

## Immune response of bovine milk somatic cells to endotoxin in healthy quarters with normal and very low cell counts

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Low somatic cell count (SCC) is a reliable indicator of high-quality milk free of pathogenic microorganisms. Thus, an important goal in dairy practice is to produce milk with low SCC. Selection for cows with low SCC can sometimes lead to extremely low SCC in single quarters. The cells in milk are, however, predominantly immune cells with important immune functions. To investigate the mammary immune competence of quarters with very low SCC, healthy udder quarters of cows with normal SCC of  $(40\text{--}100) \times 10^3$  cells/ml and very low SCC of  $<20 \times 10^3$  cells/ml were challenged with lipopolysaccharide (LPS) from *Escherichia coli*. In the first experiment, SCC and cell viability after a challenge with 50 ng of LPS/quarter was investigated. In the second experiment, tumour necrosis factor  $\alpha$  (TNF- $\alpha$ ) concentration and lactate dehydrogenase (LDH) activity in milk, and mRNA expression of various innate immune factors in milk cells were measured after a challenge with 100  $\mu$ g LPS/quarter. LPS challenge induced an increase of SCC. SCC levels reached were higher in quarters with normal SCC and maximum SCC was reached 1 h earlier than in very low SCC quarters. The increase of TNF- $\alpha$  concentrations in milk in response to LPS challenge was lower in quarters with very low SCC than in quarters with normal SCC. The viability of cells and the LDH activity in milk increased in response to LPS challenge, however, without a difference between the groups. The mRNA expression of IL-1 $\beta$  and IL-8 was increased in milk cells at 12 h after LPS challenge, whereas that of TNF- $\alpha$  and lactoferrin was not increased at the measured time points (12, 24 and 36 h after LPS challenge). No differences of mRNA expression of measured immune factors between normal and very low SCC samples were detected. The study showed that udder quarters with very low SCC responded with a less marked increase of SCC compared with quarters with normal SCC. This difference corresponded with simultaneously lower TNF- $\alpha$  concentrations in milk. However, the immune competence of the cells themselves based on mRNA expression of TNF- $\alpha$ , IL-8, IL-1 $\beta$ , and lactoferrin, did not differ. The results may indicate that very low SCC can impair the immune competence of udder quarters, because the immune response in udder quarters with lower SCC is less efficient as fewer cells contribute to the production of immunoregulators.

**Keywords:** Bovine somatic cells, immune response, endotoxin, SCC.

Somatic cell count (SCC) is used worldwide as a hygienic parameter of milk. Low SCC is a reliable indicator of healthy mammary gland quarters because increasing SCC is mostly the result of pathogen invasion (Schukken et al. 2001). In dairy production exceeding a fixed SCC limit invokes penalties. Thus, it is economically advantageous to reduce SCC to very low levels, and a goal in dairy practice can be

the breeding of cows that produce milk with very low SCC (Shook, 1989).

The cells in milk are, however, predominantly immune cells. These somatic cells play, evidently, important roles in the defence of the mammary gland. They recognize invading pathogens and initiate the innate immune reaction through liberation of immunomodulators (Rainard & Riollot, 2006). Each cell type (macrophages, neutrophils, lymphocytes, and epithelial cells) has a vital role in the immunity of the mammary gland e.g. phagocytosis, antigen presentation, secretion of antibacterial factors, immunological memory, or regulation of the immune response

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(Sordillo & Streicher, 2002). An excessive reduction of SCC might therefore have a negative effect on the mammary immune competence.

There are numerous reports concerning low SCC and clinical mastitis incidence. Green et al. (1996) and Suriyasathaporn et al. (2000) showed an association between low herd SCC and increasing mastitis incidence. Beaudeau et al. (2002) found a greater risk of clinical mastitis in herds containing a high proportion of cows with low SCC ( $<50 \times 10^3$  cells/ml). In addition, Deluyker et al. (1993) showed that in a low-SCC herd, cows with clinical mastitis had a higher SCC ( $>245 \times 10^3$  cells/ml) prior to mastitis than control cows ( $<90 \times 10^3$  cells/ml). Furthermore, Sarikaya et al. (2005) found that cells from milk with very low SCC ( $<12 \times 10^3$  cells/ml) had lower mRNA expression levels of inflammatory factors compared with cells from milk with higher SCC. No previous investigation had, to our knowledge, the specific focus of studying the immune competence of mammary gland quarters with very low SCC (VLS) and with normal SCC (NS) in response to a defined challenge of the mammary immune system.

