



Biotecnia

ISSN: 1665-1456

Universidad de Sonora, División de Ciencias Biológicas y de la Salud

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Biotecnia, vol. 24, no. 1, 2022, January-April, pp. 15-22

Universidad de Sonora, División de Ciencias Biológicas y de la Salud

DOI: <https://doi.org/10.18633/biotecnia.v24i1.1353>

Available in: <https://www.redalyc.org/articulo.oa?id=672974928005>

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Characterization of native plant growth-promoting bacteria (PGPB) and their effect on the development of maize (*Zea mays* L.)

Caracterización de bacterias promotoras de crecimiento vegetal (BPCV) nativas y su efecto en el desarrollo del maíz (*Zea mays* L.)

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ABSTRACT

Maize (*Zea mays* L.) is one of the most important cereals for Mexico and humanity. The Yaqui Valley -located at the northwest of Mexico- is one of the most important agricultural regions worldwide, it is characterized by an indiscriminate use of inorganic fertilizers for improving crop yields, leading to an increased environmental and economic cost of maize production. This work carried out a morphological, metabolic, and molecular characterization of native bacteria associated with maize rhizosphere, evaluating the positive effects of bacterial inoculation on plants under greenhouse conditions. The evaluated strains were taxonomically affiliated -based on the 16S rRNA gene- as *Bacillus* sp. (13B41), *Advenella incenata* (22A67), *Pantoea dispersa* (22B45), and *Rhizobium pusense* (31B11). All these strains were able to synthesize indoles, produce siderophores, and solubilize phosphates. The individual inoculation of these strains to maize plants showed a significant increment (compared to un-inoculated plants) in height (35-40 %), shoot dry weight (244-289 %), root dry weight (99-137 %), and SPAD values (40- 47 %). The native bacteria associated with maize in the Yaqui Valley are a promising alternative to promote the growth of their host plant and contribute to a sustainable maize production.

Keywords: microbial inoculants, plant growth promotion, soil, greenhouse.

RESUMEN

El maíz (*Zea mays* L.) es uno de los cereales más importantes para México y la humanidad. El Valle del Yaqui -ubicado en el noroeste de México- es una de las regiones agrícolas más importantes a nivel mundial, se caracteriza por un uso indiscriminado de fertilizantes inorgánicos para mejorar el rendimiento de los cultivos, lo que genera un mayor costo ambiental y económico de la producción de maíz. En este trabajo se realizó una caracterización morfológica, metabólica y molecular de bacterias nativas asociadas a la rizósfera del maíz y se evaluaron los efectos positivos de la inoculación bacteriana en plantas de maíz bajo condiciones de invernadero. Basándose en la secuencia del gen del 16S ARNr las cepas evaluadas se afiliaron taxonómicamente como *Bacillus* sp. (13B41), *Advenella incenata* (22A67), *Pan-*

toea dispersa (22B45) y *Rhizobium pusense* (31B11). Todas estas cepas pudieron sintetizar indoles, producir sideróforos y solubilizar fosfatos. La inoculación individual de estas cepas en maíz mostró un incremento significativo (en comparación con plantas no inoculadas) en altura (35-40 %), peso seco de brotes (244-289 %), peso seco de raíces (99-137 %) y valores SPAD (40-47%). Las bacterias asociadas al maíz nativas del Valle del Yaqui son una alternativa prometedora para promover el crecimiento de su planta hospedante y contribuir a una producción de maíz sostenible.

Palabras clave: inoculantes microbianos, promoción del crecimiento vegetal, suelo, invernadero.

INTRODUCTION

Maize (*Zea mays* L.) belongs to the group of cereals with worldwide importance, due to its applications in animal consumption, including human. In 2018, world maize production reached 1147.62 million tons, the highest amount among the main cereals (FAO, 2020). In Mexico, during 2018, around 7.3 million hectares were cultivated with this cereal, with a total production of 27.1 million tons (SIAP-SADER, 2020). On the other hand, maize production in the main agricultural zones of northwestern Mexico, such as the Yaqui Valley, reached 449,000 tons in the year 2018, while in 2019 its production increased up to 567,000 tons (SIAP-SADER, 2020). However, this global and national food demand has led to the indiscriminate use of agrochemicals, in areas of intensive agriculture such as the Yaqui Valley, where high amounts of nitrogen and phosphate fertilizers have been used, as well as pesticides, to improve the yield of several crops such as wheat and maize (McCullough and Matson, 2016). Nevertheless, the excessive use of agrochemicals contributes to the contamination of soils and underground water supplies, leading to potential health risks and environmental degradation (Meza-Montenegro *et al.*, 2012). Thus, the need for the development of sustainable strategies for food production, to diminish the damage caused by the use of large quantities of agrochemicals, is evident.

