



23 **Abstract**

24 *Enterobacteriaceae* comprise food spoilage organisms as well as food-borne pathogens including  
25 *Escherichia coli*. Heat resistance in *E. coli* was attributed to a genomic island called the locus of  
26 heat resistance (LHR). This genomic island is also present in several other genera of  
27 *Enterobacteriaceae*, but its function in the enteric pathogens *Salmonella enterica* and  
28 *Enterobacter cloacae* is unknown. This study aimed to determine the frequency of the LHR in  
29 food isolates of *E. coli*, and its influence on heat resistance in *S. enterica* and *Enterobacter* spp.  
30 Cell counts of LHR-positive strains of *E. coli*, *S. enterica* and *E. cloacae* were reduced by less  
31 than 1, 1, and 4 log (cfu/mL), respectively, after exposure to 60°C for 5 min, while cell counts of  
32 LHR-negative strains of the same species were reduced by more than 7 log (cfu/mL).  
33 Introducing an exogenous copy of the LHR into heat-sensitive enteropathogenic *E. coli* and *S.*  
34 *enterica* increased heat resistance to a level that was comparable to LHR-positive wild type  
35 strains. Cell counts of LHR-positive *S. enterica* were reduced by less than 1 log(cfu/mL) after  
36 heating to 60°C for 5 min. Survival of LHR-positive strains was improved by increasing the  
37 NaCl concentration from 0 to 4%. Cell counts of LHR-positive strains of *E. coli* and *S. enterica*  
38 were reduced by less than 2 log (cfu/g) in ground beef patties cooked to an internal core  
39 temperature of 71°C. This study indicates that LHR-positive *Enterobacteriaceae* pose a risk to  
40 food safety.

41

42 **Keywords**

43 Heat resistance; LHR; *Escherichia coli*; *Salmonella enterica* Senftenberg; food-borne pathogens;  
44 meat, undercooked burger

45

## 46 1. Introduction

47 Heat resistance of *Enterobacteriaceae* is highly variable. Heat resistance in species of  
48 *Enterobacteriaceae* results from the activity of alternative sigma factors (Dodd and Aldsworth,  
49 2002; Noor, 2015), the presence of specialized heat shock chaperones and proteases (Arsène *et*  
50 *al.*, 2000) and the accumulation of compatible solutes (Hengge-Aronis *et al.*, 1991; Li and  
51 Gänzle, 2016; Pleitner *et al.*, 2012). In *Escherichia coli*, elevated temperatures induce the  
52 expression of major heat shock proteins and molecular chaperones involved in protein folding,  
53 refolding and degradation (Noor, 2015). Additionally, the heat resistance of cultures increases  
54 upon entry to the stationary phase as a result of the activity of the alternative sigma factor  $\sigma^S$   
55 (Dodd and Aldsworth, 2002). Similar mechanisms of heat resistance have been identified in  
56 related enteric species, including *Salmonella enterica* (Dodd *et al.*, 2007). In addition to the  
57 inducible gene expression in response to heat shock or entry into the stationary phase of growth,  
58 *E. coli* stably adapts to growth at high temperature (Rudolph *et al.*, 2010) and acquires increased  
59 resistance to lethal heat challenge through genomic adaptation (Vanlint *et al.*, 2011).

60 *E. coli* AW1.7 is a food isolate with exceptional resistance to heat (Dlusskaya *et al.*,  
61 2011). The heat resistance of *E. coli* AW1.7 is not mediated by the  $\sigma^S$  regulon (Ruan *et al.*, 2011)  
62 but was attributed to the 14-kb genomic island termed locus of heat resistance (LHR) (Mercer *et*  
63 *al.*, 2015). The LHR encodes several putative heat shock proteins, proteases, and transport  
64 proteins, and is present in the genomes of 2% of all *E. coli* for which genome sequence data is  
65 available (Mercer *et al.*, 2015). Fragments of the LHR were also linked to increased heat  
66 resistance in *Klebsiella pneumoniae* and *Cronobacter sakazakii* (Bojer *et al.*, 2010; Gajdosova *et*  
67 *al.*, 2011). Bioinformatic analyses also identified the LHR in *Yersinia enterocolitica*, *Citrobacter*  
68 *sp.* and *Enterobacter cloacae* (Mercer *et al.*, 2015). LHR sequences from diverse

69 *Enterobacteriaceae* exhibit > 99% sequence identity. The high GC content and the presence of  
70 flanking mobile elements support the hypothesis that diverse species of the *Enterobacteriaceae*  
71 acquired this genomic island by horizontal gene transfer (Mercer *et al.*, 2015).

72 Horizontal gene transfer of the LHR may allow transfer of the genomic island to  
73 pathogenic *Enterobacteriaceae* for which presence of the LHR has not yet been reported. Food-  
74 borne pathogens in the *Enterobacteriaceae* include *S. enterica* and Shiga-toxin producing  
75 *Escherichia coli* (STEC) (Scallan *et al.*, 2011). In Canada, *Salmonella* and O157 STEC are  
76 estimated to be responsible for 30% of hospitalizations and 24% of deaths associated with food-  
77 borne illness, annually (Government of Canada, 2015). In food production and food preparation,  
78 enteric pathogens are controlled by pasteurization, steam or hot water intervention steps that are  
79 applied in production of meat, or domestic cooking of meat to an internal core temperature of  
80 71°C (Health Canada, 2015; Minihan *et al.*, 2003; Rajic *et al.*, 2007; Yang *et al.*, 2015). Strains  
81 of *E. coli* harbouring the LHR resist thermal interventions that are lethal to LHR-negative strains  
82 (Dlusskaya *et al.*, 2011). The heat resistance of *E. coli* and *S. enterica* increases with increasing  
83 NaCl concentrations (Blackburn *et al.*, 1997; Juneja *et al.*, 2003; Pleitner *et al.*, 2012). The effect  
84 of NaCl on heat resistance of the LHR positive *E. coli* AW1.7 was linked to increased  
85 accumulation of compatible solutes (Pleitner *et al.*, 2012); however, the effect of NaCl on heat  
86 resistance of other LHR-positive *Enterobacteriaceae* has not been described.

