








CS

6/9/2022

**REVIEWED**  
By Tamara Salazar at 2:14 pm, Jun 10, 2022

**Worklist: 5974**

<u>LAB CASE</u>	<u>ITEM</u>	<u>ITEM TYPE</u>	<u>DESCRIPTION</u>	
M2022-1179	2	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	
M2022-1322	2	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	
M2022-1889	3	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	
P2022-0879	4	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	
P2022-0997	2	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	
P2022-1440	1	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	
P2022-1466	1	UCK	AM 27 Urine Cannabinoids Confirmation by LC-QQQ	

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

Extraction Date: 05/25/2022

Plate lot#: 220309

**Mobile phase A:** 0.1% Formic Acid in LCMS Water

**Blank Blood Lot:** Lampire 20L20723

**LCMS-QQQ ID:** 069901

Analyst: Celena Shrum

Plate Retest Date: 09/09/2022

**Mobile phase B:** 0.1% Formic acid in Acetonitrile

**Column:** UCT Selectra DA 100 x 2.1mm 3um

**Blank Urine Lot:** POC021022

### Pre-Analytic:

- 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. Using a calibrated pipette, add **1000µl blood and urine (if applicable) (calibrated pipette)** into the appropriate wells of analytical (standards) plate. **Pipette ID: #42**
- 3. Place on shaking incubator at ambient temp., 900rpm for 15 minutes.
- 4. Pipette **500µL 0.1% formic acid in water blood sample** of analytical plate.
- 5. Place on shaking incubator at ambient temp., 900rpm for 15 minutes.
- 6. Transfer **800µL of blood+acid or urine+acid** mixture to corresponding wells of SLE+ plate.
- 7. 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)**
- 8. Wait 5 minutes.
- 9. Add **2.25mL MTBE. (Add in 3 increments of 750uL)**
- 10. Wait 5 minutes.
- 11. Apply positive pressure for approx. 15 seconds. **(10-15 PSI- Selector to the left).**
- 12. Add **2.25mL Hexane. (Add in 3 increments of 750uL)**
- 13. Wait 5 minutes.
- 14. Apply positive pressure for approx. 15 seconds. **(10-15 PSI- Selector to the left).**
- 15. Remove plate containing eluate. Place on SPE Dry and evaporate to dryness at approx. 35°C.
- 16. Reconstitute in **100µL 100% MeOH** and heat seal plate with foil. Place in autosampler and run worklist.

### Post-Analytic

- 1. Create batch and process data.
- 2. Make any necessary integration changes, Curve weighting of Linear 1/x with  $r^2$  values  $\geq 0.98$  for each analyte
- 3. 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
- 4. Case sample response for THC 1ng/mL and OH-THC 3ng/mL (quantitative), Carboxy-THC: 5ng/mL (qualitative only) will be reported. Samples with a THC or OH-THC response over 50 ng/mL will be reported out as greater than 50 ng/mL.
- 5. Did all QCs pass for each analyte? (if not, describe in comments section)
- 6. Enter QCs into control charting.
- 7. Central File Packet to include: LIMS Worklist, Method Checklist, Calibration and Control Reports

COMMENTS: Only THC and THC-COOH evaluated. Urine QC had to be reinjected due to it not injecting properly the first time. The sample was reinjected on 5/31/22 with no issues. A blood sample from a previous batch was also included in this run (M2022-1304-1).

	1	2	3	4	5	6
a	cal 1ng	QC 2	P2022-1440-1			
b	cal 3 ng	NEG Blood	P2022-1466-1			
c	cal 5 ng	Neg Urine	M2022-1304-1			
d	cal 10ng	M2022-1179-2				
e	cal 25 ng	M2022-1322-2				
f	cal 50 ng	M2022-1889-3				
g	cal 100 ng	P2022-0879-4				
h	QC 1	P2022-0997-2				

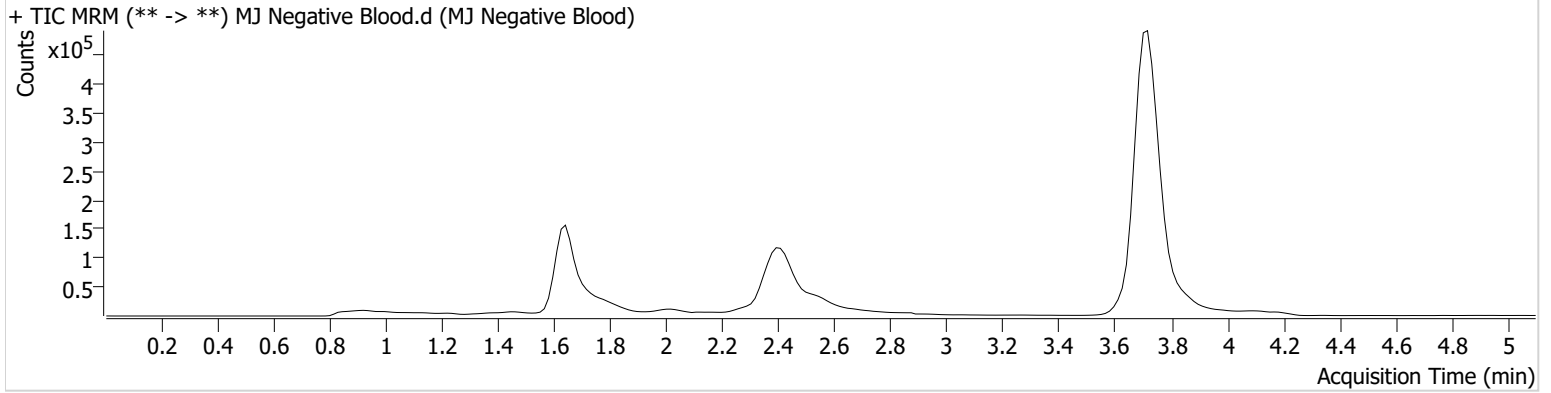
# AM #27 Cannabinoid Quant. Results



**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	MJ Negative Blood.d
<b>Type</b>	Sample	<b>Sample</b>	MJ Negative Blood
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-B2	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 3:35:47 PM		
<b>Sample Info.</b>			

