

High-Performance Liquid Chromatographic Quantification of Rifampicin in Human Plasma: Method for Therapeutic Drug Monitoring

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Abstract. A high performance liquid chromatography (HPLC) method has been developed that allows quantification of Rifampicin in human plasma. The method is based on the precipitation of proteins in human plasma with methanol. Optimal assay conditions were found with a C18 column and a simple mobile phase consisting of 0.05 M dipotassic hydrogen phosphate buffer and acetonitrile (53/47, V/V) with 0.086 % diethylamin, pH = 4.46. The flow-rate was 0.6 mL/min and the drug was monitored at 340 nm. Results from the HPLC analyses showed that the assay method is linear in the concentration range of 1-40 µg/mL ($r^2 > 0.99$). The limit of quantification and limit of detection of Rifampicin were 0.632 µg/mL and 0.208 µg/mL, respectively. Intraday and interday coefficient of variation and bias were below 10% for all samples, suggesting good precision and accuracy of the method. Recoveries were greater than 90% in a plasma sample volume of 100 µL. The method is being successfully applied to therapeutic drug monitoring of Rifampicin in plasma samples of tuberculosis and staphylococcal infections patients.

Keywords: rifampicin, liquid chromatography, therapeutic drug monitoring

Introduction

Rifampicin (RIF), a semi-synthetic macrocyclic complex antibiotic derived from *Streptomyces mediterranei*, is member of rifamycin class antibiotics (Maggi *et al.*, 1966) used for the treatment of tuberculosis and other infectious diseases such as staphylococcal infections (Tsankov and Angelova, 2003). It is categorised as one of the first line antituberculous agents; however various side effects such as hepatotoxicity, allergic rashes, lack of appetite and nausea or immunological disturbances have been reported with its administration (Gallieni *et al.*, 1999). The therapeutic drug monitoring (TDM) of RIF is necessary if the cultures are still positive for *Mycobacterium* after two months of treatment. TDM is important when it is a risk of under dosing, especially (i) in the infection with *Mycobacterium avium*, and the tuberculosis meningitis (Yew, 1998) (ii) in immune deficiency syndrome patients in which the gastrointestinal absorption of antibacterial drugs is reduced and when the incidence of infection by *Mycobacterium avium* is increased (Peloquin, 1997) (iii) drug interaction (with imidazole antifungal and/or pyrazinamide). The TDM is also necessary in situations where the risk of

toxic accumulation is increased particularly in liver dysfunction and renal failure (Peloquin, 2002; Yew, 1998; Peloquin *et al.*, 1996).

TDM of RIF is also useful in case of serious illness associated with gastrointestinal disease, cystic fibrosis, diabetes and in children regardless of the clinical situations. The variability in RIF's concentration and absorption due to concomitant food supported RIF TDM (Zent and Smith, 1995).

Tuberculosis is considered as a public health problem in Tunisia (affects about 17 Tunisian people /100,000 every year according to The National League against Tuberculosis and Lung Disease) and the increasing risks of toxicity with using high RIF doses in antistaphylococcal infections justify the necessity of RIF TDM.

Therefore, it was necessary to develop a sensitive, accurate and reproducible analytical method to analyse these samples. Numerous methods have been published previously for the analysis of RIF in biological fluids (plasma, serum, urine). The most common described procedures include microbiological assays (Bruzzese *et al.*, 2000), HPLC, thermospray mass spectrometry and electrospray mass spectrometry (Korfmacher *et al.*,

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1993); HPLC/tandem mass spectrometry method (Stefan *et al.*, 2011; Lorena *et al.*, 2010; Fang *et al.*, 2010; De Velde *et al.*, 2009; Song *et al.*, 2007) and HPLC methods (Deirder *et al.*, 2011; Marina *et al.*, 2010; Allanson *et al.*, 2007; Calleja *et al.*, 2004; Hemanth *et al.*, 2004; Moussa *et al.*, 2002; Walubo *et al.*, 1994; Weber *et al.*, 1983). Most of them needed relatively large volume of samples (≥ 0.5 mL) (Conte *et al.*, 2000; Walubo *et al.*, 1994; El-Yazigi and Raines, 1992; Swart and Pappis, 1992; Jamaluddin *et al.*, 1990; Ishii and Ogata, 1988; Woo *et al.*, 1987; Chan, 1986; Ratti *et al.*, 1981) and involve complex or very long extraction procedures, which notably increase the analysis time (Woo *et al.*, 1987; Ratti *et al.*, 1981). The present study was undertaken to develop and validate a simple and rapid HPLC method with UV detection, which could be useful for TDM of RIF in routine setting in patients as well as to evaluate pharmacokinetic of RIF.

