

Testing commercial biopreservative against spoilage microorganisms in MAP packed Ricotta fresca cheese

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3 1 Short Communication

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5 2 **Testing commercial biopreservative against spoilage microorganisms in MAP packed Ricotta**  
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7 ***fresca* cheese.**

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27 12 **Abstract**

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29 13 Ricotta *fresca* cheese is susceptible to secondary contamination and is able to support the growth of  
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31 14 pathogens or spoilage psychotrophic bacteria during storage. The aim of the present study was to  
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33 15 evaluate which among three commercial biopreservatives was suitable to be used to control the  
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35 16 growth of spoilage microorganisms in sheep's milk MAP ricotta *fresca* cheese. 144 Ricotta *fresca*  
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37 17 cheese samples were inoculated either with the bioprotective culture Lyofast FPR 2 (including  
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39 18 *Enterococcus faecium*, *Lactobacillus plantarum* e *Lactobacillus rhamnosus*) or Lyofast CNBAL  
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41 19 (*Carnobacterium* spp) or the fermentate FERM 430D. Not inoculated control and experimental  
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43 20 ricotta were MAP packed and stored at 4°C. Triplicate samples were analyzed after 5 h and 7, 14  
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45 21 and 21 days after inoculation for total bacterial count, mesophilic lactic acid bacteria,  
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47 22 *Enterobacteriaceae*, *Pseudomonas* spp, *Listeria monocytogenes*, moulds and yeasts.  
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50 23 *Carnobacterium* spp reduced the concentration of *Pseudomonas* spp and *Enterobacteriaceae* of at  
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52 24 least 1 log<sub>10</sub> at the end of the shelf-life. Therefore, *Carnobacterium* spp was selected as the culture  
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54 25 of choice to conduct a challenge study against *Pseudomonas* spp.

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27 Keywords: *Carnobacterium* spp.; protective cultures; *Pseudomonas* spp; ricotta; MAP.

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## 29 **1. Introduction**

30 *Ricotta fresca* is a traditional whey cheese produced by heat coagulation of sheep's milk whey. In  
31 Sardinia (Italy) it is generally manufactured from the whey remaining after the production of hard  
32 semi-cooked cheeses (Pecorino Romano PDO and Pecorino Sardo PDO). The industrial production  
33 follows the traditional batch production process (Pala *et al.*, 2016). *Ricotta fresca* intended for  
34 large-scale retail are commercialized in modified atmosphere packaging (MAP), under refrigeration  
35 temperature, with a shelf life determined by Food Business Operators up to 21 days. The batch  
36 production process exposes *Ricotta fresca* to post-process contamination originating from the dairy  
37 plant environment (Greenwood *et al.*, 1991). Due to its naturally poor competitive microflora  
38 (Pintado *et al.*, 2002), to its composition, inherent physical and chemical properties and the absence  
39 of preservatives, *Ricotta fresca* is an excellent substrate for the growth of pathogens or spoilage  
40 psychotropic bacteria during refrigerated storage (De Santis and Mazzette, 2002). The use of bio  
41 preservatives (i.e. nisin, other bacteriocins, fermentates or bioprotective cultures) alone or combined  
42 with other treatments, have been proposed to preserve the quality and safety of dairy products and  
43 other foods (Sobrino-López & Martín-Belloso, 2008; Elsser-Gravesen, & Elsser-Gravesen, 2013).  
44 Shelf life extension of whey cheeses using bio preservatives have been previously tested against  
45 *Listeria monocytogenes* (Davies *et al.*, 1997; Samelis *et al.*, 2003; Martins *et al.*, 2010). However, to  
46 date no available studies investigated the use of biopreservatives against psychotropic spoilage  
47 microorganism in sheep ricotta cheese. The present study was conducted as a preliminary  
48 investigation to assess the potential use of biopreservatives to control the growth of spoilage  
49 microorganism during refrigerated storage of MAP *ricotta fresca*. The main objective was to select  
50 which among two commercial bioprotective cultures and a fermentate was suitable to be used for a

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121 51 successive validation study. The selection of the biopreservative to be used was based on the  
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123 52 adaptation to *ricotta fresca* substrate and on the reduction of psychotropic microorganism's growth.

