

MICROBIAL POOL FOR BIOLOGICAL CONTROL OF MILK QUALITY AND PATHOGEN PROFILE OF MILK FROM A DAIRY FARM IN SOUTHERN BRAZIL

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Abstract

The objective of this study was to assess the health and environmental profile milk from a local dairy farm by applying a microbial pool of Bacillus cereus, Bacillus subtilis, Bacillus licheniformis, Bacillus amyloliquefaciens, Lactococcus lactis subsp. lactis in different environments. The inoculum used was diluted with water and then sprinkled on the surface of predetermined sites for characterization. Five swab samples from the ground were collected at random from the room or environment. Populations of the total and fecal coliforms, Salmonella spp., Staphylococcus coagulase positive and sulfite reducing Clostridium were set as indicators of hygiene and sanitary conditions. Milk samples were also collected from the cooling tank. Somatic cell counts (SCC), Somatic cell counts (SCC), fat, protein, lactose and milk total solids were determined parameters were determined, we used the methods set by ANVISA. A statistically significant reduction of SCC ($P < 0.05$) was observed from 2.8 to 2.0 log SSC.mL⁻¹, significant interactions were considered when the probability was lower than 5%. The inoculum also showed a highly significant effect on pathogen populations, especially on Salmonella spp. and sulfite reducing Clostridium, while a lower reduction on the other two pathogens. In this way, the inoculum was effective in reducing pathogen populations, improved the sanitary quality, thus it might be used as a prevention tool for the diseases with environmental origin in dairy farms.

Keywords: Mastitis. Enterobacteria. Sanitation control

1 Introduction

Milk is a relatively cheap source of nutrients having a high intake in developing countries (MÜLLER, 2002) therefore, milk quality needs to be assessed in order to ensure consumer safety. In Brazil, milk quality related to the livestock health has great impact on the productivity. Somatic cell count (SCC) is a useful predictor of intra mammary infection (IMI) that includes 75% leucocytes (neutrophils, macrophages, lymphocytes, erythrocytes) and 25% epithelial cells. High SCC has a negative influence on the quality of raw milk and is related to changes in milk consistency (density), low protein and high risk for milk hygiene, since it might even contain pathogenic organisms (SHARMA, SHINGH & BHADWAL, 2011). In addition, high SCC also results in reduced productivity of a herd, as an animal cannot be milked until it is completely recovered. The impact of this problem in the Brazilian herds is about 17.45% in clinical and 72.56% in subclinical settings (LADEIRA, 2001). These mastitis episodes are regularly associated with the presence of *Enterobacteriaceae* and the main source of this strain is the gastrointestinal tract (LEVY, 2004). Thermotolerant coliforms, *Staphylococcus* and *Salmonella* spp., are important indicators of hygiene and sanitary conditions since they are directly linked to good hygiene practices and presence of infectious bacterium (SANTOS et al., 2010).

Dairy facilities are at a greater risk for environmental mastitis than the cows on pasture at the time of milking. Sources of environmental pathogens include manure, bedding, feedstuffs, dust, dirt, mud, and water. Even in semi-confinement regime, the flock is exposed to various threats. The lack of proper sanity at the milking and feeding sites associated with high density of animals promotes the occurrence and spread of the diseases caused by the pathogens normally found in animal excreta. Thus, it is essential to ensure good hygienic and sanitary conditions, so that the health of the herd can be assured (RADOSTITS et al., 2002).

Usually the household cleaning products are used to perform sterilization. However, these chemicals are detrimental to the environment, since they are mostly not biodegradable (BRASIL, 1988). In consequent, water resources lose capability of self-purification when contaminated by these products.

An alternative to reduce or eliminate such pathogens in dairy environment is the use of an inoculum in pavements sites. The inoculum contains one or more kinds of non-pathogenic bacteria and dispersing agents, which indirectly control the population of pathogens mainly by increasing competition for subsistence. This method is usually implied in waste biological treatment systems with an objective to eliminate organic matter, and thus promotes an environmental competition for the substratum between the microorganisms naturally present in the waste. This biological control may also produce secondary metabolites that are bactericidal and/or bacteriostatic helping in control of certain microbial species (XAVIER & LUCAS, 2010).

