Separation of 11-Hydroxy-THC Metabolites and Quantitation of 18 Total Cannabinoids in Whole Blood by UHPLC-MS/MS

INTRODUCTION

Despite changes in legal status of medical and recreational $\Delta 9$ -Tetrahydrocannabinol ($\Delta 9$ -THC), employer restrictions and regional accessibility continue to reinforce the popularity of $\Delta 8$ -Tetrahydrocannabinol ($\Delta 8$ -THC) among cannabis users. While toxicology testing for $\Delta 9$ - and $\Delta 8$ -Carboxy-THC-metabolites has become more routine, in many cases the determination of the psychoactive metabolites, 11-Hydroxy- $\Delta 9$ -THC and 11-Hydroxy- $\Delta 8$ -THC, is of higher importance. The method developed by our laboratory provides a detailed analysis of whole blood specimens, separating $\Delta 9$ - and $\Delta 8$ -11-Hydroxy-THC-metabolites, and quantitatively evaluating 18 cannabinoids at concentrations from 0.500-200 ng/mL.

OBJECTIVE

Develop an analytical method for the extraction, detection, and quantitation of (-)-Δ9-THC, (-)-Δ8-THC, 11-Hydroxy-Δ9-THC (11-OH-Δ9-THC), 11-Hydroxy-Δ8-THC (11-OH-Δ8-THC), Δ9-Carboxy-THC (Δ9-COOH-THC), Δ8-Carboxy-THC (Δ8-COOH-THC), Δ9-Tetrahydrocannabivarin (Δ9-THCV), Δ8-Carboxy-Tetrahydrocannabivarin (Δ9-COOH-THCV), Δ8-Carboxy-Tetrahydrocannabivarin (Δ8-COOH-THCV), Cannabidiol (CBD), 7-Carboxy-Cannabidiol (7-COOH-CBD), 7-Hydroxy-Cannabidiol (7-OH-CBD), Cannabidiolic Acid (CBDA), Cannabinol (CBN), Cannabigerol (CBG), Cannabicyclol (CBL), and Cannabichromene (CBC) in whole blood by LC-MS/MS for a controlled-dosing research study.

