

**REVIEWED**

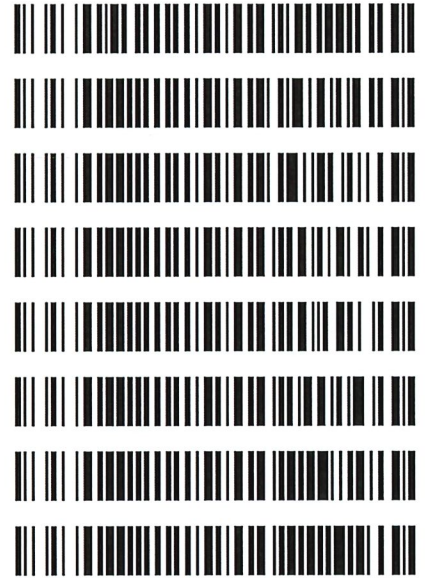
By Tamara Salazar at 1:19 pm, Aug 20, 2020

5 D

8/20/2020

**Worklist: 4435**

<u>LAB CASE</u>	<u>ITEM</u>	<u>ITEM TYPE</u>	<u>DESCRIPTION</u>
M2020-2912	2	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2194	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2274	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2304	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2308	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2324	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2368	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2020-2369	1	BCK	AM 27 Blood THC Quant by LC-QQQ



55  
P

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

Extraction Date: 08/14/20  
Plate lot#: IDP-108-2-200303

Analyst: Sarah Pickle  
Plate Expiration: 09/03/20

**Mobile phase A:** 0.1% Formic Acid in LCMS Water  
MTBE LCMS Methanol  
**Mobile phase B:** 0.1% Formic acid in Acetonitrile  
Hexane  
**Blank Blood Lot:** Hemostat 445283-4  
**LCMS-QQQ ID:** 069901  
**Column:** UCT Selectra DA 100 x 2.1mm 3um

## 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. Pipette **1000 µL blood (calibrated pipette)** in wells of analytical (standards) plate. **Pipette ID: 3382167**
- 3. Place on shaking incubator at ambient temp., 900rpm for 15 minutes.
- 4. Pipette **500 µL 0.1% formic acid in LCMS water** in wells of analytical plate.
- 5. Place on shaking incubator at ambient temp., 900rpm for 15 minutes.
- 6. Transfer **800 µL of blood+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 85-95 PSI- Selector to the right)*
- 8. Wait 5 minutes.
- 9. Add **2.25 mL MTBE. (Add in 3 increments of 750 µL)**
- 10. Wait 5 minutes.
- 11. Apply positive pressure for approx. 15 seconds. *(12-15 PSI- Selector to the left).*
- 12. Add **2.25 mL Hexane. (Add in 3 increments of 750 µL)**
- 13. Wait 5 minutes.
- 14. Apply positive pressure for approx. 15 seconds. *(12-15 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 **100 µL 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<sup>2</sup> values ≥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 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? Y / N
- 6. Enter QCs into control charting.
- 7. Central File Packet to include: LIMS Worklist, Method Checklist, Calibration and Control Reports

COMMENTS: *Hands of the analyst Sophie Jackson.*

*THC-OH was not evaluated.*

*Samples were inadvertently left on lab bench overnight unrefrigerated (8/12/20-8/13/20).*

# AM #27 Cannabinoid Quant. Results

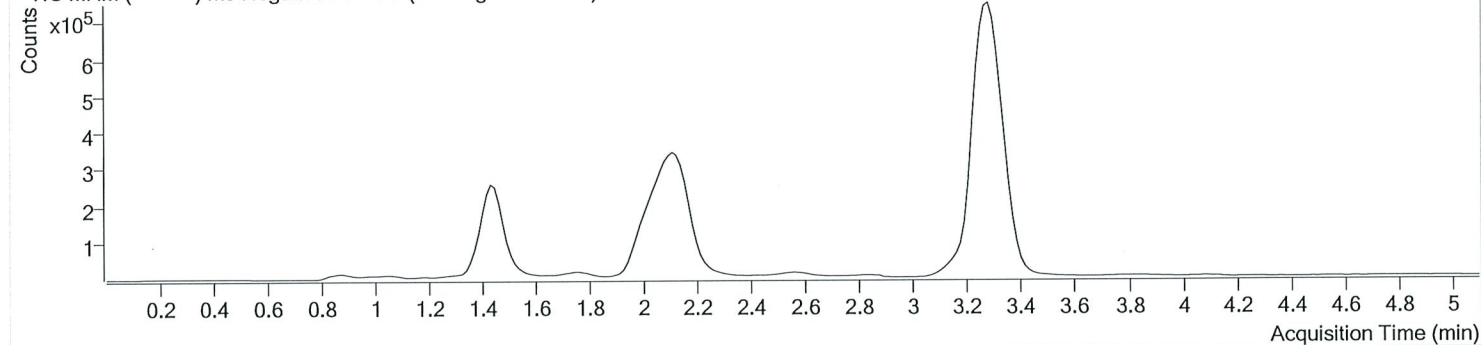


**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Calibration Last Update** 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Negative Blood.d
<b>Type</b>	Sample	<b>Sample</b>	MJ Negative Blood
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-H5	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 12:14:06 PM		
<b>Sample Info.</b>			

