

1 **Antimicrobial Activity of Bioactive Starch Packaging Films Against *Listeria monocytogenes***  
2 **and Reconstituted Meat Microbiota on Ham**

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21 **Abstract**

22 Contamination with spoilage organisms and *Listeria monocytogenes* are major concerns  
23 for quality and safety of cooked ready-to-eat (RTE) meat products. Thus, the objective of this  
24 study was to investigate the use of antimicrobial starch packaging films to control competitive  
25 microbiota and *L. monocytogenes* growth on a RTE ham product. Starch packaging films were  
26 prepared with different bioactives, gallic acid, chitosan, and carvacrol, using subcritical water  
27 technology. The viability of the incorporated strains on ham in contact with different antimicrobial  
28 starch packaging films was examined during 28-day storage period at 4 °C. Starch films with gallic  
29 acid had the least effect on ham antimicrobial activity; starch films with chitosan and carvacrol  
30 fully inhibited *L. monocytogenes* growth throughout 4 weeks of storage. RTE meat microbiota was  
31 more resistant to the antimicrobials than *L. monocytogenes*. Starch films loaded with chitosan or  
32 chitosan and carvacrol did not fully inhibit growth of RTE meat microbiota but delayed growth of  
33 RTE meat microbiota by one to two weeks. Moreover, competitive meat microbiota fully inhibited  
34 growth of *L. monocytogenes*. Therefore, antimicrobial starch packaging films prepared by  
35 subcritical water technology used in this study showed a promising effect on inhibiting *L.*  
36 *monocytogenes* in RTE ham.

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38 **Keywords:** Antimicrobial starch films; *Carnobacterium*; *Leuconostoc*; *Brochotrix*; *Listeria*  
39 *monocytogenes*; ready-to-eat meat; chitosan.

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## 42 **1. Introduction**

43 Ready to eat (RTE) foods including RTE meats represent a growing segment of the overall  
44 food market, owing to their convenient use by consumers (Alberta Agriculture and Forestry, 2017).  
45 The main food safety concern related to RTE meat products is contamination with *Listeria*  
46 *monocytogenes*, which may grow to high cell counts during refrigerated storage (Yousef & Lou,  
47 1999), and cause life-threatening infections in at-risk individuals (Farber & Peterkin, 1991; WHO,  
48 2004). Because RTE meats are typically consumed without further cooking, the risk of infection  
49 depends on the cell counts of *L. monocytogenes* on the product. Cell counts ranging from 0.04 to  
50 100 CFU *L. monocytogenes* / g are considered an acceptable risk (FSIS, 1989; WHO, 2004). The  
51 contamination of RTE meats is primarily attributed to post-cooking contamination. In addition to  
52 process hygiene, the addition of preservatives to RTE meats to prevent growth of *L.*  
53 *monocytogenes* is a key measure to reduce the risk of foodborne listeriosis (Mejlholm et al., 2010).

54 Microbiota of RTE meats predominantly consists of *Brochothrix thermosphacta* (Miller et  
55 al., 2014), *Carnobacterium* spp. (Horita et al., 2017), psychrotrophic lactobacilli (Giello et al.,  
56 2018) and *Leuconostoc* spp. (Maksimovic et al., 2018). These bacteria can cause discoloration, gas  
57 and slime production, or produce off-odors and off-flavors (Borch et al., 1996; Pothakos et al.,  
58 2014a). However, many strains of *Lactobacillus* spp. and *Carnobacterium* grow to high cell counts  
59 without negatively affecting product quality. Some strains are used as (bacteriocin-producing)  
60 biopreservatives to inhibit growth of *Listeria* during refrigerated storage (Drider et al., 2006;  
61 Nilsson et al., 2005; Schillinger et al., 1991).

62 Common methods used to control microbial contamination of RTE meats include in-  
63 package thermal pasteurization, high pressure processing, and product-reformulation with  
64 preservatives (Murphy et al., 2003; Seman et al., 2002; Teixeira et al., 2016). In-package thermal

65 pasteurization eliminates *L. monocytogenes* but also increases shrinkage and drip loss in the  
66 products (Murphy et al., 2003). Current commercial high pressure processes reduce cell counts of  
67 *L. monocytogenes* only by 4 log (CFU/g) (Teixeira et al., 2016) and thus require combination with  
68 high hygienic processing standards, or with other antimicrobial agents, such as nisin or essential  
69 oils (de Oliveira et al., 2015; Hereu et al., 2012). Antimicrobials such as sodium lactate, sodium  
70 diacetate, and potassium benzoate have extensively been used to extend the shelf-life and ensure  
71 the safety of meat products (Seman et al., 2002). Some new natural derived antimicrobial agents  
72 for use in meat products include phenolic compounds (Starčević et al., 2015), essential oils  
73 (Sirocchi et al., 2017) and chitosan (Arslan & Soyer, 2018). Preservatives, however, also affect  
74 the sensory quality of the products.

75         Microbial contamination of RTE meats occurs at the surface, therefore, the use of natural  
76 antimicrobials in packaging films can control spoilage and pathogenic microorganisms on the  
77 product. Chitosan is a film-forming cationic polysaccharide with antimicrobial activity, which is  
78 suitable for production of antimicrobial packaging films. The use of chitosan-based active  
79 packaging films reduced cell counts of *Listeria*, or inhibited growth of spoilage microbiota on RTE  
80 meats and salmon (Benabbou et al., 2018; Guo et al., 2014; Zhao et al., 2018). Also, the addition  
81 of rosemary and licorice extract to packaging films delayed growth of *L. monocytogenes* on cooked  
82 ham (Zhang et al., 2009). Preliminary studies that assessed the antimicrobial activity of chitosan-  
83 gelatine films on microbiota of cod demonstrated differential activity of the film against different  
84 groups of bacteria (Gómez-Estaca et al., 2010); however, studies that document the differential  
85 activity of chitosan-starch based films on *L. monocytogenes* and spoilage or protective RTE  
86 microbiota are currently unavailable. Therefore, the aim of this study was to investigate the effect  
87 of bioactive starch packaging films, containing gallic acid, or chitosan and gallic acid or carvacrol,

88 for RTE ham on growth of *L. monocytogenes* and reconstituted meat microbiota. The RTE ham  
89 was produced according to current commercial practice in Canada (Teixeira et al., 2016), cut  
90 aseptically, and inoculated with a 5 strains cocktail of *L. monocytogenes* and/or a 5 strain cocktail  
91 representing microbiota of RTE meats.

