

**Worklist: 3769**

<u>LAB CASE</u>	<u>ITEM</u>	<u>ITEM TYPE</u>	<u>DESCRIPTION</u>
M2019-4464	1	BCK	AM 27 Blood THC Quant by LC-QQQ
M2019-4621	2	BCK	AM 27 Blood THC Quant by LC-QQQ
M2019-4654	2	BCK	AM 27 Blood THC Quant by LC-QQQ
P2019-3186	1	BCK	AM 27 Blood THC Quant by LC-QQQ
P2019-3208	1	BCK	AM 27 Blood THC Quant by LC-QQQ



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

Extraction Date: 10/23/19

Analyst: Sarah Pickle

Plate lot#: Lot # 190716 Item # IDP-108

Plate Expiration: 1/16/20

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

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

**Blank Blood Lot:** 445283-3

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

**LCMS-QQQ ID:** 069901

### 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: #3**
- 3. Place on shaking incubator at ambient temp., 900rpm for 15 minutes. *Shaker ID: 067105*
- 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)* Manifold ID: 067104
- 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.  
*SPE Dry ID: 067103*
- 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.  
Worklist path: D:\MassHunter\Data\2019\AM 27\102219 MDQ THCQ SP Batch Name: THCQ
- 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 and OH-THC 3ng/mL (quantitative), Carboxy-THC: 10ng/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 Ranges: THC: 1-100, THC-COOH 5-250, THC-OH 3-100*

S

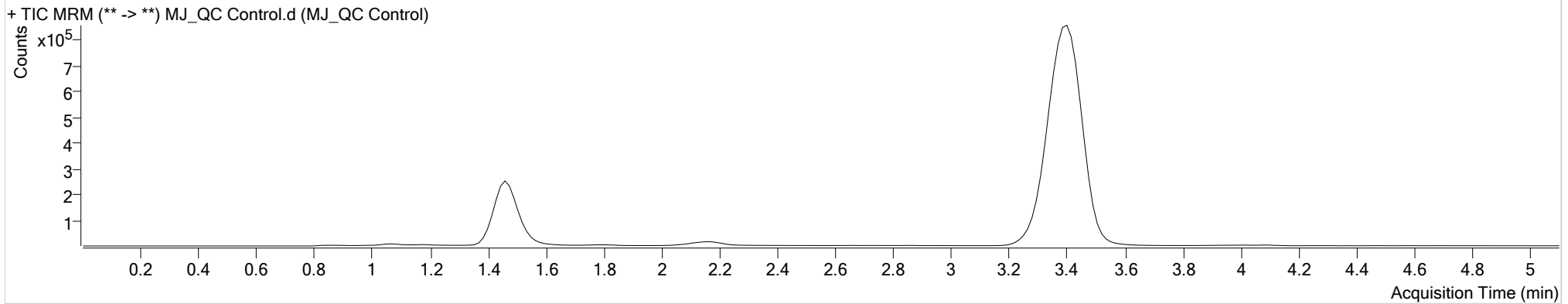


# AM #27 Cannabinoids Quant. Results

**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

**Instrument** Falco **Data File** MJ\_QC Control.d  
**Type** Sample **Sample** MJ\_QC Control  
**Acq. Method** AM 27 THC quant.m  
**Sample Position** P3-A6 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 10/23/2019 12:29:41 PM  
**Sample Info.**

### Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.405	223830	1469.96	27.0	282.29	6870181	4.2290 ng/ml
THC-COOH	1.504	105990	250.84	56.9	702.40	303163	14.3486 ng/ml
THC-OH	1.468	72493	94.12	10.5	122.79	936480	4.4085 ng/ml

