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6 **Root inoculation of strawberry with the entomopathogenic fungi *Metarhizium robertsii* and *Beauveria***
7 ***bassiana* reduce incidence of the twospotted spider mite and selected insect pests and plant diseases in**
8 **the field**

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29

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40

41 **Key message**

42

- 43 • Few studies have investigated the potential of plant inoculated entomopathogenic fungi as microbial
44 control agents under natural field conditions.
- 45 • First report of reduced *Tetranychus urticae* numbers on strawberry plants receiving root inoculation
46 with the entomopathogenic fungi *Metarhizium robertsii* and *Beauveria bassiana* under commercial
47 cultivation regimes.
- 48 • Reduction in foliar plant pathogenic fungi and no harmful effects on naturally occurring predatory
49 mites were also observed.
- 50 • This represent a new tool and an innovative biocontrol strategy that may be implemented in IPM and
51 organic strawberry production.

52

53 **Abstract**

54

55 The effect of inoculation of strawberry roots by two entomopathogenic fungal isolates, *Metarhizium robertsii*
56 (ESALQ 1622) and *Beauveria bassiana* (ESALQ 3375), on naturally occurring arthropod pests and plant
57 diseases were investigated in four commercial strawberry fields during two growing seasons in Brazil. Three
58 locations represented open field production while strawberries were grown in low tunnels at the fourth location.

59 Population responses of predatory mites to the fungal treatments were also assessed. Plants inoculated by the
60 fungal isolates resulted in significantly fewer *Tetranychus urticae* adults compared to control plants at all four
61 locations. The mean cumulative numbers \pm SE of *T. urticae* per leaflet were: *M. robertsii* (225.6 \pm 59.32), *B.*
62 *bassiana* (206.5 \pm 51.48) and control (534.1 \pm 115.55) at the three open field locations, while at the location with
63 tunnels numbers were: *M. robertsii* (79.7 \pm 10.02), *B. bassiana* (107.7 \pm 26.85) and control (207.4 \pm 49.90). Plants
64 treated with *B. bassiana* had 50% fewer leaves damaged by Coleoptera, while there were no effects on numbers
65 of whiteflies and thrips. Further, lower proportions of leaflets with symptoms of the foliar plant pathogenic
66 fungi *Mycosphaerella fragariae* and *Pestalotia longisetula* were observed in the *M. robertsii* (4.6% and 1.3%)
67 and *B. bassiana* (6.1% and 1.3%) treated plots compared to control plots (9.8% and 3.7%). No effect was seen
68 on numbers of naturally occurring predatory mites. Our results suggest that both isolates tested may be used
69 as root inoculants of strawberries to protect against foliar pests, particularly spider mites, and also against foliar
70 plant pathogenic fungi without harming naturally occurring and beneficial predatory mites.

71

72 **Keywords:** Endophytic entomopathogenic fungi; Microbial control; Plant-microbe interactions; *Tetranychus*
73 *urticae*; Integrated pest management (IPM).

74

75 **1. Introduction**

76

77 Strawberry is an important fruit throughout the world and in 2016 approximately 9.2 million tons of
78 fruits were produced worldwide, with a yield of 22.690 kg/ha (FAOSTAT 2018). Cultivated strawberry,
79 *Fragaria x ananassa* (Duch; Rosales: Rosaceae), is attacked by a large complex of arthropod pests and plant
80 diseases that may reduce the yield (Solomon et al. 2001). The twospotted spider mite, *Tetranychus urticae*
81 Koch (Acari: Tetranychidae), is an important pest of many crops throughout the world (Greco et al. 2005),
82 including strawberries (Raworth 1986; Easterbrook et al. 2001; Solomon et al. 2001). *Tetranychus urticae* feed
83 mainly on the underside of leaves and this feeding may lead to reduced photosynthesis and increased
84 transpiration as well as injection of phytotoxic substances when feeding on mesophyll and parenchyma plant
85 cells (Sances et al. 1979, 1982; Attia et al. 2013). The feeding damage therefore decreases foliar and floral
86 development causing reductions in quality and quantity of fruits (Rhodes et al. 2006).

87 Other important pest of strawberries worldwide includes the western flower thrips, *Frankliniella*
88 *occidentalis* Pergande (Thysanoptera: Thripidae) which causes damage by the feeding of nymphs and adults
89 resulting in flower abortion, fruit bronzing and malformation, and consequently yield loss (Solomon et al.
90 2001; Coll et al. 2007). Strawberries are also attacked by aphids of different species such as *Chaetosiphon*
91 *fragaefolli* Cockerell, *Aphis forbesi* Weed, *A. gossypii* Glover and *Mizus persicae* Sulzer (Hemiptera:
92 Aphididae) (Solomon et al. 2001; Bernardi et al. 2015; Dara 2016). The whitefly *Trialeurodes vaporariorum*
93 (Westwood) (Hemiptera: Aleyrodidae) is also a significant pest of strawberry crop in many regions (Solomon
94 et al. 2001; Bernardi et al. 2015; Dara 2016). Moreover, *Neopamera bilobata* Say (Hemiptera:
95 Rhyarochromidae) and the spotted wing fruit fly, *Drosophila suzukii* Matsumura (Diptera: Drosophilidae)
96 have recently invaded and caused economic losses in the production of many strawberry fields in Brazil (Kuhn
97 et al. 2014; Andreazza et al. 2016). High incidence of plant pathogens, especially fungal pathogens, is another
98 challenge faced by strawberry farmers in all producing countries and cause problems throughout the crop cycle,
99 from the newly planted seedlings to the final fruit producing stage (Garrido et al. 2011).

100 The main pest control strategy in strawberries throughout the world is the use of synthetic chemical
101 pesticides (Solomon et al. 2001; Garrido et al. 2011). Dependency of these chemicals for pest control in
102 strawberries is associated with undesirable effects on environment and human health (e.g. Attia et al. 2013;
103 Barzman et al. 2015; Czaja et al. 2015). Outbreaks of *T. urticae* are often observed following continuous
104 pesticide treatments (Klingen and Westrum 2007; Van Leeuwen et al. 2009, 2010) due to the emergence of
105 pest resistance to the particular pesticides and destruction of the pests' natural enemies (Solomon et al. 2001;
106 Sato et al. 2005). The use of invertebrate predators, parasitoids and microbial control agents in biological
107 control is considered a sustainable alternative to synthetic chemical pesticides for control of arthropod pests
108 (Garcia et al. 1988; Eilenberg et al. 2001). Except from the application of predatory phytoseiid mites to control
109 *T. urticae*, biological control is not widely used in strawberry production, and more development of macro-
110 and microbial control agents and application strategies is therefore necessary (Solomon et al. 2001; Attia et al.
111 2013).

112 Entomopathogenic fungi within the order Hypocreales are used in microbial control and many species
113 are known to have a quite wide host range (Goettel et al. 1990; Rehner 2005). The species *Beauveria bassiana*
114 (Balsamo-Crivelli) Vuillemin (Cordycipitaceae) and several species of *Metarhizium* (Clavicipitaceae) have
115 been considered promising microbial control agents in strawberries (Sabbahi et al. 2008; Castro et al. 2018)

116 and may be implemented in programs for integrated pest management (IPM) (Hajek and Delalibera 2010).
117 There are, however, constraints in the use of entomopathogenic fungi as microbial control agents, such as non-
118 consistent effects against pests, short survival time of the fungal propagules in the environment, quality of
119 commercial products, shelf life and costs (Lacey et al. 2015). These aspects are influenced by abiotic factors
120 such as temperature, light intensity and quality, humidity and rainfall (Meyling and Eilenberg 2007; Castro et
121 al. 2013) and by biotic factors such as multitrophic interactions with plants, invertebrates, other
122 microorganisms and plant pathogens (Klingen and Haukeland 2006; Meyling and Eilenberg 2007; Meyling
123 and Hajek 2010). In order to optimize pest control by entomopathogenic fungi, it is important to understand
124 how these factors and their interactions affect the efficacy of the microbial control agent in question.