## 92 **2. Materials and Methods**

### 93 **2.1 Bacterial strains and growth conditions**

94 The human disease cocktail containing 5 strains of *L. monocytogenes* (FSL J1-177, FSL  
95 C1-056, FSL N3-013, FSL R2-499, and FSL N1-227) (Fugett et al., 2006) and a “reconstituted  
96 meat microbiota” cocktail containing *Brochothrix thermosphacta* FUA3558, *Carnobacterium*  
97 *maltaromaticum* FUA3559, *Leuconostoc gelidum* FUA3560 and FUA3561 and *Lactobacillus*  
98 *sakei* FUA3562 (Teixeira et al., 2018) were used in this study.

99 Strains of *L. monocytogenes* were aseptically streaked from -80 °C stock cultures onto  
100 Tryptic Soy (TS) agar (Difco, Becton–Dickinson, Sparks, MD, USA), followed by inoculation into  
101 TS broth (TSB) and incubation overnight at 37 °C. Fresh broth was inoculated with 1% (v/v) of  
102 the overnight culture and incubated at 37 °C to the stationary growth phase. Strains of reconstituted  
103 meat microbiota were prepared in the same manner but grown on All Purpose Tween (APT) agar  
104 and broth at 25 °C. For preparation of cocktails, an equal volume of each individual culture was  
105 mixed to form a 5-strain cocktail of *L. monocytogenes* or reconstituted meat microbiota. These  
106 cocktails were harvested by centrifugation (7000 × g for 10 min), re-suspended in saline solution  
107 containing 8.5 g / L NaCl and diluted. Media and incubation conditions for the organisms are  
108 summarized in Table 1.

### 109 **2.2 Antimicrobial compounds**

110 Gallic acid (GA) (97.5-102.5% titration), chitosan (75-85% deacetylated) with medium  
111 molecular weight of 190-310 kDa and carvacrol (food grade, >99%) were obtained from Sigma  
112 Aldrich (Oakville, ON, Canada). Gallic acid stock solution (22.5 g/L) was prepared in sterilized  
113 distilled water. Chitosan stock solution (11.25 g/L) was prepared in 2% (w/w) citric acid solution  
114 and carvacrol stock solution (56.56 g/L) was prepared in 0.8% (w/w) lecithin solution.

### 115 **2.3 Determination of the combined activity of gallic acid or carvacrol and chitosan with the** 116 **checkerboard method**

117 The checkerboard procedure was carried out to determine the combination of inhibitory  
118 and bactericidal activity of gallic acid or carvacrol and chitosan against *L. monocytogenes* and  
119 reconstituted meat microbiota. Briefly, 100  $\mu$ L of TS or APT broth was added to each well of a  
120 96-well microplate. Combinations of gallic acid + chitosan or carvacrol + chitosan stock solutions  
121 (100  $\mu$ L) were added to separate wells and serially 2-fold diluted across the plate in a two-  
122 dimensional way. Stationary phase cultures of *L. monocytogenes* or reconstituted meat microbiota  
123 were diluted in TS or APT broth to obtain a cell count of about  $10^8$  CFU/mL. Each well of the  
124 microplates were inoculated with 50  $\mu$ L of these diluted cultures. Plates were incubated for 24 h  
125 at 37 °C for *Listeria* or 25 °C for reconstituted meat microbiota.

### 126 **2.4 Preparation of antimicrobial starch films**

127 Bioactive starch packaging films were prepared as described by Zhao et al. (2018). Briefly,  
128 antimicrobials (gallic acid, chitosan, and carvacrol essential oil), cassava starch, potato cull (15.2%  
129 starch purity, wet basis), glycerol, and water were loaded into the subcritical fluid reactor (270  
130 mL). The mixture was homogenized for 5 min before the desired temperature and pressure were  
131 reached. Then, the reaction was performed for 10 min, followed by cooling. After unloading and  
132 degassing, the solution was transferred into a plastic petri dish of 15 cm diameter and dried at

133 40 °C for 48 h. Subsequently, the dried film was conditioned at 40% RH and 25 °C for at least  
134 48 h. Formulations used for antimicrobial film formation are shown in Table 2.

## 135 **2.5 Sample preparation and inoculation**

136 Previously manufactured experimental cooked ham, with a known formulation and sodium  
137 chloride concentration of 3% (w/w), was used in this study (Teixeira et al., 2016). The ham was  
138 sliced aseptically. Un-inoculated slices of ham had a total aerobic plate count of less than 100  
139 CFU/cm<sup>2</sup> after slicing. Individual slices of ham (50 cm<sup>2</sup> surface area with 3 mm thickness) were  
140 surface inoculated with the cocktail of *L. monocytogenes* and/or the cocktail of reconstituted meat  
141 microbiota to achieve cell counts of about 10<sup>3</sup> CFU *Listeria*/cm<sup>2</sup> and/or 10<sup>4</sup> CFU reconstituted  
142 meat microbiota/cm<sup>2</sup>. Experimental groups were categorized as: (i) *L. monocytogenes*, (ii)  
143 reconstituted meat microbiota, and (iii) *L. monocytogenes* combined with reconstituted meat  
144 microbiota. Each of the three experimental groups was covered with the antimicrobial films with  
145 2 cm<sup>2</sup> surface area (Table 2). Samples were aseptically packed, sealed and stored at 4 °C for up to  
146 28 days. Un-inoculated ham served as the control and surface plating of control samples on APT,  
147 TS and PALCALM agars verified that the plate counts of control samples remained below the  
148 detection limit of 100 CFU/cm<sup>2</sup> throughout 28 days of storage. Detection of surviving cells was  
149 determined by surface plating as described below. Experiments were performed in triplicate.

## 150 **2.6 Sampling and quantification of surviving cells**

151 The presence or absence of *L. monocytogenes* and/or reconstituted meat microbiota was  
152 monitored after 0, 7, 14, 21 and 28 days of storage at 4 °C. Also, un-inoculated ham samples were  
153 prepared and stored for 28 days at 4 °C to ensure the absence of contaminating microbiota from  
154 the meat prior to the experiment and after storage. Samples were opened aseptically, film and ham  
155 were collected by coring with a sterile corer. The cores with a 2 cm<sup>2</sup> surface area were transferred

156 to a sterile 50 mL centrifuge tube and diluted with sterile saline (0.85% NaCl). Samples were  
157 homogenized for 60 s prior to serial 10-fold dilutions in sterile saline.