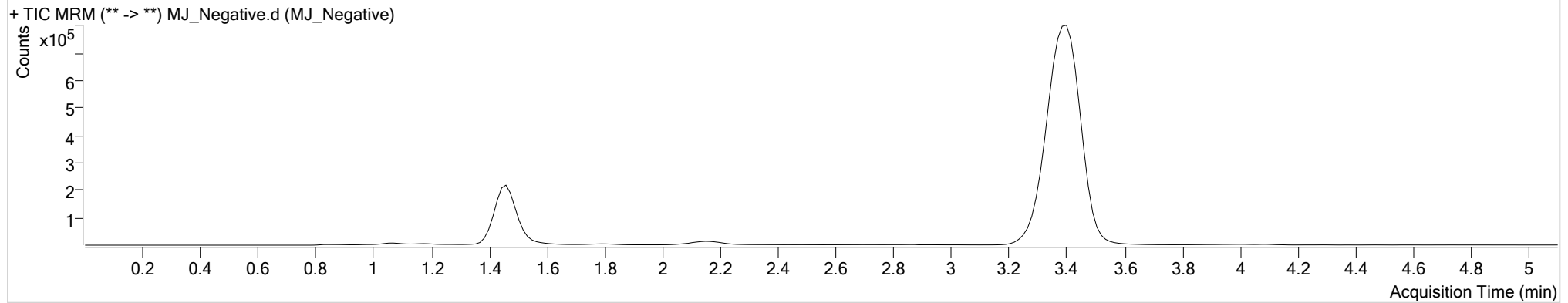
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ_Negative.d
<b>Type</b>	Sample	<b>Sample</b>	MJ_Negative
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Comment</b>	
<b>Sample Position</b>	P3-H5		
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	10/23/2019 12:44:53 PM		
<b>Sample Info.</b>			

## Sample Chromatogram





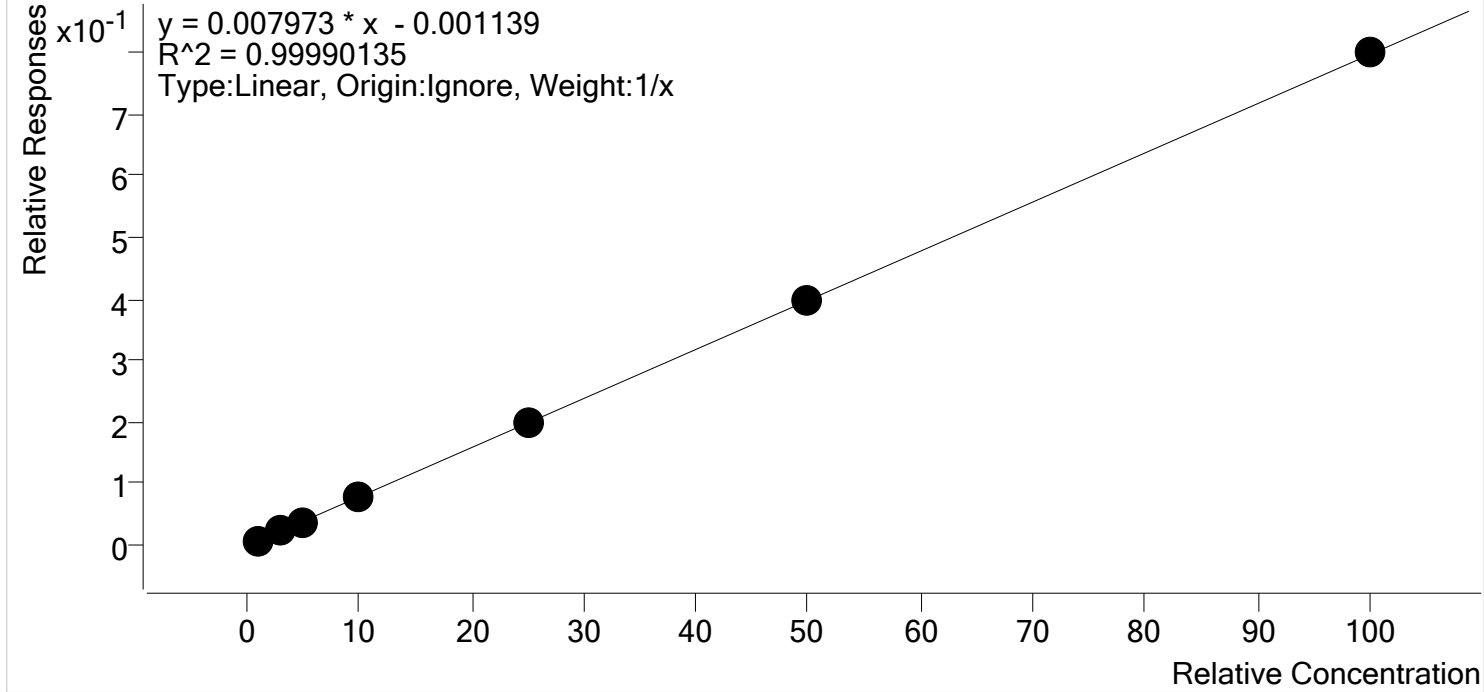
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# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Last Cal. Update** 10/23/2019 2:11 PM  
**Analyst Name** ISP\datastor  
**Analyte** THC **Internal Standard** THC-D3

