

Amino acids other than glutamate affect the expression of the GAD system in Listeria monocytogenes enhancing acid resistance

Article

Accepted Version

Creative Commons: Attribution-Noncommercial-No Derivative Works 4.0

Paudyal, R., O'Byrne, C. P. and Karatzas, K.-A. (2020) Amino acids other than glutamate affect the expression of the GAD system in Listeria monocytogenes enhancing acid resistance. Food Microbiology, 90. 103481. ISSN 0740-0020 doi: https://doi.org/10.1016/j.fm.2020.103481 Available at http://centaur.reading.ac.uk/92900/

It is advisable to refer to the publisher's version if you intend to cite from the work. See <u>Guidance on citing</u>. Published version at: https://www.sciencedirect.com/science/article/pii/S0740002020300708 To link to this article DOI: http://dx.doi.org/10.1016/j.fm.2020.103481

Publisher: Elsevier

All outputs in CentAUR are protected by Intellectual Property Rights law,



including copyright law. Copyright and IPR is retained by the creators or other copyright holders. Terms and conditions for use of this material are defined in the <u>End User Agreement</u>.

www.reading.ac.uk/centaur

CentAUR

Central Archive at the University of Reading

Reading's research outputs online

1	Amino acids other than glutamate affect the expression of the GAD system in Listeria
2	monocytogenes enhancing acid resistance
3	
4	Ranju Paudyal ¹ , Conor P. O'Byrne ² and Kimon-Andreas Karatzas ¹ *
5	¹ Department of Food and Nutritional Sciences, The University of Reading, PO Box 226,
6	Whiteknights, Reading RG6 6AP, UK.
7	
8	² Bacterial Stress Response Group, Microbiology, School of Natural Sciences, National
9	University of Ireland, Galway, Galway, Ireland
10	
11	
12	* Corresponding author
13	Email address: k.karatzas@reading.ac.uk
14	Corresponding author address:
15	School of Chemistry, Food and Pharmacy
16	Department of Food & Nutritional Sciences,
17	University of Reading,
18	Reading
19	RG6 6AD
20	UK
21	
22	Tel. +44 118 378 6678
23	Fax. +44 118 931 0080
24	Email: <u>k.karatzas@reading.ac.uk</u>
25	
26	

27 ABSTRACT

28 The Glutamate Decarboxylase (GAD) system is important for survival of L. monocytogenes and other microorganisms under acidic conditions. Environmental conditions influence the 29 function of the GAD system. Until now, the only conditions known to lead to increased 30 transcription of the GAD system are the stationary phase in rich media and anoxic 31 32 conditions. Previously, we showed that transcription of the GAD system requires unidentified compounds other than glutamate present in rich media. Following a test looking at various 33 compounds we identified for first time that peptone, tryptone and casamino acids activate the 34 35 GAD system under oxic conditions suggesting that amino acid(s) other than glutamate and/or peptides are important for the above process. The defined medium, where the GAD 36 37 system is inactive, once it is supplemented with the above compounds results in an active intracellular and extracellular GAD system and increased acid resistance. Through functional 38 genomics we show that these compounds are required for GadD2 activity and although we 39 40 previously showed that GadD3 is active part of the intracellular GAD system, the supplementation did not activate this gene. The above is explained by the fact that only gadD2 41 42 transcription was upregulated by these compounds while the transcription of gadD1 and gadD3 remained unaffected. Together our results show that the L. monocytogenes GadD2 43 44 decarboxylase is activated in the presence of amino acids or peptides other than glutamate, 45 a finding that has important implications for acid tolerance and food safety.

46

47 Keywords

48 <u>Glutamate</u>

decarboxylase

monocytogenes; <u>peptone</u>;

L.

system;

49 tryptone; casaminoacids; γ -aminobutyrate; acid tolerance; GABA

50

51

52

53 INTRODUCTION

54

The Glutamate Decarboxylase System (GAD) system is present in a wide variety of living 55 56 organisms from all kingdoms of life, playing different roles ranging from brain functions in 57 mammals, to response under stress conditions such as temperature shock and hypoxia in plants 58 (Shelp et al., 1999) and to acid resistance in microorganisms. The main role of the GAD system in microorganisms is protection from acidic conditions (Capitani et al., 2003; Cotter et al., 59 2001a; Su et al., 2011) and in some of them such as L. monocytogenes (Cotter et al., 2001a) 60 and Escherichia coli (Capitani et al., 2003; Foster 2004) it is the most important acid resistance 61 mechanism. The GAD system is present in many other bacteria such as Shigella flexneri 62 (Waterman and Small, 2003), Mycobacterium tuberculosis (Cole et al., 1998; Cotter et al., 63 2001a), Lactobacillus reuteri, Lactococcus lactis, Lactobacillus plantarum (Su et al., 2011) 64 65 and Brucella spp. (Damiano et al. 2014). L. monocytogenes normally possesses three 66 decarboxylases GadD1, GadD2 and GadD3 and two antiporters GadT1 and GadT2 (Cotter et 67 al., 2005; Karatzas et al., 2012). The five corresponding genes are arranged in three separate 68 operons namely gadD1T1, gadT2D2 and gadD3 (Cotter et al., 2005). The gadD1T1 operon has been shown to enhance growth under mild acidic conditions (Cotter et al., 2005), while 69 gadD2T2 plays important role in survival under severe acidic conditions (Cotter et al., 2001a; 70 Cotter et al., 2005). Furthermore, we have shown that the GAD system comprises two distinct 71 parts i) the extracellular glutamate decarboxylase system (GADe) and ii) the intracellular 72 73 glutamate decarboxylase (GAD_i) system (Karatzas et al., 2012). The GAD_e is the part of the 74 GAD system that imports glutamate from the extracellular environment through the antiporters and the decarboxylases convert it to γ -amino butyric acid (GABA) and CO₂. This reaction 75 consumes a proton resulting in an increase of intracellular pH. Subsequently, the GABA is 76

exported via the antiporters GadT in exchange for a further glutamate molecule. The GAD_i has 77 previously been described converting intracellular pools of glutamate to intracellular GABA 78 (GABA_i; (Karatzas et al., 2010; Karatzas et al., 2012). Subsequently, GABA_i can directed 79 through the GABA shunt and converted to succinate semi aldehyde and succinate (Feehily et 80 al., 2013). Most strains of L. monocytogenes such as 10403S and LO28 utilize both GADi and 81 GADe (Karatzas et al., 2010) whereas EGD-e which is the most studied L. monocytogenes 82 83 strain, utilizes only the GAD_i system resulting in a highly acid sensitive phenotype (Feehily et 84 al., 2013).

