

Rhizobacterium *Serratia* sp. and phosphorus fertilization rates affect aerobic rice development

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Abstract

Beneficial microorganisms participate in different processes that affect transformation of soil nutrients, including phosphorus (P), making them available for plants. However, there are few studies about the use of beneficial microorganism and P fertilization in the development of aerobic rice. The objective was to determine the effect of *Serratia* spp. isolate BRM 32114 and doses of P at sowing on the production of biomass, gas exchange, nutrient content in leaves and grain, grain yield and its components of aerobic rice. The field experiments were conducted over two growing seasons, 2015/16 and 2016/17, under a no-tillage system in the Brazilian Cerrado. The experimental design was a complete randomized block in a factorial 4x2x2 scheme, with four replications. The treatments consisted of four P₂O₅ doses (0, 40, 80 and 120 kg ha⁻¹) with or without rhizobacterium (2) in two growing seasons (2). Phosphorus rates increased the number of panicles and grain yield of aerobic rice, and the grain yield was higher in *Serratia* inoculation compared to no inoculation. The *Serratia* spp. strain BRM 32114 inserted in the production system of aerobic rice, under no-tillage, promoted gas exchange activity; enriched the K⁺, Ca⁺² and Mg⁺² content in leaves; enhanced biomass accumulation (13%), and resulted in a higher number of panicles per plant (11%) and grain yield (19%) than untreated plants. Therefore, this beneficial microorganism is a promising technology for potential incorporation into cropping systems in Tropical regions of the world.

Keywords: bioagent, *Oryza sativa*, fertilization, growth promoter, rhizobacteria, sustainable development.

Abbreviations: GPR_growth promoting rhizobacteria; PGPM_plant growth promoting microorganisms; A_photosynthesis rate; E_transpiration rate; gs_stomatal conductance; Ci_internal CO₂ concentration; Tleaf_leaf temperature; PPFD_photon flux density photosynthetic active.

Introduction

Rice is a staple food in many countries of the world (Kumar and Ladha, 2011). As the population is increasing, there is a need to increase rice grain yield to meet the people's demands. Average grain yields of aerobic rice are generally less than 2000 kg ha⁻¹ in most of the aerobic rice producing regions due to many environmental stresses and use of low inputs by farmers (Fageria and Baligar, 2003). However, aerobic rice is increasing worldwide as it is reducing the available water for irrigated rice (Kumar and Ladha, 2011). The soil fertility of tropical regions, such as those found in the Brazilian Cerrado, African Savannas and Asia, is limited mainly by the low natural availability of phosphorus (P) due to low nutrient contents in the forms available to plants or by high nutrient complexation rates, due to the chemical fixation reactions with soil particles (Fageria and Nascente, 2014). Almost 75 to 90% of added P fertilizer in agricultural soils is sorption and precipitated by iron, aluminum and calcium complexes present in soils (Turan et al., 2006). Thus, cultivation systems require constant and relatively large

amounts of P application to ensure the crops productivity. However, it also increases production costs. Several studies around the world have shown the positive effect of P fertilization in growth, yield components, and grain yield of rice (Crusciol et al., 2005; Fageria et al., 2013; Nascente et al., 2014).

Although the economic benefits of phosphorus fertilization on crop production is well described, excessive soil phosphorus is a potential threat to water quality. Phosphorus lost from agricultural soils, together with nitrogen, are usually, the nutrients that controls eutrophication of fresh waters. Eutrophication permits algal blooms, fish kills, harm wildlife and livestock by causing anoxia, or through the production of toxins, difficulty of drinking water purification (Mullins, 2009).

Soil microorganisms considered plant growth promoting microorganisms (PGPM) participate in different processes that affect the transformation of soil nutrients and play an essential role on soil nutrients cycle (Chen et al., 2006), such

as solubilizing unavailable soil P into soluble forms (Viverk and Singh, 2001; Sudhakara et al., 2002).

Among the bacterial genera with this capacity are *Achromobacter*, *Aereobacter Agrobacterium*, *Azotobacter Bacillus* ssp, *Bacillus circulans*, *Bradyrhizobium*, *Burkholderia*, *Enterobacter*, *Erwinia*, *Flavobacterium*, *Kushneria*, *Micrococcus*, *Paenibacillus*, *Pseudomonas*, *Ralstonia*, *Rhizobium*, *Rhodococcus*, *Salmonella*, *Serratia*, *Sinomonas* and *Thiobacillus* (Alori et al., 2017; Goldstein, 1986; Krishanaraj and Goldstein, 2001; Rodriguez and Fraga, 1999). Besides, previous studies showed that *Serratia* ssp, strain BRM 32114 promoted an increase in gas exchange, nutrients accumulation and biomass production, which differed from the control (untreated plants) in the aerobic rice plants, in greenhouse conditions (Nascente et al., 2017).

