

# Microbial characteristics of *Conciato Romano*: an artisanal cheese made from raw sheep's milk

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## **Abstract**

The aim of this study was to assess the microbial characteristics of a batch of Conciato Romano during manufacturing and ripening. Conciato Romano is a traditional cheese made from raw sheep's milk without starter cultures in the province of Caserta (Southern Italy) using traditional methods. A total of 7 samples (raw milk, curd and cheese wheels taken after 25, 60, 120 and 180 days of ripening) were screened for hygiene indicators microorganisms counts (total viable Enterobacteriaceae, total coliforms, E. coli, clostridia sulphite reducing, yeasts, coagulasepositive staphylococci, enterococci), for autochthonous lactic acid flora counts (mesophilic and thermophilic lactococci and lattobacilli), and also for Salmonella spp. and Listeria monocytogenes presence. In raw milk, low values were detected for total aerobic flora (3.2 log cfu/mL), Enterobacteriaceae and total coliforms (2 cfu/mL), and the autochthonous starter lactic flora was predominant (3.2 log cfu/mL). During ripening, total aerobic flora was constant (107-108 cfu/g); total coliforms, E. coli. Enterobacteriaceae and veasts were not detected starting from the 60th day of ripening. Enterococci ranged from 4.2 to 6.2 log cfu/g. The mesophilic lactic flora was dominant with values always >6 log cfu/g during the whole ripening period. Pathogens were never detected. The results of this study highlighted how the raw milk indigenous lactic flora, the traditional production techniques and the cheesemaker's experience are essential to guarantee the unique nature of *Conciato Romano*.

# Introduction

Conciato Romano is an artisanal cheese produced in the province of Caserta, Southern Italy. The technique of aging it in clay jars probably dates back to Samnite civilisation. The traditional process of production involves the use of raw sheep milk without starter cultures,

so the maturation of cheese is a direct consequence of the metabolic activity of the lactic flora coming from the environment and milk whose organoleptic properties are influenced by the aromatic plants of the mountains around Caserta where the sheep graze (Buchin et al., 1999). The microbial ecology of artisanal cheeses widely affects the production process and its quality. The study of the microbiological characteristics of these cheeses is fundamental in order to improve their quality in accordance with current legislation and to preserve, at the same time, the microbial biodiversity and their characteristic nature. Therefore, the purpose of this study was to evaluate the microbial ecology of Conciato Romano during production and ripening.

#### **Materials and Methods**

The experiment began in February 2012 at the Le Campestre farm located in the municipality of Castel di Sasso, Caserta Province, Southern Italy. In accordance with the traditional methods used in the area and the production capacity of small processing dairy farm, one batch of cheese was made by transforming about 10 L of raw milk derived exclusively from the morning milking of about 50 sheep of different breeds (mainly meat breeds). The milk was transported in a plastic container to the production laboratory where was processed in 15 min. In accordance with the unusual production methods characterised by the use of a tanning made from olive oil, casavecchia wine, chili and piperna. At the end of processing, the batch of cheese was left to season for six months in earthenware closed jars that were rotated every ten days to allow a uniform distribution of tanning.

A total of 7 samples was collected and transported at 4°C to the Laboratory of Food Microbiology, Department of Veterinary Medicine, University Federico II, Naples, Italy, and directly used for evaluation of bacterial populations. In particular, the sampling plan provided the analysis of bulk milk (L) and the curd (C) on the day of production, the *primosale* (*i.e.* fresh, lightly salted cheese) 24 h after production (Cf 0), 1 cheese wheel after 25 days of drying (Cf 1) and 3 cheese wheels after 60 (Cf 2), 120 (Cf 3) and 180 (Cf 4) days of ripening respectively.

