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8 **Probiotic potential of autochthone microbiota from dry-cured sheep ham**

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10 *Probiotisches Potenzial der autochthonen Mikrobiota aus trocken gereiftem Schafschinken*

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14

15 **Summary**

16 This study assessed the potential of probiotic characteristics of bacterial strains isolated from
17 dry-cured sheep ham. It is one of the most common autochthonous processed meat products
18 made in a traditional way on the Pešter plateau (Western Serbia). Isolates were identified as
19 *Lactobacillus curvatus* (9 strains), *Lactobacillus sakei* (3 strains), and *Enterococcus faecium*
20 (4 strains) using MALDI-TOF mass spectrophotometry. The study of probiotic characteristics
21 of 16 dry-cured sheep ham isolates included survival rate through the gastrointestinal tract (GI),
22 the possibility of biogenic amine synthesis, growth on medium with different concentrations of
23 phenol, and antimicrobial activity. The results showed that in simulated gastric juice
24 conditions, the cell number decreased after the first hour of incubation in the tested strains of
25 *Lb. curvatus*, *Lb. sakei* and *En. faecium* except in the case of *Lb. curvatus* Ios19 where the
26 number of cells remained approximately the same. After the second hour of incubation, the
27 number of cells generally remained at the level of the first hour except in the case of the
28 following isolates: *Lb. sakei* Ios12, *Lb. curvatus* Ios18 and *En. faecium* Ios24, where an
29 increase in the number of cells was noticed after the second hour of incubation.

30 In simulated small intestine conditions, an increase in the number of vital cells after 4 and 6
31 hours of incubation was observed in the isolates *Lb. curvatus* Ios4, *Lb. sakei* (Ios12,
32 Ios13), and *En. faecium* Ios1a. Synthesis of biogenic amines was not observed in
33 investigated lactobacilli and enterococci. Analyzed isolates exhibited growth on media with
34 0.1% and 0.2% phenol, while 5 isolates exhibited decarboxylase activity. Six *Lactobacillus*

35 strains, *Lb. curvatus* (IIos6, IIos17, and IIIos1), *Lb. sakei* (IIIos16, Ios12, and IIIos13) and
36 *En. faecium* Ios4 inhibited the growth of tested pathogens, including *Escherichia coli* ATCC
37 25922, *Staphylococcus aureus* ATCC 25923, *Listeria monocytogenes* ATCC 19115,
38 *Pseudomonas aeruginosa* ATCC 27853, and *Bacillus cereus* ATCC 14579.

39 Keywords: Lactic acid bacteria (LAB); probiotic properties; simulated gastric juice
40 conditions; bile salt tolerance.

41 **Zusammenfassung**

42 Diese Studie untersuchte das Potenzial von probiotischen Eigenschaften von Stämmen, die
43 aus trocken geheilten Schafschinken isoliert wurden. Es ist eines der am meisten verbreiteten
44 autochthonen Fleischprodukte, die traditionell auf dem Pester Plateau (Westserbien)
45 hergestellt werden. Die Untersuchung der probiotischen Eigenschaften von 16 Isolaten von
46 trocken geheilten Schafen beinhaltete die Überlebensrate durch den Gastrointestinaltrakt
47 (GI), die Möglichkeit einer biogenen Aminsynthese, das Wachstum auf einem Medium mit
48 verschiedenen Konzentrationen von Phenol und antimikrobielle Aktivität. Die Ergebnisse
49 zeigten, dass unter simulierten Magenbedingungen alle Stämme von *Lactobacillus curvatus*
50 und *Lactobacillus sakei* eine hohe Resistenz gegen 0,3% Konzentration von Gallensalzen
51 aufwiesen, während *Enterococcus faecium* Isolate einen signifikant geringeren Grad an
52 Resistenz zeigten. Unter simulierten Dünndarmbedingungen wurde in den Isolaten *Lb.*
53 *curvatus* IIos4, *Lb. sakei* (Ios12, IIIos13) und *En. faecium* Ios1a eine Zunahme der Anzahl
54 vitaler Zellen nach 4 und 6 Stunden Inkubation beobachtet. Die Synthese von biogenen
55 Aminen wurde in untersuchten Laktobazillen und Enterokokken nicht beobachtet. Die
56 analysierten Isolate zeigten ein Wachstum auf Medien mit 0,1% und 0,2% Phenol, während 5
57 Isolate Decarboxylase-Aktivität zeigten. Acht Milchstämme, *Lb. curvatus* (IIos6, IIos17 und
58 IIIos1), *Lb. sakei* (IIIos16, Ios12 und IIIos13) und *En. faecium* Ios4 hemmten das Wachstum
59 getesteter Pathogene, einschließlich *Escherichia coli* ATCC 25922, *Staphylococcus aureus*
60 ATCC 25923, *Listeria monocytogenes* ATCC 19115, *Pseudomonas aeruginosa* ATCC 27853
61 und *Bacillus cereus* ATCC 14579.

