

Article

Development and Validation of a Reliable UHPLC-MS/MS Method for Simultaneous Quantification of Macroyclic Lactones in Bovine Plasma

Gemechu Zeleke ^{1,2,†}, Siegrid De Baere ^{1,†}, Sultan Suleman ² and Mathias Devreese ^{1,*}

¹ Laboratory of Pharmacology and Toxicology, Department of Pathobiology, Pharmacology and Zoological Medicine, Faculty of Veterinary Medicine, Ghent University, B-9820 Merelbeke, Belgium; Gemechu.Carii@UGent.be (G.Z.); Siegrid.DeBaere@UGent.be (S.D.B.)

² Institute of Health, School of Pharmacy, Jimma University, 378 Jimma, Ethiopia; sultan.sulemanl@gmail.com

* Correspondence: mathias.devreese@ugent.be; Tel.: +32-(0)9-264-73-47

† These authors contributed equally to this work.

Abstract: A fast, accurate and reliable ultra-high performance liquid chromatography–tandem mass spectrometry (UHPLC-MS/MS) method was developed for simultaneous quantification of ivermectin (IVER), doramectin (DORA), and moxidectin (MOXI) in bovine plasma. A priority for sample preparation was the eradication of possible infectious diseases to avoid travel restrictions. The sample preparation was based on protein precipitation using 1% formic acid in acetonitrile, followed by Ostro® 96-well plate pass-through sample clean-up. The simple and straightforward procedure, along with the short analysis time, makes the current method unique and suitable for a large set of sample analyses per day for PK studies. Chromatographic separation was performed using an Acquity UPLC HSS-T3 column, with 0.01% acetic acid in water and methanol, on an Acquity H-Class ultra-high performance liquid chromatograph (UHPLC) system. The MS/MS instrument was a Xevo TQ-S® mass spectrometer, operating in the positive electrospray ionization mode and two multiple reaction monitoring (MRM) transitions were monitored per component. The MRM transitions of m/z 897.50 > 753.4 for IVER, m/z 921.70 > 777.40 for DORA and m/z 640.40 > 123.10 for MOXI were used for quantification. The method validation was performed using matrix-matched calibration curves in a concentration range of 1 to 500 ng/mL. Calibration curves fitted a quadratic regression model with $1/x^2$ weighting ($r \geq 0.998$ and $GoF \leq 4.85\%$). Limits of quantification (LOQ) values of 1 ng/mL were obtained for all the analytes, while the limits of detection (LOD) were 0.02 ng/mL for IVER, 0.03 ng/mL for DORA, and 0.58 ng/mL for MOXI. The results of within-day (RSD < 6.50%) and between-day (RSD < 8.10%) precision and accuracies fell within acceptance ranges. No carry-over and no peak were detected in the UHPLC-MS/MS chromatogram of blank samples showing good specificity of the method. The applicability of the developed method was proved by an analysis of the field PK samples.

Keywords: bovine; plasma; method development; macrocyclic lactones; UHPLC-MS/MS; bioanalysis

Citation: Zeleke, G.; De Baere, S.; Suleman, S.; Devreese, M. Development and Validation of a Reliable UHPLC-MS/MS Method for Simultaneous Quantification of Macroyclic Lactones in Bovine Plasma. *Molecules* **2022**, *27*, 998. <https://doi.org/10.3390/molecules27030998>

Academic Editor: Wilfried Rozhon

Received: 27 December 2021

Accepted: 28 January 2022

Published: 1 February 2022

Publisher's Note: MDPI stays neutral with regard to jurisdictional claims in published maps and institutional affiliations.



Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (<https://creativecommons.org/licenses/by/4.0/>).

1. Introduction

Malaria, a vector-borne disease caused by the *Plasmodium* parasite, continues to have a devastating impact on people's health and lives around the world [1]. Malaria cases and deaths remain high in Africa, especially in children under the age of five [2]. The current increased insecticide resistance and zoophilic behavior in *Anopheles arabiensis* necessitate looking for new complementary vector control tools [3–6]. Zoo-prophylaxis of bovines with Macroyclic Lactones (MLs) against *An. Arabiensis*, as it preferentially feeds on cattle,

is a novel and complementary approach for malaria control and elimination, especially in East and Central Africa [7–11].

The avermectins (ivermectin (IVER) and doramectin (DORA)) and milbemycins (moxidectin (MOXI)) are 6-membered MLs endectocides commonly used against veterinary helminths, ectoparasites. They are semi-synthetic fermentation products of soil-dwelling bacteria of the genus *Streptomyces* with an excellent safety profile [12]. IVER is the prototype, and the commonly used endectocide, which is a mixture of 22, 23-dihydroavermectin B1a (>90%) and 22, 23-dihydroavermectin B1b (<10%). DORA (25-cyclohexyl-5-O-demethyl-25-de(1-methylpropyl)-avermectin A1a.) and MOXI (semi-synthetic methoxime derivative of nemadectin) are the newer generation MLs that have a potent action. MLs act by binding to glutamate-gated chloride channels in nerve and muscle cells of invertebrates, and cause paralysis in the neuromuscular system [13]. Currently, the use of MLs has been extended to humans, especially IVER, against onchocerciasis (river blindness) and is a candidate drug for malaria vector control when given to cattle targeting *An. Arabiensis* [14]. Pharmacokinetic (PK) data is available for the major breeds, including Holstein-Friesian [15]. However, PK data are lacking in cattle breeds in endemic regions, particularly in Ethiopia. To that aim, a fast, economic, and sensitive bioanalytical method is mandatory to allow accurate quantification of MLs in bovines. Furthermore, restrictions usually apply to ship samples from endemic regions to other countries based on the possible transmission of infectious agents, such as foot-and-mouth disease (FMD).

In an extensive review described by Danaher et al., 2006 [16], high-performance liquid chromatography-fluorescence (LC-FL) and high-performance liquid chromatography interfaced to tandem mass spectrometry (LC-MS/MS) have been utilized for the quantification of MLs in biological matrices. In LC-FL methods, MLs bioanalysis was only possible just after sample derivatization to produce fluorescent derivatives and requires a large matrix volume per analysis. The sample derivatization prior to LC-FL bioanalysis is more laborious, time-consuming, and not suitable in the case of large sets of PK sample analysis. On top of this, the rapid degradation of the MLs just after derivatization, as described by de Montigny et al., 1990 [17], Cerkvenik, 2001 [18], Danaher et al., 2006 [16], and Kolberg et al., 2009 [19], makes LC-FL methods less suitable, and consequently, majorly limits its use for PK studies unless online derivatization is devised. In fact, this may also be dependent on the protocol used during sample derivatization.

However, high-performance liquid chromatography interfaced to tandem mass spectrometry (LC-MS/MS) has exhibited multiple advantages, facilitating fast, more sensitive, and accurate analysis of drugs and metabolites in biological matrices. LC-MS/MS also allows accurate simultaneous analysis of mixtures in biological matrices, such as plasma. A few studies have been reported for quantification of MLs using LC-MS/MS methods in biological matrices, such as in milk [20], lamb tissues [21,22], and lamb serum [23], human [24], dog [25], and calf [26] plasma.

To date, LC-MS/MS analysis of MLs in cattle plasma has only been described by Croubels et al., 2002. Even though this method has been successfully applied for IVER, it demands a large volume matrix and further sample clean-up to apply the method to the other MLs [26]. As a result, a novel sample preparation technique is mandatory for the fast and accurate quantification of MLs in bovine plasma. One of the new approaches is the use of an Oasis Ostro® Protein Precipitation & Phospholipid Removal 96-well plate, which combines protein precipitation using an organic solvent with the removal of phospholipids to reduce the matrix effects via a pass-through principle. Due to its simplicity, reproducibility, and highly efficient phospholipid removal, the Oasis Ostro™ 96-well plate pass-through technique is superior and attractive for obtaining high-quality data in LC-MS/MS analysis [27–29].

The objective of this study, therefore, was to develop, optimize and validate a method for fast, accurate, and reliable simultaneous quantification of the MLs (IVER, DORA, and MOXI) in bovine plasma using UHPLC-MS/MS, deploying a straightforward sample clean-up procedure for its application in PK-PD studies. Special attention was given to a

sample preparation that inactivates infectious agents that would otherwise hamper sample shipment to several countries.

2. Results and Discussion

2.1. Method Development

The simplicity and suitability of the procedure for large sample set analysis were taken into consideration during the UHPLC-MS/MS method development and optimization. The aim was to develop a very short, simple, and inexpensive sample preparation procedure to allow the extraction of ≥ 96 samples in one batch. Moreover, a UHPLC-MS/MS method with a short run time (~12 min) was developed to allow the analysis of the MLs in the large sets of plasma samples ($n \geq 96$) within a day. Finally, the developed UHPLC-MS/MS method was fully validated in-house to verify its specificity, reliability, and sensitivity for quantification of the MLs (IVER, DORA, and MOXI) in bovine plasma.

2.1.1. Optimization of Sample Preparation and Clean-Up

The sample preparation was initiated based on previous studies described by Croubels et al., 2002 (IVER in calf plasma) [26] and Morbidelli et al., 2018 (IVER in dog plasma) [25]. The protein precipitation solvents acetonitrile, methanol, and ethanol were evaluated and were added to the plasma samples in a ratio of 75/25 (*v/v*). Even though protein precipitation using acetonitrile showed good extraction recovery, a fluctuating and low signal intensity was observed for MOXI and DORA, resulting in poor method accuracy and precision (results not shown). This signified the need for a further clean-up of the plasma extract using solid-phase extraction (SPE) to remove fats, phospholipids, and other co-eluting components.

Two SPE sorbents using a simple and fast pass-through protocol were evaluated during preliminary experiments, i.e., Oasis® PriME HLB and OstroTM (both from Waters). Veterinary drugs pass through the Oasis® RriME HLB column, while the sorbent holds back interferences [30]. The OstroTM sorbent removes proteins, particulates, and more than 95% of phospholipids from the sample matrix. The Ostro™ 96-well plate utilizes protein precipitation solvent in combination with a single, rapid, pass-through method [29]. Three protein precipitation solvents were tested for the extraction of standard solutions (100 ng/mL) i.e., 1% formic acid (FA) in acetonitrile, methanol, and ethanol. As can be seen in Table S1-Supplementary Materials, good extraction recovery was obtained while using Ostro™ 96-well plate (IVER, DORA, and MOXI were 84.8, 97.0, and 98.1%, respectively) relative to the Oasis® PriME 96-well μ -Elution plate. Moreover, 1% FA in € was the good deproteinisation solvent along with OstroTM 96-well plate pass-through SPE and was therefore selected in the final procedure. A schematical overview of the final sample preparation procedure for quantification of IVER, DORA, and MOXI in bovine plasma is indicated here (see Figure 1).

