

***Rahnella* spp. are commonly isolated from onion (*Allium cepa*) bulbs and are weakly pathogenic.**

Journal:	<i>Applied Microbiology</i>
Manuscript ID	JAM-2019-0362.R1
Journal Name:	Journal of Applied Microbiology
Manuscript Type:	JAM - Original Article
Date Submitted by the Author:	15-May-2019
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Key Words:	Plant diseases, Plant pathology, PCR (polymerase chain reaction), Pathogenesis, Infection

This is a post-peer-review, pre-copyedit version of an article published in Journal of Applied Microbiology. The final authenticated version is available online at: <http://dx.doi.org/10.1111/jam.14340>

1 *Rahnella* spp. are commonly isolated from Onion (*Allium cepa*) bulbs and are weakly
2 pathogenic
3

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7 Abbreviated running title: *Rahnella* spp. in onion (*Allium cepa*)
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20 **Abstract**

21 **Aims:** Bacterial decays of onion bulbs have serious economic consequences for
22 growers, but the etiologies of these diseases are often unclear. We aimed to determine
23 the role of *Rahnella*, which we commonly isolated from bulbs in the USA and Norway, in
24 onion disease.

25 **Method and Results:** Isolated bacteria were identified by sequencing of housekeeping
26 genes and/or fatty acid methyl ester (FAME) analysis. A subset of *Rahnella* spp. strains

27 were also assessed by multilocus sequence analysis (MLSA); most onion strains
28 belonged to two clades that appear closely related to *R. aquatilis*. All tested strains
29 from both countries caused mild symptoms in onion bulbs but not leaves. PCR primers
30 were designed and tested against strains from known species of *Rahnella*. Amplicons
31 were produced from strains of *R. aquatilis*, *R. victoriana*, *R. variigena*, *R. inusitata*, and
32 *R. bruchi*, and from one of the two strains of *R. woolbedingensis*.

33 **Conclusions:** Based on binational testing, strains of *Rahnella* are commonly
34 associated with onions, and they are capable of causing mild symptoms in bulbs.

35 **Significance and Impact of the Study:** While *Rahnella* strains are commonly found
36 within field-grown onions and they are able to cause mild symptoms, the economic
37 impact of *Rahnella*-associated symptoms remains unclear.

38
39 **Keywords:** Plant diseases, Plant pathology, PCR (polymerase chain reaction), Pathogenesis,
40 Infection

41 42 Introduction

43 Onions (*Allium cepa*) are susceptible to damage by a number of pests and
44 pathogens, including insects, nematodes, fungi, and bacteria. Decays of onion bulbs
45 caused by bacteria can cause serious economic losses. Bacterial decays can develop
46 in the field during the growing season or in post-harvest storage of the bulbs. Numerous
47 bacteria have been described as onion bulb pathogens, including strains from genera
48 *Burkholderia*, *Enterobacter*, *Pantoea*, *Pseudomonas*, *Pectobacterium*, *Lactobacillus*,
49 and *Leuconostoc* (Schwartz and Mohan, 2007; Bonasera *et al.* 2017).

50 Onion bulbs with bacterial decay may have any combination of discoloured,
51 water-soaked, macerated, or shrunken scales. In disease caused by macerating
52 bacteria, for example *Burkholderia* spp. or *Dickeya* sp. (Mahenthiralingam *et al.* 2005;

53 **Palacio-Bielsa et al. 2007**), rotten bulbs can often be identified by visual inspection of
54 intact bulbs or by manually assessing bulb firmness, especially at the bulb neck. In New
55 York State and in Norway, growers often employ skilled workers to hand-sort bulbs and
56 cull any with discernible symptoms of decay.

57 While macerating bacteria often cause significant damage to bulbs and affect
58 bulb integrity, non-macerating bacteria, **for example *Pantoea ananatis* or *Enterobacter***
59 **sp. (Carr et al 2010; Schroeder and du Toit 2010)**, may cause internal discolouration of
60 scales. They may slightly reduce the firmness of the bulb neck, but often cause no
61 external symptoms, making them indistinguishable from healthy bulbs during grading.
62 When shipments of bulbs are received by potential buyers, a random sample of bulbs
63 typically is cut and inspected. If inspection reveals unacceptable numbers of
64 symptomatic bulbs, the entire shipment may be rejected. Manual sorting and rejected
65 lots add to the economic impact of bacterial decays of onions on grower profits.

66 In both New York State and Norway, onion bulbs may be stored for several
67 months after harvest before they are sorted and marketed. In 2010, bacteria were
68 recovered from more than 500 bulbs that had been culled during hand-sorting from cold
69 storage in western New York State. Strains putatively identified as *Rahnella* spp. were
70 recovered from **more than 25%** of culled bulbs. Also, in Norway, similar surveys yielded
71 *Rahnella* spp. from more than 20% of symptomatic bulbs. In the current work, we
72 determined that strains of *Rahnella* spp. were widely distributed geographically as
73 onion-associated bacteria, and they elicited mild symptoms in artificial inoculation
74 experiments. We isolated several species of *Rahnella* from onions. **Most strains**
75 **clustered into two clades that appear to be closely related to *R. aquatilis*.** To facilitate

76 further work detecting *Rahnella* strains, we developed specific primers and an
77 associated protocol for a polymerase chain reaction (PCR) test.

78

79 **Materials and methods**

80

81 **Bacterial growth and maintenance (USA)**

82 Bacteria were routinely grown on Luria-Bertani (LB) agar plates and incubated for
83 1-2 days at 28°C for use in colony PCR or for inoculations of bulbs. For storage of
84 strains, bacteria were transferred from freshly grown plates using sterile cotton-tipped
85 applicators into sterile-filtered 15% glycerol. Bacteria were stored at -80°C.

86

87 **Bacterial growth and maintenance (Norway)**

88 Bacteria were routinely grown on Nutrient Glucose Agar (NGA) (Lelliott and
89 Stead, 1987). NGA plates were incubated for 1-3 days at room temperature for colony
90 PCR or inoculations of bulbs. For storage of identified strains, bacteria were transferred
91 from freshly grown plates to “protect” vials (Technical Service Consultants, Lancashire,
92 UK) containing ceramic beads. Bacteria were then stored at -80°C.

93

94 **Isolation of bacteria from onion and environmental samples (USA)**

95 In 2010, growers in western New York State set aside onions with suspected
96 bacterial decay during hand-sorting in cold storage prior to marketing. Approximately
97 500 bulbs, mostly symptomatic, were sampled at this time. In the winter of 2011-2012,
98 one wooden crate of onions (approximately 400 kg) was selected for sampling from

99 each of three growers' cold storages in the Elba, NY region. Approximately 100 onions
100 were randomly chosen from those crates three times over the storage season, in
101 October, January, and March. In New York, onions are typically harvested in late
102 August through mid-October. Bulbs were refrigerated until processed by lab personnel.
103 Other samples were occasionally received from onion growers suspecting rot in growing
104 onion plants or recently harvested or stored bulbs. Plants were typically sent to the lab
105 by overnight mail and processed immediately or refrigerated and processed within a few
106 days of arrival.

107 For onion plants from the field, roots were trimmed, and plants were rinsed with
108 distilled water to remove soil particles. Symptomatic tissues or disease margins were
109 probed with sterile wooden applicators and streaked onto onion extract medium (OEM)
110 (Zaid *et al.* 2012) directly. When dealing with bulbs, they were bisected longitudinally,
111 photographed and assessed for symptoms, and bacterial isolations were made from
112 each bulb. Representatives of the various colony types growing on OEM plates were
113 dilution streaked to purity on LB agar. All incubations were carried out at 28°C.

114 Strain FC061912-K was isolated from a creek flowing adjacent to an onion field
115 in Western New York. A volume of 400 ml of creek water was centrifuged at 5500 x g
116 for 15 minutes. The resulting pellet was resuspended in 1/100 volume of autoclaved
117 high-purity water, and 100 µl were plated on OEM agar. Colonies of different
118 morphologies were picked and purified by dilution streaking.

119

120 **Isolation of bacteria from onions (Norway)**

121 The majority of putatively diseased onions were collected from the southeastern
122 part of Norway, in the counties Vestfold, Østfold and Oppland. A smaller number of
123 samples originated from the counties Hedmark, Rogaland and Nord-Trøndelag.
124 Samples were collected from the field during the growing season, directly after harvest,
125 or after storage. In addition, samples were collected from field trials where pathogen
126 control measures with various compounds were being investigated. A total of 368
127 samples, each consisting of one to 20 onions, typically three to five, were collected
128 during the project period (2012 to 2015), and stored at 5°C until processed.

129 For onion plants from the field, roots were trimmed, and plants were rinsed with
130 distilled water to remove soil particles. For both growing plants and mature bulbs,
131 symptomatic tissues or the margins between symptomatic and healthy tissue were
132 sampled. Bacteria were released from the sampled tissue by either soaking for 30
133 minutes in sterile 10 mM phosphate buffered saline, pH 7.2 (PBS) (Anonymous, 2006)
134 or crushing in sterile water. Resulting suspensions were dilution streaked onto **NGA**.
135 **Onion** tissue samples were homogenized in 10-15 ml SPCB buffer (120 mM
136 sodium phosphate, 2 % CTAB, 1.5 M NaCl, pH 8.0) using a Bioreba homogenizer. DNA
137 was isolated from the crude extract using the Kingfisher Duo Prime with KingFisher Cell
138 and Tissue DNA kit, according to the manufacturer's (Thermo Fischer Scientific,
139 Waltham, MA) instructions.

