

AM# 30: THC and Metabolites Screen in Blood by LC-QTOF

Extraction Date: 11/4/2020
Plate lot#: 200723

Analyst: Britany Wylie
Plate Expiration: 1-23-2021

Mobile phase A: 0.1% Formic acid in Water
0.1% Formic Acid in Water
1N KOH Saturated Phosphate Buffer

Mobile phase B: 0.1% Formic acid in MeOH
MTBE Hexane

Blank Blood Lot: 20G20792

Urine Blank: 10120

Column: Phenomenex Phenyl Hexyl (4.6x50mm: 2.6 um)

LCMS-QTOF ID: 70044

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. Urine hydrolysis: add 1.5 ml urine to blank plate, add 250 ul 1N KOH mix and incubate at 40 degrees for 15 minutes.
Pipette **1000 µL blood (calibrated pipette)** in wells of analytical (standards) plate. **Pipette ID: K52558g**
Pipette 1000 ul urine to analytical (standards) plate.
- 3. Place on shaking incubator at ambient temp., 900 rpm for 15 minutes.
- 4. Pipette **500 µL 0.1% formic acid in blood** wells **500 ul saturated phosphate buffer in urine** wells of analytical plate.
- 5. Place on shaking incubator at ambient temp., 900 rpm for 15 minutes.
- 6. Transfer **800 µL 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) Manifold ID: 66792
- 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. 10-15 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. 10-15 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: 66819
- 16. Reconstitute in **100 µL 100% LCMS MeOH** and heat seal plate with foil. Place in autosampler and run worklist.

Post-Analytic

- 1. Create batch and process data.
- 2. Calculated sample concentration of 3 ng/mL or greater for THC and THC-OH, a calculated sample concentration of 10 ng/mL or greater for Carboxy-THC.
- 3. Retention time within +/- 2% or +/-0.100 min whichever is greater of the average retention time of the calibrators.
- 4. Did all QCs pass for each analyte? Yes
- 5. Central File Packet to include: LIMS Worklist, Method Checklist, Calibration and Control Reports

COMMENTS: *samples were ran on the QTOF due to variability in low end calibrators only c-THC was evaluated, the samples were ran on the QTOF and then reconstituted and run on the LCQQQ.*

Hydroxy-THC and THC not evaluated in this run. Low response and variability in the lower end calibrators.

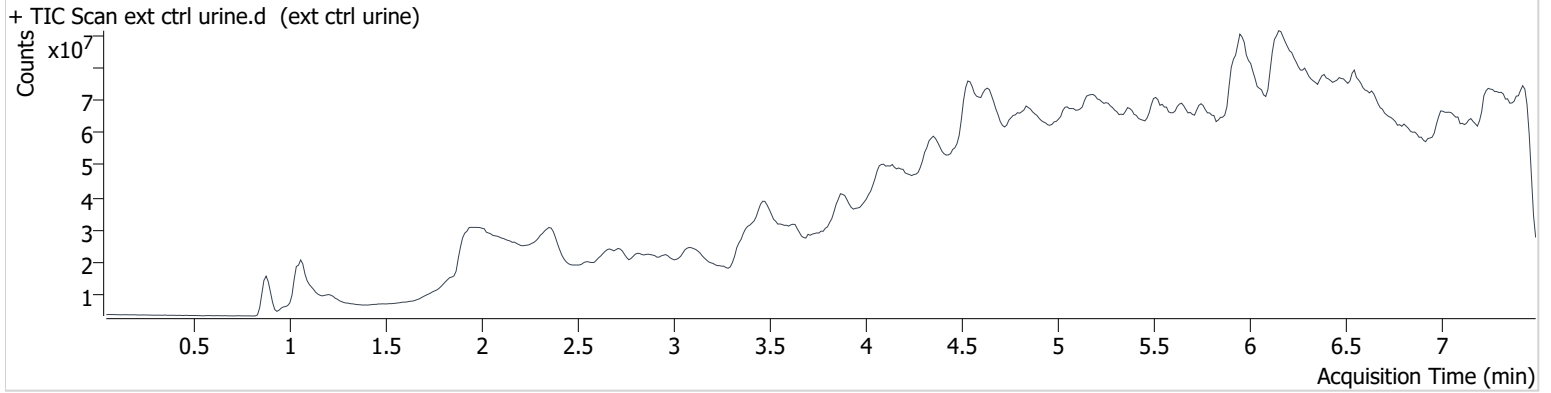
AM #30 Cannabinoids

Batch results D:\MassHunter\Data\2020\am 30\11420\QuantResults\thcs_qtof.batch.bin
Calibration Last Update 11/6/2020 11:31:15 AM

