

Compatibility and antimicrobial activity of *Trichoderma* spp. combined with diazotrophs and growth-promoting bacteria

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Abstract: The study evaluated the compatibility, *in vitro*, of diazotrophs *Bradyrhizobium* spp., and plant growth-promoting bacteria (PGPB), *Azospirillum brasilense* and *Bacillus* spp., widely used as biofertilizers and biostimulation soybean culture combined with as biofungicides/bionematicides *Trichoderma harzianum* and *Trichoderma asperellum*. Compatibility was evaluated through antagonistic activity and the minimum inhibitory concentration (MIC), using the techniques of dual cultures and disc diffusion, respectively. The experiment was conducted at the soil microbiology laboratory out at the Embrapa Temperate Climate, Pelotas, RS, Brazil. We adopted a completely randomized design (CRD) with three replications and the data were submitted to ANOVA followed by Tukey's test ($p \leq 0.05$) for compare of the mycelial growth variables, that showed significant difference. *B. elkanii* strain SEMIA.587 was compatible with all accessions of *Trichoderma* spp. *A. brasilense* strains Ab-V5 and Ab-V6, were compatible with the *T. asperellum* accession UFRA.T09. *B. elkanii* strain SEMIA.5019, *B. japonicum* strain SEMIA.5079 and *B. diazoefficiens* strain SEMIA.5080, *B. subtilis* strain CNPMS.B2084 and *B. megaterium* strain CNPMS.B119 inhibited from 47% to 83% the growth of *Trichoderma* spp. The MIC demonstrated that diazotrophs and PGPB are not sensitive to *Trichoderma* spp. *B. elkanii* strain SEMIA.587 can be used in combination with all *Trichoderma* spp. accessions, as well as the Ab-V5 and Ab-V6 strains of *A. brasilense* with the *T. asperellum* accession UFRA.T09 in the inoculation via a seeding furrow.

Keywords: Biofungicide; Dual Cultures; MIC; PGPB; Strains.

Introduction

The use of biological agents in soybean *Glycine max* (L.) has shown relevant results in the uptake of nitrogen and phosphorus, accumulation of plant biomass, synthesis of phytohormones, and increase in nodulation (Chagas et al., 2017; Moretti et al., 2020; Duré et al., 2022; Braccini et al., 2023), in addition to mitigating the environmental impacts caused by the use of chemical inputs (Hungria et al., 2015). The diazotrophs *Bradyrhizobium* spp. are widely used in biological nitrogen fixation (BNF) (Kaschuk et al., 2016; Mirriam et al., 2022) and its effectiveness supplies the totality of the nitrogen demand necessary for the crop (Hungria et al., 2020; Mahmud et al., 2020).

Brazilian legislation contains records of four strains of *Bradyrhizobium* spp. (*B. japonicum* SEMIA.5079, *B. diazoefficiens* SEMIA.5080, *B. elkanii* SEMIA.5019 and SEMIA.587) available for soybean crops (MAPA, 2011). However, some studies have shown that the association of these microorganisms with PGPB, through the so-called co-inoculation technique, has shown benefits for plant development (Ilićević et al., 2017; Moretti et al., 2019; Garcia et al., 2021).

The main combinations involve the co-inoculation of *Bradyrhizobium* spp. and *A. brasilense* strains Ab-V5 and Ab-V6. Currently, this technique is applied in 25% of the soybean cultivation area (Santos et al., 2021) and results in phytohormone production and increases in root growth, benefiting the performance of *Bradyrhizobium* spp. in nodulation and, consequently, in BNF (Hungria et al., 2015; Ferri et al., 2017; Barbosa et al., 2021).

The genus *Bacillus* has been successfully employed with *Bradyrhizobium* spp. for inoculation, showing improvement in nodulation, growth, and shoot and root length (Atieno et al., 2012; Sibponkrung et al., 2020; Miljaković et al., 2022). Simultaneously, there has been an increase in the use of disease biocontrol agents based on *Trichoderma* spp., which in addition to controlling root pathogens and being a more sustainable

alternative (Iturralde et al., 2020; Barbosa et al., 2022; Conte et al., 2022), show promising results in promoting growth (Conte et al., 2022), synthesizing indole acetic acid and auxins (Chagas et al., 2017), and increasing tolerance to abiotic stress (Tyśkiewicz et al., 2022).

Inoculation of soybean seeds with *Trichoderma* spp. via a seeding furrow is an increasingly used practice (Meyer et al., 2022) because, in addition to being efficient biocontrol agents (Conte et al., 2022), they have potential as plant growth promoters (Chagas et al., 2017; Junior et al., 2022). Using co-inoculation technology via seeding furrows, *Trichoderma* spp., when used in conjunction with PGPB and diazotrophs, enables increased efficiency in the inoculation process (Ferri et al., 2017; Brignoli et al., 2023). When using the co-inoculation technique, it is important that no organism is inhibitory to one another or interferes too much with the growth of the other (Whipps, 2001; Thomloui et al., 2019).

