Effects of addition of somatic cells to caprine milk on cheese quality

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**A B S T R A C T**

Milk quality criteria and limits for somatic cell count (SCC) established in many countries make it difficult to maintain SCC of bulk tank goat milk below the threshold, due to non infection factors linked to goat physiology. The aim of this study was to objectively verify the effects of SCC on fresh caprine milk cheese. Somatic cells were recovered from pooled healthy goat milk, and added to low SCC raw or pasteurised goat milk. Miniature cheeses were made and evaluated after 1 and 7 d. Somatic cells had a major effect on lipolysis, increasing free fatty acids regardless of whether milk was raw or pasteurised. The effect of somatic cells on proteolysis was specific for caseins and the effects were different if cheeses are made from raw or pasteurised milk. It is concluded that somatic cells themselves in caprine milk can directly affect some cheese parameters.

1. **Introduction**

Currently, there is an increased demand for caprine and diverse products. Ribeiro and Ribeiro (2010) and Silakinove, Leitner, Merin, and Prosser (2010) reviewed the healthy properties of caprine milk and products. According to Food and Agriculture Organization of the United Nations (FAOSTAT, 2012), the worldwide dairy goat population reached 191 million in 2010, and caprine milk production surpassed 16.7 million tons, representing significant increases of 25 and 31%, respectively, compared with a decade ago. The populations of dairy goats in the USA and the European Union in 2010 were 356,000 and 7,732,631, respectively.

Milk quality criteria are part of payment systems for milk, thereby ensuring good quality finished products (Raynal LJutovac, Gaborit, & Lauret, 2005). In the USA, the limit maximum for SCC (somatic cell count) established by the Food and Drug Administration is 750,000 cells mL\textsuperscript{-1} and 1,000,000 cells mL\textsuperscript{-1}, respectively for bovine and caprine milk. In the European Union (Directive 92/ 46ECC Council, 1992), the legal limit for bovine milk is 400,000 cells mL\textsuperscript{-1} and there is no legal limit for caprine and ovine milk. Some authors (Paape et al., 2007; Raynal LJutovac, Pirisi, de Crémoux, & Gonzalo, 2007) have reported that North American goat producers have difficulty maintaining bulk tank goat milk SCC below the threshold limit due to non infectious factors, such as breed, parity, stage of lactation, oestrus, diurnal, monthly and seasonal variations (Fernández, Rodriguez Alcala, Abilleira, de la Fuente, & Fontecha, 2009; Haenlein & Hinckley, 1995; Paape, Poutrel, Contreras, Marco, & Capuco, 2001) that have major impact on counts in caprine milk; this should be taken into account when the European Union establishes limits for SCC in caprine milk. Numerous studies have shown that increased SCC in bovine milk results in decreased milk production and affects milk composition, which causes a reduction in the ability to produce cheese and decreases the shelf life of consumer milk (Erdem, Atasever, & Kul, 2010; Ma, Barbano, & Santos, 2003; Politis & Ng Kwai Hang, 1988). The main component of the economic loss associated with SCC is the reduced milk production in cows with mastitis (Huijs, Lam, & Hogeveen, 2008). Raynal LJutovac et al. (2005) concluded that somatic cells, although they are an indicator of the hygienic quality of goat milk, do not seem to necessarily impair cheese making. However, Raynal LJutovac et al. (2007) intensively reviewed analytical, sanitary, productive and technological effects of somatic cells on caprine milk and cheese, and controversies among the conclusions of the studies were found, probably due to the different variables, such as the variety of cheese making or using different animals in each experimental group.

Despite these differences, it is not clear what the effects of somatic cells are on milk properties, cheese quality, lipolysis and
proteolysis. Marino, Considine, Sevi, McSweeney, and Kelly (2005) recovered somatic cells from mastitic bovine milk and added them to good quality milk, then evaluated their contribution to proteolysis during Cheddar cheese ripening; these authors concluded that proteolytic enzymes of somatic cells from milk contributed directly to proteolysis in milk and cheese. Caroprese et al. (2007) studied the effect of macrophages on proteolysis and plasmnin activity in bulk ovine milk. Somatic cells contain a range of lysosomal enzymes, such as cathepsin B, cathepsin D, cathepsin G and elastase (Le Roux, Laurent, & Moussaoui, 2003), many of which pass into the milk, where they cause increased proteolysis and lipolysis (Albenzio et al., 2005; Le Roux, Colin, & Laurent, 1995). Many somatic cell enzymes have been found or measured in fractions of milk other than the cells themselves, suggesting that leakage or secretion of enzymes occurs, or that the cells lyse and release their intracellular enzymes (Kelly & Fox, 2006). Moreover, increasing SCC is usually accompanied by imbalances in other parameters in milk, so it is very difficult to determine if somatic cells are directly responsible for changes in milk and cheese.

