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2 **Internalisation of mycobacteria in plants confirmed by microscopy, culture and**
3 **quantitative real time PCR examination**

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21

22 Abstract

23

24 The environment is a reservoir of non-tuberculous mycobacteria and is considered as
25 a source of infection for animals and humans. Mycobacteria can persist in different
26 types of environments for a relatively long time. We have studied their possible
27 internalisation into plant tissue through intact as well as damaged root systems in
28 different types of plants grown *in vitro* and under field conditions. The substrate in
29 which plants were seeded was previously contaminated with different strains of
30 *Mycobacterium avium* (10^8 to 10^{10} cells/gram of soil), and faeces from animals
31 infected with paratuberculosis. We detected *M. avium* subsp. *avium*, *hominissuis* and
32 *paratuberculosis* in the stems and leaves of the plants using both culture and
33 quantitative real time PCR (qPCR). The presence of mycobacteria in the plant tissues
34 was confirmed by microscopy. The number of mycobacteria found inside plant tissue
35 was several orders of magnitude lower (up to 10^4 cells/gram of tissue) than the initial
36 amount of mycobacteria present in the culture media or substrate. These findings
37 lead us to the hypothesis that plants may play a role in the spread and transmission
38 of mycobacteria to other organisms in the environment.

39

40 Introduction

41

42 Non-tuberculous mycobacteria cause a wide range of diseases in animals and
43 immunocompromised individuals. Mycobacterial infection is acquired mainly through
44 the respiratory and the gastrointestinal tract. Mycobacteria are ubiquitously
45 distributed and some members are present in high numbers in natural and man-
46 made environments; thus, they pose a constant risk to susceptible species of animals
47 and immunocompromised humans. The diversity of mycobacteria in the environment
48 was studied using combination of molecular biology methods (1). This enabled
49 qualitative and quantitative analysis and detection of sequences of pathogenic
50 mycobacteria in all types of tested soil. Non-tuberculous mycobacteria have been
51 described as causal agents in different types of diseases, most often pulmonary, skin
52 and soft tissue infections (2). Although members of the *Mycobacterium avium*
53 complex are usually associated with pulmonary disease, colonization and infection of
54 the gastrointestinal tract in AIDS patients has also been described (3). However, the
55 route of transmission is usually unclear. Water has been proposed as a main
56 reservoir (4), while infections through aerosol from soil have also been described (5).
57 Little information is available regarding the possibility of plants or food of vegetable
58 origin being contaminated with mycobacteria. Likewise, only a small number of
59 studies were concerned with food as a source of infection in humans. A study of the
60 food as a source of exposure of mycobacteria to HIV positive patients detected
61 mycobacteria in seven out of 121 samples examined (6). A later study compared the
62 genotypes of *M. avium* isolates from patients and foods using PCR- Restriction
63 fragment length polymorphism and demonstrated a link between them (7). Non
64 tuberculous mycobacteria were isolated from from salads, leak, lettuce, mushrooms,

65 and other vegetables as well as apple juice. Twenty nine isolates were obtained from
66 46 samples, with the predominant isolated species being *M. avium* (8).

67 Studies investigating the contamination of vegetables with mycobacteria have not
68 proven whether mycobacteria can be present inside the plant tissue. In a few studies
69 mycobacteria were identified in or on the surface of different plants (9; 10; 11).

70 Zwieler et al. (11) studied the microbial communities present in the phyllosphere
71 of lettuce leaves. After Denaturing Gradient Gel Electrophoresis and sequencing
72 analyses, sequences from the genus *Mycobacterium* were found on leaves as well as
73 soil samples. The sequence obtained from conventionally grown lettuce was most
74 similar to *M. alvei*. Also, *M. avium* subsp. *paratuberculosis* (*M. a. paratuberculosis*)
75 was detected in grass samples using qPCR (12).

76 Plants in aquatic environment are also known to harbour mycobacteria. *M. avium*
77 was detected in a red beed samples from a constructed wetland, and plants were
78 also selected as possible reservoirs of *M. ulcerans* in the environment (9;13).

79 The penetration of bacteria, namely *Salmonella* and *E. coli* into plant tissues has
80 been studied previously (14; 15; 16; 17). It was shown that motile bacteria can enter
81 the plant through roots or even hydathode on the leaves of tomato plants (18).