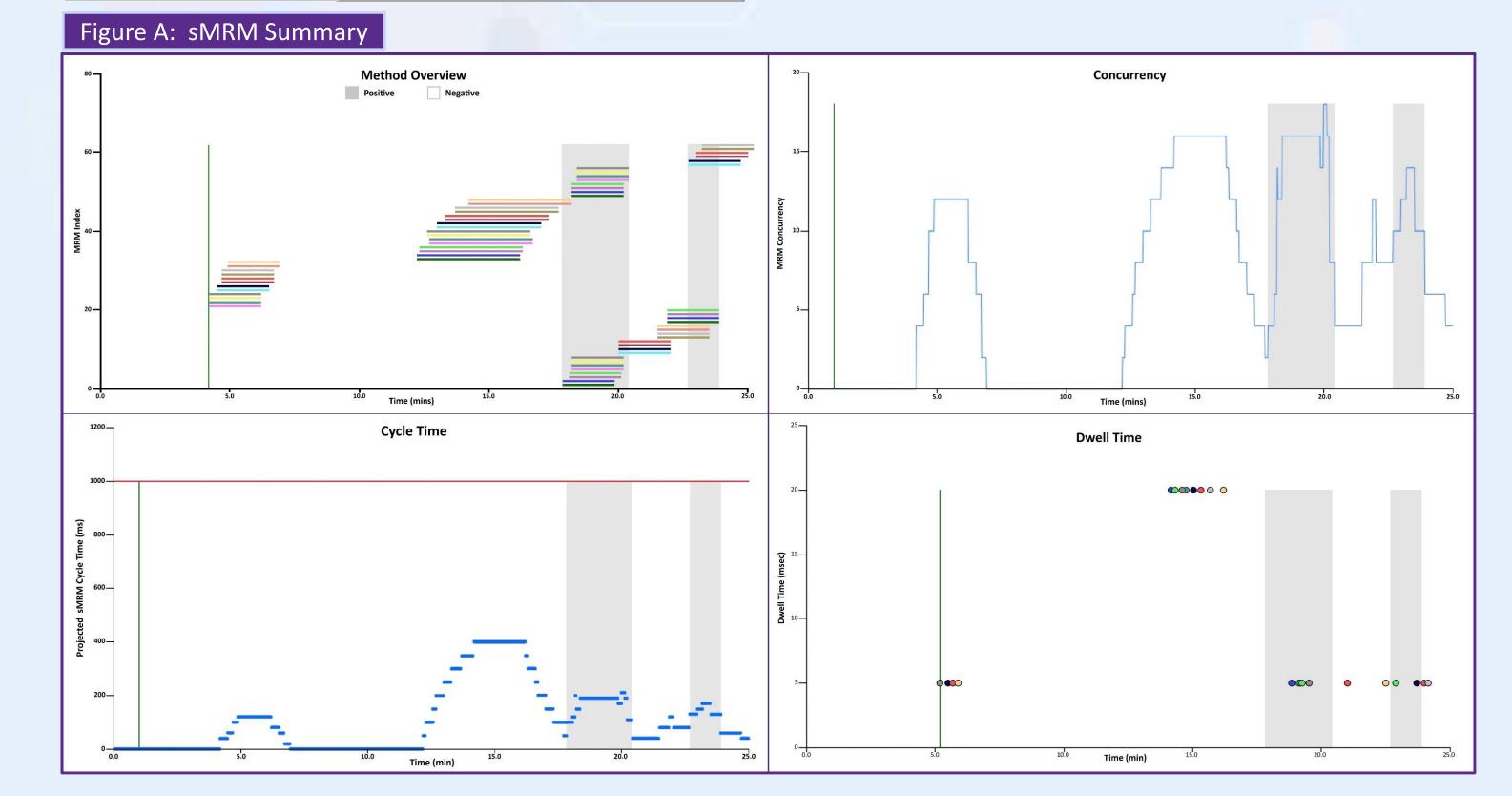
EXTRACTION METHOD

Specimens were prepared by mixing a 0.200 mL aliquot of whole blood sample with 20 μ L of internal standard solution and 0.1M Sodium Phosphate in appropriately labeled silanized glass culture tubes. While vortexing, 500 μ L of ice-cold Acetonitrile was added to each tube. Samples were then centrifuged and the supernatant was decanted from the blood protein pellet. A liquid-liquid extraction was performed using this supernatant and 9:1 Hexanes: Ethyl Acetate, and the organic components were subsequently dried and reconstituted with 0.1% Formic Acid in 50:50 DI H2O: Methanol.

