

# Evaluation of Microbiological Quality of Raw Milk Produced at Two Properties in the Far West of Santa Catarina, Brasil

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**Abstract** Milk is a food that inherently favors microbial growth and due to its characteristics several precautions must be taken to prevent contamination in its production, processing, marketing and consumption, which are routinely subject to changes. The aim of this study was to evaluate microbiological contamination in milk produced at two farms in the Far West of Santa Catarina, before and after the application of Good Manufacturing Practices (GMP). Initially, samples of the milk, surfaces of equipment and utensils for milking, the teats of animals, disinfectants, and water were tested. Next, we conducted training of the farmers in microbiological analysis of milk samples. The analyses included counts of mesophilic aerobes (MA), *Staphylococcus coagulase positive* (SA), total coliform (TC), and thermotolerant (FC). The methods used for analyses were those described by the Regulation number 62 of August 26th, 2003 published by the Brazilian Ministry of Agriculture and Food Supply (MAPA) that follows methods recommended by the Compendium of Methods for the Microbiological Examination of Foods - APHA. The mean values for MA, SA, TC, and FC in milk obtained before and after the training were, respectively: 4.88 and 3.69 log colony-forming-unit (CFU)/ ml, 3.04 and 2.37 log CFU / ml, 61.19 Most probable number (MPN) and 17.89 MPN/ ml, and 40.26 and 8.71 MPN/ ml. Thus, according to these results, including training in GMP can improve the quality of milk, with immediate results for MA, TC, and CF. But, beyond the procedures employed, the control and prevention of mastitis could help to avoid contamination by SA.

**Keywords** Milk, Contamination, Good Manufacturing Practices

## 1. Introduction

Milk, being a complex mixture, nutritious, with a high level of water and a pH close to neutral, is highly perishable. It is a product highly conducive to microbial growth, especially bacterial pathogens[1]. Depending on the manipulations it is subject to, milk can have its physical, chemical and biological properties easily altered by the actions of micro-organisms. Thus the number of bacteria in milk, directly influences the quality and safety of dairy products[2]. Due to its characteristics, milk deserves special attention in its production, processing, marketing and consumption. Several factors, such as the health of the herd, sterility of the cleaning equipment and utensils used to obtain it, the health conditions of the milking place, the excretion from the udder of an infected animal and quality of water used on the farm,

may influence the microbiological quality of milk products[3-4].

Brazil stands out as one of the largest milk producers in the world. However, despite the fact that present production is growing, many milk producers still use non-specialized methods, resulting in raw material of poor quality[5]. The milk contaminated by high levels of bacteria usually becomes unsuitable for further processing since it does not meet the consumer's expectations in terms of health (nutritional value), safety (hygienic quality) and satisfaction (sensory attributes)[6].

The globalization of markets, including the availability of a variety of imported dairy products, has led the Brazilian consumer to become more demanding about the quality and safety of the products offered[7]. The dairy industry has undergone major transformations in an attempt to become more competitive, with benefits to the producer in terms of quality[8]. The parameters physico-chemical, microbiological, hygienic, and sanitation measures have been deployed by the industry to test and verify the quality of milk[9].

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The pathogens that have been involved in foodborne outbreaks associated with the consumption of milk include *Listeria monocytogenes*, *Salmonella*, *Campylobacter*, *Staphylococcus aureus*, *B. cereus* and *Costridium botulinum* and thermotolerant coliforms, especially *Escherichia coli* that is the most common contaminant of raw and processed milk[1],[10]. According to Mhone, Matope, Saidi[10], the total count of bacteria also became one of the criteria to evaluate the classification and processing of dairy products.

In aiming to meet the need for improvements in the dairy sector, the Ministry of Agriculture, Livestock, and Supply published Normative Instruction No. 51 (IN 51) on September 18, 2002, establishing definitive standards for the quality of refrigerated raw milk commencing in January of 2012[11].

In this context, there arose the need to implement changes and improvements in the milk production chain in order to meet the requirements of NI 51. The adoption of quality programs is also aimed at reducing costs in the process, with consequent increases in quality and product differentiation in the marketplace, and in meeting consumer demands.

