



Co-inoculation of two symbiotically efficient *Bradyrhizobium* strains improves cowpea development better than a single bacterium application

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Abstract

The co-inoculation of *Bradyrhizobium* with other non-bradyrhizobial strains was already assessed on cowpea, but the co-inoculation of two *Bradyrhizobium* strains was not tested up to now. This study aimed to evaluate the cowpea growth, N accumulation, and *Bradyrhizobium* competitiveness of the elite strain *B. pachyrhizi* BR 3262 when co-inoculated with other efficient *Bradyrhizobium* from the Brazilian semiarid region. Three potted-plant experiments were carried out. In the first assay, 35 efficient *Bradyrhizobium* isolates obtained from the semiarid region of Brazil were co-inoculated with the elite strains *B. pachyrhizi* BR 3262. The experiment was conducted in gnotobiotic conditions. The plant growth, nodulation, N nutritional variables, and nodular occupation were assessed. Under gnotobiotic and non-sterile soil conditions, ten selected bacteria plus the elite strain *B. yuanmingense* BR 3267 were used at the second and third experiments, respectively. The cowpea was inoculated with the 11 bacteria individually or co-inoculated with BR 3262. The plant growth and N nutritional variables were assessed. A double-layer medium spot method experiment was conducted to evaluate the interaction among the co-inoculated strains in standard and diluted YMA media. The co-inoculation treatments showed the best efficiency when compared to the treatments inoculated solely with BR 3262. This strain occupied a low amount of cowpea nodules ranging from 5 to 67.5%. The treatments with lower BR 3262 nodule occupancy showed the best results for the shoot nitrogen accumulation. The culture experiment showed that four bacteria inhibited the growth of BR 3262. In contrast, seven strains from the soils of Brazilian semiarid region were benefited by the previous inoculation of this strain. In the second and third experiments, the results indicated that all 11 co-inoculated treatments were more efficient than the single inoculation, proofing the best performance of the dual inoculation of *Bradyrhizobium* on cowpea.

Keywords Biological nitrogen fixation · Dual *Bradyrhizobium* inoculation · Inoculant · Strain selection · *Vigna unguiculata* (L.) walp

Introduction

Cowpea [*Vigna unguiculata* L. (Walp)] is an important crop in the tropics. In Brazil, this species is grown mainly in family-based rainfed agricultural systems, mainly in the North and Northeast regions (Freire Filho 2011). In the last

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few years, the crop spread to Central Brazil, grown after the soybean in large and high technological farms (Batista et al. 2017; Silva Júnior et al. 2018). In the Brazilian semiarid region, cowpea is grown without fertilizer application during the short rainy season, and the average production is low, below 400 kg ha⁻¹ below than those in the North (around 800 kg ha⁻¹) and Central Brazil (above 1300 kg ha⁻¹) (IBGE 2019). Nevertheless, the development of low-cost and environmentally safe technologies is needed to improve cowpea yield in the Northeast Brazilian region.

The inoculation of cowpea rhizobia is a promising technology to improve cowpea production in the Brazilian northeast region (Marinho et al. 2014). Isolation and selection of native rhizobia have been reported in the field conditions in Brazilian drylands (Martins et al. 2003; Fernandes Júnior et al. 2012; Marinho et al. 2014, 2017; Xavier et al. 2017). More recently, the bioprospection of new cowpea rhizobia from the same region indicated the existence of high efficient *Bradyrhizobium* and *Microvirga* strains, showing better performance than those officially recommended strains for inoculant production (Oliveira et al. 2020; Sena et al. 2020).

Four *Bradyrhizobium* strains are authorized for inoculant production in Brazil (Brasil 2011). Among that highly efficient N₂-fixing and competitive strains, two have been extensively studied by our research group in the last few years: *B. yuanmingense* BR 3267 native from the semiarid region of Pernambuco state (Martins et al. 2003); and *B. pachyrhizi* BR 3262 isolated from an agroecological production system in Rio de Janeiro state. Both strains are highly efficient in Brazilian drylands (Marinho et al. 2014; Xavier et al. 2017). Besides its efficiency, BR 3262 strain also presents in vitro the ability to produce auxins (Menezes et al. 2016), a remarkable characteristic of plant growth promoter bradyrhizobia (Ferreira et al. 2020b). Although, the interaction of the elite strain *B. pachyrhizi* BR 3262 with members of the bradyrhizobial community of the soils of Brazilian drylands was not studied.

Inoculation of non-rhizobial plant-growth-promoting bacteria and rhizobia can increase the plant growth and nodulation of several legumes such as cowpea (Rodrigues et al. 2012), soybean (*Glycine max*) (Hungria et al. 2013, 2015; de Naoe et al. 2020; Moretti et al. 2020; Rondina et al. 2020), peanuts (*Arachis hypogaea*) (Ibáñez et al. 2009; Vicario et al. 2016), common bean (*Phaseolus vulgaris*) (Figueiredo et al. 2008; Hungria et al. 2013; Ferreira et al. 2020a) among others. Recently, the growth promotion and nitrogen fixation were improved by the co-inoculation of *Bradyrhizobium elkanii* 29w or *B. diazoefficiens* USDA 110^T and *Rhizobium tropici* CIAT 899^T in common bean (Jesus et al. 2018; de Carvalho et al. 2020). In these studies, the authors co-inoculated an agronomically efficient common bean *R. tropici* CIAT 899^T with

its non-preferred symbionts (*Bradyrhizobium* spp.). These results support that *Bradyrhizobium* could act as a plant-growth promoter, helping the common bean nodulation by its preferential microsymbiont (CIAT 899^T).

