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# Quantification of Ticlopidine in Human Plasma Using Protein Precipitation and Liquid Chromatography Coupled with Tandem Mass Spectrometry

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## Abstract

A simple and rapid method has been developed and validated for determination of ticlopidine in human plasma using protein precipitation and liquid chromatography coupled with tandem mass spectrometry (LC/MS/MS). Ticlopidine was extracted from 20- $\mu$ L aliquots of human plasma by one-step protein precipitation with 980  $\mu$ L of acetonitrile containing 10 ng/mL clopidogrel as an internal standard (IS). Chromatographic separation was performed on a reverse-phase Gemini C<sub>18</sub> column (50 mm x 2.0 mm, 5  $\mu$ m) with an isocratic mobile phase (acetonitrile: 1 mM ammonium acetate in water = 75:25, v/v). Ticlopidine and IS were detected and quantified by tandem mass spectrometry with positive electrospray ionization using multiple reaction monitoring of the transition m/z 264.04 to m/z 154.20 for ticlopidine and m/z 322.40 to m/z 212.20 for IS. This method was linear over the concentrations ranging from 2 to 2000 ng/mL. The accuracy in the inter-batch assay was 92.4-95.6% and the precision was within 6.4% coefficient of variation. The validated method was successfully applied to the human pharmacokinetic study of ticlopidine.

**Keywords:** Ticlopidine; Human plasma; Protein precipitation; LC/MS/MS; Validation

## Introduction

Ticlopidine (5-[ (2-chlorophenyl) methyl] -4, 5, 6, 7 - tetrahydrothieno [3, 2-c] pyridine hydrochloride), an inhibitor of platelet aggregation, is used therapeutically for the prevention of stroke and myocardial infarction in high-risk patients [1-3]. Its mechanism of action involves inhibiting the binding of adenosine diphosphate (ADP) to receptor P2Y<sub>12</sub> on the platelet membrane and modifying glycoprotein IIb/IIIa, which inhibits the binding of fibrinogen, resulting in the inhibition of platelet aggregation [4].

Several methods have been described for the quantification of ticlopidine in biological fluid, including gas chromatography coupled with mass spectrometry [5-7], high performance liquid chromatography (HPLC) with ultraviolet light [8, 9], HPLC coupled with mass spectrometry (LC/MS) [10, 11] and HPLC coupled with tandem mass spectrometry (HPLC/MS/MS)[12]. The previous methods using HPLC coupled with mass spectrometry employed manual solid-phase extraction (SPE) [10] or liquid-liquid extraction [11, 12]. The solid-phase extraction often involves multi-step purification and evaporation, and a plasma volume of at least 0.5 to 1 mL plasma is usually required. The liquid-liquid extraction often needs high-purity organic solvents, is troublesome to handle, and requires an evaporation step. It can be expensive and time-consuming.

In the present study, we developed and validated a high throughput, selective HPLC/MS/MS method for the determination of ticlopidine in human plasma with a micro-sample volume using one-step protein precipitation.

## Experimental Procedure

Other than the sample preparation method, we referred to the previously reported HPLC/MS/MS method by Borges et al. [12], and modified it slightly as follows.

## Materials

Ticlopidine hydrochloride (Lot No. WST 06016; purity, > 98%) was provided by Yuyu Pharma. Inc. (Seoul, Korea). (+)-Clopidogrel hydrogen sulfate (purity, > 98%), an internal standard (IS), was purchased from Toronto Research Chemicals Inc. (North York, Canada). Ammonium acetate was purchased from Sigma-Aldrich (St. Louis, MO, USA), and HPLC grade acetonitrile and methanol were purchased from Fischer Scientific (Fair Lawn, NJ, USA). A Milli-Q<sup>®</sup> (Millipore Co, Milford, MA, USA) water purification system was used to obtain the purified water for the analysis. Blank human plasma was provided by the Blood Bank in Seoul National University Hospital and stored at approximately -70°C until used.

## Chromatographic conditions

A Symbiosis Pharma HPLC system (Spark Holland, Emmen, Netherlands) consisting of a binary solvent pump, degasser, and autosampler was used for setting the reverse-phase liquid chromatographic conditions. The chromatographic separation was achieved on a Gemini C<sub>18</sub> column (50 x 2.0 mm, 5 $\mu$ m; Phenomenex, Torrance, CA, USA) with a Gemini C<sub>18</sub> guard column (4 x 2.0 mm, 5 $\mu$ m; Phenomenex) at room temperature. The isocratic mobile phase consisted of acetonitrile and 1 mM ammonium acetate in water (75:25, v/v) and eluted at a flow rate of 0.30 mL/min. The temperature of the autosampler was maintained at 4°C.