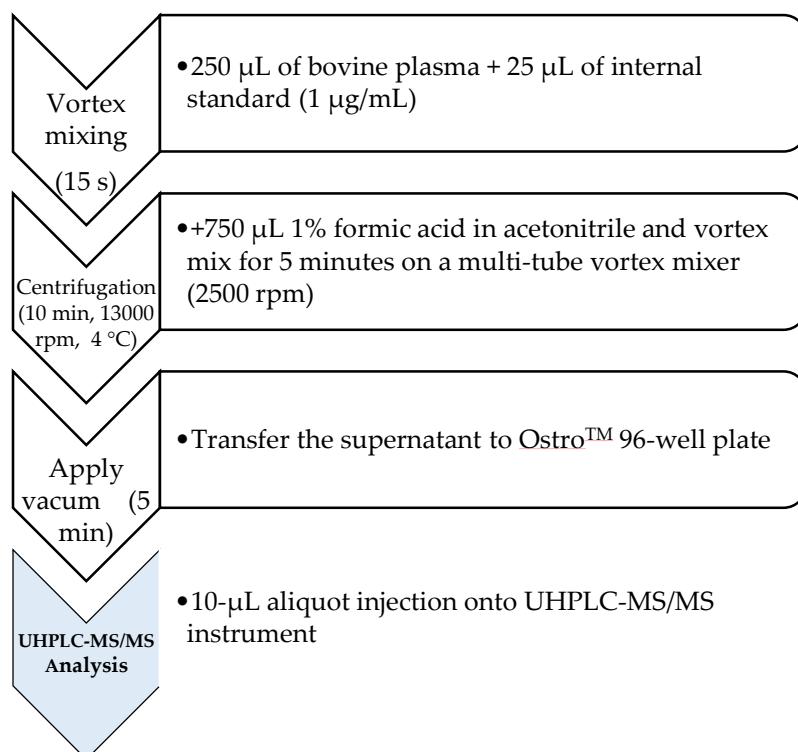


Figure 1. Schematical overview of the final sample preparation procedure for quantification of ivermectin (IVER), doramectin (DORA), moxidectin (MOXI) in bovine plasma.

Compared to other methods in the literature, the current method is fast and simple to apply since drying and reconstitution steps were absent with less solvent consumption; moreover, the procedure is rather inexpensive, since Ostro™ 96-well plates are substantially cheaper than conventional SPE columns [23,25] and samples do not need to be filtrated using 0.22 μ m filters or transferred into autosampler vials, and hence, can be directly injected from the 96-well collector plate onto the LC-MS/MS instrument. As a result, the developed sample preparation procedure is suitable for large sample set extraction within a short period of time. Moreover, Ostro™ 96-well plate was effectively applied for the analysis of other veterinary drugs and mycotoxins in plasma samples of different animal species, such as chicken, turkey, cattle, and pig (Lauwers et al., 2019) [31], De Baere et al., 2018 [32], De Baere et al., 2015 [33].

Finally, the acidification using formic acid was essential to reduce the pH < 6, effectively inactivating viral agents such as FMD. This has a significant impact on the shipping restrictions of the samples.

2.1.2. Internal Standardization

Deuterated internal standards that display similar properties to the analytes, were used to enhance method performance and reduce analytical variations, majorly resulting from analyte loss during sample preparation and from matrix effects due to the co-eluting endogenous components during UHPLC-MS/MS analysis. Stable isotopically labeled internal standards, which could only be commercially obtained at a reasonable price for IVER and MOXI (i.e., ivermectin-d2 and moxidectin-d3), were used. These isotopically labeled internal standards have the same molecular and physicochemical properties as the analytes of interest and elute at the same retention time [34,35].

Ivermectin-d2 (for IVER and DORA) and moxidectin-d3 (for MOXI) were used with success as internal standards, as can be seen from the validation results (see Section 2.2).

2.1.3. Optimization of Chromatographic Conditions

Chromatographic conditions were optimized to develop a simple and reliable method for simultaneous quantification of IVER, DORA, and MOXI in bovine plasma within a total run time of not more than 12 min. Two reverse phase UPLC columns (Acquity BEH C18 column (50×2.1 mm i.d., dp: $1.8 \mu\text{m}$) and an Acquity HSS-T3 column ($(100 \times 2.1$ mm), dp: $1.8 \mu\text{m}$), both from Waters, were evaluated for the chromatographic separation of the analytes in plasma. The Acquity HSS T3 100×2.1 mm UPLC column ($1.8 \mu\text{m}$ particle size) in combination with an Acquity HSS T3 $1.8 \mu\text{m}$ Vanguard pre-column showed an optimal peak separation with good signal intensity and peak shape compared to the Acquity BEH C18 column and was therefore selected for further experiments (results not shown). Similarly, in a previous study, an Acquity HSS T3 100×2.1 mm UPLC column ($1.8 \mu\text{m}$ particle size) in combination with an Acquity HSS T3 $1.8 \mu\text{m}$ Vanguard pre-column was also used for the multi-residue analysis of veterinary drugs, including ivermectin in pond water as described by Goessens et al., 2020 [36].

A mixture of ULC/MS grade aqueous and organic solvents (acetonitrile and methanol) in combination with organic modifiers (formic acid (FA) and acetic acid (AA)) were evaluated in several compositions and gradient programs to evaluate the chromatographic baseline separation, signal intensity, peak shape, and retention time. The addition of 0.1% AA to both the aqueous and organic phase (acetonitrile) resulted in a better signal for DORA, compared to water and 0.1% FA, whereas for IVER the best results were obtained by using no organic modifier (see Figure S1A). MOXI was not tested during the initial experiment, but Chhonker et al. 2018 [24] also previously reported 0.1%AA as an organic modifier for MOXI. By reducing the AA concentration from 0.1% to 0.01%, the signal intensity for IVER and MOXI increased, whereas a decreased signal intensity for DORA was observed (see Figure S1B). As a compromise result, 0.01% acetic acid was selected as the final concentration of the organic modifier since this was also successfully applied by Croubels et al., 2002 [26]. Next, the influence of the organic phase on signal intensity was evaluated. As can be seen in Figure S1C, good signal intensity was obtained for all components while 0.01% AA in methanol was used as an organic phase, relative to 0.01% AA in acetonitrile.

Moreover, the retention time of the analytes increased using methanol as an organic phase, compared to acetonitrile (see Figure S2). The mobile phase gradient was further optimized to achieve a total run time of a maximum 12 min, which is much shorter than previous methods reported by Inoue et al., 2009 [22], analysis of tissues (30 min); Moshou et al., 2019 [37], analysis of fish tissue (20 min); however, comparable with the methods reported by Morbidelli et al., 2018 [25], analysis in milk (11 min); Croubels et al., 2002 [26], analysis of calf plasma (6 min).

2.1.4. Optimization of MS/MS and Multiple Reaction Monitoring (MRM) Parameters

Optimization of the MS/MS parameters was performed by infusion of $1 \mu\text{g/mL}$ standard solutions of each analyte and IS's into the Xevo TQ-S® mass spectrometer at a flow rate of $10 \mu\text{L/min}$, in combination with the mobile phase (20% A/80% B, flow rate: 0.2 mL/min) using the IntelliStart Fluidics system.

Good sensitivity for all analytes and their respective ISs was obtained when the mass spectrometer was operated in the positive electrospray ionization (ESI) mode. While operating in the full scan MS mode, the mass spectra for IVER, IVER-d2, and DORA showed the sodium adduct ions $[\text{M} + \text{Na}]^+$ as the base peak, which were selected as the precursor ions (at m/z 897.50, 899.50, and 921.70, respectively). For MOXI and MOXI-d3, the $[\text{M} + \text{H}]^+$ ion was chosen as the precursor ion (at m/z 640.40 and 643.50, respectively) having similarity with Baptista et al., 2017 [23], Michele et al., 2018 [22], and Hofmann et al., 2021 [38]. The precursor ion of each analyte was fragmented and monitored for product ions at different collision energies and cone voltages. For each precursor ion, the two most abun-

dant product ions were monitored and used for quantification and identification purposes, respectively (see Table 1). The product ions for IVER, DORA, and MOXI were in accordance with other reports in the literature (Croubels et al., 2002 [26], Wang et al., 2011 [39], and Li et al., 2017 [40]). For MOX, the most intense product ion with $m/z = 622$ corresponds with the $[M-H_2O + H]^+$. However, the product ion with $m/z = 123$ was used as a quantifier ion since method performance was better compared to the use of the product ion with $m/z = 622$ as quantifier ion (results not shown).

Table 1. Analyte specific MS/MS parameters obtained in the positive electrospray ionization mode.

Analyte	Precursor Ion (m/z)	Product Ion (m/z)	Dwell Time (Second)	Cone Voltage (V)	Collision Energy (eV)	Retention Time (min)
Ivermectin	897.50	329.20 ^b	0.050	50.00	46.00	7.25
	$[M + Na]^+$	753.40 ^a	0.050	50.00	40.00	
Doramectin	921.70	353.20 ^b	0.025	50.00	48.00	6.70
	$[M + Na]^+$	777.40 ^a	0.025	50.00	37.00	
Moxidectin	640.40	123.10 ^a	0.025	50.00	18.00	6.65
	$[M + H]^+$	622.20 ^b	0.025	50.00	14.00	
Ivermectin-d2	899.50	183.1 ^b	0.050	50.00	48.00	7.23
	$[M + Na]^+$	331.3 ^a	0.050	50.00	48.00	
Moxidectin-d3	643.50	123.00 ^a	0.025	50.00	20.00	6.61
	$[M + H]^+$	625.20 ^b	0.025	50.00	11.00	

Note: a = product ion used for quantification, b = product ion used for confirmation.

2.2. Method Validation

In this study, the method validation parameters including linearity, the limit of detection (LOD), limit of quantification (LOQ), precision, accuracy, carry-over, specificity, and stability were evaluated to determine the UHPLC-MS/MS method performance. The current UHPLC-MS/MS method with a straightforward sample preparation (protein precipitation with 750- μ L 1% formic acid in acetonitrile followed by Ostro® 96-well plate pass-through clean-up), met all requirements.