Compared to the single inoculation, the co-inoculation of two *Bradyrhizobium* strains was positively related to the growth, chlorophyll content (Vargas-Díaz et al. 2019), and nitrogen accumulation on soybean (de Carvalho et al. 2005) but not on cowpea (Xavier et al. 2017; Silva Júnior et al. 2018). In cowpea, the strains probably compete with each other and occupy the nodulation sites and are efficient in N fixation, whatever the nodule-occupying bacteria. In this case, it is not expected that the co-inoculation of two efficient and competitive *Bradyrhizobium* could increase the cowpea nodulation, growth, and N fixation. The plant growth promotion abilities of *Bradyrhizobium* on legumes (Jesus et al. 2018; Ferreira et al. 2020a) and non-legumes (Machado et al. 2016; Cavalcanti et al. 2020; Ferreira et al. 2020b) indicate that when co-inoculated with two *Bradyrhizobium*, the lower nodulating bacteria could act as an efficient plant growth promoter.

Therefore, we hypothesized that the co-inoculation of two efficient *Bradyrhizobium* could promote cowpea growth and nodulation better than a single strain inoculation. This study aimed to evaluate the cowpea growth, N accumulation, and *Bradyrhizobium* competitiveness of the elite strain *B. pachyrhizi* BR 3262 when co-inoculated with other efficient *Bradyrhizobium* from the Brazilian semiarid region.

Materials and methods

Bradyrhizobium strains and cowpea material

Bradyrhizobium spp. from the semiarid region of Brazil were used in this study. The strains were previously isolated, identified, and confirmed to be symbiotically efficient in different cowpea genotypes. The strains ESA 124, ESA 125, ESA 132, ESA 138, ESA 144, ESA 147, ESA 151, ESA 158, ESA 162, ESA 163, ESA 166, ESA 167, ESA 168, ESA 173, ESA 180, and ESA 192 were isolated and selected by Oliveira et al. (2020). The bacteria ESA 366, ESA 369, ESA 371, ESA 372, ESA 373, ESA 376, ESA 378, ESA 379, ESA 380, ESA 381, ESA 382, ESA 383, ESA 384, ESA 385, ESA 386, ESA 387, ESA 388, ESA 389 and ESA 390 were isolated and characterized by Sena et al. (2020). The strains *B. pachyrhizi* BR 3262 (Zilli et al. 2009) and *B. yuanmingense* BR 3267 (Martins et al. 2003) are elite strains recommended to the production of cowpea [*Vigna unguiculata* (L.) Walp.] inoculants in Brazil (Brasil 2011). The cowpea cv. BRS Pujante was used in the three plant experiments.

First plant experiment: co-inoculation of 35 *Bradyrhizobium* spp. and the elite strain *B. pachyrhizi* BR 3262 under gnotobiotic conditions

In the first experiment, co-inoculation of *B. pachyrhizi* BR 3262 with 35 strains *Bradyrhizobium* spp. obtained from the semiarid region of Brazil ("ESA" isolates above-mentioned) were evaluated in a greenhouse under gnotobiotic conditions.

The cowpea seeds were surface disinfected with ethanol 96% ($v v^{-1}$) for 30 s, sodium hypochlorite 2.5% ($v v^{-1}$) for five minutes followed by eight washes in distilled and autoclaved water (DAW) (Somasegaran and Hoben 1994). The experiment was implemented in 500 mL polystyrene pots filled with around 600 g of twice-autoclaved sand (120 °C and 1.5 atm for 1 h, with no less than 72 h between the sterilizations). The pots were disinfected by rinsing with sodium hypochlorite 2.5% ($v v^{-1}$). After disinfection, the pots were washed three times, with DAW, and filled carefully with the sterile sand. Then, three seeds were sowed per pot.

The bradyrhizobia grew in YM medium (Vincent 1970) in the constant stirring of 120 rpm for six days at room temperature (25 ± 3 °C) in an orbital shaker (Tecnal, TE-145, Brazil). After growth, the optical density was adjusted to 0.6 at 600 nm of wavelength ($OD_{600} = 0.6$) in a spectrophotometer (Thermo-Scientific, Multiskan GO, USA). After sowing, one milliliter of culture broth was inoculated over each seed. In the co-inoculated treatments, the seeds received 1 mL of *B. pachyrhizi* BR 3262 and 1 mL of the other bradyrhizobial strain. In the treatments inoculated only with *B. pachyrhizi* BR 3262 or *B. yuanmingense* BR 3267, 2 mL of the culture broths were inoculated over each seed. Besides the inoculated treatments, two uninoculated treatments were assessed: the "negative control" treatment, without inoculation or N application, and the "N-fertilization" treatment, with the application of 80 mg of $N-NH_4NO_3$ week $^{-1}$, from the second to the fifth week. In the sowing, the uninoculated treatments received 2 mL of sterile YM medium per seed.

The pots were supplied with 50–100 mL of DAW daily up to the cotyledon fall off, at 10 days after the emergence (DAE). At 10 DAE, the spare plants were cut and removed, leaving one plant per pot. From the 14 DAE to the end of the experiment, the plants received around 50–100 mL of DAW daily, and 100 mL of a nitrogen-free nutrient solution (Norris and Mannelje 1964) was applied once a week. The experiment was conducted between September to October of 2018. The plants were harvested at 45 DAE.

The roots were separated from the shoots and washed carefully with current tap water. The nodules were detached and counted. Roots, shoots, and nodules of each plant were placed separately in paper bags and left to dry in an air-flow chamber at 65 °C for 7 days and weighted for determination of shoot dry mass (SDM), root dry mass (RDM) and nodule dry mass (NDM). Shoots were also milled and

sieved (2 mm) for determination of the nitrogen concentration in the shoots (NCS) by the dry combustion method in a TruSpec CN elemental analyzer (Leco, USA) following the manufacturer instructions. These values were used to calculation of the nitrogen accumulation in the shoots (NAS) by the multiplication of NCS ($mg N g plant^{-1}$) to the SDM.

