



# Meat starter cultures: Isolation and characterization of lactic acid bacteria from traditional sausages

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**Abstract.** Fermented meat products represent an important segment of our alimentation. Obtaining these products is based on beneficial microorganism activity. In the case of traditional food products, these are commercial starters or autochthonous microflora. Fermentation of raw materials is mainly done by sugar metabolization of lactic acid bacteria (LAB). In addition, these microorganisms can have other beneficial properties too such as probiotic properties, antimicrobial compound production abilities, etc.

In order to meet consumer demands, starter cultures are continuously developed to produce high-quality, healthy, and tasty products, thus contributing to guaranteeing microbiological safety and to improving one or

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**Keywords and phrases:** lactic acid bacteria, starter culture, fermentation, sausage

more sensory characteristics, technological, nutritional, or health properties of the fermented products. The aim of our research is to determine the technological properties of autochthonous lactic acid bacteria originated from commercial fresh sausages in order to select and use them as potential starter cultures in the meat industry. In our work, we determined the relevant characteristics (such as salt tolerance, proteolytic activity, antimicrobial activity, and antibiotic resistance) of bacteria isolated from 16 fresh sausages. Based on our results, the studied bacterial isolates originated from sausages could be potentially used as autochthonous meat starter cultures.

## 1 Introduction

Some of the important preservation methods of meat products are fermentation and drying. Traditional sausage production has a long history, originating from the Mediterranean region. Worldwide, Europeans are the main consumers and producers of this type of fermented meat products. Over centuries, the environmental conditions, certain intrinsic factors, and tradition had an impact on the production of sausage and fermented meat (Zeuthen, 1995).

Traditional sausages are defined (Leroy *et al.*, 2015) as meat products containing different meat mixtures with added spices and nitrite/nitrate salts, stuffed in casing, eventually smoked. The final product is obtained by a ripening process including fermentation and drying.

Depending on the sausage type, the fermentation is initiated by commercial starters or autochthonous microflora. Fermented meat products have several benefits (Singh *et al.*, 2012; Bourdichon *et al.*, 2012) such as enhanced nutritional and quality characteristics, are rich in essential amino and fatty acids, and are less perishable with an extended shelf life.

In fermentation, two groups of microorganisms play a key role, namely lactic acid bacteria (LAB) (counts of  $10^3$ – $10^9$  CFU/g) and coagulase-negative Gram-positive staphylococci (counts up to  $10^8$  CFU/g). In addition to these, less significant are eukaryotic microorganisms, yeasts, and moulds. Microbes and their enzymes are responsible for the complex biochemical changes in the meat matrix. LAB have an impact on the technological characteristics of the product and also influence microbiological stability. In sausages, bacterial starter culture strains determine the functional properties and safety aspects of the product. Acidification, flavour development, proteolytic activity, amino acid metabolism, antioxidant enzymes (catalase and superoxide dismutase), and ni-

trate reduction all contribute to the formation of the final product. Regarding the safety of the starter culture, it should not possess biogenic amine production and acquired antimicrobial resistance. It also has to be free of enterotoxin determinants (Cocconcelli & Fontana, 2014). During the metabolic activities, lactic acid (resulted from the fermentation of carbohydrates by LAB) reduces pH-level and thus contributes to safety, texture transformation, and acid taste development. Depending on the microflora, raw materials, and additives, the following metabolites are released in varying amounts: acetoin, pyruvic and acetic acid, ethanol, and carbon dioxide. Safety aspects result also from the bioprotective character of some of the LAB (Garriga & Aymerich, 2015) as different antimicrobial natural peptides (bacteriocins) are produced by them. Furthermore, owing to probiotic properties, LAB may enhance the functional value of the fermented food.