125 Recent studies have reported that entomopathogenic fungi in the Hypocreales, mainly *Metarhizium*
126 spp. and *Beauveria* spp., may also interact with plants as endophytes (Vega 2008, 2018; Vega et al. 2009;
127 Greenfield et al. 2016). Endophytic fungi are able to colonize the internal tissues of a host plant and cause no
128 apparent negative effect on the plant (Carroll 1988; Stone et al. 2004; Vega 2008). This relationship between
129 entomopathogenic fungi and their host plant may protect the plant against arthropod pests and plant diseases
130 (Bing and Lewis 1991; Ownley et al. 2010; Jaber and Ownley 2018). Furthermore, endophytic fungi are
131 protected inside the plant tissues from the effect of ambient abiotic factors (Vega 2008, 2018) and the challenge
132 of short survival time of fungal propagule in the environment due to abiotic factors may therefore be reduced.
133 The mechanisms responsible for any plant protection capacity of plant associated entomopathogenic fungi
134 against arthropod pests and plant pathogens remains uncertain (Vidal and Jaber 2015; McKinnon et al. 2017).

135 Most of the published studies on entomopathogenic fungi as plant inoculants were carried out under
136 controlled experimental conditions, and so far, only few studies have investigated the pest control potential of
137 entomopathogenic fungi as inoculants of plants under field conditions while no field studies have evaluated
138 effects against plant pathogens (Jaber and Ownley 2018). Field studies have been carried out with inoculation
139 of common beans, *Phaseolus vulgaris* L. (Fabales: Fabaceae) with *B. bassiana* against *Liriomyza* leafminers
140 (Diptera: Agromyzidae) (Gathage et al. 2016); of *Sorghum bicolor* L. (Moench) (Poales: Poaceae) with *B.*
141 *bassiana*, *Metarhizium robertsii* Bisch., Rehner & Humber, and *Isaria fumosorosea* (Wize) Brown & Smith
142 (Cordycipitaceae) (Mantzoukas et al. 2015); and of cotton *Gossypium* spp. (Malvales: Malvaceae) with *B.*
143 *bassiana* against *Aphis gossypii* Glover (Homoptera: Aphididae) (Castillo-Lopez et al. 2014). These recent
144 field studies report of significant effects against foliar arthropod pests under field conditions suggesting that

145 implementation of entomopathogenic fungi as plant inoculants into outdoor IPM programs has a major
146 potential (Lacey et al. 2015; Jaber and Ownley 2018). Few field studies have been conducted on strawberry.
147 One study was conducted on soil drench granulate or root dipping application of Met52® *Metarhizium*
148 *brunneum* (reported as *M. anisopliae* (Metsch.) Sorokin) to strawberry against the soil living larvae of the
149 black vine weevil *Otiorhynchus sulcatus* in a temperate region (UK) and was suggested to be a potential strategy
150 (Ansari and Butt 2013). Further, the persistence of locally adapted isolates of *M. brunneum* Petch and
151 *Beauveria pseudobassiana* Rehner & Humber applied as granulates close to strawberry roots were confirmed
152 in studies in Norway (Klingen et al. 2015). However, none of these studies evaluated the potential of these
153 fungi for improving plant productivity or controlling pests above-ground in strawberry.

154 The aim of the present study was therefore to evaluate the potential of two selected isolates of
155 entomopathogenic fungi as root inoculants of strawberry plants for above-ground pest management under field
156 conditions in Brazil. The fungal species used were *M. robertsii* and *B. bassiana* and the origin of the isolates
157 were from Brazil. They were selected based on the ability to reduce *T. urticae* numbers on strawberry (F.
158 Canassa, unpubl.) and on common beans *P. vulgaris* (Canassa et al. 2019), in greenhouse experiments. The
159 effects on natural predatory mite populations were also assessed to evaluate the effect of the fungal inoculation
160 strategy on natural enemies of *T. urticae* in the strawberry foliage. Further, prevalence of insect pests and
161 important strawberry foliar pathogens were also monitored.

162

163 2. Material and Methods

164

165 2.1. Fungal isolates

166

167 Based on earlier efficacy studies (F. Canassa, unpubl.), two entomopathogenic fungal isolates *M.*
168 *robertsii* ESALQ 1622 and *B. bassiana* ESALQ 3375, identified to species level by molecular techniques
169 according to Rezende et al. (2015) and Rehner and Buckley (2005), were selected. Isolates were kept at - 80°C
170 in the entomopathogen collection "Prof. Sérgio Batista Alves" in the "Laboratory of Pathology and Microbial
171 Control of Insects" at Escola Superior de Agricultura "Luiz de Queiroz" at University of São Paulo
172 (ESALQ/USP), Piracicaba, São Paulo, Brazil. The *M. robertsii* ESALQ 1622 isolate originated from soil of a
173 corn field in Sinop City (11°51'47"S; 55°29'01"W), Mato Grosso State, Brazil and the *B. bassiana* ESALQ

174 3375 isolate was obtained from soil of a strawberry field in Senador Amaral City (22°33'12"S; 46°13'41"W),
175 Minas Gerais State, Brazil.

176

177 2.2. Experimental set up

178

179 The experiments were conducted in four different commercial strawberry fields (Fig. 1). The roots of
180 the strawberry seedlings were immersed in one of the following treatments before planting: A) *M. robertsii*
181 ESALQ 1622 in water + 0.05% Tween 80; B) *B. bassiana* ESALQ 3375 in water + 0.05% Tween 80; C) Water
182 + 0.05% Tween 80 (control). A randomized block design was used in all four field experiments.

183 Three experiments were conducted in Atibaia City, São Paulo State, Brazil, from March to September
184 2018 in three separate open commercial strawberry fields with black plastic mulching and drip irrigation (Open
185 field locations 1, 2, 3 are shown in Fig. 1). At all three locations, an experimental strawberry block was 60 m
186 long (20 m for each treatment), 1.1 m wide and contained 600 plants (200 plants for each treatment).
187 Experiments at location 1 (23°04'14.32''S; 46°40'58.2''W) and location 2 (23°04'33.5''S; 46°40'30.1''W)
188 had 6 blocks (=strawberry beds), where the three treatments A), B), C) were randomized inside each block,
189 totaling 3.600 plants, while at location 3 (23°08'00.7''S; 46°37'04.5''W) there were 4 blocks (=strawberry
190 beds), where the three treatments A), B), C) were also randomized inside each block, totaling 2.400 plants.
191 Strawberry cultivars of locations 1, 2 and 3 were Camarosa (University of California, 1993), Camino real
192 (University of California, 2001), and Oso grande (University of California, 1989), respectively. At these three
193 locations, bare root strawberry plants (*Fragaria x ananassa*) were planted at the 4 leaves stage in three rows
194 per bed with a distance of 0.27 cm between rows.

195 The experiment at location 4 was conducted in Senador Amaral City (22°33'12.1''S; 46°13'41.8''W),
196 Minas Gerais State, Brazil from July 2017 to January 2018, in low tunnels (short hoop structures covered with
197 white plastic), with black plastic mulching and drip irrigation (Tunnel -location 4 in Fig. 1). This field
198 experiment was established in 18 low tunnels representing four blocks, each with three strawberry beds of each
199 treatment, i.e. 12 strawberry beds per treatment. Each bed was 20 m long, 1.1 m wide and contained 250 plants,
200 totaling 3,000 plants per treatment. At location 4, bare root strawberry plants, cultivar Albion (University of
201 California, 2006) were planted at the 4 leaves stage individually in three rows with a distance of 0.27 cm
202 between rows.

