



Potentials of Lactic Acid Bacteria in Enhancing Nodulation of *Bradyrhizobium daqingense* and Yield in Soybean

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ABSTRACT

Background: The ferments of lactic acid bacteria (LAB) are used since decades in agricultural practice to control diseases, to promote plant growth and also to improve soils. However the functional roles of LAB in phytomicrobiome need to be discovered, which would result in understanding of the symbiotic relationship between LAB and plants and that could be exploited to improve agricultural production.

Methods: In this study, the scientific investigation was carried out for pot culture evaluation of six efficient LAB isolates from soybean rhizosphere, on nodulation and yield of soybean in green house condition, which were proven positive for IAA and GA production and PGPR traits.

Result: Among the eight different treatment combination with *Bradyrhizobium daqingense* the treatment which received consortium of all six LAB isolates had significant impact on plant growth characters viz. plant height, root length, number of branches and chlorophyll content at 30 and 60 DAS. The LAB consortia also showed significantly high nodule number (47.67), nodule dry weight (117 mg plant⁻¹) and leg haemoglobin content (6.27 mg g⁻¹ fresh nodule) at 30 DAS. The yield and yield related traits was also highest in consortium treated plants. The property of plant to produce more nodules and healthy root growth can be attributed to IAA producing ability of inoculated LAB isolates.

Key words: Co-inoculation, Lactic acid bacteria, Nodulation.

INTRODUCTION

The plant growth-promoting rhizobacteria (PGPR) that live in rhizosphere occupy approximately 5 to 17% of the total root surface (Gray and Smith, 2005). The most widely studied genera include *Rhizobium*, *Pseudomonas*, *Bacillus* and *Azospirillum*. These microorganisms have beneficial effects on seed germination, seedling emergence and plant growth (Ahmad *et al.* 2008). According to Richardson *et al.* (2009), the interactions between plant roots and soil microbiota play a significant role in plant nutrition. Soil microorganisms promote plant growth through synthesis of plant hormones (auxins and gibberellins), nitrogen fixation, solubilization of inorganic phosphate and mineralization of organic phosphate, thus making these elements available to plants (Rodriguez and Fraga, 1999). These organisms also have beneficial effects on legume growth and some strains enhance nodulation and nitrogen fixation by affecting interactions between plant and rhizobia (Parmar and Dadarwal, 1999). With this background scientific study in improving nodulation efficiency of symbiotic *Bradyrhizobium* sp. in legumes by co-inoculation was carried to combat excess use of chemical N fertilizers, for cost effective cultivation. The above mentioned context of improved nodulation is favoured by production of flavonoid like compounds or by stimulating the host legume to produce flavonoid signal molecule (Pankaj *et al.*, 2009). As per the scientific evidences one of the important phytohormone that is IAA has positive significant influence over lateral root formation and nodulation when produced at lower doses. This is proven in case of *Arabidopsis thaliana* (Overvoorde *et al.*, 2010).

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Studies have also shown that many rhizosphere microorganisms are capable of synthesizing plant growth regulators *in vitro*. Sarwar and Kremer (1995) studied microorganisms in the rhizospheres of different plants and found that root-associated isolates were more efficient in terms of auxin production than non-associated isolates. It is estimated that 80% of bacteria isolated from the rhizosphere are able to produce IAA (Pereira *et al.* 2012).

The LAB represent a group of gram positive genera like *Lactobacillus*, *Lactococci*, *Leuconostoc*, *Pediococcus* and *Streptococcus*. They are facultative anaerobic bacteria, which reside on substrates that are rich in carbohydrates and ferment them into organic acids. As per the scientific evidences, lactic acid bacteria (LAB) are being extensively

used in food industries in preparing fermented food products. The property of producing organic acids and antimicrobial compounds by LAB is used in bio preservation of food products in maintaining quality product. Studies related to application of LAB in agriculture are relatively less; however, *in vitro* studies related to growth stimulation and antimicrobial studies are being carried out.