Microbial ecology of fermented sausages could be considered as a complex and rich microbiological niche. Bacterial strains involved in fermentation and biochemical transformation could be characterized by robustness and flexibility. Acidification is a functional characteristic of the most frequently present species of the *Lactobacillus* genus. *Lactobacillus sakei* was the predominant species among the LAB isolated from different sausages prepared with altered technologies. Different genotypes of this bacteria were isolated, and they represent 55% of all LAB (Garriga & Aymerich, 2015). The main explanation of the fact that these bacteria harboured and adapted to the conditions is that they are able to use as energy source both carbohydrates and amino acids (Cocconcelli & Fontana, 2015). Other species of the *Lactobacillus* genus were also detected: *Lactobacillus plantarum*, *Lb. curvatus*, and *Lb. rhamnosus*. Unlike in Europe, *Pediococcus* was a relevant member of the microbial ecology of sausages in the United States as it is mostly added as starter culture. *Pediococcus acidilactici* and *P. pentosaceus* are responsible also for acidification and are able to produce bacteriocins (Cocconcelli, 2007; Cocconcelli & Fontana, 2014). The predominant Gram-positive, catalase-positive cocci are *Staphylococcus saprophyticus*, *S. xylosus*, *S. succinus*, and *S. equorum*. Different technological functions are related to this species due to the involvement of the fatty acid metabolism and amino acid catabolism in flavour development. Nitrate reduction leads to the red colour formation – for example, *Kocuria varians* is added for nitrate reduction (Selgas & García, 2014) –, and catalase activity prevents lipid oxidation (Macedo *et al.*, 2017).

Safer and functional foods are the current trends in food consumption. To satisfy consumer demands, there are certain innovative attempts, for example, in starter culture development. Besides product processing, environmen-

tal factors also influence the metabolic activity of starter cultures. A current strategy in meat industry is the application of wild strains for standardized sausage fermentation. To meet this demand, selection criteria focus on naturally occurring strains from meat ecology with desired technological aspects and stress resistance (*Corbo et al.*, 2017; *Pereira et al.*, 2019).

The aim of the present study was to determine the technological properties of autochthonous lactic acid bacteria originated from commercial fresh sausages in order to select and use as potential starter cultures in the meat industry.

## 2 Materials and methods

### *Isolation of lactic acid bacteria*

In the course of our work, more than 60 lactic acid bacteria were isolated on de Man, Rogosa, and Sharpe (MRS) agar from 16 different commercially available, traditionally processed, fresh sausages. Surface-sterilized (2 min in 72% ethanol) 10 g sausages were smashed in a sterile mortar and homogenized with 90 ml physiological solution. From these samples, dilution series were prepared and an amount of 0.1 ml from the homogenized mixture was spread on the surface of MRS agar medium and incubated at 37 °C for 24 hrs in aerobic conditions. The isolated bacteria were analysed for the most representative technological characteristics of LAB; the experiments were done in two replicates.

### *Bacterial growth at different salt concentrations*

LAB isolates 0.1% (v/v) were inoculated in MRS broth with different salt concentrations: 6.5%, 8%, and 10% (*Papamanoli et al.*, 2010). Bacterial growth was evaluated after incubation at 30 °C for 48 hrs as absorbance values at 630 nm. Salt tolerance was expressed as growth index ( $G_i$ ) (*Speranza et al.*, 2014), calculated from the formula:

$$G_i = 100 \cdot \frac{A_s}{A_c},$$

where:  $A_s$  – sample absorbance in the present of salt and  $A_c$  – absorbance of the control.

### *Bacterial growth at different temperatures*

LAB isolates 0.1% (v/v) were inoculated in MRS broth and incubated at three different temperatures: 20 °C, 30 °C, and 37 °C. Bacterial growth was evaluated after incubation as absorbance values at 630 nm.

### *Bacterial growth at different pH values*

LAB isolates 0.1% (v/v) were inoculated in MRS broth adjusted to different pH values: 3, 4, 5 (*Papamanoli et al.*, 2010). Bacterial growth was evaluated after incubation at 30 °C for 48 hrs as absorbance values at 630 nm.