The use of soil microorganisms (mainly bacteria) as inoculants with beneficial traits for plant development and soil health represents an attractive alternative to conven-

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Recibido: 28 de junio del 2021

Aceptado: 28 de octubre de 2021

tional agriculture, named Plant Growth Promoting Bacteria (PGPB) (Valenzuela-Aragon *et al.*, 2019). PGPB includes a large group of microorganisms that can be found in the rhizosphere and bulk soil, as well as within and on aerial plant tissues (de Souza *et al.*, 2015). PGPB applied to plants, either on soil or seeds, improve plant crops growth and protection against diseases and abiotic stress, by improving the biological nitrogen fixation, solubilization of phosphates, reduction of stress through diverse enzyme production, synthesis of phytohormones, production of siderophores, among others (Glick, 2012; de Souza *et al.*, 2015). Thus, the most studied genera of PGPB are *Acetobacter*, *Acinetobacter*, *Alcaligenes*, *Arthrobacter*, *Azospirillum*, *Azotobacter*, *Bacillus*, *Bradyrhizobium*, *Burkholderia*, *Enterobacter*, *Gluconacetobacter*, *Klebsiella*, *Methylobacterium*, *Ochrobactrum*, *Paenibacillus*, *Pantoea*, *Pseudomonas*, *Serratia*, *Stenotrophomonas*, and *Rhizobium* (Glick, 2012; Ahemad and Kibret, 2014). Several studies have demonstrated the ability of these bacterial genera to promote the growth of different crops such as rice, tomato, wheat, maize, chickpea, peanuts, among others, showing significant changes in physiological parameters, *i.e.*, dry weight, chlorophyll content, and plant height (Kumar *et al.*, 2014; Sharon *et al.*, 2016).

PGPB can be classified as native or indigenous when those are endemic to the site where they will be applied and as exogenous when they are isolated from other locations. Several research, mainly related to pollutants biodegradation, have shown that for an exogenous PGPB to properly perform its functions, not only must adapt to the local region edaphoclimatic conditions but also there must be compatibility with the native microorganisms (*i.e.* to avoid competence for nutrients or a specific niche). For example, Festa *et al.* (2016) found that the inoculation of exogenous microorganisms caused significant changes in the diversity of the microbial community, which can compromise the subsequent functionality of the soil. On the other hand, Kaur and Reddy (2013) demonstrated that native bacterial strains associated with maize crop have the ability to facilitate microorganism-plant interaction in order to enhance plant growth parameters, such as shoot length, germination, shoot, and root dry weight, as well as an increase in crop yield, by the production of phytohormones, siderophores, and solubilization of phosphates. However, the bacterial inoculant success in the field is not assured, due to the influence of several important factors such as climatic conditions, native microbiota, and its formulation (Dutta and Podile, 2010; de los Santos-Villalobos *et al.*, 2018). The introduction of exogenous PGPB could induce at least a transient perturbation of the equilibrium of soil microbial communities, this may be undesirable if important native species are lost, affecting subsequent crops (Trabelsi and Mhamdi, 2013). However, this effect could be buffered by ecosystem resilience, which is driven by the level of diversity and interactions of the plant-soil-biota (Kennedy, 1999).

At present, research works on native microbiota in the Yaqui Valley are very scarce (de los Santos-Villalobos *et*

al., 2018), even when this is the birthplace of the Green Revolution, and there are no microbial inoculants developed with native strains from this area, which have adapted to the edaphoclimatic conditions (high temperatures, alkaline and low content of organic matter). The aim of this study was to identify and characterize promising native PGPB associated with maize in the Yaqui Valley, as well as to quantify their inoculating impact on morphometric parameters of maize, grown under greenhouse conditions.

MATERIALS AND METHODS

Bacterial strains

The studied bacterial strains (22A67, 13B41, 22B45, and 31B11) were selected according to their ability to promote *in vitro* maize growth (unpublished data). These strains are cryopreserved in Colección de Microorganismos Edáficos y Endófitos Nativos (COLMENA, www.itson.mx/COLMENA) (de los Santos-Villalobos *et al.*, 2018). Glycerol-frozen (-80 °C) bacterial strains were pre-cultured in Petri dishes containing Nutrient Agar (NA) as a culture medium, and incubated for 24 h at 28 °C.

