Determination of Antibiotics Residues in Milk Using a QuEChERS Method Using Full Factorial Design and Liquid Chromatography-Tandem Mass Spectrometry

Andressa H. A. Grabsk, João Raul B. de Souza, Francilaine E. De Marchi, Rodolpho M. do Prado, Geraldo T. dos Santos, Carla Porto and Eduardo J. Pilau

Laboratório de Biomoléculas e Espectrometria de Massas (LaBioMass), Departamento de Química (DQI), Universidade Estadual de Maringá, 87020-900 Maringá-PR, Brazil

Núcleo de Pesquisa e Estudo em Pecuária Leiteira (NUPEL), Departamento de Zootecnia (DZO), Universidade Estadual de Maringá, 87020-900 Maringá-PR, Brazil

Mestrado em Ciência, Tecnologia e Segurança Alimentar, Instituto Cesumar de Ciência, Tecnologia e Inovação (ICETI), Centro Universitário de Maringá (UniCesumar), Av. Guedner, 1610, 87050-900 Maringá-PR, Brazil

This study evaluated different methods of extraction based on quick, easy, cheap, effective, rugged and safe (QuEChERS) method for analysis of the antibiotics ceftiofur, cloxacillin and enrofloxacin in milk using ultra-performance liquid chromatography-tandem mass spectrometry. The optimized QuEChERS acetate method presented excellent recoveries, from 95 to 99%. A complete factorial design was used to evaluate the effects of variables of the clean-up step: anhydrous octadecylsilane (C18), primary secondary amine (PSA) and sodium acetate (NaAc). Linearity (R²) above 0.96 was achieved for all compounds. Accuracy and precision were assessed by recovery. Accuracy was 91-99%. Intraday precision with relative standard deviations (RSD) lower than 12.3% and interday precision lower than 12.4% were obtained. Limits of detection (LOD) and quantification (LOQ) were obtained between 1.4-6.8 and 1.5-8.7 µg L⁻¹, respectively. The applicability was evaluated using 91 real milk samples.

Keywords: enrofloxacin, cloxacillin, ceftiofur, cow milk, UPLC-MS/MS

Introduction

Antimicrobial resistance is a major global threat with human mortality rates of 10 million per year predicted by 2050.¹ Increasing incidence of antimicrobial resistance is largely attributed to intensive use of antibiotics for humans and livestock production.² Close to two-thirds of the global production of antibiotics is attributed to agricultural use,³ where antibiotics are mainly used in livestock production to control infectious and common diseases, and to enhance animal growth.⁴ The antibiotic prescription and the intensive use can result on persistence of antibiotics residues (ARs) in biologic tissues and fluids of animals, and the inappropriate consumption can promote antimicrobial resistance, may also interfere with fermented milk product processing.⁵ Moreover, the presence of ARs in livestock products may affect human health and interfere with industrial processes. For example, there is evidence that ARs affects the growth of desirable bacteria for cheese production or dairy beverages.⁶ Thus, maximum residue limits (MRLs) for various veterinary drugs in foods, milk included, are established by governmental agencies. Each antibiotic has a MRL considered safe, above which there is potential to cause harm to human health.⁶ However, extraction of ARs for analysis in milk is challenging due to the complex nature of milk, which is a matrix enriched in fats, sugars and proteins.⁷

Anastassiades and Lehotay⁸ proposed an extraction method for the simultaneous analysis of pesticides in various agricultural food matrices, which was termed as QuEChERS (quick, easy, cheap, effective, rugged and safe).

This method is based on the difference of affinities between reagents and analytes in a liquid-liquid extract improved by salting out effect. For cleaning purposes, dispersive solid phase extraction (d-SPE) is the most relevant, in which different sorbents with different affinities are used for specific analytes.⁹ Yet, modifications of the QuEChERS acetate method, described by Lehotay et al.¹⁰ were proposed to consider nature of the analytes and

*e-mail: epilau@gmail.com
matrices. The authors added octadecylsilane (C18) to the usual sorbents in the clean-up step to improve extraction on high fat food matrices, such as milk, eggs, avocado and animal tissues. More recently, used sodium sulfate (Na₂SO₄) as the drying reagent, and the reagents C18, primary secondary amine (PSA), and sodium acetate for the clean-up step for multi-residue analysis of antibiotics in milk.