### *Gas production*

Gas production of LAB isolates was detected in MRS broth containing inverted Durham tubes (*Patil et al.*, 2010).

### *Proteolytic activity*

Proteolytic activity of LAB isolates was determined with well diffusion method on modified skimmed milk agar medium (0.5% casein, 0.25% yeast extract, 0.1% dextrose, skim milk powder 2.8%, and agar 1.5%). 50 µl of supernatant LAB isolates were inoculated in the 8-mm-diameter hole in the modified skimmed milk agar medium and incubated at 37 °C for 48 hrs. Proteolytic activity was determined in accordance with the measured clear zone diameter surrounding each culture.

### *Antibacterial activity*

The antibacterial activity of LAB isolates was determined with agar diffusion method on different pathogenic and spoilage indicator strains (from the microbiological laboratory of the University) as *Micrococcus luteus*, *Bacillus cereus*, *Bacillus subtilis*, *Escherichia coli*, *Staphylococcus aureus*, and *Proteus vulgaris*. These bacterial cultures were grown for 24 hrs at 28 °C and 37 °C on Nutrient agar. 0.1 ml bacterial suspension (with OD = 1) was spread in the case of each bacteria with surface streaking on Nutrient agar (meat extract 1 g, yeast extract 2 g, peptone 5 g, NaCl 5 g, agar 15 g, distilled water 1000 ml), and 50 µl of the cell-free supernatant of LAB isolates (centrifuged at 14,000 rpm, for 5 min) was dropped in the 8-mm-diameter hole cut with the help of a sterile test-tube. The inoculated Petri dishes were incubated at 28 °C

and 37°C. The antibacterial effect of the tested LAB isolates was expressed in accordance with the diameter of the inhibition zone.

### *Antibiotic susceptibility*

Determination of antimicrobial susceptibilities of LAB isolates was realized according to the guidelines reported in EFSA (2012). For the assessment of the susceptibility to ampicillin, gentamicin, streptomycin, erythromycin, chloramphenicol, and kanamycin, two-fold serial dilutions were realized ranging from 0 up to 128 µg/mL, with the exception of erythromycin (1–2 µg/mL) in MRS broth (*Laslo et al.*, 2015).

## 3 Results and discussions

Enumeration of viable LAB counts were done from 16 different, commercially available, traditionally processed sausage samples. The viable count of LAB in tested samples ranged between  $3.3 \cdot 10^1$  and  $5.8 \cdot 10^5$  CFU/g. In three samples, the viable count of LAB was very low, perhaps because these samples underwent heat treatment during the production process. Based on colony size and morphology, 35 colonies were picked from MRS agar for further characterization.

Salt content is one of the major environmental factors that has an impact on bacterial strains involved in fermentation. Salt is used in processed meat to develop the taste and extend shelf life. Almost all of the tested LAB isolates were able to grow in the presence of 6.5% salt. According to *Cruxen et al.* (2019), salt tolerance is a selection criterion in the case of native isolates. Our results regarding the tolerance of 6.5% NaCl are in concordance with those reported by *Almeida et al.* (2015). A higher salt concentration decreased the growth of the isolates, with an exception, as found also by *Aina* (2019). *Lb. plantarum* originated from artisanal sausage could survive the presence of 7% NaCl, whereas *Staphylococcus* spp. tolerated salt concentration up to 15% (*Cruxen et al.*, 2019).

The growth index (*Fig. 1*) varied in the range of 4–99%, and 74% of the isolates showed excellent growth. With the exception of 11% of LAB isolates, all of them had a higher growth index. In the presence of 8% salt, 26% of the isolated LAB were able to grow well, 37% of the isolated bacteria were inhibited, and 37% of bacterial isolates showed medium growth. The growth index ranged between 2.17% and 92.83%. In the presence of 10% salt, most

of the isolates were inhibited. Only 20% of the isolates showed mild growth, with a growth index between 2 and 22.64%.