Milk pasteurisation is commonly practiced before making cheese. This heat treatment inactivates microorganisms, activates—-inactivates indigenous milk pro enzymes and enzymes, partially denatures whey proteins, and modifies starter bacteria activity. It also concomitantly reduces fermentation and degradation reactions (Buchin et al., 1998; Grappin & Beuvier, 1997). Marino et al. (2005) reported that proteolysis during ripening of cheese increased with addition of somatic cells, although this effect was reduced by pasteurisation after cell addition. A relationship between proteolysis in milk and SCC was described by Hachana and Kraiem (2009).

In the Canary Islands, most caprine milk cheese is made with raw milk, using traditional methods (Fresno et al., 2008). The islands have an exceptional sanitary situation, due to being officially free of caprine and ovine brucellosis. This fact allows approximately 500 artisan producers to sell raw milk cheese after less than 60 days of ripening. Overall, few studies about the direct effect of somatic cells from healthy animals on cheese ripening have been reported in the literature, and no objective study has been reported about the contribution of somatic cells to cheese made following a simple rennet cheese making process using raw caprine milk. The aims of this study were to objectively verify the effects of somatic cells on chemical composition, proteolysis and lipolysis profile of fresh cheese made with raw or pasteurised caprine milk.

2. Material and methods

2.1. Recovery of somatic cells

Somatic cells were recovered by centrifugation (280 × g, 20 min, 4 °C) (Beckman, J2 CI Centrifuge, Beckman Coulter Inc., FL, USA) from high SCC raw milk (approximately 1.78 × 10^6 cells mL^-1) from two healthy Majorera breed goats from the Animal Science Unit of Universidad de Las Palmas de Gran Canaria. The resulting pellet was resuspended in phosphate buffered saline at pH 6.8, as described by Verdi and Barbano (1991), until a pool with approximately 2.1 × 10^7 SCC mL^-1 was obtained. The SCC of this stock preparation, before and after dilution in phosphate buffered saline (PBS), was determined using a DCC Device counter (DCC, DeLaval International AB, Tumba, Sweden). The type of somatic cell in the pool was not specifically determined, but is likely that the somatic cell population in caprine milk is dominated by polymorphonuclear neutrophils (45–74%), followed by macrophages (15–41%) and lymphocytes (9–20%) (Dulin, Paape, Schultz, & Weinland, 1983). This process was repeated every day before making cheese from the same goats.

2.2. Cheese production

Miniature cheese systems have been proposed as the best alternative to pilot plant experiments, being closer to cheese than any other model proposed to date; moreover, they can be prepared under controlled conditions and are more economical, reproducible and easier to obtain (Shakeel Ur Rehman, Fox, McSweeney, Madkor, & Farkye, 2001).

2.2.1. Cheese formulation

Raw goat milk with low SCC (approximately 2 × 10^5 SCC mL^-1) was used to make the miniature cheeses. Over 10 L of milk was obtained from four goats from a healthy experimental herd of Majorera dairy goat breed. The four batches of experimental cheeses were produced using six different experimental treatments (three levels of SCC and pasteurisation or non pasteurisation). Each day the raw caprine milk was separated into two groups: raw milk or pasteurised milk (63 °C for 30 min). Both groups were then divided into three subgroups with different levels of somatic cells added: low SCC milk (without somatic cells added), medium SCC milk (mean value approximately 1 × 10^6 mL^-1), and high SCC milk (mean value approximately 2.5 × 10^6 mL^-1). Somatic cells were added after pasteurisation of milk. The finished cheeses were ripened for 1 and 7 d in duplicate, resulting in a total of 96 cheese samples (4 batches × raw or pasteurised milk × 3 level of SCC × 2 ripening times × duplicate).

