

## Inoculation of nitrogen-fixing halobacteria in the contribution to tolerance to salt stress in bean tepary

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### Abstract

Drought tolerant tepary bean (*Phaseolus acutifolius* A. Gray) is a microbiologically unexplored culture in arid regions of Mexico where it is consumed as a source of dietary protein. However, soil salinity hinders the productivity of this crop. The isolation of specific halotolerant nitrogen-fixing bacteria (BFN) could be a novel practice to improve salt tolerance and improve crop productivity. In the present study, 24 BFN were isolated from seeds of three tepary ecotypes exposed to high salinities (0, 0.25 and 0.5 M NaCl) *in vitro*. Nitrogen-fixing halotolerant strains were characterized. Among the halotolerant isolates, only one showed high nitrogenous activity of  $6.97 \pm 1.1$  nmol of culture h<sup>-1</sup>. From the 16S analysis of rRNA, the halotolerant microorganism exhibited 99% sequence homology with the known bacterium *Bacillus amyloliquefaciens*. The results show that the inoculation with the identified halotolerant isolate had stimulating effects on growth parameters, seed germination, seed emergence, shoot length, root length, biomass, foliar chlorophyll and protein content, ash and crude fibers.

**Keywords:** *Azospirillum halopraeferens*, *Bacillus amyloliquefaciens*, *Phaseolus acutifolius*, nitrogen fixation, saline stress.

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## Introduction

The growing problem of the increase of salts in the areas of cultivation together with the low rainfall and high temperatures of arid-saline environments, requires integral solutions for their productivity (Ungar, 1982). The tepary bean (*Phaseolus acutifolius* A. Gray) is a legume adapted to the high temperatures and droughts of the Sonoran Desert (Southwest of the United States of America and Northwest of Mexico).

The seeds of this species are characterized by a high content of dietary protein (21.1 to 32.49%) compared to other *Phaseolus* species; for example, *P. vulgaris* (19.1-29.7%), *P. lunatus* (19.7-24.9%), *P. coccineus* (20-27.4%), *P. polyanthus* (21.6-25.6%), *P. filiformis* (24.2%) and *P. Angustissimus* (25.9%) (Nabhan, 1985). In general, tepary yields have been estimated between 200 and 900 kg ha<sup>-1</sup>. Yields of up to 2000 kg ha<sup>-1</sup> have been reported (Hamama and Bhardwaj, 2002; Ahmad *et al.*, 2013) depending on the sowing season and the appropriate supplementation of nitrogen fertilizers (N).

In northwestern Mexico, large amounts of chemical fertilizers are applied to the soils to compensate for N deficiency and to increase the yields of tepary beans, unfortunately, the excessive and continuous use of these fertilizers increases salinity, affecting the physical and soil chemistry, which as a whole alter microbial activities, which could be beneficial for crops (Rueda-Puente *et al.*, 2003, 2007, 2010). To reduce inputs of nitrogen fertilizers in order to slow or stop further increases in soil salinity and reduce fertilizer costs, rhizobacteria that promote plant growth, particularly N-fixing bacteria (RPCP-BFN), are an attractive and sustainable alternative for the cultivation of tepary beans in saline soils of semi-arid and arid regions, such as those located in the state of Sonora, Mexico.

Studies on bacteria associated with tepary bean yields are scarce. Studies related to RPCP-BFN have shown the biopromotor and bioprotective effect against environmental stress in plants, for example, drought, osmotic stress and flood (Mayak *et al.*, 2004), extreme temperatures (Terre *et al.*, 2007), nutrient deficiency (Cassan *et al.*, 2009) and toxic metals (Sandhya *et al.*, 2010; Rokhzadi and Toashih, 2011).