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



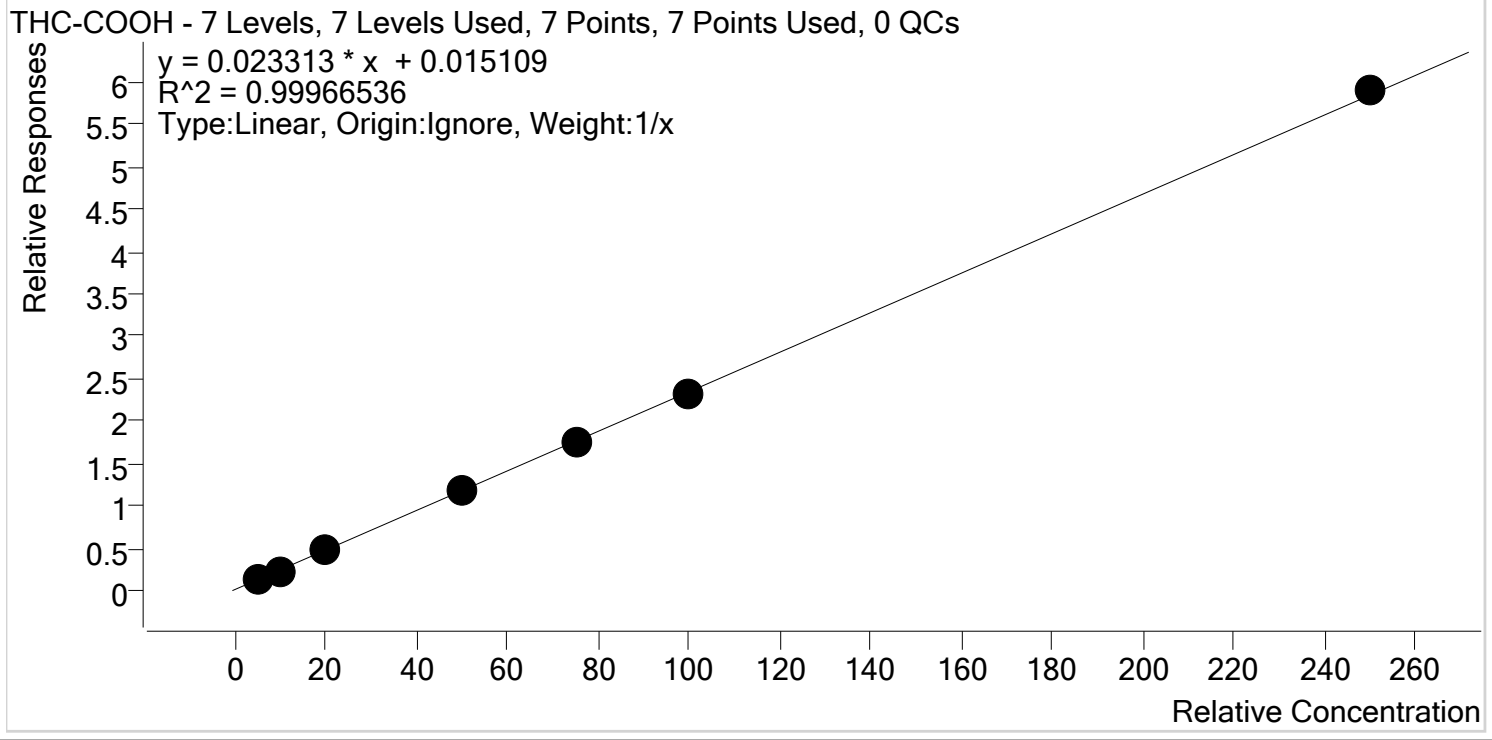
Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJ_Cal 1	1	✓	1.0	1.1	106.8
MJ_Cal 2	2	✓	3.0	2.9	97.7
MJ_Cal 3	3	✓	5.0	4.8	96.5
MJ_Cal 4	4	✓	10.0	9.9	99.3
MJ_Cal 5	5	✓	25.0	24.8	99.1
MJ_Cal 6	6	✓	50.0	50.1	100.2
MJ_Cal 7	7	✓	100.0	100.4	100.4