87 The role of the LHR in heat resistance of *S. enterica*, *Enterobacter* spp. and pathogenic  
88 strains of *E. coli* has not been reported. Therefore, this study aimed to investigate the effect  
89 of the LHR on heat resistance of several members of *Enterobacteriaceae*, the effect of NaCl on  
90 heat resistance of LHR-positive *Enterobacteriaceae*, and their survival after cooking in ground  
91 beef.

92 2. **Material and Methods**

93 2.1. Bacterial strains, plasmids and culture conditions

94 Strains of *E. coli*, *S. enterica*, and *Enterobacter* spp. used in this study are listed in Table  
95 1. For this study, we selected the heat resistant *S. enterica* Senftenberg (Ng et al., 1969) and a  
96 heat-sensitive reference strain of *S. enterica*, and strains of *Enterobacter cloacae* from a  
97 collection of coliforms previously isolated from a beef processing plant (Aslam et al. 2004).  
98 Additionally, a total of 92 DNA samples of *E. coli* from a meat processing facility were screened  
99 for the LHR genotype. All four LHR-positive and four LHR-negative strains were obtained for  
100 further experimental analysis. Unless otherwise noted, strains were cultured at 37°C in Luria-  
101 Bertani (LB) media, which contains 1% NaCl (w/v). Media were supplemented with 15 µg/mL  
102 tetracycline-HCl when necessary for plasmid selection. For experiments determining the effect  
103 of NaCl on LHR-mediated resistance, LB media with addition of 0, 2 or 4% NaCl were also  
104 used. Plasmids and primers are listed in Table 2. The recombinant plasmids pRK767, pLHR and  
105 pLHR1-2 were transformed into wild type strains by electroporation and the transformed strains  
106 were plated on LB media containing 15 mg/L tetracycline-HCl (Mercer et al., 2015). The  
107 taxonomic position of *E. coli* strains was confirmed by PCR targeting the β-glucuronidase gene  
108 for *E. coli* (Yang et al, 2011; Table 2). The identity of other species was confirmed by PCR  
109 amplification and Sanger sequencing of genes coding for 16S rRNA by service of MacroGen  
110 (Rockville, MD), followed by sequence analysis using the ribosomal database project release 11  
111 (<http://rdp.cme.msu.edu/>).

112 2.2. PCR screening to determine the presence of the LHR in *Enterobacteriaceae*

113 To identify LHR-positive strains, 3 target regions of the LHR were amplified by PCR as  
114 previously described (Mercer *et al.*, 2015). Primer pairs HR-F1/HS-R1, HR-F2.2/HR-R2 and  
115 HS-F1 and HR-R3 (Table 2) were used in PCR reactions with a recombinant Taq DNA  
116 polymerase (Invitrogen, Burlington, Ontario). Genomic DNA from 92 strains of *E. coli* was used  
117 as templates for screening. These strains were selected to represent the diversity of more than  
118 400 isolates that were previously obtained from a beef-processing facility (Yang *et al.*, 2015). *E.*  
119 *coli* AW1.7 and *E. coli* AW1.7 $\Delta$ pHR1 (Pleitner *et al.*, 2012) were used as LHR-positive and -  
120 negative controls, respectively. Colony PCR with the same primers was used to confirm the LHR  
121 genotypes for strains of *S. enterica* and *E. cloacae*.

### 122 2.3. Heat inactivation in laboratory media

123 Heat inactivation was used to determine the level of resistance for each strain as  
124 previously described, using 60°C as challenge temperature that allows straightforward  
125 differentiation of heat resistant and heat sensitive strains (Dlusskaya *et al.* 2011; Mercer *et al.*,  
126 2015). Cultures were grown overnight in LB broth containing 0, 1, 2 or 4% NaCl at 37°C with  
127 200 rpm agitation (Pleitner *et al.*, 2012). Heat treatments were performed at 60°C for 5, 10 or 20  
128 min. For each experiment, *E. coli* AW1.7 and *E. coli* AW1.7 $\Delta$ pHR1 were used as LHR-positive  
129 and -negative controls, respectively. The reduction in cell counts was determined in three  
130 biological replicates. Statistically significant differences (p-value < 0.05) were determined by  
131 analysis of variance (ANOVA).

### 132 2.4. Heat inactivation in ground beef patties

133 Ground beef was aseptically prepared from beef rounds with 4% fat that were obtained  
134 from a federally inspected beef-processing facility. Cell counts of uninoculated beef patties were

135 determined by mixing 200 g of ground beef with 200 mL of 0.1% buffered peptone water  
136 (composition per litre: 10 g peptone, 3.5 g Na<sub>2</sub>HPO<sub>2</sub>, 1.5 g KH<sub>2</sub>PO<sub>4</sub>, 5 g NaCl) in a stomacher  
137 and plating 100 µL on plate count agar [PCA; BD Difco™, Mississauga, Ontario, Canada] and  
138 violet-red bile agar (VRBA; BD Difco™), followed by incubation at 37°C for quantification of  
139 total aerobic plate counts and coliform bacteria, respectively. Total aerobic plate counts and  
140 counts of coliform bacteria were less than 3000 and 50 cfu/g, respectively.

