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Published in final edited form as:

Title: Interspecific cooperation: enhanced growth, attachment and strain-specific distribution in biofilms through *Azospirillum brasilense*-*Pseudomonas protegens* co-cultivation.

Authors: Pagnussat LA, Salcedo F, Maroniche G, Keel C, Valverde C, Creus CM

Journal: FEMS microbiology letters

Year: 2016 Oct

Issue: 363

Volume: 20

DOI: [10.1093/femsle/fnw238](https://doi.org/10.1093/femsle/fnw238)

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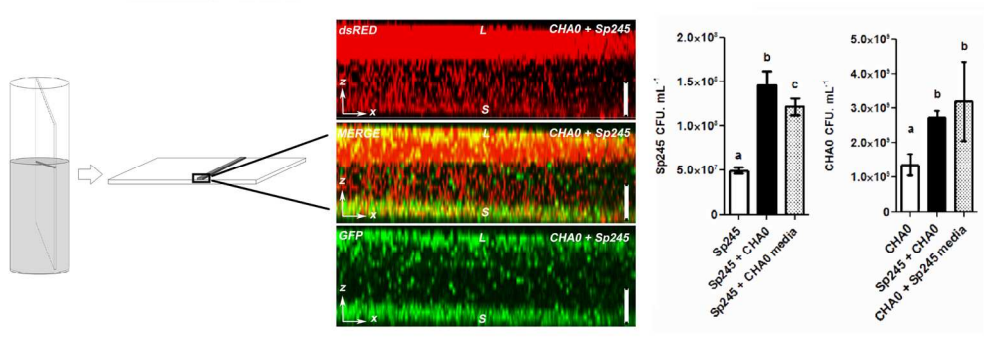
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Interspecific cooperation: enhanced growth, attachment and strain-specific distribution in biofilms through *Azospirillum brasilense*-*Pseudomonas protegens* co-cultivation

Journal:	<i>FEMS Microbiology Letters</i>
Manuscript ID	FEMSLE-16-04-0331.R3
Manuscript Type:	Research Letter
Date Submitted by the Author:	n/a
Complete List of Authors:	Pagnussat, Luciana; Universidad Nacional de Mar del Plata, Fac. Cs. Agrarias Salcedo, Florencia; Universidad Nacional de Mar del Plata, Facultad de Ciencias Agrarias; CONICET, Lab. Bioquímica Vegetal y Microbiana Maroniche, Guillermo; Universidad Nacional de Mar del Plata, Facultad de Ciencias Agrarias; Consejo Nacional de Investigaciones Científicas y Técnicas, Lab. Bioquímica Vegetal y Microbiana, FCA Keel, Christoph; Université de Lausanne, Département de microbiologie fondamentale Valverde, Claudio; Universidad Nacional de Quilmes, Departamento de Ciencia y Tecnología Creus, Cecilia; Universidad Nacional de Mar del Plata, Fac. Cs. Agrarias
Keywords:	plant-growth-promoting-bacteria, biofilm structure, colony phenotype
All articles in FEMS Microbiology Letters are published under one of eight subject sections. Please select the most appropriate subject category for your submission from the drop down list:	Environmental Microbiology (Editor: Tim Daniell)

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167x57mm (300 x 300 DPI)

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3 1 **Interspecific cooperation: enhanced growth, attachment and strain-specific distribution in**
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5 2 **biofilms through *Azospirillum brasilense*-*Pseudomonas protegens* co-cultivation**
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49 22 **One sentence summary:** Cooperative interaction between *Azospirillum brasilense* and
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51 23 *Pseudomonas protegens* enhances growth and produces structured mixed biofilms.
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56 25 **Keywords:** plant-growth-promoting-bacteria; biofilm structure; colony phenotype
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26 ABSTRACT

27 Plant growth promoting bacteria belonging to *Azospirillum* and *Pseudomonas* genera are major
28 inhabitants of the rhizosphere. Both are increasingly commercialized as crops inoculants. Inter-
29 specific interaction in the rhizosphere is critical for inoculants aptness. The objective of this
30 work was to evaluate *Azospirillum* and *Pseudomonas* interaction in mixed biofilms by co-
31 cultivation of the model strains *A. brasilense* Sp245 and *P. protegens* CHA0. The results
32 revealed enhanced growth of both strains when co-cultured in static conditions. Moreover,
33 Sp245 biofilm formed in plastic surfaces was increased 2-fold in the presence of CHA0.
34 Confocal microscopy revealed highly structured mixed biofilms showing Sp245 mainly on the
35 bottom and CHA0 towards the biofilm surface. In addition, *A. brasilense* biofilm was thicker
36 and denser when co-cultured with *P. protegens*. In a colony-colony interaction assay, Sp245
37 changed nearby CHA0 producing small colony phenotype, which accounts for a diffusible
38 metabolite mediator; though CHA0 spent medium did not affect Sp245 colony phenotype.
39 Altogether, these results point to a cooperative interaction between *A. brasilense* Sp245 and *P.*
40 *protegens* CHA0 in which both strains increase their static growth and produce structured mixed
41 biofilms with a strain-specific distribution.

43 INTRODUCTION

44 The rhizosphere constitutes a complex and dynamic environment. As an outcome of co-
45 evolution, interaction between rhizospheric microorganisms involve morphological changes and
46 adjustments in secondary metabolism (Philippot, *et al.*, 2013). In consequence, the fitness of
47 bacteria is largely determined by their capabilities to bear up secondary metabolites released by
48 other rhizospheric organisms.

49 Biofilms, defined as matrix-enclosed bacterial population adhered to each other and/or to
50 surfaces, constitute a micro-niche where the exchange of metabolites between bacterial

