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7
8

9 **Abstract**

10 In recent years, biocontrol of foodborne pathogens has become a concern in the food industry,
11 owing to safety issues. *Listeria monocytogenes* is one of the foodborne pathogens that causes
12 listeriosis. The major concern in the control of *L. monocytogenes* is its viability as it can survive
13 in a wide range of environments. The purpose of this study was to isolate lactic acid bacteria
14 with antimicrobial activity, evaluate their applicability as a cheese starter, and evaluate their
15 inhibitory effects on *L. monocytogenes*.

16 *Lactococcus lactis* strain with antibacterial activity was isolated from raw milk. The isolated
17 strain was a low acidifier, making it a suitable candidate as an adjunct starter culture. The
18 commercial starter culture TCC-3 was used as a primary starter in this study. Fresh cheese was
19 produced using TCC-3 and *La. lactis* CAU2013 at a laboratory scale. Growth of *L.*
20 *monocytogenes* (5 log CFU/g) in the cheese inoculated with it was monitored during the storage
21 at 4°C and 10°C for 5 days. The count of *L. monocytogenes* was 1 log unit lower in the cheese
22 produced using the lactic acid bacteria strain compared to that in the cheese produced using the
23 commercial starter.

24 The use of bacteriocin-producing lactic acid bacteria as a starter culture efficiently inhibited
25 the growth of *L. monocytogenes*. Therefore, *La. lactis* can be used as a protective adjunct starter
26 culture for cheese production and can improve the safety of the product leading to an increase
27 in its shelf-life.

28 **Keywords:** *Lactococcus lactis*, bacteriocin, *Listeria monocytogenes*, cheese starter culture,
29 foodborne pathogen

30 **Introduction**

31 Listeriosis is a foodborne disease caused by *Listeria monocytogenes*. It can lead to sepsis,
32 meningitis, encephalitis, and even death (de Noordhout *et al.*, 2014). Despite its low incidence
33 compared with that of other foodborne illnesses, listeriosis is one of the major issues in the food
34 industry because of its high fatality rate. *L. monocytogenes* is found in dairy products,
35 particularly in ready-to-eat cheese products. As *L. monocytogenes* survives in various
36 environments such as those with a wide range of temperature (0–45°C) and pH (4.1–9.6), it
37 can contaminate cheese at several stages of production; therefore, its growth is difficult to
38 control (Lungu *et al.*, 2008; Melo *et al.*, 2014).

39 Several methods have been used to control the growth of *L. monocytogenes* in cheese,
40 including using bacteriocin or bacteriocin-producing lactic acid bacteria (LAB). Bacteriocins
41 are peptides or proteins, ribosomally synthesized by bacteria, which have antimicrobial ability
42 against closely related species. The application of bacteriocin-producing bacteria is
43 advantageous as they are stable, cost-effective, and safe. Anti-listerial activity of LAB in cheese
44 have also been reported (Coelho *et al.*, 2014; Dal Bello *et al.*, 2012; Kondrotiene *et al.*, 2018).

45 In the present study, we aimed to determine the effects of bacteriocin-producing LAB isolated
46 from raw milk on the growth of *L. monocytogenes* in milk broth and cheese.

47

48 **Materials and Methods**

49 **Isolation of bacteriocin-producing lactic acid bacteria**

50 Potential bacteriocin-producing LAB were isolated from raw bovine milk, obtained from a
51 Chung-Ang University-affiliated farm (Anseong, Republic of Korea). The sample was serially
52 diluted ten-fold and plated on MRS agar (BD Difco, USA). The plates were incubated at 37°C
53 for 24–48 h, and a total of 90 well-isolated colonies were collected. Each colony was inoculated
54 into MRS broth for 24 h at 37°C.

55 To screen for antimicrobial activity, the cell-free supernatant (CFS) was obtained after
56 neutralization with 1N NaOH, centrifugation at 13,000 rpm for 10 min at 4°C, and filtered
57 through 0.45 µm filters to remove bacterial cells. Then, each supernatant was spotted on the
58 tryptic soy agar (TSA, BD Difco, USA) plate inoculated with a lawn of *Listeria monocytogenes*
59 ATCC 19115 as an indicator strain. The plates were incubated at 30°C for 12 h, and antibacterial
60 activity was confirmed with the presence of inhibition zone. The strains with antibacterial
61 activity were routinely cultured in MRS broth at 37°C overnight and were preserved in 10%
62 skim milk supplemented with 25% (v/v) glycerol, stored at -80°C for further use.

