

1 **Effect of copy number of the *spoVA*^{2mob} operon, sourdough and reutericyclin on**
2 **ropy bread spoilage caused by *Bacillus* spp.**

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21 **ABSTRACT**

22 *Bacillus* spp. cause ropy bread spoilage of bread, which is characterized by a rotten
23 fruity odor, followed by discoloration and degradation of the crumb. *Bacillus* spp. are
24 wheat grain endophytes and form heat resistant endospores, therefore, process hygiene
25 and heating during baking do not prevent ropy spoilage. This study used 8 strains of
26 *Bacillus subtilis* and *Bacillus amyloliquefaciens* to determine whether the presence and
27 the copy number of *spoVA^{2mob}* operon influences survival after baking; in addition, the
28 spoilage phenotype was correlated with the presence of amylolytic enzymes in genomes
29 of *Bacillus* spp.. The presence and copy number of the *spoVA^{2mob}* operon had only a
30 minor effect on survival of *Bacillus* endospores. Strains of *B. amyloliquefaciens* caused
31 ropy spoilage faster than strains of *B. subtilis*, this difference correlated to the number
32 and type of extracellular amylases encoded in the genomes of the strains of
33 *B. amyloliquefaciens* and *B. subtilis*. The inhibitory effect of sourdough on ropy
34 spoilage was determined by addition of 3 – 24% sourdough fermented with *L. reuteri*
35 TMW1.656. Addition of 12% and 24% sourdough, corresponding to a bread pH of
36 5.93 ± 0.041 and 5.53 ± 0.040 , respectively, delayed ropy spoilage for 2 and more than
37 5 d, respectively. The comparison of addition of 12% sourdough fermented with the
38 reutericyclin producing *L. reuteri* TMW1.656 and the isogenic reutericyclin-negative
39 strain *L. reuteri* TMW1.656 Δ gtfA Δ rtcN demonstrated that reutericyclin produced in
40 sourdough inhibits growth of *Bacillus* in bread. In conclusion, sourdough inhibits
41 germination of *Bacillus* spores in bread and the effect of sourdough is enhanced by
42 reutericyclin.

43 **KEY WORDS:** *Bacillus* spp., Ropy spoilage, Extracellular amylase, Sourdough,
44 Reutericyclin

45 1. INTRODUCTION

46 Bread or steamed bread is a staple of the daily diet in most regions of the world with a
47 temperate climate (Kourkouta et al., 2017). Ropy spoilage of bread is caused by strains
48 of *Bacillus* spp. including *B. amyloliquefaciens*, *B. subtilis* and *B. licheniformis* (Pepe
49 et al., 2003; Sorokulova et al., 2003; Valerio et al., 2015, 2012). *Bacillus* endospores
50 are present in flour and survive after baking. Their growth and production of amylases
51 and proteases initially generate a typical fruity odor, subsequently result in a sticky and
52 stringy crumb and slime formation, making the bread inedible (Corsetti et al., 2000;
53 Pepe et al., 2003).

54 *Bacillus* endospores have been isolated from bakery environment and also from raw
55 materials, such as wheat flour, yeast and bread improvers (Bailey and von Holy, 1993;
56 Pepe et al., 2003). Contamination of wheat flour with *Bacillus* endospores relates to the
57 stable occurrence of these organisms as part of commensal microbiota of plants
58 including wheat (Chen et al., 2007; Fan et al., 2011; Jamal et al., 2019; Tilak and Reddy,
59 2006; Vessey and Buss, 2002), and to the presence of *Bacillus* as member of seed-borne
60 endophytic microbial community (Robinson et al., 2016; Shahzad et al., 2016). *Bacillus*
61 endospores remain viable throughout storage and processing of grains (Fangio et al.,
62 2010; Needham et al., 2005) which makes the contamination of flour not accidental but
63 unavoidable.

64 Endospores of *Bacillus* species are heat resistant (Setlow, 2006, 2003) and not
65 inactivated during baking, where the crumb is heated to a maximum temperature of
66 100 °C for a few minutes. Strains of *B. amyloliquefaciens*, *B. subtilis* and *B. pumilus*
67 were reported to exhibit comparable heat resistance, amylase activity and spoilage
68 ability (Setlow, 2006; Valerio et al., 2012). However, a large intra-species variation of
69 heat resistance of *Bacillus* endospores was attributed to the presence the *spoVA*^{2mob}

70 operon (Berendsen et al., 2016). Strains of *B. amyloliquefaciens* and *B. subtilis* contain
71 up to three copies of the *spoVA*^{2mob} operons and the heat resistance of spores increased
72 with an increase of the copy number (Berendsen et al., 2016; Wang et al., 2018). A
73 relationship between the *spoVA*^{2mob} operon and the potential to cause ropy spoilage is
74 further suggested by the observation that all strains of *B. subtilis* and
75 *B. amyloliquefaciens* that were isolated by Röcken and Spicher (1993) from ropy bread
76 were later identified to harbor multiple copies of the *spoVA*^{2mob} operons (Li et al., 2019).
77 A contribution of the *spoVA*^{2mob} operon to the ability of bacilli to survive baking,
78 however, has not been determined experimentally.

79 Ropy spoilage of bread develops rapidly at warm and humid conditions (Vaičiulytė-
80 Funk et al., 2015), and is prevented by chemical preservatives (Saranraj and Geetha,
81 2012) or the use of sourdough. In past years, the use of sourdough or sourdough
82 products in baking has been re-established as the default method of bread production
83 (Anonymous, 2019a, 2019b). Sourdough prevents ropy spoilage (Pepe et al., 2003;
84 Sadeghi, 2008) through acidification and the inhibitory activity of undissociated
85 organic acids (Rosenquist and Hansen, 1998).

86 Sourdough isolates of lactic acid bacteria were reported to produce bacteriocins with
87 activity against rope-forming bacilli (Digaitiene et al., 2012; Pepe et al., 2003), however,
88 their activity in bread is limited (Rosenquist and Hansen, 1998), likely because the
89 bacteriocins are inactivated by proteases and thiol exchange reactions in sourdough
90 (Gänzle, 2014). Some strains of *Limosilactobacillus reuteri*, previously named
91 *Lactobacillus reuteri* (Zheng et al., 2020), are stable components of type II sourdoughs
92 (Zheng et al., 2016) and produce reutericyclin, a heat stable and antimicrobial tetramic
93 acid with activity against a wide range of gram-positive bacteria including rope-forming
94 bacilli (Gänzle et al., 2000). Reutericyclin is produced in active concentration in

95 sourdough fermentation (Gänzle and Vogel, 2003), but a possible contribution to bread
96 preservation has not been demonstrated. Therefore, the aim of this study was to
97 determine whether ropy spoilage of bread is inhibited by the use of sourdough
98 fermented with reutericyclin-producing *L. reuteri* TMW 1.656. To identify relevant
99 spoilage organisms, strains of *Bacillus* spp. that were previously isolated from ropy
100 bread or other cereals products were characterized with respect to the copy number of
101 the spoVA^{2mob} operon, their heat resistance during the baking process, and with respect
102 to the presence of extracellular amylases that may relate to the spoilage phenotype.

103 **2. MATERIALS AND METHODS**

104 **2.1. Bacterial strains and culture conditions**

105 The strains used in this study and their origin are listed in Table 1. Strains were
106 maintained at -80 °C in 20% glycerol. Strains of *Limosilactobacillus reuteri* were
107 incubated in modified de Man, Rogosa, and Sharpe (mMRS, Gänzle et al., 2000)
108 medium at 37 °C for 24 h. The mutant strain of *L. reuteri* TMW1.656 Δ rtcN Δ gtfA was
109 constructed by disruption of the reuteransucrase GtfA in *L. reuteri* TMW1.656 Δ rtcN
110 as described (Chen et al., 2016; Lin et al., 2015). Strains of *Bacillus* were grown
111 aerobically on Luria-Bertani (LB) agar plates at 37 °C for 18 h.

112 **2.2. Preparation of *Bacillus* endospore suspensions**

113 Strains of *Bacillus* were surface plated on LB plates and incubated at 37 °C for 5 d to
114 ensure the sporulation of more than 90% of cells (Li et al., 2019; Margosch et al., 2004).
115 Sporulation was confirmed by staining with the Schaeffer-Fulton method (malachite
116 green solution of 5 % and 0.1 % safranin) which results in green staining of spores and
117 red staining of vegetative cells. Microscopic examination of spore suspensions
118 confirmed that 90 to 99 % of the cells contained spores. Cells from 5 d old cultures on
119 LB plates were collected by flooding the surface of the plates with 5 ml sterile distilled

120 water twice. The spore suspensions were washed with sterile water for four times by
121 centrifugation at 3,000×g for 15 min at 5 °C and resuspended in sterile distilled water.
122 Between the second and third wash cycles, the spore suspensions were treated at 80 °C
123 for 10 min to kill vegetative cells. The spore suspensions were stored at −80 °C until
124 further use (Li et al., 2019; Paidhungat et al., 2002).