Antagonistic bacteria have been reported as a potential alternative to inhibit pathogens instead of administering antibiotics. In related studies, *Bacillus cereus*, *Bacillus subtilis* and *Bacillus amyloliquefaciens* proved as excellent antagonistic agents of gram-positive bacteria, playing the role of inhibitors for pathogenic species, producing antibiotics like surfactin, fungycin e iturin (LEE et al., 2010). Another bacterium that had its antagonistic effects studied is the *Bacillus licheniformis*, a lactic-acid bacterium considered as a probiotic due to a production of inhibitory peptidic substances for pathogens and spoilage bacterium (CLADERA-OLIVEIRA, CARON & BRANDELLI, 2004; SHOBHARANI, PADMAJA & HALAMI, 2015). Another antimicrobial agent that can act as a probiotic for producing metabolites such as lactic acid, diacetyl, carbon dioxide, hydrogen peroxide and bacteriocins, is the *Lactococcus lactis* subsp. *lactis* which showed a potential for opposing the growth of bacteria such as *Salmonella* spp., *Staphylococcus aureus* (KHALID et al., 2011).

In this way, the aim of this study was to apply a pool of microorganisms on the floor of a milk cattle farm, evaluating its effects on the sanitation quality and overall quality of the milk produced on the property.

2 Materials and Methods

2.1. Microbial Pool Application

The microbial pool consisted of *Bacillus cereus*, *Bacillus subtilis*, *Bacillus licheniformis*, *Bacillus amyloliquefaciens*, *Lactococcus lactis* subsp. *lactis* in a concentration higher than $1,25 \times 10^8$ CFU.g⁻¹, sodium bicarbonate, sodium chloride and wheat bran; every component of a commercial product granted to the Laboratory of Wastes from the Universidade Federal de Pelotas (UFPel) to evaluate their antagonistic effect.

The application was carried out using a simple dilution with deionized water to reach a concentration of 71.4 mg.L⁻¹.cow⁻¹, followed by direct spraying onto the farms flooring, weekly reaching a total of 60 days.

2.2. Sampling

The milk and flooring samples were collected in a milk cattle farm, with a lot of 20 cows with a two milkings routine each day. The experiment included the calf stables with three animals with approximately 30 days of age each. The farm was located

in south of Rio Grande do Sul State and the cattle-management system was characterized by the pasture feed during summer. Three collection spots were defined to application and sampling: feeding square, pre-milking room and heifer rearing. The flooring was a slick concrete paving.

Five environmental samples per site were collected from random points by using a swab dipped in sterilized saline solution. Milk samples were also collected directly from milk cooling tank in the bronopol milk sample preservative (2-bromo-2-nitropropane-1,3-diol), with a 3 hours period between the milking and sample collection. All samples were stored and transported in insulated box with ice to the laboratory of wastes where the analytical procedures began within two hours of sampling.

2.3. Microbiological Analysis

For hygienic and sanitary characterization populations of total and thermotolerant coliforms, *Salmonella* spp., *Staphylococcus* coagulase positive and *Clostridium* were evaluated using a methodology recommended by the American Public and Health Association (APHA, 2001).

2.4. Milk Analysis

The data were analyzed for normality by the Shapiro-Wilk test, homoscedasticity by Hartley test and the independence of the waste by graphical analysis. For the somatic cell count and physicochemical characterization of the milk, we used the methods set by ANVISA (BRASIL, 2001). The determination of fat, solids, protein and lactose was carried out by absorption spectrometry method in the mid-infrared (MID) region; wherein the transmittance rate is measured to identify the organic compound.

2.5. Statistical Analysis

The experiment followed a fully randomized design with three repetitions and one factorial arrangement in which the factors were inoculum applications (before and after application). The milk samples were evaluated for fat, protein, lactose, milk total solids, and SCC. The manure samples were evaluated for microbial profiles.

After that, the data were subjected to analysis of variance ($P \leq 0.05$). In case of statistical significance T-test was applied ($P \leq 0.05$). Significant interactions were considered when the probability was lower than 5%.