During the last decades, the RPCP-BFN have been isolated and cultivated as a promising alternative (Pathak and Keharia, 2013). Based on the above described, the hypothesis that arises is whether the inoculation with RPCP-BFN, especially those of the halotolerant type (HBFN) contribute to improve the effect of salinity stress in crops such as tepary beans. Therefore, the objectives of the present study were: i) to isolate a nitrogen-fixing halotolerant bacteria from the rhizosphere of *P. acutifolius*; and ii) to identify the contribution of the isolate (*Bacillus amyloliquefaciens* RP22) on tolerance to stress by salinity in seeds and seedlings of *P. acutifolius*.

## Materials and methods

### Seeds

The seeds evaluated in the present study were those ecotypes of San Judas (SJ), Indio Yumi (IY) and Elena-Mora Property (EMP), which were collected from the Sierra de Sonora-Chihuahua (29° 05' north latitude 110° 57' west longitude) in the state of Sonora.

## Halobacteria nitrogen fixers

### Isolation

For the isolation of bacteria, the technique of Rueda *et al.* (2003). The seeds were crushed with mortar and pestle to obtain the final weight of 4 g and serial dilutions (up to  $10^{-6}$ ) in 0.85% sterile saline were seeded on OAB free of N in agar medium according to (Bashan *et al.*, 1993) with three different concentrations of NaCl (0, 0.25 and 0.5 M) (Rueda *et al.*, 2003), agar plates were incubated at 30 °C for four days, obtaining 24 individual colonies on agar, based on the phenotype (color, brightness, shape, elevation and margin) according to Smibert and Krieg (1994), later the purified isolates were stored at -80 °C (15% glycerol).

### Nitrogen fixation (acetylene reduction technique)

The acetylene reduction assay (ERA) to the selected strains was performed as described by Reinhold *et al.* (1987). The halotolerant isolates were cultured in serum flasks containing 25 ml of liquid medium OAB free of N and 0.5 M NaCl. The results were compared with a known halobacteria (HBFN) *Azospirillum halopraeferens* AU10 (Reinhold *et al.*, 1987). The isolate that showed the highest acetylene reduction activity was selected for genetic identification and for further evaluation with inoculation tests.

### Identification

The selected halotolerant bacterium was identified at the genus level by partial sequencing of 16S rRNA genes by Acculab, Inc. (USA). The sequence obtained was compared with sequences in the GenBank database of the National Center for Biotechnology Information (NCBI) using the BLAST program (<http://blast.ncbi.nlm.nih.gov/>; <http://www.megasoftware.net/>) and then it was deposited in the GenBank under the access number KM652480. In addition, a phylogenetic tree was built using MEGA6 freeware (<http://www.megasoftware.net/>).

### Inoculation tests-experiment design

The inoculation tests were carried out in a growth chamber and inside a greenhouse. Both trials were randomized in a complete block design with a three-dimensional design ( $3 \times 3 \times 3$ )= 27 treatments with n repetitions; 3 bean ecotypes (San Judas= SJ =, Indian Yumi= IY = and Elena-Mora Property= EMP=; 3 levels of NaCl (0, 0.25 and 0.5 M) and 3 levels of bacteria (Ba= *Bacillus amyloliquefaciens*, Ah= *Azospirillum halopraeferens* AU10, U= the non-inoculated ones (Table 3) Once seeds were sterilized, they were inoculated with the HBFN isolated and characterized in this study, using the air vacuum pump technique described by Carrillo *et al.* (1998).

### Tests in growth chamber

The germination of 675 seeds (25 seeds per treatment) was performed in sterilized Petri dishes, sealed with paraffin tape to maintain humidity conditions and placed in a growth chamber with a 16-hour light photoperiod ( $32 \pm 0.5$  °C) and 8 h at night ( $25 \pm 0.5$  °C) with a relative humidity of  $36 \pm 1\%$  (HR). After 7 days, the germination rates of the seed, the growth (length of stem and root) and the dry weights were recorded.