There are still few studies under field conditions for the use of rhizobacteria in crop development, especially in rainfed places and fewer in the Cerrado region. Therefore, the objective was to determine the effect of the *Serratia* spp. isolated BRM32114 and doses of P at sowing fertilization on the production of biomass, gas exchange, nutrient content in leaves and grain yield components and grain yield of aerobic rice, under a no-tillage system in a Cerrado Region.

Results

Leaf gas exchange evaluation

The presence of *Serratia* spp. strain BRM32114 provided changes in the gas exchange parameters in aerobic rice plants. However, there was no difference of these parameters among P rates (Table 2). For g_s , there was interaction between *Serratia* spp. strain BRM32114 and evaluation time and between *Serratia* spp. strain BRM 32114 and growing season. Hence, the plants showed greater ability to perform gas exchange in the early stages of their development, mainly in growing season 2016/17 (Table 3). For A, E, Ci and Tleaf, there was interaction between evaluation time and growing season. Aerobic rice plants in reproductive stage presented higher values of A, E and Tleaf in 2015/16; while, A and Tleaf values were higher in plants, at the vegetative stage, in 2016/17. Ci presented a significant difference only in the 2015/16 crop, and higher values were obtained in plants at the vegetative stage. The interaction between *Serratia* spp. strain BRM32114 and growing season showed that at the growing season 2015/16, treated plants presented higher E, Ci and Tleaf, and untreated plants presented higher Ci. In the growing season 2016/17, treated plants presented increase in g_s and Ci parameters. Tleaf was higher in untreated plants.

Nutrient content in rice leaves

The treatments with *Serratia* spp. strain BRM32114 positively affected K, Ca and Mg content in rice leaves (Table 4), and did not affect N, P, Cu, Fe, Mn and Zn content. Considering treated and untreated plants together, in growing season 2015/2016 rice leaves presented higher values of P, K, Mg, Fe and Zn. In the growing season 2016/17, rice leaves presented higher values of N, Ca, Cu and Mn. There was no effect of P rates in the nutrient content in rice leaves.

Plant biomass, yield components and grain yield of aerobic rice plants

The use of *Serratia* spp. strain BRM32114 provided increases in plant biomass, number of panicle per plant and grain yield of aerobic rice (Table 5). Number of grains per panicle and mass of 1000 grains was not affected by Rhizobacterium treatment.

Regarding growing season effect, in 2016/17, plants presented higher biomass, number of panicle per plant, number of grains per panicle and grain yield than growing season 2015/2016 (Table 5). There was no effect of P rates in the plant biomass, number of grains per panicle and mass of 1000 grains. On the other hand, there was an effect of P rate on the number of panicles (Fig 2) and an interaction between *Serratia* spp. strain BRM32114 and P rate for grain yield (Fig 3).

Nutrient content in rice grains

Plants not treated with *Serratia* spp. had a higher N, Cu and Zn content in rice grains (Table 6). In growing season 2015/16, the contents of P, K, Mg, Cu and Mn were higher.

Discussion

One of the environmental conditions that could positively affect the grain yield is the rain distribution (Heinemann and Stone, 2009). Total rainfall for the growing season 2015/2016 (692.8 mm) was higher than growing season 2016/2017 (678.2 mm). However, it was not equally distributed. There was less rain in the period from flowering to grain filling in the growing season 2015/2016 (195.4 mm) than in the growing season 2016/2017 (370.0 mm) (Fig 1). Associated with rainfall distribution, the greater soil fertility at 2016/17 (Table 1) could explain the higher rice grain yield in this season and other differences between these seasons. However, independent of rain and fertility conditions, there were an increase in plant biomass (13%), number of panicle per plant (11%) and grain yield (19%), in *Serratia* spp. strain BRM 32114-treated aerobic rice plants. The observed gain seems to be determined by A (photosynthetic rate) and g_s (stomatal conductance), which was increased by 6 and 28 %, respectively, in treated plants. The treatment with *Serratia* spp. strain BRM 32114 affected g_s more than A in aerobic rice plants, indicating that the gas exchange capability was potentiated in treated plants. The processes of grain and biomass production depends on the sink size, pre-anthesis dry matter, translocation of pre-anthesis assimilates to grain, nutrient accumulated and remobilized as well as assimilate source (light capture and photosynthetic rate) (Alvarez et al., 2014; Sandhu et al., 2015). Augé (2001), Bandara et al. (2006), Guo et al. (2008), Pérez-García et al. (2011), Barzana et al. (2012) and Nascente et al. (2017) have also shown that microorganisms can provide indirect stimulation in crop plants by promoting greater gas exchange activity. Regarding the nutritional status of the plants, it was observed that *Serratia* spp. strain BRM 32114 significantly increased the K, Ca and Mg content in rice leaves over the average of two growing seasons. Microorganisms can also provide indirect plant growth stimulation by improving the availability and absorption of nutrients (Pérez-García et al., 2011; Zhang et al., 2011). One of the hypotheses is that beneficial microorganisms modify

