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Cite this: *Anal. Methods*, 2024, 16, 7255

Development of a novel and useful methodology for the simultaneous analysis of multiclass contaminants in bovine fat†

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The instrumental development of tandem mass spectrometers fosters the actual trend in the trace analysis of organic compounds to the development of methods that allow the analysis of contaminants of the most diverse origin in a single analytical sample. The multiclass methods are aligned with the Who's One Health initiative while accomplishing the concepts of green chemistry. However, there are few reports of wide scope multiclass methods for the analysis of contaminants in this matrix. In this work, a method for the simultaneous determination of 62 compounds in bovine fat, was developed and validated following DG-SANTE/11813/2021 guidance. Liquid nitrogen milled fat was extracted with toluene and acetonitrile. Then a clean-up in a cryogenic bath, followed by dispersive solid phase extraction was performed. Residue determination was done using liquid chromatography for 44 compounds and gas chromatography for 18 compounds, both coupled to tandem mass spectrometry in MRM mode. The method was first developed and validated for two ectoparasiticides, then the scope was expanded for the analysis of 13 veterinary drugs and 49 pesticides. Recovery percentages were in the range of 60–134%, high matrix effect was observed in 50% of the scope of the method. Most compounds presented limits of quantification of 10 $\mu\text{g kg}^{-1}$ in compliance with international requirements. The method was applied to monitor 49 commercial samples to evaluate its performance. Eighty percent of samples contained ethion and 10% had fluazuron, both within MRLs, highlighting the need for proper withdrawal times.

Received 22nd March 2024
Accepted 12th September 2024

DOI: 10.1039/d4ay00516c

rsc.li/methods

Introduction

Pest control in agriculture and livestock involves treatments with a wide variety of synthetic compounds such as pesticides and veterinary drugs. Pesticides can be transferred from plants to animals through the food chain, subsequently contaminating by-products used for human consumption (muscle, milk, liver, and others).¹ Veterinary drugs employed for animal disease control can also contaminate the final products. To frame the contaminant contents within Good Agricultural and Veterinary Practices, the Codex Alimentarius Commission, multinational organizations such as the European

Commission, and regulation bodies of several countries, have established Maximum Residue Levels (MRLs) in different food matrices, including, the bovine ones.^{2–5}

Distinct regulations govern foods of animal and plant origin, labeled as “pesticides” and “veterinary drugs,” respectively, despite often being the same molecule (codex, EU). Compounds exclusively used in agriculture, lack established MRLs, defaulting to 10 parts per billion (ppb) for enforcement. This unharmonized regulations, has shaped diverse approaches to food safety. Residue analysis for animal-derived foods focuses on specific compound families (antibiotics, antiparasitics, among others), while plant origin analysis relies on multi-residue methods for fungicides, herbicides, and insecticides.

Given the potential for contamination through feed intake and sanitary management, it is essential to consider pesticide exposure in animals within the overall analytical scope of potentially dangerous contaminants in food of animal origin.

The complexity of analyzing contaminants in animal products, particularly in fat matrices, has led to the development of various analytical methods. Fat is a challenging matrix due to its ability to bioaccumulate lipophilic compounds, posing a risk of exposure through food.¹ Particularly, organophosphorus and pyrethroids pesticides, antibiotics such as chloramphenicol, parasiticides like ivermectin, with high K_{ow} tend to accumulate

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† Electronic supplementary information (ESI) available. See DOI: <https://doi.org/10.1039/d4ay00516c>

in fat tissues and partitionate between the extracting solvent and the fat itself.⁶

In addition, fat co-extractives are known to pollute the chromatographic system, poisoning electron capture and mass detectors, lowering the overall sensibility of the method; and forcing exhaustive clean-up steps to decrease the levels of co-extractives in the final extract.^{7,8}

Moreover, there are at least two types of fat samples with different water content and lipid composition, the one composed of 100% triglycerides, which is used to fry purposes, and fat tissues, with variable fat content. The U.S. Food and Drug Administration (FDA) defines fatty foods as those with fat content over 2%,⁹ but Lehotay *et al.*, considered that matrices with a fat content over 20% should be cataloged as high fatty foods.⁹

Sample treatment can influence the final concentration expressed for each compound. For fatty tissues, old methodologies preferred to melt the sample before extraction, and report the findings on the basis of the melted fat.¹⁰ The assumption made is that all the lipophilic compounds are concentrated in fat and their occurrence in other tissues is negligible. This is not the actual trend in the analysis of veterinary drugs and pesticide residues analysis. During the analysis of high fat matrices, it is recommended to express the concentration levels on a lipid adjusted basis, to obtain more representative results in comparison with unadjusted values.^{10,11} To compare results, the concentration might be expressed as milligrams or micrograms per kilogram of fat.¹⁰

Several methods have been developed to address this issue. Akre and Macneil, 2006 utilized Solid Phase Extraction (SPE) and gas chromatography electron capture detection for analyzing synthetic pyrethroids.¹² Sartarelli *et al.*, 2014 adapted the QuEChERS method for analyzing pesticides in bovine fat and meat, involving an extraction with acetonitrile and hexane, followed by a clean-up step with PSA and MgSO₄, and GC-MS analysis.¹³ Also, Hmadamin and Hassan developed a variation of the QuEChERS method for the analysis of six pesticides in bovine fat.¹⁴ Unterluggauer *et al.*, introduced the FATChers method, a QuEChERS-based approach suitable for fat and tricky matrices, which is widely used in the European Union Reference Laboratory for Food of Animal Origin.¹⁵ This method allows the determination of a wide range of compounds with varying polarity, using a combination of liquid–liquid extraction and dispersive solid phase extraction. Other methods, such as those developed by Castillo *et al.* (2012) and Hoyos *et al.* (2013), also focus on specific families of contaminants, typically those that are highly lipophilic and GC-amenable.^{16,17} However, these methods often concentrate on a limited range of compounds, leaving out other relevant contaminants.

