

REVIEWED

By Anne Nord at 1:30 pm, Mar 12, 2020


3/10/2020

Worklist: 4075

<u>LAB CASE</u>	<u>ITEM</u>	<u>ITEM TYPE</u>	<u>DESCRIPTION</u>
C2020-0441	1	UCK	AM 3 Urine Carboxy-THC



AM 3: Carboxy-THC Urine Extraction

Extraction Date: 3/10/2020

Analyst: Britany Wylie

Negative Urine Lot: 11420

GC/MS ID: 65198

Positive Control Working Solution Lot: 31020

Methanol Lot: 156796

Hexane Lot: 142712

Ethyl Acetate Lot: 020419

Silylating Agent Lot: FN08181601

Pre-Analytic:

- 1. *Positive Control Working Solution Preparation Instructions:*
Add 180uL of 100ug/mL 11-nor-9-carboxy- Δ 9-THC Stock Solution to 9.82mL Methanol. Other volumes may be prepared. Solution is stable for 1-year or the expiration of the stock reference material (whichever is sooner). Store under refrigeration
- 2. Verify Tune and Tune evaluation completed within the previous 7 days. Tune and Tune evaluation reports initialed and filed.
- 3. Create GCMS sequence to include controls, case blanks and case samples.

Analytic:

- 1. Remove working solutions, external control, negative urine and case samples from cold storage.
- 2. To each labeled round bottomed tube add 3mL sample, using negative urine sample for both negative and positive control. Positive control: spike negative urine with 100uL positive control working solution.
- 3. Add 500uL 1N KOH to all tubes. Check pH. *(If pH <12, add additional 500ul 1N KOH). (Note: put a mark on the tube or separate the tubes that have a pH<12 as you will need to know this in step 5).*
- 4. Place tubes in 40C water bath for ~15 min. Remove and allow to cool.
- 5. If original pH was >12, add 1.5mL pH 1.8 Saturated Phosphate Buffer and 3mL Hexane/Ethyl Acetate (87:13)
If original pH was <12, add 3mL pH 1.8 Saturated Phosphate Buffer and 4mL Hexane/Ethyl Acetate (87:13)
- 6. Rock at ambient temp for ~ 10 minutes.
- 7. Centrifuge for ~ 10 min at ~3500rpm.
- 8. Transfer solvent to tapered bottom tube and evaporate to dryness under nitrogen @ 37C.
- 9. Add 50uL Ethyl Acetate and 50uL silylating reagent, cap and vortex. Heat @ 95C for 15min, then allow to cool.
- 10. Transfer sample to labeled ALS vial with insert.
- 11. Place ALS Vials in appropriate location on GCMS rack and run using appropriate GCMS method.

Post-Analytic

- 1. Complete Data analysis on all samples and corresponding sample blanks
- 2. Did positive and negative control samples provide intended response? Y / N
- 3. **Criteria for ID:** RT +/- 0.1 min., Ion Ratio of 347:473 & 371:488 within +/- 20%.
Sample response greater than Min Corrected Area, Diluted samples
- 4. Central File Packet to include: LIMS Worklist, Method Checklist, Working solution prep sheet, and Control sample GCMS data printouts

Comments:

Area Percent Report

Handwritten signature

Data Path : D:\DATA\2020\am 2\31020 am 2 and am3\
Data File : 00701011.D
Acq On : 10 Mar 2020 13:55
Operator : Instrument 65198
Sample : NCMJ
Misc :
ALS Vial : 7 Sample Multiplier: 1

Integration Parameters: rteint.p
Integrator: RTE
Smoothing : ON Filtering: 5
Sampling : 1 Min Area: 3 % of largest Peak
Start Thrs: 0.2 Max Peaks: 100
Stop Thrs : 0 Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >
Peak separation: 5

Method : D:\MassHunter\GCMS\1\METHODS\DEFAULT.M
Title :

Signal : EIC Ion 371.00 (370.70 to 371.70): 00701011.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.246	3	12	31	rVB	10056	46250	100.00%	53.465%
2	4.520	54	63	70	rBV	24992	40255	87.04%	46.535%