2.2.2. Manufacture of miniature cheese

Cheeses were made with six different milk treatments: raw milk low SCC (RL) cheese, raw milk medium SCC milk (RM) cheese, raw milk high SCC (RH) cheese, pasteurised milk low SCC (PL) cheese, pasteurised milk medium SCC (PM) cheese, and pasteurised milk high SCC (PH) cheese. Milk was salted with 4 g NaCl for 100 mL milk. The same procedure was used to make each type of miniature cheese: 40 mL of salted milk was poured into four 50 mL centrifuge tubes, which were then placed in a water bath at 32 °C. Commercial animal rennet (Marshall rennet powder, Rhône Poulenc TTeXel, Dangé Saint Romain, France), comprising 50% pepsin and 50% chymosin, was used. No starter cultures were added. Curd was cut 45 min after rennet addition with a stainless steel spatula and allowed to stand for additional 5 min. The curds were then centrifuged at 3400 × g for 20 min and the whey was discarded. The non packaged cheeses were divided randomly into two groups and allowed to ripen 1 or 7 d at 10–12 °C and 75–80% relative humidity.

2.3. Gross composition, pH and colour measurement of cheese

Cheese pH was measured was determined using a Crison 507 pH meter at 3 internal (in the cheese centre) locations for each cheese. After that, cheese samples were cut into small pieces, weighed and freeze dried. The difference in weight before and after the freeze drying was used to calculate the percentage moisture content. The rest of the analysis was performed with the resultant dried matter. Soxhlet extraction was performed to quantify the fat content in dry basis using a Büchi 810 Soxhlet fat extraction apparatus (Büchi Labortechnik, Flawil, Switzerland). Protein content was determined in duplicate as %N × 6.28 using a LECO CNS 2000 nitrogen analyser (Leco Corporation, St. Joseph, MI, USA).

2.4. Extraction of proteins and electrophoretic analysis

All chemical reagents were of analytical grade from Panreac (Barcelona, Spain), Merck (Darmstadt, Germany), and Bio Rad Laboratories (Hercules, CA, USA). Water soluble proteins were
extracted from the cheeses after 1 and 7 d of ripening using the method described by Tunick, Malin, Smith, and Hilsinger (1995) as modified by Sánchez Macías et al. (2011).

Lyophilised samples were prepared, separated by SDS PAGE (Laemmli, 1970), and stained as described by Sánchez Macías et al. (2011). Images of stained gels were captured with a Bio Rad imaging device (Gel Doc EQ; Bio Rad Laboratories), and bands were quantified using Quantity One Quantitation Software (Bio Rad Laboratories). Densitometry analysis of the gels was performed using lane based background subtraction followed by measurement of the bands by the area under the intensity profile curve, and the values were then used for statistical analysis. Each sample was analysed on duplicate gels. Individual protein species from cheese samples were identified by comparing their relative mobilities with those of standard proteins (Bio Rad Laboratories) from a previous SDS PAGE gel.

2.5. Total free fatty acids (copper soap method)

The total free fatty acid (FFA) content was determined using the copper soap method (Mellilli et al., 2004; Shipe, Senyk, & Fountain, 1980), modified for cheese analysis as described by Sánchez Macías et al. (2011). The final results were expressed as milligrams of FFA per 100 g of dry matter (DM) of cheese and were also converted to milligrams of FFA per g of fat in cheese.

2.6. Statistical analysis

Statistical analyses were performed using the SAS program package (Version 9.00, SAS Institute Inc., Cary, NC, USA). A PROC MIXED procedure (analysis of variance with repeated measures) was used to evaluate the effect of different SCC level and ripening time on the chemical composition, pH, lipolysis level, and level of primary proteolysis of the raw or pasteurised goat milk cheeses. Significantly different means were identified using the Tukey's test.

3. Results and discussion

3.1. Gross chemical cheese analysis and pH values

The gross chemical composition and pH of cheeses after 1 and 7 d of ripening are shown in Table 1. At day 1 of ripening, the moisture was higher in control pasteurised milk cheese than its counterpart made with raw caprine milk, though no differences were found between RH and PH cheeses. These data concur with the findings of Pears, Linklater, Hall, and Mackinlay (1985), who reported an adverse effect of heat treatment on syneresis of rennet induced milk gels using model systems. This effect on syneresis is probably due to the presence of denatured whey protein on the surface of the micelles, which retard aggregation and fusion of the rennet treated micelles (Rynne, Beresford, Kelly, & Guine, 2004). Henry, Mollé, Morgan, Fauquant, and Bouhallab (2002) reported that more water is bound in cheese when β-lactoglobulin–κ casein complexes are formed after pasteurisation of milk. Moisture content decreased after 7 d of ripening in all cheeses because of surface water evaporation, and no differences were found between cheeses. Marino et al. (2005) suggested in a similar study that somatic cells themselves may contribute to the increase in moisture content in Cheddar type cheeses ripened for 15 days, and they proposed that the inclusion of cells may physically interfere with the process of curd syneresis of this type of cheese. Fat percentage in DM was within the ranges required by Spanish regulation for full fat (60–45% fat in DM) cheese designation, and very similar to those found by Sánchez Macías et al. (2010) in cheese made from milk from the same breed.