## Greenhouse tests

For the tests inside the greenhouse, the substrate used in the plastic pots was sterile vermiculite without nutrients. The 540 seeds (20 seeds per treatment) were planted in pots and the emergence rates were recorded at day 9. The seedlings were cultivated under control conditions with a photoperiod 16 h day ( $32 \pm 0.5$  °C) and 8 h night ( $25 \pm 0.5$  °C) and with  $36 \pm 1\%$  HR. The plants were irrigated every third day, considering spilling the water to prevent an increase in salinity with the corresponding concentrations of NaCl and maintained until 21 days.

## Variables evaluated

Germination and emergency seed rates were calculated using the formula described by Maguire (1962):  $M = n_1/t_1 + n_2/t_2 + \dots + n_7/t_7 + \dots + n_9/t_9$ , where  $n_1, n_2, \dots, n_7, \dots, n_9$  are; 1 are the number of germinated seeds;  $t_1, t_2, \dots, t_7, \dots, t_9$  is the time in days. The chlorophyll foliar content was measured with a chlorophyll meter (SPAD-502, Minolta, Japan). Root and shoot lengths were measured separately with a digital calibrator (General No. 143, General Tools Manufacturing Co., Inc., New York, USA). The plants were dried at 110 °C for 36 h to estimate the dry weight. The micro-kjeldahl method was used to determine the total N and crude protein ( $N \times 6.25$ ), while the crude fiber and the ash contents were determined gravimetrically (Snedecor, 1956).

## Statistical analysis

Statistical analyzes were performed using a one-way analysis of variance (Andeva) followed by Tukey's minimum significant difference test when statistical differences were observed. The values are expressed as means  $\pm$  standard error and the significant values were with  $p \leq 0.05$ .

## Results and discussion

### Halobacteria nitrogen fixer

The total counts of nitrogen-fixing bacteria (BFN) are shown in Table 1. It could be seen that the isolates on control agar plates (N-free OAB agar medium without NaCl) were  $8.6$  to  $12.6 \times 10^6$  forming units of colonies (UFC  $g^{-1}$  and from  $7.6$  to  $10.6 \times 10^6$  UFC  $g^{-1}$  for seeds with and without seed coat, respectively.) Isolates in the SJ and EMP ecotypes decreased significantly ( $p \leq 0.05$ ) in agar plates supplemented with NaCl 0.5 and 0.75 M. In the case of control seeds in the SJ ecotype ( $8.3$ - $9.3 \times 10^6$  CFU  $g^{-1}$ ), the counts showed a NaCl tolerance of 45-54% ( $3.7$ - $5 \times 10^6$  CFU  $g^{-1}$  seeds) and 29-33% ( $2.4$ - $3.1 \times 10^6$  CFU  $g^{-1}$  seeds) of total culturable BFN at 0.5 and 0.75 M NaCl, respectively.

Similarly, counts in the EMP ecotype showed a salt tolerance of 28-41% ( $3.6$ - $4.4 \times 10^6$  CFU  $g^{-1}$  seeds) and 18-22% ( $2.3 \times 10^6$  CFU  $g^{-1}$  seeds) of BFN total arable to 0.5 and 0.75 M NaCl, while in the control the values were  $10.6$ - $12.6 \times 10^6$  CFU  $g^{-1}$ .

**Table 1. Colony forming units of HBFN associated with seed of different ecotypes of *Phaseolus acutifolius* on agar OAB free N supplemented with different concentrations of NaCl.**

Seeds	Ecotypes	Count ( $\times 10^6$ UFC $g^{-1}$ seed)			
		0 M NaCl	0.25 M NaCl	0.5 M NaCl	0.75 M NaCl
with seed coat	SJ	9.3 $\pm$ 1.2a*	8.3 $\pm$ 0.6ab	5 $\pm$ 0.5bc	3.1 $\pm$ 0.5a
	IY	8.6 $\pm$ 2.6a	8 $\pm$ 1.1a	5 $\pm$ 1.1ab	2.6 $\pm$ 0.8ab
	EMP	12.6 $\pm$ 2a	8.6 $\pm$ 0.6b	3.6 $\pm$ 0.3c	2.3 $\pm$ 0.3b
without seed coat (endosperm)	SJ	8.3 $\pm$ 0.4a	6.7 $\pm$ 0.3b	3.7 $\pm$ 0.9c	2.4 $\pm$ 0.3b
	IY	7.6 $\pm$ 1.2a	7.4 $\pm$ 1.2a	5 $\pm$ 0.6ab	3.3 $\pm$ 0.4a
	EMP	10.6 $\pm$ 0.9a	7.6 $\pm$ 0.4b	4.4 $\pm$ 0.3c	2.3 $\pm$ 0.7b

