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1 **Rhizosphere microorganisms enhance *in vitro* root and plantlet development of *Pyrus* and *Prunus***
2 **rootstocks**

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8 **Main conclusion:** The *in vitro* application of rhizosphere microorganisms led to a higher rooting
9 percentage in *Pyrus* Py12 rootstocks and increased plant growth of *Pyrus* Py170 and *Prunus* RP-20.

10 **Abstract**

11 The rooting of fruit tree rootstocks is the most challenging step of the *in vitro* propagation process.
12 The use of rhizosphere microorganisms to promote *in vitro* rooting and plant growth as an alternative to the
13 addition of chemical hormones to culture media is proposed in the present study. Explants from two *Pyrus*
14 (Py170 and Py12) rootstocks and the *Prunus* RP-20 rootstock were inoculated with *Pseudomonas*
15 *oryzihabitans* PGP01, *Cladosporium ramotenellum* PGP02 and *Phoma* sp. PGP03 following two different
16 methods to determine their effects on *in vitro* rooting and plantlet growth. The effects of the microorganisms
17 on the growth of fully developed Py170 and RP-20 plantlets were also studied *in vitro*. All experiments
18 were conducted using vermiculite to simulate a soil system *in vitro*. When applied to Py12 shoots, which
19 is a hard-to-root plant material, both *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 fungi were able to
20 increase the rooting percentage from 56.25% to 100% following auxin indole-3-butyric acid (IBA)
21 treatment. Thus, the presence of these microorganisms clearly improved root development, inducing a
22 higher number of roots and causing shorter roots. Better overall growth and improved stem growth of
23 treated plants was observed when auxin treatment was replaced by co-culture with microorganisms. A root
24 growth-promoting effect was observed on RP-20 plantlets after inoculation with *C. ramotenellum* PGP02,
25 while *P. oryzihabitans* PGP01 increased root numbers for both Py170 and RP-20 and increased root growth
26 over stem growth for RP-20. It was also shown that the three microorganisms *P. oryzihabitans* PGP01, *C.*
27 *ramotenellum* PGP02 and *Phoma* sp. PGP03 were able to naturally produce auxin, including indole-3-acetic
28 acid (IAA), at different levels. Overall, our results demonstrate that the microorganisms *P. oryzihabitans*

29 PGP01 and *C. ramotenellum* PGP02 had beneficial effects on *in vitro* rooting and plantlet growth and could
30 be applied to *in vitro* tissue culture as a substitute for IBA.

31 **Keywords:** Auxins, Fruit tree rootstocks, Micropropagation, Plant-growth promoting fungi, Plant-growth
32 promoting rhizobacteria.

33 **Abbreviations**

34 CRD, completely randomized design; NYDA, nutrient yeast dextrose agar; PDA, potato dextrose agar;
35 PDB, potato dextrose broth; PGPMs, plant growth-promoting microorganisms; REM, root elongation
36 media; REM-V, root elongation media with vermiculite; RIM, root induction media; TSB, tryptone soy
37 broth.

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44 **Author contribution statement**

45 DC, NT and RDS contributed to the design of the study. All the experiments, as well as the data
46 collection, were conducted by DC, MC, and GS. The data analysis and interpretation were performed by
47 DC, NT, RT, and RDS. The writing and revision of the manuscript were performed by DC, NT, RT, and
48 RDS. All the authors have read the article and made critical contributions to improve the quality of the
49 manuscript.

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Introduction

53 Of the different steps of *in vitro* plant propagation, the rooting of micropropagated shoots of
54 different fruit tree species belonging to the *Prunus* genus is not easy (Quambusch et al. 2016; Wiszniewska
55 et al. 2016; Arab et al. 2018), and rooting enhancement is crucial to ensure plant production and survival
56 in soil conditions. The main procedures used to stimulate the formation of roots in *in vitro* environments
57 involve a reduction in the concentration of macronutrients, as well as the addition of exogenous auxin
58 (Iglesias et al. 2004; Dobránszki and Teixeira da Silva 2010; Goel et al. 2018; Lucchesini et al. 2019). The
59 effectiveness of the process depends on several factors, including the genotype, the type of auxin and the
60 dose of hormone applied (Magyar-Tábori et al. 2002; Ruzic and Vujovic 2007).

61 In recent years, restrictions imposed by the European Commission concerning the use of chemicals
62 in plant production (including auxin) have led to the development of new strategies to improve *in vitro*
63 rooting using more ecological sources to avoid the application of exogenous auxin (Pacholczak et al. 2012;
64 Elmongy et al. 2018). It is well known that some plant growth-promoting microorganisms (PGPMs) have
65 an impact on *in vitro* plant development, increasing plant growth or the efficacy of the propagation and
66 rooting of explants (Contesto et al. 2010; Trinh et al. 2018). Bacteria and fungi are able to produce hormones
67 such as auxin or gibberellins (Waqas et al. 2012; Iqbal and Hasnain 2013; Meents et al. 2019), making the
68 use of microorganisms a promising alternative to the use of chemical compounds. In a study conducted by
69 our research group, the plant growth-promoting effects of two fungi (*Cladosporium ramotenellum* PGP02
70 and *Phoma* sp. PGP03) and one bacterium (*Pseudomonas oryzihabitans* PGP01) isolated from *Pyrus* and
71 *Prunus* endogenously contaminated *in vitro* cultures were reported (Cantabella et al. 2020).

72 On this basis, the aim of the present study involves the evaluation of the effects of these three
73 microorganisms on root induction and development and on the growth of micropropagated plantlets from
74 different rootstocks belonging to species of the *Pyrus* and *Prunus* genera. This application has generated
75 universal interest in agricultural research as an instrument to increase abiotic stress tolerance or disease
76 resistance (Asín et al. 2011; Elias-Roman et al. 2019; Riaz et al. 2019; Silva et al. 2019), and the use of *in*
77 *vitro* tissue culture techniques serves as a way to study tolerance to abiotic stresses, such as tolerance to
78 lime-induced chlorosis (Dolcet-Sanjuan et al. 1992, 2004a, 2008; Donnini et al. 2009). To ensure better
79 plant-microbe coexistence, an *in vitro* culture system using media comprising vermiculite instead of the
80 traditional approach in which semisolid agar-containing media are used was used in this study.

81 **Material and methods**

82 **Plant materials and *in vitro* culture conditions**

83 Rootstocks for which there is commercial interest for fruit production were used in this study. A
84 *Prunus* rootstock marketed for commercial use named Rootpac® 20 (RP-20) (Agromillora Group,
85 Barcelona, Spain) and two *Pyrus* rootstocks named “Py12” and “Py170”, which are under agronomic
86 evaluation by the IRTA, were propagated by axillary branching through *in vitro* shoot tip cultures. RP-20
87 is a natural hybrid between Myrobalan plum (*Prunus cerasifera* Ehr.) and almond (*Prunus dulcis*) and can
88 be used as a rootstock for Japanese plum, peach, nectarine, almond and several apricot cultivars (Pinochet
89 2010). Py170 is a hybrid between OH11 (Simard and Michelesi 2002) and *Pyrus amygdaliformis*, while
90 Py12 is derived from the open pollination of *Pyrus communis* cv. Williams. Both clones are in the last
91 selection phase of an IRTA pear rootstock breeding programme that aims to obtain tolerance to both lime-
92 induced chlorosis and reduced vigour.