82 The persistence of bacteria inside the plant tissue depends most probably on the
83 conditions inside the plant. The survival of *Salmonella* in basil was limited to a few
84 days (17). To the authors' knowledge, there is no record of internalisation of
85 mycobacteria through the intact root system of plants and their distribution inside the
86 plant itself.

87

88 The aims of this study were to investigate whether mycobacteria present in culture
89 media, or in faeces from infected cattle, can penetrate into the intact or damaged

90 plant tissue of two different plant species through their root system. To this end, we
91 first analysed the presence of mycobacteria into *in vitro*-grown plants under sterile
92 conditions. Subsequently, we performed a field experiment, in which plants were
93 grown in a phytotron. We used beans and tomatoes, as both produce edible parts
94 which are not in contact with contaminated soil. Moreover, both plants can be
95 routinely cultured in laboratory conditions.

96

97 Materials and Methods

98

99 *In vitro* experiment

100 *Mycobacterium avium* subsp. *hominissuis* (*M. a. hominissuis*, field isolate obtained
101 from infected swine) was grown under laboratory conditions on Middlebrook broth
102 (M7H9) with enrichment (OADC, Beckton and Dickinson) with constant shaking for
103 two weeks. *M. a. hominissuis* was chosen due to its rapid availability and fast growth
104 *in vitro*. The concentration of IS1245 was quantified using qPCR (19). To 20 ml of
105 Murashige and Skoog agar (Duchefa, Haarlem, The Netherlands) at a temperature of
106 50° C 10⁹/ml of the suspension were added. Cultivation was performed in 250 ml
107 glass cylindrical vessels. Surface sterilisation of the seeds of beans (Pinto beans;
108 *Phaseolus vulgaris*) was performed by submerging them in 3% Sodium hypochlorite
109 for three minutes, followed by three subsequent washes in sterile water (each 5
110 minutes). Sterilised seeds were planted on the agar and grown under laboratory
111 conditions for two weeks. Samples for microscopy, mycobacterial culture and qPCR
112 were collected from leaves and stems, using sterile forceps and scissors.

113

114 Field experiment

115 For the field experiment, we used *Mycobacterium avium* subsp. *paratuberculosis*
116 (*M. a. paratuberculosis*, reference strain CAPM 6381) and *Mycobacterium avium*
117 subsp. *avium* (*M. a. avium*, reference strain CAPM 5889). These subspecies were
118 chosen due to their possible presence in the soils after fertilization with faeces from
119 infected animals. Both strains were grown on Middlebrook broth (M7H9) with
120 enrichment (OADC, Beckton and Dickinson) and mycobactin J (Allied Monitor) with
121 constant shaking for one month.

122

123 Tomato seeds (*Solanum lycopersicum*) were planted in pots containing one litre of a
124 commercially available substrate of potting soil. The substrate was tested using
125 qPCR and it was negative for the presence of *M. a. avium* and *M. a. paratuberculosis*
126 prior to inoculation. Twenty four plants were tested in total; nine of them were grown
127 in soil contaminated with *M. a. avium*, twelve were grown in soil contaminated with
128 *M. a. paratuberculosis* and three served as negative controls. Three groups of plants
129 were grown in soil contaminated with *M. a. avium* and four groups were grown in soil
130 contaminated with *M. a. paratuberculosis*. Each group contained three plants. In the
131 first group, the substrate was contaminated with *M. a. avium* (10^8 cells /L of potting
132 soil) or *M. a. paratuberculosis* (10^{10} cells /L of potting soil) suspension immediately
133 before the seeding. Ten millilitres of the suspension was added to the soil and the
134 soil was mixed using glass spatula. In the second and third groups, the suspensions
135 were added two weeks after seeding, in the proximity of the main roots, with the
136 difference that the roots of the third group were mechanically damaged a syringe
137 needle. The fourth group consisted of plants seeded in the substrate mixed with
138 faeces from a cow clinically ill with paratuberculosis. The amount of cells in the

139 substrate after the addition of faeces was 10^6 cells/ L of potting soil (quantified using
140 qPCR).

141 The plants were grown in a phytotron, where day temperature was adjusted to 22 °C,
142 humidity at 29% and CO₂ concentration at 630ppm. There were three collection
143 points. Samples from leaves, stems, and fruits (where available), were collected two,
144 four, and eight weeks after seeding (for group 1) and after contamination (for groups
145 2 and 3).