Protein content on a DM basis was similar among the different cheeses in the study; however, an increase of this parameter was found after 7 d of ripening. van Vliet and Walstra (1994) reported that about 90% of the water present in milk gels is mechanically enclosed between the casein aggregates, which determines the ease of removal of water from the gel. The increased protein content measured on a DM basis can be explained due to the more possible water that is released and available after 7 d of ripening, which can be extracted from cheeses after the casein network is broken down due to proteolysis. As cheeses were not packaged, some available water evaporated from their surfaces. Cheese pH was approximately 6.6 at 1 d of ripening, similar to the typical values reported for Majorera caprine milk cheese (Álvarez et al., 2007; Sánchez Macías et al., 2010). During the first week of ripening, pH values decreased in all cheese groups, though PL cheese had a significantly higher value than the others. Pirisi et al. (2000) and Viana, Mazal, Santos, Bolini, and Gigante (2008) reported that higher milk SCC resulted in increased pH in cheese, while other authors (Jaeggi, Govindasamy Lucey, Berger, & Johnson, 2003) did not observe significant variations, possibly due to other changes in the milk that accompanied the increasing SCC. Marino et al. (2005), on adding somatic cells from mastitic milk to control milk, did not find significant differences of pH in cheeses in

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Time</th>
<th>Treatment</th>
<th>RL</th>
<th>RM</th>
<th>RH</th>
<th>PL</th>
<th>PM</th>
<th>PH</th>
<th>SEM</th>
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</thead>
<tbody>
<tr>
<td>Moisture (%)</td>
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<td>48.86&lt;sup&gt;a&lt;/sup&gt;</td>
<td>49.65&lt;sup&gt;a&lt;/sup&gt;</td>
<td>51.98&lt;sup&gt;a&lt;/sup&gt;</td>
<td>55.27&lt;sup&gt;a&lt;/sup&gt;</td>
<td>55.02&lt;sup&gt;a&lt;/sup&gt;</td>
<td>53.17&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0.57</td>
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</tr>
<tr>
<td>Fat in DM (%)</td>
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<td>44.19</td>
<td>44.72</td>
<td>44.18</td>
<td>43.43</td>
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<td>0.61</td>
<td></td>
</tr>
<tr>
<td>Protein in DM (%)</td>
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<td>36.58&lt;sup&gt;b&lt;/sup&gt;</td>
<td>36.30&lt;sup&gt;b&lt;/sup&gt;</td>
<td>35.50&lt;sup&gt;b&lt;/sup&gt;</td>
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<td></td>
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<tr>
<td>pH</td>
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<td>38.81&lt;sup&gt;a&lt;/sup&gt;</td>
<td>40.13&lt;sup&gt;a&lt;/sup&gt;</td>
<td>39.61&lt;sup&gt;a&lt;/sup&gt;</td>
<td>38.05&lt;sup&gt;a&lt;/sup&gt;</td>
<td>38.21&lt;sup&gt;a&lt;/sup&gt;</td>
<td>39.48&lt;sup&gt;a&lt;/sup&gt;</td>
<td>0.43</td>
<td></td>
</tr>
</tbody>
</table>

<sup>a</sup> Values are means; values with different lower case superscript letters within a row and values with different upper case superscript letters within a column differ (P < 0.05) for the parameter; SEM, standard error of the mean.