Table 1. Chemical and physical properties of the soil at the beginning of the trial.

Growing season 2015/2016							
Depth (m)	Ca	Mg	H+Al	SOM ¹	P	K	pH (H ₂ O)
0-0.20	18.0	13.3	24.0	32.7	9.9	101.3	5.9
Growing season 2016/2017							
Depth (m)	Ca	Mg	H+Al	SOM ¹	P	K	pH (H ₂ O)
0-0.20	10.7	10.9	31.0	42.22	12.6	97	5.5
Depth (m)	Cu	Zn	Fe	Mn	Clay	Silt	Sand
0-0.20	0.8	4.6	20.5	10.1	480	226	294
Depth (m)	Cu	Zn	Fe	Mn	Clay	Silt	Sand
0-0.20	1.2	8.1	31.0	12.2	520	180	300

¹SOM – soil organic matter.**Table 2.** Phosphorus rates and *Serratia* spp. isolated BRM 321145 with or without inoculation in rice seeds as affecting photosynthesis (A), transpiration (E), stomatal conductance (gs), internal CO₂ concentration (Ci) and leaf temperature (Tleaf) of rice plants in two growth stages at vegetative and reproductive phase of rice development. Santo Antônio de Goiás, growing season 2015/2016 and 2016/2017.

Factors	A	E	gs	Ci	Tleaf
<i>Serratia</i> BRM32114	μmol CO ₂ m ⁻² s ⁻¹	mmol H ₂ O m ⁻² s ⁻¹	mol H ₂ O m ⁻² s ⁻¹	vpm	°C
With	25.67 a*	6.00	1.27 a	287	29.92
Without	24.21 b	5.95	0.99 b	284	29.83
Growth stage					
Vegetative	26.37 a	6.03	1.19	295 a	29.01 b
Reproductive	23.50 b	5.91	1.06	276 b	30.74 a
Growing season					
2015/2016	23.32 b	6.01	0.51 b	256 b	31.89 a
2016/2017	26.55 a	5.94	1.75 a	315 a	27.86 b
ANOVA (F Probability)					
<i>Serratia</i> (S)	<0.001	0.7231	0.0119	0.2817	0.6771
Phosphorus rates (P)	0.0507	0.4303	0.9025	0.6709	0.9957
Growth stage (GSt)	<0.001	0.4438	0.2410	<0.001	<0.001
Growing season (GS)	<0.001	0.6603	<0.001	<0.001	<0.001
S x P	0.0525	0.4491	0.7360	0.6456	0.9947
S x GSt	0.8448	0.9490	0.0088	0.8498	0.5926
S x GS	0.4044	0.0302	0.0215	0.0002	0.0060
P x GSt	0.0501	0.9505	0.1322	0.9284	0.9433
P x GS	0.0612	0.8398	0.8200	0.7570	0.9099
GSt x GS	<0.001	<0.001	0.1263	<0.001	0.0094
S x P x GSt	0.5577	0.3154	0.8288	0.9252	0.9828
S x P x GS	0.0519	0.3703	0.7344	0.4490	0.9491
P x GSt x GS	0.7864	0.5527	0.1973	0.5308	0.9842
S x GSt x GS	0.0761	0.3444	0.0519	0.3578	0.3313
S x GS x GS x P	0.0524	0.5490	0.8342	0.6235	0.9858

* Means followed by the same letter do not differ by Tukey's test (P<0.05).

Table 3. Interaction between growing season and growth stage for photosynthesis (A), transpiration (E), internal CO₂ concentration (Ci) and leaf temperature (Tleaf) of rice plants at vegetative and reproductive phase of rice development, and with or without the use of rhizobacterium *Serratia* spp. strain BRM32114 with transpiration, stomatal conductance (gs), internal CO₂ concentration and leaf temperature of aerobic rice plants. Santo Antônio de Goiás, growing season 2015/16 and 2016/17.

