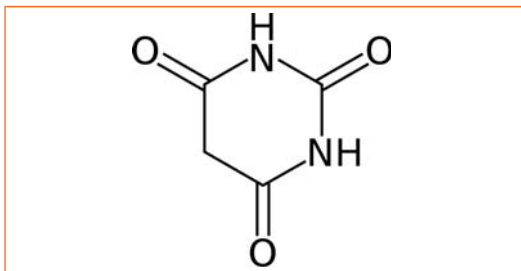


# Extraction of Barbiturates From Human Urine Using ISOLUTE® SLE+ Columns with GC-MS Analysis

## Introduction

This application note describes the extraction of a range of barbiturates from human urine using ISOLUTE SLE+ supported liquid extraction columns followed by GC-MS analysis.



**Figure 1.** Structure of Barbituric acid, the basic structure of all barbiturates

This method describes the use of ISOLUTE SLE+ supported liquid extraction 1 mL sample volume columns to extract a range of barbiturates from human urine. The analysis of these analytes was carried out by GC-MS. This simplified and efficient extraction method has significant analyte recoveries ranging from 103-108% with LOQs of 10 ng/mL and RSDs <10%.

ISOLUTE SLE+ Supported Liquid Extraction plates and columns offer an efficient alternative to traditional liquid-liquid extraction (LLE) for bioanalytical sample preparation, providing high analyte recoveries, no emulsion formation, and significantly reduced sample preparation time.

## Analytes

Butabarbital, butalbarbital, amobarbital, pentobarbital, secobarbital, hexobarbital, phenobarbital.

## ISOLUTE SLE+ procedure

### ISOLUTE SLE + 1 mL sample volume column, part number 820-0140-C

**Sample Pre-treatment:** To 500 µL of urine add 100mM ammonium acetate pH 5 (500 µL, 1:1, v/v).

**Sample Load:** Load pre-treated sample (1 mL) to column followed by a pulse of vacuum to initiate flow and allow to absorb for five minutes.

**Analyte Extraction:** Elute with dichloromethane (2.5 mL). Leave to flow under gravity for 5 minutes, then follow with a further aliquot of dichloromethane (2.5 mL) and allow to flow under gravity for a further five minutes, to complete extraction apply a short pulse of vacuum.

**Post extraction:** Evaporate to dryness at room temperature (80 L/min) and reconstitute in ethyl acetate (200 µL).

## Reagents

Ethyl acetate from Fisher Scientific, Loughborough.

Ammonium acetate from Sigma-Aldrich, Gillingham.

Trimethylphenylammonium hydroxide (TMAH) from Sigma-Aldrich, Gillingham.

## GC Conditions

<b>Carrier:</b>	Helium 2 mL min <sup>-1</sup> (constant flow)
<b>Inlet:</b>	Splitless, 150 °C
<b>Injection:</b>	In-port flash alkylation: 1 µL sample + 1 µL 0.2M TMAH (trimethylphenylammonium hydroxide) in MeOH
<b>Oven:</b>	120 °C to 290°C at 15 °C min <sup>-1</sup> , hold 2min
<b>Transfer Line:</b>	280 °C

## Mass Spectrometry Conditions

<b>Source temp:</b>	230 °C.
<b>Quadropole temp:</b>	150 °C.
<b>Solvent delay:</b>	7 min.
<b>MSD mode:</b>	SIM.
<b>SIM Groups:</b>	1 - 7.4 min to 7.8 min / 2 - 7.8 min to 8.1 min / 3 - 8.1 min to 8.4 min / 4 - 8.4 min to 9.8 min / 5 - 9.8 min to 10.4 min / 6 - 10.4 min to 16.0 min

Table 1. SIM parameters

Scan function	Compound	Quant Ion	1 <sup>st</sup> Qual Ion	2 <sup>nd</sup> Qual Ion	Dwell / ms
1	Butalbarbital	196	181	25	100
1	Butabarbital	169	211	37	100
2	Amobarbital	169	225	29	100
3	Pentobarbital	169	225	33	100
4	Secobarbital	196	181	25	100
5	Hexobarbital	235	81	27	100
6	Phenobarbital	232	175	33	100

## Results

Figure 2 shows the mass chromatograms for all the extracted barbiturates spiked at 10 ng/mL. Figure 3 shows average analyte recoveries ranging from 103-108% for barbiturates spiked at 10 ng/mL, with RSDs < 10% (n=3).