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## Mass spectrometric conditions

Mass spectrometric detection employed an API 4000QTRAP (Applied Biosystems/MDS SCIEX, Foster City, CA, USA) equipped with an electrospray ionization (ESI) source operated in the positive ion mode. The curtain, nebulizer, and turbo gas (nitrogen) were set at 10, 40, and 50 psi, respectively. The ion spray voltage was adjusted to 5500 V and the source temperature was set at 600°C. The mass spectrometer was operated in the multiple reaction monitoring (MRM) mode at unit resolution for both Q1 and Q3 with a dwell time of 200 ms per MRM channel. The transition of the protonated precursor to the product ion was monitored at  $m/z$  264.04 to  $m/z$  154.20 for ticlopidine and  $m/z$  322.40 to  $m/z$  212.20 for clopidogrel. Declustering potentials were set at 71 and 40 eV for ticlopidine and clopidogrel, respectively, and collision energies were set at 25 and 23 eV, respectively. Data acquisition and integration were processed by Analyst version 1.4.1 software (Applied Biosystems).

## Preparation of calibration standards and quality control samples

Stock solutions of ticlopidine and clopidogrel (IS) were prepared by dissolving 1 mg/mL in methanol-water solution (50:50, v/v) and stored at 4°C. Ticlopidine stock solution was diluted serially with 50% methanol to obtain working standard solutions (0.04, 0.1, 0.4, 1, 4, 10, 40, 100 µg/mL). To prepare calibration standards, 50 µL of each working standard solution was spiked to 950 µL of blank human plasma, yielding final concentrations of 2, 5, 20, 50, 200, 500, and 2000 ng/mL. The concentrations of ticlopidine and clopidogrel were calculated in the salt free form. The lower limit of quantification (LLOQ, 2 ng/mL) for ticlopidine in human plasma was defined as the lowest concentration with a signal-to-noise ratio over 10 and acceptable accuracy (80-120%) and precision (< 20%).

Quality control (QC) samples at low, medium, and high concentration (6, 100, and 1600 ng/mL, respectively) were prepared by spiking the working standard solutions of ticlopidine into blank human plasma, and aliquots (100 µL) were stored in the freezer below -70°C until analysis. The IS stock solution was diluted with acetonitrile to a final concentration of 10 ng/mL. The calibration standards and QC samples were extracted for each analytical batch with the unknown samples.

## Sample preparation

All frozen human plasma samples were thawed at room temperature. Twenty microliters of each unknown samples, standards and QC samples were transferred to 1.5 mL micro-centrifuge tubes, and 980 µL of acetonitrile containing IS (10 ng/mL) were added. For the blank sample, 20 µL of blank plasma was pipetted to 980 µL of 100% acetonitrile. Then, the tubes were vortexed for 1 min and followed by centrifugation at 4°C, 13,000 rpm for 5 min to the precipitate protein. A 200-µL aliquot of each supernatant was transferred into the injection vial and a 2-µL aliquot was injected into the LC/MS/MS system. All prepared samples were kept at 4°C until injection.

## Method validation

**Stability:** The stability of ticlopidine in human plasma was evaluated using six replicates of low and high QC samples under three different test conditions as follows. For the short-term stability test, QC samples were kept at room temperature for 18 h before analysis. For the freeze-thaw stability test, QC samples were analyzed after 3 successive

freeze-thaw cycles. The post-preparative stability was assessed using the processed QC samples which were kept at 4°C on the autosampler for 36 h before analysis.

**Selectivity, carryover and linearity:** Selectivity was investigated by comparing six blank plasma samples from different origins with the plasma spiked with ticlopidine (at LLOQ) and IS. The carryover effect on the LC/MS/MS system was evaluated by injecting consecutively blank samples after a standard sample at the highest concentration of the calibration range.

