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1	Inhibitory activity of phenolic acids against Listeria monocytogenes:
2	deciphering the mechanisms of action using three different models
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4	
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12 Abstract

13 Phenolic compounds are well known for their antimicrobial activity. They may provide an interesting solution to ensure food safety by preventing the growth of foodborne pathogens 14 15 while addressing the wishes of consumers for the use of natural preservatives in food and favoring the reuse of agro-industry byproducts. However, their mechanism of action is still 16 not very well known. Here, we aimed to decipher the complex mechanism of action of eight 17 18 phenolic acids by decomposing their effects, such as the general effect of the decrease of extracellular pH (γ (pH)) and specific inhibitory effects of the undissociated (γ (A_u)) and 19 dissociated $(\gamma(A_d))$ forms. We thus developed three different models and applied them to a 20 dataset of Listeria monocytogenes growth rates experimentally obtained in the presence of 21 various concentrations of phenolic acids at several pHs. The model that best fits the dataset 22 23 was selected for each phenolic acid to explore the potential mechanisms. The results show that the antimicrobial activity is mainly due to the effect of the undissociated forms, except 24 25 for chlorogenic and gallic acids, for which the antimicrobial activity is mainly due to a 26 decrease in extracellular pH. In addition, the dissociated forms of p-coumaric and ferulic acids show significant inhibitory activity. 27

28

29 Keywords

30 phenolic, antimicrobial, Gamma concept, undissociated form, dissociated form

32 Introduction

Listeria monocytogenes is a ubiquitous, wide spread, highly environmentally resistant (soil, 33 lakes, rivers, etc.) soil bacterium that can contaminate food at all stages of the food chain 34 (Anses, 2011). It causes listeriosis, which affects humans and animals and occurs mainly as an 35 invasive form (Anses, 2011). Since the 90's, the prevalence of L. monocytogenes in many 36 food categories has been reduced due to improved control measures (Buchanan et al., 2017). 37 38 However, the rate of illness has remained constant over the last decade and recent outbreaks have challenged control measures (Allerberger and Wagner, 2010; Buchanan et al., 2017). 39 Ready-to-eat food, meat, fish, and dairy products, as well as fruits and vegetables, are the 40 predominant vehicles involved in the main listeriosis outbreaks (Rodriguez-Lopez et al., 41 42 2018).

43 The use of preservatives can help to inhibit microbial growth or inactivate pathogenic bacteria. However, this approach is increasingly challenging due to the emergence of bacterial 44 45 resistance, mistrust of consumers towards chemical additives, and regulatory constraints that 46 reduce the list of protective ingredients. Thus, there is renewed interest in several families of natural antimicrobials (Sorrentino et al., 2018; Weiss et al., 2015). Simple phenols, such as 47 eugenol, thymol, and carvacrol, are found in aromatic plants and recovered in high 48 49 concentrations in essential oils and hydrosols. They intercalate into the phospholipid layers of the bacterial membrane and disturb the van der Waals interactions between the lipid acyl 50 chains, leading to the disruption of phospholipid packing and membrane integrity (Burt, 51 2004). Consequently, ion gradients are disrupted and vital constituents, such as ions and 52 53 macromolecules, are released, leading to bacterial death.

Organic acids act by a very well-known antimicrobial mechanism through the penetration of
their undissociated form into the cell and acidification of the cytoplasm leading to cell death.
Phenolic acids are of interest due to their natural plant-based origin, their presence in large

quantities in byproducts of the fruit, wine and cereal industries and their demonstrated global
antimicrobial activity. We previously evaluated the inhibitory activity of several phenolic
compounds against *Listeria monocytogenes* as a function of their total concentration (Pernin
et al., 2018). However, very few studies have focused on deciphering the various effects of
their inhibitory activity, which can be similar to those of organic acids and/or simple phenols
(Sorrentino et al., 2018).

We reasoned that the total inhibitory activity is a combination of the following effects: *i*) decreased pH in the presence of phenolic acids which directly inhibits growth, *ii*) the inhibitory effect of the undissociated acid form, and *iii*) the inhibitory effect of the dissociated acid form. Each acid form could potentially either intercalate into the phospholipid membrane or cross the membrane and decrease the intracellular pH and/or interact with cellular

68 constituents.

