

METHOD DEVELOPMENT AND VALIDATION FOR THE ESTIMATION OF AMPRENAVIR IN HUMAN PLASMA BY LIQUID CHROMATOGRAPHY-ELECTROSPRAY IONIZATION-TANDEM MASS SPECTROMETRY

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ABSTRACT

Objective: The main aim of the present study was to develop a sensitive liquid chromatography–electrospray ionization–tandem mass spectrometric technique for the quantitation of amprenavir in human plasma.

Methods: Chromatographic separation was achieved on a reversed-phase Symmetry C₁₈ (50 mm×4.6 mm, 3.5 μm) column with isocratic elution by acetonitrile and 0.1% v/v formic acid in the ratio of 90:10 v/v as mobile phase. Chromatographic peaks were resolved with 0.7 ml/min flow rate. Drug was extracted with ethyl acetate solvent by liquid–liquid extraction method. Monitoring of transition of m/z 506.2 and 71.0 for amprenavir and 628 and 421 for methyl-indinavir was made on multiple reaction monitoring.

Results: Calibration curve of amprenavir was linear over 1–600 ng/ml concentration range with regression coefficient (r²) value of >0.99. The % relative standard deviation values were <8.5% for interday and intraday precision and accuracy. The method has excellent recovery, and the percentage recovery values of lower quality control (QC), median QC, and higher QC samples were 101.86%, 102.8%, and 99.28%, respectively.

Conclusion: The drug was stable for more time at variable stability conditions, and method was successfully applicable to regular analysis of amprenavir in biological matrices.

Keywords: Amprenavir, Protease inhibitor, Liquid chromatography tandem-mass spectrometry, Food and Drug Administration guidelines and dilution integrity.

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INTRODUCTION

Amprenavir is a protease inhibitor with activity against human immunodeficiency virus type-1 (HIV-1). Protease inhibitors block the part of HIV called protease. HIV-1 protease is an enzyme required for the proteolytic cleavage of the viral polyprotein precursors into the individual functional proteins found in infectious HIV-1. Amprenavir chemically designated as (3S)-oxolan-3-yl N-[(2S,3R)-3-hydroxy-4-[N-(2-methylpropyl)(4-aminobenzene)sulfonamido]-1-phenylbutan-2-yl] carbamate with molecular formula C₂₅H₃₅N₃O₆S (Fig. 1) and molecular weight 505.628 g/mol [1-3]. Amprenavir binds to the protease active site and inhibits the activity of the enzyme. This inhibition prevents cleavage of the viral polyproteins resulting in the formation of immature non-infectious viral particles. Protease inhibitors are almost always used in combination with at least two other anti-HIV drugs. Amprenavir inhibits the HIV viral proteinase enzyme which prevents cleavage of the gag-pol polyprotein, resulting in noninfectious, immature viral particles. Hepatic amprenavir is metabolized in the liver by the cytochrome P450 3A4 (CYP3A4) enzyme system. The two major metabolites result from oxidation of the tetrahydrofuran and aniline moieties. Glucuronide conjugates of oxidized metabolites have been identified as minor metabolites in urine and feces [4,5].

Drug literature review discloses only few analytical quantification methods for the amprenavir in bulk, formulations, and biological matrices. The reported analytical techniques were high-performance liquid chromatography (HPLC) [6], spectroscopic [7], and liquid chromatography tandem-mass spectrometry (LC-MS/MS) [8]. Goal of the research was to develop a fast and sensitive LC-MS/MS technique for the quantification of amprenavir in human plasma samples and application of method validation as per the regulatory guidelines.

METHODS

Chemicals and reagents

Amprenavir (purity: 99.87%) was obtained from MSN Laboratories, India. Internal standard (IS) (methyl-indinavir) of 99.81% was acquired from hetero drugs Pvt. Ltd., Hyderabad, India. Acetonitrile of HPLC grade and formic acid of analytical grade were bought from J.T.Baker, Hyderabad, India. In the present research work, water used from Milli-Q water purification system installed in the laboratory obtained from Bengaluru, India.

LC-MS/MS system

A modular LC system (Shimadzu, Japan) equipped with a DGU-20A3 solvent degasser, binary LC-20AD prominence pump, column temperature ovens (CTO)-ASVP-oven for column, and high-throughput SILHTEC autosampler were utilized for the present research.