## 125 53 **2. Materials and methods**

### 127 128 54 *2.1. Biopreservatives*

129  
130 55 The protective cultures and the fermentate were selected, among available products on the market,  
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132 56 based on the proven activity against spoilage and pathogen microorganisms, their ability to grow at  
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134 57 refrigeration temperature and the low development of acidity and aroma in the product. Of the two  
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136 58 commercial protective cultures tested, one was Lyofast FPR 2 (Clerici-Sacco Group, Como, Italy)  
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138 59 consisting of *Enterococcus faecium*, *Lactobacillus plantarum* and *Lactobacillus rhamnosus* in the  
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140 60 ratio 1:1:1 with an optimum growth temperature of 37 °C. The second was Lyofast CNBAL  
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142 61 (Clerici-Sacco Group, Como, Italy) consisting of a selected strain of *Carnobacterium spp* producing  
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144 62 bacteriocins with an optimum growth temperature between 25-45 °C. The fermentate, the microbial  
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146 63 fermentation complex FERM 430D (Danisco), like other fermented products has a complex and  
147  
148 64 undefined composition.

### 150 151 65 *2.2. Samples*

152  
153 66 144 Ricotta *fresca* cheese samples were randomly selected from 3 different batches (48 from each  
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155 67 batch), manufactured in a local industrial sheep cheese making plant. The day after production  
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157 68 ricotta fresca samples were packed in rigid polypropylene trays sealed with lidding films and  
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159 69 transported refrigerated to the laboratory. Culture one samples (FRP) were ricotta *fresca* treated  
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161 70 with Lyofast FPR 2, culture two samples (CNBAL) were ricotta *fresca* treated with Lyofast  
162  
163 71 CNBAL and Fermentate samples (FERM) were ricotta *fresca* treated with FERM 430D. Blank  
164  
165 72 samples (BS) were untreated ricotta *fresca*. According to manufactures instruction protective  
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167 73 cultures were individually diluted in distilled water to a final concentration of  $10^6$  cfu mL<sup>-1</sup> while  
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169 74 the fermentate was resuspended in distilled water in order of 0.5-1% of the samples weight. Then  
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171 75 2.5 mL of Lyofast FPR 2 and Lyofast CNBAL were sprayed respectively on the surface of FPR and  
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180 76 CNBAL samples and 4 mL of FERM 430D final suspension distributed on the surface of FERM  
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182 77 samples. After the inoculum all *Ricotta fresca* cheese samples were repacked in MAP (30% CO<sub>2</sub>  
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184 78 and 70% N<sub>2</sub>) using the FP Basic Sec tray sealer (Ilpra, Vigevano, Italy).

### 186 79 2.3. Microbiological profile intrinsic properties and composition analysis

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189 80 For each batch, triplicate samples of ricotta *fresca* were analyzed for the determination of  
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191 81 microbiological profile, intrinsic properties and composition 5 h (T<sub>0</sub>), 7, 14 and 21 days (T<sub>7</sub>, T<sub>14</sub>,  
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193 82 T<sub>21</sub>) after the addition of the biopreservatives. Microbiological analysis were conducted for the  
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195 83 enumeration of aerobic mesophilic bacteria (ISO 4833:2003), for the enumeration of mesophilic  
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197 84 lactic acid bacteria (ISO 15214: 1998), for the enumeration of *Pseudomonas* spp (ISO/TS  
198  
199 85 11059:2009), for the detection and enumeration of *Enterobacteriaceae* (ISO 21528-1:2004) and for  
200  
201 86 the enumeration of yeast and molds (ISO 6611/IDF094:2004). Detection and enumeration of  
202  
203 87 *Listeria monocytogenes* was also conducted (ISO 11290-1: 1996, ISO 11290-2:1998). Samples  
204  
205 88 inoculated with Lyofast CNBAL at T<sub>0</sub> were also analyzed for the enumeration of *Carnobacterium*  
206  
207 89 spp using MRS modified by increasing the pH to 8.5, omitting acetate, and substituting glucose for  
208  
209 90 sucrose (Hammes et al., 1992).

