

Simultaneous Quantification of Multi-Class Antimicrobials in Chicken Kidney and Liver by New Validated UPLC-MS/MS Method

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DOI: 10.52794/hujpharm.1599484

ABSTRACT

A novel and significant method was developed and validated with a sensitive, rapid, and simultaneous analytical method to determine antimicrobials in chicken tissues such as the kidney and liver. The process involved a unique approach to precipitation extraction. This method has not been widely used in this context, followed by the evaporation of the supernatant and reconstitution with the mobile phase. Antimicrobials, including Azithromycin, Clarithromycin, Erythromycin, Clavulanic acid, Ciprofloxacin, Clofazimine, Fluconazole, Linezolid, and Moxifloxacin were meticulously considered for development and validation in the chicken tissues. These antimicrobials were chosen based on their everyday use in poultry farming and their potential impact on human health. We used Ultra-Performance Liquid Chromatography with triple quad Mass Spectrometry and employed multiple reaction monitoring to detect the analytes of interest. All the compounds were well separated using Atlantis T3, 4.6x50mm, 3 µm. The linear range was set between 25 to 1000 ng/gm. The method was validated following linearity, extraction recovery, matrix effect impact, limit of detection, sensitivity, autosampler and benchtop stability, ensuring the results' reliability and our method's robustness.

Keywords: Antimicrobials, Antimicrobial resistance, Chicken tissues, Method development, Validation.

Received Date: 16.12.2024

Accepted Date: 21.02.2025

1. Introduction

Antimicrobials play a pivotal role in modern agriculture, safeguarding the health and well-being of livestock populations while ensuring food safety for consumers. Their judicious use is paramount, as their indiscriminate application can lead to the emergence of antimicrobial-resistant strains, posing a significant threat to both animal and human health [1]. Consequently, precise and consistent analytical methods for quantifying antimicrobial residues in animal tissues are imperative.

The development of antimicrobial resistance via poultry animals is crucial as the consumption of poultry meat has increased predominantly in recent years. One or more antibiotic residues are frequently found in the meat samples. This is an alarming situation to regulate the usage of antibiotics in poultry farms [2,3]. The primary cause of using antibiotics in poultry farms is to save the birds from illness and promote their growth. The most frequently used antibiotics to treat infections that spread across animals are Aminoglycosides, Bambermycin, Beta-lactams, Macrolides, Quinolones, and Sulfonamides. These antibiotics develop resistance among farm animals and humans who consume them [4]. Antimicrobials in poultry must be regulated to fight and eradicate antimicrobial resistance. As an initiative, the Australian government has released a guideline on prescribing antimicrobials for poultry that describes the appropriate usage and best practices in handling antimicrobials [5].

A country-wide residue monitoring program is in place to monitor the residue levels in the meat, which guides the safe usage of antimicrobials, random monitoring of the medicines used in the animals, and appropriate sampling and analysis of the same [6-8]. Post-administration, the drugs are absorbed into the intestine and transported to other tissue parts via blood. This will increase the risk of deposition of residues in the tissues, such as the liver, kidney, spleen, and muscle [9]. To safeguard the consumers, regulatory agencies perform safety evaluations to determine the antibiotic concentrations in the edible tissues. This will ensure the consumers are not exposed to the residual compounds [10].

The analytical methodologies that are being followed currently and in the past decade to determine the antibiotics either qualitatively or quantitatively suggest various technologies that include micro-

biological assay, immunoassay, physical and chemical assays, and biosensors [11]. Different detection methods, such as UV, PDA, GC, and MS, are used based on sensitivity requirements. The extraction of antibiotics from the meat is widely achieved by the SPE technique, which includes the sample cleanup procedure, and the resultant sample is analysed using LC-MS/MS [12]. pH of the sample plays a vital role in the analysis as the compounds are more stable in acidic conditions. Unlike blood and plasma sample preparation, tissue sample processing requires a buffer medium for homogenising the samples. Extracting the residues from the animal tissues is quite challenging as an appropriate sample pre-treatment procedure must be followed to avoid the loss of recovery. Also, multiple steps of the extraction process are involved, such as homogenising the tissue using the suitable buffer, incubation, processing steps, etc., to keep the sample intact. The sensitivity may be affected by various other parameters like mobile phase composition and pH, column chemistry, flow rate, signal enhancers, matrix interferences, and sample compatibility [13-15].

Many studies have reported the development and validation of antibiotics in chicken muscles according to European Commission Decision 2002/657/EC. European Commission and Codex Alimentarius pointed out the MRLs of frequently used antibiotics in animal tissues [16-17]. Extensive research has been done in various chicken tissues to detect and determine antibiotics. However, there was not much focus on the other two antimicrobials, i.e., antivirals and antifungals, as these classes of compounds also trigger resistance [18-23]. There is a need to determine all three types of antimicrobials in a simultaneous method. The antimicrobial residues reach animal tissues via feed and environmental resources such as surface water, soil, etc., as the environmental resources are contaminated with all these classifications of antimicrobials [24-26]. Hence, developing a separate method to detect the antimicrobials will not solve the purpose.