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# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Last Cal. Update** 10/23/2019 2:11 PM  
**Analyst Name** ISP\datastor  
**Analyte** THC-COOH **Internal Standard** THC-COOH-D9



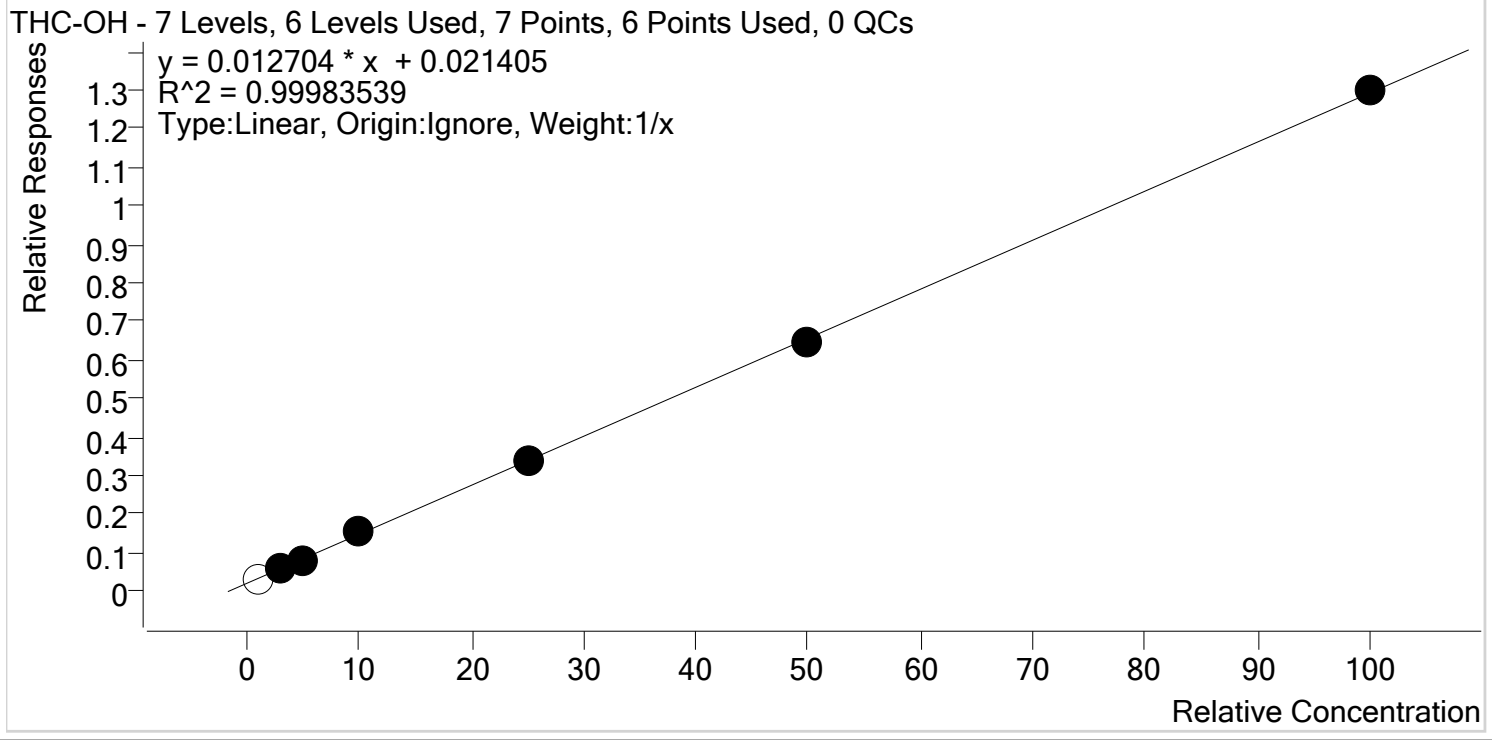
Sample	Level	Enabled	Expected Concentration	Final Concentration	Accuracy
MJ_Cal 1	1	✓	5.0	5.4	107.8
MJ_Cal 2	2	✓	10.0	9.5	94.6
MJ_Cal 3	3	✓	20.0	19.7	98.3
MJ_Cal 4	4	✓	50.0	50.6	101.3
MJ_Cal 5	5	✓	75.0	73.8	98.3
MJ_Cal 6	6	✓	100.0	98.8	98.8
MJ_Cal 7	7	✓	250.0	252.3	100.9

