

Mouse milk somatic cell count in coagulase negative *Staphylococcus* species induced mastitis



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Abstract

Bovine mastitis is an economically important disease of bovines and caused by multi etiological factors in which bacteria is the major cause. In the present study, milk somatic cell count (SCC) in mouse mastitis induced by *Staphylococcus epidermidis*, *S. chromogenes*, *S. haemolyticus* and *S. aureus* isolated from apparently normal bovine milk was studied. The 2×10^4 cfu organisms in 50 μ l per teat were inoculated through intramammary route in 4th and 5th pairs of abdominal mammary gland in mice. The squeeze method, a technique was developed and standardized for milk collection from mice at 6, 12, 24, 48, 72 and 96 hr after intramammary inoculation (IMI). The milk somatic cell count was estimated using NucleoCounter SCC-100[®]. The mouse milk ranging from 50 to 200 μ l per mice was collected successfully. The milk somatic cell count showed significant increase in *S. aureus* inoculated mice at 6, 12, 24 hr and increase was two to four folds when compared to PBS control. The three coagulase negative staphylococcus species showed increased SCC levels at 24, 48, 72 and 96 hr after IMI. Thus, Coagulase negative staphylococci (C) species can increase the mice milk somatic cell count moderately but two to four fold increase was observed in *S. aureus* which indicated the subclinical nature of CNS infections in animals. Mouse is a suitable model to study coagulase negative staphylococcus species induced bovine subclinical mastitis.

Introduction

Bovine mastitis is one of the major disease conditions affecting dairy cattle worldwide. Subclinical mastitis is difficult to detect due to the absence of any visible indications and has major cost implications. Early detection of subclinical mastitis will help in reducing the economic losses to the dairy farmers. Among the bacteria isolated in bovine mastitis, *Staphylococcus* species occupies an important place. The staphylococcal genus has been divided into coagulase positive and coagulase negative based on their ability to coagulate rabbit plasma. Coagulase negative staphylococci (CNS) have become more important as bovine mastitis causing agents during recent years. More than ten different coagulase negative staphylococcal species have been isolated from mastitis bovine milk samples and most commonly reported are *Staphylococcus chromogenes*, *Staphylococcus simulans*, *Staphylococcus haemolyticus*, *Staphylococcus hyicus* and *Staphylococcus epidermidis* (Trinidad *et al.*, 1990; Matthews *et al.*, 1992; Myllys, 1995; Thorberg *et al.*, 2006). These bacteria are sometimes referred to as environmental staphs and most frequent organisms isolated from milk samples from herds that have controlled the major mastitis pathogens (Janus, 2009). The study of mastitis in bovines is costly and involves various ethical and social issues, especially, in India and also keeping the bovines in controlled environment is difficult. Hence, the bovine mastitis is mostly studied in laboratory animal models like mice, rat and rabbits. Various studies have been carried out using the animal models with major mastitis causing pathogens like *Staphylococcus aureus*, *Streptococcus* species and Coliforms. Simojoki *et al.*, (2009) suggested that studying of host response of CNS in experimental mastitis model is necessary to understand the host pathogen interaction. Keeping this information in background, the present study was undertaken to know milk somatic cell count in three CNS and *S. aureus* in mouse mastitis model which may help in early diagnosis of subclinical mastitis.

Materials and Methods

Animals and housing

Three coagulase negative staphylococci (CNS) namely *Staphylococcus epidermidis*, *Staphylococcus chromogenes* and *Staphylococcus haemolyticus* and one coagulase positive *Staphylococcus aureus* were isolated from apparently healthy bovine milk from dairy farms in Karnataka. The timed pregnant (Day 12 to 15), one hundred and sixty eight Swiss albino mice were procured from National Centre for Laboratory Animal Science, National Institute of Nutrition, Hyderabad. The mice were grouped in to seven groups (six time points and one control) of six mice each containing 42 mice per organism. The mice were housed in individually ventilated cages (IVC) during the experiment. The temperature and humidity of the animal room were maintained at $23 \pm 3^\circ\text{C}$ and 50 to 70 per cent respectively. Mice were provided with standard pellet feed and purified water *ad libitum*, and autoclaved paddy

husk as bedding material. The animal experiments was approved by Institutional Animal Ethics Committee (IAEC) of Veterinary College, Bangalore and carried out as per the guidelines of Committee for the Purpose of Control and Supervision of Experiments on Animals (CPCSEA), Ministry of Environment and Forests, New Delhi.