INSTRUMENT PARAMETERS

Table 1: UHPLC-MS/MS Parameters

Table 1. UHPLC-IVIS/IVIS Parameters								
	Shimadzu Nexera	LC-40D X3 Pumps	Positive Ionization					
UHPLC System		SIL-40C X3 Auto Sampler	Analysta	Internal Characterist	Precursor	Product Ion	Product Ion	Retention Time
		SCL-40 System Controller	Analyte	Internal Standard	lon	Quantifier	Qualifier	(± 0.8 mins)
			Δ9-THCV	THCV-D5	287.2	165.1	123.0	19.90
		CTO-40C Column Oven	THO	CV-D5	292.2	170.1	123.1	19.84
		DGU-405 Degassing Unit	Δ8-THCV	THCV-D5	287.2	165.1	123.0	20.16
Injection Volume	25 μL		CBD	CBD-D3	315.2	193.1	135.0	20.31
•				D-D3	318.2	196.1	135.0	20.28
Analytical Column	(2) Waters CORTECS C18+, 90Å, 2.7 μm, 2.1 mm x 150 mm (Waters Part No. 186007398)		CBN	CBN-D3	311.2	208.0	223.0	22.24
			(-)-Δ9-THC	N-D3 (-)-Δ9-THC-D3	314.2 315.2	208.0 193.1	195.0 123.0	22.19 23.66
	Waters CORTECS C1	8+ VanGuard, 90Å, 2.7 μm,		-J-Z9-111C-D3	318.2	196.0	123.0	23.60
Guard Column			(-)-Δ8-THC	(-)-Δ8-THC-D9	315.2	193.1	123.0	24.08
	2.1 mm x 5 mm (Waters Part No. 186007685)		\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	-THC-D9	324.2	202.1	123.0	23.91
Column Temp.	40°C	7						
	Aqueous: 0.1% Ace	tic Acid in DI H2O	Negative Ionization					
Mobile Phase	Organic: 0.1% Acet	ic Acid in Acetonitrile	A so also de a		Precursor	Product Ion	Product Ion	Retention Time
EL D.			Analyte	Internal Standard	lon	Quantifier	Qualifier	(± 0.8 minutes)
Flow Rate	0.700 mL/min		7-COOH-CBD	7-COOH-CBD-D3	343.1	297.1	231.1	5.17
Run Time 27.00 minutes			7-COOH	H-CBD-D3	346.2	300.2	234.2	5.13
Mana Construction		7500 Triple Quad	Δ8-COOH-THCV	7-COOH-CBD-D3	315.2	271.1	163.0	5.51
Mass Spectrometer	Sciex API7500 Triple Quad		7-OH-CBD	7-OH-CBD-D3	329.2	299.2	268.1	5.66
Ionization	ESI Positive and Negative			CBD-D3	332.2	302.2	271.1	5.61
Source Temp.	550°C		Δ9-COOH-THCV	7-COOH-CBD-D3	315.2	271.1	163.0	5.92
120-240-sec detection window		11-OH-Δ9-THC	11-OH-Δ9-THC-D3	329.2	173.0	268.1	13.48	
Scheduled MRM			11-OH-Δ8-THC	19-THC-D3	332.2	173.0	271.1	13.34
	Target cycle time	1000 milliseconds		11-OH-Δ8-THC-D3 Δ8-THC-D3	329.2 332.2	173.0 173.0	268.1 271.1	13.91 13.76
UHPLC Mobile Phase Gra	dient 100		Δ8-COOH-THC	Δ8-COOH-THC-D6	343.1	245.1	191.1	14.50
Time Aqueous (%) Orga	nic (%)			H-THC-D6	349.2	251.1	191.1	14.14
	48.0	1-1	Δ9-COOH-THC	Δ9-COOH-THC-D9	343.1	299.1	245.1	15.54
	48.0 Aqueous	, , , ,	Δ9-COO	H-THC-D9	352.2	308.1	254.2	14.98
	48.0		CBG	CBG-D3	315.0	191.1	136.0	20.17
	70.0		СВ	G-D3	318.0	194.0	136.0	20.14
	72.0 40 Organic		CBDA	CBDA-D3	357.2	245.1	179.0	20.74
	0.0 Organic		CBDA-D3		360.2	248.1	182.0	20.71
	30.0		CBL	CBC-D9	313.2	191.0	203.0	24.66
	18.0 ¹⁰ %		СВС	CBC-D9	313.2	191.0	203.0	25.17
27.00 STOP	0 3 6	9 12 _{Minutes} 15 18 21 24 27	СВ	C-D9	322.3	200	212	25.06