69 Here, we aimed to decipher the complex mechanism of action of phenolic acids by

70 decomposing the various inhibitory effects. We thus developed three different models using

the Gamma concept approach (Lambert and Bidlas, 2007; Zwietering et al., 1992). These

72 models aim to describe the proportion of inhibition due to the three potential effects. The

raight phenolic acids (*p*-hydroxybenzoic, protocatechuic, gallic, vanillic, *p*-coumaric, caffeic,

74 ferulic and chlorogenic acids) studied were chosen based on their efficiency in our previous

study (Pernin et al., 2018). The model that best fits the dataset was selected for each phenolicacid to explore the potential mechanisms.

77

78 2. Materials and methods

79 2.1. Chemical reagents

Hydrochloric acid (1 mol/L), sodium hydroxide (1 mol/L), and acetone were purchased from
Carlo Erba (Fontenay-aux-Roses, France).

82

83 2.2. Bacterial strain

The strain used in this study was *Listeria monocytogenes* CNL 895805, serotype ½ a, isolated
from sheep brain. It was graciously provided by P. Velge (INRA, Nouzilly) (Van
Langendonck et al., 1998). Before each experiment, the strain, stored in cryovials at -80°C,
was resuscitated in two successive subcultures in tryptic soy broth (TSB, Biomérieux, France)
at 30°C.

89

90 2.3. Phenolic compounds

91 *p*-Hydroxybenzoic acid, protocatechuic acid, vanillic acid, gallic acid, *p*-coumaric acid,

92 caffeic acid, ferulic acid, and chlorogenic acid were purchased from Sigma-Aldrich (St

93 Quentin Fallavier, France). The chemical structures and physicochemical parameters (pKa,

94 logP) are provided in Table 1.

trace acetone after evaporation.

95 Stock solutions of the phenolic compounds were prepared according to specific protocols

96 depending on their ability to be solubilized in the culture medium. *p*-Hydroxybenzoic,

97 protocatechuic, gallic, and chlorogenic acid powders were directly dissolved in the culture

98 medium. Vanillic, *p*-coumaric, and ferulic acids were first dissolved in acetone, which was

99 evaporated under nitrogen flow after addition to the culture medium. Caffeic acid was first

dissolved in acetone/distilled water (80/20 (v/v)) and the acetone evaporated under nitrogen

101 flow. Bacterial growth controls were carried out to ensure the absence of inhibitory activity of

103

102

104 2.4. Growth in the presence of phenolic compounds at various pHs

105 The growth of *L. monocytogenes* was followed in TSB containing one of the eight phenolic

acids at various pHs. The phenolic acids were added at various concentrations in either neutral

107 TSB pH 7.2 (7.1-7.4) or acidic TSB pH 5.5 (5.5-5.7). pH 7.2 is the native pH of TSB and it is

108 optimal for *L. monocytogenes* growth (Anses, 2011). pH 5.5 allows to increase the

109 concentrations of undissociated forms of acids without decreasing too much the growth rate of

110 *L. monocytogenes.* After solubilization of the phenolic acid at the target concentration in TSB

111 7.2, the pH was adjusted to 7.2 (7.2a) with 1 mol/L sodium hydroxide or not adjusted (7.2na)

112 (Pernin et al., 2018). Similarly, TSB 5.5 was adjusted to a pH of 5.5 (5.5a) or not adjusted

113 (5.5na). The pH of each culture medium was measured with an SI Analytics lab 870 pH-meter

114 (Mainz, Germany) in an independent experiment. Gallic, caffeic, and chlorogenic acids were

not tested in TSB 7.2a because the medium became brown after pH adjustment, probably due

116 to oxidation.

117 The final concentration of the phenolic compounds in TSB varied from 0 to at least 30

118 mmol/L. For some compounds, the minimal inhibitory concentration was not achieved due to

solubility issues. Eight to 64 concentrations prepared from at least two different stock

solutions were tested per compound for each series of experiments and a total of 47 to 146

assays were conducted for each compound.