Interaction between growing season and growth stage				
Growing season 2015/16				
Growth stage	A	E	Ci	Tleaf
Vegetative	22.79	5.64 b*	272 a	30.71 b
Reproductive	23.86	6.38 a	239 b	33.06 a
Growing season 2016/2017				
Growth stage	A	E	Ci	Tleaf
Vegetative	29.96 a	6.43 a	318	27.30 b
Reproductive	23.15 b	5.45 b	313	28.43 a
Interaction between growing season and rhizobacterium BRM32114				
Growing season 2015/16				
rhizobacterium	E	gs	Ci	Tleaf
With	6.21 a	0.52	250 b	32.26 a
Whitout	5.81 b	0.49	261 a	31.52 b
Growing season 2016/17				
rhizobacterium	E	gs	Ci	Tleaf
With	5.79	2.02 a	324 a	27.59 b
Whitout	6.08	1.48 b	307 b	28.14 a

* Means followed by the same letter do not differ by Tukey's test (P<0.05).

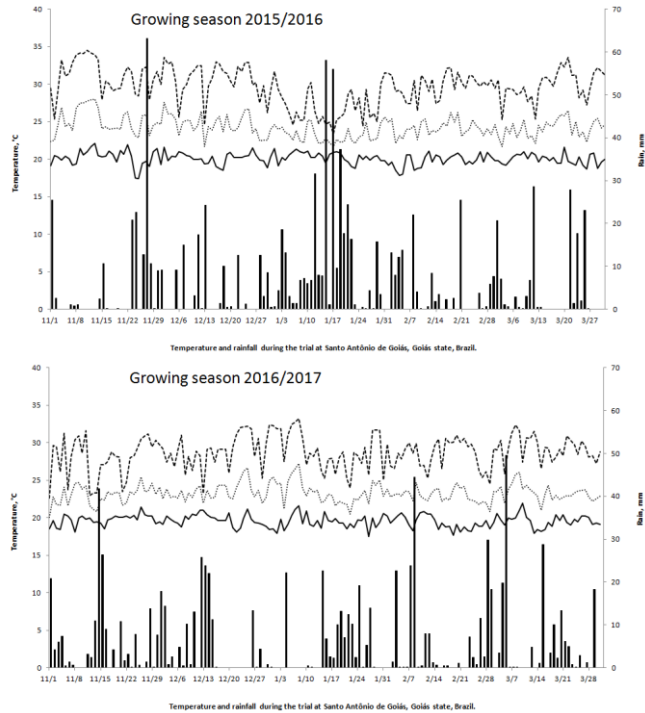


Fig 1. Maximum (T maximum), minimum (T minimum) and average (T average) temperatures and rainfall during the trial period of aerobic rice grown under a no-tillage system in the experimental fields of Santo Antônio de Goiás in the 2015/16 and 2016/17 growing seasons.

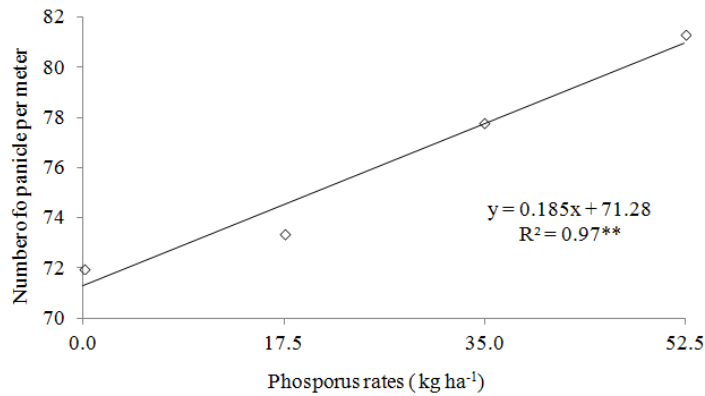


Fig 2. Number of panicles per meter of aerobic rice as a function of phosphorus rates. Average of growing seasons 2015/16 and 2016/17.