158 Surviving cells were determined by surface plating on selective PALCAM (Becton-  
159 Dickinson) agar (*L. monocytogenes* combined with reconstituted meat microbiota) and on non-  
160 selective TS (*L. monocytogenes*) or APT agar (reconstituted meat microbiota and *L.*  
161 *monocytogenes* combined with reconstituted meat microbiota). Appropriate dilutions were plated  
162 and incubated at 37 °C (PALCAM and TS agar) or 25 °C (APT agar) for 48 h. The limit of  
163 detection was 100 cfu/cm<sup>2</sup>.

## 164 **2.7 Extraction of total DNA and PCR**

165 For microbial analysis, 1 mL aliquot of the homogenate wash from samples stored for 28  
166 days at 4 °C was centrifuged (5000 × *g* for 10 min) to collect bacterial cells, and total DNA was  
167 extracted from the pellet using DNeasy Blood and Tissue Kit (Qiagen, ON, Canada) following the  
168 Gram-positive bacteria protocol provided by the manufacturer. The DNA was amplified by PCR  
169 with Taq DNA polymerase and dNTPs from Invitrogen (Burlington, ON, Canada).

170 Species-specific primers for characterization of meat microbiota were purchased from  
171 Integrated DNA Technologies (IDT; Coralville, IA, USA) and are listed in Table 3. Species-  
172 specific primers for *Leuconostoc gelidum*, LMG4-F and LMG4-R, were identified by alignment  
173 of reference genomes using Mauve (Darling et al., 2004). Species-specific primers LMG4-F and  
174 LMG4-R were designed targeting unique sequences using PrimerQuest Tool (IDT). The  
175 specificity of the candidate primers was confirmed by Nucleotide BLAST  
176 (<https://blast.ncbi.nlm.nih.gov/Blast.cgi>) and 1% agarose gel after PCR. The PCR products were  
177 visualized after electrophoretic separation on agarose gels.

## 178 **2.8. Statistical analysis**

179 All experiments were performed in triplicate. The RStudio software (Version 0.99.903,  
180 RStudio, Inc., Boston, MA, USA) was used to conduct the analysis of variance (ANOVA).  
181 Significant differences were identified with Tukey's test as post-hoc analysis at an error probability  
182 of 5% ( $p < 0.05$ ).

### 183 **3. Results**

#### 184 **3.1. Inhibitory activity of gallic acid or carvacrol as a function of chitosan concentration** 185 **against *L. monocytogenes* and reconstituted meat microbiota**

186 To determine the relative activity of gallic acid and carvacrol against the 10 strains of  
187 *Listeria* and RTE microbiota, their inhibitory effect was determined alone and in combination with  
188 chitosan. At 1.875 g/L, chitosan alone inhibited all strains of *L. monocytogenes*. Gallic acid showed  
189 higher MIC values (15 g/L) than carvacrol (0.61 g/L). Carvacrol and chitosan acted synergistically  
190 in *Listeria* inhibition as shown in Figure 1B by the pronounced convex shape of the curve, while  
191 synergistic activity of gallic acid and chitosan was less pronounced as observed in Figure 1A.

192 Reconstituted meat microbiota was less sensitive to all antimicrobial compounds (Figure  
193 2). Chitosan alone inhibited meat microbiota at 7.5 g/L, which is four times higher than the MIC  
194 against *L. monocytogenes* (1.875 g/L, Figure 1). The MIC of carvacrol (1.22 g/L, Figure 2B) and  
195 gallic acid (15 g/L, Figure 2A) did not inhibit the reconstituted meat microbiota. Even in  
196 combination with 3.75 g/L chitosan, gallic acid at the highest concentration did not inhibit all  
197 strains representing meat microbiota (Figure 2A). But, carvacrol exhibited additive activity with  
198 chitosan (Figure 2B).

#### 199 **3.2. Inhibition of *L. monocytogenes* or reconstituted meat microbiota on ham by bioactive** 200 **starch films**

201 Chitosan was incorporated at 0.025 or 0.150 g/g starch as antimicrobial agent in cassava  
202 starch films to provide antimicrobial activity. Films containing 0.1 g gallic acid/g starch or up to  
203 0.195 g carvacrol/g starch were also prepared; the addition of gallic acid or carvacrol was based  
204 on their *in vitro* antimicrobial activity. Packaging films were also produced from cull potatoes,  
205 a starch-rich by-product of the potato processing, alone or with addition of gallic acid (Zhao &  
206 Saldaña, 2019). The inhibition of *L. monocytogenes* on ham is shown in Figure 3. Cell counts on  
207 TS (Figure 3A) and PALCAM (Figure 3B) agar were not different, indicating that *L.*  
208 *monocytogenes* on ham were not sublethally injured. Cell counts of un-inoculated ham remained  
209 below the detection limit of 100 cfu/cm<sup>2</sup> throughout 4 weeks of storage, confirming that the aseptic  
210 ham was free of contaminants that would interfere with interpretation of results. For ham packaged  
211 with cassava starch films or films from cull potatoes, *L. monocytogenes* grow to high cell counts  
212 after 21 d of storage at 4°C. Addition of up to 0.3 g gallic acid/g starch delayed growth of *L.*  
213 *monocytogenes* by one week (Figures 3a and 3b). Starch films containing chitosan and gallic acid  
214 inhibited growth throughout four weeks of refrigerated storage (Figure 3a and 3b); however, *L.*  
215 *monocytogenes* was detected on at least one of the three replicates in all samples. Starch films with  
216 chitosan and carvacrol also inhibited growth of *L. monocytogenes* throughout 4 weeks of storage.  
217 Incorporation of carvacrol at 0.195 g/g starch reduced initial cell counts by 0.5 log CFU/cm<sup>2</sup> but  
218 *L. monocytogenes* remained detectable in one of the three replicates throughout 4 weeks of storage.

219 Consistent with the *in vitro* MIC data, reconstituted meat microbiota was more resistant to  
220 starch films containing gallic acid, or chitosan with gallic acid or carvacrol (Figure 4). For the ham  
221 covered with starch films without antimicrobials, reconstituted meat microbiota grew to high cell  
222 counts after two weeks of refrigerated storage. The growth of reconstituted meat microbiota on  
223 ham covered with 0.1 g gallic acid/g starch packaging film was comparable to the cull potato

224 control but addition of 0.3 g gallic acid/g starch to the packaging film delayed growth of meat  
225 microbiota. Adding 0.1 g gallic acid/g starch in combination with 0.025 or 0.15 g chitosan/g starch  
226 delayed growth of reconstituted meat microbiota by one or two weeks. For ham covered with films  
227 containing both carvacrol and chitosan, the initial cell counts of reconstituted meat microbiota  
228 were reduced by 1 – 2 log (CFU/cm<sup>2</sup>) and re-growth of the organisms was delayed. Cell counts on  
229 ham covered with film containing chitosan and 0.195 g carvacrol/g starch remained below 7 log  
230 (CFU/cm<sup>2</sup>). The antimicrobial packaging film, however, did not completely eliminate or fully  
231 inhibit reconstituted meat microbiota during refrigerated storage of 28 days.