125 **2.3. Determination of survival of strains of *Bacillus* spp. with different copy**
126 **number of the *spoVA*^{2mob} operons after baking.**

127 To determine the contribution of the *spoVA*^{2mob} operon or multiple copies of the operons
128 on survival after baking, we inoculated wheat dough with spores of 8 strains of *Bacillus*
129 with a different copy number of the *spoVA*^{2mob} operon, and determined the survival
130 after baking. The recipe of wheat bread is shown in Table 2. Ingredients were mixed
131 with a KitchenAid 3 Qt mixer for 1 min (slow) and 14 min (fast), followed by a bulk
132 proof at 25 °C for 60 min. The dough was divided into pieces of 150 g and each piece
133 was rolled into a rectangular dough sheet, and inoculated by spreading 1.5 ml of spore
134 suspensions of *Bacillus* spp. with a viable spore count of 10⁸ spores/ml. Subsequently,
135 the dough was rolled, shaped, proofed for 60 min at 32 °C, and baked at 180 °C for 20
136 min. The internal temperature of bread during baking was monitored by a thermocouple
137 that was inserted to the bread during baking. The temperature profile during baking
138 process is shown in Figure 1. Samples were taken from the dough before baking, after
139 baking and cooling at room temperature for 1 h. Total aerobic plate counts of bread
140 samples were determined by mixing 10 g of bread with 90 ml peptone in a stomacher.
141 Serial dilutions in peptone water were surface plated on LB plates and were incubated
142 at 37 °C for 18 h.

143 **2.4. Determination of the spoilage phenotype of different strains of *Bacillus***

144 **inoculated on dough**

145 To monitor the typical characteristics of ropy spoilage including appearance, aroma and
146 bacterial growth, two slices of each bread were placed in Petri dishes and stored at 30 °C
147 for 7 days in a sealed container with 10 % NaCl solution to maintain the water activity
148 (aw) at 0.95 throughout storage. The slices were examined daily. Two pieces of bread
149 of each sample were analyzed each day and the experiment was repeated 3 times.
150 Representative images of slices of bread inoculated with *B. subtilis* FAD109 or *B.*
151 *amyloliquefaciens* FUA2154 are shown in Figure 2.

152 **2.5. Determination of the spoilage phenotype of different strains of *Bacillus***
153 **inoculated on bread**

154 Uncontaminated bread was prepared following a similar procedure as indicated above.
155 After baking and cooling at room temperature, the bread was sliced. Then 30 µl of spore
156 suspensions from 5 strains of *Bacillus* were inoculated at 5 different spots on one bread
157 slice, and 3 replicates of slices were prepared under the same procedure. Uninoculated
158 slices that were incubated at the same conditions served as control.

159 The inoculated bread slices were placed in petri dishes in containers with 10 % NaCl
160 solution (same as before) and incubated at 30 °C for 5 days. The bacterial growth and
161 characteristics of ropy spoilage such as discoloration, visible lines, soft and sticky
162 crumb, and slime were identified throughout five days of storage (Figure 2). The
163 experiment was repeated 3 times.

164 **2.6. Analysis of amylases encoded in the genomes of *Bacillus* spp.**

165 Sequences of extracellular amylases and glucoamylases in *Bacillus* species were
166 retrieved from the Universal Protein Resource (UniProt) (<https://www.uniprot.org/>)
167 database. These protein sequences were used as query sequences for BLAST analysis

168 of whole genome sequences from the strains of *Bacillus* spp. (Li et al., 2019) used in
169 this study. A different complement of amylases was found in *B. amyloliquefaciens* and
170 *B. subtilis*. Therefore, we further blasted these sequences of amylases against genome
171 sequences of additional *Bacillus* strains, including 5 strains of *B. amyloliquefaciens*, 2
172 strains of *B. subtilis* and 2 strains of *B. velezensis* in total (Li et al., 2019).

173 **2.7. Preparation of sourdough**

174 Anti-rope activity has been considered as an important characteristic of some lactic acid
175 bacteria (Pepe et al., 2003). The reutericyclin positive *L. reuteri* TMW1.656 was used
176 as reference strain and its anti-rope activity was compared to *L. reuteri*
177 TMW1.656 Δ *rtcN* Δ *gtfA*. After incubating the two strains in mMRS medium at 37 °C
178 for overnight, 10 ml of culture were added into a 15 ml centrifuge tube, respectively.
179 The culture was washed by centrifugation at 5000 rpm for 3 min and resuspended in 10
180 ml sterile tap water. The 10 ml of washed strain culture was mixed thoroughly with 10
181 g of reweighed white flour. The mixture was fermented at 30 °C for 24 h, corresponding
182 to a sourdough pH of 4.60 \pm 0.02 and 4.57 \pm 0.03 for *L. reuteri* TMW1.656 and
183 TMW1.656 Δ *rtcN* Δ *gtfA*, respectively. The fermented sourdough was placed at -20 °C
184 overnight and then freeze-dried. After freeze drying, the dry sourdough was ground and
185 stored at -20 °C until use.

186 **2.8. Evaluation of the effect of sourdough on ropy spoilage**

187 On the basis of slime production and heat resistance, *B. amyloliquefaciens* FAD 99 was
188 selected as an indicator to determine the antagonistic activities of different sourdough
189 dosage. Sourdough bread was prepared following the recipe in Table 2 by adding 0, 3,
190 6, 12 or 24 % freeze-dried sourdough made with *L. reuteri* TMW1.656, respectively
191 (substitute white flour with freeze-dried sourdough). Each dough was inoculated with
192 1 ml spore suspension of *B. amyloliquefaciens* FAD 99 as described above. Sourdough

193 bread was prepared as described above after inoculation on dough with the spore
194 suspension of *B. amyloliquefaciens* FAD 99. This strain was chosen because, among
195 the strains of *Bacillus* investigated in this study, *B. amyloliquefaciens* FAD99 showed
196 the greatest spoilage potential. The bacterial growth and ropy characteristics were
197 observed over 7 d to determine the dosage of sourdough that inhibits growth of bacilli
198 in bread. This experiment was carried out in three independent experiments.

199 **2.9. Detection of the inhibition effect of reutericyclin on ropy bread spoilage** 200 **caused by *Bacillus* species**

201 The reutericyclin-positive *L. reuteri* TMW1.656 and the isogenic reutericyclin-
202 negative derivative *L. reuteri* TMW1.656 Δ *rtcN* Δ *gtfA* were used to compare the
203 inhibition effect of reutericyclin on ropy spoilage that caused by *B. amyloliquefaciens*
204 FAD 99. The same procedure was applied to produce freeze-dried sourdoughs
205 fermented with *L. reuteri* TMW1.656 and TMW1.656 Δ *rtcN* Δ *gtfA*, respectively. One
206 ml spore suspensions from *B. amyloliquefaciens* FAD 99 was inoculated on the dough
207 containing 12 % freeze-dried sourdough. The sourdough bread was baked as described
208 before and bacterial growth and ropy spoilage characteristics were monitored daily. The
209 experiment was performed in three independent replicates (independent baking trials).

210 **2.10. Statistical analysis**

211 Experiments were carried out at least in three replicates with different batches of spore
212 suspensions and the same batch of freeze-dried sourdough. Statistical analysis of cell
213 counts of *Bacillus* spp. after baking was carried out by ANOVA with Tukey's Honest
214 Significant Difference test in RStudio. Significant differences were assessed with an

215 error probability of 5% ($P < 0.05$).

216 **3. RESULTS**

217 **3.1. Variations of heat resistance of spores from strains of *Bacillus* with different** 218 **copy number of the *spoVA*^{2mob} operons in bread**

219 Heat resistance of *Bacillus* endospores is dependent on the copy number of the
220 *spoVA*^{2mob} operon (Berendsen et al., 2016; Li et al., 2019). Since ropy spoilage of bread
221 is dependent on spore survival during the baking process, bread dough was inoculated
222 with spores of strains of *Bacillus* with a different copy number of the *spoVA*^{2mob} operon
223 and their survival after baking was determined (Figure 3). Overall, the lethality of the
224 baking process, which exposed spores of *Bacillus* spp. to a temperature of more than
225 90 °C for about 10 min (Figure 1), was limited. Spore counts of *Bacillus* strains that did
226 not encode for a *spoVA*^{2mob} operon were reduced by more than 1 log (CFU/g) after
227 baking; spores of *Bacillus* strains that encoded three copies of the *spoVA*^{2mob} operons
228 were reduced by less than 0.5 log (CFU/g); spores of *Bacillus* strains with two copies
229 of the *spoVA*^{2mob} operons showed strain-specific survival (Figure 3). Surviving spores
230 of *B. subtilis* and *B. amyloliquefaciens* caused visible bread spoilage after 1 – 2 days of
231 incubation (Figure 2 and data not shown).