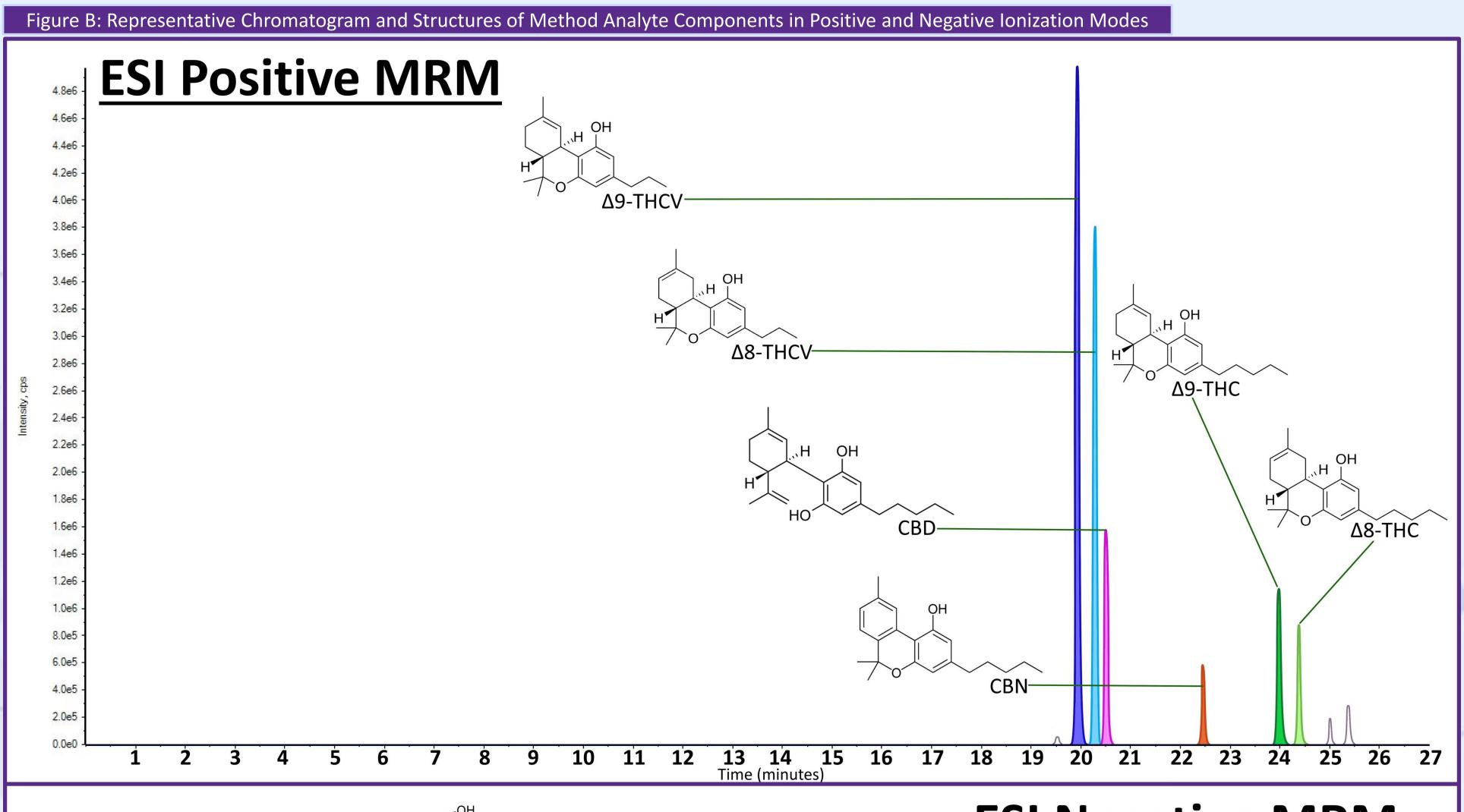


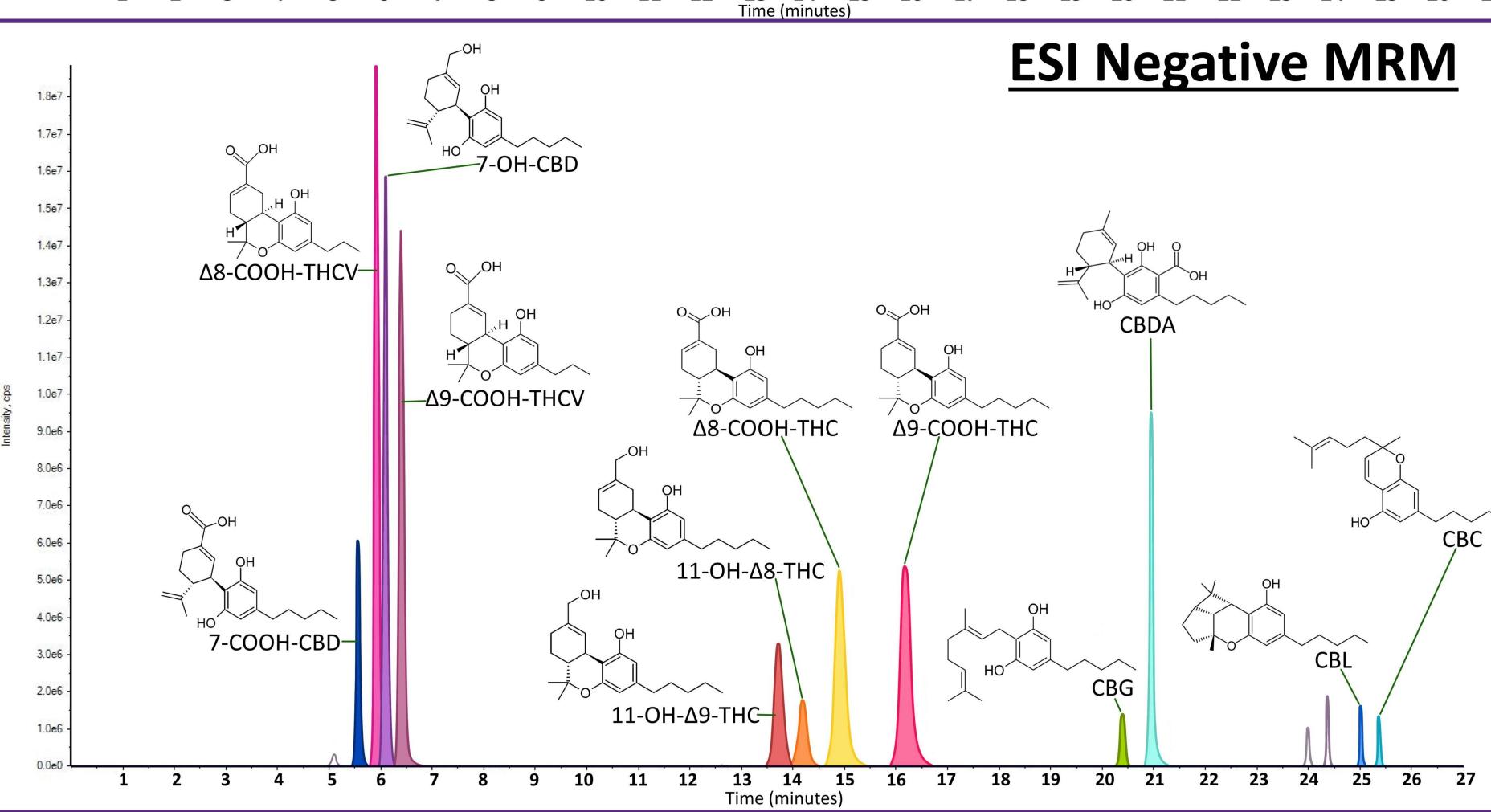
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RESULTS / DISCUSSION

A single-point calibrator at 5.0 ng/mL was used for quantitation. A low control at 2.0 ng/mL (40% of the calibrator), two positive controls at 6.25 ng/mL (125% of the calibrator), and two negative controls were run with each analytical batch, with one of the negative controls and one of the positive controls injected at the end of the batch to bracket donor samples. In addition to the low and positive controls, a conversion control was included in each batch to monitor the potential conversion of CBD and its metabolites to $\Delta 9$ -THC and $\Delta 8$ -THC and corresponding metabolites; the conversion control contained CBD, 7-OH-CBD, 7-COOH-CBD, and CBDA at 5.0 ng/mL.

