

# Extraction of Polar Basic Drugs from Plasma with Polymeric SPE Cation Exchange, Bond Elut Plexa PCX

# **Application Note**

BioPharma

#### **Authors**

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#### Introduction

Basic pharmaceutical drugs are ideal for a cation exchange sorbent. Analytes are easily charged in an acidic solution and readily interact with the ion exchange function of the sorbent. Polar basic compounds can be problematic for reversed phase sorbents due to their poor hydrophobic interaction and water solubility.

Agilent Bond Elut Plexa PCX is a new addition to the Plexa family and uses a polymeric cation exchange technique. Plexa PCX uses a generic and simplified method to remove neutral and acidic interferences from the matrix and concentrate basic analytes, resulting in improved analytical performance and analytical sensitivity in the quantification of basic compounds.

In addition, Plexa PCX offers faster and highly reproducible flow rates, resulting in excellent tube-to-tube and well-to-well performance. Plexa PCX significantly reduces ion suppression because its highly polar, hydroxylated surface is entirely amide-free. The particle exterior minimizes strong binding of proteins and phospholipids. Efficient removal of phospholipids from plasma is ensured. A simple generic method was developed for the extraction of polar basic drugs in human plasma.



#### **Materials and Methods**

Table 1. SPE Reagents and Solutions

2% Phosphoric Acid	Add 20 $\mu$ L of concentrated $H_3PO_4$ to 1 mL of DI water	
Methanol	Reagent grade or better	
2% Formic Acid	Add 20 µL of concentrated formic acid to 1 mL of DI water	
Methanol:acetonitrile (1:1, v/v)	Add 1 mL of methanol to 1 mL of acetonitrile	
5% NH <sub>3</sub> Methanol:acetonitrile (1:1, v/v)	Add 50 µL of concentrated ammonia to 1 mL of methanol:acetonitrile (1:1, v/v)	

Bond Elut Plexa 10 mg 96 well plate (part number A4968010)

Table 2. SPE Method

Sample Pre-treatment	100 µL human plasma. Dilute 1:3 with 2% H <sub>3</sub> PO <sub>4</sub> .
Condition	1. 500 μL CH <sub>3</sub> OH 2. 500 μL DI H <sub>2</sub> O
Load	Sample with the drug mixture at the flow rate of 1 mL/min
Wash 1	500 μL 2% formic acid
Wash 2	500 μL acetonitrile:methanol (1:1, v/v)
Elution	500 µL 5% NH <sub>3</sub> methanol:acetonitrile

All samples are evaporated to dryness and reconstituted in 100 µL of 80:20 0.1% aqueous formic acid: CH,OH.

## **Results and Discussion**

LC Conditions

Mobile Phase: A: 0.1% Formic acid

B: Methanol

Gradient: t = 0 min80% A: 0% B  $t = 0-2 \min$ 20% A: 80% B

t = 3.5-5 min80% A: 20% B

Column: Agilent Pursuit C18 3 µm,

 $2.0 \times 50 \text{ mm}$ 

(part number A3051050X020)

MS Conditions

Transition ions and collision energy were:

Compound	<b>Q</b> 1	<b>Q</b> 3	CE				
Albuterol	240.1	148.0	-23.5V				
Lamotrignine	256.0	256.0	-5.0V				
Atenolol	267.0	145.0	-34.0V				
Sumatriptan	296.1	201.1	-14.0V				
Capillary = 25 V, Dry gas temp = 400 °C, 30 psi,							
CID = Argon							

Polarity:

Positive

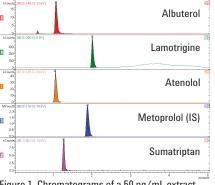


Figure 1. Chromatograms of a 50 ng/mL extract

This LC/MS method describes the quantitative determination of polar basic compounds in human plasma using Bond Elut Plexa PCX for SPE (Figure 1). The limit of detection (LOD) of the solid phase extraction and LC/MS/MS analysis was 1.0 ng/mL. Recoveries were calculated from a 2nd order regression with RSD values based on a sampling of n = 6.

Excellent recoveries were achieved, which demonstrated good retention and elution, as well as minimal ion suppression. Response for all the compounds evaluated was linear up to 3 orders of magnitude from 1.0 ng/mL to  $1.0 \mu\text{g/mL}$  with correlation coefficients all above 0.999. To demonstrate reproducibility, samples were analyzed at two different concentrations (n = 6). As shown in Table 3, reproducibly high recoveries were obtained according to the generic standard protocol.

Table 3. Recoveries of polar basic compounds from human plasma

Analyte	log P	рКа	% Rec	% RSD <sup>2</sup>	% Rec	% RSD <sup>2</sup>
			(500 ng/mL)		(1000 ng/mL)	
Sumatriptan	0.96	9.6	95	5	97	4
Atenolol	1.30	9.6	94	3	91	2
Albuterol	1.30	10.3	95	5	100	7
Lamotrigine	1.50	5.7	92	3	97	4

<sup>1</sup>Recoveries calculated as % of signal intensity of an extracted sample compared to that calibration curve. <sup>2</sup>RSD = standard deviation/average recovery x 100; n = 6.

### **Conclusions**

With Agilent Bond Elut Plexa PCX, a generic drug extraction protocol from plasma can be applied to polar analytes with basic amino functional groups. Under acidic conditions, the charged analyte binds to the cation exchange groups of the sorbent (see Table 3 for pKa). Polar interferences and proteins are washed away with an acidic, aqueous solution. A neutral wash with relatively strong solvents, such as 50% methanol:acetonitrile, is possible without any loss of analyte. The wash elutes neutral compounds retained in the hydrophobic cores of the sorbent. Finally, a mixture of organic solvents with ammonia is used to disrupt the cation exchange interaction, resulting in the elution of the basic drugs.

Flow rate all over the 96-well plate is fast because Plexa PCX particles have a much narrower particle size distribution with no fines to cause blockages. thus resulting in excellent well-to-well reproducibility. Automated 96-well technology is easily possible, which opens up new opportunities to maximize efficiency. Bond Elut Plexa PCX is therefore a useful tool for high throughput SPE applications, which require analysis at low analyte levels, validated reproducibility and quick implementation, with minimal method development. It is therefore highly recommended for bioanalytical research work in pharmaceutical clinical trials, including contract research.

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