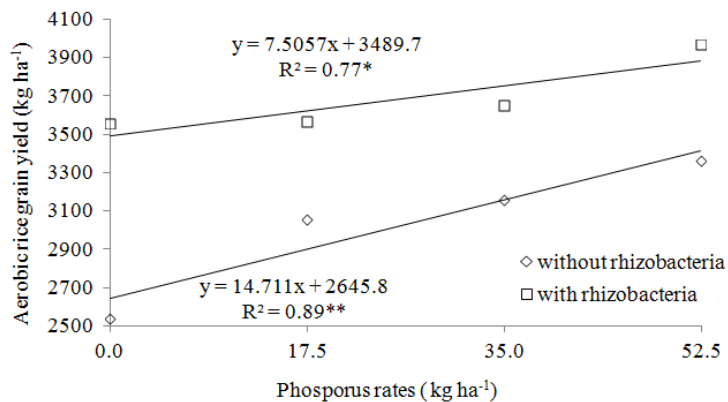


Fig 3. Aerobic rice grain yield as affected by use of Rhizobacteria BRM 32114 and phosphorus rates. Average of growing seasons 2015/16 and 2016/17.

Table 4. Phosphorus rates and *Serratia* spp. isolated BRM 32114 with or without inoculation in rice seeds as affecting nutrient content in rice leaves at full flowering stage. Santo Antônio de Goiás, growing season 2015/2016 and 2016/2017.

Factors	N	P	K	Ca	Mg	Cu	Fe	Mn	Zn
<i>Serratia</i> BRM32114	-----g kg ⁻¹ -----			-----mg kg ⁻¹ -----					
With	21.37	2.79	16.34 a*	3.26 a	2.21 a	6.60	112.18	69.85	35.55
Without	21.16	2.92	15.76 b	2.94 b	2.04 b	6.63	114.67	66.44	33.63
Growing season									
2015/2016	12.70 b	3.83 a	17.59 a	1.93 b	2.39 a	6.23 b	169 a	34.76 b	33.17 a
2016/2017	29.83 a	1.89 b	14.51 b	4.27 a	1.88 b	7.00 a	58 b	101.53 a	16.01 b
Factors	ANOVA (F probability)								
<i>Serratia</i> R-235 (S)	0.5849	0.3254	0.0437	0.0009	0.0003	0.8520	0.8098	0.4886	0.4106
Phosphorus rates (P)	0.9517	0.4819	0.5376	0.9587	0.6306	0.2404	0.9755	0.7641	0.2331
Growing season (GS)	<0.001	<0.001	<0.001	<0.001	<0.001	0.0002	<0.001	<0.001	<0.001
S x P	0.5304	0.8213	0.2014	0.2360	0.6979	0.3183	0.2685	0.6367	0.8397
S x GS	0.9760	0.2337	0.2724	0.0502	0.0512	0.4969	0.3562	0.2118	0.1930
P x GS	0.1187	0.6462	0.0602	0.4440	0.2132	0.1941	0.2043	0.6169	0.2899
S x P x GS	0.5591	0.8884	0.5305	0.7785	0.8065	0.5830	0.5731	0.5823	0.8851

* Means followed by the same letter do not differ by Tukey's test (P<0.05).

Table 5. Phosphorus rates and *Serratia* spp. isolated BRM 32114 with or without inoculation in rice seeds as affecting plant biomass (PB), number of panicles per plant (NPP), number of grain per panicle (NGP), mass of 1000 grains (1000M) and grain yield (GY) of aerobic rice. Santo Antônio de Goiás, growing season 2015/2016 and 2016/2017.

Factors	PB	NPP	NGP	1000M	GY
<i>Serratia</i> BRM32114	g m ⁻²	unit	unit	g	kg ha ⁻¹
With	197 a*	80 a	134	23.63	3691 a
Without	172 b	72 b	132	23.20	2990 b
Growing season					
2015/2016	143 b	69 b	122 b	23.22	2303 b
2016/2017	225 a	83 a	145 a	23.60	4379 a
Factors	ANOVA (F probability)				
<i>Serratia</i> R-235 (S)	0.0088	0.0071	0.6236	0.1648	<0.001
Phosphorus rates (P)	0.5371	0.0419	0.8639	0.3415	0.0428
Growing season (GS)	<0.001	<0.001	<0.001	0.2086	<0.001
S x P	0.6112	0.9475	0.4431	0.7248	0.0306
S x GS	0.2474	0.9656	0.9519	0.1760	0.0965
P x GS	0.7354	0.2527	0.1028	0.7550	0.3651
S x P x GS	0.9850	0.8306	0.611	0.8669	0.5976

* Means followed by the same letter do not differ by Tukey's test (P<0.05).

Table 6. Nutrient content in rice grains affected by phosphorus rates and *Serratia* spp. isolated BRM 32114 with or without inoculation in rice seeds during the growing season 2015/2016 and 2016/2017.