A general thought in the past was that this type of methods should be developed only to monitor highly lipophilic compounds, such as organochlorine, organophosphate or pyrethroids insecticides, all GC amenable compounds, leaving out other families of contaminants.⁹ Nowadays, new lipophilic compounds that are heat sensitive and which also tend to bioaccumulate in fat tissues,^{18,19} can only be analysed by HPLC

(ivermectins, spinosads, spirotetramat and metabolites) and must be determined as well.⁹

In line with green chemistry concepts, the present work aims to develop and validate a simple and comprehensive multiclass method for the simultaneous analysis of 62 pesticides and veterinary drugs in fat. The proposed method seeks to address the limitations of previous approaches by including a broader range of contaminants, particularly those that are lipophilic, heat-sensitive, and tend to bioaccumulate in fat tissues. By integrating various extraction and clean-up steps, the method offers a more holistic approach to food safety, providing a more representative analysis of potentially hazardous compounds in animal-derived products. This comprehensive approach allows for the simultaneous determination of multiple contaminants at trace levels after a single sample preparation step, enhancing the overall assessment of food safety in the production chain.

Experimental

Materials

Samples of perirenal fat for method development and validation were obtained from local butcheries, then stored at −20 °C until analysis. Analytical standards in a range of purity of 96–99.9%, were purchased from Dr Ehrenstorfer (Augsburg, Germany). All solvents were of pesticide analysis grade, EtOAc was obtained from J.T Baker (NJ, USA), dichloromethane (CH₂Cl₂), *n*-hexane, MeCN and methanol (MeOH) from Merck (Darmstadt, Germany) and toluene was purchased from Pharmco (CT, USA). Water used for HPLC-MS/MS analysis was obtained from a Barnstead EASY pure RoDi from Thermo Scientific (Waltham, USA). Ammonium Formate (HCOONH₄) and MgSO₄ were acquired from J.T Baker (NJ, USA). Sodium Chloride (NaCl), PSA and 40–60 µm RPC-18 were purchased from Scharlau (Barcelona, Spain). Alumina and Florisil were provided by Carlo Erba (Sabadell, Spain). Individual stock standard solutions of the targeted compounds at 2000 mg L^{−1} were prepared according to solubility properties in EtOAc, MeCN, and MeOH, and stored at −18 °C. Working solutions of 100 and 10 mg L^{−1} for the spiking procedure and matrix-matched calibration, were prepared by dilution of the stock standard solutions.

Instrumental analysis

Liquid chromatography analysis was conducted with an Agilent 1200 series coupled to a Sciex 4000 Qtrap (Concord, Canada). Quadrupole-linear ion trap (QTrap®) was operated in triple quadrupole MS/MS mode, in positive and negative ESI polarity. Two transitions for the correct identification of each compound were selected. Source voltages were set at 4000 and −4500 V for the positive and negative ionization modes, respectively. Solvent evaporation in the source was assisted by heated nitrogen as drying gas (500 °C per 50 psi). Collision Energies (CE) and Declustering Potentials (DPs) for each investigated compound were optimized, after direct infusion, using Analyst Software v 1.5.1. Scheduled™ MRM mode was selected to increase compound sensitivity, as it is shown in Table ESI 1 and 2.†

For the analysis of pesticides and veterinary drugs the positive MS/MS mode was performed using a mobile phase (MP) of 0.1% formic acid and 5 mM of HCOONH_4 in water (A) and 0.1% formic acid and 10 mM of HCOONH_4 in MeOH (B). The gradient was 0 min, 98% A, 12 min, 0% A (4 min hold time), and 21 min, 98% A. The re-equilibration time was 5 min.

For negative MS/MS mode, the MP was a mixture of 0.1% of formic acid in water (A) and MeCN (B). The chromatographic method held the initial mobile phase composition 0 min, 50% B; 5 min, 100% B (4 min hold time); 10 min, 50% B. The re-equilibration time until the next injection took 5 min. The flow rate used was 0.6 mL min^{-1} for positive and negative mode, and the injection volume was $5 \mu\text{L}$.