Sum of corrected areas: 86505

Signal : EIC Ion 473.00 (472.70 to 473.70): 00701011.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.499	53	59	69	rVB	718	1954	100.00%	47.531%
2	4.585	71	75	91	rVB2	268	734	37.56%	17.855%
3	5.031	152	158	165	rVB2	692	1267	64.84%	30.820%
4	5.171	173	184	185	rBV2	45	156	7.98%	3.795%

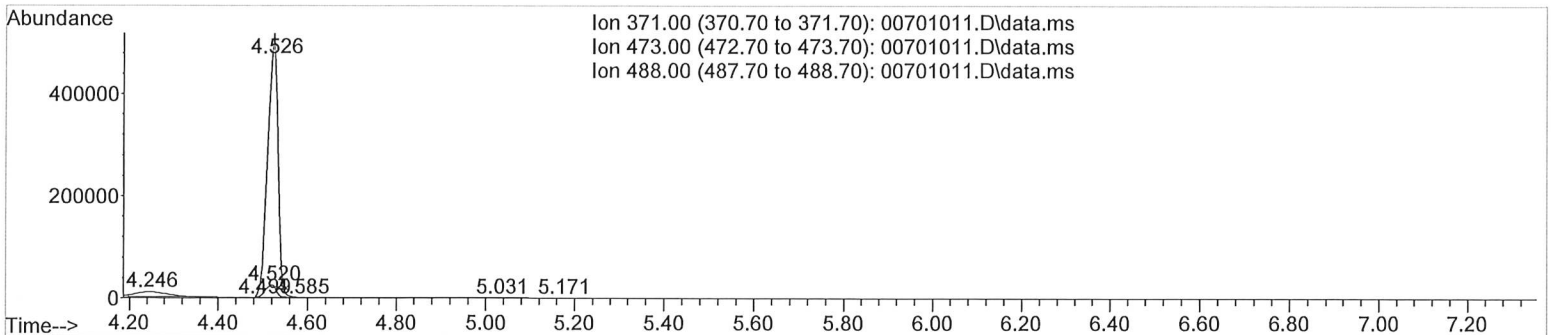
Sum of corrected areas: 4111

Signal : EIC Ion 488.00 (487.70 to 488.70): 00701011.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.526	53	64	98	rBV	517832	810158	100.00%	100.000%

Sum of corrected areas: 810158

DEFAULT.M Tue Mar 10 14:30:29 2020



Area Percent Report

Data Path : D:\DATA\2020\am 2\31020 am 2 and am3\
 Data File : 00601010.D
 Acq On : 10 Mar 2020 13:46
 Operator : Instrument 65198
 Sample : PC MJ
 Misc : am 3
 ALS Vial : 6 Sample Multiplier: 1

Integration Parameters: rteint.p
 Integrator: RTE
 Smoothing : ON
 Sampling : 1
 Start Thrs: 0.2
 Stop Thrs : 0

Filtering: 5
 Min Area: 3 % of largest Peak
 Max Peaks: 100
 Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >
 Peak separation: 5

Method : D:\MassHunter\GCMS\1\METHODS\DEFAULT.M
 Title :

Signal : EIC Ion 371.00 (370.70 to 371.70): 00601010.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.251	3	13	31	rVB	7355	35994	11.34%	9.108%
2	4.520	54	63	67	rBV	25504	41698	13.13%	10.552%
3	4.579	67	74	94	rVB	183119	317491	100.00%	80.340%

Sum of corrected areas: 395183

Signal : EIC Ion 473.00 (472.70 to 473.70): 00601010.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.579	66	74	95	rVB	88367	149252	100.00%	100.000%