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3 51 populations is favored. The biofilm mode of growth has several advantages for soil bacteria
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5 52 compared to the planktonic lifestyle. Biofilm provides protection against desiccation and toxic
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7 53 compounds. Furthermore, an organization of multispecies bacterial consortia into biofilms may
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9 54 enable the coexistence of species that would otherwise outcompete each other and facilitate
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11 55 synergistic interactions and gene transfer (for reviews see: (Burmølle, *et al.*, 2007, Karatan &
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13 56 Watnick, 2009, López, *et al.*, 2010, Karunakaran, *et al.*, 2011)). Recent findings suggest that
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15 57 when both competitive and cooperative genes are transferred, cooperators are favored because
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17 58 their transfer increases allelic assortment (Dimitriu, *et al.*, 2014).
18
19 59 Plant growth-promoting rhizobacteria (PGPR) enhance growth of many terrestrial crop plants
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21 60 by a wide variety of mechanisms (Lugtenberg & Kamilova, 2009, Hayat, *et al.*, 2010,
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23 61 Bhattacharyya & Jha, 2012). *Azospirillum brasilense* improves crop growth mainly by a direct
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25 62 effect on the plant, through the production of several phytohormones such as indole acetic acid
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27 63 (Tien, *et al.*, 1979, Ona, *et al.*, 2005), and nitric oxide (Creus, *et al.*, 2005, Molina-Favero, *et al.*,
28
29 64 2008). Conversely, fluorescent pseudomonads, other major inhabitants of the rhizosphere,
30
31 65 display mostly indirect beneficial effects on plant growth (Kloepper, *et al.*, 1980). *Pseudomonas*
32
33 66 strains produce a wide range of compounds with antimicrobial activity, being 2,4-
34
35 67 diacetylphloroglucinol (DAPG) one of the most studied (Keel, *et al.*, 1990). These PGPR are
36
37 68 currently commercialized as inoculants containing single or combined strains (i.e. *Azospirillum*
38
39 69 *brasilense* and *Pseudomonas fluorescens*) (Bhattacharyya & Jha, 2012). PGPR efficiency
40
41 70 depends considerably on their capabilities to survive and establish effective root colonization.
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43 71 Therefore, interaction with other bacteria in the rhizosphere constitutes a critical bottleneck in
44
45 72 PGPR aptness. An important concern is that pseudomonads may secrete antibiotics that can
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47 73 negatively affect *A. brasilense* proliferation in mixed inoculants (Combes-Meynet, *et al.*, 2010,
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49 74 Couillerot, *et al.*, 2011). It has been reported that, *in vitro*, *A. brasilense* Cd strain was sensitive
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51 75 to the addition of the synthetic antimicrobial metabolite DAPG, which induced carotenoids
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3 76 accumulation, formation of poly- β -hydroxybutyrate-like granules, cytoplasmic membrane
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5 77 damage and growth inhibition (Couillerot, *et al.*, 2011). The major rhizosphere inhabitants, *A.*
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7 78 *brasilense* and fluorescent pseudomonads, are expected to be found in mixed consortia in
8
9 79 nature. Therefore, studying their interaction is of particular interest for agronomical applications
10
11 80 in inoculants technology. The aim of this work was to assess whether interspecific interaction
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13 81 between the model strains *A. brasilense* Sp245 and *Pseudomonas protegens* CHA0 results in
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15 82 altered static growth, attachment to polystyrene surfaces and strain distribution within mixed
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17 83 biofilms.
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85 MATERIALS AND METHODS

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87 Bacterial strains and culture conditions

88 Fluorescent derivatives of strains *A. brasilense* Sp245 and *P. protegens* CHA0 were obtained by
89 introducing plasmid pMP2444 (Gm^R , $P_{lac-egfp}$) into Sp245 by biparental mating (Arruebarrena
90 Di Palma *et al.* 2013) or plasmid pME7134 (Tc^R , $P_{lac-dsRed}$) by electroporation. Plasmid
91 pME7134 was constructed as follows. A 0.7-kb *SmaI-HindIII* fragment from pDsRed.T3_S4T
92 containing the *dsred.T4_S4T* gene (Sørensen *et al.*, 2003) was cloned under the control of the
93 P_{lac} promoter into pME6552 (Wenner *et al.*, 2014). Following this, a 1.5-kb *MluI-HindIII*
94 fragment from the resulting plasmid containing the $P_{lac-dsred.T3_S4T}$ fusion was subcloned
95 into pME6031 (Heeb *et al.*, 2000) to give pME7134. This reporter vector provides bright red
96 fluorescence and can be used in a wide variety of Gram-negative bacteria.
97 Starter single-species cultures of *Azospirillum brasilense* Sp245/pMP2444 and *Pseudomonas*
98 *protegens* CHA0/pME7134 were grown in Luria-Bertani medium (LB) (Sambrook, *et al.*, 1989)
99 at 30°C for 18 h with orbital shaking (100 rpm). When required, final antibiotic concentrations
100 were Gm 25 $\mu g mL^{-1}$ for *A. brasilense* and Tc 125 $\mu g mL^{-1}$ for *P. protegens*. For growth,

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3 101 biofilm and colony morphology assays, Nfb-NO₃⁻-iron-enriched media (Nfb-Fe-NO₃⁻) was used
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5 102 according to (Döbereiner & Day, 1976) with the following modifications: 27.6 mM malic acid,
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7 103 13.8 mM KNO₃, 0.17 mM FeCl₃ and 0.015 mM Fe-EDTA. For preparation of stationary-phase
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9 104 cell free media, bacteria were grown overnight or for three days (overgrown culture) in Nfb-Fe-
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11 105 NO₃⁻ without antibiotics, pelleted and the supernatant were filter-sterilized (0.22 µm pore size).
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13 106

107 **Bacterial growth and biofilm quantification**

108 Static growth of Sp245/pMP2444 and CHA0/pME7134 was analyzed in axenic conditions on
109 polystyrene microtiter 96 flat bottom-well plates. Single cultures were initiated with 200 µL of
110 fresh Nfb-Fe-NO₃⁻ medium containing 10⁵ cells.mL⁻¹ based on optical density (OD₆₀₀; Spectra
111 MR; Dynex Technologies). Mixed cultures were initiated with 5.10⁴ cells of each strain in a
112 total volume of 200 µL in order to reach a final concentration of 10⁵ total cells.mL⁻¹. To test the
113 effect of CHA0 cell free spent medium on the growth of Sp245, 100 µL of Nfb-Fe-NO₃⁻ cell
114 filtered media were added to 100 µL of fresh medium containing 5.10⁴ Sp245 cells.mL⁻¹. The
115 opposite combination was also tested. When necessary, different combinations of bacterial
116 densities were tested varying the initial number of Sp245 cells: 10⁴, 10⁵ or 10⁶ but keeping 10⁵
117 cells of CHA0. Plates were statically incubated for 2 days at 30°C. For total cell growth
118 quantification, biofilms developed in the wells were vigorously pipetted with a sterile tip for
119 mechanical disaggregation and mixing with planktonic cells. OD₆₀₀ and fluorescence (ABI 7500
120 in fluorimeter mode: Ex: 538 nm, Em: 605 nm for CHA0/pME7134 and Ex: 460, Em: 525nm
121 for Sp245/pMP2444) were registered. Cells in the biofilm were quantified in the same way after
122 discarding planktonic cells of paralleled wells. Percentage of biofilm-associated cells was
123 calculated as Biofilm-associated cells/ Total cells × 100. Biofilms formed in polystyrene
124 paralleled wells were also quantified by crystal violet dye staining (OD₅₅₀) and

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3 125 normalized to total bacterial growth (OD_{600}) as previously described (Arruebarrena Di Palma *et*
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5 126 *al.* 2013).

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7 127 The experimental design consisted of three to nine replicated plates depending on the measured
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9 128 variable, in which three independent starter cultures were each sown in triplicate wells. Results
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11 129 were statistically analyzed by ANOVA and Tukey test. Correspondence between the
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13 130 fluorescence values and the $CFU.mL^{-1}$ of bacterial suspensions was determined by linear
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15 131 regression for each strain (Sp245/pMP2444 R square: 0.9869 and CHA0/pME7134 R square:
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17 132 0.9855; Graphpad Prism 5.03 software). Then, $CFU.mL^{-1}$ values were indirectly estimated by
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19 133 measuring fluorescence and interpolating values from the curve (Fig S3). In order to ensure the
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21 134 correspondence of fluorescent measurements to $CFU.mL^{-1}$, CFU counts in total and biofilm
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23 135 fractions were analyzed by the drop method on Nfb-Fe- NO_3^- medium supplemented with $25\mu g$
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25 136 mL^{-1} Gm or $125\mu g mL^{-1}$ Tc for Sp245/pMP2444 or CHA0/pME7134, respectively. Controls to
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27 137 evaluate the level of fluorophore crosstalk were done by registering values of fluorescence (red:
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29 138 Ex: 538 nm Em: 605 nm and green: Ex: 460nm Em: 525nm) for different concentrations of
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31 139 CHA0/pME7134. A regression curve between green and red fluorescence (R^2 : 0.9905) was
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33 140 done. Correction of crosstalk was made by subtracting the value of green fluorescence obtained
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35 141 from the curve to the fluorescence measured in co-cultures.
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38 39 40 41 42 43 **Colony morphology experiments**

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45 144 Suspensions of Sp245 and CHA0 obtained from overnight cultures in LB with agitation (100
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47 145 rpm) were adjusted to 10^8 cells. mL^{-1} and 10^9 cells. mL^{-1} , respectively. Serial dilutions of each
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49 146 strain were spotted onto Nfb-Fe- NO_3^- agar plates that had been previously inoculated with 1 mL
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51 147 of either CHA0 or Sp245 lawn, or 1 mL of CHA0 or Sp245 cell-free culture filtrate (spent
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53 148 medium).