With intramammary injections of the *Escherichia coli* lipopolysaccharide (LPS), it is possible to mimic bacterial invasion and to induce a mammary inflammation very similar to natural mastitis (Schmitz et al. 2004). In contrast to a bacterial infection, the intensity of the inflammatory stimulus is exactly defined by the LPS dose, and the immune response can be compared between animal groups. SCC augmentation following stimulation with pathogen or endotoxin is due to the release of immunomodulators from resident somatic cells and mammary epithelial cells after recognition of the antigen (Sordillo & Streicher, 2002). TNF- $\alpha$  is a rapidly responding mediator of inflammatory reactions that increases in milk after an immune stimulation with LPS (Paape et al. 2002). IL-1 $\beta$  is a cytokine that is rapidly up-regulated in milk cells during infection to initiate the immune response (Lee et al. 2006). IL-8 is an important chemokine that is involved in the further recruitment of immune cells into the milk during the immune response (Persson-Waller et al. 2003). Lactoferrin is known to have antibacterial effects and during acute mastitis neutrophil leucocytes are besides the epithelial cells an important source (Harmon & Newbould, 1980). Therefore, the production of the factors TNF- $\alpha$ , IL-1 $\beta$ , IL-8 and lactoferrin indicates the immune activity of the cells. Parallel to the release of these factors and the induced immune response, an increased recruitment of new leucocytes from blood into the milk is induced. Finally the enhanced population of young leucocytes increases the viability of the total milk cell population (Mehrzhad et al. 2004).

The present study was performed to compare the response to LPS of udder quarters with NS and VLS milk in relation to the SCC increase, changes of TNF- $\alpha$  and LDH in milk and the mRNA expression of TNF- $\alpha$ , IL-1 $\beta$ , IL-8 and lactoferrin in milk cells. This could give an idea of whether the immune protection of udder quarters differs

where the SCC is below  $20 \times 10^3$  cells/ml compared with the quarters with SCC of  $(40-100) \times 10^3$  cells/ml.

## Materials and Methods

### Experiment 1

Thirty-three quarters of 19 lactating dairy cows (8 Holstein, 11 Red Holstein  $\times$  Simmental) from udders free of clinical signs of mastitis were used for this experiment. All cows were chosen randomly and were in months 2–15 of lactations 1–7. Cows were kept in a tethered barn and were fed twice daily with hay, corn and concentrate. Water was available ad libitum. Milking was performed routinely twice daily.

One or two quarters of each cow were randomly selected based on foremilk SCC determined with the DeLaval cell counter (DCC, DeLaval, Tumba, Sweden). Quarters were divided into two groups: very low SCC (VLS; SCC  $<20 \times 10^3$  cells/ml; SCC =  $16.2 \pm 3.0 \times 10^3$  cells/ml;  $n=16$ ) and normal SCC [NS;  $(40-100) \times 10^3$  cells/ml; SCC =  $56.3 \pm 4.4 \times 10^3$  cells/ml;  $n=17$ ). The other quarters of each used udder did not show SCC above 300 000 cells/ml. The number of lactations of the cows was normally distributed between the two groups (5, 8, 2 and one quarter of cows in the 1<sup>st</sup>, 2<sup>nd</sup>, 3<sup>rd</sup> and 6<sup>th</sup> lactation in VL group, and 6, 7, 3, and one quarter of cows in the 1<sup>st</sup>, 2<sup>nd</sup>, 3<sup>rd</sup> and 7<sup>th</sup> lactation, respectively). In each group, 7 control quarters (C quarters) were randomly chosen and all other quarters were treated with LPS (LPS quarters).

Cows were milked during routine milking procedures using a quarter milking device for separate quarter milk collection (Sarikaya et al. 2005). Immediately after morning milking LPS, isolated from a mastitis-inducing *Esch. coli* strain by the University of Constance group (Prof. Hartung and Dr von Aulock), dissolved in sterile saline solution (9 g/l), was intramammarily infused into LPS quarters (50 ng/10 ml saline solution). Into the C quarters 10 ml saline solution was infused. Before injection teat openings were carefully disinfected with 70% alcohol. To measure the change of SCC after treatment, strict foremilk samples (first 3 squirts) were collected hourly from all quarters until the evening milking and daily (except day 6) before morning milking until day 7 post challenge (p.c.). Samples were obtained by hand milking without udder preparation to prevent any milk ejection, and therefore alveolar milk, in the samples (Bruckmaier & Hilger, 2001). SCC was measured with the DCC. From 26 of the quarters (6 C and 6 LPS for VLS and 7 C and 7 LPS for NS group) 1 l of milk at the morning milking (0 h) and at the evening milking (10 h p.c.) was collected and filtered ( $\varnothing$  100  $\mu$ m). Milk samples were diluted 1 : 1 with cold (4 °C) sterile PBS and centrifuged at 1000 g at 4 °C for 15 min. Fat layer and supernatant were removed (aspiration) and cell pellets resuspended and centrifuged twice in 50 ml PBS at (400 g and 300 g at 4 °C for 10 min). Pellets were resuspended in cold PBS (2–10 ml, depending on SCC). Cell suspension

