Section Three Blood Toxicology

3.10 SPE Methods for GC/MSD Confirmation

3.10.1 Extraction of 11-nor- Δ^9 -THC-9-COOH (Carboxy-THC) from Blood Employing the United Chemical Technologies (UCT) 200 mg CLEAN SCREEN THC Column

3.10.1.1 BACKGROUND

 Δ^9 -THC (Figure 1) is the chief psychoactive cannabinoid resulting from exposure to marijuana. Δ^9 -THC has a peak blood concentration within 5 to 15 minutes following smoking of a marijuana cigarette. This blood concentration drops rapidly after cessation of smoking.^{3,4} The level may fall to less than 5µL within 30 to 60 minutes, although longer detection times have been reported.^{3,4} Detection of low dose (1.75% post-smoking Δ^9 -THC has been reported to vary from 3 to 12 hours. 4 This detection window was based on a limit of quantitation of 0.5ng/mL. The number, duration, and spacing of puffs, hold time, and inhalation volume all impact the degree of drug exposure and thus bioavailability tonger detection times have been observed for frequent users. The Δ° THC metabolite 11-nor- Δ° -THC-9-COOH (Carboxy-THC), concentration gradually increases and may plateau for several hours.⁴ There is poor correlation between blood Δ^9 -THC and psychoactive effects since the 29-THC concentrations begin to decline prior to the time of peak effects. 3,4,5 Research continues on models using the relative amounts of A-THC and Carboxy-THC to assist with establishing recent drug use.

Negative behavioral effects reported from exposure to marijuana include altered time perception, lack of concentration, impaired learning and memory which can lead to impairment of cognitive and performance tasks.⁴ Establishing impairment in an individual is based on evaluation of all available information in conjunction with quantitative blood levels.

For additional background, refer to analytical method 2.4.4 and provided references.

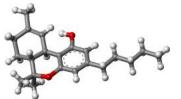


Figure 1.

3.10.1.2 PRINCIPLE & SCOPE

This procedure outlines the use of the 200mg United Chemical Technologies (UCT) CLEAN SCREEN® THC Column for the extraction of carboxy-THC from blood. The CLEAN SCREEN® THC column utilizes a copolymeric sorbent which combines a cationic exchanger and a hydrophobic functionality (reverse phase) to interact effectively, physically and chemically, with analytes of interest while interacting minimally with interfering substances in the blood sample. The cation exchanger utilizes an anionic sorbent to bind to cations. Additional retention mechanisms include hydrophobic interactions and polar adsorption.

For the extraction of carboxy-THC, a deuterated internal standard is added Blood proteins are precipitated with 10% Methanolto the sample. Acetonitrile solvent mix and are removed (via centrifugation. The supernatant is transferred to a clean tube; nigogen is used to reduce the volume by one-half. The sample pH is then adjusted with an acetate buffer (pH 4.5) and loaded onto a pre-conditioned SPE column The conditioning of the SPE column creates an environment which allows for optimal interaction between the sorbent and the analyte of interest. The column is subsequently washed to selectively emove matrix components and interfering substances from the column. Next, the column is dried to remove traces of aqueous and organic solvents. After drying, the analyte of interest is eluted from the SPE doumn with an organic solvent mixture. Following elution and evaporation of the solvent, the extract is derivatized for compound confirmation using a gas chromatograph equipped with a mass selective detector (GC-MSD).

3.10.1.3 EQUIPMENT AND SUPPLIES

	EQUIPMEN	I AND SUPPLIES
	3.10.1.3.1	200mg CLEAN SCREEN® THC Extraction Column
	3.10(1)3.2	Disposable inserts for SPE manifold ports
	3.10.1.3.3	Drybath or laboratory oven capable of 70°C
C	3.10.1.3.4	Evaporative concentrator equipped with nitrogen tank.
Ò,	3.10.1.3.5	Tube rocker
) `	3.10.1.3.6	Vortex mixer
	3.10.1.3.7	Laboratory centrifuge capable of 3400- 3500rpm
	3.10.1.3.8	Vacuum Manifold/ Vacuum pump
	3.10.1.3.9	Fixed and adjustable volume single channel air displacement
		pipetters, and appropriate tips, capable of accurate and
		precise dispensing of volumes indicated.
	3.10.1.3.10	16 x 100mm glass tubes (silanized recommended)
	3.10.1.3.11	Screw Cap for 16mm O.D. tubes
	3.10.1.3.12	GC/MS Automated Liquid Sample (ALS) vials
	3.10.1.3.13	GC/MS Vial Microinsert (silanized recommended)
	3.10.1.3.14	Gas Chromatograph equipped with a Mass Selective Detector
		and a nonpolar capillary column with a phase composition

comparable 100%-dimethylpolysiloxane 95%to or dimethylpolysiloxane with 5%-diphenyl.