In this context, we present the development and validation of a cutting-edge UPLC-MS/MS method tailored to simultaneously determine antimicrobial compounds within the complex matrix of chicken kidney and liver tissues. This method is a significant advancement in the field, offering high sensitivity, selectivity, and efficiency in quantifying a diverse range of antimicrobials. In the best of the search, no

methods are available to determine antibiotics and antifungals in chicken tissues simultaneously using LC-MS/MS. The intended method can be applied to determine various frequently used antimicrobials in the chicken liver and kidneys with a minimal cost compared with other sophisticated processing methodologies.

Given their physiological significance in drug metabolism and excretion, including kidney and liver tissues in our analytical approach is essential. Ensuring accurate quantification in these matrices is critical for comprehensive residue monitoring and regulatory compliance. Moreover, this study addresses a crucial gap in the analytical methodologies, particularly in simultaneous determination. The ability to assess multiple antimicrobials concurrently provides a comprehensive snapshot of residue levels, reflecting real-world scenarios of complex drug regimens and potential interactions. This research contributes to the analytical toolkit for veterinary drug residue analysis by establishing a robust and validated UPLC-MS/MS method. It holds implications for food safety, public health, and regulatory enforcement. The potential applications of this method are far-reaching, encompassing routine surveillance of antimicrobial residues and investigative studies on drug pharmacokinetics and pharmacodynamics in avian species.

As per the Codex Alimentarius and the National Pharmaceutical Regulatory Agency, the maximum residue levels of some key chemical classifications in poultry products are macrolides with 600-800 µg/kg, quinolones with 30-80 µg/kg, azole derivatives with 50 µg/kg, antibacterials with 100 µg/kg, etc.,

In the subsequent sections, we elaborate on the methodology employed in developing and validating this analytical approach, present our study's key findings, and discuss their broader implications for animal husbandry and food safety.

2. Material and Methods

2.1. Chemicals and reagents:

Methanol and acetonitrile were procured from Merck (Germany), and water, an LC-MS grade, was purchased from the RCI labscan limited. Formic acid, an analytical grade, was procured from Sigma Aldrich. Ammonium acetate was procured from Merck.

Centrifuge tubes for processing were from Abdos (Eppendorf, Germany) and the weighing balances were purchased from Mettler Toledo (Ohio, USA). The tube vortexer was purchased from Remi (Maharashtra, India). Atlantis T3, 4.6x50mm, 3 µm was procured from Waters Inc. Other solutions and reagents were equivalent to analytical grade or higher than that and were purchased from renowned suppliers. UPLC and Xevo TQD were from Waters Corporation, USA. Naïve chicken tissues were procured from a farm grown in an antibiotic-free environment. Phosphate-buffered saline tablets were purchased from Sigma Aldrich, USA. The Incubator was procured from Thermo. The homogeniser/blender was procured from Pro Scientific Inc., USA. Reference standards were procured from Clearsynth, Sigma Aldrich, TCI chemicals, and BLD pharm, India. The reference standards are Azithromycin (AZI), Clarithromycin (CLAR), Clavulanic acid (CLAV), Ciprofloxacin (CIPRO), Clofazimine (CLOF), Erythromycin (ETH), Fluconazole (FLU), Linezolid (LIN), Moxifloxacin (MOXI), and Sulphaphenazole (Internal Standard).

2.2. Preparation of stock and working solutions of standards and quality control samples

Two individual reference standards were weighed for standards and quality control samples and dissolved using dimethyl sulfoxide to achieve a stock concentration of 1 mg/mL. The intermediate stock solutions for standards and quality controls were made using acetonitrile/water (80: 20 ratio) as a diluent. The final spiked concentration of standards ranged from 25, 50, 100, 200, 250, 500, 750, and 1000 ng/gm and three levels of QCs with working concentrations of 45, 475, and 775 ng/gm, respectively, in the chicken tissues.

Sulphaphenazole was used as an internal standard in the assay. The compound was weighed and dissolved using dimethyl sulfoxide for a final 1mg/mL concentration. The final working solution concentration for ISTD was 500 ng/mL, prepared by diluting 100µL of the stock into a 200 mL container containing acetonitrile with 0.1% formic acid. The prepared stock and working solutions were kept in a refrigerator at 2-8°C until the subsequent use.