Chromatography was achieved on a reversed-phase (RP) Symmetry C₁₈ (50 mm×4.6 mm, 3.5 μm) with isocratic elution by acetonitrile and 0.1% v/v formic acid in the ratio of 90:10 v/v as mobile phase. Chromatographic peaks were resolved by the mobile phase with a flow rate of 0.7 ml/min. Amprenavir and methyl-indinavir internal standard were separated in the total run time of 5 min. The autosampler temperature and analytical column temperatures were kept at 5°C and 35°C, respectively.

The eluents of the LC system were infused into the electrospray ionization source operated with positive ionization method. Starting 0.5 min eluent was avoided from the chromatographic system to evade unnecessary impurities from the various salts existed in the human plasma samples.

In the mass system, following conditions were applied: Gas-1 – nitrogen (40 psi), gas-2 – nitrogen (40 psi), temperature of ion source – 400°C, curtain gas – nitrogen (25 psi), and voltage of ion spray – 5000 V. Monitoring of transition of m/z 506.2 and 71.0 for amprenavir and 628 and 421 for methyl-indinavir was made on multiple reaction monitoring. The mass conditions are presented in Table 1.

Sample preparation

To 250 µl of spiked plasma, 50 µl of methyl-indinavir (1 µg/ml) was mixed and sonicated for 15 s. To the resulting solution, 500 µl of ethyl acetate was added and vortexed for 5 min, followed by centrifuged at 5000 rpm for 10–20 min at 5°C. The organic phase was dried in a lyophilizer. The final residue was dissolved in 200 µl of mobile phase and transferred into a pre-labeled autosampler vials and infused into an LC-MS/MS system.

Preparation of standard stock and calibration standards (CCs)

Amprenavir and IS stock solutions were processed in 60% methanol at concentration level of 1000 µg/ml. Quality control (QC) and CC solutions were processed by spiking blank human plasma sample from the amprenavir stock solution. CC solutions of eight concentration levels were prepared to produce the final concentrations of 2.0, 4.0, 20.0, 40.0, 100.0, 200.0, 300.0, and 600.0 ng/ml. Lower QC (LQC) standard, median QC (MQC) standard, and higher QC (HQC) standards were QC sample solutions and were prepared to produce the concentrations of 4, 200, and 500 ng/ml, respectively. All the stock, CC and QC solutions were stored at –20°C till the method of analysis.

Validation

The method of analysis was assessed by validation parameters such as sensitivity, precision, linearity, recovery, dilution integrity, accuracy, matrix effect (ME), and stability. Three QC samples of LQC, MQC, and HQCs as well as lower limit of quantification (LLOQ) were employed and analyzed in method validation [9-12].

Precision and accuracy (PA)

Intraday and interday PA were examined as a part of PA parameter. Intraday PA was evaluated by injecting QC solutions (2, 200, and 500 ng/ml) and LLOQ (1.0 ng/ml) in 5 replicates in a day arbitrarily. Interday PA was evaluated by injecting the same QC and LLOQ solutions once in a day for 5 different days. The % relative standard deviations (RSDs) for LQC, MQC, and HQCs should be ≤20% for LLOQ and ≤15% for the remaining QC standards [13].

Linearity

CC standards (non-zero) of 8 different concentrations at 2.0, 4.0, 20.0, 40.0, 100.0, 200.0, 300.0, and 600.0 ng/ml were prepared and analyzed in three separate runs. Linearity curve (peak area ratio of amprenavir and methyl-indinavir peaks against nominal concentration) was plotted by least squares linear regression and reciprocal of the squared concentration (1/x²) used as a weighting factor. Deviation should be within ±20% for LLOQ and ±15% for remaining concentrations.

Specificity and selectivity

Method selectivity was analyzed by comparing the chromatograms obtained from blank and spiked solutions. Method specificity was analyzed by infusing six different lots of blank plasma solutions to ensure no endogenous compounds interfere with amprenavir and IS.

