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Impact of a thermisation treatment on oxytetracycline spiked ovine milk: Fate of the molecule and technological implications

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ABSTRACT

A few studies have been published on the distribution of oxytetracycline (OTC) residues present in milk among cheese, whey, and milk protein fractions, throughout cheese-making, most of them, focused on the effect of pasteurization and Ultra High Temperature (UHT) treatments, and carried out on cow milk. This study aimed to investigate the impact of a thermisation treatment of ovine milk spiked with oxytetracycline at MRL (maximum residue limit) and half MRL, on the fate of the molecule, and the effect of OTC residues on starter culture development and cheese composition. The antibiotic recovery and partition from milk into whey and cheese were assessed by liquid chromatography-high resolution mass spectrometry (LC-HRMS). Starter and non-starter microflora development was monitored by viable plate counts. Milk thermisation did not affect OTC recovery, partition and cheese chemical composition. On a dry matter basis, an OTC reduction between 15 and 19% was calculated in 60-day cheese, at MRL and half MRL, respectively. OTC caused a dose-dependent difference in the time required to reach pH 5.60, which was significantly higher ($P < 0.05$) at MRL level compared to half MRL and control, allowing coliform bacteria to reach $6 \log \text{CFU g}^{-1}$ in 1-day MRL OTC cheeses.

1. Introduction

Antimicrobial agents are an important tool to treat and prevent diseases and as metaphylaxis in food producing animals. However, their presence as residues in foods of animal origin may lead to the possible induction of allergic reactions in consumers (Adetunji, 2011), and contribute to the development of antimicrobial resistance phenomena (AMR) (European Food Safety Authority EFSA, 2017).

Antibiotics residues occur in milk mainly as a result of incorrect veterinary treatments, such as the adoption of uncorrected withdrawal periods, or the unregulated use of extra label molecules (Beyene, 2016). Sometimes, these wrong practices can lead to overcome the maximum residue limit (MRL) adopted for milk by the international authorities in order to protect the consumers.

Moreover, some authors (Berruga, Beltrán Martínez, Novés, Molina, & Molina, 2011) reported that residual sub-MRL concentrations of antibiotics may be able to affect the manufacturing process of dairy products. Microorganisms continuously exposed to antibiotics at sub-in-

hibitory concentrations can undergo a specific transcriptional change that will modify their metabolic activity (Broszat & Grohmann, 2014). For instance, residues of OTC in yogurt and cheese may cause a significant delay in the pH decrease rate and defects in the development of lactic acid bacteria (Berruga, Battacone, Molina, Román, & Molina, 2008; Cabizza et al., 2017; Suhren, 1998), and interfere with the coagulation and ripening phases affecting the sensory properties of cheeses (Berruga, Molina, Althaus, & Molina, 2016; Nagel et al., 2009).

A few studies have been published on the distribution of antibiotics residues among cheese, whey, and milk protein fractions, during cheese-making. Shappell et al. (2017), working in tubes, at a laboratory scale, found that only 14% of oxytetracycline (OTC) added to the milk was recovered in the curd, and, based on lipophilicity characteristics of various type of animal drugs molecules, provided an empirical model for predicting their distribution between cow skim milk curd, whey and associations with proteins. However, the actual process conditions and the cheese-making technology applied should be taken into account, since they could affect the molecules distribution. Indeed,

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Cabizza et al. (2017), who studied the transfer of oxytetracycline added in ovine raw milk at MRL level ($100\mu\text{gkg}^{-1}$), showed that the cheese was able to retain most of the molecule added to the milk and the presence of the molecule produced a delay during cheese-making in the early acidification phase.

The thermal treatment of milk is a common practice before cheese-making. Indeed, most of the ovine milk undergoes thermisation or pasteurization in order to contrast the development of pathogen and spoilage bacteria, preventing possible technological defects in cheese. Therefore, the ability of these treatments to reduce the antibiotics concentrations should be also assessed taking into account the specific technological process. Some authors conducted studies on the effect of heat treatment on the stability of β -lactams, quinolones, aminoglycosides in milk (Zorraquino, Althaus, Roca, & Molina, 2011, 2009; Roca, Castillo, Marti, Althaus, & Molina, 2010; Roca, Villegas, Kortabitarte, Althaus, & Molina, 2011; Zorraquino, Roca, Castillo, Althaus, & Molina, 2008a; Zorraquino, Roca, Fernandez, Molina, & Althaus, 2008b).

The available studies on the effect of milk thermal treatments on OTC degradation were mainly carried out on cow milk and focused on pasteurization, while no data are available on ovine milk and the effect of thermisation. Shahani (1958) reported a 23.6% degradation of OTC, in milk spiked at $320\text{--}900\mu\text{g L}^{-1}$, after a long time low temperature (LTLT) treatment (61.7°C , 30 min). Kellnerová, Navrátilová, and Borkovcová (2015) reported a 15.3% degradation of OTC in 1.5 MRL spiked milk, after high pasteurization treatment (85°C , 3 s), while an ultra-high-temperature sterilization (UHT) was shown to be effective to completely degrade the OTC spiked in McIlvaine buffer (Hassani, Lázaro, Pérez, Condón, & Pagán, 2008). Zorraquino, Althaus, Nagel, Roca, and Molina (2010) investigated the effect of different heat treatments on the antimicrobial activity of milk samples spiked with OTC by antimicrobial assay. The thermal treatments produced a high loss of antimicrobial activity with classic sterilization (120°C , 20 min), while LTLT treatment and UHT produced a little loss of antimicrobial activity. However, most of these studies were focused on hardly comparable OTC concentrations ranges and time-temperature combinations, performed in laboratory conditions, and often on cow milk or other matrices. Moreover, the impact on the microbiota is even less studied and, to date, a study assessing the effect of low-dosage antibiotics on starter lactic acid bacteria development during cheese-making lacks.

