



Agronomía Colombiana

ISSN: 0120-9965

ISSN: 2357-3732

Universidad Nacional de Colombia, Facultad de Agronomía

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Agronomía Colombiana, vol. 39, no. 1, 2021, January-April, pp. 78-89
Universidad Nacional de Colombia, Facultad de Agronomía

DOI: <https://doi.org/10.15446/agron.colomb.v39n1.88907>

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Rhizospheric rhizobia with potential as biofertilizers from Cuban rice cultivars

Rizobios rizosféricos con potencial como biofertilizantes a partir de cultivares cubanos de arroz

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ABSTRACT

Rice biofertilization with *Rhizobium* increases the growth and yield of the crop. However, evidence for this has not been observed in Cuban rice cultivars. This research aimed to typify two *Rhizobium* isolates, considering the use of different carbon sources, their tolerance to stress conditions, and the ability to promote the growth and development of rice. Two *Rhizobium* sp. isolates (Rpr11 and 5P1) were used and their facility to grow on different carbon sources, pH, and salinity levels was determined. The effect of the inoculation of these bacteria on the growth and yield of rice was evaluated under controlled, greenhouse, and field conditions. Both isolates grew on mannitol, glycerol, maltose, and fructose at the highest concentrations of NaCl (1.0, 1.5 and 2.0%). The isolate 5P1 grew at all evaluated pH levels, especially at pH 5.0 and pH 8.0. The inoculation of both isolates increased the plant biomass and the potassium content. The plants inoculated with 5P1 had the highest contents of nitrogen, total chlorophyll, carbohydrates and proteins, and grain yield. This study is the first in Cuba that shows the beneficial effect of *Rhizobium* inoculation on the physiology and yield of rice.

Key words: *Rhizobium*, grass, salinity, acidity, growth, yield.

RESUMEN

La biofertilización de arroz con *Rhizobium* incrementa el crecimiento y rendimiento del cultivo. Sin embargo, en cultivares cubanos de arroz no se han observado tales evidencias. El objetivo de esta investigación fue tipificar dos aislamientos de *Rhizobium* considerando el uso de diferentes fuentes de carbono, su tolerancia a condiciones de estrés y la capacidad de promover el crecimiento y el desarrollo del arroz. Se emplearon dos aislamientos de *Rhizobium* sp. (Rpr11 y 5P1) y se determinó su capacidad para crecer en diferentes fuentes carbonadas, pH y niveles de salinidad. Se evaluó el efecto de la inoculación de estas bacterias sobre el crecimiento y rendimiento del arroz en condiciones controladas, de invernadero y de campo. Ambos aislamientos crecieron en manitol, glicerol, maltosa y fructosa y en las mayores concentraciones de NaCl (1.0, 1.5, y 2.0%). El aislamiento 5P1 creció en todos los niveles de pH, especialmente en pH 5.0 y pH 8.0. La inoculación de ambos aislamientos incrementó la biomasa y el contenido de potasio en las plantas. Las plantas inoculadas con 5P1 mostraron un mayor contenido de nitrógeno, clorofilas totales, carbohidratos y proteínas, y rendimiento del grano. Este estudio es el primero en Cuba que demuestra el efecto benéfico de la inoculación de *Rhizobium* en la fisiología y el rendimiento del arroz.

Palabras clave: *Rhizobium*, gramínea, salinidad, acidez, crecimiento, rendimiento.

Introduction

Rhizobia are bacteria that establish a symbiotic association with leguminous plants. As a result of complex molecular communication between the bacteria and the plant, nodules are formed on the roots or stem where rhizobia perform biological nitrogen fixation (Taiz *et al.*, 2015). However, some studies reveal the beneficial effects of rhizobia on non-leguminous plants such as corn (*Zea mays* L.), lettuce (*Lactuca sativa*), tomatoes (*Solanum lycopersicum*) and wheat (*Triticum* spp.) (García-Fraile *et al.*, 2012; Flores-Félix *et al.*, 2013). Furthermore, rhizobia have been also found associated with rice (*Oryza sativa* L.)

as rhizospheric and endophytic bacteria. The rhizobia-rice interaction differs in many ways from that established with leguminous plants. These differences are fundamentally related to gene induction, cell-cell signaling, the infection process, and bacteria distribution inside the vegetable tissue (Chen *et al.*, 2015; Wu *et al.*, 2018).

Bradyrhizobium, *Sinorhizobium*, *Mesorhizobium*, *Azorhizobium* and *Rhizobium* are rhizobia genera that have been mostly associated with rice. These microorganisms increase the growth of grass mainly by phytostimulation (production of indole acetic acid and gibberellins) and enhance physiological mechanisms such as photosynthesis (Yanni

Received for publication: 6 July, 2020. Accepted for publication: 5 March, 2021.

Doi: 10.15446/agron.colomb.v39n1.88907

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et al., 2001; Chi *et al.*, 2005; Chen & Zhu, 2013). Previous studies report that rhizobia inoculation increases rice yield, even with lower doses of nitrogen than those recommended (Osorio Filho *et al.*, 2016, Lemes dos Santos *et al.*, 2019). However, studies about the potential of rhizobia strains to obtain inoculants that increase rice yields in Cuban cultivars and that decrease mineral fertilization are scarce.

In Cuba, rice is a prioritized crop since annual consumption is around 72 kg per capita, one of the highest in Latin America (Galán, 2017). However, imports are the main source for supplying the cereal in this country (FAO, 2019). Cuban rice cultivars INCA LP-5 and INCA LP-7 are widely distributed in the country (Galán, 2017) due to their positive characteristics that ensure high grain yields and resistance to diseases.