GC amenable analytes were determined using a GC-MS/MS with a Shimadzu GC-2010 plus coupled to MS/MS TQ 8050 spectrometer in Electron Impact ionization (EI) mode. The target analytes were separated in an RTX-5MS (5% diphenyl/95% dimethyl polysiloxane) analytical column ($30 \text{ m} \times 0.25 \text{ mm i.d.} \times 0.25 \text{ m film thickness}$), from Restek, (Bellfonte, PA, USA). Tandem mass spectrometry detection was performed in the multiple reaction monitoring (MRM) mode using transitions and collision energies previously selected for each compound using Smart Pesticide Database (SPDB) provided by Shimadzu Corporation.²⁰

The injector temperature was set at 240°C , interface temperature at 300°C , and Helium at 14 mL s^{-1} as carrier gas. Oven temperature: 0 min, 80°C (2 min hold); 7 min, 180°C ($20^\circ\text{C min}^{-1}$ rate); 31 min, 300°C (5°C min^{-1} rate, 3 min hold). The injection volume was $1 \mu\text{L}$ in splitless mode. The retention times for each pesticide were set using the Automatic Adjustment Retention Time function (AART)TM (Shimadzu). The AART function establish the retention times of the target components based on Linear Retention Indices (LRI) and the retention times of *n*-alkanes. The final MRM method parameters are shown in Table ESI 1.†

Sample preparation

Pieces of frozen perirenal fat samples of about $1 \times 1 \text{ cm}$ were placed in a mixer with liquid nitrogen and the mixture milled to obtain a fine powder. N_2 was allowed to evaporate and the comminuted fat sample was transferred to polypropylene flasks and stored at -18°C . Blank samples for all fortification experiments and matrix-matched calibration curves were analyzed using the same analytical method, confirming that no pesticides or veterinary drugs from the scope, were present above the quantification limits.

Method development and validation

Preliminary tests: Three different strategies were evaluated in terms of recovery percentages and relative standard deviations during method development: QuEChERS, Ethyl Acetate method and Matrix Solid Phase Dispersion (MSPD).^{15,21,22} For every combination, two sample pre-treatments were tested; cutting the sample in small pieces with a knife or cryogenic freezing with liquid N_2 . All the combinations are summarized in Table 1.

Table 1 Methods tested, critical steps and the variables tested and compared during method optimization

Method	Critical step	Variables
MSPD	Sample homogenization Dispersant Elution	With or without N_2 Alumina or florisil Type of solvent Volume amount Concentration factor
QuEChERS	Sample homogenization Sample amount Extraction Purification	With or without N_2 2 or 5 g Type of solvent Amount and type of sorbent
Ethyl acetate	Sample homogenization Sample amount Extraction Purification	With or without N_2 2 or 5 g Type of solvent Amount and type of sorbent

QuEChERS and ethyl acetate methods

For QuEChERS and ethyl acetate, some variations of the original methods were evaluated. Two sample amounts (2 and 5 g) were tested. As extraction agents EtOAc, MeCN, and MeCN : Toluene (2 : 1, *v/v*) were assessed. Moreover, two different clean-up sorbents, (PSA and RPC-18) were selected, in two amounts (25 and 50 mg mL^{-1}).

MSPD

For MSPD the selection of an appropriate dispersive agent and the types of solvents for the elution step are critical. Therefore, to obtain the best extraction conditions, method comparison was performed selecting alumina and florisil as sorbents and two different volumes (10 and 15 mL) of EtOAc and hexane saturated in MeCN as elution solvent.

Selected sample preparation methodology

The final method consisted of weighting (2 ± 0.5) g of liquid N_2 milled perirenal fat previously homogenized in a 50 mL-polypropylene falcon tube followed by the addition of 5.0 mL of toluene and 5 minutes vortex. Then, 10.0 mL of MeCN were added and shaken in a vortex for 1 minute. Then the tubes were centrifuged at $4400 \times g$ for 4 minutes, placed in a cryogenic bath at -42°C for 15 minutes and centrifuged again for 2 minutes. Finally, eight mL of the supernatant were transferred to a 15 mL-falcon tube containing 360 mg of PSA, 360 mg of RPC-18 and 750 mg of anhydrous MgSO_4 , vortexed for 1 minute and centrifuged at $4400 \times g$ for 4 minutes (Fig. 1). For GC-MS/MS analysis, 2 mL aliquot was concentrated under nitrogen stream at 55°C for 20 minutes and re-dissolved in one mL of EtOAc. For HPLC-MS/MS analysis, 2 mL aliquot was concentrated in the same conditions, re-dissolved in one mL of MeCN, and stored overnight at -20°C . The extract was filtered with a $0.45 \mu\text{m}$ PVDF filter and transferred to an amber vial for injection.

Total lipidic content determination was made by gravimetry, as in the method developed by Folch *et al.*²³ This percentage was