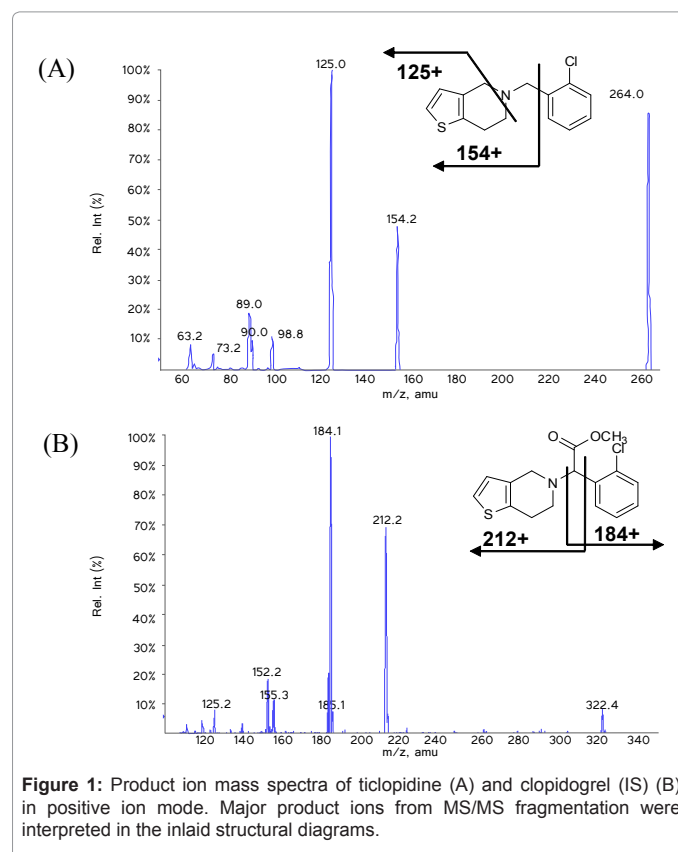
A calibration curve was prepared by 7 standard samples covering the entire range of 2-2000 ng/mL of ticlopidine. The linearity of each calibration curve was assessed by a weighted ( $1/x^2$ ) linear regression method based on the peak area ratios ( $y$ ) of ticlopidine to IS versus the nominal concentration ( $x$ ) of ticlopidine.

**Precision, accuracy and recovery:** The intra-batch precision and accuracy were assessed using 6 replicates at three different concentrations (6, 100 and 1600 ng/mL) within a batch. The inter-batch precision and accuracy were determined using 6 replicates of three different concentration levels on 5 different days. Precision was expressed as relative standard deviation,  $RSD (\%) = 100 \times (\text{standard deviation}/\text{mean})$  and accuracy was calculated as  $\text{accuracy} (\%) = 100 \times (\text{mean observed concentration}/\text{spiked concentration})$ . Recovery was calculated by comparing mean peak areas obtained from extraction of QC samples and extracted blank plasma spiked with the corresponding concentration of ticlopidine.

## Results

### Mass spectrometry and chromatography

Mass spectra of ticlopidine and clopidogrel (IS) in the positive ion mode showed the protonated molecular ions  $[M+H]^+$  as the major ion