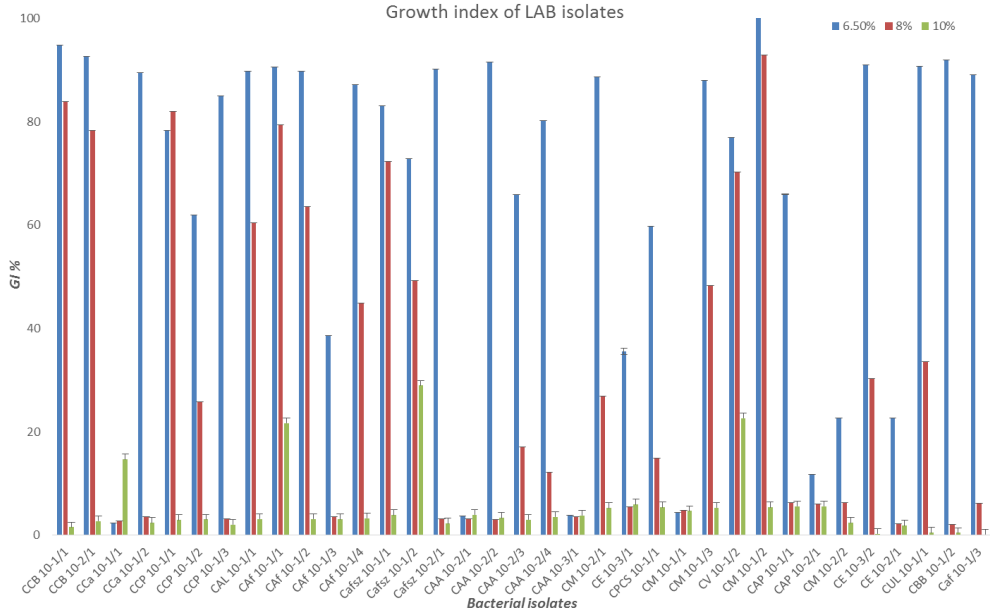


Figure 1: Growth index of LAB isolates in the presence of different salt concentrations

Further, the most beneficial strains were selected. The results of proteolytic activity for the most representative strains are presented in *Table 1*. All selected LAB showed proteolytic activity on the assayed medium agar. The clear zone diameter ranged in the interval of 13–17.33 mm. The benefits of proteolytic activity consist in texture and flavour development, contribution to water release through enhancing the drying process, and that via hydrolysis different peptides and volatile compounds are released, which serves as aroma precursor. It was demonstrated that different LAB, such as *Pediococcus pentosaceus* or *Lactobacillus curvatus*, originated from sausages hydrolysed sarcoplasmic proteins, generating different volatile compounds (*Cruzen et al.*, 2019). Different bioactive peptides derived from the peptidase and aminopeptidase activity of starter cultures are involved in functional food development (*Arihara*, 2014).

Table 1: Proteolytic activity of the isolates

<i>Bacterial isolates</i>	<i>Clear zone mm</i>
CCB 10-1/1	13.33 ± 1.5
CCB 10-2/1	14.33 ± 0.6
CCP 10-1/1	14.33 ± 1.5
CAL 10-1/1	13.2 ± 1
CAf 10-1/1	15.67 ± 3.2
CAf 10-1/2	16.33 ± 2.1
CAf 10-1/4	15 ± 1
Cafsz 10-1/1	15 ± 1
Cafsz 10-1/2	16 ± 2
CM 10-2/1	16.07 ± 2.1
CM 10-1/3	17.33 ± 3.06
CV 10-1/2	14.33 ± 1.53
CM 10-1/2	16.67 ± 3.1
CE 10-3/2	13 ± 1
CUL 10-1/1	14 ± 2