2.2.1. Linearity

The linearity of the method was evaluated by the correlation coefficient (r), goodness-of-fit coefficients (GoF), and the back-calculated concentrations. During the validation, matrix-matched calibration curves were freshly prepared on 3 different analysis days from blank cattle plasma that was spiked with the analytes of interest at 10 concentrations (1, 2.5, 5, 10, 25, 50, 75, 100, 250, and 500 ng/mL). The selected concentration range of the calibration curve was based on the concentrations that were determined in incurred bovine plasma samples.

Calibration curves were best constructed using weighted ($1/x^2$) quadratic regression analysis ($y = ax^2 + bx + c$, with y = peak area ratio analyte/IS for IVER, DORA, and MOXI). The correlation coefficient (r) values ranged from 0.9985 to 0.9997 and goodness-of-fit coefficients (GoF) were between 1.91 to 4.85% (see Table 2), which complied with the acceptance criteria of $r > 0.99$ and GoF < 10% [41]. The back-calculated concentrations of the analytes in each calibrator sample fulfilled the criterion for accuracy (results not shown). Calibration curves were broader than those reported elsewhere by Croubels et al., 2002 [26], Baptista et al., 2017 [23], and Morbidelli et al., 2018 [25].

Table 2. Results of the evaluation of the calibration model (correlation coefficient (r) and goodness-of-fit coefficient (GoF), mean + standard deviation, ($n = 3$), the limit of quantification (LOQ), and limit of detection (LOD) for IVER, DORA, and MOXI in bovine plasma.

Analyte	Calibration Range (ng/mL)	r (Mean \pm SD)	gof (Mean \pm SD, %)	LOQ (ng/mL)	LOD (ng/mL)
IVER	1–500	0.9997 ± 0.00026	1.91 ± 0.89	1	0.02
DORA	1–500	0.9985 ± 0.00117	4.58 ± 1.97	1	0.03
MOXI	1–500	0.9990 ± 0.00045	3.86 ± 0.91	1	0.58

Note: Acceptance criteria: $r \geq 0.99$ and GoF $\leq 10\%$.

2.2.2. LOQ and LOD

As can be seen in Table 2, LOQ values of 1 ng/mL were reached for IVER, DORA, and MOXI which was the same as the previous report by Croubels et al., 2002 [26] and was more sensitive than the previous studies reported in lamb tissues [21,22] and lamb serum [23]. This LOQ was low enough to allow quantification of the MLs after 4 h and until 35 days after a single subcutaneous administration of 0.2 mg/kg BW of IVER, DORA, and MOXI to cattle. The LOD was calculated based on the signal-to-noise ratio (S/N) of the analyte peak in the LOQ samples. The theoretical concentration that corresponded with an S/N ratio of 3/1 was set as the LOD. In this study, the calculated LOD values were 0.02 ng/mL, 0.03 ng/mL and 0.58 ng/mL for IVER, DORA, and MOXI, respectively [41].

2.2.3. Precision and Accuracy

Within-day precision and accuracies were evaluated at the limit of quantification level (1 ng/mL) and at a low (5 ng/mL), middle (50 ng/mL), and high (250 ng/mL) concentration level (in 6 replicates each). Similarly, between-day precision and accuracy were evaluated by combining the results of the three data sets obtained for within-run precision and accuracy. The results of within-day and between-day precision (RSD, %) ranged between 1.1% to 6.50 and 2.3% to 8.10%, respectively, which comply with VICH GL49 acceptance criteria, whereas accuracies all fell within the acceptance limit at the specified concentration levels [41, 42] (see Table 3). From the one-way ANOVA analysis using STATA software, there was no statistically significant difference between means of accuracies ($p = 0.1480$) and precisions ($p = 0.1594$) for each of the drug groups during the three different days of the validation experiment. In pairwise comparison, no significant differences were also found for MOXI ($p > 0.315$), DORA ($p > 0.712$), and IVER ($p = 1.000$) with respect to accuracy and precision ($p = 1.000$) showing the repeatability, reproducibility, and accurateness of the analytical method.

Table 3. Validation results for within-day and between-day precision and accuracy of ivermectin (IVER), doramectin (DORA), moxidectin (MOXI) in bovine plasma.

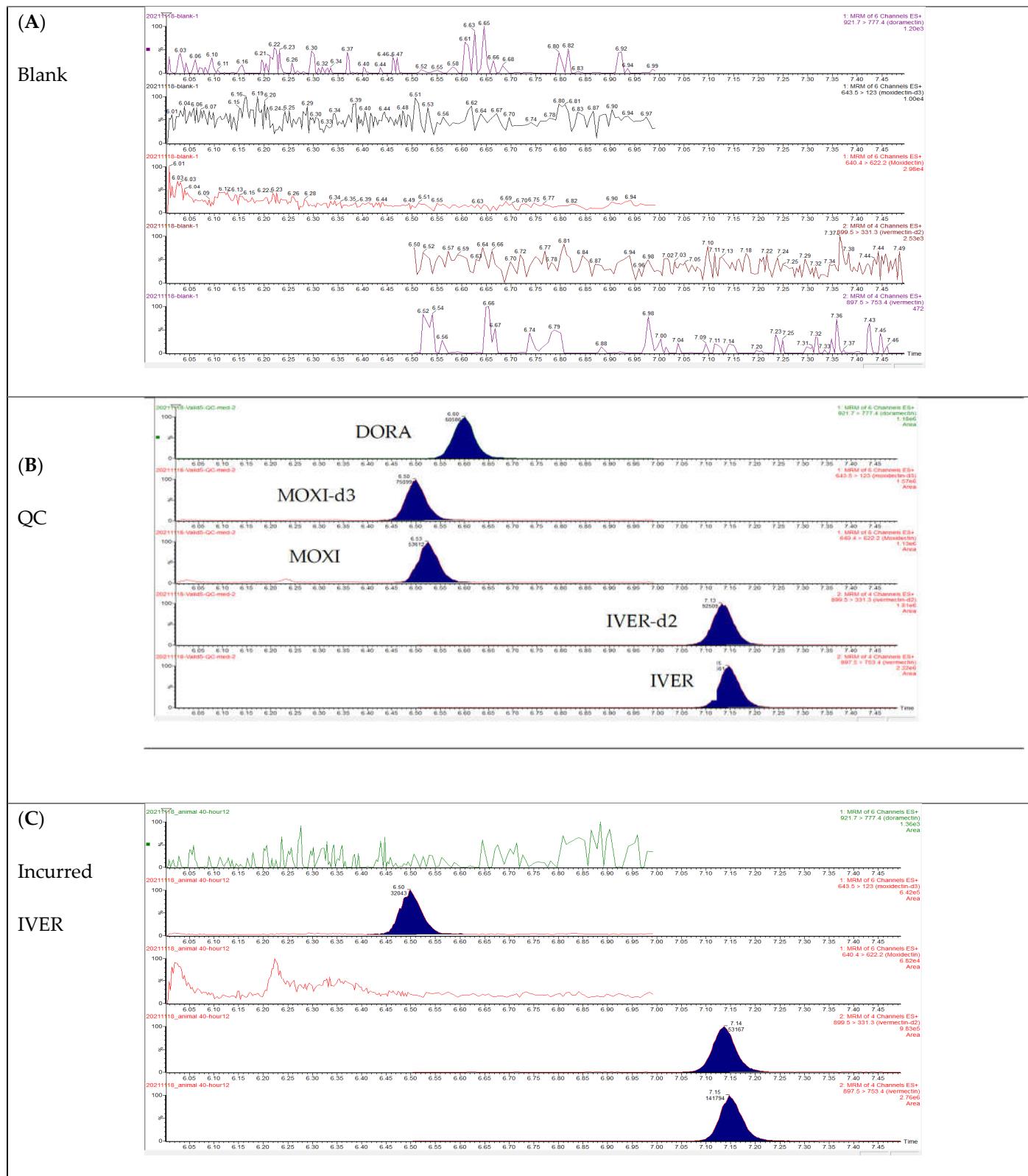
Within-Run Accuracy and Precision					
Analyte	Theoretical Conc.(ng/mL)	Mean Conc.(ng/mL)	SD (ng/mL)	Precision (RSD, %)	Accuracy (%)
IVER	1	0.99	0.04	4.5	-1.2
	5	4.98	0.09	1.8	-0.4
	50	48.89	0.52	1.1	-2.2
	250	250.27	5.92	2.4	0.1
DORA	1	1.01	0.05	4.7	0.8
	5	5.09	0.33	6.5	1.8
	50	47.86	2.41	5.0	-4.3
	250	235.00	11.45	4.9	-6.0
MOXI	1	1.13	0.03	3.1	12.7
	5	5.037	0.30	6.0	0.7
	50	49.257	2.36	4.8	-1.5
	250	247.271	6.31	2.6	-1.1
Between-Run Accuracy and Precision					
Analyte	Theoretical Conc. (ng/mL)	Mean Conc. (ng/mL)	SD (ng/mL)	Precision (RSD, %)	Accuracy (%)
IVER	5	5.09	0.16	3.1	1.7
	50	49.17	1.13	2.3	-1.7
	250	258.42	11.56	4.5	3.4
DORA	5	5.20	0.26	5.1	4.0
	50	49.35	2.36	4.8	-1.3
	250	251.30	18.15	7.2	0.5
MOXI	5	5.15	0.26	5.0	3.0
	50	50.75	2.37	4.7	1.5
	250	248.57	20.19	8.1	-0.6

Note: SD = standard deviation; RSD = relative standard deviation (%); acceptance ranges for within-run precision: RSD < 25%, <15% and <10% for analyte concentrations ranging ≥ 1 to <10 ng/mL, ≥ 10 to <100 ng/mL and ≥ 100 ng/mL, respectively; between-run precision: RSD $\leq 32\%$, 23% and 16% for analyte concentrations ranging ≥ 1 to <10 ng/mL, ≥ 10 to <100 ng/mL and ≥ 100 ng/mL, respectively; acceptance ranges for accuracy: -20% to +10%, -30% to +10%, -40 to +20 for the analyte concentration of ≥ 100 ng/mL, ≥ 10 ng/mL to <100 ng/mL, ≥ 1 to <10 ng/mL (VICH, GL49).

2.2.4. Carry-Over and Specificity

The carry-over on the LC-MS/MS instrument was evaluated by the injection of a solvent sample after the highest calibrator sample. No carry-over was observed at or near the retention time of each of the analytes and respective internal standards (see Figure 2B).