62 Schlüsselwörter: Milchsäurebakterien (LAB); probiotische Eigenschaften; Simulierte Magen-
63 und Darmsäfte

64

65

66 **Introduction**

67 In the food industry, starter cultures that have probiotic properties are used in a large number of
68 products to preserve quality, sustainability, and organoleptic characteristics. The addition of
69 selected strains of probiotic bacteria to food products, often along with fibers and prebiotics,
70 enhances the beneficial probiotic effects on human health. Probiotic strains should be present in
71 the product in at least 10^6 CFU/g to have an impact on the health of consumers (Kołodziej-
72 Krajewska and Dolatowski, 2012). Foods containing probiotic bacteria fall within the category
73 of functional foods (Pavli et al., 2016). Probiotic bacteria are mainly lactic acid bacteria (LAB),
74 safe organisms that have a beneficial effect on both humans and animals (Gerez et al., 2012). It
75 should be emphasized that the preventive role of probiotics is far more important than the
76 therapeutic one. Lactic acid bacteria (LAB) that has probiotic features is added to fermented
77 milk products, used as a part of the bread-making process, vegetable and fruit juices, and other
78 foods, including fermented meat. Probiotic meat products are obtained by the addition of
79 probiotic microorganisms to fermented meat products. It is known that meat is an excellent
80 medium for the growth of probiotic microorganisms and could be a suitable carrier to support
81 and deliver probiotics to the host (Gänzle et al., 1999; Khan et al., 2011).

82 Research on LAB from meat products as possible sources of probiotic cultures is becoming
83 more and more interesting today, therefore special attention must be paid to autochthonous
84 isolates as potential probiotics. Bacterial strains that can be used in the manufacturing of
85 fermented meat products should be capable of surviving in conditions found within those
86 products as well as dominate other microorganisms found in the finished product (Kołodziej-
87 Krajewska and Dolatowski, 2012). The acid and bile tolerance, resistance to degradation by
88 hydrolytic enzymes, and bile salts in the small intestine are fundamental properties that indicate
89 the ability of probiotic microorganisms to survive the GI (gastrointestinal) tract (Pieniz et al.,
90 2014). The LAB genera that have been identified from meat products include *Lactobacillus*,
91 *Pediococcus*, *Leuconostoc*, *Weissella*, and *Enterococcus* and are well adapted to the ecological
92 niche of meat fermentation (Olaoye and Idowu, 2010). Dry-cured sheep ham is among the
93 traditional fermented food products with autochthonous microbial populations.

94 Dry-cured sheep ham or Sjenica sheep ham is one of the oldest and most popular meat
95 products in Western Serbia. This product is prepared in an exceptionally complex way from
96 mutton carcasses of an autochthonous breed of sheep.

97 An essential part of its production is the sanitary safety of raw materials which must meet all
98 veterinary and sanitary conditions of production. Production of dry-cured sheep ham includes

99 several phases: a selection of raw materials, salting and brining, smoking, drying, and aging,
100 in which the product does not change and receives its trademark smell, consistency, and
101 texture (Stamenković and Dević, 2006).