232 **3.2. Appearance of ropy spoilage phenotype of different strains of *Bacillus*** 233 **inoculated on dough**

234 *B. subtilis* and *B. amyloliquefaciens* are recognized as the causative agents of ropy
235 bread spoilage. To determine whether the spoilage phenotype of different strains of *B.*
236 *subtilis* and *B. amyloliquefaciens* differs, spore suspensions of 5 different strains of
237 *Bacillus* were inoculated at 5 different spots on one bread slice. The inoculation of bread
238 rather than dough was chosen to eliminate the influence of spore survival after baking.

239 Spoilage characteristics were monitored for 5 d (Figure 4). The spores from 2 strains of
240 *B. amyloliquefaciens* spoiled the bread more rapidly when compared to spores from 3
241 strains of *B. subtilis*. *B. amyloliquefaciens* FAD 99 visibly degraded the crumb after 1
242 day of incubation and formed slime after 3 – 4 d; crumb degradation and slime
243 formation by spores from *B. subtilis* was observed only after 3 and 5 d, respectively
244 (Figure 4). *B. amyloliquefaciens* FAD 99 was chosen as spoilage organism for the
245 subsequent experiments.

246 **3.3. Analysis of amylases encoded in genomes of *Bacillus* spp.**

247 To identify putative genetic determinants for the different spoilage phenotypes of *B.*
248 *amyloliquefaciens* and *B. subtilis* strains, genes coding for extracellular amylases and
249 proteases were identified in the genomes of 5 strains of *B. amyloliquefaciens*, 2 strains
250 of *B. subtilis* and 2 strains of *B. velezensis* (Li et al., 2019). Amylases were identified
251 by BLAST analysis with protein sequences of extracellular amylases from *Bacillus* spp.
252 that were available in the UniProt database as query sequences. Strains of
253 *B. amyloliquefaciens* and *B. subtilis* differed with respect to the amylases that were
254 identified in the genomes. Strains of *B. subtilis* and *B. velezensis* encode for AmyE as
255 sole extracellular amylase while five extracellular amylases including a
256 hyperthermostable α -amylase are encoded in genomes of each strain of
257 *B. amyloliquefaciens* (Figure 5). All genomes encode for two glucoamylases, but
258 β -amylases were not identified. The different complement of amylases likely relates to
259 the different spoilage phenotype of strains of *B. amyloliquefaciens* and *B. subtilis*.

260 **3.4. Effect of sourdough dosage on ropy spoilage**

261 To determine the effect of sourdough on ropy spoilage, bread doughs were prepared
262 with 0, 3, 6 12 and 24 % of freeze-dried sourdough fermented with the reutericyclin-

263 producing *L. reuteri* TMW1.656. Bread dough was inoculated with a spore suspension
264 of *B. amyloliquefaciens* FAD 99 and the spoilage phenotype was monitored daily
265 (Figure 6). The addition of 3 % sourdough did not affect spoilage, but spoilage was
266 delayed by 2 – 3 days after the addition of 6 or 12 % sourdough corresponding to a
267 bread pH of 6.2 and 5.9, respectively (Figure 6). The bread made with 24 % sourdough
268 did not spoil during the 5 days of observation.

269 **3.5. Detection of the inhibition effect of reutericyclin produced by *L. reuteri* on** 270 **ropy bread spoilage caused by *Bacillus***

271 To compare the inhibitory effect of reutericyclin on ropy spoilage caused by FAD 99
272 *B. amyloliquefaciens*, sourdoughs were fermented with the reutericyclin positive
273 *L. reuteri* TMW1.656 and its isogenic, reutericyclin negative *L. reuteri*
274 TMW1.656 Δ *rtcN* Δ *gtfA* (Chen et al., 2016, this study). The GtfA – RtcN double mutant
275 was used to avoid exopolysaccharide formation during sourdough fermentation, which
276 may confound the assessment of slime formation during ropy spoilage. Bread was
277 prepared with 12 % sourdough, a dosage that delays spoilage by 1 – 2 d (Figure 6), and
278 bread dough was inoculated with spores of *B. amyloliquefaciens* FAD 99. Bread
279 without reutericyclin showed visible line on the third day and became slimy on the
280 fourth day (Figure 7). Bread with reutericyclin showed visible lines only on day five
281 and slime was not observed until day seven (Figure 7).

282 **4. DISCUSSION**

283 Ropy spoilage of bread is frequent in warm and humid climate including Mediterranean
284 countries, Africa as well as Australia (Voysey and Hammond, 1993). *Bacillus* spp.,
285 which produce extracellular slimy polysaccharides and possess proteolytic and
286 amylolytic enzymes, are the major cause of ropiness (Corsetti et al., 2000; Pepe et al.,

287 2003). *Bacillus* endospores that are present in raw materials survive during baking
288 (Pepe et al., 2003; Valerio et al., 2012) and lead to ropy spoilage if storage conditions
289 support germination and growth (Vaičiulytė-Funk et al., 2015).

290 The *spoVA*^{2mob} operon plays an essential role in the heat resistance of *Bacillus* spores;
291 its effect on heat resistance is dependent on the copy number of the operon in genomes
292 of *Bacillus* spp. (Berendsen et al., 2016; Wang et al., 2018). This study used 8 strains
293 of *Bacillus* isolated from ropy bread and daqu (Röcken and Spicher, 1993; Wang et al.,
294 2018) to determine the impact of the copy number of the *spoVA*^{2mob} operon on heat
295 resistance during baking process, where the temperature reaches up to 100 °C for
296 several minutes. Viable counts of spores with two or three copies of *spoVA*^{2mob} operons
297 were reduced by less than 1 log (CFU/g); however, the lethality of the baking process
298 towards spores of strains that did not harbor a *spoVA*^{2mob} operon was also rather limited.
299 Owing to this limited inactivation of endospores that do not harbour multiple copies of
300 the *spoVA*^{2mob} operon, the presence of multiple copies of the operons did not appreciably
301 improve survival during baking and is not a prerequisite for the ability of strains to spoil
302 bread.

303 Strains of *B. subtilis* were most frequently isolated from ropy bread; however, several
304 past studies misidentified *B. amyloliquefaciens* as *B. subtilis* (Pepe et al., 2003; Röcken
305 and Spicher, 1993; Valerio et al., 2012) because the two species show a high level of
306 similarity of their 16S rRNA gene sequence (Reginensi et al., 2013). This study
307 compared the spoilage phenotype of 3 strains of *B. subtilis* and 2 strains of *B.*
308 *amyloliquefaciens* and demonstrated that *B. amyloliquefaciens* caused spoilage of bread
309 2-3 days earlier than strains of *B. subtilis*. We further investigated the complement of
310 extracellular amylases and proteases that are encoded in genomes of 5 strains of
311 *B. amyloliquefaciens*, 2 strains of *B. subtilis* and 2 strains of the closely related species

312 *B. velezensis* (Li et al., 2019). Strains of *B. amyloliquefaciens* were differentiated from
313 *B. subtilis* and *B. velezensis* by encoding for multiple extracellular α -amylases and
314 gluco-amylases. Moreover, a hyperthermostable α -amylase was identified in each
315 genome of *B. amyloliquefaciens* but not in *B. subtilis* and *B. velezensis*; this enzyme
316 may remain active after baking and initiate starch degradation in bread. The comparison
317 of the genotype of *B. subtilis* and *B. amyloliquefaciens* with the spoilage phenotype
318 suggests that extracellular amylases and proteases contribute to ropy spoilage of bread,
319 however, the contribution of individual enzymes remains subject to future studies.

320 Rope-forming bacilli are endophytes of plants including wheat, and their presence in
321 seeds mediates vertical transmission of commensal plant microbiota to the offspring
322 (Robinson et al., 2016; Shahzad et al., 2016). Therefore, contamination of wheat flour
323 with *Bacillus* spores is unavoidable (Pepe et al., 2003; Saranraj and Geetha, 2012). The
324 use of sourdough as a biopreservative has only limited effect against fungal bread
325 spoilage (Axel et al., 2017; Quattrini et al., 2018) but inhibits growth of rope-forming
326 bacilli (Röcken and Spicher, 1993; Rosenquist and Hansen, 1998; Valerio et al., 2008).
327 The preservative effect of sourdough was mainly attributed to formation of organic
328 acids and acidification of the bread crumb (Rosenquist and Hansen, 1998; Valerio et
329 al., 2008). Acidification to a pH of 3.7 to 4.5 inhibits ropy spoilage (Pepe et al., 2003)
330 and undissociated acetic acid enhances the preservative effect of sourdough
331 (Rosenquist and Hansen, 1998).