With regard to specificity, no peaks were observed at the elution zone of each analyte of interest and respective internal standards, demonstrating the specificity of the developed method for the analysis of IVER, DORA, and MOXI, as can be seen in Figure 2A.



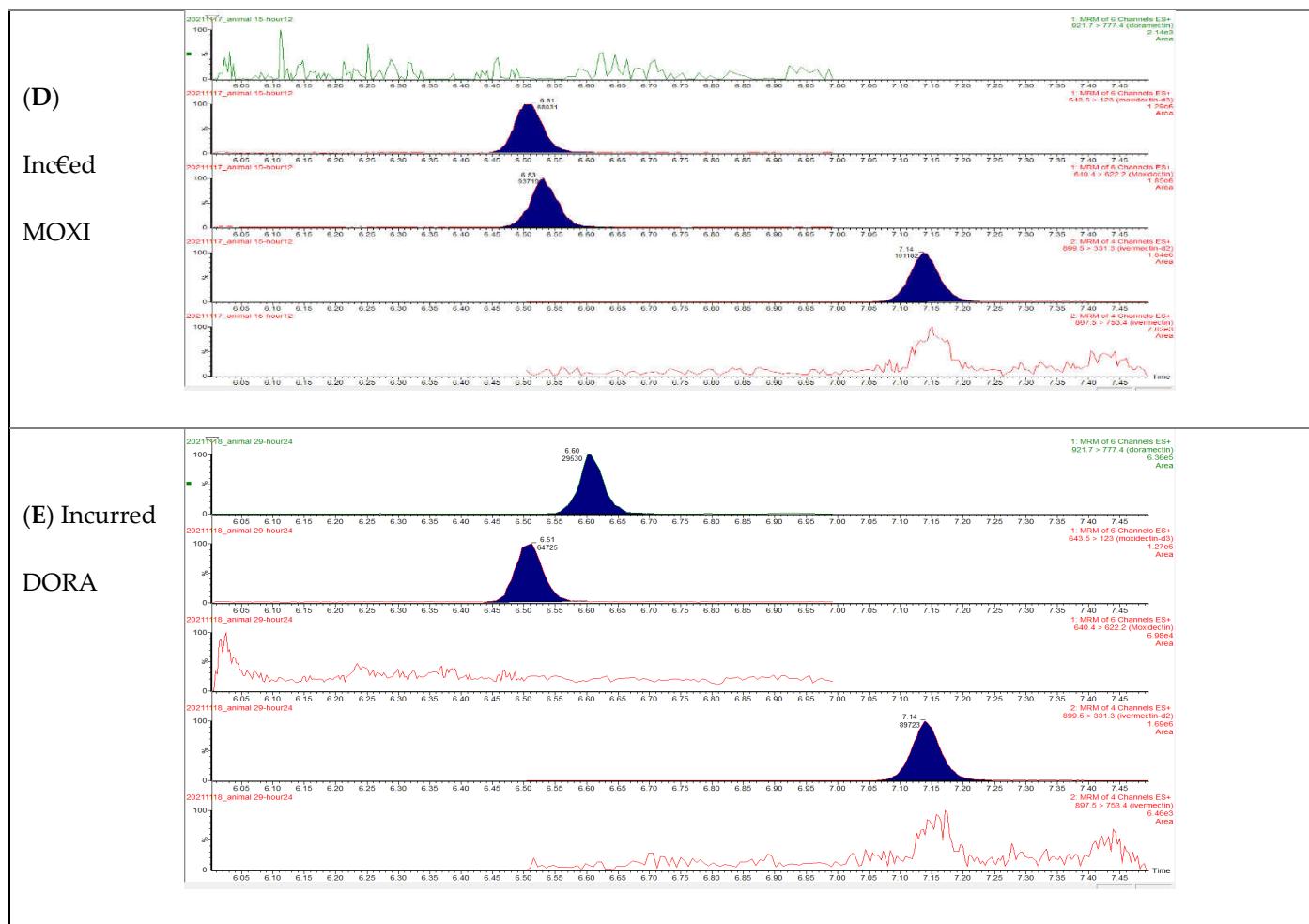


Figure 2. UHPLC-MS/MS chromatogram of (A) a blank plasma, (B) a quality control (QC) sample spiked with all analytes at a concentration of 50 ng/mL and incurred plasma samples containing (C) IVER (concentration: 128.89 ng/mL), (D) MOXI (concentration: 99.84 ng/mL), and (E) DORA (concentration: 14.60 ng/mL) that were extracted using 1% formic acid in acetonitrile as deproteinization solvent, followed by OstroTM 96-well plate clean-up.

2.2.5. Matrix Effect (ME) and Extraction Recovery (RE)

The efficiency to precipitate proteins and extract the analytes of interest from the plasma matrix (RE) and the signal enhancement or suppression due to matrix effect (ME) of the sample extracts on the LC-MS/MS instrument were evaluated for the analytes of interest IVER, DORA, and MOXI, using the final procedure, and the results are shown in Figure 3. As can be seen, RE ranged between 37.0 and 52.8% (see Figure 3A), whereas the ME was between 105.2–123.7% (see Figure 3B). Despite the low extraction recovery (RE), the isotopically labeled internal standards (IVER-d2 and MOXI-d3) effectively compensated the analyte losses in all three drugs (IVER, DORA, and MOXI) and maintained reproducibility and accuracy of the method. In this final method, the matrix effect (signal suppression or enhancement) due to co-eluting undetected matrix components was sufficiently minimized by using the OstroTM 96-well plates clean-up procedure. Furthermore, matrix-matched calibrator samples and the isotope-labeled internal standards used in the present method have also contributed to this minimum matrix effect. With respect to matrix effects, the coefficient of variation (CV) for IVER, DORA, and MOXI was 13%, 5%, and 4%, respectively, which was in line with the acceptance limit VICH GL49 [41] and EU recommendation [42].

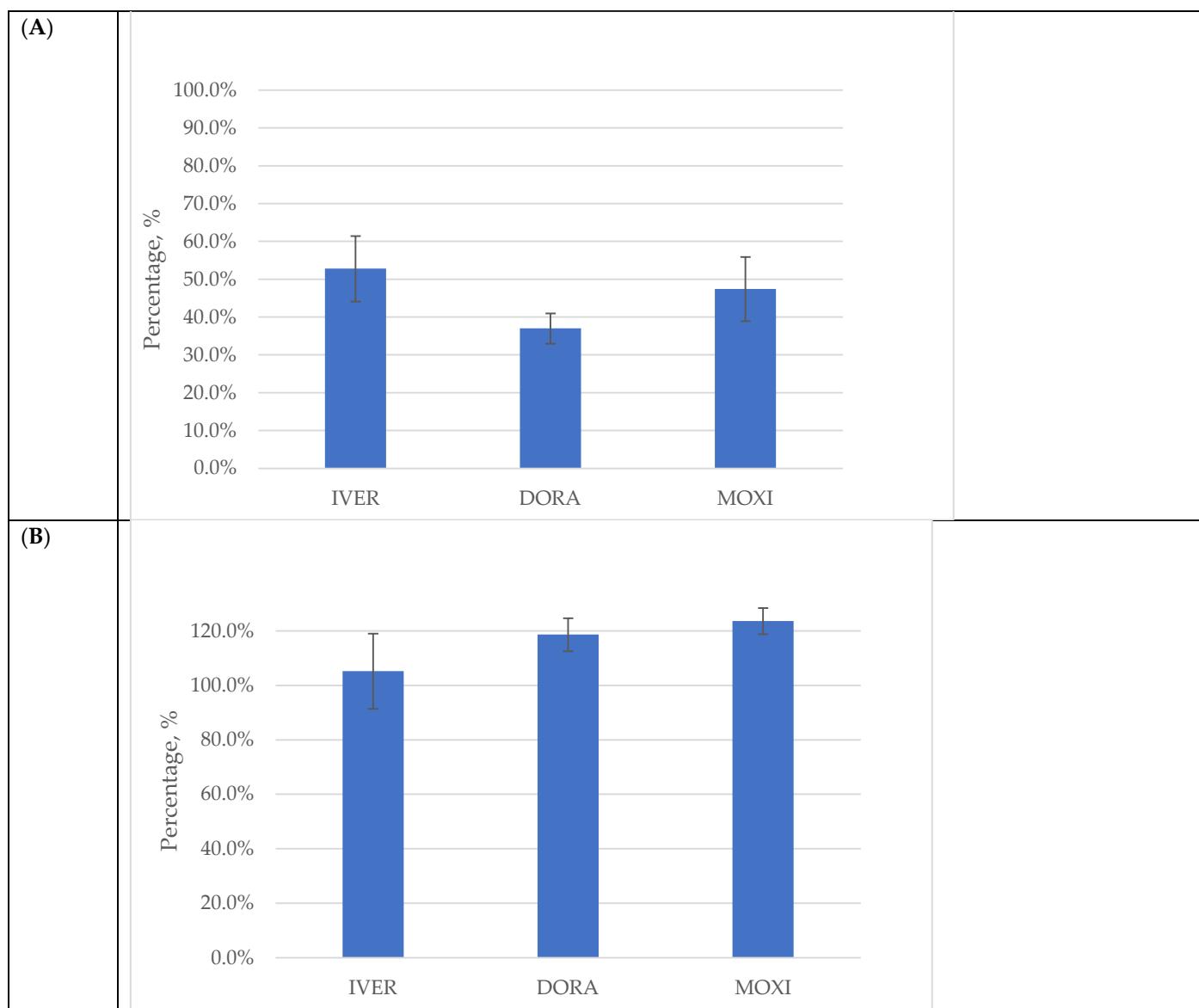


Figure 3. Results ($n = 3$) of the evaluation of extraction recovery (RE, panel A) and signal suppression or enhancement due to the matrix effect (ME, panel B) in bovine plasma spiked with 100 ng/mL ivermectin (IVER), doramectin (DORA), and moxidectin (MOXI) in using three different biological samples in each series ($n = 3$) after sample deproteinisation using 1% formic acid (FA) in acetonitrile (ACN) followed by OstroTM 96-well plate pass-through clean-up.