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3 149 For the drop assay, 1000 CFU of Sp245 were spread onto RC agar plate and 10 μ l of a
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5 150 suspension of CHA0 (10^9 cells.mL⁻¹) or Sp245 (10^8 cells.mL⁻¹) were spotted at the center of the
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7 151 plate, and incubated for five days at 30°C. When required, plates were incubated for 23 days.
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11 153 **Confocal microscopy**

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14 154 Polystyrene covered slides were placed in test tubes containing 3 mL Nfb-Fe-NO₃⁻ with 10⁵
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16 155 Sp245/pMP2444 cells, 10⁵ CHA0/pME7134 cells, or mixed cultures, and statically incubated
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18 156 for 2 days at 30°C. Slides were removed, smoothly washed and the biofilms formed in the
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20 157 liquid-air interface zone were directly observed with a confocal laser scanning microscope
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22 158 Nikon C1, using 40.0x/1.30/0.22 Oil spring-loaded lens. GFP and dsRed proteins were excited
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24 159 at 488 and 543 nm, and detected at 550–650 nm and 650–750 nm, respectively. Images were
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26 160 analyzed with Nikon EZ-C1 Freeviewer.
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31 162 **RESULTS**

32 163 33 34 164 **Bacterial growth is enhanced in static mixed cultures**

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37 165 Interspecific relationships between *A. brasilense* and *P. protegens* were evaluated in single and
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39 166 mixed-species static cultures of Sp245 and CHA0 strains carrying the plasmid pMP2444 (*Gm*^r,
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41 167 *P*_{lac-egfp}) or pME7134 (*Tc*^r, *P*_{lac-dsRed}), respectively. After two days of cultivation, overall
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43 168 bacterial growth was greater when both species grew together (Fig. 1A). Moreover,
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45 169 fluorescence measurements revealed that Sp245 and CHA0 cell number exhibited a 3 and 2 fold
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47 170 increase, respectively, when co-cultured (Fig. 1B and 1C). This was highly dependent on the
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49 171 initial number of cells from each strain. If the initial inoculum of Sp245 was 10⁶ cells.mL⁻¹ the
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51 172 promoting effect on growth was not significant for neither of the species (Fig. S1). Moreover,
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53 173 CHA0 cell-free spent media was sufficient to exert Sp245 growth promotion, suggesting that a
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3 174 soluble factor released by CHA0 was responsible for Sp245 growth stimulation. Furthermore,
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5 175 Sp245 cell-free stationary-phase supernatants were also able to enhance CHA0 growth (Fig.
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7 176 1C). No pH changes were detected between single and mixed cultures, so a change in media pH
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9 177 was not responsible for this effect (Fig. S2). These results suggest that a non-direct bacterial
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11 178 interaction between CHA0 and Sp245 is sufficient to improve the growth of both species.
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16 180 **Bacterial biofilm on artificial surfaces**

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18 181 The amount of biofilm in individual cultures of Sp245 or CHA0 was significantly lower than
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20 182 the amount of biofilm produced in co-cultures (Fig. 2A). Noticeably, whereas Sp245 showed a
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22 183 2-fold increase in the percentage of biofilm formed in mixed cultures, no difference was
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24 184 observed for CHA0 (Fig 2B). This implies that the overall biofilm increase in mixed cultures
25
26 185 was due to a higher proportion of Sp245 cells in the biofilm. The fact that these strains produced
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28 186 biofilm without compromising each other's growth (Fig 1B, C) suggests that they may occupy
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30 187 different spatial niches within the biofilm. As shown in Fig 2C, mixed biofilm was localized on
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32 188 the entire surface covered by CHA0 (the liquid/air interface) and by Sp245 (deeper on the wall
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34 189 and the base of the well).
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40 191 **Strain distribution within single and mixed-species biofilms**

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42 192 The structure and strain distribution within the biofilm developed in the air-liquid interface zone
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44 193 was studied on polystyrene covered slides and observed by confocal microscopy. Single-species
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46 194 Z-projections demonstrated that the Sp245 biofilm was thinner than that formed by CHA0 (Fig.
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48 195 3C, D). This is in agreement with the location of the biofilm ring observed in microplate
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50 196 experiments using crystal violet staining (Fig. 2C). We also observed during washing steps that
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52 197 the Sp245 biofilm was loosely attached to the surface, while CHA0 single-species biofilm was
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54 198 as tightly attached as the mixed biofilm. CHA0 single species biofilm also showed a higher cell
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3 199 density and homogeneity on the layers closer to the slide (not shown) and a sponge-like
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5 200 structure with channels in the upper layers (Fig. 3C). Surprisingly, CHA0 single species biofilm
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7 201 was much thicker than the mixed-species biofilm, reaching a depth of about 600 μm (25 μm for
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9 202 Sp245 and 200 μm for Sp245+CHA0). As it was previously demonstrated, quantification
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11 203 performed on microtiter plates revealed higher levels of bacterial biofilm in mixed than in
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13 204 single-species cultures. This apparent contradiction might be explained by the fact that Sp245
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15 205 biofilm formed all over the microplate well (surface, walls and bottom) during static co-culture
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17 206 with CHA0, which would account for the observed increase in total biofilm mass (Fig. 2C).
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19 207 Mixed biofilms were multilayered and complex in structure and showed a particular strain-
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21 208 specific distribution within the biofilm. Sp245 cells were mainly placed towards the slide
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23 209 forming the bottom layers of the biofilm, whereas CHA0 cells were located towards the biofilm
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25 210 surface, in contact with the liquid medium (Fig. 3A, B). Nevertheless, a small fraction of Sp245
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27 211 cells was also located close to the surface (Fig. 3A, B).
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213 **Sp245 colony diameter is reduced by CHA0**

