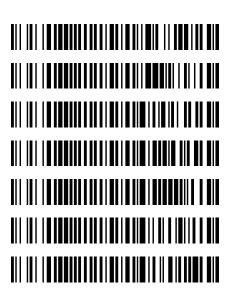
6/15/2023

#### Worklist: 6406

LAB CASE	ITEM	ITEM TYPE	DESCRIPTION
P2023-1258	1	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ
P2023-1343	1	ВСК	AM 27 Blood THC Quant by LC-QQQ
P2023-1459	1	ВСК	AM 27 Blood THC Quant by LC-QQQ
P2023-1505	1	ВСК	AM 27 Blood THC Quant by LC-QQQ
P2023-1506	1	ВСК	AM 27 Blood THC Quant by LC-QQQ
P2023-1575	1	ВСК	AM 27 Blood THC Quant by LC-QQQ
P2023-1591	1	ВСК	AM 27 Blood THC Quant by LC-QQQ



### AM# 27: Quantitation of THC and Metabolites in Blood and Urine by LC-MS/MS

Extraction Date: 06/07/2023 Plate lot#: 230113 Mobile phase A: 0.1% Formic Acid in LCMS Water Blank Blood Lot: Lampire 23A52594 Column: UCT Selectra DA 100 x 2.1mm 3um Analyst: <u>Tamara Salazar</u> Plate Retest Date: 07/13/2023 Mobile phase B: 0.1% Formic acid in Acetonitrile Blank Urine Lot: POC021022 LCMS-QQQ ID: 069901

### **Pre-Analytic:**

- $\boxtimes$  1. Check levels of mobile phases and needle wash refill as needed. Ensure waste is not full.
- ☑ 2. Ensure correct column is installed and begin mobile phase flow allow to equilibrate ~ 30 minutes.

### Analytic:

- ☑ 1. Remove standards, plate, controls, and samples from cold storage. Allow to reach room temperature.
- ☑ 2. Urine hydrolysis: add 1.5mL urine to blank plate, add 250µl 1N KOH. Shake and incubate at 40 degrees for 15 minutes.
- ☑ 3. Using a calibrated pipette, add 1000µl blood and urine (if applicable) (calibrated pipette) into the appropriate wells of analytical (standards) plate. Pipette ID: 42
- ☑ 4. Place on shaking incubator at ambient temp., 900rpm for 15 minutes.
- 5. Add 500µL of 0.1% formic acid in water to blood samples, and 500µL of saturated phosphate buffer to urine samples-in the wells of the analytical plate.
- $\boxtimes$  6. Place on shaking incubator at ambient temp., 900rpm for 15 minutes.
- ⊠ 7. Transfer **700-800µL of blood+acid or urine+acid** mixture to corresponding wells of SLE+ plate. Amount transferred:  $750\mu L$
- ☑ 8. Apply positive pressure for approx. 10-15 seconds (or until no liquid remains on top of sorbent). (Load at 85-100 PSI- Selector to the right)
- $\boxtimes$  9. Wait 5 minutes.
- ⊠ 10. Add 2.25mL MTBE. (Add in 3 increments of 750uL)
- $\boxtimes$  11. Wait 5 minutes.
- ☑ 12. Apply positive pressure for approx. 15 seconds. (10-15 PSI- Selector to the left).
- ☑ 13. Add 2.25mL Hexane. (Add in 3 increments of 750uL)
- $\boxtimes$  14. Wait 5 minutes.
- ☑ 15. Apply positive pressure for approx. 15 seconds. (10-15 PSI- Selector to the left).
- $\boxtimes$  16. Remove plate containing eluate. Place on SPE Dry and evaporate to dryness at approx. 35°C.
- ☑ 17. Reconstitute in **100µL 100% MeOH** and heat seal plate with foil. Place in autosampler and run worklist.