## Sample Chromatogram

+ TIC MRM (\*\* -> \*\*) MJ Negative Blood.d (MJ Negative Blood)



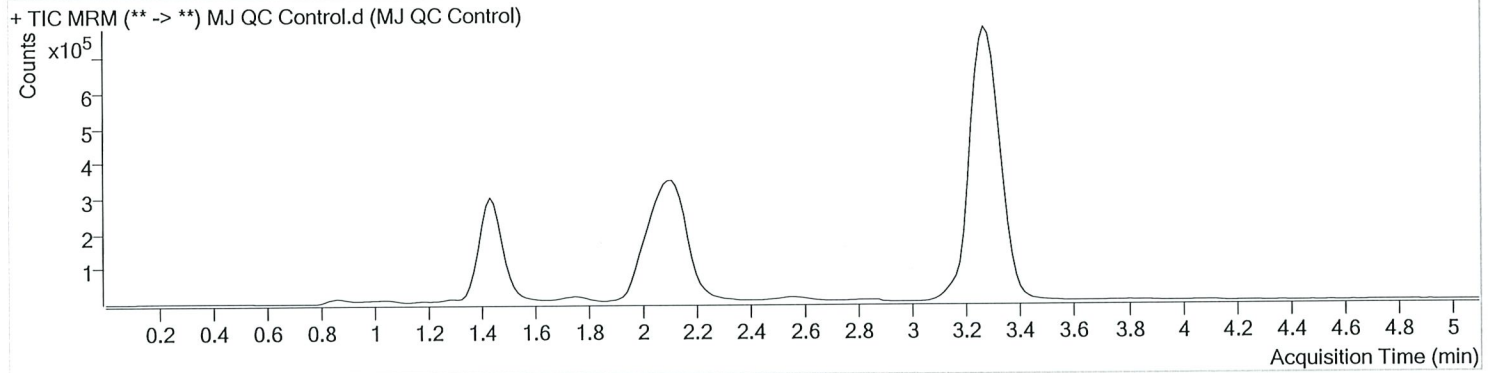


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Calibration Last Update** 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ QC Control.d
<b>Type</b>	Sample	<b>Sample</b>	MJ QC Control
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-A6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:58:54 AM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
* THC-OH	1.453	157468	∞	8.9	∞	1072021	3.0876 ng/ml
THC-COOH	1.459	133856	∞	50.7	235.76	371522	15.4933 ng/ml
THC	3.285	232977	1200.17	26.8	∞	5924427	4.2898 ng/ml