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35 214 Sp245 colonies were noticeably smaller nearby a CHA0 spot, whereas there was no difference
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37 215 in Sp245 colony diameter close to the control spot (Fig. 4). The diameter of Sp245 colonies was
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39 216 restored at distances higher than 7.5 mm away from the CHA0 spot. This result suggests that a
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41 217 diffusible metabolite produced by CHA0 cells is responsible for the observed phenotype. When
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43 218 plates were cultivated for longer periods (23 d), the small colony phenotype of Sp245 in the
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45 219 proximity of CHA0 spot remained unaltered (data not shown). Small colonies picked from this
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47 220 region of the plate were re-streaked onto fresh medium and they displayed a normal sized
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49 221 phenotype (Fig 4C). To determine whether the reduction of Sp245 colony size requires the
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51 222 presence of CHA0 cells, serial dilutions of both strains were placed over a CHA0 or Sp245
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53 223 strains lawn (1 mL of a 10^6 CFU.mL⁻¹ suspension), or over their overnight spent cell-free media,
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3 224 and incubated for 3d at 30°C. Sp245 developed small colonies in the presence of a CHA0 lawn
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5 225 (Fig. 5A, D), while CHA0 colonies remained unaffected when grown over Sp245 lawn (Fig.
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7 226 5A, inset). However, Sp245 colonies grown in plates with CHA0 cell-free media remained
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9 227 unaffected (Fig. 5A, E). This indicates that the metabolite responsible for the reduction in Sp245
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11 228 colony size in co-culture with CHA0 in agar plates (Fig. 4A, B), was either absent or in
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13 229 inadequate concentration in this experimental setup. A CHA0 cell-free filtrate from overgrown
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15 230 cultures (3 days, DO₆₀₀ = 6) was used to test this possibility; however, Sp245 colonies again
16
17 231 exhibited a normal size, discarding a dilution effect (data not shown). Taken together, the results
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19 232 indicate that the development of Sp245 small colonies requires the presence of CHA0 cells in
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21 233 their proximity, or alternatively, a metabolite that is only secreted in cells growing in solid
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23 234 medium and that is induced by the presence of Sp245.
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236 **DISCUSSION**

237 PGPR have been studied for their ability to stimulate the growth and health of economically
238 important crops (Lugtenberg & Kamilova, 2009, Hayat, *et al.*, 2010, Beneduzi, *et al.*, 2012,
239 Bhattacharyya & Jha, 2012, Drogue, *et al.*, 2013). However, cell-to-cell communication
240 between PGPR and other rhizosphere-inhabiting microorganisms has received much less
241 attention. Static co-cultivation of strains Sp245 and CHA0 results in a clear interspecific
242 cooperation, as overall growth was enhanced for both species (Fig. 1 and S4). Moreover, cell-
243 free stationary-phase supernatants from CHA0 and Sp245 cultures were sufficient to exert
244 interspecies growth promotion (Fig. 1C). Bacteria are found to interact synergistically, by either
245 providing nutrients, removing some inhibitory products, or stimulating each other through
246 physical or biochemical mechanisms (Burmølle *et al.*, 2006). Growth enhancement upon CHA0
247 and Sp245 interaction can be explained by several mechanisms like pH changes,
248 exopolysaccharide (EPS) production, microaerophilic conditions and/or release of soluble

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3 249 metabolites that could modify medium redox conditions, among others. However, experiments
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5 250 conducted with spent media (where neither microaerophilic conditions nor EPS production can
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7 251 be determinant factors) also increased interspecies growth (Fig 1 B and C). Thus, these
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9 252 observations support the hypothesis that soluble metabolites released by CHA0 and Sp245 are
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11 253 responsible for this growth stimulation. The secondary metabolite DAPG is one of the most
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13 254 important antimicrobial compounds produced by certain *Pseudomonas* strains and can act as a
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15 255 relevant interspecific signal molecule (Notz, *et al.*, 2001, Combes-Meynet, *et al.*, 2010, Yang &
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17 256 Cao, 2012, Powers, *et al.*, 2015). In this regard, sub-inhibitory concentrations of DAPG promote
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19 257 *A. brasilense* growth *in vitro* (Couillerot, *et al.*, 2011), suggesting that DAPG can contribute to
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21 258 the observed *A. brasilense* growth promotion. Further experiments with CHA0 strains unable to
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23 259 produce DAPG might be interesting to unravel this issue.

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27 260 The position of the biofilm in the well was different for both species in single cultures, in
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29 261 agreement with their preferences for O₂ concentration: it was located deep in the well for Sp245,
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31 262 where microaerobiosis is generated, and in the upper part for CHA0, a strictly aerobic
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33 263 microorganism. However, in mixed cultures the biofilm was developed at the same superficial
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35 264 location than for strain CHA0 alone (Fig. 2C). We propose that in co-culture, O₂ consumption
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37 265 by *Pseudomonas* biofilm allowed *Azospirillum* to exploit the generated microaerophilic zones
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39 266 located in the well at an upper location than when cultivated alone. In line with this hypothesis,
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41 267 confocal images showed multilayered mixed biofilms located on the air-liquid interface zone,
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43 268 with Sp245 strain mainly restricted to the bottom of the biofilm in contact with the polystyrene
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45 269 covered slide, and with CHA0 mainly located in contact with culture medium (Fig. 3A, B).