Due to the relevance of the dairy sector for the national economic scenario, and considering the requirements of the major rule concerning federal sanitary legislation (the IN 51), this study aimed to assess the microbiological quality of refrigerated raw milk produced at two properties in the western end of Santa Catarina, and propose applying techniques of Good Manufacturing Practice (GMP) to the manufacturing production process in order to enhance quality and promote continuous improvement of the dairy sector.

## 2. Materials and methods

### 2.1. Collection of Samples

The study was conducted in two farms located in the Far West of Santa Catarina from November 2010 to May 2011. The state of Santa Catarina has great national highlights in the industry livestock and milk production of the major activities in the economy of the states occupying the 5<sup>th</sup> place in the national production of bovine milk in the 2010. The far west region of Santa Catarina is responsible for most of that milk production through small farms.

The samples of refrigerated raw milk and water were collected aseptically into sterilized bottles. Samples of disinfectant solution used for the realization of “pre-dipping” (means immersing the teats in disinfectant solution before milking) were aseptically collected 15 minutes after preparation of disinfectant solution in sterilized bottles. However, samples from the surface equipment, utensils, and teats of milking animals were collected with sterile swabs, by friction in the area of the sample. The swabs were dipped and kept in test tubes containing 10 mL of sterile peptone water 0.1%. All samples were transported in isothermal boxes with ice to the Microbiology Laboratory at the University of West of Santa Catarina- UNOESC, for analysis.

After mixing of the samples, decimal dilutions were performed by serial transfer of aliquots of 1 mL from sample tubes into tubes containing 9 mL of sterile water and 0.1% peptone (Difco, France), then the microbiological analyzes were performed.

### 2.2. Data Collection and Microbiological Analyses

Initially, a profile of each property was prepared by means of a questionnaire that was used for data collection. The first phase of the study aimed to identify points of contamination of milk during milking, and evaluate the microbiological quality of the milk produced on farms with conventional hygienic practices adopted by producers.

To achieve this objective, microbiological analyses were carried out on 20 samples of milk and two samples of disinfectants, two samples of the water used in properties, eight samples from the surfaces of teats of animals, and seven samples from equipment surfaces and milking utensils (two samples from milking equipment, two samples from storage tanks of milk samples, two from milk filters, and one from milk sampling of milk storage cans).

Subsequently, in the second phase of the study, after GMP techniques were applied in the production process, additional samples were collected, together with the collection and microbiological analysis of 40 samples of refrigerated raw milk, in order to evaluate the effects of adopting GMP.

The samples of refrigerated raw milk, disinfectants, and teat surfaces, milking equipment and utensils, were submitted to microbiological analysis for quantification of mesophilic aerobes (MA), *Staphylococcus coagulase positive* (SA), total coliform (TC), and thermotolerant (FC). Water samples were subjected to microbiological analysis to quantify microorganisms MA, TC and FC, according to the parameters required by Ordinance No. 518/2004 from the Ministry of Health, published on March 25, 2004[12]. All microbiological analyses were performed in triplicate on plates of the same dilution. The methods used for analyses were those described by the Regulation number 62 of August 26<sup>th</sup>, 2003 published by the Brazilian Ministry of Agriculture and Food Supply (MAPA)[13] that follows methods recommended by the Compendium of Methods for the Microbiological Examination of Foods - APHA[14].

The counts of MA were performed using a pour-plate technique. Aliquots of 1 ml of each dilution were placed on sterile petri dishes, followed by the addition of Plate Count Agar (Merck, Germany). Then the plates were gently mixed and incubated at  $36 \pm 1^\circ \text{C}$  for 48 hours. Plates which grew in the range of 25-250 colonies were counted. The results were expressed as CFU / ml.

Determinations of TC and FC in refrigerated raw milk and water were performed using Multiple Tube Fermentation. Aliquots of 1 ml of milk samples at dilutions of  $10^0$ ,  $10^{-1}$  and  $10^{-2}$ , and aliquots of 10 ml, 1 ml and 0.1 ml of the water samples, were transferred to tubes containing 10 ml of broth containing Lauryl Sodium Sulfate (Merck, Germany). The tubes were then homogenised and incubated at  $36 \pm 1^\circ \text{C}$  for