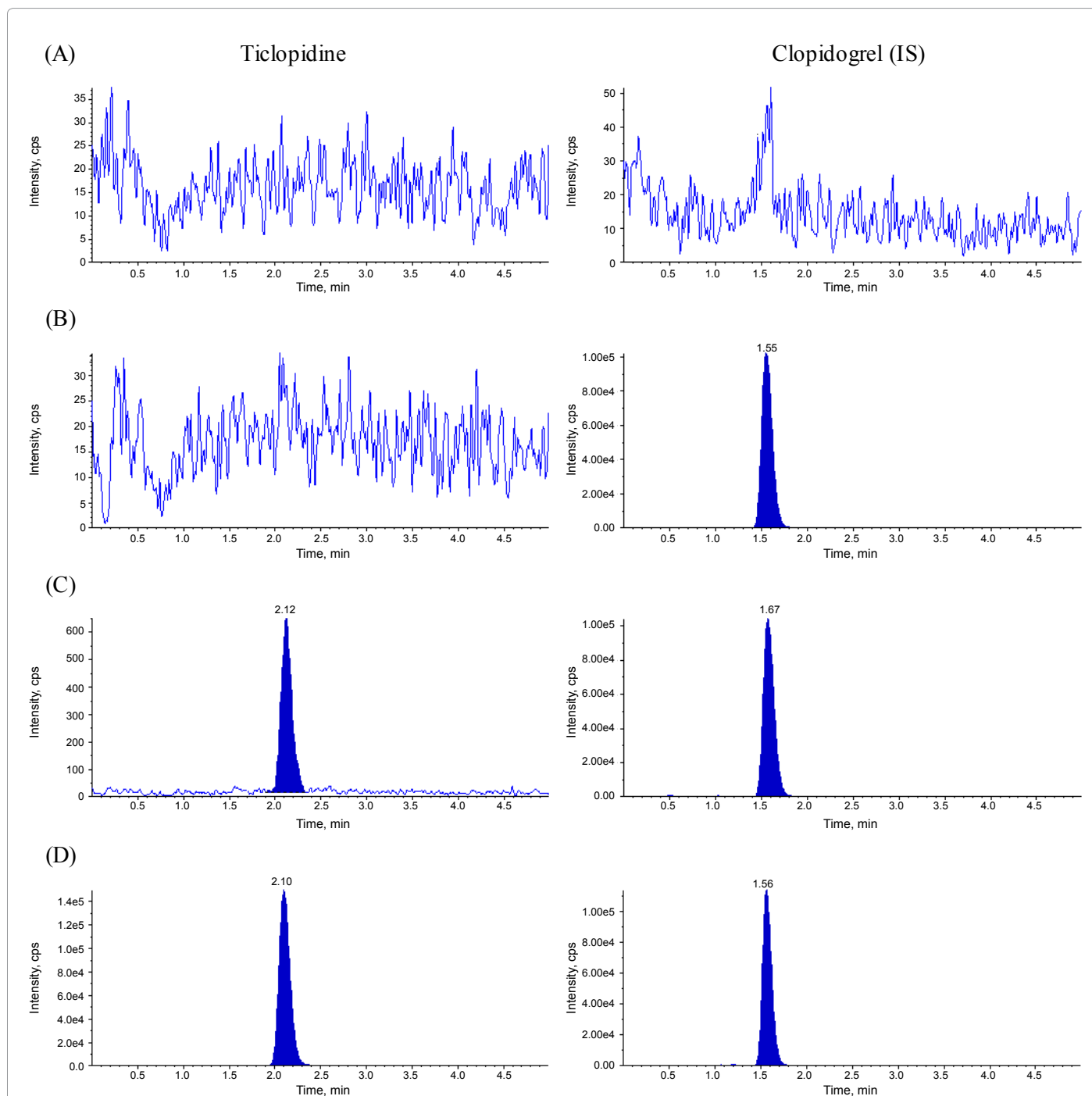


species,  $m/z$  264.04 and 322.40, respectively. The chemical structures of ticlopidine and clopidogrel with proposed fragmentation patterns are shown in Figure 1 based on the full-scan product ion spectra of  $[M+H]^+$  ions. The MS/MS parameters were optimized to maximize the response for precursor to product ion in the positive ion mode. Consequently, the mass transition was monitored at  $m/z$  264.04 to 154.20 for the analyte and  $m/z$  322.40 to 212.20 for IS. The MRM ion chromatograms were obtained after simple preparation of human plasma samples with

acetonitrile showed that total chromatographic run time was 5.0 min and retention times were 2.1 min for ticlopidine and 1.6 min for IS (Figure 2).

### Selectivity, carryover and linearity

No interfering peaks at the elution times for the analyte or IS were observed in the six blank human plasma samples from different origins. Figure 2 shows typical chromatograms for blank plasma, blank plasma with IS, plasma spiked with ticlopidine at 2 ng/mL (LLOQ), and



**Figure 2:** MRM ion chromatogram of ticlopidine ( $m/z$  264.04 to  $m/z$  154.20) and clopidogrel (IS,  $m/z$  322.40 to 212.20) in (A) a blank plasma sample, (B) a zero sample, (C) a LLOQ (2 ng/mL) sample, and (D) a plasma sample from a healthy volunteer at 1.5 hours after oral administration of ticlopidine 250 mg.

plasma obtained from a volunteer at 1.5 hours after oral administration of ticlopidine 250 mg. LLOQ of ticlopidine was 2 ng/mL, using 20  $\mu$ L of human plasma. There was no carryover effect observed; no enhancement in the response was observed in the chromatogram of a blank sample after subsequent injection of the highest calibration standard at the retention time of analyte and IS (data not shown). The calibration curves for ticlopidine were established in the range of 2-2000 ng/mL and exhibited good linearity with the correlation coefficients ( $r$ ) ranging between 0.9979 and 0.9999 in five batches.

### Precision, accuracy and recovery

The precision and accuracy of the calibration curves ranged 1.0-3.7% and 95.5-103.2%, respectively. The intra- and inter-batch precision of this method was below 2.1 and 6.4%, respectively, and the intra- and inter-batch accuracy was 101.0-102.9% and 92.4-95.6%, respectively (Table 1).

The mean recovery of ticlopidine at the 6, 100, and 1600 ng/mL levels were 102.7%, 100.8%, and 100.7%, respectively. The recovery of IS at the concentration of 10 ng/mL was 84.0%.

### Stability

The stability of ticlopidine in human plasma under various conditions is shown in Table 2. Ticlopidine was stable after short-term storage at room temperature for 18 h, three freeze-thaw cycles, or storage in an autosampler at 4°C for 36 hr, with a difference of less than 5.4% between the reference and test concentrations observed. The long-term stability test was not performed because Borges et al. reported that ticlopidine in human plasma was stable for 269 days at -20°C [12].