2.2.6. Stability

All the analytes were stable in stock and working solutions during storage for at least 21 months (results not shown). As can be seen in Table 4, all the analytes were found to be stable in the sample extract and during three freeze-thaw cycles ($\leq -15^{\circ}\text{C}$ to room temperature). Sample extracts were stored in capped 96-well collector plates for 12 days at $2-8^{\circ}\text{C}$, which corresponds with the autosampler temperature of the LC-MS/MS instrument. The mean found concentration fell within the acceptance ranges for accuracy and precision for all analytes at the tested concentration levels (5 and 50 ng/mL, see Table 4). These results demonstrate that analyte concentrations were not significantly affected by the applied extraction method, thus showing the practicability of the method for quantification of the analytes in bovine plasma.

Table 4. Evaluation of the stability of ivermectin (IVER), doramectin (DORA), and moxidectin (MOXI) in bovine plasma during three freeze/thaw cycles and in extracted samples ($n = 3$ replicates).

Freeze-Thaw Stability Samples ($\leq -15^{\circ}\text{C}$ to Room Temperature)				
	Theor. Conc. (ng/mL)	Mean Conc. \pm SD (ng/mL)	RSD %	Acc %
IVER	5.00	5.18 \pm 0.08	1.5	3.5
	50.00	48.26 \pm 0.39	0.8	-3.5
DORA	5.00	5.24 \pm 0.05	1.0	4.9
	50.00	52.37 \pm 0.19	0.4	4.7
MOXI	5.00	4.95 \pm 0.21	4.2	-0.9
	50.00	49.91 \pm 2.82	5.6	-0.2
Stability in Extracted Samples (for 12 Days at 2–8 °C)				
	Theor. Conc. (ng/mL)	Mean Conc. \pm SD (ng/mL)	RSD %	Acc %
IVER	5.00	5.14 \pm 0.07	1.3	2.9
	50.00	48.24 \pm 0.66	1.4	-3.5
DORA	5.00	5.23 \pm 0.04	0.7	4.6
	50.00	50.71 \pm 0.28	0.6	1.4
MOXI	5.00	5.23 \pm 0.04	0.7	4.6
	50.00	49.12 \pm 0.33	0.7	-1.8

Note: standard deviation, SD; Relative standard deviation, RSD; Accuracy, Acc.

The long-term stability of the analytes of interest was studied in acidified plasma (containing 0.5% FA, pH~4), since it was the aim to ship these acidified plasma samples from Ethiopia to Belgium, where the final sample preparation and UHPLC-MS/MS analysis were performed.

The blank bovine plasma that was acidified with 0.5% formic acid and spiked at an analyte concentration level of 100 ng/mL was extracted and analyzed after a storage period of 12 and 28 days and 9 months at $\leq -15^{\circ}\text{C}$ ($n = 3$ per time point). At each time point, mean peak area ratios (analyte/IS) were determined and compared with peak area ratios at the time of preparation (day 0). As can be seen from Figure 4, the calculated % recoveries fell within $90 \pm 10\%$ for all analytes, indicating that no significant degradation occurred within a storage period of 28 days. Over a long-term storage period (9 months) at $\leq -15^{\circ}\text{C}$, analyte concentrations decreased between 15–20% of the initial value, which is still acceptable.

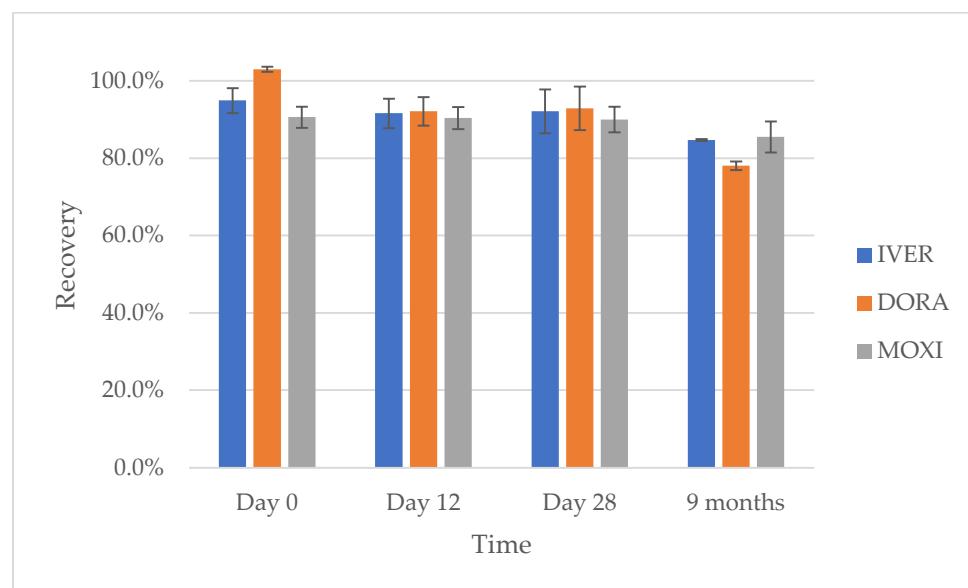


Figure 4. Analyte recovery of ivermectin (IVER), doramectin (DORA), and moxidectin (MOXI) in acidified bovine plasma (0.5% formic acid in plasma matrix, pH ~4) after long-term storage at $\leq -15^{\circ}\text{C}$.

2.2.7. Analysis of Biological Samples

The applicability of the current method was proved by UHPLC-MS/MS analysis of field PK samples, along with quality control samples using a matrix-matched calibration curve (concentration range from 1–500 ng/mL). As can be seen in Figure 5, analyte concentrations above the LOQ level (1 ng/mL) could be detected for all analytes until 21 days after a single subcutaneous administration of 0.2 mg/kg BW of IVER, DORA, and MOXI, showing the applicability of the developed method for future PK studies with these components in bovine animal species. This section may be divided into subheadings. It should provide a concise and precise description of the experimental results, their interpretation, as well as the experimental conclusions that can be drawn.

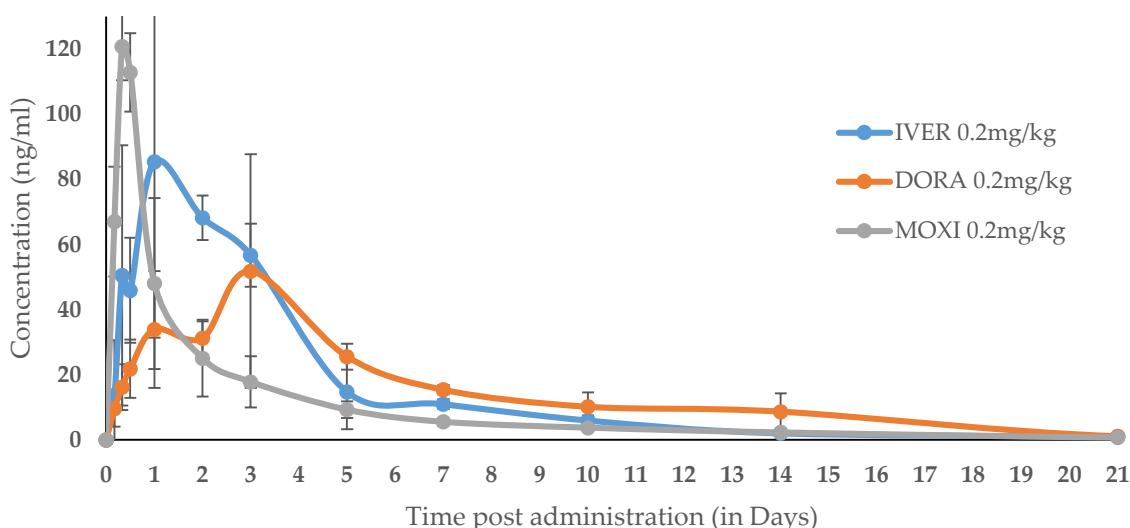


Figure 5. Plasma concentration versus time curves of IVER, DORA, and MOXI in bovine plasma after the subcutaneous administration of Ethiopian zebu cattle bovines ($n = 3$) with a single dose of 0.2 mg/kg BW.

3. Materials and Methods

3.1. Chemicals, Products and Reagents

All the standards (ivermectin (IVER), doramectin (DORA), moxidectin (MOXI), ivermectin-d2 (IVER-d2), and moxidectin-d3 (MOXI-d3)) were obtained from Sigma-Aldrich (Bornem, Belgium). All the standards were stored at $\leq -15^{\circ}\text{C}$. UPLC-MS grade acetonitrile, methanol, acetic acid, and formic acid obtained from Biosolve BV (Valkenswaard, The Netherlands) were used for the preparation of mobile phases. While, the ultrapure water used in the mobile phase was from a Milli-Q-SP reagent water system (Merck Millipore, Overijse, Belgium). Oasis Ostro® Protein Precipitation & Phospholipid Removal 96-well plates used for sample clean-up were purchased from Waters (Zellik, Belgium).

3.2. Blank Plasma Samples

Blank cattle plasma used for matrix-matched calibrator and quality control samples was obtained from donor cattle maintained at the Department of Large Animal Internal Medicine, Faculty of Veterinary Medicine, Ghent University. The animal was healthy and had no past history of drug therapy with macrocyclic lactones. The blank plasma was acidified with formic acid to achieve a final 0.5% formic acid in blank bovine plasma with a pH < 6 . The 0.5% formic acid in blank bovine plasma was prepared by the addition of 10% formic acid in a water solution into a 15-mL tube containing 10 mL blank plasma, followed by vortex mixing.

3.3. Incurred Plasma Samples

For a demonstration of the applicability of the UHPLC-MS/MS method, plasma samples from a pharmacokinetic (PK) study with MLs in bovines were analyzed. The drugs IVER, DORA, and MOXI were administered to local Ethiopian bovines (*Bos indicus*) once subcutaneously (dose: 0.2 mg/kg body weight) in a parallel study which was performed in Ethiopia after ethical review and approval (Ref. No. IUC-JU/M45/12) by the Institutional Animal Care and Use Committee at the veterinary facility of Jimma University, Ethiopia. Blood samples were collected from each animal before and periodically post-administration (p.a.), i.e., at 4, 8, 12, 24, and 48 h, and at 3, 5, 7, 10, 14, 21, 28, and 35 days, using EDTA vacutainer tubes. About 5 mL of a blood sample was collected from the left jugular vein. The collected blood samples were immediately put in an ice box and then centrifuged (within two hours) for 20 min at a maximum of 3000 rpm. The plasma was transferred into plastic tubes and stored in a deep freezer at $\leq -15^{\circ}\text{C}$.

To inactivate animal-borne viruses, plasma samples were treated with formic acid (0.5%) to lower the pH < 6 . Therefore, 1000 μL of plasma was transferred into a tube followed by the addition of 50 μL of a 10% FA solution in water. After vortex mixing, the samples were stored again in the freezer at $\leq -15^{\circ}\text{C}$ until shipment to Belgium (within two weeks).