### 232 **3.3. Inhibition of combined inocula of *L. monocytogenes* and reconstituted meat microbiota** 233 **on ham**

234 To understand the influence of antimicrobial packaging films on the interaction of  
235 reconstituted meat microbiota and *L. monocytogenes*, ham was inoculated with the mixture of a  
236 cocktail of 5 *L. monocytogenes* strains and a cocktail of 5 reconstituted meat microbiota (Figure  
237 5). Cell counts on ham were predominantly attributable to reconstituted meat microbiota. An initial  
238 cell count reduction of 1.5 log (CFU/cm<sup>2</sup>) was observed on ham with films containing carvacrol  
239 or chitosan. Total cell counts on ham covered with starch film containing 0.1 g gallic acid/g starch  
240 showed no difference to cull potato control after 14 days of storage, while the addition of 0.3 g  
241 gallic acid/g starch delayed bacterial growth (Figure 5A). The use of 0.1 g gallic acid/g starch  
242 combined with 0.025 or 0.15 g chitosan/g starch film and films with 0.025 g chitosan/g starch and  
243 0.048 g carvacrol/g starch reduced total viable plate counts by 1-1.5 log (CFU/cm<sup>2</sup>). The most  
244 pronounced inhibitory effect was observed on ham covered with starch films containing both  
245 chitosan and carvacrol. In these products, the cell counts of *L. monocytogenes* were at or below  
246 the limit of detection (100 cfu/cm<sup>2</sup>) (Figure 5B) or about 7 log (CFU/cm<sup>2</sup>) lower when compared

247 to the growth of *L. monocytogenes* to  $10^9$  cfu/cm<sup>2</sup> on ham packaged out addition of antimicrobials  
248 or competing microbiota (Figure 3B).

249 Reconstituted meat microbiota reduced growth of *L. monocytogenes* even in the absence  
250 of antimicrobials in the packaging films (Figure 5b). Growth of *L. monocytogenes* was also  
251 delayed on ham covered with cull potato starch film containing 0.1 g gallic acid/g starch. When  
252 combined with the reconstituted meat microbiota, gallic acid (0.3 g/g starch), chitosan or carvacrol  
253 completely inhibited growth of *L. monocytogenes* but *L. monocytogenes* remained detectable in  
254 one of the three replicates throughout 4 weeks of storage.

### 255 **3.4. Prevalence of individual strains of reconstituted meat microbiota on ham**

256 Because different bacterial species differ with respect to their impact on product quality,  
257 dominant meat microbiota on ham at different storage times was therefore identified after isolation  
258 of community DNA from the surface of the ham, followed by species-specific or genus-specific  
259 PCR (Table 4). The primers readily differentiated *B. thermosphacta*, *C. maltaromaticum*, *Lc.*  
260 *gelidum* and *Lb. sakei*, however, the two strains of *Lc. gelidum* were not differentiated from each  
261 other. *Lc. gelidum* was predominant in all populations collected from ham covered with different  
262 antimicrobial packaging films. Consistent with the MIC and cell counts data, packaging films with  
263 gallic acid had little impact on the composition of meat microbiota. On the ham covered with starch  
264 / gallic acid film, all four species included in the strain cocktail were detected after 28 d of storage;  
265 however, the starch / gallic acid film inhibited *C. maltaromaticum* in one of the three replicates  
266 (Table 4). In contrast, inclusion of chitosan into starch films inhibited all meat microbiota with  
267 exception of *Lc. gelidum*. After 28 d of storage of ham covered with any of the films containing  
268 chitosan in combination with gallic acid or carvacrol, *Lc. gelidum* was the only organism detected.

## 269 **4. Discussion**

270 RTE ham is processed prior to final packaging, and is consumed without further cooking;  
271 therefore, contamination with spoilage organisms and pathogens prior to packaging determines the  
272 storage life and the safety of the product. Antimicrobial packaging provides an additional hurdle  
273 for inhibition of contaminants. Laboratory tests of packaging films with culture media or model  
274 foods may not accurately predict the *in situ* inhibitory effect (Dutta et al., 2009; Ramos et al., 2012;  
275 Sun et al., 2014). This study therefore evaluated the antimicrobial efficiency of bioactive starch  
276 packaging films on a meat product. Various natural bioactive agents were effective in laboratory  
277 applications but did not show antibacterial activity in food because they were rendered inactive by  
278 the specific characteristics of the food and storage conditions (Malhotra et al., 2015). The  
279 antimicrobial activity of essential oils relates to their hydrophobicity, which enables them to pass  
280 through the cell membrane (Dorman & Deans, 2000). The fat content of the food matrix, however,  
281 strongly influences their activities by increasing the diffusion path length or sequestering (Weiss  
282 et al., 2015). Other components in food, e.g. proteins, may bind phenolic compounds, lowering the  
283 amount available for controlling microbial growth (Tassou et al., 2000). Also, chitosan activity  
284 may be compromised through ionic interactions with food components (Hu and Gänzle, 2018).

285 To account for intra-species differences of pathogenic bacteria in resistance to intervention  
286 technologies, novel food preservation technologies are generally validated with strain cocktails  
287 and are considered effective only if all strains are inhibited or eliminated (Hoque et al., 2008;  
288 Solomakos et al., 2008). Moreover, antimicrobial interventions differentially affect the  
289 competitiveness of non-pathogenic meat microbiota (Teixeira et al., 2018), which may influence  
290 spoilage of RTE meats. The strain cocktail used in the present study to reconstitute meat microbiota  
291 represents the diversity of microorganisms that are normally found in RTE meat products. Among

292 150 bacterial isolates from commercially available RTE meats, *Lc. gelidum*, *C. maltaromaticum*,  
293 *Lb. sakei* and *B. thermosphacta* accounted for more than 90% of the isolates (Miller et al., 2014).

