Simplified Liquid-Liquid Extraction Method Coupled with LC/DAD by Quantitative Determination of Risperidone in Human Serum

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ABSTRACT

Risperidone, an antipsychotic drug, was determined in human serum, using an optimized liquid-liquid extraction for isolation from biological fluid, with ethyl acetate as extraction solvent. This sample extraction method was chosen to increase extraction efficiency and reduce matrix interferences. Quantitation was achieved by LC/DAD at 294 nm over the concentration range of 1 ng/mL to 60 ng/mL (r=0.999). Desipramine was used as internal standard (tR for Risperidone: 4.23; tR Desipramine: 7.31). Mobile phase was water: acetonitrile: triethilamine: glacial acetic acid (70:29.16:0.52:0.33 v/v), using a Purosphere STAR RP-18e 250 x 4.5 mm (5µ) column. RSD, for the intra-assay study, was between 0.52 % - 2.09 % (n=3), and between 0.79 % - 3.25 % (n=9) for the inter-assay. LOD was 0.40 ng/mL and LOQ was 0.87 ng/mL. Recovery percentage for the accuracy study was between 90.82% and 98.56% ((RSD ≤ 3.93).

The method is simple, fast, precise, accurate, sensible and selective. This method was successfully applied to quantify risperidone in patient serum samples. In conclusion, the method is useful for quantitative determination of risperidone in human serum.

Key words: Risperidone, human serum, liquid chromatography, quantitative determination.

1. INTRODUCTION

Risperidone (Fig. 1) is an antipsychotic drug used for the symptomatic management of psychotic disorders, especially in patients with schizophrenia, and, generally, is required for long-term stabilization to sustain symptom remission or control and to minimize the risk of relapse.1-2

![Chemical structure of risperidone](image)

**Figure 1:** Chemical structure of risperidone.

Risperidone is administered orally or by intramuscular injection. About 90% of patients receive the drug orally. Oral administration can be in a once-daily dose or in two equally divided doses daily, with a target dosage of 4 mg daily in most of patients. After normalization to a dose of 4 mg/day, median concentrations had been of 9 ng/mL (80% range, 0.9-27.9 ng/mL) for the parent compound and 24.1 ng/mL (80% range, 12.0-57.6 ng/mL) for the metabolite, 9-hydroxy-risperidone.3-4

Although antipsychotic drugs have been effective in reducing symptoms of schizophrenia, issues with adherence to these agents continue to be a barrier to the implementation and delivery of a successful treatment plan. Non-adherence lies on a spectrum, is often covert, and is underestimated by clinicians, but affects between 50 %- 60 % of patients with schizophrenia. It increases the risk of relapse, re-hospitalization, self-harm and inpatient costs, leads to lower quality of life, greater use of emergency psychiatric services, functional decline, and increased risk of death. Recognizing that all patients with schizophrenia are at risk for medication non-adherence is important. No one technique has been shown to be most effective; therefore, the risk for non-adherence should continually be assessed.5-9

There are some indirect ways to evaluate the pharmacotherapy adherence of the patients, but these are subjective, depending on confident of patient to response with trustiness.10

The determination of drug levels in biological fluids provides objective information. Therefore, drug monitoring of patients biological fluids can be a useful tool to evaluate adherence by relating the serum or plasma levels of drugs with pharmacotherapy compliance.10-11

For determination of risperidone in blood, some HPLC12-14 and spectrophotometric methods have been reported. All of these methods used human plasma.

In view of the clinical and analytical challenge, the present study describes the development of environment friendly, simple, rapid, sensitive, accurate and precise LC/DAD method. Among all different detectors used to analyze risperidone (evaporative light scattering detector: ELSD, refractive index detector: RID, UV, MS), UV-DAD is simplest and suitable for the purpose of this work. Moreover, serum instead plasma as blood sample allows a simple extraction of analyte from matrix.

Therefore, the aim of this study was to develop and validate a simple, rapid sensitive, accurate and precise LC/DAD method for quantification of risperidone in human serum, suitable for pharmacotherapy adherence studies.

2. EXPERIMENTAL

2.1. Chemicals and reagents

USP standards of risperidone and desipramine were purchased from Sigma-Aldrich, St. Louis, MO. Methanol, acetonitrile, triethylamine, glacial acetic acid and ethyl acetate, were obtained from Merck, Darmstadt, Germany. All of the reagents were pro-analysis quality.

2.2. Preparation of standard solutions

Stock solutions of 1 mg/mL of risperidone were prepared by dissolving in methanol. Different concentrations required for preparation of spiking serum calibration and quality control samples were subsequently prepared, to obtain working solutions of 5, 30 and 60 ng/mL. All of these solutions were stored at 2-8°C for about two days.