146 Additionally, we performed distribution analysis on two selected plants, one grown on
147 *M. a. avium* contaminated soil and the other on *M. a. paratuberculosis* contaminated
148 soil. After eight weeks, we analysed stems and leaves from every 5 cm height of the
149 plant, as well as roots, fruit and pollen samples (the pollen sample was collected from
150 several plants during flowering phase).

151

152 Sample examination

153

154 Light and fluorescence microscopy

155 Microscopy was performed on histological cuts of stem (15-20 µm using a cryostat),
156 root and leaf tissue, which were stained according to the Ziehl Neelsen method.

157 Slides were analysed under 1000x magnification on an Olympus microscope BX41.

158 Slides for fluorescent microscopy were prepared using a primary polyclonal rabbit
159 anti-*Mycobacterium* antibody, and a secondary Cy3-labelled anti-rabbit antibody

160 (ExBio, Czech Republic). Briefly, the protocol included fixation in acetone, followed
161 by incubation with a blocking solution (Dako). The primary antibody was diluted to

162 10 µg/ml and incubation was performed overnight (12-16 hours) at 4 °C. After three
163 washes in PBS, the secondary antibody was applied for one hour at room

164 temperature. After the final wash in PBS, slides were mounted with mounting medium
165 and analysed under 1000x magnification on an Olympus BX41 microscope
166 (Olympus, Japan). For the samples stained with fluorescent antibody the 510-550 nm
167 filter was used.

168

169 Electron microscopy

170 Ultrathin sections of plant tissue were fixed in 3% glutaraldehyde in cacodylate buffer,
171 post-fixed in 1% OsO₄ solution in cacodylate buffer, dehydrated in 50,70,90,100%
172 acetone and embedded in an Epon-Durcupan mixture (Epon 812, Serva, Germany;
173 Durcupan, ACM Fluka, Switzerland). The sections were stained with 2% uranyl
174 acetate and 2% lead citrate and observed at 80 kV under a Philips EM 208
175 transmission electron microscope (Phillips, The Netherlands).

176

177 Cultivation of mycobacteria

178 Samples from stems and leaves were washed in 3 % Sodium hypochlorite for 3 min
179 and sterile water (3 times for 5 minutes) prior to culture in order to avoid possible
180 surface contamination. Samples from tomatoes were cut aseptically and the inside
181 part of the fruit was cultured. Cultivation of 1g of each sample (homogenized in 2 ml
182 of phosphate buffered saline) was performed without any decontamination, and from
183 an additional 1 g with decontamination according to Fischer et al. (20). Briefly, the
184 homogenised sample was treated with 1 M HCl for 20 minutes, and subsequently
185 was neutralised using 2M NaOH. The material was inoculated onto four different
186 culturing media (Table 1): Herrold egg yolk medium, Leslie medium, Middlebrook
187 MH711 with antibiotic mixture PANTA (Polymyxin B, Amphotericin B, Nalidixic acid,
188 Trimethoprim and Azlocillin) and without antibiotics. The Middlebrook MH711 was

189 obtained from BD Diagnostics (Denmark). The rest of the media were prepared in our
190 laboratory, as described (21). When culturing the samples from the tomato plants, it
191 was necessary to perform a decontamination step because of the high rate of
192 contamination.

193

194 DNA isolation and qPCR

195 Several commercially available kits for DNA isolation from plant material were tested
196 for their efficiency. Due to the expected low number of bacterial cells in the plant
197 tissue, we attempted to use as much starting material as possible. The best results
198 for 0.25g of tissue were achieved using the commercially available PowerFood
199 Microbial DNA isolation kit (MoBio, USA) with certain modifications to the original
200 protocol. Initial homogenization of the samples was done in a MagnaLyser (Roche,
201 Germany) at 6400 rpm/2 min, after the addition of four 3.2 mm chrome steel beads
202 (Biospec, USA) as well as the beads provided in the kit. An increased volume of the
203 lysis buffer (700 μ l) was used and an additional step of heating at 65° C for 10 min
204 with shaking at 1400 rpm was included. The remaining steps were performed
205 according to the manufacturer's recommendations. DNA was eluted in 100 μ l of pre-
206 heated TE buffer (Amresco, USA) and used subsequently in the qPCR reaction.
207 Triplex qPCR for simultaneous detection of IS1245 and IS901 was performed for
208 every sample in duplicate for the plants contaminated with *M. a. avium* and
209 *M. a. hominissuis*, according to Slana et al. (19). For the tomato plants contaminated
210 with *M. a. paratuberculosis*, qPCR for detection of IS900 was performed as described
211 earlier (22). The qPCR results were transformed to cells/g by calculating the mean copy
212 number of insertion sequences per cell (25 copies of IS1245 in *M. a. hominissuis*, 15 copies
213 of IS901 in *M. a. avium* and 15 copies of IS900 in *M. a. paratuberculosis*).