294 The cell counts on ham conformed to MICs data and reconstituted meat microbiota were  
295 less sensitive to all antimicrobials when compared to *Listeria*. Among the three antimicrobials  
296 evaluated, gallic acid showed the least antimicrobial activity against both *L. monocytogenes* and  
297 reconstituted meat microbiota due to the presence of 3 hydroxyl groups in gallic acid, these  
298 increase gallic acid polarity and reduce its capacity to cross the cell membrane (Sánchez-  
299 Maldonado et al., 2011). High MIC values (>5 mM) for gallic acid were also reported at different  
300 pH values of 5 to 7 (Miyague et al., 2015).

301 Adding chitosan to starch films achieved complete inhibition of *L. monocytogenes* and cell  
302 counts remained below 100 CFU/cm<sup>2</sup> throughout the storage life of the products. This low cell  
303 counts meet the requirements of the regulation to guarantee food safety and extend storage shelf  
304 life (WHO, 2004). The few studies that used chitosan in packaging films to control pathogen in  
305 food demonstrate that their lethality is limited to a reduction of viable cell counts by less than 1 –  
306 2 log (CFU/cm<sup>2</sup>) (Hu and Gänzle, 2018). For example, *L. monocytogenes* exposed to 0.3% chitosan  
307 impregnated LDPE films recovered to levels of control films after 12 h exposure (Park et al., 2010).  
308 Chitosan films reduced cell counts of *Listeria innocua* or *L. monocytogenes* by only 0.8 or 1 log  
309 CFU/cm<sup>2</sup> on RTE deli turkey meat or black radish, respectively (Guo et al., 2014; Jovanović et al.,  
310 2016). The current use of preservatives, however, does not aim to eliminate *L. monocytogenes*;  
311 potassium lactate and sodium diacetate addition to processed meats in combination with process  
312 hygiene inhibits growth and thus maintains low cell counts throughout the storage life of the  
313 products (Stekelenburg & Kant-Muermans, 2001). Therefore, the use of chitosan in starch films

314 in our study demonstrated the potential application on RTE meat to inhibit *Listeria* without use of  
315 preservative additives to the RTE meat product.

316 Gram-positive bacteria are generally more sensitive to essential oils than gram-negative  
317 bacteria, and *L. monocytogenes* was among the most sensitive organisms (Gutierrez et al., 2008).  
318 Even at minimum carvacrol concentration (0.048 g/g starch) used in the film formulation,  
319 carvacrol essential oil completely inhibited *L. monocytogenes*. Concentrations of carvacrol that  
320 exceed the flavor threshold, however, may negatively impact sensory properties of the ham.  
321 Rosemary and thyme essential oils released from the sachet restricted the growth of *L.*  
322 *monocytogenes* on mozzarella cheese, resulting in a 2.5 log CFU/g reduction on day 9 at 10 °C  
323 (Han et al., 2014). Chitosan films with 1% and 2% oregano essential oil decreased the cell count  
324 of *L. monocytogenes* on bologna slices by 3.6 to 4 logs (Zivanovic et al., 2005). But, none of them  
325 showed complete inhibition of *L. monocytogenes*.

326 In our study, reconstituted meat microbiota competed with *L. monocytogenes* and inhibited  
327 its growth. Inhibition of *L. monocytogenes* by microbial antagonism of lactic acid bacteria in meat  
328 was previously reported (Balay et al., 2017; Chaillou et al., 2014; Woraprayote et al., 2016). Fast  
329 growth rates at refrigeration temperatures; nutrient depletion, acid production and the strain-  
330 specific production of bacteriocins contribute to inhibition of *L. monocytogenes* by lactic acid  
331 bacteria on meat (Cornu et al., 2011; Woraprayote et al., 2016). These factors make lactic acid  
332 bacteria promising biopreservatives for replacement of chemical preservatives, however, some of  
333 the lactic acid bacteria also contribute to spoilage by formation of off-odours or slime. Depending  
334 on the type of organism growing on RTE ham, a cell count of  $10^6$  to  $10^7$  CFU/cm<sup>2</sup> may lead to  
335 spoilage (Fung, 2009). Rot or acid odours produced by *B. thermosphacta* decrease consumer  
336 acceptance (Vermeiren et al., 2005). *Leuconostoc* species spoil RTE meats by slime production

337 when sucrose is present (Pothakos et al., 2014b). In contrast, *Lb. sakei* and *C. maltaromaticum* did  
338 not impair sensory attributes or consumer acceptance of RTE meat products (Bredholt et al., 2001;  
339 Vermeiren et al., 2005). The present study demonstrates that reconstituted meat microbiota in  
340 combination with antimicrobial starch packaging films inhibited *L. monocytogenes* during 28 d of  
341 refrigerated storage. In these products, cell counts of *L. monocytogenes* remains below 100  
342 CFU/cm<sup>2</sup>. However, chitosan-starch films with gallic acid or carvacrol also selected for *Lc.*  
343 *gelidum* as dominant organism on meat. Because strains of this species spoil meat products by  
344 slime production based on its dextransucrase activity (Pothakos et al., 2014a and 2014b), the use  
345 of chitosan based packaging films may accelerate spoilage if the product formulation includes  
346 sucrose.

347 In conclusion, this challenging antimicrobial test on ham demonstrated the successful use  
348 of antimicrobial starch packaging as an important strategy to control reconstituted meat microbiota  
349 and foodborne pathogens, particularly for RTE meat products. The cell count test data were  
350 coherent with the MIC assay data, where antimicrobial starch packaging films with gallic acid  
351 were the least effective antimicrobial. Among all formulations, starch packaging films with  
352 chitosan and carvacrol exhibited strong effects against *L. monocytogenes* and meat reconstituted  
353 meat microbiota. *L. monocytogenes* growth was successfully inhibited during the storage period  
354 of 4 weeks. However, reconstituted meat microbiota was less sensitive, especially using the gallic  
355 acid incorporated films. Bioactive starch films produced by subcritical water technology showed  
356 potential use as antimicrobial packaging films of ham.

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522

523 **Figure legends**

524 **Figure 1.** Minimal inhibitory concentration (g/L) of gallic acid (**Panel A**), and carvacrol (**Panel**  
525 **B**) as a function of chitosan concentration (g/L) for *L. monocytogenes* strains FSL J1-177 (○), FSL  
526 C1-056 (●), FSL N3-013 (□), FSL R2-499 (■), and FSL N1-227 (Δ). Data are means ± standard  
527 deviations of triplicate independent experiments.

528 **Figure 2.** Minimal inhibitory concentration (g/L) of gallic acid (**Panel A**), and carvacrol (**Panel**  
529 **B**) as a function of chitosan concentration (g/L) for *Brochothrix thermosphacta* FUA3558 (○),  
530 *Carnobacterium maltaromaticum* FUA3559 (●), *Leuconostoc gelidum* FUA3560 (□) and  
531 FUA3561 (■), and *Lactobacillus sakei* FUA3562 (Δ). Data are means ± standard deviations of  
532 triplicate independent experiments.

