

# LC-MS/MS Method for Determination of **Colistin in Human Plasma: Validation** and Stability Studies

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# Abstract

A simple and reliable high performance liquid chromatography tandem mass spectrometry (LC-MS/MS) assay for the determination of colistin A and colistin B in human plasma was developed and validated. Clarithromycin was used as an internal standard (IS). Plasma extraction was performed using C-18 cartridges and methanol containing 0.1% formic acid. Analysis was performed using Atlantis dC18 (2.1 × 100 mm, 3 µm) column at room temperature and a mobile phase of 0.2% formic acid in acetonitrile and water (50:50, v:v), delivered at a flow rate of 0.2 ml/minute. Eluent was detected in the positive ion mode using electrospray ionization at the following transitions of mass to charge (m/z): colistin A, 585.6  $\rightarrow$  101.4; colistin B, 578.7  $\rightarrow$ 101.3; and IS, 748.6  $\rightarrow$  158.4. No interference by components of blank plasma or commonly used drugs was observed. The relationship between colistin A colistin B concentrations and their corresponding peak height ratios to the IS was linear over the range of  $0.05 - 10 \,\mu\text{g/ml}$ . Inter-day coefficient of variation and bias were, respectively, ≤11.5% and −3.0 to 6.0 for colistin A and  $\leq$  9.9 and -4.7 to 3.0 for colistin B. Mean extraction recovery of colistin A, colistin B, and the IS were 97%, 94%, and 97%, respectively. The method was applied to assess the stability of colistin A and colistin B in processed samples (24 hr. at room temperature, 48 hours at -20°C) and unprocessed samples (24 hr. at room temperature, 8 weeks at -20°C) and after three cycles of freeze and thaw found to be  $\geq 87\%$ .

# **Keywords**

Colistin, Clarithromycin, Human Plasma, LC-MS/MS

1

## **1. Introduction**

Colistin A and colistin B are part of polmyxcin E, together with other polypeptides. Commonly, they constitute more than 85% of polymyxcin E by weight. They have similar chemical structure with colistin A having one extra methylene moiety [1]. Colistin is one of the oldest antibiotics but is used as last-line treatment for infections caused by multi-drug-resistant gram-negative bacteria [2] [3]. It is commercially available as colistin methanesulfonate for intravenous administration and as colistin sulfate for oral use. Pharmacokinetic of colistin is very variable and it has narrow therapeutic window [4], necessitating careful drug monitoring.

Several analytical methods are reported for measurement of colistin in various biological matrices, including human plasma and urine [5]-[17]. These include thin layer chromatography [6], isotachophoresis [7], capillary electrophoresis [8], and high performance liquid chromatography with ultraviolet [9] or fluorescence detection after dervatization with 9-fluorenylmethyl chloroformate [10] [11] [12] [13] [14]. Liquid chromatography coupled with tandem mass spectrometry (LC-MS/MS) is becoming more frequently used because of its higher selectivity and sensitivity [15] [16] [17]. Most LCMS/MS assays use polymyxin B1 as internal standard [15] [17]. However, in one report, colistin B level was over estimated using this internal standard [17].

In this paper, we report a simple, precise, and rapid LCMS/MS assay for measurement of colistin A and colistin B in human plasma using clarithromycin as an internal standard in order to produce reliable measurement of both colistin A and colistin B.

# 2. Materials and Methods

## 2.1. Instrument

Sep-Pak classic C-18 cartridges (Waters Corporation, Milford, MA, USA) was used for analyte extraction from plasma samples. Chromatography was performed on an LCMS/MS system consisting of Water Alliance 2695 separation module (Waters Associates, Inc Milford, MA, USA) for solvent delivery and sample introduction, and a Micromass Quattro micro API bench-top triple quadruple mass spectrometer (Micromass, Manchester, UK) interfaced with a Z-spray ESI source as detector. Analysis was performed using reversed phase Atlantis dC<sub>18</sub> column (2.1 × 100 mm, 3  $\mu$ m) protected by guard column Symmetry C<sub>18</sub> (3.9 × 20 mm, 5  $\mu$ m) at room temperature. Mass Lynx software (Ver 4.0) working under Microsoft Window XP professional environment was used to control the instrument, data acquisition, peak integration, peak smoothing, and signal-tonoise ratio measurement.

### 2.2. Chemical and Reagents

Colistin sulfate and clarithromycin pure standards were purchased from Sigma-Aldrich, MO, USA. Formic acid, methanol, and acetonitrile (HPLC grade) were purchased from Fisher Scientific, NJ. USA. Water HPLC grade was prepared by reverse osmosis and further purified by passing through a Milli-Q, Synergy System (Millipore, Bedford, MA, USA). Drug-free human plasma was obtained from the blood bank of King Faisal Specialist Hospital & Research Centre (KFSHRC) Riyadh, Saudi Arabia.

