

Ciência Rural

ISSN: 0103-8478

cienciarural@mail.ufsm.br

Universidade Federal de Santa Maria Brasil

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Ciência Rural, vol. 46, núm. 12, diciembre, 2016, pp. 2257-2263

Universidade Federal de Santa Maria
Santa Maria, Brasil

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ISSN 1678-4596 FOOD TECHNOLOGY

Microbial quality of industrial and retail market grated parmesan cheese in the State of São Paulo, Brazil

Comparação da qualidade microbiológica de queijos tipo parmesão ralados industrialmente e no comércio varejista no Estado de São Paulo, Brasil

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ABSTRACT

With the objective to evaluate the hygienic and sanitary conditions of grated parmesan cheese acquired from the retail business, total 120 cheese samples were acquired: 60 of which were obtained from four different brands of cheese that were grated in factories, and the other 60 samples were obtained from another four brands of cheese that are normally acquired by retailers (supermarkets) in blocks and grated at the moment of sale. The population of heterotrophic mesophilic microorganisms ranged from 1.2×10^3 to 1.1×10^7 colony-forming units (CFU)·g⁻¹. All samples analyzed contained Staphylococcus spp. with populations varying from 1.2×10³ to 8.7×10⁶CFU·g⁻¹, from which 60.0% were classified as coagulase-positive Staphylococcus and 52.5% of the samples possessed populations above the permissible limit set by legislation. Staphylococcus aureus was detected in 57.5% of the samples. Population of molds and yeasts varied from <10 to 1.8×106CFU·g¹. Salmonella spp. was not isolated in this research. A difference was observed between the parmesan cheese grated in factories and that grated in supermarkets, where the former presented better microbiological quality than the latter. Thus, procedures must be proposed to minimize the presence of pathogenic agents reported in grated parmesan cheeses evaluated in the present study because of the public health risk associated with food bacterial contamination.

Key Words: dairy, food microbiology, public health, Staphylococcus aureus.

RESUMO

Com o objetivo de avaliar as condições higiênicas e sanitárias de queijos tipo parmesão ralados, vendidos no comércio varejista, foram realizadas análises microbiológicas em 120 amostras, sendo 60 destas de quatro marcas comerciais, as quais foram raladas industrialmente e outras 60 de quatro marcas, cujos queijos foram adquiridos pelos comércios varejistas em grandes peças e ralados no ato da venda. As populações de microrganismos heterotróficos mesófilos variaram de 1,2×10³ a 1,1×107UFC.g¹. Todas as amostras analisadas apresentaram Staphylococcus spp., com populações que variaram de 1,2×10³ a 8,7×106UFC.g¹, sendo que 60,0% dessas apresentaram **Staphylococcus** coagulase positivo e 52,5% delas com populações acima do limite estabelecido pela legislação. A presença de Staphylococcus aureus foi identificada em 57,5% das amostras. As populações de bolores e leveduras apresentaram variação de <10 a 1,8×106UFC.g1. Salmonella spp. não foi isolada no presente trabalho. Houve diferença entre os queijos ralados industrialmente e os queijos ralados nos supermercados, sendo que o primeiro grupo apresentou resultados de qualidade microbiológicas melhores do que o segundo grupo. Assim, medidas visando minimizar as populações de agentes contaminantes e deteriorantes encontradas nos queijos parmesão analisados no presente trabalho devem ser realizadas, visto que estas podem representar um risco à saúde pública.

Palavras-chave: laticínio, microbiologia de alimentos, saúde pública, Staphylococcus aureus.

INTRODUCTION

Cheese is a nutritive food, which can facilitate the multiplication of pathogenic microorganisms that cause foodborne diseases (MARINHEIRO et al., 2015). Food contamination can occur because of several factors, such as poor microbiological quality

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of raw material, inadequately sanitized facilities and equipment, and improper manipulation and storage of products (MELLO & ARMACHUK, 2013).

Among several types of cheese commonly sold in Brazil, parmesan cheese is one of the most consumed and is usually grated at the time of sale. Parmesan cheese is a hard, semi-fat, cooked, low-moisture cheese; it has a tough and granular texture with a thick, smooth, and straw-yellow crust (JASTER et al., 2014).

However, this cheese can pose a risk to public health due to several processing steps that are required before the final product is obtained. Consequently, production under improper conditions can have serious consequences on the population, becoming a public health threat (GAULIN et al., 2003).

Therefore, assessing the presence of certain microorganisms, such as coagulase-positive Staphylococci, *Salmonella* spp., mold, yeast, and enterobacteria, in parmesan cheese is of utmost importance because of the danger it poses to human health when consumed (SEIXAS, 2009).