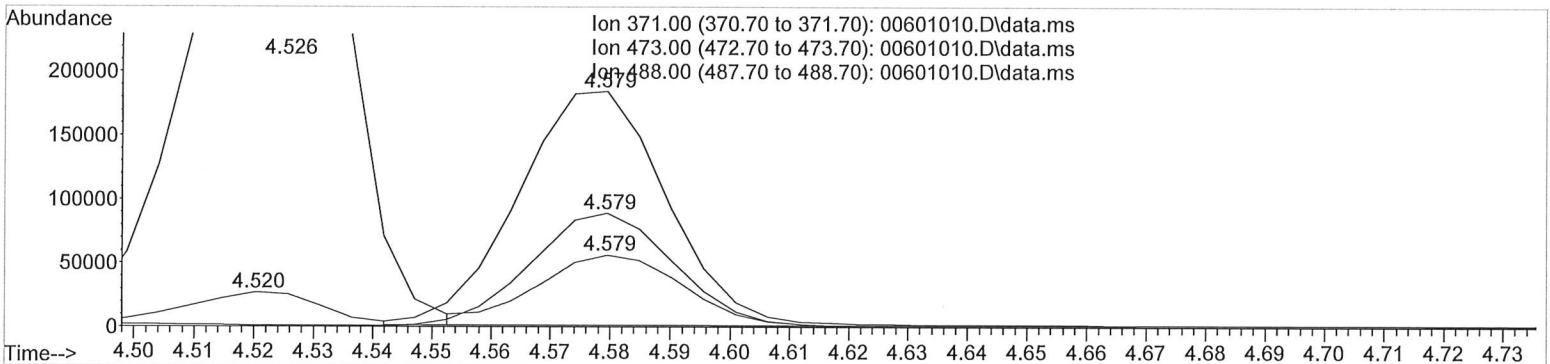
Sum of corrected areas: 149252

Signal : EIC Ion 488.00 (487.70 to 488.70): 00601010.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.526	52	64	69	rBV	535861	835709	100.00%	89.695%
2	4.579	69	74	85	rVB	55696	96017	11.49%	10.305%

Sum of corrected areas: 931726

DEFAULT.M Tue Mar 10 14:30:58 2020



Area Percent Report

BW

Data Path : D:\DATA\2020\am 2\31020 am 2 and am3\
 Data File : 00601015.D
 Acq On : 10 Mar 2020 14:32
 Operator : Instrument 65198
 Sample : PC MJ
 Misc : am 3
 ALS Vial : 6 Sample Multiplier: 1

Integration Parameters: rteint.p
 Integrator: RTE
 Smoothing : ON Filtering: 5
 Sampling : 1 Min Area: 3 % of largest Peak
 Start Thrs: 0.2 Max Peaks: 100
 Stop Thrs : 0 Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >
 Peak separation: 5

Method : D:\MassHunter\GCMS\1\METHODS\DEFAULT.M
 Title :

Signal : EIC Ion 371.00 (370.70 to 371.70): 00601015.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.521	56	63	67	rBV	30122	47221	12.92%	11.443%
2	4.580	67	74	96	rVB	207062	365455	100.00%	88.557%

Sum of corrected areas: 412676

Signal : EIC Ion 473.00 (472.70 to 473.70): 00601015.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.580	66	74	94	rVB	99300	170892	100.00%	100.000%

