



NEW DETERMINATION AND QUANTIFICATION OF POSACONAZOLE IN HUMAN PLASMA BY LIQUID CHROMATOGRAPHY WITH TANDEM MASS SPECTROMETRY: IT'S APPLICATION TO A PHARMACOKINETIC STUDY

Venkata Ramu Derangula¹ and Venkateswarlu Ponneri^{2*}

¹Research Studies, Rayalaseema University, Kurnool-518 002, India.

^{2*}Analytical and Environmental Chemistry Division, Department of Chemistry, Sri Venkateswara University, Tirupati-517502, India.

***Corresponding Author: Dr. Venkateswarlu Ponneri**

Research Studies, Rayalaseema University, Kurnool-518 002, India.

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ABSTRACT

A simple, rapid and sensitive liquid chromatography with tandem mass spectrometry (LC-MS/MS) assay method has been proposed for the determination of Posaconazole in human plasma samples using Posaconazole D₅ as internal standard (IS). Analyte and the IS were extracted from the 100 µL of K₂ EDTA human plasma by simple Protein Precipitation extraction. The chromatographic separation was achieved on a Zorbax Eclipse XDB-C₁₈ column by using a mixture of Acetonitrile and 5mM ammonium acetate in 0.1% formic acid buffer (85:15, v/v) as the mobile phase at a flow rate of 1.0 mL/min. The calibration curve obtained was linear ($r^2 \geq 0.99$) over the concentration range of 5.01 – 2502.29 ng/mL. The Mass detection of Posaconazole involves m/z - 701.20 (parent) and 683.30 (product) and Posaconazole D₅ involves m/z - 706.30 (parent) and 687.30 (product) as internal standard in Positive ion mode. Method validation was performed as per FDA guidelines and the results met the acceptance criteria. The intra-day and inter-day precision (%CV) and accuracy results in three validation batches across six concentration levels were well within the acceptance limits. A run time of 2.40 min for each sample made it possible to analyze more number of samples in short time, thus increasing the productivity. The proposed method successfully applied to a pharmacokinetic study of Posaconazole 40 mg/mL oral suspension formulation in healthy, adult, human male subjects under fed condition.

KEYWORDS: Posaconazole, human plasma, Protein Precipitation extraction, LC-MS/MS.

INTRODUCTION

Posaconazole is used to treat invasive aspergillosis and candidiasis and fungal infections caused by *Scedosporium* and *Fusarium* species, which may occur in immunocompromised patients. It is also used for the treatment of oropharyngeal candidiasis (OPC), including OPC refractory to itraconazole and/or fluconazole therapy.^[1-3] It is also used to treat invasive infections by *Candida*, *Mucor* and *Aspergillus* species in severely immunocompromised patients.^[4,5] Posaconazole works by disrupting the close packing of acyl chains of phospholipids, impairing the functions of certain membrane-bound enzyme systems such as ATPase and enzymes of the electron transport system, thus inhibiting growth of the fungi. It does this by blocking the synthesis of ergosterol by inhibiting of the enzyme lanosterol 14 α -demethylase and accumulation of methylated sterol precursors. Posaconazole is significantly more potent at inhibiting 14- α demethylase than itraconazole.^[6-9] Few methods were reported on estimation of

Posaconazole by different analytical methods. Störzinger, D et.al reported Development and validation of a high-performance liquid chromatography assay for posaconazole utilizing solid-phase extraction.

Groll AH et.al, shown clinical pharmacology and potential for management of fungal infections. Chhun, S. et. al reported Simultaneous quantification of voriconazole and posaconazole in human plasma by high-performance liquid chromatography^[10-17] with ultra-violet detection. Among the methods present method involve less retention time and protein precipitation extraction which is modern and less reported method. The analytical method should satisfy the scientists in terms of simplicity, sensitivity, runtime, time consumption, sample volume and efficient extraction procedure.^[18-20]

MATERIALS AND METHODS

A Reference sample of Posaconazole (99.42%) was obtained from Indoco Remedies Limited (Mumbai,

India), while Posaconazole D₅ (99.37%) was from VIVEN Life Sciences Pvt. Limited (Mumbai, India). Their chemical structures are shown in Fig. 1. Water used for the LC-MS/MS analysis was prepared by using Milli Q water purification system procured from Millipore (Bangalore, India). HPLC grade Acetonitrile was purchased from J.T. Baker (Phillipsburg, USA), while analytical grade formic acid was from Merck Ltd (Mumbai, India). The control K2-human plasma sample was procured from Deccan's Pathological Lab's (Hyderabad, India).

