

LISTERIA SPP. ASSOCIATED TO DIFFERENT LEVELS OF AUTOCHTHONOUS MICROBIOTA IN MEAT, MEAT PRODUCTS AND PROCESSING PLANTS

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Submitted: September 15, 2006; Returned to authors for corrections: July 04, 2007; Approved: September 20, 2007.

ABSTRACT

High levels of microbial contamination, commonly found in animal origin foods and food processing environments, are able to hinder the growth of pathogens in these products and interfere in the results of laboratory analyses for detection of these pathogens. With the aim of verifying the possible interference of the autochthonous microbiota encountered in meat and meat products and processing plants over the presence of *Listeria* spp., 443 samples, collected from 11 meat retail establishments, were submitted to microbiological analysis to determine the levels of mesophilic aerobes, total coliforms and *Escherichia coli* and the presence of *Listeria* spp., according to the methodology proposed by the USDA. The results did not show evident interference of the autochthonous microbiota over *Listeria* spp., once the genus was detected even in the meat, meat products and environmental samples with high levels of contamination by mesophilic aerobes and coliforms.

Key words: meat; *Listeria* spp., autochthonous microbiota

INTRODUCTION

Listeria monocytogenes emergence as a foodborne pathogen dates from 1980, with the occurrence of many outbreaks and sporadic cases of listeriosis associated with the consumption of contaminated foods (3,7,20,28). The invasive form of this pathogen is considered too risky especially to newborns, elderly and immune-compromised individuals, who may present septicemia, encephalitis, meningitis and abortion, and high levels of mortality (11,12,29). Among *Listeria* species, only *L. monocytogenes* is recognized as a human pathogen (7). Most foods are tested for the presence of non-pathogenic *Listeria* spp only, as an indication of the possible presence of *L. monocytogenes* (5,6,26,30).

The occurrence of *Listeria* spp. and *L. monocytogenes* in foods is highly variable worldwide (7,9,24). In many regions the reported occurrence is low, especially in foods with high levels

of microbial contaminants (15,16,24). According to Jay (15,16), levels of contaminants higher than 10⁶ CFU/g create unsatisfactory conditions for the growth of pathogens in foods. The dynamic of the interference depends on the presence of specific antagonistic microorganisms and also on the ability of producing inhibitory substances.

In meat and meat products, the autochthonous microbiota, which includes the natural microbiota of beef and the contaminants determined by the hygienic procedures during slaughter and processing (1,10,13,16,23,33), may include strains capable of producing a variety of antimicrobial compounds that may impair the survival or inhibit the growth of *L. monocytogenes* (2,4,18,22). Moreover, the contaminants can also interfere on the effectiveness of the laboratory isolation and detection methods, leading to erroneous results about the presence of *L. monocytogenes* in the product (31).

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Several methodologies are available for isolation of *Listeria* spp. in foods. All methods employ enrichment phases at different temperatures using different supplements, selective agents and incubation conditions (6,19,27,31). Selective enrichment steps are necessary to inhibit the growth of the competitive microbiota and optimize the multiplication of *Listeria* spp., leading to their amplification to required level for proper detection. As an example, the University of Vermont *Listeria* Enrichment broth (UVM), recommended by the United States Department of Agriculture (32), is an enrichment medium which contains nalidixic acid and acriflavine as supplements to inhibit Gram negative and Gram positive microorganisms, respectively. This culture medium is considered the most suitable for testing foods where high levels of competitive microbiota are expected (25,30).

In beef, despite the occurrence of antagonistic microorganisms in the autochthonous microbiota (2,4,16), there is no evidence of the direct interference of these microbiota on *Listeria* spp. isolation procedures (6,21). The aim of this study was to verify the influence of the autochthonous microbiota in meat and meat products and in the meat processing environment on the detection of *Listeria* spp., using the detection protocol recommended by USDA (32).