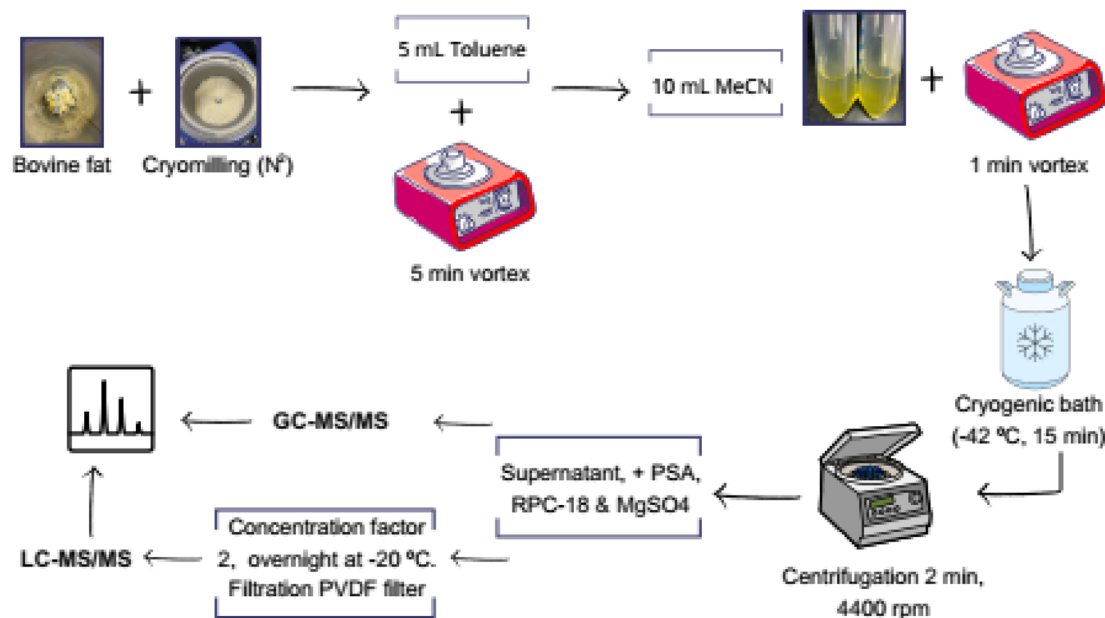


Fig. 1 Scheme of the validated methodology for multiclass analysis of veterinary drugs and pesticides in bovine fat.

used in the eqn (1) to express the concentration of each sample in 100% fat.

$$\text{Fat content} = \frac{\text{Experimental concentration} \times 100}{\text{Lipid content}} \quad (1)$$

Method validation

Identification criteria. Retention times, two characteristic transitions and the area relation between them (ion ratio), were selected as identification criteria according to Document N° SANTE/11813/2021.²⁴

The evaluation of the selectivity was performed through the injection of blank matrix extractions spiked with a mix of the selected compounds, to check the presence of potential interferences.

Validation parameters. Trueness, precision, linearity, matrix effect and limit of quantification, were determined based on the specifications established in Document N° SANTE/11813/2021.²⁴

Trueness and precision were studied through the determination of recoveries (%Rec) and relative standard deviations (%RSD), at 10, 50 and 100 µg kg⁻¹ by quintuplicate.

Recoveries were calculated based on the comparison between the concentration calculated with the matrix-matched calibration curve and the true concentration for the spiked analytes at each fortification level.

Repeatability and intermediate precision were tested on different days, a %RSD < 20 was considered acceptable.²⁴

The linearity was studied by preparing six-point calibration curves; in pure solvent and in matrix extract, in the range of 5–100 µg kg⁻¹ for HPLC and GC-MS/MS. Coefficient of variations (*r*²) and Back Calculated Concentration (BCC) residuals of each

calibration curve were determined for all the compounds being less than ± 20%.

Matrix effect (ME) was calculated by comparing the slopes of the curves in pure solvent and in the matrix-matched calibration curve. SANTE guidelines suggest that for every compound with a %ME > 20 in absolute value, matrix-matched calibration is mandatory.²⁴ Compounds were classified in three ranges of ME percentages according to Kmellar *et al.*, (2008).²⁵

LOQs were determined as the lowest concentration of each analyte that has been validated with acceptable trueness and precision by applying the complete analytical method for both instruments.²⁴

Results and discussion

As stated above, lipophilic contaminants in animal origin foods, might come either from the feed, which is normally some agricultural waste, or from veterinary drugs used to prevent diseases and pests. It is well known that adipose tissue is a relevant reservoir of lipophilic compounds.²⁶ These contaminants, at relatively high concentrations, can bioaccumulate within the fatty tissues. When consumed by human they can be released by the enzymes of the digestive tract. To perform an efficient and comprehensive evaluation of food safety, the development of multiclass methods to analyze multiple contaminants of diverse origin in a single chromatographic run is of great importance. Also, the use of multiclass methods enhances the productivity of routine laboratories.

Scope of the validated method

Being fat a highly lipidic matrix, non and middle-polar compounds with middle to high *K_{ow}*, tend to accumulate in this tissue.^{7,9,27} Assuming that compounds with high *pK_{ow}*

Table 2 List of selected analytes: results of validation parameters (instrumental method, trueness, precision and LOQ) and EU MRLs. nd: not detected