533 **Figure 3.** Growth of a 5 strain cocktail of *L. monocytogenes* strains FSL J1-177, FSL C1-056, FSL  
534 N3-013, FSL R2-499, and FSL N1-227 on the surface of RTE ham during storage at 4 °C. Bacteria  
535 were enumerated on TSB agar (**Panel A**) or on PALCAM agar (**Panel B**). The ham was covered  
536 with cull potato film (Potato control), cull potato films containing 0.1 g or 0.3 g gallic acid/g starch,  
537 cassava starch film (Cassava control), cassava starch films containing 0.1 g gallic acid/g starch  
538 and 0.025 g or 0.15 g chitosan/g starch, or cassava starch films containing 0.048 g or 0.195 g  
539 carvacrol and 0.025 g chitosan/g starch. Data are means ± standard deviations of triplicate  
540 independent experiments. The dotted line indicates the detection limit of 2 log CFU/cm<sup>2</sup>. Cell  
541 counts of un-inoculated ham remained below the detection limit throughout the 4 weeks of storage.  
542 Data points are shown without error bars with a value of 1.5 log(CFU/cm<sup>2</sup>) when viable cell counts  
543 for one or two of the three replicates were below the detection limit.

544 **Figure 4.** Growth of a 5 strain cocktail of reconstituted meat microbiota containing *Brochothrix*  
545 *thermosphacta* FUA3558, *Carnobacterium maltaromaticum* FUA3559, *Leuconostoc gelidum*

546 FUA3560 and FUA3561, and *Lactobacillus sakei* FUA3562 on the surface of cooked ham during  
547 storage at 4 °C, bacteria were counted on APT agar. The ham was covered with cull potato film  
548 (Potato control), cull potato films containing 0.1 g or 0.3 g gallic acid/g starch, cassava starch film  
549 (Cassava control), cassava starch films containing 0.1 g gallic acid/g starch and 0.025 g or 0.15 g  
550 chitosan/g starch, or cassava starch films containing 0.048 g or 0.195 g carvacrol and 0.025 g  
551 chitosan/g starch. Data are means  $\pm$  standard deviations of triplicate independent experiments. The  
552 dotted line indicates the detection limit of 2 log CFU/cm<sup>2</sup>.

553 **Figure 5.** Growth of the mixture of a 5 strain cocktail of reconstituted meat microbiota containing  
554 *Brochothrix thermosphacta* FUA3558, *Carnobacterium maltaromaticum* FUA3559, *Leuconostoc*  
555 *gelidum* FUA3560 and FUA3561, and *Lactobacillus sakei* FUA3562, and a 5 strain cocktail of *L.*  
556 *monocytogenes* strains: FSL J1-177, FSL C1-056, FSL N3-013, FSL R2-499, and FSL N1-227 on  
557 the surface of cooked ham during storage at 4 °C. Bacteria were enumerated on APT agar (**Panel**  
558 **A**) and PALCAM agar (**Panel B**). The ham was covered with cull potato film (Potato control), cull  
559 potato films containing 0.1 g or 0.3 g gallic acid/g starch, cassava starch film (Cassava control),  
560 cassava starch films containing 0.1 g gallic acid/g starch and 0.025 g or 0.15 g chitosan/g starch,  
561 or cassava starch films containing 0.048 g or 0.195 g carvacrol and 0.025 g chitosan/g starch. Data  
562 are means  $\pm$  standard deviations of triplicate independent experiments. The dotted line indicates  
563 the detection limit of 2 log CFU/cm<sup>2</sup>. Cell counts of un-inoculated ham remained below the  
564 detection limit throughout the 4 weeks of storage. Data points are shown without error bars with a  
565 value of 1.5 log(CFU/cm<sup>2</sup>) when viable cell counts for one or two of the three replicates were  
566 below the detection limit.

**Table 1.** Bacterial strains and growth conditions used in this study.

| Strains                                      | Growth conditions           | Reference           |
|--|-----------------------------|---------------------|
| <i>L. monocytogenes</i> FSL J1-177           | Tryptic Soy Broth,<br>37 °C | Fugett et al., 2006 |
| <i>L. monocytogenes</i> FSL R2-499           |                             |                     |
| <i>L. monocytogenes</i> FSL C1-056           |                             |                     |
| <i>L. monocytogenes</i> FSL N1-227           |                             |                     |
| <i>L. monocytogenes</i> FSL N3-013           |                             |                     |
| <i>Brochothrix thermosphacta</i> FUA3558     | All Purpose Tween,<br>25 °C | Miller et al., 2014 |
| <i>Carnobacterium maltaromaticum</i> FUA3559 |                             |                     |
| <i>Leuconostoc gelidum</i> FUA3560           |                             |                     |
| <i>Leuconostoc gelidum</i> FUA3561           |                             |                     |
| <i>Lactobacillus sakei</i> FUA3562           |                             |                     |

**Table 2.** Formulations of antimicrobial films based on potato by-products and cassava starch.

| <b>Sample name</b>                           | <b>Weight of potato cull (g)</b>    | <b>Glycerol/cull starch ratio (g/g)</b> | <b>Gallic acid/starch ratio (g/g)</b> |                                    |
|--|-------------------------------------|---|---------------------------------------|------------------------------------|
| <b>Potato by-product control<sup>1</sup></b> |                                     |   | 0:1                                   |                                    |
| <b>GA1</b>                                   | 36                                  | 1:1                                     | 0.1:1                                 |                                    |
| <b>GA2</b>                                   |                                     |   | 0.3:1                                 |                                    |
|  | <b>Weight of cassava starch (g)</b> | <b>Glycerol/starch ratio (g/g)</b>      | <b>Gallic acid/starch ratio (g/g)</b> | <b>Chitosan/starch ratio (g/g)</b> |
| <b>Cassava starch control<sup>2</sup></b>    |                                     |   | 0:1                                   | 0:1                                |
| <b>CH1</b>                                   | 13                                  | 0.5:1                                   | 0.1:1                                 | 0.025:1                            |
| <b>CH2</b>                                   |                                     |   | 0.1:1                                 | 0.15:1                             |
|  |                                     | <b>Glycerol/starch ratio (g/g)</b>      | <b>Carvacrol/starch ratio (g/g)</b>   | <b>Chitosan/starch ratio (g/g)</b> |
| <b>Carv1</b>                                 |                                     |   | 0.048:1                               |                                    |
| <b>Carv2</b>                                 |                                     | 0.5:1                                   | 0.195:1                               | 0.025:1                            |

GA: gallic acid, CH: chitosan, Carv: carvacrol.