Only four reports about the rhizobia/non-leguminous crop interactions have been reported for Cuba. Two of them showed the beneficial effect of rhizobia inoculation on the growth and yield of corn. However, the rhizobia used in these trials were isolated from legume nodules and not from the crop's rhizosphere itself (Pérez-Pérez *et al.*, 2019). The other two studies explored the rhizobia-rice interaction but did not show the effect of rhizobia inoculation on the physiology and yield of rice (Hernández Forte & Nápoles García, 2017, 2019). Therefore, Cuban rice cultivar biofertilization with *Rhizobium* could be an alternative for reducing contamination and improving rice yield production and soil fertility in the country.

About 30% of Cuban soils and 15% of the country's agricultural area are affected by acidity and salinity, respectively. Around 104,000 ha dedicated to rice cultivation in Cuba are affected by salinity (Mesa, 2003). To increase productivity, mineral fertilizers are irrationally used (Goulding, 2016; Toledo, 2016). The use of tolerant microorganisms for stressful soil conditions could be an alternative for decreasing the application of mineral fertilizers and increasing rice yields. This research aimed to typify two *Rhizobium* isolates, considering the use of different carbon sources as a nutrient, their tolerance to stress conditions, and the ability to promote the growth and development of rice.

Materials and methods

Biological material

We used two isolates, Rpr11 and 5P1, belonging to the genus *Rhizobium* (accession numbers: MT387213 and MT759831, respectively) from the bacteria collection of the Laboratory

of Microbiology of the Department of Plant Physiology and Biochemistry at the National Institute of Agricultural Sciences (INCA), Cuba. Isolate Rpr11 was obtained from rice cultivar INCA LP-5 rhizoplane (Hernández Forte & Nápoles García, 2017) and isolate 5P1 from rhizospheric soil of rice cultivar INCA LP-7 plants. Both cultivars were cultivated under flood conditions, in a petroferic nodule ferruginous gleysol soil from Pinar del Río, Cuba (Hernández Jiménez *et al.*, 2015). The isolates were inoculated on 5 ml of yeast-mannitol (YM) medium (Vincent, 1970) and kept under shaking conditions at 150 rpm for 16 h at 30°C. The optical density (OD) ($\lambda = 600$ nm) of the inoculum was adjusted to 0.05.

Certified rice seed cultivars INCA LP-5 and INCA LP-7 were used in inoculation tests under controlled and greenhouse conditions. Seeds were disinfected and pre-germinated with 70% ethanol for 5 min and 6% (v/v) sodium hypochlorite for 30 min. Then, they were washed ten times with sterile distilled water, put onto plates with water agar medium (0.8%, m/v), and incubated at 30°C for 3 d in the dark (Hernández Forte & Nápoles García, 2019).

Rhizobium sp. growth on different carbon sources

Several multiplication tactics were carried out with the *Rhizobium* isolates. Microbial growth was determined in tubes with 4.5 ml of YM medium and in four variants where mannitol was replaced by maltose, lactose, glycerol and fructose (10 g L⁻¹). Every medium was inoculated with 500 μ l of inocula and incubated in a thermostated shaker (HEIDOLPH-UNIMAX-2010, Schwabach, Germany) at 150 rpm and 30°C for 24 h. The OD ($\lambda = 600$ nm) was measured every 2 h for 24 h. Five replicates of each isolate were used in each culture medium with different carbon sources.

Rhizobium sp. growth at different pH and salinity levels

Tubes containing 4.5 ml of YM medium with three pH levels (4.0, 5.0, and 8.0) and three concentrations of sodium chloride (1.0%, 1.5%, and 2.0%) were used. The tubes were inoculated with 500 μ l of inocula and incubated at 150 rpm at 30°C. Yeast-mannitol medium with pH 6.8 and 0.01% sodium chloride (Vincent, 1970) was used as a positive control. The inocula OD ($\lambda = 600$ nm) with different pH levels was determined every 2 h for 16 h, whereas the readings of inoculum with different concentrations of sodium chloride were performed every 2 h for 24 h. The pH and salinity of the culture medium were not controlled during the evaluation period. Three replicates were used for each isolate and for each pH and salinity condition.

Effect of *Rhizobium* sp. inoculation on rice growth

In vitro growth conditions

Disinfected and pre-germinated rice seeds were placed in pots with 0.21 kg of non-sterilized petroferric nodule ferruginous gleysol soil. Three seeds were placed in each pot. The chemical characterization of the soil showed a slightly acidic pH, medium organic matter content, high levels of phosphorus, adequate contents of calcium, magnesium, and sodium (Hernández *et al.*, 2015), and a low potassium level (Paneque Pérez *et al.*, 2010) (Tab. 1).

Rice seedlings of cultivars INCA LP-5 and INCA LP-7 were inoculated with 300 µl of *Rhizobium* sp. isolates Rpr11 and 5P1 at 5×10^9 colony-forming units (CFU) ml⁻¹. Rice seedlings inoculated with sterile YM medium were considered as a negative control. Ten pots were used for each treatment in a completely randomized design. The pots were placed in trays containing diluted Hoagland nutrient solution (1:2). The plants were grown in a 12 h light/12 h dark photoperiod at 26°C/22°C (day/night) and 70% relative humidity. At 7 d after inoculation (DAI), two plants were removed leaving only one plant per pot.

At 50 DAI the following variables were evaluated: plant height (cm), root length (cm), shoot dry weight (g), and root dry weight (g). The content of N, P and K in shoots and roots (%) was also determined from three samples of 0.2 g of the dry weight of each plant part per treatment. The samples were digested with sulfuric acid (H₂SO₄) and the color was subsequently developed with Nessler's reagent to determine the N content and with molybdenum blue for P and K (Paneque Pérez *et al.*, 2010).

Greenhouse conditions

Pots containing 1.2 kg of non-sterilized petroferric nodule ferruginous gleysol soil were used. The inoculation of the seedlings was carried out similarly to the test under controlled conditions. Two control treatments were used in the experiment. The negative control consisted of seedlings inoculated with sterile YM medium. As a positive control, the *Herbaspirillum seropedicae* Z67 strain was inoculated at the same volume and concentration as *Rhizobium*. Six pots were used for each treatment in a completely randomized design.