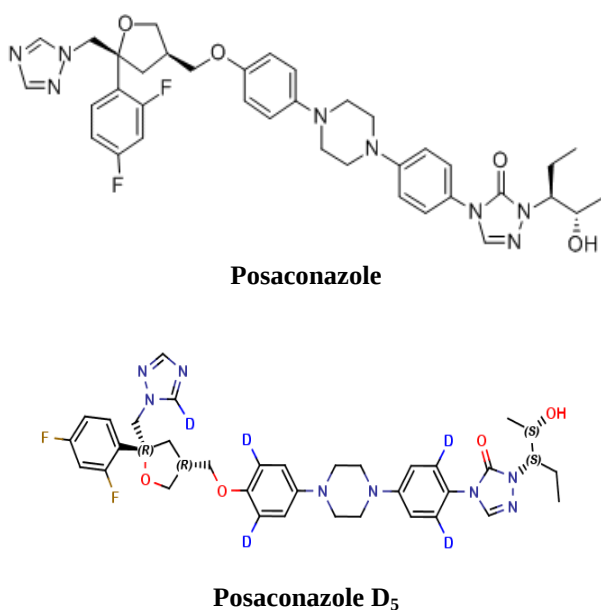


Fig 1: Chemical structures of Posaconazole (A) and Posaconazole D₅ (B) (IS).

LC-MS/MS instrument and conditions

An HPLC system consisting of a Eclipse XDB-C₁₈ column 4.6 mmx150 mm x5µm; Agilent Technologies, Santa Clara, CA, USA), a binary LC-20AD prominence

pump, an auto sampler (SIL-HTc) and a solvent degasser (DGU-20A₃) was used for the study. Aliquot of 10 µL of the processed samples were injected into the column, which was kept at 40°C. An isocratic mobile phase consisting of a mixture of Acetonitrile and 5mM ammonium acetate in 0.1% formic acid buffer (85:15, v/v) was used to separate the analyte from the endogenous components and delivered at a flow rate of 1.0 mL/min into the electrospray ionization chamber of the mass spectrometer. Quantification was achieved with MS-MS detection in positive ion mode for the analyte and the IS using an MDS Sciex API-4000 mass spectrometer (Foster City, CA, USA) equipped with a Turboionspray™ interface at 500°C. The ion spray voltage was set at 4500 V. The source parameters viz. the nebulizer gas (GS1), auxiliary gas (GS2), curtain gas and collision gas were set at 40, 40, 30 and 8 psi, respectively. The compound parameters viz. the declustering potential (DP), collision energy (CE), entrance potential (EP) and collision cell exit potential (CXP) were 24,46,10,14 V for Posaconazole and 24,46,10,14 V for the IS. Detection of the ions was carried out in the multiple reaction monitoring mode (MRM), by monitoring the transition pairs of *m/z* 701.20 parent ion to the *m/z* 683.30 for product ion of Posaconazole and *m/z* 706.30 parent ion to the *m/z* 687.30 product ion for the IS were shown in Fig.2. Quadrupoles Q1 and Q3 were set on unit resolution. The analysis data obtained were processed by Analyst Software™ (version 1.6.1). The MRM technique provided intrinsic selectivity and sensitivity, hence chosen for the study.^[20, 21]