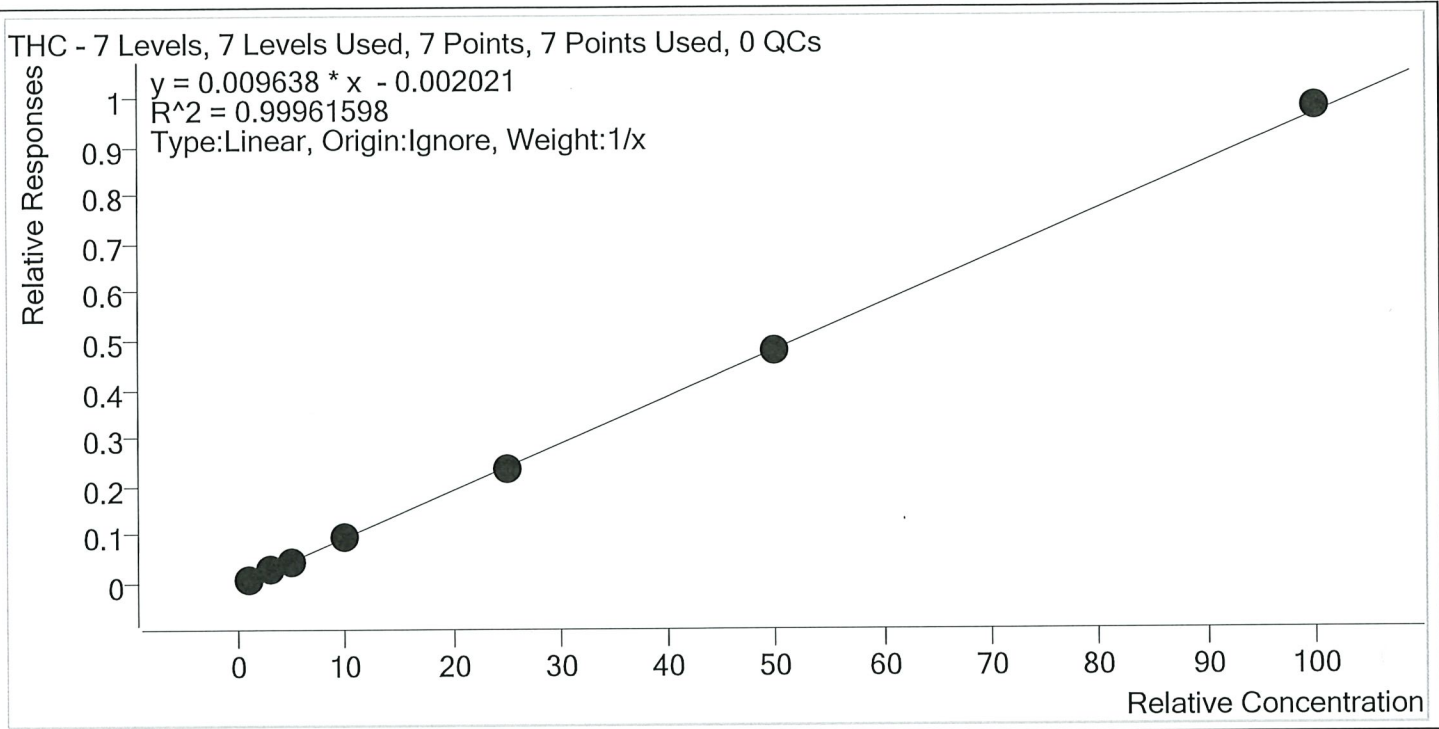
\*Not evaluated



BP

# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Last Cal. Update** 8/20/2020 8:23 AM  
**Analyst Name** ISP\datastor  
**Analyte** THC **Internal Standard** THC-D3



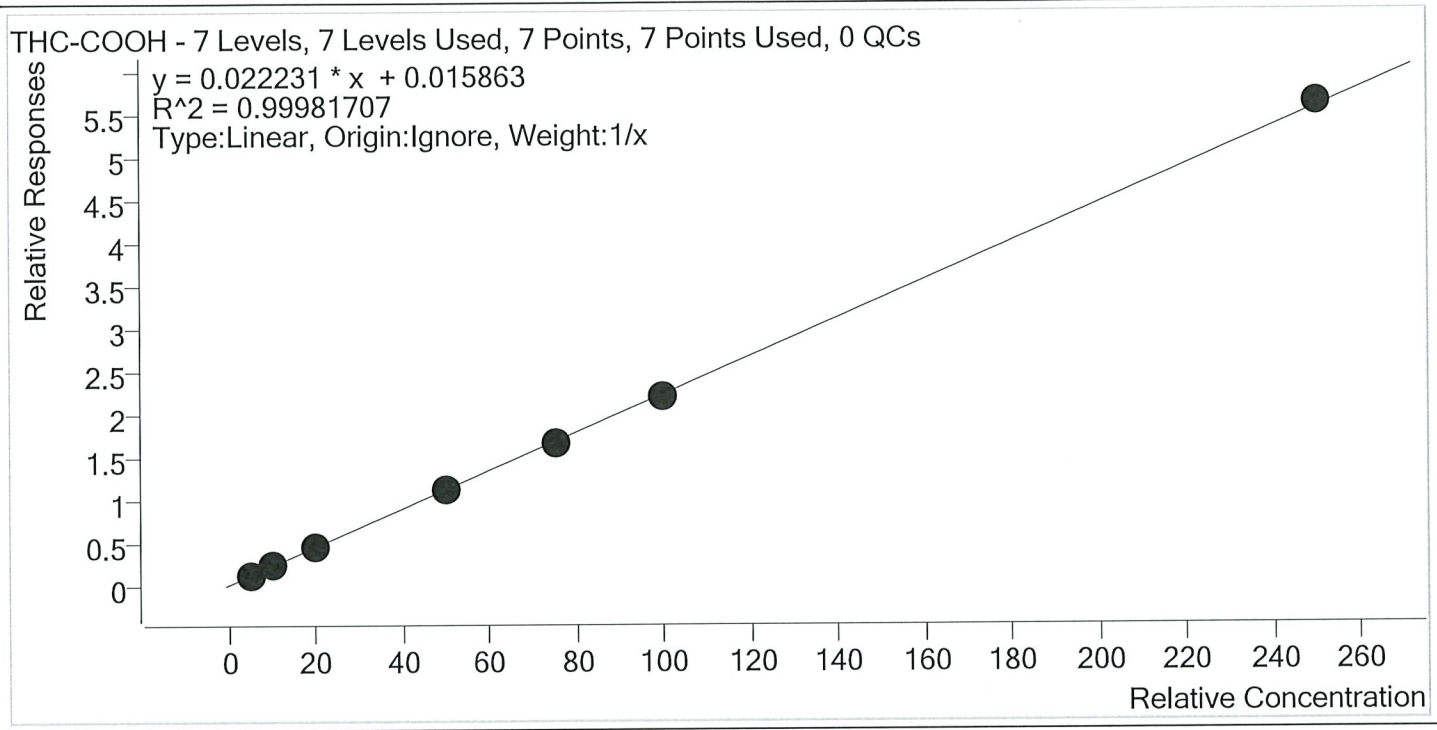
Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJ Cal 1	1	✓	1.0	1.1	109.7
MJ Cal 2	2	✓	3.0	2.9	96.7
MJ Cal 3	3	✓	5.0	4.9	98.0
MJ Cal 4	4	✓	10.0	9.8	97.6
MJ Cal 5	5	✓	25.0	24.4	97.6
MJ Cal 6	6	✓	50.0	49.5	98.9
MJ Cal 7	7	✓	100.0	101.5	101.5