140

141 **Preliminary Identification of bacteria (USA)**

142 **In New York, bulbs harvested from the same field and sampled at the same time**
143 **were treated as batches. Strains from the same batch of bulbs were grouped based on**

144 similar colony morphologies, digest patterns of amplicons from the DNA gyrase subunit
145 B gene (*gyrB*) as described by Bonasera *et al.* (2014), and by results of indole tests,
146 nitrate reductase and oxidase activities (Schaad *et al.* 2001) and by fluorescence on
147 King's B agar (King *et al.* 1954), modified to contain 0.4 g instead of 1.5 g of
148 MgSO₄·7H₂O per liter. Representative strains were chosen from each group, and *gyrB*
149 amplicons obtained by using the 1480F/2242R primer pair (Bonasera *et al.* 2014) were
150 sequenced: amplicons were cleaned using the Clean & Concentrator-5 kit (Zymo
151 Research Corp., Irvine, CA) and sequenced using the *gyrB* 1480F primer, at the Cornell
152 University Biotechnology Resource Center. Resulting sequences were used to search
153 the NCBI Nucleotide collection (nr/nt) database via blastn
154 (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>). Strains with *gyrB* fragment sequences that were
155 most similar to *Rahnella* strains were pursued further.

156

157 **Preliminary Identification of bacteria (Norway)**

158 Isolates were identified initially by fatty acid methyl ester (FAME) analysis
159 (Sasser, 1990). Of 431 isolates, 130 isolates were also identified by sequencing of a
160 hypervariable region of the 16S ribosomal gene, using primers F985PTO and R1378
161 and conditions as described previously (Heuer *et al.* 1999, Table 1). Templates were
162 from bacterial colonies suspended in 500 µl sterile H₂O and incubated for 10 min at
163 96°C. Purified PCR amplicons were sequenced in both directions at GATC Biotech,
164 Germany, using the same primer set as for the PCR amplification. Sequences were
165 assembled, manually edited and aligned using the CLC Main Workbench.

166

167 **Onion bulb inoculations with *Rahnella* strains**

168 Yellow onion bulbs were purchased from a local grocery store and prepared as
169 described by Schroeder *et al.* (2009). Strains C1b and A66, isolated in North America,
170 were grown on LB agar for 1-2 days at 28°C and swabbed from plates using sterile
171 cotton-tipped applicators into autoclaved high-purity water. Bacterial suspensions were
172 adjusted to OD₆₀₀ of 0.2. Four to five bulbs per strain were injected with 100-500 µl of
173 inoculum using a syringe and 18 gauge needle. Bulbs were incubated at 28-30°C for
174 10-17 days, after which they were cut longitudinally and assessed for symptoms.
175 Bacteria were recovered from inoculated onion bulbs using LB agar and assessed with
176 Rah 3783 F1/R1 primers or by production of a PCR amplicon using gyrB1480 F/R
177 primers followed by sequencing of the amplicons. Inoculation and re-isolation
178 experiments were completed for strain C1b and A66 three times each, with bacteria
179 recovered from one or two bulbs per assay.

180 To compare pathogenicity of *Rahnella* strains isolated from Europe and North
181 America, eight isolates of *Rahnella spp.* that had also been included in MLSA (four from
182 Norway and four from the USA) were compared in a pathogenicity test as described
183 above (Figure 2), and scored based on the degree of symptoms (Figure 3). Data were
184 analysed by analysis of variance, and significant differences were separated using
185 Tukeys pairwise comparison (Minitab).

186

187 **Onion leaf inoculations**

188 Onion plants were grown in an environmental growth chamber as described
189 previously (Bonasera *et al.* 2017). Six leaves of twelve plants were inoculated with

190 strains C1b or A66 or water by dipping sterile toothpicks in bacterial suspensions or
191 water. These strains were chosen as strains that were isolated early in the study and,
192 based on preliminary analysis of sequencing data from their *gyrB* 1480F/2242R
193 amplicons, both were *Rahnella* and were clearly distinct from each other. Bacterial
194 suspensions were prepared as for onion bulb inoculations. Six plants inoculated with
195 each strain or sterile water were placed in an incubator set to 30°C and six others were
196 placed at room temperature in the laboratory.

197

198 **Partial *gyrB* sequencing 1480F/2242R**

199 In order to place the Norwegian *Rahnella* strains in context with strains isolated
200 from New York, six strains from Norway were sequenced using the *gyrB* 1480F/2242R
201 primers. Additionally, twelve strains from five different species of *Rahnella* and the
202 closely related bacterium *Ewingella americana* were sequenced with the same primers
203 for use as references. For most strains, these sequences were generated by a single
204 sequencing reaction. Sequences were aligned in Megalign (DNASStar, Madison, WI),
205 and were trimmed to eliminate ambiguous base calls and gaps resulting from poor-
206 quality sequence occurring at the beginning or end of amplicons. Quality of the
207 remaining sequences were then assessed by viewing trace files using FinchTV 1.4.0
208 (Geospiza, Inc.; Seattle, WA, USA; <http://www.geospiza.com>). For several strains,
209 additional PCR and sequencing was performed to obtain good-quality sequence over
210 the whole alignment. Sequences generated for this work were deposited in Genbank
211 under accession numbers MK391682-MK391746 and MK408759 (Table S1).

212

213 **Multilocus sequence analysis (MLSA)**

214 Seven strains from the USA and four strains from Norway that were putatively
215 identified as *Rahnella* spp. were further analysed by MLSA, using partial sequences of
216 four conserved housekeeping genes, *gyrB*, *rpoB* (RNA polymerase β subunit), *infB*
217 (translation initiation factor IF-2), and *atpD* (ATP synthase subunit beta). In an effort to
218 place *Rahnella* onion isolates in context with existing sequence data of *Rahnella* spp.,
219 amplicons with coverage that included the sequence positions used in the MLSA
220 published by Brady *et al.* (2014) were obtained. Consequently, a combination of
221 previously published and new primers were used (Table 1), as not all primer pairs from
222 Brady *et al.* (2014) worked well with the conditions used in this study. PCRs to generate
223 amplicons for sequencing were generally performed in 24 μ l volumes, using 12.18 μ l of
224 water, 4.8 μ l 5x OneTaq GC buffer (New England Biolabs, Ipswich, MA), 2.4 μ l 2.5 mM
225 dNTPs, 1.25 μ l each of the forward and reverse primers, 0.12 μ l of OneTaq (New
226 England Biolabs), and 2 μ l template. For each novel sequence used in the MLSA
227 (GenBank accession nos. MK387392-MK387415, Table S2), two amplicons produced in
228 separate reactions were sequenced as described above. Contigs were assembled using
229 SeqMan Pro version 12.2.0 or 13.0.0 (DNASStar). Groups of sequences were aligned in
230 MegAlign or using the “align two or more sequences” option for the blastn tool
231 (<https://blast.ncbi.nlm.nih.gov/>). Sequences were trimmed to match the coverage of
232 previously reported *Rahnella* MLSA sequences (Brady *et al.* 2014) and concatenated in
233 the following order: *gyrB*, *rpoB*, *infB*, and *atpD*. The *gyrB* sequences used in MLSA are
234 upstream of and do not overlap with the sequences obtained from the *gyrB*
235 1480F/2242R amplicons.

236

237 **Generation of phylogenetic trees**

238 Separate phylogenetic trees were constructed for the *gyrB* 1480F/2242R
239 amplicon sequences (Table S1, Figure S1) and for the concatenated MLSA sequences
240 (Table S2, Figure 1). Sequences were aligned using the ClustalW method according to
241 default parameters, and phylogenetic trees were generated using MEGA version 7.0.26
242 (Kumar *et al.* 2016). There were no gaps in the alignments. The evolutionary history
243 was inferred using the Maximum Likelihood method based on the Tamura-Nei model
244 (Tamura and Nei, 1993). Initial tree(s) for the heuristic search were obtained
245 automatically by applying Neighbor-Join and BioNJ algorithms to a matrix of pairwise
246 distances estimated using the Maximum Composite Likelihood (MCL) approach, and
247 then selecting the topology with superior log likelihood value.

248

249 **Design of *Rahnella*-specific primers**

250 The genomes of *Rahnella aquatilis* CIP 78.65 = ATCC 33071 (GenBank
251 accession no. CP003244.1) (Martinez *et al.* 2012a) and *Serratia proteamaculans* 568
252 (CP000826.1) were aligned using Progressive Mauve, Mauve version 2.3.1 build 173
253 (Darling *et al.* 2010). Strains of *Serratia* are relatively close relatives to *Rahnella*, and
254 are occasionally isolated from onions. The *Serratia* strain was included in the
255 comparison in order to exclude genes that are conserved outside of the genus
256 *Rahnella*. Genes annotated as “hypothetical proteins” and present in the *Rahnella* strain
257 but not in the *Serratia* strain were used to search the NCBI Genomes database using
258 blastn. Genes present in the three *Rahnella* genomes available at the time, *R. aquatilis*

259 ATCC 33071, *Rahnella* sp. Y9602, and *R. aquatilis* HX2, but not in other available
260 genomes were used to search the NCBI Whole Genome Shotgun (WGS) database and
261 filtered based on length (at least 300 bp). Putative genes that appeared to be unique to
262 the three sequenced *Rahnella* strains were considered good target regions for
263 designing specific primers. Similar sequences from strains ATCC 33071, Y9602, and
264 HX2 were aligned using MegAlign, and well-conserved portions of three genes,
265 Rahaq2_0130, Rahaq2_3783, and Rahaq2_3707, were selected for primer design.
266 Target regions were manually chosen, and annealing temperature and predicted
267 annealing sites within the target genes were assessed using PrimerSelect (DNASar).
268 Potential primers were checked for specificity to *Rahnella* by searching specifically
269 genomes from the Enterobacteriaceae (taxid:543), Pseudomonadales (taxid:72274),
270 and Burkholderiaceae (taxid:119060) using the Primer-BLAST tool from NCBI
271 (<https://www.ncbi.nlm.nih.gov/tools/primer-blast/index.cgi>). Primer pair Rah 3783 F1/R1
272 (Table 1), designed to amplify part of the gene designated Rahaq2_3783 in strain ATCC
273 33071, yielded single amplicons of the expected size, 525 base pairs (bp), from six
274 *Rahnella* strains in preliminary experiments. This pair was further assessed for
275 specificity and sensitivity.