## Sample Chromatogram



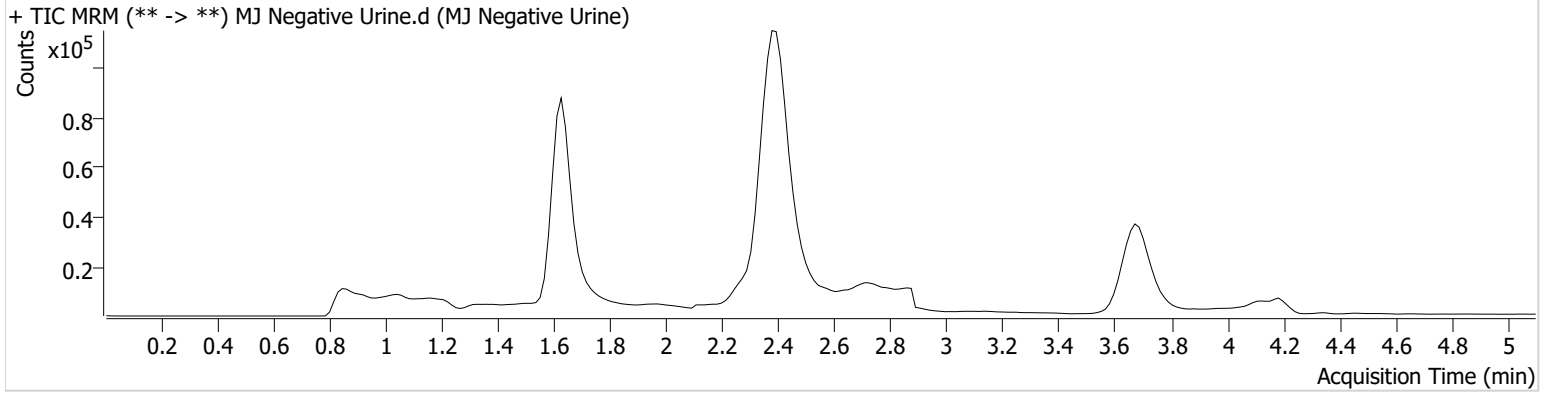
# AM #27 Cannabinoid Quant. Results



**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	MJ Negative Urine.d
<b>Type</b>	Sample	<b>Sample</b>	MJ Negative Urine
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-C2	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 3:43:24 PM		
<b>Sample Info.</b>			

## Sample Chromatogram



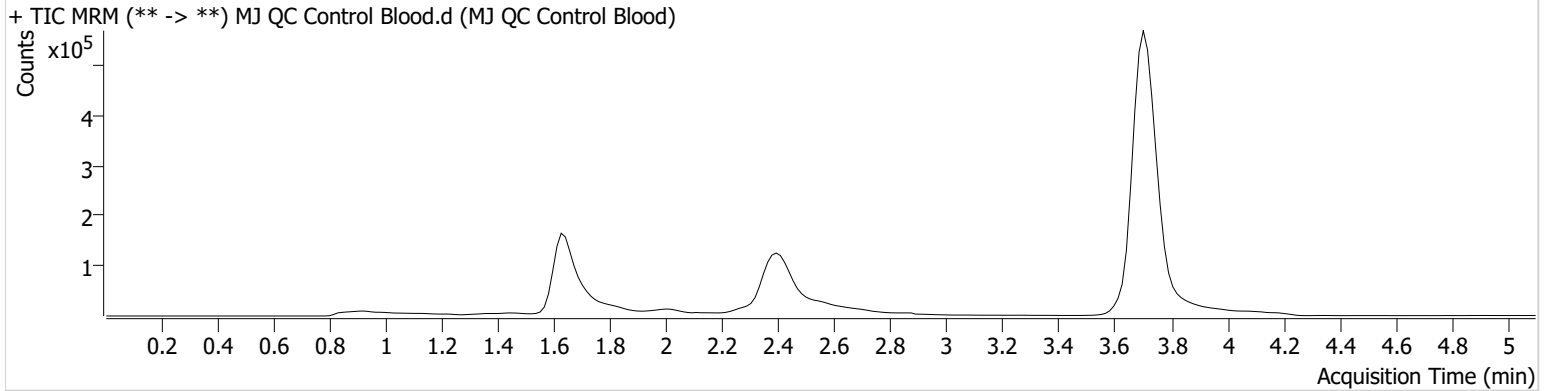


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

**Instrument** Falco (069901) **Data File** MJ QC Control Blood.d  
**Type** QC **Sample** MJ QC Control Blood  
**Acq. Method** AM 27 THCQ.m **Operator** Celena Shrum  
**Sample Position** P5-H1 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 5/25/2022 3:20:35 PM  
**Sample Info.**