The aim of this work was to assess the effect of a thermisation treatment of ovine milk spiked with oxytetracycline, both at half MRL and MRL, on the recovery and partition of OTC residues in resulting cheese and whey, and investigate their possible effect on starter culture development, and composition of 1- and 60-day-old cheeses.

2. Material and methods

2.1. Chemicals and reagents

Oxytetracycline hydrochloride (OTC, purity $\geq 96.7\%$) was purchased from Sigma-Aldrich (St. Louis, MO, USA); 4-epi-oxytetracycline (EOTC, purity $\geq 97\%$), α -apo-oxytetracycline (α -apo-OTC) and β -apo-oxytetracycline (β -apo-OTC) were purchased from Acros Organics (Geel, Belgium). All solvents used were liquid chromatography-mass spectrometry (LC-MS) grade from Carlo Erba (Milan, Italy), and LC-MS grade water was produced with an Advantage System (Millipore, Billerica, MA, USA). Stock and spiking solutions were prepared as previously described (Cabizza et al., 2017).

2.2. Cheese-making process

Ovine milk was collected from an experimental flock of Sarda breed sheep at AGRIS Sardegna Research Agency (Olmedo, Sassari, Italy). Sheep were in good health and did not undergo antibiotics treatments. The milk was divided into 3 jacketed vats of 12 kg each. The first one was the control (not spiked), the second and the third one were spiked with OTC in order to obtain milk at half MRL ($50\mu\text{gkg}^{-1}$) and at MRL level ($100\mu\text{gkg}^{-1}$), respectively. Subsequently, raw milk was heated up to 63°C , and immediately cooled until 38°C . Thermal treatment was performed in batch using circulating water jacket at 68°C and 12°C during the increasing and the decreasing step, respectively. The temperature of milk during the treatment was simultaneously measured every minute in each vat by a thermocouple. The average heat penetration curve is reported in Fig. 1.

Then, the starter culture was inoculated ($6\log\text{CFU g}^{-1}$, CHOOZIT® Su Casu LYO, Danisco, Denmark) and calf rennet added (500IMCU kg^{-1} , Bellucci, Italy), following the cheese-making process described in Fig. 2, in order to obtain uncooked, hard, pecorino cheeses ripened until 60 days. The experiment was replicated three times in a short period (3 consecutive weeks, in January 2016) to minimize the possible effect of milk composition.

Data from the heat penetration curves obtained in the present work were used to calculate the cook value or C value (Auwah, Ramaswamy, & Economides, 2007), as reported in Equation (1):

$$C = \int_0^t 10^{(T-T_r)/z_r} dt \quad (1)$$

where t is the duration of treatment (min), T is the temperature, T_r is the reference temperature and z_r is the reference temperature resistance coefficient. The trapezoidal rule was used for integration. The same procedure was used to estimate C values of thermal treatment performed by Shahani (1958) and Kellnerová et al. (2015).

2.3. Milk and cheese composition, mass balance and component recoveries (fat and protein)

The parameters of milk composition were evaluated by using MilkoScan FT+ (Foss, Hillerød, Denmark) as described in ISO 9622 (ISO, 2013). pH was measured by pH meter (Knick 911, Knick, Berlin, Germany), equipped with a InLab® Solids electrode (Mettler Toledo, Ohio, USA). The pH of cheeses was measured every 60 min until 300 min and then every 30 min until pH 5.5. Total solids were measured according to the ISO method 6731 (ISO, 2010). Cheeses, at 1 and 60 days of ripening, were analysed for many parameters: pH (pH meter Crison Basic 20+, equipped with a 5232 Puncture electrode); dry matter in accordance with ISO 5534 (ISO, 2004); fat (Soxhlet, 1879), lipolysis index (Nuñez, García-Aser, Rodríguez-Martin, Medina, & Gaya, 1986); protein in accordance with ISO 8968-1 (ISO, 2014); pH 4.6-soluble N, 12% trichloroacetic acid soluble N, 5% phosphotungstic acid-soluble N as described by Gripon, Desmazeaud, Le Bars, and Bergere (1975); total ash by gravimetric analysis after ashing the sample at 550°C for 12 h according to ISO 27:1964 (ISO, 1964); NaCl according to ISO 5943:2006 (ISO, 2006). Each analysis was performed in duplicate. Mass balance and components recoveries (fat and protein) in 1-day cheese were expressed as indicated in Cabizza et al. (2017).

