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Comparison of Three Analytical Methods for the Determination of Quinolones in Pig Muscle Samples

Inmaculada Jiménez–Díaz¹, María Pilar Hermo², Oscar Ballesteros^{*1}, Alberto Zafra–Gómez¹, Dolores Barrón², José Barbosa², Alberto Navalón¹

¹Research Group of Analytical Chemistry and Life Sciences, Department of Analytical Chemistry, Campus of Fuentenueva, University of Granada, E–18071 Granada, Spain

²Research Group of Bioanalysis, Department of Analytical Chemistry, University of Barcelona, Avda. Diagonal 647, E–08028 Barcelona, Spain

14 Abstract This work presents a comparison between three analytical methods 15 developed for the simultaneous determination of eight quinolones regulated by the 16 European Union (marbofloxacin, ciprofloxacin, danofloxacin, enrofloxacin, difloxacin, 17 sarafloxacin, oxolinic acid and flumequine) in pig muscle, using liquid chromatography with fluorescence detection (LC-FD), liquid chromatography-mass spectrometry (LC-18 19 MS) and liquid chromatography-tandem mass spectrometry (LC-MS/MS). The 20 procedures involve an extraction of the quinolones from the tissues, a step for clean-up 21 and preconcentration of the analytes by solid-phase extraction (SPE) and a subsequent 22 liquid chromatographic analysis. The limits of detection of the methods ranged from 0.1 to 2.1 ng g⁻¹ using LC-FD, from 0.3 to 1.8 using LC-MS and from 0.2 to 0.3 using LC-23 MS/MS, while inter- and intra-day variability was under 15% in all cases. Most of those 24 25 data are notably lower than the maximum residue limits (MRL) established by the 26 European Union for quinolones in pig tissues. The methods have been applied for the 27 determination of quinolones in six different commercial pig muscle samples purchased 28 in different supermarkets located in the city of Granada (South-East Spain).

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31 Keywords Liquid chromatography · Fluorescence detection · Mass spectrometry
 32 detection · Quinolones · Pig muscle · Methods comparison
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- 41 *Corresponding author: Tel.: +34 958243326; fax: +34 958 243328.
- 42 *E-mail address:* <u>oballest@ugr.es</u> (O. Ballesteros)
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44 Introduction

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46 The research in the field of contamination in foods has extended in the last years beyond 47 classical contaminants-pesticides, biocides, polyaromatic hydrocarbons, dioxins or 48 polychlorinated biphenyls-to other compounds such as pharmaceuticals or personal care 49 products [1]. Since pharmaceuticals are produced and applied with the aim of being 50 biologically active and stimulate a physiological response in human and animals even at 51 low concentrations, there is a growing concern in relation to these substances and their 52 recognition as contaminants, mainly due to the adverse effects that their wide use and 53 disposal have on human health [2]. European consumption of pharmaceuticals is known 54 to be increasing on a yearly basis, and today more than 5000 products are being used as 55 painkillers, contraceptives, tranquilizers, lipid regulators, beta-blockers or antibiotics 56 [1]. Antibiotics and their degradation metabolites rank among the most used drugs in 57 human and veterinary medicine. Resistance to antibiotics and other anti-infective agents 58 constitutes a major threat to public health and ought to be recognized as such more 59 widely than it is currently. Therefore, the European Union (EU) recommends the 60 prudent use of antimicrobial agent in human medicine.

61 One of the most important groups of antibiotics is quinolones. They are a family 62 of highly potent antibiotics with a broad spectrum of activity against both Gramnegative and Gram-positive pathogens. They are widely used in human and veterinary 63 64 medicine in the treatment of infections and represent an expanding class of broad-65 spectrum antibacterials [3]. Quinolones have become an integral part of the livestock 66 production industry and can be used therapeutically to treat disease or to prevent it as 67 well as for promoting growth [4]. Their use in veterinary applications can result in the 68 appearance of residues of the compounds and metabolites in edible animal meats and 69 may give rise to public health concerns, including development of resistant bacterial 70 strains, toxic effects or allergic hypersensitivity [5]. Some international organizations 71 such as the World Health Organization (WHO) have recommended a higher attention 72 and control in the use of antimicrobial growth promoters that belong to an antimicrobial 73 class used in humans. The EU agreed to reduce the use of all antimicrobial growth 74 promoters from 2002. To ensure safety, it has been established maximum residue limits 75 (MRLs) for veterinary drugs in those animal tissues that enter the human food chain [6-76 9]. The MRLs values of quinolones in pig muscle are lower than the ones established in 77 other tissues as kidney or liver. So, the MRL in pig muscle for enrofloxacin plus ciprofloxacin, danofloxacin and oxolinic acid are fixed at 100 ng g^{-1} , for marbofloxacin 78 and flumequine at 150 and 200 ng g^{-1} respectively, while for difloxacin the MRL is 400 79 ng g^{-1} . The MRL of sarafloxacin, major metabolite of difloxacin is not vet established. 80 81 Therefore, more analytical methodology is demanded to quantify and confirm the 82 identity of these compounds in food-producing animal. In the scientific literature, some 83 analytical methodologies have been described for the determination of fluoroquinolone 84 residues in animal derived foods. Given the complexity of these samples, the majority 85 of these methodologies are based in the use of liquid chromatography (LC) with ultraviolet (UV) [10], fluorescence (FD) [11] or mass spectrometric (MS) detection [12] 86 87 after sample clean-up by SPE [10-12]. Owing to its specificity, mass spectrometry is the 88 most powerful confirmatory technique; however, it is expensive and thus not available to all laboratories. In the case of fluorescent drugs, as quinolones, because of its 89 90 selectivity and sensitivity, FD is a very good detection approach.