Table 2: Antimicrobial effect of bacterial isolates on Gram-positive bacteria

	<i>LAB isolates</i>	<i>Inhibition diameter (mm)</i>			
		<i>M. luteus</i>	<i>B. cereus</i>	<i>B. subtilis</i>	<i>S. aureus</i>
1	CCB 10-1/1	-	11.33±0.58	12.53±0.5	11.67±0.58
2	CCB 10-2/1	13.53±0.5	13.20±0.2	14.00±0.2	13.73±0.64
3	CCP 10-1/1	-	14.33±1.53	-	16.33±1.53
4	CAL 10-1/1	-	13.00±1	-	14.33±1.53
5	CAf 10-1/1	-	13.07±0.9	11.67±0.58	12.40±0.53
6	CAf 10-1/2	12.33±0.58	14.33±1.53	14.53±0.5	15.60±0.53
7	CAf 10-1/4	-	14.00±2	8.80±0.2	11.67±0.58
8	Cafsz 10-1/1	-	13.07±1.01	8.60±0.2	11.67±0.58
9	Cafsz 10-1/2	-	-	10.60±0.53	10.60±0.53
10	CM 10-2/1	-	-	13.87±0.31	13.53±0.5
11	CM 10-1/3	13.00±1	14.40±0.87	12.67±3.06	14.00±2
12	CV 10-1/2	-	11.00±1.0	-	10.07±0.31
13	CM 10-1/2	-	-	-	13.93±2.10
14	CE 10-3/2	-	11.60±0.53	11.67±0.58	15.00±1.0
15	CUL 10-1/1	20.13±1.28	13.00±1.0	13.33±1.15	16.00±2.0

The tested LAB isolates displayed different levels of bacteriostatic effects against the studied bacterial strains. Concerning Gram-positive bacteria, the most suppressive effect was detected in the case of *Staphylococcus aureus*, with a large inhibition zone of 16.63 mm in the case of the CCP 10-1/1 iso-



late. The characterized LAB isolates from sausages exerted greater inhibitory effect against *S. aureus* in contrast with bacterial isolates from home-made fermented foods (Ren *et al.*, 2018). In the case of the two *Bacillus* strains, the inhibition zone ranged between 11.40 and 14.40 (*B. cereus*) and between 8.6 and 14.53 (*B. subtilis*). The cell-free supernatant of CAf 10-1/2 inhibited with a larger inhibition zone both *Bacillus* bacterial strains. The most effective bacteriostatic effect was observed in the case of CAf 10-1/2. That isolate suppressed all the four tested Gram-positive bacteria with a large inhibition zone. LAB isolates exerted weak or no antibacterial activity against *Micrococcus luteus*. In concordance with our results, it has been shown that *Micrococcus luteus* was the most resistant bacteria (Evurani *et al.*, 2018) and was not inhibited by *Lactobacillus casei* and *Lb. brevis*.

The antibacterial effect against the tested Gram-negative bacteria was not promising. The LAB isolates exerted weak suppressive effect against *P. vulgaris*. Only five isolates displayed antibacterial effect against these tested bacteria. According to Aruna *et al.* (2016), *P. vulgaris* – tested by them – is resistant to bacteriocins. The largest inhibitory zones were observed in the case of CM 10-2/1. The inhibition zone was 13.00 mm in the case of *E. coli* and 14.33 in the case of *P. vulgaris*. The largest inhibition zone appears for *E.coli*: it was 14.47 mm.

Table 3: Antimicrobial effect of bacterial isolates on Gram-negative bacteria

	<i>LAB isolates</i>	<i>Inhibition diameter (mm)</i>	
		<i>E. coli</i>	<i>P. vulgaris</i>
1	CCB 10-1/1	12.33 ± 0.58	-
2	CCB 10-2/1	13.67 ± 0.42	-
3	CCP 10-1/1	13.33 ± 0.58	-
4	CAL 10-1/1	13.67 ± 1.15	-
5	CAf 10-1/1	10.00 ± 1.00	-
6	CAf 10-1/2	11.73 ± 0.64	11.67 ± 0.58
7	CAf 10-1/4	-	-
8	Cafsz 10-1/1	-	-
9	Cafsz 10-1/2	-	11.00 ± 1.0
10	CM 10-2/1	13.00 ± 1.00	14.33 ± 1.53
11	CM 10-1/3	-	-
12	CV 10-1/2	-	-
13	CM 10-1/2	10.60 ± 0.4	-
14	CE 10-3/2	14.47 ± 0.5	10.53 ± 0.64
15	CUL 10-1/1	11.00 ± 1.0	13.67 ± 0.58