Ten-fold serial dilutions in Ringer's solution were prepared for the microbiological analysis from which inocula of 0.1 and 1 mL were seeded in order to enumerate total viable count (TVC) at 30°C (NF EN ISO 4833:2003; ISO, 2003): the group of hygiene indicators germs as total coliforms (ISO 4832:2006; ISO, 2006), *E. coli* (ISO 16649-2:2001; ISO, 2001a), Enterobacteriaceae (ISO 21528-2:2004; ISO,

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2004a), coagulase-positive staphylococci (NF EN ISO 6888-1/A1:2004; ISO, 2004b), yeasts on rose-bengal chloramphenicol agar containing chloramphenicol supplement (Oxoid) at 22°C for 5 days, sulphite-reducing clostridia on SPS Agar (Oxoid) at 43°C for 24 h, enterococci on Kanamycin Easculin Azide Agar Base (Oxoid) + Kanamycin Selective Supplement (Oxoid) at 37°C for 24-48 h, the autochthonous lactic acid bacteria such as mesophilic and thermophilic lactococci (Mlc; Tlc) on M17 Agar (Oxoid) at 30 and 42°C respectively for 72 h under anaerobic conditions, and mesophilic and thermophilic lactobacilli (Mlb;Tlb) on Rogosa agar (Oxoid) at 30°C for 5 days and 42°C for 72 h respectively under anaerobic conditions (5% CO<sub>2</sub>). Finally, each sample was screened for the presence of Salmonella spp. (ISO 6785:2001 IDF 93:2001; ISO, 2001b) and Listeria monocytogenes (ISO 11290-1:1996; ISO, 1996). All analyses were performed in duplicate, and the mean value of each count was expressed in Log cfu/mL-g.

# Results

The counts of TVC at 30°C, Enterobacteriaceae, total coliforms, *E. coli*, clostridia sulphite reducing, yeasts, enterococci and lactic flora in milk and cheese are shown in Figures 1 and 2.

In raw milk, TVC at  $30^{\circ}\text{C}$  was  $3.2~\text{log}\,\text{cfu/mL}$  and in cheese it ranged from 6.5~to~8.5~log





cfu/g. Enterobacteriaceae and total coliforms, from loads lower than 2 log cfu/mL in raw milk, reached 6 log cfu/g in *primosale*, then decreased until becoming undetectable from the 60<sup>th</sup> ripening day. *E. coli* was not detected in raw milk and in cheese showed the same performance of total coliforms but with values lower than 1 log cfu/g. Yeasts were detected only in C, *primosale* e Cf1 samples with loads of 4.8, 6.1 and 5.6 log cfu/g, respectively (Figure 1). Clostridia sulphite reducing and coagulase positive were never detected.

Enterococci were absent in raw milk while in cheese they reached a maximum of 7 log cfu/g 25 days after production, then ranged from 5 to 6.2 log cfu/g up to 120 days and were absent from the  $180^{\rm th}$  ripening day on. Mesophilic lactobacilli (Mlb), were not revealed in raw milk, while in cheese they reached 7.5 log cfu/g after 60 ripening days and then always demonstrated values above 6 log cfu/g up to the end. Termophilic lactobacilli (Tlb) were present only in C and the cheese sample examined on the  $60^{\rm th}$  ripening day with values of 1.2 e 7 log cfu/g, respectively (Figure 2).

Mesophilic and termophilic lactococci (Mlc; Tlc) had the same values of 3.2 log cfu/g in raw milk which, on the contrary, showed a different trend during production and ripening; Mlc reached 8.4 log cfu/g after 25 days from production and then always showed values above 6.2 log cfu/g until the end. Tlc showed a maximum of 6.3 log cfu/g in C, then decreased and was absent from 120th ripening day (Figure 2). Listeria monocytogenes and Salmonella spp. were never detected.

### **Discussion**

This work studied the microbial ecology of *Conciato Romano* cheese for the first time. The raw milk flora was made from total viable count levels conforming to limits established by European Regulation regarding milk of different species from that vaccine intended for the production of raw milk products (All . III, sect. IX of EC Regulation 2004/853; European Commission, 2004), a small number of Enterobacteriaceae and total coliforms and a predominant indigenous starter lactic flora (Mlc and Tlc).