REAGENTS 3.10.1.4

Refer to manual section 5.12 for solution preparation instructions.

Rejer to man	uai section 5.12 for solution preparation instructi
3.10.1.4.1	Deionized/distilled (DI) water
3.10.1.4.2	Methanol (Certified ACS Grade)
3.10.1.4.3	Hexane (Certified ACS Grade)
3.10.1.4.4	Ethyl Acetate (Certified ACS Grade)
3.10.1.4.5	Acetonitrile (Certified ACS Grade)
3.10.1.4.6	10% Methanol in Acetonitrile
3.10.1.4.7	100mM Acetate Buffer (pH 4.5)
3.10.1.4.8	100mM HCl
3.10.1.4.9	70:30 Hexane:Ethyl Acetate
3.10.1.4.10	70:30 100mM HCl:Acetonitrile
3.10.1.4.11	BSTFA + 1% TMCS

QUALITY ASSURANCE MATERIAL 3.10.1.5

Calibrator and Control Solutions 3.10.1.5.1

Corresponding calibrator and control reference material must be obtained from different vendors or be from different lot numbers if suitable second vendors are not available.

Reference Material Stock Solutions

Concentration: 100µg/mL or 1mg/mL

Carboxy-THC

Store remaining solution stock as recommended by vendor.

Reference Material Working Solutions

Working solutions are stable for 6 months when stored under refrigeration.

Concentration: 1.0ng/µL

As appropriate, add 100µL of 100µg/mL or 10µL of 1mg/mL Stock Solution to approximately 9mL Methanol in a 10mL volumetric class A flask. QS to 10mL.

Concentration: 0.1ng/µL

Add 1000µL of 1ng/µL working calibration solution to approximately 8mL Methanol in 10mL volumetric class A flask. QS to 10mL.

3.10.1.5.2 Internal Standard Solutions

3.10.1.5.2.1 **Stock Solution**

Concentration: 100µg/mL or 1mg/mL Carboxy-THC-D₉

Store remaining stock solution as recommended by vendor.

3.10.1.5.2.2 Working Internal Standard Solution

Working internal standard solution is stable for 6 months when word under refrigeration.

Concentration: 1.0ng/pl

Add $100\mu L$ of $100\mu g/mL$ or $10\mu L$ of 1mg/mL stock solution to approximately 9mL Methanol in a 10mL volumetric class A flask. QS to 10mL.

3.10.1.5.3 Whole Blood Negative Control Negative Whole Blood

3.10.1.6 PROCEDURE

3.10.1.6.1 <u>Initial set ap</u>

Label extraction tubes (x3), 200mg CLEAN SCREEN[®] extraction columns (x1), and GC/MSD vials (x1) with microinsens for calibrators, controls and case samples.

3.10.1.6.2 <u>Calibrator Preparation</u>

Use the same lot of negative blood for all controls and calibrators.

3.10.1.6.2.1

Add 1mL of negative whole blood to six extraction tubes.

3.10.1.6.2.2 Add the volume of 0.1ng/μL carboxy-THC calibrator working solution as indicated in the following table:

Level	Desired ng/mL	μL Working RM
1	2.5	25
2	5	50
3	10	100

3.10.1.6.2.3 Add the volume of $1.0 ng/\mu L$ carboxy-THC calibrator working solution as indicated in the following table:

Level	Desired ng/mL	μL Working Reference Material	
4	25	25	
5	50	50	
6	100	2 000	

3.10.1.6.3 <u>Positive Control Sample Preparation</u>

Use the same lot of negative blood for all controls and calibrators.

3.10.1.6.3.1 Add 1mL of negative whole blood to two extraction tobes.

3.10.1.6.3.2 Add indicated amount of 0.1ng/μL working control solution.

Des	ired	ng/mL	μL Working Control
.0	6.0	77,	60.0

3.10.1.6.3.3 Add indicated amount of 1.0ng/µL working control solution.

$ar{\Xi}$	Desired ng/mL	μL Working Control
Y	60.0	60.0

3.10.1.6.4 Negative Control Sample Preparation

Add mL of negative whole blood into an extraction tube.

3.10.1.6.5 Case Sample Preparation

Place sample container on tube rocker for a minimum of five minutes. If sample is clotted, homogenize as necessary. Transfer 1mL of blood into a labeled extraction tube.