Among the antibiotics most used by producers of dairy cattle in the northern region of Paraná are enrofloxacin (ENRO), ceftiofur (CEFT) and cloxacillin (CLOX) (data not shown). Enrofloxacin is a fluoroquinolone, nalidixic acid derivative with broad-spectrum activity against Gram-negative bacteria. Ceftiofur is a cephalosporin semisynthetic antibiotic, a class of β-lactams, and it has activity against both Gram-positive and Gram-negative bacteria. Cloxacillin is a β-lactam antibiotic of the penicillin group. Brazilian milk production was 33.7 billion liters in 2016, the Southern Region participated with 12.5 billion liters (37.1%), the largest producing region of the country. The state of Paraná produced 4.730 billion and increased 1.5% compared to 2015. The acquisition of raw milk in the first quarter of 2018 was 6.10 billion liters, 4.1% higher than in the first quarter of 2017. Studies aimed at improving animal welfare, reducing somatic cell counts, improving cow health and minimizing waste management expenses are constantly being developed to that milk production in these regions will grow to face competition from the world market.

The development and optimization of analytical methods is laborious and resource-consuming. Thus, the use of statistical methods and the experimental design for systematic optimization (e.g., factorial design and response surface analysis) can be applied to different systems. The advantage of such approaches is that they provide more information about the variables and their interactions with fewer experiments compared to traditional univariate studies. Full factorial design is a statistical multivariate optimization, which is widely applied to identify significant variables and the best conditions of an experiment.

The objective of this research was to optimize and develop a high throughput method for routine analysis to detect trace levels of three major ARs in whole milk (ENRO, CEFT and CLOX) using the QuEChERS acetate and modified QuEChERS methods. Furthermore, experimental design was used to optimize the conditions used for sample preparation. Finally, the optimized method was used to evaluated the three ARs on 91 bulk samples of dairy cattle milk from the north region of the State of Paraná, Brazil.

### Experimental

#### Chemical and reagents

Ceftiofur (purity ≥ 95.0%), cloxacillin (purity ≥ 98.0%) and enrofloxacin (purity ≥ 98.0%) were purchased from Sigma-Aldrich (St. Louis, USA). Acetonitrile (ACN), glacial acetic acid and methanol from Panreac AppliChem (Darmstadt, Germany) were purchased high-performance liquid chromatography (HPLC) grade. Formic acid (98%) from Panreac AppliChem (Darmstadt, Germany, grade PA), anhydrous sodium sulfate (Na₂SO₄), anhydrous sulfate magnesium (MgSO₄) from Fmaia (São Paulo, Brazil) and anhydrous sodium acetate (NaAc) from Dynamics (São Paulo, Brazil), were of analytical grade. Ultrapure water was purchased with Gehaka water purification system, model OS10LXE (Curitiba, Brazil). Octadecylsilane (C18) and secondary primary amine (PSA) were obtained from Sigma-Aldrich (St. Louis, USA).

#### Preparation of standards

Stock standard solutions were prepared at concentration of 1.0 mg mL⁻¹, using ethanol (97%) with CEFT; ultrapure water with CLOX; and methanol with ENRO. Diluted working solutions were prepared using ultrapure water at concentrations of 1.0 to 50.0 µg mL⁻¹. Stock and diluted solutions were prepared monthly and stored at −20 °C.

#### Milk samples

Pasteurized dairy milk samples obtained from local supermarkets (Maringá city, Brazil) were used for the validation. For the study of validation, dairy milk samples were first analyzed by ultra-performance liquid chromatography-tandem mass spectrometry (UPLC-MS/MS) to verify the absence of the studied antibiotics. After that, known concentrations of standards were added to the milk samples. For the study of occurrence of AR in Paraná, dairy milk samples were sampled from bulk milk tanks from various dairy farms in Paraná State, Brazil. The quantification was performed by external calibration by matrix overlap.