DM, dry matter.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Low SCC</th>
<th>Medium SCC</th>
<th>High SCC</th>
<th>Low SCC</th>
<th>Medium SCC</th>
<th>High SCC</th>
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<tbody>
<tr>
<td>RL</td>
<td>48.86&lt;sup&gt;a&lt;/sup&gt;</td>
<td>49.65&lt;sup&gt;a&lt;/sup&gt;</td>
<td>51.98&lt;sup&gt;a&lt;/sup&gt;</td>
<td>55.27&lt;sup&gt;a&lt;/sup&gt;</td>
<td>55.02&lt;sup&gt;a&lt;/sup&gt;</td>
<td>53.17&lt;sup&gt;a&lt;/sup&gt;</td>
</tr>
<tr>
<td>RM</td>
<td>44.19</td>
<td>44.72</td>
<td>44.18</td>
<td>43.43</td>
<td>43.83</td>
<td>44.32</td>
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<tr>
<td>RH</td>
<td>35.53&lt;sup&gt;a&lt;/sup&gt;</td>
<td>36.58&lt;sup&gt;b&lt;/sup&gt;</td>
<td>36.30&lt;sup&gt;b&lt;/sup&gt;</td>
<td>35.50&lt;sup&gt;b&lt;/sup&gt;</td>
<td>35.34&lt;sup&gt;b&lt;/sup&gt;</td>
<td>36.72&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>PL</td>
<td>38.81&lt;sup&gt;a&lt;/sup&gt;</td>
<td>40.13&lt;sup&gt;a&lt;/sup&gt;</td>
<td>39.61&lt;sup&gt;a&lt;/sup&gt;</td>
<td>38.05&lt;sup&gt;a&lt;/sup&gt;</td>
<td>38.21&lt;sup&gt;a&lt;/sup&gt;</td>
<td>39.48&lt;sup&gt;a&lt;/sup&gt;</td>
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<tr>
<td>PM</td>
<td>56.74&lt;sup&gt;ab&lt;/sup&gt;</td>
<td>56.86&lt;sup&gt;ab&lt;/sup&gt;</td>
<td>55.37&lt;sup&gt;ab&lt;/sup&gt;</td>
<td>62.36&lt;sup&gt;ab&lt;/sup&gt;</td>
<td>58.34&lt;sup&gt;ab&lt;/sup&gt;</td>
<td>57.25&lt;sup&gt;ab&lt;/sup&gt;</td>
</tr>
<tr>
<td>PH</td>
<td>0.12</td>
<td>0.12</td>
<td>0.15</td>
<td>0.06</td>
<td>0.12</td>
<td>0.13</td>
</tr>
</tbody>
</table>

Values are means; values with different lower case superscript letters within a row and values with different upper case superscript letters within a column differ (P < 0.05) for the parameter; SEM, standard error of the mean.

DM, dry matter.

Treatments: RL, raw goat milk with low SCC; RM, raw goat milk with medium SCC; RH, raw goat milk with high SCC; PL, pasteurised goat milk with low SCC; PM, pasteurised goat milk with medium SCC; PH, pasteurised goat milk with high SCC.
this study, somatic cells from healthy goats were used, and cheese pH decreased as somatic cells were added to the caprine milk.

3.2. Lipolysis during cheese ripening

Mean levels of FFA are summarised in Table 2. The indigenous lipoprotein lipase (LPL) hydrolyses the 1,3 position in mono, di, and triglycerides to give free fatty acids. Large proportions of caprine milk LPL are bound to cream, and not to caseins, unlike in bovine milk, which could explain why lipolysis in milk is well correlated with milk LPL activity in this species (Chilliard, Selselet Attou, Bas, & Morand Ferh, 1984). FFA levels did not differ significantly among the cheeses after 1 d of ripening; however, slight increases in FFA content were found as SCC increased in cheese milk.

Quantitative analysis of the total FFA content showed a general increase in all cheeses after 7 d of ripening, and the levels were higher as SCC increased in milk. Moreover, raw caprine milk cheese had higher levels of FFA than pasteurised caprine milk cheese, for both low SCC samples. Other authors have found lower levels of FFA in cheese on comparing pasteurised and raw milk cheeses: 38% and 50% lower FFA content in Manchego Cheese (Gaya, Medina, Rodríguez Marin, & Nuñez, 1990) and Cheddar cheese (McSweeney, Fox, Lucey, Jordan, & Cogan, 1993), respectively. Morgan and Gaborit (2001) reported that the processing of the caprine milk (cheese, yoghurt), physical treatment, and cold storage increased the lipolysis level in the milk, whereas pasteurisation decreased the level of lipolysis. Medium and high SCC milk cheeses had similar FFA content for both raw and pasteurised milk. Chen, Wang, Van Kessel, Ren, and Zeng (2010) reported that SCC in pasteurised goat milk cheese did not significantly affect FFA content, and similar results were obtained by Jaeggi et al. (2003) for hard ewe milk cheese. In both cases, cheese milk was collected from animals grouped according to SCC and, thus, not only somatic cells themselves were involved in lipolysis in cheese, but other changes in milk. Moreover, in the present study, somatic cells were added to the milk after pasteurisation, which means that somatic cell enzymes were not affected by heating, and lipolysis clearly depended on somatic cells. Lipolysis only occurs when the membrane of the fat globules is broken, and the fat is accessible to the enzyme. Azzara and Dimick (1985) found that macrophages from mastitic bovine milk secreted lipolytic enzymes which may bind to the fat globule membrane and damage it, exposing the fat to degradation by LPL.