SP



# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Last Cal. Update** 8/20/2020 8:23 AM  
**Analyst Name** ISP\datastor  
**Analyte** THC-COOH **Internal Standard** THC-COOH-D9



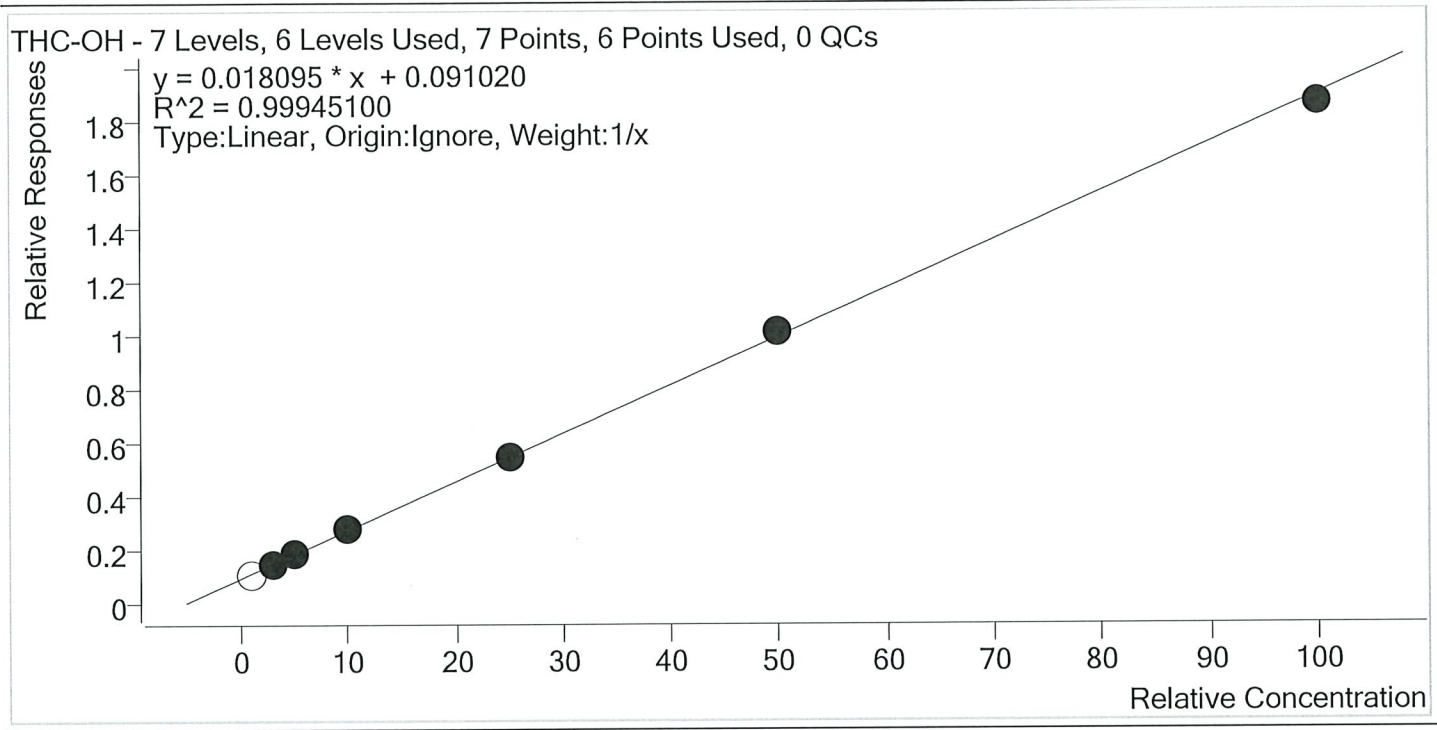
Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJ Cal 1	1	✓	5.0	5.0	100.6
MJ Cal 2	2	✓	10.0	10.3	102.8
MJ Cal 3	3	✓	20.0	19.7	98.6
MJ Cal 4	4	✓	50.0	49.8	99.7
MJ Cal 5	5	✓	75.0	74.2	98.9
MJ Cal 6	6	✓	100.0	98.5	98.5
MJ Cal 7	7	✓	250.0	252.5	101.0