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	56012	142.50	53.2	349.17	174457	15.7948 ng/ml
THC	3.721	130047	1431.24	29.4	∞	3517016	4.7324 ng/ml

# AM #27 Cannabinoid Quant. Results

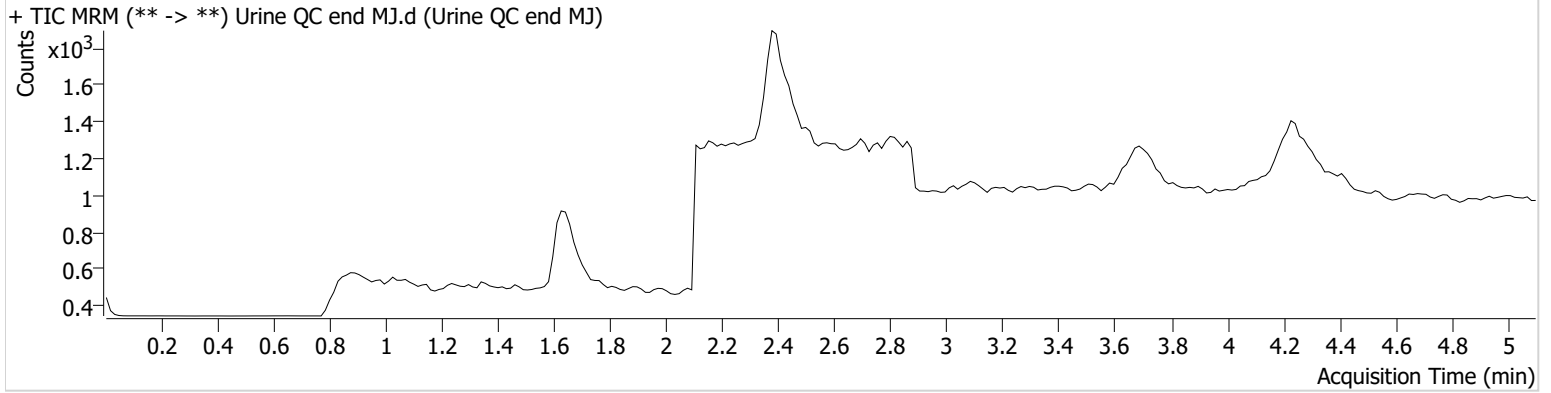


**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Urine QC end MJ.d
<b>Type</b>	QC	<b>Sample</b>	Urine QC end MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-A2	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 6:00:26 PM		
<b>Sample Info.</b>			

QC was reinjected due to an injection issue.

## Sample Chromatogram



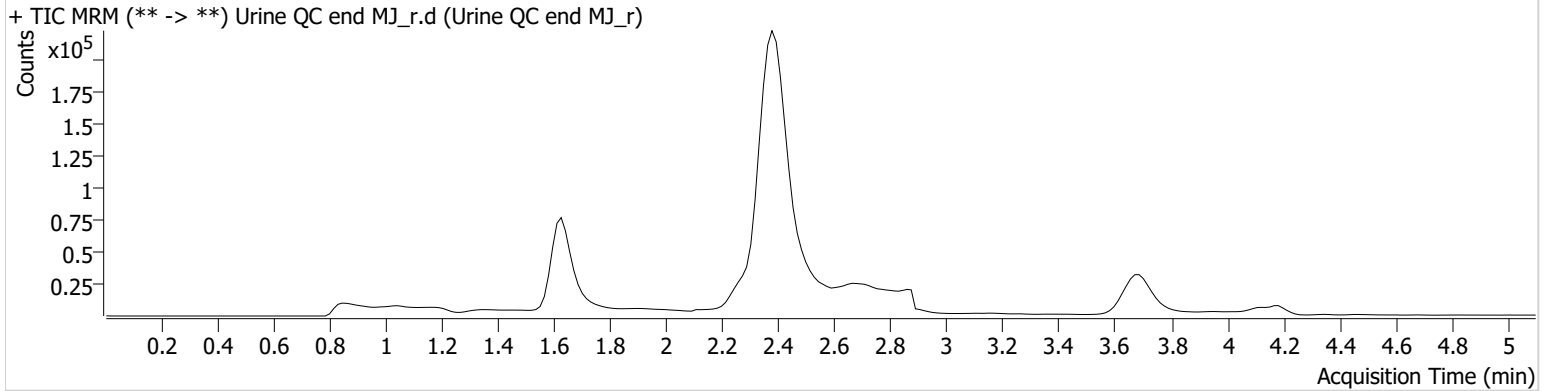


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

**Instrument** Falco (069901) **Data File** Urine QC end MJ\_r.d  
**Type** QC **Sample** Urine QC end MJ\_r  
**Acq. Method** AM 27 THCQ.m **Operator** Celena Shrum  
**Sample Position** P5-A2 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 5/31/2022 1:10:11 PM  
**Sample Info.** Reinjection data.