### 212 91 2.4. Intrinsic properties, composition and headspace gas analysis

214 92 PH and a<sub>w</sub> were measured using pH meter GLP22 (Crison Instruments SA, Barcelona, Spain) and  
215  
216 93 water activity meter Aqualab 4TE (Decagon, Pullman, WA, USA) respectively. Fat, moisture,  
217  
218 94 protein and total solids were analyzed using a near infrared transmittance (NIT) compositional  
219  
220 95 analyzer (FOSS, Eden Prairie, MN, USA). The composition of the headspace gas mixture was  
221  
222 96 conducted on ricotta *fresca* samples on the sealed packages prior to other analysis. Measure of  
223  
224 97 combined residual O<sub>2</sub> % and CO<sub>2</sub> % were obtained piercing the lid using a sterile needle connected  
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226 98 to the Dansensor gas analyser (PBI Dansensor, Ringsted, Denmark).

## 231 99 3. Results and discussion

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### 3.1. Microbiological profile

*Ricotta fresca* cheese total bacterial count in control samples at  $T_0$  was  $< 3 \log_{10}$  cfu  $g^{-1}$  and increased after 21 days of refrigerated storage above  $7 \log_{10}$  cfu  $g^{-1}$  while the mesophilic lactic acid bacteria were below the detection limit at  $T_0$  and ca  $5 \log_{10}$  cfu  $g^{-1}$  at  $T_{21}$ . During refrigerated storage a significant increase ( $P < 0.01$ ) of spoilage microorganisms to level as high as  $6 \log_{10}$  and  $8 \log_{10}$  was observed for *Enterobacteriaceae* and *Pseudomonas* spp, respectively. Yeast and molds were occasionally reported, with maximum values around  $4 \log_{10}$  at  $T_{21}$ . The complete microbiological profile with mean counts ( $\log_{10}$  cfu  $g^{-1}$ ;  $\bar{x} \pm SD$ ) over time is reported in table 1. *L. monocytogenes* was never detected on either blank samples and ricotta inoculated with biopreservatives. *Carnobacterium* spp.  $\log_{10}$  counts were  $6.28 \pm 0.35$  at  $T_0$ ,  $6.64 \pm 1.56$  at  $T_7$ ,  $8.03 \pm 0.39$  at  $T_{14}$  and  $8.59 \pm 0.47$  at  $T_{21}$  showing a significant increase after  $T_{14}$  ( $P < 0.05$ ).

### 3.2. Physico-chemical characteristics and MAP gas composition

In blank samples the pH showed a slight decrease over time, from 6.67 at  $T_0$  to 6.52 at  $T_{21}$  ( $P < 0.05$ ) while no significant difference was observed in the  $a_w$ . In blank samples the  $O_2$  content in the headspace increased from the initial level of 0.87% up to 1.80% at  $T_7$ , to decrease again as low as 0.42 at  $T_{21}$ . Instead, the  $CO_2$  content decreased from  $T_0$  to  $T_{21}$  respectively from 13.05% to 6.78%. Intrinsic properties, composition and gas composition in the headspace ( $\bar{x} \pm SD$ ) during the refrigerated storage are reported in table 2.