The competition of the BR 3262 strain was evaluated by the PCR amplification using strain-specific primers according to the methodology proposed by Osei et al. (2017) briefly described below. Ten nodules of the crown region of each cowpea root (each pot) were randomly selected in all inoculated treatment, totalizing 1480 nodules. The DNA of the nodules was extracted individually for all nodules. The extraction was conducted by manual maceration of each nodule in sterile 1.5 mL centrifuge tubes using sterilized plastic pestles. The DNA was stored at -20 °C until the performance of PCRs.

The PCRs were adjusted to 15 μ L with 1X buffer, $MgCl_2$ 1.5 mM, dNTP 0.75 mM, 0.10 μ M of the primers 2645-F (TAGAGGGCTGCTATCATGTC) and 2645-R (GAGATGATTACCGCAATGAG), 1 U of Taq DNA polymerase, and 2 μ L of nodule DNA as the template. To the PCR, an initial denaturation step of 94 °C for 5 min, followed by 35 cycles of denaturation (30 s, 95 °C), annealing (30 s, 60 °C), and extension (30 s, 72 °C) followed by a final cycle of extension of five minutes at 72 °C. The reactions were performed in a Veriti 96 well thermocycler (Applied Biosystems, USA). The PCR products were subjected to horizontal electrophoresis in 1% ($w v^{-1}$) agarose gel at 100 V for 60 min.

Each nodule was considered occupied by the BR 3262 strain by a clear amplicon around 200 bp, characteristic of the strain-specific amplification. In all reactions, one sample with the DNA of BR 3262 strain and other with DNA extracted by a nodule induced by BR 3262 in gnotobiotic conditions were as positive controls. The negative controls were the DNA of BR 3267 strain and other with DNA extracted by a BR 3267 nodule, in addition to the non-template control. As a quality control, in all rounds of DNA extraction, two nodules of cowpea inoculated with BR 3262 or BR 3267 (gnotobiotic conditions) were used, as positive and negative controls, respectively, to assure the correct extraction procedure.

Second and third plant experiments: co-inoculation of 11 *Bradyrhizobium* spp. and the elite strain *B. pachyrhizi* BR 3262 under gnotobiotic conditions and in non-sterile soil

In the second and third experiments, the strains ESA 124, ESA 162, ESA 168, ESA 173, ESA 192, ESA 369, ESA 376, ESA 380, ESA 386, ESA 387, and BR 3267 were selected to be assessed in single inoculation and under co-inoculation with the elite strain BR 3262. These strains were selected

because they had the best performance when co-inoculated with BR 3262 in first experiment. For both experiments, the bacteria grew in the YM medium and were $OD_{600} = 0.6$ adjusted, as described above. Besides, co-inoculated treatments received 1 mL of OD_{600} broth for each bacteria per seed. The treatments inoculated with one strain received 2 mL of the broth. For both assays, the cowpea seeds were surface disinfected, as described in the first experiment. The second experiment was conducted in a greenhouse under gnotobiotic conditions, as described for the first assay. The pots, substrate, and inoculation procedures were the same above-described. The experiment was conducted between January to February of 2019. The plants were harvested at 45 DAE.

The third experiment was conducted in forest-nursery using non-sterile soil as a substrate. The soil was collected in 0–0.2 m of a Ultisol in the Bebedouro Experimental Field (09°08'13" S; 40°18'24" W), in the dependencies of Embrapa Semiárido, in Petrolina, Pernambuco state. The chemical composition of the soil was assessed, according to Teixeira et al. (2017) (Table S1). 3-L polypropylene pots were filled with the soil (around 3.5 kg pot⁻¹), and the cowpea seeds were sowed after that. The plants were irrigated daily with 200–300 mL for each day of distilled water. This experiment was conducted between March and May of 2019, and the plants were harvested at 52 DAE. The N-fertilization treatment plants in this experiment received 100 mg of N-NH₄NO₃ week⁻¹, from the second to the fifth week.

The variables assessed in the second and third experiments were the shoot (SDM), root (RDM) and nodule number per plants (NN), nodule dry mass (NDM), the nitrogen concentration in the shoots (NCS) nitrogen accumulation in the shoots (NAS) according to above-mentioned.

Assessment of the *Bradyrhizobium* spp. growth pattern in a double-layer medium experiment

The synergy/antagonism between the 10 *Bradyrhizobium* spp. from the semiarid region of Brazil and the strain *B. yuanmingense* BR 3267 were assessed against the elite strain *B. pachyrhizi* BR 3262. We used the spot test method for this assay, according to Schwinghamer (1971), with modifications, briefly described below.

All bacteria grew in the YM medium for 5 days. Then, 1 mL of each bacterial broths were centrifuged at 6000 g for five minutes. The supernatants were discarded, and the pellets were resuspended in DAW. This procedure was repeated twice. Three aliquots with 10 µL were dropped in three equidistant points of YMA dishes. We used both original YMA and 1/5 strength YMA medium (reducing all nutrients and carbon source by 1/5). Therefore, this medium will be called oligotrophic YMA (oYMA). The use of the oYMA medium was applied to evaluate the bacterial behavior under standard

culture conditions (YMA medium) and oligotrophic conditions, close to the bacterial conditions when just inoculated in the sowed seeds.

The YMA and oYMA dishes inoculated solely with all bacterial broths (10 *Bradyrhizobium* spp., BR 3267, and BR 3262) were incubated at room temperature for 6 days. For the second layer, we produced YMA and oYMA media using a low melt point agar. All bacteria grew in YM medium, and the $OD_{600} = 0.6$ adjusted as described before, and the broths were mixed in the low-temperature YMA and oYMA media. After the mixture (OD_{600} adjusted broth + medium), the second layers were overlaid in the Petri dishes.