In cheese, lipolysis results in the formation of FFA that may contribute to cheese flavour and also serve as substrates for further reactions, with the consequence of producing highly flavoured catabolic end products (Collins, McSweeney, & Wilkinson, 2003). Jaubert, Bodin, and Jaubert (1996) observed a positive correlation between lipolysis and SCC in goat milk, and a more intense caprine flavour with increasing SCC was found in caprine milk. Sensory studies are needed to evaluate if somatic cells themselves are able to affect the sensory characteristics of cheese made from caprine milk.

3.3. Protein profile and proteolysis during cheese ripening

A densitometer derived protein distribution analysis is summarised in Table 3. Primary proteolysis essentially measures the extent of breakdown of caseins, which occurs mainly as a result of the action of residual coagulant, plasmin (PL), and somatic cell proteases. However, microorganisms are also active in the hydrolysis of caseins (Fox, 1993; Fox & McSweeney, 1996). Harmon (1994) and Moslehiyad and Ezzatpanah (2010) described that the increasing SCC of milk affects casein micelle microstructure and properties due to hydrolysis of casein. Due to the increasing proteolytic activity of enzymes such as PL, neutral, and acidic proteases, therefore, casein could be more susceptible to hydrolysis.

During ripening, β CN hands had the highest intensity values, followed by αs2 CN and αs1 CN. Caprine milk contains relatively little αs1 CN, and β CN is the major protein (Tziboula Clarke, 2003). Quantitative analysis showed a general reduction in casein fractions as ripening progressed, and hydrolysis was higher for β CN (26–43% reduction), than for para κ CN (23–37%), αs2 CN (20–30%), and αs1 CN (14–25%). Pasteurised caprine milk cheese had higher levels of caseins compared with raw caprine milk cheese when no somatic cells were added. On the other hand, different patterns of casein breakdown were found as SCC was increased in raw or pasteurised goat milk cheeses.

Regarding αs2 CN, this was degraded faster in RM and RH cheese than in RL cheese, when expressed in absolute values. However, when expressed as percentage breakdown from 1 to 7 d, a faster breakdown of this casein was observed in RM and RH compared to RL cheese. On the other hand, hydrolysis of αs2 CN was accelerated in pasteurised caprine milk cheese when somatic cells were added.

The level of αs1 CN in cheese was not affected by heat treatment or somatic cell addition; however a lower content was found after 7 d of ripening in all cheese types. Marino et al. (2005) found that adding somatic cells from mastitic milk to cow milk resulted in faster hydrolysis of αs1 CN in cheese at 15 d of ripening than in control cheese, indicating accelerated proteolysis of this casein.

In this study, β CN content was similar in all cheese groups after 1 d of ripening, and a significant reduction of this casein was found at 7 d of ripening. Proteolysis of β CN was similar in RL and PL. There were no differences in β CN breakdown between raw milk cheeses. In contrast, the rate of β CN breakdown increased with increasing SCC in pasteurised milk cheese. It is notable that, in pasteurised milk cheeses, the breakdown of β CN was higher than its respective cheeses made with raw caprine milk. When isolated and lysed macrophages from bulk ewe milk were incubated with sodium caseinate, changes in percentage composition of intact casein were observed: β CN did not undergo appreciable hydrolysis, unlike α CN (about 20%) (Caroprese et al., 2007).