93 Shoot-tip cultures of both *Pyrus* rootstocks (Py12 and Py170) and commercial *Prunus* (RP-20)
94 rootstock were the source of shoot explants used to induce rooting and produce full plantlets. The three
95 plant materials were propagated by axillary branching in MS media (Murashige and Skoog 1962)
96 supplemented with 5 µM benzylaminopurine (BAP), as described by Iglesias et al. (2004). For shoot
97 elongation, 50 ml of liquid MS media without hormones was added to each flask after a 4-week-long culture
98 period in 100 ml of semisolid multiplication media. Root induction on *in vitro* elongated shoots was
99 conducted in ½-strength MS media supplemented with 10 µM indole-3-acetic acid (IAA), henceforth
100 referred to as RIM-10IBA, for a 7-day-long period in darkness, followed by a transfer to ½-strength MS
101 auxin-free media, henceforth referred to as root elongation media (REM). The *in vitro* coexistence of shoot
102 explants or full plantlets with the microorganisms took place in ½-strength MS media without hormones
103 but amended with vermiculite (50:40, v/v, vermiculite:medium), as described by Dolcet-Sanjuan et al.
104 (2004b), henceforth referred to as root elongation media with vermiculite (REM-V). The pH of the different
105 liquid media was adjusted to 5.7 using NaOH before the addition of gelling agar (8 g l⁻¹ in RIM and REM
106 and 6 g l⁻¹ in REM-V). For semisolid agar-containing media, 15-mm-diameter tubes, each with 15 ml of
107 media, were used. When REM-V was used, 38-mm-diameter tubes, each containing 50 ml of vermiculite
108 and 40 ml of semisolid media, were used. The media were autoclaved at 121°C for 20 min, and once they
109 had cooled, they were stored at 14°C. All *in vitro* cultures, including those in co-culture with the

110 microorganisms, were maintained in a culture chamber set at 24°C and providing 100-120 $\mu\text{mol m}^{-2} \text{s}^{-1}$
111 cool-white fluorescent light under a 16-h-light photoperiod.

112 ***In vitro* root induction and development in the presence of microorganisms**

113 The effects of co-culture with microorganisms during root induction and development were
114 evaluated using 3-cm-long elongated shoots of RP-20, Py12 and Py170. Two different protocols were
115 followed to study the impact of the three microorganisms on *in vitro* rooting. In the first protocol (method
116 1), elongated shoots were transferred to RIM-10IBA for 7 days in dark conditions. Afterward, the shoots
117 were transferred to REM-V and immediately inoculated with 1 ml of the microorganism suspension. In the
118 second protocol (method 2), elongated shoots were immersed for 5 min in 1 ml of microorganism
119 suspension and then placed in a sterile 2-cm-diameter well, which enabled immersion of 3 mm of the shoot
120 base. Afterward, the shoots were cultured in tubes with REM for one week to ensure the microorganisms
121 grew in direct contact with the shoot base. After this period, the shoots were transferred to REM-V to favour
122 root development, as was done for method 1. For both methods, a total of fifteen shoots per treatment were
123 used. Four treatments per experiment were used, namely, inoculation with PGP01 (*P. oryzihabitans*),
124 inoculation with PGP02 (*C. ramotenellum*), inoculation with PGP03 (*Phoma* spp.) and inoculation with a
125 control. The control treatment involved the addition of 1 ml of sterile water for method 1 or immersing the
126 shoot base in 1 ml of sterile water for method 2. After 8 weeks of co-culture, the rooting percentage was
127 calculated by dividing the number of rooted shoots by the total number of shoots. Morphometric and
128 photosynthetic parameters, including the number of leaves, root number, stem length (cm), root length (cm),
129 total plant fresh weight (FW, in g), root fresh weight (g) and stem fresh weight (g) were measured for each
130 plantlet, as described by Cantabella et al. (2020).

131 ***In vitro* plant development in the presence of microorganisms**

132 To test whether *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 had plant
133 growth-promoting effects, *in vitro* fully developed RP-20 and Py170 plantlets were used. In previous
134 experiments (Cantabella et al. 2020), the three microorganisms were able to increase the FW, stem length
135 and root length of rooted pear seedlings derived from *in vitro* embryo rescue.

136 For these assays, a total of ten 3-cm-long micropropagated shoots per treatment were induced for
137 rooting in RIM-10IBA for 7 days, followed by a 1- to 2-week-long culture period in REM to favour root
138 elongation and development. After this period, the shoots that displayed visible roots were removed from

139 the agar-containing semisolid media and transplanted into REM-V, followed by inoculation with 1 ml of
140 each microorganism suspension or the same volume of sterile double distilled water for the control
141 treatment. After 8 weeks of co-culture, morphometric and photosynthetic parameters, including the number
142 of leaves, root number, stem length (cm), root length (cm), total plant fresh weight (FW, in g), root fresh
143 weight (g) and stem fresh weight (g) were measured for each plantlet. The root fresh weight (g)/stem fresh
144 weight (g) ratio was calculated for each plantlet and used as an indicator of differential plant growth
145 distribution.

146 **Preparation of microorganism inocula**

147 In the present study, the bacterium *P. oryzihabitans* PGP01 and fungi *C. ramotenellum* PGP02 and
148 *Phoma* sp. PGP03 belonging to the IRTA postharvest programme PGPMs collection, were used to test their
149 potential effects on the *in vitro* rooting and growth of *Prunus* and *Pyrus* rootstocks. These microorganisms
150 were isolated from *Pyrus* and *Prunus* embryos germinated in an aseptic environment and preliminarily
151 tested in *in vitro* rooted pear plantlets, the results of which showed beneficial effects (Cantabella et al.
152 2020). *P. oryzihabitans* PGP01 cultures preserved at -80°C were grown in nutrient yeast dextrose agar
153 (NYDA: nutrient broth, 8 g l⁻¹; yeast extract, 5 g l⁻¹; anhydrous glucose, 10 g l⁻¹; and agar, 15 g l⁻¹) plates
154 at 25°C, while *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 cultures were grown in potato dextrose agar
155 (PDA: potato tissue, 200 ml; glucose, 20 g l⁻¹; and agar, 20 g l⁻¹) plates at the same temperature. Forty-
156 eight-hour-old plates of bacteria and 14-d-old plates of both fungi were used to prepare the microorganism
157 suspensions. *P. oryzihabitans* PGP01 cells were collected in phosphate buffer (70 ml KH₂PO₄ 0.2 M; 30
158 ml K₂HPO₄ 0.2 M; 300 ml of deionized water) and washed by centrifugation at 6164 × g. After suspension
159 in phosphate buffer, the bacterial concentration was adjusted with sterile distilled water at 2x10⁸ CFU ml⁻¹
160 by measuring the absorbance at 420 nm with a spectrophotometer (SP-2000 UV, Shanghai Spectrum
161 Instruments Co., Ltd., Shanghai, China). The conidia of both *C. ramotenellum* PGP02 and *Phoma* sp.
162 PGP03 were measured using a haemocytometer, and the concentration was ultimately set to 2x10⁶ spores
163 ml⁻¹ with sterile distilled water (sp ml⁻¹). In both cases, the plate dilution technique on solid PDA and NYDA
164 media was applied to calculate the true colony forming units (CFU ml⁻¹).