Similarly in case of Effective microorganisms (EM), which is a commercial biofertilizer that contains a mixture of co-existing beneficial microorganisms, predominantly it consists of species of lactic acid bacteria, (Arshad and Rukhsana, 2010) up on application in mung bean (*Vigna radiata* L.) and soybean (Javaid and Mahmood, 2010), showed a significant increase in nodule number and biomass in legumes (Javaid, 2011).

The characteristic feature of LAB to produce indole acetic acid, plant growth promoting activities and their significant role in consortia gives us a new dimension to work over their efficiency in augmenting nodulation in legumes as co-inoculant.

MATERIAL AND METHODS

The present experiment was conducted at Department of Agricultural Microbiology, UAS, Dharwad and the entire study was conducted under green house condition during the year 2019-2020. From our previous studies, six LAB isolates which showed positive result for IAA and GA production *viz*, AL-44, AL-45, AL-48, AL-49, AL-58 and AL-59 were selected for the present investigation.

Compatibility testing

The compatibility among the above mentioned microbial strains was confirmed by following multi strain streaking method and then used in consortia (Fig 1).

Germination test (roll towel technique)

This was done by following the roll towel technique as per ISTA (Tinnin and Kirkpatrick, 1985). Seeds of soybean were washed with distilled water and used. As many as 400 seeds were used for each treatment. The seeds were dipped in cultures of selected strains of the LAB for 10 min and then placed on the germination papers and rolled.

Pot culture and treatment details

Earthen pots of 30 cm top diameter was filled with 10 kg soil. The soil in each pot was mixed with 0.26g urea, 1.5 g SSP and 0.12g MOP to supply 40:80:25 Kg N:P₂O₅:K₂O per ha on soil weight basis as per the package of practices. Half of the N was applied at the time of sowing and remaining half was applied as top dressing after 30 days of sowing. Soybean cultivar used was Dsb-21 in three replication and eight treatment combination using CRD design (Panse and Sukhatme, 1985). The levels of significance used in the 'F' and 't' test was P=0.01.

- 1) Treatment 1: Absolute control (POP only).
- 2) Treatment 2: POP + Lactic acid bacteria AL-44.
- 3) Treatment 3: POP + Lactic acid bacteria AL-45.

- 4) Treatment 4: POP + Lactic acid bacteria AL-48.
- 5) Treatment 5: POP + Lactic acid bacteria AL-49.
- 6) Treatment 6: POP + Lactic acid bacteria AL-58.
- 7) Treatment 7: POP + Lactic acid bacteria AL-59.
- 8) Treatment 8: POP + Lactic acid bacteria Consortium (AL-44+ AL-45+ AL-48+ AL-49+ AL-58+ AL-59).

POP : NPK @ 16 kg/ac N, 32 kg/ac P₂O₅, 10 kg/ac K₂O₅, ZnSO₄ - 5 kg/ac, sulphur- 8 kg/ ac, *Bradyrhizobium daqingense* - 500 g/30 kg seeds. PSB (*Pseudomonas striata*) - 500 g/30 kg seeds.

Seed treatment

The 100 ml of Mann, De Rogosa and Sharpe broth was inoculated with pure culture of LAB isolates for mass multiplication and kept on shaker for 2 days at 100 rpm, at 32°C and after attaining a population of 10⁸ cfu per ml. This one ml broth culture was mixed with lignite powder at 1:3 ratio, which had *Bradyrhizobium daqingense* used for seed treatment along with gum for effective and uniform seed coating. In case of consortium, one ml of all six isolates containing 10⁸ CFU/ml were properly assorted and then used in carrier material in the above mentioned ratio for seed treatment.

