

Analysis: 6DaT-1

Path: Home/YL Research

Based on method: Triplets Method

Created	24-May-2019 13:45:13	visionCATSuser
Modified	01-Jun-2019 18:11:45	visionCATSuser
Last HPTLC log	01-Jun-2019 18:11:45	Analysis modified
Explorer notes		

Track	Vial ID	Description	Volume	Position	Type
1	MeOH blank	MeOH Blank	2.0 µl	A1	Sample
2	Mixture 100	Mixture 500ng	5.0 µl	B1	Sample
3	9-THC 100	D9-THC 500ng	5.0 µl	C1	Reference
4	CBD 100	CBD 500ng	5.0 µl	D1	Reference
5	CBN 100	CBN 500ng	5.0 µl	E1	Reference
6	CBG 100	CBG 500ng	5.0 µl	F1	Reference
7	CBC 100	CBC 500ng	5.0 µl	A2	Reference
8	THCV 100	THCV 500ng	5.0 µl	B2	Reference
9	CBDV 100	CBDV 500ng	5.0 µl	C2	Reference
10	8-THC 100	D8-THC 500ng	5.0 µl	D2	Reference
11	THCA-A 100	THCA-A 500ng	5.0 µl	E2	Reference
12	CBDA 100	CBDA 500ng	5.0 µl	F2	Reference
13	CBGA 100	CBGA 500ng	5.0 µl	A3	Reference
14	Mixture 100	Mixture 500ng	5.0 µl	B1	Sample
15	MeOH blank	MeOH Blank	2.0 µl	A1	Sample

Sequence table notes

A track marked with 🚩 means: the application type is overridden in some evaluation(s).

System setup:

Software	Server User-PC, version 2.5.18072.1
ATS4	S/N:080713
Chamber	N/A
Derivatization dip	N/A
Scanner3	S/N:031025
Visualizer	S/N:230515

Chromatography

Plate layout:

Stationary phase	Merck, HPTLC plates silica gel 60 F 254
Plate format	200.0 x 100.0 mm
Application type	User
Application	Position Y: 10.0 mm, length: 8.0 mm, width: 0.0 mm
Track	First position X: 20.0 mm, distance: 11.4 mm
Solvent front position	70.0 mm
Notes	

Take image clean plate 1a - Visualizer (S/N: 230515):

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Quality	Enhanced
RT White	auto capture, Auto, level 85 %, Band
R 254	auto capture, Auto, level 85 %, Band
Instrument diagnostics	Valid diagnostics
Documentation step label	
Notes	

Application 1 - ATS 4 (S/N: 080713):

Spray gas	NI
Sample solvent type	Methanol
Filling speed	15 µl/s
Predosage volume	200 nl
Retraction volume	200 nl
Dosage speed	150 nl/s
Filling quality	User
Rinsing cycles / vacuum	1 / 4 s
Filling cycles / vacuum	1 / 4 s
Rinsing solvent name	Methanol
Nozzle temperature	Unheated
Rack in use	Standard
Instrument diagnostics	Valid diagnostics
Notes	

Development 1 - Chamber:

Tank	TTC 20x10
Mobile phase	
Saturation time	20 min
Use saturation pad	true
Use smartALERT	false
Volume front through	10 ml
Volume rear through	20 ml
Drying time	5 min
Drying temperature	Room temperature
Notes	

Take image developed plate 1a - Visualizer (S/N: 230515):

Quality	Enhanced
RT White	auto capture, Auto, level 85 %, Band
R 254	auto capture, Auto, level 85 %, Band
R 366	auto capture, Auto, level 85 %, Band
Instrument diagnostics	Valid diagnostics
Documentation step label	
Notes	

Scan developed plate 1b - Scanner 3 (S/N: 031025):