2.3. LC-MS/MS conditions

The Waters UPLC system was equipped with a pump, column oven, degasser, and autosampler. It is connect-

ed with a Xevo triple quad mass spectrometer. Positive electrospray ionization was used as a primary ionization technique to detect all the compounds. Multiple reaction monitoring mode was used to detect the ions at a unit resolution. The chromatographic separations were achieved using a C18 column, i.e., Atlantis T3, 4.6x50mm, 3 μ m. Mobile phase A consists of ammonium acetate buffer with 0.1% formic acid, whereas mobile phase B consists of acetonitrile with 0.1% formic acid with a flow rate of 0.250 mL/min. A gradient flow was followed to achieve the desired compound separation and elution. Setting the column oven temperature to 40°C and the autosampler temperature to 15°C leads to an appropriate peak shape. The column was loaded with a 5 μ L injection volume, ensuring the volume was sufficient to achieve the desired peak response. All the analytes were eluted within a 4-minute run time. Mass parameters such as gas, temperature, collision energy, cone voltage, etc., were determined and optimised using an auto-tune mode. However, the tune parameters were set manually. Mass Lynx software was used to perform the data acquisition and regression. The mass and mass spectrometer parameter details are given in Table 1.

2.4. Sample preparation and extraction procedure

One phosphate buffer tablet was dissolved in a specified quantity of deionised water. The saline solu-

tion's pH needed to be maintained between 7.2 and 7.4 to prevent any possible interactions with tissues. A 1 gm sample of chicken tissues was weighed and placed in an individual tube. Subsequently, 5 mL of saline solution was added, and the entire content was homogenised using a hand blender. The homogenised tissues were spiked with intermediate standards and quality controls to achieve the desired concentration. The spiked samples were vortexed to ensure thorough mixing. They were incubated in tubes in a controlled environment at 37 °C for 30 minutes, allowing the analytes to bind effectively with the chicken tissues.

Following incubation, 500 μ L of the spiked samples were aliquoted into another tube. 2.5 mL of the internal standard working solution was transferred to precipitate the sample. The mixture was vortexed for 5 minutes. Subsequently, the samples were centrifuged at 13000 rpm for 10 minutes at 4 °C. The supernatant was aliquoted into another tube and dried by evaporation under nitrogen pressure at 40 °C. The dried samples were resuspended using a reconstitution solution containing acetonitrile: water mixture with 0.1% formic acid and loaded into autosampler vials for analysis. A zero-concentration sample was also processed to identify and assess potential analyte contamination. Figure 1 shows a schematic extraction technique.

Table 1. Summary of the compound parameters

S. no.	Antimicrobials	MW	Chemical formula	Parent ion/daughter ion (m/z)	MS Collision Energy (eV)	MS Cone Voltage (V)
1	AZI	749.00	C ₃₈ H ₇₂ N ₂ O ₁₂	749.7/158.1	30	44
2	CLAR	747.95	C ₃₈ H ₆₉ NO ₁₃	748.7/83.0	45	32
3	CLAV	199.16	C ₈ H ₉ NO ₅	200.0/76.9	32	86
4	CIPRO	331.34	C ₁₇ H ₁₈ FN ₃ O ₃	332.1/288.2	18	38
5	CLOF	473.40	C ₂₇ H ₂₂ C ₁₂ N ₄	473.2/283.4	72	64
6	ETH	733.93	C ₃₇ H ₆₇ NO ₁₃	734.7/158.1	22	32
7	FLU	306.27	C ₁₃ H ₁₂ F ₂ N ₆ O	307./238.1	16	28
8	LIN	337.35	C ₁₆ H ₂₀ FN ₃ O ₄	338.2/296.2	18	36
9	MOXI	401.43	C ₂₁ H ₂₄ FN ₃ O ₄	402.3/110.0	22	44
10	Sulphaphenazole	314.36	C ₁₅ H ₁₄ N ₄ O ₂ S	315.0/91.9	38	32

3. Results and Discussion

3.1. UPLC-MS/MS method optimization

3.1.1. Optimization of mass spectrometry

A 20 µg/mL tuning solution containing all the analytes, including ISTD, was prepared using a mixture of 80% acetonitrile in water with 0.1% formic acid as a diluent. Formic acid was added to increase the mass signal during the tuning process. The parent and the daughter ions were determined through the infusion process for ESI+ mode. The positive mode was selected as all the compounds exhibit greater signal strength in this mode of ionization to determine the analytes in the chicken tissues quantitatively. The predominant daughter ions were selected to quantify the analytes. The parent and daughter ions were given in Table 1. All the compounds exhibit various collision energy and cone voltage based on the individual compounds' response. However, the gas parameters, such as cone and desolvation gas, and

temperature parameters, such as source and desolvation temperature, remain the same. The various optimized instrument parameters were selected based on the molecules' intensity. The optimum parameters show no ion enhancement and or suppression. The representative chromatograms are given in Figure 2.