24–48 hours. After incubation, the degree of turbidity was observed, and production of gas from lactose fermentation was evaluated in Durham tubes. From confirm TC aliquots of positive tubes from the previous test were then transferred to tubes containing 10 ml broth Bile 2% Brilliant Green (Merck, Germany), incubated at  $36 \pm 1^\circ \text{C}$  for 24–48 hours to and to confirm CF put into tubes containing 10 mL of EC broth (Merck, Germany) and incubated at  $45 \pm 0.2^\circ \text{C}$  for 24–48 hours. Tubes showing evidence of gas or effervescence when shaken gently were considered positive. From the combination of numbers corresponding to the tubes that had a positive result in each of the confirmatory tests, the numbers of microorganisms of Most Probable Number (MPN) were determined according to the table contained in the IN 62. The results were expressed in MPN/mL for samples of milk, and in MPN/ 100ml for water samples.

To count of TC and FC in disinfectants and on teat surfaces, milking equipment and utensils, the pour-plate overlay technique with agar crystal violet neutral red bile (VRBA) (Difco, France) was used. After homogenization, the plates were incubated at  $36 \pm 1^\circ \text{C}$  for 24–48 hours. Dilutions on plates which produced in the range of 25–250 colonies were counted and the results were expressed as CFU/ml. To count and confirm TC and FC, three colonies characteristic of each plate of VRBA were inoculated into tubes containing 10 mL Bile 2% Brilliant Green (Merck, Germany), incubated at  $36 \pm 1^\circ \text{C}$  for 24–48 hours, and inoculated into tubes containing 10 mL of EC broth (Merck, Germany) and incubated at  $45 \pm 0.2^\circ \text{C}$  for 24–48 hours. Tubes with gas or effervescence when gently shaken were considered positive.

The numbers of SA were determined by inoculating 0.1 ml from each sample, and seeding with the aid of handle Drigalski on the surface of Petri dishes containing Baird-Parker agar (Oxoid, England). The plates were incubated at  $36 \pm 1^\circ \text{C}$  for 30–48 hours. To make the evaluation easier, plates with 25-250 colonies were selected for reading. Typical colonies were counted (colonies with a shiny black opaque halo surrounded by a clear halo or transparent) and atypical colonies (bright gray or black, without halo or with one halo). For confirmation, we selected an average of three colonies of each type (typical and atypical colonies). Then, the selected colonies were transferred to tubes containing broth and agar Brain Heart Infusion (BHI) (Oxoid, England), followed by incubation for  $36 \pm 1^\circ \text{C}$  for 24 hours. From each subculture,

the strains were subjected to coagulase testing, catalase, and fermentation of mannitol salt agar.

### 2.3. Statistical Analysis

To evaluate the existence of statistical differences between the parameters evaluated before and after the adoption of GMP procedures for handling, the Practice T-Students test was used. The T-Student analysis was used to compare the average counts of two different groups in order to verify statistically significant differences between them. Data analyses were performed on the SPSS (Statistical Package for Social Sciences) release 17.0. A p-value of  $< 0.05$  was considered statistically significant.

## 3. Results and Discussion

The main points of contamination by MA, TC, and FC were the milking equipment for the animals and the teats of animals. SA was only found on teats (Table 1). The disinfectant also showed high counts of MA (mean 5.9 log CFU / mL) and TC (mean 3.3 log CFU / ml). Thus, it can be stated that the practice of “pre-dipping”, conventionally used on these properties (reusable towels soaked in a 30% solution of diaminopropil laurilamina) was not effective in reducing microbial contamination of the teats to acceptable levels.

In most cases, disinfectants are chosen by habit of use, ease of application, or price, which, coupled with the lack of access to laboratory tests to evaluate the efficacy of the disinfectant solutions used in the routine milking processing, compromise the quality of the milk[15].

The water analysed at the two properties were in accordance with the standards required for MA contamination by Ordinance No. 518/2004, from the Ministry of Health; however, the samples showed the presence of TC and FC, and may have been a likely source of contamination by TC and FC of teats and milking utensils, and consequently of the milk produced. The result found results corroborate the statement of Amaral et al.[3], which emphasized the importance of the water used in order to obtain products of good microbiological quality; because, according to the author, the water used in production has great influence on the contamination of the milk, and, being a vehicle for transmission of pathogens, must have characteristics of potability.