Observations

a) Plant physiological observations were taken at 30 and 60 DAS, number of nodules and their weight at 30 DAS, dry matter accumulation at harvest by placingsamples in hot air oven at 70°C for 48 h, number of branches per plant and Chlorophyll content at 30 and 60 DAS. Later using SPAD (Soil Plant Analysis Development) instrument. The leg

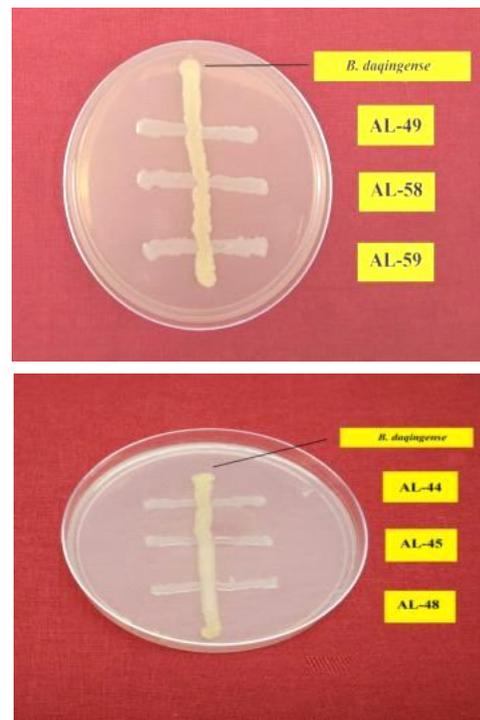


Fig 1: Compatibility test of efficient nodulation stimulating LAB isolates with *Bradyrhizobium daqingense*.

haemoglobin content in nodules was quantified using drapkin solution at 254 nm at 30 DAS and yield parameters at harvest.

b) LAB, total free living N₂ fixers and P-solubilizers count was taken by serial dilution of ten gram of soil up to 10⁻⁵ and plated on MRS agar media, Norris N free medium and Pikovskaya's medium respectively. Bacterial colonies thus formed were taken as total count for LAB and N₂ fixers on respective media and zone of solubilization for P-solubilizers.

c) Rhizosphere soil enzyme activities like Urease activity (Tabatabai and Bremner, 1972), Phosphatase activity (Evazi and Tabatabai, 1979) and Dehydrogenase activity (Casida *et al.* 1964) at peak stages.

d) Plant nutrient content at harvest was estimated for total Nitrogen by kjeldhal process (Tandon, 1998), phosphorus content by vanadomolybophosphoric yellow colour method (Tandon, 1998), potassium content by flame photometry method (Sparks, 1996).

RESULTS AND DISCUSSION

Seedling vigour and compatibility

Seed inoculation significantly enhanced seed germination and seedling vigour of soybean. The bacterial strain AL-59 increased seed germination up to 11% over nontreated

Table 1: Effect of selected efficient nodulation stimulating LAB isolates on seedling vigour of soybean.

| Isolate | Germination (%) | Shoot length (cm) | Root length (cm) | SVI |
|---------------|-----------------|-------------------|------------------|--------|
| Control | 84.67 | 4.0 | 3.5 | 0633.9 |
| AL-44 | 86.33 | 4.5 | 3.6 | 0704.8 |
| AL-45 | 93.33 | 6.1 | 4.8 | 1011.6 |
| AL-48 | 94.00 | 6.0 | 5.0 | 1037.7 |
| AL-49 | 87.00 | 4.7 | 4.2 | 0781.3 |
| AL-58 | 89.67 | 5.1 | 4.1 | 0825.8 |
| AL-59 | 95.67 | 6.7 | 5.4 | 1160.4 |
| S.Em. ± | 1.19 | 0.18 | 0.28 | 40.62 |
| C.D. (p=0.01) | 4.92 | 0.73 | 1.16 | 167.77 |

Table 2: Effect of efficient nodulation stimulating LAB isolates on plant growth characters and chlorophyll content at 30 and 60 DAS.