It is well known that the GAD system is expressed in rich and complex media like Brain Heart 85 Infusion (BHI; Hersh et al., 1996; Penfound et al., 1999). However, we have shown that the L. 86 monocytogenes GAD system is not being upregulated under acidic conditions, as somebody 87 would expect and its expression depends on unknown signalling compounds present in rich 88 growth media. To date, little is known about the regulation and expression of the GAD system 89 in L. monocytogenes. GadD2, the most potent part of the GAD system is known to be 90 91 upregulated under anoxic conditions (Feehily et al., 2013; Jydegaard-Axelsen et al., 92 2004), during stationary phase (Cotter et al., 2001b) and in gastric fluid (Cotter et al., 2001a) 93 while we are not aware of any other conditions leading to its upregulation. L. monocytogenes 94 requires specific media for growth and for the GAD system to be expressed (Cotter et al., 2001b; Penfound et al., 1999). The inability of the GAD system to be expressed in a defined 95 medium (DM) supplemented with glutamate (Glt; DMG) has been previously shown by 96 97 Karatzas et al., (2010). In the same study, it was shown that the transcription of the GAD genes 98 was minimal in a defined medium supplemented with glutamate (DMG; Karatzas et al., 2010). Therefore, this medium could be used to study environmental conditions and compounds that 99 trigger the expression of the GAD system, in order to gain insights into how this important acid 100 resistance mechanism is regulated in L. monocytogenes. 101

The main aim of this study was to identify compounds that are essential for the expression and function of the GAD system through the use of DMG medium as a basis. During these experiments a number of compounds and mixtures of components were tested and tryptone, peptone and casamino acids were identified as activators of the GAD system. Furthermore, we identified the impact of these mixtures of compounds on survival under acidic conditions, on the GAD_e and GAD_e system activity, the role of each one of the GAD genes and on transcription of the GAD genes.

Knowledge of the environmental conditions that activate or inhibit the GAD system can help 109 to understand the acid resistance of this pathogen since the GAD system is the most important 110 111 mechanism of acid resistance. This knowledge could help us predict the survival of L. monocytogenes in various acidic foods or environments or even help us manipulate its acid 112 resistance through interventions that could affect the availability of these activators. These 113 findings could have a major economic impact on the food industry and lead to new methods 114 reducing the incidence of listeriosis whose prevalence in the EU and UK has increase 115 116 significantly the last years (EFSA and ECDC, 2018). Furthermore, since L. monocytogenes is a model microorganism for the study of the GAD system in Gram-positive 117 118 bacteria, that is distinctly different from that of Gram-negative bacteria such as E. coli, it could contribute to the wider understanding of acid resistance in Gram-positive organisms. 119

120

121

122 MATERIALS AND METHODS

123

124 Bacterial strains and growth conditions

125	Mutants in the three GAD decarboxylases of 10403S and their isogenic WT were used in this
126	study (Table 1). All strains were stored at -80°C in cryovial tubes with 7% DMSO. Stock

cultures from -80°C were grown in BHI Agar (LAB M, Lancashire, UK) and plates incubated 127 at 37°C overnight. Three colonies from each plate were transferred into 3 ml of sterile BHI 128 (LAB M, Lancashire, UK) and incubated overnight at 37°C with shaking at 140 rpm. DMG 129 (Amezaga et al., 1995) was prepared by supplementation of 6.8% of either tryptone (Thermo 130 Scientific Oxoid, LP0042, Basingstoke, UK), peptone (Thermo Scientific Oxoid, CM009, 131 Basingstoke, UK) and casamino acids (Difco, Michigan, USA). Similarly, for other 132 133 experiments DMG was prepared with 3.6% peptone or 10% casamino acids. Subsequently, the overnight cultures were used to inoculate 20 ml of sterile DMG medium (1% inoculum) either 134 with or without supplementation in 250 ml conical flasks and incubated at 37°C with shaking 135 at 140 rpm overnight (~24 h). Subsequently, these overnight cultures were used in all other 136 experiments. As described below in GABase Assay, GABAe was measured in DMG under 137 oxic and anoxic conditions in presence of 0.5 mg/ml of sodium chloride, 0.5 138 mg/ml maleic acid, 0.5 mg/ml succinic acid, 0.5 mg/ml acetic acid, 0.5 mg/ml lactic acid, 0.1% 139 sodium deoxycholate, 5% oxgall, 5% bile salts, 2%, 10% casamino acids, 3.6% peptone and 140 141 6.8% tryptone was also used to measure GABAe.

142

143 Survival under acidic conditions

Acid survival experiments were performed in DMG. Strains were grown as described above in 144 145 the presence or absence of 6.8 % tryptone, or peptone, or casamino acids. The 10403S WT strain and its isogenic gad mutants $\Delta gadD1$, $\Delta gadD2$ and $\Delta gadD3$ were grown in the presence 146 147 or absence of 6.8%, 3.6%, 10% tryptone, peptone and casamino acids respectively. Different 148 concentrations were used depending on the gad mutants and to achieve measurable rate of death at similar pH. To achieve death of cells at measurable rate, the pH of the overnight 149 cultures was adjusted to 2.4 and 2.5 depending on the experiments performed and challenged 150 with HCl (1M). Samples were obtained prior to the pH adjustment and thereafter every 20 min 151

up to 60 min decimal serial dilutions were prepared from those samples and plated onto BHI
agar. Subsequently, plates were incubated at 37°C overnight, and subsequently, colonies were
counted to assess survival under lethal acidic conditions. All experiments were performed in
triplicate.