2.3. Spiking procedure for calibration and quality control (QC) samples

Calibrator samples were prepared just prior to extraction by spiking 2 mL of pooled human blank serum with 1 mL of a convenient working solution of fluoxetine in methanol, to give concentrations of 1, 5, 20, 30, 45 and 60 ng/mL. Quality control samples were prepared at concentrations of 5, 30 and 60 ng/mL, and used to determine the intra and inter-assay precision and accuracy of the method.

Drug-free serum used for the validation of the method was obtained from healthy volunteers, with informed consent.
2.4. Sample preparation
Human blank serum and human serum samples were taken out from deep freezer maintained at -20 °C, and kept at room temperature for 45 minutes for thawing. Then, 2 mL of sample was processed by adding initially 25 µL of desipramine (internal standard) (0.5 ng/mL) and 3 mL of ethyl acetate to the solution, which was subsequently vortexed for 5 min and then centrifuged at 13200 rpm for 5 min. The supernatant was transferred to an Eppendorf tube and evaporated to dryness at 80 min at 45°C. The dry residue was reconstituted in 200 µL of mobile phase, and a 20 µL mL aliquot of this solution injected for analysis.

2.5. Instrumentation and chromatographic conditions
HPLC analyses were carried out on a Purosphere STAR RP-18e 250 x 4.5 mm column (5µ) (Darmstadt, Germany).

Separation was achieved using a mobile phase consisting of water: acetonitrile: triethylamine: glacial acetic acid (70:29.16:0.52: 0.33 v/v), at pH= 5.5 and a flow rate of 1 mL/min. The eluted compounds were monitored at 294 nm. The column was maintained at ambient temperature, and an injection volume of 20 µL was used. Chromatographic run was 10 minutes.

The mobile phase was filtered through 0.45 micron membrane filter and ultrasonicated for 10 minutes prior to use. Peak homogeneity was expressed in terms of peak purity values and was obtained directly from spectral analysis report obtained using the instrument software.

2.6. Application of the method
This assay was carried out in accordance with Ethical Committee of the University, and all of patients signed informed consent forms. Blood of ten patient volunteers who were in treatment with risperidone (4 mg daily), were used for quantification of the drug from serum. The collection of the samples was done in the morning, before the next administration of the drug.

Serum was separated from plasma by heating to 37 °C and centrifugation of blood samples at 3000 rpm for ten minutes. The volume of serum used for experiment was 1 mL.

2. RESULTS AND DISCUSSION

3.1. Method optimization
Different conditions for sample extraction of risperidone and chromatographic conditions was tried to achieve optimal results. A Design of Experiments was performed (3-level factorial design: 3^3) to optimize the chromatographic parameters, choosing as study parameters the mobile phase flow, pH of mobile phase, and the column temperature. A pool of standard risperidone extracts with concentration of 30 µg/mL was used. By Statgraphics ® Software, area and height of chromatographic peaks, as its purity were the more important parameters. Quantifications were done in triplicate, by LC/DAD.

The selection of the mobile phase was carried out on the basis of polarity i.e., with a satisfactory separation of risperidone and the internal standard desipramine with good peak symmetry (system suitability test). Moreover, it allows a fast running time of 7 minutes.

Several trials for optimization of the mobile phase were taken and finalized as water: acetonitrile: triethylamine: glacial acetic acid (70:29.16:0.52: 0.33 v/v).

Several wavelengths were tried, chosen 294 nm as working wavelength. Complete resolution of the peaks with clear baseline separation was obtained of this way.

Sample extraction was optimized to eliminate the laborious extraction steps, with minimal losses of risperidone and very good recoveries from spiked serum samples and from real samples.

Optimization of chromatographic parameters for human serum as matrix: the optimal conditions found taking as response the peak purity, and the optimal chromatographic conditions: mobile phase flow 1.0 mL/min, water phase pH: 5.5, and column temperature: 25 °C.

3.2. Calibration curves
Calibration curves were constructed for the range of 1-60 ng/mL. Each solution (1.5, 20, 30, 45 and 60 ng/mL) was injected three times. This range of solution concentrations include the risperidone concentrations expected in serum: 2.9 ng/mL (80 % range: 0.9-27.9 ng/mL) a.

Standard curves were constructed by plotting the peak area ratio of the analyte to the internal standard as a function of the concentration added.

The mean equation (curve coefficients ± standard deviation) for the calibration curve (n=5), obtained from five points, was

\[ y = 0.0074x + 0.067, \text{ with a correlation coefficient, } r = 0.999. \]

3.3. Precision and accuracy
The intra-assay precision of the assay was measured by the percent coefficient of variation (RSD) over the concentration range of low (L), medium (M), and high (H) quality control levels (1-30-60 ng/mL) for three replicates, each in the same analytical run (each replicate was spotted three times), and intra-assay accuracy was measured from percentage recoveries obtained by the analyzes of the quality control samples, determined by linear regression equation of peak area vs. drug concentration. Inter-assay precision and inter-assay accuracy were calculated after repeated analysis in three different analytical runs. Each experiment included the sample extraction step.