332 Although sourdough has re-claimed its place as the standard process of bread-making,
333 the use of sourdough or sourdough products may not always suffice to inhibit ropy
334 spoilage. Dried sourdough products are standardized on the basis of the concentration
335 of organic acids but typically much of the volatile acetic acid is lost during drying
336 (Brandt, 2007). Moreover, not all sourdoughs are effective at inhibition of rope-forming

337 *Bacillus* spores due to their high pH and low concentration of organic acids (Katina et
338 al., 2002). The concentration of organic acids in sourdoughs and bread dough fermented
339 in bakeries is not as readily standardized as for stabilized sourdough products (Brandt,
340 2007). In addition, low pH and undissociated organic acids may alter the flavor and
341 taste of bread (Hansen and Schieberle, 2005). In this study, addition of 24 % sourdough
342 to a bread recipe effectively inhibited the growth of *B. amyloliquefaciens* but the
343 combined use of organic acid with specific antimicrobial compounds may allow bread
344 preservation without excessive acidity.

345 Bacteriocins of lactic acid bacteria were evaluated with respect to their activity against
346 *Bacillus* spp. (Digaitiene et al., 2012; Pepe et al., 2003) even though nisin is the only
347 bacteriocin which has a general inhibitory effect against strains of *Bacillus* (Rosenquist
348 and Hansen, 1998). The use of nisin as additive or the use of nisin-producing starter
349 cultures in sourdough, however, had no effect on growth of *Bacillus* in bread
350 (Rosenquist and Hansen, 1998). This may relate to inactivation of nisin by proteases
351 and thiol exchange reactions at the dough stage, or to their heat inactivation during
352 baking (Gänzle, 2014). Reutericyclin is produced by *Streptococcus mutans* and few
353 strains of *L. reuteri* (Lin et al., 2015; Tang et al., 2019). Reutericyclin is a tetramic acid
354 derivative which resists proteolysis in dough and heat inactivation during baking
355 (Gänzle et al., 2000). Reutericyclin exhibits a bactericidal mode of action against
356 *Bacillus* spp., and inhibited spore germination (Gänzle et al., 2000). We used some of
357 the same strains of *Bacillus* that were previously characterized with regards to their
358 sensitivity to reutericyclin (Gänzle et al., 2000). Fermentation of sourdoughs with
359 isogenic strains that produce comparable levels of organic acids but differ with respect
360 to reutericyclin production (Lin et al., 2015) demonstrated that reutericyclin
361 substantially delays growth of *Bacillus* spp. in bread. The use of reutericyclin-

362 producing strains thus allows the prevention of ropy spoilage with a reduced dosage of
363 sourdough and corresponding reduced levels of acidity. The reutericyclin producing
364 strain *L. reuteri* TMW1.656 was isolated from an industrial sourdough and also
365 beneficially impacts bread texture through production of reuteran (Chen et al., 2016;
366 Gänzle and Vogel, 2003).

367 In conclusion, multiple copies of the *spoVA*^{2mob} operon show only a limited effect on
368 the heat resistance of *Bacillus* spores during baking. The different spoilage phenotype
369 observed after growth of *B. amyloliquefaciens* and *B. subtilis* likely relates to the
370 differential presence and expression of extracellular amylases and proteases in these
371 species. Moreover, this study confirmed the inhibitory effect of sourdough and
372 demonstrated that reutericyclin produced at the sourdough stage contributes to
373 inhibition of growth of rope-forming bacilli in bread.

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377

378 **REFERENCES**

- 379 Anonymous, 2019a. Bread market-growth, trends and forecasts [WWW Document].
380 URL <https://www.mordorintelligence.com/industry-reports/bread-market>
381 (accessed 10.29.19).
- 382 Anonymous, 2019b. Global sourdough market-segmented by type, source, application
383 and geography [WWW Document]. URL
384 <https://www.mordorintelligence.com/industry-reports/sourdough-market>
385 (accessed 10.31.19).
- 386 Axel, C., Zannini, E., Arendt, E.K., 2017. Mold spoilage of bread and its
387 biopreservation: A review of current strategies for bread shelf life extension. *Crit.*
388 *Rev. Food Sci. Nutr.* 57, 3528–3542.
389 <https://doi.org/10.1080/10408398.2016.1147417>
- 390 Bailey, C.P., von Holy, A., 1993. *Bacillus* spore contamination associated with
391 commercial bread manufacture. *Food Microbiol.* 10, 287–294.
392 <https://doi.org/10.1006/fmic.1993.1033>
- 393 Berendsen, E.M., Boekhorst, J., Kuipers, O.P., Wells-Bennik, M.H.J., 2016. A mobile
394 genetic element profoundly increases heat resistance of bacterial spores. *ISME J.*
395 10, 2633–2642. <https://doi.org/10.1038/ismej.2016.59>
- 396 Brandt, M.J., 2007. Sourdough products for convenient use in baking. *Food Microbiol.*
397 24, 161–164. <https://doi.org/10.1016/j.fm.2006.07.010>
- 398 Chen, X.H., Koumoutsis, A., Scholz, R., Eisenreich, A., Schneider, K., Heinemeyer, I.,
399 Morgenstern, B., Voss, B., Hess, W.R., Reva, O., Junge, H., Voigt, B., Jungblut,
400 P.R., Vater, J., Süßmuth, R., Liesegang, H., Strittmatter, A., Gottschalk, G.,
401 Borriss, R., 2007. Comparative analysis of the complete genome sequence of the

402 plant growth-promoting bacterium *Bacillus amyloliquefaciens* FZB42. Nat.
403 Biotechnol. 25, 1007–1014. <https://doi.org/10.1038/nbt1325>

404 Chen, X.Y., Levy, C., Gänzle, M.G., 2016. Structure-function relationships of bacterial
405 and enzymatically produced reuterans and dextran in sourdough bread baking
406 application. Int. J. Food Microbiol. 239, 95–102.
407 <https://doi.org/10.1016/j.ijfoodmicro.2016.06.010>

408 Corsetti, A., Gobbetti, M., De Marco, B., Balestrieri, F., Paoletti, F., Russi, L., Rossi,
409 J., 2000. Combined effect of sourdough lactic acid bacteria and additives bread
410 firmness and staling. J. Agric. Food Chem. 48, 3044–3051.
411 <https://doi.org/10.1021/jf990853e>

412 Digaitiene, A., Hansen, Å.S., Juodeikiene, G., Eidukonyte, D., Josephsen, J., 2012.
413 Lactic acid bacteria isolated from rye sourdoughs produce bacteriocin-like
414 inhibitory substances active against *Bacillus subtilis* and fungi. J. Appl. Microbiol.
415 112, 742. <https://doi.org/10.1111/j.1365-2672.2012.05249.x>

416 Fan, B., Chen, X.H., Budiharjo, A., Bleiss, W., Vater, J., Borriss, R., 2011. Efficient
417 colonization of plant roots by the plant growth promoting bacterium *Bacillus*
418 *amyloliquefaciens* FZB42, engineered to express green fluorescent protein. J.
419 Biotechnol. 151, 303–311. <https://doi.org/10.1016/j.jbiotec.2010.12.022>

420 Fangio, M.F., Roura, S.I., Fritz, R., 2010. Isolation and identification of *Bacillus* spp.
421 and related genera from different starchy foods. J. Food Sci. 75, 218–221.
422 <https://doi.org/10.1111/j.1750-3841.2010.01566.x>

423 Gänzle, M.G., 2014. Enzymatic and bacterial conversions during sourdough
424 fermentation. Food Microbiol. 37, 2–10. <https://doi.org/10.1016/j.fm.2013.04.007>

425 Gänzle, M.G., Hölzel, A., Walter, J., Jung, G., Hammes, W.P., 2000. Characterization

426 of reutericyclin produced by *Lactobacillus reuteri* LTH2584. Appl. Environ.
427 Microbiol. 66, 4325–4333. <https://doi.org/10.1128/AEM.66.10.4325-4333.2000>

428 Gänzle, M.G., Vogel, R.F., 2003. Contribution of reutericyclin production to the stable
429 persistence of *Lactobacillus reuteri* in an industrial sourdough fermentation. Int.
430 J. Food Microbiol. 80, 31–45. [https://doi.org/10.1016/S0168-1605\(02\)00146-0](https://doi.org/10.1016/S0168-1605(02)00146-0)

431 Hansen, A., Schieberle, P., 2005. Generation of aroma compounds during sourdough
432 fermentation: applied and fundamental aspects. Trends Food Sci. Technol. 16, 85–
433 94. <https://doi.org/10.1016/J.TIFS.2004.03.007>

434 Jamal, Q., Monkhung, S., Munir, S., Cho, J.Y., Moon, J.H., Khattak, B.U., Malik, M.S.,
435 Younis, F., Kim, K.Y., 2019. Identification of cyclo(L-Pro-D-Tyr) from *Bacillus*
436 *amyloliquefaciens* Y1 exhibiting antifungal activity against *Fusarium*
437 *graminearum* to control crown rot in wheat. Appl. Ecol. Environ. Res. 17, 6299–
438 6314. https://doi.org/10.15666/aeer/1703_62996314

439 Katina, K., Sauri, M., Alakomi, H.L., Mattila-Sandholm, T., 2002. Potential of lactic
440 acid bacteria to inhibit rope spoilage in wheat sourdough bread. LWT - Food Sci.
441 Technol. 35, 38–45. <https://doi.org/10.1006/fstl.2001.0808>