### **Post-Analytic**

- $\boxtimes$  1. Create batch and process data.
- $\boxtimes$  2. Make any necessary integration changes, Curve weighting of Linear 1/x with r<sup>2</sup> values  $\ge 0.98$  for each analyte
- RT +/- 3% or 0.100 min, whichever is greater, +/- 20% Accuracy for greater than (+/- 30% for 10ng/ml or less). Ion ratios must be within +/- 20% of the averaged calibrators
- $\boxtimes$  4. Did all QCs pass for each analyte? (if not, describe in comments section)
- $\boxtimes$  5. Enter QCs into control charting.
- 8 6. Central File Packet to include: LIMS Worklist, Method Checklist, Calibration and Control Reports

### COMMENTS:

	1	2	3	4	5	6
А	IS + Cal. 1	IS + QC_1	Neg Urine	IS + Sample	IS + Sample	IS + QC_1
В	IS + Cal. 2	Neg Blood	P2023-1258-1	IS + Sample	IS + Sample	IS + Cal. 7
с	IS + Cal. 3	P2023-1343-1	IS + Sample	IS + Sample	IS + Sample	IS + Cal. 6
D	IS + Cal. 4	P2023-1459-1	IS + Sample	IS + Sample	IS + Sample	IS + Cal. 5
E	IS + Cal. 5	P2023-1505-1	IS + Sample	IS + Sample	IS + Sample	IS + Cal. 4
F	IS + Cal. 6	P2023-1506-1	IS + Sample	IS + Sample	IS + Sample	IS + Cal. 3
G	IS + Cal. 7	P2023-1575-1	IS + Sample	IS + Sample	IS + Sample	IS + Cal. 2
н	IS + QC_1	P2023-1591-1	IS + Sample	IS + Sample	IS + QC_1	IS + Cal. 1

All wells to contain 100  $\mu l$  of residual DMSO

	1	2	3	4	5	6
А	IS + Cal. 1	IS + QC_1	Neg Urine			
В	IS + Cal. 2	Neg Blood	P2023-1258-1			
с	IS + Cal. 3	P2023-1343-1	P2023-1459-1*			
D	IS + Cal. 4	P2023-1459-1*				
E	IS + Cal. 5	P2023-1505-1				
F	IS + Cal. 6	P2023-1506-1				
G	IS + Cal. 7	P2023-1575-1				
н	IS + QC_1	P2023-1591-1				

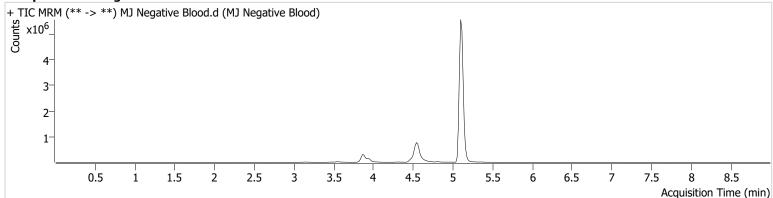
\*Sample moved during step 7 of the extraction process due to a clot



 Batch results
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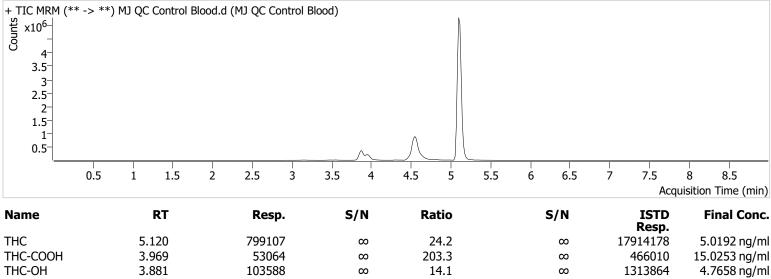
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Batch resultsD:\MassHunter\Data\2023\AM 27 28\060623 AM 27 28 TS\QuantResults\AM 27.batch.binCalibration Last Update6/8/2023 8:16:48 AM