\* = Values represent an average  $\pm$  standard error (average of three repetitions). Different letter in the same line denotes significant difference ( $p \leq 0.05$ , Tukey test). Ecotypes SJ = San Judas; IY = Indian Yumi; EMP = Elena-Mora Property.

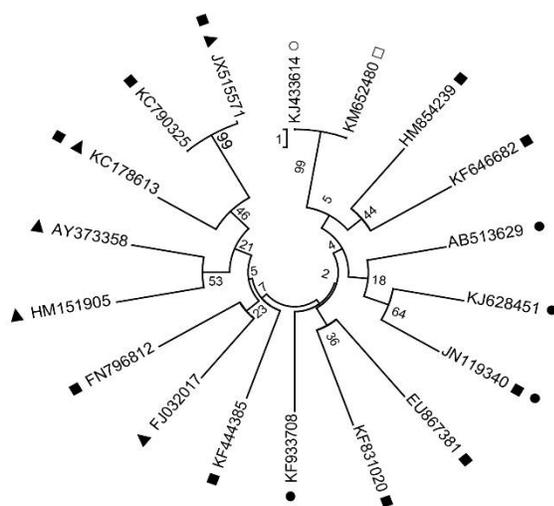
In relation to the capacity of N fixation (acetylene reduction), the strain encoded as RP22 had the highest acetylene reduction activity ( $6.97 \pm 1.1$  nmol culture  $h^{-1}$ ), which is similar to reference *A. halopraeferens* AU10 ( $7.83 \pm 1.3$  nmol of culture  $h^{-1}$ ). This result shows that growth improved by 28 and 43% with bacteria isolated at NaCl concentrations between 0.5 M and 0.75 M NaCl.

Analogously, Rueda *et al.* (2003) proposed that 5% of the halotolerant strains of saline-flat areas reduced acetylene  $\sim 6.26 \pm 0.56$  nmol of culture  $h^{-1}$  (*Klebsiella pneumoniae*) with the control *A. halopraeferens*, which produced a value of  $7.1 \pm 1.7$  nmol $^{-1}$   $h^{-1}$ . This could be attributed to the codependency of other bacteria, a phenomenon that is quite common among microorganisms (Diby and Harshad, 2015). The genetic identification of the bacterial isolate in the present study, based on the 16S rRNA gene sequence, showed a high similarity (99%) with the plant growth promoting bacterium *Bacillus amyloliquefaciens* (Figure 1).

The phylogenetic analysis based on the 16S rRNA gene isolated from the NFB halotolerant showed 99% similarity with a *B. amyloliquefaciens* (accession number KJ433614), a bacterium isolated from the cold desert of the northwest Himalayas by PGPR analyzes. As far as is known, the species *B. amyloliquefaciens* has never been reported for *P. acutifolius*, which is indicative of the association of beneficial bacteria and the plant under study. Other species of the genus *Bacillus* have been isolated and identified in other leguminous plants (*B. pumilus*, *B. cepacia*, *B. japonicum*, *B. vallismortis*, *B. mojavensis*, *B. atrophaeus*, *B. megaterium* and *B. Stearothermophilus*) (Bashan *et al.*, 1993; Dobbelaere and Okon, 2007ab).

### Inoculation tests

When the seeds were inoculated with bacterial strains, the negative effects of 2% NaOCl on germination of the seeds decreased (Table 2). Thus, higher germination rates were obtained under growth chamber conditions with inoculated seeds (53.3-100%) than with non-inoculated seeds (46.7 - 86.7%) in the three ecotypes of tepary beans treated with 0 M NaCl and 0.25 M. In addition, the germination rate was strongly affected when 0.50 M NaCl was applied, showing ranges of 0-13.3% and 6.7-26.7% for the non-inoculated and inoculated seeds, respectively (Table 2).