<sup>1</sup>Zhao Saldaña (2019), <sup>2</sup>Zhao et al. (2018).

**Table 3.** Primers and PCR conditions.

| <b>Species</b>                       | <b>Sequence (5'-3')</b>  | <b>Amplicon size / T<sub>m</sub></b> | <b>Reference or target</b>    |
|--------------------------------------|--|--------------------------------------|-------------------------------|
| <i>Brochothrix thermosphacta</i>     | Bcr3r – GTTGTCCGGAATTATTGGG<br>Bcr3f – CTCCTCTTCTGTCCTCAAG         | 121 bp / 58 °C                       | Pennacchia et al. (2009)      |
| <i>Carnobacterium maltaromaticum</i> | Cpis – TTTATTTTTAATTAAATACCC<br>23S-7 – GGTACTTAGATGTTTCAGTTC      | >500 bp / 48 °C                      | Cailliez-Grimal et al. (2007) |
| <i>Leuconostoc gelidum</i>           | LMG4-F – GTCTACCTTCTTTGCCCTTACA<br>LMG4-R – TTCCAAACGAACCTGGAGATAG | 431 bp / 60 °C                       | 23S rRNA (This study)         |
| <i>Lactobacillus sakei</i>           | 16S – GCTGGATCACCTCCTTTC<br>Ls – ATGAAACTATTAATTGGTAC              | 220 bp / 52 °C                       | Bertheir and Ehrlich, (1999)  |

T<sub>m</sub>, annealing temperature

**Table 4.** Detection of individual strains in reconstituted meat microbiota stored for 28 days.

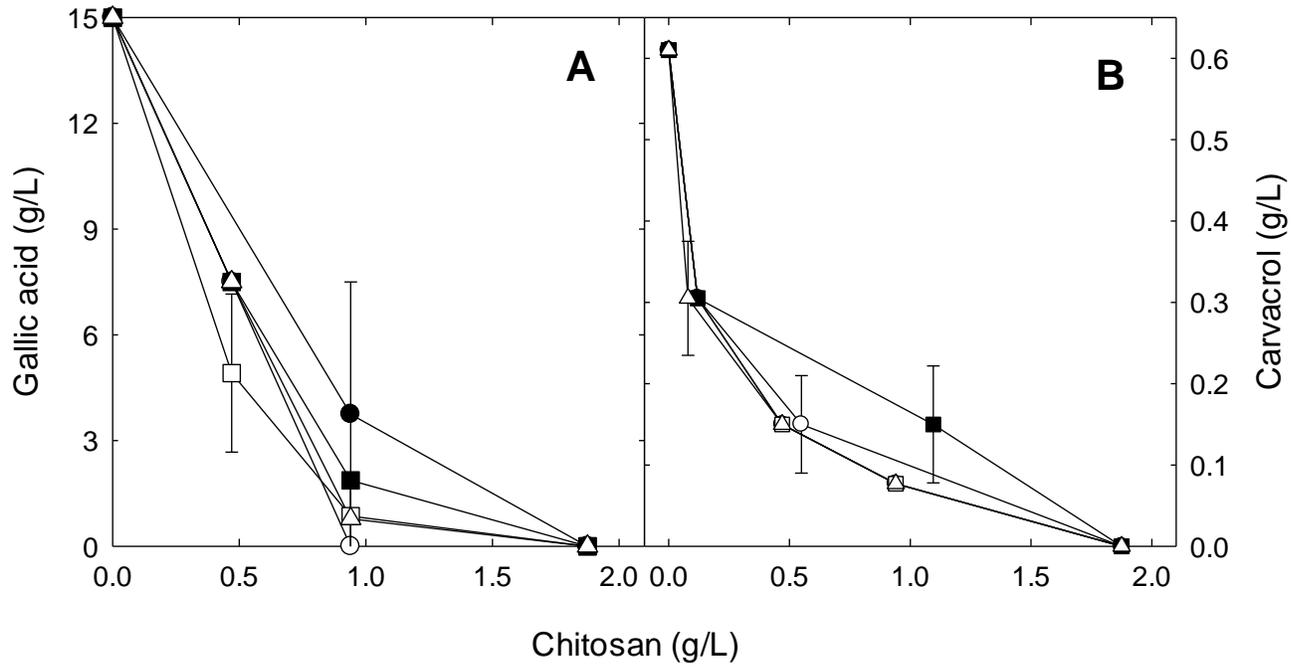
| Species / Films<br>[antimicrobial]           | Cull<br>potato<br>control | Cassava<br>starch<br>control | Gallic acid (g/g) |     | Chitosan (g/g) <sup>1</sup> |      | Carvacrol (g/g) <sup>2</sup> |       |
|--|---------------------------|------------------------------|-------------------|-----|-----------------------------|------|------------------------------|-------|
|  |                           |                              | 0.1               | 0.3 | 0.025                       | 0.15 | 0.048                        | 0.195 |
| <i>Brochothrix thermosphacta</i> FUA3558     | +                         | +                            | +                 | +   | -                           | -    | -                            | -/+   |
| <i>Carnobacterium maltaromaticum</i> FUA3559 | +                         | +                            | +                 | -/+ | -                           | -    | -                            | -     |
| <i>Leuconostoc gelidum</i> FUA3560           | +                         | +                            | +                 | +   | +                           | +    | +                            | +     |
| <i>Leuconostoc gelidum</i> FUA3561           | +                         | +                            | +                 | +   | +                           | +    | +                            | +     |
| <i>Lactobacillus sakei</i> FUA3562           | +                         | +                            | +                 | +   | -                           | -    | -                            | -/+   |

Abbreviations: (+) present; (-) absent; (-/+) positive in one of the triplicates.