203

204 **2.3. Preparation of fungal inoculum**

205

206 The two fungal isolates (*M. robertsii* ESALQ 1622 and *B. bassiana* ESALQ 3375) were retrieved from
207 the -80°C culture collection and plated onto Petri dishes (90 x 15 mm) containing 20 ml Potato Dextrose
208 Agar (PDA; Merck, Darmstadt, Germany). The cultures were then kept in darkness at 25°C for 10 days until
209 harvesting of conidia. This was done by adding 10 ml sterile 0.05% Tween 80 (Oxiteno, São Paulo, Brazil) to
210 the culture and scraping off the conidia with a sterile spatula. Conidial concentrations were estimated using a
211 Neubauer hemocytometer (Merck, Darmstadt, Germany) and adjusted to 1×10^8 conidia ml⁻¹. Later, 10 ml of
212 each suspension was inoculated with a pipette into individual polypropylene bags (35 cm length x 22 cm width)
213 containing 300 g autoclaved (121°C, 20 min) parboiled rice, inside an aseptic laminar flow chamber.

214 The fungal inoculated rice kernels were mixed in the plastic bags and incubated in darkness at 25°C
215 for 10 days. The bags were gently shaken every two days to ensure evenly distributed fungal growth on rice
216 kernels. Prior to use in the experiment, the conidial viability was checked by preparing a conidial suspension
217 by adding 1 g of rice with sporulating fungi from the plastic bag to 10 ml sterile 0.05% Tween 80. From the
218 third dilution, 150 µl of the conidial suspension were transferred with a pipette onto PDA. The percentage of
219 conidia germination was then evaluated according to Oliveira et al. (2015). Suspensions were only used if
220 germination rates were higher than 95%.

221

222 **2.4. Fungal inoculation of strawberry roots**

223

224 Rice kernels colonized with the two isolates (*M. robertsii* ESALQ 1622 and *B. bassiana* ESALQ 3375)
225 were added into water plus 0.05% Tween 80 as described below. For the Open Field experiments at locations
226 1, 2, 3, the original conidia concentration per gram of rice kernels for each isolate was estimated to 2.5×10^8
227 /g rice for *M. robertsii* and 1.3×10^9 /g rice for *B. bassiana*. The concentration was then adjusted to 1.5×10^{12}
228 conidia of *M. robertsii* on 3.0 kg rice and *B. bassiana* on 0.56 kg rice. The rice was mixed with 100 L of well
229 water plus 50 ml 0.05% Tween 80, resulting in 1.5×10^6 conidia/ml. The control consisted of 100 L of well
230 water plus 50 ml 0.05% Tween 80. The final suspensions for the experiments contained 1.5×10^6 conidia/ml.

231 For the Low Tunnel experiment at location 4, the original conidia concentration per gram of rice
232 kernels for each isolate was estimated to 1.8×10^8 /g rice for *M. robertsii* and 7.5×10^8 /g rice for *B. bassiana*.
233 The concentration was then adjusted to 1.5×10^{12} conidia of *M. robertsii* on 8.3 kg rice and *B. bassiana* on 2.0
234 kg rice. The rice was mixed with 750 L well water plus 375 ml 0.05% Tween 80, resulting in 2.0×10^6
235 conidia/ml. The control consisted of 750 L of well water plus 375 ml 0.05% Tween 80.

236 Strawberry roots were inoculated by immersing the root system of each plant completely into the
237 respective treatment suspensions for 2 min. The inoculated plants were transported to the correct position in
238 the rows inside plastic trays to avoid dripping suspension and then the plants were immediately planted into
239 the row. The suspensions were continuously mixed with a wooden stick during the strawberry root inoculation
240 to ensure homogenized concentrations.

241

242 **2.5. Evaluations: arthropod pests, natural enemies and plant pathogens**

243

244 All four field experiments were evaluated each 30 day for six months. However, the results obtained
245 at location 4 (Low Tunnel experiment) are only reported up to 120 days after inoculation, because the producer
246 applied a synthetic chemical pesticide at this time, which may have influenced the following observations at
247 150 and 180 days after inoculation.

248 In the Open Field experiments at locations 1, 2, and 3, we observed 15 leaflets (= one leaf from a
249 triplet) and 15 flowers representing 15 plants in each of the central rows of the strawberry beds as indicated in
250 Fig. 1. In the Low Tunnel experiment at location 4, we observed 15 leaflets (= one leaf from a triplet) and 15
251 flowers from six plants (i.e. 2 or -3 leaflets per plant) in each of the central rows per strawberry bed as indicated
252 in Fig. 1. Each leaflet was destructively sampled by hand and visually observed, and the arthropod pests were
253 identified to species level and counted in the field.

254 The predatory mites were transferred to plastic vials (500 ml, 8.5 cm high, 10 cm diameter) containing
255 70% ethanol and taken to the laboratory for identification by observing each specimen under microscope. Each
256 predatory mite was collected with a fine brush from the vial with 70% ethanol and mounted in Hoyer's medium
257 for identification to species by comparing their morphology with information from original descriptions and
258 redescriptions provided in Rowell et al. (1978), Chant and Yoshida-Shaul (1991), Moraes et al. (2004) and
259 Tixier et al. (2008).

260 Leaflets with characteristic symptoms of the plant pathogenic fungi *Mycosphaerella fragariae* Tul.
261 (Lindau), *Dendrophoma obscurans* (Ell & Ev.) and *Pestalotia longisetula* (Guba) were recorded and the
262 percentage of leaflets with the diseases was calculated.

263

264 **2.6. Evaluation of colonization of strawberry leaves and soil**

265

266 Sampling of strawberry leaves and soil adjacent to plant roots was done 180 days after inoculation to
267 evaluate the presence of entomopathogenic fungi. One strawberry leaf (= three leaflets) was randomly and
268 destructively collected from one plant per plot in the center row of each replicate plot treatment at each of the
269 four locations. Collected leaves were placed in separate plastic bags and transferred to the laboratory for
270 evaluation of endophytic colonization. The leaves were cut in sections of 4 cm x 1 cm, and they were then
271 surface sterilized by following the method described by Greenfield et al. (2016). Three sections of leaves were
272 plated on one Petri dish (90 x 15 mm) with the following selective media: 20 ml of PDA, 0.5 g.L⁻¹ of
273 cycloheximide, 0.2 g.L⁻¹ of chloramphenicol, 0.5 g.L⁻¹ of Dodine (65%) and 0.01 g.L⁻¹ of Crystal Violet (Behie
274 et al. 2015). The sterilization efficiency was confirmed by plating 100 µl of the last rinsing water of the
275 sterilization onto PDA (Parsa et al. 2013). Further, imprints of sterilized leaves were used as an additional
276 method to confirm whether the sterilization was successful. This was done by gently pressing the leaf section
277 with the cut edge onto the PDA medium (Greenfield et al. 2016) before placing sections in selective media
278 plates. The Petri dishes were incubated at 25°C for 15 days before visually observed for fungal outgrowth of
279 *Metarhizium* or *Beauveria* on each plant fragment. The frequency of occurrence was estimated as the number
280 of plant fragments with entomopathogenic fungi present in relation to the total number of plant fragments.

281 Soil samples adjacent to plant roots were collected with a garden spade, from the same plants where
282 leaves were sampled, without removing the plants. Then soil with roots were placed into individual plastic
283 bags and brought back to the laboratory. Here, the soil was mixed, and subsequently 1 g was sampled and
284 added to 10 ml of sterile 0.05% Tween 80, and vigorously vortexed for 30 s and serially diluted into distilled
285 water + 0.05% Tween 80 to obtain the following concentrations: 1x10⁰, 1x10⁻¹, 1x10⁻² and 1x10⁻³. Petri dishes
286 (90 x 15 mm) containing selective agar medium as described above were divided into four equal quarter
287 sections by marking the bottom part of the Petri dishes with a permanent marker. Then 100 µl from each soil
288 dilution suspension was pipetted onto the selective media in each of the four sections. After the 100 µl was

289 dried up inside a laminar flow chamber, the Petri dishes were incubated in darkness at 25°C for 15 days, and
290 the presence of *Metarhizium* or *Beauveria* was detected according to fungal growth morphology in each plate.
291 The frequency of occurrence was estimated as the number of soil samples with entomopathogenic fungi in
292 relation to the total number of samples.