## 4. Discussion

*Ricotta fresca* cheese as consequence of high temperature applied during manufacturing has poor competing microbiota, which reflects on the growth of psychotropic microorganisms such as *Pseudomonas* spp., *Enterobacteriaceae*, *Listeria monocytogenes*, *B. cereus* and *Arcobacter* spp (De Santis and Mazzette, 2002; De Santis *et al.* 2008; Ibba *et al.*, 2013; Scarano *et al.*; 2014; Spanu *et al.*, 2016). However, a large part of Ricotta cheese microflora at the end of the shelf-life is

296  
297  
298 126 generally represented by *Pseudomonas* spp, that could exert a competitive activity against other  
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300 127 species, including pathogens (Pala *et al.*, 2016). As the improvement of the hygiene management  
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302 128 procedures is a measure that could only reduce the level of initial contamination, the use of bio  
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305 129 preservatives to compete with contaminants is an interesting perspective in Ricotta cheese. The  
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307 130 fermentate showed no activity against the growth of microbiota in ricotta during refrigerated  
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309 131 storage. In fact, total bacterial counts, LAB, *Enterobacteriaceae*, *Pseudomonas* spp., yeast and  
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311 132 molds showed no significant differences between blank samples and samples inoculated with  
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313 133 FERM. The higher counts at T<sub>0</sub> of mesophilic LAB (ca 5 log<sub>10</sub>) in *Ricotta fresca* cheese samples  
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315 134 inoculated with FRP as compared to control samples and ricotta inoculated with the other bio  
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317 135 preservatives was expected. FRP cultures demonstrated, despite the refrigeration, a slight increase  
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319 136 during storage (ca 1 log<sub>10</sub>). However, FRP showed no control against *Enterobacteriaceae* and  
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322 137 *Pseudomonas* spp which, at the end of the storage, were ca 1 log<sub>10</sub> higher respect to blank samples.  
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324 138 In ricotta samples inoculated with CNBAL mesophilic LAB counts were always lower as compared  
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326 139 to the other samples. This result could be explained with the fact that for the isolation and  
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328 140 cultivation of LAB the De Man, Rogosa and Sharpe (MRS) agar is generally used, in which it has  
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330 141 been observed that most of the *Carnobacterium* spp are not able to growth (Hammes *et al.*, 1992).  
331  
332 142 This could lead to a significant underestimation of its concentration in foods. In the present study  
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334 143 *Carnobacterium* spp showed a good adaptive response to the experimental condition of inoculum  
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336 144 and storage, showing an increase in its mean counts of approximately of 2 log<sub>10</sub> from T<sub>0</sub> to T<sub>21</sub>. The  
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338  
339 145 competitive activity of CNBAL was effective in reducing *Pseudomonas* spp and  
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341 146 *Enterobacteriaceae* at the end of the shelf-life of at least 1 log<sub>10</sub>. However, it should be noticed that  
342  
343 147 the effect of CNBAL was greater after 14 days were the difference with blank samples was  
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345 148 respectively of 2 log<sub>10</sub> for *Pseudomonas* and almost 3 log<sub>10</sub> for *Enterobacteriaceae*. It is worth to  
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347 149 note that the growth of *Carnobacterium* spp did not lowered Ricotta fresca pH, which may have had  
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349 150 an impact of the sensory characteristics. The possible adoption of CLAB as protective culture  
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151 requires the determination of the changes in the sensory profile of *Ricotta fresca*. However, sensory  
152 analysis could not be performed in the present research since the level of *Pseudomonas* spp  
153 contamination at T<sub>14</sub> was already as high as 6 log<sub>10</sub>, compatible with alteration of the product, and  
154 yet beyond the acceptability of consumers.

155 The gas mixture chosen for MAP packaging of ricotta *fresca* (30% CO<sub>2</sub> and 70% O<sub>2</sub>) is the  
156 composition generally used in sardinian industrial cheesemaking plants. As previously  
157 demonstrated (Pala *et al.*, 2016), the concentration of CO<sub>2</sub> in the head space at T<sub>0</sub> differs from the  
158 level used during packaging, as a result of gas solving in the product, while the further reduction  
159 during the successive storage is attributable to gas permeability of packaging materials used.  
160 Instead, the reduction of O<sub>2</sub>% during storage is associated with the growth of aerobic mesophilic  
161 microorganisms.