276

277 **Assessment of *Rahnella*-specific primers**

278 Bacterial suspensions for use in colony PCR were prepared by touching sterile
279 wooden applicators five times to ribbons of bacterial growth on LB agar plates and
280 swirling the applicator into 200 µl of sterile high-purity water. The resulting suspensions
281 were slightly cloudy and used as templates in PCR directly.

282 Each 12 µl reaction contained 5.09 µl water, 2.4 µl 5x OneTaq GC buffer (New
283 England Biolabs, Ipswich, MA), 1.2 µl 2.5 mM dNTPs, 0.625 µl 10 µM Rah 3783 F1
284 primer, 0.625 µl 10 µM Rah 3783 R1 primer, 0.06 µl OneTaq DNA Polymerase (New
285 England Biolabs), and 2 µl of template. Amplification was performed with one cycle at
286 95°C for 10 min; 45 cycles of: 95°C for 30 s, 57°C for 45 s, 72°C for 50 s; and a cycle of
287 72°C for 10 min. PCR products were analysed following electrophoresis through a 1%
288 agarose gel.

289 To assess specificity of the Rah3783 F1/R1 primer pair, 19 strains of *Rahnella*
290 spp. isolated from onions, selected from different branches of a phylogenetic tree based
291 on partial *gyrB* sequence, were used for testing. In addition, 11 strains from 8 other
292 genera documented as onion pathogens (*Xanthomonas axonopodis* pv. *allii*,
293 *Pseudomonas viridiflava*, *Pectobacterium carotovorum* subsp. *carotovorum*, *Pantoea*
294 *ananatis*, two strains of *Pantoea agglomerans*, *Erwinia rhapontici*, *Enterobacter* sp.,
295 *Dickeya dadantii*, *Burkholderia gladioli* pv. *alliicola*, and *Burkholderia cepacia*) were
296 included in the primer testing, as well as 10 reference strains of *Rahnella* spp. and two
297 of *E. americana*. This experiment was repeated three times.

298 Primer sensitivity of the Rah3783 F1/R1 primer pair was determined against
299 bacterial suspensions of *Rahnella* sp. Y9602 in sterile water. Bacterial suspensions
300 were adjusted to an optical density at 600 nm (OD₆₀₀) of 0.2 (approximately 10⁸ CFU/ml)
301 and were serially diluted in 10-fold steps ranging approximately 10⁸ to 10¹ CFU/ml.
302 Volumes of 5 µl from each dilution were spotted five times each onto LB agar and
303 incubated overnight at 28°C to obtain colony counts of viable bacteria. PCRs of 12 µl
304 were prepared as stated above. Undiluted, 10⁻¹, 10⁻², 10⁻³, 10⁻⁴, and 10⁻⁵ dilutions of

305 bacterial suspension and water were used as templates. This experiment was repeated
306 three times.

307

308 Results

309 Recovery of *Rahnella* spp. from naturally infected onion bulbs and plants

310 *Rahnella* strains were frequently recovered from onions in both the USA and
311 Norway. In western NY in 2010, *Rahnella* strains were recovered from 136 of 508 culled
312 bulbs assessed (27%). This initial survey prompted an additional study of bacteria from
313 onion bulbs. In the winter of 2011-2012, onion bulbs were randomly sampled from
314 growers' overwinter storage in western NY. Bulbs were sampled early (late October),
315 midway (late January), and late (mid-March) in the storage season. *Rahnella* strains
316 were recovered from both healthy and symptomatic bulbs at low levels. Bulbs from
317 which *Rahnella* strains were isolated ranged from nonsymptomatic to completely
318 discoloured with severe maceration (Figure S2). *Rahnella* strains were recovered from 9
319 of 748 (1%) of healthy-appearing and 25 of 150 (17%) of symptomatic bulbs (Table S3).
320 *Rahnella* strains were often isolated together with other genera of bacteria. In Norway
321 *Hafnia* sp. and *Serratia* sp. were most often isolated together with *Rahnella*, while in the
322 USA *Pseudomonas* spp. were most often isolated together with *Rahnella* strains.
323 In a 4-year Norwegian survey of 368 samples of groups of one to twenty
324 symptomatic bulbs, 109 isolates were identified as *R. aquatilis* by FAA and/or 16S
325 sequencing. The FAA similarity index was > 0.8 for *R. aquatilis*. These were confirmed
326 with 16S rRNA sequences, which were 100% similar to a number of different *R.*
327 *aquatilis* isolates.

328

329 Pathogenicity of *Rahnella* strains

330 Attempts to infect onion leaves using *Rahnella* strains C1b and A66 were not
331 successful. Currently, there is no evidence that *Rahnella* strains are capable of causing
332 leaf lesions (data not shown).

333 Artificially inoculated yellow onion bulbs showed symptoms ranging from mild
334 discolouration along the inoculation site to water-soaking and discolouration of one or a
335 few internal scales, but the bulbs generally remained firm and without signs of
336 maceration. Symptoms were distinct from sterile water-injected negative controls |
337 Severity of symptoms were not completely consistent, sometimes resulting in more
338 severe symptoms (Figure S3). Bacteria recovered produced an amplicon of appropriate
339 size with *Rahnella*-specific primers or produced *gyrB* 1480F/2242R amplicons with
340 identical sequences to those of the inoculated strains.

341 Additional inoculations were performed to compare pathogenicity of *Rahnella*
342 strains recovered from the USA and Norway. The results showed water-soaking and
343 discolouration (from light to dark brown); in some cases, scale shrinkage was observed.
344 In a side-by-side comparison of strains from USA and Norway, there were no significant
345 differences in virulence (Figure 3).

346

347 Phylogenetic analysis

348 In the routine course of identifying bacteria from onions, we generated sequence
349 for the *gyrB* 1480F/2242R amplicon from numerous strains of *Rahnella*. To assess the
350 utility of these sequences in identifying strains of *Rahnella* to species level, partial *gyrB*

351 sequences generated for verified strains of *R. aquatilis*, *R. victoriana*, *R. variigena*, *R.*
352 *inuitata*, *R. bruchi*, *R. woolbedingensis*, and *E. americana*, or downloaded from
353 GenBank. Phylogenetic trees were generated and showed that most isolates from
354 onions (originating from both North America and Europe) formed a group containing
355 three major clades. Six strains clustered tightly with *R. aquatilis*, 17 strains formed a
356 separate clade with *R. aquatilis* as its nearest neighbor, and 27 strains clustered with
357 *Rahnella* sp. Y9602 (Figure S1).

358 MLSA was performed on a subset of *Rahnella* strains isolated from the USA and
359 Norway to conclusively identify them. Strains were placed into context with different
360 *Rahnella* species based on sequence data available in GenBank (Brady *et al.* 2014),
361 using the MLSA scheme designed by Brady *et al.* (2008). Of the ten *Rahnella* strains
362 isolated from onions and used in MLSA, one strain (AR25a) clustered tightly with *R.*
363 *aquatilis*, three additional strains (L57-1-12, SL6, and A66) formed a separate clade
364 near *R. aquatilis*, four strains (L31-1-12, L172-1A, C1b, F57b) clustered with *Rahnella*
365 sp. Y9602, one strain (G37d) clustered with *R. victoriana* strains, and one (H11b) did
366 not cluster well with any of the reference strains. An additional strain (FC61912-K) was
367 isolated from a creek flowing adjacent to an onion field; it clustered loosely with *R.*
368 *inuitata* (Figure 1). Strains from onion that were represented in both the MLSA and
369 *gyrB* tree grouped to the same previously-characterized *Rahnella* strains in both trees.

370

371 ***Rahnella*-specific primers**

372 The Rah3783 primer pair produced amplicons of approximately 500 bp (expected
373 size 525 bp) from 22 of 23 *Rahnella* strains isolated from onions, including all strains

374 from the **clades** containing most onion isolates. Among these 23 strains, six were from
375 Norway, fourteen from New York, and three from Oregon. The strain (H11b) that did not
376 produce an amplicon with the Rah3783 primer pair did not cluster with the majority of
377 onion isolates and did not cluster tightly with any reference strains of *Rahnella*.
378 Additionally, reference strains from *R. aquatilis*, *R. victoriana*, *R. variigena*, *R. inusitata*,
379 *R. bruchi*, *R. woolbedingensis*, and *E. americana* were tested. All *Rahnella* strains
380 produced a fragment of the expected size except one of the two strains of *R.*
381 *woolbedingensis*. The *E. americana* strains did not produce a fragment (Figure 4, Table
382 2).