63

64 **Identification of bacteriocin-producing strain**

65 The bacteriocin-producing strains were identified by Gram staining, carbohydrate
66 fermentation profile (analytical profile index (API) test), and 16S rRNA gene sequencing
67 analysis. Gram staining and API analysis was performed using a Gram-stain kit (BD Difco,
68 USA) and API 50 CHL kit (Biomérieux, France), respectively, according to the manufacturer's
69 instructions.

70 For 16S rRNA analysis, the genomic DNA was extracted using QIAamp PowerFecal DNA
71 Kit (Qiagen, Germany) and amplified using 2X H-star Taq PCR Master Mix (BioFACT,
72 Republic of Korea). Polymerase chain reaction (PCR) was performed using the universal
73 bacterial primers 27F (5'-AGAGTTTGATCMTGG CTCAG-3'), 1492R (5'-
74 TACGGYTACCTTGTTACGACTT-3'), 785F (5'-GGATTAGA TACCCTGGTA-3'), and 805R
75 (5'-GACTACCAGGGTATCTAATC-3'). The PCR products were purified using a PCR
76 purification kit (Qiagen, Germany) and sequenced by SolGent Co. Ltd. (Daejeon, Republic of
77 Korea).

78 The analyzed sequences were confirmed using the EzTaxon-e server (www.ezbiocloud.net/)
79 (Kim *et al.*, 2012) and NCBI GenBank database using the Basic Local Alignment Search Tool
80 (BLAST) algorithm (<https://blast.ncbi.nlm.nih.gov/Blast.cgi>) (Altschul *et al.*, 1990).

81

82 **Antibacterial activity of bacteriocin**

83 Bacteriocin activity was assessed using a spot-on-lawn method as described previously
84 (Phumisantiphong *et al.*, 2017) with minor modifications. Briefly, each indicator strain was
85 inoculated to 4 mL of molten TSA and overlaid on the base TSA plate. After solidification, 20
86 μ L of the neutralized CFS of LAB strains was spotted onto the indicator lawn. After incubation
87 at 30°C for 12 h, a clear inhibition zone was observed. The foodborne pathogens used as
88 indicator strains were cultured in tryptic soy broth (TSB, BD Difco, USA) at 37°C overnight
89 before use. The experiment was conducted in triplicates.

90

91 **Evaluation of acid production**

92 The acid production of the *La. lactis* CAU2013 was evaluated and compared with that of the
93 commercial starter TCC-3 (Chr. Hansen, Denmark), which consisted of *Lactobacillus*

94 *delbrueckii* subsp. *bulgaricus* and *Streptococcus thermophilus*. The cultures were grown in
95 MRS broth at 37°C overnight. Then, individual cultures and mixture of *La. lactis* CAU2013
96 and TCC-3 (1:1 ratio) were inoculated in 10% skim milk broth (Harrington and Hill, 1991) and
97 whole milk. The pH and titratable acidity (TA) were measured every three hours for 12 h while
98 incubating at 30°C. To determine TA, 0.1% phenolphthalein was used as an indicator, and 0.1
99 N sodium hydroxide (NaOH) for titration.

100

101 **Anti-listerial activity of strain CAU2013 as an adjunct starter in milk**

102 To determine the anti-listerial properties of strain CAU2013 when used as an adjunct starter
103 in milk, 10% skim milk broth and whole milk media were inoculated with an overnight culture
104 of CAU2013 and 1:1 ratio of CAU2013 and TCC-3 starter (final concentration of 7 log
105 CFU/mL). Additionally, the overnight culture of *L. monocytogenes* ATCC 19115 was
106 inoculated to each setup (final concentration of 5 log CFU/mL). The inoculated milk media
107 were incubated at 30°C for 12 h. The viable cell count of *L. monocytogenes* was determined
108 every three hours. Samples were diluted serially in ten-fold increments using 1×phosphate-
109 buffered saline (PBS, pH7.5) and plated on Oxford agar (BD Difco, USA).

110 **Manufacture of laboratory-scale cheese**

111 The lab-scale cheese was manufactured following the methods of Mills *et al.*(2011) with
112 some modifications. TCC-3 was used as the primary starter and *La. lactis* CAU2013 as an
113 adjunct culture. The starter cultures were initially grown in MRS broth at 37°C for 24 h before
114 inoculation into 10% skim milk broth and incubated for 18 h at 37°C before use. Additionally,
115 *L. monocytogenes* ATCC19115 was cultured in TSB for 18 h at 37°C before use.