3.10.1.6.6 Internal Standard Addition

3.10.1.6.6.1 To calibrators, controls and case samples, add $25\mu L$ of internal standard.

3.10.1.6.6.2 Vortex tube briefly and allow to stand 15 to 30 minutes for sample equilibration.

3.10.1.6.7	Protein Precipita 3.10.1.6.7.1	While vortexing, add 2mL cold 10% methanol in acetonitrile dropwise to case, calibrator and control samples.
	3.10.1.6.7.2	Cap tube and continue vortexing tube for approximately 30 seconds.
	3.10.1.6.7.3	Allow tube to stand for approximately five minutes.
	3.10.1.6.7.4	Centrifuge at approximately 3500 rpm for 10 minutes.
	3.10.1.6.7.5	Decant organic supernatant into second labeled glass tube.
	3.10.1.6.7.6	Transfer tube to Evaporative Concentrator and evaporate under nitrogen at ≤40°C to approximately ImL. Do not allow extract to go to dryness.
	3.10.1.6.7.7	To the evaporated extract add 2mL 100mM acetate buffer (pH 4.5). Vortex briefly to mix.
£108	SPE Column Pr	Linecessary, centrifuge buffered solution for an additional 5 minutes at approximately 3500 rpm to remove blood fragments or foam.
310168	SPE Column Pr	enaration
Property 10.1.6.8	SPE Column Pr 3.10.1.6.8.1	Insert valve liners and labeled SPE columns into appropriate location on vacuum manifold. For each following SPE step, allow to gravity flow or aspirate at ≤ 3 in. Hg to prevent sorbent drying.
	3.10.1.6.8.2	To each SPE column, add 3mL 70:30 Hexane:Ethyl acetate .
	3.10.1.6.8.3	To each SPE column, add 3mL methanol to the column.
	3.10.1.6.8.4	To each SPE column, add 3mL deionized or distilled water to the column.

Forensic Services

	3.10.1.6.8.5	To each SPE column, add 1mL 100mM
	3.10.1.0.6.3	HCl.
3.10.1.6.9	Blood Extract L	oading I blood extract onto the SPE column.
	Decam buriered	i blood extract onto the SPE column.
3.10.1.6.10	Column Wash	
	3.10.1.6.10.1	To each SPE column, add 2mL of deionized or distilled water .
		S
	3.10.1.6.10.2	To each SPE column, add 2mL 70:30
		100mM HCl:Acetonitrile.
3.10.1.6.11	Dry Disc	co,
	Turn on/increas	se vacuum to ≥ 10 in. Hg (≥ 34 kPa) for ≥ 5
	minutes.	2816
3 10 1 6 12	Compound Elut	ion (e) of o
3.10.1.0.12	3.10.1.6.12.1	Open vacuum manifold, wipe collection
		tips, and insert the collection rack containing
		the labeled glass tibes.
	3.10.1.6.12.2	To each SPE column, add 200uL hexane
	ש	(important for elution solvent reception).
	CXOL C	Graving flow only. Do not allow column to
	2,116	dry.
×	3.10.1.6.12.3	To each SPE column, add 3mL 70:30
		Hexane: Ethyl Acetate elution solvent.
8/0	~CV	Collect eluate with gravity flow or apply

3.10.1.6.13 Eluate Evaporation

Transfer centrifuge tube to Evaporative Concentrator. Evaporate eluates to dryness, under a gentle stream of nitrogen at $\leq 40^{\circ}$ C.

3.10.1.6.14 <u>Derivatization</u>

3.10.1.6.14.1 Add 40µL ethyl acetate and vortex.

minimal vacuum.

3.10.1.6.14.2 Add 40µL BSTFA (1% TMCS) to each extract. Cap tubes and vortex briefly.

3.10.1.6.14.3 Heat tubes at 70°C for 15 minutes.

3.10.1.6.14.4	Remove tubes from heat and allow to cool to
	room temperature.

3.10.1.6.14.5 Transfer derivative to labeled GC/MSD ALS vial with microinsert.

3.10.1.6.15 Preparation for Analysis Run

3.10.1.6.15.1 Into Sequence log table, enter the sample case numbers, blanks and controls.

3.10.1.6.15.2 Load samples, standards, blankand controls into the quadrant rack as noted in the sequence table.

3.10.1.6.16 GC-MS Calibration Curve

3.10.1.6.16.1 The calibration curve must be established with a minimum of four data points.

3.10.1.6.16.2 Calibrators should be analyzed in order of increasing concentration.

3.10.1.6.16.3 The least squares line resulting from the analysis of the calibrators must have a coefficient of correlation of ≥0.98.