Fermented meat microbiota is made up of bacterial strains of *Lactobacillus* and *Lactococcus* genus with bacteriocinogenic features and biopreservative role (Castilho *et al.*, 2019). Different strains from fermented meat are called bacteriocin producers: various *L. sakei* and *L. curvatus* strains, *Pediococcus acidilactici* strain MCH14, and *S. xyloso* strain SX S03/1 M/1/2 (Laranjo *et al.*, 2017).

LAB exhibits a broad-spectrum antimicrobial activity due to various mechanisms. The antagonistic mechanisms of LAB are diverse due to the altered gene expression and molecular structure of bacterial strains. The inhibitory activity of LAB strains is associated with their primary metabolites such as organic acids, alcohol, and carbon dioxide. LAB also produce different compounds with antagonistic effect. It was shown that formic and benzoic acids, hydrogen peroxide, diacetyl, acetoin, and bacteriocins have a role in the suppression of different microorganisms. For the suppression of Gram-negative bacteria, the organic acids and hydrogen peroxide are presumed to be responsible, while in the case of Gram-positive bacteria proteinaceous compounds are the potential causes. Reduced pH contributes to the inactivation of bacterial cell as well. The weak acids entering the cell may cause the acidification of the cytoplasm, leading to different disorders in the metabolism, structure, and function of the bacterial cell (Olorunjuwon *et al.*, 2013; Gao *et al.*, 2019). The mode of action of the nisin is dual, consisting of pore formation in the membrane or disrupting cell wall synthesis, leading to bacterial cell death (Jozala *et al.*, 2015).

From the isolated bacteria, 6 isolates showed CO<sub>2</sub> production. In the manufacture of different meat products, homofermentative strains are used (Cocconcelli, 2007).

Almost all tested LAB isolates showed varying levels of growth at pH 4 and pH 5 (data not shown). None of the tested bacterial strains showed activity at pH 3. According to Erkkilä and Petäjä (1999), most of the LAB isolated from meats has shown no decrease at pH 4. Our results also showed that the isolated strain was not able to grow at 3 pH. In the case of pH 4 and 5, isolates showed the highest growth. The highest growth was detected at pH 5, with the exception of two bacterial isolates: Cafs 10-1/1 and CCB 10-1/1.

Based on the results of the growth assay carried out at different temperatures, it can be said that almost all isolates grew at 20 °C (data not shown), with an OD value of 1.92. As for 30 and 37 °C, the same results were obtained. At these two temperatures, the highest OD value was 2.02. From the assayed bacterial isolates, two LAB isolates (CAf 10-1/3, CCA 10-1/1) did not show growth at 20 °C. In the case of one isolate, the growth was observed at 30 °C.

It was shown that pH and temperature influence the growth of lactic acid bacteria in different ways (*Adamberg et al.*, 2003).

Table 4: Minimal inhibitory concentration in the selected LAB isolates in the case of the tested antibiotics

<i>Bacterial isolates</i>	<i>Ampicillin</i>	<i>Gentamicin</i>	<i>Streptomycin</i>	<i>Kanamycin</i>	<i>Chloramphenicol</i>	<i>Erythromycin</i>
CCB 10-2/1	4	8	64	>128	8	2
CCP 10-1/1	4	8	32	>128	8	2
CAL 10-1/1	4	4	32	>128	8	2
CAf 10-1/2	4	64	>128	>128	8	>2
CAf 10-1/4	4	8	64	>128	8	2
Cafsz 10-1/2	4	64	>128	>128	16	>2
CM 10-2/1	4	16	64	>128	8	>2
CV 10-1/2	8	8	64	>128	8	2
CE 10-3/2	2	16	128	>128	8	2
CUL 10-1/1	4	16	64	>128	8	2

LAB resistance towards antibiotics represents an emerging and serious food safety concern. The World Health Organization suggests that LAB used in food industry should be free of antibiotic resistance (*Álvarez-Cisneros & Ponce-Alquicira*, 2018).