Morphological and molecular characterization

All bacterial strains were cultured in Nutrient Agar (NA) and incubated for 24 h at 28 °C to characterize them by macro (color, shape, margins, and elevation) and microscopical (Gram stains and shape) traits. DNA from each strain axenic bacterial culture was extracted using the UltraClean Microbial DNA Isolation Kit (QIAGEN®), the DNA integrity analyzed by 1% agarose electrophoresis while the quality and concentration by spectrophotometry (NanoMicro-Spectrophotometer Jenway®). To amplify the 16S rRNA gene (1500bp) we used the FD1 (5'-CCGAATTCGTCGACAACAGAGTTTGATCCTGGCTCAG-3') and RD1 (5'-CCCGGGATCCAAGCTTAAGGAGGTGATCCAGCC-3') primers (Weisburg *et al.*, 1991), in a 50 µL PCR reaction mixture containing 100 ng DNA, 1X PCR buffer, 0.2 µM primers, and 4U MyTaq DNA polymerase (Bioline). The PCR cycle was as follows: 5 min at 95 °C, followed by 30 cycles of 30 s at 95 °C, 40 s at 57 °C, 2 min at 72 °C, and a final elongation step of 5 min at 72 °C. PCR products were verified by electrophoresis gel agarose/TAE (2 %), and the purified amplicons sequenced by the Sanger platform. The edition and analysis of the obtained DNA sequences was with the software FinchTV 1.4.0 (Geospiza, Seattle, WA), CLC Sequence viewer 7 (QIAGEN, Denmark), and BLAST (NCBI, www.ncbi.nlm.nih.gov) used to identify the reported sequence with greater similarity to the studied bacterial strains. The phylogenetic analysis was generated with the MEGA software version 7.0 (Kumar *et al.*, 2016) using the neighbor-joining method, the evolutionary distances computed using the Tamura 3-parameter model. The partial sequences of the 16S rRNA gene were deposited in the Genbank database (MG807394, MG807395, MG807396, MG807397 accession numbers).

Plant growth promoting traits by bacterial strain

Indoles production. For quantification of indoles production, 1×10^3 Colony Forming Units (CFU) of each bacterial strain were inoculated, in triplicate, into 30 mL of nutrient broth (NB), supplemented with 100 ppm of tryptophan, and incubated in a rotary shaker (Yamato ®) at 150 rpm and 28 °C for 3 days. After incubation, the bacterial culture was centrifuged at 10,000 rpm for 10 min, and 1 mL of the supernatant was placed in a 1.5 mL Eppendorf tube and centrifuged again at 8000 rpm for 10 min according to de los Santos-Villalobos *et al.* (2013). Quantification of indoles was performed by the spectrophotometric method using Sal-kowski reagent according to Glickmann and Dessaux (1995). The concentrations of produced indoles were estimated with a standard curve of indole-3-acetic acid (Sigma®), in a range of 2-50 ppm. The tests were carried out in triplicate.

Phosphate solubilization. Each bacterial strain (1×10^3 CFU) was inoculated, in triplicate, on Petri dishes containing Pikovskaya (PVK) medium and incubated at 25 °C for 10 days. The PVK medium was composed of 2 solutions (salts and bromophenol blue Stock) according to Onyia and Anyanwu (2013). The ability of the studied strains to solubilize phosphorus (P) was observed by a transparent halo around the bacterial colony. The Phosphate solubilization index (PSI) was obtained according to Sharon *et al.* (2016), following the Equation 1:

$$\text{Phosphate solubilization or siderophore production index} = \frac{(\text{Halo zone} + \text{Colony diameter})}{\text{Colony diameter}}$$

Equation 1. Phosphate solubilization index or Siderophore production index

Siderophore production. To determine the ability of studied bacterial strains to produce siderophores, 1×10^3 CFU of each strain was inoculated on Petri dishes containing chrome azurol S medium (CAS), and incubated at 28 °C for 10 days, by three independent replicates (García-Meléndez *et al.*, 2017). The CAS medium preparation was following the method described by Alexander and Zuberer (1991). Strains that showed a transparent or colored halo around the bacterial colony were positive for this test. The siderophore production index was obtained following the Equation 1.

1-Aminocyclopropane-1-Carboxylate(ACC) deaminase activity. The ability of strains to produce ACC deaminase was screened in Petri dishes containing minimal medium according to Dworking and Foster (1958). 1×10^3 CFU of each strain was inoculated, by triplicate, in Petri dishes containing minimal media with ACC as its sole nitrogen source; then incubated at 28 °C for 3 days. Bacterial growth in the minimal medium was considered positive in ACC deaminase activity (nutrient medium was used as control).

In all the characterization tests mentioned above, *Bacillus* sp. P41B2 whose plant growth promoting traits are known was used as technical control.