## 2.3. Chromatographic Conditions

The mobile phase consisted of 0.2% formic acid in acetonitrile and water (50:50, v:v). It was degassed before use and delivered at a flow rate of 0.2 ml/minute at ambient temperature with a run time of 4 minutes. The electrospray ionization source was operated in the positive-ion mode at a capillary voltage of 3.5 kV and a cone voltage of 35 V. Nitrogen was used as nebulizing and desolvation gas at a flow rate of 50 and 600 L/hr, respectively. Argon was used as the collision gas at a pressure of  $1.28 \times 10^{-3}$  mbar. The optimum collision energy for colistin and the IS was 25 eV. The ion source and the desolvation temperatures were maintained at 120°C and 350°C, respectively. Colistin A, colistin B, and the IS were detected in the positive ion using multiple reactions monitoring mode at the following transitions of mass to charge (m/z): 585.6  $\Rightarrow$  101.4, 578.7  $\Rightarrow$  101.3, and 748.6  $\Rightarrow$  158.4, respectively.

# 2.4. Preparation of Standard and Quality Control Samples

Stock solution of colistin (A and B) and IS (1.0 mg/ml) were prepared in methanol. They were further diluted with plasma (colistin A and B) and water (IS) to produce working solutions of 10 µg/ml for colistin (A & B) and 0.1 µg/ml for IS. Ten calibration standards in the range of 0.05 - 10.0 µg/ml and four quality control (QC) samples (0.05, 0.15, 5.0, and 9.0 µg/ml) were prepared in human plasma. Calibration standards and QC samples were vortexed for one minute and 1.0 ml aliquots were transferred into Teflon-lined, screw-capped, borosilicate glass (13 × 100 mm) culture tubes and stored at  $-20^{\circ}$ C until used.

## 2.5. Sample Preparation

Aliquots of 1 ml blank plasma, calibration curve, or QC samples were allowed to equilibrate to room temperature. To each tube, 150  $\mu$ l of the IS 0.1  $\mu$ g/ml solution was added and vortexed for 20 seconds. Before loading samples, Sep-Pak C-18 cartridges were conditioned with 1 ml methanol followed by 2 ml water. Then the sample (1 ml) was loaded followed by 1 ml water. Samples were eluted with 1 ml methanol containing 0.1% formic acid. The eluates were then evaporated to dryness under a gentle nitrogen stream in a heating block at 45°C, the residues were reconstituted in 100  $\mu$ l of 0.1% formic acid in water, transferred into auto-sampler vial, and 10  $\mu$ l were injected into the LC-MS/MS system.

# 2.6. Stability Studies

A total of 40 aliquots of four QC samples (0.05, 0.15, and 9.0  $\mu$ g/ml) were used

for stability studies. Five aliquots of each QC sample were extracted and immediately analyzed (baseline), five aliquots were allowed to stand on the bench-top for 24 hours at room temperature before being processed and analyzed (counter stability, 24 hours at room temperature), five aliquots were stored at  $-20^{\circ}$ C for eight weeks before being processed and analyzed (long term freezer storage stability), and five aliquots were processed and stored at room temperature for 24 hours or 48 hours at  $-20^{\circ}$ C before analysis (autosampler stability). Finally, fifteen aliquots of each QC sample were stored at  $-20^{\circ}$ C for 24 hours, they were then left to thaw completely at room temperature unassisted. Five aliquots of each QC sample were extracted and analyzed and the rest returned to  $-20^{\circ}$ C for another 24 hours. The cycle was repeated three times (freeze-thaw stability).

#### 2.7. Method Validation

The method was validated according to standard procedures described in the US Food and Drug Administration bioanalytical method validation guidance [18]. The validation parameter included specificity, linearity, accuracy, precision, recovery, and stability.

## 3. Results and Discussion

#### 3.1. Optimization of LC and MS/MS Conditions

The product and precursor ions of colistin A, colistin B, and clarithromycin (IS) were determined by infusing a standard mixture (1.0 µg/ml in methanol) in the mass spectrometer using a syringe pump at a flow rate of 20 µl/ml. Mass spectrometry acquisition was performed in multiple reaction monitoring (MRM) with positive ion transition mode set at (precursor  $[M-2H]^{2+}$  to product ion) mass to charge (m/z): 585.6  $\rightarrow$  101.4, 578.7  $\rightarrow$  101.3, 748.6  $\rightarrow$  158.4, for colistin A, colistin B, and the IS, respectively. The optimum peak height response were obtained by applying capillary voltage of 3.5 KV, cone voltage of 35V, and collision energy of 25 eV. The optimal chromatographic condition consisted of 0.2% formic acid in acetonitrile and water (50:50, v:v), delivered at a flow rate of 0.2 ml/minute. Analysis was completed within 4 minutes run time. The retention time of colistin A, colistin B and the IS were around 1.4, 1.4, and 2.8 minutes, respectively. **Figure 1** represents total ion chromatogram (TIC) of colistin A, colistin B and IS extracted from plasma.