156

157 GABase assay

A commercial preparation known as GABase was used to determine the intracellular (GABA_i)
and extracellular (GABA_e) concentrations of GABA. GABA_i was quantified as described by
O'Byrne et al., (2011), while GABA_e was quantified according to the method of Tsukatani et
al., (2011) as modified by Karatzas et al. (2010). The pH of the cultures was adjusted to 4.2 to
quantify the levels of GABA_i and GABA_e. This pH was chosen as it does not cause any cell
death, which is essential when activity of a cellular system is measured, while it is low enough
to ensure GAD system activation (Karatzas et al., 2012).

The GABase reaction was monitored by measurement of the absorbance at 340 nm every 2 min for 3 h at 37°C using a Sunrise spectrophotometer (Tecan, Mannedorf, Switzerland) operated by Magellan software (Tecan, Mannedorf, Switzerland). All reagents used for the GABase assay were obtained from Sigma-Aldrich (Steinheim, Germany).

169

170 Real-time PCR determination of gad gene expression

Transcription of the *gad* genes (*gadD1*, *gadD2* and *gadD3*) was assessed as described previously by Karatzas et al., (2010) by real time reverse transcription-PCR (RT-PCR). Transcription of the antiporter-encoding genes (*gadT1* and *gadT2*) was not assessed because their transcription is similar to the corresponding glutamate decarboxylases (*gadD1* and *gadD2* respectively) that belong to the same operon, as shown previously (Karatzas et al., 2012). Efficiencies of the primer pairs gadD1F-gadD1R, gadD2F-gadD2R, gadD3F-gadD3R and

16SF-16SR were 2.12, 2.09, 2.03 and 2.27 respectively (Karatzas et al., 2010). Overnight 177 cultures of 10403S WT were grown for ~24 h in DMG in the presence or absence of 6.8% 178 tryptone or peptone or casamino acids. Samples were taken at 28 h and relative expression of 179 the data was calculated as a ratio between expression of each of the target genes and that of 180 16S rRNA which was used as the reference gene for each cDNA sample. The advanced relative 181 182 quantification settings of the Light Cycler 480 SW 1.5.1 software programme were used, with 183 PCR efficiency as described previously (Karatzas et al., 2010). Relative expression of each gene was calculated by dividing the values obtained for this gene with those for the 16S rRNA 184 gene. Subsequently the relative expression values in the presence of either peptone, tryptone 185 or casamino acids in DMG were compared to the controls without those compounds and fold 186 187 changes were calculated.

188

189 **RESULTS**

190

191 gad genes in 10403S play no role in survival under acidic conditions in DMG

We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of We previously showed that glutamate supplementation of DM had no effect on survival of

199 Discovery of compounds resulting in GAD_e activity and GABA_e export

DMG was supplemented with different compounds or mixes of compounds and following theDMG was supplemented with different compounds or mixes of compounds and following the

DMG was supplemented with different compounds or mixes of compounds and following the growth of *L. monocytogenes* 10403S, GAD_e activity was assessed through GABA_e measurements to identify potential GAD activators (Fig. 1).

None of the individual compounds or mixes of compounds tested resulted in major GAD_e activity, with the exception of an array of undefined mixes of amino acids and/or peptides such as casamino acids, tryptone and peptone. The supplementation of DMG with these mixes resulted in significant levels of $GABA_e$ (**Fig. 1**A). All these experiments were also performed under <u>anoxic</u> conditions (**Fig. 1**B)

under <u>these</u> conditions the presence of casamino acids, tryptone and peptone activated the GAD_e system. It seems that casamino acids resulted in higher levels of GABA_e compared to other compounds tested. <u>Under anoxic</u> conditions, some levels of GABA_e were observed in presence of lactic acid (**Fig. <u>1</u>B**).

similarly,

215

210

and

216 Acid resistance of 10403S WT as a result of GAD system activation

217 <u>Strain</u>

218 10403S WT grown overnight in 6.8% peptone or tryptone or casamino acids was challenged 219 10403S WT grown overnight in 6.8% peptone or tryptone or casamino acids was challenged with HCl at 2.4 pH. We settled doing experiments with one of the three 220 concentrations used above (6.8mM) to allow comparisons between the different mixtures of 221 222 compounds and because this concentration provided a clear difference with the negative 223 control. In all cases, 10403S WT grown in DMG with peptone, tryptone and casamino acids 224 was more resistant under acidic conditions compared to DMG alone (Fig. 2A). During acid challenge for 40 min, cells of 10403S WT in DMG supplemented with peptone 225

were the most resistant while those in DMG with tryptone or casamino acids were more

sensitive, but still more resistant than those in DMG alone. Interestingly, during the course of
the experiment, inactivation rate of cells in DMG with peptone or DMG with tryptone increased
while that of cells in DMG with casamino acids decreased, at 60 min the latter had higher
numbers than those grown in peptone and tryptone (Fig. 2A).

231 GABAe measurements were performed in the presence of 6.8% of each one of these mixtures in pH 4.2. In the presence of these compounds, GABAe levels increased (P<0.05, paired t-test) 232 233 steadily within the 60 min of the experiment while no GABA was observed in DMG. Tryptone and peptone similarly resulted in the highest levels of GABAe reaching close to 10 mM after 234 60 min whereas casamino acids resulted in intermediate levels of GABAe close to 4 mM (Fig. 235 236 **2B**). As expected, the non-supplemented control did not produce $GABA_e$ in response to 237 acidification. Furthermore, upon supplementation with these three mixes of compounds we also observed statistically significant (P<0.05, paired t-test) higher levels of GABA_i production 238 239 in comparison to the DMG control (Fig. $\underline{2}C$).

240

241 The role of GAD genes in GAD_e and GAD_i activation and survival under acidic conditions 242 Acid survival was assessed in DMG with supplementation of 3.6%, 6.8% and 10% peptone, Acid survival was assessed in DMG with supplementation of 3.6%, 6.8% and 10% peptone, 243 244 Acid survival was assessed in DMG with supplementation of 3.6%, 6.8% and 10% peptone, Acid survival was assessed in DMG with supplementation of 3.6%, 6.8% and 245 10% peptone, tryptone and casamino acids individually respectively. In the presence of 246 peptone, WT and its three gad mutants showed increased resistance compared to DMG alone 247 248 (Fig. 3A). WT, $\Delta gadD1$ and $\Delta gadD3$ had very similar increased levels of GABA_e in presence 249 of peptone (Fig. 3B). The $\Delta gadD2$ was the most sensitive strain either in the presence or absence of peptone suggesting that GadD2 is the most important component of the GAD 250

251	system for survival (Fig. $\underline{3}A$) and this coincides with no detectable levels of GABA _e being
252	found in the $\Delta gadD2$ (Fig. <u>3</u> B) either in presence or absence of peptone.