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Scanner type	Single λ
Optimization for	Resolution
Measurement mode	Absorption
Filter	n/a
Detector mode	Automatic
Scanning speed	20 mm/s
Data resolution	100 $\mu\text{m}/\text{step}$
Slit	5 x 0.2 mm, micro
Partial scan	No
Lamp	Deuterium & Tungsten
Wavelength(s)	254 nm
Instrument diagnostics	Valid diagnostics
Documentation step label	
Notes	

Derivatization 1 - dip:

Reagent name	Fast Blue B salt
Dipping speed	3
Dipping time	5 s
Reagent preparation	1g Fast Blue B salt in 200mL water
Heating	none
Notes	Air dry for 5 minutes

Take image derivatized plate 1a - Visualizer (S/N: 230515):

Quality	Enhanced
RT White	auto capture, Auto, level 85 %, Band
R 366	auto capture, Auto, level 85 %, Band
Instrument diagnostics	Valid diagnostics
Documentation step label	
Notes	

System suitability tests:

SST settings:

SST tracks	
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Data acquisition

Application 1 - ATS 4 (S/N: 080713):

Executed	24-May-2019 13:47:24 visionCATSuser
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Development 1 - Chamber:

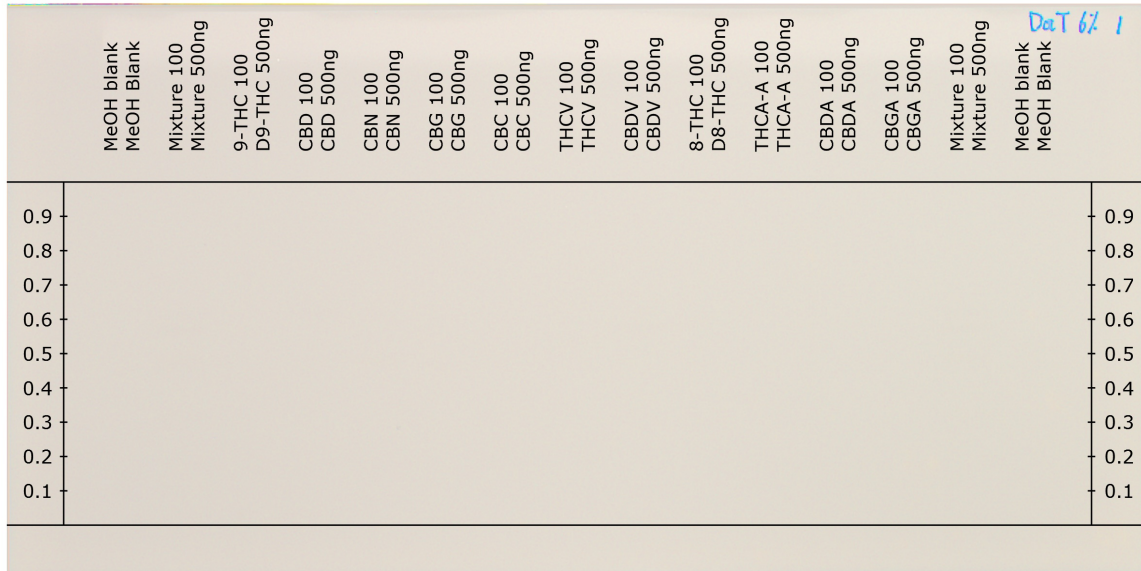
Executed	24-May-2019 14:27:29 visionCATSuser
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Take image developed plate 1a - Visualizer (S/N: 230515):

Executed	24-May-2019 15:04:29 visionCATSuser
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RT White