## 5. Conclusion

162 The present study was specifically designed to provide preliminary information on the possible use  
163 of biopreservatives to control the growth of psychotropic spoilage microorganism's in MAP  
164 packaged ricotta fresca. Since no information was previously available on the adaptation of  
165 biopreservatives on sheep's milk ricotta *fresca*, the primary objective of the study was to select  
166 among three commercial products which one was suitable as biopreservative. *Carnobacterium* spp.  
167 inoculated on the finished product showed a good adaptation to grow in ricotta *fresca* and  
168 promising results in controlling spoilage microorganisms. However, the present investigation was  
169 conducted on naturally contaminated ricotta samples. Therefore, CNBAL was the protective culture  
170 of choice to conduct a challenge test specifically designed to assess the effect of *Carnobacterium*  
171 spp against *Pseudomonas* spp.

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229 Table 1. Microbiological profile ( $\log_{10}$  cfu  $g^{-1}$ ;  $\bar{x} \pm SD$ ) of ricotta fresca by time (days after) and sample type.

Parameters	sample unit	T <sub>0</sub>	T <sub>7</sub>	T <sub>14</sub>	T <sub>21</sub>
Aerobic mesophilic bacteria	BS	2.72 ± 0.44 <sup>a</sup> ( <i>n</i> = 9/9)	4.90 ± 1.36 <sup>b</sup> ( <i>n</i> = 9/9)	6.01 ± 0.56 <sup>c</sup> ( <i>n</i> = 9/9)	6.90 ± 0.86 <sup>d</sup> ( <i>n</i> = 9/9)
	CNBAL	3.18 ± 1.95 <sup>a</sup> ( <i>n</i> = 9/9)	6.53 ± 0.85 <sup>b</sup> ( <i>n</i> = 9/9)	6.87 ± 1.13 <sup>b</sup> ( <i>n</i> = 9/9)	8.87 ± 0.38 <sup>c</sup> ( <i>n</i> = 9/9)
	FRP2	5.11 ± 0.62 <sup>a</sup> ( <i>n</i> = 9/9)	6.63 ± 0.91 <sup>b</sup> ( <i>n</i> = 9/9)	7.35 ± 0.51 <sup>c</sup> ( <i>n</i> = 9/9)	7.85 ± 0.22 <sup>c</sup> ( <i>n</i> = 9/9)
	FERM	2.91 ± 0.51 <sup>a</sup> ( <i>n</i> = 9/9)	5.21 ± 1.05 <sup>b</sup> ( <i>n</i> = 9/9)	6.11 ± 0.82 <sup>c</sup> ( <i>n</i> = 9/9)	6.92 ± 0.51 <sup>d</sup> ( <i>n</i> = 9/9)
mesophilic lactic acid bacteria	BS	ND	3.55 ± 0.49 <sup>a</sup> ( <i>n</i> = 9/9)	4.33 ± 0.71 <sup>b</sup> ( <i>n</i> = 9/9)	4.92 ± 0.67 <sup>b</sup> ( <i>n</i> = 9/9)
	CNBAL	2.13 ± 0.76 <sup>a</sup> ( <i>n</i> = 4/9)	3.30 ± 0.93 <sup>b</sup> ( <i>n</i> = 9/9)	3.76 ± 0.62 <sup>b</sup> ( <i>n</i> = 9/9)	3.32 ± 0.60 <sup>b</sup> ( <i>n</i> = 9/9)
	FRP2	5.01 ± 0.72 <sup>a</sup> ( <i>n</i> = 9/9)	5.35 ± 0.78 <sup>ab</sup> ( <i>n</i> = 9/9)	5.16 ± 0.44 <sup>a</sup> ( <i>n</i> = 9/9)	5.77 ± 0.40 <sup>b</sup> ( <i>n</i> = 9/9)