#### Sample extraction-QuEChERS acetate

CEFT, CLOX and ENRO were extracted from spiked milk samples according to Lehotay et al. Briefly, 15.0 mL of spiked milk was mixed with 15.0 mL of the extraction solvent (ACN with 1% acetic acid v:v), 6.0 g anhydrous MgSO₄ and 1.5 g anhydrous NaAc, and shaken by hand.
vigorously for 1 min. The volume was centrifuged for 10 min at 6,000 rpm at 4 °C. An aliquot of 1 mL of the ACN supernatant was transferred to a 2.0 mL microfuge tube for the d-SPE (containing 50.0 mg PSA + 50.0 mg C18 + 150.0 mg anhydrous MgSO₄). The tube was vortexed for 20 s and centrifuged. The aliquot was filtered and transferred to an auto sampler vial for UPLC-MS/MS analysis.

Sample extraction-modified QuEChERS

CEFT, CLOX and ENRO were also extracted from spiked milk samples according to Wang et al. Briefly, 10.00 mL of spiked milk was mixed with 10.0 mL of the extraction solvent (ACN with 1% acetic acid; v:v) and 6.0 g of anhydrous Na₂SO₄, and the volume was vortexed for 2 min. The volume was centrifuged for 10 min at 6,000 rpm at 4 °C. The ACN layer was transferred to a 25 mL volumetric flask. 10.0 mL of ACN with 1% ammonium hydroxide was added to the remaining tube and vortexed for 2 min. The volume was centrifuged for 10 min at 6,000 rpm at 4 °C. The ACN layers were transferred to a 25 mL volumetric flask diluted with ACN to a volume of 25 mL. An aliquot of 5.0 mL top layer extract was transferred to a 15 mL vial and added C18, PSA, and anhydrous NaAc (100.0 mg, 50.0 mg and 1.0 g, respectively) to the extraction solution. The tube was vortexed for 2 min and centrifuged for 10 min at 6,000 rpm at 4 °C. All the supernatant of the aliquot was evaporated to dryness on a vacuum-rotary evaporator and 2.0 mL of water/ACN solution (95:5/v:v, with 0.1% acetic acid) was added and vortexed to dissolve the analyte. The aliquot was filtered and transferred to an auto sampler vial for UPLC-MS/MS analysis.

Liquid chromatography-tandem mass spectrometry determination

The liquid chromatography analysis was performed on an ACQUITY UPLC System with an ACQUITY UPLC® BEH C18 column (50 × 2.1 mm; 1.7 µm particle diameter) from Waters (Massachusetts, USA) at 30 °C. Mobile phase component A was ultrapure water and component B was ACN, both with 0.1% formic acid. The optimized gradient used was 98% of phase A for 0.5 min; then it decreased linearly by 70% of phase A until 2.0 min; 50% A until 3.0 min; 30% A until 4.0 min; 2% A until 5.0 min and maintained for 4 min. Finally, phase A increased linearly until 10 min to achieve 98% of phase A.

The MS/MS measurements were performed in an ACQUITY TQD triple quadrupole mass spectrometer (Waters, MA, USA). The ionization source was an electrospray probe operated in positive mode. Acquisition was performed in multiple reaction monitoring (MRM) mode to obtain sufficient quantification points to confirm each analyte (CEFT, CLOX and ENRO). Ionization and mass spectrometric conditions were optimized for each AR by infusion at a flow rate of 5 µL min⁻¹ using methanol:water (50:50/v:v) with 0.1% formic acid as mobile phase. The specific MS/MS parameters for each AR are shown in Table 1. The following spectrometer parameters were similar for the three analytes: source temperature at 150 °C, capillary voltage of 2.0 kV, nitrogen as desolvation gas at a rate of 600 L h⁻¹, nitrogen as nebulizer gas at a flow rate of 50 L h⁻¹, desolvation temperature at 350 °C and argon was used as collision gas.