116 Milk (400 mL) (Seoul milk, Republic of Korea) was heated to 31°C before the inoculation of
117 starter culture. The starter cultures were inoculated as follows: TCC-3 and *La. lactis* CAU2013

118 and TCC-3 (1:1 ratio), both at a final concentration of 7 log CFU/mL. Subsequently, 0.01% *L.*
119 *monocytogenes* at a level of 5 log CFU/mL was inoculated into both treatments. After 30 min,
120 0.2 g/L of rennet was added, and the mixture was stirred for 2 min. Once coagulum formed
121 firmly, the curd was cut into cubes, and the mixture was stirred for 10 min. Then, the mixture
122 was heated to 36°C for 10 min and stirred for 20 min. The whey was drained off, and curd was
123 distributed into the sterile dish. The samples were stored at 4°C and 10°C for 5 days. The
124 procedure of cheese production is illustrated in Figure 1.

125

126 **Microbial analysis of laboratory-scale fresh cheese**

127 The viable cell counts of LAB and *L. monocytogenes* in the lab-produced cheese were
128 determined in duplicate every day during storage at 4°C and 10°C. For microbial analysis, 1 g
129 of cheese was homogenized in 9 mL of PBS buffer and were serially diluted ten-fold in the
130 same buffer and plated on the appropriate agar plate. The LABs were enumerated on MRS agar
131 after incubation at 37°C for 3 days, and *L. monocytogenes* on Oxford agar after incubation at
132 37°C for 24 h. All of the experiments were conducted in triplicates.

133 **Results and discussion**

134 **Isolation and identification of bacteriocin-producing strains**

135 Among the 90 colonies isolated from raw milk, one isolate exhibited antibacterial activity
136 against *L. monocytogenes*. The strain CAU2013 was characterized as a gram-positive, coccus-
137 shaped bacterium. The biochemical characteristics determined using the API 50 CHL kit are
138 described in Table 1. 16S rRNA gene sequence analysis revealed that strain CAU2013 is most
139 likely a strain of *Lactococcus lactis* (Table 2), which commonly produce nisin (Shin *et al.*,
140 2016). Neighbor-joining (NJ) phylogenetic tree of the strain CAU 2013 and related type strains
141 based on 16S rRNA gene sequences also clearly show that this strain belongs to *Lactococcus*
142 *lactis* (Supplementary Figure 1). *La. lactis* strains are historically used in the fermentation and
143 preservation of food and are generally recognized as safe (GRAS) (Cook *et al.*, 2018).
144 Therefore, *La. lactis* CAU2013 was selected for downstream applications in the study.

145 *L. monocytogenes* ATCC 19115 was used as an indicator strain for all experiments because it
146 belongs to the serotype 4b, which causes most cases of listeriosis.

147

148 **Antibacterial activity of bacteriocin**

149 The bacteriocin produced by *La. lactis* CAU2013 had antibacterial activity against all *Listeria*
150 strains as well as *Staphylococcus aureus*, which are common foodborne pathogens (Yoon,
151 2020). However, no antibacterial activity was observed against other gram-positive foodborne
152 pathogens, such as *Salmonella enteritidis* and *Escherichia coli* (Table 3). Generally, nisin is
153 highly effective against gram-positive bacteria by binding to lipid II, which leads to the
154 inhibition of cell wall biosynthesis or pore formation in the membrane. However, nisin cannot

155 bind to its target lipid II in gram-negative bacteria, because of the presence of the outer
156 membrane (Li *et al.*, 2018).

157

158 **Characterization of acid production**

159 The changes in pH and TA values in 10% skim milk broth and in whole milk are presented
160 in Figure 2. *La. lactis* CAU2013 reduced the pH of skim milk broth from 6.41 to 5.77 and that
161 of whole milk from 6.65 to 6.20. Additionally, TA value increased to 0.25 in both broths. Ayad
162 *et al.* (2004) described fast, medium, or slow-acidifying strains as $\Delta\text{pH}(=\text{pH}_{\text{at time}}-\text{pH}_{\text{zero time}})$
163 of 0.4 U achieved after 3 h, 3–5 h, and > 5 h, respectively. Also, Raquib *et al.* (2003) classified
164 strains with titratable acidity as low, moderate, or fast when the TA values were < 0.5, between
165 0.5 and 0.6, and > 0.6, respectively. Therefore, *La. lactis* CAU2013 can be classified as a low
166 acidifier strain. This result is consistent with other studies that reported poor acid production
167 from *La. lactis* strains (Ayad *et al.*, 2004; Coelho *et al.*, 2014).