Factors	N	P	K	Ca	Mg	Cu	Fe	Mn	Zn	
<i>Serratia</i> BRM32114	-----g kg ⁻¹ -----			-----mg kg ⁻¹ -----						
With	13.15 b*	3.24	2.67	0.33	1.51	5.18 b	29	23	36 b	
Without	14.52 a	3.47	2.66	0.37	1.51	5.73 a	32	26	39 a	
Growing season										
2015/2016	13.80	3.81 a	3.03 a	0.36	1.64 a	5.61 a	32	20 b	38	
2016/2017	13.88	2.74 b	2.18 b	0.33	1.34 b	5.25 b	29	31 a	37	
Factors	ANOVA (F probability)									
<i>Serratia</i> R-235 (S)	0.0040	0.1486	0.9658	0.1450	0.9330	0.0014	0.1654	0.0269	0.0022	
Phosphorus rates (P)	0.1377	0.5171	0.2847	0.8177	0.2992	0.2084	0.7049	0.4691	0.3390	
Growing season (GS)	0.9893	<0.001	<0.001	0.2221	<0.001	0.2993	0.0506	<0.001	0.4927	
S x P	0.0887	0.6956	0.6549	0.5566	0.3335	0.3698	0.7795	0.5294	0.4522	
S x GS	0.0457	0.2830	0.2421	0.2401	0.1272	0.2460	0.3728	0.0034	0.1319	
P x GS	0.2820	0.8989	0.2539	0.7585	0.1617	0.7454	0.1951	0.8182	0.7016	
S x P x GS	0.3841	0.6807	0.8683	0.7438	0.6558	0.1819	0.4841	0.6390	0.2868	
Interaction between growing season and Rhizobacteria										
Growing season 2015/16										
rhizobacteria	N					Mn				
With	g kg ⁻¹					mg kg ⁻¹				
Without	13.55					20				
Without	14.04					19				
Growing season 2016/17										
rhizobacteria	N					Mn				
With	g kg ⁻¹					mg kg ⁻¹				
Without	12.62 b					27 b				
Without	15.15 a					35 a				

* Means followed by the same letter do not differ by Tukey's test (P<0.05).

the structure of rice root (Rêgo et al., 2014). According to the authors, rice seeds treated with *T. asperellum* pool, *Burkholderia* sp. and *Pseudomonas* sp. showed changes in root architecture by increasing the length and the volume of the root. Both greater activity in gas exchange and nutrient enrichment in aerobic rice leaves are factors that most contributed to the greater accumulation of biomass in aerobic rice plants and number of panicle per plant, which had a significant effect on crop grain yield (Fageria et al., 2011).

Phosphorus rates provided increases in number of panicles and grain yield. Increasing rates of P, provided increases in the number of panicles and in grain yield. Several studies around the world have shown the positive effect of P fertilization in growth, yield components, and grain yield of rice (Crusciol et al., 2005; Fageria et al., 2013a; Nascente et al., 2014). This could be due to the levels of P in the soil (Table 1). However, the low response of nutrients to increase P may be due to the adequate level in 2015/16 and high level in 2016/17 (Sousa and Lobato, 2003). The number of panicle per plant was higher in treated plants (Table 5, Figure 3). It seems that the number of formed and fertile tillers made the difference. The beginning of tiller phase, when rice plants emits the 3rd leaves does not depend on the environmental conditions. However, the emergence and development until 65 to 70 days, depend on nutritional and temperature conditions. To achieve vigorous tillering, N, P and S are necessary, together with carbohydrate (Fageria 1984). According to our data (Fig 2), there is a positive influence of different P rates on tillage numbers m⁻², until 60 days after sowing (Fageria et al., 1982). BRM 32114 is an Indoleacetic acid (IAA) and phosphatase producer (Nascente et al., 2017), which may enhance the tillering successful ensuring a higher number of fertile tillers and formed panicles (Fig 2 and Fig 3). Our results showed that using rhizobacterium *Serratia* spp. strain BRM 32114 with phosphorus allowed more increase in grain yield than using only P rates. In this sense, it is likely that the rhizobacterium BRM32114 positively interacted with P fertilization.

Our research seems promising and showed that using a beneficial microorganism could bring many benefits to the crops systems. In this sense, using aerobic rice treated with beneficial microorganism could contribute to improve the profitability of the crop for the farmers. The adequate to high level of P in the soil, along with rhizobacteria seem to improve the plant capacity to uptake nutrients to the plants and allowed higher grain yield than without rhizobacterium. The use of microbial for integrated nutrient management for rice production could contribute to a sustainable agriculture (Lavakush et al., 2014). Besides, our research confirmed the improvement of aerobic rice plants development allowed by the rhizobacterium under greenhouse condition (Nascente et al., 2017) in the field conditions.