165 **Determination of auxin production via spectrophotometry**

166 The indole-3-acetic (IAA) production ability of *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02
167 and *Phoma* sp. PGP03 was tested spectrophotometrically according to the methods described by Gordon

168 and Weber (1951). *P. oryzihabitans* PGP01 bacteria and *C. ramotenellum* PGP02 and *Phoma* sp. PGP03
169 fungi were grown in flasks containing 50 ml of tryptone soy broth (TSB) and potato dextrose broth (PDB)
170 supplemented or not supplemented with 0.5 and 1 g l⁻¹ tryptophan for 24 h and 168 h, respectively, and
171 incubated at 25°C and 150 rpm. Bacterial and fungal cultures were pelleted by centrifugation for 10 min at
172 6164 × g, and 1 ml of the supernatants was tested for the presence of indole compounds using 2 ml of
173 Salkowski reagent (1 ml of 0.5 M FeCl₃ in 50 ml of 35% HClO₄). After 25 min of incubation in darkness,
174 the absorbance at 530 nm was measured with an SP-2000 UV spectrophotometer (Shanghai Spectrum
175 Instruments Co., Ltd., Shanghai, China). The content of auxin was determined via a standard curve of
176 synthetic IAA (Duchefa Biochemie, Haarlem, The Netherlands) at different concentrations (from 0 to 20
177 µg ml⁻¹) and treated in the same way as were the bacterial and fungal supernatants.

178 **Data analysis**

179 The different experiments were designed in accordance with a completely randomized design
180 (CRD), and the data were analysed by one- or two-way factorial ANOVA using JMP Pro Software (version
181 13.1.0, SAS Institute, Cary, NC, USA). Different letters were used to denote significant differences
182 according to Student's t-test ($p \leq 0.05$). Statistical significance was judged at the level $P < 0.05$, and Tukey's
183 test was used to separate the means within one factor when the differences were statistically significant.
184 Significant differences in *in vitro* rooting percentages between treatments and the control were analysed
185 via Fisher's exact test ($P \leq 0.05$).

186 **Results**

187 **Effects of the applications of three microorganisms on *in vitro* root induction, development** 188 **and biometric parameters**

189 To study the effect of the three microbes *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and
190 *Phoma* sp. PGP03 on *in vitro* root induction and plantlet development, microorganisms were applied
191 following two different methods: inoculation after root induction in the medium supplemented with 10 µM
192 IBA (RIM-10IBA, Method 1), and immersion of the shoot basis in microbe suspensions without root
193 induction with IBA (REM, Method 2). Using these methods, the effects of these microorganisms were
194 tested in three rootstock genotypes: the easy-to-root *Prunus* RP-20, the easy-to-root *Pyrus* Py170 and the
195 hard to root *Pyrus* Py12. To ensure the coexistence of plants and microorganisms, all the experiments were
196 conducted in REM supplemented with vermiculite (REM-V). After 8 weeks of co-culture, the *in vitro*

197 rooting percentage, as well as several parameters regarding root induction and development (number of
198 roots and root length) and plantlet development (plant, shoot and root FW) were measured.

199 As expected, the plant material had an important effect on the rooting response. The control
200 treatment of the shoots, with neither inoculation with microorganisms (method 1) nor immersion into
201 suspensions of microorganisms (method 2), induced to root in RIM-10IBA (method 1) or by a 7-day-long
202 culture in REM (method 2), followed by an 8-week-long culture in REM-V, induced 100% rooting on RP-
203 20 and Py170, while it was only 56.3% for Py12, a hard-to-root *Pyrus* clone (Table 1).

204 Inoculation with *P. oryzihabitans* PGP01 or *C. ramotenellum* PGP02 through either methodology
205 had no detrimental influence on rooting percentage, which remained at 100%, for the easy-rooting clones
206 RP-20 or Py170 (Table 1). For Py12 in co-culture with *C. ramotenellum* PGP02 and *Phoma* sp. PGP03, the
207 rooting percentage doubled, increasing to 100% when root elongation (method 1) or root induction and
208 elongation (method 2) were performed in co-culture with the tested microorganisms (Table 1).

209 The effect of the microorganisms on the number of developed roots ($P < 0.001$) and the length of
210 rooted shoots of RP-20, Py170 and Py12 (Fig. 1a) were highly influenced ($P < 0.001$) by the method of
211 inoculation. After the first protocol, we observed that *P. oryzihabitans* PGP01 was able to induce a 35%
212 significant increase in the number of roots in *Prunus* RP-20 shoots but not in either the *Pyrus* Py170 or
213 Py12 rootstocks (Figs. 1a and 2). However, this increase in the number of roots was not accompanied by a
214 greater root length for either of the three tested plant genotypes compared with non-treated plantlets (Figs.
215 1b and 2). Inoculation with *C. ramotenellum* PGP02 led to a higher number of roots for the three plant
216 genotypes tested compared with their respective controls, with increases of 57, 54 and 312% for RP-20,
217 Py170 and Py12, respectively (Figs. 1a and 2). Together with the positive effect of inoculation with this
218 microorganisms on the number of roots, a significant (135%) increase in root length was observed only in
219 *Pyrus* rootstock Py170 after 8 weeks of co-culture (Figs. 1b and 2). *Phoma* sp. PGP03 drastically affected
220 this parameter in the hard-to-root Py12 rootstock compared with the non-treated plantlets; the root length
221 increased 225% after 8 weeks of co-culture (Figs. 1a and 2). However, a significant inhibition in root
222 elongation was reported for the RP-20 and Py12 genotypes after inoculation with *Phoma* sp. PGP03 in
223 comparison to that of the control plants (Figs. 1b and 2).

224 Overall, the number of roots observed on RP-20 and Py170 shoots was similar or lower than that
225 on the control plants when they were immersed in the microorganism suspensions (method 2) with no

226 previous exogenous auxin treatment (Fig. 1a). However, with the Py12 clone, inoculation with *C.*
227 *ramotenellum* PGP02 significantly increased the number of roots (almost 50% higher than those of the
228 control) (Figs. 1a and 2). Regarding root length, inoculation by root immersion in the suspension of *P.*
229 *oryzihabitans* PGP01 induced significantly longer roots (110% increase) from shoots of Py170 (Figs. 1b
230 and 2). As observed with method 1, *Phoma* sp. PGP03 inhibited root elongation in RP-20 and Py12, and
231 this reduction was significant only in RP-20 (58% lower than that of the control) (Figs. 1b and 2).

232 Regardless of the genotype, the effects of the three microorganisms on the total plant ($P < 0.001$),
233 shoot ($P < 0.001$) and root FW ($P = 0.003$, $P = 0.004$ and $P < 0.001$ for RP-20, Py170 and Py12,
234 respectively) were affected by the method used for their application (Fig. 3a, b, c). After inoculation, during
235 root elongation (method 1) of RP-20 and Py170 with the suspension of *P. oryzihabitans* PGP01, while plant
236 FW was not affected, an important reduction in this parameter was reported in the hard-to-root Py12
237 genotype (Fig. 3a). The reduction in shoot FW occurred in the RP-20 and Py12 plantlets in response to
238 inoculation with *P. oryzihabitans* PGP01 (48 and 67% lower than that of the control, respectively).
239 Inoculation with *P. oryzihabitans* PGP01 following this method induced a nearly 70% increase in RP-20
240 root FW (Fig. 3c). However, the plant, shoot and root FWs remained unchanged or were lower than those
241 of the control plants after 8 weeks of co-culture with both *C. ramotenellum* PGP02 and *Phoma* sp. PGP03
242 inoculated according to method 1 (Fig. 3a, b, c).