Compound	Instrumental method	10 $\mu\text{g kg}^{-1}$		50 $\mu\text{g kg}^{-1}$		100 $\mu\text{g kg}^{-1}$		LOQ ($\mu\text{g kg}^{-1}$)	Matrix effect	MRL ($\mu\text{g kg}^{-1}$)
		% rec	% RSD	% rec	% RSD	% rec	% RSD			
Abamectin	HPLC-MS/MS	88	6	84	10	80	9	10	109	10
Alachlor	HPLC-MS/MS	81	14	94	2	90	5	10	1.3	10
Ametryn	HPLC-MS/MS	101	4	113	4	106	4	10	−8.2	10
Atrazine	HPLC-MS/MS	97	5	116	3	103	3	10	−6.3	10
Azinphos methyl	GC-MS/MS	91	16	104	15	94	18	10	92	10
β -Cyfluthrin	GC-MS/MS	nd	nd	112	7	101	9	50	71	200
Byfenthrin	GC-MS/MS	102	54	116	12	96	14	10	49	3000
Boscalid	HPLC-MS/MS	143	13	117	5	98	2	10	16	300
Carbofuran	HPLC-MS/MS	67	14	115	6	110	10	10	15	10
Cypermethrin	GC-MS/MS	77	19	90	15	97	11	10	66	2000
Clomazone	HPLC-MS/MS	96	4	87	3	84	3	10	−1.3	10
Chlorantraniliprole	HPLC-MS/MS	104	10	115	5	100	5	10	7.1	200
Chlorfenvinphos	GC-MS/MS	79	11	114	7	106	9	10	54	10
Chlorpyrifos methyl	GC-MS/MS	72	8	105	8	99	12	10	54	10
Coumaphos	HPLC-MS/MS	108	4	106	6	97	12	10	74	10
Diazinon	HPLC-MS/MS	81	6	112	8	96	10	10	38	700
Difenoconazole	HPLC-MS/MS	106	13	133	13	128	19	10	−6.3	50
Deltamethrin	GC-MS/MS	70	12	107	10	99	12	10	66	500
Doramectin	HPLC-MS/MS	88	6	69	10	73	11	10	104	150
Emamectin benzoate	HPLC-MS/MS	84	10	80	18	78	9	10	−3.9	20
Endosulfan sulfate	GC-MS/MS			120	3	98	15	50	35	50
Eprinomectin	HPLC-MS/MS	92	13	89	10	76	13	10	107	250
Ethion	HPLC-MS/MS	90	8	101	3	90	7	10	83	10
Fenhexamid	HPLC-MS/MS	63	19	108	16	66	19	10	−11	50
Fenthion	HPLC-MS/MS	70	4	87	18	88	11	10	59	50
Fenvalerate	GC-MS/MS	70	10	106	10	100	9	10	62	250
Fipronil	GC-MS/MS	86	4	134	7	111	11	10	56	30
Fluazuron	HPLC-MS/MS	130	12	103	8	96	25	10	66	7000
Flutriafol	HPLC-MS/MS	116	7	127	3	104	1	10	−7.2	10
Fluvalinate	GC-MS/MS	70	7	120	15	109	12	10	53	300
Haloxypop methyl	HPLC-MS/MS	100	5	119	4	97	10	10	7.2	10
Hexythiazox	HPLC-MS/MS	96	18	87	7	82	9	10	105	50
Iprodione	GC-MS/MS	67	19	105	10	104	11	10	69	10
Isoprothiolane	HPLC-MS/MS	100	4	116	4	110	4	10	1.8	10
Ivermectin	HPLC-MS/MS	86	8	73	16	60	8	10	59	100
Kresoxim methyl	HPLC-MS/MS	96	7	119	2	111	8	10	−1.8	50
Lambda cyhalothrin	GC-MS/MS	104	19	124	8	112	14	10	49	3000
Linuron	HPLC-MS/MS	91	4	111	3	108	5	10	−12	10
Malaoxon	HPLC-MS/MS	111	3	115	3	98	2	10	−3.5	20
Malathion	HPLC-MS/MS	108	5	111	6	97	2	10	5	20
Mebendazole	HPLC-MS/MS	104	4	117	3	106	4	10	−2.3	60
Metconazole	HPLC-MS/MS	123	11	81	10	80	6	10	−1.8	20
Methiocarb	HPLC-MS/MS	89	6	104	4	96	5	10	−12	30
Metolachlor	HPLC-MS/MS	94	5	101	3	98	4	10	−4.3	10
Mirex	GC-MS/MS	60	17	81	7	68	11	10	34	10
Monensin	HPLC-MS/MS	83	15	71	10	74	13	10	101	10
Moxidectin	HPLC-MS/MS	72	19	60	20	50	21	10	101	500
<i>p,p'</i> -DDD	GC-MS/MS	64	24	97	14	110	12	10	90	10
<i>p,p'</i> -DDT	GC-MS/MS	67	12	65	12	61	17	10	39	10
<i>p,p'</i> -DDE	GC-MS/MS	61	10	79	5	77	8	10	38	10
Parathion methyl	GC-MS/MS	82	5	117	11	106	10	10	74	10
Pendimetalin	HPLC-MS/MS	91	11	79	5	76	9	10	109	200
Permethrin	GC-MS/MS	80	29	114	16	94	13	10	57	500
Pyraclostrobin	HPLC-MS/MS	106	9	138	5	102	19	10	−12	50
Pirimifos methyl	HPLC-MS/MS	76	17	115	10	105	12	10	41	10
Propiconazole	HPLC-MS/MS	96	8	92	5	95	8	10	−0.73	10
Spinosyd A	HPLC-MS/MS	63	27	84	14	73	7	10	−24	3000
Spinosyd D	HPLC-MS/MS	96	23	87	20	104	16	10	−35	3000
Tebuconazole	HPLC-MS/MS	115	12	109	8	114	11	10	0.6	100
Tebufenozide	HPLC-MS/MS	116	8	128	3	106	4	10	−11	50

Table 2 (Contd.)