3.1.2. Chromatography

A mobile phase with a gradient flow was selected for the elution of the compounds. The gradient is as follows- mobile phase A: 90% till 0.8 minutes and switch to 10% from 0.8 to 3 minutes and back to 90% from 3 to 4 minutes. More aqueous in the initial run time elutes all matrix impurities, and switching the run to more organic separates the non-polar compounds. Individual samples were initially eluted to ensure the respective retention time. After that, a mix of standard solutions was prepared for simultaneous elution. Since pH plays a significant role, the intermediate stock dilutions, mobile phase, and the

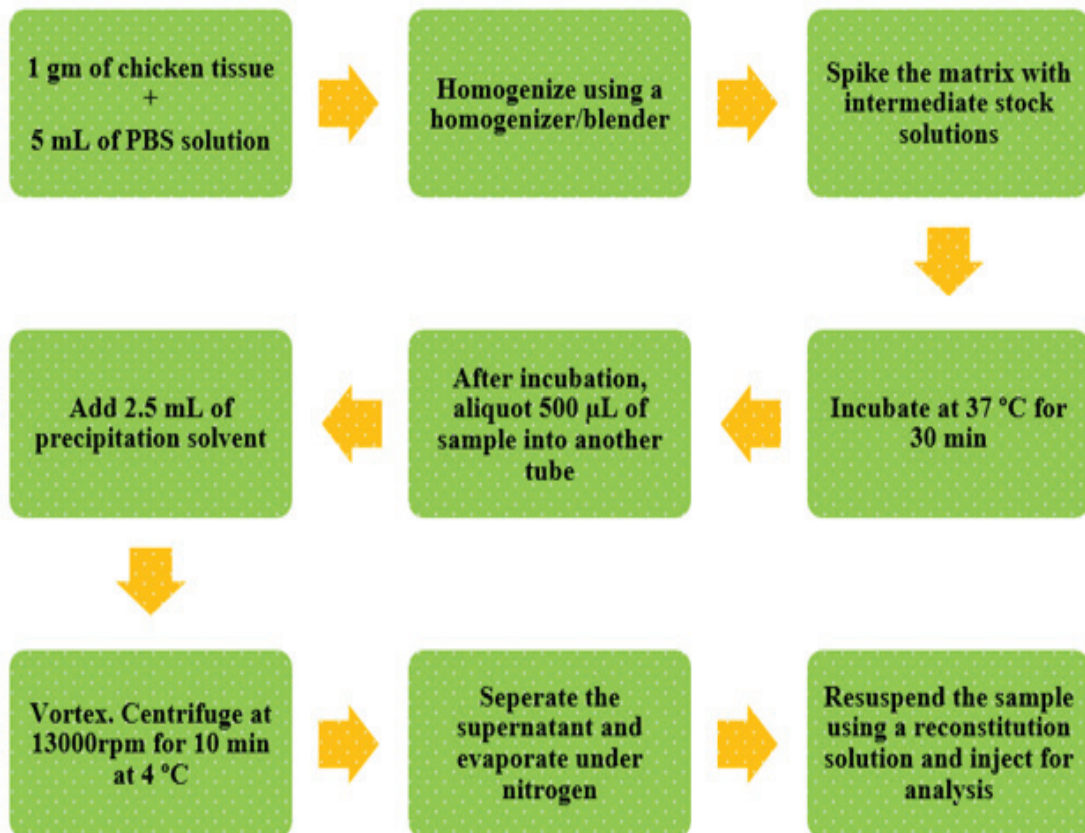


Figure 1. Schematic diagram of the sample preparation and extraction procedure

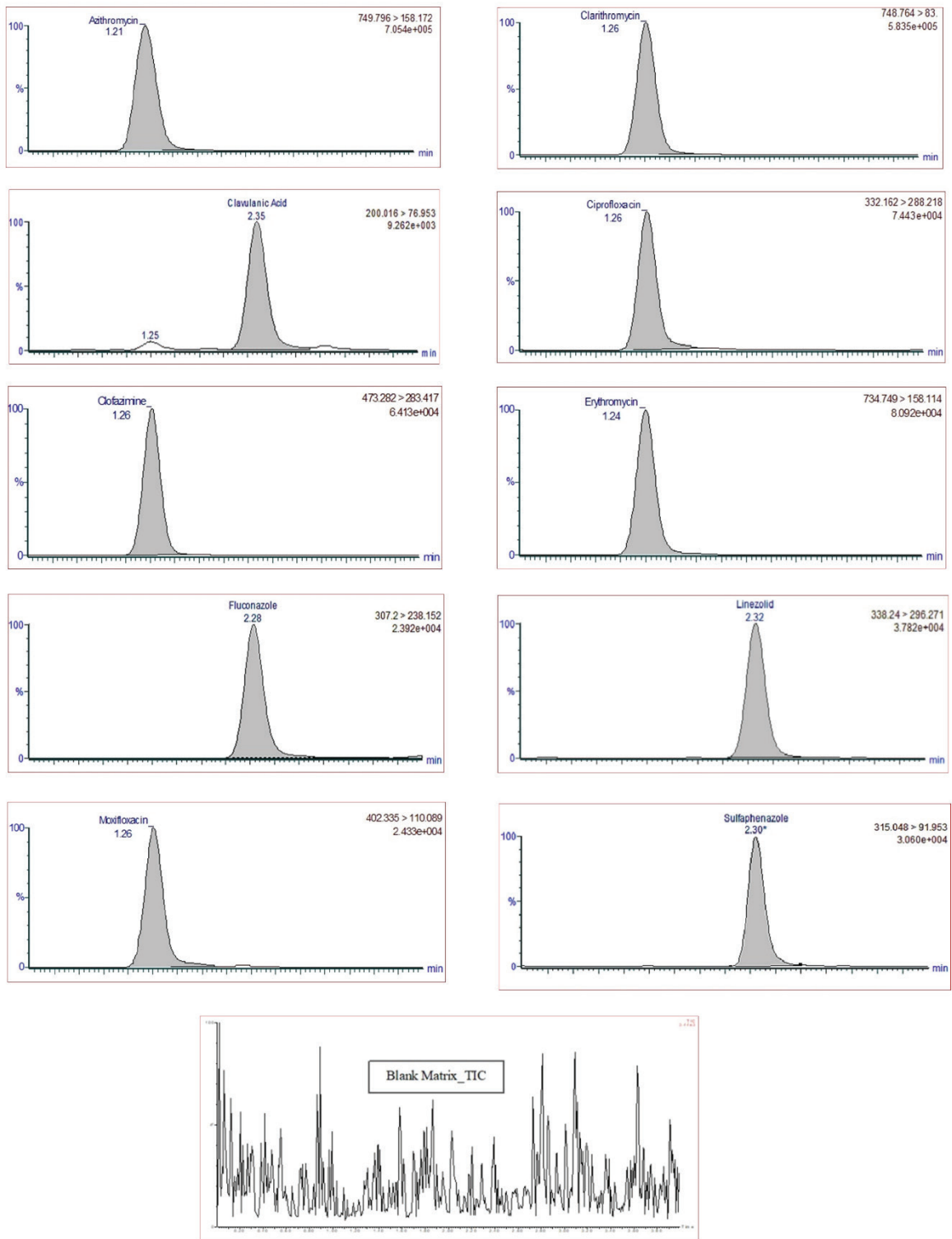


Figure 2. Representative chromatograms for antimicrobials

internal standard working solution were acidified with 0.1% formic acid to make the sample compatible for an appropriate ionization. The proposed for-

mic acid strength gives an enhanced signal intensity compared with 0.05% and 0.2% formic acid, as these two formic acid strengths suppressed the ionization.

A non-polar column was selected to ease the compounds' elution as the C18 column exhibits an appropriate separation. The 0.250 mL/min flow rate enhances the sensitivity of compounds compared with 0.3 and 0.4 mL/min. A solid organic wash containing 10% water in acetonitrile with 0.1% formic acid and a weak organic wash containing 90% water in acetonitrile with 0.1% formic acid ensures no carryover. Each compound was well separated when the column temperature was set at 40 °C compared with various other temperatures at 25 °C and 30 °C, as peak tailing was observed. Using deuterated internal standards will be expensive for individual compounds. Hence, Sulphaphenazole was used, which was compatible with the analytical method and had little to no variation.

3.2. Method validation

With the proposed sample processing method and instrument conditions, method validation was performed with parameters such as linearity, matrix factor, recovery, sensitivity, limit of detection, autosampler stability, and bench-top stability.