243 In method 2, inoculation with *P. oryzihabitans* PGP01 significantly increased the plant and shoot
244 FW of *Pyrus* Py170 rootstock (225% higher than that of the control) but not in RP-20 or Py12 rootstock
245 (Fig. 3a). When this bacterium was applied via this method, we reported an increase in root FW for all three
246 genotypes: RP-20, Py170 and Py12 (68, 120 and 80% more than that of the non-treated plantlets) (Fig. 3c).
247 In this sense, it is important to highlight that this increase in root FW observed for RP-20 in response to
248 inoculation with *P. oryzihabitans* PGP01 was quite similar to that obtained via method 1. Surprisingly, *C.*
249 *ramotenellum* PGP02 significantly promoted plant, shoot and root FW of *Pyrus* rootstocks Py170 and Py12,
250 whereas no effects on the *Prunus* RP-20 rootstock occurred after 8 weeks of co-culture with this microbe
251 (Fig. 3a, b, c). Finally, while a negative effect of the application of *Phoma* sp. PGP03 was observed on RP-
252 20 shoots, the plant, shoot and root FW decreased (74, 86 and 171% lower than those of control,
253 respectively), but no significant changes in these parameters were found in Py170 or Py12 (Fig. 3a, b, c).

254 In general, inoculation with the three microorganisms following method 1 stimulated root
255 development, mainly increasing the number of roots of RP-20, Py170 and Py12 explants. On the other
256 hand, increased growth of plantlets was observed when the inoculation was carried out by immersing the
257 shoots in *P. oryzihabitans* PGP01 and *C. ramotenellum* PGP02 suspensions, since better values of plant,
258 shoot and root FW were observed in response to these microbes in the three genotypes tested in the study.

259 ***In vitro* plantlet development in the presence of microorganisms**

260 In this experiment, we focused our efforts in elucidating the effect of these three microorganisms
261 *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 on *in vitro* plant development of
262 the two rootstock genotypes *Prunus* RP-20 and *Pyrus* Py170. For that purpose, *in vitro* micropropagated
263 explants were inoculated with the three microorganisms in REM-V after root induction and development
264 in RIM-10IBA and REM, respectively. Again, several biometrical parameters including the number of
265 roots, root and shoot FW and root/shoot FW ratio were measured after 8 weeks of co-culture with microbes.

266 Plant material had a significant effect on root and plantlet development after inoculation with
267 microorganisms, and this interaction was significant for the number of roots ($P < 0.001$), shoot FW ($P =$
268 0.002), root FW ($P = 0.001$) and root/shoot FW ratio ($P = 0.002$). In this sense, we observed that *P.*
269 *oryzihabitans* PGP01 positively affected all the parameters concerning root growth for the RP-20 plants,
270 such as the number of roots, root FW and the root/shoot FW ratio compared with those of the non-treated
271 plantlets (Fig. 4a, b and d). This root-localized effect was also observed when the bacterium was applied to
272 Py170 *in vitro* explants, since a significant increase in the number of roots was recorded after 8 weeks of
273 co-culture (Fig. 4a and 5). On the other hand, both plant genotypes behaved differently after inoculation
274 with *C. ramotenellum* PGP02. The application of the fungus favoured the root growth of RP-20 *in vitro*
275 plantlets, increasing their root FW (129% compared with that of the control plants). This fungus also
276 increased shoot length compared with that of the non-treated plantlets (data not shown). Nevertheless, only
277 shoot growth increased when the application was carried out on the *Pyrus* Py170 explants (Fig. 4b, c and
278 5). In this sense, the shoot FW significantly increased by only 32% compared with that of the control
279 plantlets. Again, no significant plant growth promotion was observed in response to inoculation with *Phoma*
280 sp. PGP03, except for the higher number of roots observed for Py170 plantlets after 8 weeks of co-culture
281 with this fungus (Fig. 4a and 5). Other parameters, including root length and number of leaves, were not
282 significantly affected by either of the three microbes (data not shown).

283 **Determination of IAA in *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp.**
284 **PGP03 cultures.**

285 The content of IAA was determined in *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and
286 *Phoma* sp. PGP03 cultures. The ability to produce auxins was suggested in the three microorganisms, as
287 significant increases in the IAA content were detected when the different culture media were supplemented
288 with Trp. In this context, it is noteworthy to mention that the concentration of IAA detected in *P.*
289 *oryzihabitans* PGP01 cultures was higher than that observed for *C. ramotenellum* PGP02 and *Phoma* sp.
290 PGP03. In the case of *P. oryzihabitans* PGP01, an increase in the IAA content was observed when the
291 tryptophan was added to the medium in a dose-dependent manner, registering a 4.35- and a 7.07-fold
292 increase at 0.5 and 1 g l⁻¹, respectively (Fig. 6). On the other hand, IAA was also found in *C. ramotenellum*
293 PGP02 cultures when fungal cells were grown in PDB without Trp (Fig. 6), and these levels remained
294 unchanged when the lowest concentration of tryptophan was tested. However, the addition of 1 g l⁻¹ induced
295 a 3.57-fold increase in IAA levels in comparison to those recorded in PDB without tryptophan (Fig. 6).
296 Regarding PGP03, the addition of tryptophan at 0.5 or 1 g l⁻¹ to the PDB medium induced significant
297 increases on IAA content in comparison to non-supplemented medium (3.2 and 3.7-fold), but, in that case,
298 no differences between both doses of tryptophan were registered (Fig. 6).

299 **Discussion**

300 Rootstocks have important agronomic value in fruit tree crops since they are used for grafting
301 commercial varieties, providing desirable fruit production and quality and tree tolerance to edaphic factors
302 associated with abiotic and biotic stresses (Prodhomme et al. 2019). Some clones selected for their desirable
303 agronomic characteristics, especially those clones that result in low vigour, have poor rooting efficiencies
304 and root development, such as the following: apple rootstock M9 (*Malus x domestica* Borkh) (Sun and
305 Bassuk 1991; Amiri and Elahinia 2011); pear rootstocks OHxF 333, Brossier P.2274 and P.227 (Mielke
306 and Turner 2008; Necas and Kosina 2008); and peach rootstocks Lovel and *Prunus americana* (Mayer et
307 al. 2015). As a consequence, *in vitro* propagation is not feasible. Herein, owing to its tolerance to lime-
308 induced chlorosis and reduced vigour (Asín et al. 2011), Py12, a pear rootstock clone selected in an IRTA
309 breeding programme, also has low rooting efficiency compared to that of another *Pyrus* rootstock, Py170,
310 or the commercial *Prunus* rootstock Rootpac® 20 (RP-20). When *in vitro* rooting is the main obstacle to
311 obtaining complete viable plants in woody plant species (Wiszniewska et al. 2016), research to improve the

312 efficiency of this step frequently emphasizes determining the hormone type and level (Dobránszki and
313 Teixeira da Silva 2010) to be added during *in vitro* growth, though the results are highly dependent on
314 genotype. In three apple rootstocks (M.26, MM.106 and JTE-H) in which three different levels of IBA were
315 used, different responses in terms of rooting percentage and root length were observed for the different
316 genotypes (Magyar-Tábori et al. 2002).