168 The pH values measured corresponded with the calculated TA and were generally similar for
169 skim milk broth and whole milk. The mixed starter, consisting of TCC-3 and *La. lactis*
170 CAU2013, accelerated the acidification in milk. Nevertheless, bacteriocin-producing strains
171 delay acidification (Garde *et al.*, 1997); however, the strain CAU2013 did not show similar
172 properties. The accelerated acidification might be because of the interaction between the strains;
173 however, the underlying mechanisms need further research. Ávila *et al.* (2005) observed that
174 enterocin-producing adjunct starter enterococci enhanced milk acidification, which may be
175 stimulated by the low-molecular-weight nitrogen compounds produced by primary starter,
176 *Lactobacillus helveticus* LH92.

177 The rapid decline in pH during the initial stage of cheese production is crucial for curd

178 formation and prevention of the growth of undesirable microorganisms. Therefore, the fast-
179 acidifying strains can be used as primary starters, while the slow-acidifying bacteria can be used
180 as adjunct starters. As the strain CAU2013 has antibacterial property but has low acid
181 production ability, it is better suited as an adjunct starter culture.

182

183 **Anti-listerial activity of strain CAU2013 as an adjunct starter in milk**

184 The growth of *L. monocytogenes* was monitored in skim milk broth and whole milk during
185 incubation at 30°C. In skim milk broth with *La. lactis* CAU2013, the concentration of *L.*
186 *monocytogenes* count was reduced by 3 log units more compared with that of other samples
187 after 3 h and not detected following 6 h of fermentation (Figure 3a). In the whole milk with the
188 strain CAU2013, *L. monocytogenes* count was reduced by 0.5 log unit after 6 h, and 1 log unit
189 after 9 h compared with that of other samples (Figure 3b).

190 The results support the findings from several studies that reported that the addition of
191 bacteriocin affects the biocontrol of spoilage bacteria. Muñoz *et al.* (2007) investigated that *E.*
192 *faecalis*-produced enterocin in milk and found that it could control the growth of
193 *Staphylococcus aureus*. In addition, according to Arqués *et al.* (2011), the addition of nisin in
194 milk decreased *L. monocytogenes* count by 3 log units after 4 h.

195 The efficiency of the combined starter cultures in the inhibition of *L. monocytogenes* was
196 lower in whole milk than in skim milk. The difference in the composition between the two milk
197 media could be a factor responsible for the difference. In addition, Muñoz *et al.* (2007) stated
198 that low effectiveness in foods could be attributed to higher retention of the bacteriocin
199 molecules by milk components, resulting in slower diffusion. However, in both cases, inhibition
200 of *L. monocytogenes* growth was observed. The results suggest the potential application of *La.*
201 *lactis* CAU2013 in various food systems to control *L. monocytogenes* growth.

202

203 **Inhibition of *L. monocytogenes* in laboratory-scale fresh cheese**

204 The cell count of the starter cultures was determined during the storage at 4°C and 10°C
205 (Figure 4). In both the cases, LAB reached a final concentration of 9 log CFU/g during cheese
206 manufacture.

207 During the storage at 4°C, the cheese treated with TCC-3 starter culture maintained *L.*
208 *monocytogenes* count at 7.5 to 7.7 log CFU/g. In contrast, the cheese treated with TCC-3 and
209 *La. lactis* CAU2013 had less *L. monocytogenes* count, approximately 0.5 log unit at 0 h and 1
210 log unit after 5 days with a final concentration of 6.4 log CFU/g. Besides, during the storage at
211 10°C, the cheese treated with TCC-3 starter culture maintained the bacterial count between 6.86
212 and 7.31 log CFU/g (Figure 4a). within contrast, the cheese treated with TCC-3 and CAU2013
213 had less *L. monocytogenes* count, approximately 1 log unit at 0 h and 1.5 log unit after 5 days,
214 with a final concentration of 5.76 log CFU/g (Figure 4b). This result is consistent with a study
215 that reported that 2 log unit reduction was observed in cheese with *La. lactis* strain (Coelho *et*
216 *al.*, 2014). Moreover, Kondrotiene *et al.* (2018) showed that nisin-producing *La. lactis* strains
217 decreased the growth of *L. monocytogenes* in fresh cheese during 7 days of storage at 4°C.

218 Therefore, the results support that manufacturing cheese using a bacteriocin-producing starter
219 reinforced the inhibition of growth of *L. monocytogenes*, and it would be effective in controlling
220 contamination during cheese production. Additionally, after storage at temperatures of 4°C and
221 10°C, *L. monocytogenes* count was reduced, which may confirm the potential of LAB in
222 controlling the growth of *L. monocytogenes* during storage at refrigeration temperature.

223

224 **Conflicts of interest**

225 The authors declare no potential conflicts of interest.

226

227 **Author contributions**

228 Conceptualization: Kim GB. Data curation: Yoon SH, Kim GB. Investigation: Yoon SH, Kim
229 GB. Writing - original draft: Yoon SH. Writing - review & editing: Yoon SH, Kim GB.