**Table 1.** Average and standard deviations total count of mesophilic aerobes (MA), total coliform (TC), fecal coliform (FC) and *Staphylococcus* coagulase positive (SA) in samples from the surface of the teats, equipment and utensils of milking before training

Collection site	Mean $\pm$ standard deviations of the microorganisms			
	MA (log UFC/cm <sup>2</sup> )	TC (log UFC/cm <sup>2</sup> )	FC (log UFC/cm <sup>2</sup> )	SA (log UFC/cm <sup>2</sup> )
Milking (equipment)	4.1 $\pm$ 5.29	3.4 $\pm$ 3.38	3.4 $\pm$ 3.38	Not detectable#
Teats	3.8 $\pm$ 3.82	3.4 $\pm$ 2.94	3.1 $\pm$ 3.4	2.37 $\pm$ 2.81
Milk filters	5.3 $\pm$ 5.3	0.7 $\pm$ 3.39	0.7 $\pm$ 3.39	Not detectable#
Expansion tanks	4.2 $\pm$ 5.28	2.1 $\pm$ 2.13	1.7 $\pm$ 1.94	Not detectable#
Milk cans	0.1*	Not detectable*	Not detectable*	Not detectable#

\* Minimum detection of 1 logCFU/ml

# Minimum detection of 2 log CFU/ml

• Do not have a standard deviation, as only a farm possessed milk cans

Through observation of the practices adopted by the producers, negligence was noted on the judicious use of cleaning systems and sanitizing of equipment, especially in relation to concentrations of chemicals used, as well as the temperature and circulation time of the solutions. In some checks we found inadequate temperatures in milk storage (over 4 ° C) in addition to a delay time, on average, for initiating the cooling process. According to Guimarrães[16], the rate of cooling after obtaining the milk is the first step in maintaining bacterial counts at low levels. The more quickly the temperature is reduced, the better the milk storage because it prevents the multiplication of microorganisms[1]. Thus, expansion tanks should provide conditions for accelerated cooling of milk to 4 ° C in a time equal to or less than three hours, and then maintain this temperature.

For contribute towards improving the quality of milk we suggested to some producers handling practices during milking, such as: monitoring and recording procedures for cleaning / sanitizing of equipment; control of concentration, temperature, circulation time, and pH of cleaning solutions; control of drainage waste water equipment; replacement of milk filters between milkings; replacement of pre-dipping with teat dipping in a solution of iodine-based disinfectant and drying teats with paper towels; discarding the first three streams of milk and using a strip cup for diagnosis of clinical mastitis, including diagnosis and monitoring of subclinical mastitis with the California Mastitis Test (CMT test), often fortnightly; milking of animals by batch; and biannual cleaning of water reservoirs; as well as monitoring and control of cooling milk to 4 ° C within three hours after milking.

The average counts of MA, TC, and FC in raw milk obtained before and after training the producers were, respectively: 4.88 versus 3.69 log CFU / ml, of 1.78 log MPN/ml versus 1.25 log MPN/ml, and 1.60 log MPN/ml versus 0.94 log MPN/ml, respectively (table 2).

The results found in our study are similar to Andrade, Hartmann and Masson[17] and Fagan *et al.*[18] who found MA counts of 3.86 log CFU/ml and 4.21 log CFU/ml, respectively with the definitive limits set by IN 51 (5 log CFU/ml). However, several authors have reported poor quality of milk, with MA contamination levels above the levels recommended by IN 51, ranging from 1x10<sup>6</sup> CFU/mL to 1.68 X10<sup>7</sup> CFU/ml[16,19-23].

The high rates of contamination by mesophiles, TC, and FC in refrigerated raw milk are associated with failures in hygiene and sanitary procedures used in obtaining the milk, such as insufficient cleaning and disinfection of teats; and inadequate hygiene in the production system, including the deposition of waste on equipment surfaces which, in turn, act as nutrients that support the growth of contaminant microorganisms[22,24-25]. Moreover, it is important to note that TC and FC are not part of the natural microflora of milk, and represent a potential risk of contamination by pathogens[5,26]. According to Chye, Abdullah, Ayob[1], *E. coli* and coliform bacteria are often used as indicator microorganisms, and the presence of this bacteria implies a risk that

other enteric pathogens may be present in the sample. Thus, the reduction in MA, and statistically significant reductions in TC and FC counts observed after training of producers, showed that adoption of GMP standards at milking were effective in controlling these contaminations (table 2). Thus, we can say that counts of MA, TC, and FC can be controlled in a short time, when there is specific technical assistance and quality controls. These results indicate the need for immediate implementation of training programs for producers, to encourage improvements in the microbiological quality of the milk and milk products offered to consumers.