383 None of the 11 strains from other bacterial genera documented as onion
384 pathogens (*X. axonopodis* pv. *allii*, *P. viridiflava*, *P. carotovorum* subsp. *carotovorum*, *P.*
385 *ananatis*, *P. agglomerans*, *E. rhapontici*, *Enterobacter* sp., *D. dadantii*, *B. gladioli* pv.
386 *alliicola*, and *B. cepacia*) produced amplicons (Table 2).

387 The minimum amount of *Rahnella* sp. strain Y9602 that could be reliably
388 amplified using the Rah3783 primer pair was an average of 7,600 CFU/reaction. A ten-
389 fold dilution of that template yielded no band or only faintly discernible bands.

390 The PCR assay was also tested on onion samples from Norway that had varying
391 degrees of symptoms. Of 88 samples tested, 64 were positive, 5 were weakly positive
392 and 19 were negative for *Rahnella* spp. Samples with no symptoms **were used as**
393 **controls and** did not give any PCR product with the *Rahnella*-specific primers. The
394 assay successfully detected *Rahnella* sp. in onion samples, and hence may prove to be
395 a valuable tool for identification, detection and epidemiological studies of the bacterium.

396

397 Discussion

398 “Bacterial decay” in onions is an umbrella term describing onion bulb disease
399 symptoms consistent with bacterial infection, in the absence of detectable fungal or
400 insect problems. The symptoms caused by the various known bacterial decay
401 pathogens are not easily distinguishable, with many pathogens causing water-soaking
402 and discolouration of bulb scales and several causing maceration (Schwartz and
403 Mohan, 2007). Similar conditions are favorable for multiple bacterial decay pathogens,
404 such as wounded leaves, high relative humidity, free water, and high temperatures
405 (Schwartz and Mohan, 2007). Loss of plant tissue integrity associated with infection can
406 also make onions more vulnerable to additional colonization by secondary invaders
407 (Brewster, 2008). Finally, endophytic bacteria that may exist in relatively low numbers in
408 otherwise healthy onion bulbs may grow more rapidly in stressed or compromised
409 tissue, resulting in opportunistic infection (Cother and Dowling, 1986). Examples of
410 opportunistic bulb diseases are known: *Enterobacter* bulb decay and internal brown rot
411 of onions (caused by *Pseudomonas aeruginosa*) have been described as opportunistic
412 infections or as only occurring under special conditions. (Bishop and Davis 1990; Cother
413 *et al.* 1976)

414 Growers describe bacterial decays in growing onions and harvested bulbs as a
415 problem that has caused increasing losses in the last 15-20 years. The reasons behind
416 the increased losses are unknown but may involve a combination of factors, including
417 emergence of new pathogens, changing cultural practices, the introduction of new onion
418 cultivars, and changing climate. Because of the increased problems with bacterial
419 decays and because of the possibility of identifying emerging pathogens in onion-

420 growing regions, researchers in the USA and Norway separately investigated which
421 bacteria were commonly associated with diseased onion bulbs in their regions and
422 whether these commonly-detected bacteria represented substantial threats to onion
423 production.

424 *Rahnella* strains were some of the most commonly isolated bacteria from
425 diseased onion bulbs in both the USA and Norway. A subsequent survey of randomly
426 chosen onion bulbs from growers' storage revealed that *Rahnella* strains could be
427 isolated from both symptomatic and healthy-appearing bulbs. Recently, researchers in
428 Nova Scotia, Canada also detected *Rahnella* strains from both healthy and symptomatic
429 bulbs from growers' storage (Yurgel *et al.* 2018). In our study, the frequency with which
430 *Rahnella* strains were isolated from symptomatic bulbs was 17% versus only 1% for
431 healthy bulbs. The relatively greater abundance of *Rahnella* strains suggests a
432 relationship between the growth of *Rahnella* in onion bulbs and the presence of disease
433 symptoms. However, it was unclear whether *Rahnella* strains were involved in the
434 disease process directly or whether *Rahnella* strains are particularly capable of
435 colonizing or multiplying within diseased onion bulbs.

436 In this study, *Rahnella* strains were isolated from onion bulbs exhibiting a range
437 of symptoms, from mild discolouration of one or a few scales to water soaking and
438 maceration of entire bulbs. However, in laboratory inoculations of healthy-appearing
439 bulbs, pure cultures of *Rahnella* strains typically caused mild symptoms, indicating that
440 additional bacteria or fungi were probably responsible for the most severe symptoms in
441 bulbs from which *Rahnella* strains were isolated. *Rahnella* strains may therefore exist as
442 endophytes that are opportunistically pathogenic to onion bulbs, and their ability to

443 cause mild symptoms, including scale discolouration and shrinkage, may predispose
444 bulbs to disease caused by other pathogens. **Alternatively**, *Rahnella* strains may be
445 particularly adept at colonizing bulbs with symptoms caused by other pathogens. More
446 work is needed to tease apart these possibilities, which are not mutually exclusive.

447 A number of factors may influence the composition of the viable microbes in an
448 onion bulb, including **susceptibility of the host to infection**, inter-species competition,
449 antibiosis, and external environmental factors. Onion storage facilities are designed to
450 keep bulbs at low temperature, either by refrigeration, or by use of louvers that allow
451 cold winter air into the storage facility. Onion bulb storage at low temperature may be
452 particularly favorable for *Rahnella* strains compared to other bacteria. Strains of
453 *Rahnella* are considered psychrotrophic and have previously been described as
454 spoilage bacteria for foods stored under refrigerated conditions of 4-5°C (Jensen *et al.*
455 2001; Ercolini *et al.* 2006). During cold growing seasons or in growers' storage during
456 the winter months, *Rahnella* strains might be expected to survive or multiply better than
457 other bacteria, including virulent onion pathogens. The conditions under which onion
458 bulbs were stored may have contributed to the frequent isolation of *Rahnella* strains
459 from bulbs in this study.

460 In addition to being tolerant of a wide range of growth temperatures, strains of
461 the genus *Rahnella* are able to occupy many niches successfully. *Rahnella* spp. strains
462 have been isolated from many different substrates, including soil, water, insects, plants,
463 and people (Brady *et al.* 2014). Some species of *Rahnella* have also previously been
464 described as commonly associated with diseased plant tissue. *R. victoriana* in particular
465 is commonly associated with trees suffering from acute oak decline in the UK, but the

466 disease appears to be caused by a complex of species and the particular role of *R.*
467 *victoriana* in the disease is not clear (Denman *et al.* 2017). This acute oak decline
468 situation bears some resemblance to observations in this study, in which strains of
469 *Rahnella* were more commonly isolated as a component of the bulb microbiome from
470 diseased, rather than healthy plant tissue, yet *Rahnella* strains produced only mild
471 symptoms in pathogenicity tests in the laboratory. Indeed, the difficulty in isolating
472 known virulent pathogens from many symptomatic bulbs, and the existence of onion
473 diseases that become problematic under particular storage conditions and are caused
474 by bacteria that can frequently be isolated from healthy bulbs (for example,
475 *Enterobacter* bulb decay), suggests that bulb decays **may** sometimes **be** caused by
476 complexes of opportunistically pathogenic endophytic bacteria.

477 While multiple species of *Rahnella* were isolated from onions in the course of
478 these studies, the majority of strains belonged to a **monophyletic group consisting of**
479 **three clades** represented by the type strain of *R. aquatilis*, **the genome-sequenced**
480 **strain** *Rahnella* sp. Y9602, **and a branch that may represent an undescribed species**
481 **cluster**. This **group** of strains from onions collected in different years and from across
482 vast geographic distances suggests that these strains share features that allow
483 successful colonization and survival within onions that strains outside of this group lack.
484 This work suggests that this particular group of *Rahnella* strains, specifically *R. aquatilis*
485 and **two** closely-related **species**, have diseased onion bulbs as a niche. In the future,
486 comparison of genomes from onion-associated *Rahnella* strains might suggest suites of
487 genes involved in successful colonization of onion tissues. Primers developed in this

488 work should help to advance future studies by aiding in the rapid screening for onion-
489 associated *Rahnella* strains.

490

491 **Acknowledgements**

492 We are grateful to Jean Bonasera for technical support, including propagation of onion
493 plants. We are grateful to the Norwegian Agricultural Extension Service for providing
494 samples of onions; I.-L. Akselsen, Eva Borowski. E. Gauslå and M. Skogen for
495 assistance in sample preparation and PCR. The Norwegian study was supported by the
496 Research Council of Norway, and Norwegian onion growers. We would also like to
497 thank Robert Martinez and Carrie Brady for generous gifts of strains. Studies in the USA
498 were supported by the Onion Research and Development Program of New York State,
499 the New York Specialty Crops Block Grant Research Program, and the New York Farm
500 Viability Institute. We appreciate the gifts of onions from many growers in New York
501 State and the assistance of Cornell Cooperative Extension Associates who assisted in
502 gathering samples, and River Point Farms in Oregon.