	FERM	1.52 ± 0.24 <sup>a</sup> ( <i>n</i> = 3/9)	2.86 ± 0.05 <sup>ab</sup> ( <i>n</i> = 2/9)	4.05 ± 0.68 <sup>b</sup> ( <i>n</i> = 7/9)	4.58 ± 1.68 <sup>b</sup> ( <i>n</i> = 7/9)
<i>Enterobacteriaceae</i>	BS	2.20 ± 1.02 <sup>a</sup> ( <i>n</i> = 4/9)	4.05 ± 0.86 <sup>b</sup> ( <i>n</i> = 5/9)	4.43 ± 0.99 <sup>b</sup> ( <i>n</i> = 7/9)	5.34 ± 0.97 <sup>b</sup> ( <i>n</i> = 8/9)
	CNBAL	ND	1.95 ± 0.00 <sup>ab</sup> ( <i>n</i> = 1/9)	1.77 ± 1.15 <sup>a</sup> ( <i>n</i> = 6/9)	3.90 ± 0.42 <sup>b</sup> ( <i>n</i> = 5/9)
	FRP2	2.03 ± 0.00 <sup>a</sup> ( <i>n</i> = 2/9)	3.79 ± 0.67 <sup>b</sup> ( <i>n</i> = 9/9)	5.41 ± 0.75 <sup>c</sup> ( <i>n</i> = 9/9)	6.29 ± 0.47 <sup>d</sup> ( <i>n</i> = 9/9)
	FERM	3.79 ± 1.24 <sup>a</sup> ( <i>n</i> = 4/9)	3.21 ± 0.82 <sup>a</sup> ( <i>n</i> = 6/9)	4.24 ± 0.91 <sup>a</sup> ( <i>n</i> = 6/9)	5.84 ± 0.59 <sup>b</sup> ( <i>n</i> = 7/9)
<i>Pseudomonas</i> spp	BS	2.64 ± 0.59 <sup>a</sup> ( <i>n</i> = 5/9)	4.89 ± 1.21 <sup>b</sup> ( <i>n</i> = 9/9)	6.52 ± 0.99 <sup>c</sup> ( <i>n</i> = 9/9)	6.83 ± 0.91 <sup>c</sup> ( <i>n</i> = 9/9)
	CNBAL	2.43 ± 0.18 <sup>a</sup> ( <i>n</i> = 4/9)	2.59 ± 0.67 <sup>a</sup> ( <i>n</i> = 9/9)	4.59 ± 0.65 <sup>b</sup> ( <i>n</i> = 9/9)	5.27 ± 0.64 <sup>b</sup> ( <i>n</i> = 9/9)
	FRP2	2.53 ± 0.51 <sup>a</sup> ( <i>n</i> = 5/9)	5.89 ± 0.64 <sup>b</sup> ( <i>n</i> = 9/9)	6.81 ± 0.82 <sup>c</sup> ( <i>n</i> = 9/9)	7.01 ± 0.53 <sup>c</sup> ( <i>n</i> = 9/9)
	FERM	2.69 ± 0.27 <sup>a</sup> ( <i>n</i> = 6/9)	5.02 ± 0.81 <sup>b</sup> ( <i>n</i> = 9/9)	6.33 ± 0.94 <sup>c</sup> ( <i>n</i> = 9/9)	7.26 ± 0.31 <sup>d</sup> ( <i>n</i> = 9/9)
Yeast and molds	BS	ND	2.78 ± 0.40 <sup>a</sup> ( <i>n</i> = 4/9)	3.62 ± 0.33 <sup>b</sup> ( <i>n</i> = 3/9)	3.43 ± 0.76 <sup>ab</sup> ( <i>n</i> = 5/9)
	CNBAL	2.36 ± 0.10 <sup>a</sup> ( <i>n</i> = 3/9)	2.15 ± 0.21 <sup>a</sup> ( <i>n</i> = 2/9)	3.00 ± 0.00 <sup>b</sup> ( <i>n</i> = 1/9)	ND
	FRP2	2.00 ± 0.00 <sup>a</sup> ( <i>n</i> = 1/9)	3.01 ± 0.49 <sup>ab</sup> ( <i>n</i> = 3/9)	3.52 ± 0.38 <sup>b</sup> ( <i>n</i> = 5/9)	3.64 ± 0.73 <sup>b</sup> ( <i>n</i> = 3/9)
	FERM	ND	3.97 ± 0.42 <sup>a</sup> ( <i>n</i> = 3/9)	3.88 ± 0.68 <sup>a</sup> ( <i>n</i> = 9/9)	3.19 ± 1.14 <sup>a</sup> ( <i>n</i> = 8/9)

BS indicates blank sample units used as negative control; C1, C2 and FERM indicates samples units inoculated respectively with protective culture CNBAL and FRP2 and the fermentate. Means in the same row with different superscript were significantly different ( $P < 0.05$ ); Values within brackets indicate the prevalence of positive samples.