| Treatments | Plant height (cm) | | Root length (cm) | | No of branches | | Relative chlorophyll content (SPAD values) | |
|----------------|-------------------|--------|------------------|--------|----------------|--------|--|--------|
| | 30 DAS | 60 DAS | 30 DAS | 60 DAS | 30 DAS | 60 DAS | 30 DAS | 60 DAS |
| T ₁ | 37.0 | 49.3 | 16.3 | 17.7 | 6.33 | 9.2 | 38.8 | 39.0 |
| T ₂ | 42.7 | 53.0 | 23.0 | 23.3 | 8.00 | 10.2 | 42.1 | 42.0 |
| T ₃ | 60.3 | 62.0 | 48.7 | 50.3 | 13.00 | 15.3 | 42.6 | 41.4 |
| T ₄ | 62.3 | 64.7 | 32.7 | 36.3 | 9.67 | 13.8 | 42.0 | 40.7 |
| T ₅ | 40.3 | 51.0 | 33.0 | 35.0 | 6.33 | 8.3 | 40.9 | 41.7 |
| T ₆ | 61.7 | 65.7 | 44.7 | 51.3 | 10.00 | 14.1 | 40.3 | 41.9 |
| T ₇ | 61.0 | 62.3 | 45.7 | 46.3 | 10.00 | 14.2 | 42.2 | 43.0 |
| T ₈ | 72.0 | 74.3 | 77.3 | 82.7 | 11.67 | 17.8 | 42.4 | 44.0 |
| S.Em. ± | 1.7 | 1.0 | 1.7 | 5.2 | 0.42 | 0.9 | 0.50 | 0.50 |
| C.D. (p=0.01) | 6.9 | 4.1 | 6.8 | 21.6 | 1.76 | 3.5 | 2.09 | 2.00 |

*UIC- Un -inoculated control. *DAS: Days after sowing.

control. The highest was noticed in AL-59 and AL-48 treated seeds, which recorded 1160.4 and 1037.7 vigour index respectively (Table 1).

Plant physiological observations

The co-inoculation of LAB had significant influence on plant physiological parameters at 30 and 60 DAS. The observations recorded in Table 2, clearly depicts that the treatment T₈ (POP+Consortium) showed highest plant height, root length, number of branches and chlorophyll content by 72 cm, 77.3 cm, 11.67 and 42.4 respectively (Table 2) and similar trend was noticed at 60 DAS (Fig 2 and 3).

The phenomenon of enhanced root and shoot growth can be attributed to production of auxins and nutrient mineralisation by PGPR (Steenhoudt and Vanderleyden, 2000). Similar findings were observed by Overvoorde *et al.* (2010), who opined that IAA when produced at low concentration showed positive significant influence over lateral root formation.

Nodulation

Co-inoculation of *Bradyrhizobium daqingense* with LAB isolates significantly increased the number of nodule; Leg haemoglobin content and nodule dry weight per plant as compared to control. The highest number of nodules per plant was recorded in treatment T₈(POP+Consortium) which scored 74.3 nodules per plant, which is 33.67% more than control (49.3), followed by T₆ (POP+ AL-58) with 65.7 nodule number (Table 3). The root scanning images (Fig 1) of uprooted plants at 30 DAS confirms the differential ability of consortium treated compared to other treatments in rooting and nodule bearing. Consequently the analysis for nodule dry weight is in accordance with nodule number, that highest nodule bearing consortium treated plants showed higher nodule dry weight of 117 mg. The analysis of leghaemoglobin content was done for one gram of fresh nodule extract using drapkin solution. The spectrophotometric values at 254 nm reveals that, consortium treated plants produced more leghaemoglobin (6.21 mg g⁻¹ fresh weight nodules) compared to control (2.98 mg g⁻¹ fresh weight nodules) (Table 3).

Many studies have shown that simultaneous infection with rhizobia and rhizospheric bacteria increases nodulation and growth in a wide variety of legumes (Bolton *et al.*, 1990). Accumulation of lactic acid in root nodules at the time of nodule development in response to symbiotic *B. japonicum* (Laurent *et al.*, 2010) might favour the nodule development, this accumulation of lactic acid was previously reported to be abundant in alfalfa (Swaraj and

Bishnoi 1999; Barsch *et al.*, 2006) and some lactic acid polymers have also shown to stimulate plant growth in corn and soybean (Kinnersley *et al.*, 1990; Chang *et al.*, 1996).