503

504 **Conflict of Interest**

505 No conflict of interest declared.

506

507 **References**

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- 598

599

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Table 1 Primers used in this study

	Sequence	Annealing temperature used	Amplicon size	Reference
Primers for multilocus sequence analysis (MLSA)				
gyrB 01-Fs	TAA RTT YGA YGA YAA CTC YTA YAA AGT	45°C	971 bp	Brady <i>et al.</i> (2008)
gyrB 02-R	CMC CYT CCA CCA RGT AMA GTT			Brady <i>et al.</i> (2008)
infB 01-F	ATY ATG GGH CAY GTH GAY CA	48°C	1124 bp	Brady <i>et al.</i> (2008)
infB 02-R	ACK GAG TAR TAA CGC AGA TCC A			Brady <i>et al.</i> (2008)
rpoB 2522-2543 R	TCA GGC CCT AAC TTG GTG TCA C	53°C	1514 bp	this study
rpoB 1030-1049 F	GGC GCG TAC ATG TCC GAG AC			this study
atpD 922-943 R	GAG CGA AGG TGG TAG CTG GAG A	53°C	907 bp	this study
atpD 37-57 F	GTG GTG GAC GTC GAG TTC CCT			this study
atpD 58-81 F	CAG GAT GCA GTA CCG AAC GTG TAC	N/A	N/A	this study
atpD 900-921 R	TGG GTC AGT CAA GTC ATC CGC A	N/A	N/A	this study
rpoB 1307-1325 F	GTA ACG GCC AGG GCG AAG T	N/A	N/A	this study
rpoB 2138-2159 R	CGT TTG GCT ACG GCA GTC ACA C	N/A	N/A	this study
infB 1236-1257 F	CTC ATT GCT TGA CTA CAT TCG T	N/A	N/A	this study
infB 2092-2116 R	CCT GAA CGT CTG ACT TCA GAA CAA T	N/A	N/A	this study
Primers to amplify <i>gyrB</i> fragments for preliminary identification of strains				
gyrB 1480F	GGC ATC ATC ATC ATG ACC GA	50°C	788 bp	Bonasera <i>et al.</i> (2014)
gyrB 2242R	GTS GTT TCC CAS AGC TG			Bonasera <i>et al.</i> (2014)
<i>Rahnella</i>-specific primers				
Rah 3783 F1	CGG GAT CGT CCG TTA TAA AGG CA	57°C	524 bp	this study
Rah 3783 R1	ACG GTG CGT CCG TTC AGA TCA CC			this study
16S rDNA primers				
F985PTO	AAC GCG AAG AAC CTT AC	55°C	434 bp	Heuer <i>et al.</i> (1999) - modified
R1378	CGG TGT GTA CAA GGC CCG GGA ACG			Heuer <i>et al.</i> (1999)

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602

603 **Table 2** Detection of strains with *Rahnella*-specific primers

Species	Strain	Isolated from	Received from / Reference	Rah 3783 F1/R1 amplicon
<i>Rahnella</i> strains isolated from onions				
<i>Rahnella</i> sp.	A66	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	A78	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	AG1a	isolated from onion tplant, NY USA	this study	+
<i>Rahnella</i> sp.	AP10b	isolated from onion plant NY, USA	this study	+
<i>Rahnella</i> sp.	AR16b	isolated from onion bulb grown in OR, USA	this study	+
<i>Rahnella</i> sp.	AR20	isolated from onion bulb grown in OR, USA	this study	+
<i>Rahnella</i> sp.	AR25a	isolated from onion bulb grown in OR, USA	this study	+
<i>Rahnella</i> sp.	C10	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	C1b	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	E32Ma	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	F57b	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	G37d	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	G4	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	G42	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	H11b	culled onion from storage NY, USA	this study	-
<i>Rahnella</i> sp.	H23	culled onion from storage NY, USA	this study	+
<i>Rahnella</i> sp.	I50b	freshly harvested onion bulb NY, USA	this study	+
<i>Rahnella</i> sp.	L151-1a	onion from county of Østfold, Norway	this study	+
<i>Rahnella</i> sp.	L172-1A	onion from county of Vestfold, Norway	this study	+
<i>Rahnella</i> sp.	L173-1B	onion from county of Vestfold, Norway	this study	+
<i>Rahnella</i> sp.	L31-1-12	onion from county of Vestfold, Norway	this study	+
<i>Rahnella</i> sp.	L57-1-12	onion from county of Oppland, Norway	this study	+
<i>Rahnella</i> sp.	SL6	onion from county of Hedmark, Norway	this study	+
<i>Rahnella</i> strains isolated from other sources				
<i>Rahnella</i> sp.	FC61912-K	Creek water, NY, USA	this study	+
<i>R. victoriana</i>	FRB 225 ^T	<i>Quercus robur</i> , symptomatic inner bark, Suffolk, UK	Brady <i>et al.</i> (2014)	+
<i>R. victoriana</i>	USA 13	<i>Quercus kelloggii</i> , symptomatic inner bark, California, USA	Brady <i>et al.</i> (2014)	+
<i>R. variigena</i>	FOD 20/8	<i>Quercus robur</i> , wound response fluid, Gloucestershire, UK	Brady <i>et al.</i> (2014)	+
<i>R. variigena</i>	PFK 1/1C2a	<i>Quercus robur</i> , symptomatic inner bark, Sussex, UK	Brady <i>et al.</i> (2014)	+
<i>R. inusitata</i>	FOD 9/5a	<i>Quercus robur</i> , symptomatic inner bark, Gloucestershire, UK	Brady <i>et al.</i> (2014)	+
<i>R. inusitata</i>	FOD 9/21	<i>Quercus robur</i> , symptomatic inner bark, Gloucestershire, UK	Brady <i>et al.</i> (2014)	+
<i>R. bruchi</i>	FRB 226 ^T	<i>Agrilus biguttatus</i> , gut, Shropshire, UK	Brady <i>et al.</i> (2014)	+

<i>R. bruchi</i>	ALN 45	<i>Alnus glutinosa</i> , inner bark, Surrey, UK	Brady <i>et al.</i> (2014)	+
<i>R. woolbedingensis</i>	FRB 227 ^T	<i>Alnus glutinosa</i> , inner bark, Surrey, UK	Brady <i>et al.</i> (2014)	+
<i>R. woolbedingensis</i>	WAL 10	<i>Juglans regia</i> , inner bark, Surrey, UK	Brady <i>et al.</i> (2014)	-
Other bacteria				
<i>Ewingella americana</i>	FOD 24/3b	<i>Quercus robur</i> , symptomatic inner bark, Gloucestershire, UK	Brady <i>et al.</i> (2014)	-
<i>Ewingella americana</i>	AT 14b	<i>Quercus robur</i> , symptomatic inner bark, Shropshire, UK	Brady <i>et al.</i> (2014)	-
<i>Burkholderia cepacia</i>	ATCC 25416	Onion, 1948	type strain of <i>B. cepacia</i>	-
<i>Burkholderia gladioli</i> pv. <i>allii</i> cola	ATCC 19302	Onion bulb rot, USA	type strain of <i>B. gladioli</i>	-
<i>Dickeya dadantii</i>	Dickey 151			-
<i>Enterobacter</i> sp.	EcWSU1	Onion, USA	Humann <i>et al.</i> (2011)	-
<i>Erwinia rhapontici</i>	ATCC 29283	Rhubarb, England	type strain of <i>E. rhapontici</i>	-
<i>Pantoea agglomerans</i>	SUH1	Onion, South Africa	Hattingh and Walters (1981)	-
<i>Pantoea agglomerans</i>	ATCC 27155	Knee laceration	type strain of <i>P. agglomerans</i>	-
<i>Pantoea ananatis</i>	ATCC 33244; LMG 2665	Pineapple, Brazil	type strain of <i>P. ananatis</i>	-
<i>Pectobacterium carotovorum</i>	ATCC 15713	Potato, Denmark	type strain of <i>P. carotovorum</i>	-
<i>Pseudomonas viridiflava</i>	LMG 2352	Dwarf or runner bean, Switzerland	type strain of <i>P. viridiflava</i>	-
<i>Rahnella aquatilis</i>	ATCC 33071	Drinking water, France	type strain of <i>R. aquatilis</i>	-
<i>Xanthomonas axonopodis</i> pv. <i>allii</i>	O274	Onion, CO, USA	H. Schwartz, Colorado State University	-

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607 **Figure 1** Multilocus sequence analysis tree of 11 *Rahnella* strains isolated from onion or
608 a creek running adjacent to an onion field shown in context with published *Rahnella*
609 strains. Strains isolated from onions are highlighted in yellow. The tree with the highest
610 log likelihood (-11128.21) is shown. The percentage of trees in which the associated
611 taxa clustered together is shown next to the branches. The tree is drawn to scale, with
612 branch lengths measured in the number of substitutions per site. The analysis involved
613 27 strains. There were a total of 2635 positions in the final dataset. *Xenorhabdus*
614 *nematophila* ATCC is used as an outgroup. Analysis was performed with concatenated
615 sequences from *gyrB* (gyrase subunit B gene), *rpoB* (RNA polymerase β subunit), *infB*
616 (translation initiation factor IF-2), and *atpD* (ATP synthase subunit beta) genes. There
617 were a total of 2636 positions in the dataset. N: Isolated from onions in Norway; OR:
618 Isolated from onions grown in OR, USA; NY: Isolated from onions grown in NY, USA;
619 W: Isolated from creek water adjacent to onion field, NY, USA.

620 **Figure 2** Five onion bulbs each were inoculated with 8 *Rahnella* strains recovered from
621 Norway and the USA. Symptoms were generally mild. Strains from Norway (L31-1-12,
622 L57-1-12, SL6, and L172-1a,) and USA (A66, AR25a, F57b, and C1b).