3.4. Preparation of Standard Stock and Working Solutions

The standard stock solutions were prepared using acetonitrile as a solvent. IVER, DORA, MOXI, IVER-d2, and MOXI-d3 stock solutions were prepared at a concentration of 1 mg/mL based on the percent purity information from the manufacturer and stored at $\leq -15^{\circ}\text{C}$. The stock solutions of all analytes were stable for at least one year under these storage conditions.

Mixed working solution (WS_{mix}) of IVER, DORA, and MOXI (10 $\mu\text{g}/\text{mL}$) in acetonitrile was prepared by transferring 100 μL of each 1 mg/mL stock solution in a volumetric flask of 10.0 mL and addition of acetonitrile up to the mark, followed by gently mixing and equilibration for 5 min at room temperature. Further, WS_{mix} solutions with concentrations of 5000, 2500, 1000, 750, 500, and 250 ng/mL were prepared by appropriate dilution of WS_{mix} 10 $\mu\text{g}/\text{mL}$ with acetonitrile. Similarly, the WS_{mix} solutions with concentrations of

100, 50, 25, and 10 ng/mL were prepared from the WS_{mix} 1000 ng/mL by appropriate dilution with acetonitrile.

Individual working solutions (WS_{ind}) of 100 µg/mL ivermectin-d2 and moxidectin-d3 were prepared separately by diluting 100 µL of 1 mg/mL stock solution with 900 µL acetonitrile in an Eppendorf cup and vortex mixed. Further, a mixed working solution containing 1 µg/mL moxidectin-d3/ ivermectin-d2 (WS_{IS_mix}) was prepared by transferring 100 µL of each WS_{ind} 100 µg/mL of moxidectin-d3 and WS_{ind} 100 µg/mL of ivermectin-d2 in a volumetric flask of 10.0 mL and with an addition of acetonitrile up to the mark, followed by gently mixing. All working solutions were stored at ≤-15 °C.

3.5. Sample Preparation

The sample preparation was based on a deproteinisation with 750-µL of a 1% formic acid in acetonitrile solution, followed by an Ostro® 96-well plate pass-through clean-up.

For deproteinisation, 250-µL of the plasma sample was transferred to an Eppendorf cup, followed by the addition of 25-µL of the WS_{IS_mix} (1 µg/mL) and vortex mixing for 15 s. For the preparation of calibration/quality control samples with analyte concentrations of 1, 2.5, 5, 10, 25, 50, 75, 100, 250, and 500 ng/mL, and 225 µL of blank plasma was transferred to an Eppendorf cup and spiked with 25 µL of WS_{mix} 10, 25, 50, 100, 250, 500, 750, 1000, 2500 and 5000 ng/mL, respectively. Then, 25 µL of the WS_{IS_mix} (1 µg/mL) was added to each calibrator and quality control sample, followed by vortex mixing for 15 sec and equilibration for 5 min at room temperature.

To each of the above (spiked) samples (including the blank sample), 750 µL of a 1% formic acid in acetonitrile solution were added and vortex mixed for 5 min on a multi-tube vortex mixer (2500 rpm, BenchMixer™ XLQ, Sigma-Aldrich, Overijse, Belgium). Further, the samples were centrifuged at 13,000 rpm for 10 min. at 4 °C using a microcentrifuge (Biofuge Fresco, Sysmex, Hoeilaart, Belgium).

For further sample clean-up, the supernatant of each of the above samples was carefully transferred to an Ostro™ 96-well plate and closed by a polypropylene cap-mat. Then, vacuum was applied for 5 min allowing the transfer of each sample to a 2 mL square 96-well collector plate. The polypropylene cap-mat was replaced by a pre-slotted silicone/PTFE treated cap-mat before transfer to the autosampler of the UPLC-MS/MS instrument. Finally, a 10-µL aliquot was injected onto the UPLC-MS/MS instrument for analysis.

3.6. UHPLC-MS/MS Instrumentation

An Acquity H-Class ultra-high performance liquid chromatograph (UHPLC) system consisting of an Acquity UPLC H-Class Quaternary Solvent Manager and Flow-Through-Needle Sample Manager with temperature-controlled tray and column oven from Waters (Zellik, Belgium) was used. Chromatographic separation was achieved on an Acquity UPLC HSS T3 column (100 mm × 2.1 mm i.d., dp: 1.8 µm) in combination with an Acquity HSS T3 1.8 µm Vanguard pre-column, both from Waters.

Mobile phase A consisted of 0.01% acetic acid in water, while mobile phase B was 0.01% AA in methanol. A gradient elution was performed: 0–0.5 min (20% A, 80% B), 6.0 min (linear gradient to 99% B), 6.0–7.7 min (1% A, 99% B), 8.0 min (linear gradient to 20% A), 8.0–12.0 min (20% A, 80% B). The flow rate was 0.3 mL/min. The temperatures of the column oven and autosampler tray were set at 40 °C and 8 °C, respectively.

The UPLC column effluent was interfaced to a Xevo® TQ-S triple quadrupole tandem mass spectrometer system (MS/MS), equipped with an electrospray ionization (ESI) probe operating in the positive mode. A divert valve was used and the UPLC effluent was directed to the mass spectrometer from 5.5 to 8.0 min to avoid contamination and pollution of the mass spectrometer and to maintain sensitivity for a longer period.

Instrument parameters were optimized by direct infusion of working solutions of 100 ng/mL of all analytes and the ISs at a flow rate of 10 µL/min and in combination with the mobile phase (20% A, 80% B, flow rate: 200 µL/min).

The following Xevo® TQ-S mass spectrometer settings were used: capillary voltage: 3.5 kV; source offset: 60 V; source temperature: 150 °C; desolvation temperature: 500 °C; desolvation gas: 800 L/h; cone gas: 150 L/h; nebuliser pressure: 6.9 bar; LM resolution 1 and 2: 2.8; HM resolution 1 and 2: 15; ion energy 1 and 2: 0.2 and 0.8, respectively; collision gas flow: 0.15 mL/min.

MS/MS acquisition was performed, simultaneously for all the MLs in the plasma sample, in the multiple reaction monitoring (MRM) mode. The MRM transitions that were monitored are shown in Table 1. Data acquisition and processing were performed using the MassLynx® and TargetLynx® software (all from Waters).

3.7. Determination of Extraction Recovery (RE) and Matrix Effects (ME)

Extraction recovery and matrix effects were evaluated based on the post-extraction spike method described by Matuszewski et al., 2003 [43] and Chambers et al., 2007 [44]. Nine cattle plasma samples were prepared in three series' (A-series, B-series, and C-series) using three different biological samples in each series. The C-series samples were neat standard solutions prepared by spiking 250 µL of water with 25 µL of WS_{mix} 25 ng/mL and 25 µL of WS_{IS_mix} 1 µg/mL, followed by the addition of 750 µL of 1% FA in acetonitrile.

The B-series samples consisted of blank plasma that was subjected to the sample preparation procedure described above. Just after passing the supernatant of blank extract through the Oasis Ostro® 96-well plate, 25 µL of WS_{mix} 25 ng/mL and 25 µL of WS_{IS_mix} 1 µg/mL were added to the sample extract in the collector plate.

The A-series samples consisted of blank plasma that was spiked prior to the sample preparation with 25 µL of WS_{mix} 25 ng/mL and 25 µL of WS_{IS_mix} 1 µg/mL. After vortex mixing, the samples were subjected to the sample preparation procedure, as described above in Section 3.5.

The RE and ME were determined based on the absolute peak areas of the analytes in the different samples using the following formulas:

$$RE (\%) = (A\text{-series peak area}/B\text{-series peak area}) \times 100 \quad (1)$$

$$ME (\%) = (B\text{-series peak area}/C\text{-series peak area}) \times 100 \quad (2)$$

3.8. Evaluation of Stability of IVER, DORA and MOXI in Acidified Plasma

To allow the import of plasma samples from Ethiopia to Belgium, the pH of the plasma samples had to be <6. Therefore to assess the stability of the analytes in acidic pH, formic acid was added to blank plasma at different percentages (3%, 1%, 0.5%, 0.1%, and 0%) and the pH was recorded. In addition, the short-term stability of IVER, DORA, and MOXI in the acidified plasma was evaluated.

To 2700 µL of (acidified) blank plasma (% FA ranging from 0% to 3%), 300 µL of WS_{mix} 1 µg/mL was added, followed by vortex mixing for 15 sec and equilibration for 5 min at room temperature (final concentration of IVER, DORA, and MOXI: 100 ng/mL). The spiked (acidified) plasma was divided into 250-µL aliquots into Eppendorf cups and stored at <-15 °C until the moment of sample preparation and UHPLC-MS/MS analysis. Three 250-µL aliquots of each spiked (acidified) plasma were analyzed the same day of preparation (day 0). The remaining aliquots were thawed, extracted, and analyzed at day 12, day 28 and after 9 months, along with a freshly prepared spiked plasma sample (25 µL WS_{mix} 1.0 µg/mL added to 225 µL blank plasma, vortex mixed and equilibrated for 5 min at room temperature). The peak area ratios (analyte versus internal standard) in the spiked (acidified) plasma after storage at ≤-15 °C (test) were compared with the corresponding peak area ratios in freshly spiked plasma samples that were not acidified with formic acid (reference). The percentage analyte recovery was calculated as follows:

$$\% \text{ recovery} = (\text{peak area ratio test}/\text{peak area ratio reference}) \times 100 \quad (3)$$

3.9. Method Validation

The UHPLC-MS/MS method developed for the quantification of IVER, DORA, and MOXI in bovine plasma was validated based on international guidelines [41, 42, 45]. The following parameters were evaluated: linearity, within-day precision, and accuracy, between-day precision and accuracy [41], the limit of quantification (LOQ), limit of detection (LOD), carry-over, specificity, and stability. All the parameters were evaluated using spiked blank acidified (containing 0.5% formic acid) bovine plasma obtained from a healthy, untreated animal.