| | | | |
|-------------------------|----------------------|------------------|------------------|
| Instrument | 70044 | Data File | ext ctrl urine.d |
| Type | Sample | Sample | ext ctrl urine |
| Acq. Method | THC Screen 1122.m | Operator | Britany Wylie |
| Sample Position | P2-B2 | Comment | |
| Injection Volume | 10 | | |
| Acq. Date-Time | 11/4/2020 4:36:39 PM | | |

Sample Info.

Sample Chromatogram



| Name | RT | Resp. | Mass Accuracy | Mass Abundance Score | ISTD Resp. | Final Conc. |
|----------|-------|--------|---------------|----------------------|------------|---------------|
| THC-COOH | 6.613 | 787962 | -0.18 | 83.5 | 1799934 | 41.5956 ng/ml |

Toxicology AM method 27/26 external prep information

BW

working solution 15 ug/ml in meoh C-THC, THC-OH, 7.5 ug/ml THC

Stock solution 1mg/ml 7.5 ul each THC, 100 ug/ml 150 ul C-THC, 150 ul THC-OH in 9692.5 ul meOH

Ppd 8/26/20 Exp: 7/1/21 lot 82620 by AMN

| Drug | lot | expiration |
|--------|------------|------------|
| C-THC | FE01061702 | 3/1/2022 |
| THC-OH | FE07221601 | 7/1/2021 |
| THC | FE01041701 | 3/1/2022 |

AM 27/26 blood control 100 ul working solution lot () in 9900 ul blood lot ()

| | | | |
|--|--|--|--|
| | | Concentration 7.5 ng/ml THC, 15 ng/ml C-THC, THC-OH | |
|--|--|--|--|

AM 27/26 urine control 400 ul working solution lot (82620) in 9600 ul urine

out of use

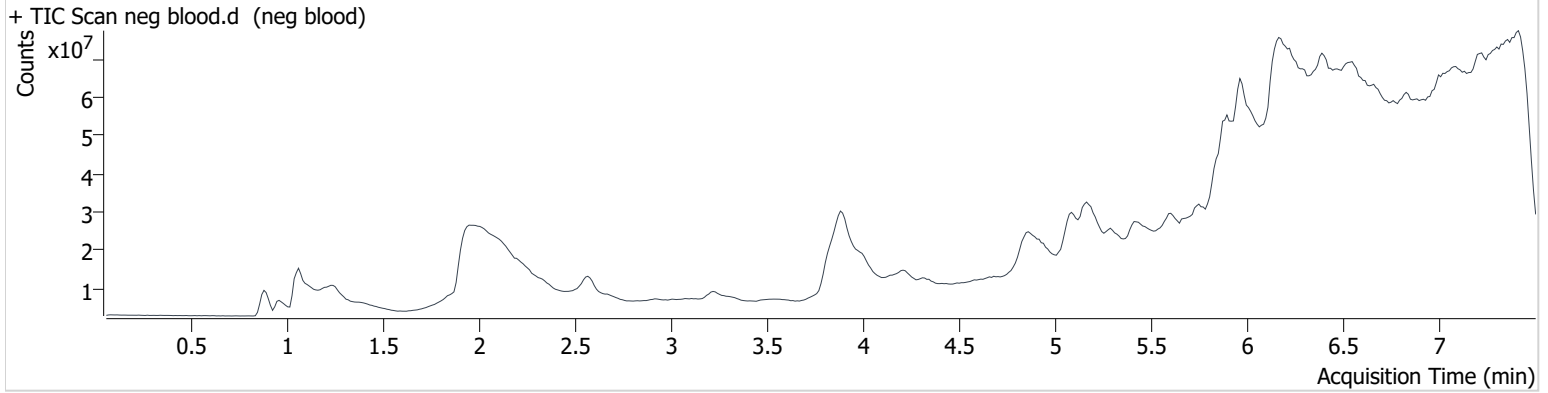
| | | | | |
|---|------------|---|--------|-----------|
| ppd 8/26/20 Exp 7/1/21 neg urine lot 73020 | lot u82620 | Concentration 30 ng/ml THC, and 60 ng/ml C-THC, THC-OH | by amn | 10/4/2020 |
| ppd 10/5/20 Exp 7/1/21 neg urine lot 10120 | lot 10520 | Concentration 30 ng/ml THC, and 60 ng/ml C-THC, THC-OH | by amn | |
| | | | | |
| | | | | |