442 Kourkouta, L., Koukourikos, K., Tsaloglidou, 2017. Bread and Health. J. Pharm.
443 Pharmacol. 5, 821–826. <https://doi.org/10.17265/2328-2150/2017.11.005>

444 Li, Z., Schottroff, F., Simpson, D.J., Gänzle, M.G., 2019. The copy number of the
445 *spoVA2mob* operon determines pressure resistance of *Bacillus* endospores. Appl.
446 Environ. Microbiol. 85, e01596-19. <https://doi.org/10.1128/aem.01596-19>

447 Lin, X.B., Lohans, C.T., Duar, R., Zheng, J., Vederas, J.C., Walter, J., Gänzle, M., 2015.
448 Genetic determinants of reutericyclin biosynthesis in *Lactobacillus reuteri*. Appl.
449 Environ. Microbiol. 81, 2032–41. <https://doi.org/10.1128/AEM.03691-14>

450 Margosch, D., Gänzle, M.G., Ehrmann, M.A., Vogel, R.F., 2004. Pressure inactivation
451 of *Bacillus* endospores. *Appl. Environ. Microbiol.* 70, 7321–7328.
452 <https://doi.org/10.1128/AEM.70.12.7321-7328.2004>

453 Needham, R., Williams, J., Beales, N., Voysey, P., Magan, N., 2005. Early detection
454 and differentiation of spoilage of bakery products. *Sensors Actuators, B Chem.*
455 106, 20–23. <https://doi.org/10.1016/j.snb.2004.05.032>

456 Paidhungat, M., Setlow, B., Daniels, W.B., Hoover, D., Papafragkou, E., Setlow, P.,
457 2002. Mechanisms of induction of germination of *Bacillus subtilis* spores by high
458 pressure. *Appl. Environ. Microbiol.* 68, 3172–3175.
459 <https://doi.org/10.1128/AEM.68.6.3172-3175.2002>

460 Pepe, O., Blaiotta, G., Moschetti, G., Greco, T., Villani, F., 2003. Rope-producing
461 strains of *Bacillus* spp. from wheat bread and strategy for their control by lactic
462 acid bacteria. *Appl. Environ. Microbiol.* 69, 2321–2329.
463 <https://doi.org/10.1128/aem.69.4.2321-2329.2003>

464 Quattrini, M., Liang, N., Fortina, M.G., Xiang, S., Curtis, J.M., Gänzle, M., 2018.
465 Exploiting synergies of sourdough and antifungal organic acids to delay fungal
466 spoilage of bread. *Int. J. Food Microbiol.*
467 <https://doi.org/10.1016/j.ijfoodmicro.2018.09.007>

468 Reginensi, S., González, M.J., Gorgoroso, F., Reginensi, S.M., Olivera, J.A., Bermúdez,
469 J., 2013. Polyphasic identification of closely related *Bacillus subtilis* and *Bacillus*
470 *amyloliquefaciens* isolates from dairy farms and milk powder. *J. Microbiol.*
471 *Biotechnol. Food Sci.* 2, 2326–2331.

472 Robinson, R.J., Fraaije, B.A., Clark, I.M., Jackson, R.W., Hirsch, P.R., Mauchline, T.H.,
473 2016. Endophytic bacterial community composition in wheat (*Triticum aestivum*)

474 is determined by plant tissue type, developmental stage and soil nutrient
475 availability. *Plant Soil* 405, 381–396. <https://doi.org/10.1007/s11104-015-2495-4>

476 Röcken, W., Spicher, G., 1993. Fadenziehende Bakterien-Vorkommen, Bedeutung
477 Gegenmaßnahmen. *Getreide Mehl Brot* 47, 30–35.

478 Rosenquist, H., Hansen, Å., 1998. The antimicrobial effect of organic acids, sour dough
479 and nisin against *Bacillus subtilis* and *B. licheniformis* isolated from wheat bread.
480 *J. Appl. Microbiol.* 85, 621–631. [https://doi.org/10.1046/j.1365-](https://doi.org/10.1046/j.1365-2672.1998.853540.x)
481 [2672.1998.853540.x](https://doi.org/10.1046/j.1365-2672.1998.853540.x)

482 Sadeghi, A., 2008. The secrets of sourdough; a review of miraculous potentials of
483 sourdough in bread shelf life. *Biotechnology* 7, 413–417.
484 <https://doi.org/10.3923/biotech.2008.413.417>

485 Saranraj, P., Geetha, M., 2012. Microbial spoilage of bakery products and its control
486 by preservatives. *Int. J. Pharm. Biol. Arch.* 3, 38–48.

487 Setlow, P., 2006. Spores of *Bacillus subtilis*: their resistance to and killing by radiation,
488 heat and chemicals. *J. Appl. Microbiol.* 101, 514–525.
489 <https://doi.org/10.1111/j.1365-2672.2005.02736.x>

490 Setlow, P., 2003. Spore germination. *Curr. Opin. Microbiol.* 6, 550–556.
491 <https://doi.org/10.1016/j.mib.2003.10.001>

492 Shahzad, R., Waqas, M., Khan, A.L., Asaf, S., Khan, M.A., Kang, S.-M., Yun, B.-W.,
493 Lee, I.-J., 2016. Seed-borne endophytic *Bacillus amyloliquefaciens* RWL-1
494 produces gibberellins and regulates endogenous phytohormones of *Oryza sativa*.
495 *Plant Physiol. Biochem.* 106, 236–243.
496 <https://doi.org/10.1016/j.plaphy.2016.05.006>

497 Sorokulova, I.B., Reva, O.N., Smirnov, V.V., Pinchuk, I.V., Lapa, S.V., Urdaci, M.C.,
498 2003. Genetic diversity and involvement in bread spoilage of *Bacillus* strains
499 isolated from flour and ropy bread. *Lett. Appl. Microbiol.* 37, 169–173.
500 <https://doi.org/10.1046/j.1472-765X.2003.01372.x>

501 Tang, X., Kudo, Y., Baker, J., LaBonte, S., Jordan, P.A., McKinnie, S.M.K., Guo, J.,
502 Huan, T., Moore, B.S., Edlund, A., 2019. Cariogenic *Streptococcus mutans*
503 produces strain-specific antibiotics that impair commensal colonization. *bioRxiv*
504 755025. <https://doi.org/10.1101/755025>

505 Tilak, K.V.B.R., Reddy, B.S., 2006. *Bacillus cereus* and *B. circulans* – novel inoculants
506 for crops. *Curr. Sci.* 90, 642–644. <https://doi.org/10.2307/24089111>

507 Vaičiulytė-Funk, L., Žvirdauskienė, R., Šalomskienė, J., Šarkinas, A., 2015. The effect
508 of wheat bread contamination by the *Bacillus* genus bacteria on the quality and
509 safety of bread. *Zemdirbyste-Agriculture* 102, 351–358.
510 <https://doi.org/10.13080/z-a.2015.102.045>

511 Valerio, F., De Bellis, P., Di Biase, M., Lonigro, S.L.L., Giussani, B., Visconti, A.,
512 Lavermicocca, P., Sisto, A., 2012. Diversity of spore-forming bacteria and
513 identification of *Bacillus amyloliquefaciens* as a species frequently associated with
514 the ropy spoilage of bread. *Int. J. Food Microbiol.* 156, 278–285.
515 <https://doi.org/10.1016/j.ijfoodmicro.2012.04.005>

516 Valerio, F., De Bellis, P., Lonigro, S.L., Visconti, A., Lavermicocca, P., 2008. Use of
517 *Lactobacillus plantarum* fermentation products in bread-making to prevent
518 *Bacillus subtilis* ropy spoilage. *Int. J. Food Microbiol.* 122, 328–332.
519 <https://doi.org/10.1016/j.ijfoodmicro.2008.01.005>

520 Valerio, F., Di Biase, M., Huchet, V., Desriac, N., Lonigro, S.L., Lavermicocca, P.,

521 Sohier, D., Postollec, F., 2015. Comparison of three *Bacillus amyloliquefaciens*
522 strains growth behaviour and evaluation of the spoilage risk during bread shelf-
523 life. Food Microbiol. 45, 2–9. <https://doi.org/10.1016/J.FM.2014.07.006>

524 Vessey, J.K., Buss, T.J., 2002. *Bacillus cereus* UW85 inoculation effects on growth,
525 nodulation, and N accumulation in grain legumes: Controlled-environment studies.
526 Can. J. Plant Sci. 82, 283–290.