The plants were irrigated every other day with running water. Two plants were removed 7 DAI, leaving one plant per pot. At 70 DAI, the following variables were evaluated: plant height (cm), root length (cm), shoot dry weight (g), and root dry weight (g). The relative index of total chlorophyll content (SPAD) was measured in the flag leaf and other randomly chosen leaves, using a chlorophyll reader (SPAD-502, Konica Minolta, China). The total soluble carbohydrates (mg g⁻¹) were determined by the anthrone technique (Leyva *et al.*, 2008) and proteins (µg g⁻¹ fresh weight) were quantified by the microLowry method (Sun, 1994) in leaves and roots. In all cases, six replicates were evaluated per treatment.

Field trial

A field experiment was carried out at the Basic Technological-Scientific Unit "Los Palacios" in Pinar del Rio, Cuba (22°44' N, 83°45' W). The experimental area has petroferric nodule ferruginous gleysol soil (Hernández *et al.*, 2015) with the following chemical properties: pH (in water) of 6.46, 2.86% organic matter, 46.80 mg kg⁻¹ of P₂O₅ and 0.18 cmol_c kg⁻¹ of K⁺.

Rice plants were obtained from certified INCA LP-7 rice seeds. The seeds were sown in plastic trays (60 cm length x 30 cm width x 3 cm depth) with a 5400 cm³ mixture of petroferric nodule ferruginous gleysol soil and organic matter (1:1). Triple superphosphate (27 g m²), urea (7 g m²) and potassium chloride (4 g m²) were applied to the mixture. After 5, 10 and 15 d, urea and potassium chloride were applied again at the same concentration. One thousand six hundred plants were cultivated in each tray and irrigation was carried out with permanent watering.

Fifty plants were collected at 28 d, and the roots were embedded into the inoculum base of the selected *Rhizobium* strain (5×10^9 CFU ml⁻¹) for 10 min. Non-inoculated plants were used as a negative control of the experiment. The plants were taken to the plots (total area of 9 m²) and sowing was carried out with one plant per node, leaving 25 cm between plants.

Before transplantation and 10 d after transplantation, 20% of the recommended nitrogen fertilization (Ministerio

TABLE 1. Chemical characteristics of petroferric nodule ferruginous gleysol soil used in inoculation of rice plants.

pH (KCl)	OM (%)	P ₂ O ₅ (mg 100 g ⁻¹ soil)	Ca ²⁺	Mg ²⁺			K ⁺
				(cmol _c kg ⁻¹)			
6.57 ± 0.27	3.05 ± 0.70	75.1 ± 6.5	11.62 ± 0.97	4.75 ± 0.1	Traces	0.80 ± 0.02	

OM - organic matter.

de la Agricultura, 2014) was applied to the plants. Plants with 100% mineral fertilizer (Ministerio de la Agricultura, 2014) were used as positive controls and non-inoculated and non-fertilized plants were used as an absolute control. A randomized block design with three replicates for each treatment was used.

Fifteen plants were randomly selected and the tiller number was determined at 35, 42, 50, 57, 63, 71, 78, 85, 92, 99 and 105 d after transplanting. At 105 d, plants were harvested and the number of filled grains per panicle, the weight of 1000 grains (g), and yield ($t\ ha^{-1}$) (14% grain moisture) were determined.

Statistical analysis

Absorbance values obtained in the growth tests and multiplication dynamics at different carbon sources, pH levels, and salinity, as well as the data from the inoculation tests under controlled and greenhouse conditions were subjected

to the normality test (Bartlett test) and homogeneity of variance (Kolmogorov-Smirnov test). A simple classification analysis of variance was applied with the Tukey HSD (inoculation assay under controlled and greenhouse conditions) or Duncan (assay under field conditions) mean comparison tests for $P < 0.05$. The Statgraphic Plus program version 5.0 was used for statistical processing of the data and Microsoft Excel 2010 for its representation.

Results

Rhizobium uses different carbon sources as nutrients

The results showed that isolate Rpr11 growth in glycerol was lower than in the rest of the carbon sources at 8 h. However, the isolate showed increased growth in this alcohol after 20 h. The isolate Rpr11 showed lower growth in maltose and fructose than in mannitol from 12 h to 20 h after which no differences were found between mannitol and fructose (Fig. 1A).