Another important issue is the contamination of these products at the point-of-sale, mainly due to poorly sanitized equipment and utensils, which can contaminate food, making it unsafe for consumption. Lack of proper sanitization of equipment and utensils is responsible for up to 16% of foodborne disease outbreaks (ABREU et al., 2010a); thereby, highlighting that some markets and retails markets frequently have poor sanitization programs (SILVA, 2011; SOTO et al., 2009).

To assure the identity and safety of commercially sold grated cheeses, the Brazilian Ministry of Agriculture, Livestock and Food Supply created ordinance n. 357 on September 4, 1997 (BRASIL, 1997), which established the Technical Regulation of Identity and Quality of Grated Cheese. This ordinance sets a maximum limit of 10³ colonyforming units (CFU)·g-1 of coagulase-positive Staphylococcus and the absence of Salmonella spp. in 25g of sample and a maximum mold and yeast count of 5×10³CFU·g⁻¹. However, this ordinance only applies to industrially produced grated cheese; cheese grated at retail markets falls under RDC n. 12 (BRASIL, 2001) regulation act, which states that for low-humidity cheeses, a maximum limit of 10³CFU·g⁻¹ for coagulase-positive *Staphylococcus* is allowed with the absence of Salmonella spp. in 25g of sample.

Because of the lack of information regarding the microbial integrity of grated parmesan cheese in the industry and at the point-of-sale in retail markets, this study focused on assessing and comparing the microbiological quality of currently available commercialized parmesan cheeses in the municipality of Ribeirão Preto, state of São Paulo.

MATERIALS AND METHODS

A set of 60 samples of parmesan cheese grated in industries under federal inspection were evaluated. The samples were collected from four different commercial brand (brands A, B, C, and D); 15 samples of each brand from the same batch. Another 60 samples of parmesan cheese of four different brands (E, F, G. and H) grated in retail markets were also collected with 15 samples per brand from the same batch. All 120 samples were obtained during 2013 in the municipality of Ribeirão Preto, state of São Paulo, Brazil. Markets purchased the parmesan cheese in large pieces that were kept refrigerated and were grated in the presence of the consumer during sale. The industry samples of grated parmesan cheese were maintained at room temperature (as sold) until laboratory analyses were performed; whereas, retail market samples were transported refrigerated to the laboratory in boxes containing recyclable ice (as sold).

A 25g amount of each sample of grated cheese from its original packing was placed in sterile flasks containing 225mL of 0.1% buffered peptone water and serially diluted till 10-4. The methodology described by the American Public Health Association (APHA, 2001) was adopted to assess the most probable number (MPN) of total coliforms · g-1, MPN of thermotolerant coliforms and Escherichia coli per gram, standard count of aerobic or facultative heterotrophic mesophilic microorganisms, presence of Salmonella spp., and count of Staphylococcus spp., positive-coagulase Staphylococcus, and Staphylococcus aureus. Mesophilic microorganisms were counted using standard plate count agar (SPC; Oxoid), and plates were incubated at 35°C for 48h.

To analyze the presence of *Salmonella* spp., initial samples were pre-enriched and incubated at 37°C for 24h. Selenite cystine broth (Difco, Detroit, USA) and Rappaport–Vassiliadis broth (Difco, Detroit, USA) were used for selective enrichment. Subsequently, 0.1mL of a 0.4% novobiocin solution (INLAB, São Paulo, Brazil) was added to the two broths. Selective plating was performed on brilliant green and MacConkey agars (Difco, Detroit, USA).

Counting of *Staphylococcus* spp. was performed using Petri dishes containing Baird–Parker agar (BPA; Difco Laboratories) supplemented

with egg yolk-potassium tellurite, and the plates were incubated at 35°C for 48h. A coagulase test was performed using the isolates from brain–heart infusion broth and rabbit plasma to assess the presence of coagulase-positive strains. *S. aureus* biochemical characterization was performed according to MacFaddin (1976).

The Kruskal-Wallis (P<0.05) statistical test was used to compare the mean count of mesophilic bacteria, *Staphylococcus* spp., positive coagulase *Staphylococcus*, *S. aureus*, mold, and yeast (THEODORSSON-NORHEIM, 1986). Whenever significant differences were observed, the Dunn's *post hoc* test for multiple comparisons was performed to detect the differences reported. All statistical analyses were performed using SAS statistical software 9.1.3 release (SAS Corporation, Cary, NC).