Maize growth promotion by studied PGPB under greenhouse conditions

Bacterial production. A bacterial pre-inoculum was prepared for each strain (1×10^5 UFC), in tubes with 25 mL of Nutrient Broth (NB), incubated at 120 rpm for 24 h at 28 °C. Then, the pre-inoculum optical density adjusted to $OD_{630} = 0.5$ (Valenzuela-Aragon *et al.*, 2019). For the bacterial inoculum, 1 mL of pre-inoculum was inoculated to 3 L of NB, and incubated for 72 h, at 28 °C with shaking at 120 rpm. The bacterial suspension was pelleted by centrifugation for 15 min at 4000 rpm, washed twice and finally re-suspended in sterile distilled water. The bacterial concentration (CFU/mL) was adjusted to 1×10^9 CFU/mL.

Greenhouse assay. The greenhouse assay was carried out in 4 L pots under greenhouse conditions at the Instituto Nacional de Investigaciones Forestales, Agrícolas y Pecuarias (INIFAP) located at Cajeme, Sonora (27°N, 110°W). The max/min temperature during assay was 30 °C/8 °C and 12-hour photoperiod. CARIBU® (ASGROW®) commercial hybrid maize was used. A non-sterile mixture of perlite/agricultural soil/Pro-Mix GTX® substrate (1:3:2) was used as a plant substrate. The soil mixture was classified as clay-loam, with the following characteristics: organic matter = 5.48 %, pH in $CaCl_2 = 7.07$, electrical conductivity = 2.903 dS/m, total salts = 1858 ppm, N = 217 Kg/Ha, P = 15 ppm. In each pot, 300 mL (1×10^9 CFU/mL) of each bacterial strain were applied separately to each plant, 10 replicates per treatment and an un-inoculated control were established. Plants were kept well-watered, and no chemical fertilizers were applied during the assay (Parra-Cota *et al.*, 2014).

Measurement of maize growth parameters. Data collection was weekly over 6 weeks and at the end of the assay (42 days after sowing). The first measurement was 14 days after sowing between stages V3 and V4 of maize growth. Plant height, leaf chlorophyll level (SPAD 502 Minolta®), and leaves number were measured, and at the end, height, plant and root dry weight, and stem thickness.

Statistical analysis

A completely randomized design was applied in all experimental assays with three replicates per treatment, in the greenhouse assay a replicate was constituted by 10. Data analysis was by one-way analysis of variance (ANOVA) test and Tukey–Kramer test ($p=0.05$), using Statgraphics Centurion XVI.II (Statgraphics Technologies, Inc., Virginia).

RESULTS AND DISCUSSION**Morphological and molecular characterization of studied bacterial strains**

The macroscopic traits of all strains were similar in shape (circular), margin (entire), and elevation (flat) but different colors. Cell shape varied from coccus for strain 22A67 to rod-shaped for the other strains. Similarly, Gram stain was different between the bacterial strains (Table 1). Based on the 16S rRNA gene sequence phylogenetic analysis (Figure 1), the identifications were as follows: strain 22A67 as *Advenella incenata* (MG807395, accession number), strain 22B45

Table 1. Morphological characteristics of studied PGPB.**Tabla 1.** Características morfológicas de las BPCV estudiadas.

Strain	Macroscopic traits		Microscopic traits	
	Morphological traits	Color	Cell shape	Gram reaction
13B41	Circular with entire margin and flat	Brown	Rod-shaped	Positive
22A67	Circular with entire margin and flat	Light Brown	Coccus	Negative
22B45	Circular with entire margin and flat	Yellow	Rod-shaped	Negative
31B11	Circular with entire margin and flat	Brown	Rod-shaped	Negative

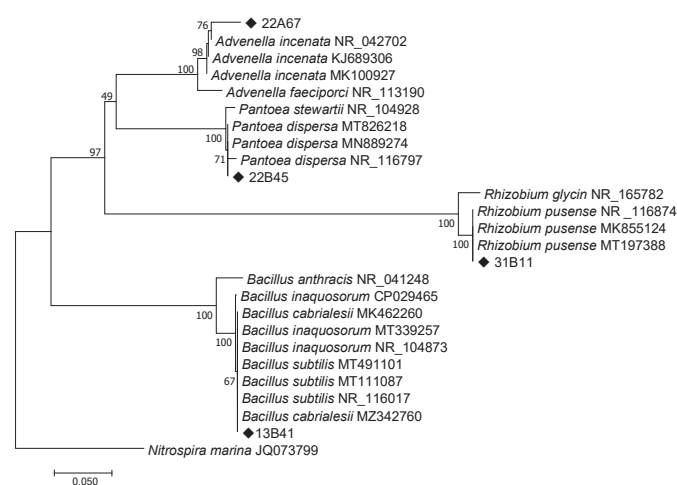
**Figure 1.** Phylogenetic Tree inferred using the Neighbor-Joining method. Numbers at nodes indicate values of bootstrap support based on an analysis of 1,000 resampled data sets. *Nitrospira marina* was used as an outgroup sequence. The scale bar indicates the number of base substitutions per site.