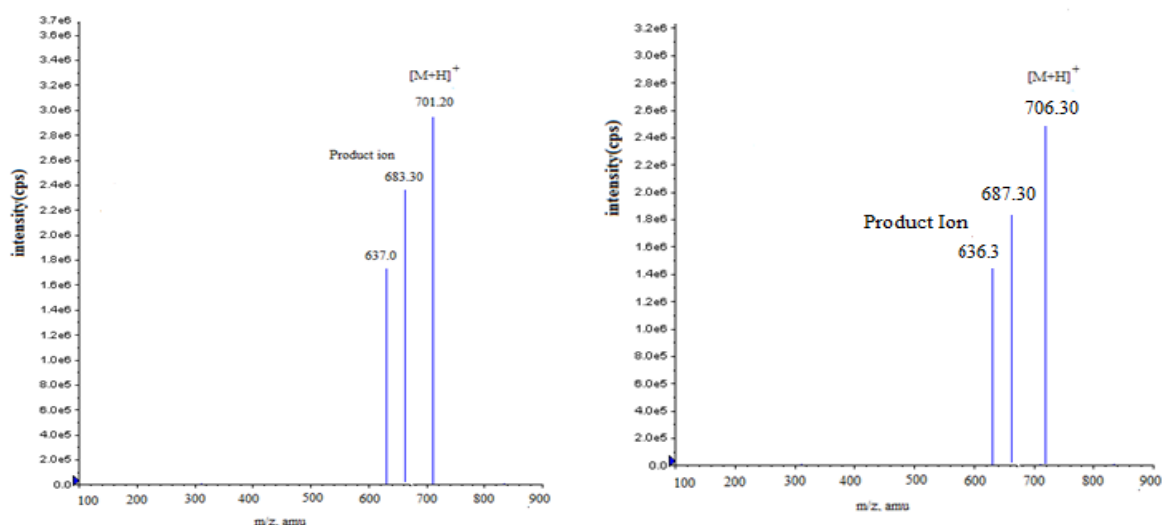


Fig 2: Product ion mass spectra of [M+H]⁺ of Posaconazole and Posaconazole D₅ (IS).

Preparation of plasma standards and quality controls

Standard stock solution of Posaconazole and IS (1mg/mL) were prepared in Acetonitrile. Working solutions for calibration and controls were prepared by appropriate dilution in Acetonitrile and water (60:40, v/v; diluent). The IS working solution (30000 ng/mL) was prepared by diluting its stock solution with diluent. Stock solutions of Posaconazole and IS were found to be stable for 7 days at 2-8 °C.

Calibration samples were prepared by spiking 950 µL of control K₂ EDTA human plasma with the 50 µL working standard solution of the analyte as a bulk, to obtain Posaconazole concentration levels of 5.01, 10.02, 40.11, 125.36, 250.73, 501.46, 1002.92, 1501.37, 2001.83 and 2502.29 ng/mL as a single batch at each concentration. Similarly, quality control (QC) samples were also prepared as a bulk based on an independent weighing of standard drug, at concentrations of 5.064 (LLOQ), 15.116 (LQC), 251.926 (MQC1), 1259.630 (MQC2) and 1952.91ng/mL (HQC) as a single batch at each concentration. The calibration and control bulk samples were divided into aliquots in micro centrifuge tubes (Tarson, 2 mL) and stored in the freezer at -70 ± 10 °C until analyses.

Sample processing

All frozen subject samples, calibration standards and quality control samples were thawed and allowed to equilibrate at room temperature prior to analysis. Withdraw one set of calibration curve standards, one or more sets of quality control samples and subject plasma

samples from the deep freezer and allow to thaw at room temperature. Vortex the thawed samples to ensure complete mixing of the contents. Pipette out 100 µL of the sample into pre-labelled RIA vial tubes. Add 10 µL of internal standards dilution (30000.000ng/mL of Posaconazole D5) except in blank and subject pre dose sample wherein add 10 µL of diluent and vortex. Also process 100 µL of each pre-dose sample with addition of 10 µL of internal standard dilution and vortex. To this 1.0 mL of mobile phase will be added and vortex. The samples will be centrifuged at 4000 rpm for 20 minutes at 4°C. Sample was transferred into loading vials and would be loaded into the auto sampler and it was injected into the LC-MS/MS system. The typical chromatograms of Posaconazole were shown in Fig.3.

Method development

The objective of the present work was to develop and fully validate an LC-MS/MS method for the determination of Posaconazole in human plasma with high sensitivity to monitor the concentration of Posaconazole for pharmacokinetic/bioequivalence studies. To develop a sensitive and selective bioanalytical method requires the judicious selection of chromatography column, mobile phase and organic solvent. These parameters should be carefully monitored to produce the required resolution from endogenous components which in turn affect sensitivity and reproducibility of the analytical method by ion suppression.

