

1 **Neutrophil and CD4⁺ milk cell count related to natural incidence of mastitis in Jersey**
2 **cattle**

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4 Supplementary file

5
6 **Material & Methods**

7 *Reagents and milk samples*

8 ATTO620NHS ester, CD4FITC antibody conjugate, ethylenediaminetetraacetic acid
9 disodium salt dehydrate (EDTA) were purchased from Sigma-Aldrich (Germany).

10 Raw cow milk was obtained from quarters of 40 cows from local farms. The tests were
11 performed during the summer. The regulation in force while the analyses were performed was
12 Commission regulation EC No 1662/2006. Approximately 10 ml of each milk sample were
13 collected aseptically. Milk was transported in sterile tubes under refrigerated conditions. All
14 of the analyses were performed on the day of the collecting of the milk samples.

15
16 *Microbial analyses*

17 First of all, after collecting the row milk samples, microbiological examinations were
18 performed. The analyses were made by dairy laboratory “St. George” (Burgas, Bulgaria),
19 conforming to standard **ISO/IEC 17025:2017**. For total bacteria count (TBC) in milk was used
20 horizontal method, skim milk plate count agar with colony count at 30 °C by the pour plate
21 technique (ISO 4833-1:2013). Coliforms were counted by a horizontal method for
22 enumeration and colony-counting technique based on their ability to ferment lactose with

23 production of acid and gas (ISO 4832:2006). For beta-glucuronidase-positive *Escherichia coli*
24 (*E. coli*) was used a colony-count technique at 44 °C and 5-bromo-4-chloro-3-indolyl-β-D-
25 glucuronide (ISO 16649-2:2014). Coagulase-positive staphylococci (CPS) were determined
26 by technique using Baird-Parker medium (ISO 6888-1:1999). The samples were tested for
27 *Salmonella* by four successive stages including pre-enrichment, enrichment, plating out on
28 selective solid media, and confirmation (ISO 6579-1:2017).

29

30 *Somatic cell counting – total count, neutrophils and CD4⁺ cells*

31 Neutrophil cells and CD4⁺ cells were analyzed by immunofluorescence staining. Milk sample
32 was gently mixed and 50 µl of it was loaded in a microcentrifuge tube. After that, 10 µl of the
33 anti-neutrophil antibody – ATTO620 conjugate (1 mg/ml) and 2 µl of the CD4FITC antibody
34 conjugate (1 mg/ml) were added, and 40-min incubation was performed at 4°C. Then, 8 µl of
35 the well-mixed sample were loaded in a chip for automatic cell counting by Lactoscan SCC.
36 After the counting, the rest of the sample was diluted properly with Saline-EDTA (0.9 %
37 NaCl and 7.5 mM EDTA) to obtain about 100 000 cells/ml and 150 µl of it was loaded into a
38 well of polystyrene flat bottom microplate. The analysis was performed by a flow cytometer
39 Guava easyCyte™ 8HT.

40

41 *Lactoscan SCC and Guava easyCyte™ – parameters*

42 All measurements were performed on the automatic cell counter Lactoscan SCC and the flow
43 cytometer Guava easyCyte™ 8HT. Optimization of the parameters for both apparatus was
44 performed prior the analyses. The automatic cell counter Lactoscan SCC was equipped with

45 two lasers, blue laser at 470 nm and red at 627 nm. The parameters of the instrument are
46 shown in Table S1.

47 Table S1. Lactoscan SCC parameters for neutrophil and CD4⁺ cells counting in cow milk.

48 The values are for adjustment of light source 627 nm (for anti-neutrophil antibody –

49 ATTO620 conjugate) and light source 470 nm (for CD4FITC antibody conjugate).

	Laser 627 nm	Laser 470 nm
Exposition	2.110	2.110
Gain	33	3
Power	100%	100%
Focus	2880	2720

50

51 The samples were also analyzed by the flow cytometer (Guava easyCyte™ 8HT), program
52 guava® ExpressPlus. The program allowed adjustment up to three fluorescence parameters
53 (GRN – green, YLW – yellow, RED – red) in combination with forward scatter (FSC) and
54 side scatter (SSC). The samples were run at medium speed and mixing. The counted events
55 for each analysis was 3 000. The instrument used software for the analysis of the data. Target
56 cells – neutrophils and CD4⁺ were gated on dot plots representing cell size based on FSC and
57 granularity based on SSC.