102 Dry-cured sheep ham fermentation is a long-lasting process caused and helped by
103 autochthonous LAB, which defines the taste of texture as well as the nutritional properties of
104 the product. As the probiotic potential of this product so far has not been explored, the goal of
105 this study was to determine the probiotic characteristics of 16 autochthonous LAB isolates
106 from dry-cured sheep ham through the research of survival rate through the GI tract, the
107 possibility of biogenic amine synthesis, growth on medium with different concentrations of
108 phenol and antimicrobial activity towards pathogens *Escherichia coli* ATCC 25922,
109 *Staphylococcus aureus* ATCC 25923, *Listeria monocytogenes* ATCC 19115, *Pseudomonas*
110 *aeruginosa* ATCC 27853 *Bacillus cereus* ATCC 14579. These observations present a first step
111 towards establishing logical criteria for screening and selecting foodborne microorganisms that
112 exhibit probiotic properties beneficial to humans.

113

114 1. **Materials and methods**

115

116 **Bacterial strains and growth conditions**

117 The research used 16 isolates of LAB isolated from 9 samples of dry-cured sheep ham.
118 Isolation of LAB samples was performed using serial 10-fold dilution technique. 1 gram of
119 meat sample was mixed in 9 mL of quarter strength ringer solution (Hemofarm, Vršac,
120 Serbia) and further serially diluted. Then, 0.1 ml of sample was placed on selective medium
121 de Man Rogosa Sharpe (MRS) agar (Torlak, Belgrade, Serbia). After the incubation period of
122 24-48 hours, initial growth at 37°C was checked. Enumerated *Enterococcus* spp. colonies
123 were screened for growth on modified MRS (6.5% NaCl, 9.6 pH), at 10 and 45°C as well
124 their growth on kanamycin aesculin azide (KAA) agar (Merck, Darmstadt, Germany). The
125 purity of these isolates was confirmed by microscopic observation of homogeneity and their
126 biochemical reactions were evaluated for: arginine hydrolysis, growth ability on MRS agar at
127 different temperatures (15°C and 45°C), growth ability on MRS agar with 4 and 8% NaCl,
128 CO₂ production from glucose, lipolytic activity and proteolytic activity. Gram-positive,
129 catalase-negative, non-spore-forming isolates were considered as LAB and further
130 distinguished by their carbohydrate fermentation profile using an API 50CH and API 20
131 Strep tests, bacterial identification system (Bio Merieux, S.A., France). An analysis of
132 biochemical profiles of isolates was performed to species and subspecies level using

133 identification software (API WEB, V 1.1.0, Biomerieux, Marcy l'Etoile, France). LAB
134 isolates were identified as *Lb. curvatus*, *Lb. sakei*, and *En. faecium*. Identification of isolates
135 was confirmed with MALDI-TOF mass spectrophotometry, using the Bruker Microflex LT
136 instrument (Bruker Daltonics, Bremen, Germany) 34/5000 which is equipped with a nitrogen
137 laser (337 nm) under control of Flexcontrol software ver. 3.1 (Bruker Daltonics). Isolates of
138 LAB were analyzed using the protein extraction method with modifications described in
139 detail in Muruzović et al. (2018). Measurement results for each isolate were expressed by
140 results from MALDI-biotyper (from 0.000 to 3.000), and comparisons were made with
141 similarities to known bacterial profiles available on the MALDI-biotyper software database,
142 with values ≥ 2.000 taken as the correct identification of species levels.

143

144 **Survival in the gastrointestinal (GI) tract**

145 **Tolerance to simulated gastric juice conditions.** Overnight cultures of isolates were
146 inoculated in 1:10 ratio into the solution of 0.5% NaCl and 0.22% pepsin with pH value 2.0
147 and incubated at 37°C for 1 and 2 hours (Radulović et al., 2008). Growth was monitored at
148 OD₆₂₀ (Bassyouni et al., 2012). Isolates that showed tolerance to simulated gastric juice
149 conditions were further subjected to bile salt testing.

150 **Bile salt tolerance.** Overnight cultures of the isolates were inoculated into a solution of
151 0.2% pancrelipase, 0.4% bile salts and 0.5% NaCl with pH 8. Tubes were incubated for 4 and 6
152 h at 37°C (Radulović et al., 2008). Growth was monitored at OD₆₂₀ (Bassyouni et al., 2012).