623 **Figure 3** Symptoms in bulbs from experiments with strains from the USA and Norway.
624 A. Three scoring categories were established (from left to right): 0 = no symptoms; 1 =
625 weak discolouration; 2 = darker discolouration and scale shrinkage. B. Results from
626 pathogenicity test of isolates from the USA and Norway.

627 **Figure 4** Example of agarose gel with PCR products amplified using *Rahnella* specific
628 primers Rah 3783 F1/R1. L: 2-log ladder (New England Biolabs), 1: *R. victoriana* FRB

629 225T, 2: *R. victoriana* USA 13, 3: *R. variigena* FOD 20/8, 4: *R. variigena* PFK 1/1C2a, 5:
630 *R. inusitata* FOD 9/5a, 6: *R. inusitata* FOD 9/21, 7: *R. bruchi* FRB 226T, 8: *R. bruchi*
631 ALN 45, 9: *R. woolbedingensis* FRB 227T, 10: *R. woolbedingensis* WAL 10, 11: *E.*
632 *americana* FOD 24/3b, 12: *E. americana* AT 14b, 13: *Rahnella* sp. C1b, 14: water
633 control.

634

635 Supporting Information

636 **Table S1** Accession numbers for partial gyrase B sequence derived using the *gyrB*
637 1480F/2242R primers

638 **Table S2** Accession numbers for sequences used in multilocus sequence analysis

639 **Table S3** Numbers of bulbs from which particular genera of bacteria were recovered
640 from surveys of diseased and healthy onion bulbs in USA

641 **Figure S1** Samples of bulbs from which *Rahnella* strains were recovered in screen of
642 random bulbs from growers' cold storage in NY. A. Examples of bulbs from which only
643 *Rahnella* strains were recovered. B. Examples of bulbs from which both *Rahnella*
644 strains and other bacteria were recovered.

645 **Figure S2** Maximum Likelihood tree using partial *gyrB* sequence. Strains isolated from
646 onion are highlighted in yellow. The percentage of trees in which the associated taxa
647 clustered together is shown next to the branches. The tree is drawn to scale, and the
648 units of branch lengths are the number of substitutions per site. The analysis involved
649 nucleotide sequences from 71 strains. There were a total of 625 positions in the final
650 dataset. N: Isolated from onions in Norway; OR: Isolated from onions grown in OR,

651 USA; NY: Isolated from onions grown in NY, USA; W: Isolated from creek water
652 adjacent to onion field, NY, USA.

653 **Figure S3** Bulbs syringe-inoculated with water (A), *Rahnella* sp. C1b (B), and
654 *Enterobacter* sp. EcWSU1 (C). Symptoms elicited by *Rahnella* are typically mild. The
655 bulbs presented in (B) had severe symptoms compared to other repetitions of the assay
656 (see Figure 2). The reasons for between-assay variations in severity are unknown but
657 could be due to variations in host susceptibility due to bulb age or genotype. Variation
658 in the symptom severity between assays adds to the difficulties in assessing the real-
659 world impacts of *Rahnella* spp. bacteria on onion production. Strains inoculated with
660 *Enterobacter* sp. EcWSU1 are included for comparison with a known opportunistic
661 pathogen of onion bulb.

662

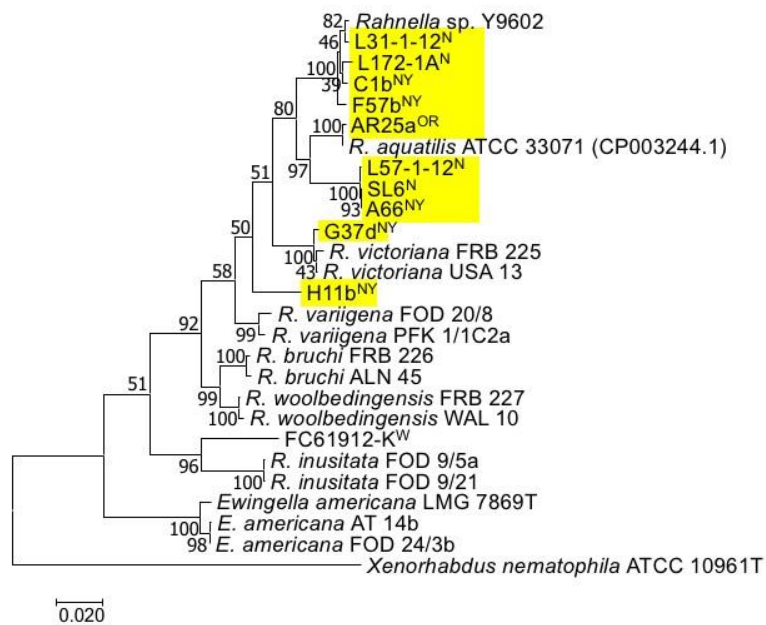


Figure 1 Multilocus sequence analysis tree of 11 *Rahnella* strains isolated from onion or a creek running adjacent to an onion field shown in context with published *Rahnella* strains. Strains isolated from onions are highlighted in yellow. The tree with the highest log likelihood (-11128.21) is shown. The percentage of trees in which the associated taxa clustered together is shown next to the branches. The tree is drawn to scale, with branch lengths measured in the number of substitutions per site. The analysis involved 27 strains. There were a total of 2635 positions in the final dataset. *Xenorhabdus nematophila* ATCC is used as an outgroup. Analysis was performed with concatenated sequences from *gyrB* (gyrase subunit B gene), *rpoB* (RNA polymerase β subunit), *infB* (translation initiation factor IF-2), and *atpD* (ATP synthase subunit beta) genes. There were a total of 2636 positions in the dataset. N: Isolated from onions in Norway; OR: Isolated from onions grown in OR, USA; NY: Isolated from onions grown in NY, USA; W: Isolated from creek water adjacent to onion field, NY, USA.

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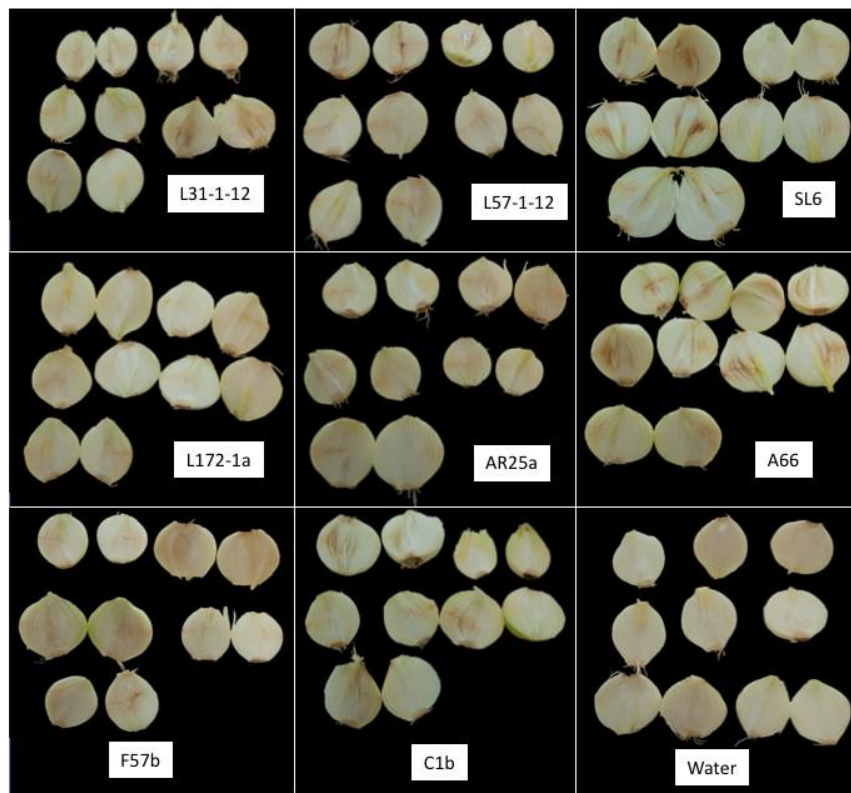


Figure 2 Five onion bulbs each were inoculated with 8 *Rahnella* strains recovered from Norway and the USA. Symptoms were generally mild. Strains from Norway (L31-1-12, L57-1-12, SL6, and L172-1a,) and USA (A66, AR25a, F57b, and C1b).

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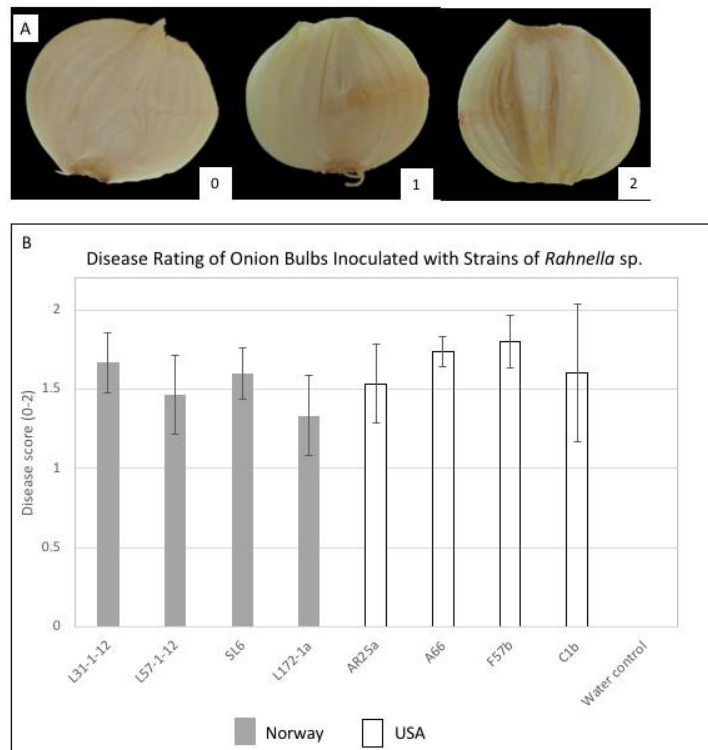


Figure 3 Symptoms in bulbs from experiments with strains from the USA and Norway. A. Three scoring categories were established (from left to right): 0 = no symptoms; 1 = weak discoloration; 2 = darker discoloration and scale shrinkage. B. Results from pathogenicity test of isolates from the USA and Norway.