3.2.1. Calibration curve

Three precision and accuracy batches were performed individually for the chicken tissues on two

consecutive days. The r^2 is the square of the correlation coefficient and provides information about how close the calculated points are to the line. The R-value, which is near to 1, is desirable. All the antimicrobials exhibit r^2 values between 0.9900 and 0.9925 in chicken kidney and liver tissues. The results are provided in Table 2. The calibration curve for both tissues is given in Figure 3.

3.2.2. Precision and accuracy, carry-over impact

The carry-over impact was determined by injecting a blank extracted sample before and after the lower and higher concentrations. No significant peak response was detected in the blank extracted samples, which confirms no potential carry-over for the proposed method. To establish the precision and accuracy of the extraction method in both tissues, five replicates of three concentrations, i.e., LQC, MQC, and HQC, were injected, followed by a standard curve. The precision and accuracy of the standard curve were observed at 0.7 to 12.2% and 87.2 to 108.6%, respectively, for chicken kidney tissue. The same for chicken liver tissue was observed at 0.3 to 9.5% and 87.7 to 108.7%, respectively. The precision and accuracy of the three QCs for chicken kidneys were observed at 2.7 to 9.9% and 90.7 to 108.6%. The same for chicken liver was observed at 3.9 to 10.7% and 91.1 to 108.7%, respectively. The results met the

Table 2. Summarised standard curve correlation coefficient (r^2)

S. no.	Drug	CC range (ng/gm)	Average r^2	
			Kidney	Liver
1	AZI	25 - 1000	0.9910	0.9925
2	CLAR	25 - 1000	0.9920	0.9913
3	CLAV	25 - 1000	0.9930	0.9913
4	CIPRO	25 - 1000	0.9900	0.9913
5	CLOF	25 - 1000	0.9900	0.9907
6	ETH	25 - 1000	0.9900	0.9907
7	FLU	25 - 1000	0.9910	0.9907
8	LIN	25 - 1000	0.9923	0.9923
9	MOXI	25 - 1000	0.9917	0.9910

^aAverage r^2 was derived from at least three precision and accuracy batches.