230

231 **Ethics Approval**

232 This article does not require IRB/IACUC approval because there are no human and animal
233 participants.

234

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293

294 **Table 1.** Carbohydrate fermentation patterns of the two isolated bacteriocin-producing lactic
 295 acid bacteria. The test was performed with API 50 CHL kit and all data are from this study. +,
 296 positive; -, negative

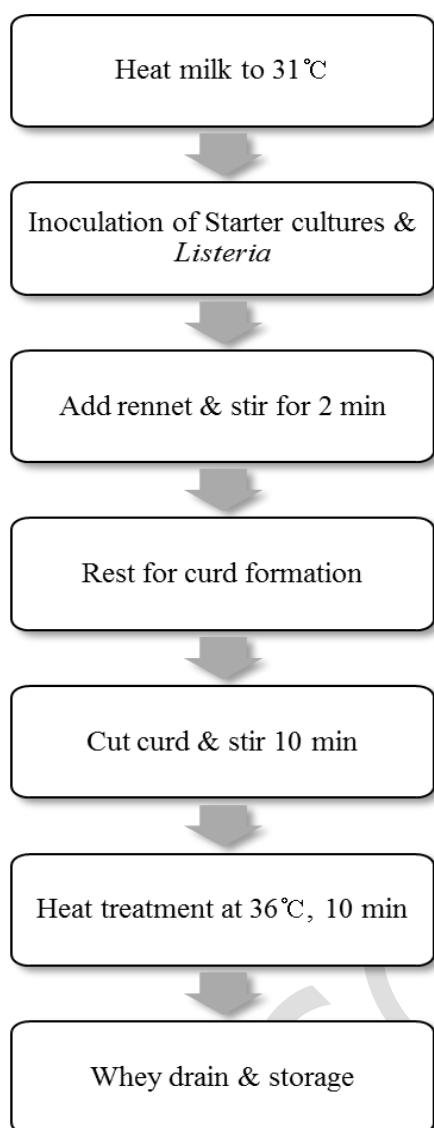
| Carbohydrates | | Carbohydrates | |
|--------------------------------|---|---------------------------|---|
| Glycerol | - | Salicin | + |
| Erythritol | - | D-cellobiose | + |
| D-arabinose. | - | D-maltose | + |
| L-arabinose | + | D-lactose (bovine origin) | + |
| D-ribose | + | D-melibiose | - |
| D-xylose | + | D-saccharose (sucrose) | + |
| L-xylose | - | D-trehalose | + |
| D-xylose | - | Inulin | - |
| Methyl-beta-D-xylopyranoside | - | D-melezitose | + |
| D-galactose | + | D-raffinose | - |
| D-glucose | + | Amidon (starch) | + |
| D-fructose | + | Glycogen | - |
| D-mannose | + | Xylitol | - |
| L-sorbose | - | Gentiobiose | + |
| L-rhamnose | - | D-turanose | - |
| Dulcitol | - | D-lyxose | - |
| Inositol | - | D-tagatose | + |
| D-mannitol | + | D-fucose | - |
| D-sorbitol | - | L-fucose | - |
| Methyl-alpha-D-mannopyranoside | - | D-arabitol | - |
| Methyl-alpha-D-glucopyranoside | - | L-arabitol | - |
| N-acetylglucosamine | + | Potassium gluconate | - |
| Amygdalin | + | Potassium 2-ketogluconate | - |
| Arbutin | + | Potassium 5-ketogluconate | - |
| Esculin ferric citrate | + | | |

298 **Table 2.** Identification of bacteriocin-producing strains by BLAST and Ez-Taxon.

| Strain | BLAST | | EzTaxon | |
|---------|--|----------------|--|----------------|
| | Taxon name | Similarity (%) | Taxon name | Similarity (%) |
| CAU2013 | <i>Lactococcus lactis</i> subsp. <i>lactis</i> | 100 | <i>Lactococcus lactis</i> subsp. <i>lactis</i> | 100 |
| | <i>Lactococcus lactis</i> subsp. <i>hordniae</i> | 99.86 | <i>Lactococcus lactis</i> subsp. <i>hordniae</i> | 99.86 |
| | <i>Lactococcus lactis</i> subsp. <i>tractae</i> | 99.39 | <i>Lactococcus lactis</i> subsp. <i>tractae</i> | 99.39 |