As in other pecorino cheeses (Gerasi *et al.*, 2003; Pisano *et al.*, 2006) total coliforms, *E. coli*, Enterobacteriaceae were absent from the 60<sup>th</sup> day of ripening. The same can be said for yeasts; these are part of the secondary or residual flora of ripened cheeses (Corbo *et al.*, 2001) and only in some cheeses their presence on the crust play an important role in the maturation of the product.

A key role in the maturation of *Conciato Romano* could be performed by the metabolic

activity of lactic acid bacteria: the flora of this cheese was dominated by mesophilic lactococci and lactobacilli. Mlc predominated in the first 25 days after production carrying out the acidification of the product. From the initial phase of ripening, Mlc decreased by about 1 log and non starter lactic flora (Mlb) predominated with values never below 6 log cfu/g until the end as reported by Coda *et al.* (2006) who studied various Italian sheep's cheeses. The non-starter lactic acid bacteria, although normally present in low numbers in milk, become the distinctive element of lactic flora of the ripened cheeses due to their ability to grow in

particularly hostile environments characterised by high salt concentrations, low temperatures, low concentrations of fermentable carbohydrates, acid pH, low a<sub>w</sub> and by the presence of bacteriocins. These microorganisms, being also widespread in the processing environments, are often isolated from artisanal dairy productions (Macedo *et al.*, 1995; Pedonese *et al.*, 2002). As described in numerous studies on cheeses made from raw sheep's and goat's milk (Macedo *et al.*, 2004; Medina *et al.*, 1992; Psoni *et al.*, 2003), enterococci were present with values of between 4.2 and 7 log cfu/g up to 120 days of ripening (Figure 2).

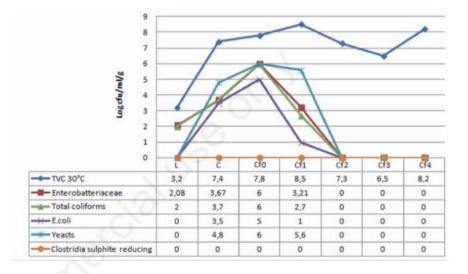


Figure 1. Performance of total viable count (TVC) at 30°C, Enterobacteriaceae, total coliforms, *E. coli*, yeasts and Clostridia sulphite reeducing in bulk milk (L), curd (C), *primosale* (*i.e.* fresh, lightly salted cheese) (Cf 0), cheese wheels after 25 drying days (Cf 1) and 60 (Cf 2), 120 (Cf 3), 180 (Cf 4) ripening days.

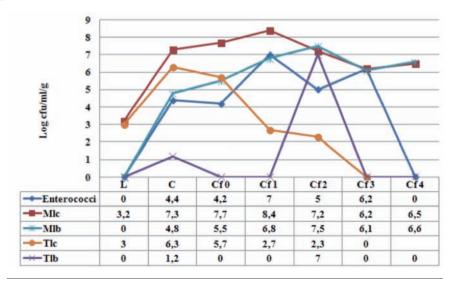


Figure 2. Performance of mesophilic and termophilic lactococci (Mlc; Tlc), mesohpilic and termophilic lactobacilli (Mlb; Tlb), and enterococci in bulk milk (L), curd (C), primosale (i.e. fresh, lightly salted cheese) (Cf 0), cheese wheels after 25 drying days (Cf 1) and 60 (Cf 2), 120 (Cf 3), 180 (Cf 4) ripening days.



Enterococci certainly were shown to be an important part of *Conciato Romano* flora because they are now recognised by several authors (Cenci Goga and Di Antonio, 1995; Giraffa *et al.*, 1997; Mormile *et al.*, 2012) as microorganisms characterised by an important starter and non-starter activity.

## **Conclusions**

This survey was conducted to define the evolution of the microflora of *Conciato Romano* in order to examine its health quality by means of objective criteria. The present study certainly highlighted how *Conciato Romano* is an artisanal cheese characterised by unique organoleptic characteristics and good sanitary qualities at the same time. It is therefore desirable to promote the development of such artisanal products from Campania which, despite being small scale, are unique.

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