The inoculation strategy consisted of: (1) YMA medium at both layers—BR 3262 inoculated in the spots at the bottom layer overlaid separately with the 12 strains (BR 3262, BR 3267, and the ten strains isolated from the semiarid region of Brazil); (2) oYMA medium at both layers—BR 3262 inoculated in the spots at the bottom layer overlaid separately with the 12 strains; (3) YMA medium at both layers—the 12 strains inoculated in the spots at the bottom layer overlaid with BR 3262; and (4) oYMA medium at both layers—the 12 strains inoculated in the spots at the bottom layer overlaid with BR 3262.

The dishes were incubated for 6 days after the inoculation of the second layer. The evaluation, we observed the growth pattern of the bacteria in the second layer: Inhibition of the bacterial growth by the bacteria in the bottom layer (translucent zone surrounding the colonies of the bottom layer); No interference in the bacterial growth; Stimulation of the bacterial growth (intense growth in the edges or on the colonies of the bacteria in the bottom layer).

The experiment was conducted in three replications in a completely randomized design. The results were assessed in the three replications. The experiment was repeated twice, and the data compared. No discrepancies were observed in both assays.

Statistical analysis

The potted-plant experiments were set up in a completely randomized design with for replications per treatment. The normal distribution of data from greenhouse experiments was verified through of Shapiro–Wilk test. Data were evaluated through one-way ANOVA using the transformation $(x + 1)^{0.5}$ to NN, NDM, and TNC. The data of BR 3262 nodule occupancy in the first experiment, the transformation $\arcsin(x/100)^{0.5}$, was used. Following the ANOVA, the Scott–Knott average range test ($p < 0.05$) was applied.

The statistical analysis was carried out using the Sisvar software v. 5.0 (Ferreira 2011). The principal component analysis (PCA) carried with the second and third experiment compiled data was conducted with the correlation matrix in the PaSt v. 4.02 software (Hammer et al. 2011). NN and

NDM variables were not used in PCA to avoid bias due to the absence of nodules (gnotobiotic assay) and low nodulation (non-sterile soil assay) as intrinsic characteristics of the non-inoculated treatments. Before PCA, the data were normalized by subtracting the average of the treatment on each experiment.

Results

Co-inoculation of *B. pachyrhizi* BR 3262 and 35 *Bradyrhizobium* spp. in gnotobiotic conditions

The non-inoculated treatments did not nodulate, while all the plants of the 37 inoculated treatments nodulated equally (Table 1). The co-inoculation of BR 3262 and 28 *Bradyrhizobium* spp. improved the dry mass of cowpea shoots (15–64%) compared to the single elite strains BR 3262 and BR 3267. For the dry mass of roots, 30 co-inoculation treatments and the single inoculation of BR 3262 and BR 3267 were clustered in the highest statistical cluster, based on the Scott-Knott mean range test, highest than the N-fertilized and non-inoculated treatments.

The co-inoculation of BR 3262 with the strains ESA 151, ESA 158, ESA 167, ESA 168, and ESA 380 (74–82 mg N g plant⁻¹), in addition to the single inoculation of the elite strains BR 3262 (74 mg N g plant⁻¹) and BR 3267 (70 mg N g plant⁻¹) resulted in the same nitrogen concentration in the shoots then the N-fertilization treatment. The co-inoculation treatments of BR 3262 with the strains ESA 151, ESA 167, or ESA 380 showed the same averages for the variable N accumulation (143–166 mg N plant⁻¹) in the shoots when compared to the N fertilization (202 mg N plant⁻¹). To the same variable, in 13 co-inoculation treatments, the averages ranged from 100–166 mg N plant⁻¹, being superior to the single inoculation treatment with BR 3262 that reached 93 mg N plant⁻¹.

The co-inoculation of the native strains from the Brazilian semiarid region reduced the nodule occupancy of *B. pachyrhizi* BR 3262 in the co-inoculated treatments. The co-inoculation of the isolates ESA 162, ESA 163, ESA 167, ESA 382, and ESA 389 resulted in the lower occupation of the BR 3262 (5.0–12.5%). The co-inoculation of the isolates ESA 147 and ESA 373 allowed the highest nodule occupancy of the strain BR 3262, with an average value of 67.5% for both treatments.

Individual inoculation of 11 *Bradyrhizobium* isolates and their co-inoculation with BR 3262 under gnotobiotic and non-sterile conditions

The single inoculation of the elite strains BR 3262, BR 3267, and 10 selected cowpea *Bradyrhizobium* from the Brazilian

semiarid region induced high nodulation in all experimental replications (Table 2). The single inoculation of the bacteria ESA 124, ESA 162, ESA 168, ESA 173, BR 3262, and BR 3267 resulted in the same dry mass of shoot of the cowpea plants with mineral N. The same result was observed in all 11 co-inoculated treatments. The single inoculation of 10 *Bradyrhizobium* and 10 co-inoculated treatments improved the root dry mass of the cowpea plants when compared to both non-inoculated controls (improvements until 104%).

The co-inoculation treatments ESA 192 + BR 3262, ESA 387 + BR 3262, and BR 3267 + BR 3262 resulted in the same N in the shoots observed in the N fertilized treatment under gnotobiotic conditions. Furthermore, all inoculated and co-inoculated treatments resulted in a higher nitrogen concentration in cowpea shoots than the negative control. The plants of the treatments ESA 124 + BR 3262, ESA 162 + BR 3262, ESA 192 + BR 3262, ESA 380 + BR 3262, ESA 387 + BR 3262, and BR 3267 + BR 3262 accumulated high amounts of N in their shoots, with the averages ranging from 153 to 185 mg N plant⁻¹, and were comparable to the mineral N treatments (166 mg N plant⁻¹). All treatments were higher than the negative control treatment.