Compound	Instrumental method	10 $\mu\text{g kg}^{-1}$		50 $\mu\text{g kg}^{-1}$		100 $\mu\text{g kg}^{-1}$		LOQ ($\mu\text{g kg}^{-1}$)	Matrix effect	MRL ($\mu\text{g kg}^{-1}$)
		% rec	% RSD	% rec	% RSD	% rec	% RSD			
Tetraconazole	HPLC-MS/MS	115	18	117	4	104	3	10	−12	200
Triclabendazole	HPLC-MS/MS	101	6	97	9	99	3	10	−0.7	100
Tryfloxystrobin	HPLC-MS/MS	110	6	128	4	105	8	10	−11	60
Triflumuron	HPLC-MS/MS	133	32	99	12	71	38	10	−11	100
Triticonazole	HPLC-MS/MS	107	27	100	18	86	10	10	−5	10
Vinclozolin	GC-MS/MS	81	9	129	9	99	10	10	49	10

would tend to bioaccumulate, two criteria were considered for the selection of the scope. First, the Uruguayan National Program of Biologic Residues (PNRB), which establishes the directives for local residue monitoring.²⁸ Second, other compounds used in cattle breeding, as well as those agricultural productive chains whose by-products are used as feed for cattle.^{18,19} The final list was checked with the international requirements of Codex Alimentarius and EU regulations to comprehensively include important MRLs.^{2–5} Based on these

criteria, we selected 70 compounds (Table 2) either classified as pesticides or veterinary drugs to be analyzed in a single run. Fig. 2 shows the HPLC-MS/MS MRM transitions at 100 $\mu\text{g kg}^{-1}$ of a fat extract and the Extracted Ion Chromatograms (XIC) at 10, 50 and 100 $\mu\text{g kg}^{-1}$ of three compounds.

Sample preparation

The particle size of the sample is critical for the quantitative extraction of the analytes from the fatty matrix.²⁹ The cryo-

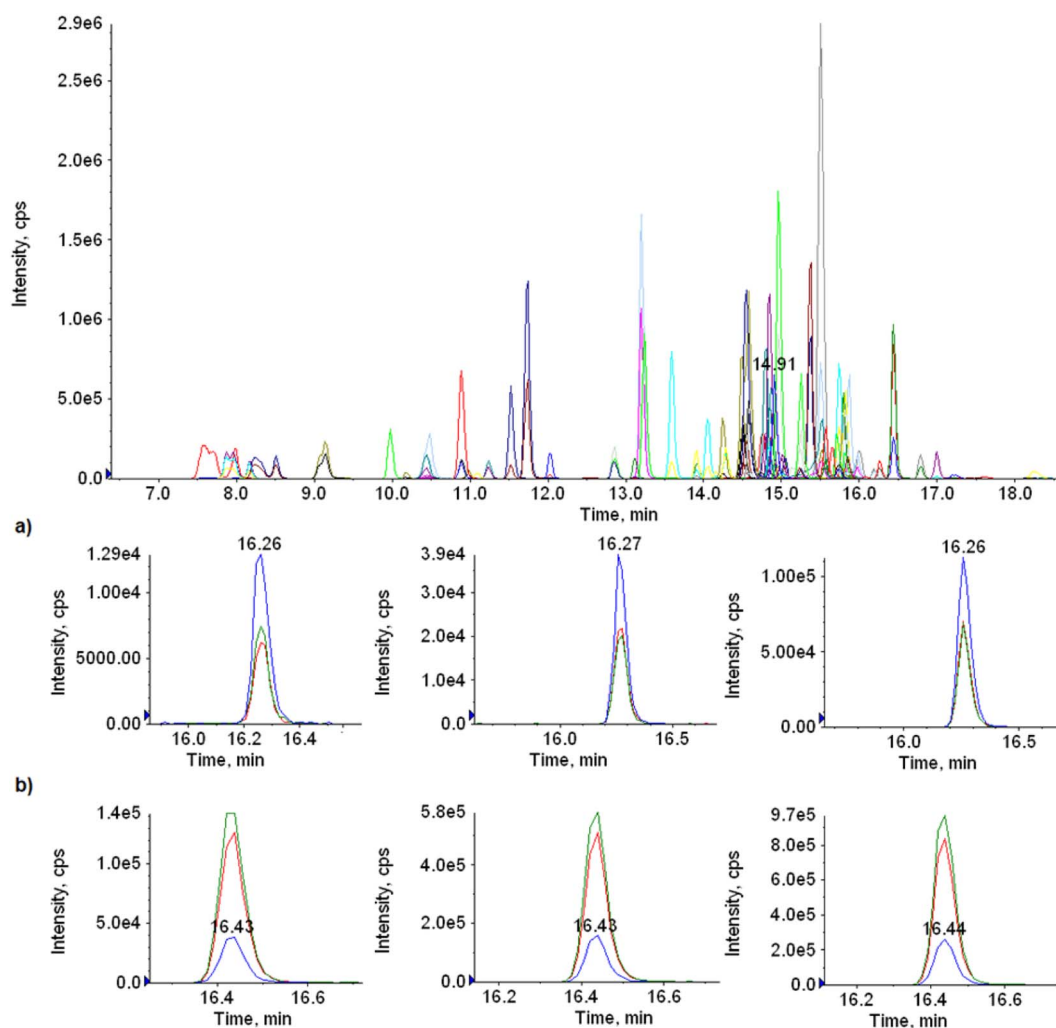


Fig. 2 HPLC-MS/MS MRM transitions at 100 $\mu\text{g kg}^{-1}$ of a fat extract and the Extracted Ion Chromatograms (XIC) of (a) fluazuron and (b) ethion at, from left to right, 10, 50 and 100 $\mu\text{g kg}^{-1}$.

analysis. So, the solution was kept at $-18\text{ }^{\circ}\text{C}$ overnight, filtered and analyzed.