Materials and methods

Site description

The experiments were conducted at Capivara Experimental Station of Embrapa Rice and Beans, located at Santo Antônio de Goiás, GO, Brazil, 16°28'00"S and 49°17'00"W coordinates, and at 823 m of elevation. The climate is Tropical Savanna and is considered Aw (tropical with wet summer and dry winter) according to Köppen classification (Köppen, 1918). Therefore, there are two well-defined

seasons: a usually dry season from May to September (autumn/winter) and a wet season from October to April (spring/summer). The average annual rainfall is between 1500 and 1700 mm, and the average annual temperature is 22.7°C, ranging annually from 14.2 °C to 34.8 °C. During the period of this study, the temperature and the amount of rainfall data were recorded (Fig 1).

The experimental soil is classified as a clay loam (kaolinitic, thermic Typic Haplorthox) acidic soil (Embrapa, 2006). Prior to the study, in November 2015 and 2016, chemical and physical analyses were performed according to the methodology by Donagema et al. (2011) for the initial characterization of the area (Table 1). Therefore, P and K were extracted by Mehlich 1 extracting solution (0.05M HCl in 0.0125M H₂SO₄). From the extracted, phosphorus content was determined colorimetrically and K by flame photometry. Ca, Mg, and Al were extracted with 1M KCl. Aluminum was determined by titration with NaOH and Ca and Mg by titration with EDTA from the extracted solution. Micronutrients were determined in a portion of the extract for P by atomic absorption spectrophotometry. Soil organic matter was determined by Walkley and Black's method.

Experimental design and treatments

Trials were conducted in rainfed conditions using the rice genotype Primavera CL. Trials were arranged in a complete randomized block in a 4x2x2 factorial scheme, with four replications in different areas very close to each other. The plots had the dimension of 3.5 m x 6 m. The usable area of the plot (place where we sampled the data) was composed of the four central rows of rice, disregarding one row of each side of the plot and 0.50 m to each side of the rows in each plot. The treatments consisted of four P₂O₅ doses (0, 40, 80 and 120 kg ha⁻¹ as triple superphosphate) with or without *Serratia* spp in two growing seasons (2015/16 and 2016/17). The strain BRM32114 was applied by seed microbiolization and sprayed at plant/soil in the field at the 7 and 15 DAS (days after sowing) in all plots with rhizobacteria treatment. In the plots with no microorganisms, we applied only water. *Serratia* spp. strain BRM32114 belongs to the Microorganism Culture Collection at the Embrapa Rice and Bean Research Center, with KX378746 GenBank code.

Seed microbiolization and plant/soil spray pulverization

Serratia spp. strain BRM32114 bacterial suspensions were prepared with water, from cultures that had been growing for a 24 hour period on solid medium 523 (Kado and Heskett, 1970), at 28 °C, and the concentration was set in a spectrophotometer to A540 = 0.5 (10⁸ CFU). Rice seeds were immersed in the bacterial suspension, during 24 hours under constant agitation at 25 °C, and control seeds (no bacteria) were immersed only in water, during 24 hours, in the same conditions. We performed plan (7 days after sowing) and soil (15 days after sowing) pulverization with bacterial suspension prepared as described above, in the same concentration (10⁸ CFU ml⁻¹), at 600 l ha⁻¹, with a costal atomizer.

Rice crop management

Cover crops were desiccated with a glyphosate application (1.8 kg ha⁻¹ acid equivalent) 30 days before sowing of the aerobic rice. The sowing was performed mechanically using

200 seeds per m^{-2} of rice. The seeds were sown on December 15th, 2015 and on November 23rd, 2016. Seven days before the sowing of the rice, the seed microbiolization was carried out. Rice plant emergence occurred five days after sowing for both growing seasons. The base fertilization, to be applied in the sowing furrows, was calculated according to the soil's chemical characteristics and followed the recommendations of Sousa and Lobato (2003). Therefore, sowing fertilization was P rates according each treatment and 50 kg ha^{-1} of N as urea. Topdressing fertilization was performed 18 days after sowing with 45 kg ha^{-1} of N as urea, in both years. Cultural practices were performed according to standard recommendations for a rice crop to keep the area free from weeds, diseases and insects.