Accuracy values were between 90.82 % and 98.56 % of recovery percentage.

Precision and accuracy did not exceed 3.93 % of RSD at any level. The results are presented in Table 1 and Table 2.

\[ \text{Table 1. Precision of the method.} \]
\[
\begin{array}{l|c|c}
\text{Concentration (ng/mL)} & \text{Intra-assay}^a & \text{Inter-assay}^b \\
1.00 & 2.09 & 3.25 \\
30.00 & 1.59 & 1.77 \\
60.00 & 0.52 & 0.79 \\
\end{array}
\]

\[ a_n = 3; \text{analyzed on the same day (three solutions of each concentration)} \]

\[ b_n = 9; \text{analyzed on three different days (three solutions of each concentration prepared for 3 days)} \]

\[ \text{Table 2. Method accuracy.} \]
\[
\begin{array}{l|c|c|c}
\text{Actual concentration (ng/mL)} & \text{Measured concentration (ng/mL)} & \text{Accuracy, %} & \text{RSD, %} \\
1.00 & 0.97 \pm 0.04 & 90.82 & 3.93 \\
30.00 & 29.46 \pm 0.71 & 98.44 & 2.44 \\
60.00 & 61.05 \pm 1.11 & 98.56 & 1.82 \\
\end{array}
\]

3.4. Detection and quantification limits
The limit of detection (LOD) and limit of quantification (LOQ) were calculated preparing solutions at three concentrations (0.2-0.5-1.0 ng/mL) in the lower range of linear regression curve.

LOD was 0.40 ng/mL and LOQ was 0.87 ng/mL, determined using the equations: LOD = 3.3 σ/b; LOQ = 10 σ/b, where σ is the standard deviation of the responses, and “b” corresponds to the slope obtained from the curve peak area ratio of the analyte to the internal standard versus concentration of the analyte. These values were experimentally verified applying the method to prepared solution with these analyte concentrations.

3.5. Selectivity
The method selectivity was demonstrated on three blank serum samples obtained from healthy volunteers; the chromatograms were found to be free of interfering peaks. Risperidone and the internal standard desipramine were well resolved, with IR = 3.82 for risperidone, and 7.55 for desipramine (Fig. 2). Selectivity between risperidone and its major metabolite was not assay because the aim of the study was obtained a LC/DAD method useful to evaluate pharmacotherapy adherence, therefore, it is not important that the parent drug is well resolved from its metabolites.

3.6. Extraction recovery
The extraction recovery of risperidone from serum and that of the internal standard were quantified using the concentrations of 1, 30 and 60 ng/mL for the drug and concentration of 0.5 ng/mL for the internal standard. The extraction recoveries were calculated by comparing the observed concentrations obtained from the processed standard samples to the concentrations obtained from the standards solutions added to the serum after the extraction, which represented 100 % recovery. The extraction recovery of risperidone from serum ranged from 90.5 % to 96.5 %. The internal standard extraction recoveries were found between 91.5 % and 95.8 %.

3.7. Application of the method
The risperidone concentration ranged between 1.52 ng/mL to 91 ng/mL.
Risperidone plasma concentrations had been of 2.9 ng/mL (80% range, 0.9-27.9 ng/mL) in chronic use, therefore, all of values found in the samples studied were at therapeutic range.

**Figure 2:** Selectivity between risperidone and the internal standard desipramine. Peak nº1: risperidone; peak nº2: desipramine.

### 4.- CONCLUDING REMARKS

The developed LC/DAD method is simple, rapid, precise, sensible, selective and accurate for quantitative determination of risperidone in human serum. The method works with a simple liquid-liquid extraction step and it was designed to cover the usual serum concentration level of patients taking the drug.

The most significant advantages of the present LC/DAD method, as difference of other analytical methods founded in bibliography to quantify risperidone in biological fluids, are the small sample volume needed, few running time, few injection volume, and better sensitivity and selectivity. Among all different detectors used to analyze risperidone (ELSD, RID, UV, MS, etc), UV-DAD is simplest and suitable for the purpose of this work. Moreover, serum instead plasma as blood sample allows an easy and fast extraction of analyte from matrix. The limit of quantification and recovery are more than adequate for use in pharmacokinetic studies. Our results indicate that this LC/DAD procedure is a reproducible method that provides consistent quantification of risperidone in human serum.

Therefore, the developed LC/DAD method is useful for quantification of risperidone in human serum, suitable for pharmacotherapy adherence studies.

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