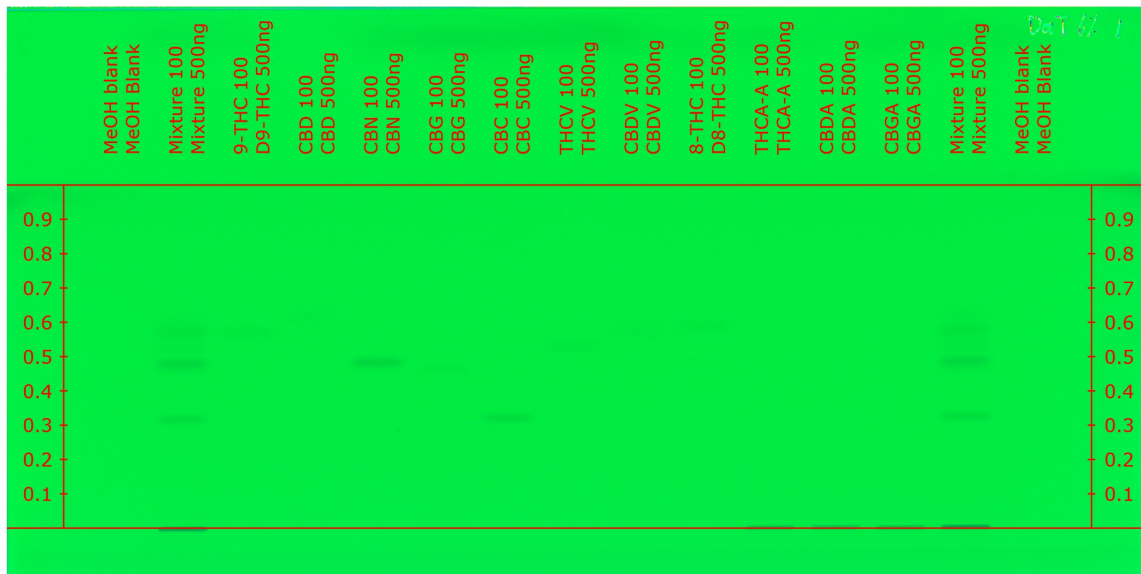
visionCATS
Developed, RemTransVis



Exposure	0.090 s
Contrast	1
Normalized exposure	Disabled
Clarify	Disabled
White balance	1.00, 1.00, 1.00

R 254

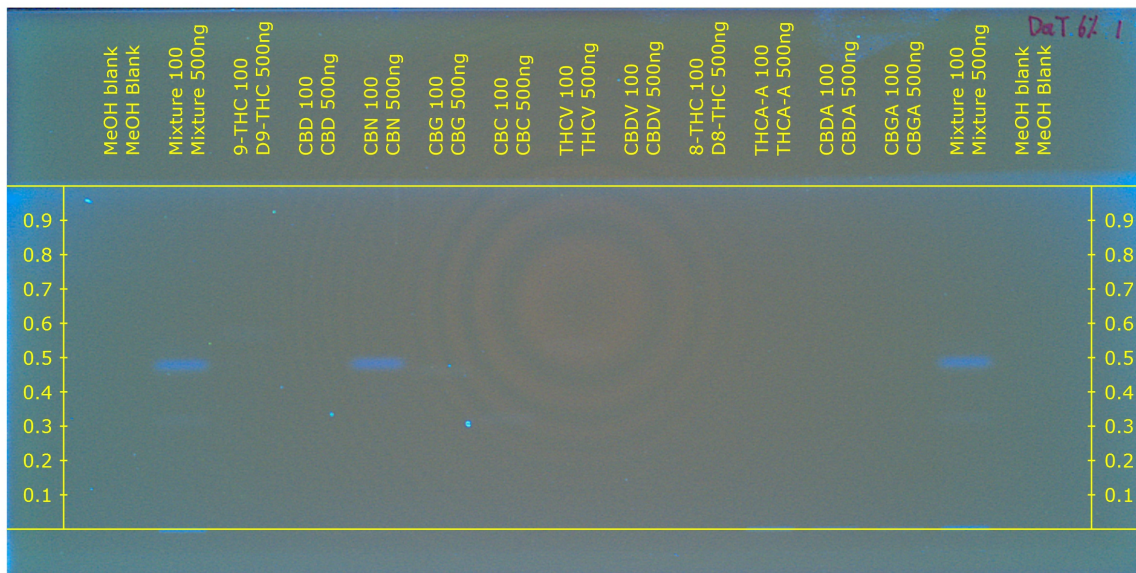
Developed, Remission254



Exposure	0.282 s
Contrast	1
Normalized exposure	Disabled
Clarify	Disabled
White balance	1.00, 1.00, 1.00