In the presence of tryptone, WT and all its *gad* mutants showed similar increased resistance compared to those ones without tryptone (**Fig. 4A**). This coincides with increased levels of GABA_e, however $\Delta gadD2$ did not produce GABA_e compared to other strains even in presence of tryptone (**Fig. 4B**). If we follow the trend, we find that $\Delta gadD2$ with and without tryptone is the most sensitive during survival suggesting that GadD2 is important under acidic conditions.

259 The $\Delta gadD2$ was the most sensitive strain whereas WT, $\Delta gadD1$, $\Delta gadD3$ were the most 260 resistant in the presence of casamino acids. <u>A</u>lthough the presence of casamino

261

262 10403S WT gad gene transcription

263 Overall transcription of all GAD decarboxylases was low at basal conditions in DMG with264 Overall

transcription of all GAD decarboxylases was low at basal conditions in DMG with gadD3The presence of these mixtures resulted in gadD2 having the highest transcription levels, followed by gadD3 while gadD1 had the lowest transcription. <u>Remarkably</u>, the supplementation with these three mixtures did not result in a significant change in the transcription of gadD1 (**Fig. 6A**) and gadD3 (**Fig. 6C**) but only in that of gadD2. From all three mixtures, peptone had the strongest effect on gadD2 transcription followed by casamino acids and tryptone (**Fig. 6B**).

- 272
- 273

274 **DISCUSSION**

275

The GAD system plays an important role in L. monocytogenes under acidic conditions and it 276 277 functions by converting extracellular glutamate to GABA, a reaction that removes protons from the intracellular milieu and helps to maintain pH homeostasis (Karatzas et al., 2010; O'Byrne 278 and Karatzas 2008; Waterman and Small 2003). Although work has increased our knowledge 279 on the function of the GAD system in this organism and other Gram-positive bacteria, the 280 281 effects of environmental conditions and signals on its expression and function are still poorly 282 understood. The only known environmental conditions known to result in increased transcription of the GAD system in this organism are growth in rich media under stationary 283 284 phase (Cotter et al., 2001a; Cotter et al., 2001b; Penfound et al., 1999), anoxic conditions (Feehily et al., 2013; Jydegaard-Axelsen et al., 2004) and gastric fluid (Cotter et al., 285 286 2001a). No other conditions are currently known to activate the system and even the first 287 condition is vague as the specific compounds, other than glutamate, that lead to activation in 288 the rich media are unknown.

289 In our study, we used the chemically defined medium (Amezaga et al., 1995), which was previously shown to prevent functioning of the GADe system in this organism, even upon 290 291 supplementation with glutamate under acidic conditions (Karatzas et al., 2010). This suggests 292 that unknown compounds are essential for GAD activity in L. monocytogenes (Karatzas et al., 2010) and we attempt to identify these compounds in this study. We used this DMG medium 293 as a basis for our screening experiments since L. monocytogenes is unable to export GABA in 294 this medium. Supplementation of this medium with various compounds could give an 295 indication of which compounds are required for GAD system upregulation and/or function. 296 297 Since we also planned to see which part of the GAD system is affected by the supplemented 298 compound(s) we initially assessed if the removal of each one of the decarboxylases in 10403S

had an effect on survival in DMG. No difference was observed (Fig. 15) confirming that each
of the GAD system components is completely inactive in this medium. Subsequently, a variety

of different compounds was tested for their ability to activate the GAD system of L. 301 302 monocytogenes. None of the individual compounds resulted in increased GAD activity 303 however, supplementation of DMG while all three crude mixtures of aminoacids and/or peptides namely, tryptone, peptone and casamino acids resulted in significant GADe 304 305 activity (Fig. 1A). As anoxic conditions have been shown to activate expression of 306 the GAD system in rich media (Jydegaard-Axelsen et al., 2004) we repeated the above 307 experiments under anoxic conditions. Similar results were obtained in oxic conditions however, levels of GABA looked higher (Fig. 1B). These mixes also activated 308 309 the GAD_i system (Fig. <u>1</u>C) and together with the increased GAD_e activity, the overall 310 increased GAD activity resulted in increased acid resistance (Fig. 1A). It should be stated that 311 this increased resistance might also be the result of other mechanisms however, most probably 312 it is majorly the result of GAD system activation since this is the most dominant acid resistance mechanism. 313

It could be suggested that the above observed GAD system activation and increased acid 314 315 resistance is the result of osmotic effects. To clarify this, we could compare the osmotic pressure in DMG supplemented with 0.5 M NaCl with DMG supplemented with the amino 316 317 acid and peptide mixes. However, it is difficult to accurately calculate the levels of osmotic pressure in the DMG supplemented with the amino acid and peptide mixes since they are crude 318 and not defined. However, it is known from chemical analyses of these mixes that their average 319 molecular weight is above 250 Daltons (>250 g/mol). Based on this value a 2, 3.6, 6.8 and 10% 320 solution is equivalent to 0.08, 0.14, 0.27, 0.40 M respectively. Therefore, since 0.5 M NaCl has 321 322 absolutely no effect on GABA export and given that the molarity of all these mixtures is 323 significantly lower than 0.5 M, we could conclude that the effects observed were not related to 324 osmotic pressure.

Interestingly, all three activators identified here were crude mixtures of amino acids and 325 peptides. Casamino acids are a mixture of free amino acids and some very small peptides 326 resulting from the acid hydrolysis of casein (Mueller and Johnson, 1941) whereas tryptone is 327 an assortment of peptides formed by the digestion of casein by the protease trypsin (Fraser and 328 Powell, 1950). Peptone is derived from animal milk or meat digested by proteolysis. It contains 329 peptides, salts, vitamins and many other biological compounds (Payne, 1976). Peptone and 330 331 tryptone are rich in tryptophan whereas casamino acids lack tryptophan because of acid 332 treatment during casein hydrolysis.