The main objective of this work is to compare the quality control parameters of three different analytical methodologies developed using LC–FD, LC–MS or LC– MS/MS for the determination of quinolones in pig muscle samples in order to provide the method that has the best analytical characteristics. The three analytical methodologies were satisfactorily applied for the quantification of compounds in samples picked up at different supermarkets of Granada (Spain).

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99 Experimental

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- 101 Chemical and reagents
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Pure quinolone standards were purchased from different pharmaceutical companies.
Ciprofloxacin (CIP) from Ipsen Pharma (Barcelona, Spain); danofloxacin (DAN) from
Pfizer (Karlsruhe, Germany); difloxacin (DIF) and sarafloxacin (SAR) from Abbott
(Madrid, Spain); enrofloxacina (ENR) from Cenavisa (Tarragona, Spain); flumequine
(FLU), norfloxacina (NOR) and oxolinic acid (OXO) form Sigma–Aldrich (Madrid,
Spain) and marbofloxacin (MAR) from Vetoquinol (Lure, France).

Acetonitrile, MeCN (LC–grade), *o*–phosphoric and citric acids were obtained from Panreac (Barcelona, Spain). Methanol, ethanol, hexane, ammonia, formic acid, trifluoroacetic acid and *m*–phosphoric acid were supplied by Merck (Darmstadt, Germany). Isolute ENV+ (200 mg/3 mL) solid–phase extraction (SPE) adsorbent cartridges were purchased from Isolute Sorbent Technologies (Mid Glamorgan, UK).

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115 Instrumentation and software

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LC-FD analysis were performed using an HP Agilent Technologies (Palo Alto, CA, USA) 1100 series liquid chromatography system with fluorescence detector connected on-line. ChemStation for LC 3D software (Agilent) was used for instrument control and for data acquisition and analysis. LC-MS and LC-MS/MS analysis were performed using an API 3000 (Applied Biosystems, Foster City, CA, USA) triple quadrupole mass spectrometer system. In order to obtain data, the Analyst 1.4 software was used.

All pH measurements were made with a Crison (Crison Instruments S.A, Barcelona, Spain) combined glass–Ag/AgCl (KCl 3 M) electrode using a previously calibrated Crison 2000 digital pH–meter. A Branson digital sonifier (Danbury, CT, USA) and a Hettich Universal 32 centrifuge (Tuttlingen, Germany) were also used. SPE was performed on a Supelco (Madrid, Spain) vacuum manifold for 12 columns connected to a Supelco vacuum tank and to a vacuum pump. Statgraphics software was used for statistical and regression analysis.

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131 Preparation of standard and stock solutions

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For LC–MS and LC–MS/MS analyses individual stock solutions of CIP, DAN, DIF, ENR, MAR, NOR, and SAR (100 μ g mL⁻¹), were prepared in 50 mM acetic acid aqueous solution. FLU and OXO (100 μ g mL⁻¹) were prepared in MeCN. Individual working solutions were prepared by diluting the initial standard solutions with MeCN.

137 For LC–FD analysis, individual stock solutions of CIP, DAN, DIF, ENR, MAR, 138 NOR and SAR (100 μ g mL⁻¹) were prepared in ethanol (99.9% ν/ν). Individual stock 139 solutions of FLU and OXO (100 μ g mL⁻¹) were prepared in MeCN. Individual working 140 solutions were prepared by diluting suitably with a MeCN–water mixture (12:88, ν/ν). 141 All solutions were stored at 4 °C in the dark for not longer than 2 months.