Evaluations of traits

Leaf gas exchange and photosynthetic capacity

Aerobic rice gas exchange, at 37 and 68 days after emergence (DAE) for the first growing season (2015/16) and 49 and 79 DAE for the second growing season (2016/17), corresponding to the vegetative and reproductive stages of plants, were evaluated. The assessments were taken in the middle third of the upper leaves (completely expanded and with good exposure to the sun), as following: photosynthesis rate - A ($\mu\text{mol CO}_2 m^{-2} s^{-1}$), transpiration rate - E ($\text{mmol H}_2\text{O } m^{-2} s^{-1}$), stomatal conductance - gs ($\text{mol H}_2\text{O } m^{-2} s^{-1}$), internal CO_2 concentration - Ci (vpm) and leaf temperature - Tleaf ($^{\circ}\text{C}$). The gas exchange measures were determined by the portable infrared gas analyzer (LCpro+, ADC BioScientific) in the period from 8:00 to 10:00 in the morning. The equipment was set to use concentrations of 370-400 $\text{mol mol}^{-1} \text{CO}_2$ in the air, which is the reference condition used in the IRGA photosynthesis chamber. The photon flux density photosynthetic active (PPFD) used was 1200 $\mu\text{mol [quanta]} m^{-2} s^{-1}$. The minimum equilibration time set for performing the reading was 2 minutes.

Samples were taken in the middle third of the first fully expanded leaf (top to base) during the two evaluation periods in the two-year experiment. The equipment was set to use concentrations of 370-400 $\text{mol mol}^{-1} \text{CO}_2$ in the air, which is the reference condition used in the IRGA photosynthesis chamber. The photon flux density photosynthetic active (PPFD) used was 1200 $\mu\text{mol [quanta]} m^{-2} s^{-1}$. The minimum equilibration time set for performing the reading was 2 minutes.

Nutrients in leaves and grains of rice

Rice leaf samples were collected from the flag leaf at the full blooming stage. Leaves from 50 plants per plot were collected as proposed by Malavolta (2006), washed and then dried under forced-air circulation at 65 $^{\circ}\text{C}$ for 72 h before grinding and analyzing the samples for chemical composition. In each plot, one sample of 100 g of grains was taken for nutrient analyses. In leaves and grains, the concentrations of N, P, K, Ca, Mg, Cu, Zn, Fe and Mn were determined using methods described by Malavolta et al. (1997).

Shoot dry biomass

Rice shoots of 1.0 m from one of the rows, in each plot were sampled at the full blooming stage. The shoots were washed in water and then dried in an oven with forced air circulation at 65 $^{\circ}\text{C}$, then weighed to determine the shoot dry mass.

Yield components and grain yield

Rice harvest was carried out by hand after physiological maturity (March 30th, 2016 and March 21st, 2017) of the grains in the usable area of each plot. Plots were evaluated for the number of panicles m^{-1} , which was determined by counting the number of panicles within 1.0 m of one of the rows in the useful area of each plot; number of grains per panicle, which was determined by counting the number of grains in 10 panicles random sampled in the usable area and divided by 10; the mass of 1000 grains, which was randomly evaluated by collecting and weighing 1000 grains from each plot, corrected to 13% of water content; and the grain yield, which was determined by weighing the harvested grain of each plot, corrected to 13% of the water content and converted to $\text{kg } ha^{-1}$.

Statistical analysis

For the statistical analysis, the SAS Statistical Software, SAS Institute, Cary, NC, USA (SAS, 1999) was used. In the qualitative factor (with or without *Serratia* spp. strain BRM32114), data was subjected to an analysis of variance, and when the F test proved significant, the data was compared by a Tukey test at $p < 0.05$. In the quantitative factor (P rates), results were submitted to regression analysis when $p < 0.05$. The blocks and all block interactions were considered as random effects. The P rates, use of rhizobacteria (*Serratia* spp. strain BRM32114) and year were considered as fixed effects.

Conclusions

The trial was conducted to evaluate beneficial microorganism on the development of aerobic rice, under field conditions in tropical region. This is one of the first studies that connected microorganism and P rates in the crop aerobic rice. The results showed that phosphorus rates increased the number of panicles and grain yield of aerobic rice, and the grain yield was higher with *Serratia* inoculation than without. Under no-tillage, *Serratia* spp. strain BRM 32114 was inserted in the production system of aerobic rice and promoted higher gas exchange activity, enriched the K, Ca and Mg content in leaves, enhanced biomass accumulation, and resulted in a higher number of panicles per plant and grain yield than untreated plants. Therefore, this beneficial microorganism is a promising technology for potential incorporation into cropping systems in Brazilian Cerrado region and could be evaluated under different aerobic conditions worldwide.

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