	4.73 <sup>fg</sup>	1.09 <sup>efghi</sup>	4.00 <sup>fg</sup>
T8 (40% of field capacity + SIR1)	1.42 (1.55) <sup>c</sup>	0.39 (1.18) <sup>c</sup>	1.31 (1.52) <sup>cd</sup>	6.39 <sup>efg</sup>	1.76 <sup>de</sup>	5.85 <sup>ef</sup>
T9 (40% of field capacity + KUR2)	1.40 (1.55) <sup>cde</sup>	0.38 (1.17) <sup>cd</sup>	1.25 (1.50) <sup>e</sup>	5.79 <sup>efgh</sup>	1.55 <sup>efg</sup>	5.15 <sup>efgh</sup>
CD <sub>0.05</sub>	0.05	0.02	0.04	3.81	0.92	2.91

Within columns, means followed by same letter are not significantly different [critical difference (CD) at  $P=0.05$ ] T1: 80% of field capacity; T2: 80% of field capacity + SIR1; T3: 80% of field capacity + KUR2; T4: 60% of field capacity; T5: 60% of field capacity + SIR1; T6: 60% of field capacity + KUR2; T7: 40% of field capacity; T8: 40% of field capacity + SIR1; T9: 40% of field capacity + KUR2; number of replications per treatment (n=03); number of repeats (n=03)

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## Compliance with ethical standards

**Conflict of Interest** The authors declare that there are no conflicts of interest.

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