Idaho State Police Forensic Services Toxicology Section

Section Two

- 2.4 Liquid-Liquid Extraction Methods for GC/MSD Confirmation
 - 2.4.1 General Extraction of Urine Samples for Qualitative Confirmation of Basic and Neutral or Acidic Drugs

2.4.1.1 BACKGROUND

These extraction procedures are extensions of the TOXI-LAB® TOXI-A and TOXI-B thin layer chromatography (TLC) drug detection systems. The samples are extracted as with the TLC system, however, instead of concentrating the extract onto a disc, the solvent extract is concentrated and placed into an automated liquid sampler (ALS) vial for analysis by a gas chromatograph equipped with a mass selective detector (GC/MSD).

2.4.1.2 PRINCIPLE

This procedure describes the extraction of drug compounds from urine. Depending upon the pK_q of a drug compound, either Toxi-A or Toxi-B tubes are used. Basic compounds are extracted with a Toxi-A tube. Addition of urine to the Toxi-A tubes results in the urine becoming alkaline (pH=9) into 1,2-Dichloroethane, dichloromethane, heptane and isopropanol. Acidic compounds are isolated from an acidified solution (pH=4.5) into methylene chloride and heptane with zinc chloride to facilitate the extraction process. The extraction is achieved with an Ansys Toxi-B extraction tube. Either resulting extract is analyzed by full scan GC/MS in El mode

2.4.1.3 EQUIPMENT AND SUPPLIES

A MACHINE	THE SCITCES		
2.4.1.3.1	Tube Rocker (Fisher Scientific or equivalent)		
2.4.1.3.2	Electric Warmer with Omega-12 extraction solvent		
	concentrator (Ansys 118/153)		
2.4.1.3.3	Laboratory Centrifuge (Fisher Marathon or equivalent)		
2.4.1.3.4	Disposable Aluminum Concentration Cups (Ansys 152)		
2.4.1.3.5	Glassware GC/MS vials (HP 5182-0865 or equivalent)		
	GC/MS vial microinsert (HP 5183-2088 or equivalent)		
2.4.1.3.6	Gas Chromatograph equipped with a mass selective		
	detector (HP 6890/5973) and a HP-5MS Ultra low bleed		
	(5%-Diphenyl-95%-Dimethylsiloxane copolymer) capillary		
	column (25M).		

2.4.1.4	REAGENTS 2.4.1.4.1	ANSYS TOXI-TUBES A and B (Ansys 109A-100/ 109B-100)
2.4.1.5	CONTROLS 2.4.1.5.1	Toxi-Control No. 19 — Morphine, amphetamine, imipramine, methadone, propoxyphene, phenobarbital, secobarbital and benzoylecgonine (Ansys 191AB).
	2.4.1.5.2	Toxi-Control No. 2 — Amphetamine, methamphetamine, nicotine and cotinine. (Ansys 170B).
2.4.1.6	STANDARI	os co
	2.4.1.6.1	Run necessary analytical standards as indicated by examination of GC/MSD data. RE Initial set-up
2 4 1 7	PROCEDUI	
2.4.1.7	2.4.1.7.1	Initial set-up
	2.4.1.7.1	Label TOXI-TUBES & or B, and GC/MS vials with microinserts with negative control, TC-19 and or TC-2 and appropriate laboratory numbers.
	2.4.1.7.2	Extraction Procedure Toxi-A Extraction (Basic or Neutral Compounds) Fransfer 5 mL of urine specimen, negative urine or appropriate Foxi-Control to a TOXI-TUBE A (pH=9).
	24102	Rock TOXI-TUBE A for 15 minutes.
	2.4.1(7.3	Centrifuge tube at 2500 rpm for 15 minutes. Transfer solvent from tube into concentration cup in
. 6	2 4.3.7. 4	Omega-12 extraction solvent concentrator. Allow cups to warm prior to the addition of extract.
8kg	2.4.1.7.5	Evaporate solvent to approximately 200μL on electric warmer.
	2.4.1.7.6	Transfer solvent to labeled GC/MS ALS vial with microinsert.
	2.4.1.7.7	Extraction Procedure Toxi-B Extraction (Acidic Compounds)
	2.4.1.7.8	Transfer 4.5 mL of urine specimen, negative urine or Toxi-Control 19, to a TOXI-TUBE B (pH=4.5).
	2.4.1.7.9	Rock TOXI-TUBE B for 15 minutes.
	2.4.1.7.10	Centrifuge tube at 2500 rpm for 15 minutes.
	2.4.1.7.11	Transfer solvent from tube into concentration cup in Omega-12 extraction solvent concentrator.

- 2.4.1.7.12 Evaporate solvent to approximately 200µL on electric warmer.
- Transfer solvent to labeled GC/MS ALS vial with 2.4.1.7.13 microinsert.
- 2.4.1.7.4 Gas Chromatography/Mass Spectrometry (GC/MS) Analysis 2.4.1.7.4.1 Inject 1 µL into GC/MS using the ALS.
 - Analyze sample extract in full scan acquisition. Refer to attached GC/MSD method printout for
- The presence of a drug compound can be established if there are no significant differences in the retention time and mass spectra for the sample