Materials and Methods

Chemicals and reagents. Lyophilic Rifadine[®] was provided from Sanofi Aventis (Tunis, Tunisia). The internal standard (IS) clonazepam (Clonazepam[®]) was provided by Pharmascience (Montreal, Canada). Rifamycin[®]1000 000UI/100 (887 IU/mg) was provided by Thea (France).

Acetonitrile HPLC grade, Methanol HPLC grade, Water HPLC grade, K₂HPO₄ buffer HPLC grade and Diethylamine HPLC grade were obtained from Scharlau (Tunis, Tunisia). Orthophosphoric acid H₃PO₄ was provided by Prolabo (Tunis, Tunisia). Bidistilled water was prepared in the Laboratory of Clinical Pharmacology of the National Centre of Pharmacovigilance (Tunis, Tunisia).

Preparation of solutions. RIF solution SRF1 (60 mg/mL) was prepared from dilution of 600 mg of lyophilic RIF powder dissolved in 10 mL of water for injection and stored at -20 °C (for maximum 6 months). A RIF solution SRF2 (600 µg/mL) was prepared by diluting of SRF1 in the bidistilled water. Clonazepam solution (125 µg/mL) was prepared in methanol and stored at -20 °C in dark.

Two plasmatic solutions of RIF PRF1 (100 µg/mL) and PRF2 (12 µg/mL) were prepared by diluting in free human plasma. These solutions were freshly prepared before each experiment. They were used for preparation of standards (STD) of RIF (1, 2, 3, 5, 6, 9, 10, 12, 20 and 40 µg/mL) used for validation procedure by diluting

in blank human plasma. Blank human plasma for analytical development and validation was obtained from Blood Bank (HUC Charles Nicolle, Tunisia).

HPLC equipment. The HPLC system manufactured by Varian[®] (Australia) was used for the present study. For the data acquisition and integration, Galaxie chromatography data system version 1.9.3.2 software operated by Pentium 4HT was used.

Chromatography conditions. RIF was separated using a LiChocart cartridge column (150 × 4.6 mm) filled with Lichrospher 100 C18, having a 5 µm practical size (Merck, France) maintained at 25 °C. The samples were eluted in a mobile phase consisting of 0.05 M dipotassic hydrogen phosphate buffer and acetonitrile (53/47, V/V) with 0.086% diethylamin, pH = 4.46 at a flow rate of 0.6 mL/min. The mobile phase was filtered through a 0.22 µm filter (Millipore) and degassed under vacuum prior to use. The detector wavelength was set at 340 nm.

Sample extraction. Aliquots of 100 µL plasma samples (blank, standard or patient sample) and 20 µL of IS solution were vortexed for 1 min. Methanol 200 µL was added to the mixture, re-vortexed for 3 min and let stand for 10 min in dark. Each sample was centrifuged at 4 °C (4000 g × 12 min), and then 50 µL of the supernatant was injected into the HPLC column for quantification.

Method validation. Validation study was performed according to two references; (a) US Food Drug Administration (US FDA, 2001) guidelines for bioanalytical methods for evaluation of precision, accuracy, recovery and linearity and (b) US Pharmacopoeia (USP, 1999) for the determination of limit of detection and limit of quantification.

Linearity. The calibration curves were prepared (n=6) with seven increasing concentrations of RIF (STD; µg/mL). The final concentrations of RIF in plasma samples were 1, 3, 6, 9, 12, 20 and 40 µg/mL. They were built with the peak areas ratios of each STD and IS, and fitted using linear regression.

Precision and accuracy. Replicate analysis (n=5) of STDs at the 4 concentrations (2, 5, 10 and 20 µg/mL) was used for the precision and accuracy determination. All four concentrations were chosen to encompass the whole range of the calibration curve corresponding to the drug levels frequent in the majority of patients samples. Inter-assay accuracy and precision were determined by repeated analysis performed in three different

days. The concentration in each sample was determined using calibration curve prepared on the same day. The precision was calculated as the coefficient of variation (CV %) within a single run (intraday) and between different assays in different days (interdays). The analytical series were considered valid and accepted only if the percentage of deviation (bias) between theoretical and back-calculated (experimental) concentrations for each calibration level samples were less than 10%.