.10.1.6.16.4 If calibrators are run in duplicate, it is not required that duplicate calibration points be included as long as the linearity requirement is met.

3.10.1.7 GC and MSD ACQUISITION PARAMETERS

Critical parameters are specified below. Parameters not specified are at the discretion of the analyst and should be optimized for the particular GC-MSD instrument. Each laboratory should maintain a centrally stored printed or electronic copy of current and past GC-MSD methods. The data supporting the GC-MSD method should be stored centrally.

3.10.1.7.1 <u>GC Temperature Parameter</u>

Injection Port: 250°C or 260°C

3.10.1.7.2 MSD Instrument Parameters

Detector/Transfer Line: 280°C

3.10.1.7.3 ALS Parameters

Injection Volume: 1µL (1 stop)

Viscosity Delay: A minimum of 1 second

Solvent Washes (A & B): A minimum of 3 pre- and post-wash rinses.

3.10.1.7.4 MS SIM Parameters

Analyte	Target Ion	Qualifier	Qualifier
		Ion 1	Ion 2
Carboxy-THC	371	473	488
Carboxy-THC-D9	380	482	4 97

3.10.1.8 REPORTING CRITERIA

3.10.1.8.1 Qualitative Chromatographic Criteria

Acceptable retention time window established by calibrators is ± 0.1 minutes.

3.10.1.8.2 Qualitative Mass Spectral SIM Criteria

Ion ratios for the analyte and its corresponding internal standard, established by calibrators for target and qualifier ions, must not differ by more than $\pm 20\%$ (relative). Refer to section 3.10.1.8.4 5for cutoff criteria.

Cut-off for Carboxy THC is the lowest calibrator (2.5ng/mL), or the lowest calibrator that meets quality assurance requirements. Any analyte with a quantitative value below this cut-off will be reported as "none detected." If the concentration exceeds the calibration range, the sample can either be appropriately diluted with negative whole blood for reanalysis or qualitatively confirmed in full scan mode (refer to section 3.10.1.8.3 for confirmation criteria).

3.10.08.3 Qualitative Mass Spectral Full Scan Criteria

Analytes may be confirmed from full scan data if the retention time for the sample versus applicable reference material does not differ by more than ± 0.1 minutes and there are no significant differences in the mass spectral data.

3.10.1.8.4 Quantitative Mass Spectral Criteria

3.10.1.8.4.1 Refer to Section 3.10.1.9.2 for determination of when this method may be used for quantitative purposes.

3.10.1.8.4.2 Quantitative results can be accepted if the calculated concentrations of all calibrator and control samples are within $\pm 20\%$ of their respective concentrations (relative).

3.10.1.8.4.3 Quantitation is achieved through the plotting of the target ion response ratio versus the concentration for each calibrator.

3.10.1.8.4.4 Quantitative values for case samples, calibrators and controls will be truncated for reporting purposes.

3.10.1.8.4.5 Administrative limit of detection (LOD) for carboxy-THC is 2.5ng/mL, the lowest calibrator level. Results < this LOD should be reported as negative noises there are extenuating circumstances. The Toxicology Discipline Leader fairst be consulted to evaluate exceptions.

3.10.1.8.4.6 If the concentration exceeds the calibration range, the sample must be appropriately diluted with negative whole blood for reanalysis. Alternatively, the analyte may be reported using full scan data; refer to section 3.10.1.8.3 for criteria.

3.10.1.9 REPORTING OF RESULTS

3.10.1.9.1 Qualitative Confirmation

If Carboxy-PHC meets confirmation criteria, it may be reported. The administrative cut-off of 2.5ng/mL, or the lowest calibrator meeting quality assurance requirements, will be used to determine if the analyte is detected.

3.10(1.9.2 Quantitative Value

This method is currently only approved for the qualitative dehtification of drugs. Quantitative values are not to be reported or expressed. They are currently being used to establish an administrative cut off. Once the uncertainty of measurement is established for this method, it will be evaluated for quantitative reporting.

3.10.1.10 QUALITY ASSURANCE REQUIREMENTS

3.10.1.10.1 General

3.10.1.10.1.1 Blood samples are to be stored under refrigeration after aliquots are removed for analysis.

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3.10.1.10.1.2 Refer to toxicology manual section 5.1, 5.2, 5.8, and 5.10 for quality assurance and material reference authentication requirements.