122 The largest possible range of conditions was tested to allow the best adjustment of the models: total acid concentration, concentration of undissociated or dissociated forms, and pH. Series 123 7.2na covered pH ranges from 7.2 to 3.2 (0 to 10 mmol/L of the undissociated form and 0 to 124 20 mmol/L of the dissociated form) and series 5.5na, 5.5 to 4.0 (0 to 15 mmol/L of the 125 undissociated form and 0 to 10 mmol/L of the dissociated form). Series 7.2a and 5.5a were set 126 to the pH of the initial TSB batch, around 7.2 and 5.5, respectively. The concentration of the 127 undissociated form was close to zero and that of the dissociated form from 0 to 30 mmol/L for 128 series 7.2a. The corresponding values for the series 5.5a were from 0 to 1.5 mmol/L and 0 to 129 25 mmol/L. 130

TSB at pH adjusted from 3 to 10 with 1 mol/L hydrochloric acid or 1 mol/L sodium 131 132 hydroxide solutions was prepared to evaluate the effect of pH on bacterial growth in the absence of the phenolic acids. In total, 19 pHs were tested in two independent experiments. 133 TSB of each series was inoculated at 1% (v/v from the second subculture; approximately 10^6 134 CFU/mL) with a standardized inoculum. Two hundred microliters of inoculated TSB from 135 each series was added to the wells of 100-well microplates which were incubated at 30°C 136 with slow continuous shaking. Bacterial growth was followed in an automatic 137 spectrophotometer (Bioscreen C, Labsystems, Helsinki, Finland) by measuring the optical 138 density (OD) at 600 nm for 72 h. In total, 1408 growth kinetics were acquired. 139 140

141 2.5. Determination of the growth rate

Maximum specific growth rates (µmax) of *L. monocytogenes* were estimated from the growth
kinetics by fitting the modified Gompertz model (Guillier et al., 2007; Pernin et al., 2018).
The standard deviations of the model parameters and sum of squares were calculated using
the complementary macro SolverAid (de Levie, 2012).

146

147 2.5. Modelling the effect of different acid forms on the growth rate

The effect of phenolic acids on the growth rate of *L. monocytogenes* was modelled using the
Gamma concept approach (Lambert and Bidlas, 2007; Zwietering et al., 1992). The effects of
pH and undissociated and dissociated forms of the acid were taken into account (eq. (1)).

151
$$\mu_{max} = \mu_{ref} \cdot \gamma(pH) \cdot \gamma(A_u) \cdot \gamma(A_d) \qquad \text{eq. (1)}$$

152 where:

153 μ_{ref} is the *L. monocytogenes* maximum specific growth rate at 30°C in TSB at optimal pH and 154 without phenolic acid and γ (pH) describes the pH effect as described in eq. (2) (Presser et al., 155 1997).

156
$$\gamma(pH) = 1 - \frac{10^{pH_{min}}}{10^{pH}}$$
 eq. (2)

157 The pH_{min} value was set to 4.24 (Augustin et al., 2005).

158 $\gamma(A_u)$ describes the effect of the undissociated form as described in eq. (3) (Presser et al.,

159 1997).

160
$$\gamma(A_u) = \left(1 - \frac{A_{tot}}{MIC_u \cdot (1 + 10^{pH - pKa})}\right)^{\alpha} \qquad \text{eq. (3)}$$

where MIC_u is the minimum inhibitory concentration of the undissociated form of the acid, α is a shape parameter and A_{tot} is the concentration of total acid.

163 $\gamma(A_d)$ describes the effect of the dissociated form as described in eq. (4) (Presser et al., 1997).

164
$$\gamma(A_d) = \left(1 - \frac{A_{tot}}{MIC_d \cdot (1 + 10^{pKa - pH})}\right) \qquad \text{eq. (4)}$$

165 where MIC_d is the minimum inhibitory concentration of the dissociated form of the acid.

166

167 Three different models are proposed.

168 In model #1, pH is the only factor taken into account and μ_{ref} is the only parameter to estimate 169 (eq. (5)).

170
$$\mu_{max} = \mu_{ref} \gamma(pH) \qquad \text{eq. (5)}$$

171 In model #2, pH and A_u are the factors taken into account and μ_{ref} , MIC_u and α are the

172 parameters to estimate (eq.(6)).

173
$$\mu_{max} = \mu_{ref} \cdot \gamma(pH) \cdot \gamma(A_u) \qquad \text{eq. (6)}$$

In Model #3, pH, A_u and A_d are the factors taken into account and μ_{ref} , MIC_u, α and MIC_d are the parameters to estimate (eq.(7)).