SR



# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Last Cal. Update** 8/20/2020 8:23 AM  
**Analyst Name** ISP\datastor  
**Analyte** THC-OH **Internal Standard** THC-OH-D3



Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJ Cal 1	1	x	1.0	0.8	81.5
MJ Cal 2	2	✓	3.0	2.9	95.1
MJ Cal 3	3	✓	5.0	5.1	101.1
MJ Cal 4	4	✓	10.0	10.1	101.2
MJ Cal 5	5	✓	25.0	25.5	101.9
MJ Cal 6	6	✓	50.0	51.2	102.4
MJ Cal 7	7	✓	100.0	98.3	98.3

303

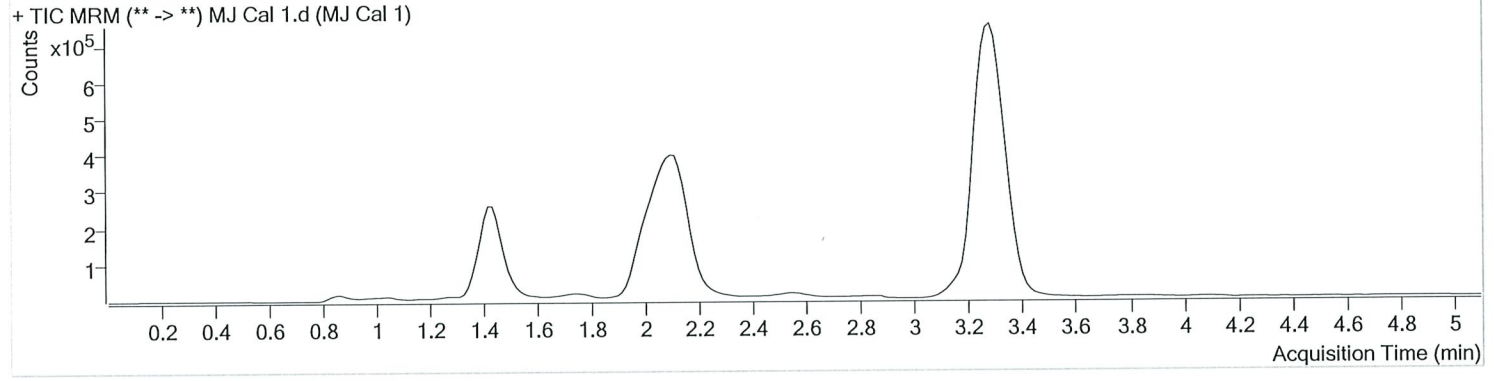


# AM #27 Cannabinoid Quant. Results

Batch results D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
Calibration Last Update 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Cal 1.d
<b>Type</b>	Cal	<b>Sample</b>	MJ Cal 1
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-H6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:05:40 AM		

### Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.483	113631	∞	4.5 <b>Low</b>	11.95	1074285	0.8153 ng/ml <b>Low</b>
THC-COOH	1.459	48135	∞	47.4	∞	377093	5.0284 ng/ml
THC	3.285	53744	46.51	29.5	106.73	6283457	1.0971 ng/ml <b>Low</b>



