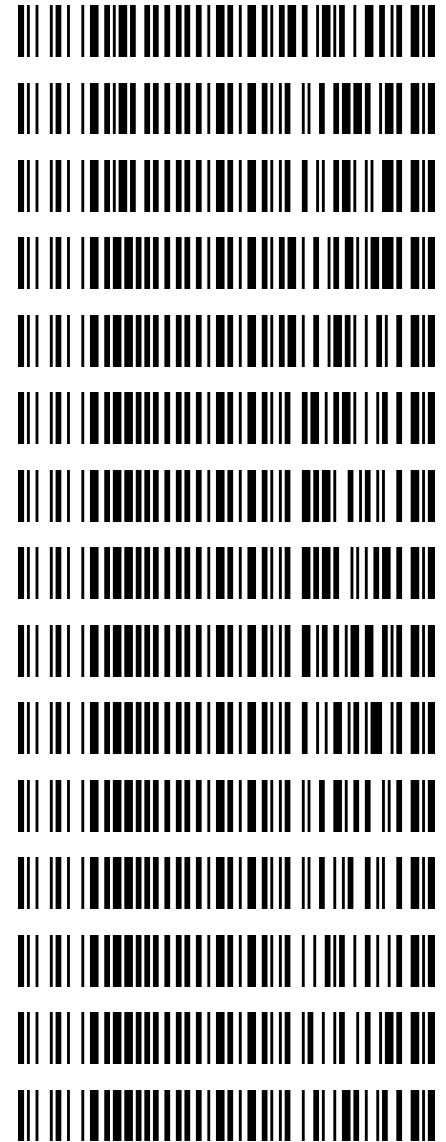


SC

2/24/2021

**Worklist: 4802**

<u>LAB_CASE</u>	<u>ITEM</u>	<u>ITEM_TYPE</u>	<u>DESCRIPTION</u>
M2021-0182	1	BCK	AM 27 Blood THC Quant by LC-QQQ
M2021-0404	2	BCK	AM 27 Blood THC Quant by LC-QQQ
M2021-0434	3	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0095	2	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0101	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0245	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0278	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0279	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0282	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0288	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0290	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0294	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0315	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0317	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2021-0325	1	BCK	AM 27 Blood THC Quant by LC-QQQ



8C

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

Extraction Date: 02/23/21  
Plate lot#: IDP-108-2-201206

Analyst: Sarah Collins  
Plate Expiration: 06/06/21

**Mobile phase A:** 0.1% Formic Acid in LCMS Water  
MTBE  
**Mobile phase B:** 0.1% Formic acid in Acetonitrile  
Hexane  
**Blank Blood Lot:** Lampire 20L20724  
**LCMS-QQQ ID:** 069901  
**Column:** UCT Selectra DA 100 x 2.1mm 3um  
LCMS Methanol

### 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: 16**
- 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  $r^2$  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? Y / N
- 6. Enter QCs into control charting.
- 7. Central File Packet to include: LIMS Worklist, Method Checklist, Calibration and Control Reports

COMMENTS: *Curve range limited: THC 3-100 Reconstituted and reinjected Calibrator 6 due to low internal standard response in initial injection. Reinject data used.*

Calibrator 1 dropped for THC due to high ratio

8C

**Idaho State Police  
Forensic Services  
Toxicology Discipline**

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**Request for Departure from an Analytical Method**

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Date of Request

02/24/2021

Forensic Scientist

Anne Nord

Analytical Method

Toxicology AM #27: Quantitative Analysis of THC and Metabolites in Blood and Urine by LCMS-QQQ

Request

The method currently reads:

4.3.2.5 If any points are dropped from the approved quantitative range of the curve, the compound will be reported qualitatively. For calibrators and controls 10 ng and below, the accuracy must be within 30%, for calibrators and controls greater than 10 ng/mL the accuracy must be within 20%. If a control falls outside the accuracy range, at the analyst's discretion, the compound may be reported qualitatively.

I would like to add in the following exception:

If the 1ng/ml point is dropped for THC. If the 1 ng/ml point is dropped the quantitative range will be 3-50 ng/ml.

**Discipline Leader Review**

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Departure approved

Comments: This deviation is approved and will remain in effect until it is changed in the actual method.

Departure Not Approved

Comments:



Celena Shrum

Toxicology Discipline Lead

Date: 02/24/2021

8C

	1	2	3	4	5	6
A	IS + Cal. 1	negative	p2021-0279-1			IS + QC_1
B	IS + Cal. 2	m2021-0182-1	p2021-0282-1			IS + Cal. 7
C	IS + Cal. 3	m2021-0404-2	p2021-0288-1			IS + Cal. 6
D	IS + Cal. 4	m2021-0434-3	p2021-0290-1			IS + Cal. 5
E	IS + Cal. 5	p2021-0095-2	p2021-0294-1			IS + Cal. 4
F	IS + Cal. 6	p2021-0101-1	p2021-0315-1			IS + Cal. 3
G	IS + Cal. 7	p2021-0245-1	p2021-0317-1			IS + Cal. 2
H	IS + QC_1	p2021-0278-1	p2021-0325-1			IS + Cal. 1