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47 270 Alternatively, the enhanced attachment of Sp245 in co-cultures might be the result of its co-
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49 271 aggregation with CHA0. *Azospirillum* cells aggregate and flocculate under diverse stress
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51 272 conditions and in the presence of various carbon and nitrogen sources (Burdman, *et al.*, 1998,
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53 273 Joe, *et al.*, 2009). The ability of *Azospirillum* to co-aggregate with other species endures its
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3 274 tolerance to desiccation, heat, and osmotic shock, at different degrees depending on the identity
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5 275 of the species involved (Joe, et al., 2009). Recently, Ren, et al. (2015) showed the prevalence of
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7 276 synergistic effects in biofilm formation among isolates from different soils when co-cultured in
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9 277 combinations of four species. Moreover, absolute individual strain cell numbers were
10
11 278 significantly enhanced when compared with those of single-species biofilms, indicating that all
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13 279 the individual strains benefit from inclusion in the multispecies community (Ren et al., 2015).
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15 280 The fact that the *A. brasilense* biofilm is thicker (Fig. 2A) and denser (Fig. 3) when co-cultured
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17 281 with *P. protegens* suggests that this particular interspecific relationship could be an adaptation
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19 282 to complex environments where microbial communities co-exist. This response may result in an
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21 283 increased ability of *A. brasilense* to colonize roots in the rhizosphere when both microorganisms
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23 284 are present. On the contrary, the outcome of the interaction depends on the species partner, as it
24
25 285 has been reported that biofilms of *Bacillus subtilis* are inhibited by *P. protegens* and *P. putida*
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27 286 (Powers, et al., 2015). To our knowledge, our work is the first to demonstrate a true
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29 287 collaboration between statically co-cultured *A. brasilense* and *P. protegens* that is reflected not
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31 288 only in an increase in the total cell numbers, but also in a distribution of niche occupancy within
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33 289 an interspecies biofilm.
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37
38 290 On the other hand, interaction on agar plates revealed that Sp245 colonies are non-pigmented
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40 291 and noticeably smaller nearby a CHA0 spot, but they are unaffected when spread onto agar
41
42 292 plates containing a cell-free CHA0-spent-medium (Fig. 5A, D and E). The smaller Sp245
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44 293 colonies around a CHA0 spot cannot be interpreted as slowly-growing colonies since their size
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46 294 remained unaltered even after 23 days of culture (data not shown). An inoculum of Sp245
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48 295 produced ca. 10^8 CFU. mL⁻¹ when plated either on fresh media or on a lawn of CHA0 (Fig. 5B).
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50 296 The fact that the number of CFU.mL⁻¹ of the same Sp245 inoculum is not diminished, or even
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52 297 showed a statistically non-significant tendency to augment, revealed that CHA0 did not affect
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54 298 Sp245 viability but only its colony phenotype. A nutrient depletion effect cannot be discarded in
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3 299 this experimental approach in which a lawn of CHA0 coexists with colonies of Sp245. Although
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5 300 Nfb-Fe-NO₃⁻ is a complete rich medium supplemented with nitrate to favor biofilm formation
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7 301 (Arruebarrena Di Palma, *et al.*, 2013) and with iron to limit siderophores production that can
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9 302 impair Sp245 growth, a different experimental set up in which CHA0 was spotted in the center
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11 303 of the plate showed a radial negative effect on Sp245 colony development (Fig. 4A, B). This
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13 304 undoubtedly accounts for one or more soluble factors that diffuse and affect the size of Sp245
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15 305 colony. The small non-pigmented colony phenotype is usually observed in pathogenic bacteria
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17 306 under stress conditions. Such variants are called *small colony variants* (SCV) and have been
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19 307 extensively studied in several genera including *Pseudomonas* (Häußler, *et al.*, 1999). SCV
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21 308 constitute a subpopulation of non-pigmented colonies that are ca. 10 times smaller than their
22
23 309 counterparts on agar plates and less susceptible to antibiotics (Proctor, *et al.*, 2006). The
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25 310 observed effect of CHA0 on Sp245 colonies might be a result of exposition to a stressful
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27 311 metabolite that induces a SCV-like phenotypic response. The fact that Sp245 small colonies
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29 312 restore to a normal colony phenotype upon being streaked in fresh medium (Fig 4C), implies
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31 313 that the probability of plasmid loss or genetic rearrangement is negligible in the SCV.
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33 314 Moreover, the radial and dose-dependent SCV phenotype observed in our experiments (Fig 4A,
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35 315 B) is neither consistent with genetic rearrangements. Antibiotic production by *P. protegens*
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37 316 CHA0 is tightly regulated by the post-transcriptional cascade Gac/Rsm. This signal transduction
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39 317 pathway regulates the production of important secondary metabolites for the biocontrol of root
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41 318 pathogens (Lapouge, *et al.*, 2008). Since the SCV-like phenotypic response of Sp245 might be
42
43 319 the result of exposition to stressful metabolites such as those released by CHA0, future studies
44
45 320 using CHA0 Gac/Rsm mutants, which are impaired in antibiotic production, could be useful to
46
47 321 elucidate the participation of metabolites secreted by CHA0 in the development of Sp245 SCV.
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49 322 In conclusion, the observations presented here suggest that interspecies co-culture forming
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51 323 mixed biofilms is a favorable way of cooperation, in which the growth of interacting strains is
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3 324 enhanced, and the development of spatially structured biofilms may be a cooperative solution
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5 325 for better exploitation of available resources. The two species used in this work are major
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7 326 inhabitants of rhizosphere and their mutually cooperative interaction could probably prevail in
8
9 327 this habitat. There is a need for integrative studies in soil microbiology, and a holistic
10
11 328 consideration of the interactions between the various species at play in the rhizosphere would
12
13 329 undoubtedly improve management of the rhizosphere microbiota and PGPR performance.
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16 330

17 331 **FUNDINGS**

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19
20 332 This work was supported by ‘Secretaría de Ciencia y Técnica’, UNMdP, Argentina (AGR
21
22 333 411/15). LP and CC are researchers from UNMdP, Argentina. CK is researcher from University
23
24 334 of Lausanne, Switzerland. GM, LP and CV are researchers from CONICET, Argentina. FS is
25
26 335 doctoral fellow from CONICET, Argentina.
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29 336

30 337 **ACKNOWLEDEMENTS**

31
32
33 338 We thank Dr. M. Pechy-Tarr and Dr. E. Baehler (Département de Microbiologie Fondamentale,
34
35 339 Université de Lausanne, Switzerland) for help with the construction of pME7134 plasmid, Dr. J.
36
37 340 Carella (Departamento de Ciencia e Ingeniería de Polímeros, FI, UNMdP) for valuable
38
39 341 assistance with polystyrene slides and S. Larraburu for technical assistance.
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Fig 1

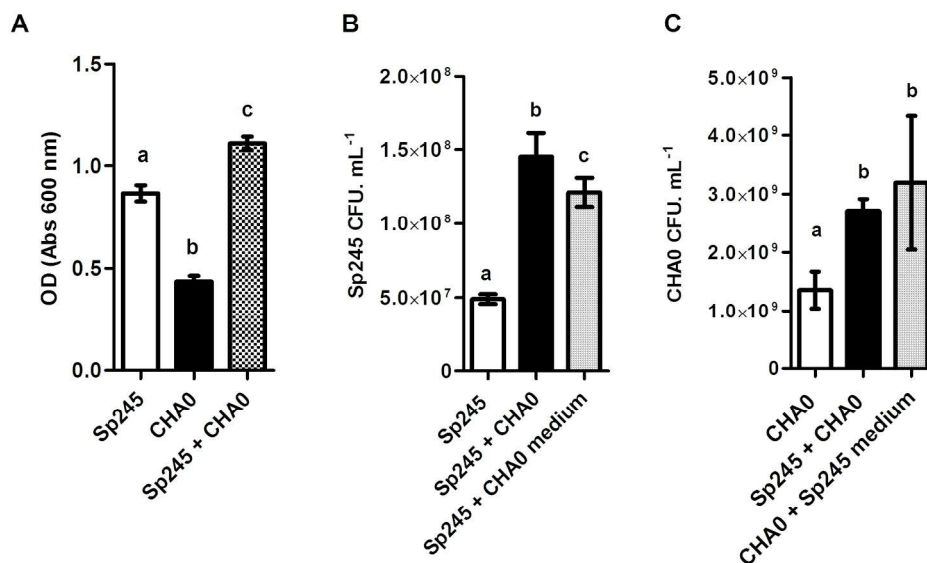
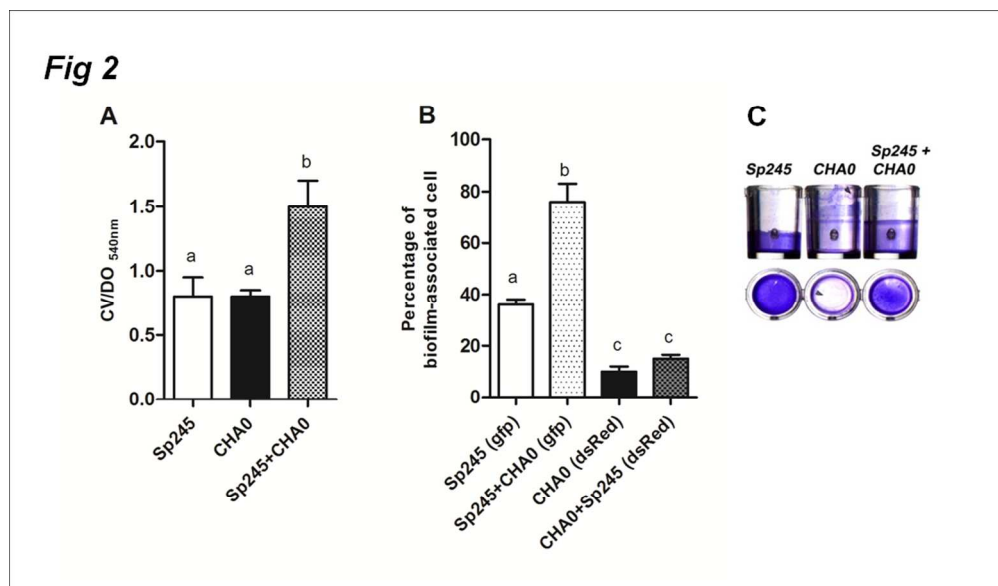