Information related to the compatibility of bacteria and fungi beneficial to the development of the soybean crop is still limited (Karuppiyah et al., 2019; Mattos et al., 2020) and, for the most part, is limited to greenhouse evaluations (Ayoubi et al., 2012; Cadore et al., 2020; El-Nahrawy et al., 2020). Thus, the aim of this study was to evaluate the *in vitro* compatibility of diazotrophs (*B. japonicum*, *B. diazoefficiens* and *B. elkanii*) and PGPB (*A. brasilense*, *B. subtilis*, *B. megaterium*) with biocontrol fungi, *T. asperellum* and *T. harzianum*, by evaluating their antagonistic effects.

Results

The results obtained in the antagonism test, through the dual culture technique, demonstrated that all evaluated bacteria, diazotrophs and PGPB, had an inhibitory effect against the isolates of *T. asperellum* and *T. harzianum* (Table 2). As an exception, *B. elkanii* strain SEMIA.587 did not form an inhibition halo with the evaluated *Trichoderma* spp. isolates (Fig.1A), nor

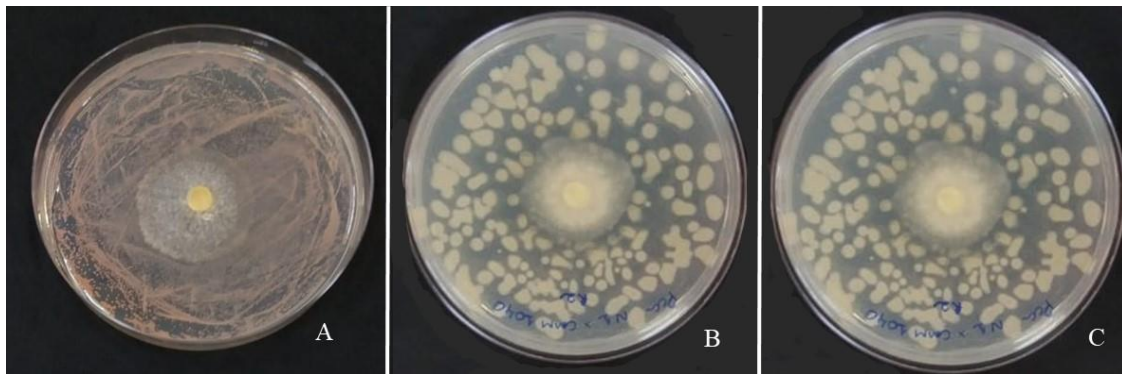


Fig 1. Antagonistic activity through the dual culture technique, without occurrence of formation of inhibition halo, after 120 hours of inoculation. *Bradyrhizobium elkanii* strain SEMIA.587 vs *Trichoderma harzianum* strain Esalq.1306 (A); *Azospirillum brasilense* strain Ab-V6 vs *Trichoderma asperellum* strain UFRA.T09 (B); and *Azospirillum brasilense* Ab-V5 vs *Trichoderma asperellum* strain UFRA.T09 (C). Photograph: Gobbi, PC.

Table 1. Selective medium and incubation conditions in a microbiological oven for the growth of different microorganisms used in the evaluation of antagonism and minimum inhibitory concentration. Embrapa Clima Temperado, Pelotas-RS, Brazil.

Microorganism	Culture medium	Incubation conditions
<i>Azospirillum</i> spp.	Red Congo	24 hours in a bacteriological oven at 28°C
<i>Bacillus</i> spp.	Casein-Peptone Dextrose Yeast Agar	24 hours in a bacteriological oven at 28°C
<i>Bradyrhizobium</i> spp.	Yeast Mannitol Agar	24 hours in a bacteriological oven at 28°C
<i>Trichoderma</i> spp.	Potato Dextrose Agar	7 days in BOD at 27°C

did *A. brasilense* strains Ab-V5 (Fig.1B) and Ab-V6 (Fig.1C) when faced with *T. asperellum* strain UFRA.T09.

When assessing PGPB compatibility of *A. brasilense* strains Ab-V5 and Ab-V6 with *Trichoderma* spp. isolates, the smallest halos of mycelial growth inhibitions occurred for *T. asperellum* strain UFRA.T09 (47.04% (F=6.976; df=4; p=0.006) and 62.85% (F=6.778; df=4, p=0.007), respectively; however, the latter did not differ significantly from *T. harzianum* strain Esalq.1306 (65.73%; Table 3).

In contrast, for PGPB, *B. megaterium* strain CNPMS.B119 and *B. subtilis* strain CNPMS.B2084, the highest percentage of inhibition occurred when confronted with *T. asperellum* strain UFRA.T09 (77.96% (F=71.28; df=4; p<0.05) and 68.44% (F=30.00; df=4; P<0.05), respectively) and did not differ for other evaluated isolates (Table 3).