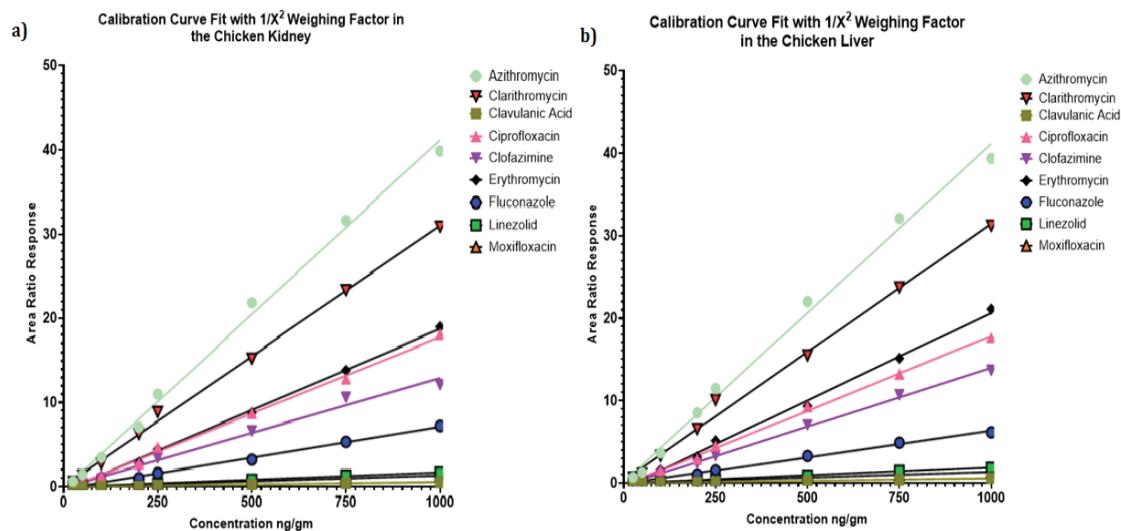


Figure 3. Calibration curve of antimicrobials in a) chicken kidney tissue, b) chicken liver tissue

acceptance criteria with $\pm 15\%$ accuracy and $\leq 15\%$ precision. Summarised results are shown in Table 3.

3.2.3. Recovery from matrix, limit of detection, and matrix effect

Recovery, i.e., extraction efficiency and matrix effect, was determined at LQC and HQC levels in the respective chicken tissues. While the recovery experiment was performed to confirm the extraction efficiency, the impact of the matrix effect was established to verify that no suppression and enhancement of the ions were detected in the intended method. In chicken kidney tissues, the recovery ranged between 85.16 and 102.14 %, and the matrix effect was observed from 84.15 to 119.64 %, respectively. Meanwhile, in chicken liver, the recovery was between 59.18 and 103.17 %, and the matrix effect was between 76.53 and 109.50 %, respectively. The recovery of Ciprofloxacin, fluconazole, and moxifloxacin in chicken liver tissues was reduced by less than 70%. This may be due to the irreversible binding nature of compounds in particular tissues.

The signal-to-ratio produced by the mass spectrometer was used to determine the limit of detection. The limit of detection is the lower limit of concentration, making the S/N ratio at three values. The results of the matrix effect, recovery, and limit of detection are given in Table 4 and Table 5. The average recovery from matrix and matrix effect of antimicrobials in various chicken tissues are presented in Figure 4 a, b, c, and d.

3.2.4. Sensitivity

Samples with a concentration of lower limit quantitation were determined under the calibration standards. The average accuracy for the lower limit of quantitation for all the antimicrobials was observed between 80 and 120% of the actual concentration, as shown in Table 6.

3.2.5. Autosampler and benchtop stability

The autosampler and benchtop stability of the antimicrobials extracted from the respective chicken tissues were established for at least 23 hrs 14 min and 7 hrs 18 min, respectively. For the autosampler stability, the autosampler temperature was maintained at 15°C. The spiked samples were stored on a benchtop at a controlled room temperature for benchtop stability. Both stability experiments were established at low and high QC samples against freshly prepared calibration curve standards and quality control samples. The resulting bias is given in Table 7 and Table 8.