## Sample Chromatogram



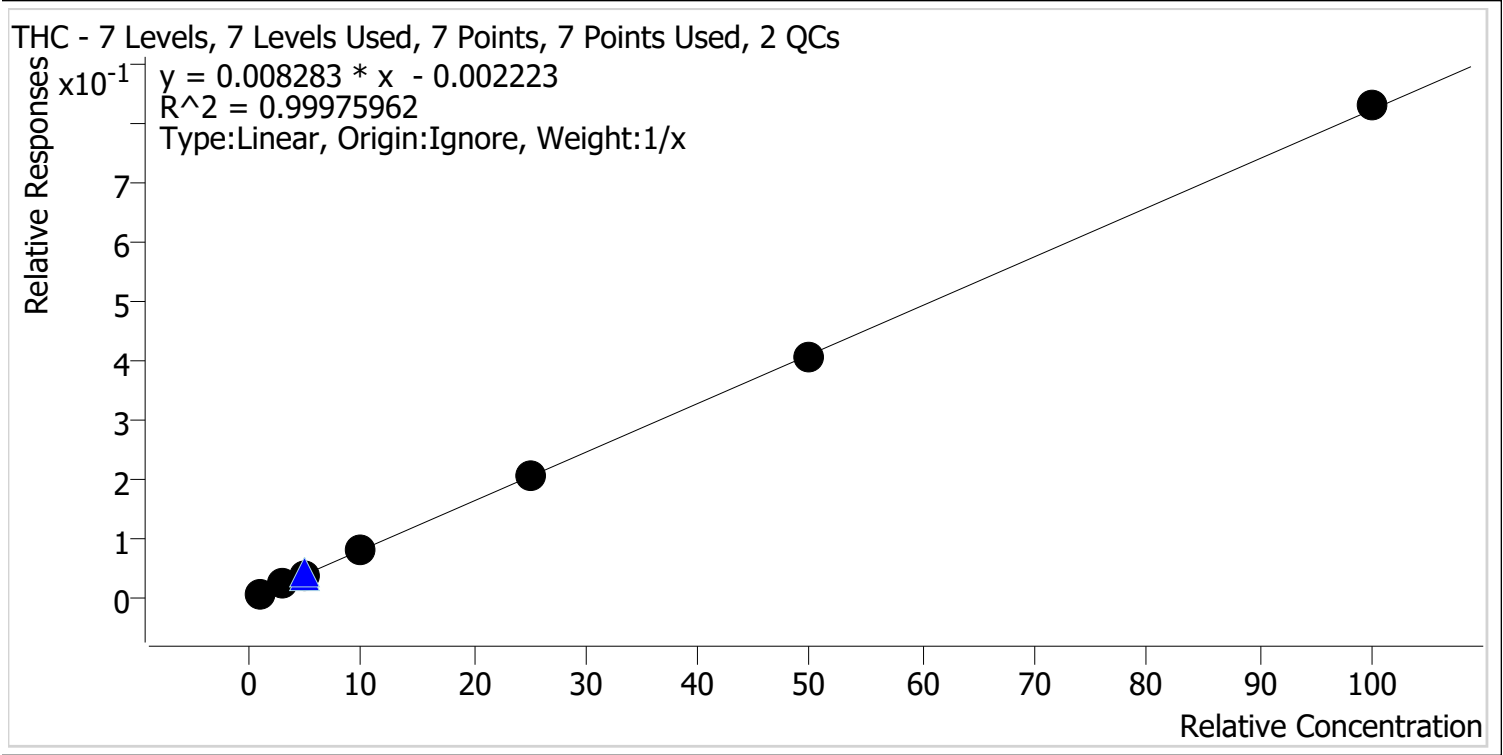
Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.655	26379	122.04	50.7	97.09	76139	16.9646 ng/ml
THC	3.706	8997	∞	32.8	10.18	221272	5.1771 ng/ml





# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Last Cal. Update** 6/9/2022 9:16 AM  
**Analyst Name** ISP\datastor  
**Analyte** THC **Internal Standard** THC-D3