333 As seen in Fig. 2B and by comparing the overall effect shown in Fig. 3B, 4B and 5B, supplementation with peptone and tryptone resulted in similarly high levels of GAD_e activity. 334 Furthermore, the effect of casamino acids on GADe activity seemed to be weaker compared to 335 the effect of the above compounds (Fig. 2B) although the effect on GABAi was similar to 336 337 tryptone and peptone (Fig. $\underline{2}$ C). We speculate that some higher peptides contained in these 338 compounds but not in casamino acids might be activating the GADe system. However, it 339 should be stated that under anoxic conditions casamino acids seem to have a higher 340 effect on GADe activity than the other two compounds (Fig. 1). Although supplementation of 341 DMG with these compounds increases GABAe, GABAi levels and acid resistance, the overall 342 GABAe levels achieved by each of these mixes of compounds did not completely correspond to the acid resistance. For example, although supplementation with peptone and tryptone 343 344 resulted in similar GABA_e levels (Fig. 2B), peptone conferred a higher acid resistance than 345 tryptone (Fig. 2A). Furthermore, although supplementation with casamino acids provided a 346 much weaker GAD_e activity compared to tryptone and peptone (Fig. 2B), it conferred the 347 highest acid resistance (Fig. 2A). As stated before, other acid resistance and survival mechanisms might be affected by the supplementation by these compounds. However, it should 348 be noted that the general trend in GADe activity corresponded with the trends in survival. It 349

350 could be seen that upon supplementation with casamino acids, deployment of both GADe and 351 acid resistance was delayed as both curves seem to have an upward trend during the course of 352 the experiment (Fig. 2A, B). In contrast, supplementation with tryptone and peptone resulted in a rapid deployment of GADe (~70% of GABAe is produced within the first 20 min) which 353 354 corresponded well with a higher acid resistance in the initial stages (Fig. 2A, B). As the 355 experiment progressed, GADe activity of cultures supplemented with tryptone or peptone 356 seemed to reduce, which correlated well with a reduction in acid resistance at later stages (Fig. (2A, B). The above suggests that the time of the GAD_e deployment is critical for survival. 357

Furthermore, it should be stated that acid survival at 60 min (Fig.3A) correlated well with the 358 359 $GABA_i$ levels (Fig. <u>2</u>C), with casamino acids having the highest effect followed by peptone 360 and tryptone. However, GABA_i levels are not only the result of GAD_i activity but are also affected by GABA catabolism through the GABA shunt (Feehily et al., 2013). It should be 361 stated though that, high levels of GABAi are possibly associated with higher GADi activity as 362 363 GABA catabolism might not be affected significantly, but this is an area that requires further work. Overall it could be said that peptone and tryptone resulted in higher GADe activity, while 364 365 casamino acids most probably resulted in higher GAD_i activity.

366 Similarly, in previous work casamino acids enhanced prolonged survival of Streptococcus lactis by supplying amino acids and minimizing break down of essential components needed 367 for survival (Thomas and Batt, 1968). However, we are not aware if this strain possessed a 368 GAD system. Another study showed that various E. coli strains and several other bacteria had 369 370 less acidic shock in Tryptone Soy Broth when supplemented with casamino acids, however 371 there was decreased survival in presence of glutamate alone (Park and Diez-Gonzalez, 2004). These authors have suggested that additional amino acids might enhance the glutamate-372 dependent acid resistance, which agrees with what we demonstrate here. Recent work has also 373 demonstrated that the presence of amino acids enhanced the ability of L. monocytogenes to 374

grow under mild acidic conditions although no explanation was offered for this (Muchaambaet al., 2019).

Furthermore, we investigated the effect of 6.8% tryptone 3.6% peptone and 10% casamino 377 acids on strains carrying deletions of the GAD decarboxylase genes. We used different 378 379 concentrations of these compounds in order to obtain measurable levels of inactivation during 380 these survival experiments. The results showed that the $\Delta gadD2$ was the most sensitive either 381 in the presence or absence of peptone or tryptone (Fig. <u>3</u>A, <u>4</u>A). Similarly, in presence of casamino acids $\Delta gadD2$ was the most sensitive (Fig. 5A). Even in presence of either of those 382 383 compounds $\Delta gadD2$ exported less GABA_e than the other mutants and the WT (Fig. <u>3B</u>, <u>4B</u>, 384 **5B**). The results suggest that GadD2 is the GAD component most affected by 385 supplementation. This is expected as GadD2 is the main component of the GADe system that is responsible for GABA export (Cotter et al., 2001a; Cotter et al., 2005) and and highly 386 387 important for survival under acidic conditions (Cotter et al., 2001a). WT, AgadD1 388 and $\Delta gadD2$ were able to export GABA_e in the presence of peptone, tryptone, or casamino acids which also suggests that these compounds might activate the GAD system. This might 389 390 be the reason for the ability of the strains to export GABAe.

Subsequently, to identify how these compounds activate the GAD system, we performed RT-391 qPCR to quantify the transcription of gadD1, gadD2 and gadD3 that encode the three different 392 decarboxylases of the GAD system. We looked at transcription during stationary phase as it is 393 well-known that the GAD system is mainly expressed at this stage of growth. Furthermore, we 394 have shown previously that L. monocytogenes does not respond to acidic conditions by 395 396 significant increase in the transcription of the GAD system genes gadD1, gadD2 and gadD3 (Karatzas et al., 2010; Karatzas et al., 2012). It seems that the GAD system needs to be 397 deployed prior to acidification and therefore we just assessed transcription at stationary phase 398 and not following an acid drop. Furthermore, we did not quantify the transcription of 399

antiporters *gadT1* or *gadT2* as it has been previously confirmed that they follow the
transcription of their corresponding GAD decarboxylases (*gadD1* and *gadD2*) with which they
belong to the same operon (Karatzas et al., 2012).