Validation procedure

The validation procedure was performed based on the Manual de Garantia de Qualidade Analítica, of Ministério da Agricultura, Pecuária e Abastecimento (MAPA). The evaluated parameters for quantitative methods were recovery (REC) and matrix effect (ME), selectivity, linearity, intraday and interday precision, accuracy, limits of detection (LODs) and limits of quantification (LOQs). The analytes included in the method comprise substances with different MRLs, thus, this study was based in the specific MRL of each drug. Concentration levels corresponding to 0.25 × MRL, 0.50 × MRL, 1.00 × MRL, 1.50 × MRL and 2.00 × MRL were used (MRLs: 100 µg L⁻¹ to ENRO and CEFT; 30 µg L⁻¹ to CLOX). The MRL for ENRO is based on the sum of enrofloxacin and its marker residue ciprofloxacin. Spiked pasteurized milk samples with CEFT, CLOX and ENRO were prepared using the stock solutions at various concentrations.

The REC and ME were evaluated using samples prepared in mobile phase (A), samples spiked with the standards after the extraction procedure (i.e., blank; B),

---

Table 1. Optimized values for the antibiotics residues (ARs)

<table>
<thead>
<tr>
<th>Antibiotics residues</th>
<th>Retention time / min</th>
<th>Precursor ion (m/z)</th>
<th>Product ion a (m/z)</th>
<th>Cone voltage / V</th>
<th>Collision energy / eV</th>
<th>Dwell time / s</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ceftiofur (CEFT)</td>
<td>3.38-3.50</td>
<td>524</td>
<td>210; 241</td>
<td>30</td>
<td>20</td>
<td>0.180</td>
</tr>
<tr>
<td>Clavulanicin (CLOX)</td>
<td>4.36-4.46</td>
<td>436</td>
<td>114; 160</td>
<td>20</td>
<td>30</td>
<td>0.180</td>
</tr>
<tr>
<td>Enrofloxacin (ENRO)</td>
<td>2.44-2.53</td>
<td>360</td>
<td>245; 316</td>
<td>20</td>
<td>30</td>
<td>0.180</td>
</tr>
</tbody>
</table>

aQ: transition used for quantification; I: transition employed to complete the identification.
and samples spiked with the standards (C), all at the MRLs levels and in triplicates.\textsuperscript{21} REC and ME were calculated as proposed by Matuszewski \textit{et al.}\textsuperscript{22} using the following equations:

\begin{equation}
\text{ME (\%)} = \left( \frac{B}{A} \right) \times 100
\end{equation}

\begin{equation}
\text{REC (\%)} = \left( \frac{C}{B} \right) \times 100
\end{equation}

The selectivity was evaluated using raw milk samples obtained from various dairy farms in Paraná State (Brazil) with different fat levels, in a total of 91 samples.

Linearity was analyzed using calibration curves prepared by matrix overlap at concentrations of 0.25, 0.5, 1.0, 1.5 and 2.0 $\times$ MRLs. Three calibration curves were prepared for three different days and the linearity was evaluated by linear regression using the coefficient of determination ($R^2$).

Accuracy and precision were assessed by recovery using spiked samples (six replicates on three different days) at concentrations of 0.5, 1.0 and 1.5 $\times$ MRLs. Precision was assessed through repeatability (intraday precision) and reproducibility (interday precision) and relative standard deviations (RSD) were calculated for both.\textsuperscript{19}

The LOD and LOQ were determined by the mean of the signal-to-noise ratio, in which the signal must be three times higher than the noise for LOD and ten times higher for the LOQ. For this experiment, dilutions of blank spiked with standards were considered.