MATERIAL AND METHODS

Samples and dilutions

Eleven meat retail establishments (one slaughterhouse and ten butcher shops) located in Londrina region, north of Paraná State, Brazil, inspected by Brazilian Agriculture (the slaughterhouse) or Health (butcher shops) Authorities, were randomly selected. In these establishments, a total of 443 samples were taken (Table 1), comprising 148 surface samples collected from the equipments (knives, mincers, boxes, etc), 65 samples collected from the environment (drains, floors, etc) and 230 meat and meat product samples. Surface samples were obtained with sterile swabs and sterile templates to limit the specific area shown in Table 1. Swabs were transferred to flasks containing 45 mL of 0.1% buffered peptone water (Oxoid, Basingstoke, UK). Meat samples were collected in sterile bags. The samples were kept under refrigeration during the transport to the laboratory. For testing, 10 g of the collected samples were weighted and transferred to 90 mL of 0.1% buffered peptone water (Oxoid, Basingstoke, UK), and homogenized with a stomacher. The homogenates were submitted to serial decimal dilutions using sterile NaCl 0.85%.

Table 1. Points of sampling, number of samples and surface sampling area.

Samples	Sampled area/quantity cm ² or g	Number of samples		
		Butcher shops n	Slaughterhouse n	Total n
Equipments				
Meat knives	30 cm ²	15	6	21
Meat mincers	30 cm ²	9	1	10
Meat mixers	30 cm ²	3	1	4
Meat saws	30 cm ²	10	-	10
Meat scales	30 cm ²	9	-	9
Meat tenderizers	30 cm ²	8	-	8
Plastic boxes	30 cm ²	26	2	28
Refrigerated trucks	60 cm ²	4	-	4
Sausage fillers	30 cm ²	2	1	3
Stainless steel boxes	30 cm ²	10	-	10
Stainless steel tables	60 cm ²	20	1	21
Steel hooks	Surface area	7	3	10
Wood frames	30 cm ²	10	-	10
Environments				
Drains	Surface area	18	7	25
Floors	60 cm ²	9	1	10
Meat cabinets	60 cm ²	11	-	11
Reception platform	60 cm ²	-	5	5
Refrigeration systems	60 cm ²	4	-	4
Walls	60 cm ²	10	-	10
Products				
Beef cuts	50 cm ²	51	1	52
Bovine carcasses	50 cm ²	142	9	151
Cooked sausages	5 g	-	2	2
Fresh sausages	5 g	10	-	10
Ground beef	5 g	12	-	12
Swine carcasses	50 cm ²	-	3	3
Total		400	43	443

Microbiological analysis

The samples were tested for the presence of *Listeria* spp. according to the protocol proposed by USDA (32). In the pre-enrichment step, 5.0 mL of the peptone water containing the swab or 5.0 g of product were transferred to 45.0 mL of University of Vermont *Listeria* Enrichment broth (UVM) (Oxoid, Basingstoke, UK), and incubated at 30°C for 24h. One mL of the UVM broth was transferred to tubes containing 9 mL of Fraser broth (Oxoid, Basingstoke, UK), and incubated at 35°C for 24h. Samples that presented darkening were streaked on Modified

Oxford agar (BD Diagnostic Systems, Franklin Lakes, NJ, USA) and incubated at 35°C for 24h. When *Listeria* typical colonies were present (small, grey to black, surrounded by blackening halo), they were submitted to identification by API-Listeria (bioMérieux S.A., Marcy l'Etoile, France). The final result was expressed as presence or absence of *Listeria* spp. in the sampled food or area (Table 1).

The autochthonous microbiota was determined by enumerating the following groups of microorganisms: mesophilic aerobes (MA), total coliforms (TC) and *Escherichia coli* (EC). From each sample, two decimal dilutions were selected and plated in Petrifilm™ plates (Petrifilm™ AC for MA, and Petrifilm™ EC for TC and EC) and incubated at 35°C for 24-48h. After incubation, the colonies formed in the plated area were enumerated and expressed as Colony Forming Units per gram or cm² (CFU/g or cm²).

Data analysis

For data analysis, the counts of MA, TC and EC were converted in log₁₀. The numbers of *Listeria* spp positive samples according to the source (equipments, environments or products) and mean counts of MA, TC or EC were compared by Tukey Honest Significant Difference for unequal n, at a level of significance of 0.05. The frequencies of positive and negative results for *Listeria* spp. were also calculated considering three levels of contamination by MA, TC and EC. All analyses were processed using the software Statistica 6.0 (StatSoft, Inc., Tulsa, OK, USA).