141 Ground beef was inoculated with strains of *E. coli*, *E. cloacae*, or *S. enterica* by mixing  
142 approximately 10 mL of overnight cultures in LB with 200 g of refrigerated ground beef, and  
143 massaging in a sterile bag by hand for 2 minutes. To determine the initial count of each sample,  
144 20 g were removed and diluted with 200 ml of buffered peptone water and stomached for 2 min  
145 using a Seward Lab Blender 400 (Seward Worthing, UK). The resulting solution was serially  
146 diluted in buffered peptone water, plated on LB agar and the plates were incubated overnight at  
147 37°C. The initial cell count was about 10<sup>7</sup> cfu/g. The remaining 180 g of ground beef was formed  
148 into a patty with a diameter of 11.5 cm using a Single Hamburger Press (Weston Brand  
149 Pragotrade, Strongsville, OH, USA). The patty was cooked on a grill (Cuisinart, Woodbridge,  
150 Ontario) that was preheated to medium heat for at least 20 min to a temperature of 130 - 140°C.  
151 The temperature of the patty was monitored with a Barnant Type K thermocouple thermometer  
152 (Barnant, Barrington, USA) that was inserted in the geometric centre of the patty. Once the  
153 internal temperature reached 71°C, the patty was removed, placed in 200 mL of iced buffered  
154 peptone water, and stomached for 2 min. The solution was serially diluted, plated on LB agar  
155 and incubated overnight at 37°C. Cell counts of samples before and after treatment were  
156 determined in three replicate experiments and each experiment was analysed in duplicate.  
157 Statistically significant differences (p-value < 0.05) were determined by ANOVA.

158 3. **Results**

159 **3.1. The locus of heat resistance (LHR) provides heat resistance to diverse *E. coli***

160 To determine the frequency of LHR positive *E. coli* present in beef-processing  
161 environments, 92 isolates of *E. coli* from beef were screened by PCR. The LHR was present in 4  
162 of the 92 strains (4.3%). Heat resistance of the LHR-positive strains *E. coli* 62, 68, 79, and 85  
163 was compared with four LHR-negative strains that were isolated at the same time from the same  
164 processing facility (Table 1 and data not shown). Heat treatment at 60°C for 5 min reduced cell  
165 counts by less than 1 log(cfu/mL) for all 4 LHR-positive strains of *E. coli* (Figure 1); this  
166 reduction of cell counts is similar to reductions for other LHR-positive strains (Mercer *et al.*,  
167 2015). In contrast, cell counts of LHR-negative strains of *E. coli* 40, 50 60 and 70 were reduced  
168 by more than 7 log (cfu/mL). None of LHR positive *E. coli* possessed plasmids large enough to  
169 carry the LHR (data not shown), therefore, the LHR likely exists as a chromosomally-integrated  
170 genomic island in these strains.

171 To further confirm that the LHR confers heat resistance in *E. coli*, the plasmid pLHR  
172 (Mercer *et al.*, 2015) was transformed into the LHR-negative *E. coli* MG1655 and the  
173 enteropathogenic *E. coli* (EPEC) E2348/69. The plasmid pRK767 without the LHR served as a  
174 control. The phenotype of *E. coli* carrying pLHR was consistent for both pathogenic and non-  
175 pathogenic strains of *E. coli* (Figure 1). The same strains carrying the empty vector control,  
176 pRK767, were similar in heat resistance to other LHR-negative strains.

177 **3.2. Sequence and function of the LHR is conserved in *Enterobacteriaceae***

178 LHR sequences are present in diverse species of *Enterobacteriaceae* (Mercer *et al.*, 2015);  
179 however, to date the LHR was not identified in *S. enterica*. Searching the National Center for  
180 Biotechnology Information (<https://www.ncbi.nlm.nih.gov/>) whole genome shotgun sequence

181 database retrieved short, non-contiguous fragments of the LHR belonging to *S. enterica* strain  
182 ATCC 43845. Assembly of these fragments revealed 88% coverage and >99.9% sequence  
183 identity to the LHR sequence in *E. coli* AW1.7. PCR analysis of *S. enterica* ATCC 43845 with  
184 the same primers that were used for *E. coli* demonstrated that the strain carries the full length  
185 locus (data not shown). The heat resistance of the LHR-positive *S. enterica* ATCC 43845 (Fig. 2)  
186 was comparable to *E. coli* AW1.7 (Fig. 1). Conversely, the resistance of the LHR-negative *S.*  
187 *enteria* ATCC 13311 was comparable to heat-sensitive and LHR-negative strains of *E. coli*  
188 (Figure 2). Introducing the LHR on a plasmid into the heat sensitive strain yielded *S. enterica*  
189 ATCC 13311 pLHR, which had significantly ( $p < 0.05$ ) increased heat resistance when  
190 compared to the wild type (Figure 2).

191 The PCR screening for presence of the LHR in *Enterobacteriaceae* was extended to  
192 strains of *Enterobacter* that were isolated from a beef processing plant (Aslam et al., 2004); meat  
193 isolates were compared to isolates from dairy products (Jones, 1956), or the scat of grizzly bears  
194 (Schwab et al., 2009) (Table 1). PCR screening identified 5 strains of *Enterobacter cloacae*  
195 carrying the LHR. Cell counts of LHR-positive *E. cloacae* were reduced by 2.0 to 3.4  
196 log(cfu/mL) after treatment at 60°C for 5 min. Cell counts of the LHR-negative strain *E. cloacae*  
197 FUA1067 were reduced by more than 7 log (cfu/mL) (Figure 2). Introducing pLHR into the heat  
198 sensitive *E. cloacae* FUA1067 increased heat resistance and cell counts were reduced by 5 log  
199 (cfu/mL) (Figure 2).

### 200 **3.3. Salt affects LHR-mediated resistance in *Enterobacteriaceae***

201 NaCl increased heat resistance of *E. coli* and *S. enterica* (Blackburn et al., 1997; Goepfert  
202 et al., 1970; Mattick et al., 2001; Pleitner et al., 2012). We therefore tested the effects of NaCl  
203 on resistance of LHR-positive *E. coli* and *S. enterica* to treatment in LB broth at 60°C (Figure 3).

204 NaCl was added to the pre-culture and the NaCl concentration was maintained for the heat  
205 treatment. Cell counts of LHR-negative strains were reduced to cell counts below the detection  
206 limit after 5 min of treatment at any NaCl concentration; corresponding to a reduction of cell  
207 counts by more than 5 log(cfu/mL) (data not shown). After 5 min of treatment at 60°C, cell  
208 counts of LHR-positive strains were reduced by less than 2 log(cfu/mL) at any NaCl  
209 concentration. However, addition of 4% NaCl significantly ( $p < 0.05$ ) increased the survival of  
210 LHR-positive strains after 20 min of treatment at 60°C. Their cell counts were reduced by more  
211 than 6 log (cfu/mL) after 20 min at 60°C and 0% NaCl, but cell counts were reduced by less than  
212 4 log (cfu/mL) after 20 min at 60°C and 4% NaCl (Figure 3). The plasmid pLHR1-2, which  
213 contains the native promotor of the LHR and genes coding 10 of the 16 putative proteins, also  
214 conferred heat resistance to *E. coli* AW1.7ΔpHR1, but the resistance of *E. coli*  
215 AW1.7ΔpHR1(pLHR1-2) was lower when compared to the resistance of AW1.7ΔpHR1(pLHR)  
216 at all salt concentrations (Figure 3).