527 Voysey, P.A., Hammond, J.C., 1993. Reduced-additive breadmaking technology, in:
528 Technology of Reduced-Additive Foods. Springer US, Boston, MA, pp. 80–94.
529 https://doi.org/10.1007/978-1-4615-2115-0_4

530 Wang, Z., Li, P., Luo, L., Simpson, D.J., Gänzle, M.G., Björkroth, J., 2018. *Daqu*
531 fermentation selects for heat-resistant *Enterobacteriaceae* and bacilli. Appl.
532 Environ. Microbiol. 84, e01483-18. <https://doi.org/10.1128/AEM.01483-18>

533 Zheng, J., Wittouck, S., Salvetti, E., Franz, C.M.A.B., Harris, H.M.B., Mattarelli, P.,
534 O’Toole, P.W., Pot, B., Vandamme, P., Walter, J., Watanabe, K., Wuyts, S., Felis,
535 G.E., Gänzle, M.G., Lebeer, S., 2020. A taxonomic note on the genus
536 *Lactobacillus*: Description of 23 novel genera, emended description of the genus
537 *Lactobacillus* Beijerinck 1901, and union of *Lactobacillaceae* and
538 *Leuconostocaceae*. Int. J. Syst. Evol. Microbiol. in press;
539 <https://doi.org/10.1099/ijsem.0.004107>

540 Zheng, J., Zhao, X., Lin, X.B., Gänzle, M., 2016. Comparative genomics *Lactobacillus*
541 *reuteri* from sourdough reveals adaptation of an intestinal symbiont to food
542 fermentations. Sci. Rep. 5, 18234. <https://doi.org/10.1038/srep18234>

543

544 **FIGURES AND TABLES**

545 **Table 1** Origin of strains used in this study and copy number of the *spoVA*^{2mob} operon
 546 in each strain.

Microrganism	Strain; origin	Copy # of <i>spoVA</i>^{2mob} operon/genome	Reference
<i>B. velezensis</i>	^a FUA 2155; daqu	0	(Wang et al., 2018)
<i>B. subtilis</i>	FUA 2114; malted oats	0	(Li et al., 2019)
<i>B. subtilis</i>	FAD 110; ropy bread	2	(Röcken and Spicher, 1993)
<i>B. subtilis</i>	FAD 109; ropy bread	2	(Röcken and Spicher, 1993)
<i>B. amyloliquefaciens</i>	FUA 2154; daqu	2	(Wang et al., 2018)
<i>B. amyloliquefaciens</i>	FUA 2153; daqu	2	(Wang et al., 2018)
<i>B. amyloliquefaciens</i>	FAD 99; ropy bread	3	(Röcken and Spicher, 1993)
<i>B. amyloliquefaciens</i>	FAD We; ropy bread	3	(Röcken and Spicher, 1993)
<i>L. reuteri</i> (reutericyclin positive)	TMW1.656; sourdough	n/a	(Gänzle and Vogel, 2003)
<i>L. reuteri</i> (reutericyclin negative)	TMW1.656 Δ rtcN Δ gtfA	n/a	this study

547 ^a FUA number, Food Microbiology culture collection at the University of Alberta.

548

549 **Table 2** Wheat bread formula.

Ingredients^a	Amount [g]
White wheat flour	100.0 minus weight of freeze-dried sourdough
Freeze-dried sourdough	0, 3.0, 6.0, 12.0, or 24.0
Sterile tap water	60.0
Yeast	2.0
Salt	2.0
Sucrose	2.0

550 ^a White wheat flour (Robin Hood) and baker's yeast (Fleischmann's Active Dry Yeast)

551 were obtained from a local supermarket.

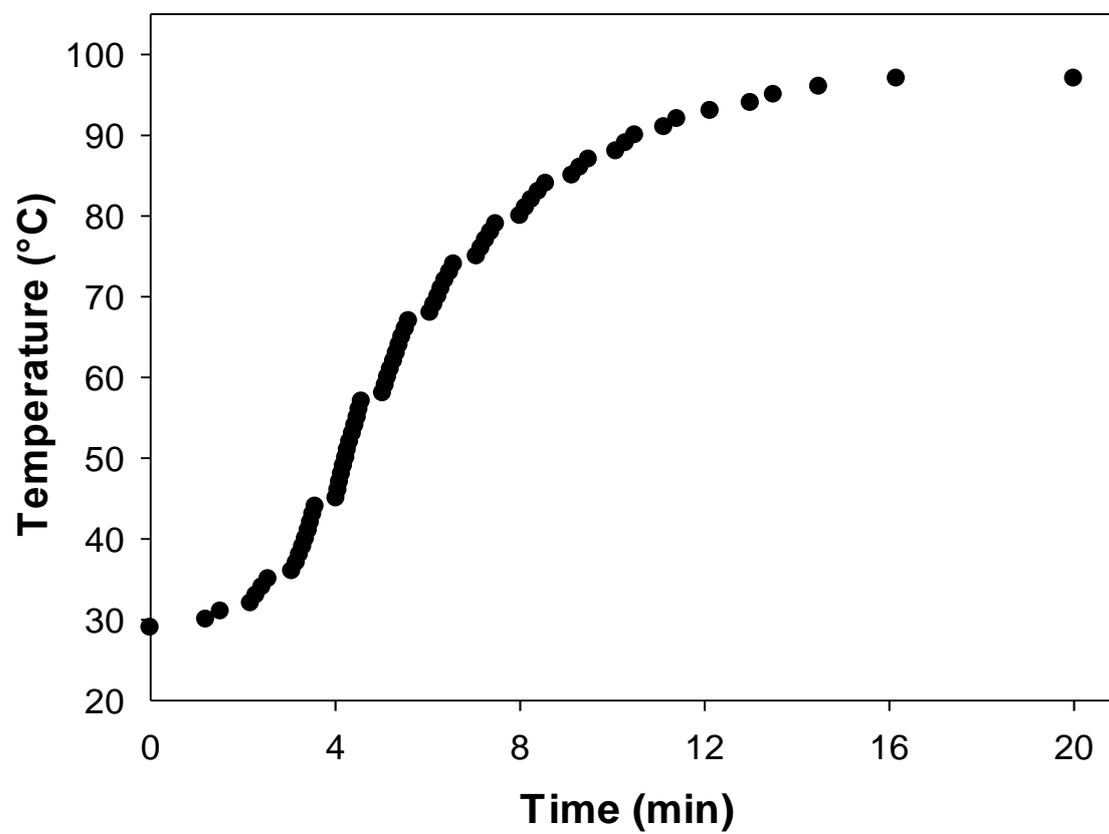


Figure 1. The temperature profile during baking process.



Figure 2. Images of bread slices that were inoculated with *B. subtilis* FAD109 (upper images) or *B. amyloliquefaciens* FUA2154 (lower images). Pictures of the same slice were taken at the day of baking (0d) and after 1 – 5 d of storage. Images are representative of three independent experiments.