Twenty-two out of 23 inoculation treatments improved the cowpea nodulation in non-sterile soil, compared to the non-inoculated treatments. No differences were observed in the dry mass of shoots. The dry mass of roots on all co-inoculated treatments (1.18–1.54 g plant⁻¹) and the single inoculation of ESA 376 (1.26 g plant⁻¹), BR 3267 (1.51 g plant⁻¹), and BR 3262 (1.58 g plant⁻¹) was higher than the other treatments. The N content in cowpea shoots inoculated with ESA 168, ESA 192, and ESA 376, plus all the co-inoculated treatments, were the same observed in N fertilized plants and higher than found in the negative control. In except to co-inoculation treatment ESA 168 + BR 3262, all co-inoculated treatments and the single inoculation of ESA 168 or ESA 376, the nitrogen accumulation in the shoots was the same as that in plants supplied with mineral N, been higher than that in the negative control treatment.

In the principal component analysis, the PC1 and PC2 explained 83.08% of the variance (Fig. 1). The biplot clustered the co-inoculated treatments partially correlated to the treatments with a single inoculation. The N-fertilization treatment with mineral N was related to the co-inoculation. The negative control treatment was positioned apart from co-inoculation and N-fertilization treatments and partially related to the single inoculation cluster.

Bradyrhizobium spp. growth pattern in the double layer medium assay

The inoculation of the *Bradyrhizobium* spp. in the top layer of the medium (overlaid) with the elite strain *B. pachyrhizi*

Table 1 Shoot (SDM), root (RDM), and nodule (NDM) dry mass, number of nodules per plant (NN), nitrogen content in the shoots (NCS), nitrogen accumulated in the shoots (NAS), and nodule occupancy of *Bradyrhizobium pachyrhizi* BR 3262 of cowpea cv. BRS Pujante inoculated or co-inoculated with symbiotically efficient *Bradyrhizobium* at 45 days after the emergence

Inoculation treatment	SDM g plant ⁻¹	RDM	NN nod plant ⁻¹	NDM mg plant ⁻¹	NCS mg N g plant ⁻¹	NAS mg N plant ⁻¹	BR 3262 nod- ule occupancy (%)
ESA 124 + BR 3262	1.61a	1.86a	154a	122a	68b	110b	35.0d
ESA 125 + BR 3262	1.87a	2.76a	131a	126a	59c	109b	17.5d
ESA 132 + BR 3262	1.71a	2.04a	113a	115a	53c	91c	50.0c
ESA 138 + BR 3262	1.62a	3.74a	129a	123a	46d	76c	15.0d
ESA 144 + BR 3262	1.33a	2.39a	133a	118a	52c	70c	55.0c
ESA 147 + BR 3262	1.17b	1.07b	107a	108a	51d	60d	67.5b
ESA 151 + BR 3262	1.85a	2.31a	155a	123a	76a	143a	40.0c
ESA 158 + BR 3262	1.55a	2.25a	201a	144a	82a	128b	20.0d
ESA 162 + BR 3262	1.49a	1.38a	100a	103a	49d	73c	7.5e
ESA 163 + BR 3262	1.71a	1.22a	107a	101a	49d	83c	12.5e
ESA 166 + BR 3262	1.11b	2.01a	138a	113a	48d	51d	50.0c
ESA 167 + BR 3262	1.86a	2.19a	153a	131a	78a	146a	5.0e
ESA 168 + BR 3262	1.52a	1.90a	152a	125a	74a	112b	50.0c
ESA 173 + BR 3262	1.91a	2.76a	110a	120a	64b	124b	32.5d
ESA 180 + BR 3262	1.82a	1.63a	141a	127a	55c	100b	55.0c
ESA 192 + BR 3262	1.67a	1.61a	143a	124a	49d	83c	35.5d
ESA 366 + BR 3262	1.54a	1.09b	147a	132a	60c	91c	42.5c
ESA 369 + BR 3262	1.69a	1.43a	161a	140a	53c	89c	42.5c
ESA 371 + BR 3262	1.49a	1.32a	117a	102a	69b	105b	32.5d
ESA 372 + BR 3262	1.07b	1.85a	109a	116a	49d	51d	22.5d
ESA 373 + BR 3262	1.70a	1.71a	125a	113a	48d	82c	67.5b
ESA 376 + BR 3262	1.77a	1.61a	118a	113a	58c	102b	17.5d
ESA 378 + BR 3262	1.92a	1.87a	144a	130a	55c	105b	25.0d
ESA 379 + BR 3262	1.20b	1.62a	106a	88a	50d	62d	25.0d
ESA 380 + BR 3262	2.02a	1.78a	135a	112a	82a	166a	17.5d
ESA 381 + BR 3262	1.92a	2.48a	137a	129a	64b	126b	15.0d
ESA 382 + BR 3262	1.55a	1.28a	128a	129a	57c	87c	7.5e
ESA 383 + BR 3262	1.20b	1.16b	96a	94a	45d	54d	27.5d
ESA 384 + BR 3262	1.76a	1.39a	110a	119a	54c	96c	57.5c
ESA 385 + BR 3262	1.97a	1.56a	101a	97a	43d	86c	55.0c
ESA 386 + BR 3262	1.26b	1.17b	132a	100a	47d	59d	27.5d
ESA 387 + BR 3262	1.84a	1.53a	144a	129a	49d	92c	25.0d
ESA 388 + BR 3262	1.47a	1.24a	118a	121a	52d	76c	22.5d
ESA 389 + BR 3262	1.82a	2.34a	108a	124a	46d	84c	10.0e
ESA 390 + BR 3262	1.31b	1.02b	99a	100a	58c	76c	15.0d
BR 3262	1.23b	2.02a	118a	134a	74a	93c	100.0a
BR 3267	1.28b	2.01a	113a	123a	70b	90c	0e
N-fertilization*	2.30a	1.19b	0b	0b	88a	202a	–
Negative control	0.73c	0.23c	0b	0b	31e	23e	–
CV (%)	18.2	16.9	12.5	10.1	14.9	13.1	11.5