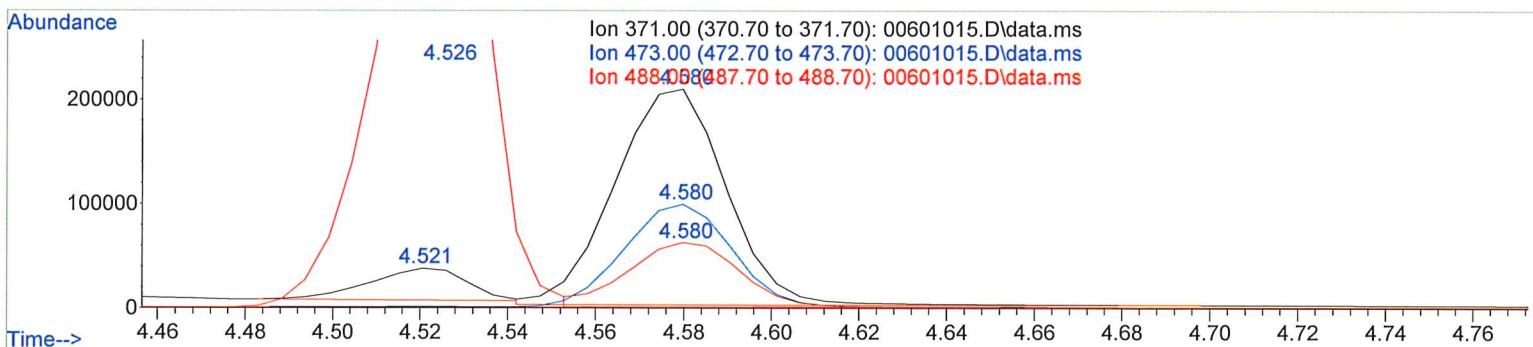
Sum of corrected areas: 170892

Signal : EIC Ion 488.00 (487.70 to 488.70): 00601015.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.526	53	64	69	rBV	576593	907383	100.00%	89.323%
2	4.580	69	74	85	rVB	61886	108459	11.95%	10.677%

Sum of corrected areas: 1015842

DEFAULT.M Wed Mar 11 07:13:27 2020



Area Percent Report

BW

Data Path : D:\DATA\2020\am 2\31020 am 2 and am3\
 Data File : 00801012.D
 Acq On : 10 Mar 2020 14:04
 Operator : Instrument 65198
 Sample : BRC3
 Misc :
 ALS Vial : 8 Sample Multiplier: 1

Integration Parameters: rteint.p
 Integrator: RTE
 Smoothing : ON Filtering: 5
 Sampling : 1 Min Area: 3 % of largest Peak
 Start Thrs: 0.2 Max Peaks: 100
 Stop Thrs : 0 Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >
 Peak separation: 5

Method : D:\MassHunter\GCMS\1\METHODS\DEFAULT.M
 Title :

Signal : EIC Ion 371.00 (370.70 to 371.70): 00801012.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.246	3	12	39	rVB2	10422	53993	56.51%	28.318%
2	4.520	54	63	67	rBV	23846	37803	39.57%	19.827%
3	4.574	67	73	99	rVB	56324	<u>95538</u>	100.00%	50.108%
4	4.779	105	111	130	rVB	1574	3330	3.49%	1.747%

Sum of corrected areas: 190664

Signal : EIC Ion 473.00 (472.70 to 473.70): 00801012.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.499	53	59	66	rBV	467	1355	3.02%	2.532%
2	4.574	66	73	97	rVB	25944	<u>44934</u>	100.00%	83.957%
3	4.779	104	111	120	rVB	1058	1822	4.05%	3.404%
4	5.031	146	158	172	rVB3	1821	5409	12.04%	10.107%