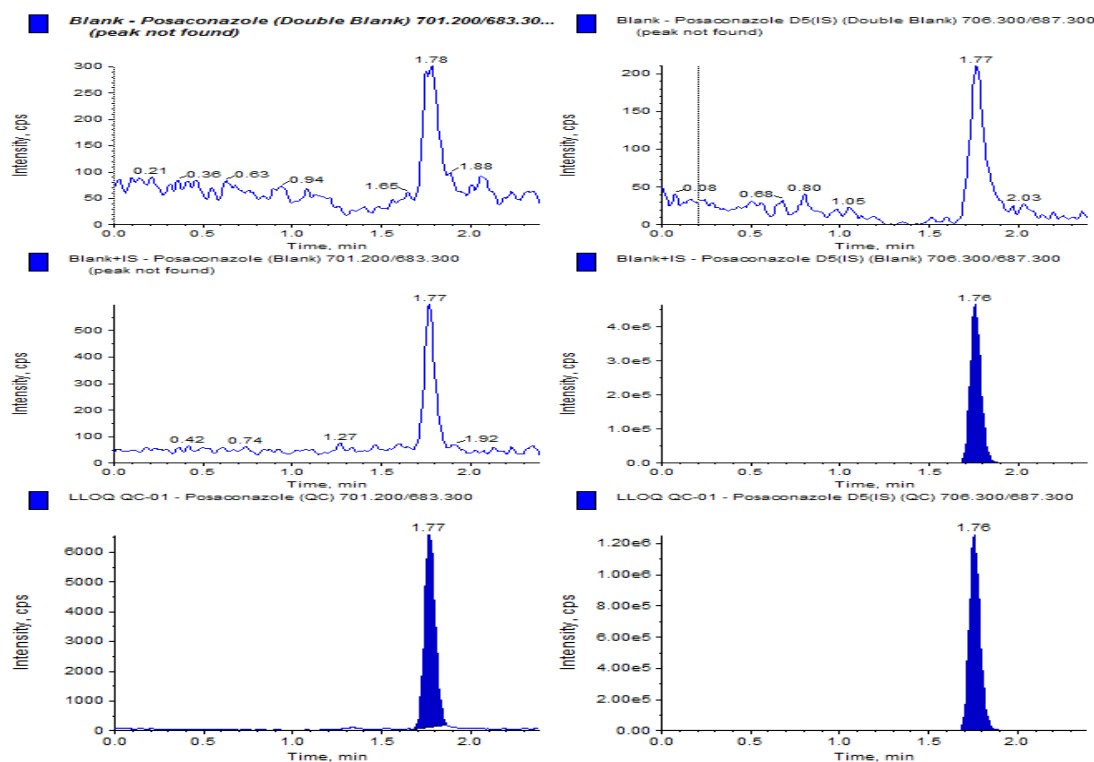


Fig 3: Typical MRM chromatograms of Posaconazole (left panel) and IS (right panel) in human blank plasma, and human plasma spiked with IS, a LLOQ sample along with IS.

Once Chromatographic column, mobile phase pH and organic solvent are set then flow rate, column temperature and buffer type and concentration can be manipulated for optimal response. An ideal internal standard should mimic the analyte in as many ways as possible. It should have a similar structure, same physicochemical properties or can be a labeled compound. For LC-MS/MS analysis, use of stable isotope-labeled drugs as internal standards proves to be helpful when a significant matrix effect is possible and to increase assay precision and limit variable recovery between analyte and the IS.

Method validation

A thorough and complete method validation of posaconazole in human plasma was carried out as per USFDA guidelines. The parameters determined were carryover test, selectivity, specificity, sensitivity, matrix effect, linearity, precision, accuracy, recovery, dilution integrity and stability. Carry over experiment was performed to verify any carryover of analyte and IS which may reflect in subsequent runs. The design of the study comprised of the following sequence of injections i.e. blank plasma sample → six samples of LLOQ → blank plasma sample → ULOQ sample → blank plasma samples to check for any interference due to carry over. Selectivity was assessed by comparing the chromatograms of six different batches of blank plasma obtained from six different sources. Sensitivity was determined by analyzing six replicates of plasma samples spiked with the lowest level of the calibration curve concentrations. Matrix effect was checked with six different lots of K₂ EDTA plasma. Three replicate samples each of LQC and HQC were prepared from different lots of plasma (18 QC samples in total). The linearity of the method was determined by analysis of standard plots associated with a ten-point (non-zero standards) standard calibration curve. In addition, blank plasma samples were also analyzed to confirm the absence of direct interferences. To determine intra-day accuracy and precision, a calibration curve and six replicates of LLOQ QC, LQC, MQC-1, MQC-2, and HQC were analyzed on the same day. Inter-day accuracy and precision were assessed by analyzing three batches of samples on two consecutive days. Recoveries of analyte and IS were determined by comparing the peak area of extracted analyte standard with the peak area of non-extracted standard. Recovery of Posaconazole was determined at a concentration of 15.116 (LQC), 1259.630 (MQC2) and 1952.91(HQC) ng/mL whereas for IS was determined at concentration of 1000 ng/mL. Dilution integrity was performed to extend the upper concentration limit with acceptable precision and accuracy. Six replicates each at a concentration of about 1.70 times of the uppermost calibration standard were diluted two and four -fold with blank plasma. The diluted samples were processed and analyzed. Stability tests were conducted to evaluate the analyte stability in stock solutions and in plasma samples under different