Method validation

After sample pretreatment and method extraction were optimized, five replicates of spiked blanks of fat at different concentration levels (10, 50, and 100 $\mu\text{g kg}^{-1}$) were analyzed to evaluate trueness (%Rec), repeatability, and intermediate precision (%RSD) of the procedure. According to Document N° SANTE/11813/2021, 62 out of the 70 compounds analyzed presented acceptable recovery percentages (70–120%) (Table 2). Regarding reproducibility all compounds showed %RSD <20. The LOQ was set as 10 $\mu\text{g kg}^{-1}$ for all compounds except for β -cyfluthrin and endosulfan sulfate which presented a limit of 50 $\mu\text{g kg}^{-1}$. As the MRLs for these compounds in fat are usually higher than in other matrices, the LOQs fairly comply with the international requirements of the EU and Codex Alimentarius.²⁻⁵

Almost all compounds showed good linearity between 5–100 $\mu\text{g kg}^{-1}$ for both instruments in pure solvent and matrix-matched, only β -cyfluthrin and endosulfan sulphate had a linear range between 10–100 $\mu\text{g kg}^{-1}$. Residuals expressed as BCC were below 20% for all the studied compounds.

Although the exhaustive clean-up of the method, matrix effects (ME) were observed for 35 out of the 62 compounds (Fig. 3). Around 37% of the analytes had strong matrix effect (over 50%), whereas 17% showed medium ME, (between 20 and 50%). No relationship between physicochemical properties and the ME profile was found, being organophosphorus pesticides, avermectins and pyrethroids the group with higher ME. As expected, in GC-MS/MS the predominant effect observed was signal enhancement. This may be since certain co-extractives are acting as analyte protectants, preventing compound degradation in the injector or along the chromatographic run.

In HPLC-MS/MS, ME alternated between signal suppression and enhancement depending on each compound. Based on these results, matrix-matched calibration was chosen for the quantification.



Real samples analysis

The validated method was checked against 49 samples obtained before the ethion restriction in Uruguay, to verify the compliance with the MRLs and generate information about the occurrence of these contaminants in local final products. Ethion was present in 80% of the selected samples, in most of them at concentrations below the LOQ ($10 \mu\text{g kg}^{-1}$) complying with both national and EU MRLs ($10 \mu\text{g kg}^{-1}$). However, it is important to state that ethion is a non-authorized compound for Codex Alimentarius, so these results enhance the importance of this method to establish correct withdrawal times. Fluazuron was observed in 10% of the samples, in all cases at concentrations below the MRL ($7000 \mu\text{g kg}^{-1}$).

All results were corrected according to the lipidic contents of the samples as reported by ref. 13. This correction is important to compare the results with classic methodologies which use melting as sample pre-treatment, to eliminate water and thus, expressing the final concentration as 100% fat, with a deviation between 20–30% compared to the proposed method.¹³

Conclusions

A simple multiclass methodology for the simultaneous analysis of pesticides and veterinary drugs was developed and validated. It consists of the extraction of the selected compounds with (MeCN/toluene), and a clean-up with a fast deep freeze-out, PSA, RPC-18, and MgSO_4 . Due to the complexity of the matrix, few methods for the analysis of these compounds are reported, so this work represents a real solution to routine and field analysis of multiclass contaminants in bovine fat. Sample treatment with liquid N_2 is much milder compared to the classical melting pre-treatments, and produces a fine powder easier to handle, improving the extraction efficiency and the recovery of heat sensible compounds. Also, lipid elimination through a fast freeze-out by a cryogenic bath (15 min), shortens the time of analysis and lowers the cost compared to lipid elimination through *n*-hexane washes, turning the whole process into a greener methodology. Finally, the presence of several classes of compounds, with different physicochemical properties, shows the flexibility of the developed method to face its scope enlargement to other polar or semi-polar contaminants.

Data availability

The data supporting the findings of this study are available within the article and its ESI.[†]

Author contributions

Souza and Gérez were responsible for sample preparation and measurement collection. Souza, Gérez and Besil were responsible for the data processing and its interpretation. Pareja, Heinzen and Cesio were responsible for ideation, funding acquisition, experimental design, data visualisation, writing and editing. Souza prepared the first draft of the manuscript and performed all the revisions.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

Authors would like to thank Programa de Desarrollo de las Ciencias Básicas (PEDECIBA). HH, VC, LP and NB thanks Sistema Nacional de Investigadores (SNI)-ANII.