# AM #27 Cannabinoid Quant. Results

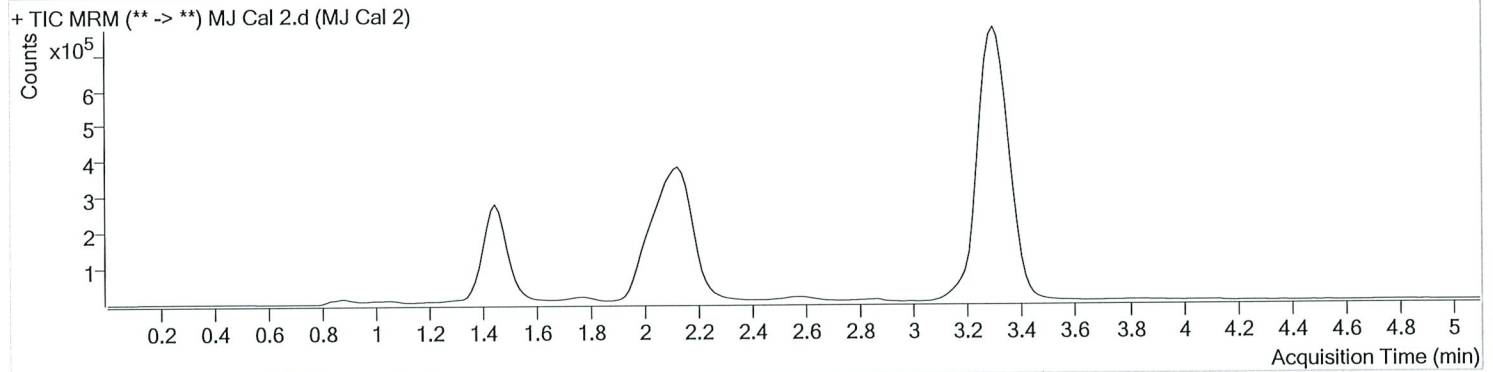
SJ  
P



Batch results D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
 Calibration Last Update 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Cal 2.d
<b>Type</b>	Cal	<b>Sample</b>	MJ Cal 2
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-G6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:13:25 AM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.498	153002	∞	6.8 <b>Low</b>	∞	1072668	2.8526 ng/ml <b>Low</b>
THC-COOH	1.474	92763	∞	52.2	∞	379473	10.2825 ng/ml
THC	3.315	158729	423.79	25.0	50.30	6115999	2.9024 ng/ml <b>Low</b>

# AM #27 Cannabinoid Quant. Results

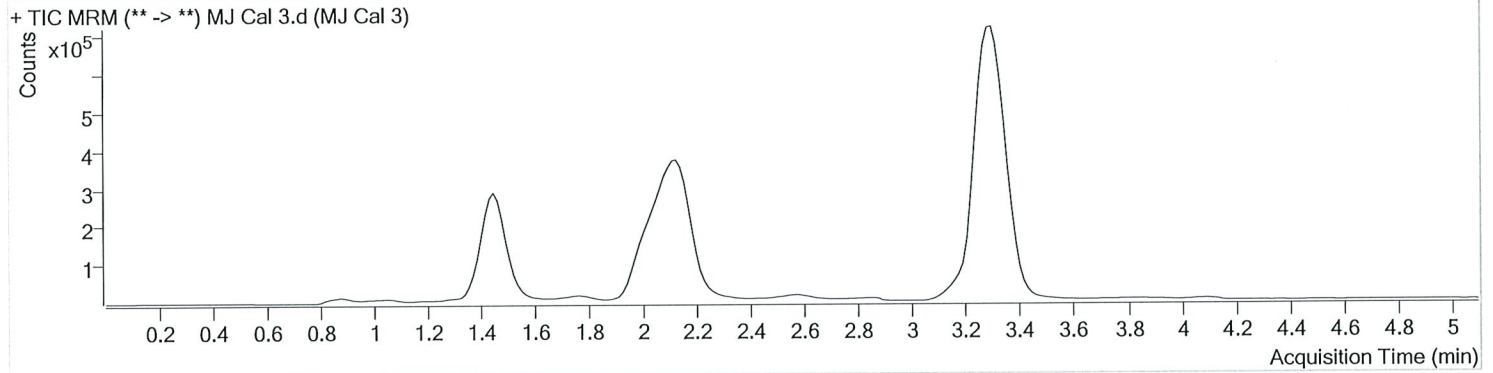


SJ  
P

**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Calibration Last Update** 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Cal 3.d
<b>Type</b>	Cal	<b>Sample</b>	MJ Cal 3
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-F6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:20:59 AM		

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.453	189497	∞	8.2	∞	1038319	5.0558 ng/ml
THC-COOH	1.474	165885	∞	56.6	∞	365221	19.7177 ng/ml
THC	3.300	254045	646.15	26.0	248.74	5621889	4.8982 ng/ml