Bradyrhizobium diazoefficiens strain SEMIA.5080 (F=4.486; df=4; p=0.025) and *B. japonicum* strain SEMIA.5079 (F=7.177, df=4; p=0.003) exhibited the lowest percentages of inhibition against the test agents *T. asperellum* strain UFRA.T12 (65.55% and 68.61%, respectively) and *T. harzianum* strain Esalq.1306 (74.63% and 66.67%, respectively). For *B. elkanii* strain SEMIA.5019, there were no significant differences between the evaluated *Trichoderma* spp. isolates (F=3.00; df=4; p=0.072; Table 3).

In the evaluation of the response of test agents to different antagonists, *B. elkanii* strain SEMIA.587 exhibited the lowest mean percentages of mycelial growth inhibition for *T. asperellum* strain UFRA.T06 (54.38%; F=20.426; df=7; p< 0.05), *T. asperellum* strain UFRA.T12 (47.02%; F=7.727; df=7; p=0.0003) and *T. asperellum* strain UFRA.T52 (49.12%; F=21.208; df=7; p<0.05) compared to other antagonists (Table 3). For *T. asperellum* strain UFRA.T09, the lowest suppression occurred in the interaction with *A. brasilense* strain Ab-V5 (F=34.05; df=7; p<0.05; Table 3).

The lowest percentage of mycelial inhibition for *T. harzianum* strain Esalq.1306 was observed with the antagonist *B. subtilis* strain CNPMS.B2084 (53.76%); however, it did not differ from *B. elkanii* strain SEMIA.587 (59.54%) or *B. megaterium* strain CNPMS.B119 (60.56%). The highest inhibition percentages for *T. harzianum* strain Esalq.1306 were verified in interactions with *B. diazoefficiens* strain SEMIA.5080 (74.63%), *B. elkanii* strain SEMIA.5019 (74.63%) and *B. japonicum* strain SEMIA.5079 (66.67%; F=16.432; df=7; p<0.05; Table 3).

Antagonists *B. diazoefficiens* strain SEMIA.5080, *B. japonicum* SEMIA.5079 and *B. elkanii* strain SEMIA.5019 also exhibited the highest inhibition values for *T. asperellum* strain UFRA.T09

(F=34.05; df=7; p<0.05) and *T. asperellum* strain UFRA.T12 (F=7.73; df=7; p=0.0004; Table 3). For *T. asperellum* strain UFRA.T06 and *T. asperellum* strain UFRA.T52, the highest percentage of inhibition occurred when confronted with *A. brasilense* strain Ab-V6 (80.28%; F=20.43; df=7; p<0.05 and 80.09%; F=21.21; df=7; p<0.05, respectively; Table 3).

The evaluation of microbial activity through the MIC demonstrated that diazotrophs and PGPB were not sensitive to the evaluated *Trichoderma* spp. isolates since there was no inhibition of bacterial growth at the different concentrations tested. Additionally, *B. subtilis* strain CNPMS.B2084 (Fig.2A), *B. megaterium* strain CNPMS.B119 (Fig.2B), *B. elkanii* strain SEMIA.5019 (Fig.2C), *B. japonicum* strain SEMIA.5079 (Fig.2D) and *B. diazoefficiens* strain SEMIA.5080 (Fig.2E) colonized the entire area of the plate, preventing the growth of *Trichoderma* spp. isolates at all evaluated concentrations.

The mycelial growth of *T. harzianum* strain Esalq.1306 and *T. asperellum* strain UFRA.T12 was observed up to a concentration of 1×10^5 conidia mL⁻¹, while that for *T. asperellum* strain UFRA.T06, *T. asperellum* strain UFRA.T09 and *T. asperellum* strain UFRA.T52 occurred up to a concentration of 1×10^6 conidia mL⁻¹ (Table 4). The evaluated *Trichoderma* spp. isolates showed reduced growth compared to the control, as shown in the antagonism test; however, for the most part, they did not differ from the control (Table 4).

Discussion

The formation of the halo of inhibition of diazotrophs and PGPB against the isolates of *T. asperellum* and *T. harzianum* did not nullify the development of the fungi after 120 hours of cultivation. However, all interactions, with and without halo formation, showed a reduction in mycelial growth in *Trichoderma* spp. isolates. Among the evaluated bacterial isolates, the diazotrophs *B. diazoefficiens* strain SEMIA.5080, *B. japonicum* strain SEMIA.5079 and *B. elkanii* strain SEMIA.5019 showed the greatest inhibitory effects against *Trichoderma* spp., and the interaction neutrality was verified for *B. elkanii* strain SEMIA.587 with *Trichoderma* spp. evaluation. Bécquer et al. (2013) found a neutral interaction between *Sinorhizobium* (Ensifer) *meliloti* and *T. harzianum* in an *in vitro* antagonism evaluation after 96 hours of cultivation, and although they did not show formation of an inhibitory halo, there was a reduction in the diameter of the *T. harzianum* colony.