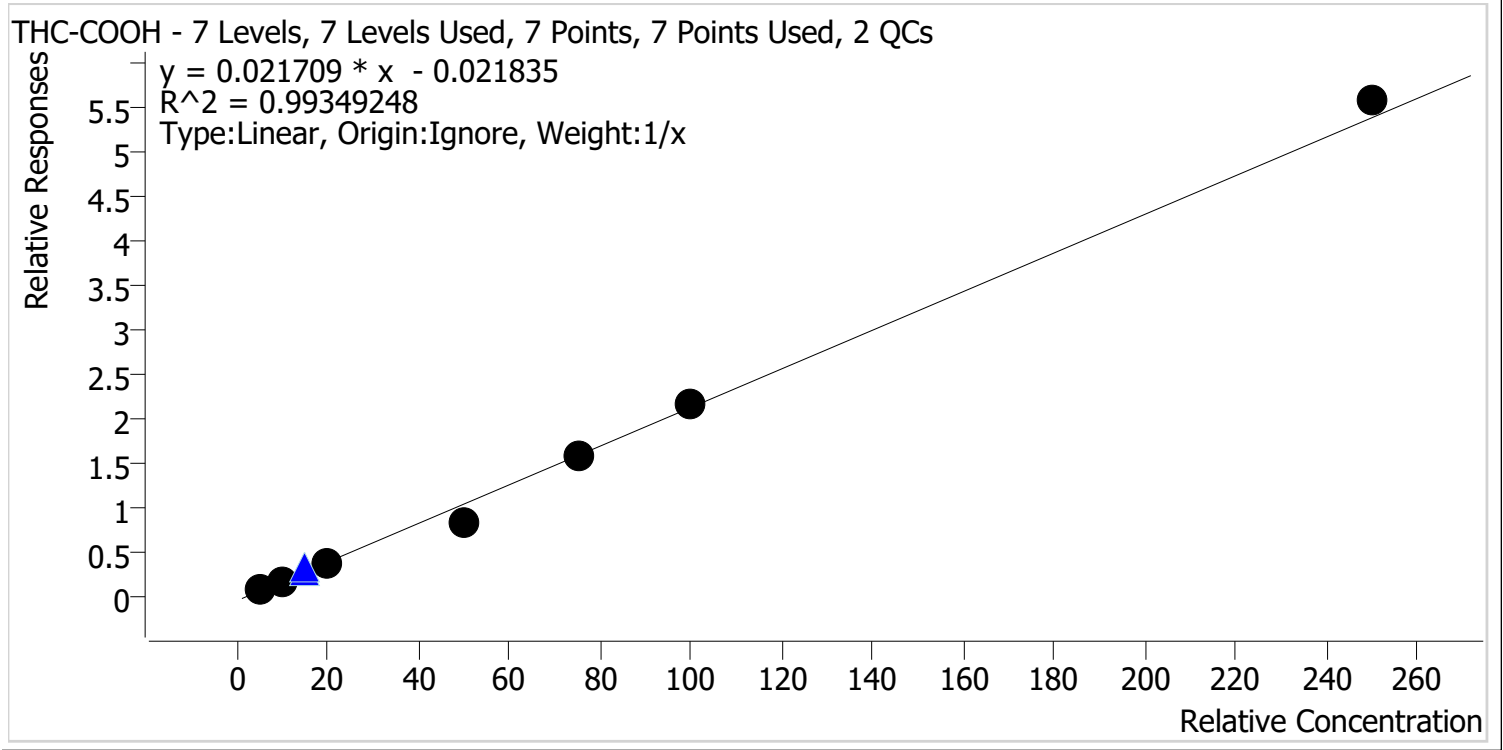


Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
Cal 1 MJ	1	✓	1.0	1.1	109.3
Cal 2 MJ	2	✓	3.0	3.0	98.4
Cal 3 MJ	3	✓	5.0	4.7	94.8
Cal 4 MJ	4	✓	10.0	9.7	97.4
Cal 5 MJ	5	✓	25.0	25.0	100.0
Cal 6 MJ	6	✓	50.0	49.6	99.3
Cal 7 MJ	7	✓	100.0	100.8	100.8



# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Last Cal. Update** 6/9/2022 9:16 AM  
**Analyst Name** ISP\datastor  
**Analyte** THC-COOH **Internal Standard** THC-COOH-D9



Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
Cal 1 MJ	1	✓	5.0	6.1	122.2
Cal 2 MJ	2	✓	10.0	9.4	93.9
Cal 3 MJ	3	✓	20.0	19.9	99.5
Cal 4 MJ	4	✓	50.0	40.2	80.4
Cal 5 MJ	5	✓	75.0	74.6	99.5
Cal 6 MJ	6	✓	100.0	101.0	101.0
Cal 7 MJ	7	✓	250.0	258.8	103.5

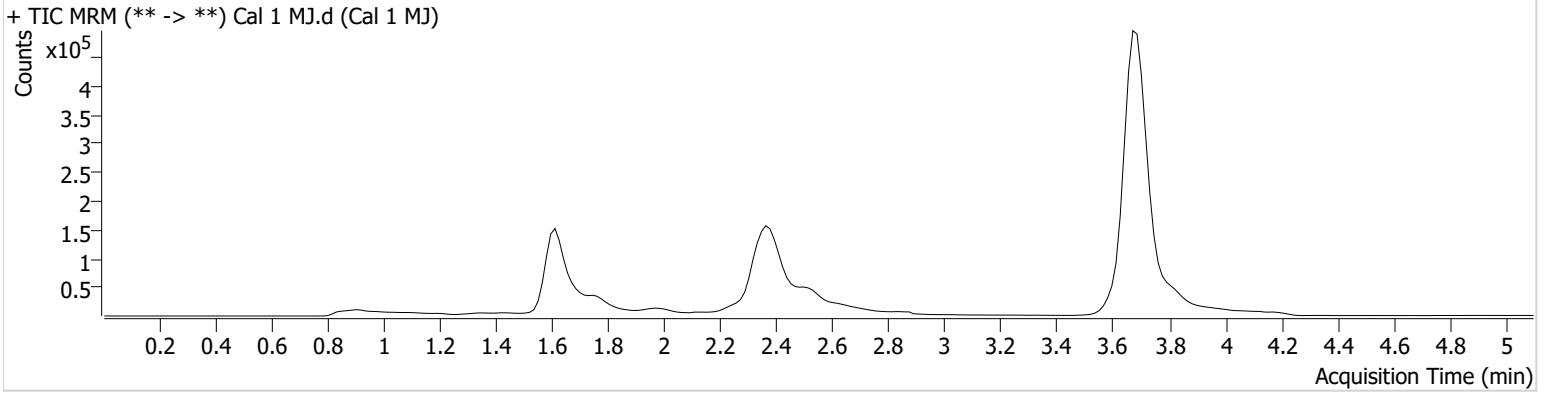


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

**Instrument** Falco (069901) **Data File** Cal 1 MJ.d  
**Type** Cal **Sample** Cal 1 MJ  
**Acq. Method** AM 27 THCQ.m **Operator** Celena Shrum  
**Sample Position** P5-A1 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 5/25/2022 2:19:34 PM  
**Sample Info.**

### Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.655	17588	∞	49.1	∞	158718	6.1100 ng/ml
THC	3.691	22676	123.18	32.0	22.03	3318620	1.0932 ng/ml