153

154 **Synthesis of biogenic amines**

155 The ability of isolates to synthesize biogenic amines from histidine and tyrosine has been
156 tested using the method of Do-Won and Jong-Hoon, (2015). Seeded and modified
157 substrates with the addition of amino acids were incubated at 37°C / 24h. The appearance of
158 a purple color in the histidine substrate or the appearance of sediment in the tyrosine
159 substrate confirms the presence of decarboxylase.

160

161 **Growth on medium with different quantity of phenol**

162 The growth capacity of isolates in the presence of phenol was determined by the inoculation
163 of overnight culture onto the MRS agar plates with the addition of 0.1%, 0.2%, and 0.3% of
164 phenol. Physiological levels of phenol compounds in human intestines are low; that is why it
165 is important to analyse the sensitivity of potential probiotics against these substances in
166 smaller concentrations of 0.1, 0.2, and 0.3%. The appearance of colonies after 48 h of

167 incubation at 37°C indicates the growth ability of isolates in the presence of a certain
168 concentration of phenol (Šusković et al., 2001).

169

170 **Antimicrobial activity**

171 Antimicrobial activity of the tested isolates has been examined using the method described in
172 Vesković-Moračanin et al. (2010). Diffusion method with small wells meant that 5 ml of soft
173 (0.7%) nutrient agar (Torlak, Belgrade, Serbia), containing indicator strains, were poured on
174 firm MRS media which was inoculated with 10^5 - 10^6 of cells of indicator culture/ml medium. In
175 the soft agar small wells of 5 mm diameter were formed into which 100 µl of partially purified
176 bacteriocin was poured. Partial purification of bacteriocin was performed in the following way:
177 after 18 h of growth, the cultures were spin-dried at 10000 revolutions for 30 minutes at 4°C.
178 After separation and neutralization of the supernatant with 10 M NaOH, up to pH 6.5 – 7.0, the
179 precipitation of bacteriocin was done by ammonium sulfate (472.2 g/l) until a saturated
180 solution was obtained. Separated bacteriocin in the form of a white deposit was dissolved in
181 25 ml 0,05 M of sodium phosphate buffer with pH 7. The sterilization of partially purified
182 bacteriocin was made by filtration through 0.22 µm microfilter (Acrodisc, Germany).
183 Antimicrobial activity was detected based on the appearance of light zones around small
184 wells as a consequence of growth inhibition in sensitive bacteria strains.

185

186 **Statistical analysis**

187 The results represent the mean ± standards deviations. Statistical analysis was conducted with
188 SPSS 11.0 Bivariate Correlation Analysis (Chicago, Illinois, USA).

189 **2. Results and Discussion**

190 The indigenous microbiota of traditional dry-cured meat products today represents a
191 significant field of research on wild strains and the product itself as functional food (Arihara,
192 2006). Our study results of dry-cured sheep ham microbiota indicated the dominance of LAB
193 strains with high probiotic potential. The results presented in Žugić-Petrović et al., (2020),
194 indicated that the number of viable LAB was in range from 2.6×10^2 to 9.2×10^3 CFU/g of
195 dry-cured sheep ham. According to the pattern of sugar fermentation, these isolates were
196 identified as *Lb. curvatus* (9 strains), *Lb. sakei* (3 strains) and *En. faecium* (4 strains) (Table
197 1). The MALDI-TOF application in the study confirmed the preliminary API identification of
198 the strains. Almost all LAB strains had a high level of identification, biotyper database and
199 software were 100% correct in assigning the isolates to species. Four enterococci were

200 identified at a level of highly probable species identification (score values ≥ 2.4), and 12
 201 isolates (*Lb. curvatus* and *Lb. sakei* strains) were identified at a level of secure genus and
 202 species identification (score values between 2.25 - 2.28). By exploring dry-cured meat
 203 products from Eastern Himalayas, Rai et al. (2010) came up with identification results
 204 showing the dominance of *Lb. curvatus* and *Lb. sakei* in the studied samples. The results of
 205 the probiotic potential of identified *En. faecium* isolates from dried Tunisian meat "Dried
 206 Ossban" were presented by Zommiti et al. (2018). They indicated the safety aspect of the
 207 strains themselves. Our study showed that MALDI-TOF can be an effective, and sustainable
 208 method in classifying *Lactobacillus* and *Enterococcus* strains.