Recovery and ME

Amprenavir recovery was evaluated by paralleling the mean peak areas of extracted and un-extracted samples at low, medium, and high QC standard levels. At each concentration level, percentage recoveries were calculated, and finally, overall mean recovery was calculated. The ME was analyzed by paralleling the un-extracted samples with post-extracted samples [14-16].

$$\% \text{ Recovery of amprenavir} = \frac{\text{Mean amprenavir peak response in extracted samples}}{\text{Mean amprenavir peak response in un-extracted samples}} \times 100$$

Stability

Stability was analyzed at LQC, MQC, and HQC levels. It includes bench-top, freeze-thaw, autosampler, and long-term stabilities. The bench-top stability was evaluated for 5 h at ambient temperature (25°C). Freeze and thaw stability was analyzed by storing the QC samples at –70°C for at least 3 h, and for thaw cycle, keep the solutions at room temperature. Repeat the freeze and thaw cycles for 3 times. The autosampler stability was analyzed by placing the QC solutions in autosampler at 10°C for 8 h. Long-term stability was evaluated by placing the QC samples in a freezer at –70°C for 3 months [17-19].

Dilution integrity

The sample solution above the upper limit of calibration range was prepared and evaluated for PA parameters. The percentage nominal concentration must be ±15%.

Ruggedness

Method ruggedness was assessed by processing QC standards for one PA batch utilizing different columns of the same composition by different analysts. The %RSDs for LQC, MQC, and HQCs should be ≤20% for LLOQ and ≤15% for the remaining QC standards.

RESULTS AND DISCUSSION

The LC-MS/MS chromatograms of amprenavir – blank and LLOQ and LQC, MQC, and HQC concentration levels – are shown in Figs. 2 and 3.

Method validation

Specificity

From the Figs. 2 and 3, system chromatographic conditions were clearly separating amprenavir and IS from endogenous and other plasma

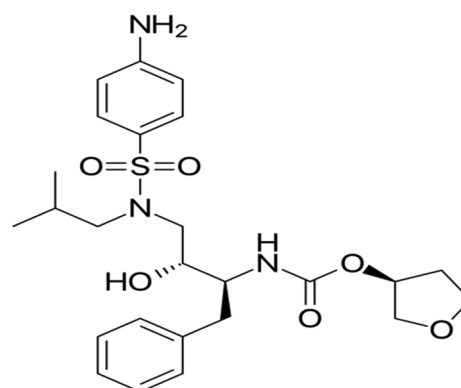


Fig. 1: Structure of amprenavir

Table 1: Mass conditions for amprenavir and IS

Component	Precursor ion (m/z)	Product ion (m/z)	DP (V)	EP (V)	CEP (V)	CE (V)	CXP (V)
Amprenavir	506.2	71.0	40	8	25	30	5
Methyl-indinavir	628	421	100	8	25	82	5

DP: Declustering potential, EP: Entrance potential, CEP: Collision cell entrance potential, CE: Collisional energy, CXP: Collision cell exit potential, IS: Internal standard

substances. The amprenavir-LLOQ peak response is more than 20% the interference peak response, and methyl-indinavir peak response is more than 5% from the interference peak response.

PA

Amprenavir interday and intraday PA were analyzed, and the %RSD values are calculated for the same and are tabulated in Table 2.

Table 2: Amprenavir precision and accuracy data

Nominal concentration (ng/ml)	Intraday		Interday	
	Accuracy (%)	%RSD ⁿ	Accuracy (%)	%RSD ⁿ
2	104.6	4.5	107.5	5.2
4	89.6	3.9	91.6	6.1
200	105.2	2.5	101.2	4.2
500	109.8	8.1	96.4	2.9

n=6 replicates (for precision), RSD: Relative standard deviation

Linearity

Amprenavir calibration graph was linear in the concentration range of 2–600 ng/ml with regression equation of $Y=0.4528X+3.145$. The regression coefficient (r^2) value is more than 0.99 which was acceptable as per the Food and Drug Administration (FDA) regulatory guidelines [18].

Recovery and ME

The method has excellent recovery and the percentage recovery values were 101.86%, 102.8%, and 99.28% for LQC, MQC, and HQC samples, respectively. The data for amprenavir recovery are tabulated in Table 3. The ME was evaluated at LQC and HQC level, and the calculated % coefficient of variation (CV) values was 4.59% and 3.68%, respectively.

Dilution integrity

Dilution integrity of amprenavir was performed and evaluated. The percentage nominal was within the limit ($\pm 15\%$), and the estimated precision was $\leq 15\%$. It shows that the drug can be dilute to 20 times and the results will be reproducible.