Table 2. Values of inhibition halos (mm) of the antagonists, diazotrophs and plant growth-promoting bacteria, against the test agents, *Trichoderma* spp. Embrapa Clima Temperado, Pelotas-RS, Brazil.

Antagonists	<i>Trichoderma harzianum</i> Esalq.1306	<i>Trichoderma asperellum</i> UFRA.T06	<i>Trichoderma asperellum</i> UFRA.T09	<i>Trichoderma asperellum</i> UFRA.T12	<i>Trichoderma asperellum</i> UFRA.T52
<i>Azospirillum brasilense</i> Ab-V6	5.00	1.67	0.00	0.87	1.33
<i>Azospirillum brasilense</i> Ab-V5	5.33	4.00	0.00	2.17	3.67
<i>Bacillus subtilis</i> CNPMS.B2084	5.33	2.66	1.67	2.00	2.33
<i>Bacillus megaterium</i> CNPMS.B119	3.00	5.67	3.00	1.67	1.33
<i>Bradyrhizobium elkanii</i> SEMIA.587	0.00	0.00	0.00	0.00	0.00
<i>Bradyrhizobium elkanii</i> SEMIA.5019	3.00	3.33	4.00	4.00	3.67
<i>Bradyrhizobium japonicum</i> SEMIA.5079	2.67	1.67	1.33	2.67	1.17
<i>Bradyrhizobium diazoefficiens</i> SEMIA.5080	3.00	1.00	2.00	1.67	1.33
Coefficient of variation (CV%)	52.57	72.33	99.39	63.08	69.30

The *in vitro* compatibility of *B. elkanii* strains SEMIA.587 and SEMIA.5019 evaluated by Mattos et al. (2020) was also shown to be the most compatible with the different isolates of *T. asperellum*. The strains belonging to *B. elkanii* produce rhizobiotoxins and present intrinsic resistance to antibiotics, unlike strains of *B. japonicum* (Minamisawa et al., 1996). The production of rhizobiotoxin limits endogenous ethylene production in soybean plants in low land soils, with poor natural drainage, favoring plant nodulation (Mattos et al., 2019). Thus, the synthesis of rhizobiotoxins can be a factor in the neutral interaction of *B. elkanii* SEMIA.587 and *Trichoderma* spp. verified in the present study.

The use of *B. elkanii* together with *T. harzianum* favored soybean production (Silva et al., 2018), and co-inoculation of *B. elkanii* and *T. asperellum* resulted in increased plant height and root dry mass production, indicating its application in cowpea, *Vigna unguiculata* (L.) Walp. (Fabaceae) (Costa et al., 2020).

Biological agents are considered compatible when they do not have a suppressive effect on each other *in vitro* co-cultivation or in rhizosphere colonization assays (Thomloui et al., 2019). Antagonistic effects *in vitro* partially reflect the response of an interaction in culture systems (Prasadd and Babu, 2017) since microorganisms can colonize different ecological niches in the rhizosphere without interfering with each other's growth (Niu et al., 2020), making the antagonistic activity notorious when space and nutrients are limited, and perhaps this is the main effect of the antagonistic action of fungi (Carvalho et al., 2014).

The use of the consortium *B. japonicum* and *T. harzianum* in the soybean crop resulted in an increase in the shoot and root length and in the N content compared to the use of *B. japonicum* alone (El-Nahrawy et al., 2020). The co-inoculation of *B. japonicum* and *T. harzianum* increased the fresh root mass without interfering with nodulation by *B. japonicum*, with soybean plants presenting nodules with the same ultrastructure as those inoculated only with *B. japonicum*. In addition, no fungal structures were observed below the root epidermis, indicating that *T. harzianum* colonized the soybean rhizosphere and remained on the surface of the roots (Iturralde et al., 2020).

The *in vitro* antagonism of *Bradyrhizobium* spp. against *T. asperellum* was evaluated by Mattos et al. (2020), and unlike the results obtained in this study, the authors observed the antagonistic action of *B. elkanii* strain SEMIA.587, *B. elkanii* strain SEMIA.5019, *B. japonicum* strain SEMIA.5079 and *B. diazoefficiens* strain SEMIA.5080 against *T. asperellum* strain UFRA.T09 and the formation of halos ranging from 1 to 5 mm diameter, demonstrating low potential for suppression of *Bradyrhizobium* spp. This occurred similarly in the present study, with halos ranging from 1.00 to 4.00 mm (Table 2) in the

evaluations of *Bradyrhizobium* spp. against *T. asperellum* and *T. harzianum* isolates.

Bacillus subtilis strain CNPMS.B2084 and *B. megaterium* strain CNPMS.B119 showed the lowest results of mycelial growth inhibition in *T. harzianum* strain Esalq.1306, when compared to other bacterial isolates, although there was formation of an inhibition halo. The antagonism of *Bacillus* spp. against phytopathogenic fungi is often related to the production of secondary metabolites with antibiotic properties and competition for nutrients and space (Harba et al., 2020; Miljaković et al., 2022); however, evaluations of the effect of these microorganisms against beneficial fungi *in vitro* are still limited.