Fig. 1. Static co-cultivation enhances growth of *A. brasilense* Sp245 and *P. protegens* CHA0 strains. Total growth (planktonic plus attached cells after disaggregation) of *A. brasilense* Sp245 and *P. protegens* CHA0 strains was analyzed after 2 days of static culture in Nfb-Fe-NO₃⁻ at 30°C in 96 wells plates. Single cultures were initiated with 10⁵ cells and mixed cultures with 5.10⁴ cells of each strain. Optical density at 600nm (A) and fluorescence of Sp245 (GFP) (B) and CHA0 (dsRed) (C) were registered. CFU. mL⁻¹ was obtained from linear regression with fluorescence values for each strain (Fig S3). Values are means ± SE of three to nine independent experiments, with three wells replicates each. Results were statistically analyzed by ANOVA and Tukey's Multiple Comparison posttests. Different letters indicate significant differences (P<0.05).

Fig 1

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Co-cultivation enhances *A. brasilense* Sp245 biofilm. Biofilm of *A. brasilense* Sp245 and *P. protegens* CHA0 strains was monitored after 2 days of static culture in Nfb-Fe-NO_3^- at 30°C in 96 wells plates. Crystal violet staining of the biofilm was quantified and normalized by the total cell growth ($\text{OD}_{550\text{nm}}/\text{OD}_{600\text{nm}}$, A). Fluorescence of disaggregated cells was also determined and normalized as percentage of biofilm-associated cells (B). Biofilm phenotypes are shown for each condition (C). Values are means \pm SE of three to nine independent experiments, with wells replicates each. Results were statistically analyzed by ANOVA and Tukey's Multiple Comparison posttests. Different letters indicate significant differences ($P < 0.05$).

Fig 2

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Fig 3

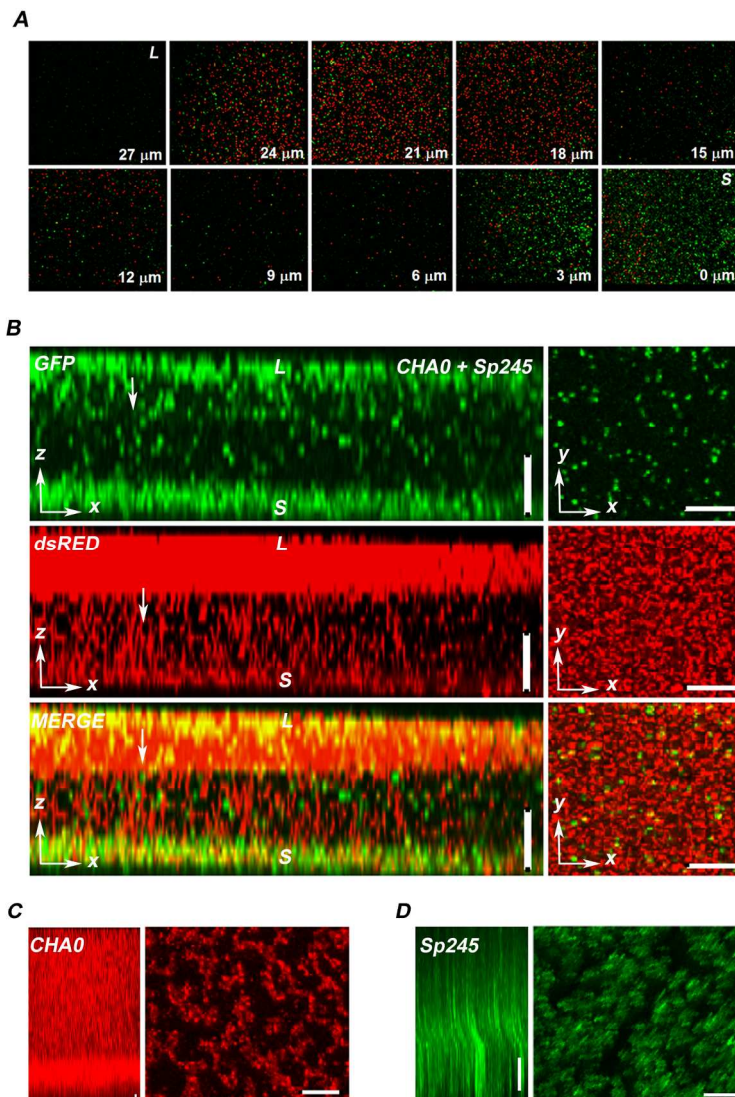


Fig. 3. Confocal laser scanning microscopy showing strain distribution within the mixed biofilm located on the air-liquid interface zone. Polystyrene covered slides were placed in 3 mL Nfb-Fe-NO_3^- with 10^5 Sp245 cells, 10^5 CHA0 cells or a combination of both strains and statically cultured for 2 days at 30°C. Fluorescence across the biofilm formed over the slides on the air-liquid interface zone was monitored, reconstruction was done with 3 μm interspace (A) and images were analyzed with a Nikon EZ-C1 Freeviewer software. Z-projections (left panels) or XY images (right panels) of Sp245, CHA0 and MERGE images on mixed (B) and individual biofilms (C and D) are shown. White arrows in the z-projection indicate the position of x-y images. Z-projection were reconstructed with 3 μm -interspace between images. L: in contact with culture media; S: in contact with slide. Bar scale: 20 μm .

Fig 3

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Fig 4

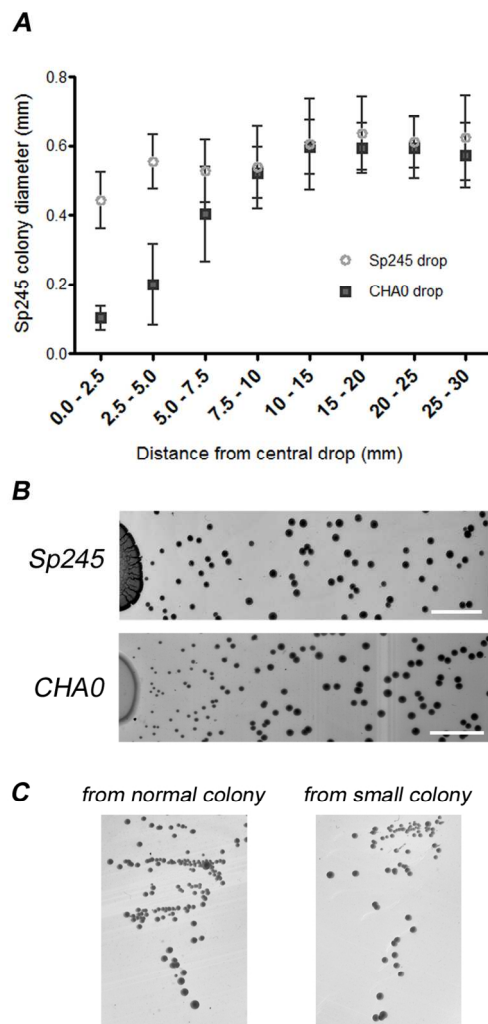


Fig. 4. Sp245 colony diameter is reduced by CHA0. 1000 CFU of Sp245 were spread onto Nbf-Fe-NO₃⁻ agar plate, and 10 μ L of a suspension of CHA0 (10^9 cells. mL⁻¹) or Sp245 (10^8 cells. mL⁻¹), was spotted at the center of the plate and incubated for five days at 30°C. Diameters of Sp245 colonies were measured (A) and colony morphology was analyzed by optical magnifier (B). Normal and small colonies phenotypes were streaked on Nbf-Fe-NO₃⁻ agar plates (C). Bar scale: 4 mm.