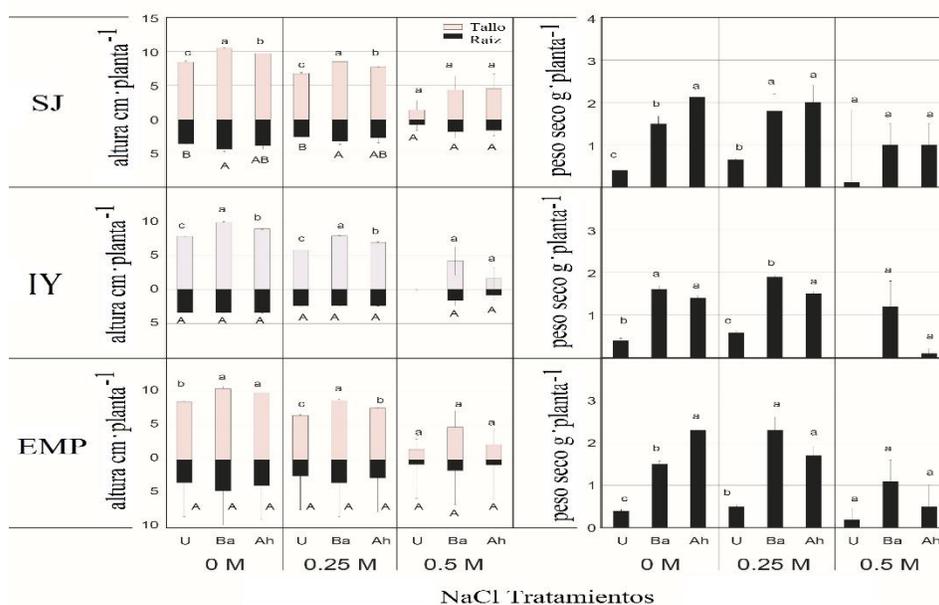
**Figure 1. Phylogenetic tree showing the similarity with the isolate *Bacillus amyloliquefaciens* RP22 (□) and the close relationship with *Bacillus amyloliquefaciens* (○) and its relationship with the halotolerant (●) fixative of N (▲) and endophyte (■) desert bacteria.** Alignment and close relationship with the tree was done using MEGA6 (<http://www.megasoftware.net/>). The bars indicate 10% divergence; a bootstrap analysis was developed with 1000 trials. KJ433614, *Bacillus amyloliquefaciens* strain IARI-AR25; KM652480, *Bacillus amyloliquefaciens* strain RP22; HM854239, *Bacillus safensis* strain KLH-14; KF646682, *Bacillus tequilensis* strain DH-10; AB513629, *Bacillus megaterium*; KJ628451, *Bacillus subtilis* strain HPCAQRKSM106; JN119340, *Halobacillus* sp. KLBMP 2429; EU867381, *Bacillus pumilus* strain CCGE2028; KF831020, *Bacillus firmus* strain L-4; KF933708, *Bacillus baekryungensis* strain QD56; KF444385, *Bacillus aryabhatai* strain HYR8(1); FJ032017, *Bacillus pumilus* strain ES4; FN796812, *Bacillus cereus* strain GP17; HM151905, *Bacillus subtilis* strain DAZ26; AY373358.1, *Bacillus megaterium* strain c5; KC178613, *Bacillus flexus* strain DNEB39; KC790325, *Bacillus methylotrophicus* strain SY33; JX515571, *Bacillus subtilis* strain H171.