209 **TABLE 1.** *Isolated species of LAB from the sheep ham*

MALDI-TOF Identification	<i>Lb. curvaus</i> (9)	<i>En. faecium</i> (4)	<i>Lb. sakei</i> (3)
Growth at:			
15°C	+	+	+
45°C	-	+	-
Growth in:			
4.0% NaCl	+	+	+
8.0% NaCl	+	+	+
Gas from glucose	-	-	-
NH ₃ from arginine	-	+	-
Lipolytic activity	-	-	-
Proteolytic activity	-	-	-
Biochemical (API)			
L-arabinose	-	+	-
Cellobiose	+	+	+
Ribose	+	+	+
Esculin	+	+	+
Galactose	+	+	+
Lactose	+	+	-
D-mannose	+	+	+
Melezitose	-	-	-
Melibiose	-	+	+
D-raffinose	-	+	-
Sucrose	+	+	+
Trehalose	-	+	+
D-xylose	-	+	-
Rhamnose	-	-	-
Mannitol	-	+	-
Maltose	+	+	+
Sorbitol	-	-	-
Salicin	+	+	+

210 "+" - positive reaction; "-" - negative reaction

211

212 **Survival in simulated gastrointestinal tract conditions**

213 Falagas et al. (2006) point out that probiotics are living microorganisms which when
 214 administered in adequate amounts confer a health benefit on the host. One of the necessary
 215 conditions that manifest their probiotic characteristics is the ability of survival through the GI
 216 tract. Gastrointestinal tract possesses very harsh conditions for the survival of
 217 microorganisms which need to remain viable in population levels of 10^6 - 10^7 CFU/g of
 218 product in order to deliver the health benefits (Pavli et al., 2016). The tested strains isolated
 219 from dry-cured sheep ham from western Balkans showed a good survival rate in the artificial
 220 gastric and bile juice conditions, and the results are presented in Table 2.

221

222 **TABLE 2.** *The survival rate of selected strains in simulated gastric and bile juice conditions*

Isolates	Initial cell	Tolerance to gastric juice conditions		Tolerance to bile juice conditions	
	Incubation time 0 h	Incubation time 1 h	Incubation time 2 h	Incubation time 4 h	Incubation time 6 h
<i>Lb. curvatus</i> IIos17	1.1 ± 0.07	0.7 ± 0.00*	0.7 ± 0.00*	0.7 ± 0.00*	0.7 ± 0.04*
<i>Lb. curvatus</i> IIos11	1.0 ± 0.00	0.8 ± 0.06*	0.8 ± 0.00*	0.7 ± 0.02*	0.7 ± 0.05*
<i>Lb. curvatus</i> IIos4	1.0 ± 0.05	0.7 ± 0.00*	0.7 ± 0.05*	0.8 ± 0.00*	1.0 ± 0.04
<i>Lb. curvatus</i> IIos3	0.9 ± 0.00	0.5 ± 0.01*	0.5 ± 0.03*	0.5 ± 0.04*	0.6 ± 0.02*
<i>Lb. curvatus</i> IIos19	1.0 ± 0.06	1.0 ± 0.02	0.9 ± 0.01*	0.8 ± 0.03*	0.5 ± 0.02*
<i>Lb. sakei</i> IIIos16	0.8 ± 0.05	0.5 ± 0.00*	0.5 ± 0.01*	0.6 ± 0.00*	0.6 ± 0.01*
<i>Lb. curvatus</i> IIos18	1.0 ± 0.00	0.8 ± 0.17*	0.9 ± 0.00*	0.9 ± 0.00*	0.6 ± 0.00*
<i>En. faecium</i> Ios4	1.0 ± 0.02	0.9 ± 0.01*	0.9 ± 0.03*	0.9 ± 0.02*	0.8 ± 0.04*
<i>Lb. curvatus</i> IIos6	1.0 ± 0.06	0.8 ± 0.02*	0.8 ± 0.00*	0.8 ± 0.04*	0.8 ± 0.00*
<i>En. faecium</i> Ios5a	1.1 ± 0.00	1.0 ± 0.04*	0.9 ± 0.05*	1.0 ± 0.07*	0.9 ± 0.02*
<i>Lb. curvatus</i> IIos17a	1.0 ± 0.05	0.9 ± 0.04*	0.8 ± 0.09*	0.8 ± 0.13*	0.8 ± 0.12*
<i>Lb. sakei</i> Ios12	1.0 ± 0.05	0.7 ± 0.02*	0.8 ± 0.03*	0.8 ± 0.09*	1.0 ± 0.04
<i>En. faecium</i> Ios1a	1.0 ± 0.05	0.6 ± 0.04*	0.6 ± 0.01*	0.6 ± 0.09*	0.8 ± 0.01*
<i>Lb. curvatus</i> IIIos1	1.0 ± 0.00	0.9 ± 0.01*	0.9 ± 0.29*	0.9 ± 0.14*	0.8 ± 0.03*
<i>En. faecium</i> IIos24	0.9 ± 0.02	0.6 ± 0.01*	0.7 ± 0.09*	0.6 ± 0.16*	0.3 ± 0.11*
<i>Lb. sakei</i> IIIos13	1.0 ± 0.00	0.9 ± 0.02*	0.8 ± 0.21*	1.0 ± 0.03*	1.0 ± 0.28*