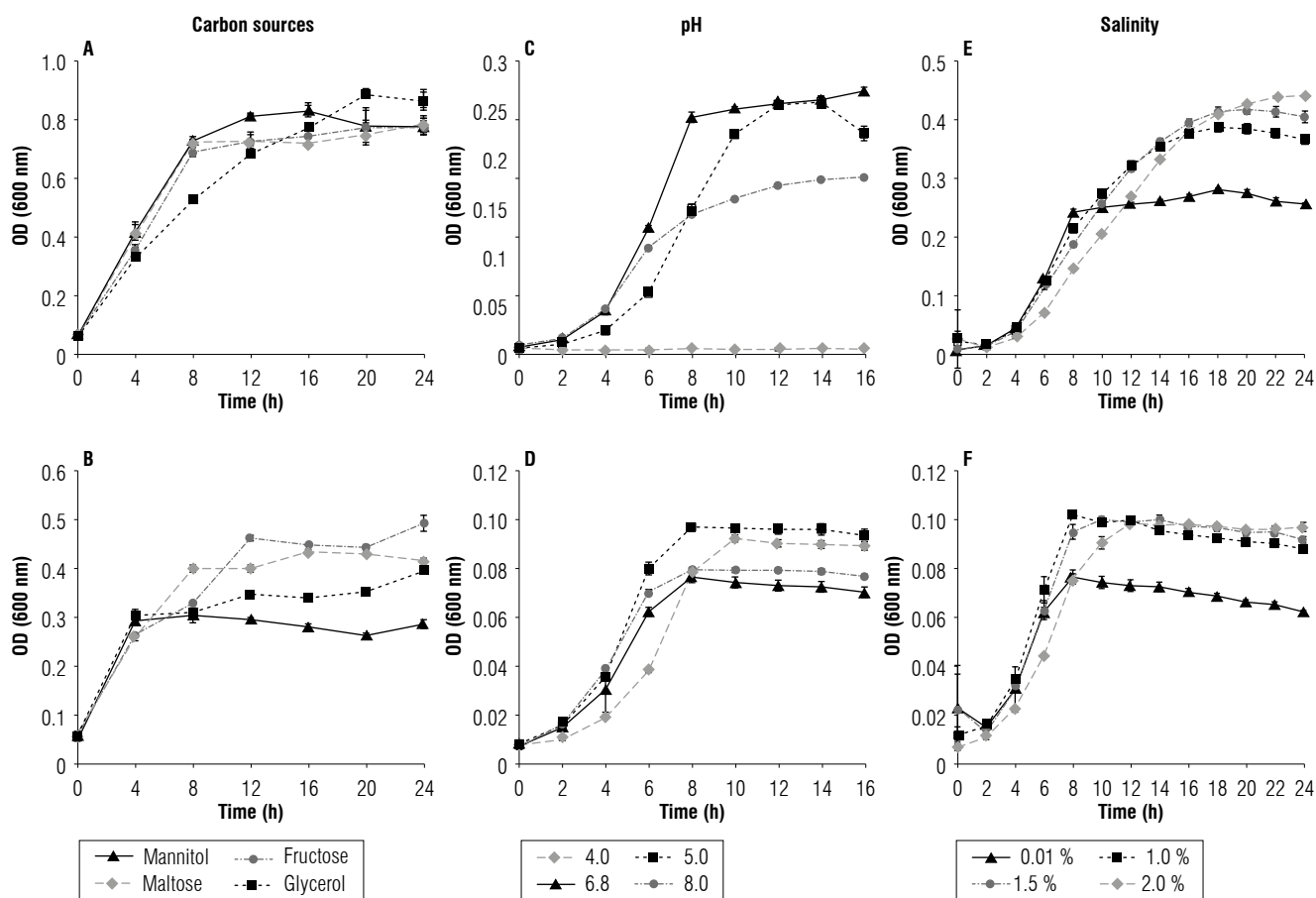


FIGURE 1. Multiplication dynamics of isolates Rpr11 (A, C, E) and 5P1 (B, D, F) in medium with different carbon sources, pH and NaCl concentrations. The data points and bars represent the means and standard errors of the mean from three replicates at each sampling time (Tukey HSD $P < 0.05$, $n = 3$). OD - optical density.

Mannitol and glycerol caused the highest growth of isolate 5P1 at 4 h. However, the bacteria increased the growth in fructose and maltose after 8 h. This isolate showed higher growth in fructose, maltose, and glycerol than in mannitol after 12 h. No differences were observed in the growth of 5P1 in fructose and maltose from 16 h to 20 h. This isolate showed greater growth in fructose than in the rest of the carbon sources at 24 h (Fig. 1B). It can be summarized that both isolates can use mannitol, glycerol, maltose, and fructose as carbon sources.

Rhizobium isolates tolerate acidity, basicity, and salinity conditions in the medium

The isolate Rpr11 showed the highest growth at pH 6.8 after 6 h. However, no differences were observed between pH 6.8 and pH 5.0 between 12 h and 14 h. The bacteria did not grow at pH 4.0 (Fig. 1C). The isolate 5P1 grew at the lowest pH levels, although this isolate could multiply at all the tested pHs. The most acidic condition allowed a better growth of this bacterium (Fig. 1D).

Regarding the dynamics at different salinity levels, isolate Rpr11 showed higher growth in 1.0% and 1.5% of NaCl than did the control treatment from 12 h. Similar behavior was observed after 14 h, when the isolate had the greatest growth at 1.0%, 1.5% and 2.0%. The lowest bacterial growth occurred at 2.0% NaCl, from 6 h to 10 h. However, this isolate had the greatest growth in the last two hours of the assay (Fig. 1E).

The isolate 5P1 showed a similar performance with increased growth in the most saline media after 10 h. No differences were displayed in the media with 1.0%, 1.5% and 2.0% of NaCl from 12 to 24 h (Fig. 1F).

Rhizobium promotes the growth of rice plants under *in vitro*, greenhouse and field conditions

Under controlled conditions, an inoculation assay was carried out to determine the effect of inoculation with isolates Rpr11 and 5P1 on rice plant growth. The results showed that plants inoculated with isolates Rpr11 and 5P1 increased the root and shoot dry weight, respectively (Tab. 2).

The inoculation of *Rhizobium* sp. isolates increased the nutrient content in the shoot of rice plants (Fig. 2). Plants inoculated with isolate Rpr11 showed an increase in the potassium content (Fig. 2A), while those treated with 5P1 showed a higher nitrogen and potassium content than non-inoculated rice plants (Fig. 2B).

On the other hand, the results of the inoculation assay with the isolate 5P1 on the cultivar INCA LP-7 under greenhouse conditions showed that the inoculation increased plant height, root length, shoot dry weight, and relative index of total chlorophyll, total soluble carbohydrates, and total soluble protein contents (Fig. 3, Tab. 3).

The plant inoculation with 5P1 and *H. seropedicae* Z67 produced a similar beneficial effect on rice growth (Fig. 3, Tab. 3). However, the plants inoculated with 5P1 showed a higher total carbohydrate content in leaves than plants inoculated with the reference strain *H. seropedicae* Z67.

Inoculation with 5P1 under field conditions showed a higher tiller number than the absolute control plants. Fertilized plants showed the highest tiller number at all evaluation levels (Fig. 4).