Fig 4

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Fig 5

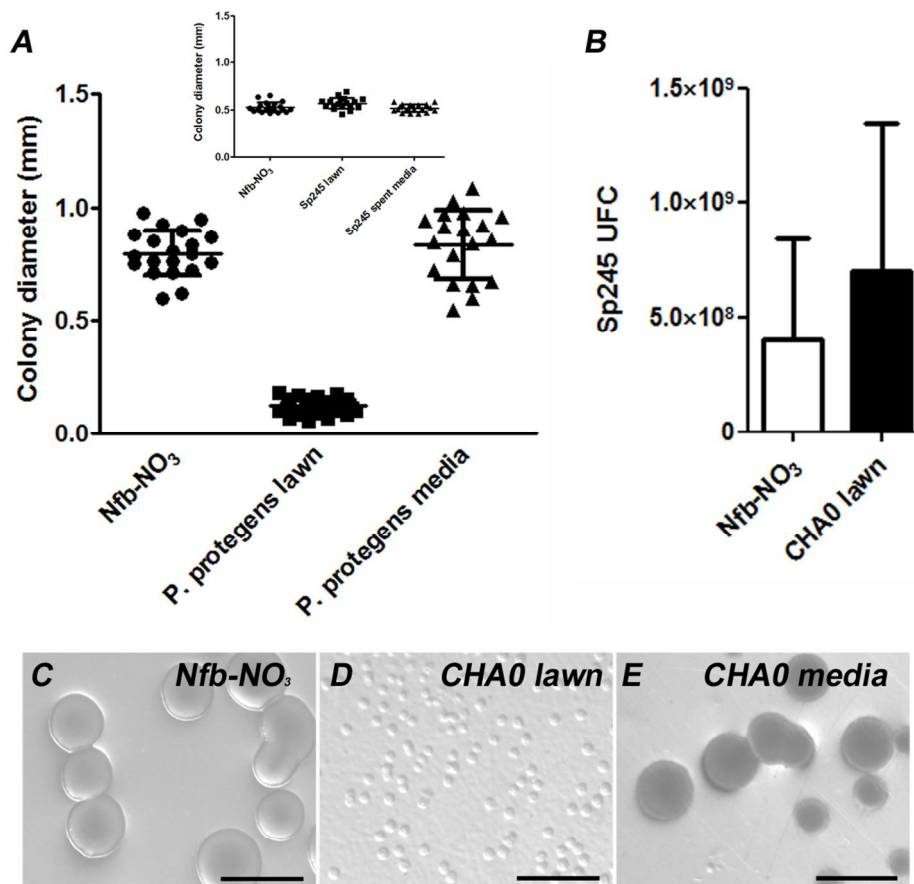
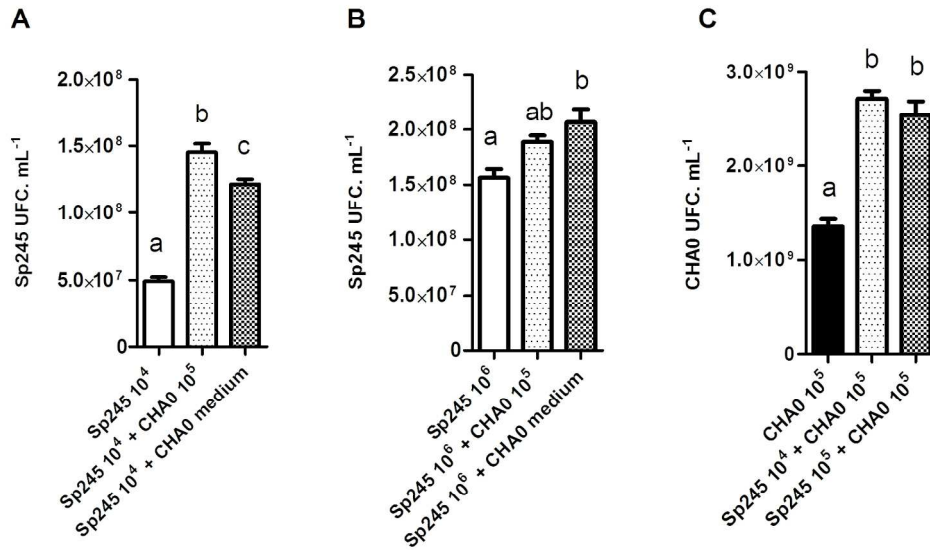


Fig. 5. Sp245 colony diameter is not affected by CHA0 stationary-phase supernatants. Serial dilutions of each strain were spotted onto Nfb-Fe-NO₃⁻ agar plates, over a CHA0 or Sp245 lawn or over cell-free culture filtrates. Sp245 (A) or CHA0 (A, inset) colonies diameter were measured, Sp245 CFU.mL⁻¹ was calculated (B) and Sp245 colony morphology was analyzed by optical magnifier (C-E). Bar scale: 1 mm.

Fig 5
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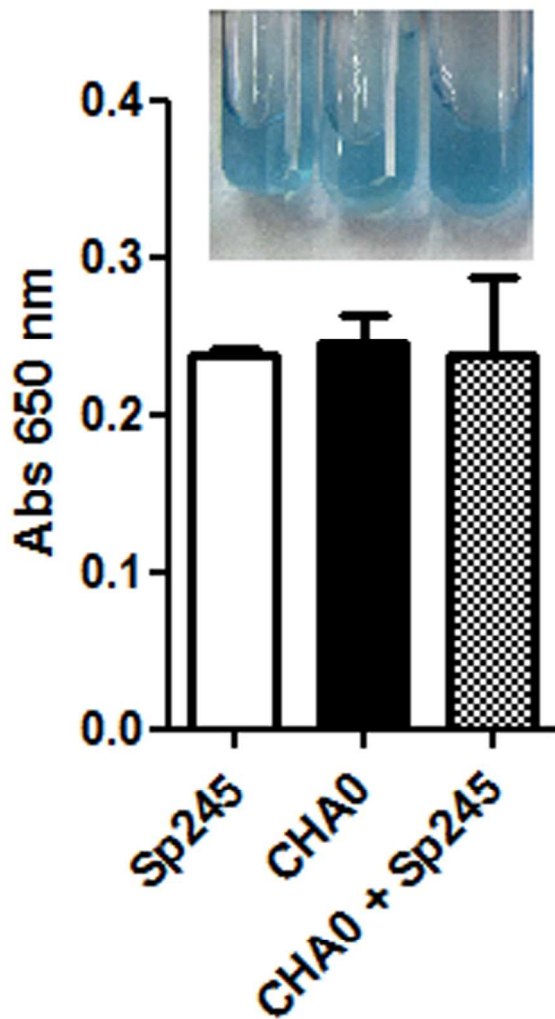
Fig S1