Previously it has been shown that GadD2 is the most important part of the GAD system under acidic stress (Cotter et al., 2001a; Cotter et al., 2005). In 10403S WT (Fig. 6B) peptone had the strongest effect on *gadD2* transcription followed by casamino acids and tryptone (Fig. 6B). Although previously GadD3 was identified as a part of the GAD_i system, it did not seem to be upregulated by any of the compounds used here and it did not seem to affect survival in any of the experiments. It might be possible that GadD3 plays a role under different conditions or activated by other environmental conditions.

Up to now there is no previous work showing environmental conditions or signals that affect 410 expression of the GAD system in L. monocytogenes and most other microorganisms with the 411 exception of E. coli. Our research contributes to our understanding of the activation of the GAD 412 413 system and the acid resistance in this organism. We show that casamino acids, peptone and 414 tryptone are major GAD system activators resulting in upregulation of the transcription of 415 gadD2, which is the most important component of the GAD system. This research would be 416 important to scientists working on the acid resistance of L. monocytogenes and other organisms. It can also help us predict the behaviour of L. monocytogenes in acidic foods and contribute in 417 the development of strategies to manipulate the acid resistance of L. monocytogenes and 418 possibly other pathogens by restricting the availability of these activators. This could lead to 419 420 the elimination of these pathogens from food and therefore contribute in the reduction of 421 foodborne infections.

422

423 Acknowledgements.

424	We would like to thank all colleagues at the Microbiology research team, Food and Nutritional
425	Sciences, University of Reading. The help of Marcia Boura in reviewing the manuscript and
426	the help and support of our deceased colleague Dr. Bernard Mackey has been invaluable. This
427	work was partly supported by a Marie Curie European Reintegration Grant (grant ERG
428	265154), a Science Foundation Ireland Starting Investigator Research Grant (SIRG; grant
429	09/SIRG/B1570) awarded to K. A. Karatzas, and funds from the University of Reading.

430 431

432 **REFERENCES**

433

434	Amezaga, MR., Davidson, I., McLaggan, D., Verheul, A., Abee, T., Booth, I.R., 1995. The
435	role of peptide metabolism in the growth of Listeria monocytogenes. Microbiol. 141,41-
436	49. doi.org/10.1099/00221287-141-1-41.

- Capitani, G., Biase, D De., Aurizi, C., Gut, H., Bossa, F., Gru, MG., Ch, È., 2003. Crystal
 structure and functional analysis of *Escherichia coli* glutamate decarboxylase. EMBO
 J. 22, 4027-4037. doi:10.1093/emboj/cdg403.
- 440 Cole, S.T., Brosch, R., Parkhill, J., Garnier, T., Churcher, C., Harris, D., Gordon, S.V.,
- 441 Eiglmeier, K., Gas, S., Barry, C.E., Tekaia, F., Badcock, K., Basham, D., Brown, D.,
- 442 Chillingworth, T., Connor, R., Davies, R., Devlin, K., Feltwell, T., Gentles, S., Hamlin,
- 443 N., Holroyd, S., Hornsb, y T., Jagels, K., Krogh, a., McLean, J., Moule, S., Murphy, L.,
- 444 Oliver, K., Osborne, J., Quail, M a., Rajandream, M a., Rogers, J., Rutter, S., Seeger,
- 445 K., Skelton, J., Squares, R., Squares, S., Sulston, J.E., Taylor, K., Whitehead, S., Barrell,
- 446 B.G., 1998. Deciphering the biology of Mycobacterium tuberculosis from the complete
- 447 genome sequence. Nature. 393,537–544. doi:10.1038/31159.
- 448 Cotter, P.D., Gahan, C.G., Hill C., 2001a. A glutamate decarboxylase system protects Listeria

449	monocytogenes in gastric fluid. Mol. Microbiol 40, 465-475. doi.org/10.1046/j.1365-
450	2958.2001.02398.x.

- 451 Cotter, PD., O'Reilly, K., Hill, C., 2001b. Role of the glutamate decarboxylase acid resistance
- 452 system in the survival of *Listeria monocytogenes* LO28 in low pH foods. J. Food. Prot.
 453 64,1362–1368. doi: 10.4315/0362-028x-64.9.1362.
- Cotter, P.D., Ryan, S., Gahan, C.G.M., Hill, C., 2005. Presence of GadD1 glutamate
 decarboxylase in selected *Listeria monocytogenes* strains is associated with an ability to
 grow at low pH. Appl. Environ. Microbiol. 71,2832-2839. doi: 10.1128/AEM.
 71.6.2832-2839.
- Damiano, M. A., Bastianelli, D., Al Dahouk, S., Köhler, S., Cloeckaert, A., De Biase, D.,
 Occhialini, A. 2014. Glutamate decarboxylase-dependent acid resistance in *Brucella*spp.: distribution and contribution to fitness under extreme acid conditions. Appl.
 Environ. Microbiol. 81(2):578-86. https://doi.org/10.1128/AEM.02928-14

462 European Food Safety Authority and European Centre for Disease Prevention and Control 463 (EFSA and ECDC). 201<u>8</u>. The European Union 464 summary report on trends and sources of zoonoses, zoonotic agents and food-borne 465 outbreaks 2017 in Journal, 466 EFSA 16

467 (<u>12</u>). <u>doi.org/10.2903/j.efsa.2018.5500</u>.

Feehily, C., Finnerty, A., Casey, P.G., Hill, C., Gahan, C.G.M., O'Byrne, C.P., Karatzas, K.A.G., 2014. Divergent evolution of the activity and regulation of the glutamate
decarboxylase systems in *Listeria monocytogenes* EGD-e and 10403S: Roles in
virulence and acid tolerance. PloS one 9,e112649. doi:10.1371/journal.pone.0112649.

Feehily, C., O'Byrne, C.P., Karatzas, K.-A.G., 2013. Functional γ-aminobutyrate shunt in
 Listeria monocytogenes: role in acid tolerance and succinate biosynthesis. Appl.