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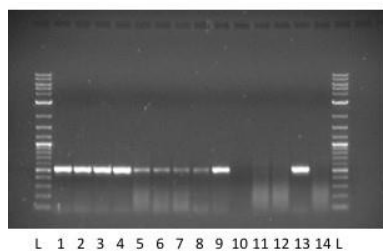


Figure 4 Example of agarose gel with PCR products amplified using *Rahnella* specific primers Rah 3783 F1/R1. L: 2-log ladder (New England Biolabs), 1: *R. victoriana* FRB 225T, 2: *R. victoriana* USA 13, 3: *R. variigena* FOD 20/8, 4: *R. variigena* PFK 1/1C2a, 5: *R. inusitata* FOD 9/5a, 6: *R. inusitata* FOD 9/21, 7: *R. bruchi* FRB 226T, 8: *R. bruchi* ALN 45, 9: *R. woolbedingensis* FRB 227T, 10: *R. woolbedingensis* WAL 10, 11: *E. americana* FOD 24/3b, 12: *E. americana* AT 14b, 13: *Rahnella* sp. C1b, 14: water control.

190x254mm (96 x 96 DPI)

Table S1 Accession numbers for partial gyrase B sequence derived using the gyrB 1480F/2242R primers

Species	Strain	Isolated from	Received from / Reference	Accession
<i>Rahnella</i> sp.	Y9602	soil at U.S. Dept. of Energy Oak Ridge Reservation in Oak Ridge, TN	Martinez <i>et al.</i> (2007)	CP002505.1
<i>Rahnella</i> sp.	Q73b	symptomatic red onion bulb from cold storage NY, USA	this study	MK391682
<i>Rahnella</i> sp.	R27c	symptomatic yellow onion bulb recovered from storage NY, USA	this study	MK391683
<i>Rahnella</i> sp.	R92a	symptomatic yellow onion bulb from storage NY, USA	this study	MK391684
<i>Rahnella</i> sp.	T11a	symptomatic yellow onion bulb from storage NY, USA	this study	MK391685
<i>Rahnella</i> sp.	T100a	symptomatic yellow onion bulb from storage NY, USA	this study	MK391686
<i>Rahnella</i> sp.	A12a	culled red onion from storage NY, USA	this study	MK391687
<i>Rahnella</i> sp.	A66	culled onion from storage NY, USA	this study	MK391688
<i>Rahnella</i> sp.	A78	culled onion from storage NY, USA	this study	MK391689
<i>Rahnella</i> sp.	AG6b	symptomatic bulb tissue from growing onion NY, USA	this study	MK391690
<i>Rahnella</i> sp.	AR20	onion bulb grown in OR, USA	this study	MK391691
<i>Rahnella</i> sp.	AR25a	onion bulb grown in OR, USA	this study	MK391692
<i>Rahnella</i> sp.	B18	culled onion from storage NY, USA	this study	MK391693
<i>Rahnella</i> sp.	C1b	culled onion from storage NY, USA	this study	MK391694
<i>Rahnella</i> sp.	C10	culled onion from storage NY, USA	this study	MK408759
<i>Rahnella</i> sp.	D36	culled onion from storage NY, USA	this study	MK391695
<i>Rahnella</i> sp.	E32Ma	culled onion from storage NY, USA	this study	MK391696
<i>Rahnella</i> sp.	F30a	culled onion from storage NY, USA	this study	MK391697

<i>Rahnella</i> sp.	F35b	culled onion from storage NY, USA	this study	MK391698
<i>Rahnella</i> sp.	F57b	culled onion from storage NY, USA	this study	MK391699
<i>Rahnella</i> sp.	FC61912- K	creek water, NY, USA	this study	MK391700
<i>Rahnella</i> sp.	G4	culled onion from storage NY, USA	this study	MK391701
<i>Rahnella</i> sp.	G29a	culled onion from storage NY, USA	this study	MK391703
<i>Rahnella</i> sp.	G33b	culled onion from storage NY, USA	this study	MK391704
<i>Rahnella</i> sp.	G37d	culled onion from storage NY, USA	this study	MK391705
<i>Rahnella</i> sp.	G42	culled onion from storage NY, USA	this study	MK391702
<i>Rahnella</i> sp.	G43a1	culled onion from storage NY, USA	this study	MK391706
<i>Rahnella</i> sp.	H11b	culled onion from storage NY, USA	this study	MK391707
<i>Rahnella</i> sp.	H23	culled onion from storage NY, USA	this study	MK391708
<i>Rahnella</i> sp.	I50b	freshly harvested symptomatic onion bulb NY, USA	this study	MK391709
<i>Rahnella</i> sp.	J9a	non-symptomatic yellow onion from storage NY, USA	this study	MK391710
<i>Rahnella</i> sp.	J55b	non-symptomatic yellow onion from storage NY, USA	this study	MK391711
<i>Rahnella</i> sp.	K60d	symptomatic red onion from storage NY, USA	this study	MK391712
<i>Rahnella</i> sp.	L50a	symptomatic red onion from storage NY, USA	this study	MK391713
<i>Rahnella</i> sp.	L52e	symptomatic red onion from storage NY, USA	this study	MK391714
<i>Rahnella</i> sp.	L54a	symptomatic yellow onion from storage NY, USA	this study	MK391715
<i>Rahnella</i> sp.	L70b	symptomatic yellow onion from storage NY, USA	this study	MK391716
<i>Rahnella</i> sp.	L72c	symptomatic yellow onion from storage NY, USA	this study	MK391717
<i>Rahnella</i> sp.	N27b	non-symptomatic yellow onion from storage NY, USA	this study	MK391718

<i>Rahnella</i> sp.	N81a	symptomatic yellow onion from storage NY, USA	this study	MK391719
<i>Rahnella</i> sp.	N89b	symptomatic yellow onion from storage NY, USA	this study	MK391720
<i>Rahnella</i> sp.	L18a	symptomatic yellow onion from storage NY, USA	this study	MK391721
<i>Rahnella</i> sp.	AG1a	symptomatic onion transplant, NY USA	this study	MK391722
<i>Rahnella</i> sp.	P36c	symptomatic red onion bulb from storage NY, USA	this study	MK391723
<i>Rahnella</i> sp.	AP10b	symptomatic onion plant NY, USA	this study	MK391724
<i>Rahnella</i> sp.	AP11b	symptomatic onion plant NY, USA	this study	MK391725
<i>Rahnella</i> sp.	AR16b	symptomatic onion bulb grown in OR, USA	this study	MK391726
<i>Rahnella</i> sp.	C60	symptomatic red onion bulb from storage NY, USA	this study	MK391727
<i>Rahnella</i> sp.	C81a	symptomatic red onion bulb from storage NY, USA	this study	MK391728
<i>Rahnella</i> sp.	L31-1-12	onion from county of Vestfold, Norway	this study	MK391741
<i>Rahnella</i> sp.	L57-1-12	onion from county of Oppland, Norway	this study	MK391742
<i>Rahnella</i> sp.	L151-1a	onion from county of Østfold, Norway	this study	MK391743
<i>Rahnella</i> sp.	L172-1A	onion from county of Vestfold, Norway	this study	MK391744
<i>Rahnella</i> sp.	L173-1B	onion from county of Vestfold, Norway	this study	MK391745
<i>Rahnella</i> sp.	SL6	onion from county of Hedmark, Norway	this study	MK391746
<i>Rahnella aquatilis</i>	ATCC 33071	drinking water, France (type strain)	Martinez <i>et al.</i> (2012a)	CP003244.1
<i>Rahnella victoriana</i>	USA13	<i>Quercus kelloggii</i> , symptomatic inner bark, California, USA	Brady <i>et al.</i> (2014)	MK391729
<i>Rahnella bruchi</i>	ALN 45	<i>Alnus glutinosa</i> , inner bark, Surrey, UK	Brady <i>et al.</i> (2014)	MK391730
<i>Rahnella bruchi</i>	FRB 226 ^T	<i>Agrilus biguttatus</i> , gut, Shropshire, UK	Brady <i>et al.</i> (2014)	MK391740
<i>Rahnella woolbedinensis</i>	FRB 227	<i>Alnus glutinosa</i> , inner bark, Surrey, UK	Brady <i>et al.</i> (2014)	MK391731