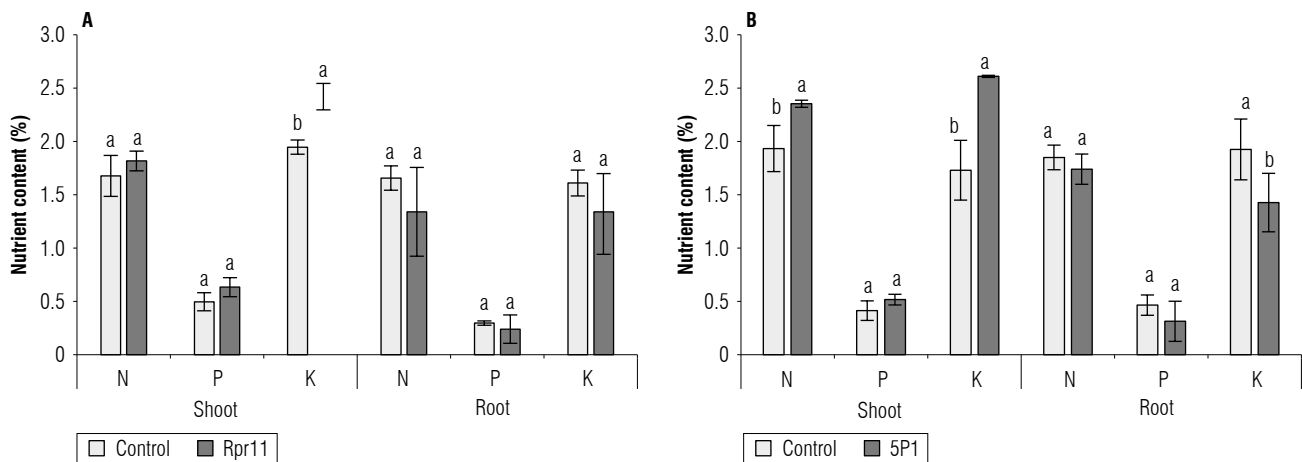


FIGURE 2. Effect of Rpr11 and 5P1 inoculation on the nutrient content of *O. sativa* L. A) cv. INCA LP-5 and B) INCA LP-7, respectively, at 50 d after inoculation (DAI) under controlled conditions. The bars represent the means \pm the standard errors of the mean from nine shoot and root nutrient content sample replicates according to the Tukey HSD test ($P < 0.05$, $n = 9$).

TABLE 2. Effect of Rpr11 and 5P1 inoculation on *O. sativa* L. cv. INCA LP-5 and INCA LP-7 growth at 50 d after inoculation (DAI) under controlled conditions.

Treatments	Height (cm)	Root length (cm)	Shoot dry weight (g)	Root dry weight (g)
INCA LP-5				
Control	53.2 ± 1.3 a	25.2 ± 1.2 a	0.61 ± 0.03 a	0.33 ± 0.02 b
Rpr11	52.5 ± 1.4 a	25.0 ± 1.1 a	0.69 ± 0.05 a	0.40 ± 0.03 a
INCA LP-7				
Control	59.3 ± 1.9 a	22.5 ± 2.0 a	0.33 ± 0.04 b	0.19 ± 0.03 a
5P1	62.1 ± 1.4 a	24.7 ± 2.1 a	0.43 ± 0.02 a	0.23 ± 0.02 a

Control plants were inoculated with sterile yeast-mannitol medium. Means with the same letter in the same column are not statistically different according to the Tukey HSD test ($P < 0.05$, $n = 10$).

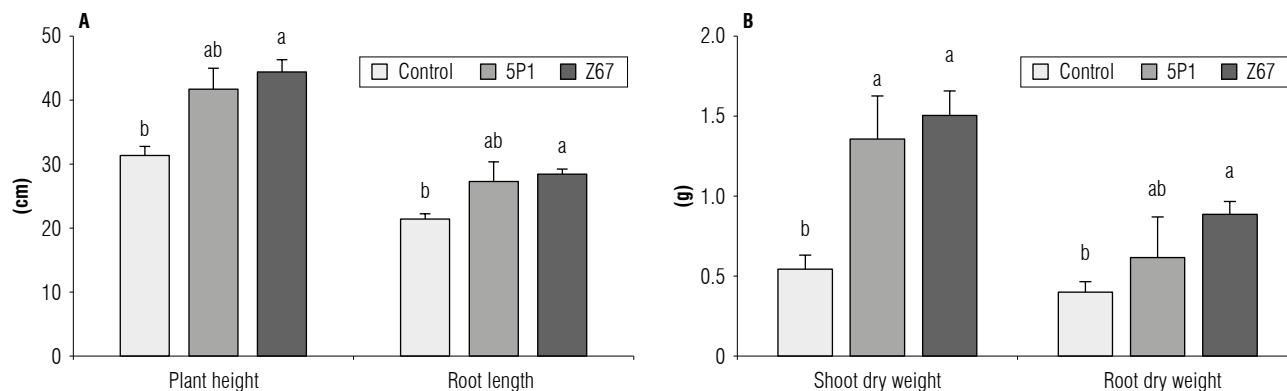


FIGURE 3. Effect of 5P1 and *H. seropedicae* Z67 (reference strain) inoculation on A) plant height and root length and B) shoot dry weight and root dry weight of *O. sativa* L. cv INCA LP-7, 70 d after inoculation (DAI) under greenhouse conditions. The bars represent the means ± the standard errors of the mean from six sample replicates (Tukey HSD $P < 0.05$, $n = 6$).

TABLE 3. Effect of 5P1 and *H. seropedicae* Z67 inoculation on chlorophyll, carbohydrate, and protein contents of *O. sativa* L. cv INCA LP-7, at 70 d after inoculation (DAI) under greenhouse conditions.