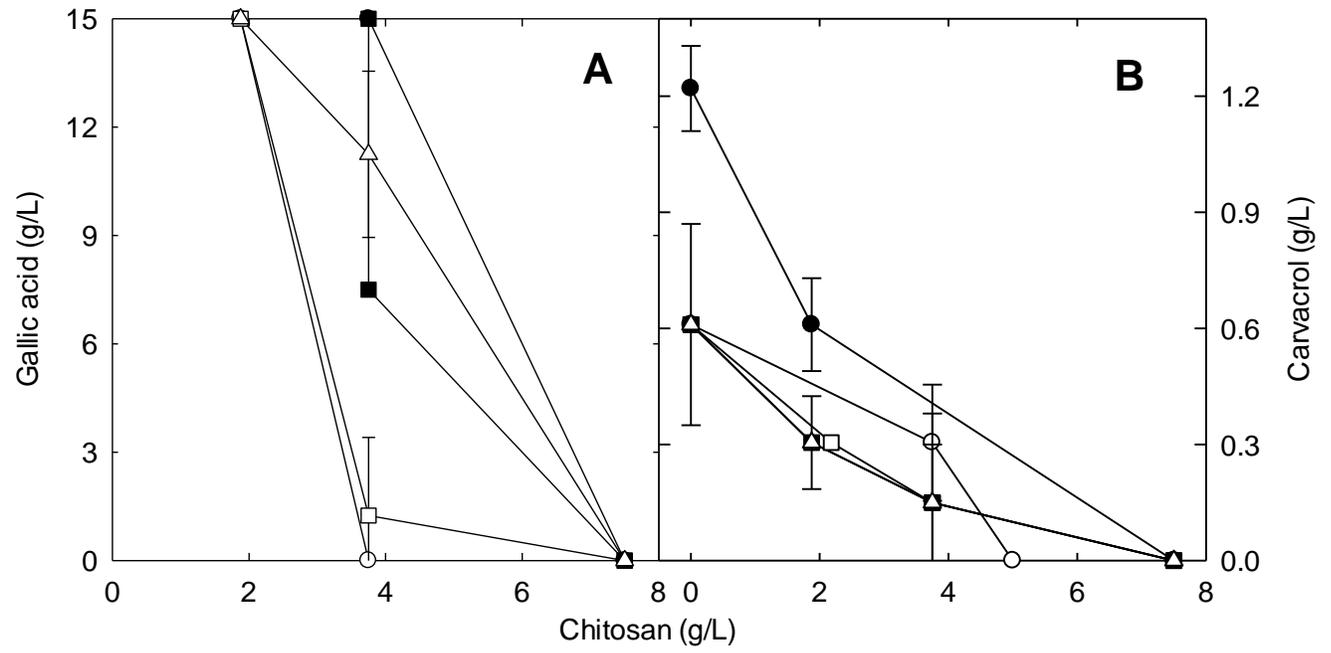
<sup>1</sup>: Cassava starch-based films containing gallic acid concentration at 0.1 g /g starch and 0.025 g or 0.15 g chitosan/g starch.

<sup>2</sup>: Cassava starch-based films containing chitosan concentration at 0.025 g /g starch and 0.048 g or 0.195 g carvacrol/g starch.

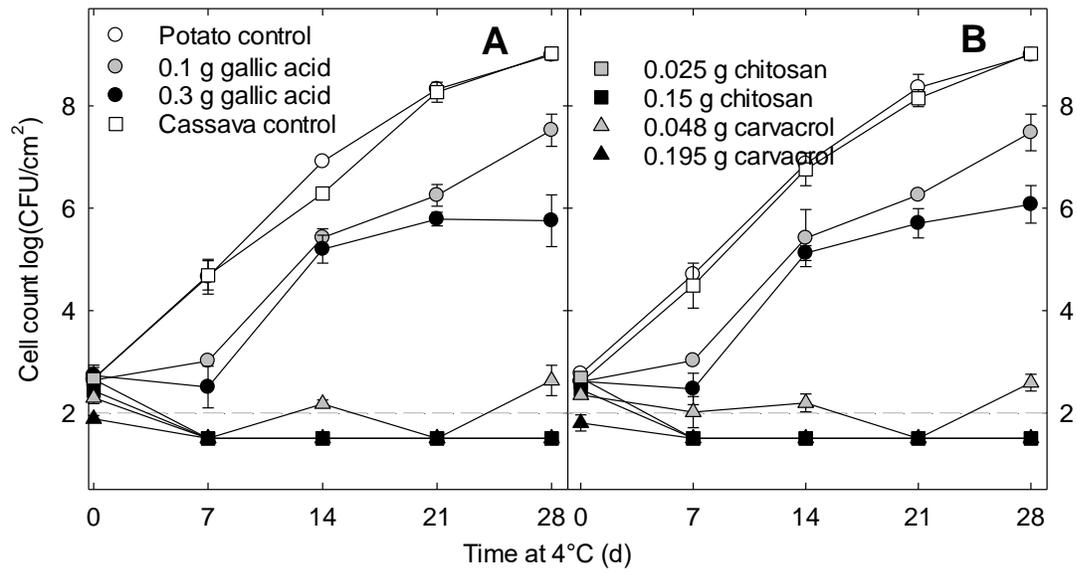
Zhao et al., Figure 1



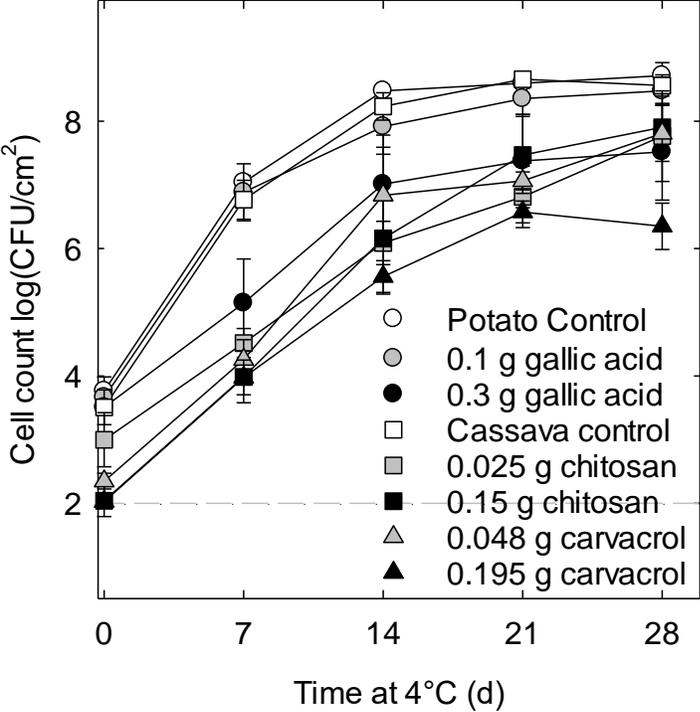
Zhao et al., Figure 2.



Zhao et al., Figure 3



Zhao et al., Figure 4



Zhao et al., Figure 5