RESULTS AND DISCUSSION

One hundred and sixty seven samples were positive for *Listeria* spp: 76 samples from equipments, 23 from environments and 68 from products. The prevalence of species was as follows: *L. innocua* (131), *L. monocytogenes* (21), *L. welshimeri* (12), *L. seeligeri* (02) and *L. grayi* (01). Fig. 1 shows the positive and negative results for *Listeria* spp in the samples, the mean counts of MA, TC and EC, and the significance of the correlations between presence of *Listeria* spp and the counts of MA, TC and EC. The frequencies of positive or negative samples for *Listeria* spp. according to the levels of contamination by MA, TC and EC are shown in Figs. 2, 3 and 4, respectively.

In general, samples with positive results for *Listeria* spp. presented higher mean counts of indicators microorganisms than the *Listeria* spp. negative samples. In equipments, 44.7% of the *Listeria* spp. positive samples presented MA counts over 5 log CFU/cm², while in the environment *Listeria* spp. positive samples, this percentage was 73.9% and in products, 41.2%. In the average, 47.3% of the *Listeria* spp. positive samples presented MA counts higher than 5 log CFU/cm², while for *Listeria* spp. negative samples this percentage was 24.3% (Fig. 2). The same situation was observed for total coliform

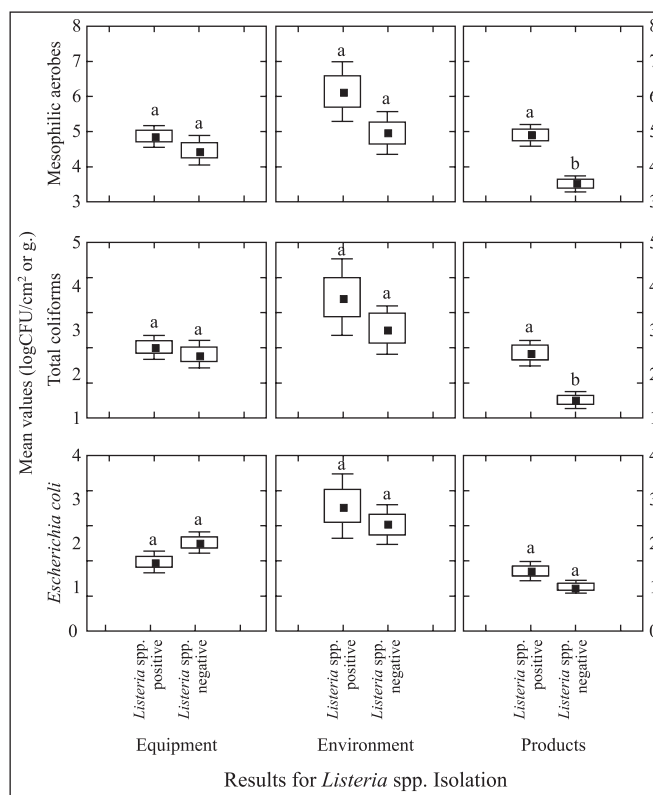


Figure 1. Counts of mesophilic aerobes, total coliforms and *Escherichia coli* and presence of *Listeria* spp. in samples collected from meat processing plants (equipment, environments and products). Distinct letters in each graph indicate significant difference ($P < 0.05$).

counts (Fig. 3) and for *E. coli* (Fig. 4). All these differences were significant ($P < 0.05$). However, it must be noted that for *E. coli* counts, the majority of results were lower than the limit of the detection method (2 log CFU per gram or cm²), so the significance of the differences between *Listeria* spp. positive and negative results could not be calculated properly.

These results suggest that the autochthonous microbiota in meat products and meat processing environment did not affect the survival or growth of *Listeria* spp. in them. The possible interferences of the autochthonous food microbiota over *Listeria* spp. and *L. monocytogenes* are widely variable due to several factors, including microbiological composition, preserving temperatures and inhibitory substances production. A common microorganism present in meat and meat processing plants, *Pseudomonas fluorescens*, was already described as improving (22), inhibiting (4) and causing no effect (8) on *L. monocytogenes* growth. Those distinct behaviors of food accompanying microbiota were also observed by Buchanan and Bagi (2).