**Table 2. Seed germination and emergence rates of three *Phaseolus acutifolius* ecotypes under growth chambers and greenhouse trials.**

NaCl treatment	Num. inoculated (control)	<i>B. amyloliquefaciens</i>	<i>A. halopraeferens</i> AU10
Germination chamber of growth			
0 M	46.7-86.7	93.3-100	86.7-100
0.25 M	53.3-66.7	73.3-100	53.3-80
0.5 M	0-13.3	20-26.7	6.7-20
Greenhouse emergency			
0 M	55.6-66.7	100	100
0.25 M	22.2-33.3	33.3-44.4	33.3-44.4
0.5 M	11.1-22.2	11.1-33.3	11.1-33.3

Regarding the parameters measured in growth chamber conditions, treatments of bean tepary in the variables of shoots and roots in seedlings, the maximum effects of bacterial inoculation with *B. amyloliquefaciens* RP22, as well as *A. halopraeferens* AU10 were for the ecotype SJ, showing significant results with  $p \leq 0.05$  to 0 M NaCl (8.9 and 10.5 cm of plant<sup>-1</sup>), between 6.9 and 8.7 cm plant<sup>-1</sup> for *B. amyloliquefaciens* and *A. halopraeferens* when adding 0.25 M NaCl, in comparison with the non-inoculated seedlings that were lower: (7.8 and 8.5 and 5.8 and 6.7 cm plant<sup>-1</sup> for 0 and 0.25 M NaCl (Figure 2).

Root lengths showed significant statistical differences ( $p \leq 0.05$ ) between the inoculated and non-inoculated seedlings in IY and EMP ecotypes, considering all the salinities, except the SJ ecotype where the plants were inoculated with *B. amyloliquefaciens* RP22, (4.4 and 2.7 cm plant<sup>-1</sup> to 0 and 2.5 M NaCl ( $p \leq 0.05$ ) (Figure 2).



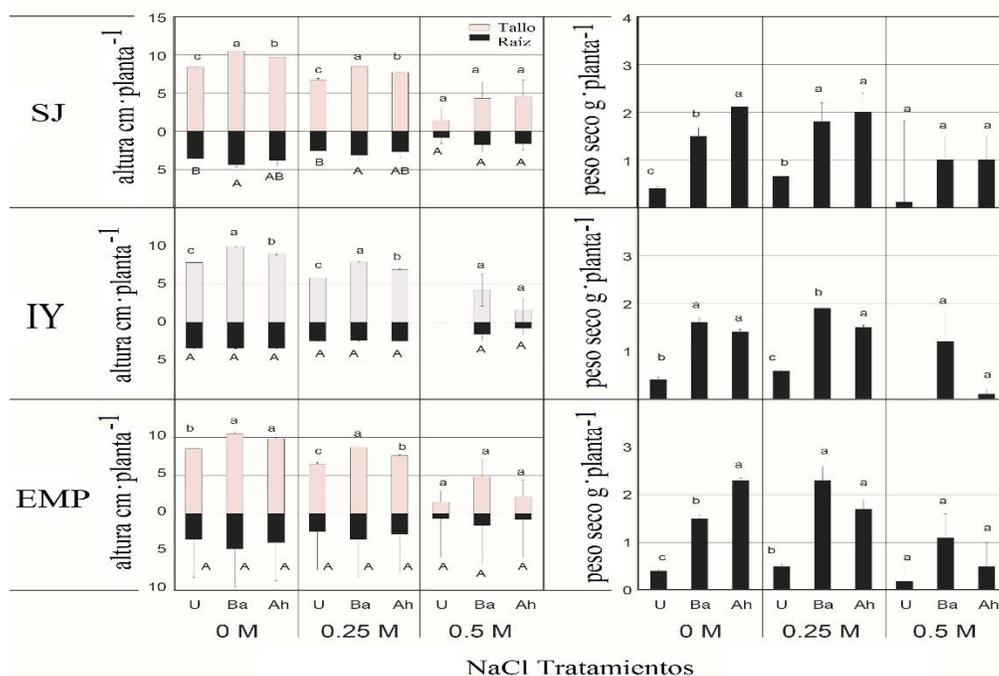
**Figure 2. Effect of NaCl concentrations and bacterial inoculation on the height and dry weight of *Phaseolus acutifolius* seedlings under growth chamber conditions.** The values represent the mean  $\pm$  standard error ( $n = 3$ ). Different letter indicates significant differences ( $p \leq 0.05$ , Tukey test) of the inoculated plants (Ba= *Bacillus amyloliquefaciens*, Ah= *Azospirillum halopraeferens* AU10) in relation to the non-inoculated controls (U). Ecotypes SJ= San Judas; IY= Indian Yumi; EMP= Elena-Mora Property.