3.9.1. Linearity

Matrix-matched calibrator samples with IVER, DORA, and MOXI concentrations of 1, 2.5, 5, 10, 25, 50, 75, 100, 250, and 500 ng/mL were prepared by spiking 225 μ L of blank acidified plasma with 25- μ L of WS_{mix} 10, 25, 50, 100, 250, 500, 750, 1000, 2500 and 5000 ng/mL, respectively. Calibration curves were prepared freshly on three consecutive days. The correlation coefficient (*r*) and goodness-of-fit coefficient (GoF) % were evaluated and had to be ≥ 0.99 and $\leq 20\%$, respectively. Moreover, the back-calculated concentration of the analytes in each calibrator sample had to fulfill the criterion for accuracy. Evaluation of the weighting factor was based on the sum of the GoF-factors of the 3 calibration curves after applying unweighted, 1/*x*, and 1/*x*² weighted regression analysis. The weighting factor that resulted in the smallest sum of GoF was considered as most appropriate.

3.9.2. Precision and Accuracy

Quality control (QC) samples with analyte concentrations at the limit of quantification level (1 ng/mL) and at a low (QC-L, 5 ng/mL), medium (QC-M, 50 ng/mL), and high (QC-H, 250 ng/mL) [41,42] concentration level were prepared to evaluate within-run precision and accuracy. Six replicates at each concentration level were prepared on three different analysis days. Similarly, between-day precision and accuracy were evaluated by combining the results of the three data sets obtained for within-run precision and accuracy. The acceptance criteria for within-day and between-day accuracy was considered being between -20% to $+10\%$, -30% to $+10\%$ and -40% to $+20\%$ for analyte concentrations of ≥ 100 ng/mL, ≥ 10 to <100 ng/mL and ≥ 1 ng/mL to <10 ng/mL in cattle plasma, respectively, according to VICH GL49 [41] and EU recommendations [42]. Within-day and between-day precisions were evaluated based on the relative standard deviation (RSD%) and maximum standard deviation (RSDmax), respectively. The acceptance criteria for within-day precision were as follows: RSD (%) $< 25\%$, $< 15\%$ and $< 10\%$ for analyte concentrations ranging ≥ 1 to <10 ng/mL, ≥ 10 to <100 ng/mL and ≥ 100 ng/mL, respectively. For between-day precision, calculated RSD (%) values had to be lower than the RSDmax value as determined by the Horwitz equation with $RSDmax = 2(1 - 0.5 \log C)$ with *C* = concentration expressed as a decimal fraction (e.g., 1 ng/mL is entered as 10^{-9}). Accordingly, the acceptance criteria for between-day precision were set at RSD (%) $\leq 32\%$, 23% and 16% for analyte concentrations ranging between ≥ 1 to <10 ng/mL, ≥ 10 to <100 ng/mL and ≥ 100 ng/mL, respectively [41,42].

3.9.3. Limit of Quantification (LOQ) and Limit of Detection (LOD)

The LOQ was the smallest measured concentration of the analytes above which the accuracy and precision were acceptable. The LOD was the smallest measured concentration of an analyte from which it was possible to deduce the presence with acceptable certainty. The LOD was determined by calculating the theoretical analyte concentration that corresponded with a signal-to-noise (S/N) ratio of 3/1, based on the S/N ratio of the analytes in the LOQ samples [41,42].

3.9.4. Carry-over and Specificity

The carry-over was evaluated by injecting a solvent sample just after the highest calibrator sample. The specificity of the method was evaluated by analyzing a blank matrix sample extract. The response of the peak that eluted eventually at the same retention time as the analyte of interest should not be more than 20% of the mean peak area of the analytes in LOQ samples [41,42].

3.9.5. Stability

Freeze/thaw stability during three cycles (≤ -15 °C to room temperature), stability in sample extracts during storage at 2–8 °C for 12 days, and long-term stability in acidified plasma matrix (containing 0.5% FA) that was stored at ≤ -15 °C for 12 and 28 days and 9 months was assessed using blank cattle plasma spiked with the analytes of interest at a low (5 ng/mL), medium (50 ng/mL) or high (100 ng/mL) concentration levels. For the freeze/thaw stability and stability in extract experiment, quantification was performed using freshly prepared matrix-matched calibration curves.

4. Conclusions

A sensitive and reliable UHPLC-MS/MS method for simultaneous quantification of IVER, DORA, and MOXI in bovine plasma was developed and validated. The sample preparation consisted of straightforward deproteinisation followed by Ostro™ 96-well plate clean-up which was high-throughput and suitable for large number PK sample studies. With respect to all the validation parameters, the results of the current method fell within the set acceptance ranges.

The applicability of the validated method for field pharmacokinetic study was proved by the analysis of part of the ongoing pharmacokinetic samples. Results showed that the LOQ values (1 ng/mL) were sufficiently low and that the calibration range (1–500 ng/mL) was appropriate to allow a proper quantification of IVER, DORA, and MOXI. Moreover, due to the simple and straightforward sample preparation procedure in combination with a short analysis time, the method was suitable for the analysis of a large number of plasma samples ($n \geq 96$) per day. This proved the importance of the current method for use in large PK studies.

Supplementary Materials: Table S1: Evaluation of recovery of the standard solutions after pass-through on an Oasis® PRiME 96-well μ -Elution plate and Ostro™ 96-well plate, Figure S1: Influence of mobile phase composition on the peak area of each analyte, Figure S2: MS/MS chromatogram of a spiked plasma sample in different mobile phases.

Author Contributions: Conceptualization, M.D., S.D.B. and S.S.; funding acquisition, M.D., G.Z. and S.S.; methodology, S.D.B. and G.Z.; investigation, S.D.B. and G.Z.; supervision, M.D. and S.D.B.; validation and analysis, G.Z. and S.D.B.; project administration, G.Z., S.D.B. and S.S.; writing—original draft preparation, G.Z.; writing—review and editing, M.D., S.D.B. and G.Z. All authors have read and agreed to the published version of the manuscript.

Funding: This study was funded by the special research fund of Ghent University (DOZA/IL/DDC/AM/082b_2019) and Jimma University.

Institutional Review Board Statement: The animal study protocol was approved by the Institutional Review Board (or Ethics Committee) by the Faculty of Veterinary Medicine, Jimma University (Ethiopia) (protocol code IUC-JU/M45/12; 13/02/2019).

Informed Consent Statement: Not applicable.

Data Availability Statement: The data presented in this study are available on request from the corresponding author.

Acknowledgments: The authors are grateful to Jelle Lambrecht and Marc Cherlet for their technical support during the experiments in UHPLC-MS/MS analysis. We authors are also grateful to the Ethiopian Biodiversity Institute for the facilitation of the incurred sample shipment.

Conflicts of Interest: The authors declare no conflict of interest.

Sample Availability: Incurred plasma samples of the compounds are available from the authors.

Reference

1. World Health Organization. *World Malaria Report 2020: 20 Years of Global Progress and Challenges*; World Health Organization: Geneva, Switzerland, 2020.
2. Dyer, O. African malaria deaths set to dwarf COVID-19 fatalities as pandemic hits control efforts, WHO warns. *BMJ* 2020, **371**, m471. <https://doi.org/10.1136/bmj.m4711>.
3. Benelli, G.; Beier, J.C. Current vector control challenges in the fight against malaria. *Acta Trop.* 2017, **174**, 91–96. <https://doi.org/10.1016/j.actatropica.2017.06.028>.
4. Killeen, G.F.; Ranson, H. Insecticide-Resistant malaria vectors must be tackled. *Lancet* 2018, **391**, 1551–1552. [https://doi.org/10.1016/S0140-6736\(18\)30844-4](https://doi.org/10.1016/S0140-6736(18)30844-4).
5. Sheridan, R.; Desjardins, L. Determination of abamectin, doramectin, emamectin, eprinomectin, ivermectin, and moxidectin in milk by liquid chromatography electrospray tandem mass spectrometry. *J. AOAC Int.* 2006, **89**, 1088–1094. <https://doi.org/10.1093/jaoac/89.4.1088>.
6. Sougoufara, S.; Ottih, E.C.; Tripet, F. The need for new vector control approaches targeting outdoor biting Anopheline malaria vector communities. *Parasites Vectors* 2020, **13**, 295. <https://doi.org/10.1186/s13071-020-04170-7>.
7. Chaccour, C.; Rabinovich, N.R. Ivermectin to reduce malaria transmission II. Considerations regarding clinical development pathway. *Malar. J.* 2017, **16**, 161.
8. Imbahale, S.S.; Lopez, J.M.; Brew, J.; Paaijmans, K.; Rist, C.; Chaccour, C. Mapping the potential use of endectocide-treated cattle to reduce malaria transmission. *Sci. Rep.* 2019, **9**, 5826. <https://doi.org/10.1038/s41598-019-42356-x>.
9. Pasay, C.J.; Yakob, L.; Meredith, H.R.; Stewart, R.; Mills, P.C.; Dekkers, M.H.; Ong, O.; Llewellyn, S.; Hugo, R.L.E.; McCarthy, J.S. Treatment of pigs with endectocides as a complementary tool for combating malaria transmission by *Anopheles farauti* (ss) in Papua New Guinea. *Parasites Vectors* 2019, **12**, 124. <https://doi.org/10.1186/s13071-019-3392-0>.
10. Ruiz-Castillo, P.; Rist, C.; Rabinovich, R.; Chaccour, C. Insecticide-Treated livestock: A potential One Health approach to malaria control in Africa. *Trends Parasitol.* 2021, **38**, 112–123. <https://doi.org/10.1016/j.pt.2021.09.006>.
11. Waite, J.L.; Swain, S.; Lynch, P.A.; Sharma, S.; Haque, M.A.; Montgomery, J.; Thomas, M.B. Increasing the potential for malaria elimination by targeting zoophilic vectors. *Sci. Rep.* 2017, **7**, 40551. <https://doi.org/10.1038/srep40551>.
12. Merola, V.M.; Eubig, P.A. Toxicology of Avermectins and Milbemycins (Macrocyclic Lactones) and the Role of P-Glycoprotein in Dogs and Cats. *Vet. Clin. N. Am. Small Anim. Pract.* 2018, **48**, 991–1012. <https://doi.org/10.1016/j.cvsm.2018.07.002>.
13. Vercruyse, J.; Rew, R.S. *Macrocyclic Lactones in Antiparasitic Therapy*; CABI: New York, NY, USA, 2002.
14. Omitola, O.O.; Umunnakwe, C.U.; Bayegun, A.A.; Anifowose, S.A.; Mogaji, H.O.; Oluwole, A.S.; Odoemene, S.N.; Awolola, T.S.; Osipitan, A.A.; Sam-Wobo, S.O. Impacts of ivermectin mass drug administration for onchocerciasis on mosquito populations of Ogun state, Nigeria. *Parasites Vectors* 2021, **14**, 212. <https://doi.org/10.1186/s13071-021-04716-3>.
15. Vercruyse, J.; Deprez, P.; Everaert, D.; Bassissi, F.; Alvinerie, M. Breed differences in the pharmacokinetics of ivermectin administered subcutaneously to Holstein and Belgian Blue calves. *Vet. Parasitol.* 2008, **152**, 136–140. <https://doi.org/10.1016/j.vetpar.2007.11.021>.
16. Danaher, M.; Howells, L.C.; Crooks, S.R.; Cerkvenik-Flajs, V.; O’Keeffe, M. Review of methodology for the determination of macrocyclic lactone residues in biological matrices. *J. Chromatogr. B* 2006, **844**, 175–203. <https://doi.org/10.1016/j.jchromb.2006.07.035>.
17. De Montigny, P.; Shim, J.-S.K.; Pivnichny, J.V. Liquid chromatographic determination of ivermectin in animal plasma with trifluoroacetic anhydride and N-methylimidazole as the derivatization reagent. *J. Pharm. Biomed.* 1990, **8**, 507–511. [https://doi.org/10.1016/0731-7085\(90\)80060-3](https://doi.org/10.1016/0731-7085(90)80060-3).
18. Cerkvenik, V. Analytics of ivermectin residues in blood plasma and food of animal origin. *Slov. Vet. Res.* 2001, **38**, 127–40.
19. Kolberg, D.; Presta, M.; Wickert, C.; Adaime, M.; Zanella, R. Rapid and accurate simultaneous determination of abamectin and ivermectin in bovine milk by high performance liquid chromatography with fluorescence detection. *J. Braz. Chem. Soc.* 2009, **20**, 1220–1226. <https://doi.org/10.1590/S0103-50532009000700004>.
20. Durden, D.A.; Wotske, J. Quantitation and validation of macrolide endectocides in raw milk by negative ion electrospray MS/MS. *J. AOAC Int.* 2009, **92**, 580–596. <https://doi.org/10.1093/jaoac/92.2.580>.
21. Inoue, K.; Yoshimi, Y.; Hino, T.; Oka, H. Simultaneous determination of avermectins in bovine tissues by LC-MS/MS. *J. Sep. Sci.* 2009, **32**, 3596–3602. <https://doi.org/10.1002/jssc.200900411>.
22. Michelle Del Bianchi, A.C.; Fernandes, M.A.; Braga, P.A.d.C.; Monteiro, A.L.; Daniel, D.; Reyes, F.G. Moxidectin residues in lamb tissues: Development and validation of analytical method by UHPLC-MS/MS. *J. Chromatogr. B* 2018, **1072**, 390–396. <https://doi.org/10.1016/j.jchromb.2017.11.041>.
23. Baptista, R.C.; Fernandes, M.A.; Gilaverte, S.; Queiroz, S.C.; Assalin, M.R.; Ferracini, V.L.; Monteiro, A.L.; Reyes, F.G. Determination of moxidectin in serum by liquid chromatography-tandem mass spectrometry and its application in pharmacokinetic study in lambs. *J. Braz. Chem. Soc.* 2017, **28**, 250–256. <https://doi.org/10.5935/0103-5053.20160171>.