AM #30 Cannabinoids

Batch results D:\MassHunter\Data\2020\am 30\11420\QuantResults\thcs qtof.batch.bin
Calibration Last Update 11/6/2020 11:31:15 AM

| | | | |
|-------------------------|----------------------|------------------|---------------|
| Instrument | 70044 | Data File | neg blood.d |
| Type | Sample | Sample | neg blood |
| Acq. Method | THC Screen 1122.m | Operator | Britany Wylie |
| Sample Position | P2-C4 | Comment | |
| Injection Volume | 10 | | |
| Acq. Date-Time | 11/4/2020 7:19:41 PM | | |

Sample Chromatogram



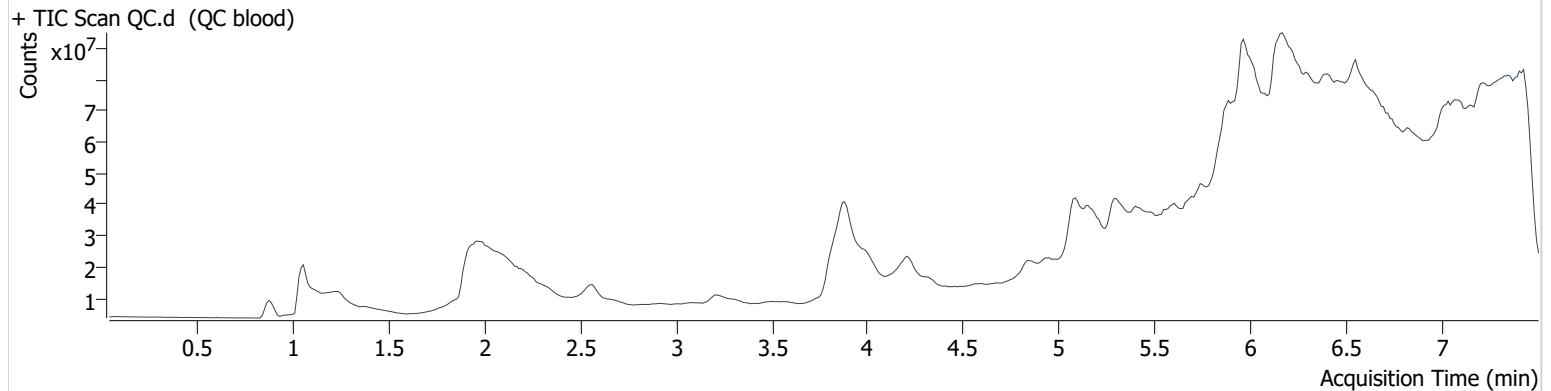
| Name | RT | Resp. | Mass Accuracy | Mass Abundance Score | ISTD Resp. | Final Conc. |
|----------|-------|--------|---------------|----------------------|------------|---|
| THC-COOH | 6.611 | 103448 | -5.58 | 0.0 | 1905010 | 5.1820 ng/ml *neg due to mass score <70 |