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234 Table 2. Intrinsic properties and composition ( $\bar{x} \pm SD$ ) of Ricotta *fresca* cheese at different testing times.

Parameters	sample unit	T <sub>0</sub>	T <sub>7</sub>	T <sub>14</sub>	T <sub>21</sub>
pH	BS	6.67 ± 0.1 <sup>a</sup>	6.58 ± 0.05 <sup>bc</sup>	6.61 ± 0.07 <sup>ab</sup>	6.52 ± 0.11 <sup>c</sup>
	CNBAL	6.66 ± 0.12 <sup>a</sup>	6.66 ± 0.11 <sup>a</sup>	6.66 ± 0.09 <sup>ab</sup>	6.54 ± 0.04 <sup>c</sup>
	FRP2	6.67 ± 0.06 <sup>a</sup>	6.59 ± 0.05 <sup>b</sup>	6.55 ± 0.07 <sup>b</sup>	6.31 ± 0.07 <sup>c</sup>
	FERM	6.68 ± 0.10 <sup>a</sup>	6.56 ± 0.04 <sup>bc</sup>	6.60 ± 0.08 <sup>ab</sup>	6.49 ± 0.14 <sup>c</sup>
a <sub>w</sub>	BS	0.990 ± 0.003 <sup>a</sup>	0.996 ± 0.006 <sup>a</sup>	0.993 ± 0.006 <sup>a</sup>	0.993 ± 0.006 <sup>a</sup>
	CNBAL	0.991 ± 0.005 <sup>a</sup>	0.989 ± 0.008 <sup>a</sup>	0.986 ± 0.001 <sup>a</sup>	0.985 ± 0.001 <sup>a</sup>
	FRP2	0.995 ± 0.004 <sup>a</sup>	0.995 ± 0.001 <sup>a</sup>	0.997 ± 0.001 <sup>a</sup>	0.997 ± 0.002 <sup>a</sup>
	FERM	0.994 ± 0.003 <sup>a</sup>	0.994 ± 0.001 <sup>a</sup>	0.992 ± 0.005 <sup>a</sup>	0.993 ± 0.008 <sup>a</sup>
Moisture (%)	BS	71.23 ± 3.52 <sup>a</sup>	73.37 ± 2.10 <sup>a</sup>	73.56 ± 2.08 <sup>a</sup>	74.70 ± 0.91 <sup>a</sup>
	CNBAL	72.02 ± 3.38 <sup>a</sup>	71.97 ± 3.82 <sup>a</sup>	71.20 ± 3.92 <sup>a</sup>	71.68 ± 2.91 <sup>a</sup>
	FRP2	77.43 ± 3.11 <sup>a</sup>	73.27 ± 2.83 <sup>a</sup>	74.35 ± 3.54 <sup>a</sup>	72.22 ± 2.09 <sup>a</sup>
	FERM	74.26 ± 2.74 <sup>a</sup>	74.61 ± 0.83 <sup>a</sup>	74.13 ± 2.01 <sup>a</sup>	73.46 ± 1.29 <sup>a</sup>
Fat (%)	BS	18.13 ± 5.80 <sup>a</sup>	14.31 ± 2.31 <sup>a</sup>	13.30 ± 2.60 <sup>a</sup>	14.66 ± 2.96 <sup>a</sup>
	CNBAL	18.03 ± 4.48 <sup>a</sup>	17.48 ± 4.46 <sup>a</sup>	17.15 ± 3.95 <sup>a</sup>	17.49 ± 4.46 <sup>a</sup>
	FRP2	11.03 ± 2.59 <sup>a</sup>	17.15 ± 1.16 <sup>c</sup>	13.09 ± 2.98 <sup>ab</sup>	15.11 ± 1.08 <sup>bc</sup>
	FERM	13.78 ± 1.62 <sup>a</sup>	12.43 ± 2.10 <sup>a</sup>	13.59 ± 1.88 <sup>a</sup>	14.67 ± 2.53 <sup>a</sup>