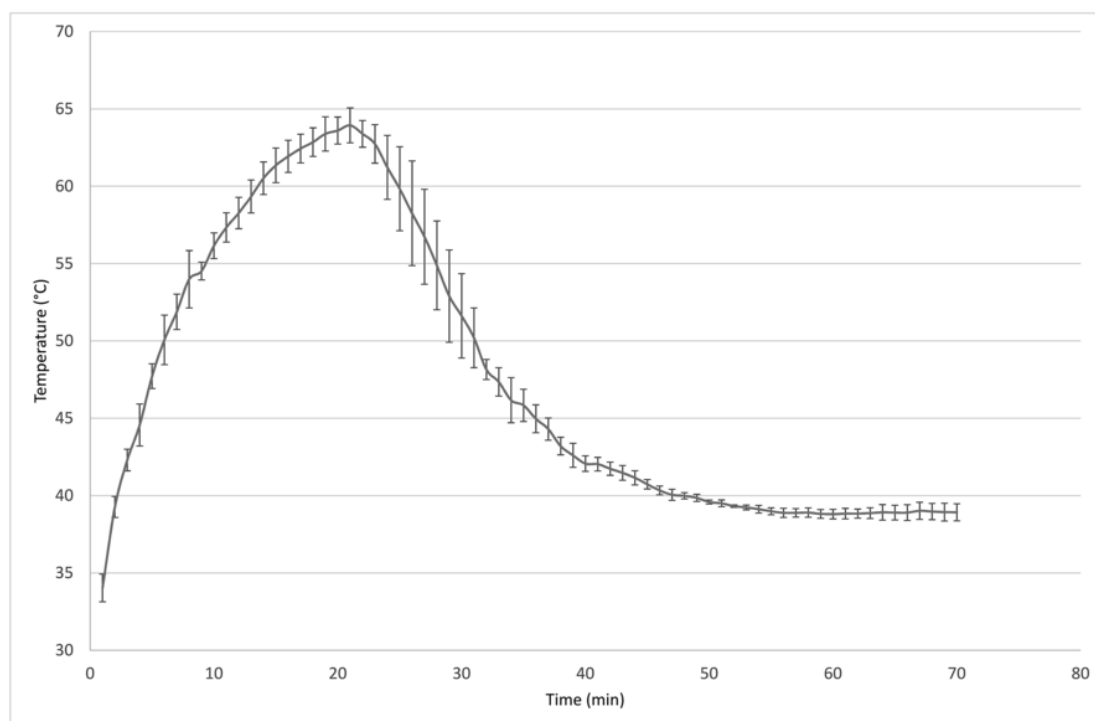


Fig. 1. Heat penetration curve of the adopted thermal treatment. The points are mean \pm standard deviation of the average temperature of the three vats during each cheese-making session ($n = 3$).

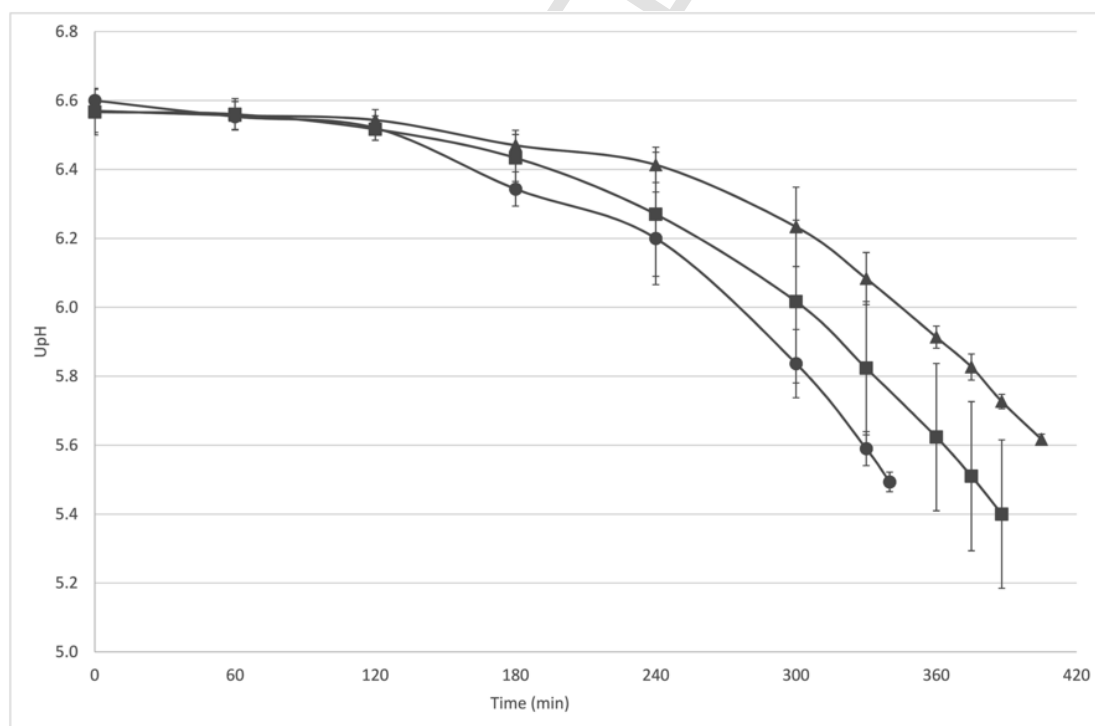


Fig. 2. Cheese-making process.