#### **3.2. Specificity**

We screened six blank plasma samples and seven commonly used drugs including aspirin, acetaminophen, ranitidine, nicotinic acid, caffeine, diclofenac and omeprazole for interferences with colistin A and colistin B, or IS. No drug or endogenous component co-eluted with colistin A, colistin B or the IS. Figure 2(A) depicts a representative chromatogram of drug free human plasma (blank) used in preparation of standards and quality control samples. Figure 2(B) depicts blank plasma spiked with the IS (0.015 µg).



**Figure 1.** Total ion current chromatogram of standard mixture containing colistin A, colistin B, and clarithromycin (IS) extracted from plasma.



Figure 2. Multiple reaction monitoring chromatogram of blank plasma and blank plasma spiked with IS.

## 3.3. Linearity and Limit of Detection and Quantification

Linearity of the assay was evaluated by analyzing a series of standard mixtures containing colistin A and colistin B in human plasma at nine concentrations over the range of 0.05 - 10.0 µg/ml. Corresponding peak height ratios to the IS and concentrations were subjected to regression analysis. Mean equations obtained were y = 0.0979x - 0.0029,  $r^2 = 0.9901$  (n = 10) and y = 0.0461x - 0.0012,  $r^2 = 0.9921$ , (n = 10) for colistin A and colistin B, respectively. The suitability of the calibration curves was confirmed by back calculating the concentration of colistin A and colistin B in human plasma from calibration curves (**Table 1**). All calculated concentrations were well within the acceptable limits. **Figure 3** represents

 Table 1. Back-calculated colistin A and colistin B concentrations from ten calibration curves.

Nominal Level (µg/ml)	Colistin A		** \	Maan	Colistin B	
	Mean (SD)	*CV (%)	(%)	(SD)	*CV (%)	**Accuracy (%)
0.05	0.05 (0.01)	8.6	107	0.05 (0.01)	7.8	108
0.1	0.10 (0.01)	8.4	102	0.10 (0.01)	4.6	104
0.2	0.19 (0.02)	9.1	100	0.21 (0.02)	8.5	105
0.5	0.51 (0.06)	11.2	100	0.51 (0.04)	7.7	103
1.0	0.98 (0.11)	11.4	99	1.01 (0.09)	8.9	101
2.0	2.02 (0.21)	10.3	101	2.20 (0.12)	6.2	101
4.0	4.10 (0.28)	6.9	102	4.01 (0.12)	3.1	100
6.0	5.90 (0.51)	8.6	98	5.94 (0.44)	7.4	99
8.0	7.68 (0.74)	9.6	96	7.68 (0.61)	8.0	96
10.0	10.27 (0.44)	4.3	103	10.27 (0.42)	4.0	103

\*Coefficient of variation (CV) = standard deviation (SD) divided by mean measured concentration × 100. \*\*Accuracy = measured level divided by nominal level × 100.





mean calibration curve of colistin A and colistin B. Limit of detection of colistin A and colistin B in plasma were 0.030  $\mu$ g/ml and 0.035  $\mu$ g/ml, respectively, whereas, limits of quantification were 0.05  $\mu$ g/ml for both compounds.

#### 3.4. Precision and Accuracy (Bias)

Precision and bias were determined for four QC concentrations (0.05, 0.15, 5.0, and 9.0 µg/ml). Intra-day precision (n = 10) was  $\leq 11.4\%$  for colistin A and  $\leq 11.2\%$  for colistin B. Inter-day precision (n = 20, over three consecutive days) was  $\leq 11.5\%$  for colistin A, and  $\leq 9.9\%$  for colistin B. Inter-day bias was in the range of -9.1% to 10.5% for colistin A and -8.6% to 10.2% for colistin B. Inter-day bias was -3.0% to 6.0% for colistin A, and -4.7% to 3.0% for colistin B. The results are summarized in Table 2. Figure 4 depicts the LC-MS/MS chromatogram of low QC (0.15 µg/ml), and high QC (9 µg/ml) both spiked with IS (0.015 µg).