Mouse milk collection

The organisms were inoculated through intramammary route at the dose of 2×10^4 cfu per teat in 4th and 5th pairs of abdominal mammary glands in mice which simulated the four quarters of bovine. The pups were allowed to suckle after one hr of inoculation to simulate the natural conditions. The pups were separated from the mother four hours before the milk collection. The milk was collected from lactating mouse at 6, 12, 24, 48, 72 and 96 hr after intramammary inoculation with bacteria and PBS. The mouse was administered with 0.5 units of oxytocin intramuscularly in the thigh region. Maximum amount of milk was collected by squeeze method from L4, L5, R4 and R5 part of mammary glands from the base of the teat. The milk samples were collected by using micro pipette and micro tips in sterile 0.2 ml collection tubes (PCR tubes).

Milk somatic cell count

The somatic cell count of mouse milk was carried out by using somatic cell counter (NucleoCounter SCC-100, Chemometec, Denmark). The mouse milk sample was diluted since the quantity of milk obtained was very less. The diluted milk sample was mixed with equal amount of lysis buffer to form cell lysate. The lysis buffer disrupted the plasma membrane and liberated the nuclei of the cells which were stained with propidium iodide stain. The number of nuclei stained with propidium iodide was counted using fluorescence microscope in combination with automatic image analyzer inside the NucleoCounter SCC-100. Five micro litre of mouse milk was diluted in 45 μl of sterile PBS in a 1.5 ml micro centrifuge tube and mixed gently. Added 50 μl of lysis buffer to it and mixed gently. Then the SCC-cassette was kept on the tube and pressed the white plunger to suck the diluted milk in to the cassette channels. Pressed the button-Analysis in the somatic cell counter and waited for few minutes. The number of somatic cells in the milk sample was shown in the machine and computer monitor, which was recorded and multiplied with the dilution factor to know the exact number of somatic cells in the mouse milk sample. The somatic cell count in the mouse milk was calculated using Somatic View software Version 1.0 (Chemometec, Denmark).

Statistical analysis

The number of somatic cells in a mouse milk sample was expressed as $\times 10^5$ cells per ml of milk. The data obtained were analyzed by one way analysis of variance (Snedecor and Cochran, 1989) and by using Statistical Analysis System (SAS) software version 9.3, Mumbai, India.

Results

The mouse milk was collected by squeeze method (Fig.1) which was developed and standardized successfully and the amount of milk collected ranged from 50 to 200 μ l. The mean $\pm$ SE of milk somatic cell count in mice infected with three CNS and *S. aureus* has been presented in Table 1. The mean mice milk somatic cell count ($\times 10^5$ per ml) were 3.05 ± 0.30 , 10.67 ± 1.38 , 7.97 ± 0.89 , 5.26 ± 0.51 and 12.46 ± 2.99 in PBS control, *S. epidermidis*, *S. chromogenes*, *S. haemolyticus* and *S. aureus* inoculated mice respectively. In the present study, the SCC of PBS inoculated mice showed mean cell count which ranged from $2.46 - 3.64 \times 10^5$ per ml of milk. The mean SCC in mice milk samples collected from mice inoculated with *S. aureus* revealed a significant ($P < 0.05$) increase at 6, 12 and 24 hr after IMI. The increase in SCC was drastic at 12 and 24 hr after IMI in the *S. aureus* infected mice. At 6 and 12 hr after IMI with three CNS, no significant difference in SCC was observed when compared to PBS control. At 48, 72 and 96 hr after inoculation, there was a significant ($P < 0.05$) increase in SCC levels in *S. epidermidis* and *S. chromogenes* infected mice when compared to the PBS control. The *S. haemolyticus* showed increase in SCC at 12, 24 and 48 hr when compared to PBS control. The overall mean SCC of all the time points revealed increase in SCC of infected mice when compared to the PBS control.