2.4. Measurement of OTC levels in milk, whey and cheese

2.4.1. Sample extraction and solid phase extraction parameters

Milk and whey samples (2.0 ± 0.1 g) were weighed into a 25 mL centrifuge tube; 10 mL of Mc Ilvaine EDTA buffer (pH 4) were then

added and mixture was agitated using a vortex (Classic Advanced Vortex Mixer, Velp Scientifica, Usmate Velate, MB) and centrifuged at $12800 \times g$ for 10 min at 0°C (Centrifuge 5810R, Eppendorf, Hamburg, Germany). The purification and extraction phases were achieved as described in Cabizza et al. (2017), with except to final volume of the eluates after reconstitution in 14% MeOH (2 mL). Similarly, the whole

preparation protocol of cheese samples was as described in Cabizza et al. (2017).

2.4.2. Liquid chromatography-tandem mass spectrometry analysis

Liquid chromatography-tandem high-resolution mass spectrometry (LC-HRMS) analysis was performed using an UPLC Ultimate 3000 (Thermo Fisher-Dionex San Jose, CA, USA) system was coupled by a HESI-II electrospray source to a Q-Exactive Orbitrap™-based mass spectrometer (all Thermo Scientific, San Jose, CA, USA). Chromatographic separation was performed on Waters Acquity BEH C18 (100 mm × 2.1 mm, 1.7 μm particles) column preceded by a guard column BEH Shield RP18 (5 mm × 2.1 mm, 1.7 μm particles), gradient elution with H₂O with 0.1% (v/v) formic acid (A) and MeCN (B). The injection volume was 2 μL, the flow rate was 0.400 mL min⁻¹, and the adopted gradient varied the % of A to B as follows: T 0.0 min, 80%; T 0.50 min, 80%; T 5 min, 0%; T 6.50 min, 0%; T 6.60 min, 80%; T 8.00 min, 80%; the target column temperature was 45 °C.

Q-Orbitrap HRMS with HESI-II electrospray source was operated in positive mode.

The following ionization parameters were applied: electrospray voltage 4 kV for positive mode, capillary temperature 300 °C, Aux gas heater temp 330 °C, sheath gas (N₂) 40 arbitrary units (arb), auxiliary gas (N₂) 10 (arb), and S-Lens RF level at 50 (arb). LTQ Velos ESI positive-ion calibration solutions (Thermo Scientific, San Jose, CA). The acquisition was achieved in full scan/dd-MS² mode. Full MS mode: mass range m/z 150–500, resolution of 70000 FWHM (m/z 200), AGC target 3.0E6, maximum injection time of 100 ms; dd-MS² mode: resolution of 17500 FWHM (m/z 200), AGC target was set at 1.0E5 ions, maximum injection time of 75 ms, isolation window of m/z 2.0.

The obtained retention time (RT) of analytes of OTC, EOTC, α-apo-OTC and β-apo-OTC were 2.56, 2.21, 3.04 min, respectively. MS monitored masses in HESI+ mode (m/z) were: OTC, precursor 461.1555, fragments 426.1 and 337.1; EOTC, precursor 461.1555, fragments 426.1 and 444.1; α-apo-OTC and β-apo-OTC, precursor 443.1449, fragment 426.1 (fragment ions are used only for qualitative purposes).

2.4.3. Method validation

The method performance was assessed by an in-house validation protocol applied to the milk, whey and cheese, as described in Cabizza et al. (2017). The values obtained for each criteria adopted are reported below. The trueness obtained, expressed as the percent of recovery, was 102 for milk and whey and 80 for cheese. The limit of quantification was 1 μg kg⁻¹ for milk and whey, and 10 μg kg⁻¹ for cheese. The repeatability, expressed as the relative standard deviation “within-lab” repeatability (RSD_{wr}; %), was 9% for milk and whey and 10% for cheese. The extended uncertainty was 22% and 25% for milk, whey and cheese, respectively.

2.5. Microbiological analysis

Samples of thermised milk, before and after the inoculum of the starter culture, 1 and 60 days ripened cheeses were prepared according to the IDF standard 122C (1996). Viable counts were carried out to enumerate: starter lactic acid bacteria (SLAB), i.e. thermophilic cocci and lactobacilli, on M17 agar and MRS agar media (Microbiol, Cagliari, Italy), respectively, incubated at 45 °C, in anaerobiosis for 72 h; non-starter lactic acid bacteria (NSLAB), i.e. mesophilic lactobacilli, on FH agar medium (Isolini, Grand, & Glaetli, 1990), incubated at 37 °C, in anaerobiosis for 72 h; Enterococci on KAA medium (Microbiol), incubated at 42 °C, in aerobiosis, for 18–24 h; coliform bacteria, on VRBA MUG (Microbiol), incubated at 37 °C, in aerobiosis for 18–24 h.

2.6. Statistical analysis

Statistical analysis was performed by Statgraphics Centurion XVI for Windows software package (version 16.2.04; Statpoint Technologies, Inc. Warrenton, Virginia, VA, USA). Analysis of variance (ANOVA) was carried out using the general linear model (GLM) to determine the effects of treatment and replicate on each response variable. Tukey's multiple comparison test was used for paired comparison of treatment means ($P < 0.05$).