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

8C

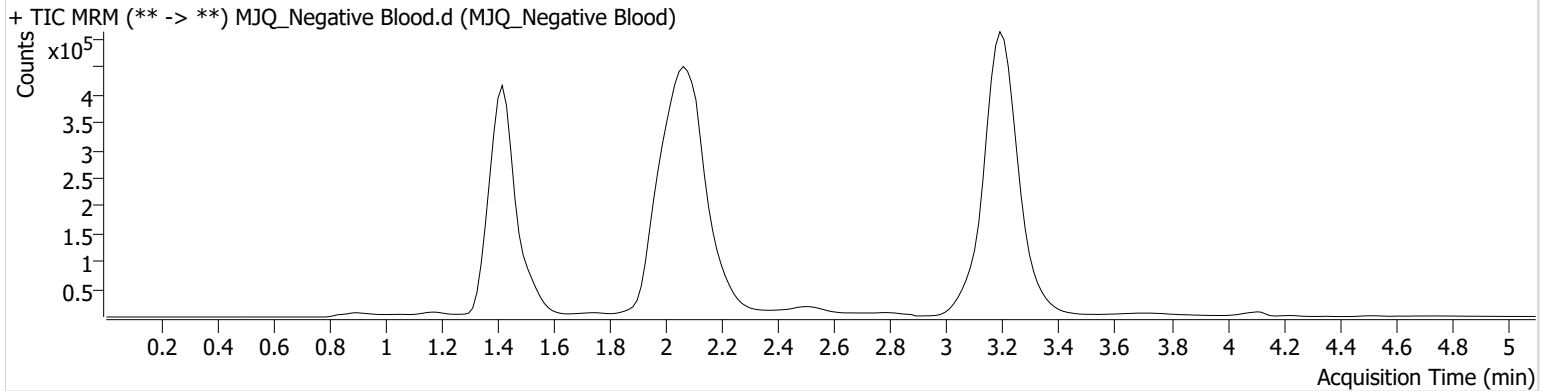


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument Type</b>	Instrument 1 Sample	<b>Data File</b>	MJQ_Negative Blood.d
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Sample Operator</b>	MJQ_Negative Blood Sarah Collins
<b>Sample Position</b>	P1-A2	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 7:11:17 PM		
<b>Sample Info.</b>			

## Sample Chromatogram



8C

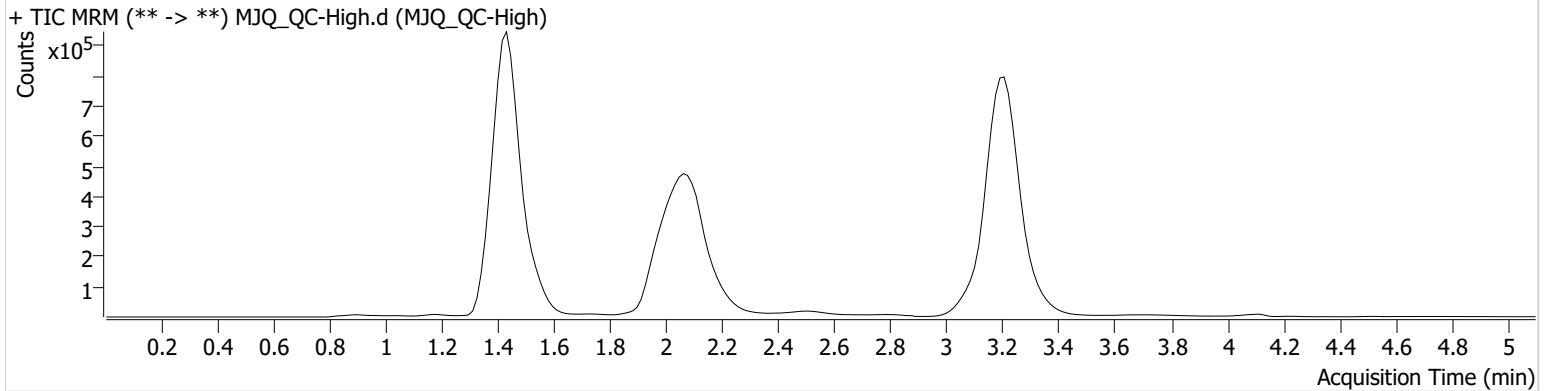


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_QC-High.d
<b>Type</b>	Sample	<b>Sample</b>	MJQ_QC-High
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-F1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 7:26:27 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	1419720	∞	11.6	∞	2188384	49.8632 ng/ml
THC-COOH	1.444	1358348	∞	59.1	5778.93	577231	99.7047 ng/ml
THC	3.209	2035183	6308.31	24.6	1312.32	4425433	50.0946 ng/ml