317 Improvement in root development in peach and pear embryo-derived plantlets inoculated with the
318 microorganisms used in the present work was reported in a previous work (Cantabella et al. 2020).
319 Therefore, an *in vitro* model in which these three microorganisms (*P. oryzihabitans* PGP01, *C.*
320 *ramotenellum* PGP02 and *Phoma* sp. PGP03) were applied to *in vitro* clonally propagated *Prunus* RP-20
321 and *Pyrus* Py12 and Py170 rootstocks at different stages of *in vitro* development was constructed to
322 determine their effects on root induction, root development, and whole-plantlet *in vitro* growth. Two
323 independent experiments were conducted: first, to study the effects of the microorganisms on the *in vitro*
324 rooting efficiency of shoots, with or without an IBA root induction treatment, and second, to study the
325 microorganisms' effects on *in vitro* whole-plantlet growth. Inoculation of microorganisms following the
326 two different methods has been previously reported in semiwoody olive microcuttings under nursery
327 conditions (Montero-Calasanz et al. 2013). Herein, to improve the coexistence between shoots or plantlets
328 and microorganisms, vermiculite was added to the culture media under *in vitro* conditions. In the first
329 method, in which IBA was used to induce rooting, which is the standard protocol (Murashige and Skoog
330 1962; Iglesias et al. 2004), *Pyrus* Py12 shoots showed 56.25% rooting, while in the presence of *C.*
331 *ramotenellum* PGP02 and *Phoma* sp. PGP03, the *in vitro*-rooted shoots completely rooted (100%). In
332 accordance with our results, other endophytic bacterial isolates demonstrated that inoculation of *in vitro*
333 microshoots of two *Prunus avium* genotypes resulted in increased rooting percentage (Quambusch et al.
334 2014). Additionally, the results obtained with method 2, without IBA treatment of the shoots, confirmed
335 that the *P. oryzihabitans* PGP01 bacterium and both *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 fungi
336 could act as a natural root-promoting source, likely due to their ability to produce IAA. In this context, it
337 has been proven that the bacterium *P. oryzihabitans* PGP01 was able to produce a higher amount of IAA
338 than both fungi. All inoculated and control shoots, including those of Py12, rooted after 8 weeks of culture,
339 except those inoculated with *Phoma* sp. PGP03, which showed a decrease in this parameter. On the other
340 hand, in the IBA-induced shoots (method 1), invariably reduced growth was observed in all plantlets
341 compared with those derived from method 2. Nonetheless, the three microorganisms increased the number

342 of roots per plantlet in the three different rootstocks in a genotype-dependent way, and these increases were
343 statistically significant for *C. ramotenellum* PGP02 in all three tested rootstocks, *P. oryzihabitans* PGP01
344 in RP-20 and *Phoma* sp. PGP03 in RP-20 and Py12.

345 In the present work, the ability of both *P. oryzihabitans* PGP01 and *C. ramotenellum* PGP02 to
346 produce IAA could explain the increase in the number of roots. *P. oryzihabitans* PGP01 and *C.*
347 *ramotenellum* PGP02, in the absence of IBA (Method 2), led to better root elongation and shoot growth –
348 more significantly in the Py170- and Py12-treated shoots. Regarding *Phoma* sp. PGP03, no important
349 effects on root length were observed in this method in spite that the ability to produce auxins was proven.
350 Studies revealing the use of microorganisms in microshoots to induce rooting while avoiding the application
351 of exogenous auxin are scarce, and some authors have reported positive effects of other natural compounds
352 such as humic acid on improving the *in vitro* rooting efficiency (Elmongy et al. 2018). Herein, though an
353 improvement in rooting percentage was proven only for Py12 shoots, better growth of the three rootstocks
354 (Py170, Py12 and RP-20) was observed after exposure to *P. oryzihabitans* PGP01 and *C. ramotenellum*
355 PGP02, and this technique is a promising alternative to the use of chemical hormones in plant production.
356 Further assays employing clonal rootstocks with a compromised rooting ability under *in vitro* conditions,
357 such as M9, OHxF 333, Brossier P.2274 and P.2276, Lovell or *P. americana*, will need to be carried out to
358 corroborate and verify this effect. The performance of rhizobacteria in the *in vitro* rooting process is highly
359 documented (Quambusch et al. 2014; Kavino and Manoranjitham 2018; Perez-Rosales et al. 2018; Burygin
360 et al. 2019); however, very little is known about the role of fungi in plant root development. It has been
361 previously established that some beneficial fungi are able to produce and accumulate auxin in their mycelia,
362 which could increase the levels of auxin as well as auxin-responsive plant gene expression in *Arabidopsis*
363 *thaliana* (Meents et al. 2019). The present research constitutes the first work showing the auxin production
364 ability of *C. ramotenellum*. In the case of *P. oryzihabitans*, Belimov et al. (2015) already reported the ability
365 of this bacterium to produce different indole compounds. In the same study, potato (*Solanum tuberosum*)
366 plants showed increased root biomass when *P. oryzihabitans* was present in the rhizosphere compared to
367 that of non-inoculated plants.

368 A different approach was conducted to analyse whether *P. oryzihabitans* PGP01 bacteria and the
369 two *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 fungi are microorganisms with potential plant growth-
370 promoting effects under *in vitro* conditions. For this purpose, fully developed Py170 and RP-20 plantlets
371 were used to evaluate whether they were able to improve root and shoot growth. The effects in response to

372 inoculation with *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 were somehow different depending on the
373 genotype. In *Prunus* RP-20 rootstocks, a greater root FW was observed in response to *C. ramotenellum*
374 PGP02, and a significantly higher number of roots was registered in Py170 after the inoculation with *Phoma*
375 sp. PGP03. These results might be explained by the ability of both fungi to produce auxins. However, the
376 clear effect of *C. ramotenellum* PGP02 on the growth of the aerial parts of Py170 and RP-20 suggested the
377 role of other compounds produced by the fungus instead of auxin. Some fungi play a crucial role in plant
378 growth and development, mostly due to the endophytic production of plant hormones such as gibberellins
379 (GAs), which are involved in important physiological processes, including stem elongation (Calvo et al.
380 2014). Previous studies have demonstrated the ability of different isolates from rhizosphere fungi belonging
381 to the *Cladosporium* and *Phoma* genera to produce GAs, which resulted in increased growth of soybean
382 and cucumber seedlings (Hamayun et al. 2009, 2010). Plant growth promotion in *Atriplex gmelinii* by a
383 gibberellin-producing strain of *Gliomastix murorum* was reported (Khan et al. 2009). Thus, considering the
384 above reference, our results could suggest that GAs play a role in the enhancement of shoot growth induced
385 by *C. ramotenellum* PGP02 in *Pyrus* Py170 rootstock. On the other hand, the bacterium *P. oryzihabitans*
386 PGP01 tended to act more specifically in the roots, increasing the number of roots for both Py170 and RP-
387 20 rootstocks, as well as the root FW and root/shoot FW ratio solely in RP-20. Reprogramming root system
388 architecture in response to rhizobacteria belonging to the *Pseudomonas* and *Bacillus* genera is mediated by
389 auxin signalling in *A. thaliana* (López-Bucio et al. 2007; Zamioudis et al. 2013). Pourjasem et al. (2020)
390 recently reported that bacteria belonging to the *Pseudomonas* genus are able to release some elements into
391 the culture media as a result of the decomposition of minerals present in vermiculite that might also favour
392 root development. These results represent a step forward compared with those of a previous publication
393 where the effects of the three microorganisms were described in *Prunus in vitro* rescued embryos for the
394 first time (Cantabella et al. 2020). However, more comprehensive studies are being undertaken to
395 corroborate that this mechanism occurs in Py170 and RP-20 plants with the studied microorganisms.

396 **Conclusions**

397 In summary, the two different procedures concerning inoculation of *P. oryzihabitans* PGP01, *C.*
398 *ramotenellum* PGP02 and *Phoma* sp. PGP03 did not reduce the *in vitro* rooting percentage of the efficient-
399 to-root *Pyrus* and *Prunus* rootstocks Py170 and RP-20. On the hard-to-root *Pyrus* Py12 rootstock, both
400 fungi increased the rooting percentage from 56.25% to 100%. As a consequence of the auxin production
401 ability by some of the microorganisms, a higher number of roots was observed with these microorganisms

402 when co-cultured after IBA induction, and increased plantlet growth was observed when no synthetic auxin
403 was used. Due to the strong potential of rootstocks in fruit tree production, this study constitutes a
404 worthwhile approach to improve the rooting efficiency of difficult-to-root genotypes such as Py12.
405 Furthermore, while *P. oryzihabitans* PGP01 primarily showed a root growth-promoting effect on both RP-
406 20 and Py170 fully developed *in vitro* plantlets, a location-dependent effect in response to inoculation with
407 *C. ramotenellum* PGP02, which positively affected the roots of RP-20 and shoots of Py170 rootstocks, was
408 observed. The present results could suggest that auxin and gibberellins play a role, and the mechanisms
409 underlying this plant growth promotion will be studied in the future.