176
$$\mu_{max} = \mu_{ref} \cdot \gamma(pH) \cdot \gamma(A_u) \cdot \gamma(A_d) \qquad \text{eq. (7)}$$

177

178 2.6. Statistical approach

- The model parameters were fitted with Excel solver according to minimization of the residual
 sum of square errors (RSS) for the tested mathematical model. The three models were
- 181 compared according to the Bayesian information criterion (BIC) (eq. (8)).

182
$$BIC = n \cdot ln\left(\frac{RSS}{n}\right) + k \cdot \ln(n) \qquad \text{eq.(8)}$$

183 where n is the number of experimental points and k the number of parameters of the model.

3. Results and Discussion

185 *3.1. Characterization of the models*

Modeling with the Gamma concept approach (Lambert and Bidlas, 2007) allowed 186 quantification of the relative importance of each impact factor. It is here difficult to show a 187 graphical representation of the whole dataset of observed and predicted growth rates because 188 of there are four relevant impact factors: pH, Atot, Au, Ad. Thus, we show in Figure 1 the 189 190 predicted μ_{max} as a function of the observed μ_{max} for each phenolic acid. It illustrates that the chosen models fit well the datasets whatever the compound. Moreover, Figure 2 shows how 191 the models and the Gamma concept approach work. It gives more details about the fitness of 192 model #3 applied to the ferulic acid dataset (Figures 2a and b). The growth rates obtained with 193 various concentrations are represented as a function of the concentrations of the undissociated 194 195 and dissociated forms for two pHs (5.6 and 7.3). A given concentration of total acid at a given pH corresponds to a concentration of the undissociated and dissociated forms, together with 196 an observed and a predicted μ_{max} (Figure 2c). The model provides a value for the reduction of 197 198 μ_{max} independently associated with the undissociated form, dissociated form, and pH. For example, when ferulic acid is introduced at 0.91 mmol/L in TSB adjusted to pH 5.6, the 199 growth rate μ_{max} is divided by 1/0.57 (that is, 1/ γ) due to the undissociated form, 1/0.99 due to 200 the dissociated form, and 1/0.96 due to the decrease in pH, relative to the growth rate in the 201 202 absence of the phenolic acid. In this example, the antimicrobial effect of the undissociated form is far more predominant than that of the dissociated form and pH (Figure 2c). When 203 ferulic acid is introduced at 30.28 mmol/L in TSB adjusted to pH 7.3, the growth rate μ_{max} is 204 205 divided by 1/0.66 due to the undissociated form and 1/0.52 due to the dissociated form. The antimicrobial effect of the dissociated form is here almost the same as that of the 206 undissociated form. The pH has no effect ($\gamma = 1.00$), as this pH is optimal for the growth of L. 207 208 monocytogenes. Such tables, which set the three Gamma values for all concentrations of each

phenolic acid help in understanding the relative impact of each factor in the inhibition ofbacterial growth and are provided in Supplementary data.

We compared the BICs obtained for each of the three models for each phenolic acid (Table 211 212 2): the lower the BIC, the better the model fits the dataset. The use of the Gamma concept approach together with the calculated BIC allows adjustment of the number of parameters to 213 214 that which is most relevant. The parsimony principle (Ross and Dalgaard, 2004) states that "models should have no more parameters than are required to describe the underlying 215 behavior studied. Too many parameters can lead to a model that fits the error in the data, *i.e.* 216 generates a model that is specific to a particular set of observations". We thus deleted 217 218 parameters that did not improve the BIC.

In models # 2 and #3, a shape parameter describing the slope of the curve $\mu = f(A_u)$ at low

220 concentrations is estimated for each phenolic acid (Table 2). It accounts for the potential non-

linear shape of the decrease in μ when the concentration of the undissociated form increases.

222 The implication of the α parameter value for each phenolic acid and its relation to the

antimicrobial mechanism will be discussed in section 3.2.2.