conditions. The stock solution stability at room temperature and refrigerated conditions (2-8°C) was performed by comparing the area response of the analyte (stability samples) with the response of the sample prepared from fresh stock solution. Bench top stability (15 h), processed samples stability (autosampler stability for 51 h, wet extract stability for 46 h and reinjection stability for 49 h.), freeze-thaw stability (4 cycles), long-term stability (50 days) were performed at LQC and HQC levels using six replicates at each level. Samples were considered to be stable if assay values were within the acceptable limits of accuracy ($\pm 15\%$ SD) and precision ($\leq 15\%$ RSD).

RESULTS AND DISCUSSION

System suitability

During method validation, the precision (% CV) of a system suitability test was found to be in the range 0.00-0.00% for retention time and 3.22-3.30% for the area response of Posaconazole and IS.

Carryover effect

Carryover evaluation was performed to ensure that it does not affect the accuracy and precision of the proposed method. No significant carryover was observed in blank sample after injection highest concentration of analyte (ULOQ) which indicates no carryover of the analyte in subsequent samples.

Selectivity and chromatography

The selectivity of the method was examined by analyzing blank human plasma extract and an extract spiked only with the IS. As shown in Fig. 3, no significant direct interference in the blank plasma traces was observed from endogenous substances in drug-free human plasma at the retention time of the analyte and the IS.

Matrix effect

Matrix effect assessment was done with the aim to check the effect of different lots of plasma on the back calculated value of QC's nominal concentration. The results found were well within the acceptable limits as shown in Tab.1. No significant matrix effect was observed in all the six batches of human plasma for the analyte at low and high quality control concentrations. Also, the extraction method was rugged enough and gave accurate and consistent results when applied to real subject samples.

Tab 1: Matrix effect of Posaconazole in human plasma ($n = 6$).

Plasma lot	LQC (ng/mL)		HQC (ng/mL)	
	Concentration found (mean $\pm$ SD; ng/mL)	% Accuracy	Concentration found (mean $\pm$ SD; ng/mL)	% Accuracy
Lot 1	14.73 $\pm$ 0.12	97.5	1984 $\pm$ 29.03	102
Lot 2	14.66 $\pm$ 0.10	97.0	1960 $\pm$ 4.97	100
Lot 3	14.65 $\pm$ 0.28	97.0	1950 $\pm$ 45.18	99.9
Lot 4	14.45 $\pm$ 0.22	95.6	1963 $\pm$ 22.91	101
Lot 5	14.69 $\pm$ 0.20	97.2	1979 $\pm$ 69.50	101
Lot 6	14.93 $\pm$ 0.21	98.8	1981 $\pm$ 27.84	101

Linearity, precision and accuracy

The ten point calibration curve was found to be linear over the concentration range of 5.01 - 2502.29 ng/mL for Posaconazole. After comparing the two weighting models ($1/x$ and $1/x^2$), a regression equation with a weighting factor of $1/x^2$ of the drug to the IS concentration was found to produce the best fit for the

concentration–detector response relationship. The mean correlation coefficient of the weighted calibration curves generated during the validation was ≥ 0.99 . The results for intra–day and inter–day precision and accuracy in plasma quality control samples are summarized in Tab.2.

Tab 2: Precision and accuracy data for Posaconazole.