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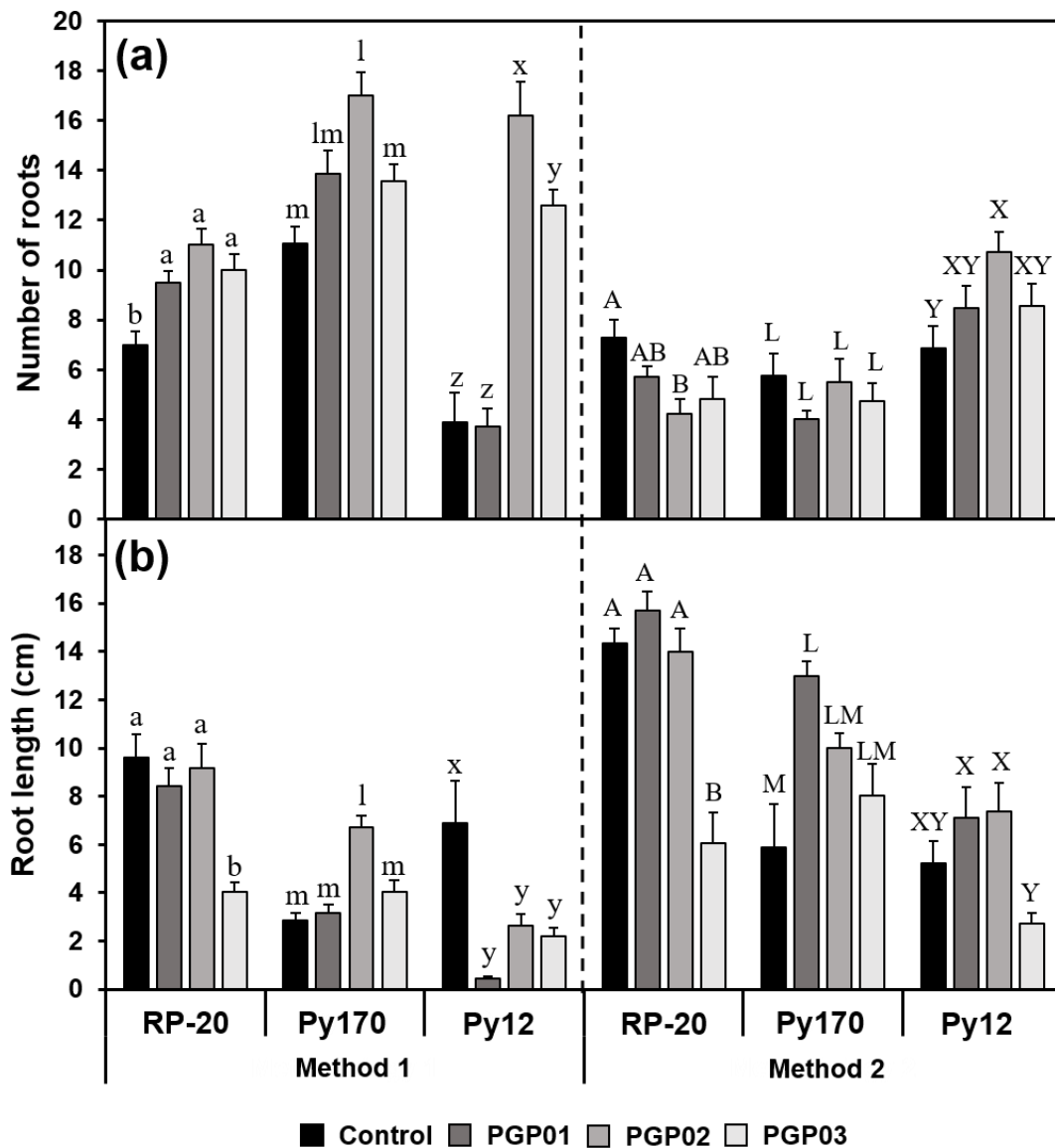
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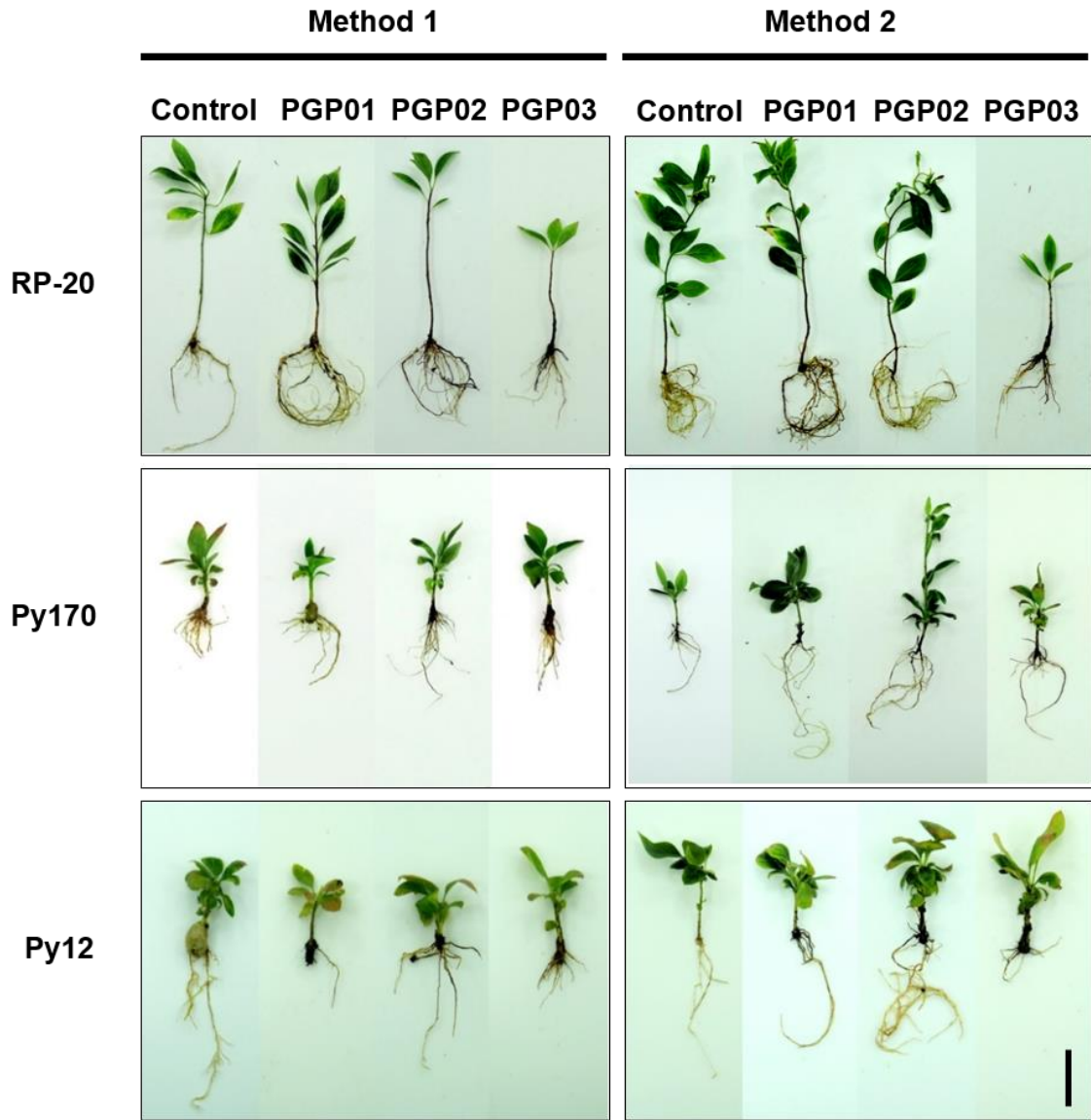