RESULTS AND DISCUSSION

Total coliforms, thermotolerant coliforms, and *E. coli* were detected in the industrially grated (IG) cheese at 10.0% (six), 1.6% (one), and 1.6% (one) of samples, respectively. The maximum populations observed in each bacterial group were 210MPN·g⁻¹, 70MPN·g⁻¹, and 70MPN·g⁻¹, respectively. Analysis of the parmesan cheese grated at the time of sale (GTS) revealed 35.0% (21), 16.6% (10), and 16.6% (10) contaminated samples, with a maximum population of 500MPN·g⁻¹, 70MPN·g⁻¹, and 23MPN·g⁻¹, respectively.

Although samples contained detectable levels of microorganisms, the amounts were below the limits established by the RDC n. 12 legislation (BRAZIL, 2001), thereby not hindering the sales of these products. One possible reason for the low count of *E. coli* is that the bacteria require more water activity than is present in parmesan cheese (ABREU et al., 2010a).

Salmonella spp. was not isolated from any of the samples. The arithmetic means and standard deviation of the mesophilic, **Staphylococcus** spp., positive-coagulase **Staphylococcus**, **S. aureus**, mold, and yeast populations are presented in tables 1 and 2.

A significant statistical difference (Dunn's test, P<0.05) was observed between the heterotrophic mesophilic microorganism population of the IG group and of the GTS group. The latter group presented a higher population of microorganisms. Detecting heterotrophic mesophilic microorganisms in food can provide information regarding the hygienic and sanitary characteristics of the product. High amounts of microorganisms suggested contamination of the

raw product, inappropriate processing conditions, or inadequate storage. Thus, the presence of these microorganisms poses a risk to the consumer's health and contributes to the deterioration of the product (MONTANHINI et al., 2013). High amounts of these microorganisms also indicated a greater possibility of pathogens being present in the food as most pathogens are mesophilic bacteria (MADIC et al., 2011).

Despite the fact that Brazilian law has no set limit for the amount of heterotrophic mesophilic microorganisms allowed in grated parmesan cheese, some groups have suggested that cheese with microbial populations larger than 10⁶CFU·g⁻¹ should not be consumed (ICMSF, 1981) due to the risk of foodborne diseases (JAY, 2005). Mesophilic bacterial counts higher than 10⁶CFU·g⁻¹ were observed in grated cheese samples from retail markets as shown in table 1.

Although errors while processing food products frequently occur, most food contamination is usually due to incorrect manipulation of products in markets (DAMER, 2015). This is in accordance with the research findings, in which 88.3% of GTS samples had bacterial counts higher than 10⁶CFU·g⁻¹; whereas, IG samples had lower counts, confirming the higher risk associated with consuming GTS cheese.

Considering that all industrial processes were performed under the same hygienic and sanitary rigor as all analyzed brands were from industries under federal sanitary inspection, the differences in the microorganism population found between the IG and GTS groups is probably due to inadequate manipulation of the blocks of parmesan cheese or even poor cleaning of the grating machines at markets.

The *Staphylococcus* population ranged from 1.2×10^3 to $1.8 \times 10^4 \text{CFU} \cdot \text{g}^{-1}$ in the IG samples and from 1.2×10^5 to $8.7 \times 10^6 \text{CFU} \cdot \text{g}^{-1}$ in the GTS samples. These results were statistically different when analyzed using the Dunn's test (P<0.05); thus, confirming that the IG samples had a lower microbial population than that in the GTS samples. The assessment of *Staphylococcus* in food is critical as over 20 its species can cause several diseases in humans (KANDARIYA et al., 2014). Moreover, *Staphylococcus* is one of the most common causes of foodborne illnesses (SCALLAN et al., 2011) and is the main source of contamination in inadequately handled food (ARGUDÍN et al., 2012).

In the USA, sources have reported that 25.3%-48.3% of staphylococcal food poisoning

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Table 1 - Arithmetic means and standard deviation of heterotrophic mesophilic bacteria, mold, and yeast found in each brand of industrially grated (IG) and grated at the time of sale (GTS) parmesan cheese collected from markets at Ribeirão Preto, São Paulo.