Protein (%)	BS	9.81 ± 0.78 <sup>a</sup>	9.97 ± 1.04 <sup>a</sup>	10.23 ± 1.36 <sup>a</sup>	8.94 ± 2.61 <sup>a</sup>
	CNBAL	9.38 ± 0.45 <sup>a</sup>	9.33 ± 0.75 <sup>a</sup>	9.46 ± 0.39 <sup>ab</sup>	10.46 ± 0.48 <sup>b</sup>
	FRP2	10.05 ± 0.02 <sup>a</sup>	11.15 ± 1.07 <sup>a</sup>	9.99 ± 0.73 <sup>a</sup>	10.86 ± 0.24 <sup>a</sup>
	FERM	9.77 ± 1.50 <sup>a</sup>	10.32 ± 1.37 <sup>a</sup>	9.87 ± 0.76 <sup>a</sup>	10.19 ± 0.81 <sup>a</sup>
O <sub>2</sub> %	BS	0.87 ± 0.49 <sup>a</sup>	1.80 ± 1.18 <sup>b</sup>	1.05 ± 0.82 <sup>a</sup>	0.42 ± 0.78 <sup>a</sup>
	CNBAL	0.99 ± 0.58 <sup>ab</sup>	1.12 ± 1.65 <sup>b</sup>	0.14 ± 0.17 <sup>ab</sup>	0.02 ± 0.01 <sup>b</sup>
	FRP2	0.99 ± 0.54 <sup>a</sup>	1.89 ± 1.47 <sup>b</sup>	0.31 ± 0.25 <sup>ac</sup>	0.004 ± 0.01 <sup>c</sup>
	FERM	1.01 ± 0.63 <sup>a</sup>	1.66 ± 0.32 <sup>b</sup>	0.75 ± 0.91 <sup>a</sup>	0.51 ± 0.84 <sup>a</sup>
CO <sub>2</sub> %	BS	13.05 ± 2.88 <sup>a</sup>	6.20 ± 2.09 <sup>b</sup>	5.50 ± 2.96 <sup>b</sup>	6.78 ± 2.89 <sup>b</sup>
	CNBAL	13.55 ± 2.42 <sup>a</sup>	5.50 ± 2.02 <sup>b</sup>	5.00 ± 2.08 <sup>b</sup>	5.18 ± 1.91 <sup>b</sup>
	FRP2	13.68 ± 2.23 <sup>a</sup>	6.96 ± 1.55 <sup>c</sup>	7.50 ± 1.73 <sup>c</sup>	10.22 ± 1.72 <sup>b</sup>
	FERM	13.24 ± 2.21 <sup>a</sup>	6.47 ± 2.10 <sup>b</sup>	5.79 ± 2.67 <sup>b</sup>	6.98 ± 4.15 <sup>b</sup>

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235 BS indicates blank sample units used as negative control; C1, C2 and FERM indicates samples units inoculated respectively with protective culture CNBAL and  
236 FRP2 and the fermentate. Means in the same row with different superscripts were significantly different ( $P < 0.05$ ).  
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## Highlights

- Ricotta fresca support psychotrophic microorganism growth during refrigerated storage
- 3 Commercial biopreservatives were tested against spoilage bacteria in ricotta fresca
- Cultures Lyofast FPR 2, Lyofast CNBAL and the fermentate FERM 430D were tested
- Inoculation of Carnobacterium spp protective culture reduced *Pseudomonas* spp growth