# AM #27 Cannabinoid Quant. Results

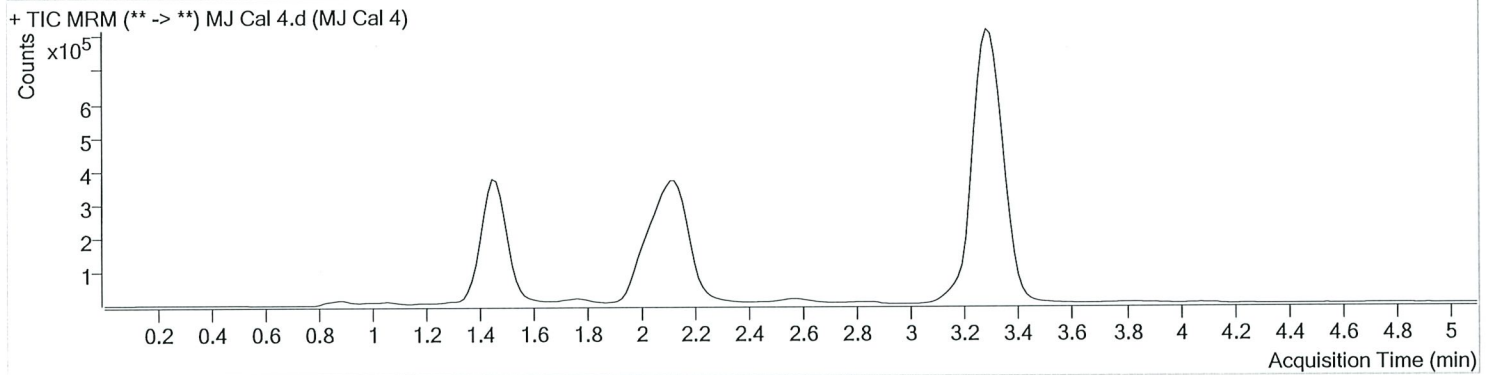
BP



**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Calibration Last Update** 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Cal 4.d
<b>Type</b>	Cal	<b>Sample</b>	MJ Cal 4
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-E6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:28:33 AM		

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.453	291819	∞	9.6	242.51	1064808	10.1155 ng/ml
THC-COOH	1.474	414965	∞	55.7	∞	369325	49.8280 ng/ml
THC	3.300	542990	3120.07	25.6	88.94	5899046	9.7600 ng/ml

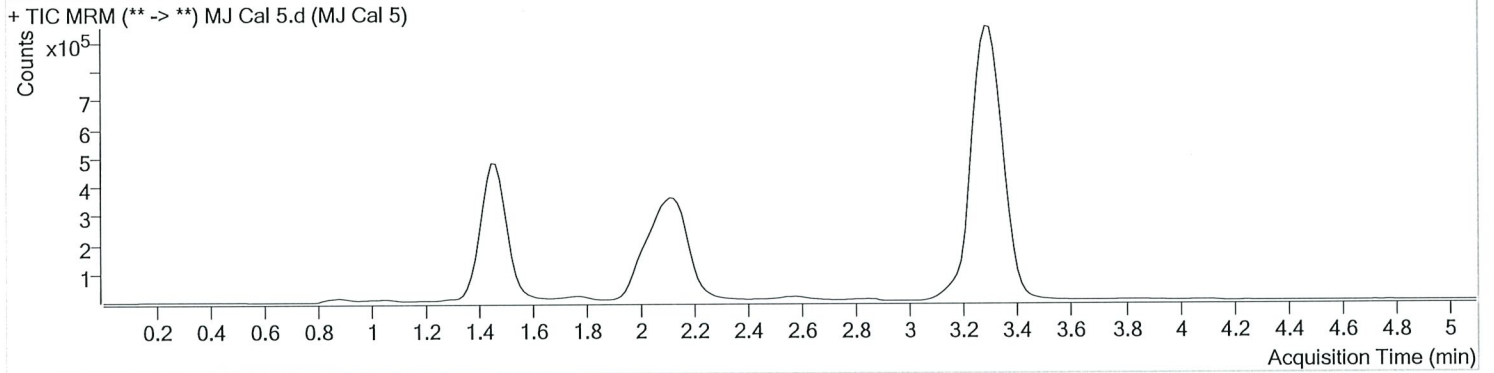
# AM #27 Cannabinoid Quant. Results



Batch results D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
Calibration Last Update 8/20/2020 8:23:52 AM

Instrument Falco Data File MJ Cal 5.d  
Type Cal Sample MJ Cal 5  
Acq. Method AM 27 THC quant.m Operator Sarah Pickle  
Sample Position P3-D6 Comment  
Injection Volume 10  
Acq. Date-Time 8/14/2020 11:36:07 AM  
Sample Info.

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.453	599444	∞	10.8	323.43	1085662	25.4839 ng/ml
THC-COOH	1.474	621144	∞	55.7	∞	373099	74.1747 ng/ml
THC	3.300	1410102	2762.79	25.9	∞	6050075	24.3918 ng/ml