3. Results and discussion

3.1. Milk and cheese composition, mass balance and components recoveries

The physical-chemical characteristics of the milk (mean ± s.d.) were: total solids 15.1 ± 0.1; protein, 4.8 ± 0.1; fat, 4.9 ± 0.1; casein, 3.5 ± 0.1; and lactose, 4.8 ± 0.1 (g 100 g⁻¹ of milk). The pH was 6.7 ± 0.1.

OTC spiked milk (half MRL and MRL) showed a dose-dependent delay during the early acidification phase (Fig. 3), until reaching pH 5.60 (35 ± 11 and 78 ± 26 min, at half MRL and MRL levels). In fact, the time required to reach pH 5.60 was significantly higher at the MRL level (406 ± 2 min), compared to the control (328 ± 24 min, $P = 0.0050$) and markedly higher, even not significantly, compared to half MRL (363 ± 30 min, $P = 0.0666$).

These results are in accordance with those previously noticed for raw milk by Cabizza et al. (2017), as milk thermisation did not affect OTC activity and its influence on starter culture acidification performance.

The delay in acidification profiles did not result in a different mass balance (weights of sweet whey, draining whey and cheese, as proportions of total processed milk weight) between control and experiments (Table 1). Control and OTC (both half MRL and MRL) cheese yields, calculated as ratio between produced cheese at 1-day and milk used for cheese-making, were comparable. Similarly, the same component recoveries in control and experimental cheeses were observed, as already noticed for cheese obtained from raw milk (Cabizza et al., 2017). Furthermore, no differences in pH, composition and soluble N levels, between control and experimental cheeses, both at 1 and 60 days of ripening, were observed (Table 2). The chemical characteristics of 60-day ripened cheeses were close to those of Pecorino Sardo cheese (Pirisi, Comunian, Urgeghe, & Scintu, 2011).

3.2. Measurement of OTC levels in fractions

The thermal treatment of spiked milk samples did not reduce the concentration of OTC both in half MRL (47 ± 6 μg kg⁻¹ and 42 ± 1 μg kg⁻¹, raw vs. thermised) and MRL (90 ± 11 μg kg⁻¹ and 90 ± 13 μg kg⁻¹, raw vs. thermised) spiked milk samples. Using for OTC a z_{ref} value of 8.88 °C and a T_{ref} of 71.1 °C, as provided by Shahani (1958) for cow milk, in the pasteurization temperature range, we obtained a C value of 1.64 ± 0.13. This value should be equivalent to about 84% of that estimated (C value of 1.84) using the conditions applied (85 °C, 3 s) by Kellnerová et al. (2015), which resulted in a 15.3% antibiotic drop. On this basis, a residual concentration of 36.4 ± 0.5 μg kg⁻¹ and 78 ± 11 μg kg⁻¹ should have been expected for samples spiked at half MRL and MRL, respectively. However, the heat degradation of oxytetracycline is reported to be also dependent from the total solid content of the milk, with a lower level of degradation as the solid content, particularly fat, increases (Moats, 1999; Tian, Khalil, & Bayen, 2017). Ovine milk, whose total solid, especially fat, content is