8C

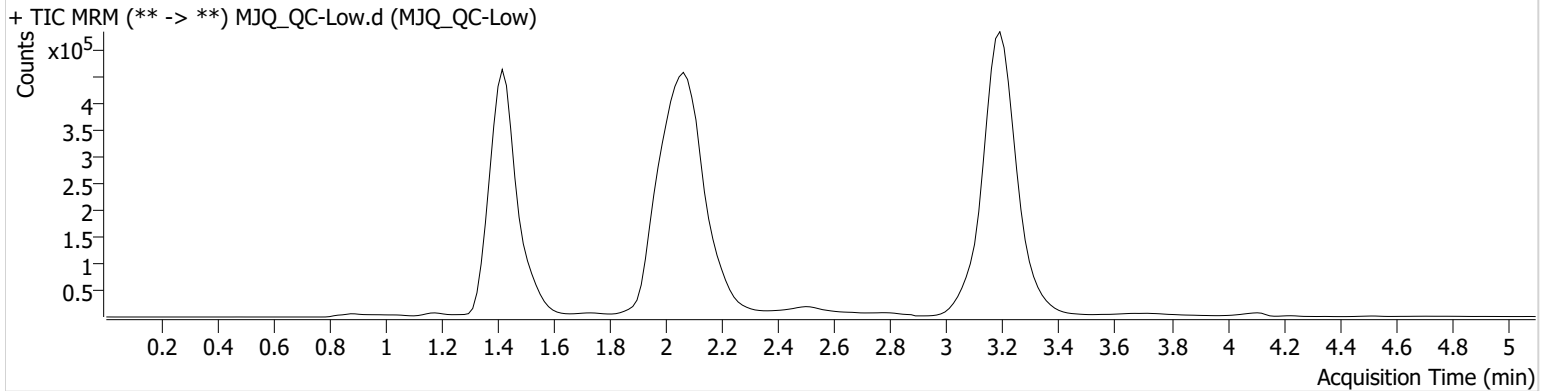


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_QC-Low.d
<b>Type</b>	Sample	<b>Sample</b>	MJQ_QC-Low
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-H1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 11:37:11 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	120409	57.40	11.3	∞	2096918	3.8769 ng/ml
THC-COOH	1.444	195750	∞	55.4	∞	576911	14.0025 ng/ml
THC	3.194	149846	296.96	30.4	53.33	4375916	3.9949 ng/ml

8C



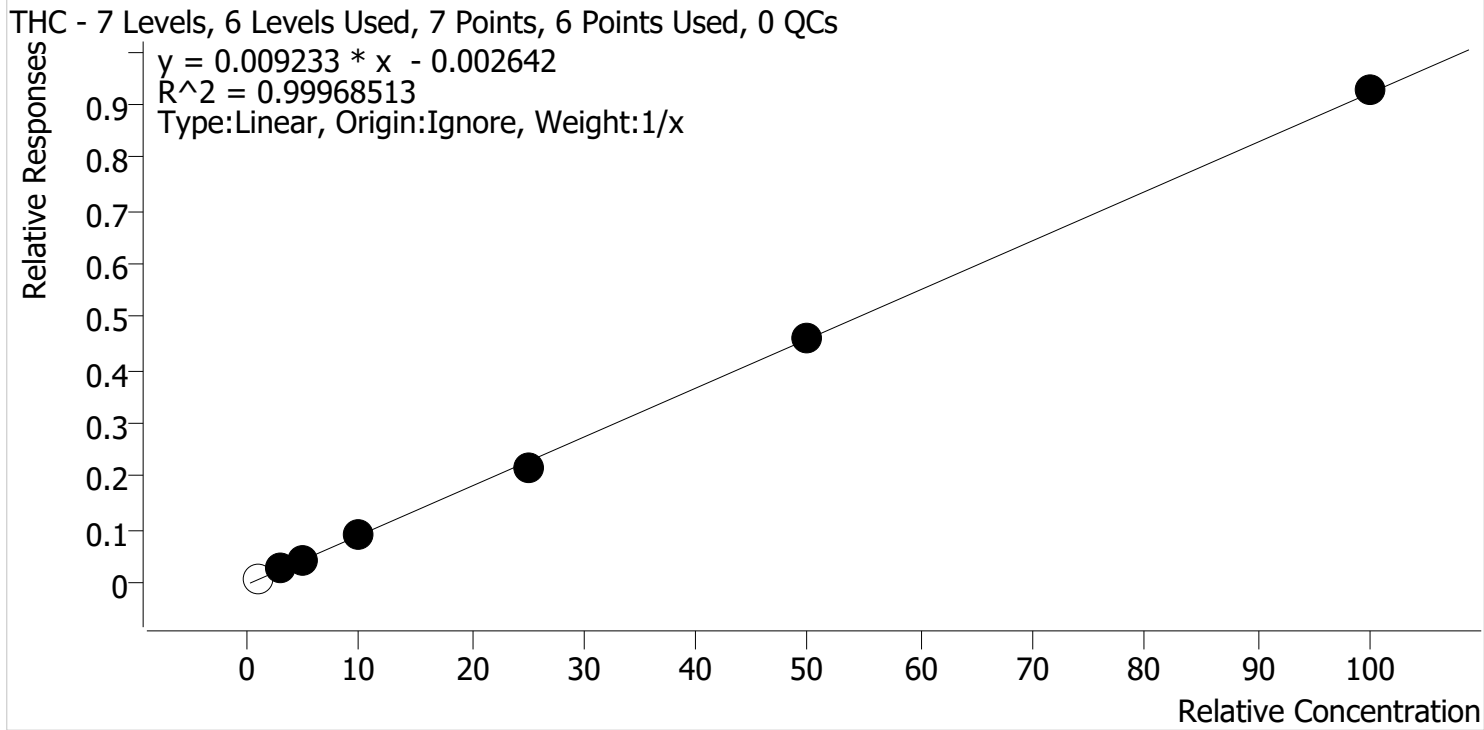
# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin  
SAEC\QuantResults\AM 27.batch.bin

**Last Cal. Update** 2/24/2021 7:48 AM

**Analyst Name** ISP\datastor

**Analyte** THC **Internal Standard** THC-D3



Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJQ_Cal 1	1	x	1.0	1.3	125.9
MJQ_Cal 2	2	✓	3.0	3.0	101.4
MJQ_Cal 3	3	✓	5.0	5.1	102.7
MJQ_Cal 4	4	✓	10.0	9.8	98.3
MJQ_Cal 5	5	✓	25.0	24.1	96.4
MJQ_Cal 6R	6	✓	50.0	50.3	100.7
MJQ_Cal 7	7	✓	100.0	100.6	100.6