Brand	Arithmetic means and standard deviation (CFU×g¹)		
	Mesophilic	Molds and Yeast	
A (IG)	$9.8 \times 10^{4bc} \pm 1.3 \times 10^4$	$3.3 \times 10^{1c} \pm 6.2 \times 10^{1}$	
3 (IG)	$1.8 \times 10^{4cd} \pm 3.5 \times 10^{3}$	$1.3 \times 10^{3bc} \pm 1.0 \times 10^{3}$	
C (IG)	$1.4 \times 10^{4d} \pm 6.9 \times 10^{3}$	$5.4 \times 10^{2bc} \pm 6.3 \times 10^2$	
O (IG)	$3.6 \times 10^{4d} \pm 1.7 \times 10^4$	$6.6 \times 10^{2bc} \pm 6.2 \times 10^2$	
E (GTS)	$5.3 \times 10^{6ab} \pm 3.7 \times 10^{6}$	$4.6 \times 10^{5a} \pm 6.2 \times 10^{5}$	
F(GTS)	$6.8 \times 10^{6a} \pm 3.9 \times 10^{6}$	$8.2 \times 10^{4ab} \pm 2.1 \times 10^{5}$	
G (GTS)	$5.1 \times 10^{6ab} \pm 3.5 \times 10^{6}$	$2.3 \times 10^{5a} \pm 2.5 \times 10^{5}$	
H (GTS)	$5.0 \times 10^{6ab} \pm 3.1 \times 10^{6}$	$3.1 \times 10^{5a} \pm 3.7 \times 10^{5}$	

Means followed by the same letters in the column do not differ according to the Dunn's test (5% significance level).

outbreaks were due to contamination by food handlers (BEAN et al., 1997); thus, confirming the impact professionals have in spreading *Staphylococcus*. This suggested an inadequate handling of cheese at the sale markets or during grating as indicated in the present research.

Although storage temperature could have influenced the results, all samples were maintained at the same temperature. Therefore, storage temperature is likely not responsible for the differences observed in the microbial populations between groups. At present, there is no ordinance that sets a maximum limit for *Staphylococcus* count allowed in parmesan cheese. However, the RDC n. 12 (BRAZIL, 2001) sets a limit for *Staphylococcus* that produce coagulase enzyme; *S. aureus* is the main coagulase-producing species. According to this ordinance, the maximum count allowed for coagulase-positive *Staphylococcus* is $10^3 \text{CFU} \cdot \text{g}^{-1}$.

In our studies, the coagulase-positive *Staphylococcus* count was <10 to $5.1 \times 10^3 \text{CFU} \cdot \text{g}^{-1}$ in the IG samples and <10 to $3.6 \times 10^6 \text{CFU} \cdot \text{g}^{-1}$ in the GTS samples. Samples in the latter group had higher count of *Staphylococcus* coagulase-positive bacteria; out of 60 samples, 88.3% had counts higher than the limits allowed by the legislation, whereas only 16.6% of the IG group had higher counts. When compared using the Dunn's test (P>0.05), no significant difference in microbial count was observed among the samples in the GTS group.

The highest count of *Staphylococcus* coagulase-positive bacteria observed in the IG samples was $5.1 \times 10^3 \text{CFU} \cdot \text{g}^{-1}$. When compared using the Dunn's test (P<0.05), no significant differences were observed within the GTS group. Using the Dunn's test (P<0.05), the B and D (IG) and E (GTS) brands were not statistically different compared with the IG and GTS groups. Overall, the

Table 2 - Arithmetic means and standard deviation of *Staphylococcus* spp., positive-coagulase *Staphylococcus*, and *S. aureus* reported in each brand of industrially grated (IG) and grated at the time of sale (GTS) parmesan cheese collected from markets at Ribeirão Preto, São Paulo.

Brand	Staphylococcus spp.	Positive-coagulase Staphylococcus	S. aureus
A (IG)	$4.7 \times 10^{3a} \pm 2.4 \times 10^{3}$	$2.6 \times 10^{2c} \pm 4.3 \times 10^2$	$2.6 \times 10^{2a} \pm 4.3 \times 10^{2}$
B (IG)	$7.4 \times 10^{3a} \pm 5.1 \times 10^{3}$	$9.6 \times 10^{2bc} \pm 1.4 \times 10^{3}$	$9.6 \times 10^{2a} \pm 1.4 \times 10^{3}$
C (IG)	$3.9 \times 10^{3a} \pm 2.6 \times 10^{3}$	$<1 \times 10^{c}$	$<1 \times 10^a$
D (IG)	$5.2 \times 10^{3a} \pm 3.7 \times 10^{3}$	$5.7 \times 10^{2bc} \pm 9.1 \times 10^2$	$3.5 \times 10^{2a} \pm 4.9 \times 10^{2}$
E (GTS)	$2.8 \times 10^{6b} \pm 1.6 \times 10^{6}$	$8.1 \times 10^{5ab} \pm 6.8 \times 10^{5}$	$7.1 \times 10^{5b} \pm 6.6 \times 10^{5}$
F (GTS)	$3.7 \times 10^{6b} \pm 1.9 \times 10^{6}$	$1.0 \times 10^{6a} \pm 8.1 \times 10^5$	$1.0\times 10^{6b} \pm 8.6\times 10^{5}$
G (GTS)	$2.9\times 10^{6b} \pm 2.2\times 10^{6}$	$1.1 \times 10^{6a} \pm 1.1 \times 10^{6}$	$1.0\times 10^{6b}\pm 1.1\times 10^{6}$
H (GTS)	$3.0 \times 10^{6b} \pm 2.3 \times 10^{6}$	$1.3 \times 10^{6a} \pm 1.0 \times 10^{6}$	$1.3 \times 10^{6b} \pm 1.0 \times 10^{6}$

Means followed by the same letters in the column do not differ according to the Dunn's test (5% significance level).