#### 217 **3.4. Survival of LHR-positive *E. coli* and *S. enterica* in beef patties cooked to a core** 218 **temperature of 71°C.**

219 To assess the effects of the LHR on survival of *Enterobacteriaceae* during cooking of  
220 meat, LHR-positive and -negative strains were inoculated into beef patties. Patties were cooked  
221 to an internal temperature of 71°C according to Health Canada's recommended safe cooking  
222 temperature (Health Canada, 2015). Cell counts of the LHR-positive *E. coli* AW1.7,  
223 AW1.7ΔpHR1 (pLHR), and *S. enterica* ATCC 43845 were reduced by 2, 2.7 and 1.7 log (cfu/g),  
224 respectively (Figure 4), confirming that the heat resistance conferred by LHR is equivalent in *E.*  
225 *coli* and *S. enterica*. In contrast, cell counts of the LHR-negative *S. enterica* ATCC 13311 were  
226 reduced by more than 5 log (cfu/g); cell counts of *E. coli* AW1.7ΔpHR1 were reduced below the

227 detection level after cooking. Remarkably, the heat resistance in beef patties of the LHR-positive  
228 *E. cloacae* FUA1140 and the LHR-negative *E. cloacae* FUA1067 did not differ and cell counts  
229 of both strains was reduced by more than 5 log (cfu/g) (Figure 4). The improved survival of  
230 LHR-positive strains was found to be consistent in ground beef patties for *E. coli* and *S. enterica*,  
231 but differed for *E. cloacae*.

#### 232 4. Discussion

233 The results of this study demonstrated that LHR-positive strains of *E. coli*, *S. enterica*  
234 and *E. cloacae* exhibit higher resistance to heat than LHR-negative strains of the same species,  
235 extending previous results obtained with *K. pneumoniae* and *C. sakazakii* (Bojer *et al.*, 2010;  
236 Gajdosova *et al.*, 2011). Moreover, the level of protection provided by the LHR is genus specific.  
237 LHR-positive *E. coli* and *S. enterica* are more resistant to heat than LHR-positive strains of *E.*  
238 *cloacae* (this study), *K. pneumoniae* (Bojer *et al.*, 2010), and *C. sakazakii* (Gajdosova *et al.*,  
239 2011).

240 The presence of *E. coli* in processing environments is used an indicator for the presence  
241 of pathogenic strains (Castillo *et al.*, 1998). The frequency of LHR-positive and heat resistant  
242 isolates that were previously obtained from a beef processing plant was determined in this study  
243 as 4.3%. Bioinformatic and genetic analysis indicate that the LHR is transferred between  
244 *Enterobacteriaceae* by horizontal gene transfer (Mercer *et al.*, 2015). This study is the first to  
245 report LHR-mediated heat resistance in *S. enterica*; and demonstrates that the LHR mediates heat  
246 resistance in pathogenic *E. coli*. EPEC O127:H6 strain E2348/69 is a prototype strain that has  
247 been widely used to study EPEC biology, genetics and virulence (Iguchi *et al.*, 2009). The wild  
248 type strain does not possess the LHR and is heat sensitive; however, transformation of the strain  
249 with a plasmid-coded LHR conferred heat resistance that is comparable to LHR-positive wild

250 type strains of *E. coli*. All wild type and LHR-positive *E. coli* identified to date maintain the  
251 element as a chromosomally-integrated genomic island (Mercer *et al.*, 2015) while strains of  
252 *Klebsiella pneumoniae* possess plasmid-borne copies of the LHR (Bojer *et al.*, 2010). Regardless  
253 of the genetic position (plasmid or chromosome), source (food or clinical isolate) or pathogenic  
254 ability (*eae*<sup>+</sup> or K-12 strain), LHR-positive *E. coli* demonstrate a similar and exceptional  
255 resistance to heat.

256 Heat resistance of *S. enterica* is highly variable from strain to strain (Lianou and  
257 Koutsoumanis, 2013). The exceptional heat resistance of the LHR-positive strain used in this  
258 study has been documented previously (Lianou and Koutsoumanis, 2013; Mañas *et al.*, 2003;  
259 Murphy *et al.*, 1999; Ng *et al.*, 1969). *S. enterica* Senftenberg ATCC 43845 was originally  
260 described as a H<sub>2</sub>S-negative strain capable of surviving 60°C for 5 min in liquid egg (Winter *et*  
261 *al.*, 1946). It was suggested that heat resistance of this strain is atypical and that it should  
262 therefore not be used when constructing strain cocktails for food safety research (Juneja *et al.*,  
263 2003; Lianou *et al.*, 2013; Ng *et al.*, 1969; van Asselt and Zwietering, 2006). Our results indicate  
264 that the exceptional heat resistance of *S. enterica* Senftenberg ATCC 43845 is conferred by the  
265 LHR. Accordingly, introduction of the LHR into the heat sensitive *S. enterica* Typhimurium  
266 ATCC 13311 resulted in heat resistance that was comparable to the LHR-positive *S. enterica*  
267 ATCC 43845 and LHR-positive strains of *E. coli*. This is the first evidence that the LHR is the  
268 genetic determinant of unusually heat resistant strains of *S. enterica*. Due to the potential  
269 horizontal acquisition of the genomic island, these strains should be considered in future thermal  
270 inactivation studies.