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# AM #27 Cannabinoids Quant. Calibration Curve Report

**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Last Cal. Update** 10/23/2019 2:11 PM  
**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	0.9	94.8
MJ_Cal 2	2	✓	3.0	3.0	100.4
MJ_Cal 3	3	✓	5.0	4.9	97.4
MJ_Cal 4	4	✓	10.0	10.3	103.3
MJ_Cal 5	5	✓	25.0	24.9	99.5
MJ_Cal 6	6	✓	50.0	49.4	98.9
MJ_Cal 7	7	✓	100.0	100.5	100.5

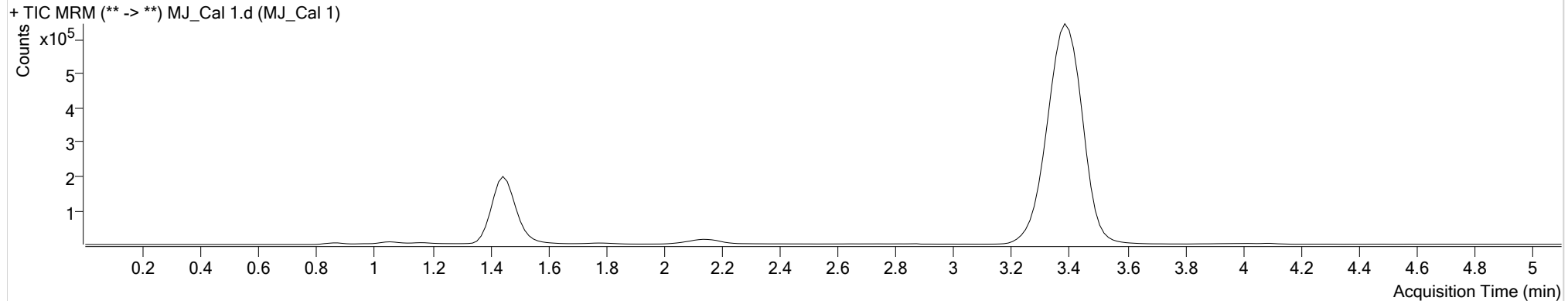
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

<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>Comment</b>	
<b>Sample Position</b>	P3-B6		
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	10/23/2019 11:28:48 AM		
<b>Sample Info.</b>			

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.405	40203	159.62	29.7	22.90	5451096	1.0678 ng/ml
THC-COOH	1.489	38514	∞	48.9	162.75	273643	5.3891 ng/ml
THC-OH	1.498	27183	∞	7.2 <b>Low</b>	18.92	812678	0.9480 ng/ml