**Experimental design to optimize the modified QuEChERS**

A full factorial design was used to evaluate the effects and optimize three variables of the sample preparation method using the QuEChERS described by Wang \textit{et al.}\textsuperscript{18} The three variables were chosen for the dispersive SPE clean-up step: the amounts of anhydrous C18, PSA, and NaAc. The MRLs were considered for method validation. A two level with three independent variables with six replicates at the central point full factorial design was performed using the software Statistica.\textsuperscript{23} The chemicals for the clean-up step C18, PSA and anhydrous NaAc, and the amounts of the chemicals for the clean-up step in the ranges of 50-150 mg, 25-75 mg, and 0.5-1.5 g, respectively, were used. The peak areas were used as the response to determine the optimal conditions for the sample preparation methods of the ARs. The design consisted of 14 experiments and six replications of the central point (Table S1, Supplementary Information (SI) section).

**Results and Discussion**

**Optimization of MS/MS and chromatographic separation**

The mass spectrometer was optimized to provide the best responses for the quantification of CEFT, CLOX and ENRO (Table 1). Each AR was characterized by the retention time and by two precursors-product ion transitions. The most abundant ion produced was used for quantification, whereas the second most abundant was used to complete the identification. With respect to the chromatographic conditions, the gradient elution was studied to determine the best separation, peak shape and sensitivity in the shortest time. Figure 1 shows the MRM transitions for each residue.

**Experimental design and optimized conditions for each antibiotic**

The extraction of ARs was optimized using a complete factorial design. In the planning, 14 experiments were conducted in triplicates with different concentrations of C18, PSA and NaAc. The analysis of variance (ANOVA) was performed with data collected from the experiments with a confidence interval of 95.0%.

The proposed model for this study was significant with $R^2$ determination coefficient appropriate to the model adequacy. In this study, the $R^2$ coefficients for ENRO, CEFT and CLOX were, respectively: 0.9844 or 98.44%; 0.9865 or 98.65%; 0.9707 or 97.07%. The coefficients indicate that about 1.56% of the variations for ENRO were not determined by the model, the same can be said for 1.35% of the variations of CEFT and 2.93% of the variations of CLOX. The adjusted $R^2$ ($R^2$ adj) coefficients for ENRO, CEFT and CLOX were respectively: 0.9710 or 97.10%; 0.9749 or 97.49%; 0.9456 or 94.56%. The coefficients $R^2$ adj propose that the model was highly significant for the simultaneous extraction of the three antibiotics studied.

The amount of NaAc was the factor that most affected the extraction of the three residues (with $p$ value from 0.000012 to 0.00017), showing a negative effect, that is, showing to be optimal at low levels, probably due to ion suppression. Ion suppression alters droplet formation efficiency or droplet evaporation, which, in turn, affects the amount of charged ions in the gas phase causing a loss of signal.\textsuperscript{24} For ENRO, factor C18 was significant and had negative effects, with $p$ value = 0.00005. For CLOX, the amount of PSA was also a significant factor in the extraction process (with $p$ value = 0.047), in which it presented a better response at low levels.
The model also showed that curvature was a significant term in the extraction of the three antibiotics shown, with $p$ value from 0.000083 to 0.00069; however, the predicted model is highly suitable with acceptable $R^2$ and $R^2$ adj coefficients, hence the terms of curvature can be ignored.

The best conditions of the three reagents (C18, PSA and NaAc) for the drug residues were 50.0 mg C18, 25.0 mg PSA and 0.5 g NaAc (Figure 2). The maximum value predicted by the complete factorial design was set for subsequent analysis, including REC and ME, set out in Table 2.

Because of the different structure, polarity, chemical and physical properties of each antibiotic, the optimized factors produce different responses for each sample, thus explaining the importance of multivariate optimization in an extraction process to know the best experimental conditions for the method in different matrices.

Figure 1. MRM chromatograms of a spiked milk samples with (a) enrofloxacin at 100 µg L$^{-1}$, (b) ceftiofur at 100 µg L$^{-1}$ and (c) cloxacillin at 30 µg L$^{-1}$. 
The QuEChERS (acetate and optimized) methods were compared evaluating the REC values of the antibiotics, as shows Table 2. The QuEChERS acetate proved to be inefficient in extracting the antibiotics studied. The residues of CEFT and CLOX had low recovery and it was not possible to extract ENRO. Acceptable recovery ranges for residue analysis are generally between 70 and 120%.\textsuperscript{25} The optimized QuEChERS was more efficient in the extraction of all the antibiotics studied, with recovery values above 95%. Thus, this method was chosen for future experiments of validation.