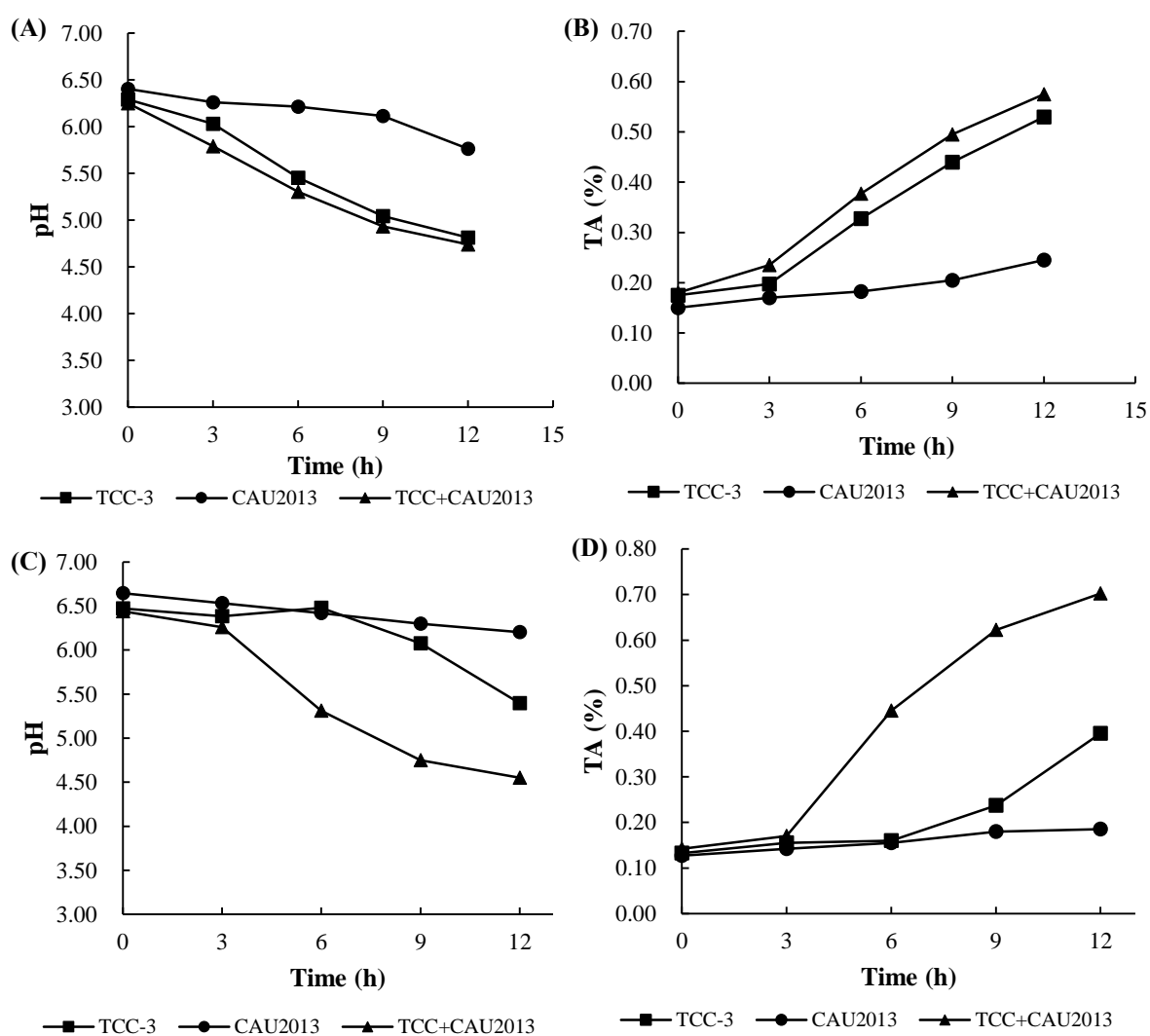
299
300
301
302 **Table 3.** Antimicrobial spectrum of bacteriocin from *L. lactis* CAU2013. +, < 10 mm; ++, >
303 10 mm; -, no inhibition zone.

| Indicator strain | Inhibition activity |
|---------------------------------------|---------------------|
| Gram positive | |
| <i>L. monocytogenes</i> ATCC 15315 | + |
| <i>L. monocytogenes</i> ATCC 7644 | ++ |
| <i>L. monocytogenes</i> ATCC 19111 | + |
| <i>L. monocytogenes</i> ATCC 19114 | ++ |
| <i>L. monocytogenes</i> ATCC 19115 | ++ |
| <i>Staphylococcus aureus</i> RN6390 | + |
| Gram negative | |
| <i>Salmonella enteritidis</i> YHS 383 | - |
| <i>Escherichia coli</i> ATCC 25922 | - |