- 143 Preparation of fortified samples
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145 Fortified samples were prepared by spiking 5 g (accurately weighed) of minced blank 146 pig muscle adding the adequate volumes of working solutions of studied quinolones and 147 norfloxacin -a forbidden veterinary quinolone- used as surrogate. Before sample 148 treatment and analysis, all samples were allowed to stand in the dark for 20 min at room 149 temperature to permit the total interaction between the antibiotics and tissues. In order 150 to evaluate recoveries, spiked samples in the same range of concentration were prepared 151 and compared with samples spiked after the SPE procedure and that were considered 152 the 100% of recovery.

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154 Basic procedure

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Two methods previously published by the authors were followed for sample treatment [11, 12]. The procedures involve an extraction of the quinolones from the tissues by shaking, a clean–up and preconcentration step by solid–phase extraction (SPE) and a subsequent liquid chromatographic analysis.

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- 162 **Results and discussion**
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- 164 Validation of the methods
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- 166 Analytical performance
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For LC-MS/MS calibration, the studied concentration levels ranged from 0.5 to 100.0 168 ng g^{-1} ; for LC LC–MS from 5.0 to 100.0 and for LC–FD from 5.0 to 50.0. In all cases 169 170 each level of concentration was made in duplicate. Calibration curves were constructed 171 using analyte/surrogate peak area ratio versus concentration of analyte. Norfloxacin (400 ng g⁻¹ for LC-MS/MS and LC-MS calibration and 20 ng g⁻¹ for LC-FD 172 173 calibration) was used as surrogate. The lack-of fit test was used to check the linearity of 174 the calibration graphs according to the Analytical Methods Committee [13]. Table 1 175 shows the calibration parameters obtained (intercepts and slopes).

176 177

Table 1

178 Methods validation parameters

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Validation of the methods was performed according to the US Food and Drugs
Administration (FDA) guideline for bioanalytical assay procedure [13] in terms of
linearity, selectivity, sensitivity and accuracy (precision and trueness).

Linearity. It was tested using the correlation coefficients (\mathbb{R}^2) and the P values of the *lack-of-fit* test. \mathbb{R}^2 values ranged from 99.6 to 99.9% for the LC–FD method, from 99.2 to 99.5% for the LC–MS method and from 99.1 to 99.7% for the LC–MS/MS method. P_{lof} values were higher than 5% in all cases. These facts indicate a good linearity within the stated ranges.

188 *Selectivity.* The specificity of the three methods was determined by comparing the 189 chromatograms of blank with the corresponding spiked pig muscle samples. No 190 interferences from endogenous substances were observed at the retention time of the 191 analytes.

192 *Sensitivity.* The limits of detection (LOD) and quantification (LOQ) were 193 calculated according with the IUPAC criterion [14] and the obtained values are shown 194 in Table 1.

195 Accuracy (precision and trueness). To evaluate the overall precision of the 196 methods, intra- and inter-day precision (as relative standard deviation, RSD) were 197 estimated at three different concentrations for each compound (25.0, 50.0 and 100.0 ng g^{-1} for LC–MS and LC–MS/MS, and 10.0, 20.0 and 40.0 ng g^{-1} for LC–FD). In the LC– 198 MS and the LC-MS/MS assess, five pig muscle samples were spiked, extracted and 199 200 analyzed; in the LC-FD assess three spiked samples were extracted and analyzed in 201 duplicate. The procedure was repeated three times on the same day to evaluate intra-day 202 variability and on three consecutive days to determine inter-day variability. Trueness 203 was evaluated by determining the recovery of known amounts of the tested compounds 204 in pig muscle samples. Samples were analyzed using the three methods and the 205 concentration of each compound was determined by interpolation in the standard 206 calibration curve within the linear dynamic range and compared to the amount of 207 analytes previously added to the samples. The results of precision and trueness, 208 summarized in Table 1, fulfill the requirements defined by the EU legislation [7].

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212 Six different pig muscle samples purchased in different markets in the area of Granada 213 (Spain), were extracted, cleaned up and analyzed according to the three methods, in 214 order to prove the presence or not of quinolones in these tissues destined to human 215 consumption. The results obtained with the three methods were similar and showed that 216 one of the analyzed samples contain residues of MAR and OXO. The found concentration of MAR was 62.0 ng g^{-1} and of OXO 20.0 ng g^{-1} . Both values are lower 217 than the MRL established by the EU for these compounds. RSDs from the mean of the 218 219 values obtained with the three methods are 2.8% for MAR and 3.6% for OXO. Figures 220 1, 2 and 3 show the chromatograms of the positive sample using LC-FD, LC-MS (SIM 221 mode) and LC-MS/MS (MRM mode).