58

59 *Statistical analysis*

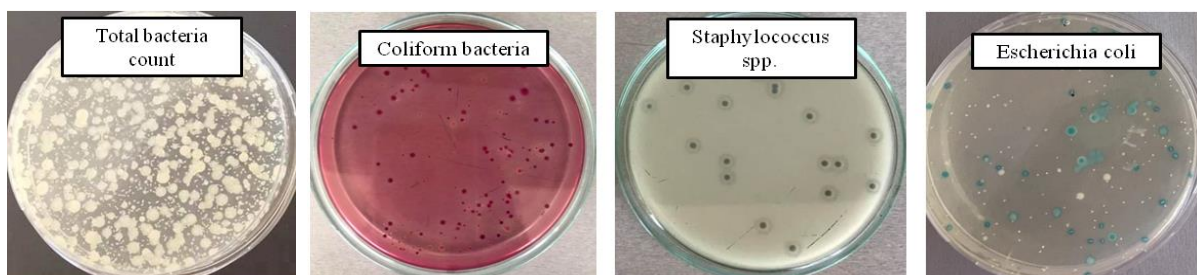
60 Each sample was analyzed triplicate. Analysis and visualization of the data were done with
61 Microsoft Office Excel 2010 (Microsoft Corporation, Redmond, WA, USA). Differences
62 were considered statistically significant at $P < 0.05$.

63

64 Results & Discussion

65 Some of the most major mastitis agents (*Staphylococcus aureus*, *Escherichia coli* and
66 *Coliform* organisms) were selected for the microbiological analyzes (Ameen *et al.* 2019;
67 Zhang *et al.* 2020). *Staphylococcus aureus* mainly produces subclinical and chronic mastitis,
68 but it also may cause per-acute mastitis and lead to gangrene of the quarters. Bacterial toxins
69 are thought to cause the appearance of mastitis and gangrene. *Escherichia coli* cause
70 inflammation of the mammary gland in dairy cows with local and sometimes severe systemic
71 clinical symptoms. It has been found that the severity of *E. coli* mastitis is mainly determined
72 by the host defence status, rather than by *E. coli* pathogenicity (Burvenich *et al.* 2003).
73 *Coliform* bacteria are found in the environment of a dairy cow (Paape & Guidry 1969; Bright
74 *et al.* 1987; Hohmann *et al.* 2020). *Coliform* mastitis is typically acute, but there are also cases
75 of chronic and subclinical infections.

76 The detection methods in this study were according to the relevant ISO standard methods
77 (Fig. S1). Each milk sample was tested for *Salmonella* but there was no sample with positive
78 result.



79

80 Fig. S1. Microbial analyzes of milk samples according to relevant ISO method (Total bacteria
81 count – ISO 4833-1:2013, Coliforms – ISO 4832:2006; Coagulase-positive staphylococci –
82 **ISO 6888-1:1999**, beta-glucuronidase-positive *Escherichia coli* – ISO 16649-2:2014).

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84 Table. S2. Bacteria and somatic cell count in three groups of milk samples.

Milk	SCC, x 10 ⁵ cells/mL	TBC, cfu/mL	<i>Staphilococcus</i> <i>spp.</i>	<i>Escherichia</i> <i>coli</i>	<i>Coliforms</i>
Healthy (n=16)	1.3 – 4.2 GM: 3.0	29 000 ± 15%	Negative (< 10 cfu/mL)	Negative (< 10 cfu/mL)	Negative (<< 1000 cfu/mL)
Dirty (n=12)	1.7 – 4.3 GM: 3.2	918 180 ± 5%	Positive (100 ± 35% cfu/mL)	Positive (1 200 ± 30% cfu/mL)	Positive (220 000 ± 30% cfu/mL)
Mastitic (n=75)	2.81 – 16.5 GM: 6.0	120 000 ± 15%	Positive	Positive	Positive

85 *GM – geometric mean; SCC – somatic cell count (determined by Lactoscan SCC); TBC –
86 total bacteria count.

87 In mastitic milk samples the neutrophil count was significantly higher than that in healthy
88 group of milks. Similar results were obtained in other studies (Riollet *et al.* 2001; Alhussien *et*
89 *al.* 2015).

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