Cal 1 dropped due to not meeting ratio requirement



8C



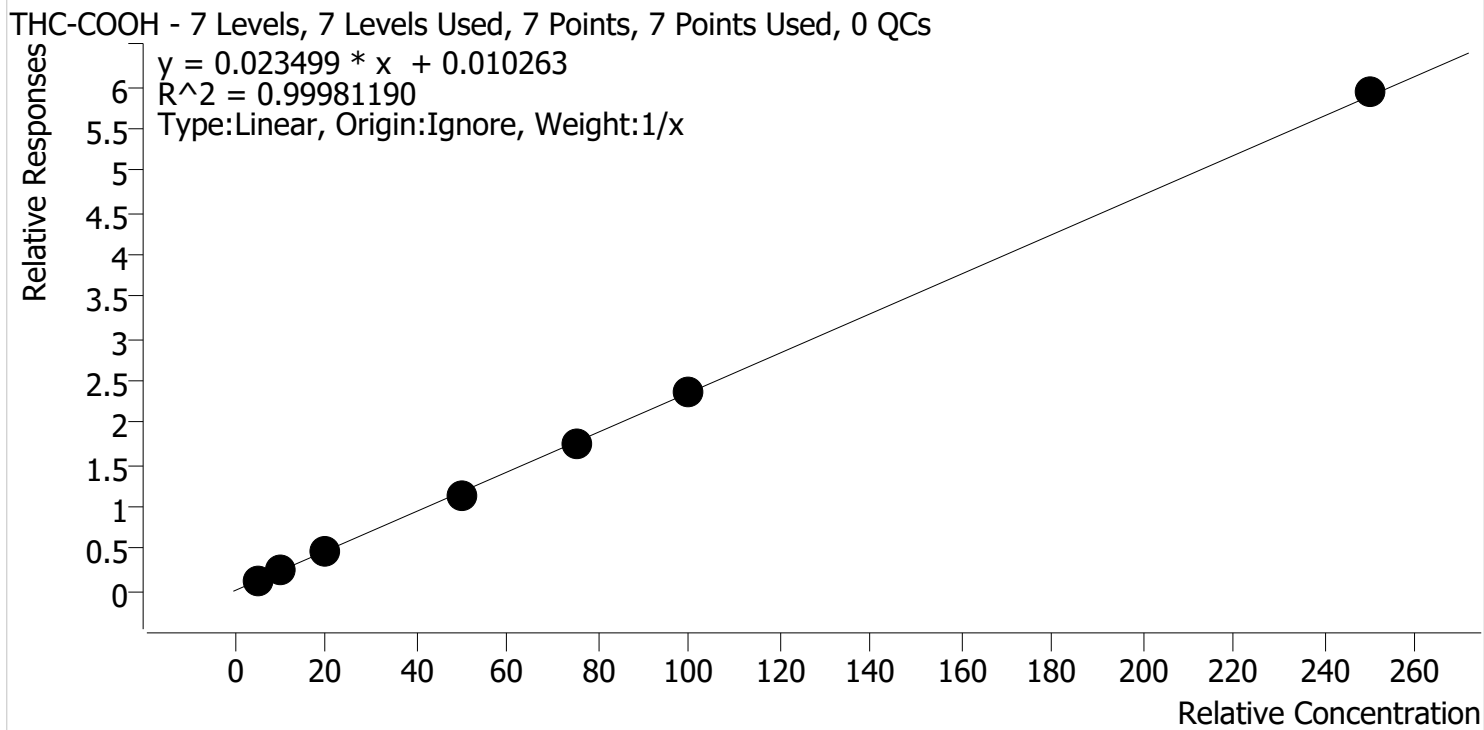
# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin  
SAEC\QuantResults\AM 27.batch.bin

**Last Cal. Update** 2/24/2021 7:48 AM

**Analyst Name** ISP\datastor

**Analyte** THC-COOH **Internal Standard** THC-COOH-D9



Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJQ_Cal 1	1	✓	5.0	5.1	101.2
MJQ_Cal 2	2	✓	10.0	10.0	100.0
MJQ_Cal 3	3	✓	20.0	20.4	102.0
MJQ_Cal 4	4	✓	50.0	48.7	97.4
MJQ_Cal 5	5	✓	75.0	73.9	98.5
MJQ_Cal 6R	6	✓	100.0	100.1	100.1
MJQ_Cal 7	7	✓	250.0	251.9	100.7