Table 2 also shows the ME of the optimized QuEChERS. No increase or suppression of signal has been observed for CEFT, so the ionization of this is not affected by the matrix. For the ENRO residue, ion suppression was observed and there was appreciation in the ionization for CLOX. An ME value above 100% is considered a signal enhancement, while ME below 100% is considered a signal suppression. The most frequent approach to avoid or minimize the matrix effect is the use of calibration curve prepared using blank matrices (matrix superposition).\textsuperscript{36}

Comparison between QuEChERS methods and validation

The QuEChERS (acetate and optimized) methods were compared evaluating the REC values of the antibiotics, as shows Table 2. The QuEChERS acetate proved to be inefficient in extracting the antibiotics studied. The residues of CEFT and CLOX had low recovery and it was not possible to extract ENRO. Acceptable recovery ranges for residue analysis are generally between 70 and 120%.\textsuperscript{25} The optimized QuEChERS was more efficient in the extraction of all the antibiotics studied, with recovery values above 95%. Thus, this method was chosen for future experiments of validation.

Table 2 also shows the ME of the optimized QuEChERS. No increase or suppression of signal has been observed for CEFT, so the ionization of this is not affected by the matrix. For the ENRO residue, ion suppression was observed and there was appreciation in the ionization for CLOX. An ME value above 100% is considered a signal enhancement, while ME below 100% is considered a signal suppression. The most frequent approach to avoid or minimize the matrix effect is the use of calibration curve prepared using blank matrices (matrix superposition).\textsuperscript{36}

Comparison between QuEChERS methods and validation

The QuEChERS (acetate and optimized) methods were compared evaluating the REC values of the antibiotics, as shows Table 2. The QuEChERS acetate proved to be inefficient in extracting the antibiotics studied. The residues of CEFT and CLOX had low recovery and it was not possible to extract ENRO. Acceptable recovery ranges for
determination coefficients ($R^2$) were above 0.96 (Table S2, SI section). The statistical evaluation of the regression residuals was performed, in which we obtained values of up to 17.0\% for the concentrations below the MRL for ENRO and 2.5\% for the remaining concentrations. The residual values for CEFT had a maximum value of 3.6\%, whereas for CLOX, the values were below 8.0\%. According to the Manual de Garantia de Qualidade Analítica, the residues should not exceed ±20\% at concentrations below the MRL and ±10\% at the MRL and above.

For the interday precision conditions, the RSD must typically be below 20\% for concentrations between 10 and 100 $\mu$g L$^{-1}$, and below 15\% for concentrations above 100 $\mu$g L$^{-1}$ to 1 mg L$^{-1}$, and to intraday precision the relative standard deviations (RSD) should typically be less than two-thirds of the values presented. Results calculated using RSD for intraday and interday precision were lower than 12.4\% (Table 3). The RSD obtained are in accordance with Manual de Garantia de Qualidade Analítica. In the absence of certified reference material, accuracy was determined by fortification of blank samples and was calculated by the REC test of each residue at the levels studied (0.5, 1.0 and 1.5 × MRL). The results obtained were above 91\% for all analytes.

The sensitivity of the method was determined by LOD and LOQ for each drug (Table 4).