AM #30 Cannabinoids

Batch results D:\MassHunter\Data\2020\am 30\11420\QuantResults\thcs qtof.batch.bin
Calibration Last Update 11/6/2020 11:31:15 AM

| | | | |
|-------------------------|----------------------|------------------|---------------|
| Instrument | 70044 | Data File | QC.d |
| Type | QC | Sample | QC blood |
| Acq. Method | THC Screen 1122.m | Operator | Britany Wylie |
| Sample Position | P2-H1 | Comment | |
| Injection Volume | 10 | | |
| Acq. Date-Time | 11/4/2020 4:17:28 PM | | |

Sample Chromatogram



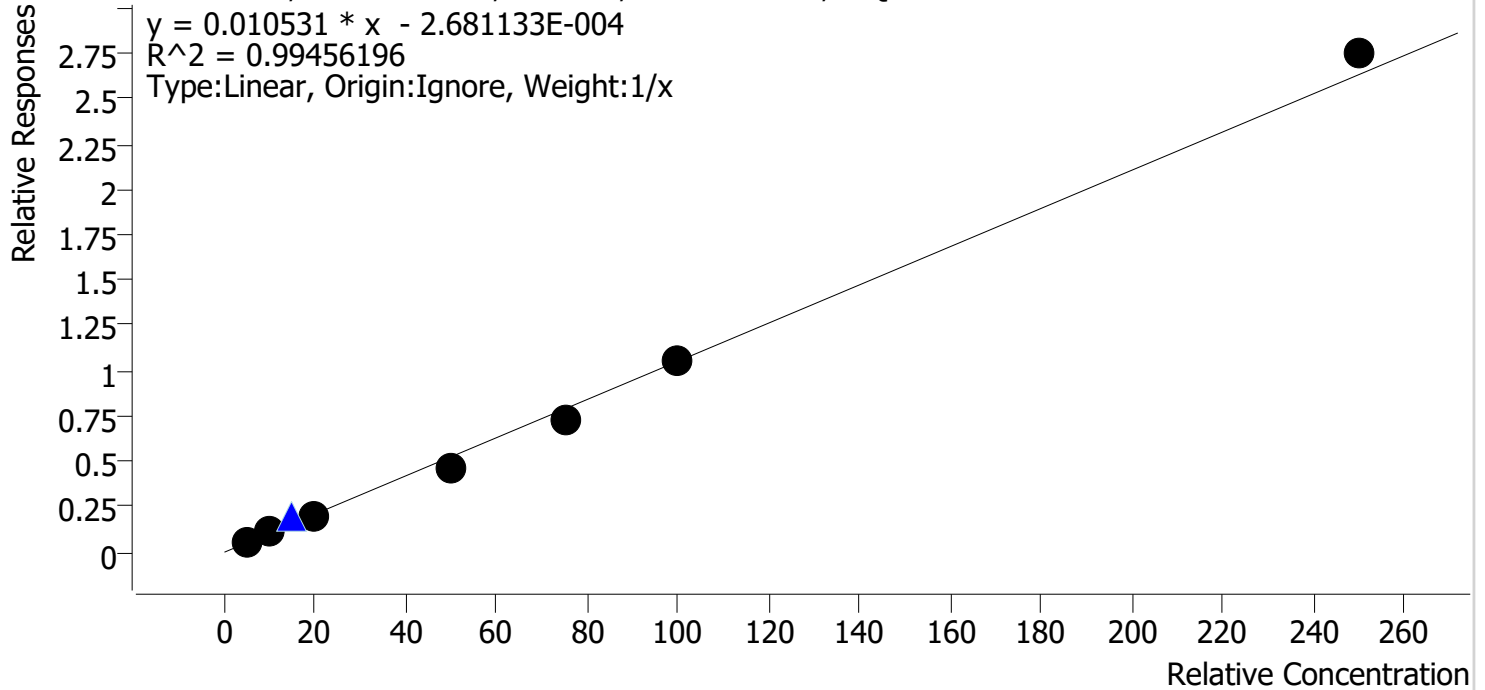
| Name | RT | Resp. | Mass Accuracy | Mass Abundance Score | ISTD Resp. | Final Conc. |
|----------|-------|--------|---------------|----------------------|------------|---------------|
| THC-COOH | 6.610 | 459630 | -0.25 | 82.9 | 2389893 | 18.2880 ng/ml |