224 To our knowledge, there is no other description in the literature of a model that takes into account the undissociated and dissociated forms in the inhibitory activity of phenolic acids. A 225 very small number of studies have attempted to model the antimicrobial activity of phenolic 226 227 acids, such as that of Ramos-Nino et al., (1996), which linked the inhibitory activity of phenolic acids with their logP and pKa (Ramos-Nino et al., 1996). In contrast, more studies 228 have attempted to model the antimicrobial activity of organic acids. They have used various 229 types of models, such as principle component analysis (Hsiao and Siebert, 1999; Nakai and 230 Siebert, 2003) or multivariate linear regression (George et al., 1996). However, only a few 231 have studied the effect of the dissociated forms (Eklund, 1985, 1983; Presser et al., 1997). 232 Eklund proposed a mathematical model to calculate the effect of the undissociated and 233

dissociated forms of sorbic, benzoic, and propionic acid on several microorganisms (Eklund, 234 235 1985, 1983). In this model, the specific effect of each form was calculated from the measured MIC of the total form of the organic acid at different pHs (from pH 4.6 to 7.6). 236 237 The model of Presser et al. (1997) is closer to ours. These authors modeled growth rates of *Escherichia coli* as a function of pH and lactic acid concentration, taking into account a 238 suboptimal pH term, an undissociated organic acid term, and a dissociated organic acid term 239 (Presser et al., 1997). A shape parameter, α, for the undissociated acid term was added in our 240 study, together with application of the parsimony principle, by combining the Gamma concept 241 approach and calculation of the BIC. 242 243

244 3.2. Hypothesis on the antimicrobial mechanism based on the best-fitting model

Model #2 fits the best for caffeic, chlorogenic, gallic, *p*-hydroxybenzoic, protocatechuic, and
vanillic acids. Model #3 fits the best for *p*-coumaric acid and ferulic acid (Table 2). Model #1
never provided a best fit. Thus, considering only decreases in extracellular pH cannot
completely describe the dataset for any of the tested phenolic acids.

249

250 *3.2.1. Decreasing extracellular pH: the main antimicrobial mechanism of action for*

251 chlorogenic and gallic acids

Although the best-fitting model for chlorogenic acid and gallic acid was model #2 (Table 2),

the calculated MIC_u are extremely high, above the solubility thresholds of these two

compounds. At pH 5.5, 1,766.72 mmol/L total chlorogenic acid and 637.06 mmol/L gallic

acid would be needed to reach the MIC_u, at least 16-fold higher than that for protocatechuic

- acid. Thus, even if the undissociated forms of chlorogenic acid and gallic acid significantly
- 257 decreased the growth rate, their effect was very small relative to that of other phenolic acids.
- In addition, the datasets of the bacterial growth rates obtained as a function of pH for various

concentrations of chlorogenic and gallic acids clearly overlap to those obtained for

260 hydrochloric acid (Figure 3a). Moreover, at adjusted pH 5.5, increasing the concentration of

261 chlorogenic or gallic acids did not have any visible impact on growth rates (Figure 3b).

262 Similarly, Wen *et al.* (2003) showed that chlorogenic acid had no antimicrobial activity on *L*.

263 *monocytogenes* at pH 5.5 adjusted (Wen et al., 2003). Indeed, the mode of action for these

two acids is mainly due to the decrease of extracellular pH.

In order to better illustrate the mechanism of action of these acids, we compared the growth

rates in the presence of gallic acid to those in presence of vanillic acid. Addition of either acid

to the culture medium to a total acid concentration of 10 mmol/L reduced the pH to

approximatively 4.6. However, according to the MIC_u and pKa values, γ (pH) = 0.54 and γ (A_u)

269 = 0.43 for gallic acid and $\gamma(pH) = 0.58$ and $\gamma(A_u) = 0.00$ for vanillic acid (Supplementary

270 data). The inhibitory effect of the undissociated form was significant for gallic acid but

271 limited to a reduction by 1/0.43. In contrast, it was far more predominant for vanillic acid, as272 it completely inhibited growth.

273 Most published studies do not separate the specific inhibitory effect due to the undissociated

form and the decrease of extracellular pH. However, chlorogenic acid and gallic acid have

been shown to be relatively ineffective antimicrobials among phenolic acids (Gutiérrez-

276 Larraínzar et al., 2012; Pernin et al., 2018; Saavedra et al., 2010; Sánchez-Maldonado et al.,

277 2011; Wen et al., 2003).