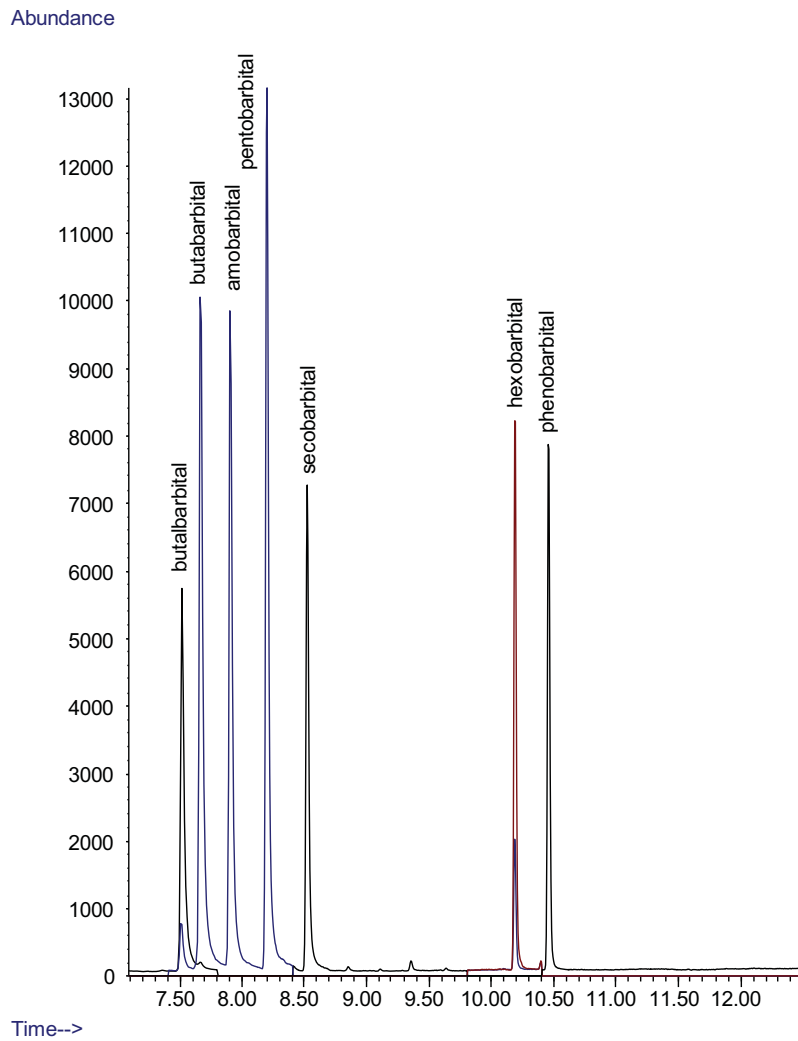


Figure 2. Typical chromatograms for all Barbiturate analytes at 10 ng/mL

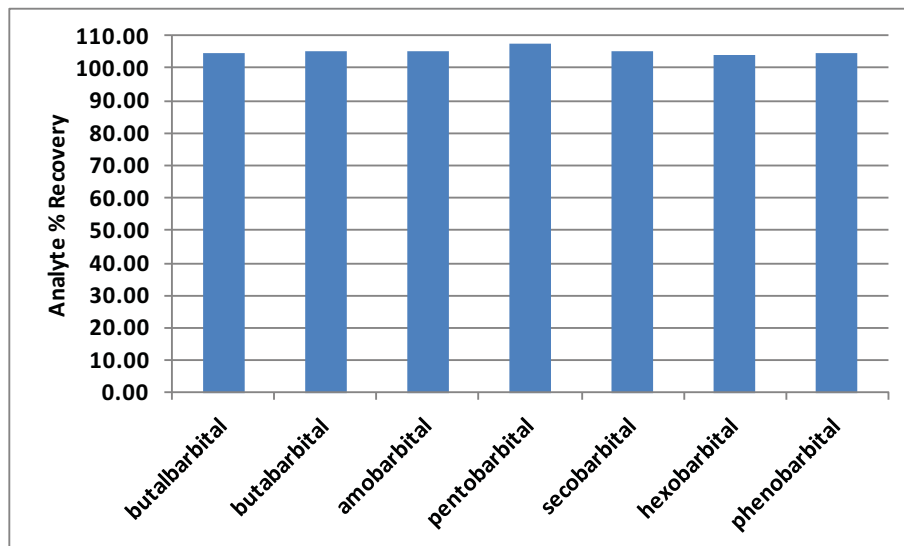


Figure 3. Average analyte recoveries of a range of Barbiturate analytes at 10 ng/mL (n=3).

### Ordering information

Part number	Description	Quantity
820-0140-C	ISOLUTE SLE + 1 mL sample	30

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