293

294 **2.7. Statistical Analysis**

295

296 We fitted Poisson generalized linear mixed models to the *T. urticae* counts obtained from locations 1,
297 2 and 3 (Open Field), including in the linear predictor the effects of block and different quadratic polynomials
298 per each treatment and location combination over time (natural log-transformed) as fixed effects, and two
299 random effects, namely, the effect of bed (since observations taken over time on the same bed are correlated)
300 and an observation-level random effect to model overdispersion. Hence, the maximal model included 32 fixed
301 effects and 2 variance components, totaling 34 parameters. We then performed backwards selection, using
302 likelihood-ratio (LR) tests to assess the significance of the fixed effects. Treatments were compared by fitting
303 nested models using grouped treatment levels and comparing them using LR tests; a significant test statistic
304 means that the treatments cannot be grouped, as they are statistically different (see e.g. Faretto et al. 2018).
305 After model selection, the effects of proportion of occurrence of each plant pathogen species present (*M.*
306 *fragariae*; *P. longisetula*; and *D. obscurans*), damage by Coleoptera (holes in the leaflets most likely caused
307 by *Colaspis* spp.), and number of thrips (*F. occidentalis*) were added, separately, as covariates in the model
308 and their significance assessed using LR tests.

309 For the other variables observed in locations 1, 2 and 3 (Open Field), we worked with the aggregated
310 values across all time points. The proportion of leaflets infected by plant pathogens present (*M. fragariae*, *P.*
311 *longisetula* or *D. obscurans*) and the proportion of leaflets damaged by Coleoptera were analysed by fitting
312 quasi-binomial models with a logit link, including the effects of block, treatment, location, and the interaction
313 between treatment and location in the linear predictor. The number of thrips was analysed by fitting quasi-
314 Poisson models, also including the effects of block, treatment, location, and the interaction between treatment
315 and location in the linear predictor. Significance of effects was assessed using F-tests, since the dispersion
316 parameter was estimated (Demétrio et al. 2014). Multiple comparisons were performed by obtaining the 95%
317 confidence intervals for the linear predictors.

318 For location 4 (Low Tunnel), Poisson generalized linear mixed models were fitted to the *T. urticae*
319 counts, including in the linear predictor the effects of block and different intercepts and slopes per each
320 treatment over time as fixed effects, and two random effects, namely, the effect of bed (since observations
321 taken over time on the same bed are correlated) and an observation-level random effect to model
322 overdispersion. Here, the maximal model included 9 fixed effects and 2 variance components, totaling 11
323 parameters. As for the models fitted for locations 1, 2, and 3 (Open Field), we then performed backwards
324 selection, using likelihood-ratio (LR) tests to assess the significance of the fixed effects. Treatments were
325 compared the same way, by fitting nested models using grouped treatment levels and comparing them using
326 LR tests. Again, after model selection, the effects of proportion of occurrence of number of pests present and
327 plant pathogens were added, individually, as covariates in the model and their significance assessed using LR
328 tests.

329 For the other variables observed at location 4 (Low Tunnel), we worked with the aggregated values
330 across all time points. The proportion of leaflets infected by plant pathogens was analysed by fitting quasi-
331 binomial models with a logit link, including the effects of block and treatment in the linear predictor. The
332 number of cucurbit beetles, white flies, thrips, and predatory mites were analysed by fitting quasi-Poisson
333 models, also including the effects of block and treatment in the linear predictor. Significance of effects was
334 assessed using F-tests, and multiple comparisons were performed by obtaining the 95% confidence intervals
335 for the linear predictors.

336 All analyses were carried out in R (R Core Team 2018). Goodness-of-fit was assessed using half-
337 normal plots with a simulated envelope, using package hnp (Moral et al. 2017). Generalized linear mixed
338 models were fitted using package lme4 (Bates et al. 2015). All plots were generated using package ggplot2
339 (Wickham 2009).

340

341 **3. Results**

342

343 **3.1. Effects of *M. robertsii* and *B. bassiana* on *T. urticae***

344

345 Root inoculation of strawberry plants with the two fungal treatments (*M. robertsii* ESALQ 1622 and
346 *B. bassiana* ESALQ 3375) significantly influenced the number of *T. urticae* adults over the six-month period

347 (180 days) in Open Field locations 1, 2 and 3 (LR = 30.31, d.f. = 2, $p < 0.0001$) (Fig. 2) and the Low Tunnel
348 location 4 (LR = 10.39, d.f. = 2, $p = 0.0055$) (Fig. 3). No difference between plants inoculated with the two
349 entomopathogenic fungi were seen in locations 1, 2 and 3 (LR = 0.07, d.f. = 1, $p = 0.3092$) nor in location 4
350 (LR = 0.02, d.f. = 1, $p = 0.8793$).

351 There was no significant three-way interaction among Open Field locations (1, 2 and 3), treatment,
352 and time (LR = 4.06, d.f. = 8, $p = 0.8516$), nor significant two-way interactions between Open Field locations
353 (1, 2 and 3) and treatment (LR = 0.69, d.f. = 4, $p = 0.9524$) and between treatment and time (LR = 3.00, d.f. =
354 4, $p = 0.5574$). However, there was a significant interaction between location and time (LR = 49.91, d.f. = 4,
355 $p < 0.0001$), which means that the population dynamics of spider mites changed differently between the
356 inoculated and control plants over time at each location, with a significantly higher number of adults on the
357 control plants in the three locations (LR = 30.31, d.f. = 2, $p < 0.0001$) (Fig. 2). For the Low Tunnel location 4,
358 there was no significant interaction between treatment and time (LR = 2.49, d.f. = 2, $p = 0.2879$), however,
359 there were significant effects of time (LR = 43.02, d.f. = 1, $p < 0.0001$) and treatment (LR = 10.39, d.f. = 2, p
360 $= 0.0055$), and hence there was a significantly higher number of *T. urticae* adults on the control plants at
361 different times of evaluation, when compared to the two fungal treatments (Fig. 3).

362 There was no significant effect of the proportion of leaflets infected by the plant pathogens *M.*
363 *fragariae* (LR = 0.20, d.f. = 1, $p = 0.6569$), *P. longisetula* (LR = 1.89, d.f. = 1, $p = 0.1693$) and *D. obscurans*
364 (LR = 1.90, d.f. = 1, $p = 0.1686$) on the number of *T. urticae* in Open Field locations 1, 2 and 3. However,
365 there was a significant effect of the proportion of leaves damaged by Coleoptera (holes in the leaflets most
366 likely caused by *Colaspis* spp.) on the number of *T. urticae* (LR = 5.13, d.f. = 1, $p = 0.0235$), suggesting that
367 numbers of *T. urticae* were lower on leaflets damaged by Coleoptera (estimate of -1.60 in the logit scale, with
368 an associated standard error of 0.72, indicating a negative relationship). Besides, in locations 1, 2, 3 there was
369 no significant interaction between numbers of *T. urticae* and thrips in flowers (LR = 1.03, d.f. = 1, $p = 0.3092$).
370 In Low Tunnel location 4, there was no significant interaction between numbers of *T. urticae* and thrips in
371 flowers (LR = 0.73, d.f. = 1, $p = 0.3929$) or whiteflies (LR = 3.74 d.f. = 1, $p = 0.0532$).

372

373 **3.2. Effects of *M. robertsii* and *B. bassiana* on other pests and diseases**

374

375 Damage caused by Coleoptera (holes in the leaflets) was significantly reduced on strawberry plants
376 inoculated with *B. bassiana* ESALQ 3375 compared to control plants in Open Field locations 1, 2 and 3 (Table
377 1). There was no significant interaction between location and treatment ($F_{4,34} = 1.68$, $p = 0.1767$), but there
378 was a significant effect of location ($F_{2,40} = 12.61$, $p < 0.0001$). The mean damage caused by Coleoptera (\pm
379 SE%) in each location were: location 1 = 10.68 ± 1.57 a; location 2 = 3.89 ± 0.84 b; and location 3 = $4.54 \pm$
380 1.15 b.