Several hypotheses can be formulated to explain the lack of antimicrobial activity of the
undissociated acid form of chlorogenic and gallic acids. First, their high steric hindrance,
especially that of chlorogenic acid, could limit their penetration through the membrane even if
Ramos-Nino *et al.* (1996) showed that this parameter was not relevant to describe the
antimicrobial efficiency of hydroxybenzoic and hydroxycinnamic acids. Second, their low
partition coefficient could limit their partition into the bacterial membrane (Table 1).

However, organic acids are highly active while they have low partition coefficient (Hirshfield 284 285 et al., 2003; Lu et al., 2011). In addition, several studies have demonstrated an antimicrobial effect of the undissociated form of chlorogenic acid and gallic acid at the membrane level 286 287 (Borges et al., 2013; Lou et al., 2011). Gallic acid was shown to modify the membrane hydrophobicity and charge of L. monocytogenes and to cause local ruptures or pore formation 288 in the cell membranes (Borges et al., 2013). Chlorogenic acid induces outer membrane 289 290 modifications and loss of membrane integrity of Streptococcus pneumoniae and Shigella dysenteriae (Lou et al., 2011). 291

292

3.2.2. The antimicrobial effect of the undissociated forms: the main antimicrobial mechanism
of action for caffeic, vanillic, protocatechuic and p-hydroxybenzoic acids.

Most of the phenolic acids tested, namely caffeic acid, *p*-hydroxybenzoic acid, protocatechuic 295 296 acid, and vanillic acid, best fit with model #2, with realistic MIC_u values (Table 2). The comparison of phenolic acids with different α shape parameters resulted in different profiles 297 298 (Figure 4). Some acids, such as protocatechuic acid and caffeic acid, display a linear antimicrobial effect when concentration of undissociated form of phenolic acids increase ($\alpha =$ 299 300 1). In contrast, phenolic compounds with $\alpha > 1$ have a steeper slope, as shown for vanillic acid ($\alpha = 8.20$) and *p*-hydroxybenzoic acid ($\alpha = 11.11$) in model #2 represented at pH 5.5 (Figure 301 302 4). The higher the value of α (>1), the higher the effect of an additional molecule at low 303 concentrations. Compounds with $\alpha = 1$, such as caffeic and protocatechuic acids, have very close chemical structures (2 OH groups on the phenolic ring; Table 1). Differences between α 304 could be due to differences in the ability of these phenolic acids to penetrate through or 305 306 accumulate in the bacterial membrane. This is the first report characterizing a shape parameter α for the effect of undissociated forms of phenolic acids. This parameter can be a key factor in 307 choosing an antimicrobial agent that will be highly active at very low concentrations. 308

310	3.2.3. Phenolic acids with a dissociated form that is significantly antimicrobial
311	Model #3 best fits the <i>p</i> -coumaric acid and ferulic acid datasets. The antimicrobial effect of
312	the undissociated form and decrease in the extracellular pH cannot explain the entire mode of
313	action. The introduction of ferulic acid at 30.28 mmol/L in TSB adjusted to pH 7.3, divided
314	the growth rate μ_{max} by 1/1.00 due to the pH, 1/0.66 due to the undissociated form, and 1/0.52
315	due to the dissociated form relative to that of the control (Figure 2c). Similarly, introduction
316	of <i>p</i> -coumaric acid at 31.55 mmol/L in TSB adjusted to pH 7.4 divided the growth rate by
317	1/0.78 due to the undissociated form and 1/0.57 due to the dissociated form relative to that of
318	the control (Supplementary data). There are several examples for <i>p</i> -coumaric acid and ferulic
319	acid in which $\gamma(A_d)$ was close to or even lower than $\gamma(A_u)$ (Supplementary data). Under
320	specific conditions, the inhibitory effect of the dissociated form predominates over that of the
321	undissociated form and pH. In contrast, $\gamma(A_d)$ never dropped below 1.00 for <i>p</i> -
322	hydroxybenzoic, caffeic, or protocatechuic acids or below 0.93 for vanillic acid when model
323	#3 was applied to other phenolic acids (Supplementary data). p-Coumaric and ferulic acids
324	are therefore the most efficient antimicrobials at pH 7.2, relative to the other phenolic acids,
325	due to their dissociated form which plays a significant role in decreasing the growth rate.
326	Nevertheless, their MIC_d values calculated for pH 7.2 are very high, 72.63 and 63.67 mmol/L,
327	respectively, which is above their solubility threshold. These results show that <i>p</i> -coumaric
328	acid and ferulic acid can be efficient antimicrobial agents, even at neutral pH, and this feature
329	is important for the choice of compounds in several applications.
330	This is the first report of the inhibitory efficacy of the dissociated form of a phenolic acid. The
331	MIC _d for <i>p</i> -coumaric acid and ferulic acid against <i>L. monocytogenes</i> are 10-times lower than
332	the average order of magnitude found in the literature for organic acids on several bacteria:
333	lactate (800-1250 mmol/L (Presser et al., 1997)), propionate (380-830 mmol/L (Eklund,