Experiment under gnotobiotic conditions. Data are an average of four replications

Means followed by the same letter, in the same variable, do not differ by the Scott–Knott mean range test ($p < 0.05$). CV coefficient of variation

*80 mg of N-NH₄NO₃ week⁻¹, from the second to the fifth week

Table 2 Shoot (SDM), root (RDM) and nodule (NDM) dry mass, number of nodules per plant (NN), nitrogen content in the shoots (NCS), nitrogen accumulated in the shoots (NAS) of cowpea cv. BRS Pujante inoculated individually with 12 *Bradyrhizobium* spp. or co-inoculated (dual inoculation) with *B. pachyrrhizi* BR 3262 and 11 *Bradyrhizobium* spp. of two experiments under the gnotobiotic condition (45 days after the emergence) and in non-sterile soil (52 days after the emergence) as substrates

Inoculation treatment	SDM g plant ⁻¹	RDM nod plant ⁻¹	NN nod plant ⁻¹	NDM mg plant ⁻¹	NCS mg N g ⁻¹	NAS mg N plant ⁻¹	SDM g plant ⁻¹	RDM nod plant ⁻¹	NN nod plant ⁻¹	NDM mg plant ⁻¹	NCS mg N g ⁻¹	NAS mg N plant ⁻¹
Non-sterile conditions experiment (soil as substrate)												
ESA 124	2.06a	1.66a	212b	307a	45c	94c	2.00a	1.06b	134c	197c	40b	82b
ESA 162	2.30a	1.29a	189b	289a	52c	119b	1.25a	0.89b	142c	219c	55b	64b
ESA 168	2.04a	1.22a	164b	208a	49c	106b	1.98a	1.12b	200c	277b	71a	150a
ESA 173	1.94a	1.30a	118b	173a	41c	79c	1.47a	0.62b	46d	96d	35b	50b
ESA 192	1.43b	1.22a	164b	222a	48c	69c	1.55a	0.93b	195c	259b	78a	121b
ESA 369	1.12b	1.14b	205b	167a	45c	51c	1.32a	0.93b	149c	221c	61b	82b
ESA 376	1.19b	1.34a	197b	302a	48c	57c	2.26a	1.26a	255b	381a	70a	165a
ESA 380	1.12b	1.21a	137b	132a	49c	55c	1.79a	0.99b	235b	378a	46b	89b
ESA 386	1.13b	1.13b	109b	166a	47c	53d	1.61a	0.81b	189c	286b	47b	77b
ESA 387	1.51b	1.23a	173b	244a	52c	78c	1.43a	0.57b	162c	298b	57b	83b
BR 3267	1.94a	1.38a	166b	172a	62b	118b	1.59a	1.51a	165c	162c	42b	67b
BR 3262	1.80a	1.50a	169b	206a	61b	111b	1.99a	1.58a	146c	205c	54b	113b
ESA 124 + BR 3262	2.42a	1.72a	170b	202a	63b	153a	1.90a	1.52a	315b	296b	76a	145a
ESA 162 + BR 3262	2.39a	1.85a	195b	190a	64b	154a	1.81a	1.54a	308b	513a	74a	132a
ESA 168 + BR 3262	2.13a	1.90a	191b	176a	65b	133b	1.21a	1.51a	268b	234c	90a	109b
ESA 173 + BR 3262	2.23a	1.33a	170b	172a	63b	141b	2.02a	1.42a	271b	270b	73a	149a
ESA 192 + BR 3262	2.48a	1.40a	172b	191a	67a	165a	1.99a	1.46a	284b	318b	77a	155a
ESA 369 + BR 3262	2.28a	2.22a	306a	333a	61b	139b	2.69a	1.26a	275b	285b	68a	194a
ESA 376 + BR 3262	2.26a	1.10b	156b	200a	56c	127b	2.18a	1.38a	291b	307b	81a	176a
ESA 380 + BR 3262	2.47a	1.26a	192b	211a	61b	152a	2.04a	1.20a	373a	417a	86a	177a
ESA 386 + BR 3262	2.18a	1.70a	192b	219a	54c	121b	2.08a	1.18a	417a	406a	95a	194a
ESA 387 + BR 3262	2.43a	1.36a	156b	186a	72a	172a	2.68a	1.29a	322b	290b	90a	247a
BR 3267 + BR 3262	2.43a	2.19a	266b	338a	76a	185a	2.44a	1.17a	149c	252b	60b	145a
N-fertilization*	2.62a	1.30b	0c	0b	64b	166a	2.08a	0.83b	6d	3d	85a	179a
Negative control	0.54c	1.09c	0c	0b	30d	34 e	1.44a	0.98b	34d	60d	28c	40c
CV (%)	13.7	16.5	18.2	13.4	12.2	14.7	13.9	12.9	16.4	10.4	13.4	18.2

Data are an average of four replications

Means followed by the same letter, in the same variable and in the same experiment, do not differ by the Scott-Knott mean range test ($p < 0.05$). CV coefficient of variation

* 80 mg of N-NH₄NO₃ week⁻¹, from the second to the fifth week; ** 100 mg of N-NH₄NO₃ week⁻¹, from the second to the fifth week