**Table 1.** Experiment 2: Sequences of PCR Primers [forward (for) and reverse (rev)], PCR product length (bp), accession number, annealing temperature (°C)

Primer		Sequence (5'→3')	Length	Accession no. 4	Annealing temperature
TNF- $\alpha$	for	CCA CGT TGT AGC CGA CAT C	155	NM_173966	60
	rev	CCC TGA AGA GGA CCT GTG AG			
Lactoferrin	for	GGC CTT TGC CTT GGA ATG TAT C	338	DQ522305	60
	rev	ATT TAG CCA CAG CTC CCT GGA G			
IL-8	for	ATG ACT TCC AAG CTG GCT GTT G	150	EU276073	60
	rev	TTG ATA AAT TTG GGG TGG AAA G			
IL-1 $\beta$	for	AGT GCC TAC GCA CAT GTC TTC	114	M37211	62
	rev	TGC GTC ACA CAG AAA CTC GTC			
GAPDH	for	GTC TTC ACT ACC ATG GAG AAG G	197	NM_00103403	60
	rev	TCA TGG ATG ACC TTG GCC AG			

(25  $\mu$ l) of the cell suspension was used for Trypan Blue staining (0.25 %) for determination of viability in Neubauer's counting chamber (Brand, Wertheim, Germany) with direct light microscopy.

### Experiment 2

Thirty-two udder quarters from 17 lactating dairy cows (6 Holstein and 11 Red Holstein  $\times$  Simmental) were grouped into a VLS group ( $n=10$ ) or into a NS group ( $n=22$ ). Udders from experimental cows did not show clinical signs of mastitis. One day before the experiment sterile milk samples were collected from all quarters and cultured over night at 37 °C on blood agar plates to assure that all quarters were free of bacterial infections.

Immediately after the morning milking 5 VLS quarters and 15 NS quarters were infused intramammarily with 100  $\mu$ g *Esch. coli* LPS (SIGMA; O26:B6; # L8274) dissolved in 10 ml saline solution, as described in Experiment 1.

Milk samples ( $\sim 5$  ml) were taken within 40 s before the start of milk ejection after 0 h, 3 h, 6 h, 9 h, 12 h, 24 h and 36 h p.c. and frozen at  $-20$  °C until analyses. The protein content of TNF- $\alpha$  was measured by radioimmunoassay according to Blum et al. (2000). In addition, LDH activity in milk was measured using a commercial kit (Roche, Rotkreuz, Switzerland).

Milking was performed immediately before LPS challenge and at 12 h, 24 h and 36 h p.c. with a special milking device for collection of quarter milk. One litre of the removed milk of each quarter was centrifuged in 500-ml flasks at 2000  $g$  at 4 °C for 20 min. The pellet was redissolved by up-and-down pipetting with 200 ml ice cold PBS and centrifuged again at 2000  $g$  at 4 °C for 10 min. The final pellet was redissolved in 1 ml of Trifast (Peglab), transferred into 1.5-ml microtubes and frozen at  $-80$  °C until the extraction of total RNA according to manufacturer's protocol.

Final RNA concentration was quantified by spectrophotometry (Biophotometer, Vaudaux-Eppendorf, Basel,

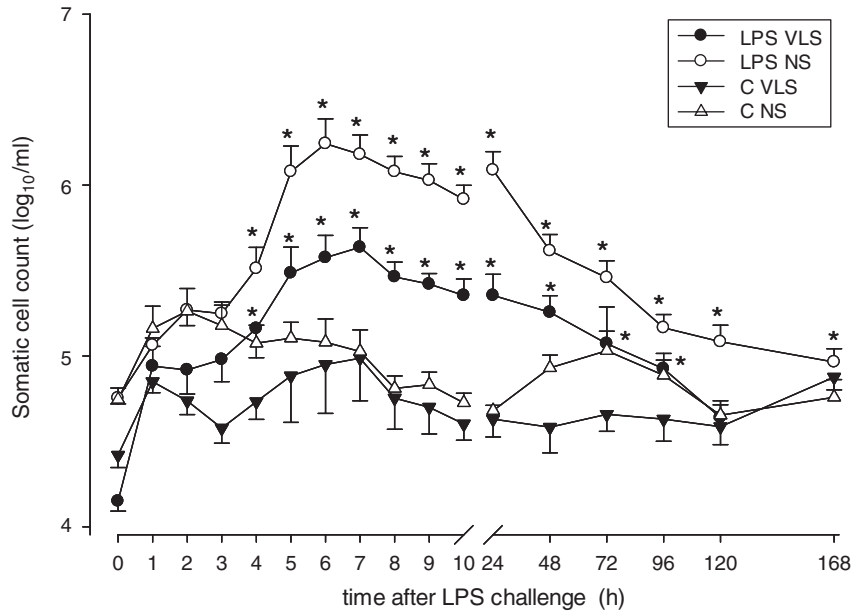
Switzerland) by measuring optical density (OD) at 260 nm. cDNA was produced using 50 ng RNA with 200 U Moleney Murine Leukemia Virus Reverse Transcriptase RNase H minus, Point Mutant (MMLV-RT, Promega, Madison WI, USA) and 100 pmol random hexamer primers (Invitrogen, Leek, The Netherlands).