3 Results and Discussions

The results of milk evaluation tests are shown in Table 1. With the exception of fat, all other milk components were affected by the inoculum application.

Table 1 - Milk constituents and SCC before and after inoculum applications.

| Treatment | BIA | AIA |
|-----------------------|--------|--------|
| Fat (%) | 4.07ns | 4.08 |
| Protein (%) | 3.52a | 3.04b |
| Lactose (%) | 4.38a | 4.23b |
| Milk Total Solids (%) | 13.20a | 12.27b |
| SCC (105 cs/mL) | 2.8a | 2.0b |

^{ns} Means (n=7) without significant difference in F test ($p < 0.05$). Means in lines followed by different letters show statistically significant difference between the treatments ($p < 0.05$). BIA means before treatment, AIA means after treatment and SCC means somatic cell count.

The protein showed a decrease of 13.64%, lactose of 3.42%, total solids of 7.04%, while the parameter, somatic cells, showed most significant decrease of 28.57%.

It's well known that few factors other than infection status have a significant impact on milk SCC and composition (HARMON, 1994), where it's possible to attribute the changes in milk profile to a reduction of the environmental pathogens. According to ICAR (2011), normal cow milk contains between 50,000 and 200,000 somatic cell/mL, which leads to categorize the milk of BIA under the hygiene-quality standards. Since the environmental pathogens often associated with SCC increase include the coliforms and environmental streptococci and enterococci (DUFOUR, 2011), it's possible to attribute the results to a competition between microorganisms (environmental pathogens versus inoculum microorganisms). High SCC milk shows a decrease in casein, fat and lactose (SABEDOT et al., 2014), thus the reduction observed in this trial can be attribute to SCC

decrease, an indirect effect of inoculum application, on the health of cows that are in milk production, allowing a better use of energy and nutrients by the animals. It is well known that a reduction in milk components affects product yield in its industrialization, being undesirable by the dairy industry. Additionally, pay for higher fat and protein is already a reality in Brazil (REIS et al., 2007) influencing the income of the milk producer.

The Normative Instruction 51, established by Agriculture, Livestock and Development Ministry (BRASIL, 2011) define the minimum values for fat, protein and defatted solids in 3%, 2% and 8.4%, respectively. In this case, both treatments were within the standards.

Table 2 - indicates the results of effectiveness on inoculum application at different places in the dairy cattle farm.

| Treatment | Total Coliforms (log MNP.mL ⁻¹) | Thermotolerant Coliforms (log MNP.mL ⁻¹) | Clostridium (log MNP.mL ⁻¹) | Staphylococcus (log CFU.mL ⁻¹) | Salmonella spp. |
|------------------------|---|--|---|--|-----------------|
| Feeding square (FS) | | | | | |
| BIA | 5.12a | 4,90a | 5.20a | N/D | A |
| AIA | 3.09b | 2,96b | 0.40b | N/D | A |
| Pre-milking room (PMR) | | | | | |
| BIA | 4.86a | 4.43a | N/D | 3.45ns | P |
| AIA | 3.14b | 3.10b | N/D | 2.71 | A |
| Heifer rearing (HR) | | | | | |
| BIA | 5.38a | 3.31ns | N/D | 4,23a | P |
| AIA | 3.21b | 3.18 | N/D | 3,13b | A |

^{ns} Means (n=7) without significant difference in F test ($p < 0.05$). Means in lines followed by different letters show statistically significant difference between the treatments ($p < 0.05$). BIA means before treatment, AIA means after treatment, P means presence and A means absence.