Figura 1. Árbol filogenético inferido mediante el método de *Neighbor-Joining*. Los números en los nodos indican valores de soporte de bootstrap basados en un análisis de 1000 conjuntos de datos remuestreados. Se utilizó *Nitrospira marina* como secuencia externa. La barra de escala indica el número de sustituciones de bases por sitio.

as *Pantoea dispersa* (MG807397), strain 31B11 as *Rhizobium pusense* (MG807394), and the strain 13B41 (MG807396) as *Bacillus* sp. In the latter, it was not possible to determine the species due to the sequences high similarity between different species of the genus *Bacillus*.

The four genera to which the studied bacterial strains belong have been previously described as PGPB. The *Bacillus* genus was first reported by Cohn in 1872 and currently includes more than 377 species, which are widely distributed in very diverse ecosystems such as aquatic, terrestrial, and those with extreme conditions (Villarreal-Delgado *et al.*, 2018; de los Santos Villalobos *et al.*, 2019). Several *Bacillus* species have been identified as plant growth-promoting bacteria since they suppress pathogens and/or improve plant development (Sansinenea, 2019). Bacteria of the *Rhizobium* genus have been extensively studied for their ability to fix nitrogen in symbiosis with legumes, but *Rhizobium* is also a

very good root colonizer of non-legumes and can produce plant hormones, siderophores, solubilize phosphorus, and exhibits adverse effects to plant pathogens (Qureshi *et al.*, 2013). At present, many research works have demonstrated the beneficial effects of *Rhizobium* inoculation to cereals, like maize, in terms of improved growth and yield (Mehboob *et al.*, 2012; Hussain *et al.*, 2016).

Pantoea is a relatively recent genus, proposed by Gavini *et al.* (1989). *Pantoea* spp. have been isolated from plants surfaces, within plant tissues, seeds, fruits, rhizosphere, water, soil, from humans and other animals; they are rarely considered as pathogens (Chauhan *et al.*, 2015). Various studies have reported strains of *Pantoea* as biological control agents and as growth promoter in crops such as corn, wheat, apple, cotton, barley, and potato (Montañez *et al.*, 2012; Kaur and Reddy, 2013; Chauhan *et al.*, 2015). Whereas the genera mentioned above has been widely studied, mainly *Bacillus* and *Rhizobium*, *Advenella* is a poorly studied genus, and reports of their role as PGPB are scarce, but these coincide in their potential role as phosphate solubilizers in soils with crops such as maize and wheat (Espinosa-Victoria *et al.*, 2009; Singh *et al.*, 2014; Wang *et al.*, 2020).

Plant growth promoting traits by studied bacterial strains

In order to have a better understanding of plant-microorganism interactions, it is important to study the PGPB potential mechanisms of action. The four studied strains were able to produce indoles, i.e., *Pantoea dispersa* 22B45 showed the higher production of indoles (11 ppm), followed by *Bacillus* sp. 13B41 (4.0 ppm), *Rhizobium pusense* 31B11 (3.8 ppm), and *Advenella incenata* 22A67 (2.8 ppm) (Table 2). Indole acetic acid (IAA) is generally considered as the most important native auxin (Glick, 2012); several species of bacteria are capable to produce this class of phytohormones, where their production vary depending on growing conditions, stage of development, availability of nutrients, among others (Duca *et al.*, 2014). The bacteria indoles production can affect the hormonal balance in plants and, therefore, can influence their growth, mainly increasing the total root surface, which leads to an enhanced mineral uptake from the soil (Spaepen and Vanderleyden, 2011). For example, in different studies, two native *Pantoea* strains associated with maize showed the ability to synthesize indoles in 121.6 ppm and 93 ppm, respectively (Montañez *et al.*, 2012; Kaur and Reddy, 2013). Similarly, Naveed *et al.* (2014) demonstrated that indoles-producing bacteria application can positively affect different growth parameters of maize, such as root dry weight and plant height, with increments of 43% and 8%, respectively, compared to un-inoculated control.

On the other hand, *Advenella incenata* 22A67 showed a high level of phosphate solubilization in PVK medium, similar to an *Advenella* strain with phytase activity reported by Singh (2014). *Pantoea dispersa* 22B45, *Bacillus* 13B41, and *Rhizobium pusense* 31B11, also showed the ability to solubilize phosphates, but at a lower level (Table 2). P is an

Table 2. Plant growth promotion traits by studied PGPB, under *in vitro* conditions.**Tabla 2.** Rasgos de promoción del crecimiento vegetal de las PGPB estudiadas, en condiciones *in vitro*.