Quantitative PCR analysis was carried out on Rotor Gene 6000 (Corbett Research, Sydney, Australia) using Sensimix DNA Kit (Quantace, Biolabo, Châtel S<sup>t</sup> Denis, Switzerland). Primers for housekeeping (GAPDH) and the target genes TNF- $\alpha$ , IL-8, IL-1 $\beta$  and lactoferrin were synthesized commercially (Microsynth, Balgach, Switzerland) (Table 1). The following 3-step programme was used: 10 min at 95 °C, 40 cycles of 95 °C for 15 s, primer specific annealing temperature for 30 s, and 20 s 72 °C, ending with a melting curve programme (60–99 °C, heating rate of 0.1 °C/s, continuous measurement). Take-off values (second derivative maximum; CP) were achieved by Rotor Gene software version 1.7.40. mRNA expression is given in  $\Delta\Delta$ CP values: to calculate the impact of treatment the gene expression was normalized to the expression of the housekeeping gene GAPDH. Then the gene expression values of non-treated control cells [ $\Delta$ CP (control cells)] and treated cells [ $\Delta$ CP (treated cells)] were set in relation, according to following equation:

$$\Delta\Delta\text{CP} = \Delta\text{CP}_{(\text{control cells})} - \Delta\text{CP}_{(\text{treated cells})}$$

### Statistical analyses

Data are presented as means  $\pm$  SEM. SCC are presented and statistically evaluated at a logarithmic scale ( $\log_{10}$ ) to ensure normal distribution. Differences of SCC, cell viability and mRNA expression of immunomodulators between groups (VLS and NS group, C and LPS quarters) were tested by analysis of variance using the PROC MIXED procedure of SAS (1999–2001). The model included time, group, and their interaction as fixed effects. Differences between means were considered significant if  $P < 0.05$ .



**Fig. 1.** Experiment 1: SCC (in  $\log_{10}/\text{ml}$ ) after intramammary injection of 0 (C) and 50 ng LPS (LPS) dissolved in 10 ml saline solution into quarters of very low SCC (VLS; SCC <20 000 cells/ml) and normal SCC (NS; 40 000–100 000 cells/ml). ● LPS VLS ( $n=9$ ), ○ LPS NS ( $n=10$ ), ▼ C VLS ( $n=7$ ), △ C NS ( $n=7$ ). Data are represented as mean  $\pm$  SEM. \*: Significant differences between LPS challenge and control within each group ( $P<0.05$ ). SCC between groups were significantly different ( $P<0.05$ ). SCC increased in response to LPS in both groups, whereas the maximum was reached 1 h earlier in the NS group.

## Results

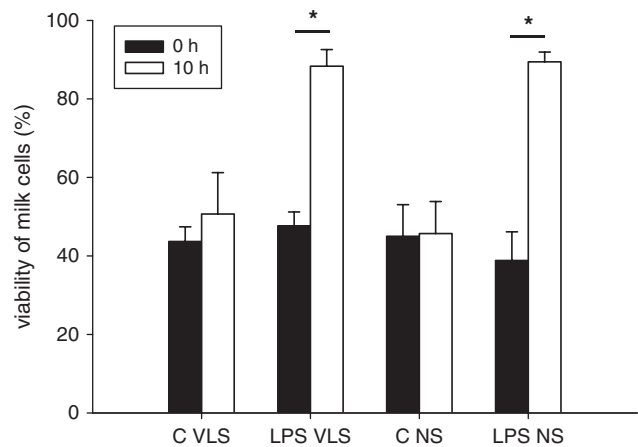
### Experiment 1

Milk SCC data from LPS and C quarters at each time point are shown in Fig. 1. Before LPS challenge (0 h) SCC of VLS ( $14.1 \pm 1.5 \times 10^3$  cells/ml) and NS group ( $56.7 \pm 4.2 \times 10^3$  cells/ml) were significantly different. SCC of C quarters from both groups did not change significantly throughout the entire experiment. At 4 h p.c. SCC began to increase ( $P<0.05$ ) in both LPS-treated groups to reach a maximum at 6 h p.c. for NS group ( $1747 \pm 696 \times 10^3$  cells/ml) and at 7 h p.c. for VLS group ( $432 \pm 130 \times 10^3$  cells/ml).