GTS brands had higher bacterial counts than those in the IG samples.

The origin of coagulase-positive Staphylococcus contamination is usually related to poor raw material quality, lack of proper sanitation of equipment and utensils, and poor personal hygiene of food handlers (NOGUEIRA et al., 2013). Results on contaminant population reported in the IG and GTS groups in this study are similar to those in other reports (NOGUEIRA et al., 2013; ABREU et al., 2010a). Several pathogenic microorganisms (including coagulase-positive Staphylococcus) can survive in food residues or water in inadequately sanitized utensils, (DEVITA et al., 2007), reinforcing the fundamental role of the food handler's hygiene and the need for implementing good food handling practices.

The GTS group showed a clear lack of equipment and personal hygiene compared with that in the IG group, although cheese in both groups was produced under sanitary conditions. Among the coagulase-producing species, *S. aureus* is one of the most frequently involved species in foodborne outbreaks. In the present study, *S. aureus* was found at counts of <10 to 5.1×10³CFU·g⁻¹ in IG parmesan and <10 to 3.6×10⁶CFU·g⁻¹ in GTS parmesan.

Among the samples tested, 15.0% and 86.6% of the cheese samples contained more than 103CFU·g-1 of bacteria in the IG and GTS groups, respectively. When the means were analyzed by Dunn's test (P<0.05), all IG group samples differed significantly from the GTS group samples. Consequently, based on our results, it is possible to affirm that the population of S. aureus was significantly higher in the GTS parmesan than in the IG parmesan, thus posing a public health concern, mainly due to the risk of consuming of toxins produced during improper food storage and handling (SOUZA, 2006). Approximately 50% of the human population is a carrier of the bacterium (BERGDOLL & BENNETT, 1989), which can lead food contaminations, such as during the grating process involved in the handling of parmesan cheese.

The amount of mold and yeast in the IG samples varied from <10 to $4.0 \times 10^3 \text{CFU} \cdot \text{g}^{-1}$ and from <10 to $1.8 \times 10^6 \text{CFU} \cdot \text{g}^{-1}$ in the GTS samples. Brand F cheese (IG group) did not statistically differ from the samples of the GTS group. The mean values of other brands differed among each other (Dunn's test, P<0.05). Therefore, in terms of mold and yeast counts, IG parmesans (with the exception of brand F) have better microbial quality than GTS parmesans.

In this study, 25% of the IG samples exhibited fungal counts higher than 10³CFU·g⁻¹. Conversely, 90% of the GTS samples had counts higher than 10³CFU·g⁻¹. The presence of a high fungal population in food can indicate deterioration, even when there no visible signs. It is crucial to note that some filamentous fungi species can produce mycotoxins, which are potential carcinogenic compounds (HYMERY et al. 2014) and can lead to serious health problems.

According to ABREU et al. (2010b), among all microorganisms, fungi most commonly benefit from low water activity situations. This explains the high count of yeast and mold obtained in this research. Moreover, producing food in poor sanitary conditions can enhance the multiplication of these microorganisms (MATTAR et al., 2011).

However, some other factors have an inhibitory effect on fungal multiplication. According to TROMBETE et al. (2012), 100.0% of food samples analyzed in their study contained sorbic acid, which is a preservative. Manufacturers frequently use sorbic acid salts in cheese and other dairy products as they reduce fungal growth and increase the shelf life of the product (SALAVESSA, 2009). Despite the inhibitory effect of this preservative, the fungal count remained quite high in the samples analyzed in our study, mainly in the GTS group, which suggested contamination due to inadequate handling at sale time.

CONCLUSION

The parmesan cheese samples of GTS group showed higher counts of contaminants than those of the IG group, resulting in a higher probability of harboring pathogenic and deteriorating microorganisms. Thus, all 60 samples of the GTS group were in disagreement with the current legislation and scientific literature in terms of microbiological quality for consumed food, suggesting an improper manipulation of the cheese at the point-of-sale.

ACKNOWLEDGEMENTS

The study was supported with Coordenação de Aperfeiçoamento de Pessoal de Nível Superior (CAPES) grant.

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