Strain	Indoles production (ppm)	Phosphate solubilization Index (PSI)	Siderophore production Index (SPI)	ACC deaminase activity
<i>Bacillus</i> sp. 13B41	4.0 ^b	1.25 ^{ab}	1.88 ^b	-
<i>Advenella incenata</i> 22A67	2.8 ^a	1.58 ^b	1.40 ^a	-
<i>Pantoea dispersa</i> 22B45	11.4 ^c	1.42 ^{ab}	1.29 ^a	+
<i>Rhizobium pusense</i> 31B11	3.8 ^b	1.15 ^a	1.30 ^a	+
<i>Bacillus</i> sp. P41B2*	3.1 ^{ab}	1.28 ^{ab}	1.37 ^a	-

Different letters indicate significant differences ($p < 0.05$, Tukey-Kramer test), $n=3$. (+)= Production, (-)= No production. *= Technical control

important nutrient of plants due to it is involved in different functions, such as photosynthesis, respiration, and nutrient movement (Viruel *et al.*, 2014). There is enough P in soil, but only a small amount is available for plants, which limits their growth (Viruel *et al.*, 2014). The P availability depends on its solubility, which is influenced by the activity of roots and microorganisms in the soil, mainly through the production of organic acids and alternate mechanisms as the production of chelating compounds and secretion of phytase enzymes (Pande *et al.*, 2017). Thus, the studied strains can be classified as phosphate-solubilizing bacteria capable of solubilizing phosphate from an insoluble tricalcium source, which has shown a positive impact on biomass weights of tomato and maize crops, compared to un-inoculated plants (Montañez *et al.*, 2012; Sharon *et al.*, 2016).

Another important micronutrient for plants is iron (Fe), which influences several important biological processes such as photosynthesis, respiration, and chlorophyll synthesis (Kobayashi and Nishizawa, 2012). However, like phosphorus, this micronutrient has low availability in soils. Soil bacteria can overcome this Fe limitation by chelating molecules called siderophores (Villarreal-Delgado *et al.*, 2018), which production capacity provides an advantage against soil-borne pathogens (de los Santos-Villalobos *et al.*, 2012). Every studied strains showed the ability to produce siderophores, observing the higher value ($SPI=1.88$) for *Bacillus* sp. 13B41, significantly different compared to the rest of the strains (Table 2). In addition, bacterial genera like *Pantoea*, *Advenella*, *Bacillus*, and *Rhizobium* have shown this biochemical trait, also related to the growth promotion effect in maize and wheat cultivars (Kaur and Reddy, 2013; Kumar *et al.*, 2014; Singh *et al.*, 2014).

The activity of the enzyme 1-aminocyclopropane-1-carboxylate (ACC) deaminase, is of a plant growth promoter

mechanism by microorganisms, which regulate plant levels of ethylene (Glick, 2012). Ethylene is produced endogenously by plants and participates in the regulation of all processes for their growth, however, under environmental stress, its production is accelerated, negatively affecting root growth, and therefore, the development of the whole plant (de Souza *et al.*, 2015). Here, *Pantoea dispersa* 22B45 and *Rhizobium pusense* 31B11 showed ACC activity, like that reported by Shahzad *et al.* (2008), where strains of the genus *Pantoea* were identified with ACC activity and used to increase the yield of chickpea of up to 54 % compared to the un-inoculated control. Hussain *et al.* (2016) showed that by the inoculation of *Rhizobium* in maize, plants grew under drought stress conditions, obtaining an increase of 16 % and 30 % in dry biomass of root and shoot, respectively, compared to un-inoculated plants.

Maize growth promotion by isolated bacterial strains, under greenhouse conditions

In this study, all inoculated treatments showed a significant difference ($p \leq 0.05$) in plant chlorophyll level compared to the un-inoculated treatment, increased up to 40 % by *Bacillus* sp., 47 % by *Rhizobium pusense*, 41 % by *Pantoea dispersa*, and 39% by *Advenella incenata* at 42 days after sowing (Figure 1). However, there was no significant difference between plants inoculated with the studied strains. At 14 days after sowing, plant chlorophyll levels increased from 13 % to 20 % for all inoculated plants compared to the un-inoculated treatment (Figure 2). Increased chlorophyll content by the application of PGPB might be attributed to the enhanced root size of plants by phytohormones production, as well as the solubilization and mobilization of nutrients (Qasim *et al.*, 2017). Furthermore, the leaf chlorophyll level is markedly related to the photosynthetic capacity, which implicates a better nutrient cycling, indicating an improvement of plant growth (Croft *et al.*, 2016).