Here in this study application of lactic acid bacteria might have created favorable conditions for the nodulation and would have assisted the *Bradyrhizobium* in the process of root hairs infection, bacteriod development and might

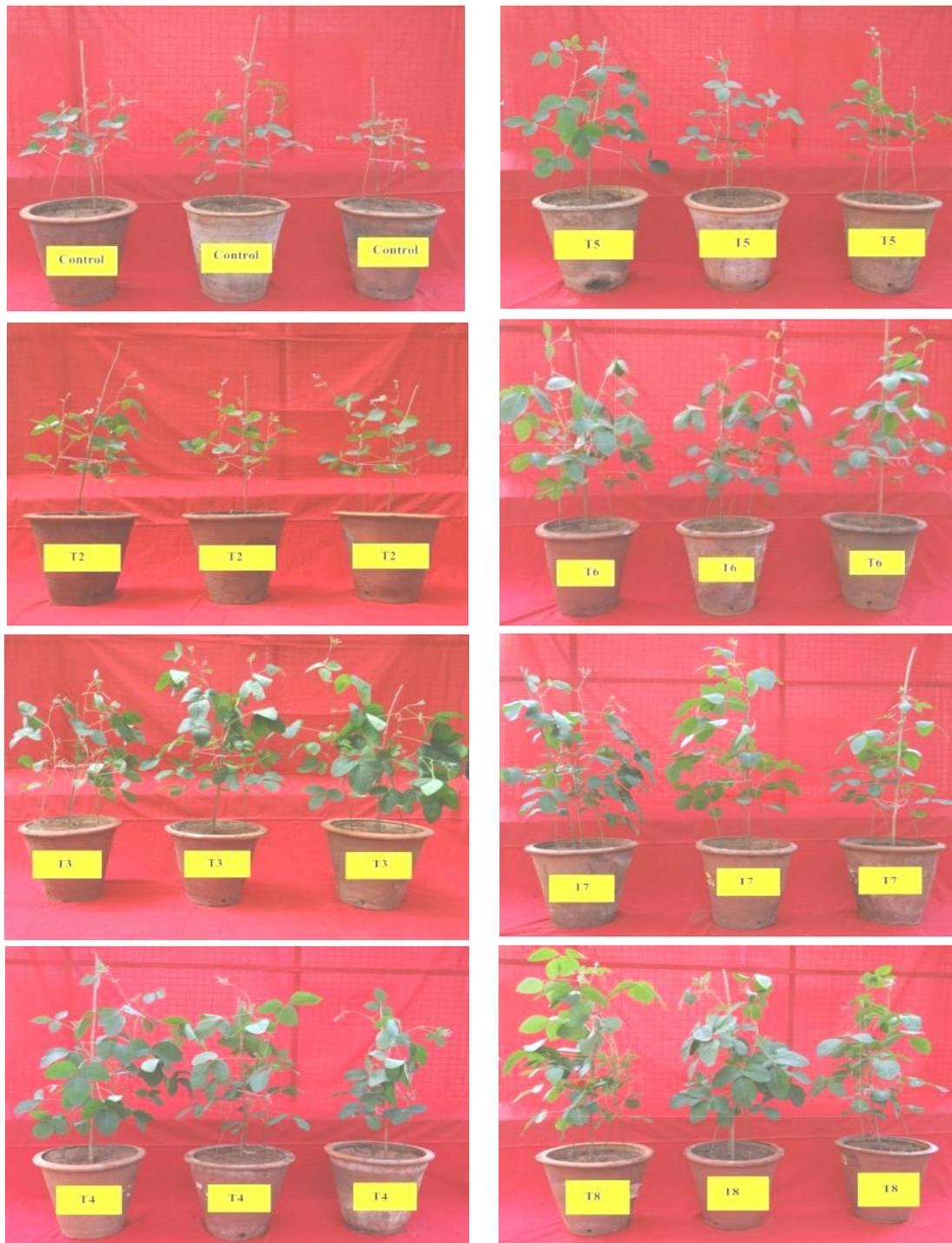


Fig 2: Comparison of effective nodulation stimulating LAB isolates with control at 30 DAS in pot culture study.

have facilitated the nodule formation and development by producing phytoalexins or flavonoid compounds.