The production of dry biomass according to the effect of the treatments were statistically different ( $p \leq 0.05$ ); the increases in dry biomass of the plants inoculated with respect to those not inoculated were 17 and 38%, respectively. Coincident with the growth chamber test, greenhouse test results (Figure 3) showed lower emergence seed rates and seedling biomass and growth values of the three ecotypes of tepary beans irrigated with 0.25 and 0.5 M NaCl with NaCl at 0 M, this assay also confirms the growth promoting effect of the plants on seeds and seedlings of both inoculated bacterial strains, which reaffirms the possibility of proposing beneficial microorganisms as biofertilizing prospects (Stefana *et al.*, 2013; Diby and Harshad, 2015).

Similar results indicate that the positive effects of this type of bacteria apparently are due to the production of growth promoting substances as reported in other studies (Yadav, 2011; Rugheim *et al.*, 2012).

Regarding the variables of growth and biomass in greenhouse conditions, significant differences were observed ( $p \leq 0.05$ ) in lengths of shoots in the SJ ecotype, considering the inoculation with *B. amyloliquefaciens* RP22 (70 cm plant<sup>-1</sup>), as well as *A. halopraeferens* AU10 (58 cm plant<sup>-1</sup>), in relation to the non-inoculated controls (U) (35 cm plant<sup>-1</sup>) in NaCl M (Figure 3).

Regarding the dry weight, the maximum values were recorded in inoculated seedlings in the three ecotypes at 0, 0.25 and 0.5 M NaCl in relation to the non-inoculated controls (U). However, it should be mentioned that salinity reduces dry weight, while it increases in all treatments (Figure 3). In particular, the values of germination, emergence, growth and biomass were higher in the seedlings inoculated with *B. amyloliquefaciens* RP22 than in those inoculated with *A. halopraeferens* AU10, particularly in the ecotype IY treated with NaCl 0 and 0.25 M (Figures 2 and 3).



**Figure 3. Effect of NaCl concentrations and bacterial inoculation on the growth and dry weight of *Phaseolus acutifolius* seedlings under greenhouse conditions.** The values represent the mean  $\pm$  standard error (n= 3). Different letters indicate significant differences ( $p \leq 0.05$ ) of the inoculated plants (Ba= *Bacillus amyloliquefaciens*, Ah= *Azospirillum halopraeferens* AU10) in relation to the non-inoculated controls (U). Ecotypes SJ= San Judas; IY= Indian Yumi; EMP= Elena-Mora Property.

Likewise, the analysis of foliar chlorophyll and the contents of proteins, ashes and crude fiber confirm the effect of NaCl treatments and bacterial inoculation in relation to controls (not inoculated) on the growth and biomass of the seedlings (Cuadro 3).

**Table 3. Effect of NaCl concentrations and bacterial inoculation on the total content of chlorophyll, protein, ash and crude fiber in *Phaseolus acutifolius* seedlings under greenhouse conditions.**