223 OD at 620nm at different time interval (hour); Values marked with asterisks are not significantly different from the control group (0 h),
 224 according to the Duncan's test (p<0.05).

225

226

227 As shown in Table 2, in simulated gastric juice conditions, the number of cells after the first
228 hour of incubation decreased in all of the tested *Lb. curvatus* and *Lb. sakei* isolates except in
229 isolate *Lb. curvatus* Ios19 where the number of cells remained approximately at the initial
230 level. After the second hour of incubation, the number of cells for the studied *Lb. curvatus*
231 and *Lb. sakei* did not decrease compared to the first hour in simulated gastric juice conditions
232 while the number of cells in isolates *Lb. curvatus* Ios18 and *Lb. sakei* Ios12 showed a slight
233 increase after the second hour of incubation. The obtained results are in accordance with the
234 work of Bacha et al. (2009) on the probiotic characteristics of lactobacilli isolated from beef
235 sausage, which showed a high rate of survival in simulated stomach conditions. The studied
236 enterococci isolates showed a decrease in the number of cells in the first hours of incubation
237 relative to their initial number in simulated gastric juice conditions. In the second hour of
238 incubation, the number of cells in the examined *En. faecium* isolates remained approximately
239 the same as in the first hour, except for *En. faecium* Ios24 in which the number of cells
240 increased slightly.

241 There is no statistically significant decrease in the number of cells in the first and second
242 hours of incubation. Hosseini et al. (2009) point out that the survival rate of *En. faecium*
243 increases with increasing pH of the environment. In simulated bile juice conditions, isolates of
244 *Lb. curvatus* mainly showed a mild decrease in the number of cells (isolates: Ios11, Ios19,
245 Ios18, Ios17, Ios6, and Ios1) or maintained approximately the same number of cells
246 (isolate Ios17), except in the case of *Lb. curvatus* Ios4 which showed an increase in the
247 number of cells after 6 h of incubation but without significant statistical differences. *Lb. sakei*
248 isolates showed the same results in relation to the number of cells under gastric juice
249 conditions. Maragkoudakis et al. (2006), researching the probiotic potential of Lactobacillus
250 strains isolated from dairy products, concluded that all strains were resistant to pancreatin, as
251 even after 4 h of exposure they retained viability. The number of viable cells of enterococci in
252 simulated bile juice conditions after incubation for 4 and 6 h decline, in all the tested isolates.
253 However, *En. faecium* isolate Ios1a showed an increase in the number of cells, but this was
254 not statistically significant. According to Ruiz-Moyano et al. (2009), *En. faecium* isolate
255 SE906 from Iberian dry-fermented sausages has a good survival capability in the simulated GI
256 tract conditions, which characterizes it as a good potential probiotic.