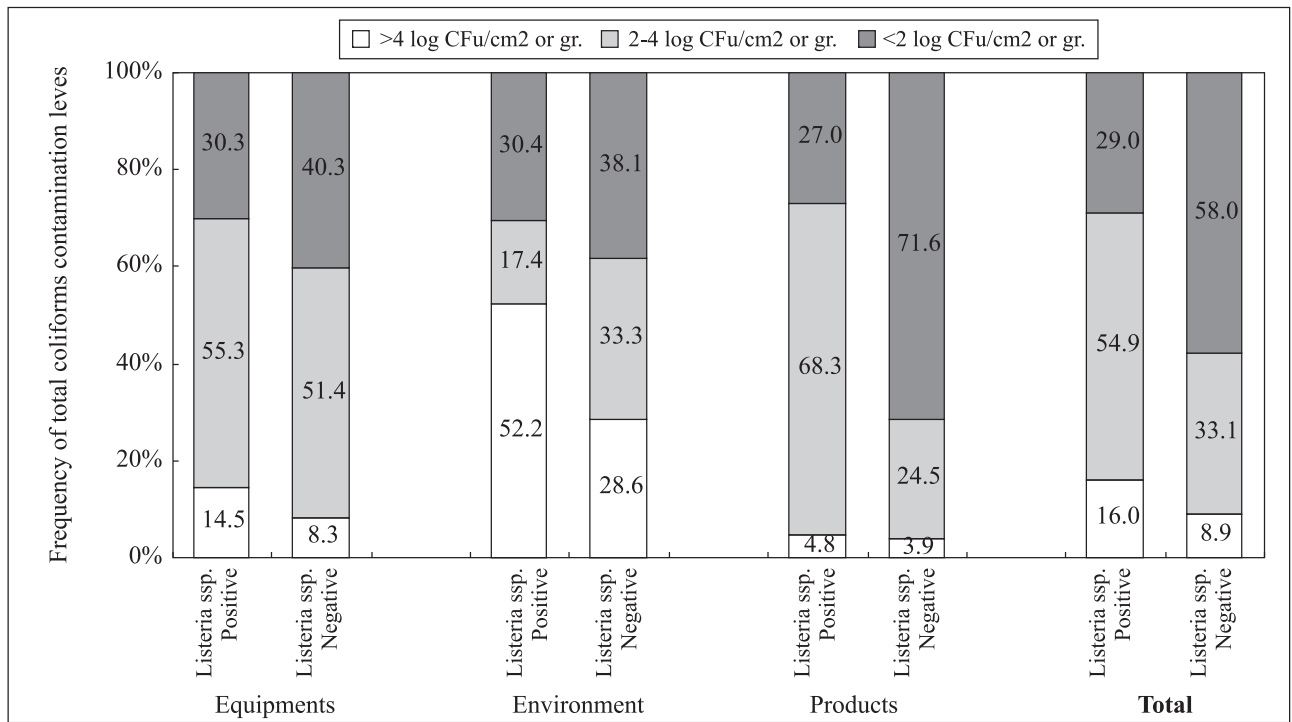


Figure 2. Occurrence of *Listeria* spp in meat products and equipment and environmental samples collected in meat processing plants located in Londrina region, Paraná State, PR, Brazil, according to the contamination level by mesophilic aerobes.