### Application of the method to human pharmacokinetic study

This simple and rapid method for determination of ticlopidine in human plasma was successfully applied to a single dose pharmacokinetic study after oral coadministration of ticlopidine 250 mg in 24 healthy male volunteers [13]. Blood samples (8 mL) were withdrawn in a

heparinized tube before dosing and at 0.5, 1, 1.5, 2, 2.5, 3, 4, 6, 8, 12, 24, and 48 hours after administration of ticlopidine 250 mg. Plasma was separated within 30 min after blood sampling by centrifugation at 4°C and 3,000 rpm for 10 min, and stored below -70°C until analysis. The LLOQ (2 ng/mL) of the assay has sufficient reliability to allow pharmacokinetic parameters of ticlopidine, given in therapeutic doses (250 mg), to be calculated. A representative chromatogram of a plasma sample obtained at 1.5 h from a subject who had administered a single oral dose of ticlopidine 250 mg is shown in Figure 2 (D) and the sample concentration was determined to be 569.4 ng/mL for ticlopidine. The mean plasma concentration-time profiles of 24 healthy volunteers are shown in the supplementary figure. The maximum ticlopidine concentrations in plasma ( $C_{max}$ ) values and the times taken to reach  $C_{max}$  ( $T_{max}$ ) after dosing were consistent with those of the literature [14, 15].

### Discussion

In this study, we developed a simple and rapid one-step method of protein precipitation to determine ticlopidine concentration in human plasma. Validation results of method indicate that this HPLC/MS/MS allows sensitive, selective, and high throughput quantification of ticlopidine so that it is considered suitable for the clinical pharmacokinetic studies of ticlopidine. It avoids the use of complicated and expensive manual solid phase extraction techniques and time-consuming liquid-liquid solvent extraction with evaporation that was used in previously reported procedures. In addition, this method requires only a small volume of plasma sample (20  $\mu$ L compared with 500  $\mu$ L for a solid phase extraction [10] and 200-500  $\mu$ L for a liquid-liquid extraction [11, 12]).

Compared to the previously reported HPLC/MS/MS method using liquid-liquid extraction by Borges et al., the LLOQ was slightly higher (2.0 ng/mL versus 1.0 ng/mL), and the run time was longer (5 min versus 3 min). However, the recovery of protein precipitation was almost 100% compared with 85.0%-91.8% for liquid-liquid extraction. Accuracy and precision were comparable with the previous study: intra- and inter-day accuracies were 101.0-102.9% and 92.4%-95.6% in this study, and 105.3%-111.3% and 101.8%-105.3% in the previous study, respectively; intra- and inter-day precision variations (CV%) were lower than 2.1% and 6.4% in this study, and 2.4% and 4.8% in the previous study, respectively. Moreover, in the previous study by Borges et al., short-term stability at room temperature for 8 h and autosampler stability at 12°C for 24 h were guaranteed, while those were established in this study for 18 h at room temperature and 36 h at 4°C, respectively.

### Conclusion

In conclusion, we report the development and validation of a rapid LC/MS/MS method with a simple preparation of samples for the determination of ticlopidine in human plasma over a concentration range of 2-2000 ng/mL using 20  $\mu$ L of plasma sample. This method showed excellent accuracy, precision, and stability. The developed method was successfully applied to the human pharmacokinetic study of ticlopidine.

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Nominal concentration (ng/mL)	Intra-batch (n=6)		Inter-batch (n=5)	
	Precision (RSD %)	Accuracy (%)	Precision (RSD %)	Accuracy (%)
6	2.1	102.9	6.4	95.6
100	1.6	101.0	5.4	94.0
1600	0.6	101.5	6.0	92.4

**Table 1:** Precision and accuracy for ticlopidine in human plasma.

Conditions	Nominal concentration (ng/mL)	Observed concentration (ng/mL)		Difference (%) between reference and test
		Mean of reference (n=6)	Mean of test (n=6)	
Storage of samples at room temperature for 18 hr	6	5.5	5.2	-5.4
	1600	1430.7	1363.0	-4.7
Three freeze-thaw cycles (-70°C)	6	6.0	5.8	-3.3
	1600	1500.8	1491.0	-0.7
Storage of extracted samples in autosampler at 4°C for 36 hr	6	6.2	6.3	1.6
	1600	1624.3	1612.0	-0.8

**Table 2:** Stability of ticlopidine in human plasma under various conditions.

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