Linearity was determined and assay limits of detection and quantitation (LOD/LOQ) and upper limit of linearity (ULOL) were established through the analysis of cannabinoid-analyte spiked UTAK negative whole blood samples at concentrations ranging from 0.250 ng/mL to 200 ng/mL. Accuracy and precision were calculated for 3 replicates of each of 14 concentration levels, including 40%, 50%, 100%, 125%, 150%, and 200% of the calibrator concentration. Qualitative acceptance was based on mean analyte concentrations within $\pm 20\%$ of target values with a coefficient of variation (CV) of <10%; due to the absence of a labeled internal standard for CBL, $\Delta 9$ -COOH-THCV, and $\Delta 8$ -COOH-THCV, accuracy within $\pm 25\%$ and CV <20% were accepted. For LOD/LOQ assessment, at the 0.250 ng/mL level, no analytes had reproducible quantitative results within $\pm 20\%$ of target with all replicates also passing qualitative acceptance criteria (see Table 2). Replicates for all analytes met both quantitative and qualitative acceptance criteria at the 0.5 ng/mL level. At the upper limit of linearity, replicates for 7-OH-CBD, $\Delta 9$ -COOH-THCV, and $\Delta 8$ -COOH-THCV met acceptance criteria at 100 ng/mL, and replicates for all other analytes met full acceptance criteria at 200 ng/mL. No carryover was observed at the highest concentrations.





Interference was investigated with 54 compounds, including over-the-counter, illicit, and commonly prescribed drugs. Throughout the study, no interference was detected with chromatography or internal standard recovery, and no erroneous peaks were observed that were greater than assay LOQ, which could create possible quantitation or identification issues. The potential of sample matrix components to interfere with the analytical method was evaluated by testing ten random negative whole blood patient samples, and continually monitored through the analysis of hundreds of study patient samples. Results showed no indication of methodic ion suppression or enhancement, and analyte and internal standard recovery was consistent. All samples passed with acceptable chromatography as no qualitative issues were observed, and no interfering peaks were present in the negative samples that could be problematic in quantitation or identification.



ble 2: Analyte LOD/LOQ and ULOL			Table 3: Quan	titative Acceptance Criter	ia			
Analyte	LOD/LOQ (ng/mL)	ULOL (ng/mL)	Relative Retention Time (RRT)	±2% of expected RRT of the analyte/internal standard pair	Apex of peak			
Δ9-THCV	0.500	200	Tille (KKI)	established by the batch calibrator	Asymmetry = <u>B</u> {larger side} A {smaller side}			
Δ8-THCV	0.500	200						
CBD	0.500	200	Internal Standard (IS)	Total IS peak area = ≥10% of	A B			
CBN	0.500	200	Internal Standard (IS)					
(-)-Δ9-THC	0.500	200	Response	calibrator IS peak area	I VIX			
(-)-Δ8-THC	0.500	200						
-COOH-CBD	0.500	200		Gaussian peaks;	Time (min) Minimum Peak Resolution = 90% (10% valley/peak ratio)			
-COOH-THCV	0.500	100	Symmetry /	asymmetry at 10% of peak height =				
7-OH-CBD	0.500	100	Peak Shape	<3.0 for IS and quant peaks				
-COOH-THCV	0.500	100		3.0 for 15 and quarte peaks	ΙΙ Λ Λ Ι			
1-OH-Δ9-THC	0.500	200						
1-OH-Δ8-THC	0.500	200		Adjacent peaks ≥90% resolved				
8-COOH-THC	0.500	200	Resolution	(≤ 10% valley/peak height ratio)				
9-СООН-ТНС	0.500	200		(ACCEPTABLE UNACCEPTABLE			
CBG	0.500	200						
CBDA	0.500	200	Ion Ratios	Ratio of abundance of quantitative				
CBL	0.500	200	(Qualifiers)	to qualifier ion = ±20% of target ratio	ACCEPTABLE 10% height			
СВС	0.500	200		established by batch calibrator	Time (min)			

CONCLUSION

This analytical method effectively separated 11-OH- Δ 9-THC and 11-OH- Δ 8-THC and demonstrated selectivity, accuracy, and reproducibility for the analysis of hundreds of samples in federally-sponsored controlled dosing research studies. Its application reliably identified and quantitated 18 cannabinoids at pg/mL levels, contributing to the scientific knowledge of cannabinoid metabolism and distribution in whole blood.

DEEEDENCE

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