Accuracy was measured as the percentage difference between measured value and nominal value, according the equation: Bias (%) = $[(C_{\text{measured}} - C_{\text{theoretical}})/C_{\text{theoretical}}] \times 100\%$, where C_{measured} and $C_{\text{theoretical}}$ are the determined (observed) concentration of RIF with the present method and their theoretical concentrations in the spiked plasma samples, respectively. Extractions were performed 3 times for each sample.

Determination of limit of detection and limit of quantification (LOQ). Limit of detection (LOD) was calculated using the equation $\text{LOD} = 3.3(\text{SD}/b)$ and limit of quantification (LOQ) was calculated using the equation $\text{LOQ} = 10(\text{SD}/b)$, where, SD is the standard deviation and b is the slope of the calibration curve (US pharmacopoeia).

The aim was to make the bias and CV % less than 10% for all quality control standards except for the LOQ where less than 20% was accepted.

Statistical analysis. Validation data and the equation of the calibration curve were calculated by regression analysis using the software-package Pythagore Biostat (Version 2.0). Data were given as mean \pm SD.

Clinical application. The method was successfully applied to the monitoring of RIF in 28 blood samples provided from patients treated by RIF for tuberculosis or staphylococcal infection diseases. Samples were accompanied with patient's clinical information (age, body weight, sex, indication, RIF's dose).

Samples (3 mL) were collected in tubes containing anticoagulant (Li heparin) 2 h (time corresponding to maximum concentration) after taking RIF oral dose. Plasma was obtained after blood samples centrifugation at $1400 \text{ g} \times 10 \text{ min}$ at 4°C and stored until HPLC analysis at -20°C .

Results and Discussion

Method development. An isocratic method was used. Different ratios, flow-rates, pHs of mobile phase and

column temperatures were studied in order to shorten the retention time of RIF and to improve peak symmetry. However, the optimal assay conditions were found when using ratio of 53/47, V/V (buffer/ACN), and 0.086% of diethylamin. The pH was adjusted to 4.46 with orthophosphoric acid, the column temperature set at 25°C and the flow-rate at 0.6 mL/min. The peaks were well resolved with λ_{max} at 340 nm. Retention time of RIF in these conditions was $5.2 \pm 0.05 \text{ min}$.

The choice of the internal standard (IS) is a crucial step in the method development, as observed in this study to quantify RIF with an internal calibration technique. The internal standard must be added at the beginning of the extraction and should follow the drug in all stages of extraction.

It must be well separated from the peak of RIF, having a retention time close to the molecule and a total analysis time as short as possible. Several internal standards are described in the literature such as papaverine hydrochloride (Allanson *et al.*, 2007), p-dimethyl (Chan, 1986), rifapentine (Panchagnula *et al.*, 1999) and sulindac (Lau *et al.*, 1996). Two molecules were tried in this study; Clonazepam ($5 \mu\text{g/mL}$) and Rifamycin ($11.27 \mu\text{g/mL}$) (derivative of Rifampicin) and these were detected at retention time of 6.46 and 5.47 min, respectively. Rifamycin was ruled out because there were impurities accompanying the peak of Rifamycin and Clonazepam was kept as internal standard at the end of this step. The total time for each sample injection was about 10 min. Retention times were approximately $5.2 \pm 0.05 \text{ min}$ for RIF and 6.3 ± 0.05 for IS for all assays.

Two extraction methods have been described for RIF in literature, a solid-liquid (Aparicio *et al.*, 2006; Conte *et al.*, 2000; Walubo *et al.*, 1994; El-Yazigi and Raines, 1992; Swart and Pappis, 1992; Jamaluddin *et al.*, 1990; Ishii and Ogata, 1988; Woo *et al.*, 1987) and a liquid-liquid (Lau *et al.*, 1996; Chan, 1986). Several extraction solvents have been described; hydrochloric acid 0.1 M (Lau *et al.*, 1996), acetonitrile (Deirder *et al.*, 2011; Marina *et al.*, 2010; Allanson *et al.*, 2007; Aparicio *et al.*, 2006; Calleja *et al.*, 2004; Hemanth *et al.*, 2004; Conte *et al.*, 2000; Walubo *et al.*, 1994; El-Yazigi *et al.*, 1992; Swart and Pappis, 1992; Jamaluddin *et al.*, 1990; Ishii and Ogata, 1988; Woo *et al.*, 1987; Ratti *et al.*, 1981). A precipitation of plasmatic proteins was carried out in this study using methanol.