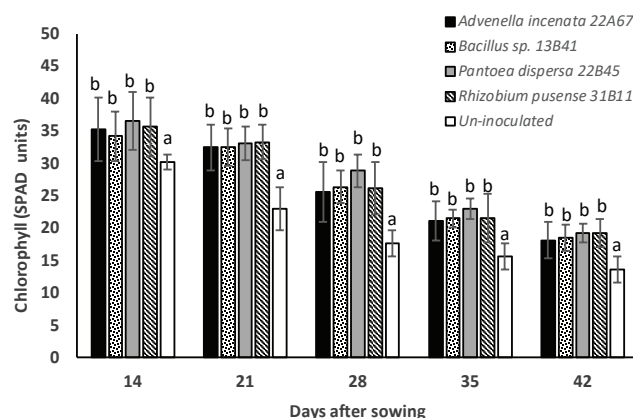
**Figure 2.** Chlorophyll level of maize plants inoculated by each studied strain and un-inoculated. Distinct letters indicate significant differences ($p < 0.05$, Tukey test), $n=10$.

Figure 2. Nivel de clorofila de plantas de maíz no inoculadas e inoculadas por cada cepa estudiada. Las letras distintas indican diferencias significativas ($p < 0.05$, prueba de Tukey), $n=10$.

Height of inoculated plants showed significant differences at 21 days after sowing, where strain *Advenella incenata* 22A67 showed a height of 37 cm, being different to the rest of the treatments [30.2 cm, 30.9 cm, 31.5 cm, and 28.2 cm for *Pantoea dispersa* 22B45, *Rhizobium pusense* 31B11, *Bacillus* sp. 13B41, and un-inoculated control, respectively]. The plant height on days 28, 35, and 42 after sowing, showed significant increases for all treatments (40, 51, and 35 %, respectively), compared to the un-inoculated treatment, and there was no difference between plants inoculated with the studied strains, as indicated in Figure 3.

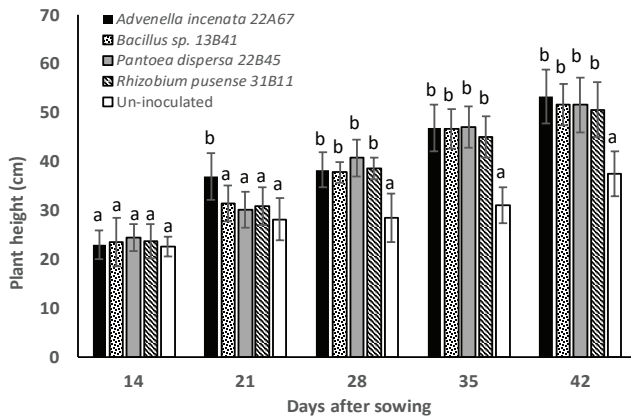


Figure 3. Height of maize plants inoculated with each strain. Distinct letters indicate significant differences ($p < 0.05$, Tukey Test), $n = 10$.

Figura 3. Altura de las plantas de maíz inoculadas con cada cepa. Las letras distintas indican diferencias significativas ($p < 0.05$, prueba de Tukey), $n = 10$.

In addition, the stem thickness showed a significant increment from 63% to 73% by studied bacterial strains (Table 3), compared to the un-inoculated treatment. This increment may relate to the accumulation of reserves in the stem during the vegetative stages, which could later increase the production of grains (Setter and Meller 1984; Seyed Sharifi *et al.*, 2017). On the other hand, the maize inoculation of the studied strains increased the number of leaves, but *Pantoea dispersa* 22B45 increased the number of leaves by 33 %, presenting at least one more leaf per plant than the rest of the strains and un-inoculated control (Table 3). Agbodjato *et al.* (2016) and Abiala *et al.* (2015) also reported an increase in the number of leaves in maize plants inoculated with PGPB under greenhouse conditions; in the first study the effect was observed by the inoculation of *P. fluorescens*; while in the second, the individual inoculation of *Bacillus*, *Lysinibacillus*, *Citrobacter* and *Enterobacter* caused this effect in plants. The increase in the number of leaves is clear evidence of the growth promoting effect generated by the PGPB.

Maize plants inoculated with *Pantoea dispersa* 22B45 also showed the highest root dry weight, 6.34 g, followed by maize inoculated with *Bacillus* sp. 13B41, *Rhizobium pusense* 31B11, *Advenella incenata* 22A67, and un-inoculated plants, as shown in Table 3. All inoculated plants showed a higher shoot dry weight and root dry weight compared to the un-inoculated treatment, with increments from 244 to 289 %,

Table 3. Effect of studied PGPB on maize growth under greenhouse conditions, at vegetative stage V8.

Tabla 3. Efecto de las PGPB estudiadas sobre el crecimiento del maíz en condiciones de invernadero, en la etapa vegetativa V8.