6DaT-1
R 366

visionCATS
Developed, Remission366



Exposure	9.999 s
Contrast	1
Normalized exposure	Disabled
Clarify	Disabled
White balance	1.00, 1.00, 1.00

Scan developed plate 1b - Scanner 3 (S/N: 031025):

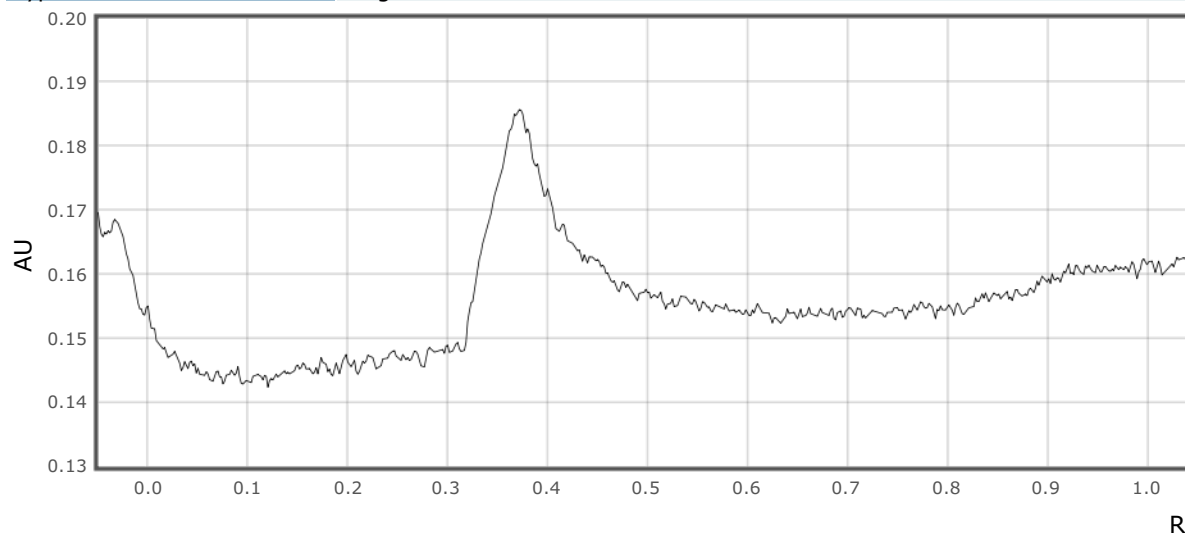
Executed	24-May-2019 15:08:19 visionCATSuser
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Scan:

Wavelength	254 nm
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Track 1:

Type	Single λ
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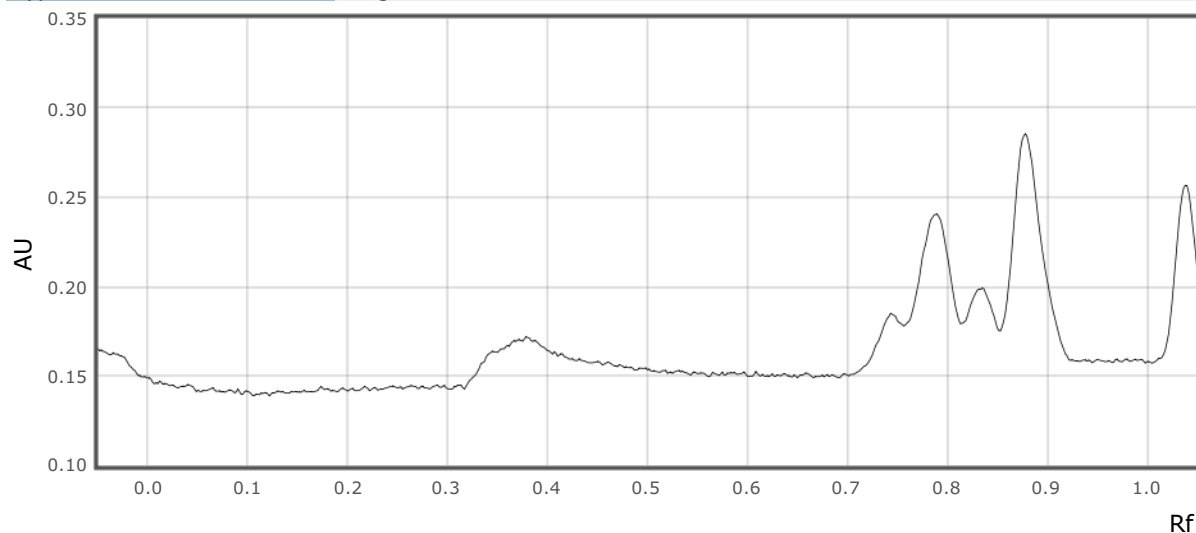