Table 3: Recovery data for amprenavir

ID	LQC			MQC			HQC		
	Un extracted (Area ratio)	Extracted (Area ratio)	%Recovery	Un extracted (Area ratio)	Extracted (Area ratio)	%Recovery	Un extracted (Area ratio)	Extracted (Area ratio)	%Recovery
1	0.213	0.251	117.85	0.648	0.634	98.451	0.897	0.885	98.66
2	0.246	0.235	95.59	0.612	0.65	106.20	0.905	0.879	97.12
3	0.254	0.249	98.03	0.598	0.635	106.18	0.856	0.985	115.07
4	0.217	0.215	99.08	0.688	0.643	93.45	0.956	0.912	95.39
5	0.214	0.219	102.34	0.675	0.684	101.33	0.965	0.845	87.56
6	0.243	0.239	98.35	0.598	0.665	111.20	0.895	0.912	101.89
Mean	0.23	0.24	101.86	0.64	0.6525	102.80	0.912	0.903	99.28
SD	0.017	0.014	7.42	0.02	0.017	5.80	0.037	0.04	8.301
%CV	7.29	5.83	7.28	5.68	2.62	5.64	4.11	4.77	8.36

LQC: Lower quality control, MQC: Median quality control, HQC: Higher quality control

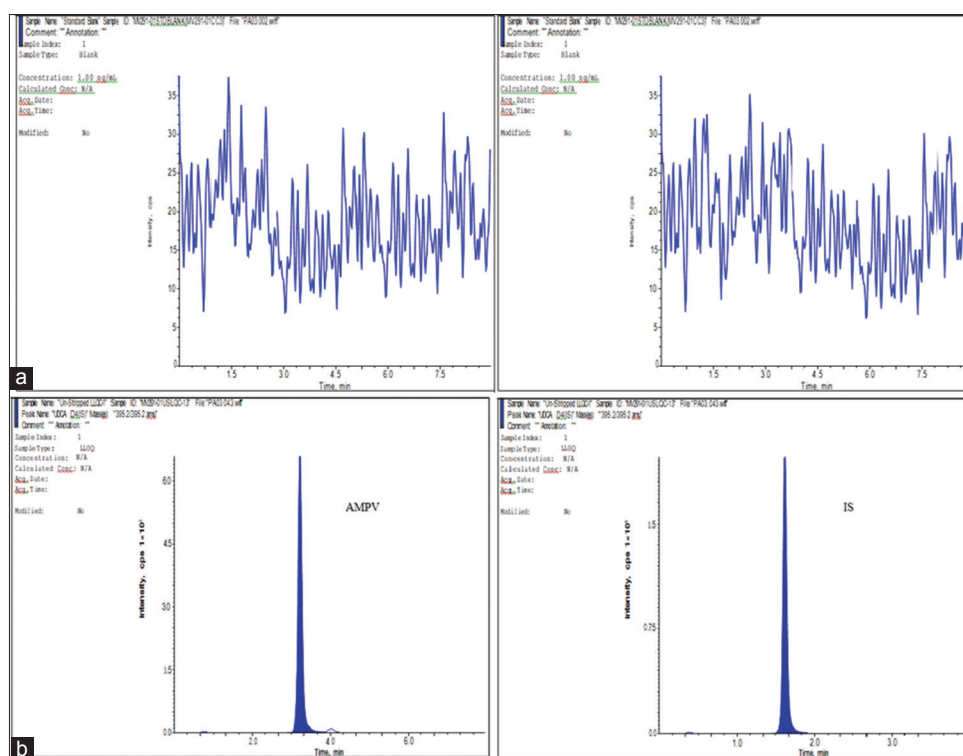


Fig. 2: Amprenavir chromatograms (a) blank and (b) spiked lower limit of quantification samples