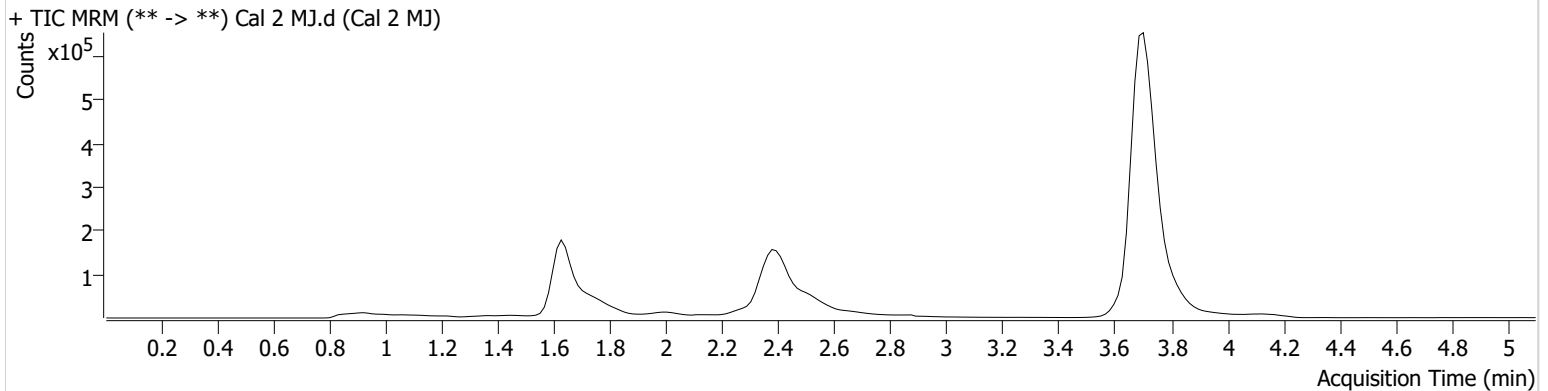
# AM #27 Cannabinoid Quant. Results



**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Cal 2 MJ.d
<b>Type</b>	Cal	<b>Sample</b>	Cal 2 MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-B1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 2:27:21 PM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	38675	75.55	58.7	331.38	212467	9.3906 ng/ml
THC	3.706	97248	1048.08	29.7	∞	4375423	2.9516 ng/ml

# AM #27 Cannabinoid Quant. Results

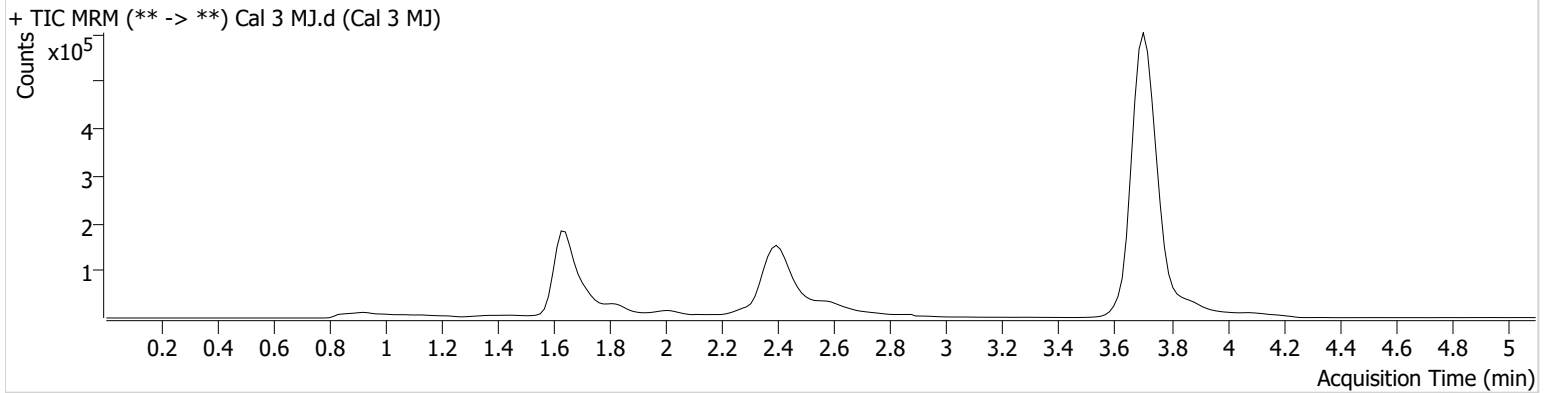


**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Cal 3 MJ.d
<b>Type</b>	Cal	<b>Sample</b>	Cal 3 MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-C1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 2:34:57 PM		

**Sample Info.**

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	74441	69.94	53.9	∞	181418	19.9067 ng/ml
THC	3.706	140741	750.43	29.1	142.31	3799766	4.7400 ng/ml