Figure 1, 2 and 3

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- 225 Comparison of methods
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227 All methods have a good linearity within the stated ranges, especially the LC-FD method that has the highest values of R^2 in all cases. In relation to the selectivity, the 228 identification of compounds in LC-FD is based on almost exclusively in its retention 229 230 time; as well the compound must be fluorescent at particular wavelengths (λ_{exc} , λ_{em}). In the case of LC-MS each compound is identified by its retention time and it 231 232 characteristic m/z (molecular ion, generally M+H⁺). On the other hand, in LC–MS/MS 233 as well as the retention time, the compounds are identified by two characteristic ions; 234 the first one is used for quantification and the second for confirmation. In this 235 technique, the ratio between quantification and confirmation ions is also used for the unequivocal identification of compounds. Therefore the LC-MS/MS method is the most 236 237 appropriate from the point of view of selectivity. Related to sensitivity, the lowest LOD 238 and LOQ were obtained when the LC-MS/MS method was used. The LODs were between 0.2 and 0.3 ng g^{-1} for the LC–MS/MS method; between 0.3 and 1.8 ng g^{-1} for 239 the LC–MS method and between 0.1 and 2.1 ng g^{-1} for the LC–FD method. In all cases, 240 these values are below of the MRL established by the EU in the Commission 241 Regulation 37/2010 amending Annexes I to IV to Council Regulation (EEC) No 242 243 2377/90 on pharmacologically active substances and their classification regarding 244 maximum residue limits in foodstuffs of animal origin. However, the values obtained 245 using the LC-MS/MS method are from 1.5 to 6 times lower than those obtained using 246 the LC-MS method and from 2 up to 10 times lower than the ones obtained using the 247 LC-FD method, except for DAN whose LOD and LOQ are lower using the LC-FD 248 method. Therefore, the LC-MS/MS method is again the best in terms of sensitivity. In 249 terms of accuracy, intra-day and inter-day precision of the methods were lower than 15% and this is within the acceptable limits proposed by the guidelines for bioanalytical 250 251 method validation ($\leq 20\%$). In all cases RSD values for the LC-FD (2–4%) method 252 were lower than those obtained for the LC-MS (5-14%) and LC-MS/MS (5-12%)253 methods. Finally, recoveries were higher than 77% in all cases with the three methods. 254 The best results were obtained when LC-MS/MS was used as analytical technique, 255 except for oxolinic acid whose recovery is higher using the LC–FD method.

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258 Conclusions

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260 In this work, three procedures which allow the extraction, identification and 261 quantification of the quinolones regulated by the EU in pig muscle samples have been 262 compared. The methods include an extraction of the quinolones from the tissue, a 263 clean-up step by SPE and separation and determination by LC-MS, LC-MS/MS and 264 LC-FD detection. The LOD and LOQ of the three methods are much lower than the 265 MRLs fixed by European Union. The lowest values were obtained when the LC-266 MS/MS method was used. Comparable values of recoveries were obtained for the three 267 methods and the best results of precision in terms of RSD were obtained for the LC-FD 268 method. Therefore, the LC-FD method and the LC-MS/MS method are the ones with 269 the best quality parameters. However, MS/MS have the important advantage of 270 allowing the possibility of confirming (selectivity) the presence of these compounds by 271 means of fragment abundance ratios at rather low concentration levels.

It could be concluded that because of its low cost, easier handling and good quality parameters the LC-FD method would be a good option for the routine analysis of quinolones in pig muscle samples and if a positive sample were found, the LC-MS/MS method should be used to confirm and ensure the result.

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322 Figure Captions

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Fig. 1 LC–FD. Contaminated sample with MAR and OXO. Concentration: MAR, 64 $ng g^{-1}$; OXO, 24 $ng g^{-1}$; and IS, 20 $ng g^{-1}$.

- **Fig. 2** LC–MS in SIM mode. Contaminated sample with MAR and OXO. Concentration: MAR, 59 ng g^{-1} ; OXO, 18 ng g^{-1} ; and IS, 400 ng g^{-1} .
- Fig. 3 LC–MS/MS in MRM mode. (A) Contaminated sample with MAR and OXO. (B)
 Confirmatory chromatograms of MAR and OXO. Concentration: MAR, 62 ng g⁻¹; OXO, 18 ng g⁻¹; and IS, 400 ng g⁻¹.