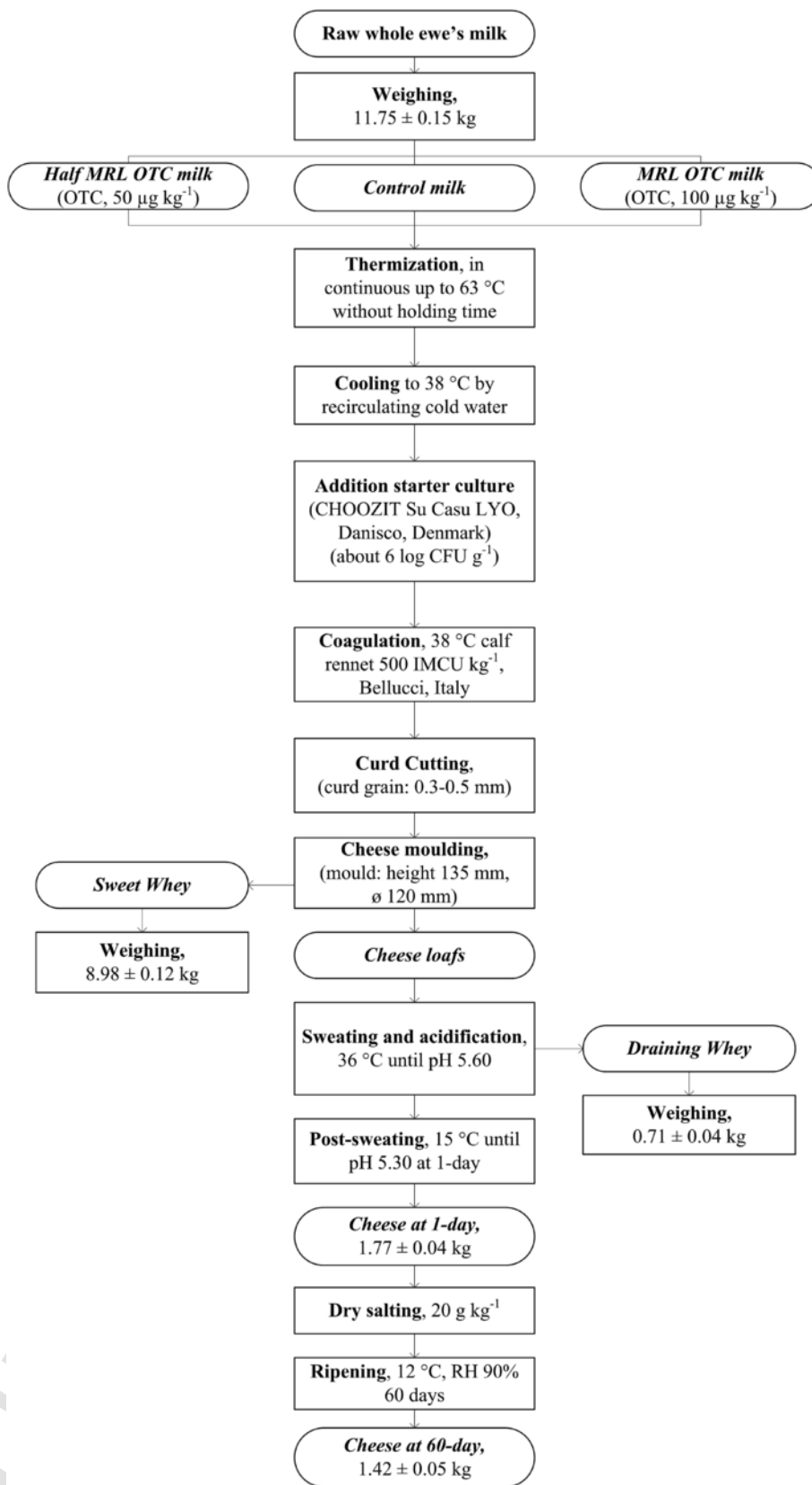


Fig. 3. Acidification curves. The points of each curve are mean values ± standard deviation (n = 3). Control (—●—), Half MRL OTC (—■—) and MRL OTC (—▲—).

Table 1
Technological parameters.

Parameter	Control			Half MRL OTC			MRL OTC		
Time to reach pH 5.60 ^A	328 ^b	±	24	363 ^b	±	30	406 ^a	±	2
Mass balance ^B									
Sweet whey	76 ^a	±	1	76.6 ^a	±	0.2	76.4 ^a	±	0.7
Draining whey	6.1 ^a	±	0.4	6.0 ^a	±	0.3	6.0 ^a	±	0.2
1-day cheese	15.1 ^a	±	0.1	15 ^a	±	0.1	15.1 ^a	±	0.5
Recovery (%) ^C									
Fat	84 ^a	±	4	84 ^a	±	3	84 ^a	±	6
Protein	73 ^a	±	1	73 ^a	±	2	74 ^a	±	2

^A Time expressed in minutes during the acidification phase to reach pH 5.60.

^B Values (g 100 g⁻¹ of milk) are means ± standard deviation (n = 3). Mass balance defined as outputs (sweet whey, draining whey, cheese) as a percentage of milk total weight.

^C Recovery in 1-day cheese, defined as percentage of fat and protein weight in cheese on fat and protein weight in milk, respectively. Level of spike with oxytetracycline (OTC): Half MRL, 50 µg kg⁻¹; MRL, 100 µg kg⁻¹. Values are means ± standard deviation (n = 3); values within a row not sharing common superscript letters were significantly different (P < 0.05).

higher than cow milk (Park, Juárez, Ramos, & Haenlein, 2007), could have protected OTC from degradation during the thermal treatment applied in this study. The adopted time-temperature profiles in cheese-making practices usually are milder than that reported in the above-cited literature, and the treatment is often performed at the plant by continuous systems. Further, as many cheeses derived from milk undergone to thermisation, particularly those made from ovine milk (Pirisi & Pes, 2011), it can expect that the thermal treatment of ovine milk, in cheese-making conditions usually applied at dairy plant, should produce a poor or negligible degradation of OTC.

The distribution between wheys (sweet and draining) and cheese of the OTC added in milk is reported in Table 3. The partition between the obtained fractions appeared to be independent from the spike level, and was similar to that previously observed in the experiments performed with raw milk (Cabizza et al., 2017), except for the cheese fraction, which was able to retain a greater amount of the added molecule (about 80%) than the cheese made from raw milk (about 60%). The overall molecule recovery, close to 100%, obtained in this study, could be ascribable to the better performances provided by the adopted analytical method, involving LC-HRMS, indicating that the unrecovered amount (17%) obtained by Cabizza et al. (2017) could be attributable to the cheese.

The concentration of OTC in cheese, for both the studied levels, did not decrease with ripening, (Table 3), differing from what previously observed for cheese obtained from raw milk (Cabizza et al., 2017). Actually, in absence of degradation phenomena, an increase in concentration of OTC should be expected, as for the other components, due to the reduction of moisture. However, considering the concentration on a dry matter basis, it was possible to calculate a reduction of OTC with ripening, from 413 ± 11 µg kg⁻¹ to 335 ± 3 µg kg⁻¹ of dry matter (19 ± 2%), and from 832 ± 57 µg kg⁻¹ to 707 ± 25 µg kg⁻¹ of dry matter (15 ± 4%), in half MRL and MRL cheeses, respectively. Some consequences of the thermal treatment of milk, such as the modification of the enzymatic pattern and the reduction of the redox potential (McSweeney & Fox, 2009) could have contributed to the stability of OTC during ripening. In our opinion this could be a reason of the lower reduction of OTC (about 50%) compared to that previously observed in raw milk cheese (Cabizza et al., 2017), even if, in that case, cheese underwent to a longer ripening period.