550

551 **Fig. 1** Effects of *P. oryzae* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 on the number
 552 of roots (a) and root length (b) of *Prunus* RP-20 and *Pyrus* Py170 and Py12 plantlets after 8 weeks of co-
 553 culture and inoculation by two different methods. In method 1, the shoots were inoculated with
 554 microorganisms after root induction in 10 mM IBA, while in method 2, the shoots were immersed in
 555 microorganism suspensions without previous root induction with IBA. In all cases, the data represent the
 556 means ± SEs (standard errors) of at least fifteen shoots each. The means with different letters within each
 557 genotype are significantly different according to Tukey's test ($P < 0.05$).

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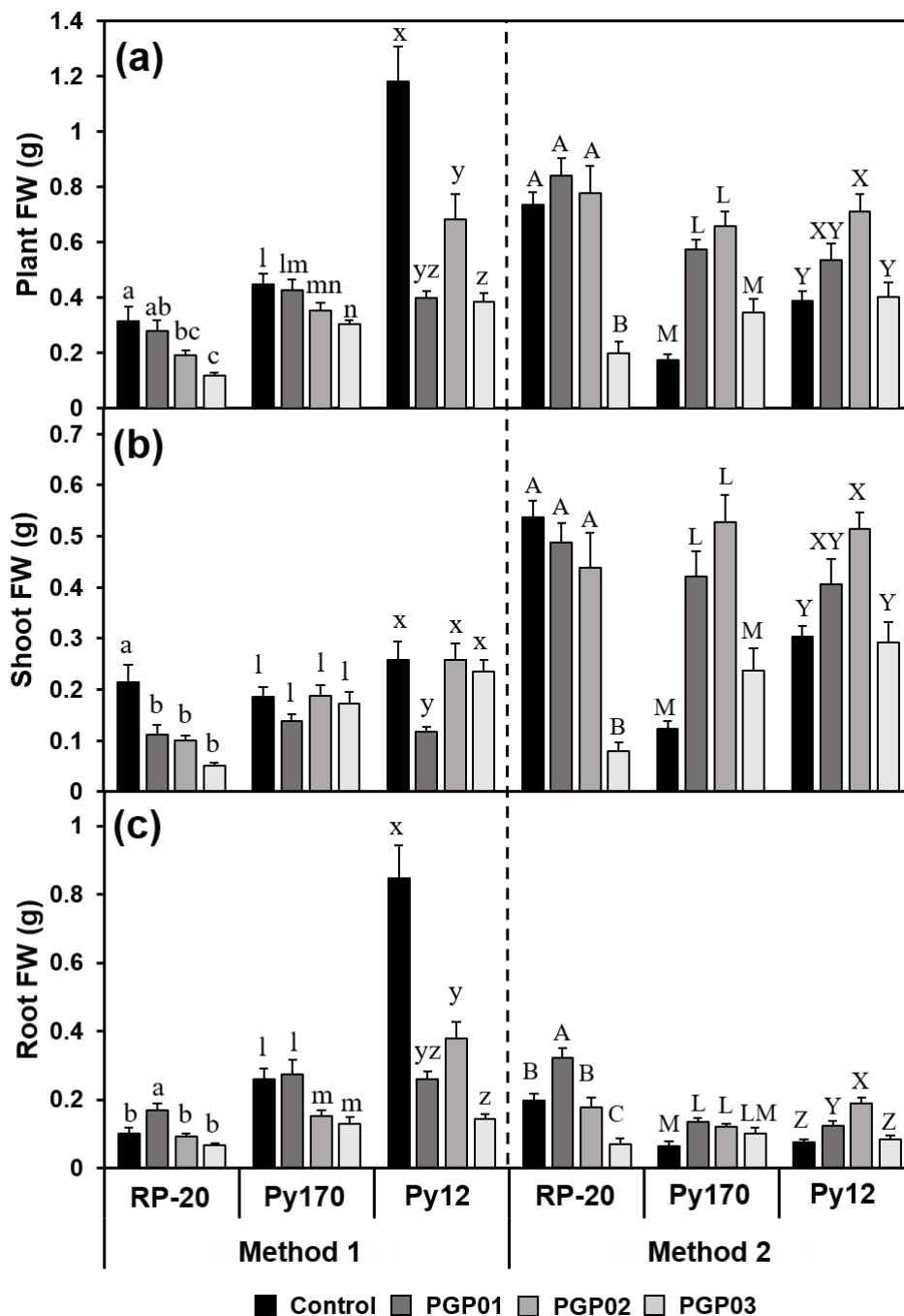
561 **Fig. 2** Plantlets of *Prunus* RP-20 and *Pyrus* Py170 and Py12 after an 8-week-long co-culture with *P.*
 562 *oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03. The shoots were inoculated with
 563 microorganisms after root induction with 10 mM IBA (method 1) or immersed in microorganism
 564 suspensions without previous IBA root induction (method 2). The black bar is equivalent to 2 cm of length.