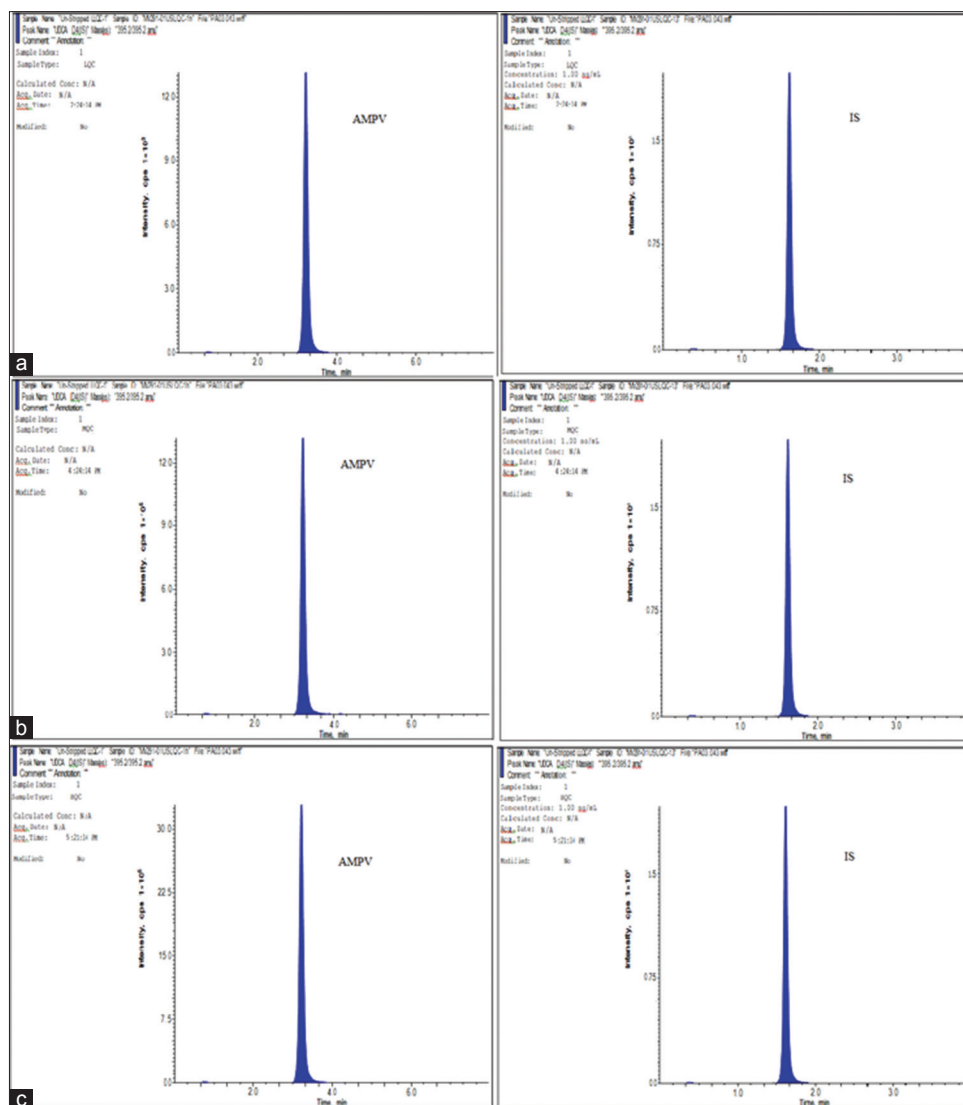


Fig. 3: Chromatograms of amprenavir spiked (a) lower quality control, (b) median quality control, and (c) higher quality control samples

Table 4: Stability for amprenavir

Drug	Concentration (ng/ml)	Bench-top stability		Autosampler stability		Freeze and thaw stability		Long-term stability	
		Mean±SD (ng/ml)	%CV	Mean±SD (ng/ml)	%CV	Mean±SD (ng/ml)	%CV	Mean±SD (ng/ml)	%CV
Amprenavir	4	4±0.5	10.7	4±0.3	6.1	4±0.2	5.3	4±0.2	6.1
	200	200±17.0	9.6	200±11.0	8.6	200±8.0	4.6	200±10.0	5.3
	500	500±20	6.6	500±8.0	4.6	500±7.3	9.4	500±11.0	8.3

SD: Standard deviation, CV: Coefficient of variance

Stability

All the QC standards were exposed to different stability conditions and evaluated to analyze the stability of amprenavir. From evaluated %CV stability data, the drug was stable for more time at variable conditions such as bench-top stability (<10.8%), freeze-thaw stability (<9.5%), autosampler stability (<8.9%), and long-term stability (<8.4%), and the values are presented in Table 4.

Ruggedness

Method ruggedness for amprenavir was performed and evaluated. The % RSD values are calculated for the same and are tabulated in Table 5.

Table 5: Amprenavir ruggedness data

Nominal concentration (ng/ml)	Analyst-1 and column-1		Analyst-2 and column-2	
	Accuracy (%)	%RSD ⁿ	Accuracy (%)	%RSD ⁿ
2	106.3	5.2	102.6	6.8
4	98.4	2.9	94.6	5.9
200	105.8	6.8	97.2	3.8
500	104.2	2.6	106.9	4.5

n: 6 replicates, RSD: Relative standard deviation