Validation. The specificity of the method was examined by comparing the chromatogram of the blank human

plasma with that of the plasma spiked with RIF. Representative chromatogram of blank plasma extracted of RIF and representative chromatogram of RIF's standard plasma (6 μg) and Clonazepam are shown in Fig. 1A-B. There were no interfering peaks in blank plasma at the retention time of RIF. No carry-over (memory effect) problem was observed in this assay as 50 μL of mobile phase was injected into the HPLC system after analyzing plasma containing RIF.

Isoniazid, pyrazinamide and ethambutol, which are often co-administered with RIF, did not interfere with the chromatograms.

Mean regression coefficient (r^2) of all RIF calibration curves was 0.9914 ± 0.008 ($n=6$) (data not shown). So the method was linear in concentration range 1 to 40 $\mu\text{g}/\text{mL}$.

The precision and accuracy of data obtained from controls have been presented in Table 1. Intra-day within day CV % of the assay varied from 1.99% at 10 $\mu\text{g}/\text{mL}$ to 8.86% at 2 $\mu\text{g}/\text{mL}$. Intra-day accuracy of estimation in different standards samples was found to be -16.30% for 2 $\mu\text{g}/\text{mL}$, -7.04% for 5 $\mu\text{g}/\text{mL}$, 1.40% for 10 $\mu\text{g}/\text{mL}$ and -1.78% for 20 $\mu\text{g}/\text{mL}$. Inter assay between day coefficient of variation was 2.78% for 20 $\mu\text{g}/\text{mL}$ and 2.95% for 2 $\mu\text{g}/\text{mL}$. The inter-day accuracy of estimation in the seeded samples varied from -1.66% to 2.1% for

20 $\mu\text{g}/\text{mL}$ and 10 $\mu\text{g}/\text{mL}$, respectively. These results suggest good precision and accuracy of the method.

The recovery of RIF was assessed by comparing the response of five replicates of extracted samples (2, 5, 10 and 20 $\mu\text{g}/\text{mL}$) to the response of pure standard at the same concentration level. The standard specimens were diluted in bidistilled water and injected directly

Table 1. Precision and accuracy of RIF in human plasma

Added RIF concentration ($\mu\text{g}/\text{mL}$)	Observed RIF concentration (Mean \pm SD; $\mu\text{g}/\text{mL}$)	Accuracy (%) Bias	Precision (%) CV%
Intra-day*			
2	1.67 \pm 0.14	-16.30	8.86
5	4.64 \pm 0.2	-7.04	4.5
10	10.04 \pm 0.2	1.40	1.99
20	19.64 \pm 0.59	-1.78	3.01
Inter-day**			
2	1.99 \pm 0.05	-0.25	2.95
5	5.02 \pm 0.18	0.46	3.61
10	10.21 \pm 0.51	2.1	4.9
20	19.66 \pm 0.54	-1.66	2.78

* $n=5$ assays in the same day for each concentration; ** $n=5$ assays per day during 3 days for each concentration. Daily (intra-day and day to day (inter-day) precisions are represented as mean values \pm SD of five different assays for each concentration.