Soil microbial analysis

The soil microbial analysis was done at the time of flowering and harvest indirectly by computing the enzyme activities like urease, phosphatase and dehydrogenase activities in soil. The result showed that microbial activity with respect to production of above mentioned enzymes was relatively higher at the time of flowering compared to harvest stage. This might be due to differential release of root exudates at different plant growth stage, which will in turn act as carbon and energy source for the soil microbes. The flowering stage is one where there is no tillage activity in soybean, in this context (Gholamreza *et al.*, 2016) showed highest microbial biomass carbon and recorded higher values of acid phosphatase, alkaline phosphatase and dehydrogenase activity. As far as treatment combinations are concerned, the one which received consortium (T₈) of all six isolates exhibited higher soil enzyme activity. Which scored 63.6 (µg NH₄-N/Soil day⁻¹) for urease activity, 700.24 (µg PNP/g soil hour⁻¹) for phosphatase activity and 233.37 (µg TPF/g soil

Table 3: Effect of LAB in augmenting nodulation of *Bradyrhizobium daqingense* in soybean at 30 DAS.

| Treatments | No. of nodules | Nodule dry weight (mg) | Leghaemoglobin content (mg g ⁻¹ fresh weight nodules) |
|----------------|----------------|------------------------|--|
| T ₁ | 14.33 | 037.3 | 2.98 |
| T ₂ | 18.33 | 044.3 | 3.87 |
| T ₃ | 32.00 | 097.9 | 5.79 |
| T ₄ | 35.00 | 095.3 | 4.69 |
| T ₅ | 18.00 | 039.7 | 4.03 |
| T ₆ | 20.33 | 075.3 | 4.49 |
| T ₇ | 39.33 | 098.3 | 4.44 |
| T ₈ | 47.67 | 117.0 | 6.21 |
| S.Em. ± | 0.93 | 3.64 | 0.35 |
| C.D. (p=0.01) | 3.83 | 15.0 | 1.43 |

*UIC- Un inoculated control. *DAS: Days after sowing.

Table 4a: Influence of co-inoculation of LAB on soil enzyme activities at flowering and harvest.

| Treatments | Urease activity (µg NH ₄ -N/Soil day ⁻¹) | | Phosphatase activity (µg PNP/g soil hour ⁻¹) | | Dehydrogenase activity (µg TPF/g soil day ⁻¹) | |
|----------------|---|------------|--|------------|---|------------|
| | At flowering | At harvest | At flowering | At harvest | At flowering | At harvest |
| T ₁ | 23.43 | 10.38 | 313.60 | 271.95 | 074.46 | 066.07 |
| T ₂ | 33.93 | 12.19 | 481.10 | 430.14 | 077.53 | 063.08 |
| T ₃ | 39.83 | 17.91 | 575.00 | 524.83 | 179.52 | 159.85 |
| T ₄ | 35.95 | 12.31 | 601.12 | 525.33 | 156.12 | 131.33 |
| T ₅ | 27.31 | 12.67 | 557.48 | 507.74 | 126.56 | 107.85 |
| T ₆ | 38.69 | 18.92 | 566.86 | 498.81 | 169.49 | 145.28 |
| T ₇ | 47.29 | 20.33 | 612.73 | 544.67 | 182.50 | 156.95 |
| T ₈ | 63.60 | 25.31 | 700.24 | 617.00 | 233.37 | 205.67 |
| S.Em. ± | 00.33 | 00.52 | 000.87 | 010.27 | 000.72 | 005.17 |
| C.D. (p=0.01) | 01.38 | 02.13 | 003.58 | 042.42 | 002.98 | 021.35 |

*UIC- Un inoculated control.