24. Chhonker, Y.S.; Ma, L.; Edi, C.; Murry, D.J. A sensitive and selective LC–MS/MS method for quantitation of moxidectin in human, mouse and monkey plasma: Clinical validation. *Bioanalysis* 2018, **10**, 1841–1852. <https://doi.org/10.4155/bio-2018-0110>.

25. Morbidelli, E.; Rambaldi, J.; Bitti, L.R.; Zagħiġi, A.; Barbarossa, A. A quick and simple method for the determination of ivermectin in dog plasma by LC–MS/MS. *MethodsX* 2018, **5**, 1503–1507. <https://doi.org/10.1016/j.mex.2018.11.011>.

26. Croubels, S.; De Baere, S.; Cherlet, M.; De Backer, P. Determination of ivermectin B_{1a} in animal plasma by liquid chromatography combined with electrospray ionization mass spectrometry. *J. Mass Spectrom.* 2002, **37**, 840–847. <https://doi.org/10.1002/jms.343>.

27. Côté, C.; Bergeron, A.; Mess, J.-N.; Furtado, M.; Garofolo, F. Matrix effect elimination during LC–MS/MS bioanalytical method development. *Bioanalysis* 2009, **1**, 1243–1257. <https://doi.org/10.4155/bio.09.117>.

28. Lahaie, M.; Mess, J.-N.; Furtado, M.; Garofolo, F. Elimination of LC–MS/MS matrix effect due to phospholipids using specific solid-phase extraction elution conditions. *Bioanalysis* 2010, **2**, 1011–1021. <https://doi.org/10.4155/bio.10.65>.

29. Wheaton, J.P.; Mantha, G.; Martin, J.; Chambers, E.E.; Diehl, D.M. *A Novel Sample Preparation Device for Improved Phospholipid Removal in Bioanalytical Assays*; Waters Corporation: Milford, MA, USA, 2010.

30. Young, M.S.; Tran, K. Oasis PRiME HLB Cartridge for Effective Cleanup of Meat Extracts Prior to Multi-Residue Veterinary Drug UPLC-MS Analysis; Waters Corporation: Milford, MA, USA, 2015.

31. Lauwers, M.; Croubels, S.; De Baere, S.; Sevastiyanova, M.; Romera Sierra, E.M.; Letor, B.; Gougoulias, C.; Devreese, M. Assessment of dried blood spots for multi-mycotoxin biomarker analysis in pigs and broiler chickens. *Toxins* 2019, **11**, 541. <https://doi.org/10.3390/toxins11090541>.

32. De Baere, S.; Croubels, S.; Novak, B.; Bichl, G.; Antonissen, G. Development and validation of a UPLC-MS/MS and UPLC-HR-MS method for the determination of fumonisin B1 and its hydrolysed metabolites and fumonisin b2 in broiler chicken plasma. *Toxins* 2018, **10**, 62. <https://doi.org/10.3390/toxins10020062>.

33. De Baere, S.; Devreese, M.; Watteyn, A.; Wyns, H.; Plessers, E.; De Backer, P.; Croubels, S. Development and validation of a liquid chromatography–tandem mass spectrometry method for the quantitative determination of gamithromycin in animal plasma, lung tissue and pulmonary epithelial lining fluid. *J. Chromatogr. A* 2015, **1398**, 73–82. <https://doi.org/10.1016/j.chroma.2015.04.022>.

34. Nilsson, L.B.; Eklund, G. Direct quantification in bioanalytical LC–MS/MS using internal calibration via analyte/stable isotope ratio. *J. Pharm. Biomed.* 2007, **43**, 1094–1099. <https://doi.org/10.1016/j.jpba.2006.09.030>.

35. Hewavitharana, A.K. Matrix matching in liquid chromatography–mass spectrometry with stable isotope labelled internal standards—Is it necessary? *J. Chromatogr. A* 2011, **1218**, 359–361. <https://doi.org/10.1016/j.chroma.2010.11.047>.

36. Goessens, T.; De Baere, S.; Troyer, N.; Deknock, A.; Goethals, P.; Lens, L.; Pasmans, F.; Croubels, S. Highly sensitive multi-residue analysis of veterinary drugs including coccidiostats and anthelmintics in pond water using UHPLC-MS/MS: Application to freshwater ponds in Flanders, Belgium. *Environ. Sci. Process. Impacts* 2020, **22**, 2117–2131. <https://doi.org/10.1039/D0EM00215A>.

37. Moschou, I.C.; Dasenaki, M.E.; Thomaidis, N.S. Ionization study and simultaneous determination of avermectins and milbemycines in fish tissue by LC-ESI-MS/MS. *J. Chromatogr. B* 2019, **1104**, 134–140. <https://doi.org/10.1016/j.jchromb.2018.11.017>.

38. Hofmann, D.; Sayasone, S.; Keiser, J. Development and validation of an LC-MS/MS method for the quantification of the anthelmintic drug moxidectin in a volumetric absorptive microsample, blood, and plasma: Application to a pharmacokinetic study of adults infected with *Strongyloides stercoralis* in Laos. *J. Chromatogr. B* 2021, **1166**, 122556. <https://doi.org/10.1016/j.jchromb.2021.122556>.

39. Wang, F.; Chen, J.; Cheng, H.; Tang, Z.; Zhang, G.; Niu, Z.; Pang, S.; Wang, X.; Lee, F.S.-C. Multi-residue method for the confirmation of four avermectin residues in food products of animal origin by ultra-performance liquid chromatography–tandem mass spectrometry. *Food Addit. Contam.* 2011, **28**, 627–639. <https://doi.org/10.1080/19440049.2011.563367>.

40. Li, X.; Guo, P.; Shan, Y.; Ke, Y.; Li, H.; Fu, Q.; Wang, Y.; Liu, T.; Xia, X. Determination of 82 veterinary drugs in swine waste lagoon sludge by ultra-high performance liquid chromatography–tandem mass spectrometry. *J. Chromatogr. A* 2017, **1499**, 57–64. <https://doi.org/10.1016/j.chroma.2017.03.055>.

41. EMA. VICH Topic GL49: Studies to Evaluate the Metabolism and Residues Kinetics of Veterinary Drugs in Human Food-Producing Animals: Validation of Analytical Methods Used in Residue Depletion Studies; Committee for Medicinal Products for Veterinary Use (CVMP): Amsterdam, The Netherlands, 2016.

42. Commission, E. Commission Decision 2002/657/EC of 12 August 2002 implementing Council Directive 96/23/EC concerning the performance of analytical methods and the interpretation of results. *Off. J. Eur. Communities* 2002, **50**, 8–36.

43. Matuszewski, B.K.; Constanzer, M.; Chavez-Eng, C. Strategies for the assessment of matrix effect in quantitative bioanalytical methods based on HPLC-MS/MS. *Anal. Chem.* 2003, **75**, 3019–3030. <https://doi.org/10.1021/ac020361s>.

44. Chambers, E.; Wagrowski-Diehl, D.M.; Lu, Z.; Mazzeo, J.R. Systematic and comprehensive strategy for reducing matrix effects in LC/MS/MS analyses. *J. Chromatogr. B* 2007, **852**, 22–34. <https://doi.org/10.1016/j.jchromb.2006.12.030>.

45. European Medicines Agency. Guideline on bioanalytical method validation. *EMEA/CHMP/EWP* 2011, **2009**, 192217.