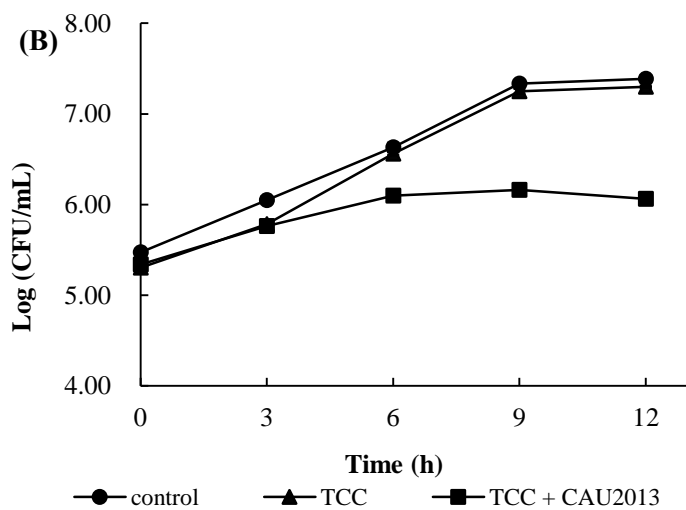
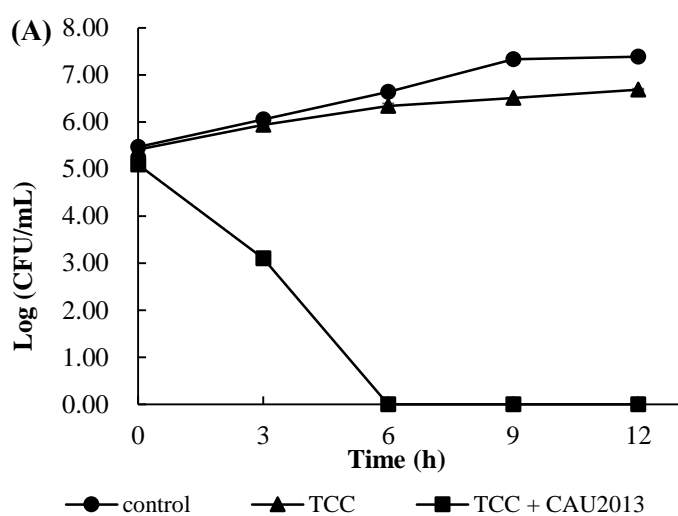
304 **Figure 1.** The procedure of lab-scale fresh cheese production.

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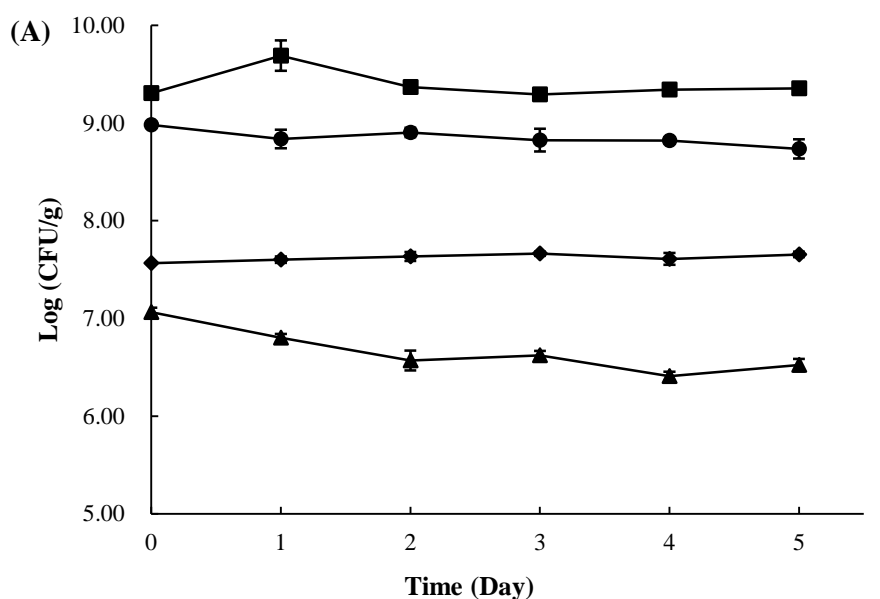
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 307 **Figure 2.** The values of pH and titratable acidity (TA) of strains grown in 10% skim milk and
 308 whole milk at 30°C. (A) pH values and (B) TA values in 10% skim milk, (C) pH values and
 309 (D) TA values in whole milk. TCC-3 (■), *L. lactis* CAU2013 (●), and the combination of
 310 TCC-3 with CAU2013 (▲).