In microbiological analysis, a decline in total and thermotolerant coliform populations as well as *Clostridium* was observed at FS site Table 2. At PMR site, we observed a decline of 39.6% ($P < 0.05$) in populations of total coliforms, and the thermotolerant coliforms were reduced from 5.12 and 4.90 MPN.mL⁻¹ log for 3.09 and 2.96 log CFU.mL⁻¹ respectively. An analysis of the sulfite reducing *Clostridium* showed the presence of 5.20 log CFU.mL⁻¹ without the addition of inoculum, and this value reduced by 92.3% ($P < 0.05$) to 0.40 log CFU.mL⁻¹ after addition of the inoculum; however, we did not detect a positive *Staphylococcus* coagulase test and *Salmonella* spp. In FS, the *Staphylococcus* population showed no significant reduction ($P > 0.05$) after addition of the inoculum; this trend was contrary to the total coliform count that decreased by 35.4% MPN.mL⁻¹. This same enclosure verified the presence of 3.1 log MPN.mL⁻¹ thermotolerant coliforms after the treatment with sulfite reducing *Clostridium* inoculum. Total coliforms and coagulase positive *Staphylococcus* presented an initial population of 5.48 log MPN.mL⁻¹ and 4.23 log CFU.mL⁻¹, respectively, before the treatment; which reduced to 3.21 log MPN.mL⁻¹ ($P < 0.05$) and 3.13 log CFU.mL⁻¹ ($P > 0.05$), respectively. On the other hand, the fecal coliforms were not significantly reduced when exposed to inoculum ($P > 0.05$).

The decrease in pathogen counting suggested that the inoculum acted efficiently at the most important places, probably by inhibiting the growth of pathogenic microorganisms by competitive mechanisms for substratum or nutrients. In addition to these factors, some studies have indicated that secondary metabolites are produced, which affect the growth of other microorganisms in the environment (CAO et al., 2013).

Bacillus subtilis is one of microorganisms used in such inoculums that impair the growth of enteric pathogens such as *Salmonella* and *Clostridium perfringens* (LEE et al., 2010; THIRAUNYANON & THONGWITTAYA, 2012; SU-YEON & HYEONG-EUM & YONG-SUK, 2017), reported a natural competition site between this bacterium and natural pathogens of dairy cattle manure, forming a competitive environment for nutrients and space for cell attachment, while it does not affect the growth of other non-pathogenic bacteria of the same genus as *B. licheniformis* and *B. amyloliquefaciens*, which satisfy the condition of synergistic interaction with other species that make up the inoculum.

Other microorganism in the inoculum that might have caused the observed effect in this study is *Lactococcus lactis*, which produces metabolites during its growth such as nisin, a bacteriocin that prevents the growth of Gram-positive bacteria such as *Clostridium* and *Bacillus* (LEE et al., 1999, ASADUZZAMAN & SONOMOTO, 2009).

It is also known that *Bacillus amyloliquefaciens*, present in the inoculum, inhibits the growth of thermotolerant coliforms, *Salmonella typhimurium* and *Staphylococcus aureus*, by producing antibacterial substances. Another producer of secondary metabolites having antagonistic effect on pathogens is *Bacillus licheniformis*, which produces bacillocina 490 (CLADERA-OLIVEIRA, CARON & BRANDELLI, 2004; HE, CHEN & LIU, 2006).

Thermotolerant coliform such as *Escherichia coli* is the microorganism that indicates specifically a contamination of fecal

origin, as it is able to thrive in adverse environment (PARUCH & MAEHLUM, 2012; SCHWAIGER et al., 2012).

The inoculum used can be considered environmentally safe because it is formed by saprophytic and non-pathogenic microorganisms widely found in the environment (CLAUS & BERKELEY, 1986).

Nevertheless, the inoculum improved the sanitary quality of the manure and promoted the maintenance of non-pathogenic microbiota. However, in FS, one of the causes of the decrease in pathogen counts might be a greater concern for health by using some more effective products for microbial elimination. The absence of sulfite reducing *Clostridium* in PMR and HR sites can be attributed to a physical removal of matter present in animal paws.

Pathogen control by enhancing the environmental pressure seems to be an alternative strategy to the use of antimicrobials in disease control in dairy cows. However, further studies are required about health aspects for successful implementation of this technique.

5 Conclusions

With the data obtained in this study, we conclude that inoculum application interferes indirectly the physicochemical characteristics of milk and directly SCC, by promoting health quality in the system. In this way, the inoculum was also effective in reducing pathogen populations and might be used as a prevention tool for environmental diseases in dairy farms.

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