Environ. Microbiol. 79,74-80. doi:10.1128/AEM.02184-12. 474

- Foster, JW., 2004. Escherichia coli acid resistance: tales of an amateur acidophile. Nature Rev. 475 Microbiol. 2,898-907. doi: 10.1038/nrmicro1021. 476
- Hersh, B.M., Farooq, F.T., Barstad, D.N., Blankenhorn, D.L., Hersh, BM., Farooq, F.T., 477
- 478 Slonczewski, J.L., 1996. A glutamate-dependent acid resistance gene in Escherichia coli . J Bacteriol. 178,3978-81. doi: 10.1128/jb.178.13.3978-3981. 479
- 480 Jydegaard-Axelsen, A.-M., Høiby, P.E., Holmstrøm, K., Russell, N., Knøchel, S., 2004. CO2and anaerobiosis-induced changes in physiology and gene expression of different 481 Listeria monocytogenes strains. Appl. Environ. Microbiol. 70,4111-4117. doi: 482 10.1128/AEM.70.7.4111-4117.2004.
- 484

483

- Karatzas, K.-AG., Brennan, O., Heavin, S., Morrissey, J., O'Byrne, C.P., 2010. Intracellular accumulation of high levels of gamma-aminobutyrate by Listeria monocytogenes 485 10403S in response to low pH: uncoupling of gamma-aminobutyrate synthesis from 486 efflux in a chemically defined medium. Appl. Environ. Microbiol. 76,3529-37. 487 doi:10.1128/AEM.03063-09. 488
- Karatzas, K.-A.G., Suur, L., O'Byrne, C.P., 2012. Characterization of the intracellular 489 490 glutamate decarboxylase system: analysis of its function, transcription, and role in the acid resistance of various strains of Listeria monocytogenes. Appl. Environ. Microbiol. 491 78,3571-9. doi:10.1128/AEM.00227-12. 492
- Muchaamba, F., Eshwar, A.K., Stevens, M.J.A., von Ah, U., Tasara T., 2019. Variable carbon 493 source utilization, stress resistance, and virulence profiles among Listeria 494 495 monocytogenes strains responsible for listeriosis outbreaks in Switzerland. Frontiers Microbiol. 10, 957. doi:10.3389/fmicb.2019.00957. 496
- Mueller, J.H., Johnson, E. R., 1941. Acid hydrolysates of casein to replace peptone in the 497 preparation of bacteriological media. J. Immunol. 40,33-38. 498

- Fraser, D., Powell, E., 1950. The kinetics of trypsin digestion. J. Biol. Chem. 187,803-820. 499
- O'Byrne, C.P., Karatzas, K.A.G., 2008. Chapter 5. The Role of Sigma B (oB) in the stress 500
- 501 adaptations of Listeria monocytogenes: Overlaps between stress adaptation and virulence. Adv. Appl. Microbiol. 65,115-140. doi: 10.1016/S0065-2164(08)00605-9
- 502
- O'Byrne, C.P., Feehily, C., Ham, R. and Karatzas, K.A.G., 2011. A modified rapid enzymatic 503 microtiter plate assay for the quantification of intracellular y-aminobutyric acid and 504 505 succinate semialdehyde in bacterial cells. J. Microbiol. Meth., 84 (1), 137-139. doi: https://doi.org/10.1016/j.mimet.2010.10.017. 506
- Park, G.W., Diez-Gonzalez, F., 2004. A novel glutamate-dependent acid resistance among 507 strains belonging to the proteeae tribe of enterobacteriaceae. FEMS Microbiol. Lett. 508 237,303-9. doi: 10.1016/j.femsle.2004.06.050. 509
- Payne, J.W., 1976. Peptides and Micro-Organisms. Adv. Microb. Physiol.13,55-113. 510
- Penfound, T.A., Elliott, J.F., Foster, J.W., Smith, D., 1999. Control of acid resistance in 511 512 Escherichia coli. J Bacteriol . 181,3525-35.
- Shelp, B.J., Bown, A.W., Mclean, M.D., 1999. Metabolism and functions of gamma-513 aminobutyric acid. Trends Plant Sci. 4, 446-452. doi.org/10.1016/S1360-514 515 1385(99)01486-7.
- Su, M.S., Schlicht, S., Gänzle, M.G., 2011. Contribution of glutamate decarboxylase in 516 Lactobacillus reuteri to acid resistance and persistence in sourdough fermentation. 517 Microb. Cell Fact. 10,S8. doi:10.1186/1475-2859-10-S1-S8. 518
- Thomas, T.D., Batt, RD., 1968. Survival of Streptococcus lactis in starvation conditions J. gen. 519 520 Microbiol. 50,367-382. doi: 10.1099/00221287-50-3-367.
- 521 Tsukatani, T., Higuchi, T., Matsumoto, K., 2005. Enzyme-based microtiter plate assay for γ -
- aminobutyric acid: application to the screening of γ -aminobutyric acid-producing lactic 522
- acid bacteria. Anal. Chim. Acta, 540, 293-297 523

524	Water	rman, S.R., Sı	nall, P	.L.C., 20	03. The glut	amate-	dependent acid	resistance system of
525		Escherichia d	coli and	Shigella	<i>i flexneri</i> is i	nhibite	d in vitro by L-	trans-pyrrolidine-2,4-
526		dicarboxylic	acid.	FEMS	Microbiol.	Lett.	224,119–125.	doi:10.1016/S0378-
527		1097(03)0042	27-0.					
528								
529								
530								
531								
532								
533								
534								
535	TAB	LES						
536								

537 Table 1. Strains used in this study

Strain	Description	Nature of mutation	Source reference
10403S	Serotype 1/2a, wild type		(Karatzas et al.,2010)
∆gadD1	10403S isogenic gadD1 mutant	In-frame deletion	(Feehily et al., 2014)
∆gadD2	10403S isogenic gadD2 mutant	In-frame deletion	(Feehily et al., 2014)
∆gadD3	10403S isogenic gadD3 mutant	In-frame deletion	(Feehily et al., 2014)

538 539

540 LEGEND TO FIGURES

541

Fig. 1. Survival after acid challenge stationary phase cultures of 10403S WT and its isogenic
 gad mutants in DMG. Cultures were grown at 37°C with shaking until stationary phase in DMG

544 Fig. 1. Cells of 10403S WT grown until stationary phase in DMG in the presence of 0.5 mg/ml 545 of sodium chloride, 0.5 mg/ml maleic acid, 0.5 mg/ml succinic acid, 0.5 mg/ml acetic acid, 0.5 546 mg/ml lactic acid, 0.1% sodium deoxycholate, 5% oxgall, 5% bile salts, 2%, 10% casamino 547 acids (CA), 3.6% peptone and 6.8% tryptone. Following growth, the pH was adjusted to pH 4 548 the extracellular GABA (GABAe) was measured under (A) oxic and (B) 549 anoxic conditions in DMG. Bars represent an average of measurements 550 performed in triplicate, and error bars represent the standard deviation. D.L. denotes the 551 detecion limit of the GABase method.