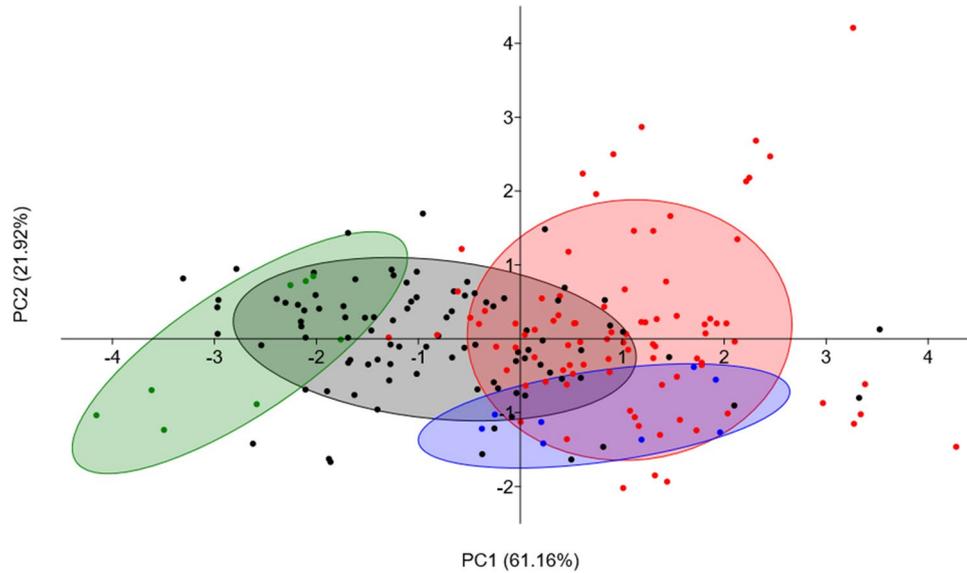


Fig. 1 Biplot of the principal component analysis (PCA) conducted with the data (all replications) of the experiments with the inoculation of 10 *Bradyrhizobium* spp. and the elite strains *B. pachyrhizi* BR 3262 and *B. yuanmingense* BR 3267, and co-inoculation of these 11 strains with and *B. pachyrhizi* BR 3262 under gnotobiotic (experi-

ment 2) and non-sterile soil (experiment 3) conditions. Red dots: co-inoculated treatments ($n=88$); Black dots: inoculated treatments ($n=96$); Blue dots: N fertilized control treatment ($n=8$); Green dots: non-inoculated and non-N fertilized treatment ($n=8$). PC1 and PC2 are the principal components one and two, respectively

BR 3262 previously inoculated in the spots of the bottom layer did not show any negative influence of BR 3262 on the 11 *Bradyrhizobium* spp. both in standard and in the oligotrophic YMA media (Table 3). In contrast, in the YMA medium, the strains ESA 124, ESA 162, ESA 168, ESA 173, ESA 192, ESA 376, and ESA 386 grew better surrounding the colonies of BR 3262 in the bottom layer. In the oYMA medium, the strains ESA 173, ESA 192, ESA 386, and BR 3267 also grew better under the influence of BR 3262 in the bottom layer. The other strains were not influenced by the strain BR 3262 in the conditions assessed.

Assessing the influence of the 11 strains in the growth pattern of BR 3262, antagonisms were caused by the strains ESA 124, ESA 162, ESA 168, ESA 173, and ESA 192 in the oYMA medium. In the traditional YMA medium, only the strain ESA 386 positively influenced the growth of BR 3262. The other ten strains did not influence the growth of the elite strain.

Discussion

The first experiment allowed us to assess the cowpea growth, nodulation, and N nutrition under the influence of two efficient rhizobial isolates, screening several combinations of efficient *Bradyrhizobium* isolates selected previously under interaction with BR 3262 and cowpea BRS Pujante. This screening showed that the co-inoculation

Table 3 Growth pattern of *Bradyrhizobium* spp. in the double layer spot test with two strains

Strain	BR 3262 in the bottom layer*		BR 3262 overlaid**	
	YMA	oYMA	YMA	oYMA
ESA 124	+	+	±	-
ESA 162	+	±	±	±
ESA 168	+	±	±	-
ESA 173	+	+	±	-
ESA 192	+	+	±	-
ESA 367	±	±	±	±
ESA 376	+	±	±	±
ESA 380	±	±	±	±
ESA 386	+	+	+	±
ESA 387	±	±	±	±
BR 3267	±	+	±	±
BR 3262	±	±		

+ Synergism (positive growth of the second layer near the first layer colonies); ± no influence of the strain in the bottom layer on the overlaid strain; - Antagonism (bacteria in the bottom layer inhibiting the growth of the overlaid strain). YMA = standard YMA medium; oYMA = 1/5 strength YMA medium

*BR 3262 in the spots of the bottom layer and the bacteria in the first collum overlaid (in this case, testing the antagonism/synergism of BR 3262 on the other strains);

**The strains in the first collum in the spots of the bottom layer and BR 3262 overlaid (in this case, testing the antagonism/synergism of the strains on BR 3262)

of two *Bradyrhizobium* improved the plant biomass and N accumulation as well as the single inoculation of elite strains of *Bradyrhizobium* BR 3262 and BR 3267, in agreement to previous reports on soybean (de Carvalho et al. 2005; Vargas-Díaz et al. 2019), but not on cowpea.

Research results showed that the co-inoculation of efficient strains of *Bradyrhizobium* spp. and *Microvirga vignae* did not result in the best plant growth and grain yield than the single inoculation of those strains in the Brazilian Semiarid region (Xavier et al. 2017). *Bradyrhizobium* + *Bradyrhizobium* co-inoculation did not improve the cowpea growth and development under organic farming systems in Minnesota, USA (Abou-Shanab et al. 2017). Our results agree with the previous soybean results reported by de Carvalho et al. (2005) in southern Brazil and Vargas-Díaz et al. (2019) in Mexico, being the first systematic report of co-inoculation benefits for cowpea with two *Bradyrhizobium* strains under potted-plant conditions.

The co-inoculation treatments with higher nodule occupancy of BR 3262 (67.5% when co-inoculated with ESA 147 and ESA 373) did not result in higher N accumulation in the shoots of cowpea when compared to the other treatments. On the other hand, in the treatments with higher N accumulation in the shoots, the BR 3262 strain nodule occupancy was lower than 40%, showing the highly competitive ability and efficiency of the isolates tested against the elite strain. Native strains of soils in semiarid regions are highly competitive to the nodule infection sites, even when competing against proven high efficient strains (Xavier et al. 1998; Martins et al. 2003; Mathu et al. 2012; Marinho et al. 2014).