3.3. Microbiological analysis

Results of viable counts, carried out to verify the effect of OTC on microflora development in 1-day and 60-day cheeses, are reported in Table 4.

No significant differences were detected between viable counts of the microbial groups searched in control and experimental thermised

milk samples, both before and after the starter inoculum, whose level was confirmed to be about 6 log CFU mL⁻¹. NSLAB, Enterococci and coliform bacteria counts were < 1 log CFU g⁻¹ in all the thermised milk samples analysed.

In 1-day cheese, SLAB (thermophilic lactobacilli and thermophilic streptococci) reached the same average log CFU g⁻¹ in control and experimental samples, despite a pronounced delay during the acidification phase. Whereas, coliform bacteria counts were significantly higher (P < 0.05) in MRL level, than in half MRL and control samples. This difference is attributable to the acidification delay, which resulted in more favourable environmental growth conditions (higher pH in OTC cheeses than in control ones) for coliform bacteria that were able to reach 6 log CFU g⁻¹. This is a crucial issue to be taken into account, since high levels of coliform bacteria could have negative technological implications during the cheese-making, resulting into cheese defects, as well as being a safety concern for human health, particularly in short-ripened cheeses.

In 60-day cheese, thermophilic lactobacilli decreased about 1 log CFU g⁻¹, in all samples, while thermophilic streptococci remained stable in both OTC spiked samples, and decreased about 1 log CFU g⁻¹ in the control, whose counts resulted significantly different (P < 0.05) from both OTC samples. In fact, as reported by Broszat and Grohmann (2014), a higher ability to survive could be acquired by microorganism as stress response to sub-inhibitory concentrations of antibiotics that could modify their metabolic activity. As expected, NSLAB, whose important role during ripening is well known, though were almost undetectable in 1-day cheese, increased in all samples, until 4 log CFU g⁻¹. Enterococci remained constant, while coliform bacteria decreased during ripening (about 4 log CFU g⁻¹), showing no significant differences among counts of OTC (both MRL and half MRL) and control 60-day cheeses.

4. Conclusions

The thermal treatment performed did not reduce the concentration of OTC, both in half MRL and MRL spiked ovine milk samples, probably because of a protective effect of the higher total solid content of the ovine milk, especially fat, compared to cow milk. No differences on mass balance, yields, chemical composition between experimental and control cheese-making processes, 1 and 60-day cheeses, were observed.

The LC-HRMS based method adopted allowed to obtain an overall recovery of the molecule spiked in milk close to 100%.

The OTC residues in cheese underwent a modest decrease during ripening. Thermisation may have caused a reduction of the redox potential and a selection of the enzymatic pattern, which contributed to the stability of the molecule, resulting in a lower decrease, compared to that observed for raw milk cheese, in a previous work.

Table 2
Physico-chemical parameters of 1-day and 60-day cheese ^A.

Parameter	1-day cheese						60-day cheese											
	Control		Half MRL OTC		MRL OTC		Control		Half MRL OTC		MRL OTC							
pH	5.4 ^a	±	0.1	5.4 ^a	±	0.1	5.4 ^a	±	0.1	5.1 ^a	±	0.1	5.2 ^a	±	0.1	5.2 ^a	±	0.1
Moisture (g 100 g ⁻¹ of cheese)	45.0 ^a	±	0.4	45.2 ^a	±	0.6	45.0 ^a	±	0.9	33.9 ^a	±	0.8	33.9 ^a	±	0.8	34 ^a	±	1
FDM (g 100 g ⁻¹ DM)	50 ^a	±	1	50 ^a	±	1	49 ^a	±	2	49.8 ^a	±	0.5	50.0 ^a	±	0.2	49.9 ^a	±	0.3
PDM (g 100 g ⁻¹ DM)	41.7 ^a	±	0.7	42.1 ^a	±	0.3	42.3 ^a	±	0.3	40.5 ^a	±	0.4	41 ^a	±	1	41 ^a	±	1
SN (g 100 g ⁻¹ TN)	6.9 ^a	±	0.8	8 ^a	±	1	7 ^a	±	2	13.7 ^a	±	0.3	15 ^a	±	1	15 ^a	±	1
TCA-SN (g 100 g ⁻¹ TN)	2.7 ^a	±	0.6	2.6 ^a	±	0.6	2.6 ^a	±	0.6	8 ^a	±	1	8.9 ^a	±	0.6	8.6 ^a	±	0.9
PTA-SN (g 100 g ⁻¹ TN)	1.4 ^a	±	0.2	1.5 ^a	±	0.2	1.5 ^a	±	0.2	4.0 ^a	±	0.2	3.9 ^a	±	0.3	3.8 ^a	±	0.3
FFA (meq 100 g ⁻¹ fat)	1.0 ^a	±	0.3	1.3 ^a	±	0.2	1.0 ^a	±	0.2	1.8 ^a	±	0.1	2.1 ^a	±	0.8	1.3 ^a	±	0.4
Ash (g 100 g ⁻¹ of cheese)	2.7 ^a	±	0.3	3 ^a	±	0.2	3.3 ^a	±	1	5.2 ^a	±	0.4	4.9 ^a	±	0.7	5.2 ^a	±	0.2
Salt (g 100 g ⁻¹ of moisture)	-			-			-			6.2 ^a	±	0.7	6.4 ^a	±	0.7	6.5 ^a	±	1.1