214

215 Results

216

217 *In vitro* experiment

218 The qPCR analyses showed that mycobacteria were present in all of the leaf and
219 stem samples from the plants grown on artificially contaminated media. The quantity
220 of mycobacteria was three order of magnitude lower than the quantity present in the
221 substrate media. The negative control gave no signal.

222 The results from culture are presented in Table 1. The decontamination method
223 clearly resulted in much lower yields in culture than samples processed without the
224 decontamination step. The comparison of four different culturing media also shows
225 that the highest recovery was obtained with the use of Middlebrook M7H11 agar. The
226 addition of PANTA antibiotics had no adverse effect on the mycobacterial growth.

227 Using electron microscopy, we observed structures similar to bacterial cells in the
228 plant tissues (Figure 1). Using Ziehl Neelsen microscopy on plant tissue sections of
229 15-20nm thickness we were able to observe acid-fast rods. We made similar
230 observations using fluorescence microscopy (Figure 1). Mycobacteria were observed
231 under microscopy only in the bean plants grown *in vitro*, were the number of cells per
232 gram exceeded 10^6 .

233

234 Field experiment

235 In the field experiment with the tomato plants, the results from qPCR are shown in
236 Figure 2 and Figure 3. *M. a. paratuberculosis* DNA was present in the entire stem
237 and leaf samples at the three sampling times in the plants from group 1. The fruit
238 sample from this group was positive only at the first sampling point two weeks post
239 inoculation. The samples from group 2 gave similar results, although leaf stem

240 samples were negative 8 weeks after inoculation, but DNA was detected in fruits at
241 four weeks post contamination. Stem samples from group 3 (with damaged roots)
242 were positive four and eight weeks after contamination, but leaf samples were
243 positive in all three time points. Fruits were positive eight weeks after contamination
244 with mycobacteria. The quantity was similar in all of the samples and ranged between
245 10^2 to 10^4 cells per gram of tissue. Samples from group 4 were positive in leaf and
246 stem tissue but not in fruits. The quantity of IS900 reached up to 10^3 /g of tissue
247 (Figure 2). The samples from tomatoes grown in soil contaminated with *M. a. avium*
248 gave similar results. Stem samples were positive in group 1 at four and eight weeks
249 after seeding, but leaf samples were positive at all sampling points. In samples from
250 group 2 all of the stem and leaf samples tested positive, but none of the fruit
251 samples. Leaf samples from group 3 were positive in all time points, fruit sample was
252 positive at two weeks after contamination. The quantity of *M. a. avium* specific DNA
253 ranged between 10^1 to 10^5 IS901 copies per gram of tissue (Figure 3).
254 We obtained also three isolates from cultivation. Two of the isolates (*M. a. avium*)
255 were from the stems in group 2 and 3, at the two week sampling time, and one
256 isolate (*M. a. paratuberculosis*) from group 2 was obtained one month after seeding.
257 The samples from the control group were negative. The qPCR results of the
258 distribution of mycobacteria inside the plant are shown in Figure 4. Mycobacteria
259 were mostly concentrated in the root samples and their quantity decreased through
260 the fruit, although we detected the target sequence also in the pollen sample.

261

262

263 Discussion

264

265 The food safety of vegetables has been of increasing concern since recent outbreaks
266 of *Salmonella* and *E. coli* were traced back to vegetables and sprouts. Much research
267 has focused on these two pathogens and their potential for penetration versus
268 surface contamination of vegetables (23, 24). Solomon et al. (25) described the
269 migration of *E. coli* O157:H7 from contaminated soil into the tissue of lettuce. There
270 have also been studies on the internalisation of *Salmonella* into tomato plants
271 through roots or even hydathodes (18). Although mycobacteria were detected in
272 vegetables previously, the present study is the first one to confirm their internalisation
273 inside the plant tissue. The large numbers of *M. avium* cells used in our experiment
274 may have biased the results; however, recent study showed that the quantity of
275 mycobacteria in soil as well as their diversity is high (1).