Cell viability (Fig. 2) of all groups at time 0 h was  $43.7 \pm 3\%$ . There was no difference between groups. At 10 h p.c. viability did not change in both C groups ( $48 \pm 6.3\%$ ) but increased significantly in LPS-treated quarters with no differences between VLS and NS groups ( $88.9 \pm 2.3\%$ ).

TNF- $\alpha$  concentrations in milk (Fig. 3A) increased within 3 h after LPS treatments in both groups and stayed elevated until 12 h p.c. Significant differences between VLS and NS groups was found from 3 h until 12 h p.c. with higher TNF- $\alpha$  concentrations in the NS group. In controls TNF- $\alpha$  concentrations did not increase.

LDH activity in milk (Fig. 3B) increased after 6 h p.c. in LPS quarters with no differences between groups and stayed elevated until the end of the experiment. In C quarters no increase of LDH activity was detectable.

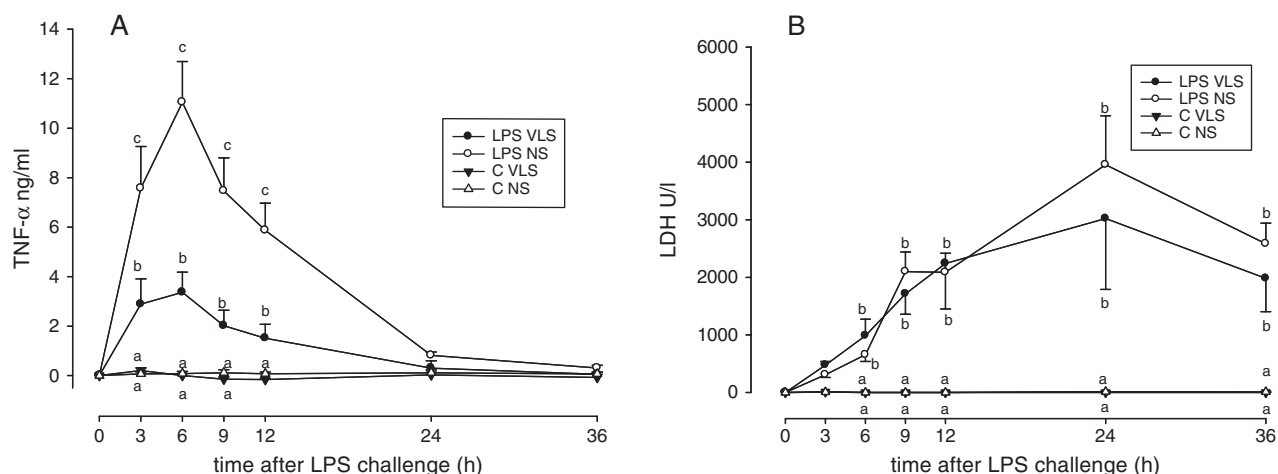


**Fig. 2.** Experiment 1: Proportion of viable cells (%) in the milk at 0 h (■) and 10 h (□) after LPS challenge in C (control without LPS challenge) and LPS quarters of very low SCC (VLS; SCC <20 000 cells/ml) and normal SCC group (NS; SCC = 40 000–100 000 cells/ml). Data are represented as mean  $\pm$  SEM ( $n=5-7$ ). \*: Significant differences between before and after LPS challenge within each group ( $P<0.05$ ). Cell viability increased after LPS challenge with no differences between the groups.

### Experiment 2

At time 0 h no differences within relative mRNA expression of milk cells of all measured genes were found between the controls and treatments and between groups.





**Fig. 3.** Experiment 1: TNF- $\alpha$  concentration (A) and LDH activity (B) in milk after LPS challenge in C quarters (control without LPS challenge) and LPS quarters of very low SCC (VL; SCC <20 000 cells/ml) ( $n=5$  for LPS- and C quarters, respectively) and normal SCC (NS; 40 000–100 000 cells/ml) ( $n=15$  and 7 for LPS- and C quarters, respectively). Data are represented as mean  $\pm$  SEM. *a, b, c*: Different letters indicate significant differences between groups and treatments within one time point ( $P<0.05$ ). If there were no differences between groups and treatments within that time point no letters are given. TNF- $\alpha$  concentration and LDH activity were not different between groups at 0 h and increased in both groups in response to LPS challenge. TNF- $\alpha$  concentration increase in response to LPS was higher in NS compared with VLS. No group differences in LDH activity increase in response to LPS were detected.