381 There was no difference in the number of thrips in flowers between fungal inoculated strawberry plants
382 and the control plants in Open Field locations 1, 2 and 3 (Table 1). There was no significant interaction between
383 location and treatment ($F_{4,34} = 0.47$, $p = 0.7651$), but there was a significant effect of location ($F_{2,40} = 11.98$, p
384 $= 0.0001$). The mean \pm SE (%) in each location were: location 1 = 27.59 ± 4.28 b; location 2 = 14.26 ± 2.23 c;
385 and location 3 = 40.09 ± 6.78 a.

386 Although there was no difference in the proportion of leaflets (n=15 leaflets per replicate) with
387 symptoms of the plant pathogenic fungus *D. obscurans* in Open Field locations 1, 2 and 3 ($F_{2,38} = 1.02$, $p =$
388 0.3710), the proportion of leaflets (n=15 leaflets per replicate) with symptoms of *M. fragariae* and *P.*
389 *longisetula* were significantly smaller on plants inoculated with *M. robertsii* ESALQ 1622 and *B. bassiana*
390 ESALQ 3375 in all fields (Table 1). Besides, for *D. obscurans*, there was no significant interaction between
391 location and treatment ($F_{4,34} = 0.79$, $p = 0.5386$), and among the three Open Field locations ($F_{2,40} = 1.54$, $p =$
392 0.2300). For *P. longisetula*, there was also no significant interaction between location and treatment ($F_{4,34} =$
393 0.58 , $p = 0.5676$), and among the three Open Field locations ($F_{2,40} = 0.04$, $p = 0.8433$). Regarding the disease
394 caused by *M. fragariae*, there was no significant interaction between location and treatment ($F_{4,34} = 0.46$, $p =$
395 0.7640), but there was a significant effect of location ($F_{2,40} = 39.84$, $p < 0.0001$). The mean \pm SE (%) in each
396 location were: location 1 = 3.83 ± 1.06 ; location 2 = 14.20 ± 1.90 ; and location 3 = 0.56 ± 0.29 .

397 In Low Tunnel location 4, in addition to *T. urticae*, the other major pests were whiteflies and thrips in
398 flowers, but there was no difference in the number of any of these among the three treatments (Table 2). In
399 this location, the density of pest was always very low and very few leaves with symptoms of plant pathogens
400 were observed. The cumulative proportion of leaflets with symptoms of all the diseases (*D. obscurans* + *P.*
401 *longisetula* + *M. fragariae*) can be viewed in Table 2.

402

403 3.3. Effects of *M. robertsii* and *B. bassiana* on predatory mites

404

405 At Open Field locations 1, 2 and 3, few arthropod natural enemies were observed, but at Low Tunnel
406 location 4 there were many predatory mites, mainly of the species *Neoseiulus californicus* McGregor (Acari:
407 Phytoseiidae). The numbers of these predatory mites at location 4 were not significantly different on plants
408 inoculated with *M. robertsii* and *B. bassiana*, compared to the control ($F_{2,30} = 0.04$, $p = 0.9642$). The mean \pm
409 SE (%) for the three treatments at location 4 were: *M. robertsii* = 14.3 ± 3.83 ; *B. bassiana* = 14.8 ± 3.06 ; and
410 control = 13.6 ± 2.57 predatory mites per leaflet accumulated for all sampling dates.

411

412 3.4. Colonization of *M. robertsii* and *B. bassiana* in strawberry leaves and soil

413

414 Low colonization levels of the plants by both *Metarhizium* spp. and *Beauveria* spp. were observed 180
415 days after inoculation of strawberry roots. At Open Field location 1, neither *Metarhizium* spp. nor *Beauveria*
416 spp. were recovered on selective media from leaf samples, but *Metarhizium* spp. was found in all soil samples
417 while *Beauveria* spp. was not recovered from soil. From samples collected at Open Field location 2, 33.3% (2
418 out of 6) of leaf sections and 16.7% (1 out of 6) of soil samples were found to harbor *Beauveria* spp., while
419 *Metarhizium* spp. was recovered from 16.7% (1 out of 6) of the soil samples but not from the leaves. At Open
420 Field location 3, *Beauveria* spp. was recovered from 25% (1 out of 4) of leaves and soil samples while
421 *Metarhizium* spp. was found in 75% (3 out of 4) of the soil samples and not in leaves. At Low Tunnel location
422 4, *Beauveria* spp. was recovered from 41.7% (5 out of 12) of leaf samples and from 8.3% (1 out of 12) of soil
423 samples. At this location *Metarhizium* spp. was not recovered from the leaves, but the recovery from soil
424 samples was 75% (9 out of 12). None of the leaf or samples from the control plots were found to contain any
425 of the target fungi at any of the four locations.

426

427 4. Discussion

428

429 Our field experiment, replicated at four locations, show that root inoculations of strawberry plants with
430 *M. robertsii* ESALQ 1622 and *B. bassiana* ESALQ 3375 resulted in lower numbers of *T. urticae* adults
431 compared to non-inoculated control plants. Few studies have investigated the potential of plant inoculated
432 entomopathogenic fungi as microbial control agents under natural field conditions (reviewed by Jaber and

433 Ownley 2018; Vega 2018) and the present study is the first report of the effect on *T. urticae* numbers on
434 strawberry plants inoculated with *M. robertsii* and *B. bassiana* evaluated in the field under commercial
435 cultivation regimes. The two fungal isolates were previously found to reduce *T. urticae* populations on bean
436 *P. vulgaris* (Canassa et al. 2019) and since our strawberry field study show a similar effect this may suggest
437 that these isolates may be used as root inoculants of other crops to control *T. urticae*. Further, predatory mite
438 populations were not negatively affected by strawberry plants inoculated with *M. robertsii* ESALQ 1622 and
439 *B. bassiana* ESALQ 3375 indicating that adverse non-target effects on arthropod natural enemies may be
440 limited or non-existing.

441 The potential of *B. bassiana* as an endophyte for pest management has been reported in field studies
442 with other crops. For example, Gathage et al. (2016) reported lower infestation levels of *Liriomyza* leafminers
443 in bean leaves (*P. vulgaris*) in a bean field experiment in Kenya where bean seeds had been inoculated with *B.*
444 *bassiana* G1LU3 and *Hypocrea lixii* Patouillard (syn. *Trichoderma lixii*) F3ST1. Further, Castillo-Lopez et al.
445 (2014) reported lower numbers of *A. gossypii* on cotton plants grown in the field in Texas, USA, from seeds
446 inoculated with the commercial product Botanigard® (BioWorks Inc, Victor, NY) based on the GHA strain of
447 *B. bassiana*. Our field experiments also suggest that strawberry plants inoculated with *M. robertsii*
448 ESALQ 1622 and *B. bassiana* ESALQ 3375 reduced the proportion of leaf damage caused by Coleopteran
449 pests, while no effects on other pest damage, such as whiteflies or thrips in flowers, were observed. Mantzoukas
450 et al. (2015) reported from field studies of *Sorghum bicolor* that *B. bassiana* and *M. robertsii* suppressed
451 tunneling *Sesamia nonagrioides* Lefébvre (Lepidoptera: Noctuidae) larvae by 60% and 87%, and increased
452 larval mortality by 80% and 100%, respectively, compared to control plants after spray inoculations of plants.