257

258 **Synthesis of biogenic amines**

259 Biogenic amines are organic compounds that may be created in the meat processing and
260 fermentation. They can cause headaches, circulatory disorders, and intoxication (Virgill et al.,

261 2007). Synthesis of biogenic amines test results on dry-cured sheep ham isolates showed that
262 in the case of histidine there is no appearance of synthesis of biogenic amines in both
263 lactobacilli and enterococci isolates. In the case of a tyrosine substrate, synthesis of biogenic
264 amines in investigated lactobacilli and enterococci was not detected, which is in disagreement
265 with the results presented by Landeta et al. (2013) that indicate that most *En .faecium* and *Lb.*
266 *sakei* strains showed the production of tyramine. Similar results of tyramine production in
267 enterococci were reported by Muñoz-Atienza et al. (2011).

268

269 **Growth on medium with different quantity of phenol**

270 According to Šušković et al. (2001), phenols can be formed in the intestines as a product of
271 bacterial deamination of some aromatic amino acids derived from foods or endogenously
272 produced. Phenols have distinct bacteriostatic properties, so phenol-tolerant bacteria have a
273 greater chance of surviving the conditions of the GI tract than those bacteria that are more
274 susceptible to this compound. As physiological levels of phenols in the human organism are
275 low, it is important to analyze the sensitivity of potential probiotics to this substance precisely
276 at concentrations in which phenols can be expected in the human body, in the range of 0.1%,
277 0.2%, and 0.3%. The results of research on the growth medium with different quantity of
278 phenol showed good growth of investigated strains in substrates with a phenolic concentration
279 of 0.1%, 0.2%, and 0.3%. Isolate *Lb. curvatus* IIIos1 showed no growth in the substrate with
280 0.3% phenol, while isolates *Lb. curvatus* IIos11 and *Lb. curvatus* IIIos1 showed no growth at
281 any tested phenol concentration. The results obtained by Aswathy et al. (2008) indicated that
282 many investigated strains of enterococci, lactobacilli, and leuconostoc have successfully
283 tolerated a low level of phenol of 0.2-0.3%, which qualified them as possible probiotics.
284 Vizoso Pinto et al. (2006) presented survival results on the phenol resistance of *Lb. plantarum*
285 strains which suggest that these are generally moderately tolerant to a phenol concentration of
286 about 0.4%.

287

288 **Antimicrobial activity**

289 Probiotic bacteria can antagonize pathogens using several mechanisms that involve the
290 production of antimicrobial compounds such as fatty acids, organic acids, hydrogen peroxide,
291 diacetyl, acetone, bacteriocins, as well as competition for the substrate and co-aggregation
292 with a pathogen (Todorov et al., 2011).

293

294

295 **TABLE 3.** Antimicrobial activity of the isolated LAB against pathogenic bacteria