In the case of the most beneficial ten LAB isolates, the MICs of six antibiotics were determined. Based on the results, the differences in MIC values did not exceed one or two orders of dilution.

In the case of ampicillin, the MIC was 4 mg/L with the exception of two isolates, where it was 2 and 8 mg/L. In the case of gentamicin, the MIC values ranged between 4 and 64 mg/L, whereas the majority was equal with 8 mg/L. The MIC values for streptomycin ranged between 32 and 128 mg/L with two exceptions, where the MIC was higher than the tested concentration. Also, with kanamycin, the tested isolates showed growth at the maximum tested concentration (128 mg/L). With the exception of one isolate, MIC was 8 mg/L for chloramphenicol. In the case of erythromycin, the MIC was 2 mg/L for seven LAB isolates, while the other isolates were growing at this concentration. Based on the results, the majority of the tested bacterial isolates showed susceptibility to ampicillin with the exception of one isolate. In general, it

is considered that bacterial strains belonging to the *Lactobacillus* genus show sensitivity to the cell wall synthesis inhibiting penicillin and  $\beta$ -lactamase antibiotics (Gueimonde *et al.*, 2013).

In the case of gentamycin, two isolates were resistant. Three of the tested isolates were found resistant to streptomycin. All tested isolates showed resistance to kanamycin and chloramphenicol. In LAB, the resistance type towards kanamycin, gentamycin, and streptomycin is intrinsic. The absence of cytochrome-mediated electron transport contributes to the formation of resistance to the aminoglycosidic antibiotics such as kanamycin and streptomycin (Narayanan & Raghavan, 2019). Enzymatic deficiency or disabled enzymatic transport of aminoglycoside-modifying enzymes, such as N-acetyltransferases, O-phosphotransferases, and O-nucleotidyltransferases, results the above mentioned resistance (Álvarez-Cisneros & Ponce-Alquicira, 2018).

All ten selected isolates showed resistance to erythromycin.

In concordance with our results, previous studies reported different LAB strains isolated from fermented sausages with multiple antibiotic resistance. *L. plantarum*, *L. fermentum*, and *L. helveticus* showed resistance to kanamycin, tetracycline, erythromycin, chloramphenicol, and to other antibacterial compounds (Patel *et al.*, 2012). LAB strains originated from Portuguese and Italian sausages showed resistance towards streptomycin, gentamycin, chloramphenicol, erythromycin, etc. (Álvarez-Cisneros & Ponce-Alquicira, 2018). Narayanan & Raghavan (2019) suggested that the LAB of starter cultures show higher resistance against antibiotics and could be the store of resistance genes. In some cases, these genes are not expressed but can be transferred to other bacterial strains. The safety characterization of LAB is essential to avoid any risk of infection by using them (Borriello *et al.*, 2003, Doron and Snyderman, 2015).

## 4 Conclusions

Based on the results, it can be concluded that the lactic acid bacterial isolates originating from the autochthonous microbial ecology of fermented sausages showed positive technological properties. Regarding the assayed technological characteristics of isolates, these strains can be involved in the fermentation of sausages, playing an important role in the development of sensorial characteristics, texture, aroma formation, and the inhibition of different pathogenic microorganisms. For a successful and safe application of these isolates, further studies are needed to assess the safety of the strains to avoid bacteremia. Also,

it is necessary to identify the isolates, determine their virulence factors, and the biogenic amine production capacity.

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