8C



# AM #27 Cannabinoids Quant. Calibration Curve Report

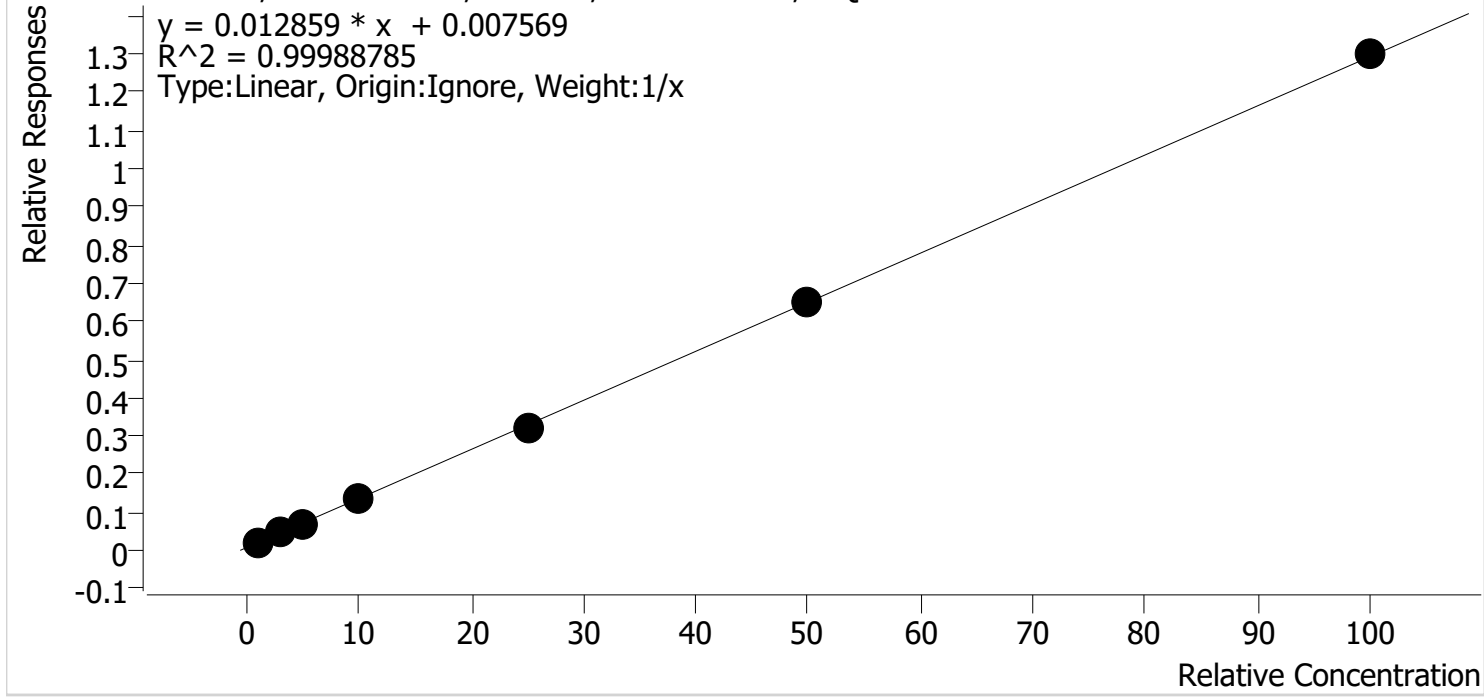
**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin  
SAEC\QuantResults\AM 27.batch.bin

**Last Cal. Update** 2/24/2021 7:48 AM

**Analyst Name** ISP\datastor

**Analyte** THC-OH **Internal Standard** THC-OH-D3

THC-OH - 7 Levels, 7 Levels Used, 7 Points, 7 Points Used, 0 QCs



Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJQ_Cal 1	1	✓	1.0	1.0	104.2
MJQ_Cal 2	2	✓	3.0	2.9	97.1
MJQ_Cal 3	3	✓	5.0	5.0	100.2
MJQ_Cal 4	4	✓	10.0	10.0	100.1
MJQ_Cal 5	5	✓	25.0	24.5	97.8
MJQ_Cal 6R	6	✓	50.0	50.1	100.2
MJQ_Cal 7	7	✓	100.0	100.5	100.5

8C

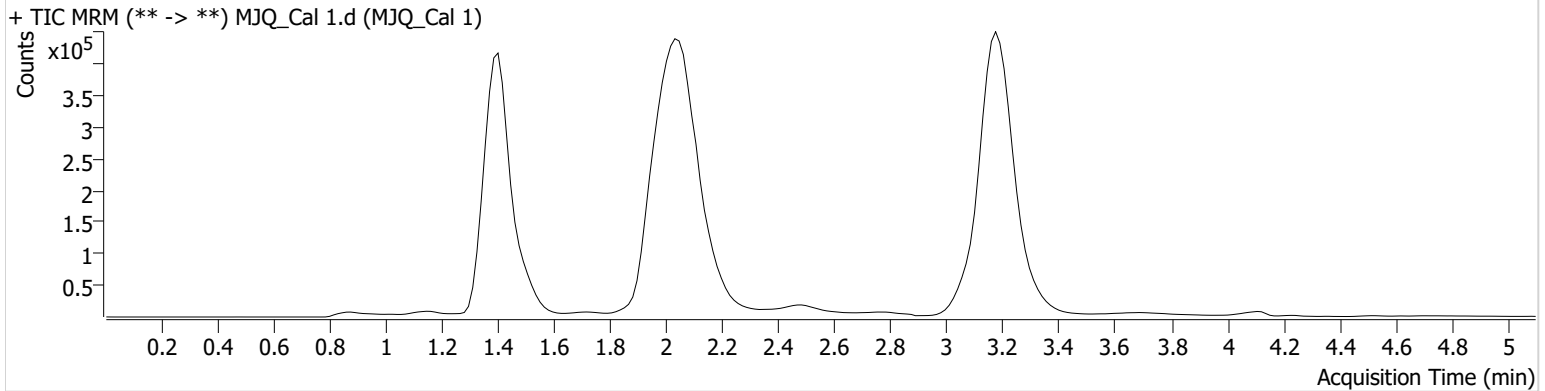


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 1.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 1
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-A1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 6:10:22 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	44018	∞	8.8	∞	2099842	1.0416 ng/ml <b>Low</b>
THC-COOH	1.429	72222	∞	46.3	∞	559021	5.0611 ng/ml
THC	3.179	35708	15.19	65.7 <b>High</b>	∞	3975176	1.2590 ng/ml