Isolates	Indicator strains				
	<i>Escherichia coli</i> ATCC 25922	<i>Staphylococcus aureus</i> ATCC 25923	<i>Listeria monocytogenes</i> ATCC 19115	<i>Pseudomonas aeruginosa</i> ATCC 27853	<i>Bacillus cereus</i> ATCC 14579
<i>Lb. curvatus</i> IIos17	10.2 ± 0.64*	12.0 ± 1.70	10.3 ± 1.10	10.0 ± 1.10	10.0 ± 1.70
<i>Lb. curvatus</i> IIos11	10.0 ± 1.90	-	8.1 ± 0.20	-	-
<i>Lb. curvatus</i> IIos4	25.6 ± 0.20	20.0 ± 1.80	18.0 ± 0.50	-	-
<i>Lb. curvatus</i> IIos3	16.1 ± 1.70	12.3 ± 1.50	-	10.0 ± 2.10	10.0 ± 0.10
<i>Lb. curvatus</i> IIos19	12.0 ± 1.00	10.0 ± 2.00	12.0 ± 0.50	-	-
<i>Lb. sakei</i> IIIos16	20.0 ± 1.70	18.3 ± 0.57	18.0 ± 1.70	20.0 ± 0.80	20.0 ± 0.30
<i>Lb. curvatus</i> IIos18	-	16.0 ± 4.10	15.0 ± 1.50	-	16.0 ± 0.50
<i>E. faecium</i> Ios4	23.0 ± 1.70	23.0 ± 1.70	25.0 ± 0.00	20.0 ± 3.00	18.0 ± 0.20
<i>Lb. curvatus</i> IIos6	25.6 ± 0.50	23.3 ± 0.50	25.0 ± 0.90	23.0 ± 0.00	22.0 ± 0.60
<i>E. faecium</i> Ios5a	20.0 ± 1.73	18.0 ± 0.00	20.0 ± 0.00	-	14.0 ± 0.80
<i>Lb. curvatus</i> IIos17a	10.0 ± 2.78	12.3 ± 0.50	10.0 ± 2.60	10.0 ± 0.00	-
<i>Lb. sakei</i> Ios12	22.0 ± 2.90	19.5 ± 2.10	20.0 ± 1.80	15.0 ± 3.20	20.0 ± 1.30
<i>E. faecium</i> Ios1a	10.0 ± 4.50	8.0 ± 1.70	-	-	8.0 ± 2.00
<i>Lb. curvatus</i> IIIos1	15.0 ± 2.17	13.1 ± 0.20	10.0 ± 0.80	10.0 ± 0.50	12.0 ± 1.70
<i>E. faecium</i> IIos24	-	14.0 ± 0.10	10.0 ± 3.00	-	10.0 ± 0.05
<i>Lb. sakei</i> IIIos13	15.0 ± 1.90	13.0 ± 0.10	14.0 ± 0.00	15.0 ± 2.20	14.0 ± 0.00

296 *Diameter of inhibitory zone (mm)

297 As shown in Table 3. *Lb. curvatus* IIos6, *En. faecium* Ios4, *Lb. sakei* IIIos16, *Lb. sakei* Ios12,
 298 *Lb. curvatus* IIos17, *Lb. curvatus* IIIos1 and *Lb. sakei* IIIos13 are isolates that showed an
 299 inhibition zone to all tested pathogens. All the isolates showed the largest inhibition zone
 300 towards *E. coli* ATCC 25922 where in the mean values of the inhibition zone ranged from
 301 25.6 to 10.0. Similar results were also obtained by Brink et al. (2006), who investigated the
 302 probiotic potential of LAB, wherein the isolates showed a strong inhibition related to *E. coli*.
 303 In the case of *L. monocytogenes* ATCC 19115, potential probiotics showed good antimicrobial
 304 properties, wherein the isolate *Lb. curvatus* IIos6 also had an inhibition zone of about 25 ± 0.9.
 305 Benito et al. (2007) in their study, highlighted the antimicrobial activity of LAB strains from
 306 Iberian dry-fermented sausages against *L. monocytogenes*. *Lb. curvatus* IIos11 had the lower
 307 antimicrobial activity, with no inhibition zone to *S. aureus*, *P. aeruginosa* and *B. cereus*. *P.*
 308 *aeruginosa* ATCC 27853 is a pathogen for which the highest number of investigated strains of
 309 LAB (43.75%) did not show an inhibition zone. Antagonistic effects against *B. cereus* had not
 310 been shown by 25% of isolates. Žugić-Petrović et al. (2020) indicated that LAB showed high

311 or moderate sensitivity to clinically relevant antibiotics (tetracycline, cephalexin, amoxicillin,
312 ceftriaxone, and erythromycin).

313 3. **Conclusion**

314 The emerging demand that traditional foods should have health benefits beyond nutritional
315 ones has provided ample opportunity to explore unexplored foods for isolation of lactic acid
316 bacteria and their potential role as probiotics. The results presented in this study demonstrated
317 that isolates of LAB from dry-cured sheep ham exhibited favorable probiotic characteristics
318 as the ability to grow and survive through the GI tract, the possibility of synthesis of biogenic
319 amines, growth on medium with different quantity of phenol and antimicrobial activity. Further
320 studies if investigated strains need to evaluate their potential health benefits, and their
321 performance as novel probiotic or starters cultures.

322 **Conflict of interest**

323 The authors declare that no conflict of interest among authors.

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