The investigation of the *in vitro* compatibility of several isolates of *Bacillus* spp. and *Trichoderma* spp. showed strong antagonistic activity, making prior analysis a necessary step for application in mixtures (Fuga et al., 2016). The combined application of *B. subtilis* and *T. harzianum* proves to be compatible, providing protection against *Rhizoctonia solani* Kühn; however, the best results were obtained when applied separately (Abeyasinghe, 2009). According to Li et al. (2005), *B. subtilis* exerts antifungal activity against the growth of *T. harzianum*, inducing the formation of chlamydospores.

The compatibility of *A. brasilense* strains Ab-V5 and Ab-V6 varied among the evaluated *Trichoderma* spp. isolates, being more inhibitory against *T. asperillum* strains UFRA.T6 and UFRA.T12 and more compatible with *T. asperillum* UFRA.T09 without the occurrence of the formation of an inhibition halo. Co-inoculation with *T. harzianum* and *A. brasilense* in beans, *Phaseolus vulgaris* L. (Fabaceae), decreased the number of nodules compared to inoculation with *T. harzianum*; however, there was an increase in the weight of nodules (Öğüt et al., 2005) and increased concentrations of micronutrients in grains (Öğüt et al., 2006).

The assessment of MIC demonstrated that diazotrophs and PGPB are not sensitive to the tested *Trichoderma* spp. isolates since they did not show reduced growth compromised in the presence of the fungus. The inhibition potential of a microorganism may be related to competition for resources available in the substrate, speed of multiplication and colonization capacity, which defines efficiency as an antagonist of *Trichoderma* spp. isolates (Benítez et al., 2004) or even to an evaluation method.

Scorzoni et al. (2007) when comparing assessment methods of MIC of an antimicrobial, observed that the agar diffusion technique was less sensitive than the microdilution technique, concluding that the effectiveness of a method may not always be adequate for an antimicrobial due to differences in physical, volatile and diffusion properties. According to Muniz et al. (2018),

Table 3. Mean percentage of inhibition of mycelial growth (\pm SE) of the antagonists, diazotrophs and plant growth-promoting bacteria, against the test agents, *Trichoderma* spp. Embrapa Clima Temperado, Pelotas-RS, Brazil.

1	<i>Trichoderma harzianum</i> Esalq.1306	<i>Trichoderma asperellum</i> UFRA.T06	<i>Trichoderma asperellum</i> UFRA.T09	<i>Trichoderma asperellum</i> UFRA.T12	<i>Trichoderma asperellum</i> UFRA.T52
<i>Azospirillum brasilense</i> Ab-V6	65.73 \pm 5.12 bB	80.28 \pm 2.05 aA	62.85 \pm 3.41 bC	80.09 \pm 5.12 aA	72.15 \pm 1.65 abABC
<i>Azospirillum brasilense</i> Ab-V5	63.47 \pm 1.63 aB	67.61 \pm 0.81 aBC	47.04 \pm 2.30 bD	64.61 \pm 3.07 aAB	64.38 \pm 5.48 aBCD
<i>Bacillus subtilis</i> CNPMS.B2084	53.76 \pm 1.54 cC	56.81 \pm 1.24 bcCD	68.44 \pm 0.48 aBC	61.87 \pm 1.39 bBC	53.42 \pm 0.79 cDE
<i>Bacillus megaterium</i> CNPMS.B119	60.56 \pm 0.81 bBC	60.09 \pm 0.81 bCD	77.96 \pm 1.07 aA	59.36 \pm 1.21 bBC	62.10 \pm 0.91 bCD
<i>Bradyrhizobium elkanii</i> SEMIA.587	59.54 \pm 0.74 aBC	54.38 \pm 0.23 bD	61.67 \pm 1.50 aC	47.02 \pm 0.54 cC	49.12 \pm 1.09 cE
<i>Bradyrhizobium elkanii</i> SEMIA.5019ns	74.63 \pm 1.49 A	62.74 \pm 4.97 CD	73.66 \pm 2.34 AB	68.46 \pm 4.63 AB	78.79 \pm 3.15 A
<i>Bradyrhizobium japonicum</i> SEMIA.5079	66.67 \pm 2.77 cAB	78.92 \pm 2.73 abAB	82.26 \pm 0.93 aA	68.61 \pm 3.42 bcAB	76.51 \pm 1.91 abcAB
<i>Bradyrhizobium diazoefficiens</i> SEMIA.5080	74.63 \pm 2.28 abA	79.71 \pm 2.79 aA	76.88 \pm 1.94 abAB	65.55 \pm 4.31 bAB	79.29 \pm 1.34 aA

Significant effect by ANOVA ($p < 0.05$; ns, $p > 0.05$). Means within a row followed by the same lower-case letter, or within a column followed by the same capital letter, do not differ significantly (Tukey test: $p \leq 0.05$).