**Table 2.** Average and standard deviations total count of mesophilic aerobes (MA), total coliform (TC), fecal coliform (FC) and *Staphylococcus coagulase positive* (SA) in samples from milk before training and after training

	Mean ± standard deviations of the microorganisms			
	MA (log UFC/ml)	TC (log MPN/ml)	TF (log MPN/ml)	SA (log UFC/ml)
Before training	4.88±5.33	1,78±1,66	1.60±1.61	3.04±2.89
After Training	3.69±3.95	1,25±1,56	0.94±1.64	2.37±2.40

The results of our study show that all microorganisms used as indicators of milk quality (MA, TC and TF) reduced their count after the training and implementation of best practices in handling properties (table 2), demonstrating the importance of maintaining programs to guide farmers to improve milk quality produced by the properties.

Another important factor to note is that SA was present in raw milk in our study in five samples (25%) before and 20 samples (50%) after training of the producers. Several authors have also observed a high frequency of *Staphylococcus aureus* (ranging from 19.5% to 71%) in milk samples from dairy herds in various regions of Brazil[2,17,27-29], demonstrating that the presence of this agent may be linked to the health of the mammary gland, since SA is a major causative agent of bovine mastitis. According to Correa, Ribas, and Madrona[5], this group of microbes is probably carried into the milk from the mammary gland infection known as mastitis. For Riekerink *et al.*[27], isolation of *S. aureus* in tank milk is a likely indicator of the prevalence of intramammary infections in dairy cattle.

The increased frequency of SA in raw milk occurring after the adoption of best practices can be linked to the fact that mastitis rates are decreasing over time, after the treatment of existing cases and prevention of new cases. According to Guimarrães[16], the control of mastitis by the use of appropriate antibiotic treatment and implementation of a program of selecting animals for disposal generates long-term results. The major difficulties in mastitis control procedures on the properties were adoption of a batch milking system, and disposal of animals suffering from recurrent mastitis, which do not respond to antibiotic treatment. The animal carriers of the infection remain in the herd, where they may contribute to the contamination of new animals in the case of contagious mastitis, in addition to increasing the contamination of milk[16]; these findings corroborate our own results. Thus,

these data indicate the need to review the process adopted on the properties, with the immediate implementation of mastitis control programs that include batch milking systems, disposing of animals suffering from recurrent mastitis, and evaluation of the *in vitro* antimicrobial susceptibility of the main causative agents of mastitis in the herd in order to assist in the choice of effective antimicrobials.

Despite the high frequency of positive results, all counts were below 3.04 Log CFU / ml (table 2). Unlike the results found in this study, some authors have found contamination by SA in refrigerated raw milk with counts above 10<sup>5</sup> CFU/mL [5,23,26,30].

It should be noted that for the occurrence of staphylococcal poisoning, SA counts in milk of over 10<sup>5</sup> CFU/mL are needed. Thus, it can be concluded that the milk samples analyzed did not show a possible risk to staphylococcal poisoning at the point of collection.

However, according to Nader Son et al. [31], although the simple presence of strains of *S. aureus* enterotoxigenic does not necessarily imply the occurrence of cases of poisoning, it is known that milk is an excellent substrate for the proliferation of these microorganisms. Thus, the use of raw milk as a raw material for the production of derivatives may carry a potential risk of poisoning in the later stages of collection, if the milk is subject to conditions and inadequate storage temperatures conducive to the multiplication of SA, with consequent production of enterotoxins.

## 4. Conclusions

Training, including adoption of Good Manufacturing Practices, are effective in improving the microbiological quality of refrigerated raw milk, with immediate results in reducing levels of contamination by MA, TC, and CF. Thus, professional monitoring of properties is essential to ensure producers adopt necessary conditions of service to fulfill the definitive parameters of IN 51. However, in addition to adopting these new procedures, control and prevention of mastitis could help to avoid contamination with SA, because this is a major causative agent of infection, and consequently may change the microbiological quality of the milk.

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