Plasmin (PL), the main indigenous proteolytic enzyme in milk, is associated with the casein micelles, and milk contains the entire PL system: PL, plasminogen (PG), PL inhibitors, PG activators (PAs), and inhibitors of PAs. PAs have been associated with somatic cells fraction in goat milk (Politis et al., 1994): PL acts mainly on β casein, which constitutes approximately 60% of the total casein in caprine milk.

Table 2

Total free fatty acid (FFA) levels in cheese made with (raw or pasteurised) caprine milk at three levels of somatic cell count (SCC) after 1 and 7 d of ripening.

<table>
<thead>
<tr>
<th>Time</th>
<th>Treatment</th>
<th>RL</th>
<th>RM</th>
<th>RH</th>
<th>PL</th>
<th>PM</th>
<th>PH</th>
<th>SEM</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Total FFA (mg) per 100 g of DM</td>
<td>52.92a</td>
<td>54.21b</td>
<td>60.96b</td>
<td>48.59a</td>
<td>56.26a</td>
<td>63.75a</td>
<td>3.14</td>
</tr>
<tr>
<td>7</td>
<td>Total FFA (mg) per 100 g of DM</td>
<td>243.5b</td>
<td>441.01b</td>
<td>641.5b</td>
<td>113.9b</td>
<td>468.10b</td>
<td>581.34b</td>
<td>35.10</td>
</tr>
<tr>
<td>SEM</td>
<td>Total FFA (mg) per 100 g of DM</td>
<td>33.09</td>
<td>55.19</td>
<td>86.11</td>
<td>11.60</td>
<td>59.52</td>
<td>72.49</td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>Total FFA (mg) per g of fat in cheese</td>
<td>1.13a</td>
<td>1.17a</td>
<td>1.31a</td>
<td>1.04a</td>
<td>1.20a</td>
<td>1.36a</td>
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</tr>
<tr>
<td>7</td>
<td>Total FFA (mg) per g of fat in cheese</td>
<td>5.22b</td>
<td>9.41b</td>
<td>13.81b</td>
<td>2.47b</td>
<td>10.14b</td>
<td>12.49b</td>
<td>0.54</td>
</tr>
<tr>
<td>SEM</td>
<td>Total FFA (mg) per g of fat in cheese</td>
<td>0.70</td>
<td>1.17</td>
<td>1.83</td>
<td>0.24</td>
<td>1.21</td>
<td>1.47</td>
<td></td>
</tr>
</tbody>
</table>

* Values are means; values with different lower case superscript letters within a row and values with different upper case superscript letters within a column differ (P < 0.05) for the parameter; SEM, standard error of the mean.

* Treatments: RL, raw goat milk with low SCC; RM, raw goat milk with medium SCC; RH, raw goat milk with high SCC; PL, pasteurised goat milk with low SCC; PM, pasteurised goat milk with medium SCC; PH, pasteurised goat milk with high SCC.

* DM, dry matter.


**Table 3**

<table>
<thead>
<tr>
<th>Treatment</th>
<th>SCC (×10^6)</th>
<th>Time</th>
<th>Percentage proteolysis of CN</th>
<th>Percentage proteolysis of CN</th>
<th>Percentage proteolysis of CN</th>
<th>Percentage proteolysis of CN</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>0.25</td>
<td>1.0</td>
<td>4.0</td>
<td>7.0</td>
</tr>
<tr>
<td>Raw milk</td>
<td>1.85 × 10^6</td>
<td>SEM</td>
<td>1.80</td>
<td>2.32</td>
<td>4.35</td>
<td>5.14</td>
</tr>
<tr>
<td>Pasteurised milk</td>
<td>2.12 × 10^6</td>
<td>SEM</td>
<td>1.75</td>
<td>2.14</td>
<td>3.85</td>
<td>4.65</td>
</tr>
</tbody>
</table>

*Values are means ± standard error of the mean (SEM) of 3 replicates.*

**References**


**Figure 4**

4. Conclusions

- Lipolysis in cheese is clearly dependent on somatic cell counts, which play a critical role in determining the quality of raw milk. The use of high-quality raw milk, with a lower somatic cell count, leads to higher-quality cheeses.
- Brie and G. lactis strains have been shown to be responsible for this process, which is important for the production of high-quality cheeses.

**Figure 5**

5. Discussions

- The influence of somatic cell count on the quality of raw milk is a critical factor in determining the quality of pasteurised milk. The use of high-quality raw milk, with a lower somatic cell count, results in a higher-quality pasteurised milk.

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The reference text is not provided.