Among the 35 bacteria assessed in this study, 34 belong to the *B. japonicum* genomic superclade (Oliveira et al. 2020; Sena et al. 2020), contrasting to the taxonomic position of *B. pachyrhizi* BR 3262 in the *B. elkanii* superclade. The biodiversity studies of cowpea in Brazil (Silva et al. 2012; Guimarães et al. 2015; Oliveira et al. 2020; Sena et al. 2020), Venezuela (Ramirez et al. 2020), and the African continent (Mohammed et al. 2018; Jaiswal and Dakora 2019) indicated the preference of cowpea to nodulate with the *B. japonicum*-like bradyrhizobia in the tropics, what probably influenced in the nodule occupancy of BR 3262.

However, BR 3262 is a strain authorized for commercial inoculants to cowpea; for this reason, this is highly efficient and competitive, highlighting to the other native *B. elkanii*-like bacteria. Despite being highly competitive, the nodulation of BR 3262, should be changed by the edaphoclimatic conditions that naturally occur in the Brazilian semiarid region. For example, the nodulation of Nepalese *B. elkanii*-like bacteria is altered by the soybean incubation and growth temperature (Suzuki et al. 2014). However, there are no data about the influence of the edaphoclimatic conditions in the Brazilian Semiarid region on the

dynamics of bradyrhizobial populations in soils, opening a new research question in this area. Nevertheless, the other two experiments data reinforce that our strains are competitive and more efficient under co-inoculation conditions on cowpea.

The compatibility of the strains in the double layer spot tests showed important findings of the co-inoculated strains interaction. Under oligotrophic conditions, somewhat those found in the seed surface just sowed in the soils, four out of 11 strains inhibited the BR 3262 growth. Some of these strains induced very low nodule occupation in the first assay. The reduction of the population of BR 3262 in the cowpea seeds should probably be partially responsible for the low nodule occupancy of the elite strains, as observed in the first experiment, and the performance of these strains observed in the other two experiments. The inhibition patterns of rhizobia observed in the double layer assays should be observed due to the production of antibiotics or bacteriocins by the strain in the bottom layer of the medium (Schwinghamer 1971; Goel et al. 1999). Further studies are needed to assure the nature of the antagonism observed in the present study.

In addition to antagonism against BR 3262, seven out of ten strains from the Brazilian semiarid region were benefited from the previous culture of BR 3262 in oligotrophic YMA medium. Together with the antagonism observations, these results indicate that, after the inoculation of two strains, even in the same concentration on cowpea seeds, both population dynamics decrease the proportional cell number of BR 3262. Nodulation of an inoculated strain in cowpea depends on the rate of active cells in the seeds and root primordia (da Silva et al. 2012). In this case, the nodulation of BR 3262, when co-inoculated with the other strains in the present study, is not favored. In addition to reducing the proportion of *B. pachyrhizi* BR 3262 active cells (by antagonism and increasing the cell density of the co-inoculated strain), this strain does not belong to the taxonomic superclade usually associated with cowpea in tropical conditions, as discussed before.

These findings support the hypothesis that, under co-inoculation conditions with the present study strains, this elite strain should act as a plant growth promoter, rather than the primary N₂-fixing strain in symbiosis with cowpea. Several *Bradyrhizobium* can promote the root growth by other mechanisms, even in non-nodulating species (Machado et al. 2016; Cavalcanti et al. 2020; Ferreira et al. 2020b). *Bradyrhizobium* also could act as an active common-bean growth promoter when co-inoculated with *R. tropici* CIAT 899^T (Jesus et al. 2018). BR 3262 as a high auxin producer in vitro (Menezes et al. 2016), a characteristic of plant-growth-promoting bradyrhizobial. The findings reported in the present study give the first evidence of a *B. elkanii*-like bacteria acting as a plant-growth promoter on cowpea.

The second and third experiments reinforced the co-inoculation advantage compared to the single inoculation

since the co-inoculation treatments induced the best plants. These observations were highlighted by the PCA given in Fig. 1 since the co-inoculation treatments were apart from the negative control treatments and closer to the N application than the single inoculation treatments. Multivariate analytical techniques are poorly exploited to experiments of inoculation of rhizobia and plant-growth-promoting experiments. These tools should help indicate inoculation and co-inoculation strategies for further studies (Vicario et al. 2016).

The production of commercial inoculants with two or more strains is required for soybean in Brazil (Brasil 2011), different from cowpea. The soybean recommendation is not to increase the inoculant performance, but to increase the probability of success of, at least, one strain in the different edaphoclimatic conditions in Brazil. According to the data observed in the present study, the recommendation to use two isolates in cowpea inoculant should be an approach to improve the field efficiency of the rhizobial inoculant. Therefore, field assays in different edaphoclimatic conditions are needed to assure the co-inoculation efficiency of cowpea.

Conclusions

Bradyrhizobium spp. isolated from soils of the Brazilian semiarid region, and the inoculant strain *B. pachyrhizi* BR 3262 interact and improve the growth and N nutrition of cowpea both in gnotobiotic and in non-sterile soil conditions, better than the inoculation of a single strain. The improved efficiency of the co-inoculation approach should be a promising strategy for developing more efficient cowpea inoculants.

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Author contributions LMVM, ADSF, and PIF-J conceived the research project. TRN, PTSS, GSO, MAMD, and TRS conducted the experimental work. TRN, ADSF, LMVM, and PIF-J analyzed the data. PIF-J wrote the first draft of the manuscript. All authors read and approved the final version of the manuscript.

Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval This paper does not contain any studies with human participants or animals.

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