Discussion

The mean mice milk SCC was increased in three CNS and *S. aureus* infected mice when compared to PBS control. However, the *S. aureus* inoculated mice showed significant increase in mice milk SCC at 6, 12 and 24 hr after intramammary inoculation. The mouse milk somatic cell count in experimental mastitis was not reported earlier. Further, the increased SCC in *S. aureus* infected mammary glands has been reported previously in dairy cows (Petzl *et al.*, 2008; Paradis *et al.*, 2010; Singh and Garg, 2011) and rabbits (Azeemullah, 2010). The *S. epidermidis* and *S. chromogenes* increased the SCC at 24, 48, 72 and 96 hr after IMI in mice moderately. The *S. haemolyticus* and *S. aureus* behaved in similar manner in different time points but SCC was increased moderately by *S. haemolyticus*. This finding concurred with previous reports which indicated the increase in geometric mean SCC in CNS infected quarters was 138,000 cells per ml and in *S. aureus* infected quarters was 357,000 cells per ml (Djabri *et al.*, 2002). Haas *et al.*, (2004) indicated the relationship between the high averages of SCC with the risk of occurrence of CNS mastitis in dairy cows. The milk SCC in dairy cows infected with CNS species has been shown to increase SCC by many researchers (De Vliegher *et al.*, 2003; Taponen *et al.*, 2007; Simojoki *et al.*, 2009; Piepers *et al.*, 2010; Malek Dos Reis *et al.*, 2011). Supre *et al.*, (2011) reported that the milk SCC in dairy cows increased due to CNS species infection which

may be comparable with *S. aureus* infected cows at times. The resident macrophages and the epithelial cells of the mammary gland after detection of invading pathogen releases several chemoattractants which trigger the migration of leukocytes, mainly PMN cells, from blood towards the inflamed mammary gland (Zhao and Lacasse, 2008). The migrated PMN cells were considered as the first line of defense in the mammary gland, but the presence of functional PMN in milk was crucial to the host defense against bacterial pathogens (Paape *et al.*, 2003). The main functions of PMN cells in the infected mammary glands is to phagocyte the pathogens and destroy them via oxygen dependent and oxygen independent systems. At the same time, PMN cells can potentially harm the mammary gland by promoting tissue injury via reactive oxygen species generation and by granular enzyme release on degranulation (Zhao and Lacasse, 2008). Thus the somatic cell count can serve as indicator for diagnosis of CNS induced subclinical mastitis. The mice might be used as suitable model to study the CNS subclinical mastitis and its therapeutic intervention strategies in future.

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Table 1: The mean mice milk somatic cell count (x 10⁵ per ml) after intramammary inoculation with three CNS species and *S. aureus* at different time points

Time points	PBS Control	<i>Staphylococcus epidermidis</i>	<i>Staphylococcus chromogenes</i>	<i>Staphylococcus haemolyticus</i>	<i>Staphylococcus aureus</i>
6 hr	1.70 ± 0.70 ^b	2.80 ± 0.40 ^b	3.15 ± 0.25 ^b	3.30 ± 0.40 ^b	8.45 ± 0.85 ^a
12 hr	3.88 ± 0.47 ^b	7.75 ± 0.35 ^b	6.85 ± 0.35 ^b	6.65 ± 0.55 ^b	25.85 ± 1.25 ^a
24 hr	3.15 ± 0.35 ^d	15.80 ± 1.60 ^b	9.65 ± 0.55 ^c	6.25 ± 0.25 ^{cd}	26.75 ± 1.15 ^a
48 hr	3.20 ± 0.30 ^c	11.05 ± 0.55 ^a	7.65 ± 0.45 ^{ab}	7.25 ± 0.85 ^b	3.64 ± 0.70 ^c
72 hr	3.15 ± 1.45 ^b	12.15 ± 0.25 ^a	7.85 ± 1.25 ^{ab}	5.00 ± 0.20 ^b	5.20 ± 0.40 ^b
96 hr	3.20 ± 0.70 ^b	15.00 ± 3.00 ^a	12.70 ± 0.20 ^a	3.10 ± 0.40 ^b	5.05 ± 1.55 ^b
Mean	3.05 ± 0.30	10.67 ± 1.38	7.97 ± 0.89	5.26 ± 0.51	12.49 ± 2.99
CI	2.46 – 3.64	7.97 – 13.37	6.23 – 9.71	4.27 – 6.25	6.63 – 18.35

Values expressed in cells x 10⁵ per ml in milk as Mean ± SE

a, b, c, d : Means with same superscript within the row do not differ significantly (P>0.05)

CI: Confidence interval at 95 per cent level

Fig. 1 Photograph to show collection of milk from mice mammary glands by squeeze method.