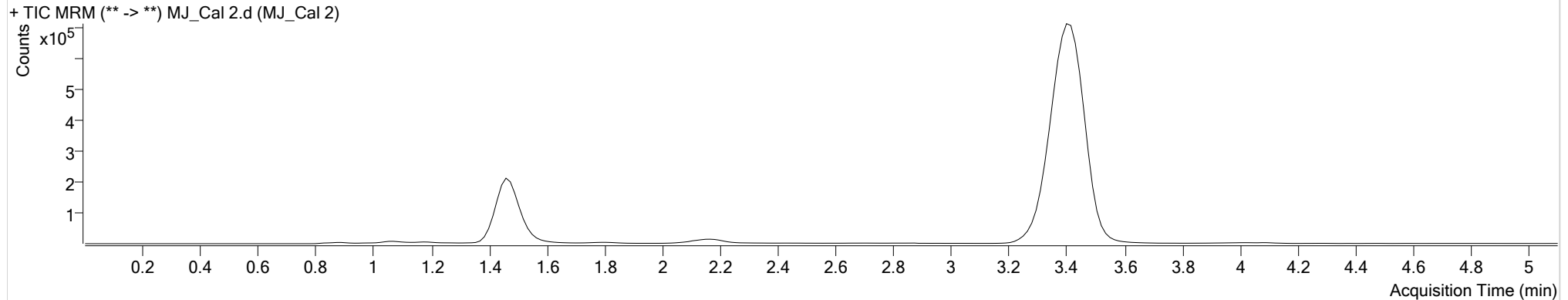
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

<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>Sample Position</b>	P3-C6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	10/23/2019 11:36:34 AM		
<b>Sample Info.</b>			

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.420	128767	348.41	27.2	130.25	5790107	2.9321 ng/ml
THC-COOH	1.504	64015	∞	57.4	334.55	271652	9.4601 ng/ml
THC-OH	1.483	48724	∞	10.1	34.64	816681	3.0113 ng/ml

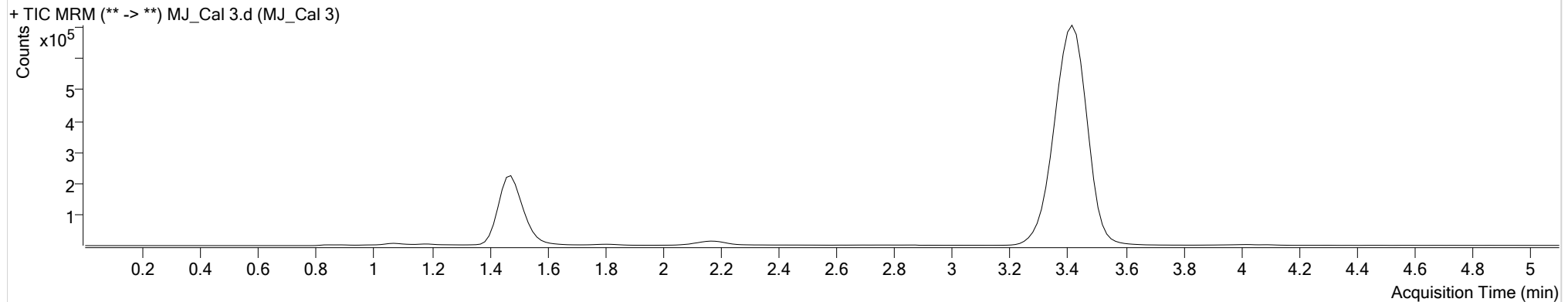
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

**Instrument** Falco **Data File** MJ\_Cal 3.d  
**Type** Cal **Sample** MJ\_Cal 3  
**Acq. Method** AM 27 THC quant.m  
**Sample Position** P3-D6 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 10/23/2019 11:44:09 AM  
**Sample Info.**

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.435	199637	1930.59	27.8	∞	5350136	4.8228 ng/ml
THC-COOH	1.519	123784	∞	58.2	657.53	261531	19.6543 ng/ml
THC-OH	1.483	65560	∞	11.1	131.36	787155	4.8712 ng/ml