Treatments	Chlorophyll (SPAD)	Total soluble carbohydrates (mg g ⁻¹)		Total soluble proteins (µg g ⁻¹)	
		Leaf	Root	Leaf	Root
Control	24.1 ± 0.4 b	1.93 ± 0.04 b	6.39 ± 0.22 b	23.20 ± 0.46 b	8.71 ± 0.46 a
<i>H. seropedicae</i> Z67	26.2 ± 0.6 a	1.90 ± 0.08 b	7.83 ± 0.33 a	30.32 ± 0.46 a	8.71 ± 0.46 a
5P1	27.2 ± 1.4 a	2.36 ± 0.07 a	7.83 ± 0.33 a	32.37 ± 0.98 a	8.71 ± 0.46 a

Control plants were inoculated with sterile yeast-mannitol medium. SPAD - relative index of total chlorophyll content. Means with the same letter in the same column are not statistically different according to the Tukey HSD test ($P < 0.05$, $n = 6$).

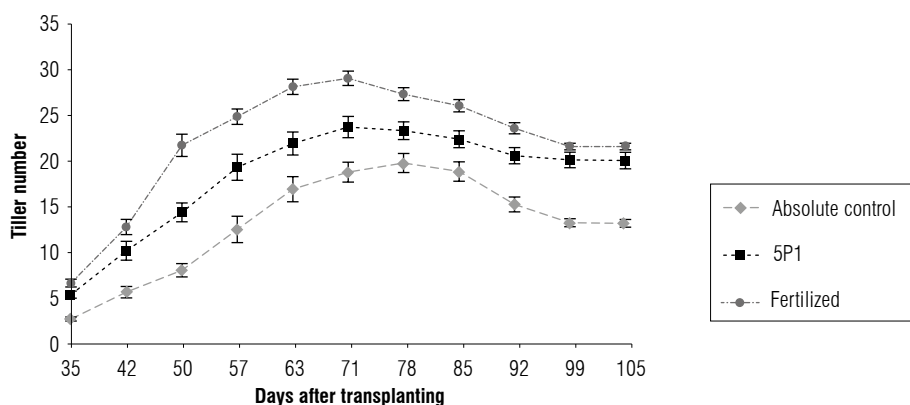


FIGURE 4. Tiller number of *O. sativa* L. cv. INCA LP-7 from 35 to 105 d after transplanting under field conditions. The absolute control treatment corresponded to non-inoculated plants with 50% mineral fertilization. The fertilized treatment corresponded to plants treated with mineral fertilization following the instructions of the Ministerio de la Agricultura (2014). The data points and bars are the means and standard errors of the mean from 15 replicates at each sampling time according to the Tukey HSD test ($P < 0.05$, $n = 15$).

TABLE 4. Effect of 5P1 inoculation and mineral fertilization on 1000 seed weight and crop yield of *O. sativa* L. cv. INCA LP-7, at 105 d after transplanting under field conditions.

Treatments	Grains filled per panicle	1000 grain weight (g)	Grain yield (t ha ⁻¹)	Increase (absolute control) (%)
Absolute control	30.8 ± 0.9 c	29.9 ± 0.8 a	1.0 ± 0.1 c	-
Fertilized	111.7 ± 2.5 a	29.2 ± 0.4 a	7.7 ± 0.2 a	87.0
5P1	103.5 ± 1.8 b	28.2 ± 0.8 a	6.4 ± 0.1 b	84.4

Absolute control - non-inoculated and non-fertilized plants. Fertilized plants - 100% mineral fertilization, according to the instructions of the Ministerio de la Agricultura (2014). Means with the same letter in the same column are not statistically different according to the Duncan test ($P < 0.05$, $n = 15$).

The maximum tiller number of fertilized plants inoculated with 5P1 was observed at 71 d after transplanting, while in the absolute control plants it occurred at 78 d. Rice tillering decreased in all treatments and remained constant from 99 to 105 d after transplanting.

No differences were observed in the weight of 1000 grains between treatments. However, plants inoculated with 5P1 increased the number of filled grains per panicle and the grain yield by more than 80% compared to the control. The fertilizer treatment allowed the greatest increases of filled grains per panicle and grain yield (Tab. 4).

Discussion

Rhizobium sp. isolates Rpr11 and 5P1 are able to use different carbon sources, tolerate stress conditions, and promote growth of rice plants. Nutritional studies of microorganisms are essential for designing culture media that meets their physiological needs, allowing a greater multiplication of the active ingredient in the inoculum. Viability, effectiveness, and efficiency are the most important characteristics of these bioproducts (Praveen Biradar & Santhosh, 2018).

The rhizobia inoculum requires a minimum bacterial concentration of 10^8 CFU ml⁻¹ or g⁻¹ (De Gregorio *et al.*, 2017). The carbon source choice is essential to increase bacterial growth and keep its viability in the inoculum. This is especially relevant for heterotrophic bacteria such as rhizobia (Zafar *et al.*, 2017).

The results showed the versatility of two *Rhizobium* sp. isolates for growing in four carbon sources. It has been reported that rhizobia strains isolated from wild legumes (*Genista microcephala* and *Argyrolobium uniflorum*) grow in different sugars as a carbon source (Dekak *et al.*, 2018). The use of different sugars as carbon and energy sources constitutes advantages for the bacteria since they can survive saprophytically to compete in the rhizosphere colonization.

The multiplication dynamics suggest that *Rhizobium* sp. isolates differ in their ability to use carbon sources, regardless of the fact that both belong to the *Rhizobium* sp. group. This is especially evident when bacterial growth was compared in the glycerol, maltose and fructose. Several bacteria prefer some sugars over others as carbon sources, since they have the necessary enzymes to oxidize them in the culture medium. Recent studies show the diversity of *Rhizobium* strains for using different carbon sources (Degefu *et al.*, 2018; Dekak *et al.*, 2018).