# AM #27 Cannabinoid Quant. Results

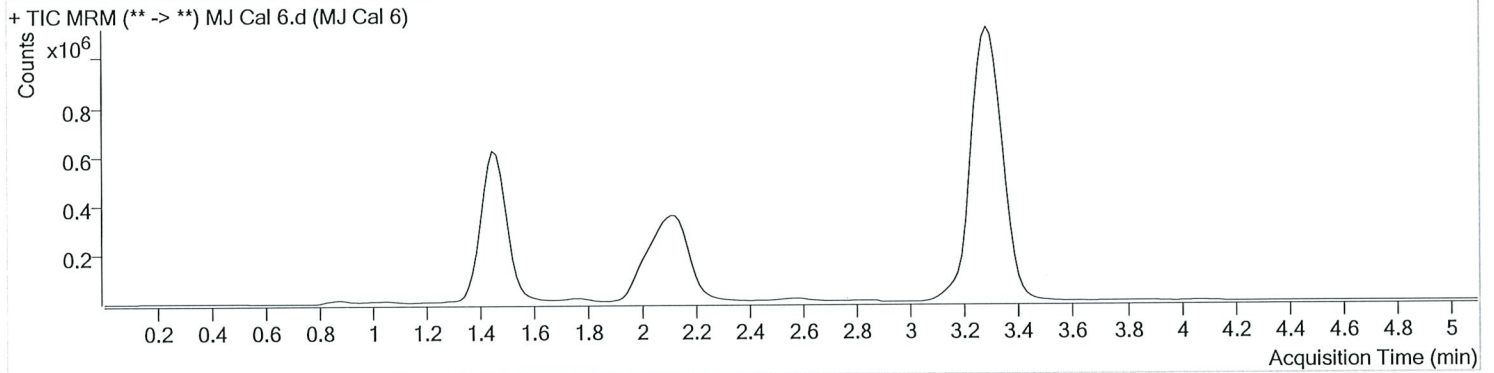
SP 55



**Batch results** D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
**Calibration Last Update** 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Cal 6.d
<b>Type</b>	Cal	<b>Sample</b>	MJ Cal 6
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-C6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:43:42 AM		
<b>Sample Info.</b>			

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.438	1087639	∞	12.0	∞	1068643	51.2166 ng/ml
THC-COOH	1.474	811683	∞	59.9	∞	368164	98.4589 ng/ml
THC	3.285	2746465	22684.21	25.7	67.28	5786333	49.4563 ng/ml

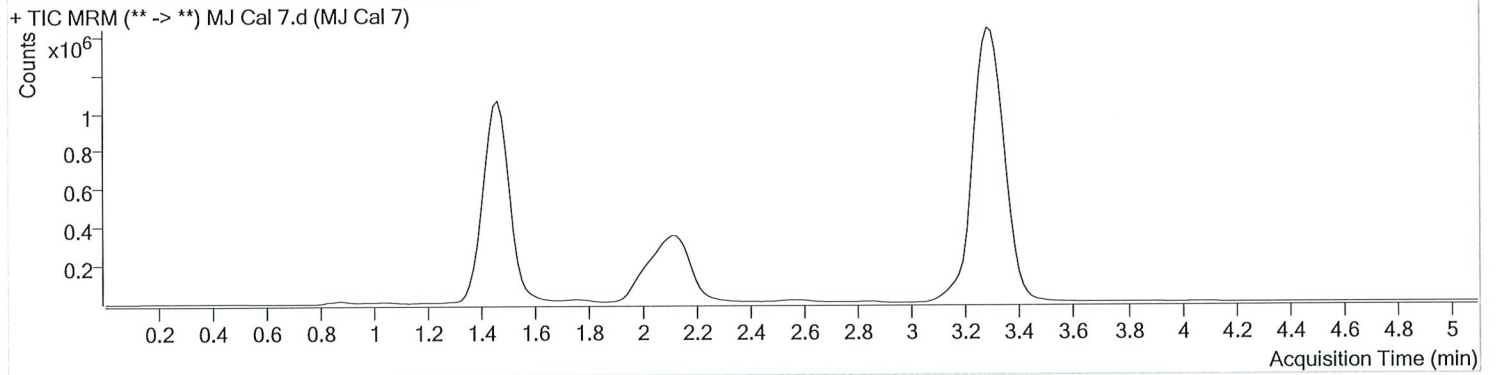
# AM #27 Cannabinoid Quant. Results



Batch results D:\MassHunter\Data\2020\AM 27-28\081420 AM 25 reinjects AM 27 SJ SP\QuantResults\AM 27 SJ SP.batch.bin  
 Calibration Last Update 8/20/2020 8:23:52 AM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ Cal 7.d
<b>Type</b>	Cal	<b>Sample</b>	MJ Cal 7
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Operator</b>	Sarah Pickle
<b>Sample Position</b>	P3-B6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	8/14/2020 11:51:17 AM		
<b>Sample Info.</b>			

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.438	1912258	∞	12.1	∞	1022979	98.2757 ng/ml
THC-COOH	1.474	1940284	∞	60.3	∞	344673	252.5098 ng/ml
THC	3.300	5235347	19891.33	25.2	2430.91	5362999	101.4942 ng/ml