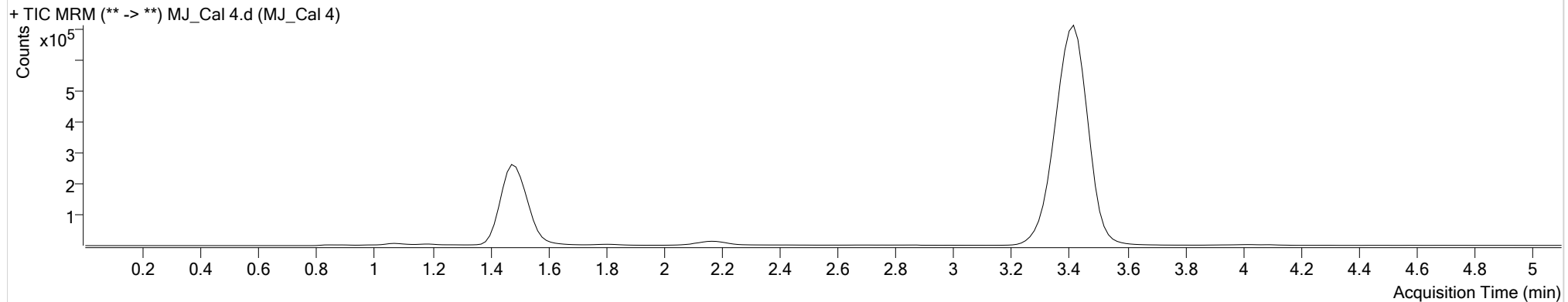
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

<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>Sample Position</b>	P3-E6	<b>Comment</b>	
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	10/23/2019 11:51:44 AM		
<b>Sample Info.</b>			

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.420	398305	1322.14	27.6	∞	5101833	9.9346 ng/ml
THC-COOH	1.519	299140	1558.22	60.6	1064.61	250194	50.6384 ng/ml
THC-OH	1.483	116127	∞	12.0	210.23	760678	10.3322 ng/ml

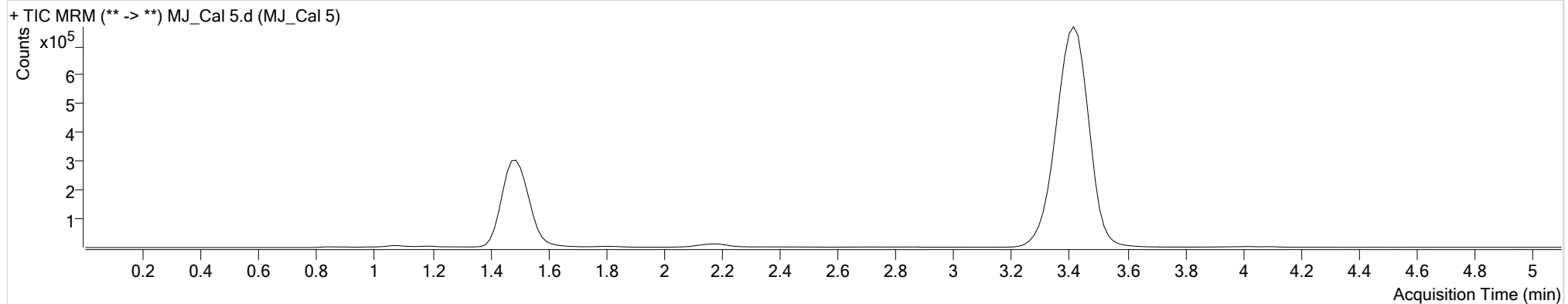
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

<b>Instrument</b>	Falco	<b>Data File</b>	MJ_Cal 5.d
<b>Type</b>	Cal	<b>Sample</b>	MJ_Cal 5
<b>Acq. Method</b>	AM 27 THC quant.m	<b>Comment</b>	
<b>Sample Position</b>	P3-F6		
<b>Injection Volume</b>	10		
<b>Acq. Date-Time</b>	10/23/2019 11:59:18 AM		
<b>Sample Info.</b>			

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.435	952737	5330.20	26.6	∞	4849981	24.7807 ng/ml
THC-COOH	1.519	409715	∞	61.2	4934.40	236192	73.7604 ng/ml
THC-OH	1.483	241469	∞	13.1	544.30	715419	24.8837 ng/ml