At 12 h p.c. an increased mRNA expression of IL-8 (Fig. 4 IL-8) and IL-1 $\beta$  (Fig. 4 IL-1 $\beta$ ) in LPS quarters from both groups was observed. In C quarters IL-8 and IL-1 $\beta$  mRNA expression was decreased at 24 h p.c. in VLS and was not changed at other time points. Lactoferrin mRNA expression (Fig. 4 Lactoferrin) was increased at 36 h p.c. in VLS and C quarters. A decrease of lactoferrin mRNA expression could be detected in LPS-treated quarters at 24 h and 36 h p.c. of the NS group and in LPS-treated quarters at 12, 24 and 36 h p.c. of the VLS group. TNF- $\alpha$  relative mRNA expression (Fig. 4 TNF- $\alpha$ ) of milk cells was decreased at 12 h in C quarters of both groups and was not changed at all time points in all treatments.

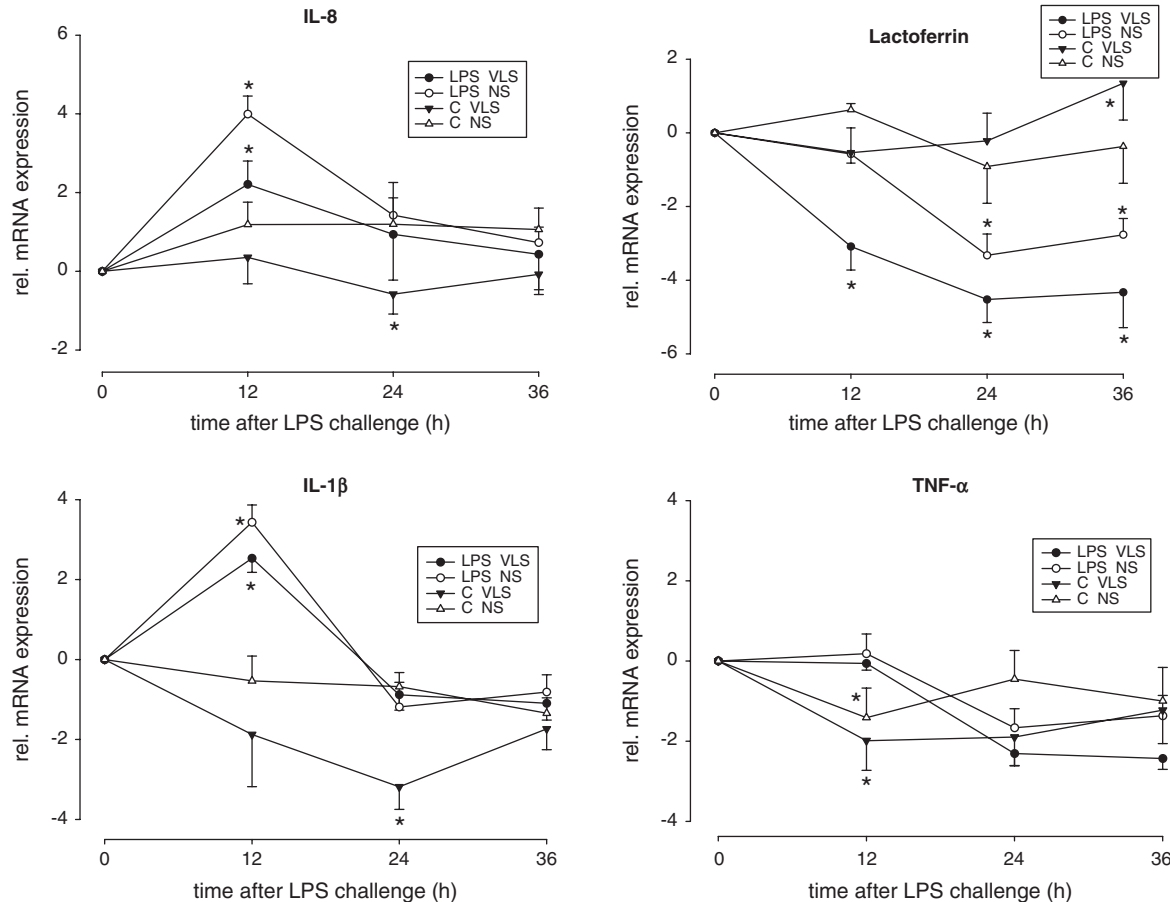
## Discussion

In the present study the immune competence of udder quarters with very low SCC (<20  $\times 10^3$  cells/ml; VLS) was compared with quarters with normal SCC (NS) of (40–100)  $\times 10^3$  cells/ml. Previous investigations already compared low SCC and high SCC but definitions of low SCC varied: 200  $\times 10^3$  cells/ml (Vangroenweghe et al. 2004; Kauf et al. 2006; Olde Riekerink et al. 2007), 150  $\times 10^3$  cells/ml (Schukken et al. 1991; Barkema et al. 1998; Schmitz et al. 2004; Werner-Misof et al. 2007), 100  $\times 10^3$  cells/ml (Paape et al. 1977; Boutet et al. 2004; Koess & Hamman, 2008), 50  $\times 10^3$  cells/ml (Beaudeau et al. 2002), 18  $\times 10^3$  cells/ml (Cheng et al. 2008) or 12  $\times 10^3$  cells/ml (Sarıkaya et al. 2006). However, a direct comparison of very low SCC levels below 20  $\times 10^3$  cells/ml with normal SCC levels of (40–100)  $\times 10^3$  cells/ml that are found very often in dairy practice, in regard to the

immune response of the cells, had not been done before as far as we were aware.

The quarters for the experiments were selected based on their SCC of <20  $\times 10^3$  cells/ml for VLS group and (40–100)  $\times 10^3$  cells/ml for NS group. The selected quarters were of experimental cows with a broad range of lactational stage and number of lactation because other cows were not available. However, all quarters were from udders without any signs of mastitis and they were grouped with normal distribution of number of lactation and lactational stages. If two quarters from one cow were used the treatments of the two quarters within this cow were randomly chosen (control and control, control and LPS treatment, or LPS and LPS treatment). Therefore, results were not influenced by a potential crosstalk between quarters.

For the measurement of responses of SCC and cell viability during an immune response in VLS and NS groups, quarters were stimulated with 50 ng LPS in 10 ml saline solution to induce a clear but not too strong immune response so as to be able to detect possible differences in cell count and viability. To investigate the immune response of the milk cells by mRNA expression of immune factors, a strong immune response is necessary to be able to detect possible differences. Therefore, it was not possible to use the same treatment for the measurements of all factors and two experiments had to be performed with a lower LPS dose for the measurement of SCC, cell viability, and TNF- $\alpha$  concentration and LDH activity in milk, and a higher LPS dose for mRNA expression of immune factors in milk cells. According to Bannerman et al. (2004), Didier & Bruckmaier (2004) and Werner-Misof et al. (2007) SCC of C quarters were stable over the time of the experiment. Therefore, comparisons between C and LPS quarters were possible.