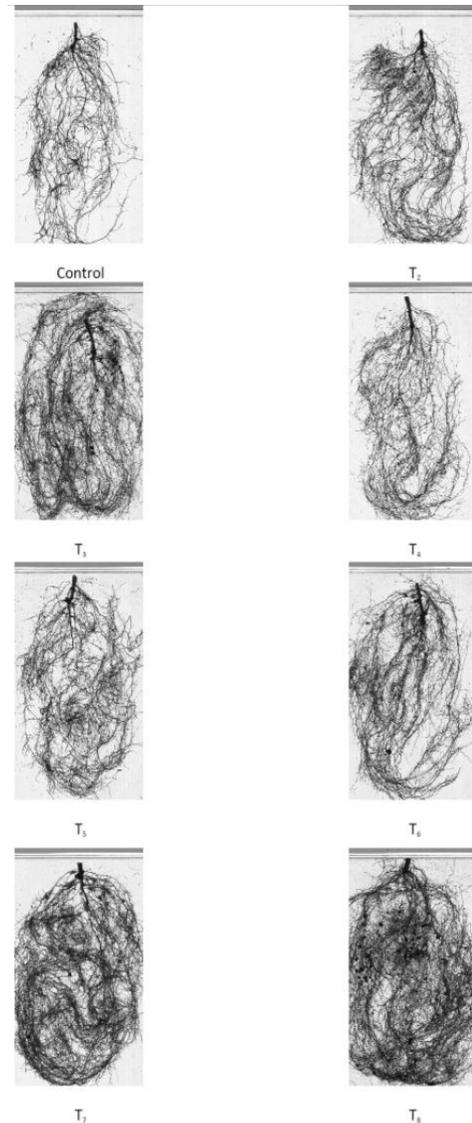


Fig 3: Root scanning images of efficient nodulation stimulating LAB isolates with control at 30DAS in pot culture study.

day⁻¹) for dehydrogenase activity. Similar tendency was noticed at harvest stage. However, enzyme activities were lower at harvest than flowering stage.

The persistence of seed inoculated microbes till to the harvest stage depicts their competitive ability to survive as free living organisms in absence symbiotic host. In this regard soil analysis was done to enumerate total free living N fixers using Norris N free agar medium, P solubilisers by

Table 4b: Soil microbial count at harvest.

| Treatments | Free living N fixers (CFU 10 ⁶ /g Soil) | PSB (CFU 10 ⁶ /g Soil) | LAB count (CFU 10 ⁶ /g Soil) |
|----------------|--|---|---|
| T ₁ | 9.7 | 11.0 | 00.6 |
| T ₂ | 10.3 | 12.3 | 19.0 |
| T ₃ | 12.7 | 18.0 | 19.7 |
| T ₄ | 11.7 | 18.1 | 17.0 |
| T ₅ | 10.0 | 12.7 | 20.7 |
| T ₆ | 12.3 | 13.3 | 20.7 |
| T ₇ | 12.7 | 12.0 | 22.3 |
| T ₈ | 13.3 | 17.3 | 27.3 |
| S.Em. ± | 0.8 | 0.8 | 1.0 |
| C.D. (p=0.01) | 3.3 | 3.4 | 4.3 |

*UIC- Un inoculated control. * PSB- Phosphorus solubilizing bacteria

Pikovaskaya agar medium and LAB count using MRS agar medium. The results depicted that, all treatments showed the existence of above mentioned microbes. However, their number varied among the treatments and the highest enumeration was recorded in consortium treated (T₈) soil (Table 4 a, b).

Plant nutrient concentration and yield

The values represented in Table 5 (a, b) clearly depicts that T₈ (POP+Consortium) showed highest N, P and K uptake, which is represented in terms of percentage as 3.68%, 0.22% and 2.03% respectively. The micronutrient analysis was done for Cu, Zn, Fe and Mn at harvest for all the treatments. In all cases T₈ (POP+Consortium) showed maximum up take for Cu, Zn, Fe and Mn up take by 94.8, 207.7, 984.7 and 95.4 ppm respectively.