References

- 1 M. LeDoux, *J. Chromatogr. A*, 2011, **1218**(8), 1021.
- 2 European Commission, *Regulation (EC) No.299/2008 of the European Parliament and of the Council of 11 March 2008 amending Regulation (EC) No. 396/2005 on maximum residue levels of pesticides in or on food and feed of Plant and animal origin*, Off J Eur Commun, L9.
- 3 European Commission, *Regulation (EU) No 37/2010 of the Commission of 22 December 2009 on pharmacologically active substances and their classification as regards the maximum residue limits in foodstuffs of animal origin*, **L15**, 1, pp. , pp. 1–72.
- 4 Codex Alimentarius Commission, *Codex Pesticides Residues in Food Online Database*, 2018, <https://www.fao.org/fao-who-codexalimentarius/codex-texts/dbs/pestres/en/>, Accessed 18.02.2024.
- 5 Codex Alimentarius Commission, *Codex Veterinary Drug Residue in Food*, 38th Session, 2015, <https://www.codexalimentarius.org/standards/vetdrugs/veterinary-drugs/en/>, Accessed 19.02.2019.
- 6 A. Valdes, A. Beltrán, C. Mellinas, A. Jiménez and M. C. Garrigós, *Trends Food Sci. Technol.*, 2018, **77**, 120.
- 7 A. D. Gil, S. M. Huertas and O. Rampoldi, <https://xdoc.mx/documents/composicin-de-la-grasa-de-cobertura-en-bovinos-5ef2669c6c8a1>, Accessed 10.08.2024.
- 8 M. Hostens, V. Fievez, J. L. M. R. Leroy, J. Van Ranst, B. Vlaeminck and G. Opsomer, *J. Dairy Sci.*, 2012, **95**, 3756.
- 9 S. J. Lehotay and K. Mastovska, *J. AOAC Int.*, 2005, **88**, 630.
- 10 Department of human health and human services centers for disease control and prevention USA (CDC), *Fourth National Report on Human Exposure to Environmental Chemicals*, 2009.
- 11 K. Lichtmannegger, R. Fischer, F. X. Steemann, H. Unterluggauer and S. Masselter, *Anal. Bioanal. Chem.*, 2015, **407**(13), 3727.
- 12 C. J. Akre and J. D. MacNeil, *J. AOAC Int.*, 2006, **89**(5), 1425.
- 13 N. C. Saratelli, A. N. de Macedo, J. P. de Sousa, A. R. de Araujo Nogueira and S. H. Govoni Brondi, *Anal. Bioanal. Chem.*, 2015, **407**, 3727.
- 14 A. Y. Hmadamin and K. I. Hassan, *Saudi J. Biol. Sci.*, 2020, **27**(3), 887.
- 15 H. Unterluggauer, *FATchers – a novel, simple and fast sample preparation in food of animal origin*, 14th *European Pesticide Residue Workshop*, 2022.
- 16 M. Castillo, C. González and A. Miralles, *Anal. Bioanal. Chem.*, 2011, **400**, 1315.

- 17 O. Hoyos, E. Duvan, O. Cuartas, A. Yudy, M. Peñuela and A. Gustavo, *Food Chem.*, 2018, **221**, 891.
- 18 University of Hertfordshire, *Pesticide properties database*, <https://sitem.herts.ac.uk/aeru/ppdb/>.
- 19 University of Hertfordshire, *Veterinary substances database*, <https://sitem.herts.ac.uk/aeru/vsdb/index.htm>.
- 20 Shimadzu Automatic Adjustment of Retention Time Function (AART), <https://www.shimadzu.eu/automatic-adjustment-retention-time-function-aart/>. Accessed 10.08.2022.
- 21 M. Anastassiades, S. J. Lehotay, D. Stajnbaher and F. Schenck, *J. AOAC Int.*, 2003, **86**, 412.
- 22 H. G. J. Mol, A. Rooseboom, R. Van Dam, M. Roding, K. Arondeus and S. Sunarto, *Anal. Bioanal. Chem.*, 2007, **389**, 1715.
- 23 J. Folch, M. Lees and G. H. Sloane-Standley, *J. Biol. Chem.*, 1957, **226**, 497.
- 24 European Commission DG-SANTE, *Method validation and quality control procedures for pesticide residue analysis in food and feed*, Document No. SANTE/12682/2019, 2021, https://ec.europa.eu/food/plant/pesticides/guidance_documents/docs/qualcontrol_en.pdf, Accessed 20 September 2023.
- 25 B. Kmellar, P. Fodor, L. Pareja, C. Ferrer, M. A. Martínez-Uroz, A. Valverde and A. R. Fernández-Alba, *J. Chromatogr. A*, 2008, **1215**, 37.
- 26 Y.-M. Lee, K.-S. Kim, D. R. Jr Jacobs and D.-H. Lee, *Obes. Rev.*, 2017, **18**, 129.
- 27 A. Pérez, G. R. González, J. González and H. Heinzen, *J. AOAC Int.*, 2010, **93**(2), 712.
- 28 Ministerio de Ganadería Agricultura y Pesca, MGAP, *Manual del Programa Nacional de Residuos Biológicos*, 2009, <https://www.mgap.gub.uy/portal/page.aspx?2,dgsg,dgsg-programa-nacional-decontrol-de-residuos-de-medicamentos-veterinarios-y-contaminantes-en-alimentos-de-origen-animal,O,es,0/>.
- 29 S. Niell, V. Cesio, J. Hepperle, D. Doerk, L. Kirsch, D. Kolberg, E. Scherbaum, M. Anastassiades and H. Heinzen, *J. Agric. Food Chem.*, 2014, **62**(17), 3675.