8C

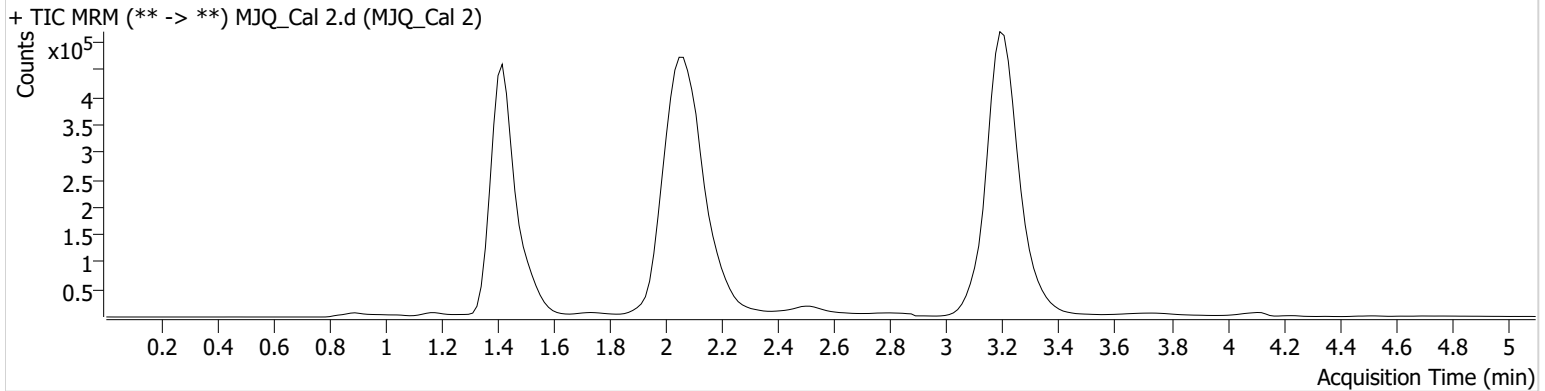


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 2.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 2
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-B1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 6:18:09 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	90282	∞	10.0	∞	2005478	2.9123 ng/ml <b>Low</b>
THC-COOH	1.444	134118	∞	51.1	243.55	546835	10.0005 ng/ml
THC	3.209	104549	265.49	30.8	∞	4109011	3.0418 ng/ml

8C

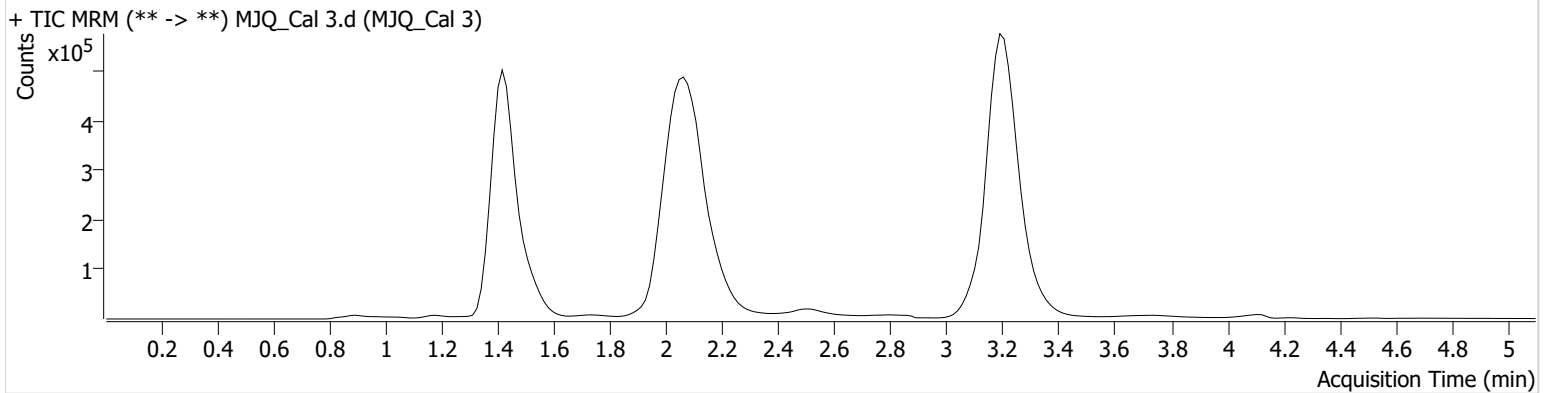


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 3.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 3
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-C1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 6:25:44 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	149361	∞	11.1	∞	2074835	5.0096 ng/ml
THC-COOH	1.444	274203	∞	55.5	1510.04	559921	20.4033 ng/ml
THC	3.209	200139	1955.98	29.6	75.38	4471712	5.1336 ng/ml