271 Strains of *E. cloacae* are regarded as opportunistic pathogens (Hart, 2006) primarily  
272 associated with nosocomial infections (Gaston, 1988) and frequently demonstrate resistance to

273 multiple  $\beta$ -lactam antibiotics (Fung-Tomc *et al.*, 1996). However, *E. cloacae* have also been  
274 frequently isolated from cattle, processing environments and retail beef (Kim and Wei, 2007).  
275 Strains used in this study were isolated from a beef processing facility and previously  
276 misidentified as *E. coli* or contaminants of strains of *E. coli* (Aslam *et al.*, 2004). The wild type  
277 LHR-positive strains of *E. cloacae* tested in this study were significantly more heat resistant than  
278 LHR-negative strains of the same species when they were heated in broth. LHR-positive wild  
279 type strain of *E. cloacae* appeared to exhibit a higher heat resistance than a LHR-negative strain  
280 of *E. cloacae* that was transformed with pLHR; this may be attributable to the copy number of  
281 pLHR in this species. However, the difference in heat resistance between LHR-positive and  
282 LHR-negative strains was not evident when they were heated in meat; probably, the severity of  
283 the challenge exceeded the heat resistance even of LHR-positive strains.

284         Accumulation of compatible solutes in response to hyperosmotic conditions has been  
285 described as ‘*passé partout*’ for resistance to diverse environmental insults (Pleitner *et al.*, 2012;  
286 Sleator and Hill, 2010). An increase of heat resistance in response to increased NaCl  
287 concentrations has been documented for both *E. coli* and *S. enterica* (Blackburn *et al.*, 1997;  
288 Jujena *et al.*, 2003; Pleitner *et al.*, 2012). Our data demonstrate that the effect of NaCl on heat  
289 resistance is also observed with LHR-positive strains of both species. The LHR encodes putative  
290 transporters and proteases (Mercer *et al.*, 2015) that may function in response to NaCl to delay  
291 thermal inactivation during prolonged periods of heat stress. Transport proteins can play a role in  
292 the accumulation of compatible substrates, while proteolytic processing of misfolded proteins is  
293 an essential part of moderating heat stress (Rosen *et al.*, 2002).

294         Pathogen intervention methods in beef processing facilities include thermal interventions  
295 and washing or spraying with solutions of antimicrobials (Yang *et al.*, 2015). These interventions

296 strongly reduce transfer of *E. coli* from the hide of animals to the carcass; however, *E. coli*  
297 nevertheless contaminate beef products, particularly ground beef (Yang et al., 2015). Moreover,  
298 improper cooking of beef or poultry provides an inadequate reduction of cell counts of  
299 pathogenic *E. coli* and *S. enterica* and may contribute to foodborne disease (Liu et al., 2015;  
300 Roccato et al., 2015). We demonstrate that LHR-positive pathogenic *E. coli* and *Salmonella*  
301 survived cooking of meat according to safe handling and cooking label instructions. Both *E. coli*  
302 and *S. enterica* that carry the LHR survived in ground beef patties cooked to an internal  
303 temperature of 71°C. This temperature is currently referred to as a safe internal temperature for  
304 ground beef to eliminate food-borne pathogens (Health Canada, 2015). LHR-positive pathogens  
305 therefore present an additional risk to food safety. Assessment of this risk, however, requires  
306 additional data on the frequency of the LHR in *E. coli* and *Salmonella*. The synergistic effect of  
307 genetics (LHR) and osmotic stress (NaCl concentration) in protecting cells from thermal  
308 inactivation also constitutes a novel risk to food safety that remains to be quantified.

309 We previously reported that the sequence of the LHR is highly conserved in  
310 *Enterobacteriaceae* (Mercer et al., 2015). The present study demonstrates that the function of the  
311 LHR is also conserved in *Enterobacteriaceae*. The LHR is a clear indicator of increased heat  
312 resistance in *Enterobacteriaceae* but the magnitude of the resistance is species-dependent. The  
313 presence of the LHR allows for novel and heat resistant pathogenic strains arising from  
314 horizontal gene transfer amongst related species of *Enterobacteriaceae*.

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319 **References**

- 320 Aslam, M., Greer, G.G., Nattress, F.M., Gill, C.O., McMullen, L.M., 2004. Genetic diversity of  
321 *Escherichia coli* recovered from the oral cavity of beef cattle and their relatedness to faecal *E.*  
322 *coli*. *Lett. Appl. Microbiol.* **39**:523-527.
- 323 Arsène, F., Tomoyasu, T., Bukau, B., 2000. The heat shock response of *Escherichia coli*. *Int. J.*  
324 *Food Microbiol.* **55**:3-9.
- 325 Bej, A.K., DiCesare, J.L., Haff, L., Atlas, R.M., 1991. Detection of *Escherichia coli* and *Shigella*  
326 spp. in water by using the polymerase chain reaction and gene probes for *uid*. *Appl. Environ.*  
327 *Microbiol.* **57**:1013-1017.
- 328 Blackburn, C.W., Curtis, L.M., Humpheson, L., Billon, C., McClure, P.J., 1997. Development of  
329 thermal inactivation models for *Salmonella enteritidis* and *Escherichia coli* O157:H7 with  
330 temperature, pH, and NaCl as controlling factors. *Int. J. Food Microbiol.* **38**:31-44.
- 331 Bojer, M.S., Struve, C., Ingmer, H., Hansen, D.S., Kroghfelt, K.A., 2010. Heat resistance  
332 mediated by a new plasmid encoded Clp ATPase, ClpK, as a possible novel mechanism for  
333 nonsocomial persistence of *Klebsiella pneumoniae*. *PLoS ONE* **5**:e15467.
- 334 Castillo, A., Lucia, L.M., Goodson, K.J., Savell, J.W., Acuff, G.R., 1998. Comparison of water  
335 wash, trimming, and combined hot water and lactic acid treatments for reducing of bacteria of  
336 fecal origin on beef carcasses. *J. Food Prot.* **61**:823-828.
- 337 Dodd, C.E.R., Aldsworth, T.G., 2002. The importance of RpoS in the survival of bacteria through  
338 food processing. *Int. J. Food Microbiol.* **74**(3):189-194.
- 339 Dodd, C.E.R., Richards, P.J., Aldsworth, T.G., 2007. Suicide through stress: A bacterial response  
340 to sub-lethal injury in the food environment. *Int. J. Food Microbiol.* **120**:46-50.