Table 4. Mean mycelial growth (\pm SE) and minimum inhibitory concentration (MIC) of five accessions of *Trichoderma* spp. against eight bacterial accessions using the disk diffusion technique. Embrapa Clima Temperado, Pelotas-RS, Brazil.

Fungi	Concentration	Control	<i>Azospirillum brasilense</i> Ab-V6		MIC	<i>Azospirillum brasilense</i> Ab-V5		MIC	<i>Bradyrhizobium elkanii</i> SEMIA.587		MIC	p-value	
<i>Trichoderma harzianum</i> Esalq.1306	1x10 ⁸	13.72 \pm 0.61	a	9.59 \pm 1.57	b	NA ⁽¹⁾	14.89 \pm 0.87	a	NA	12.97 \pm 0.50	a	NA	0.000
	1x10 ⁷	13.49 \pm 0.50	a	9.26 \pm 1.10	b	NA	11.05 \pm 1.04	b	NA	10.25 \pm 0.38	b	NA	0.000
	1x10 ⁶	11.59 \pm 0.51	a	9.10 \pm 0.33	b	NA	7.65 \pm 0.33	b	NA	8.93 \pm 0.50	b	NA	0.000
	1x10 ⁵ ns	8.93 \pm 0.27		8.44 \pm 0.83		NA	8.85 \pm 0.96		NA	7.38 \pm 0.65		NA	0.368
<i>Trichoderma asperellum</i> UFRA.T06	1x10 ⁸	14.94 \pm 0.59	a	11.08 \pm 0.87	b	NA	15.80 \pm 0.69	a	NA	13.52 \pm 1.05	ab	NA	0.008
	1x10 ⁷	13.07 \pm 0.40	a	9.70 \pm 1.24	b	NA	14.99 \pm 0.76	a	NA	13.68 \pm 0.50	a	NA	0.004
	1x10 ⁶	11.51 \pm 0.74	ab	12.89 \pm 0.68	a	NA	9.67 \pm 0.97	b	NA	12.50 \pm 0.30	ab	NA	0.033
<i>Trichoderma asperellum</i> UFRA.T09	1x10 ⁸ ns	8.49 \pm 0.44		8.92 \pm 0.40		NA	10.49 \pm 1.38		NA	9.54 \pm 0.36		NA	0.333
	1x10 ⁷	8.80 \pm 0.47	a	8.89 \pm 0.39	a	NA	9.81 \pm 0.39	a	NA	10.29 \pm 0.19	a	NA	0.043
	1x10 ⁶	9.87 \pm 0.57	a	6.53 \pm 0.22	a	NA	8.84 \pm 0.55	a	NA	6.53 \pm 0.22	b	NA	0.001
<i>Trichoderma asperellum</i> UFRA.T12	1x10 ⁸	11.47 \pm 0.10	a	8.34 \pm 0.44	b	NA	8.26 \pm 0.69	b	NA	9.11 \pm 0.28	b	NA	0.000
	1x10 ⁷ ns	9.67 \pm 0.23		8.75 \pm 0.26		NA	9.77 \pm 0.99		NA	8.94 \pm 0.80		NA	0.626
	1x10 ⁶ ns	6.87 \pm 2.31		1.83 \pm 1.83		NA	8.46 \pm 0.79		NA	4.42 \pm 2.73		NA	0.168
	1x10 ⁵ ns	7.40 \pm 0.18		6.93 \pm 0.14		NA	6.62 \pm 0.45		NA	7.06 \pm 0.21		NA	0.298
<i>Trichoderma asperellum</i> UFRA.T52	1x10 ⁸	16.50 \pm 0.29	a	15.25 \pm 0.25	ab	NA	13.19 \pm 0.94	b	NA	13.75 \pm 0.63	b	NA	0.008
	1x10 ⁷ ns	12.85 \pm 0.89		13.25 \pm 0.63		NA	13.00 \pm 0.41		NA	10.25 \pm 1.03		NA	0.060
	1x10 ⁶ ns	10.61 \pm 1.06		6.89 \pm 2.35		NA	6.22 \pm 2.10		NA	4.57 \pm 2.75		NA	0.290

⁽¹⁾No antifungal activity; Significant effect by ANOVA ($p < 0.05$; ns, $p > 0.05$); Means within a row followed by the same lower-case letter, do not differ significantly (Tukey test: $p < 0.05$).