453 We also recorded a reduction in the prevalence of the foliar plant pathogenic fungi *M. fragariae* and
454 *P. longisetula* in strawberry plants inoculated with *B. bassiana* ESALQ 3375 or *M. robertsii* ESALQ 1622.
455 According to Jaber and Alananbeh (2018), only few studies have been conducted on the effects of plant
456 inoculated entomopathogenic fungi affecting plant pathogens and so far no field studies have been carried out.
457 Jaber and Alananbeh (2018) reported, however, that sweet pepper *Capsicum annum* L. (Solanaceae)
458 endophytically colonized with *B. bassiana* (NATURALIS) and *M. brunneum* (BIPESCO5) showed
459 significantly reduced incidence and severity of three *Fusarium* species (*F. oxysporum*, *F. culmorum*, and *F.*
460 *moniliforme*) using *in planta* bioassays in controlled greenhouse settings with sterile soil. So far, *B. bassiana*
461 is the most studied entomopathogenic fungal species against plant pathogens and it has been reported to protect

462 tomato and cotton seedlings against the plant pathogens *Rhizoctonia solani* and *Pythium myriotylum* (Ownley
463 et al. 2008). Furthermore, Sasan and Bidochka (2013) reported a 59.4% inhibition of *Fusarium solani* f. sp.
464 *phaseoli* in bean, when co-cultured in pretreated sterile potting mixture with *M. robertsii*. In another study, the
465 co-inoculation of wheat seeds with *Metarhizium brunneum* Petch and the mycoparasitic fungus *Clonostachys*
466 *rosea* (Link) Schroers et al. (Hypocreales: Bionectriaceae) resulted in infections by *M. brunneum* in root-
467 feeding Coleopteran larvae and provided protection against the plant pathogen *F. culmorum* (Keyser et al.
468 2016), but *M. brunneum* did not affect the plant pathogen individually. The present strawberry field study
469 suggests that the tested isolates of *B. bassiana* and *M. robertsii* can provide long-term protection of
470 strawberries against both arthropod pests and foliar pathogens using a single root application at the time of
471 planting.

472 Our data also suggest that natural populations of predatory mites, most of them identified as *N.*
473 *californicus*, remained unaffected on strawberry plant inoculated with *M. robertsii* ESALQ 1622 or *B.*
474 *bassiana* ESALQ 3375. The field experiments therefore indicate a limited non-target effect on arthropod
475 natural enemies when the fungi are applied as root inoculants. Few studies have investigated the effects of
476 plant associated entomopathogenic fungi on arthropod natural enemies and mostly focus have been on effects
477 on parasitoids (Bixby-Brosi and Potter 2012; Akutse et al. 2014; Jaber and Araj 2018). One of the few studies
478 reporting on effects of plant-fungi interactions on predatory mites was by Schausberger et al. (2012), who
479 showed that bean (*P. vulgaris*) colonized by the mycorrhizal fungus *Glomus mosseae* and infested with *T.*
480 *urticae*, changed the composition of herbivore induced plant volatiles. This caused the fungal inoculated plants
481 to become more attractive to the predatory mites, *Phytoseiulus persimilis* Athias-Henriot (Acari: Phytoseiidae),
482 than non-mycorrhizal plants. It was suggested that the predatory mites associated the plant response with
483 presence of prey (Patiño-Ruiz and Schausberger 2014), and hence showed a higher oviposition rate on these
484 plants resulting in more efficient *T. urticae* suppression (Hoffmann et al. 2011). Canassa et al. (2019) reported
485 in short-term leaf disc experiments that *P. persimilis* showed no difference in the predation rate on spider mites
486 from inoculated plants with *B. bassiana* (ESALQ 3375) and *M. robertsii* (ESALQ 1622) compared to control
487 plants. The use of *B. bassiana* (NATURALIS) and *M. brunneum* (BIPESCO5) as inoculants in sweet pepper
488 combined with the aphid endoparasitoid *Aphidius colemani* Viereck (Hymenoptera: Braconidae) also indicated
489 compatibility in control of *Myzus persicae* Sulzer (Homoptera: Aphididae) (in a greenhouse study (Jaber and
490 Araj 2018). In another recent study, González-Mas et al. (2019) reported that the numbers of *A. gossypii*

491 parasitized by *A. colemani* was not influenced by whether the aphids had been feeding on seed inoculated
492 melon plants with *B. bassiana* (isolate EABb 01/33-Su) or not. Further, application of *B. bassiana* on melon
493 leaves did not influence the number of aphids consumed by larvae of the lacewing, *Chrysoperla carnea*
494 Stephens (Neuroptera: Chrysopidae), and *C. carnea* showed preference to feed on aphids reared on inoculated
495 rather than control plants in a choice bioassay (González-Mas et al. 2019). All these findings indicate that plant
496 inoculated entomopathogenic fungi may be used in combination with parasitoids and predators to enhance the
497 biocontrol efficacy of several plant pests in different crops.

498 In our study we were able to recover *Metarhizium* and *Beauveria* from strawberry leaves and soil
499 adjacent to the roots at the end of the experiment and cropping cycle, meaning 180 days (for location 1,2,3)
500 and 120 days (for location 4). The main aim of the present study was not to evaluate in-depth the dynamics of
501 endophytism of the inoculated fungal isolates using a close-to-practice inoculation method in strawberry
502 production systems and the use of commercial farm settings did not allow for repeated and complete
503 destructive sampling of plant material. However, Castro et al. (2016) have previously reported the persistence
504 in strawberry soil and rhizospheres in Brazil of the isolates *M. anisopliae* (ESALQ1037) and *M. robertsii*
505 (ESALQ1426) for up to 12 months after soil drench application. Further, Klingen et al. (2015) report that two
506 Norwegian isolates, one *B. pseudobassiana* and one *M. brunneum*, and an Austrian isolate of *M. brunneum*
507 had long-term persistence (>1 year) in bulk soil and rhizosphere soil of strawberries in a semi-field experiment
508 in Norway. It has previously been reported that *B. bassiana* is a more extensive colonizer of foliar tissues than
509 *Metarhizium* spp., when seed inoculations were used, while *Metarhizium* spp. have been reported as almost
510 exclusively colonizing the rhizosphere of various plant species (Ownley et al. 2008; Quesada-Moraga et al.
511 2009; Akello and Sikora 2012; Akutse et al. 2013; Behie et al. 2015), and similar results have been observed
512 in our study. Although the observed effects of the inoculation on herbivore densities were consistent,
513 endophytic colonization was not consistently detected in strawberry plants in our study. It has been previously
514 reported that endophytic establishment may be influenced by several variables, such as host plant, fungal strain,
515 environmental conditions, substrate and soil (Sánchez-Rodríguez et al. 2018). Moreover, previous research
516 has showed that the establishment of entomopathogenic fungi within plant tissues may be transient (Garrido-
517 Jurado et al. 2017) and the establishment success of fungal isolates is significantly reduced when inoculations
518 are performed in natural soils (Parsa et al. 2018), as was the case in the present study. It should therefore be

519 expected that end-point measurements of endophytic colonization will be limited in field studies, particularly
520 over the 6-month time period.

521 Given that negative effects were broadly observed against both *T. urticae* and selected plant pathogens
522 in the foliage after the single inoculation events of strawberry roots with isolates of either *B. bassiana* or *M.*
523 *robertsii*, and considering the inconsistent re-isolation of fungi from leaf samples; it seems most likely that
524 plant induced defenses were responsible for the reductions, but this will require further studies to elucidate and
525 conclude. It has been widely suggested that the mechanisms used by entomopathogenic fungi as plant
526 associates and endophytes to antagonize plant pests or pathogens may result through the production of
527 secondary metabolites by the associated fungus (Vidal and Jaber 2015; Yan et al. 2015; McKinnon et al. 2017;
528 Jaber and Alananbeh 2018). Alternatively, another mechanism could be through induced systemic defense
529 mechanisms of the inoculated plants, because the endophyte can be first recognized as a potential invader,
530 which leads the plants to trigger its immune responses and consequently synthesize specific regulatory
531 elements that may affect the arthropod pests and plant pathogen (Brotman et al. 2013; McKinnon et al. 2017).