#### 3.5. Recovery

Extraction recovery of colistin A and colistin B at four concentrations (0.05, 0.15, 5.0 and 9.0  $\mu$ g/ml) and the IS at one concentration (0.1  $\mu$ g/ml) were determined by comparing peak heights of spiked-before-extraction samples and spiked-after-extraction samples (5 sets). Mean measured extraction recovery of colistin A, and colistin B was 97%, and 94%, respectively. Recovery of IS was 97%.

#### 3.6. Matrix Effect

Matrix effect was quantitatively evaluated by comparing peak heights of colistin

Nominal Level – (µg/ml)	Measured Level Colistin A			Measured Level Colistin B			
	Mean (SD)	CV (%)	Bias (%)	Mean (SD)	CV (%)	Bias (%)	
	Intra	-day (n =	10)				
0.05	0.06 (0.01)	2.7	10.5	0.05 (0.01)	11.2	10.2	
0.15	0.14 (0.01)	3.1	-3.6	0.15 (0.02)	7.6	1.2	
5.0	4.51 (0.28)	6.3	-9.1	4.93 (0.25)	5.1	0.2	
9.0	9.09 (1.03)	11.4	-1.8	8.64 (0.45)	5.2	-8.6	
	Inter	-day (n =	20)				
0.05	0.05 (0.01)	11.5	6.0	0.05 (0.01)	9.9	3.0	
0.15	0.14 (0.01)	6.8	0.1	0.15 (0.01)	6.7	-1.0	
5.0	4.87 (0.50)	10.3	-3.0	5.09 (0.37)	7.3	-0.3	
9.0	9.25 (0.82)	8.8	3.6	8.82 (0.64)	7.2	-4.7	

Table 2. Intra- and inter-day precision and bias of colistin A colistin B assay.

SD, standard deviation. CV, standard deviation divided by mean measured concentration  $\times$  100. Bias, measured level – nominal level divided by nominal level  $\times$  100.



**Figure 4.** Multiple reaction monitoring chromatogram of two colistin quality control samples in plasma spiked with the IS (0.015  $\mu$ g).

A and colistin B (0.05, 0.15, 5, and 9  $\mu$ g/ml) and the IS (0.1  $\mu$ g/ml) prepared in human plasma to those prepared in water. Mean ion suppression effect was -4.9% for colistin A, -4.5% for colistin B and -1.0% for the IS.

#### 3.7. Stability

Colistin (A and B) and the IS stability in processed and unprocessed plasma samples at 3 concentrations (0.05, 0.15 and 9.0 µg/ml) was investigated. Colistin was stable in processed samples for at least 24 hours at room temperature ( $\geq$ 98% for colistin A and  $\geq$ 92% for colistin B) and 48 hours at  $-20^{\circ}$ C ( $\geq$ 105% for colistin A and  $\geq$ 91% for colistin B). Colistin in unprocessed plasma samples was stable for at least 24 hours at room temperature ( $\geq$ 87% for colistin A and  $\geq$ 92% for colistin B), at least eight weeks at  $-20^{\circ}$ C ( $\geq$ 90% for colistin A  $\geq$ 94% for colistin B), and after three freeze-and thaw cycles ( $\geq$ 100% colistin A and colistin B. **Table 3** summaries the results of stability studies.

Nominal	Unprocessed		Processed		Freeze-Thaw		
Level	24 hrs	8 wks	24 hrs	48 hrs	Cycle		
(µg/ml)	RT	–20°C	RT	–20°C	1	2	3
Colistin A							
0.05	91	104	99	106	117	118	113
0.15	87	101	98	105	100	103	108
9.0	104	90	106	111	110	109	107
Colistin B							
0.05	96	107	92	100	114	116	113
0.15	92	95	107	107	107	101	106
9.0	97	94	100	91	110	102	104

 Table 3. Stability of colistin A and colistin B under various clinical laboratory conditions.

Stability (%) = mean measured concentration (n = 5) at the indicated time divided by mean measured concentration (n = 5) at baseline × 100. Spiked plasma samples were processed and analyzed immediately (baseline, data not shown), after 24 hours at room temperature (24 hrs RT), after freezing at  $-20^{\circ}$ C for 8 weeks (8 wks,  $-20^{\circ}$ C), or processed and then analyzed after storing for 24 hours at room temperature (24 hrs, RT) or 48 hours at  $-20^{\circ}$ C (48 hrs,  $-20^{\circ}$ C).

# 4. Conclusion

The described LCMS/MS assay is simple, precise, and accurate, making it suitable for therapeutic drug monitoring and pharmacokinetic analysis. It requires 1 ml plasma sample, and the analysis was completed within four minutes. Assay was applied successfully to monitor stability of colistin under various conditions generally encountered in the clinical laboratories.

# **Conflicts of Interest**

The authors declare no conflicts of interest regarding the publication of this paper.

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