1985)), benzoate (90-200 mmol/L (Eklund, 1985)), and sorbate (50-400 mmol/L (Eklund,
1983)).

Only a few hypothesis has been previously proposed concerning the potential antimicrobial 336 337 mechanism of action of the dissociated form of organic acids such as effects on cell wall and membrane, inhibition of substrate transport or of key enzymes in the case of sorbates (Sofos 338 et al., 1986). Interestingly, *p*-coumaric and ferulic acids share similarities in their chemical 339 structures. They both contain a cinnamic group between the phenolic ring and the carboxylic 340 function and have only one OH group on their phenolic ring (Table 1). Thus, it is possible that 341 specific chemical structures can interact with the bacterial membrane or specific receptors. 342 343 Moreover, the logPs of p-coumaric and ferulic acids at pH 8.0 (where ~ 100% of the acid is dissociated) are the highest among phenolic compounds: -1.58 and -1.81, respectively (Table 344 1). Their « higher » partition coefficient may favor interaction with the bacterial membrane, 345 346 even if these values are far from the optimal logP values described for this activity (Ultee et al., 2002). Further studies are needed to better understand the antimicrobial mechanism of 347 348 action of the dissociated forms of both organic and phenolic acids.

349

350 Conclusion

351 We present here an original modeling approach which allows a better understanding of the mechanism of action of eight phenolic acids. It quantifies the role of the three different 352 353 antimicrobial factors (decrease in extracellular pH and the undissociated acid form and dissociated acid form). The results allowed classification of these phenolic acids into three 354 categories. First, chlorogenic acid and gallic acid mainly inhibit the growth of L. 355 356 monocytogenes through their ability to decrease extracellular pH, the molecules themselves having a low antimicrobial activity on bacterial growth. Second, caffeic acid, p-357 hydroxybenzoic acid, protocatechuic acid, and vanillic acid exert antimicrobial behavior 358

359	mainly	through	the ability	of their	undissociated	form to	o inhibit the	growth of	L.
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360 *monocytogenes*. Third, *p*-coumaric acid and ferulic acid similarly exert antimicrobial behavior

through their undissociated form, but their dissociated form also shows significant

362 antimicrobial activity. These results will help in choosing the most relevant antimicrobial

363 compounds based on product characteristics, particularly pH.

364

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367 declare no competing financial interest.

368

369 Abbreviations used

- 370 MIC = minimal inhibitory concentration
- 371 BIC = Bayesian information criterion
- 372 TSB = trypic soy broth

373

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- 465 Figure 1 Representation of the predicted μ_{max} as a function of the observed μ_{max} for the eight
- 466 phenolic acids and their best fitting models (#3 for ferulic and *p*-coumaric acids, #2 for the others)
- 467 Figure 2 Fitted complete model #3 with an example of the partial dataset of ferulic acid (a) at pH 7.3
- and (b) at pH 5.6 and (c) associated table of gamma values for the undissociated form, dissociatedform and pH.
- 470 Figure 3 *L. monocytogenes* growth rates (a) as a function of pH in the presence of various 471 concentrations of chlorogenic and gallic acids compared to those at different pHs adjusted with 472 hydrochloric acid and (b) as a function of the concentrations of chlorogenic and gallic acids at pH 5.5a 473 Figure 4 – *L. monocytogenes* growth rates as a function of the concentrations of undissociated acid at
- 474 pH 5.5, for four phenolic acids with the best fitting model #2.