276 Plants or vegetables have been suspected as possible routes for foodborne
277 mycobacterial diseases (6, 8). Typing of *M. avium* isolates from food and patients
278 showed same DNA patterns (8). Our results demonstrate that mycobacteria can be
279 taken up by the root systems of plants, even in plants with intact roots. The
280 persistence of mycobacteria in soil, manure and different parts of the environment
281 has been demonstrated previously (12, 26). *M. a. paratuberculosis* remained viable in
282 manure after 55 weeks (26). Manure from domestic animals is used as fertilizer on
283 fields where crops or vegetables are grown. Therefore, it is plausible that due to its
284 presence inside plants, there might be a risk of infection to grazing animals or to
285 humans. Although in the present study we have proven the presence of
286 mycobacteria inside the plant tissue, we have not performed any experiments
287 regarding its pathogenicity. Future research should be focused on the testing of the
288 pathogenicity of mycobacteria after the internalisation in plants. Next step would be

289 feeding animals with such plants in order to see whether this route of transmission is
290 plausible.

291 In our study, mycobacteria were present inside the plant tissue for at least two
292 months after contamination of the potting soil. This may be due to the properties of
293 mycobacteria as well as the environment inside the plant tissue. A study on the
294 survival of *Salmonella* in basil showed a decline after only three days (17), although
295 other studies have detected *Salmonella* in tomato fruits 49 days after inoculation (27).

296 Regarding the distribution of mycobacteria inside the plants, *M. a. avium* and
297 *M. a. paratuberculosis* DNA quantity was highest in the roots, and gradually
298 decreased along the height of the plant. The presence of mycobacterial DNA in the
299 fruit and pollen samples is noteworthy regarding food safety and further spread of
300 mycobacteria. However, we did not obtain an isolate from these samples using
301 culture.

302 In conclusion, we have demonstrated the internalisation of mycobacteria into different
303 types of plants; further, the distribution within the plant was found to be even.

304 However, the number of mycobacteria found inside plant tissue was several orders of
305 magnitude lower than the initial amount of mycobacteria present in the culture media
306 or substrate. Mycobacteria are probably passively taken inside the roots rather than
307 actively penetrate root epidermis. The passive intake could be also facilitated by the
308 relatively small size of mycobacterial cells. Although mycobacteria inside plant tissue
309 pose a possible risk of transmission, we suspect that the subsequent handling of
310 vegetables and secondary surface contamination with mycobacteria might play a
311 bigger role in transmission of the infectious agent.

312

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314

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412 in tomato plants from the time of inoculation at flowering and early stages of fruit
413 development through fruit ripening. *Appl. Environ. Microbiol.* **10**:4760-4764.

414
 415 Table 1. qPCR and culture examination of stem and leaf samples in *in vitro* grown beans; Input concentration of *Mycobacterium avium* subsp.
 416 *hominissuis* culture added to the agar was 10⁹ cells/ ml.
 417
 418

Part of beans	Plant No.	qPCR IS1245 (cells/g)	Cultivation without decontamination (CFU)				Cultivation with decontamination (CFU)			
			MB+PANTA	MB	HEYM	Leslie	MB+PANTA	MB	HEYM	Leslie
Stem	1	6.31 x 10 ⁶	>1000	>1000	100	0	150	0	0	0
	2	1.54 x 10 ⁷	100	0	0	200	20	0	0	0
	3	7.74 x 10 ⁶	1000	>200	>1000	200	10	10	0	0
Mean		9.81 x 10 ⁶	3700	3400	366	133.33	60	3.33	0	0
Standard deviation		4.88 x 10 ⁶	5474	5716	550.7	115.5	78.1	5.7	0	0
Leaf	1	6.70 x 10 ⁶	200	>200	1000	0	0	100	100	10
	2	1.74 x 10 ⁶	>200	>200	500	100	0	0	0	0
	3	3.49 x 10 ⁶	>100	>200	>1000	0	0	100	10	100
Mean		3.97 x 10 ⁶	166	200	3833	33.33	0	66.67	36.67	36.67
Standard deviation		2.51 x 10 ⁶	57.73	0	5346	57.7	0	57.7	55	55
Negative control		0	0	0	0	0	0	0	0	0