**Fig. 4.** Experiment 2: Relative mRNA expression ( $\Delta\Delta C_p$ ) of IL-8, lactoferrin, IL-1 $\beta$ , and TNF- $\alpha$  in milk cells in C (control without LPS challenge) and LPS quarters of very low SCC (VL; SCC <20 000 cells/ml) ( $n=5$  for LPS and C quarters, respectively) and normal SCC group (NS; 40 000–100 000 cells/ml) ( $n=15$  and 7 for LPS and C quarters, respectively). Data are represented as mean  $\pm$  SEM. \*: Significant differences within groups between LPS treatment and controls ( $P<0.05$ ). There were no considerable differences in mRNA expression of measured immune factors in response to LPS in the cells between both groups.

SCC started to increase simultaneously in both groups 4 h after the challenge with LPS. NS group reached higher SCC levels, although the highest level of SCC was reached 1 h earlier than in VLS group. The time required for SCC increase is the time needed for the recruitment of new cells from blood into milk (Harmon & Heald, 1982; Nickerson & Pankey, 1984; Kehrl & Schuster, 1994). This recruitment of cells was faster in NS quarters than in VLS quarters, which can indicate a more effective immune response.

Like SCC, milk cell viability is crucially involved in the outcome of mammary gland infection. In the present study cell viability was stable during the time of the experiment in quarters that were not stimulated. According to Mehrzad et al. (2001, 2004) the viability in LPS-treated quarters increased from  $42.9 \pm 4.3\%$  to  $88.9 \pm 2.3\%$ . This is due to the increased presence of young leucocytes that migrate from blood into milk and these cells undergo slower apoptosis than older cells (Van Merris et al. 2002). Moreover, LPS stimulation prolongs cell survival (Burvenich et al. 2003). Without stimulation and, therefore,

without this entry of new leucocytes, resident somatic cells have a long storage period (milking interval) in the cisternal cavities. During this time striking changes occur in the morphology of somatic cells owing to the ingestion of fat globules and casein (Paape et al. 1975; Paape & Guidry, 1977). Owing to these morphological and physiological changes, phagocytic and bactericidal activities, as well as cell viability are decreasing and, therefore, the mammary gland defence will be impaired (Burvenich et al. 2007; Rainard & Riollot, 2006). However, no difference in the cell viability between VLS and NS groups was observed after LPS challenge, because at 10 h p.c. the viability of the cells was already nearly 90%. Differentiation of cell populations was for technical reasons not possible in these experiments; however, it is known that the increase of SCC is due mainly to the increase of polymorphonuclear neutrophils (Sarıkaya et al. 2006).