<i>Rahnella woolbedingensis</i>	WAL 10	<i>Juglans regia</i> , inner bark, Surrey, UK	Brady <i>et al.</i> (2014)	MK391732
<i>Ewingella americana</i>	AT 14b	<i>Quercus robur</i> , symptomatic inner bark, Shropshire, UK	Brady <i>et al.</i> (2014)	MK391733
<i>Ewingella americana</i>	FOD 24/3b	<i>Quercus robur</i> , symptomatic inner bark, Gloucestershire, UK	Brady <i>et al.</i> (2014)	MK391735
<i>Ewingella americana</i>	ATCC 33852	human throat, KY, USA (type strain)		JMPJ01000067.1
<i>Rahnella victoriana</i>	FRB 225	<i>Quercus robur</i> , symptomatic inner bark, Suffolk, UK	Brady <i>et al.</i> (2014)	MK391736
<i>Rahnella variigena</i>	FOD 20/8	<i>Quercus robur</i> , wound response fluid, Gloucestershire, UK	Brady <i>et al.</i> (2014)	MK391737
<i>Rahnella variigena</i>	PFK 1/1C2a	<i>Quercus robur</i> , symptomatic inner bark, Sussex, UK	Brady <i>et al.</i> (2014)	MK391738
<i>Rahnella inusitata</i>	FOD 9/21	<i>Quercus robur</i> , symptomatic inner bark, Gloucestershire, UK	Brady <i>et al.</i> (2014)	MK391739
<i>Rahnella inusitata</i>	FOD 9/5a	<i>Quercus robur</i> , symptomatic inner bark, Gloucestershire, UK	Brady <i>et al.</i> (2014)	MK391734
<i>Xenorhabdus nematophila</i>	ATCC 19061			FN667742.1

Brady, C., Hunter, G., Kirk, S., Arnold, D. and Denman, S. (2014) *Rahnella victoriana* sp. nov., *Rahnella bruchi* sp. nov., *Rahnella woolbedingensis* sp. nov., classification of *Rahnella* genomospecies 2 and 3 as *Rahnella variigena* sp. nov. and *Rahnella inusitata* sp. nov., respectively and emended description of the genus *Rahnella*. *Syst Appl Microbiol* **37**, 545-52.

Martinez, R. J., Beazley, M. J., Taillefert, M., Arakaki, A. K., Skolnick, J. and Sobecky, P. A. (2007) Aerobic uranium (VI) bioprecipitation by metal-resistant bacteria isolated from radionuclide- and metal-contaminated subsurface soils. *Environ Microbiol* **9**, 3122-3133.

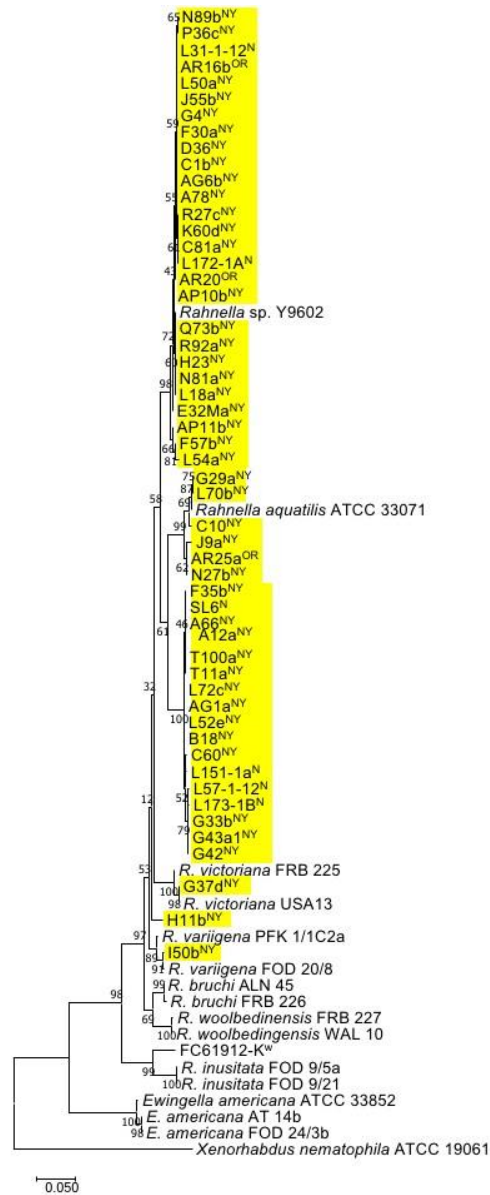
Martinez, R. J., Bruce, D., Detter, C., Goodwin, L. A., Han, J., Han, C. S., Held, B., Land, M. L., Mikhailova, N., Nolan, M., Pennacchio, L., Pitluck, S., Tapia, R., Woyke, T. and Sobecky, P. A. (2012a) Complete genome sequence of *Rahnella aquatilis* CIP 78.65. *J Bacteriol* **194**, 3020-3021.

Table S2 Accession numbers for sequences used in multilocus sequence analysis

	Strain	Source	<i>gyrB</i>	<i>rpoB</i>	<i>infB</i>	<i>atpD</i>
<i>Rahnella</i> sp.	A66	culled onion from storage NY, USA	MK387415	MK387404	MK387393	MK387382
<i>Rahnella</i> sp.	AR25a	onion bulb grown in OR, USA	MK387416	MK387405	MK387394	MK387383
<i>Rahnella</i> sp.	C1b	culled onion from storage NY, USA	MK387417	MK387406	MK387395	MK387384
<i>Rahnella</i> sp.	F57b	culled onion from storage NY, USA	MK387418	MK387407	MK387396	MK387385
<i>Rahnella</i> sp.	FC61912-K	river water, NY, USA	MK387419	MK387408	MK387397	MK387386
<i>Rahnella</i> sp.	G37d	culled onion from storage NY, USA	MK387420	MK387409	MK387398	MK387387
<i>Rahnella</i> sp.	H11b	culled onion from storage NY, USA	MK387421	MK387410	MK387399	MK387388
<i>Rahnella</i> sp.	L31-1-12	onion from county of Vestfold, Norway	MK387422	MK387411	MK387400	MK387389
<i>Rahnella</i> sp.	L57-1-12	onion from county of Oppland, Norway	MK387423	MK387412	MK387401	MK387390
<i>Rahnella</i> sp.	L172-1A	onion from county of Vestfold, Norway	MK387424	MK387413	MK387402	MK387391
<i>Rahnella</i> sp.	SL6	onion from county of Hedmark, Norway	MK387425	MK387414	MK387403	MK387392

Table S3 Genera recovered from surveys of diseased and healthy onion bulbs in USA

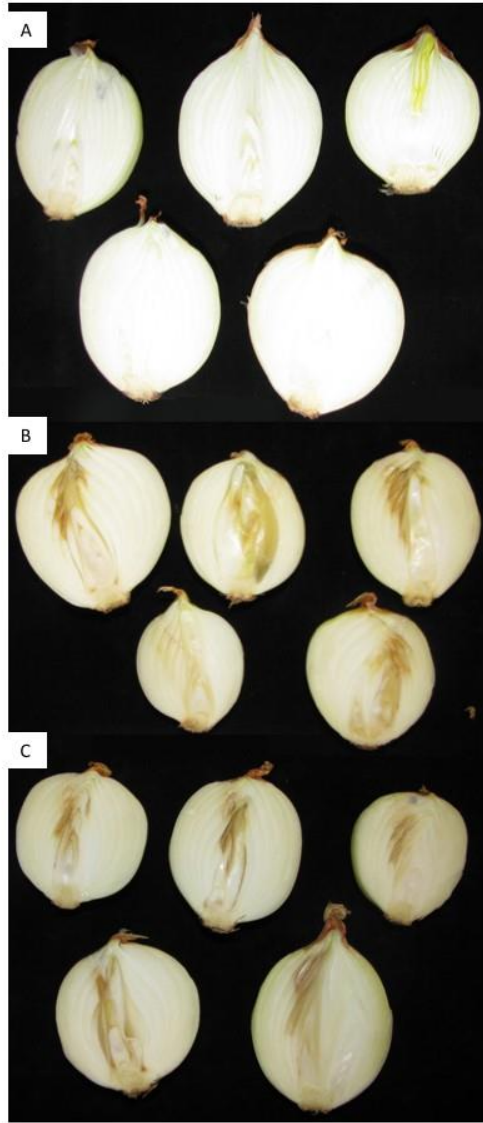
	Cull Survey		Random Survey	
	Symptomatic	Non-Symptomatic	Symptomatic	Non-Symptomatic
Total Number of Bulbs	508		898	
Total Number of Bulbs	477	31	150	748
No bacteria isolated	63	21	12	564
<i>Rahnella</i> spp.	136	0	25	9
<i>Enterobacter</i> spp.	62	1	47	34
<i>Pseudomonas</i> spp.	138	2	53	28
<i>Burkholderia</i> spp.	52	0	31	77
<i>Pantoea</i> spp.	36	0	41	55
Bacteria co-isolated with <i>Rahnella</i>				
Nothing or multiple <i>Rahnella</i> strains	55	0	7	2
Unidentified bacteria only	26	0	1	2
<i>Pseudomonas</i> spp. only	23	0	7	0
<i>Enterobacter</i> spp. only	6	0	1	1
<i>Pantoea</i> spp. only	2	0	1	0
<i>Burkholderia</i> spp. only	1	0	0	2
<i>Citrobacter</i> sp.	0	0	2	0
<i>Klebsiella</i> sp.	0	0	1	0
Multiple other bacteria Isolated	5	0	5	2



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190x254mm (96 x 96 DPI)



190x254mm (96 x 96 DPI)