341 Dlusskaya, E.A., McMullen, L.M., Gänzle, M.G., 2011. Characterization of an extremely heat-  
342 resistant *Escherichia coli* obtained from a beef processing facility. *J. Appl. Microbiol.*  
343 **110**:840-849.

344 Fong-Tomc, J.C., Gradelski, E. Huczko, E., Dougherty, T.J., Kessler, R.E., Bonner, D.P., 1996.  
345 Differences in the resistant variants of *Enterobacter cloacae* selected by extended-spectrum  
346 cephalosporins. *Antimicrob. Agents Chemother.* **40**:1289-1293.

347 Hengge-Aronis, R., Klein, W., Lange, R., Rimmel, M., Boos, W., 1991. Trehalose synthesis  
348 genes are controlled by the putative sigma factor encoded by *rpoS* and are involved in  
349 stationary phase thermotolerance in *Escherichia coli*. *J. Bacteriol.* **173**:7918-7924.

350 Gajdosova, J., Benedikovicova, K., Kamodyova, N., Tothova, L., Kaclikova, E., Stuchlik, S,  
351 Turna, J., Drahovska, H., 2011. Analysis of the DNA region mediating increased  
352 thermotolerance at 58°C in *Cronobacter* sp. and other enterobacterial strains. *Antonie Van*  
353 *Leeuwenhoek* **100**:279–289.

354 Gaston, M.A., 1988. *Enterobacter*: an emerging nosocomial pathogen. *J. Hosp. Infect.* **11**:197-  
355 208.

356 Gill, P.R. Jr., Warren, G.J., 1988. An iron-antagonized fungistatic agent that is not required for  
357 iron assimilation from a fluorescent rhizosphere pseudomonad. *J. Bacteriol.* **170**:163-170.

358 Goepfert, J.M., Iskander, I.K., Amundson, C.H., 1970. Relation of the heat resistance of  
359 salmonellae to the water activity of the environment. *Appl. Microbiol.* **19**:429-433.

360 Government of Canada, Retrieved October 6<sup>th</sup>, 2015 from: [http://healthy Canad  
ians.gc.ca/eating-  
nutrition/risks-recalls-rappels-  
risques/surveillance/illness-  
estimates-estimations-  
maladies/yearly-annuel-eng.php](http://healthy Canadians.gc.ca/eating-<br/>361 nutrition/risks-recalls-rappels-risques/surveillance/illness-estimates-estimations-<br/>362 maladies/yearly-annuel-eng.php)

363 Hart, C.A., 2006. *Klebsiella, Citrobacter, Enterobacter* and *Serratia* spp.. In S.H. Gillspie &  
364 P.M Hawkey (Eds.), *Principles and practice of clinical bacteriology*. 2<sup>nd</sup> Edition, 377-386.  
365 England, U.K.: John Wiley and Sons Ltd.

366 Health Canada, Retrieved October 28<sup>th</sup>, 2015 from: [http://www.hc-sc.gc.ca/fn-](http://www.hc-sc.gc.ca/fn-an/legislation/guide-ld/guide-cook-cuiss-meat-viand-eng.php)  
367 [an/legislation/guide-ld/guide-cook-cuiss-meat-viand-eng.php](http://www.hc-sc.gc.ca/fn-an/legislation/guide-ld/guide-cook-cuiss-meat-viand-eng.php)

368 Iguchi, A., Thomson, N.R., Ogura, Y., Saunders, D., Ooka, T., Henderson, I.R., Harris, D.,  
369 Asadulghani, M., Kurokawa, K., Dean, P., Kenny, B., Quail, M.A., Thurston, S., Dougan, G.,  
370 Hayashi, T., Parkhill, J., Frankel, G., 2009. Complete genome sequence and comparative  
371 genome analysis of enteropathogenic *Escherichia coli* O127:H6 strain E2348/69. *J. Bacteriol.*  
372 **191**:347-354.

373 Juneja, J.K., Marks, H.M., Huang, L., 2003. Growth and heat resistance kinetic variation among  
374 various isolates of *Salmonella* and its application to risk assessment. *Risk Anal.* **23**:199-213.

375 Jones, G.E., 1956. Acetone taint in milk due to *Bacterium cloacae*. *J. Dairy Res.* **23**: 21-23.

376 Kim, S.H., Wei, C.I., 2007. Expression of AmpC beta-lactamase in *Enterobacter cloacae*  
377 isolated from retail ground beef, cattle farm and processing facilities. *J. Appl. Microbiol.*  
378 **103**:400-408.

379 Li, H., Gänzle, M.G. 2016. Some like it hot: Heat resistance of *Escherichia coli* in foods. *Front.*  
380 *Microbiol. in press.*

381 Lianou, A., Koutsoumanis, K.P., 2013. Evaluation of the strain variability of *Salmonella enterica*  
382 acid and heat resistance. *Food Microbiol.* **34**:259-267.

383 Liu, Y., Gill, A., McMullen, L., Gänzle, M.G., 2015. Variation in heat and pressure resistance of  
384 verotoxigenic and non-toxigenic *Escherichia coli*. *J. Food Prot.* **78**:111-120.

385 Minihan, D., Whyte, P., O'Mahony, M., Collins, J.D., 2003. The effect of commercial steam  
386 pasteurisation on the levels of *Enterobacteriaceae* and *Escherichia coli* on naturally  
387 contaminated beef carcasses. *J. Vet. Med.* **50**:352-356.

388 Mañas, P., Pagan, R., Alvarez, I., Uson, S.C., 2003. Survival of *Salmonella senftenberg* 775W to  
389 current liquid whole egg pasteurization treatments. *Food Microbiol.* **20**:593-600.