Quality control	Run	Concentration found (mean $\pm$ SD; ng/mL)	Precision (%)	Accuracy (%)
Intra–day variations (Six replicates at each concentration)				
LLOQ	1	4.97 $\pm$ 0.38	7.74	98.1
	2	5.27 $\pm$ 0.25	4.75	104
	3	5.05 $\pm$ 0.23	4.49	96.1
	4	5.06 $\pm$ 0.08	1.59	99.9
	5	4.88 $\pm$ 0.17	3.50	96.3
LQC	1	15.6 $\pm$ 0.70	4.49	103
	2	15.3 $\pm$ 0.45	2.96	101
	3	15.6 $\pm$ 0.57	3.64	103
	4	15.6 $\pm$ 0.37	2.37	103
	5	15.6 $\pm$ 0.61	3.87	103
MQC1	1	266 $\pm$ 12.9	4.86	106
	2	259 $\pm$ 10.4	4.01	103
	3	267 $\pm$ 11.9	4.47	106
	4	262 $\pm$ 7.90	3.02	104
	5	267 $\pm$ 8.52	3.19	106
MQC2	1	1260 $\pm$ 43.3	3.44	100
	2	1226 $\pm$ 63.0	5.14	97.3
	3	1259 $\pm$ 52.7	4.19	100
	4	1251 $\pm$ 50.8	4.06	99.3
	5	1250 $\pm$ 52.2	4.18	99.2
HQC	1	1960 $\pm$ 19.5	1.00	100
	2	1944 $\pm$ 53.9	2.77	99.5
	3	1979 $\pm$ 49.2	2.49	101
	4	1954 $\pm$ 51.6	2.64	100
	5	1991 $\pm$ 38.6	1.94	102
Inter–day variations (Eighteen replicates at each concentration)				
LLOQ		5.05 $\pm$ 0.26	5.20	99.6
LQC		15.5 $\pm$ 0.52	3.38	103
MQC1		264 $\pm$ 10.3	3.89	105
MQC2		1249 $\pm$ 50.6	4.05	2.28
HQC		1966 $\pm$ 44.8	2.28	101
Spiked concentrations of LLOQ, LQC, MQC1, MQC2 and HQC are 5.06, 15.1, 251, 1260 and 1953 ng/mL, respectively.				

Stability studies and dilution integrity

In the different stability experiments carried out viz. bench top stability (15 h), autosampler stability (51 h), wet extract stability (46 h), repeated freeze–thaw cycles (4cycles), reinjection stability (49 h) and long term stability at -70°C for 50 days the mean % nominal

values of the analyte were found to be within $\pm 15\%$ of the predicted concentrations for the analyte at their LQC and HQC levels shown Tab.3. Thus, the results were found to be within the acceptable limits during the entire validation. The MRM chromatograms of Posaconazole were shown in Fig.4.

Tab. 3: Stability data for Posaconazole in plasma ($n=6$)

Stability test	QC (spiked concentration (ng/mL))	Mean $\pm$ SD (ng/mL)	Precision (%)	Accuracy/ Stability (%)
Process ^a	15.1	16.3 ± 0.41	2.50	108
	1953	2009 ± 11.2	0.56	103
Process ^b	15.1	16.6 ± 0.12	0.70	110
	1953	2083 ± 111	5.33	107
Bench top ^c	15.1	16.9 ± 0.26	1.56	112
	1953	2222 ± 14.9	0.67	114
FT ^d	15.1	16.4 ± 0.21	0.21	108
	1953	2000 ± 13.7	0.68	102
Reinjection ^e	15.1	14.9 ± 0.33	2.20	98.9
	1953	1920 ± 43.1	2.25	98.3
Long-term ^f	15.1	14.8 ± 0.56	3.76	98.2
	1953	1958 ± 20.8	1.06	99.9

^a after 51 h in autosampler at 15°C ; ^b after 46 h at 20°C ; ^c after 15 h at room temperature; ^d after 4 freeze and thaw cycles; ^e after 49 h of Reinjection; ^f at -70°C for 50 days