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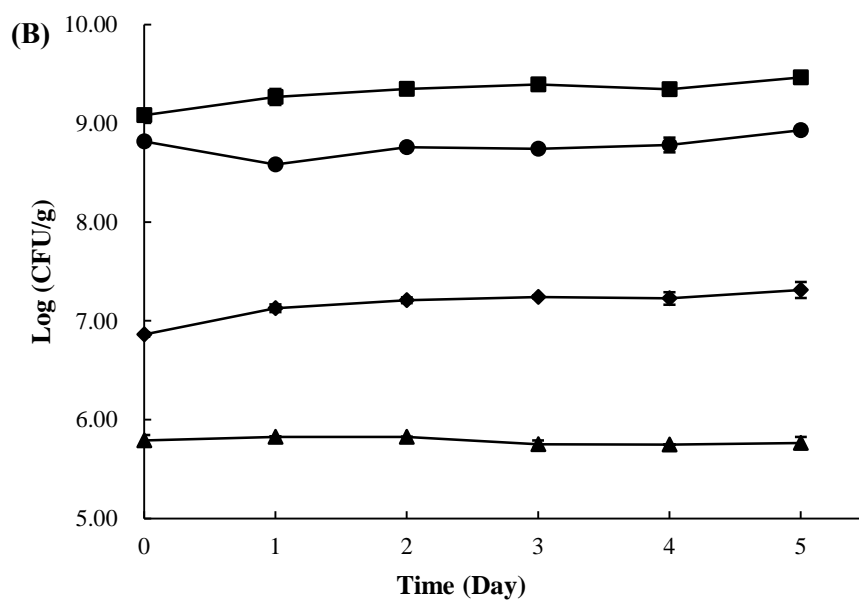


316 **Figure 3.** Biocontrol of *L. monocytogenes* in 10% skim milk broth and whole milk. *L.*
 317 *monocytogenes* was inoculated in milk broth (A) and whole milk (B), and incubated at 30°C
 318 without starter (●), or with 1 % of TCC-3 (▲), or the combination of TCC-3 and CAU2013
 319 (■).

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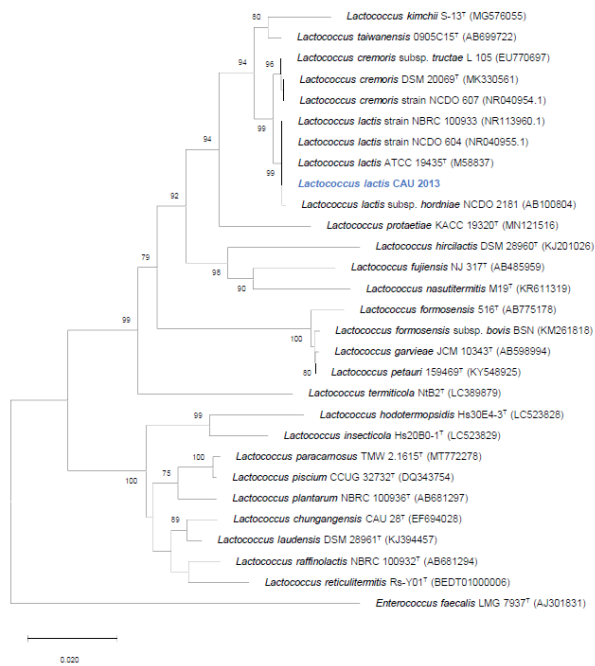
◆ TCC ▲ TCC+CAU2013 ● TCC(LAB) ■ TCC+CAU2013(LAB)



◆ TCC ▲ TCC+CAU2013 ● TCC(LAB) ■ TCC+CAU2013

323 **Figure 4.** The viable cell counts of *L. monocytogenes* and LAB in fresh cheese manufactured
 324 with TCC-3 starter and combination of *L. lactis* CAU2013 and TCC-3 and then stored for 5
 325 days. During the storage at 4°C (A) and 10°C (B), LAB in cheese produced with TCC-3 (●);
 326 TCC-3 and CAU2013 (■) were measured. Also, *L. monocytogenes* was measured in cheese
 327 with TCC-3 (◆) and cheese with mixed starter (▲).

328



Supplementary Fig 1. Neighbor-joining (NJ) phylogenetic tree of *Lactococcus lactis* strain CAU 2013 and related type strains based on 16S rRNA gene sequences (GenBank accession numbers are enclosed in parenthesis). Numbers at nodes (value >70%) are bootstrap values based on 1000 resampling datasets. *Enterococcus faecalis* LMG 7937^T was used as an outgroup. Bar, 0.02 substitutions per nucleotide position.

ACCEPTED