^A Values are means ± standard deviation (n = 3); Abbreviations are: OTC, oxytetracycline; MRL, maximum residue limit; FDM, fat in dry matter; PDM, protein in dry matter; TN, total nitrogen; SN, nitrogen soluble in water; TCA-SN, nitrogen soluble in 12% trichloroacetic acid; PTA-SN, nitrogen soluble in 10% phosphotungstic acid; FFA, free fat acids.

Table 3
Distribution of OTC in milk, wheys and cheese ^A.

	Half MRL OTC			MRL OTC		
<i>OTC partition (%)</i>						
Milk	100			100		
1-day cheese before salting	81.4	±	0.8	77	±	8
Sweet whey	23	±	9	20	±	3
Draining whey	2.9	±	0.5	2.2	±	0.4
<i>OTC Concentration ($\mu\text{g kg}^{-1}$)</i>						
Milk	41.8	±	0.6	90	±	13
1-day cheese before salting	226	±	4	457	±	34
Sweet whey	13	±	5	24	±	6
Draining whey	20	±	2	33	±	3
60-day cheese	222	±	3	470	±	18

^A Values (% , $\mu\text{g kg}^{-1}$) are means \pm standard deviation (n = 3); OTC partition (%) in component derived from milk, defined as percentage of absolute amount (μg) in 1-day cheese, sweet whey and draining whey on total absolute amount of OTC added in milk ($50\mu\text{g kg}^{-1}$, Half MRL OTC; $100\mu\text{g kg}^{-1}$, MRL OTC).

The observed delay in acidification of thermised milk spiked with OTC, compared to an antibiotic free control, proved a dose-dependent influence of low levels OTC residues on the starter culture development and metabolism, which affected lactic acid production. However, analysing 1-day cheese samples no differences in pH and SLAB counts were detected, leading to assume a temporary OTC effect, limited to the early hours of the acidification phase that has to be further investigated.

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Table 4
Thermised milk, thermised milk + starter, 1-day and 60-day cheese viable counts ^A.

Parameter	Thermised milk						Thermised milk + starter					
	Control		Half MRL OTC		MRL OTC		Control		Half MRL OTC		MRL OTC	
Thermophilic cocci	1.1 ^a	± 0.5	1.37 ^a	± 0.05	1.42 ^a	± 0.14	5.6 ^a	± 0.1	5.9 ^a	± 0.1	5.8 ^a	± 0.1
Thermophilic lactobacilli	0.6 ^a	± 0.6	1.0 ^a	± 0.2	1.0 ^a	± 0.1	4.23 ^a	± 0.08	4.49 ^a	± 0.05	4.3 ^a	± 0.1
Mesophilic lactobacilli	0 ^a	± 0	0.1 ^a	± 0.1	0.3 ^a	± 0.4	0.1 ^a	± 0.2	0 ^a	± 0	0 ^a	± 0
Enterococci	0.2 ^a	± 0.3	0 ^a	± 0	0.6 ^a	± 0.6	0 ^a	± 0	1 ^a	± 1	0.4 ^a	± 0.7
Coliform bacteria	0.3 ^a	± 0.5	0.4 ^a	± 0.5	0 ^a	± 0	0.3 ^a	± 0.5	0.3 ^a	± 0.6	0.1 ^a	± 0.2
Parameter	1-day cheese						60-day cheese					
	Control		Half MRL OTC		MRL OTC		Control		Half MRL OTC		MRL OTC	
Thermophilic cocci	8.6 ^a	± 0.4	8.6 ^a	± 0.4	8.8 ^a	± 0.5	7.7 ^b	± 0.3	8.6 ^a	± 0.4	8.6 ^a	± 0.4
Thermophilic lactobacilli	5.3 ^a	± 0.3	5.6 ^a	± 0.2	5.5 ^a	± 0.1	4.6 ^a	± 0.4	4.8 ^a	± 0.4	4.7 ^a	± 0.2
Mesophilic lactobacilli	0.2 ^a	± 0.4	0.1 ^a	± 0.1	0 ^a	± 0	4.7 ^a	± 0.2	4 ^a	± 2	5 ^a	± 2
Enterococci	3.5 ^a	± 0.6	4.4 ^a	± 0.4	4 ^a	± 1	3.6 ^a	± 0.6	4.1 ^a	± 0.5	4 ^a	± 1
Coliform bacteria	5.1 ^b	± 0.2	4.8 ^b	± 0.5	6.1 ^a	± 0.1	1.7 ^a	± 0.5	1 ^a	± 1	2 ^a	± 3

^A Values (log CFU mL⁻¹ for milk and log CFU g⁻¹ for cheese) are means ± standard deviation (n = 3). Values within a row not sharing common superscript letters were significantly different (P < 0.05).

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