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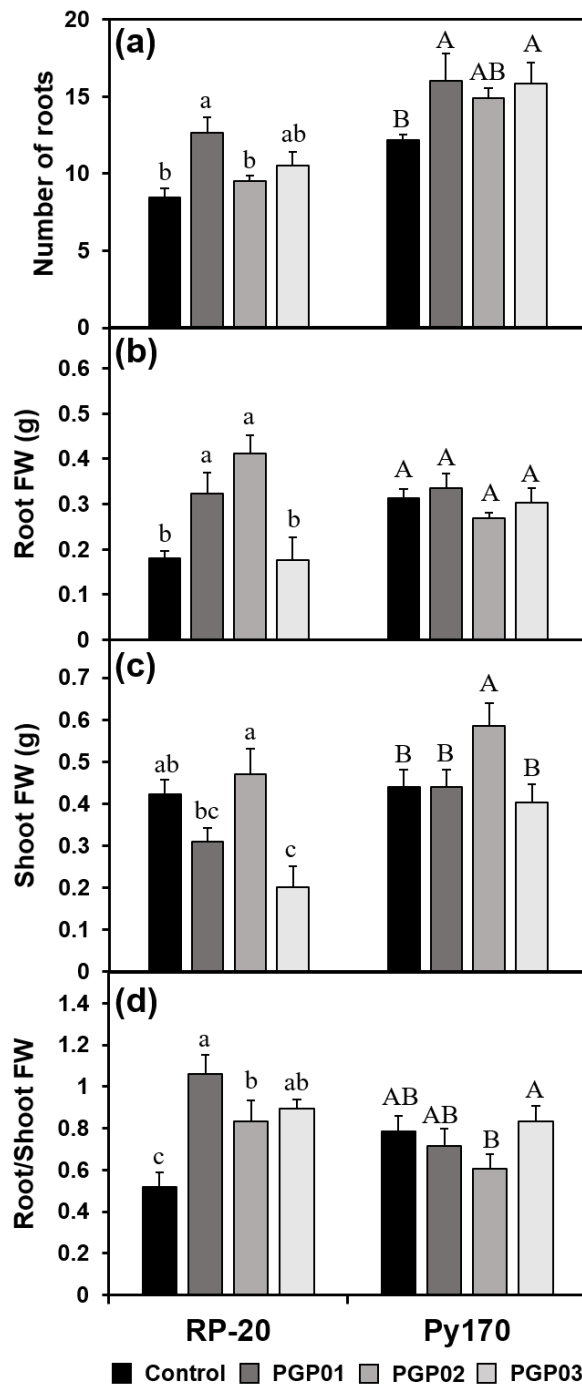
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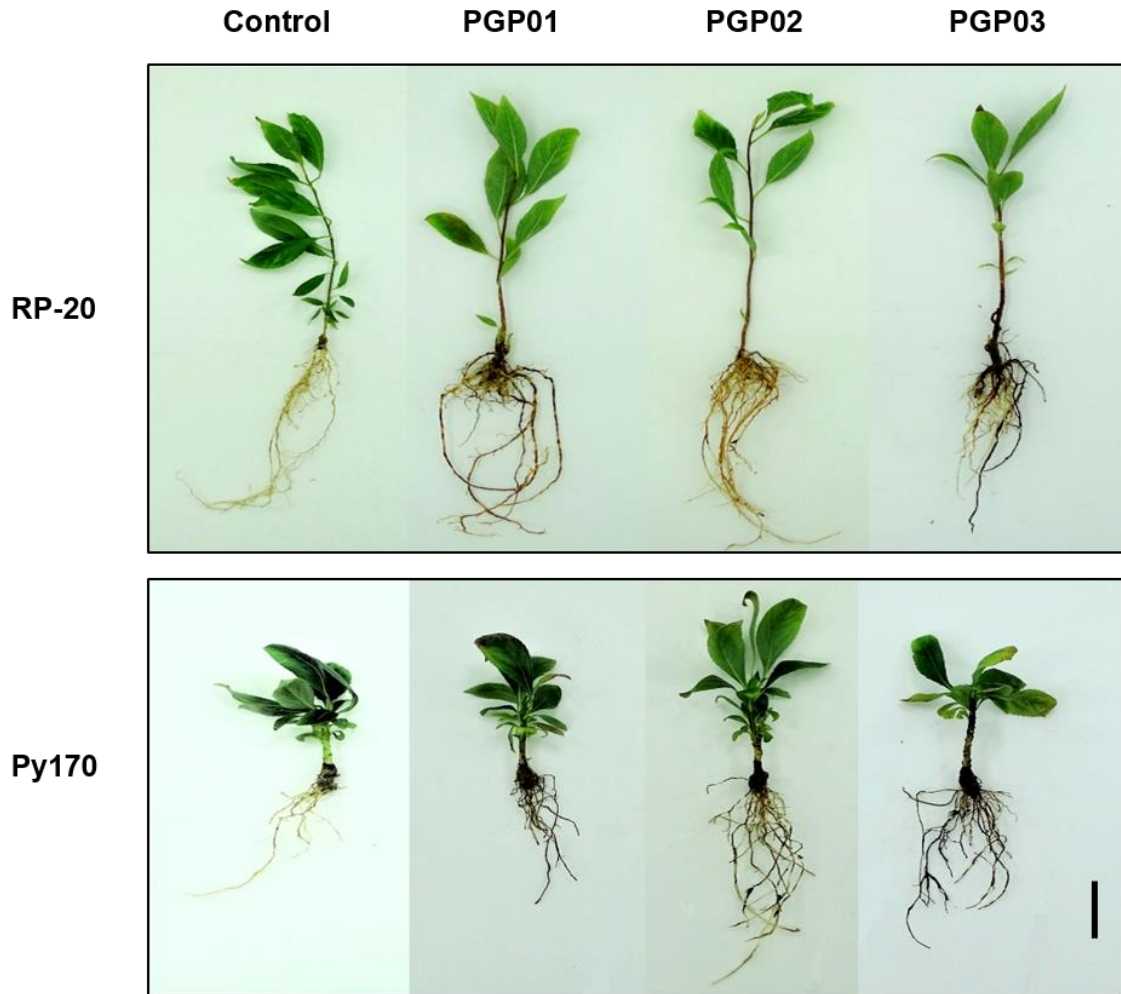
570 **Fig. 3** Effects of *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 on plant (a),
 571 stem (b) and root (c) FW of *Prunus* RP-20 and *Pyrus* Py170 and Py12 plantlets after 8 weeks of co-culture
 572 after inoculation through two different methods. In method 1, the shoots were inoculated with
 573 microorganisms after root induction in 10 mM IBA, while in method 2, the shoots were immersed in
 574 microorganism suspensions without previous root induction with IBA. In all cases, the data represent the
 575 means \pm SEs of at least fifteen plants. The means with different letters within each genotype are
 576 significantly different according to Tukey's test ($P < 0.05$).



577

578 **Fig. 4** Effects of inoculation with *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03
 579 on the number of roots (a), root FW (b), shoot FW (c) and root/stem FW ratio of *Prunus* RP-20 and *Pyrus*
 580 Py170 fully developed plantlets. The different lowercase and uppercase letters refer to significant
 581 differences between the control and treatments for RP-20 and Py170, respectively, according to Tukey's
 582 test ($P < 0.05$).

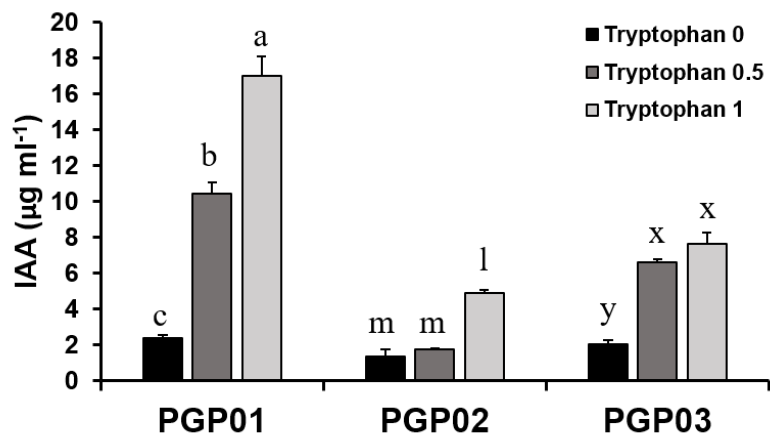
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584

585 **Fig. 5** *Prunus* RP-20 and *Pyrus* Py170 plantlets after 8-week-long co-culture with *P. oryzihabitans* PGP01,

586 *C. ramotenellum* PGP02 and *Phoma* sp. PGP03. The black bar is equivalent to 2 cm of length.



587

588 **Fig. 6** IAA production in *P. oryzihabitans* PGP01, *C. ramotenellum* PGP02 and *Phoma* sp. PGP03 cultures.

589 Bacteria and fungi were grown in TSB and PDB media, respectively, supplemented with 0.5 (dark grey

590 bars) and 1 g l⁻¹ tryptophan (light grey). Supernatants were collected after 24 and 168 h in the case of
591 bacteria and fungi, respectively, and the absorbance at 530 was measured after 25 min of incubation with
592 2 ml of Salkowski reagent. The bars represent the means ± SEs of three replicates per treatment. The
593 different lowercase and uppercase letters refer to significant differences between tryptophan-treated and
594 non-treated media according to Tukey's test ($P < 0.05$).

595