390 Mattick, K.L., Jørgensen, F., Wang, P., Pound, J., Vendeven, M.H., Ward, L.R., Legan, J.D.,  
391 Lappin-Scott, H.M., Humphrey, T.J., 2001. Effect of challenge temperature and solute type on  
392 heat tolerance of *Salmonella* serovars at low water activity. *Appl. Environ. Microbiol.*  
393 **67**:4128-4136.

394 Mercer, R.G., Zheng, J., Garcia-Hernandez, R., Ruan, L., Gänzle, M.G., McMullen, L.M., 2015.  
395 Genetic determinants of heat resistance in *Escherichia coli*. *Front. Microbiol.* **6**:932. doi:  
396 10.3389/fmicb.2015.00932

397 Murphy, R.Y., Marks, B.P., Johnson, E.R., Johnson, M.G., 1999. Inactivation of *Salmonella* and  
398 *Listeria* in ground chicken breast meat during thermal processing. *J. Food Prot.* **9**:975-1096.

399 Ng, H., Bayne, H.G., Garibaldi, J.A., 1969. Heat resistance of *Salmonella*: The uniqueness of  
400 *Salmonella* Senftenberg 755W. *Appl. Microbiol.* **17**:78-82.

401 Noor, R., 2015. Mechanism to control the cell lysis and the cell survival strategy in stationary  
402 phase under heat stress. *SpringerPlus* **4**:599.

403 Pleitner, A., Zhai, Y., Winter, R., Ruan, L., McMullen, L.M., Gänzle, M.G., 2012. Compatible  
404 solutes contribute to heat resistance and ribosome stability in *Escherichia coli* AW1.7.  
405 *Biochim. Biophys. Acta.* **1824**:1351-1357.

406 Rajic, A., Waddell, L.A., Sargeant, J.M., Read, S., Farber, J., Firth, M.J., Chambers, A., 2007.  
407 An overview of microbial food safety programs in beef, pork, and poultry from farm to  
408 processing in Canada. *J. Food Prot.* **70**:1286–1294.

409 Roccatto, A., Uyttendaele, M., Cibin, V., Barrucci, F., Cappa, V., Zavagnin, P., Longo, A., Ricci  
410 A., 2015. Survival of *Salmonella* Typhimurium in poultry-based meat preparations during  
411 grilling, frying and baking. *Int. J. Food Microbiol.* **197**:1-8.

412 Rosen, R., Biran, D., Gur, E., Becher, D., Hecker, M., Ron, E.Z., 2002. Protein aggregation in  
413 *Escherichia coli*: role of proteases. *FEMS Microbiol. Lett.* **207**:9-12.

414 Ruan, L., Pleitner, A., Gänzle, M.G., McMullen, L.M., 2011. Solute transport proteins and the  
415 outer membrane protein NmpC contribute to heat-resistance of *Escherichia coli* AW1.7. *Appl.*  
416 *Environ. Microbiol.* **77**: 2961-2967.

417 Rudolph, B., Gebendorfer, K.M., Buchner, J., Winter, J., 2010. Evolution of *Escherichia coli* for  
418 growth at high temperatures. *J. Biol. Chem.* **285**:19029-19034.

419 Schwab, C., Cristescu, B., Boyce, M.S., Stenhouse, G.B., Gänzle, M.G., 2009. Bacterial  
420 populations and metabolites in the scat of free roaming grizzly bears. *Can. J. Microbiol.*  
421 **55**:1335-1346.

422 Scallan, E., Hoekstra, R.M., Angulo, F.J., Tauxe, R.V., Widdowson, M-A., Roy, S.L., Jones,  
423 J.L., Griffin, P.M., 2011. Foodborne illness acquired in the United States - Major pathogens.  
424 *Emerg. Infect. Dis.* **17**:7–14.

425 Sleator, R.D., Hill, C. Compatible solutes: a listerial passé-partout? *Gut Microbes* **1**:77-79.

426 van Asselt, E.D., Zwietering, M.H., 2006. A systematic approach to determine global thermal  
427 inactivation parameters for various food pathogens. *Int. J. Food Microbiol.* **107**:73-82.

428 Vanlint, D., Mitchell, R., Bailey, E., Meersman, F., McMillan, P.F., Michiels, C.W., Aertsen, A.,  
429 2011. Rapid acquisition of Gigapascal-high-pressure resistance by *Escherichia coli*. *MBio*.  
430 2:e00130-10. doi:10.1128/mBio.00130-10.

431 Winter, A.R., Stewart, G.F., McFarlane, V.H., Solowey, M., 1946. Pasteurization of liquid egg  
432 products III. Destruction of *Salmonella* in liquid whole egg. *Am. J. Public Health* **36**:451-460.

433 Yang, X., Badoni, M., Gill, C.O., 2011. Use of propidium monoazide and quantitative PCR for  
434 differentiation of viable *Escherichia coli* from *E. coli* killed by mild or pasteurizing heat  
435 treatments. *Food Microbiol.* **28**:1478-1482.

436 Yang, X., Badoni, M., Tran, F., Gill, C.O., 2015. Microbiological effects of a routine treatment  
437 for decontaminating hide-on carcasses at a large beef packing plant. *J. Food Prot.* **78**:256-  
438 263.

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442 **Table 1. Bacterial strains used in this study.**