Per Analysis Run Control Requirements 3.10.1.10.2

- 3.10.1.10.2.1 Minimally, a solvent blank must follow the highest calibrator, as well as precede each case sample.
- A minimum of the spiked blood controls 3.10.1.10.2.2 described in section 3.10.16.3 must be run per batch of samples. Controls should not be grouped at the beginning of the acquisition sequence. Rather, controls should be interspersed throughout the sequence.
- If the number of case samples exceeds 10, in 3.10.1.10.2.3 addition to the two controls described in 3.10.1.6.3, and spiked or commercially obtained blood control must be run with each additional 10 case samples. Additional concentrations may be used.
- 0.1.10.3.C. O.1.10.3.C. O.1.10 Analysts may combine their samples into a single run to conserve supplies. However, each analyst with samples in the run must independently comply with the control requirements in section 3.10.1.10.2. third-party reviewer must independently review the central file packet for compliance to method requirements.

3.10.1.10.3 Monitoring of Control Values
Once the mile Once the method has been approved for quantitative purposes, the following is required: upon the completion of analysis, input blood control values on a spreadsheet used to assess uncertainty for this method.

3.10.1.11 ANALYSIS DOCUMENTATION

3.10.1.11.1 Case results are to be recorded in the LIMS system.

- 3.10.1.11.2 Original data for controls and standards will be prepared for each analysis run and stored centrally in the laboratory where the analysis was performed, until archiving or destruction.
- 3.10.1.11.3 A copy of data for controls and standards may be stored electronically in a central location and need not be included in individual case files. When necessary, a copy of the control and standard printouts can be prepared from the centrally stored document.

REFERENCES AND RECOMMENDED READING 3.10.1.12

- UCT CLEAN SCREEN® Extraction Columns Application 3.10.1.12.1 Manual.
- Standard Operating Procedure for Blood SPE Cannabinoids, 3.10.1.12.2 Montana Department of Justice Forensic Sciences Division.
- Standard Operating Procedure for Blood SPE THC and 3.10.1.12.3 Carboxy-THC GC/MSD Assay, Edmonton, Canada Office of the Chief Medical Examiners, 2003.
- Huestis, M.A. Cannabis (Marijuana) Effects on Human 3.10.1.12.4 Behavior and Performance, Forensic Science Rev. 14(1/2): 16-60, 2002.
- *Cannabis*, pp. 178-212. *in*: The Forensic Pharmacology of Drugs of Abuse, Arnold: London, 2001.
- Bock, Peter, Setting it right R&D methods for engineering, Academic Press, San Diego, 2001. Bock, Peter, Vetting it right - R&D methods for science and

Revision History

Section Three

Blood Toxicology

3.10 Manual Solid Phase Extraction (SPE) Methods

3.10.1 Extraction of Carboxy-THC from Blood Employing the United Chemical Technologies (UCT) 200 mg CLEAN SCREEN® THC Extraction Column

		5
Revision No.	Issue Date	History/Comments
0	11-22-2006	Original Issue
U	11-22-2000	Original Issue Method is approved for qualitative purposes only.
		Upon review of uncertainty determinations for
		quantitative analysis this method will be applied
		for intended use.
		0,00,00
1	07-28-2008	Clarified that negative blood used to prepare
		calibrator and positive controls is the same lot as
		used for negative control
		00 01 21
2	03-07-2011	Replaced 4°C storage with "under refrigeration",
	, O	emphasized need for sample homogeneity.
	CX CO	Refermatted for clarity.
	05.15.60	
3	05-17-2013	Removed Δ^9 -THC from method.
	05-17-2013 tr	Clarified that stock solutions are to be stored per
	K W W	vendor recommendations.
	0, 11, 8	Added provision for qualitative confirmation of
X		analytes from full scan data. Removed quantitative references – method not for
00/	\sim	quantitation.
100	8	Clarified that analytes will be deemed "none
010	O	detected" if values achieved for that analyte is
		below the lowest calibrator meeting quality
		assurance requirements.
		Added provision for shared runs by multiple
		analysts, and clarified per-analyst control
		requirements.
		Made silanized extraction tubes a
		recommendation, due to supply availability
		limitations.
		Formatting for continuity.
4	03-13-2015	Clarified that cold 10% Methanol in Acetonitrile

he added to blood for the protein precipitation step
be added to blood for the protein precipitation step.
Deleted repetitions of aspiration from the SPE
method section; made single instruction at
beginning of SPE section to replace repetitions.
Clarified vortexing procedure in the derivatization
step.
Formatting for continuity.
Consolidated quality assurance paragraphs.
Added LIMS reporting requirement.
Re-added quantitative criteria, but clarified that
method not for quantitative use at thistime.
Removed tune information from 3.10.1.6.15, and
reworded this section for method consistency.
Made retention time requirement more rigorous.

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