8C

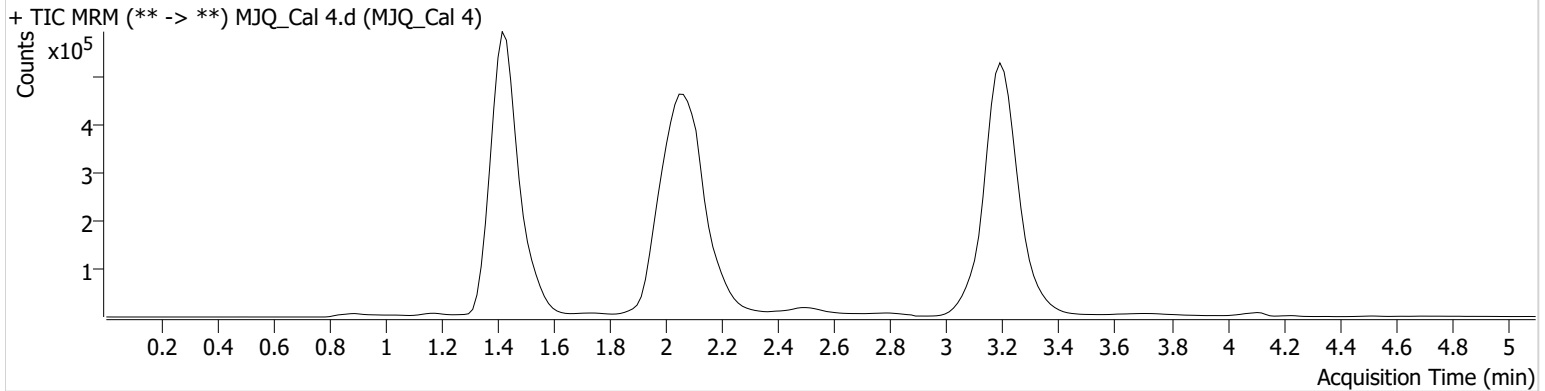


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 4.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 4
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-D1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 6:33:20 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	287465	853.85	11.1	672.52	2109451	10.0091 ng/ml
THC-COOH	1.444	649709	∞	58.7	3717.82	562401	48.7248 ng/ml
THC	3.209	359272	7132.51	25.5	336.67	4076775	9.8308 ng/ml

8C

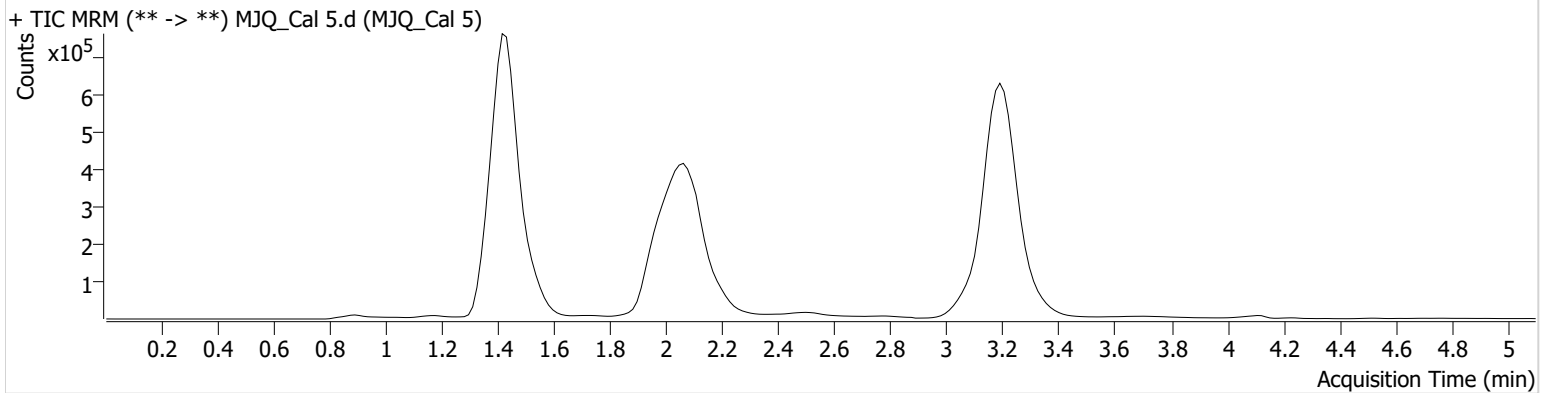


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 5.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 5
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-E1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 6:40:55 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	737192	∞	11.3	∞	2288836	24.4588 ng/ml
THC-COOH	1.444	1021053	997.27	59.1	∞	584810	73.8628 ng/ml
THC	3.194	967723	740.34	24.6	409.28	4401032	24.1012 ng/ml

8C

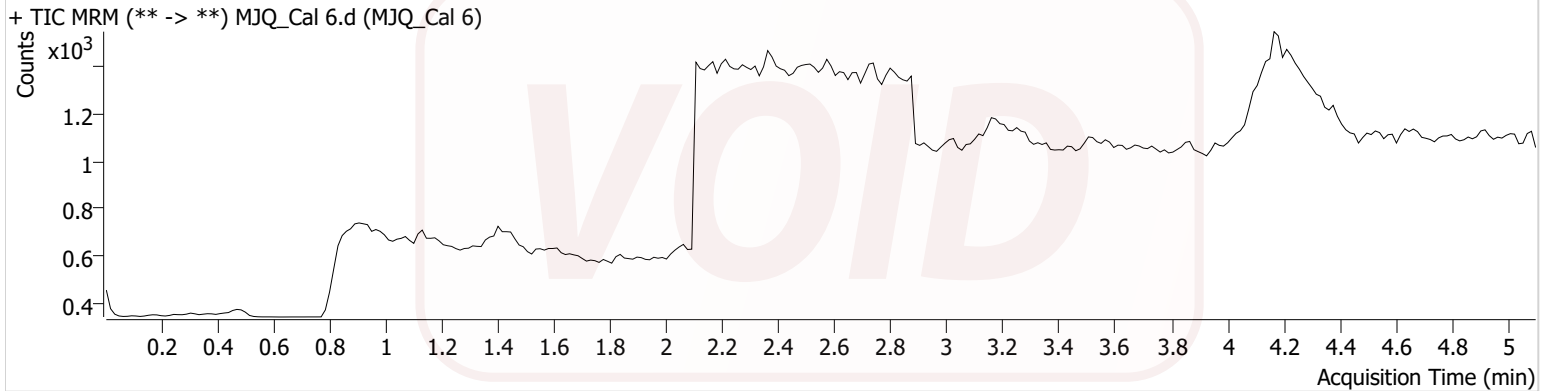


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument Type</b>	Instrument 1 Cal	<b>Data File</b>	MJQ_Cal 6.d
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Sample</b>	MJQ_Cal 6
<b>Sample Position</b>	P1-F1	<b>Operator</b>	Sarah Collins
<b>Injection Volume</b>	10	<b>Comment</b>	
<b>Acq. Date-Time</b>	2/23/2021 6:48:31 PM		
<b>Sample Info.</b>			

## Sample Chromatogram



\*Sample reconstituted and reinjected due to low internal standard response. See reinject data.