Parameter*	MAR	CIP	DAN	ENR	SAR	DIF	0X0	FLU
MS								
Calibration curve	b = 4.88	b = 0.83	b = 3.54	b = 5.39	b = 1.62	b = 2.57	b = 2.18	b = 2.89
	$a = -8.53 \cdot 10^{-3}$	$a = 6.25 \cdot 10^{-4}$	$a = 1.03 \cdot 10^{-3}$	$a = 1.53 \cdot 10^{-2}$	$a = 1.77 \cdot 10^{-2}$	$a = 3.60 \cdot 10^{-3}$	$a = 4.07 \cdot 10^{-3}$	$a = 6.68 \cdot 10^{-3}$
$LOD (ng g^{-1})$	0.6	1.8	0.6	0.6	1.3	0.6	0.3	0.3
$LOQ (ng g^{-1})$	2.0	6.0	2.0	2.0	4.3	2.0	1.0	1.0
Recovery (%)								
25.0 ng g^{-1}	92	LL	90	88	83	LL	83	92
50.0 ng g^{-1}	88	84	90	84	80	85	62	96
100.0 ng g^{-1}	90	82	93	83	83	78	78	94
Precision (%RSD)								
Intra-day $(100.0 \text{ ng g}^{-1})$	8	5	6	9	6	5	L	11
Inter-day $(100.0 \text{ ng g}^{-1})$	10	5	7	8	6	7	14	12
MS/MS								
Calibration curve	b = 3.88	b = 2.69	b = 10.09	b = 4.43	b = 2.54	b = 5.92	b = 10.94	b = 8.52
	$a = 2.74 \cdot 10^{-3}$	$a = 1.28 \cdot 10^{-3}$	$a = 1.34 \cdot 10^{-2}$	$a = 8.36 \cdot 10^{-3}$	$a = 2.02 \cdot 10^{-3}$	$a = 2.98 \cdot 10^{-3}$	$a = 1.63 \cdot 10^{-2}$	$a = 9.64 \cdot 10^{-3}$
$LOD (ng g^{-1})$	0.2	0.3	0.2	0.2	0.2	0.2	0.2	0.2
$LOQ (ng g^{-1})$	0.5	1.0	0.5	0.5	0.5	0.5	0.5	0.5
Recovery (%)								
25.0 ng g^{-1}	95	83	92	81	84	LL	LT LT	88
50.0 ng g^{-1}	93	84	88	90	90	86	78	86
100.0 ng g^{-1}	88	94	90	87	87	83	85	96
Precision (%RSD)								
Intra-day $(100.0 \text{ ng g}^{-1})$	L	8	9	5	6	9	8	10
Inter-day $(100.0 \text{ ng g}^{-1})$	12	6	11	8	11	6	6	6

Table 1. Validation parameters of the LC-MS, LC-MS/MS and LC-FD methods.

	1. 0 EN 10-3	$1 - 22 - 10^{-2}$	1- 0.21	1- 61010-2	1-22	1- 1 67 10-2	$1-0.0010^{-3}$	$1-10010^{-2}$
Calibration curve	$0 = 8.00 \cdot 10^{\circ}$	$b = 3.32 \cdot 10^{-10}$	b = 0.51	$b = 0.18 \cdot 10^{-1}$	$0 = 2.25 \cdot 10^{-3}$	$b = 4.0/ \cdot 10^{-1}$	$b = 8.30 \cdot 10^{-2}$	$b = 1.20 \cdot 10^{-10}$
	$a = 3.20 \cdot 10^{-3}$	$a = -3.10 \cdot 10^{-3}$	a = -0.10	$a = 2.30 \cdot 10^{-3}$	$a = 2.70 \cdot 10^{-3}$	$a = -1.41 \cdot 10^{-2}$	$a = -5.60 \cdot 10^{-3}$	$a = -8.30 \cdot 10^{-3}$
$LOD (ng g^{-1})$	0.9	0.8	0.1	1.2	1.6	0.4	1.6	2.1
$LOQ (ng g^{-1})$	3.1	2.5	0.4	4.1	5.4	1.5	5.3	7.0
Recovery (%)								
10.0 ng g^{-1}	62	89	78	80	86	<i>LL</i>	88	86
20.0 ng g^{-1}	83	81	85	86	<i>6L</i>	80	93	83
40.0 ng g^{-1}	87	82	LL	83	84	83	86	86
Precision (%RSD)								
Intra-day $(40.0 \text{ ng } \text{g}^{-1})$	${\mathfrak o}$	${\mathfrak o}$	4	4	5	0	2	4
Inter-day (40.0 ng g^{-1})	3	\mathcal{O}	3	3	4	3	2	3



Figure 1



Figure 2



Figure 3