PGPB	Stem thickness (mm)	Number of leaves	Root dry weight (g)	Shoot dry weight (g)
<i>Advenella incenata</i> 22A67	13.7 ± 2.50 ^b	8 ± 0.79 ^b	5.45 ± 1.88 ^b	7.90 ± 3.21 ^b
<i>Bacillus</i> sp. 13B41	13.6 ± 1.71 ^b	8 ± 0.74 ^b	5.99 ± 2.10 ^b	7.61 ± 2.13 ^b
<i>Pantoea dispersa</i> 22B45	13.6 ± 1.51 ^b	9 ± 0.52 ^c	6.34 ± 1.85 ^b	7.89 ± 2.52 ^b
<i>Rhizobium pusense</i> 31B11	12.9 ± 2.38 ^b	8 ± 0.82 ^b	5.32 ± 1.94 ^b	6.99 ± 3.11 ^b
Un-inoculated	7.9 ± 1.37 ^a	6 ± 0.48 ^a	2.67 ± 0.81 ^a	2.03 ± 0.76 ^a

Mean with different letters in the same column show significant difference at $p < 0.05$ (Tukey-Kramer) $n = 10$.

and 99 to 137 %, respectively (Table 3). Silva *et al.* (2016) reported the capacity of *Pantoea* and *Bacillus* to increase shoot and root dry matter by 28 % and 18 %, respectively, in maize plants at 40 days after emergence compared to un-inoculated control. Qureshi *et al.* (2013) showed that the genus *Rhizobium* is an effective plant growth promoter in maize cultivars, due to its ability to improve the photosynthetic rate, up to 21 %. In addition, an increase in plant height and dry matter production was observed, these traits were attributed to the ability of *Rhizobium* strains to produce phytohormones. On the other hand, although there are few reports about maize growth promotion by *Advenella* (Espinoza-Victoria *et al.*, 2009; Singh *et al.*, 2014), this study shows that it can improve maize plant height from the first stages of development (Figure 4), besides increasing the dry matter content of maize.

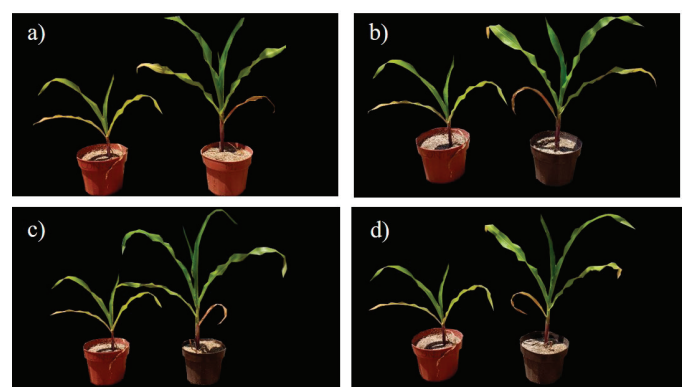


Figure 4. Effect of PGPB inoculation (right) on maize plant growth compared with un-inoculated plants (left). a) *Advenella incenata* 22A67, b) *Bacillus* sp. 13B41, c) *Rhizobium pusense* 31B11, d) *Pantoea dispersa* 22B45.

Figura 4. Efecto de la inoculación de PGPB (derecha) sobre el crecimiento de la planta de maíz en comparación con plantas no inoculadas (izquierda). a) *Advenella incenata* 22A67, b) *Bacillus* sp. 13B41, c) *Rhizobium pusense* 31B11, d) *Pantoea dispersa* 22B45.

CONCLUSION

Plant growth-promoting bacterial strains studied here showed significant plant growth promoting traits in maize plants (root weight, plant height, number of leaves, stem thickness), under greenhouse conditions compared to the un-inoculated plants; however, these beneficial effects were similar among strains. Thus, *Rhizobium pusense* 31B11, *Pantoea dispersa* 22B45, *Bacillus* sp. 13B41, and *Advenella incenata* 22A67, isolated from the Yaqui Valley, are promising bacterial strains to be evaluated in field trials in this area or in others with similar edaphoclimatic conditions, to test their ability to improve grain yield in maize, so that they can be used as native microbial inoculants for sustainable agriculture.

ACKNOWLEDGMENTS

To the Instituto Nacional de Investigaciones Forestales, Agrícolas y Pecuarias (INIFAP) for financial support through Project 2315932912 "Isolation and characterization of promising microorganisms to improve maize crop in southern Sonora and northern Sinaloa". And to the Consejo Nacional de Ciencia y Tecnología (CONACYT) for providing a fellowship (605488) to Carlos Fernando Amezquita Aviles.

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