4. Conclusion

Antibiotics in chicken tissues significantly influence the development of antimicrobial resistance, underscoring the need for a sensitive and comprehensive analytical method to detect multiple antibiotics. This study introduces a simple, accurate, sensitive, precise, and reproducible LC-MS/MS method developed and validated to detect nine different antibiotics

Table 3. Summarised precision and accuracy

S. No.	Compounds	Chicken Kidney						Chicken Liver					
		Precision (%RSD) ^a			% Accuracy ^b			Precision (%RSD) ^a			% Accuracy ^b		
		LQC	MQC	HQC	LQC	MQC	HQC	LQC	MQC	HQC	LQC	MQC	HQC
1	AZI	5.6	2.7	3.8	103.0	108.5	92.9	9.2	5.1	6.4	99.6	108.0	93.6
2	CLAR	8.5	4.6	4.7	100.9	102.7	90.7	8.6	6.0	4.4	101.2	105.2	91.1
3	CLAV	5.9	5.4	6.0	107.5	100.0	103.4	8.4	3.9	10.1	102.2	102.7	98.0
4	CIPRO	9.3	5.0	6.3	94.4	105.5	103.9	5.0	4.4	8.5	92.9	108.8	96.3
5	CLO	9.1	3.9	4.9	96.3	108.6	106.2	10.7	4.1	4.7	97.6	108.2	97.7
6	ETH	9.9	4.5	4.0	98.2	107.9	107.6	10.0	5.2	8.3	95.4	104.0	100.9
7	FLU	6.8	5.1	5.3	101.7	105.9	106.5	8.2	5.0	6.3	96.2	104.2	104.3
8	LIN	8.8	8.1	6.6	103.3	103.5	104.5	9.9	6.4	5.8	100.2	103.3	102.0
9	MOXI	9.6	6.4	7.8	97.6	99.0	100.8	10.1	4.9	8.7	95.6	100.4	98.4

^a%RSD was calculated from precision and accuracy batches. ^bAccuracy was determined from the nominal concentration of the QC concentrations, respectively.

Table 4. Summarised extraction recovery and matrix effect

S. No.	Antimicrobials	% Recovery in Chicken Kidney ^a		% Recovery in Chicken Liver ^b		% Matrix Effect in Chicken Kidney ^a		% Matrix Effect in Chicken Liver ^b	
		LQC	HQC	LQC	HQC	LQC	HQC	LQC	HQC
1	AZI	91.04	102.14	74.90	100.25	90.62	98.96	98.16	96.84
2	CLAR	92.09	104.98	83.34	98.75	96.87	101.37	98.91	98.36
3	CLAV	90.51	97.11	71.46	94.26	108.91	119.64	76.53	98.68
4	CIPRO	87.83	98.54	69.37	96.56	97.53	102.62	90.79	100.52
5	CLO	90.85	98.17	79.93	100.01	90.40	101.86	99.46	102.50
6	ETH	85.16	100.25	88.75	98.68	94.72	100.91	99.18	102.29
7	FLU	96.83	99.16	61.29	103.17	99.56	104.99	95.28	99.84
8	LIN	92.84	101.72	94.50	99.42	84.15	99.60	100.28	101.24
9	MOXI	85.76	99.1	59.18	95.14	90.20	102.68	109.50	103.04

^{a,b} Average of five replicates.

from various classes in chicken tissues, specifically the kidney and liver. The technique demonstrates precision with ≤15% variability and accuracy within

±15%. The protein precipitation technique, followed by evaporation and reconstitution using the mobile phase, effectively showcased the extraction efficien-

Table 5. Summary of the limit of detection in chicken tissues

S.no.	Antimicrobials	LOD (ng/mL)	
		Kidney	Liver
1	AZI	10.80	9.74
2	CLAR	6.60	5.49
3	CLAV	24.19	18.60
4	CIPRO	14.87	14.86
5	CLO	12.89	15.66
6	ETH	9.80	8.51
7	FLU	12.63	6.09
8	LIN	8.05	12.84
9	MOXI	8.35	10.58