### Table 3. Intraday and interday precision and accuracy for each antibiotic residue

<table>
<thead>
<tr>
<th>Antibiotic residue</th>
<th>Intraday precision (REC ± RSD) / %</th>
<th>Interday precision (REC ± RSD) / %</th>
<th>Accuracy (REC) / %</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>0.5 MRL 1.0 MRL 1.5 MRL</td>
<td>0.5 MRL 1.0 MRL 1.5 MRL</td>
<td>0.5 MRL 1.0 MRL 1.5 MRL</td>
</tr>
<tr>
<td>CEFT</td>
<td>102 ± 7.4 98 ± 1.7 97 ± 6.6</td>
<td>107 ± 10.4 100 ± 11.8 95 ± 12.4</td>
<td>98 98 91</td>
</tr>
<tr>
<td>CLOX</td>
<td>105 ± 4.4 98 ± 7.8 103 ± 6.4</td>
<td>101 ± 6.9 103 ± 10.3 98 ± 8.4</td>
<td>99 99 99</td>
</tr>
<tr>
<td>ENRO</td>
<td>100 ± 8.9 92 ± 12.3 92 ± 3.3</td>
<td>108 ± 8.1 90 ± 4.9 98 ± 7.6</td>
<td>92 97 98</td>
</tr>
</tbody>
</table>

REC: recovery; RSD: relative standard deviation; MRL: maximum residue limit; CEFT: ceftiofur; CLOX: cloxacillin; ENRO: enrofloxacin.

For the interday precision conditions, the RSD must typically be below 20\% for concentrations between 10 and 100 $\mu$g L$^{-1}$, and below 15\% for concentrations above 100 $\mu$g L$^{-1}$ to 1 mg L$^{-1}$, and to intraday precision the relative standard deviations (RSD) should typically be less than two-thirds of the values presented. Results calculated using RSD for intraday and interday precision were lower than 12.4\% (Table 3). The RSD obtained are in accordance with Manual de Garantia de Qualidade Analítica. In the absence of certified reference material, accuracy was determined by fortification of blank samples and was calculated by the REC test of each residue at the levels studied (0.5, 1.0 and 1.5 × MRL). The results obtained were above 91\% for all analytes.

The sensitivity of the method was determined by LOD and LOQ for each drug (Table 4).

### Table 4. Maximum residue limit (MRL), limit of detection (LOD) and limit of quantification (LOQ) for each antibiotic residue

<table>
<thead>
<tr>
<th>Antibiotic residue</th>
<th>MRL / $\mu$g L$^{-1}$</th>
<th>LOD / $\mu$g L$^{-1}$</th>
<th>LOQ / $\mu$g L$^{-1}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>CEFT</td>
<td>100</td>
<td>1.4</td>
<td>1.5</td>
</tr>
<tr>
<td>CLOX</td>
<td>30</td>
<td>4.9</td>
<td>5.3</td>
</tr>
<tr>
<td>ENRO</td>
<td>100</td>
<td>6.8</td>
<td>8.7</td>
</tr>
</tbody>
</table>

CEFT: ceftiofur; CLOX: cloxacillin; ENRO: enrofloxacin.

### Samples analysis

Samples of raw milk (n = 91) were processed using the validated method to provide confirmatory and quantitative analysis, only 7 of them had any of the antibiotics studied. ENRO was found in 6 analyzed samples, all below LOQ. Only one sample presented ENRO below the LOQ and CEFT, at a concentration of 5.59 $\mu$g L$^{-1}$. Only 7.69\% of the analyzed milk samples had any of the antibiotics studied under the MRL, all were in accordance with the legislation. Ciprofloxacin, the biomarker of ENRO, was also monitored by MRM and was not found in the milk samples analyzed.

The method used seemed to be quite adequate as it is based on UPLC-MS/MS, and confirms and quantifies only suspect samples, however, it should be noted that the method was used for the analysis of the active principle ceftiofur and may not be suitable for its marker residue, desfuroylceftiofur.

### Conclusions

The optimized QuEChERS method proved to be very efficient in extracting the antibiotics ENRO, CEFT and CLOX in a complex matrix such as dairy milk and was fully validated for confirmatory and quantitative purposes. It presented simplicity and applicability, fundamental characteristics for routine methods in food control, and will certainly contribute to the investigation and control of the quality of dairy milk as well as to assist the producers of dairy cattle in the best management of veterinary medicines.

### Supplementary Information

Supplementary data are available free of charge at http://jbcs.sbq.org.br as PDF file.

### Acknowledgments

The authors would like to thank CAPES for financial support.

### References


Submitted: October 14, 2018
Published online: March 26, 2019