Compound Calibration Report

Batch results D:\MassHunter\Data\2020\am 30\11420\QuantResults\thcs qtof.batch.bin
Last Cal. Update 11/6/2020 11:31 AM
Analyst Name ISP\datastor
Analyte THC-COOH

Internal Standard THC-COOH-D9

THC-COOH - 7 Levels, 7 Levels Used, 7 Points, 7 Points Used, 1 QCs



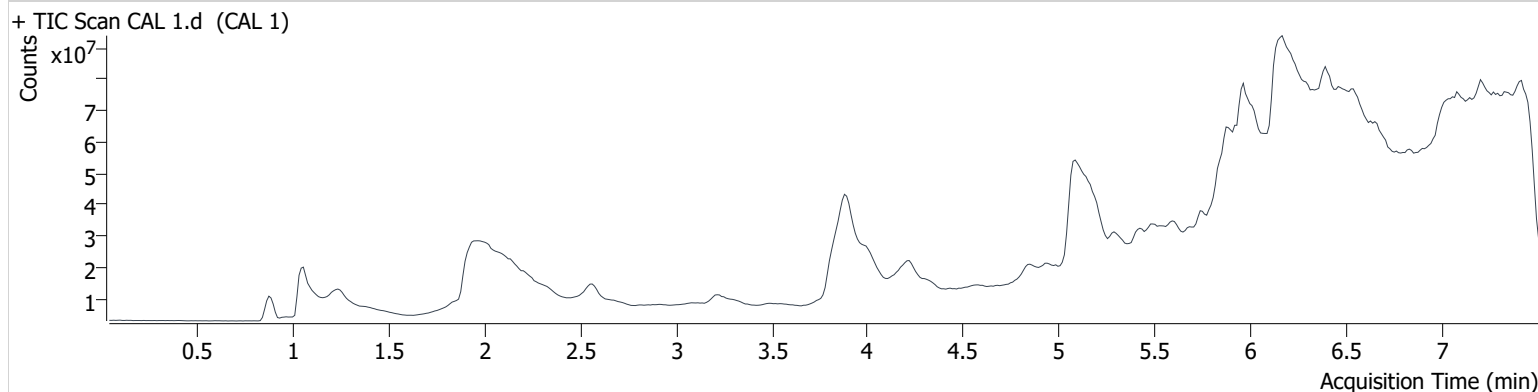
| Sample | Level | Enabled | Expected Concentration | Final Concentration | Accuracy |
|--------|-------|---------|------------------------|---------------------|----------|
| CAL 1 | 1 | ✓ | 5.0 | 5.5 | 110.2 |
| CAL 2 | 2 | ✓ | 10.0 | 11.5 | 114.6 |
| CAL 3 | 3 | ✓ | 20.0 | 17.8 | 89.0 |
| CAL 4 | 4 | ✓ | 50.0 | 44.6 | 89.2 |
| CAL 5 | 5 | ✓ | 75.0 | 68.7 | 91.6 |
| CAL 6 | 6 | ✓ | 100.0 | 101.0 | 101.0 |
| CAL 7 | 7 | ✓ | 250.0 | 260.9 | 104.4 |

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Batch results D:\MassHunter\Data\2020\am 30\11420\QuantResults\thcs qtof.batch.bin
Calibration Last Update 11/6/2020 11:31:15 AM

| | | | |
|-------------------------|----------------------|------------------|---------------|
| Instrument | 70044 | Data File | CAL 1.d |
| Type | Cal | Sample | CAL 1 |
| Acq. Method | THC Screen 1122.m | Operator | Britany Wylie |
| Sample Position | P2-A1 | Comment | |
| Injection Volume | 10 | | |
| Acq. Date-Time | 11/4/2020 3:10:13 PM | | |

Sample Chromatogram



| Name | RT | Resp. | Mass Accuracy | Mass Abundance Score | ISTD Resp. | Final Conc. |
|----------|-------|--------|---------------|----------------------|------------|--------------|
| THC-COOH | 6.610 | 117922 | -4.83 | 53.3 | 2040790 | 5.5124 ng/ml |