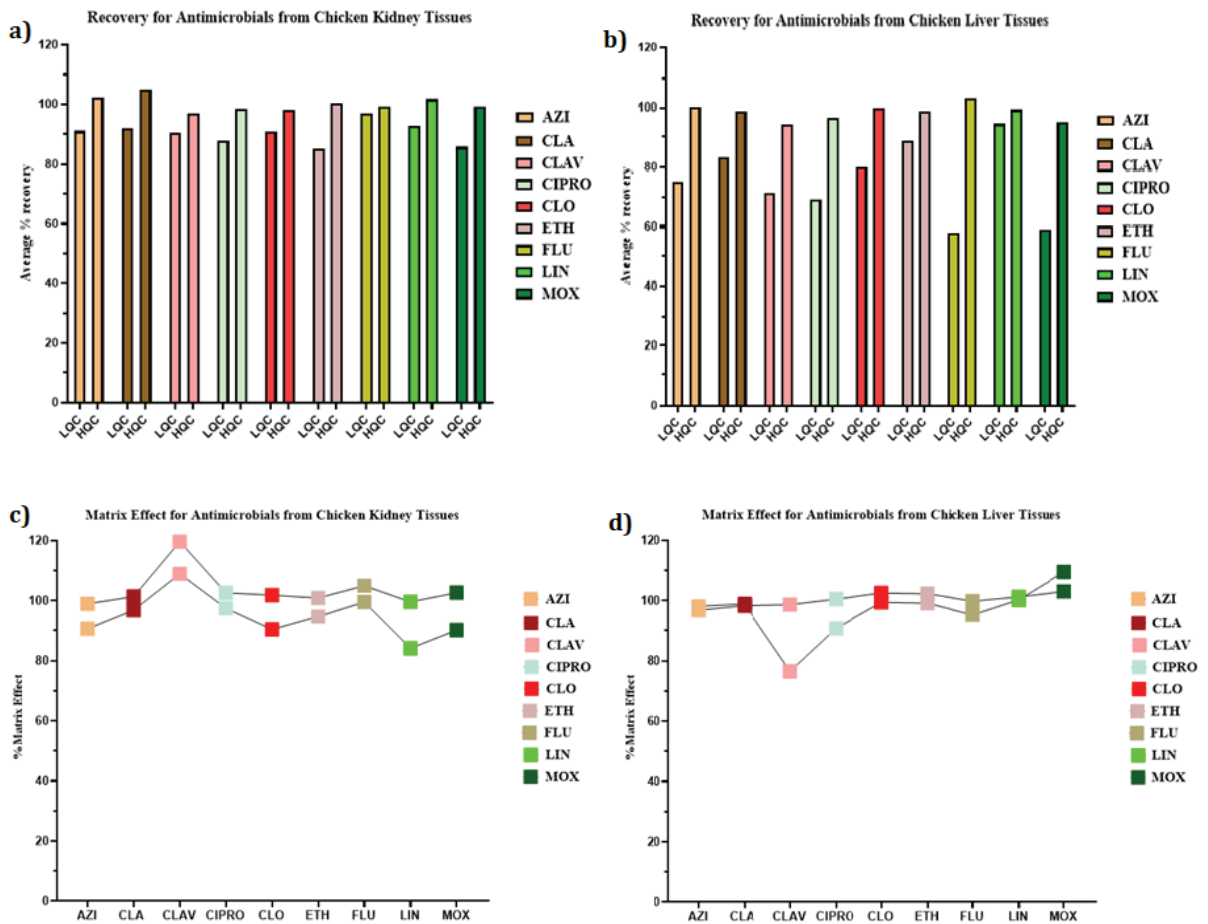


Figure 4. The average recovery of antimicrobials in a) chicken kidney b) chicken liver, Matrix effect of antimicrobials in c) chicken kidney, d) chicken liver.

Table 6. Summarised sensitivity data in chicken tissues

S. no.	Antimicrobials	The average % accuracy	
		Chicken Kidney	Chicken Liver
1	AZI	97.18	108.70
2	CLAR	104.16	96.78
3	CLAV	111.36	102.14
4	CIPRO	112.24	103.48
5	CLO	108.55	104.68
6	ETH	102.03	102.04
7	FLU	108.30	107.20
8	LIN	102.38	106.96
9	MOXI	111.36	104.94

Table 7. Summary of autosampler stability

Antimicrobials	Kidney		Liver	
	^a LQC	^b HQC	^a LQC	^b HQC
AZI	4.52	-1.56	-10.85	7.64
CLAR	0.67	0.31	-10.32	-7.41
CLAV	-10.43	-10.16	10.78	5.79
CIPRO	-5.65	-2.33	-5.09	5.28
CLO	-3.59	-2.9	-1.09	-9.85
ETH	5.02	-11.31	-2.54	-3.53
FLU	-2.19	-9.24	-12.4	-8.08
LIN	-6.28	-7.44	5.00	-1.24
MOXI	-2.60	-9.16	-3.16	-5.87

^{a,b} Average of five QCs.

cy. Both autosampler and benchtop stability tests confirmed the compounds' stability before and after processing. All compounds were eluted within the specified run time, exhibiting minimal matrix effect. The validated parameters confirm that this method is reliable for detecting antimicrobials in chicken tissues such as kidneys and liver.

Acknowledgments

I humbly thank the Foundation for Neglected Disease Research team for their support in performing this scientific activity. I also thank the Department of Pharmaceutical Sciences, Vignan's Foundation for Science Technology and Research (Deemed to be University), Guntur, for guiding my research.

Table 8. Summary of benchtop stability

Antimicrobials	Kidney		Liver	
	^a LQC	^b HQC	^a LQC	^b HQC
AZI	-5.57	1.28	-8.99	-7.59
CLAR	-5.29	-5.20	-1.4	-8.35
CLAV	0.68	-3.84	-8.88	-2.02
CIPRO	-10.92	-4.61	-2.64	0.45
CLO	-3.79	-7.19	-10.28	-4.12
ETH	-3.42	-7.37	1.86	-4.84
FLU	-12.19	-11.29	-0.26	-8.16
LIN	-5.47	-3.42	-3.60	0.66
MOXI	-9.27	0.20	-2.75	-1.8

^{a,b} Average of five QCs.

Conflicts of Interest

The authors declare no conflicts of interest, financial or otherwise, to declare.

Statement of Contribution of Researchers

Data Collection, Analysis, Interpretation, Literature Search, Draft preparation: S.S.S. Concept, Supervision, Review and Revision: S.K.K., S.N., R.K.S.

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