

Evaluation of somatic cells in milk of ewes as possible physiological level

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The physiological values of SCC in sheep's milk are still under discussion. The aim of study was to describe the frequency of distribution of ewes on the basis of their individual SCC. The ewes were divided into the five SCC groups (somatic cell count) on the basis of individual SCC (G1 = SCC $200 \times 10^3 \text{ cells.ml}^{-1}$, G2 = SCC between

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1. Introduction

A serious health and economic problem in dairy sheep is mastitis, which causes economic and breeding losses. SCC in the milk of individual dairy sheep can be used as an indicator of husbandry and milking hygiene, the welfare of animals, but especially the udder health and the presence of subclinical mastitis. In dairy sheep the mastitis leads to decrease in milk production and its quality, increase presence of pathogens inappropriate for consumption of dairy products and increase cost for treatment (Riggio and Portolano, 2015). The prevalence of subclinical forms of ewes mastitis ranges from 5 to 50% (Bergonier et al., 2003; Contreras et al., 2007; Olechnowicz and Jaskowski, 2014). From the preliminary results obtained in our dairy practice the incidence of subclinical mastitis ranged from 10–43% (Tančín et al., 2017b). In last mentioned work the criterion for subclinical mastitis was based on SCC (somatic cell count) over

The physiological values of SCC in sheep's milk are still under discussion. Pengov (2001) determined a limit value for physiological SCC in sheep milk of

The aim of study was to describe the frequency of distribution of ewes on the basis of their individual SCC per test day in selected SCC groups during whole lactation. Additionally if high level of somatic cells per lactation could affect somatic cells during following lactation.

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2 Materials and methods

This study was realized by the experimental herd of the NPPC Research Institute for Animal Production Nitra, Slovakia, during 2016 and 2017. Experimental ewes were in their first to third lactations of Slovak dairy sheep (SD) and Lacaune (LC) breeds. Machine milking was performed two times a day in milking parlour 1*16. During each milking the ewes received in parlour 0.1 kg concentrate per head. The milk yield recording and milk sampling were performed once a month during morning milking as a part of milk recording services. Samples of milk were taken in 2016 from April to September and in 2017 from April to August.

Laboratory analysis

Milk samples from each udder were transported to the certificated Central laboratory of Breeding services of the Slovak Republic (Plemenárske služby š.p. Bratislava) for milk analysis.

Categories of somatic cell count (SCC)

For evaluation only ewes with 4 and more sampling during lactation within both 2016 and 2017 were included into study. Thus minimum eight observations were available per animal. A total of 771 milk samples from 73 SD ewes and 17 LC ewes were individually collected. On the basis of individual SCC from milk recording the ewes were divided into the five SCC groups: G1 = SCC <200 × 10³ cells.ml⁻¹, G2 = SCC between 200–400 × 10³ cells.ml⁻¹, G3 = SCC between 400–600 × 10³ cells.ml⁻¹, G4 = SCC between 600–1,000 × 10³ cells.ml⁻¹ and G5 = SCC >1,000 × 10³ cells.ml⁻¹ to evaluate the distribution of individual ewes into SCC groups in different months and years of study. Additionally animals were individually divided into above mentioned SCC groups on the basis of their SCS per lactation (somatic cell score) calculated as a mean from transformed individual SCC data into SCS obtained during milk recording throughout lactation. SCS was calculated according formula $SCS = \text{LOG}_2(\text{SCC}/100)/0.693147 + 3$. Thus distribution of ewes on the basis of SCS into SCC groups was done by conversion of linear scores to somatic cell counts. Mathematical analysis was done by Microsoft Excel program.

3 Results and discussion

On the basis of SCS throughout lactation the most ewes were observed in first two SCC groups (below 400 × 10³ cells.ml⁻¹) in 2016 and 2017 (78.89% and 83.33%, respectively) (Figure 1). In 2016 13 animals (8SD, 5LC) were in SCC groups over >600 × 10³ cells.ml⁻¹, however in next lactation only 6 of them did not improve SCC in next lactation (2017) during dry period, where 5 of them belonged to Lacaune breed. Based on these results, it would be possible for farmers to select such animals for