The growth of isolates 5P1 and Rpr11 in fructose and maltose as carbon sources offers the possibility of using relatively cheap raw materials to replace mannitol, the main component of the YM medium. The use of refined sugar, which is rich in fructose, could be an alternative for the industrial production of inocula with both rhizobia isolates. Chemical or enzymatic processing of starch as the main reserve material in plants constitutes a maltose source for this purpose (Wang *et al.*, 2015).

Acidity and salinity affect rhizobia viability and the infection and colonization processes (Plá & Cobos-Porrás, 2015; Shahid *et al.*, 2018). However, inoculation with tolerant rhizobia improves the establishment of some crops in forest areas and perennial legumes such as pigeon pea (*Cajanus cajan*) (Manet *et al.*, 2016; Sethi *et al.*, 2019).

Most rhizobia grow optimally at a pH of 6-7. The methods used to isolate these bacteria confirm it (Koskey *et al.*, 2018). Therefore, it is expected that the growth of both studied *Rhizobium* isolates would be lower at acidic pH than at pH 6.8. However, the growth of isolates Rpr11 and 5P1 at pH 5.0 and the isolate 5P1 at pH 4.0 demonstrates their tolerance to acidity. Usually, *Rhizobium* is a bacterium that acidifies the culture medium as it grows, and the pH of the culture medium may decrease with its growth. Therefore, isolates such as 5P1 could have higher tolerance to acidity conditions. Recent studies show the ability of *Rhizobium* to live under acid conditions (Pádua Oliveira *et al.*, 2017; Tullio *et al.*, 2019).

The synthesis of acid shock proteins and lipopolysaccharides and the proton exclusion to extracellular space explains the ability of bacteria to survive and multiply under acidic conditions (Geddes *et al.*, 2014; Hawkins *et al.*, 2017). This constitutes an additional advantage that allows plant growth-promoting bacteria (PGPB) to compete during rhizosphere colonization. This advantage is especially relevant because of rhizosphere acidification due to organic acids and proton production from root exudates (Conte & Walker, 2011).

The *Rhizobium* genus produces acids in the culture medium (Koskey *et al.*, 2018) that could explain the growth of isolates Rpr11 and 5P1 at pH 8. However, the results showed that these bacteria displayed greater growth at pH 5.0 than at pH 8.0. Previous studies indicate a similar behavior of rhizobia strains from *Desmodium triflorum* nodules (Bécquer *et al.*, 2017).

Salinity tolerance was another trait identified in the studied *Rhizobium* isolates. Around 9.30 % of Pinar del Río soils, the origin of these isolates, are affected by salinity (1.0% of salts approximately) (Mesa, 2003). In this research, the tolerance of two *Rhizobium* isolates to different NaCl concentrations (1.0, 1.5 and 2.0%) was studied. These concentrations are lower than those found in previous studies with rhizobia (Cardoso *et al.*, 2017; Franzini *et al.*, 2019; Nohwar *et al.*, 2019) but higher than those found in the studied soils. NaCl is not the only salt that contributes to soil salinity (Shao *et al.*, 2019); however, it is used in many PGPB characterization studies to determine the salt bacteria tolerance in the culture medium (Numan *et al.*, 2018; Jiang *et al.*, 2020).

Salinity decreases the colonization of roots by rhizobia (Tewari & Sharma, 2020). However, rhizobia tolerant to salinity could survive, grow, and effectively associate with their plant hosts (Yanni *et al.*, 2016). This seems to be the case of isolate 5P1 that had the highest values of optical density when it was cultured at the highest concentration of NaCl. Tolerance to salinity may be due to a plasmid-mediated resistance and salt resistance can be rapidly transferred from tolerant to sensitive bacteria (Kajić *et al.*, 2016).

Around 8000 ha dedicated to rice production in Cuba are affected by salt excess, a factor that decreases crop yield (Lamz Piedra & González Cepero, 2013). This is an opportunity to establish biofertilization strategies with salinity-tolerant microorganisms such as 5P1 and Rpr11. Isolating *Rhizobium* strains adapted to stressful conditions and increasing their concentration in these soils from

their inoculation could have a positive ecological effect on ecosystems.

Isolate Rpr11 increases the growth of rice plants of the cultivar INCA LP-5 (Hernández Forte & Nápoles García, 2019). So, in this research the effect of inoculation of this bacterium on the nutrient content of rice plants was determined. The positive contribution of isolate 5P1 inoculation on the plant rice cultivar INCA LP-7 growth was shown for the first time. The acidity tolerance of both *Rhizobium* sp. isolates could explain their establishment on the slightly acidic petroferic nodule ferruginous gleysol soil used, which is similar to the soil where both bacterial isolates originated (Hernández Forte & Nápoles García, 2017). The non-sterilization of the soil used would suggest diverse interactions between the inoculated bacteria and the resident microbiota, which is a very important aspect in plant-microorganism interaction (Čapek *et al.*, 2018).

Growth promotion in non-leguminous plants such as rice, sorghum (*Sorghum bicolor*), and corn (*Zea mays*) when inoculated with rhizobia is already known (Solaiman *et al.*, 2011; Bécquer *et al.*, 2012; Singh *et al.*, 2013). The production of indole acetic acid, gibberellins, and vitamins of the B group are some mechanisms that rhizobacteria use to increase plant height, shoot dry weight, and root dry weight in rice plants (Gopalakrishnan *et al.*, 2015). Some of these mechanisms could explain the positive effects of inoculation with isolates Rpr11 and 5P1 in rice plants cv. INCA LP-5 and INCA LP-7, respectively.