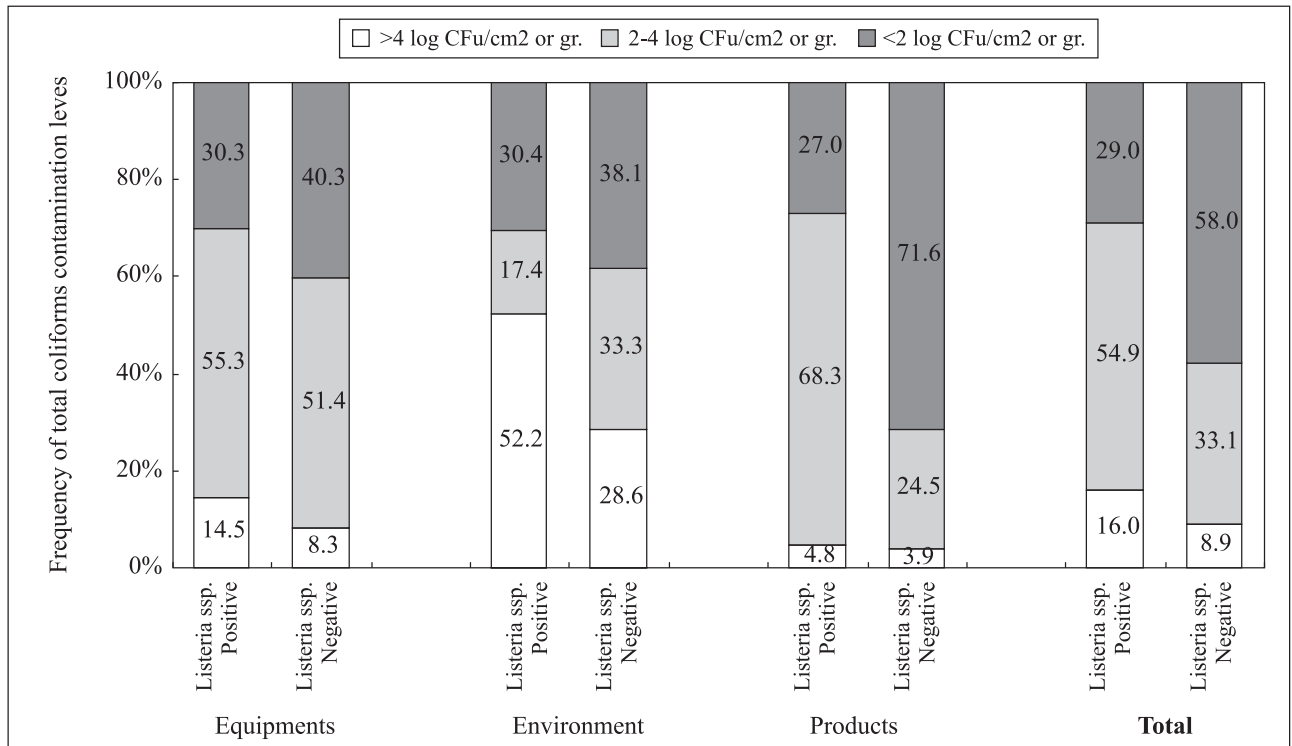


Figure 3. Occurrence of *Listeria* spp in meat products and equipment and environmental samples collected in meat processing plants located in Londrina region, Paraná State, PR, Brazil, according to the contamination level by total coliforms.