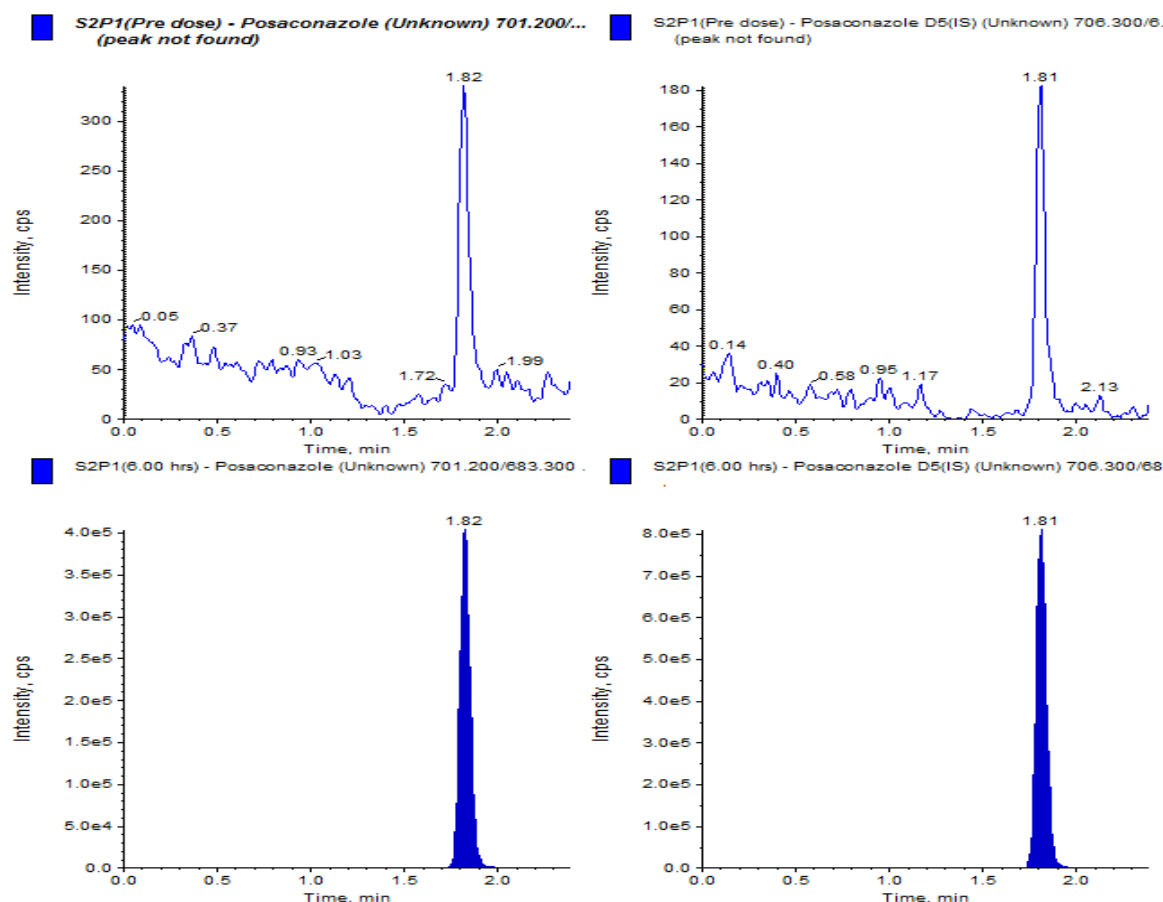


Fig 4: MRM chromatograms resulting from the analysis of subject blank plasma sample (A) and 6 h subject plasma sample (B), after the administration of a 40 mg/mL (10 mL of oral suspension of Posaconazole.

Pharmacokinetic data

A single dose pharmacokinetic study was performed in healthy South Indian male subjects ($n = 6$). The Ethics Committee (Samkshema Independent Ethics Committee, Hyderabad, India) approved the protocol and the volunteers provided with written informed consent. The subjects were fasted 12 h before administration of the drug formulation. Six healthy South Indian male subjects with an age group of 20–40 years and body-mass index (BMI) of $\geq 18.5 \text{ kg/m}^2$ and $\leq 24.9 \text{ kg/m}^2$, with body weight not less than 50 kg were chosen for the study. They were randomly assigned to one group and took a single oral dose of posaconazole 40 mg/mL oral suspension (10 mL of oral suspension i.e., 400 mg of posaconazole respectively). Blood samples were collected at pre-dose (0.00) and 0.50, 1.00, 1.50, 2.00, 2.33, 2.67, 3.00, 3.33, 3.67, 4.00, 4.33, 4.67, 5.00, 5.33, 5.67, 6.00, 7.00, 8.00, 10.00, 12.00, 14.00, 16.00, 24.00, 36.00, 48.00, 60.00 and 72.00 h, in K₂ EDTA vacutainer (5 mL) collection tubes (BD, Franklin, NJ, USA). The tubes were centrifuged at 3200 rpm for 10 min and the plasma was collected. The collected plasma samples were stored at $-70 \pm 10^\circ\text{C}$ till their use. Plasma samples were spiked with the IS and processed as per the extraction procedure described earlier. The main pharmacokinetic parameters of Posaconazole were calculated by non-compartmental model using WinNonlin Version 5.2. An incurred sample re-analysis was also conducted by selecting the 6 subject samples (2 samples from each subject) near C_{max} and the