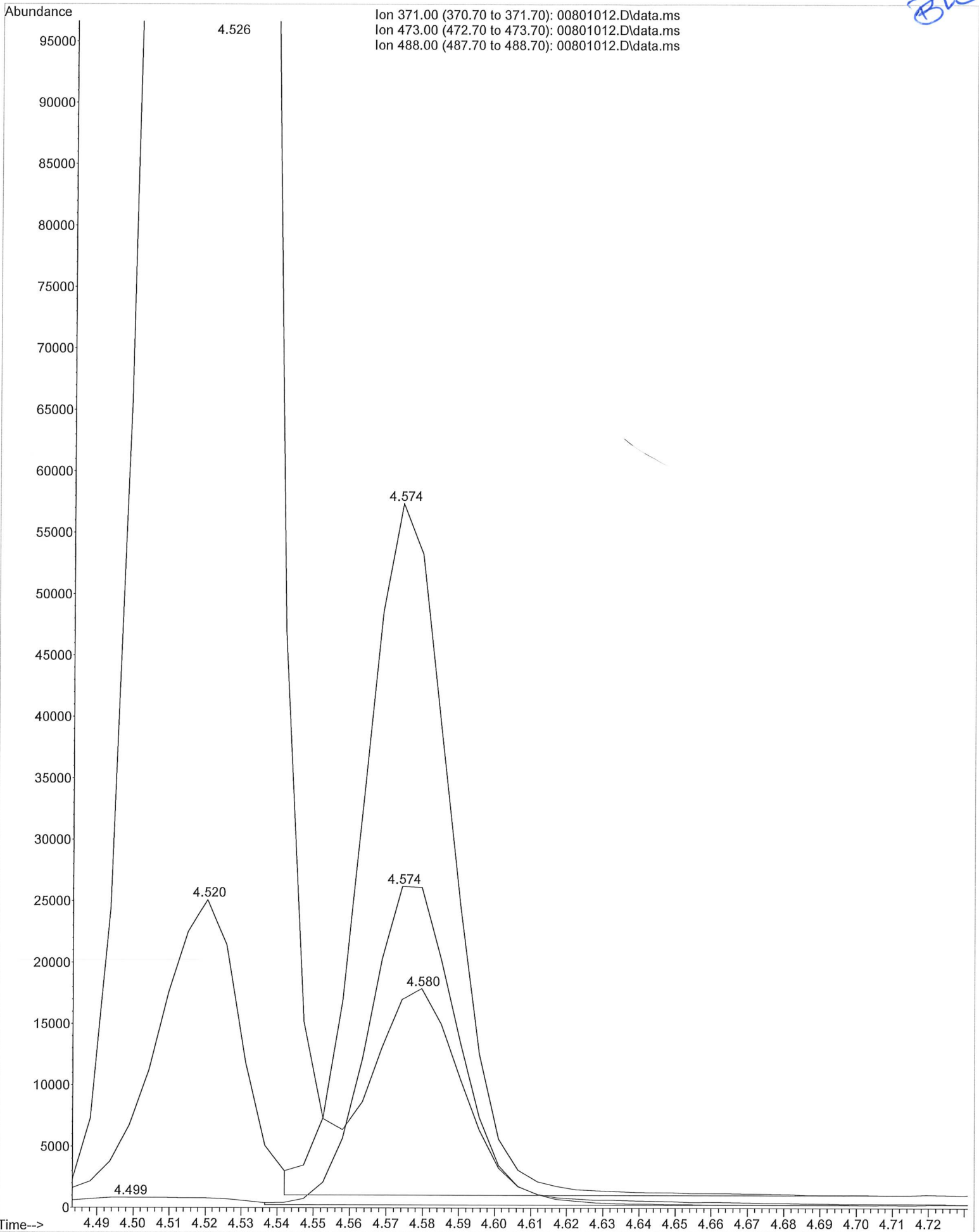
Sum of corrected areas: 53520

Signal : EIC Ion 488.00 (487.70 to 488.70): 00801012.D\data.ms

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	4.526	52	64	70	rBV	484258	768183	100.00%	96.064%
2	4.580	70	74	105	rVB	17638	<u>31477</u>	4.10%	3.936%

Sum of corrected areas: 799660

BW



Run Date: 3/10/2020

Worklist #: 4075

Laboratory Case #: BRC3 Lot 68460

Case Calculations:

EIC Ion 371 Corr. Area: 95538

EIC Ion 473 Corr. Area: 44934

EIC Ion 488 Corr. Area: 31477

473:371 Ratio: 0.470

488:371 Ratio: 0.329

Quality Control Calculations:

	1st Control	2nd Control
EIC Ion 371 Corr. Area:	317491	365455
EIC Ion 473 Corr. Area:	149252	170892
EIC Ion 488 Corr. Area:	96017	108459
Ratio of 473:371:	0.470	0.468
		Average: 0.469
Ratio of 488:371:	0.302	0.297
		Average: 0.300

Acceptable Retention Time Range: 4.480 to 4.679
~~4.479 BW 3-12-20~~ to ~~4.680 BW 3-12-20~~

Approximate Minimum Corrected Area of 371 Ion: 63498

Acceptable 473:371 Ratio Range: 0.375 to 0.563

Acceptable 488:371 Ratio Range: 0.240 to 0.360

Toxicology AM method 3 external urine preparation information

Stock solution 45 ul (100 ug/ml) C-THC in 2.455 mls neg urine lot 11420

Ppd 3/10/20 Exp: 9/1/20 lot 31020 by baw

Drug	lot	expiration
C-THC	FE07171501	9/1/2020