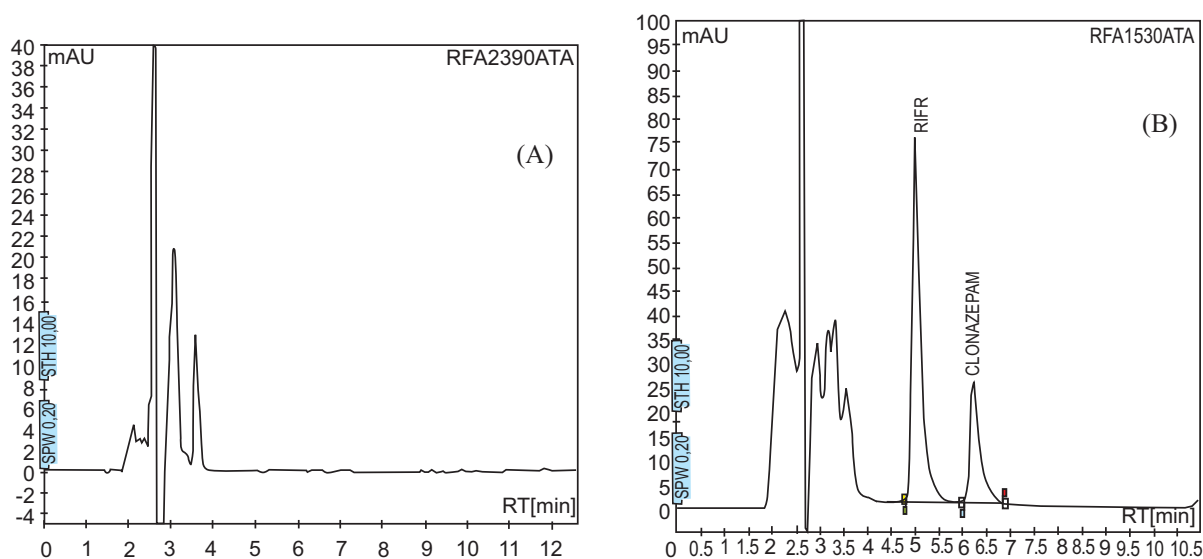


Fig. 1A-B. Representative chromatograms of blank and RIF spiked plasma (A) representative chromatogram of blank plasma extracted of RIF (B) representative chromatogram of RIF standard plasma (STD 6 $\mu\text{g}/\text{mL}$) extracted.

into the HPLC system. The recovery of RIF was greater than 90% (data not shown) for all tested concentrations.

The LOD and LOQ calculated were 0.208 $\mu\text{g/mL}$ and 0.632 $\mu\text{g/mL}$, respectively.

Application of the method. As shown by various chromatograms (chromatogram for the blank sample and chromatogram for 18.7 $\mu\text{g/mL}$ of RIF in plasma from a patient treated for a staphylococcal infection) with 900 mg/day of RIF (Fig. 2) the present HPLC/UV method seems sufficiently sensitive to be used for maximal concentration (C_{max}) RIF's plasma level determination. Plasma monitoring of RIF was realized in 28 patients (28 samples). All samples were performed 2 h after taking RIF (C_{max}).

RIF was prescribed for tuberculosis in 17 patients whose average age was 36.7 ± 15.5 yrs and average weight was 73.4 ± 18.47 kg and for staphylococcal infection in 11 patients whose average age was 59.88 ± 19.4 yrs and average weight was 67.83 ± 9.21 kg. RIF doses were varied from 5.36 mg/kg/day to 12.5 mg/kg/day, with a mean of 8.83 mg/kg/day for two indications. Found concentrations varied from 1.97 $\mu\text{g/mL}$ to 28 $\mu\text{g/mL}$ with a mean of 9.81 $\mu\text{g/mL}$. In literature target C_{max} varied from 4 $\mu\text{g/mL}$ to 24 $\mu\text{g/mL}$ according to the germ, way of action and the dose. A target concentration of 8 $\mu\text{g/mL}$ was desirable. In our population four concentrations were observed below 4 $\mu\text{g/mL}$,

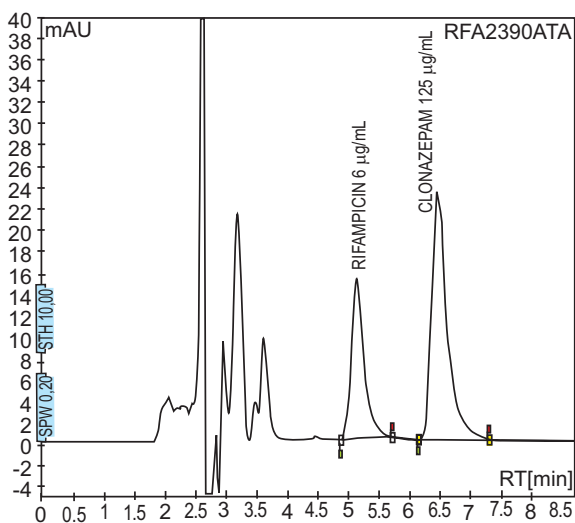


Fig. 2. Representative chromatograms for T_{2h} 10.2 $\mu\text{g/mL}$ of RIF in plasma from a staphylococcal infection patient treated with 900 mg/day of RIF.

12 concentrations between 4 $\mu\text{g/mL}$ and 8 $\mu\text{g/mL}$, and 11 concentrations were up to 8 $\mu\text{g/mL}$.

Conclusion

A novel validated method for the quantification of RIF in human plasma was developed using HPLC with UV detection. The assay is accurate, precise and fast. This simple separation method uses isocratic mobile phase conditions making the assay accessible to any laboratory in institute.

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