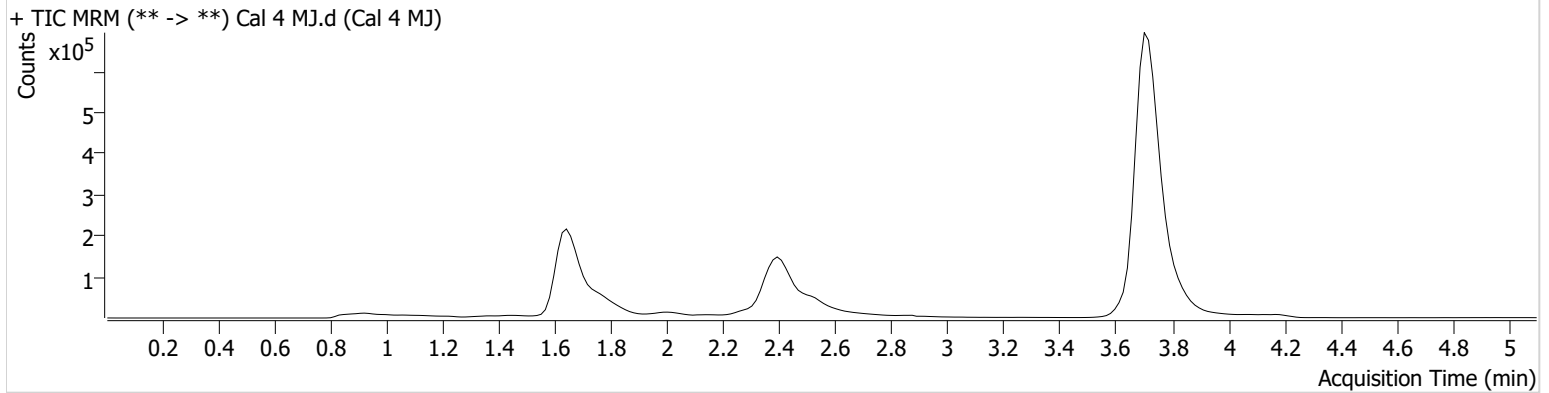
# AM #27 Cannabinoid Quant. Results



**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Cal 4 MJ.d
<b>Type</b>	Cal	<b>Sample</b>	Cal 4 MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-D1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 2:42:33 PM		

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	172888	∞	57.2	∞	203260	40.1857 ng/ml
THC	3.721	333836	∞	28.4	394.61	4257269	9.7351 ng/ml

CS

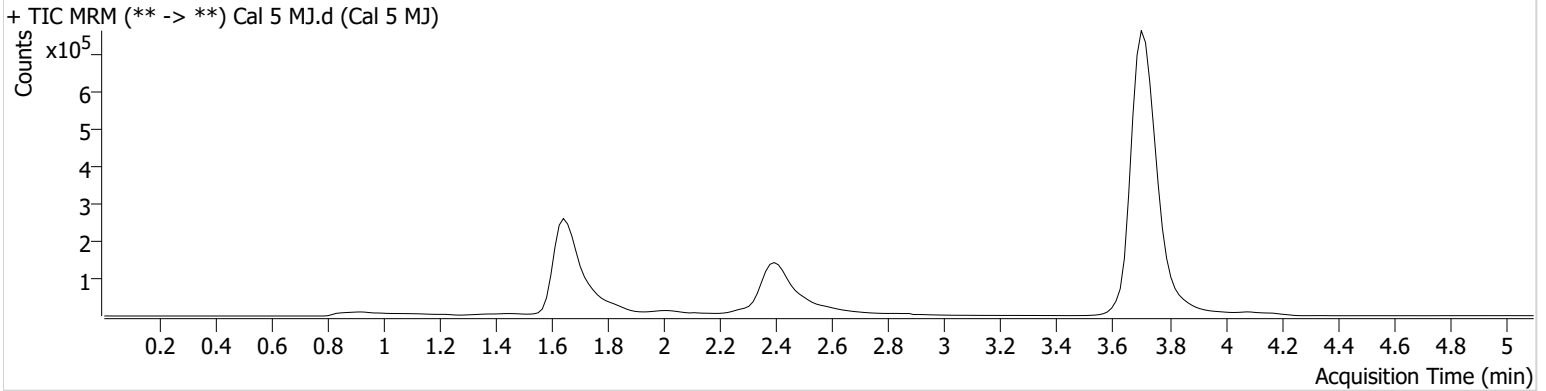


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Cal 5 MJ.d
<b>Type</b>	Cal	<b>Sample</b>	Cal 5 MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-E1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 2:50:08 PM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	304881	365.61	56.9	∞	190795	74.6121 ng/ml
THC	3.721	807837	∞	27.9	852.41	3941988	25.0089 ng/ml