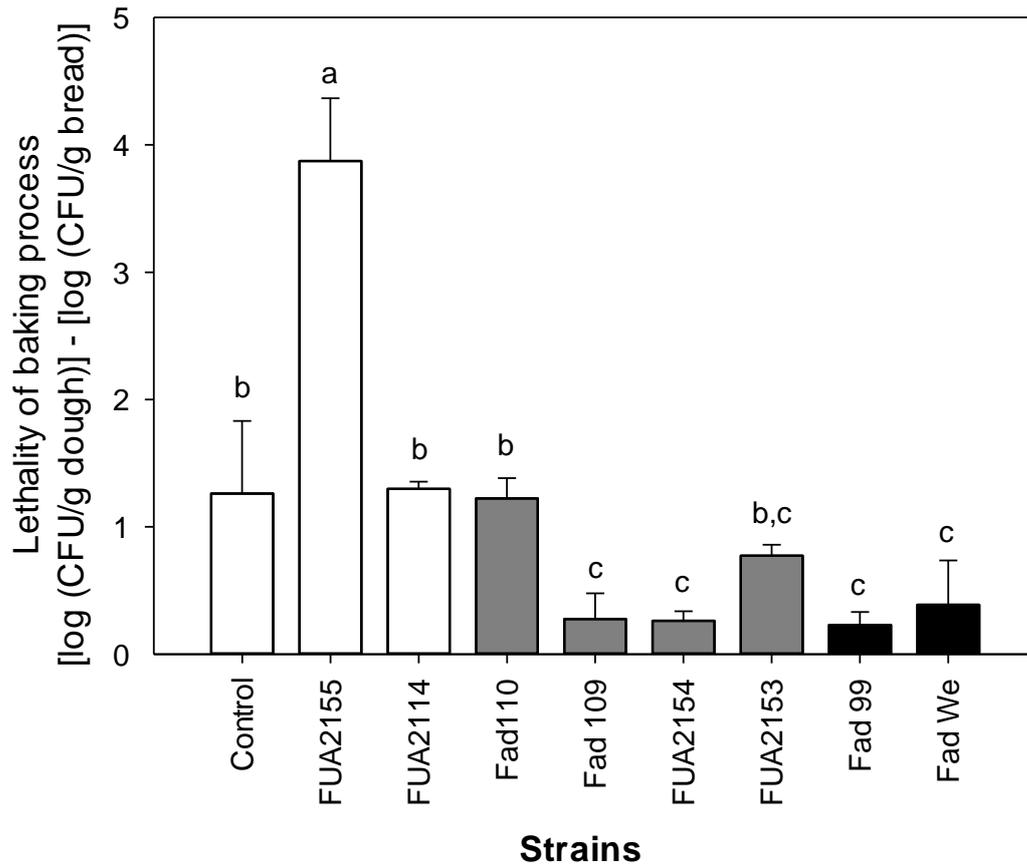


Figure 3. Lethality of the baking process [$\log(N_0/N)$] towards spores of strains of *Bacillus* with different copy number of the *spoVA*^{2mob} operons. The color coding of the bars indicates the copy number of the *spoVA*^{2mob} operon as follows: white (0 copies), gray (two copies), and black (three copies). Strain number and species are indicated as follows: *B. velezensis* FUA 2155; *B. subtilis* FUA 2114; *B. subtilis* FAD 110; *B. subtilis* FAD 109; *B. amyloliquefaciens* FUA 2154; *B. amyloliquefaciens* FUA 2153; *B. amyloliquefaciens* FAD 99; *B. amyloliquefaciens* FAD We. Control refers to bread that was not inoculated with *Bacillus* endospores. The data are shown as means of three independent experiments, error bars indicate the standard deviation. Cell counts that do not share a common superscript differ significantly ($P < 0.05$).

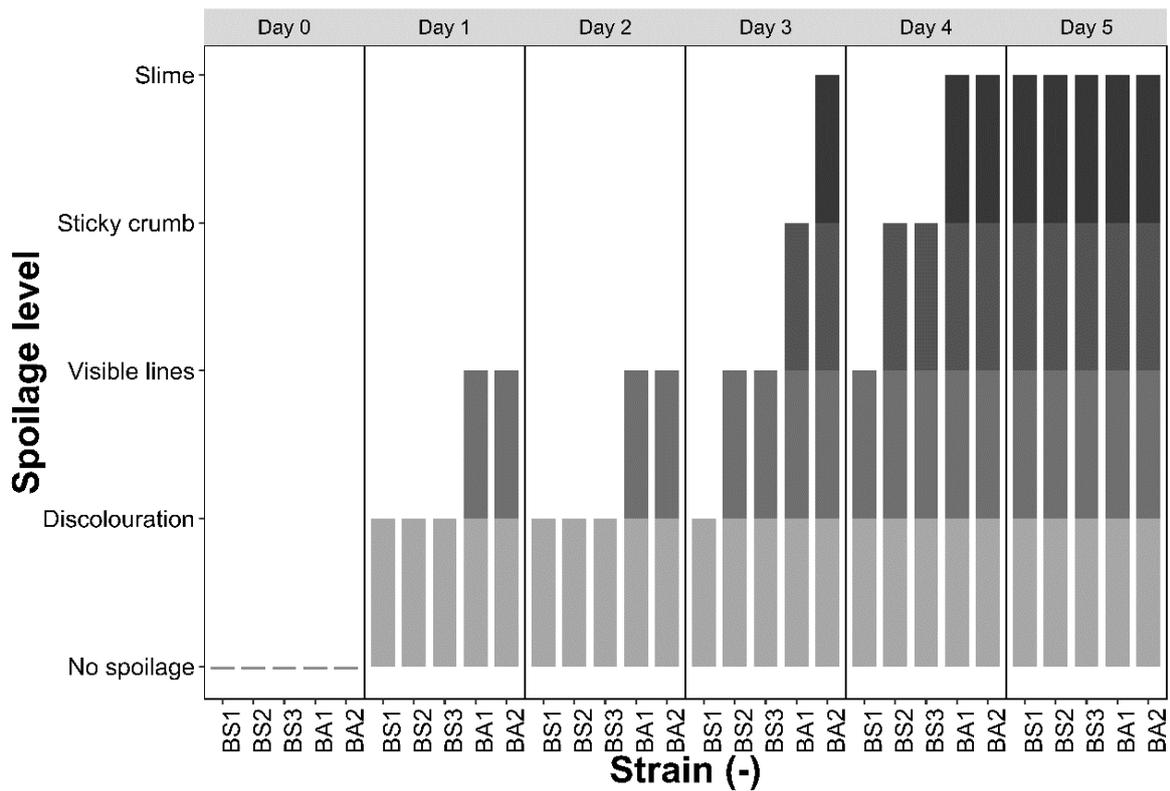


Figure 4. Kinetics of appearance of rope spoilage characteristics in breads contaminated with *B. subtilis* and *B. amyloliquefaciens* after baking. Contaminated slices were stored at 30 °C and aw 0.95 for 5 days. Different colors of gray scale represent different levels of spoilage from low to high. The strain number is labeled as follows: BS1 (*B. subtilis* FUA 2114); BS2 (*B. subtilis* FAD 110); BS3 (*B. subtilis* FAD 109); BA1 (*B. amyloliquefaciens* FUA 2154) and BA2 (*B. amyloliquefaciens* FAD 99). The data represent results from three independent experiments.

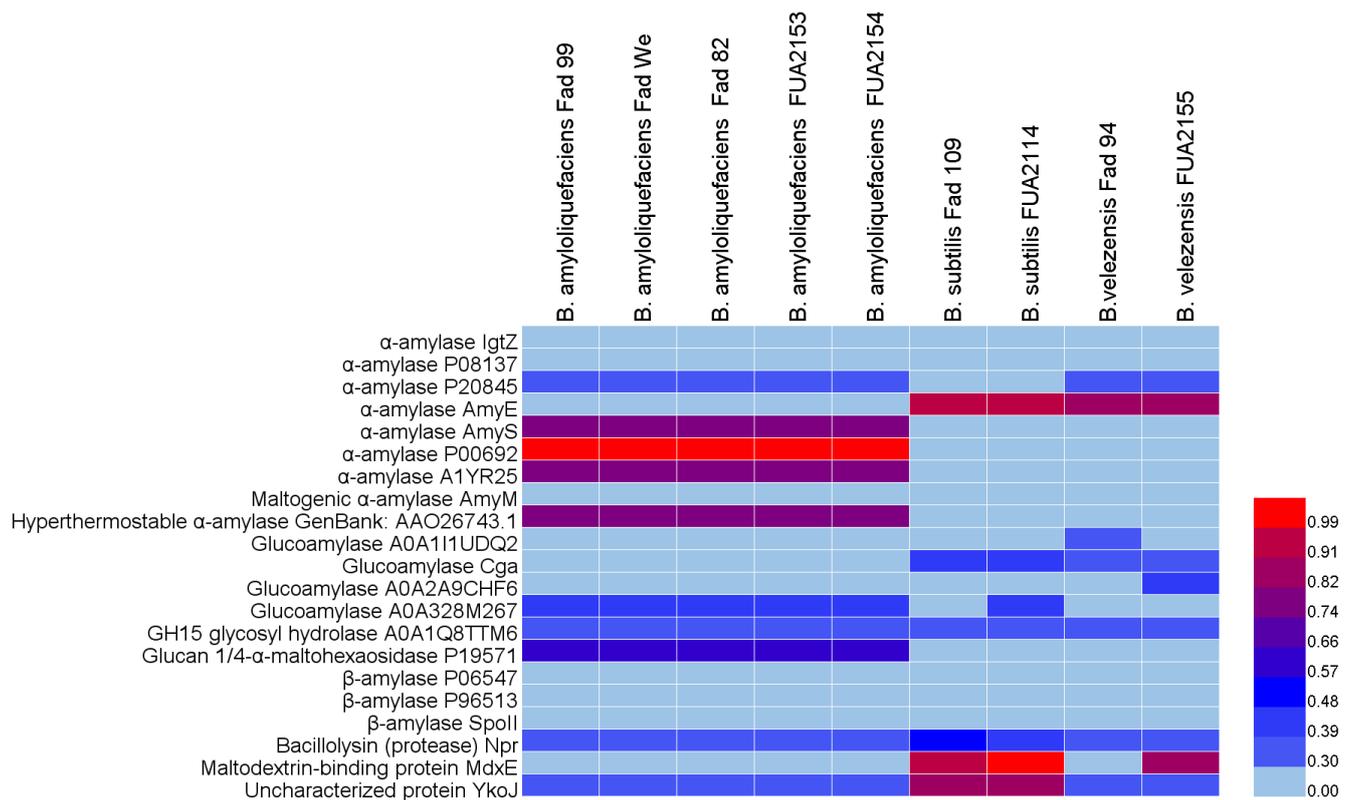


Figure 5. Identification of amylases, glucoamylases and other starch-active enzymes encoded in the genomes of 5 strains of *B. amyloliquefaciens*, 2 strains of *B. subtilis* and 2 strains of *B. velezensis*. With exception of the four glucoamylases, all proteins encode for a signal peptide that is predicted to mediate protein export. Uniprot protein accession numbers are indicated unless specified otherwise. **IgtZ** (A0P8X0) GH119 α-amylase; **P08137** GH13 α-amylase; **P20845** GH13 α-amylase; **AmyE** (P00691) GH13 α-amylase; **AmyS** (P06278 and P06279) GH13 α-amylase; **P00692** GH13 α-amylase; **A1YR25** GH13 α-amylase; **AmyM** (P19531) GH13 maltogenic α-amylase; **GenBank: AAO26743.1** GH13 hyperthermostable α-amylase; **A0A111UDQ2** GH15 glucoamylase; **Cga** A0A3S4RT35 GH15 glucoamylase; **A0A2A9CHF6** GH15 glucoamylase; **A0A328M267** GH15 glucoamylase; **A0A1Q8TTM6** GH15 glycosyl hydrolase; **P19571** GH13 glucan 1/4-α-maltohexaosidase; **P06547** GH14 β-amylase; **P96513** GH14 β-amylase; **SpoII** P36924 β-amylase; **Npr** (P29148) M4 family peptidase bacillolysin (protease); **MdxE** (O06989) maltodextrin-binding protein of the bacterial solute-binding protein 1 family. **YkoJ** (O35012) uncharacterized protein (putative protease inhibitor).