475 List of tables

- 476 **Table 1** Chemical structures and physico-chemical parameters of the eight studied phenolic acids
- 477 **Table 2** Estimated parameters and BIC for the three models and the eight phenolic acids
- 478 Unshaded areas correspond to the best fitting model for a given phenolic acid





Total	Undissociated	Dissociated			Reductio		Predicted μ_{max}	
ferulic acid (mmol/L)	acid form (mmol/L)	acid form (mmol/L)	pН	Observed μ_{max} (h ⁻¹)	undissociated form $\gamma(A_u)$	dissociated form γ(A _d)	pH γ(pH)	$\begin{array}{c} (\textbf{model #3}) \\ \mu_{\text{ref}} \gamma(A_d) \cdot \gamma(A_u) \cdot \\ \gamma(pH) \end{array}$
0.00	0.00	0.00		1.25	1.00	1.00		1.11
0.60	0.00	0.60		1.21	0.99	0.99		1.09
5.13	0.01	5.12		1.01	0.94	0.92		0.96
10.26	0.02	10.24	7.2	0.82	0.88	0.84	1.00	0.82
15.14	0.03	15.11	1.5	0.64	0.81	0.76	1.00	0.69
20.29	0.04	20.25		0.52	0.76	0.68		0.57
25.65	0.05	25.60		0.41	0.73	0.60		0.48
30.28	0.06	30.22		0.36	0.66	0.52		0.38
0.00	0.00	0.00		0.00	1.00	1.00		1.06
0.00	0.08	0.83	56	0.45	0.57	0.99	0.96	0.60
5 18	0.45	4 73	2.0	0.07	0.02	0.93	0.00	0.02
8.23	0.72	7.51		0.00	0.00	0.88		0.00





Phenolic acid	R1	R2	Name	pKa (ChemIDPlus, PubChem)	logP (ChemSpider)	logP of undissociated form, determined at pH=1.7 (Chemicalize)	logP of dissociated form, determined at pH=8.0 (Chemicalize)
Hydroxybenzoic acids Q	Н	Н	<i>p</i> -hydroxybenzoic acid	4.54	1.42	1.33	-1.95
	Н	ОН	protocatechuic acid	4.26	1.16	1.02	-2.35
ОН	Н	O-CH ₃	vanillic acid	4.51	1.33	1.17	-2.19
 R ₂	ОН	ОН	gallic acid	4.40	0.91	0.72	-2.73
Hydroxycinnamic acids O	Н	Н	<i>p</i> -coumaric acid	4.64	2.43	1.83	-1.58
	Н	ОН	caffeic acid	4.62	1.42	1.53	-1.98
	Н	O-CH ₃	ferulic acid	4.58	1.64	1.67	-1.81
	quinic acid	ОН	chlorogenic acid	2.66	-0.36	-0.28	-3.79

	pH _{min}						Mode	els				
Phenolic acids		#	1		#	\$2				#3		
		μref	BIC	μref	α	MICu	BIC	µref	α	MICu	MICd	BIC
Chlorogenic acid		0.95	- 228	1.03	6.34	2.55	- 241	1.04	6.33	2.55	6420.34	- 237
p-Coumaric acid		0.34	- 157	1.11	3.69	0.94	- 450	1.11	3.70	0.94	72.43	- 530
Caffeic acid		0.41	- 166	1.08	1.00	1.06	- 476	1.04	0.64	0.91	66 356.72	- 372
Gallic acid	1 2 1	0.84	- 173	1.10	9.39	46.88	- 230	1.11	9.35	47.10	1 000.00	- 225
Vanillic acid	4.24	0.48	- 177	1.14	8.20	3.24	- 552	1.16	7.70	3.09	424.09	- 549
Protocatechuic acid		0.76	- 169	1.12	1.00	2.12	- 309	1.18	4.21	5.07	20 718.84	- 297
<i>p</i> -Hydroxybenzoic acid		0.48	- 121	1.29	11.11	3.30	- 337	1.29	10.17	3.06	41 176.98	- 333
Ferulic acid		0.25	- 133	1.06	8.00	1.20	- 485	1.11	8.55	1.26	63.52	- 536

Mechanism 1 Mechanism 2 Decreasing Antimicrobial extracellular pH undissociated form $A_u \rightarrow A_d + H^{-1}$ → A_d + H pН pH

Chlorogenic acid Gallic acid Protocatechuic acid p-Hydroxybenzoic acid Vanillic acid Caffeic acid

Ferulic acid *p*-Coumaric acid

Mechanism 3 Antimicrobial undissociated and dissociated forms