8C

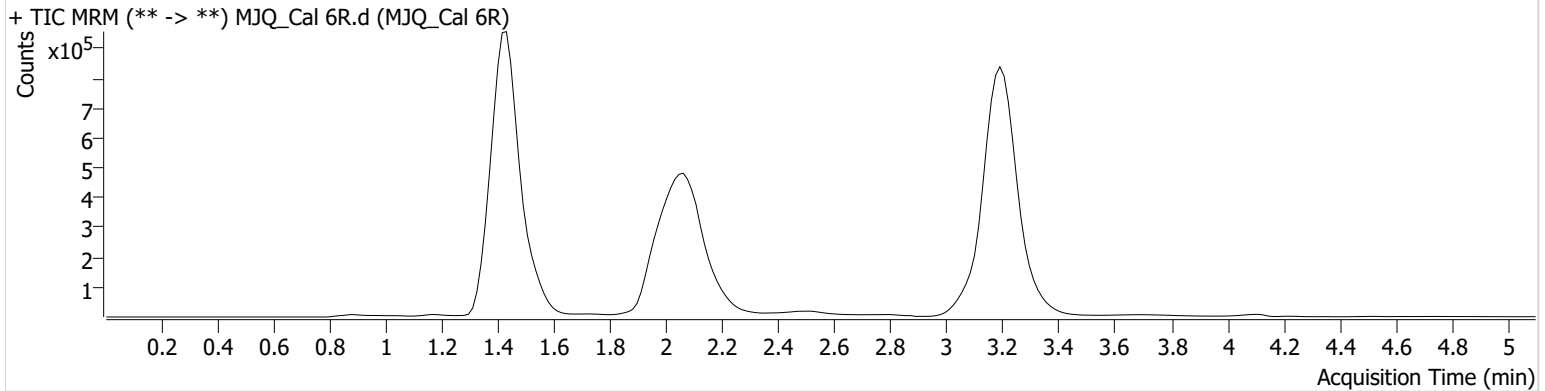


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 6R.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 6R
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-F1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 7:34:03 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	1468323	∞	11.6	∞	2253508	50.0825 ng/ml
THC-COOH	1.444	1390059	12793.43	58.7	∞	588539	100.0734 ng/ml
THC	3.194	2115972	∞	24.5	∞	4579575	50.3288 ng/ml

8C

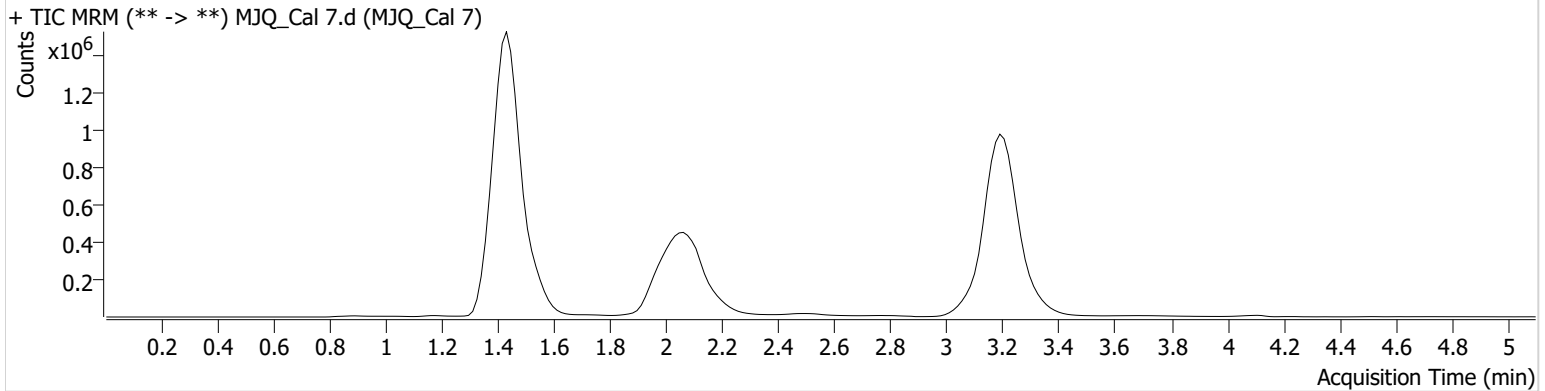


# AM #27 Cannabinoid Quant. Results

**Batch results** D:\MassHunter\Data\2021\AM 25-26\022221 MDS THCS MDS Coroner Validatoin SAEC\QuantResults\AM 27.batch.bin  
**Calibration Last Update** 2/24/2021 7:48:33 AM

<b>Instrument</b>	Instrument 1	<b>Data File</b>	MJQ_Cal 7.d
<b>Type</b>	Cal	<b>Sample</b>	MJQ_Cal 7
<b>Acq. Method</b>	AM 27 THCQ.m	<b>Operator</b>	Sarah Collins
<b>Sample Position</b>	P1-G1	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	2/23/2021 6:56:06 PM		
<b>Sample Info.</b>			

**Sample Chromatogram**



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC-OH	1.423	2683831	∞	11.7	∞	2064955	100.4862 ng/ml
THC-COOH	1.444	3037586	1731.79	58.0	9719.88	512324	251.8741 ng/ml
THC	3.194	3670633	∞	25.0	∞	3964534	100.5638 ng/ml