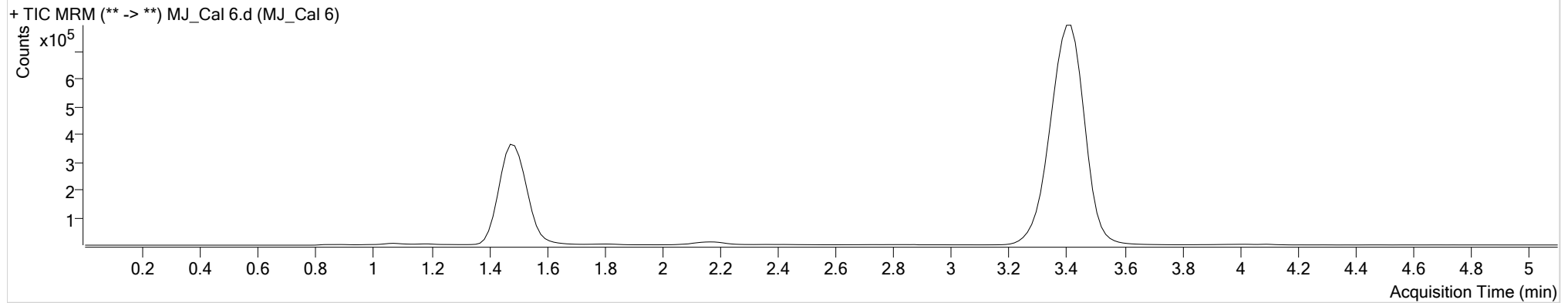
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

**Instrument** Falco **Data File** MJ\_Cal 6.d  
**Type** Cal **Sample** MJ\_Cal 6  
**Acq. Method** AM 27 THC quant.m  
**Sample Position** P3-G6 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 10/23/2019 12:06:53 PM  
**Sample Info.**

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.420	1747428	12123.15	26.7	∞	4387639	50.0932 ng/ml
THC-COOH	1.504	532904	∞	61.2	4476.51	229850	98.8029 ng/ml
THC-OH	1.468	454076	∞	13.7	872.73	699143	49.4398 ng/ml

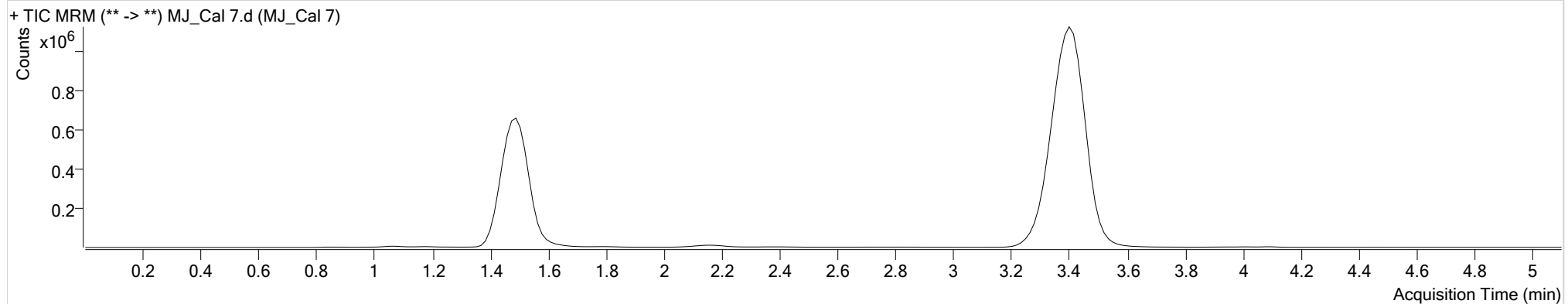
# AM #27 Cannabinoids Quant. Results



**Batch results** D:\MassHunter\Data\2019\AM 28\102219 MDQ THCQ SP\QuantResults\THCQ.batch.bin  
**Calibration Last Update** 10/23/2019 2:11:38 PM

**Instrument** Falco **Data File** MJ\_Cal 7.d  
**Type** Cal **Sample** MJ\_Cal 7  
**Acq. Method** AM 27 THC quant.m  
**Sample Position** P3-H6 **Comment**  
**Injection Volume** 10  
**Acq. Date-Time** 10/23/2019 12:14:29 PM  
**Sample Info.**

## Sample Chromatogram



Name	RT	Resp.	S/N	Ratio	S/N	ISTD Resp.	Final Conc.
THC	3.405	3679621	13614.02	26.8	1476.59	4604612	100.3688 ng/ml
THC-COOH	1.504	1297062	∞	61.5	3249.22	219960	252.2948 ng/ml
THC-OH	1.468	917305	∞	14.1	1390.48	706899	100.4618 ng/ml