elimination phase. The percent change in the value should not be more than $\pm 20\%$ Tab.5. The validated method was successfully applied for a pharmacokinetic study of Posaconazole in 6 healthy South Indian adult male subjects divided in to one group who received a single oral dose of posaconazole 40 mg / mL oral suspension (10 mL of oral suspension i.e., 400 mg of posaconazole respectively under fed condition. With the reported lowest LLOQ (5.015 ng/mL) Posaconazole was not quantifiable beyond 72 h post-dosing. The mean plasma concentration-time profile of Posaconazole was presented in Fig. 5 and the corresponding pharmacokinetic parameters were listed in Tab.4 and Tab.5.

Tab 4: Pharmacokinetic parameters of Posaconazole 40 mg/mL (10 mL of oral suspension i.e., 400 mg of Posaconazole) formulation in healthy, adult, human male subjects under fed condition ($n=6$, Mean $\pm$ SD).

PK Parameter	Mean $\pm$ SD
	40 mg/mL
t_{max} (h)	5.00 $\pm$ 0.29
C_{max} (ng/mL)	719 $\pm$ 26.2
AUC_{0-t} (ng h/mL)	18403 $\pm$ 4333
$\text{AUC}_{0-\text{inf}}$ (ng h/mL)	23585 $\pm$ 7475
$t_{1/2}$ (h)	30.5 $\pm$ 8.99
Kel (h^{-1})	0.02 $\pm$ 0.00

Tab 5: Incurred samples re-analysis data of Posaconazole.

Sample	Posaconazole 40 mg/mL (10 mL of oral suspension i.e., 400 mg of Posaconazole)		
	Initial conc. (ng/mL)	Re-assay conc. (ng/mL)	Difference ^a (%)
1	612	599	2.20
2	117	107	9.18
3	689	651	5.65
4	163	159	2.27
5	759	701	7.93
6	117	110	6.47
7	782	750	4.05
8	140	129	8.38
9	638	622	2.54
10	56	55	2.46
11	478	426	11.55
12	48	41	15.28

^a Expressed as [(initial conc.–re-assay conc.)/average] $\times 100\%$.

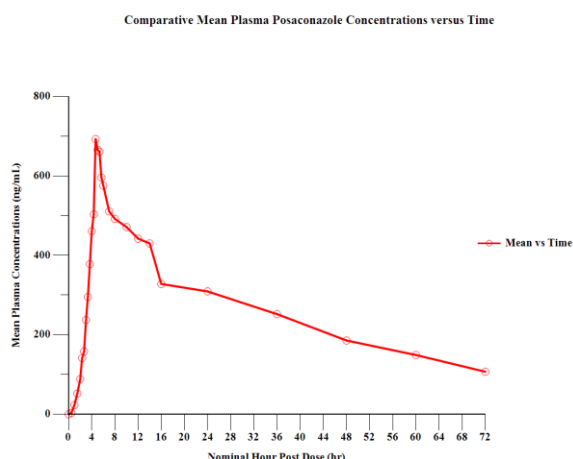


Fig 5: Mean plasma concentration-time profile of Posaconazole in human plasma following A single oral dose of Posaconazole 40 mg/mL (10 mL of oral suspension i.e., 400 mg of Posaconazole) to healthy volunteers (n=6).

CONCLUSION

The present research work involves successful method development and validation of a simple, sensitive and rapid LC-MS/MS method for the determination of Posaconazole in human plasma samples according to commonly acceptable FDA guidelines. Moreover, the total analysis time (extraction and chromatography) is the shortest. The advantage of this method is that a relatively more number of samples can be analyzed in short time, thus increasing the output. The PPE minimizes the chances of errors, saves considerable time and simplifies the sample preparation procedure. The method showed suitability for pharmacokinetic studies in humans. From the results of all the validation parameters, we can conclude that the developed method can be useful for bioavailability and bioequivalence (BA/BE) studies and routine therapeutic drug monitoring with the desired precision and accuracy.

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