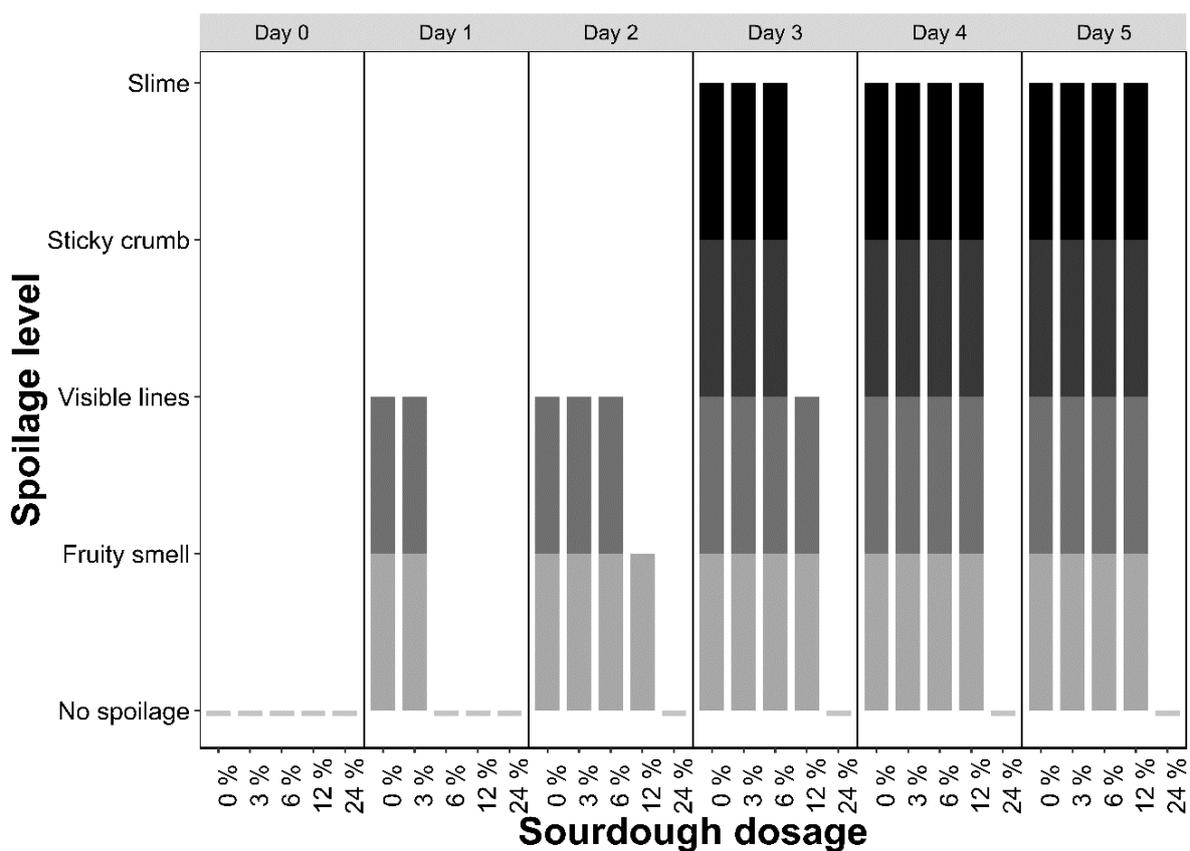


Figure 6. Kinetics of appearance of spoilage characteristics in breads baked from doughs contaminated with *B. amyloliquefaciens* (FAD 99) during storage at 30 °C and a_w 0.95 for 5 days. The doughs contained different dosage of sourdough fermented with reutericyclin-positive *L. reuteri* TMW1.656. Different colors of gray scale represent variations in levels of spoilage from low to high. Different sourdough dosage was labeled in the figure. The pH of bread produced with 0, 3, 6, 12 or 24 % sourdough was 6.40 ± 0.044 ; 6.30 ± 0.019 ; 6.17 ± 0.024 ; 5.93 ± 0.041 ; and 5.53 ± 0.040 , respectively. Data represent results from three independent experiments.

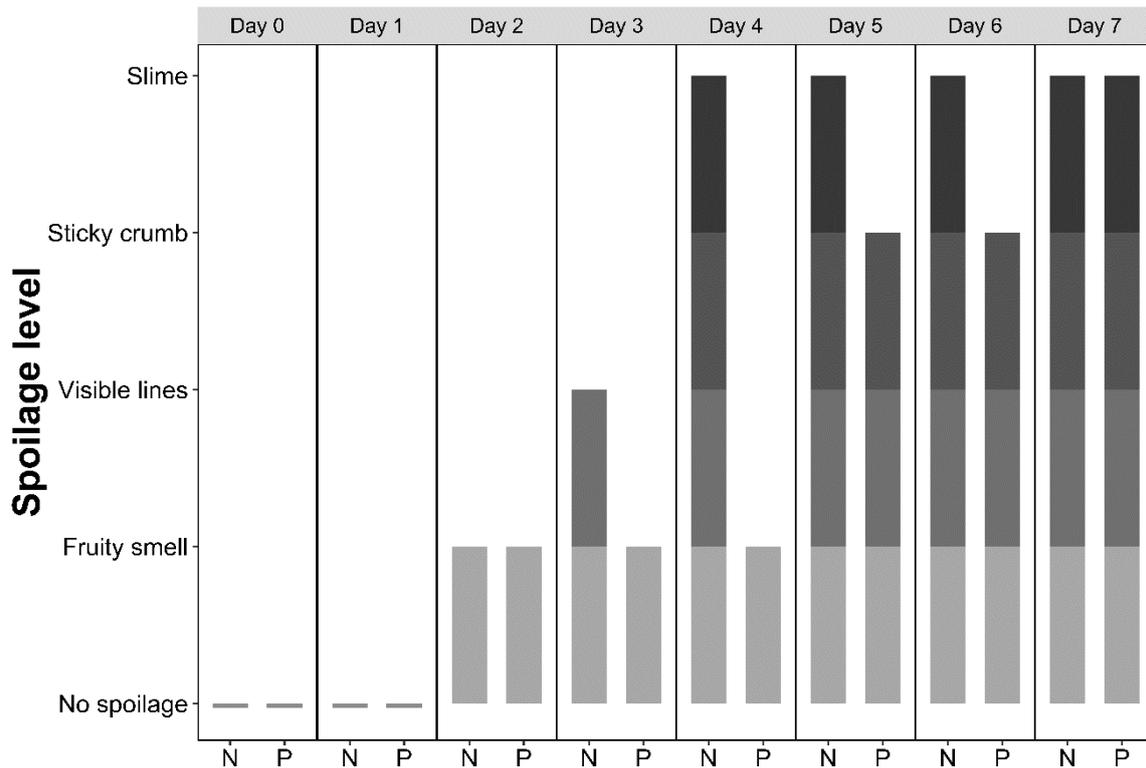


Figure 7. Effect of reutericyclin on the kinetics of appearance of ropy spoilage characteristics of breads baked from doughs contaminated with *B. amyloliquefaciens* FAD 99 during storage at 30 °C and a_w 0.95 for 7 days. Sourdough was added at a dosage of 12 %. Different colors of gray scale represent different levels of spoilage from low to high. “N” represented bread produced with sourdough fermented with the reutericyclin-negative *L. reuteri* TMW1.656 Δ *rtcN* Δ *gtfA*; “P” represented bread produced with sourdough fermented with the reutericyclin-producing *L. reuteri* TMW1.656. The pH of bread produced with sourdough fermented with *L. reuteri* TMW1.656 and *L. reuteri* TMW1.656 Δ *rtcN* Δ *gtfA* was 5.93 ± 0.041 and 5.93 ± 0.021 , respectively. Data represent results from three independent experiments.