LDH activity increased after LPS challenge in NS quarters and in VLS quarters as expected in response to intramammary LPS administration (Bogin & Ziv, 1973). LDH is

a ubiquitous enzyme that is found in all cells and is an indicator of inflammation as it is released into the extracellular fluid during cell damage and cell death (Glick, 1969). The increase of milk cells after LPS challenge led to a greater number of cells contributing to the LDH activity besides cells from the tissue. The differences in SCC increase between the NS and VLS groups were most likely not great enough to induce differences in LDH increase.

Cytokines and other pro-inflammatory factors produced after pathogen invasion or intramammary stimulation initiate the immune reaction (Rainard & Riollot, 2006). Some of the most important factors were analysed in this study. TNF- $\alpha$  is a cytokine, which serves as a rapidly responding central mediator of inflammatory functions known to play an important role in mastitis and increases in milk after LPS challenge (Paape et al. 2002). It is also involved in the endotoxin shock during an acute phase of coliform mastitis (Sordillo & Streicher, 2002). However, TNF- $\alpha$  mRNA expression in milk cells was not changed at 12 h p.c. TNF- $\alpha$  is a cytokine which is expressed in the early immune response and important for the initiation of the innate immune response. It is likely that the milk cells responded immediately within the first hours after LPS challenge with an increase of TNF- $\alpha$  transcription. Measurement of mRNA expression 12 h p.c., however, could not detect a TNF- $\alpha$  mRNA expression increase because it was already normalized to levels before stimulation. Unfortunately RNA of milk cells earlier than 12 h p.c. was not available because of an insufficient amount of milk. The protein accumulates in the milk and, therefore, an increase 12 h after LPS challenge was detectable although the increased synthesis had already stopped after 12 h. The increase of TNF- $\alpha$  concentration in milk after LPS challenge was significantly higher in NS than in VLS group. This is because more cells could contribute to the production of TNF- $\alpha$  as the SCC was higher in NS group. It is possible that the increased TNF- $\alpha$  concentration in milk could elicit the cure of a potential infection due to a faster and more effective immune response in udder quarters with slightly higher SCC.

IL-8 and IL-1 $\beta$  mRNA expression of milk cells was increased 12 h p.c. in response to LPS treatment with no differences between the groups. This increase was expected as it is known that after intramammary injection of LPS the expression of IL-8 and IL-1 $\beta$  increases (Schmitz et al. 2004; Werner-Misof et al. 2007) in mammary tissue. In addition, Lee et al. (2006) found increasing expression of cytokines in milk cells after intramammary *Esch. coli* stimulation. After 12 h the milk cells that had direct contact with the LPS are removed by milking and, for this reason milk cells at 24 h p.c. did not show an increased cytokine mRNA expression.

Surprisingly, unlike the other factors, relative mRNA expression of lactoferrin in milk cells decreased. Pfaffl et al. (2003) found a lower lactoferrin expression in quarters with SCC  $>150 \times 10^3$  cells/ml than in quarters with SCC  $<150 \times 10^3$  cells/ml. The difference between high-SCC

quarters and low-SCC quarters in Pfaffl et al. (2003) might be compared to the difference between LPS and C quarters of both groups in the present study. Thus, quarters with elevated SCC (with or without stimulation) have similar or lower lactoferrin expression in milk cells than quarters with low SCC. Why lactoferrin mRNA expression in VLS decreased earlier (12 h) than in NS group remains unclear. The increasing lactoferrin expression after intramammary injection of 100  $\mu$ g LPS in mammary tissue that was found by Schmitz et al. (2004) is due to the increased expression of lactoferrin by mammary epithelial cells, the main source of lactoferrin in milk. These cells represent only a very small population of the somatic cells.

In conclusion, the results of the present study showed differences in the immune response between quarters with normal and very low SCC. The time until SCC started to increase after LPS challenge was approximately equal in quarters with normal and with very low SCC. However, quarters with very low SCC did not reach SCC levels as high as quarters with normal SCC. Although increasing less, maximum SCC in response to LPS challenge was reached slightly later in quarters with very low SCC compared with those with normal SCC. The immune response of the milk cells themselves, based on the mRNA expression of the immune factors TNF- $\alpha$ , IL-1 $\beta$ , IL-8 and lactoferrin, was not different between cells of normal and very low SCC groups. Although the selection of experimental quarters was limited, the results suggest that the immune response of udders with slightly higher SCC might be more efficient, as more cells can contribute to the production of immune factors like TNF- $\alpha$ . It is possible that this effect can impair the immune response of udder quarters with very low SCC.

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