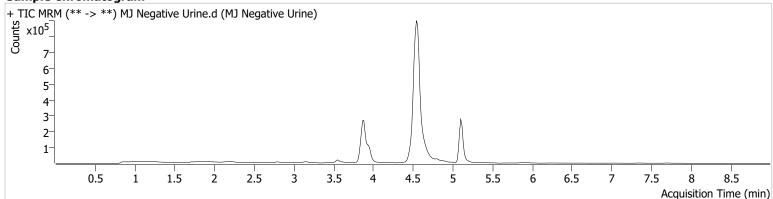
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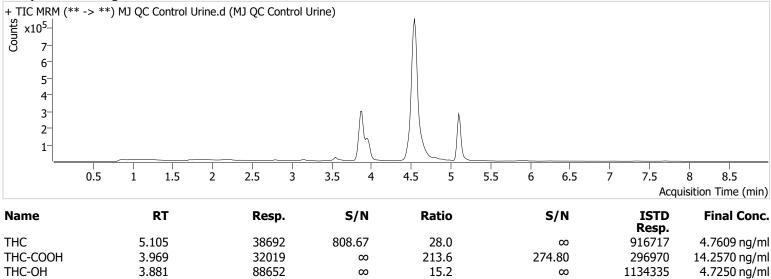
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Batch resultsD:\MassHunter\Data\2023\AM 27 28\060623 AM 27 28 TS\QuantResults\AM 27.batch.binCalibration Last Update6/8/2023 8:16:48 AM

Instrument Type Acq. Method Sample Position Injection Volume Acq. Date-Time Sample Info. Falco (069901) QC AM 27 Agilent Method.m P1-A2 10 6/7/2023 7:54:19 PM Data File Sample Operator Comment MJ QC Control Urine.d MJ QC Control Urine Tamara Salazar Only drugs and concentrations listed on the laboratory report itself are appropriate to be used for interpretation purposes. Any drugs or values included in the notes but not included on the report are used by laboratory personnel to make determinations/reach conclusions within the confines of the methods.





### AM #27 Cannabinoids Quant. Calibration Curve Report

		A	<b>′I #</b> ⊿	27 Ca	Innadi	noic	is Qua	<u>nt. (</u>	alidial		irve	керо	٦.	AENSIC SE
Batch	resu	lts		D:\Mass	Hunter\Da	ta\2023	3\AM 27 28\	060623	8 AM 27 28 T	ГS\QuantR	esults\A	M 27.bato	h.bin	
Last C	Cal. U	pdate		6/8/2023	3 8:16 AM									
Analy	st Na	me		ISP\Data	astor									
Analy	te			THC					Interr	nal Standa	rd	THC-D3		
Relative Responses H	- 7 Le 0.9- 0.8- 0.7- 0.6- 0.5- 0.4- 0.3- 0.2- 0.1- 0-	y = R^2	0.0092 = 0.9	296 * x 995371	d, 7 Point - 0.0020 3 n:Ignore, 20	49	oints Used, nt:1/x 40	, 2 QC	5 60	70	80 R	90 Relative (	100 Concentra	ition
		Sam	ole		Leve	el	Enable	d	Expected	Final	Concer	ntration	Accura	су

Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
Cal 1 MJ	1	~	1.0	1.1	108.5
Cal 2 MJ	2	~	3.0	3.0	99.6
Cal 3 MJ	3	~	5.0	4.9	98.7
Cal 4 MJ	4	~	10.0	9.5	95.3
Cal 5 MJ	5	~	25.0	24.2	97.0
Cal 6 MJ	6	~	50.0	49.7	99.5
Cal 7 MJ	7	~	100.0	101.5	101.5



### AM #27 Cannabinoids Quant. Calibration Curve Report

AI*		Inadinolus	S Quant. V	Landration	n Curve	Report	ALENSIC SE
Batch results	D:\MassHu	unter\Data\2023\	AM 27 28\06062	3 AM 27 28 TS\C	QuantResults	AM 27.batch.bin	
Last Cal. Update	6/8/2023 8	:16 AM					
Analyst Name	ISP\Datast	tor					
Analyte	THC-COO	Н		Internal S	Standard	THC-COOH-D	9
2 R^2 Type: 1.8 1.6 1.4 1.4 1.2 1- 0.8 0.6 0.4 0.2 0- 0-	evels, 7 Levels .007874 * x - = 0.99959875 Linear, Origin: 0 20 40	0.004446 Ignore, Weight		d, 2 QCs 140 160	180 200	220 240 Relative Conce	260 ntration
Samp	le	Level	Enabled	Expected	Final Conce	entration Acc	curacy

Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
Cal 1 MJ	1	~	5.0	5.0	100.7
Cal 2 MJ	2	V	10.0	9.9	98.9
Cal 3 MJ	3	~	20.0	19.7	98.4
Cal 4 MJ	4	~	50.0	52.2	104.3
Cal 5 MJ	5	~	75.0	74.7	99.6
Cal 6 MJ	6	~	100.0	97.7	97.7
Cal 7 MJ	7	~	250.0	250.8	100.3



### AM #27 Cannabinoids Quant. Calibration Curve Report

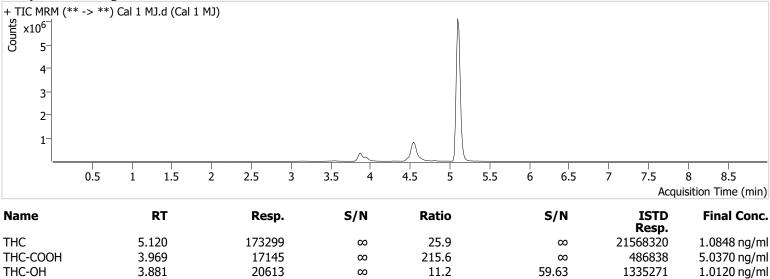
		Ar	<u>′I #Z</u>	/ Ca	innadi	noic	is Qua	<u>ni. C</u>	alibrati	on Cu	rve	Repor	MENSIC SE
	h resu			D:\Mass	Hunter\Da	ta\2023	am 27 28	060623	AM 27 28 T	S\QuantRe	sults\A	M 27.batch	n.bin
	Cal. U	-			3 8:16 AM								
Anal	yst Na	me		SP\Data	astor								
Anal	yte		-	THC-OH	4				Intern	al Standar	d	THC-OH-	D3
Relative Responses	2-OH - 1.6- 1.4- 1.2- 0.8- 0.6- 0.4- 0.2- 0-	y = 0 R^2	0.0168 = 0.99	91 * x 96284	Used, 7 P - 0.0016 2 n:Ignore, 20	56	7 Points U nt:1/x 40	Jsed, 2	QCs 60	70	80 R	90 Relative Co	100 oncentration
		Sam	ble		Leve	el	Enable	d	Expected	Final (	Concer	ntration	Accuracy

Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
Cal 1 MJ	1	~	1.0	1.0	101.2
Cal 2 MJ	2	~	3.0	2.9	96.6
Cal 3 MJ	3	~	5.0	4.9	98.6
Cal 4 MJ	4	~	10.0	10.2	101.8
Cal 5 MJ	5	~	25.0	25.9	103.7
Cal 6 MJ	6	~	50.0	49.0	98.1
Cal 7 MJ	7	~	100.0	100.0	100.0



Batch resultsD:\MassHunter\Data\2023\AM 27 28\060623 AM 27 28 TS\QuantResults\AM 27.batch.binCalibration Last Update6/8/2023 8:16:48 AM