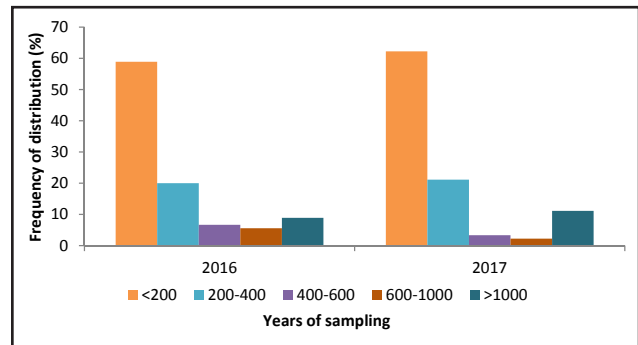


Figure 1 Frequency of ewes in SCC groups (× 10³ cells.ml⁻¹) based on SCS per lactation in 2016 and 2017

culling from the herds on the SCC data. In 2017 12 animals were in above mentioned SCC groups (4SD, 8LC).

Data shown in figure 2 presents frequency of distribution of animals in different months of milk recording in 2016 and 2017, respectively. In both years the most of the ewes were in SCC group below 400 × 10³ cells.ml⁻¹ indicating a good individual udder health of studied ewes. On the other side the percentage of ewes in both last SCC groups (mainly over 1,000 × 10³ cells.ml⁻¹) is relatively low and thus could be considered as health problem of udder. Our data contribute to the findings of Leitner et al. (2008) and Arias et al. (2012) related to physiological level of SCC in ewes.

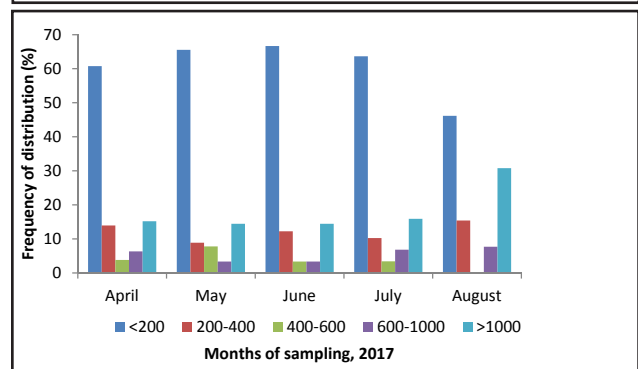
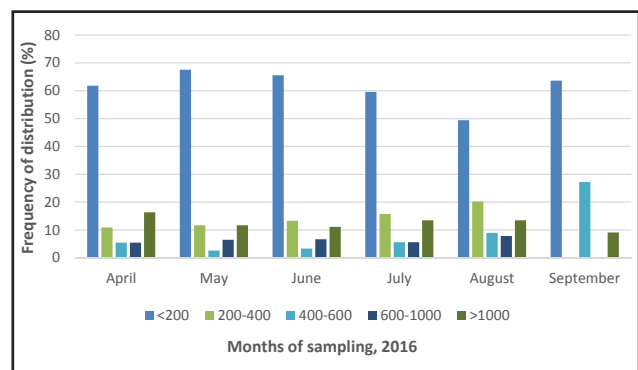


Figure 2 Frequency of ewes in SCC groups (× 10³ cells.ml⁻¹) based on individual SCC in different months of sampling and year

In our previous studies, Idriss et al. (2015) concluded that 78% of the samples were below 600×10^3 cells.ml⁻¹. In the sample group, up to 100×10^3 cells.ml⁻¹, the largest percentage of Tsigai and Improved Valachian (Idriss et al., 2015) were found out. Vršková et al. (2015) found that 76% of Tsigai had SCC below 300×10^3 cells.ml⁻¹. Tančin et al. (2017a) in their study found that 82.03% of the milk samples were below the 400×10^3 cells.ml⁻¹, 71.79% of the milk samples were below the 200×10^3 cells.ml⁻¹ and only 8.89% of the samples were above $1,000 \times 10^3$ cells.ml⁻¹ with the possible effect of breed and farm.

4 Conclusion

The results our study indicated that the most of the animals were in SCC group below 400×10^3 cells.ml⁻¹. In some ewes the high SCC during whole previous lactation negatively influenced SCC in following lactation which could be used in culling program. Possible physiological level of SCC could be taken into account but more data in dairy practice should be evaluated. Thus regular milk recording should include also SCC analysis.

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