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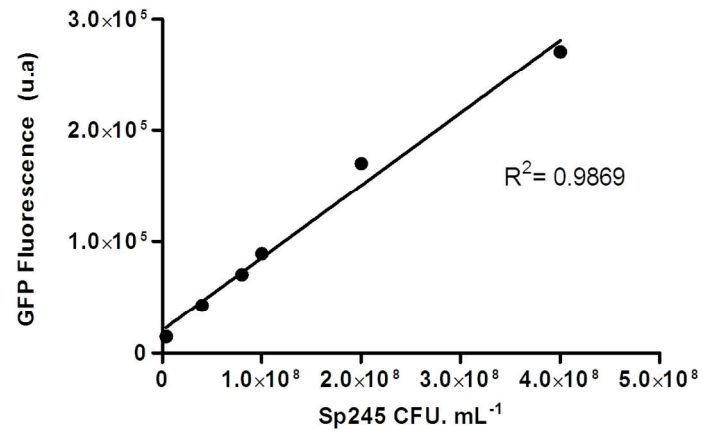
Fig S2



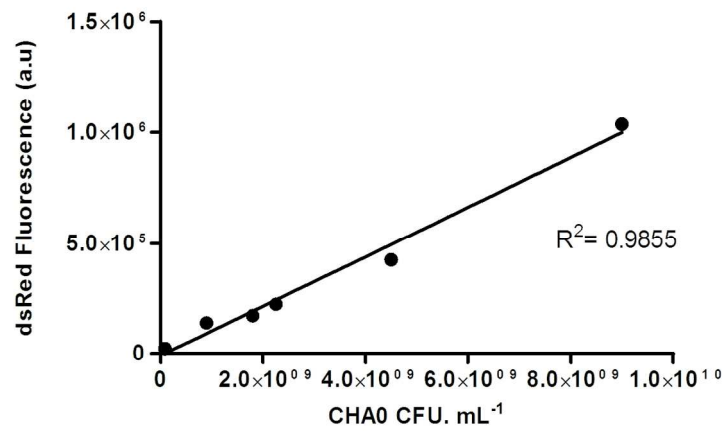
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Fig S3

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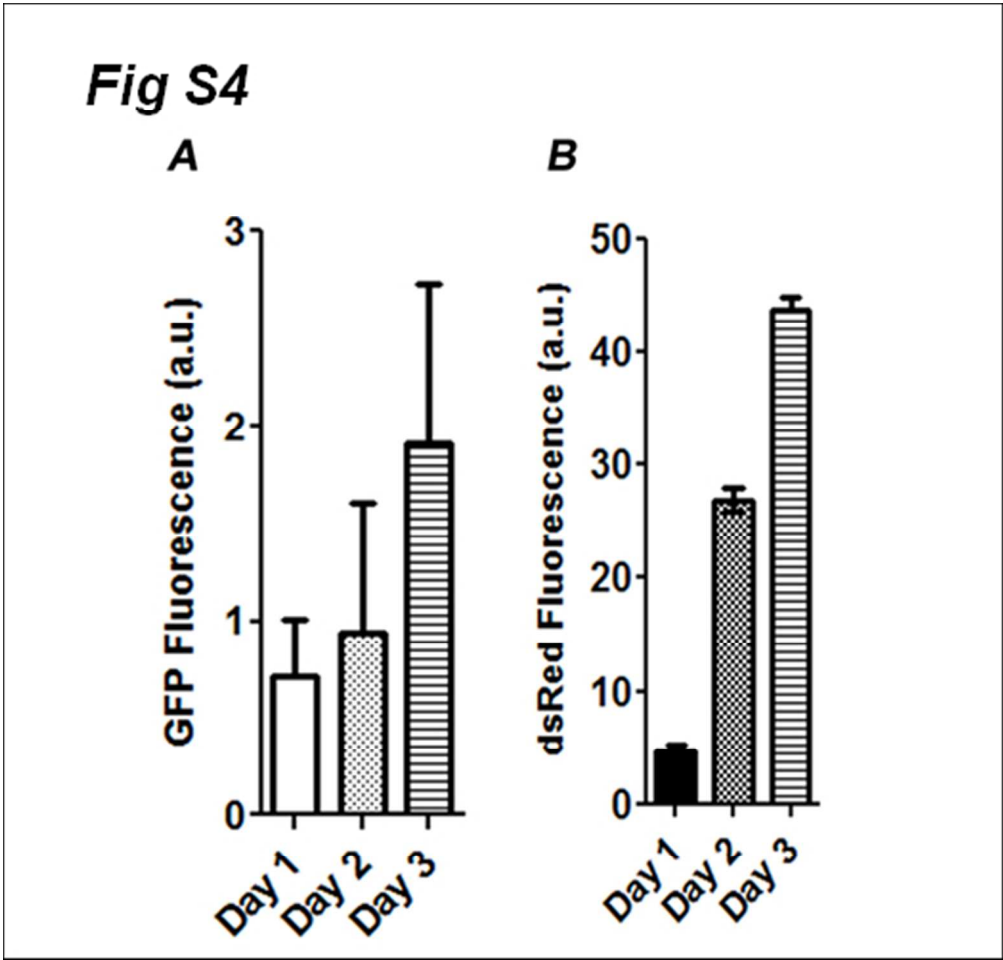


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Legends to Supplementary Figures

Fig. S1. Growth enhancement is dependent from the density of initial inocula. Total growth (planktonic plus attached cells after disaggregation) of *A. brasilense* Sp245 and *P. protegens* CHA0 strains was analyzed after 2 days of static culture at 30°C in 96 wells plates. Mixed cultures were initiated with 10^5 CHA0 cells.mL⁻¹ and 10^4 , 10^6 Sp245 cells.mL⁻¹ (A and B) or 10^4 and 10^5 Sp245 cells.mL⁻¹ (C). Fluorescence of Sp245 (A and B) and CHA0 (C) were registered. CFU.mL⁻¹ was obtained from linear regression with fluorescence values for each strain (Fig S3). Values are means \pm SE of three independent experiments, with nine replicates each. Results were statistically analyzed by ANOVA and Tukey's Multiple Comparison posttests. Different letters indicate significant differences (P<0.05).

Fig. S2. No pH changes were detected between single and mixed cultures. Media pH was analyzed after 2 days of static culture at 30°C in 96 wells with Nfb-Fe-NO³⁻ and bromothymol blue. Absorbance at 650nm of Sp245, CHA0 and mixed cultures were registered. Values are means \pm SE of three independent experiments, with nine replicates each. Results were statistically analyzed by ANOVA and Tukey's Multiple Comparison posttests.

Fig. S3. Linear regression between fluorescence values and CFU. mL⁻¹. Fluorescence measurements on serial dilutions of *A. brasilense* Sp245 (A) and *P. protegens* (B) were registered and CFU. mL⁻¹ of the same samples was determined. Linear regression was calculated between both parameters for each strain and Goodness of fit values were calculated (Graphpad Prism 5.03 software).

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3 **Fig. S4.** Total growth (planktonic plus attached cells after disaggregation) of 10^5 initial cells
4 of *A. brasilense* Sp245 (GFP) or *P. protegens* CHA0 (dsRED) strains was analyzed after
5 1, 2 or 3 days of static culture at 30°C in 96 wells plates. Fluorescence of Sp245 (A) and
6 CHA0 (B) were registered. (a.u.) arbitrary units of fluorescence.
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