532 In conclusion, the present study demonstrates that entomopathogenic fungi can be applied as root
533 inoculants in commercial strawberry fields to simultaneously control important arthropod pests, particularly
534 *T. urticae*, and plant pathogenic fungi. There were no indications that the inoculations of strawberry plant with
535 the entomopathogenic fungal isolates tested had negative non-target effects on naturally occurring predatory
536 mites, particularly *N. californicus*. Hence, inoculation of strawberry plants with entomopathogenic fungi
537 through root dipping may be used in combination with predatory mites for the control of *T. urticae*. This may
538 represent a new tool and an innovative biological control strategy that could be implemented in IPM and
539 organic strawberry production.

540

541 **Author contribution**

542

543 FC, IDJ, IK and NVM conceived and designed research. FC and FCNE conducted experiments. RAM
544 analysed data, prepared figures and wrote the statistical analysis section. FC, IK, IDJ and NV wrote the
545 manuscript. All authors reviewed and approved the manuscript.

546

547 **Compliance with Ethical Standards**

548

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

550

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789 **Tables**

790

791 **Table 1.** Means \pm SE of proportion of leaflets damaged by Coleoptera (%), cumulative number of
 792 thrips in flowers, and proportion of leaflets with symptoms of the pathogens *D. obscurans*, *P.*
 793 *longisetula* and *M. fragariae* (%) representing the differences in the Open Field locations 1, 2 and 3,
 794 with summaries of generalized linear models below. Separate analyses were performed for each
 795 response variable.

Treatments ^b	Assessment ^a				
	Locations 1, 2, 3				
	Coleoptera damage	N ^o of thrips	<i>D. obscurans</i>	<i>P. longisetula</i>	<i>M. fragariae</i>
<i>B. bassiana</i>	4.4 \pm 0.88 b	24.5 \pm 4.67 a	2.7 \pm 1.23 a	1.3 \pm 0.37 b	6.1 \pm 1.66 b
<i>M. robertsii</i>	6.6 \pm 1.15 ab	21.6 \pm 3.34 a	2.5 \pm 1.10 a	1.3 \pm 0.48 b	4.6 \pm 1.35 b
H ₂ O + Tween 80	8.7 \pm 2.02 a	30.9 \pm 6.27 a	4.5 \pm 1.58 a	3.7 \pm 1.24 a	9.8 \pm 2.69 a
Test statistic	F _{2,38} = 4.17	F _{2,38} = 1.97	F _{2,38} = 1.02	F _{2,38} = 4.92	F _{2,38} = 5.84
p-value	p = 0.0240	p = 0.1549	p = 0.3710	p = 0.0158	p = 0.0066

796 ^aData (mean \pm SE) followed by different letters within a column are significantly different (GLM,
 797 followed by *post hoc* Tukey test, *P* < 0.05).

798 ^bTreatments included root inoculations of the entomopathogenic fungal isolates *Beauveria bassiana*
 799 ESALQ 3375 (*B. bassiana*), *Metarhizium robertsii* ESALQ 1622 (*M. robertsii*), and control treatment
 800 with H₂O + 0.05% Tween 80.

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803 **Table 2.** Means \pm SE of cumulative number of whiteflies per leaflet and thrips per flower, and the
 804 mean \pm SE proportion of leaflets with symptoms of foliar pathogens (combined % incidence of *D.*
 805 *obscurans* + *P. longisetula* + *M. fragariae*) in the Low Tunnel location 4. Summaries of separate
 806 statistical analyses for each response variable using generalized linear models are presented below.

Treatments ^b	Assessment ^a		
	Whiteflies	N° of thrips	Diseases
<i>B. bassiana</i>	6.6 \pm 1.70 a	1.9 \pm 5.33 a	0.5 \pm 0.31 a
<i>M. robertsii</i>	6.0 \pm 1.54 a	1.6 \pm 3.70 a	0.5 \pm 0.31 a
H ₂ O + Tween 80	5.9 \pm 1.38 a	1.8 \pm 2.91 a	1.2 \pm 0.42 a
Test statistic	F _{2,30} = 0.07	F _{2,30} = 0.18	F _{2,30} = 0.95
p-value	p = 0.9359	p = 0.8358	p = 0.3988

807 ^aData (mean \pm SE) followed by different letters within a column are significantly different (GLM,
 808 followed by *post hoc* Tukey test, $P < 0.05$).

809 ^bTreatments included root inoculations of the entomopathogenic fungal isolates *Beauveria bassiana*
 810 ESALQ 3375 (*B. bassiana*), *Metarhizium robertsii* ESALQ 1622 (*M. robertsii*), and control treatment
 811 with H₂O + 0.05% Tween 80.

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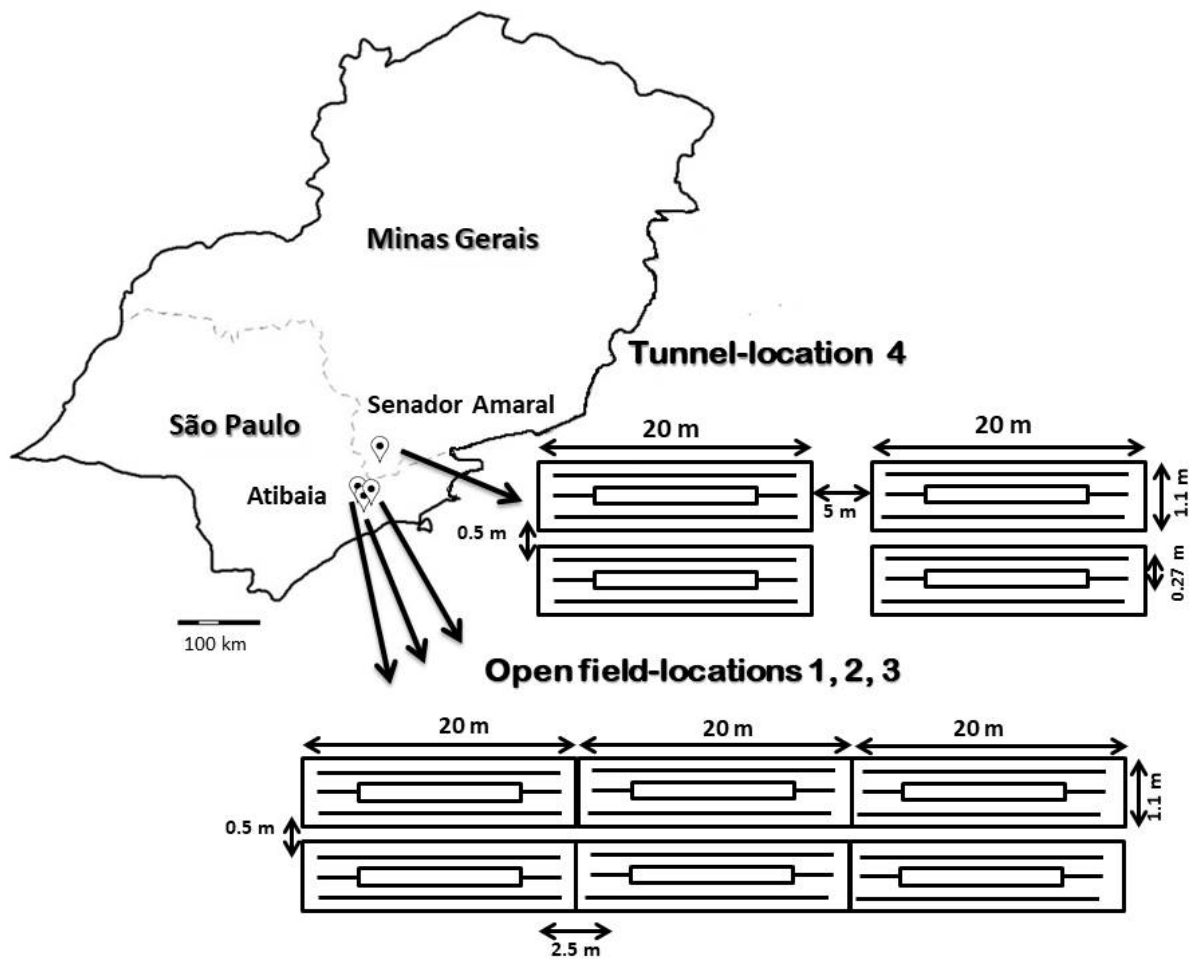
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825 **Fig. 1** Experimental field set up in Open Field locations 1, 2 and 3 in Atibaia (1: 23°04'14.32"S 46°40'58.2"W,
 826 2: 23°04'33.5"S 46°40'30.1"W, 3: 23°08'00.7 "S 46°37'04.5"W) and in Low Tunnel location 4 in Senador
 827 Amaral (22°33'12.1"S 46°13'41.8"W). Rows and area used for recording of data are indicated as a rectangle
 828 inside each bed