| Strains                       | Description   | Reference                              |
|-------------------------------|---|--|
| <i>E. coli</i>                |   |  |
| DH5 $\alpha$                  | Sub-cloning strain                                  | Invitrogen                             |
| AW1.7                         | LHR-positive wild type food isolate                 | Dlusskaya <i>et al.</i> , 2011         |
| AW1.7 $\Delta$ pHR1           | LHR-negative, heat sensitive derivative of AW1.7    | Pleitner <i>et al.</i> , 2012          |
| AW1.7 $\Delta$ pHR1(pLHR)     | Transgenic LHR-positive derivative                  | Mercer <i>et al.</i> , 2015            |
| AW1.7 $\Delta$ pHR1(pLHR1-2)  | Transgenic derivative with ORFs 1 – 10 of the LHR   | This study                             |
| MG1655                        | K-12 strain   | ATCC                                   |
| MG1655(pLHR)                  | Transgenic LHR-positive derivative of MG1655        | Mercer <i>et al.</i> , 2015            |
| MG1655(pRK767)                | Vector control for MG1655 carrying pLHR             | Mercer <i>et al.</i> , 2015            |
| EPEC O127:H6 E2348/69         | Enteropathogenic <i>E. coli</i> ; LEE-positive      | Iguchi <i>et al.</i> , 2009            |
| EPEC O127:H6 E2348/69(pLHR)   | Transgenic LHR-positive derivative of EPEC          | This study                             |
| EPEC O127:H6 E2348/69(pRK767) | Vector control for EPEC carrying pLHR               | This study                             |
| 40                            | LHR-negative beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 50                            | LHR-negative beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 60                            | LHR-negative beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 62                            | LHR-positive beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 68                            | LHR-positive beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 70                            | LHR-negative beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 79                            | LHR-positive beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| 85                            | LHR-positive beef carcass isolate                   | Yang <i>et al.</i> , 2015; this study  |
| <i>S. enterica</i>            |   |  |
| ATCC13311                     | serovar Typhimurium; LHR-negative                   | ATCC                                   |
| ATCC13311(pLHR)               | Transgenic, LHR-positive derivative of ATCC13311    | This study                             |
| ATCC43845                     | serovar Senftenberg; LHR-positive                   | ATCC                                   |
| <i>E. cloacae</i>             |   |  |
| FUA1140                       | Isolated from equipment of beef processing facility | Aslam <i>et al.</i> , 2004; this study |
| FUA1141                       | Isolated from equipment of beef processing facility | Aslam <i>et al.</i> , 2004; this study |
| FUA1144                       | Isolated from equipment of beef processing facility | Aslam <i>et al.</i> , 2004; this study |
| FUA1145                       | Isolated from equipment of beef processing facility | Aslam <i>et al.</i> , 2004; this study |
| NCDO612                       | Subsp. <i>cloacae</i> ; Dairy isolate               | NCIMB                                  |
| FUA1067                       | Isolated from bear feces                            | Schwab <i>et al.</i> , 2009            |
| FUA1067(pLHR)                 | Transgenic LHR-positive derivative of FUA1067       | This study                             |

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445 **Table 2. Plasmids and primers used in this study**

| <b>Name</b>     | <b>Description or Sequence (5' → 3')</b>                                | <b>Reference</b>            |
|-----------------|---|-----------------------------|
| <i>Plasmids</i> |   |                             |
| pUC19           | High copy plasmid   | Sigma                       |
| pRK767          | Low copy plasmid  | Gill and Warren, 1988       |
| pLHR            | Entire LHR, including promoter and <i>orfs1-16</i> , cloned into pRK767 | Mercer <i>et al.</i> , 2015 |
| pLHR1-2         | Partial LHR, including promoter and <i>orfs1-10</i> cloned into pRK767  | Mercer <i>et al.</i> , 2015 |
| <i>Primers</i>  |   |                             |
| HR-F1           | TTAGGTACCGCTGTCCATTGCCTGA   | Mercer <i>et al.</i> , 2015 |
| HS-R1           | AGACCAATCAGGAAATGCTCTGGACC  | Mercer <i>et al.</i> , 2015 |
| HR-F2.1         | AGGGTACCAGCGATATCCGTCAATTGACT   | Mercer <i>et al.</i> , 2015 |
| HR-R2           | TATCTAGAATGTCATTTCTATGGAGGCATGAATCG                                     | Mercer <i>et al.</i> , 2015 |
| HS-F1           | GCAATCCTTTGCCGCAGCTATT  | Mercer <i>et al.</i> , 2015 |
| HR-R3           | GTCAAGCTTCTAGGGCTCGTAGTTTCG   | Mercer <i>et al.</i> , 2015 |
| URL-301         | TGTTACGTCCTGTAGAAAGCCC  | Bej <i>et al.</i> , 1991    |
| URR-432         | AAAACCTGCCTGGCACAGCAATT   | Bej <i>et al.</i> , 1991    |

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## Figure Legends

**Figure 1.** Reductions of cell counts of LHR-positive and LHR-negative *E. coli* heated to 60°C for 5 min in LB broth. Source (food), pathotype (K-12 or EPEC) and presence of recombinant plasmids (pRK= pRK767 or pLHR) are indicated for each strain. *E. coli* AW1.7 and AW1.7ΔpHR1 were used as LHR-positive and LHR-negative controls, respectively. Data are shown as means ± standard deviations of 3 replicate experiments. Bars representing values greater than 7 correspond to treatments that reduced cell counts to levels below the detection limit.

**Figure 2.** Reductions of cell counts of LHR-positive and LHR-negative *S. enterica* and *E. cloacae* heated to 60°C for 5 min in LB broth. The presence (+) or absence (-) of a wild type copy of the LHR, along with the presence of exogenous copies (pLHR) are indicated below each strain. Data are shown as means ± standard deviations of 3 replicate experiments. Bars representing values greater than 7 correspond to treatments that reduced cell counts to levels below the detection limit.

**Figure 3.** Reduction of cell counts of *E. coli* and *S. enterica* harbouring the complete or a partial LHR in response to NaCl concentrations. Samples of *E. coli* AW1.7 (●), *E. coli* AW1.7ΔpHR1 (pLHR) (●), *E. coli* AW1.7ΔpHR1 (pLHR1-2) (○) and *S. enterica* ATCC 43845 (■) were grown in LB media containing 0 (A), 1 (B), 2 (C) or 4 (D) % NaCl and treated at 60°C. Data are shown as means ± standard deviations of 3 replicate experiments.

**Figure 4.** Thermal inactivation of *Enterobacteriaceae* in ground beef patties cooked to an internal temperature of 71°C. Reduction of cell counts for LHR-positive and LHR-negative *E. coli*, *S. enterica* and *E. cloacae* was determined with 3 biological replicates and means ±

standard deviations are shown. Statistically significant (p-value <0.05) differences between LHR-positive and LHR-negative strains of the same species are indicated by an asterisk (\*).