552

553 Fig. 2. (A) Cells of 10403S WT grown under oxic conditions in DMG alone (grey 554 circles) or in DMG with 6.8% tryptone (grey diamonds), peptone (black squares), casamino 555 acids (C.A.; black triangles) grown under oxic conditions until stationary phase (~24 h). Subsequently, cultures were acid challenged at pH 2.4 with the addition of 1M HCl. 556 Monitoring GABAe (B) was conducted for 0, 20, 40 and 60 min and GABAi (C) for 60 min. 557 558 GABAe and GABAi were assessed following adjustment of the pH of overnight cultures of 559 10403S WT at pH 4.2 grown until stationary phase in DMG and DMG with 6.8% tryptone, 560 peptone and casamino acids (C.A.). Markers represent an average of measurements performed in triplicate, and error bars represent the standard deviation. D.L denotes detection limit. 561 Asterisk represents statistical significant difference with the control cultures grown in DMG 562 (P<0.05). 563

564

Fig. 3. (A) Cells of 10403S WT (white boxes) and its *gad* mutants $\Delta gadD1$ (white triangles), $\Delta gadD2$ (white circles) and $\Delta gadD3$ (white diamonds) grown in DMG alone or in DMG with 3.6% peptone, with WT (black boxes) and its *gad* mutants $\Delta gadD1$ (black triangles), $\Delta gadD2$ (black circles) and $\Delta gadD3$ (black diamonds) grown <u>under oxic conditions</u> until

569	stationary phase (~24 h). Subsequently, cultures were acid challenged at pH 2.5 with the
570	addition of 1M HCl. Monitoring of $\mbox{GABA}_{e}\left(B\right)$ was conducted for 0, 20, 40 and 60 min and
571	were assessed following adjustment of the pH of overnight cultures of 10403S WT and its gad
572	mutants at pH 4.2 grown until stationary phase in DMG and in DMG with 3.6% peptone.
573	Markers represent an average of measurements performed in triplicate, and error bars represent
574	the standard deviation. D.L denotes detection limit.

575

576 Fig. 4. (A) Cells of 10403S WT (white boxes) and its gad mutants $\Delta gadD1$ (white triangles), 577 $\Delta gadD2$ (white circles) and $\Delta gadD3$ (white diamonds) grown in DMG alone or in DMG with 578 6.8% tryptone, with WT (black boxes) and its gad mutants $\Delta gadD1$ (black triangles), $\Delta gadD2$ 579 (black circles) and $\Delta gadD3$ (black diamonds) grown <u>under oxic conditions</u> until stationary phase (~24 h). Subsequently, cultures were acid challenged at pH 2.5 with the 580 addition of 1M HCl. Monitoring of GABAe (B) was conducted for 0, 20, 40 and 60 min and 581 were assessed following adjustment of the pH of overnight cultures of 10403S WT and its gad 582 583 mutants at pH 4.2 grown until stationary phase in DMG and in DMG with 6.8% tryptone. Markers represent an average of measurements performed in triplicate, and error bars represent 584 585 the standard deviation. D.L denotes detection limit.

586

Fig. 5. (A) Cells of 10403S WT (white boxes) and its *gad* mutants $\Delta gadD1$ (white triangles), $\Delta gadD2$ (white circles) and $\Delta gadD3$ (white diamonds) grown in DMG alone or in DMG with 10% casamino_acids (C.A.), with WT (black boxes) and its *gad* mutants $\Delta gadD1$ (black triangles), $\Delta gadD2$ (black circles) and $\Delta gadD3$ (black diamonds) grown <u>under oxic conditions</u> until stationary phase (~24 h). Subsequently, cultures were acid challenged at pH 2.5 with the addition of 1M HCl. Monitoring of GABA_e (B) was conducted for 0, 20, 40 and 60 min and were assessed following adjustment of the pH of overnight cultures of 10403S WT and its *gad* mutants at pH 4.2 grown until stationary phase in DMG and in DMG with 10%
casaminoacids (C.A.). Markers represent an average of measurements performed in triplicate,
and error bars represent the standard deviation. D.L denotes detection limit.

597

Fig. 6. Real-time PCR determination of transcription of gadD1, gadD2 and gadD3. Relative 598 599 expression of each gene was calculated by comparing expression relative to that of 16S rRNA 600 in 10403S WT cultures grown overnight until stationary phase in DMG or DMG supplemented 601 with 6.8% tryptone, peptone or casamino acids <u>(C.A.)</u>. 602 Markers represent an average of measurements performed in triplicate, and error bars represent 603 Markers represent an average of measurements performed in triplicate, and error bars represent 604 Markers represent average of an measurements performed in triplicate, and error bars represent standard deviations. Asterisks 605 represent statistically significant difference in the expression of each gene in the supplemented 606 media compared to that in DMG (P<0.05). Numbers above the bars indicate fold difference of 607 608 the relative expression for each gene in DMG supplemented with either tryptone, peptone or 609 casaminoacids (C.A.) compared to its expression in DMG without supplementation.

610 611

612 Supplementary data

613 **LEGEND TO FIGURES**

614

Fig. 1. Survival after acid challenge stationary phase cultures of 10403S WT and its isogenic *gad* mutants in DMG. Cultures were grown at 37°C with shaking until stationary phase in DMG
and subsequently, the medium was adjusted to pH 2.4 with the addition of HCl and survival
was assessed after 60 min through preparation of decimal dilutions and plating. Error bars

619 represent standard deviation from triplicate observations.

620