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 420
 421 1. Middlebrook M7H11+PANTA
 422 2. Middlebrook M7H11 without antibiotics
 423 3. Herrold egg yolk medium
 424 4. Leslie medium
 425

426 Figure 1. Microscopy of bean stems sections containing *M. a. avium*. A and B: Transmission electron microscopy. C: Ziehl-Neelsen
427 staining of bean stem tissue, a mycobacteria is stained in red against blue background. D: Specific-antibody labelled *M. a. avium*
428 inside bean stem tissue. Fluorescent rods were observed inside the transport plant cells.
429 Figure 2. Detection of *M. a. paratuberculosis* DNA inside tomato plants.
430 Figure 3. Detection of *M. a. avium* DNA inside tomato plants.
431 Figure 4. Distribution of *M. a. avium* and *M. a. paratuberculosis* DNA inside tomato plants.
432

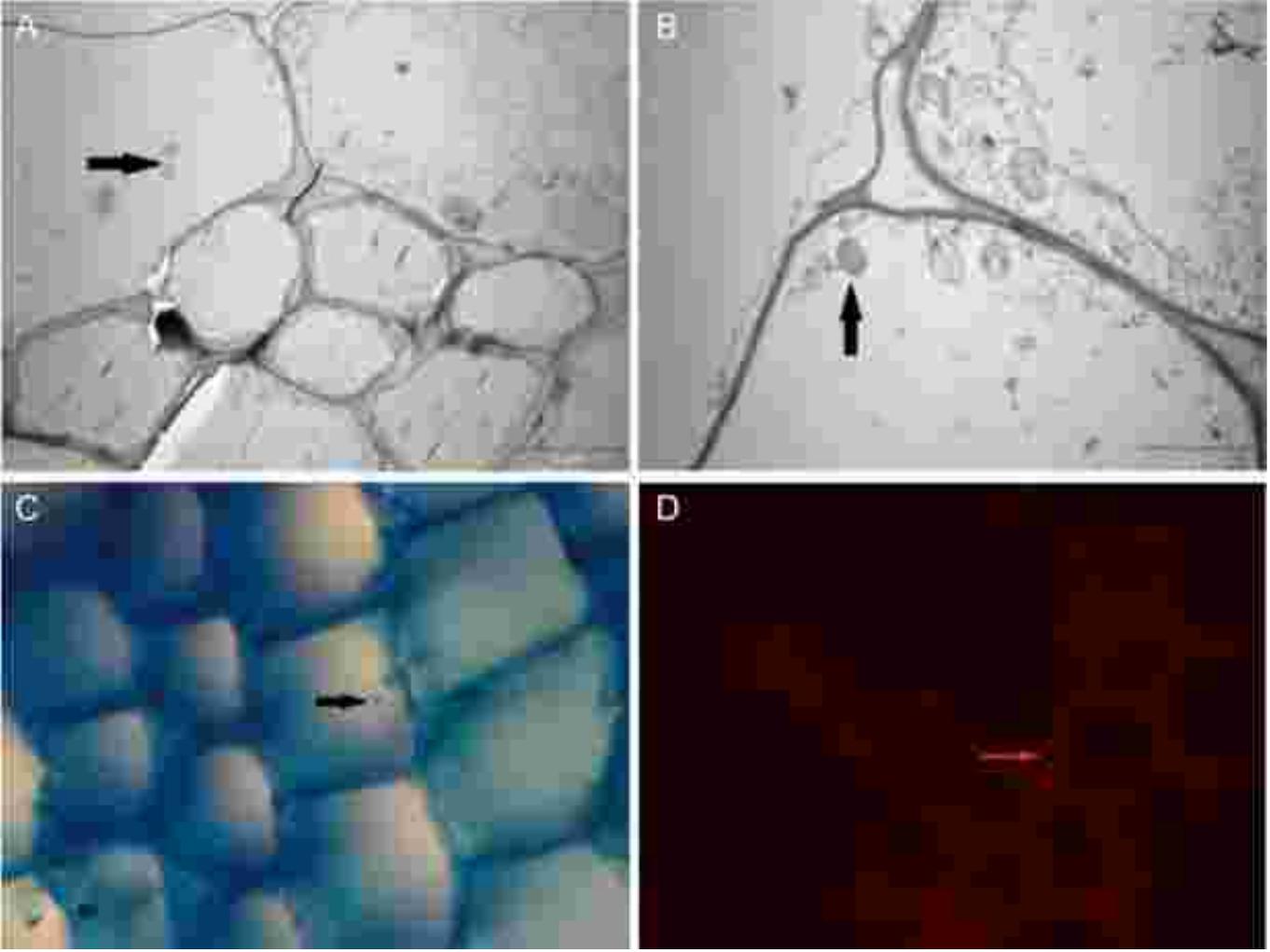


Figure 2. Detection of *M. a. paratuberculosis* DNA in tomato plants

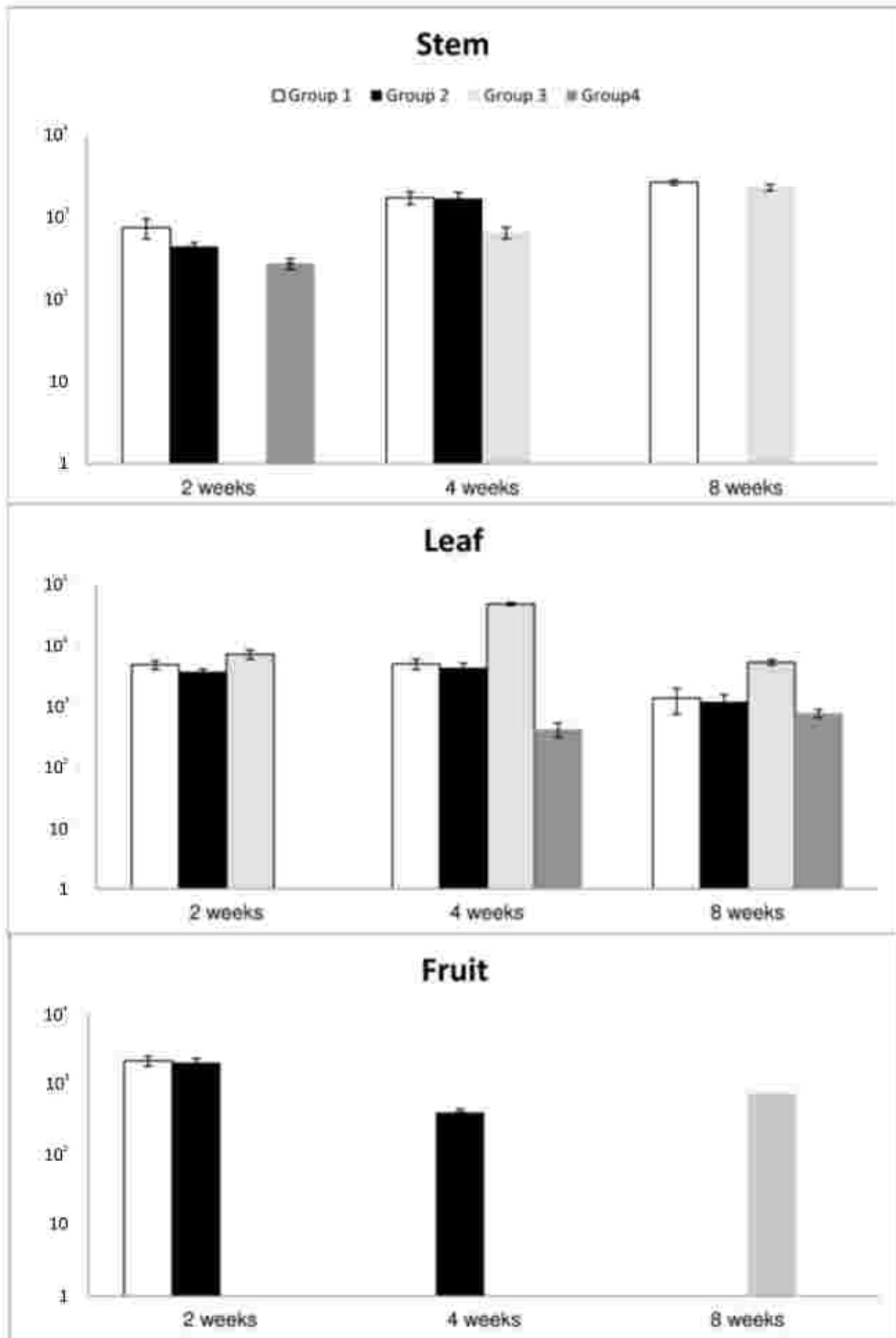


Figure 3. Detection of *M. a. avium* DNA in tomato plants

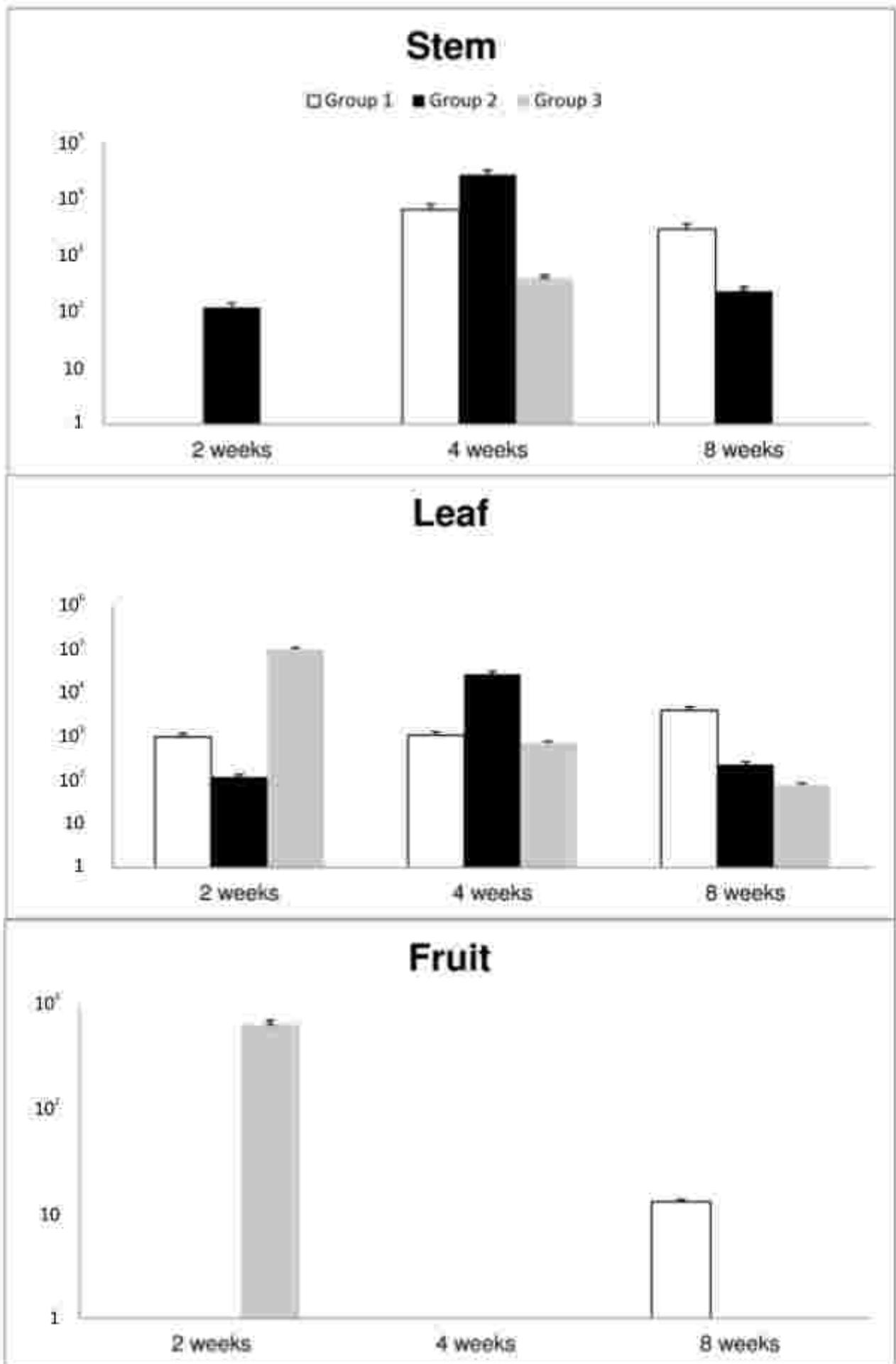


Figure 4. Distribution of *M. a. avium* and *M. a. paratuberculosis* DNA inside the tomato plants