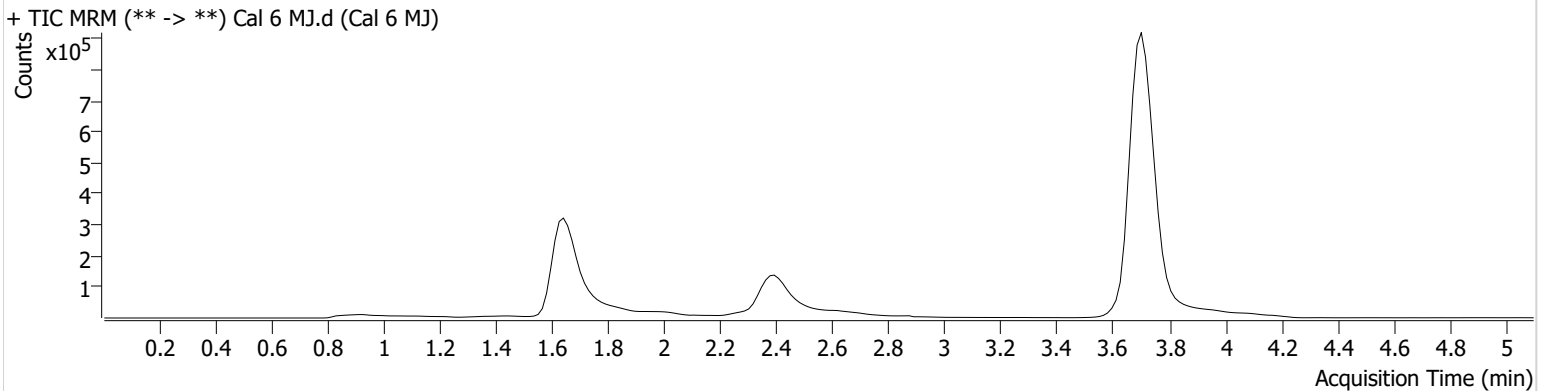
# AM #27 Cannabinoid Quant. Results



**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Cal 6 MJ.d
<b>Type</b>	Cal	<b>Sample</b>	Cal 6 MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-F1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 2:57:44 PM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	396122	846.45	57.1	2078.27	182512	100.9804 ng/ml
THC	3.706	1537977	∞	27.9	∞	3761257	49.6333 ng/ml



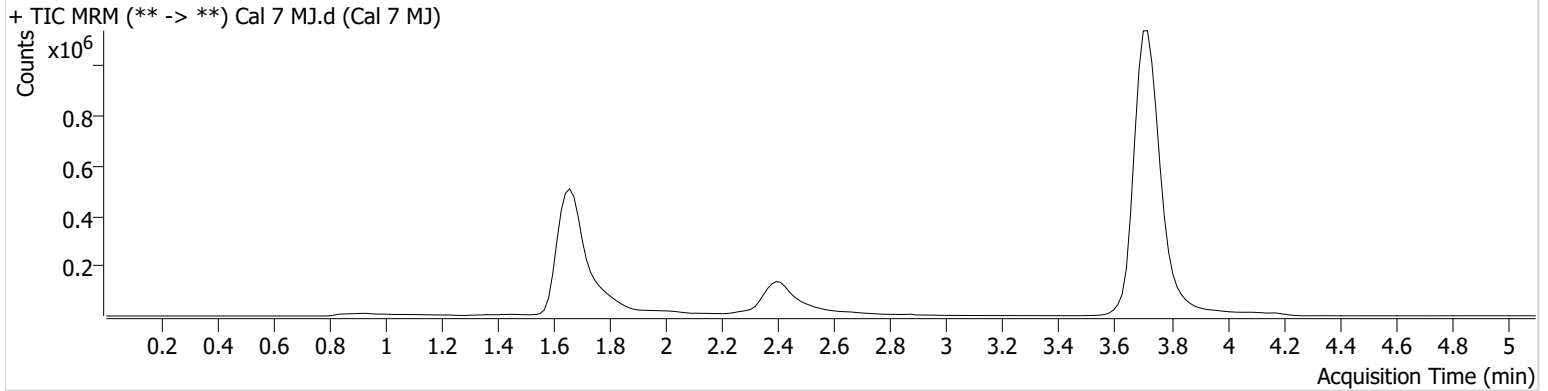
# AM #27 Cannabinoid Quant. Results



**Batch results** D:\MassHunter\Data\2022\AM 27-28\052522 AM 27 28 CS\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 6/9/2022 9:16:20 AM

<b>Instrument</b>	Falco (069901)	<b>Data File</b>	Cal 7 MJ.d
<b>Type</b>	Cal	<b>Sample</b>	Cal 7 MJ
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Celena Shrum
<b>Sample Position</b>	P5-G1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	5/25/2022 3:05:20 PM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-COOH	1.670	940617	∞	56.9	2730.20	168061	258.8145 ng/ml
THC	3.721	2966566	∞	28.4	3999.21	3561140	100.8379 ng/ml

**Idaho State Police  
Forensic Services**

**Request for Departure from an Analytical Method or Quality Standard**

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Deviation Number (assigned by QM): **TOX-22-02**

Date of Request:

03/02/2022

Requestor/Discipline:

Celena Shrum/Toxicology

Analytical Method/Quality Standard, Revision #:

Toxicology AM #25, AM #26, and AM #27, Revision 13

Temporary or Permanent Deviation:

Permanent

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**Scope of Deviation** (record specific information, e.g. affected programs, evidence types, expected end date; etc):

Deviation will remain in place until the change is made in the next method revision.

**Deviation Request** (Describe detailed instructions of the changes being made; include reference to specific section number(s) in the method manual):

Toxicology AM #25 3.3.1.1 Internal standards are prepared by the ToxBox plate manufacturer and contained on the 96 well plate. If the run contains urine samples, a positive external urine control must also be run.

Toxicology AM #26 3.3.2 A negative control will be run with each extraction. If the run contains urine samples, a negative urine control and external positive urine control must also be included.

Toxicology AM #27 3.3.2 A negative control will be run with each extraction. If the run contains urine samples, a negative urine control and positive external urine control will also be included in the run.

The deviation is to include the option of using an internal urine control in lieu of an external urine control.

**Technical Justification for Analytical Method Deviations:**

Internal controls serve the same purpose as external controls but also helps to avoid the possible issues that can occur with using external controls (incorrect spiking, incorrect preparation, evaporation of compounds, etc.). If these errors occur, runs need to be repeated and this wastes time, sample, and supplies.

**Technical Review**

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Departure approved

Comments:

Departure Not Approved

Comments:



Approver: Rachel Cutler

Date: 3/2/22

Title: Lab Manager

**Quality Review**

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Quality Approver: Jason Crowe

Title: Quality Manager

Date: 3/2/2022