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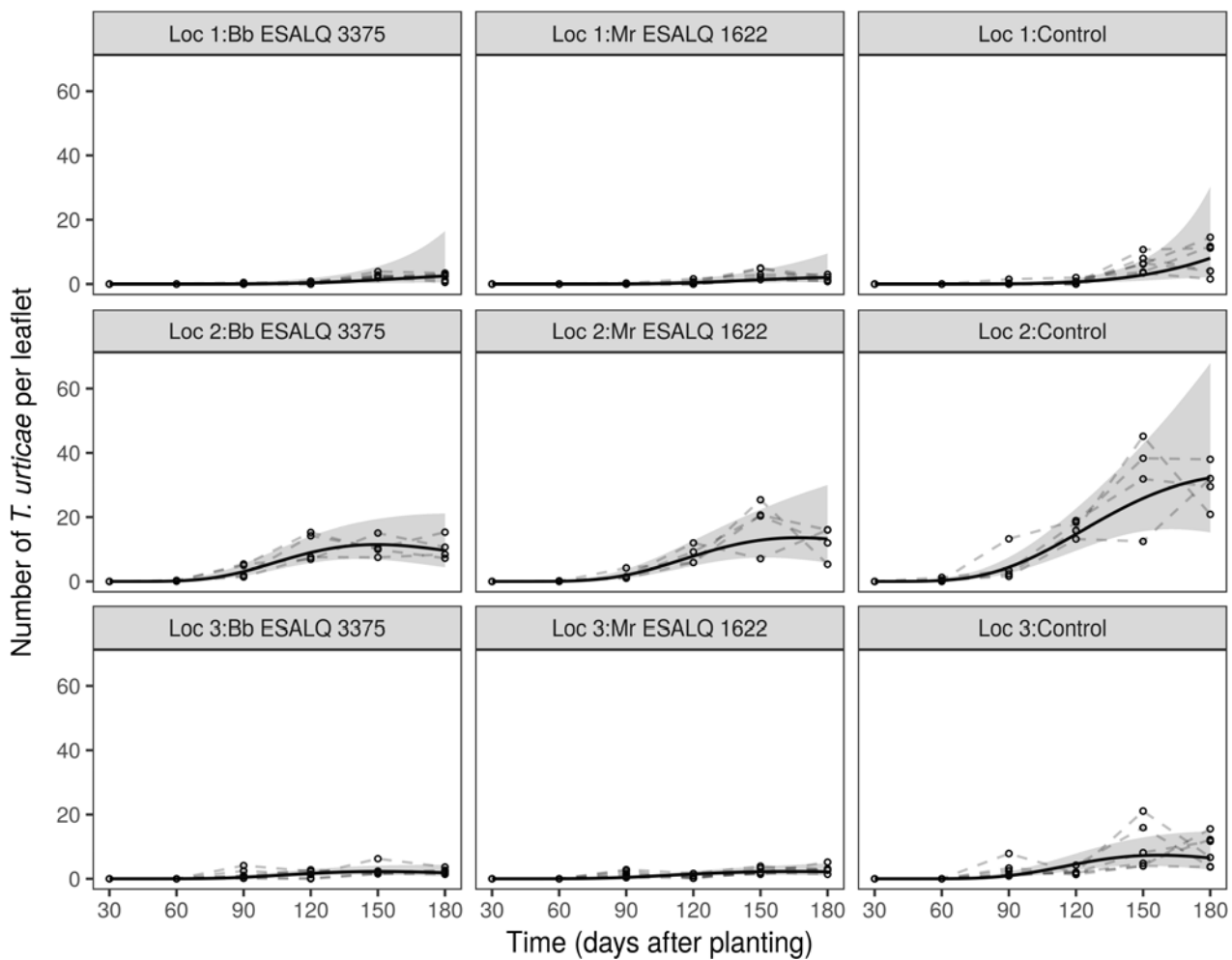
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840 **Fig. 2** Effect of inoculation of strawberry root with *Beauveria bassiana* (Bb) isolate ESALQ 3375 or
 841 *Metarhizium robertsii* (Mr) ESALQ 1622 on numbers of adult *Tetranychus urticae* per leaflet 30, 60, 90, 120,
 842 150 and 180 days after inoculation, at the Open Field locations 1, 2 and 3 in Atibaia, São Paulo State, Brazil
 843 (Loc 1: 23°04'14.32"S 46°40'58.2"W, Loc 2: 23°04'33.5"S 46°40'30.1"W, Loc 3: 23°08'00.7 "S 46°37'04.5"W).

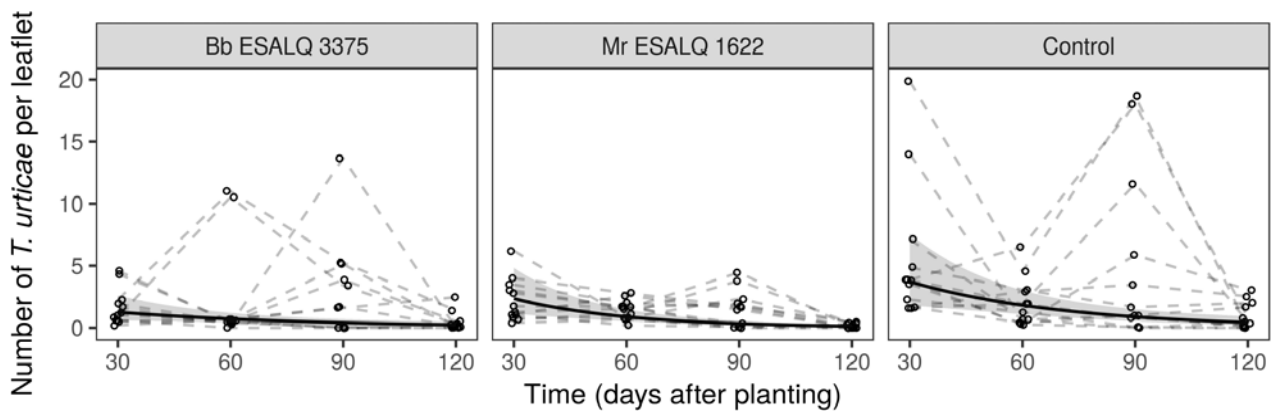
844 The dots represent the observations; the solid lines are the fitted curves for the mean number of *T. urticae* per
 845 leaflet and the gray areas represent 95% confidence intervals of the curves

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851 **Fig. 3** Effect of inoculation of strawberry root with *Beauveria bassiana* (Bb) isolate ESALQ 3375 or
 852 *Metarhizium robertsii* (Mr) ESALQ 1622 on numbers of adult *Tetranychus urticae* per leaflet from 30, 60, 90
 853 and 120 days after inoculation at the Low Tunnel location 4 in Senador Amaral, Minas Gerais State, Brazil
 854 (22°33'12.1"S 46°13'41.8"W). The dots are the observations; the solid lines are the fitted curves for the mean
 855 number of *T. urticae* per leaflet and the gray areas represent 95% confidence intervals