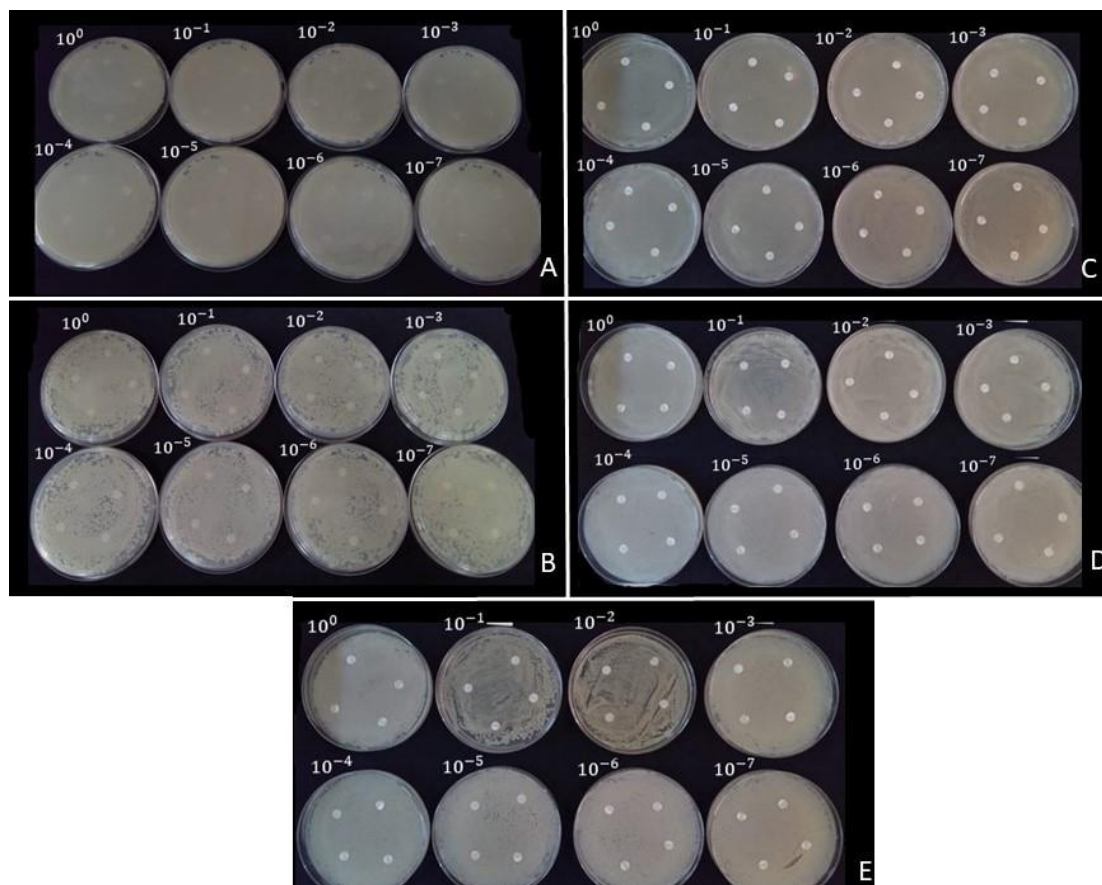


Fig 2. Antimicrobial activity of *Trichoderma* spp. assessed by determining the minimum inhibitory concentration using the disk diffusion sensitivity test, without occurrence of inhibition of bacterial growth at the different concentrations tested. Exemplars of *Trichoderma asperellum* strain UFRA.T06 evaluated with, *Bacillus subtilis* strain CNPMS.B2084 (A); *Bacillus megaterium* strain CNPMS.B119 (B); *Bradyrhizobium elkanii* strain SEMIA.5019 (C); *Bradyrhizobium japonicum* strain SEMIA.5079 (D); *Bradyrhizobium diazoefficiens* strain SEMIA.5080 (E). Photograph: Gobbi PC.

T. harzianum has the capacity for superficial colonization in a culture medium rich in sucrose, reaching 50% of growth after 48 hours. However, the standard means of *Clinical and Laboratory Standards Institute* (NCCLS, 2003), for disc diffusion sensitivity analyses, Müller–Hinton medium, does not contain a source of sugar, which could have limited the multiplication and colonization of fungi in this study.

Materials and methods

The experiment was conducted at the soil microbiology laboratory at Embrapa Clima Temperado, Pelotas, RS, Brazil (31°42' South and 52°24' West). Eight bacterial isolates were used: *A. brasilense* strains Ab-V5 and Ab-V6; *B. subtilis* strain CNPMS.B2084; *B. megaterium* strain CNPMS.B119; *B. elkanii* strains SEMIA.587 and SEMIA.5019; *B. japonicum* strain SEMIA.5079; and *B. diazoefficiens* strain SEMIA.5080. Five isolates of *Trichoderma* spp. were evaluated, with four isolates of *T. asperellum* (strains UFRA.T06, UFRA.T09, UFRA.T12 and UFRA.T52) and an isolate of *T. harzianum*, strain Esalq.1306, preserved in Coleção de Microrganismos Multifuncionais de Clima Temperado (CMMCT).

The compatibility of the microorganisms was evaluated through the antagonistic activity and the antimicrobial activity in the MIC using dual culture (Brito et al., 2018) and disc diffusion techniques (NCCLS, 2003), respectively.