Kang *et al.*, (2015) investigated the effects of three potential members of an EM consortia, *Rhodobacter sphaeroides*, *Saccharo-mycetes cerevisiae* and *Lactobacillus plantarum* on the growth and development of cucumber. All three microorganisms increased growth, nutrient uptake, amino acid content and yield. The increased availability of nutrients to plant might be due to the increased nodule number for N uptake through N fixation, production of lactic acid for P and Zn solubilisation, production of

Table 5a: Plant nutrient content at harvest as influenced by LAB inoculum.

| Treatments | N (%) | P (%) | K (%) | Cu (ppm) | Zn (ppm) | Fe (ppm) | Mn (ppm) |
|----------------|-------|-------|-------|----------|----------|----------|----------|
| T ₁ | 0.25 | 0.16 | 0.35 | 19.4 | 26.1 | 427.6 | 42.2 |
| T ₂ | 0.63 | 0.17 | 1.58 | 22.2 | 30.9 | 451.1 | 52.3 |
| T ₃ | 0.80 | 0.17 | 1.71 | 42.8 | 40.9 | 977.7 | 89.5 |
| T ₄ | 0.78 | 0.17 | 1.40 | 26.1 | 36.9 | 569.6 | 73.4 |
| T ₅ | 0.70 | 0.18 | 1.11 | 32.2 | 38.1 | 448.8 | 63.9 |
| T ₆ | 0.59 | 0.17 | 1.72 | 35.4 | 40.3 | 575.0 | 64.3 |
| T ₇ | 0.88 | 0.19 | 1.80 | 46.6 | 60.6 | 846.2 | 83.4 |
| T ₈ | 3.68 | 0.22 | 2.03 | 94.8 | 207.7 | 984.7 | 95.4 |
| S.Em. ± | 0.10 | 0.002 | 0.02 | 01.4 | 001.6 | 006.3 | 01.4 |
| C.D. (p=0.01) | 0.42 | 0.009 | 0.12 | 05.9 | 006.6 | 026.2 | 05.6 |

*UIC- Un inoculated control.

Table 5b: Effect of efficient nodulation stimulating LAB isolates on yield and yield components of soybean at harvest.

| Treatments | Pods per plant | Seeds per pod | Yield per plant in (g) | Shoot dry weight (g) | Root dry weight (g) | Total dry weight (g) |
|----------------|----------------|---------------|------------------------|----------------------|---------------------|----------------------|
| T ₁ | 05.20 | 2.00 | 03.80 | 2.95 | 0.50 | 07.25 |
| T ₂ | 06.10 | 2.33 | 07.57 | 3.43 | 0.67 | 11.67 |
| T ₃ | 09.67 | 2.23 | 11.77 | 1.26 | 0.97 | 14.00 |
| T ₄ | 11.77 | 2.33 | 10.83 | 1.41 | 0.83 | 13.07 |
| T ₅ | 10.33 | 2.00 | 08.53 | 4.03 | 0.77 | 13.33 |
| T ₆ | 07.77 | 2.10 | 14.00 | 1.74 | 0.93 | 16.67 |
| T ₇ | 13.10 | 2.20 | 13.67 | 3.36 | 1.27 | 19.30 |
| T ₈ | 18.47 | 3.00 | 18.33 | 4.36 | 1.97 | 23.67 |
| S.Em. ± | 0.58 | | 0.74 | | 0.06 | 0.50 |
| C.D. (p=0.01) | 2.39 | | 3.04 | | 0.24 | 2.05 |

*UIC- Un inoculated control.

siderophores for Fe uptake and proliferated root growth for quenching other nutrients in soil.

CONCLUSION

The LAB isolated from soybean rhizosphere have proven from the above evidences that, the property of LAB isolates to produce growth promoters like IAA and GA has interacted synergistically in augmenting nodulation of *B. daqingense*. The root scanning images clearly depicts profuse growth of root hairs in LAB treated soybean, this is more promising in LAB consortium treated plants compared to control. Healthy root and nodule growth has favoured plant to interact more with available nutrients in soil and has given more grain yield.

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