Ecotypes	NaCl treatments	Inoculants	Total chlorophyll	Protein (%)	Ashes (%)	Crude fiber (%)
SJ	0 M	U	31.5 ±0.08b	11.9 ±0.07b	5.5 ±0.03b	17.8 ±0.5b
		Ba	34.9 ±0.1a	14.4 ±0.07a	7.6 ±0.2a	21.9 ±0.9a
		Ah	34.1 ±0.03a	14 ±0.2a	6.5 ±0.2a	22.9 ±0.5a
	0.25 M	U	8.9 ±8.9b	3.1 ±3.1b	1.4±1.4b	4.7 ±4.7b
		Ba	29.4 ±0.4a	12.3 ±0.06a	7.2 ±0.2a	21.1 ±1a
		Ah	29 ±0.02a	11.8 ±0.2a	6.4 ±0.2a	22.85 ±0.3a
	0.5 M	U	4.9 ±4.9c	2.3 ±2.3b	1 ±1b	4.1 ±4.1b
		Ba	12.7 ±6.4a	5.1 ±2.5a	3.2 ±1.6a	8.4 ±4.2a
		Ah	6.6 ±6.6b	2.5 ±2.5b	1.9±1.9b	4.1 ±4.1b
IY	0 M	U	31.2 ±0.2b	11.29 ±0.04c	5.3 ±0.09b	17.5±0.9b
		Ba	34.9 ±0.8a	14.3 ±0.09a	7.6 ±0.3a	23.3 ±0.05a
		Ah	34.3 ±0.1a	13.66 ±0.2b	6 ±0.1b	23.1 ±0.2a
	0.25 M	U	8.9 ±8.9c	2.6 ±2.6c	1.3 ±1.3c	5.2 ±5.2c
		Ba	30.3 ±0.8a	12.2 ±0.07a	7 ±0.05a	23.7 ±0.6a
		Ah	19.6 ±9.8b	7.8 ±3.9b	4.2 ±2.1b	15.1 ±7.5b
	0.5 M	U	4.5 ±4.5b	2.5 ±2.5a	1 ±1a	3.7 ±3.7a
		Ba	6.6 ±6.6a	2.4 ±2.4a	1.7 ±1.7a	4.1 ±4.1a
		Ah	6 ±6a	2.6 ±2.6a	1.9 ±1.9a	4.4±4.4a
EMP	0 M	U	32.5 ±0.06b	11.5 ±0.01c	5.6 ±0.03c	22.4 ±0.03a
		Ba	35.2 ±0.4a	15 ±0.2a	8.6 ±0.4a	23.6 ±0.03a
		Ah	34.1 ±0.03a	14.6 ±0.4b	6.2 ±0.06b	23.5 ±0.19a
	0.25 M	U	9.2 ±9.2b	3.2 ±3.2b	1.5 ±1.5b	7.2 ±7.2b
		Ba	29.9 ±0.2a	12.7 ±0.2a	7 ±0.06a	21.8 ±0.2a
		Ah	29 ±0.02a	12.1 ±0.03a	6.2 ±0.2a	22.1 ±0.1a
	0.5 M	U	5.8 ±5.8b	2.5 ±2.5b	1 ±1b	4.4 ±4.4b
		Ba	18.6 ±0.61a	7.9 ±0.03a	5.8 ±0.4a	13.2 ±0.6a
		Ah	19.3 ±0.6a	7.7 ±0.1a	5 ±0.5a	12.6 ±0.3a

## Conclusions

The present study represents a first report of *Bacillus amyloliquefaciens* as a nitrogen-fixing bacterium associated with the leguminous *Phaseolus acutifolius*. Likewise, it is a first approximation in the first stages under salinity conditions, specifically with tepary beans, evaluating the isolated and halotolerant microorganism compared to a biological control *Azospirillum halopraeferens* AU10.

The halobacteriae *B. amyloliquefaciens*, isolated from ecotype tepary seeds, is able to resist high salt concentrations (0.75 M NaCl) and may facilitate the promotion of plant growth in the presence of levels of inhibition of salinity growth of soil that exceed 0.25 M NaCl.

Inoculation based on beneficial bacteria such as nitrogen fixers is a biological and reliable method to help maintain or improve the fertility of the soils that support the tepary fields. However, it is suggested to develop more evaluations in more advanced phenological stages and in other ecotypes in order to better understand the mechanisms of action and the particular reactions with each ecotype before recommending the association of bacteria at the field level.

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