Recovery and cultivation of biological isolates

Bacterial and fungal isolates were recovered and cultivated in Petri dishes (90 mm diameter) containing 25 mL of selective medium according to the characteristics of each species (Table 1). In a laminar flow chamber, smear loops of bacterial colonies, previously incubated in a microbiological oven, were removed

(Table 1), obtaining a concentration of $1-2 \times 10^8$ CFU mL⁻¹ according to tube n°0.5 of the MacFarland scale, through spectrophotometer reading (Logen Scientific SF325NM) at a wavelength of 625 nm (absorbance: 0.8-0.13).

Antagonism of diazotrophs and PGPB against *Trichoderma* spp.

For the *in vitro* evaluation of antagonism, 100 µL aliquots of the bacterial suspension were seeded using the *Drigalski* loop spreading method in Petri dishes (90 mm diameter) containing 25 mL of medium Casein-Peptide Dextrose Yeast Agar (PCA). In this evaluation, PCA was used for all bacterial isolates due to better visualization of mycelial growth and the halo of inhibition. A 6 mm diameter mycelium disc of an isolate previously grown in a Petri dish containing medium Potato-Dextrose Agar (PDA) (Table 1) was removed and deposited in the center of a Petri dish seeded with the antagonist (bacterial suspension), incubated inverted in a microbiological oven at 28°C, and evaluated after 120 hours of incubation.

The antagonistic activity was evaluated by measuring the mycelium growth diameter, including the inhibition halo, in the presence and absence of the antagonist, with the aid of a digital pachymeter. From these data, the mean percentage values of inhibition of the mycelial growth (ZI) of the test agent (Brito et al., 2018) were calculated using the following equation:

$$ZI = \frac{R1 - R2}{R1} \times 100$$

where:

R1- Diameter of the mycelium in the absence of the antagonist

R2- Diameter of the mycelium in the presence of the antagonist

The experiment was conducted in a CRD with three replications and a total of 40 interactions between fungi and bacteria.

Antimicrobial activity of *Trichoderma* spp. on bacterial isolates

The antimicrobial activity of *Trichoderma* spp. was evaluated by determining the MIC using a disc diffusion sensitivity test (NCCLS, 2003). The inoculum suspension of each accession was prepared from colonies incubated in selective medium at 28°C, 24 hours (Table 1). Each bacterium was suspended in 0.85% saline solution (autoclaved at 120°C, 20 min), and the cell concentration was adjusted to $1-2 \times 10^8$ CFU mL⁻¹, according to tube n°0.5 of the MacFarland scale. Subsequently, 100 µL of each bacterial suspension was seeded in a Petri dish (90 mm diameter) containing 25 mL of Mueller-Hinton medium using the Drigalski loop spreading method. Then, glass microfiber filter paper discs 934-AHTM (Whatman®) of 6 mm diameter that were previously autoclaved (120°C, 20 min) were placed on the surface of the agar using sterile tweezers and then inoculated with 10 µL (Rabanal et al., 2002) of the test agent (*Trichoderma* spp.) at different concentrations. For this, the fungi were cultivated in Petri dishes, as shown in Table 1, and the initial concentration of 1×10^8 conidia mL⁻¹ was determined by counting conidia in a Neubauer chamber. Serial dilutions were then performed (10^0 to 10^{-7}) in a 0.85% saline solution. The plates were incubated inverted at 35°C, 48 hours. A positive control of the test agent (*Trichoderma* spp.) was performed at the different tested concentrations. The data were evaluated after the incubation period and the MIC was determined as the lowest concentration of the test agent responsible for inhibiting the total and partial growth of bacterial isolates. The experiment was conducted in a CRD with four repetitions.

Data analysis

The normality and homoscedasticity of the ZI and mycelial growth data in the different MICs were analyzed using Shapiro-Wilk and Bartlett tests, respectively. As the statistical assumptions were met, an analysis of variance (ANOVA) was performed, followed by the Tukey test ($p < 0.05$), to compare the variables that showed significance using R v.3.4.1 software (R Development Core Team, 2022).

Conclusions

Compatibility and antimicrobial activity responses obtained in the present study confirm the importance of diazotrophs and PGPB that compose inoculants in the action of biocontrol of fungal agents. There was a neutral interaction of *B. elkanii* strain SEMIA.587 with all *Trichoderma* spp. *Bacillus subtilis* strain CNPMS.2084 and *Bacillus megaterium* strain CNPMS.B119 have the potential to inhibit the growth of *Trichoderma* spp. In contrast, *A. brasilense* strains Ab-V5 and Ab-V6 did not have suppressive mechanisms affecting *T. asperellum* UFRA.T09. Future research is needed to investigate the mechanisms of *B. elkanii* strain SEMIA.587, a producer of rhizobiotoxin, which limits endogenous ethylene production in low land soils with poor natural drainage, for greater compatibility with *Trichoderma* spp.

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