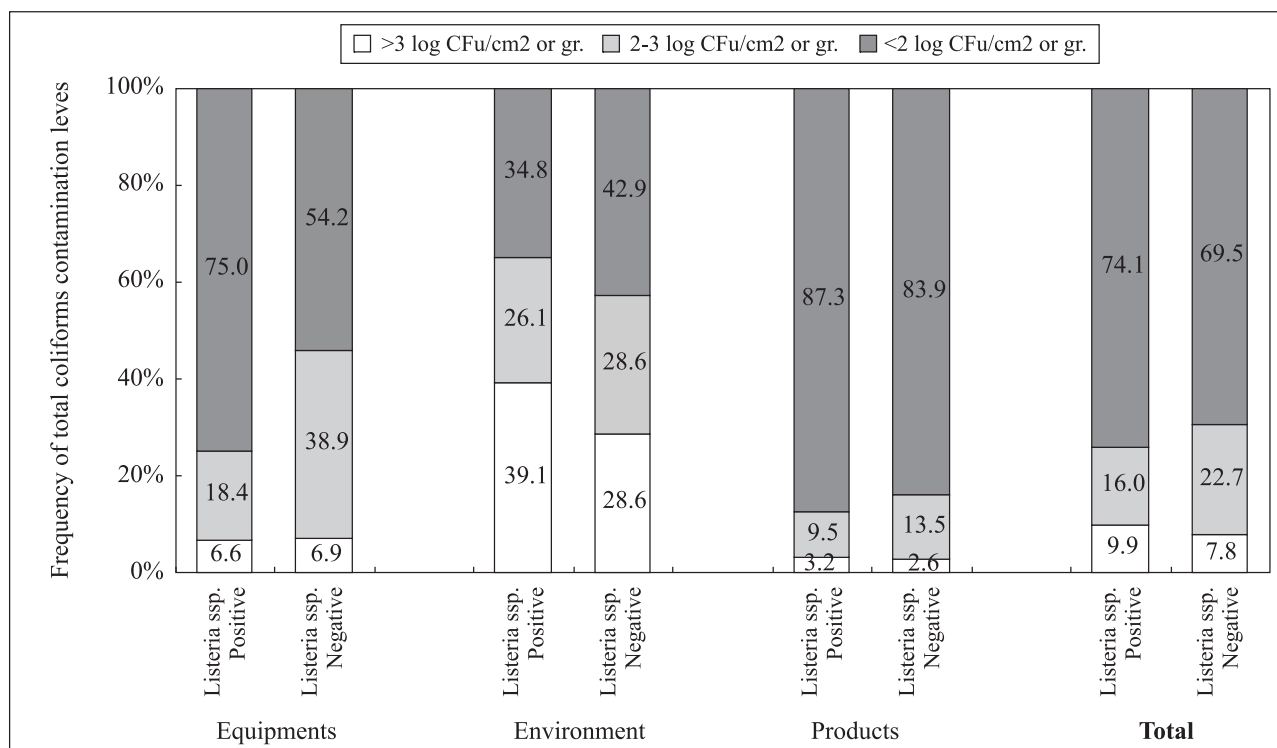


Figure 4. Occurrence of *Listeria* spp in meat products and equipment and environmental samples collected in meat processing plants located in Londrina region, Paraná State, PR, Brazil, according to the contamination level by *Escherichia coli*.

According to Jay (15,16), fresh meats with MA counts between 10^4 and 10^6 CFU/g are probably not vehicles of foodborne pathogens, due the autochthonous presence of inhibitory microorganisms. However, the results of the present study indicate that even at these levels the isolation of *Listeria* spp., including the pathogenic specie *L. monocytogenes*, is possible.

The utilization of UVM broth in the enrichment step may explain the absence of interference of the autochthonous microbiota of the meat products on the isolation of *Listeria* spp. This culture media is known as the most suitable to isolate *Listeria* spp. in foods with high levels of microbiological contamination, due to its effective inhibitory activity by the added supplements (19,30). Some authors even suggest that UVM promotes a better recovery of *L. innocua* than *L. monocytogenes* (5,26), especially due to a higher growth velocity of the first (21). However, Duffy *et al.* (6) demonstrated that this culture media did not promote significant differences in the recovery of both *Listeria* species.

In food processing surfaces and utensils, a contamination level of 10^4 CFU/cm² is considered enough to begin biofilm formation (13). In this study, several samples presented counts higher than 10^4 CFU/cm², suggesting the possibility of biofilm formation on equipment surfaces, and even in products. The importance of these findings is relevant because these contamination levels were found in several *Listeria* spp. positive

samples. The literature reports studies indicating that *L. monocytogenes* is able to grow and some autochthonous strains, such as *Flavobacterium* spp., can even improve its development (17). *Listeria* spp. are able to produce glycocalyx in stainless steel surfaces within 48 h, promoting self protection against environmental microorganisms that can produce inhibitory substances (14,33).

In conclusion, the results of the present study suggest that the autochthonous microbiota of meat and meat processing plants have very little, if any, influence on the isolation of *Listeria* spp., even when the counts of hygiene indicator microorganisms surpass 5 log CFU/cm² or g. The official methodology for the isolation of *Listeria* spp from meat and equipment surfaces was shown to be appropriate, with no interference of the autochthonous microbiota.

RESUMO

***Listeria* spp. associado a diferentes níveis da microbiota autóctone de carne, produtos cárneos e plantas de processamento**

Altos níveis de contaminação microbiana, usualmente encontrados em alimentos de origem animal e nos ambientes de

processamento, podem inibir a multiplicação de microrganismos patogênicos nesses produtos e interferir nos resultados das análises laboratoriais para o isolamento desses patógenos. Com o objetivo de verificar as possíveis interferências da microbiota autóctone encontrada na carne, produtos cárneos e plantas de processamento sobre a presença de *Listeria* spp., 443 amostras, coletadas em 11 estabelecimentos processadores, foram submetidas a análises microbiológicas para determinação dos níveis de contaminação por aeróbios mesófilos, coliformes totais e *Escherichia coli* e para verificação da presença de *Listeria* spp., de acordo com a metodologia proposta pelo USDA. Os resultados obtidos não mostraram uma interferência evidente da microbiota autóctone sobre *Listeria* spp., uma vez que esse gênero foi detectado mesmo nas amostras de carne e produtos cárneos e amostras ambientais e de superfície de equipamentos que apresentaram altos níveis de contaminação por aeróbios mesófilos e coliformes.

Palavras-chave: carne, *Listeria* spp.; microbiota autóctone

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