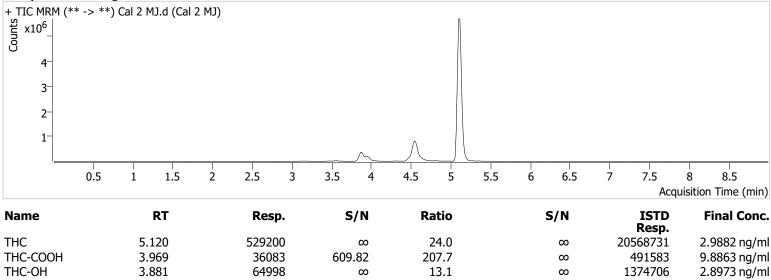
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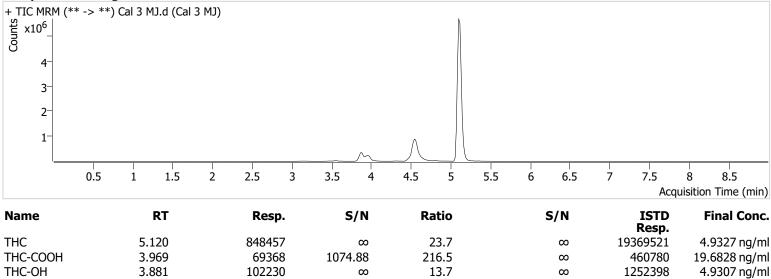
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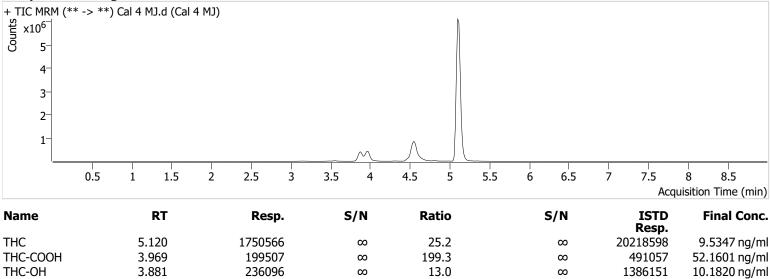
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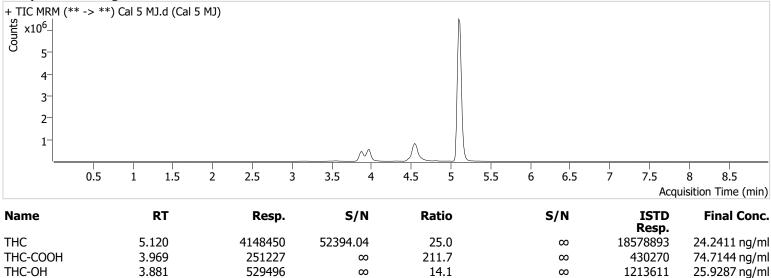
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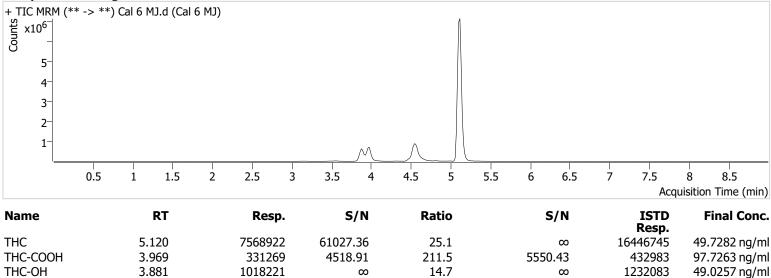
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Batch resultsD:\MassHunter\Data\2023\AM 27 28\060623 AM 27 28 TS\QuantResults\AM 27.batch.binCalibration Last Update6/8/2023 8:16:48 AM

Instrument Type Acq. Method Sample Position Injection Volume Acq. Date-Time Sample Info. Falco (069901) Cal AM 27 Agilent Method.m P1-F1 10 6/7/2023 2:53:03 PM Data File Sample Operator Comment Cal 6 MJ.d Cal 6 MJ Tamara Salazar Only drugs and concentrations listed on the laboratory report itself are appropriate to be used for interpretation purposes. Any drugs or values included in the notes but not included on the report are used by laboratory personnel to make determinations/reach conclusions within the confines of the methods.





 Batch results
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 Calibration Last Update
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Instrument Type Acq. Method Sample Position Injection Volume Acq. Date-Time Sample Info. Falco (069901) Cal AM 27 Agilent Method.m P1-G1 10 6/7/2023 3:06:07 PM Data File Sample Operator Comment Cal 7 MJ.d Cal 7 MJ Tamara Salazar Only drugs and concentrations listed on the laboratory report itself are appropriate to be used for interpretation purposes. Any drugs or values included in the notes but not included on the report are used by laboratory personnel to make determinations/reach conclusions within the confines of the methods.