The increase of root dry weight in rice plants cv. INCA LP-5 inoculated with isolate Rpr11 under controlled conditions could have favored greater potassium absorption, mainly when its concentration was low in petroferic nodule ferruginous gleysol soil. The inoculation of rice plants with strains of rhizobia causes the modification of the roots, favoring the expanded root architecture (Yanni & Dazzo, 2015). This allows us to explore a larger reservoir of nutrients from the existing resources in the rhizosphere; and, thus, increase the absorption of nutrients and the dry weight of plants. Previous research reports that *Rhizobium* inoculation increases the nutrient content in plants since it promotes root growth and enhances the plant's ability to absorb it (Osorio Filho *et al.*, 2016). This last case could be the mechanism used by isolate 5P1 which increased the potassium and nitrogen content in the rice shoot without promoting root growth.

Taiz *et al.* (2015) report that chlorophyll synthesis is closely related to nitrogen availability in soil and to the

plant's ability to absorb it. Therefore, the increase of the nitrogen content in the shoot of rice plants cv. INCA LP-7 inoculated with isolate 5P1 could explain the increase of the chlorophyll content. A higher content of these molecules enhances photosynthesis, allowing the synthesis of carbohydrates (Degiovanni *et al.*, 2010), an effect shown with the inoculation of isolate 5P1 in rice plants cv. INCA LP-7 under greenhouse conditions. The enhancement of photosynthesis is one of the main mechanisms that explains the growth promotion of rice inoculated with *Rhizobium* (Chi *et al.*, 2010).

Photosynthesis also influences the synthesis of carbohydrates to provide a reduction power for nitrogen assimilation (Degiovanni *et al.*, 2010). The positive effect of isolate 5P1 inoculation on the chlorophyll content and nitrogen absorption could explain the increase in the content of total soluble protein in rice plants. The proteomic analyses show that the inoculation of rice plants with rhizobia induces the production of plant proteins that contribute to a better yield compared to non-inoculated plants (Chi *et al.*, 2010).

The nitrogen from the decomposition of soil organic matter enhanced with 5P1 could be one possible source of nitrogen that rice plants use to increase protein synthesis in leaves and roots. Previous studies demonstrate the positive effect of rhizobia inoculation as nitrogen-fixing bacteria on plant growth and yield. The inoculation of some *Rhizobium* strains allows an increase of dry matter in rice shoots with a decrease of 40% of the nitrogen dose (Osorio Filho *et al.*, 2016; Lemes dos Santos *et al.*, 2019). Therefore, the fixed nitrogen could also be another nitrogen source for rice plants inoculated with *Rhizobium* sp. isolate 5P1.

Regarding the effects of 5P1 inoculation on rice production areas in Cuba, the number of panicles and filled grains per panicle depends on the number of effective tillers, parameters that constitute some of the components that explain rice yield (Degiovanni *et al.*, 2010). Therefore, the positive effect on the number of tillers could explain the increase in grain yield obtained in plants inoculated with isolate 5P1.

In rice, tillering requires high amounts of nitrogen (Degiovanni *et al.*, 2010). This explains why fertilized treatment had the highest number of tillers and, therefore, the highest yield. Although the application of low doses of chemical fertilizer and the inoculation with the 5P1 strain did not surpass the fertilized treatment in any of the evaluated variables, they surpassed the control treatment. Similar results are described by Yanni and Dazzo (2010) who obtain a higher tillering in plants fertilized with N-fertilizers

compared to the treatments inoculated with rhizobia. However, the combination of both treatments was even more effective. Previous research confirms that the application of PGPB such as *Bacillus*, *Pseudomonas* and *Rhizobium* to rice and wheat (*Triticum aestivum* L.) increases the tiller number and yields with low nitrogen fertilization under field conditions (Tan *et al.*, 2015; Gusain & Sharma, 2019; Saber & Qader Khursheed, 2020).

The effect of *Mesorhizobium* sp. inoculation on rice has been previously studied. One study determined that the inoculation with this bacterium does not produce statistical differences between the control and inoculation treatments in the number of grains per panicle and grain yield when 60 kg ha⁻¹ were used (Hahn *et al.*, 2016). Other authors report that *Mesorhizobium* sp. associated with the recommended nitrogen dose provides the same rice grain yield as the recommended crop dose (Lemes dos Santos *et al.*, 2019). Rice plants inoculated with isolate 5P1 allowed a higher number of grains per panicle than those obtained with *Mesorhizobium* sp. The action of PGPBs on rice plants is variable since several factors may contribute to the different responses of rice to PGPBs inoculation, such as the bacterial strain, edaphoclimatic conditions, and the specificity of the plant genotype (Buzo *et al.*, 2019).

Yanni and Dazzo (2015) reported lower means of increased yield in five rice cultivars inoculated with *Rhizobium leguminosarum* bv. *trifolii* than those obtained with isolated 5P1 in rice cv. INCA LP-7. These authors also emphasize the importance of looking for adequate concentrations of N-fertilizers and inoculum to ensure a balanced supply of nitrogen, since an excess of this element favors the overproduction of extra non-reproductive tillers that do not contribute to rice yield.

Therefore, the results of this study show the potential of the isolate 5P1 as promising bacteria for making a biofertilizer to inoculate rice, especially under acidity and salinity conditions.

Conclusions

This research revealed the potential of *Rhizobium* sp. associated with a Cuban rice cultivar to use multiple carbon sources as nutrients to tolerate acidity, basicity and salinity conditions and promote rice growth. This study is the first in Cuba to show the beneficial effect of *Rhizobium* inoculation on the physiology, growth and yield of a Cuban rice cultivar.

Author's contributions

IHF formulated the overarching research goals and aims, developed the methodology, created the models, validated the overall replication/reproducibility of results/experiments and other research outputs, conducted the research process, and created and/or presented the published work, specifically the initial draft (including substantive translation). IHF and ORH curated the data for initial and later reuse. IHF and LAML performed the experiments or data/evidence collection. RPP implemented the computer programs and the statistical, mathematical, computational, or other formal techniques to analyze or synthesize study data. MCNG and RPP performed the critical review, commentary, or revision of the manuscript. MCNG led and managed the research activity planning and execution and acquired the financial support for the project leading to this publication.

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