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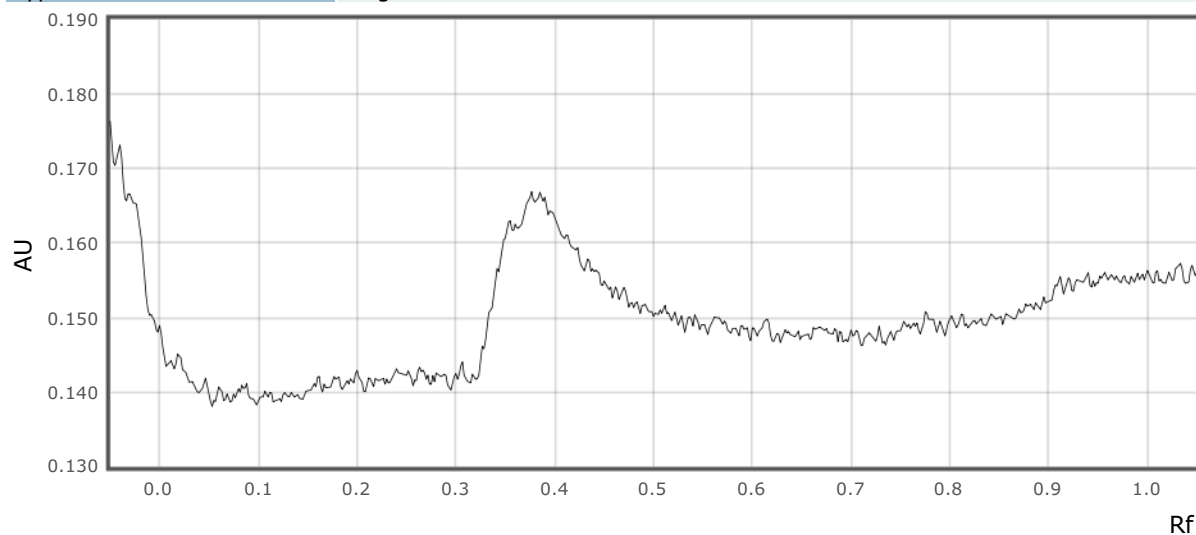
Track 2:

Type Single λ



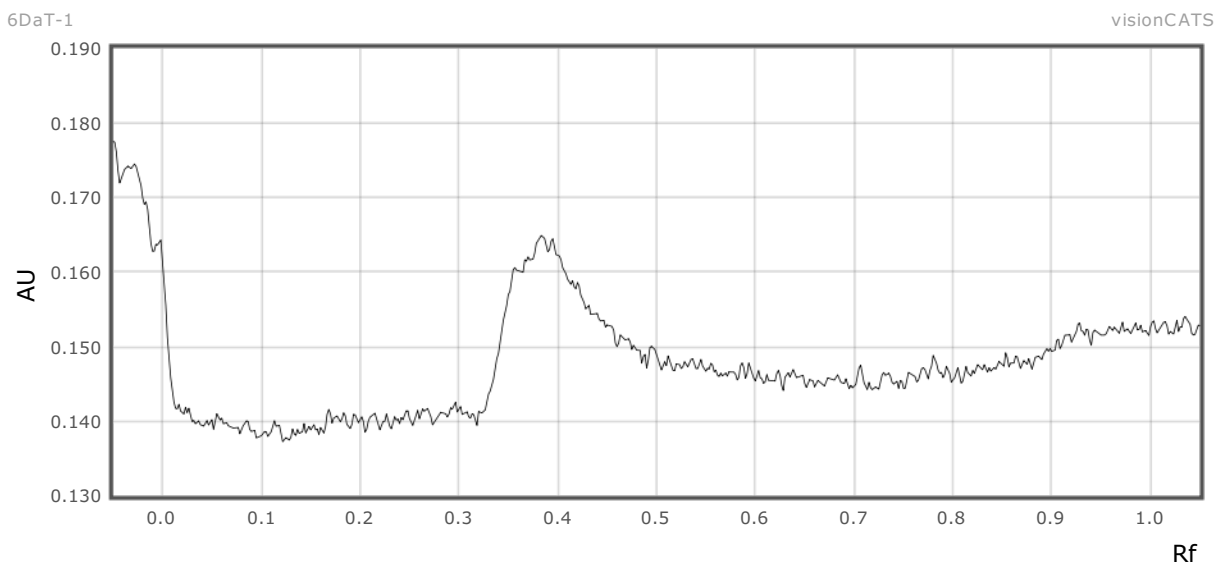
Track 3:

Type Single λ



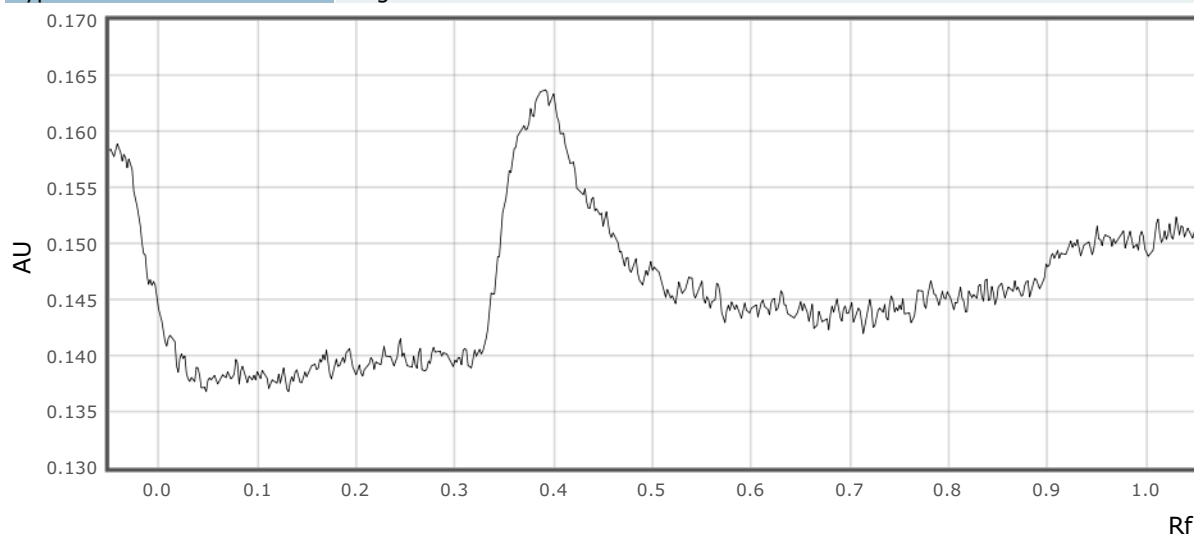
Track 4:

Type Single λ



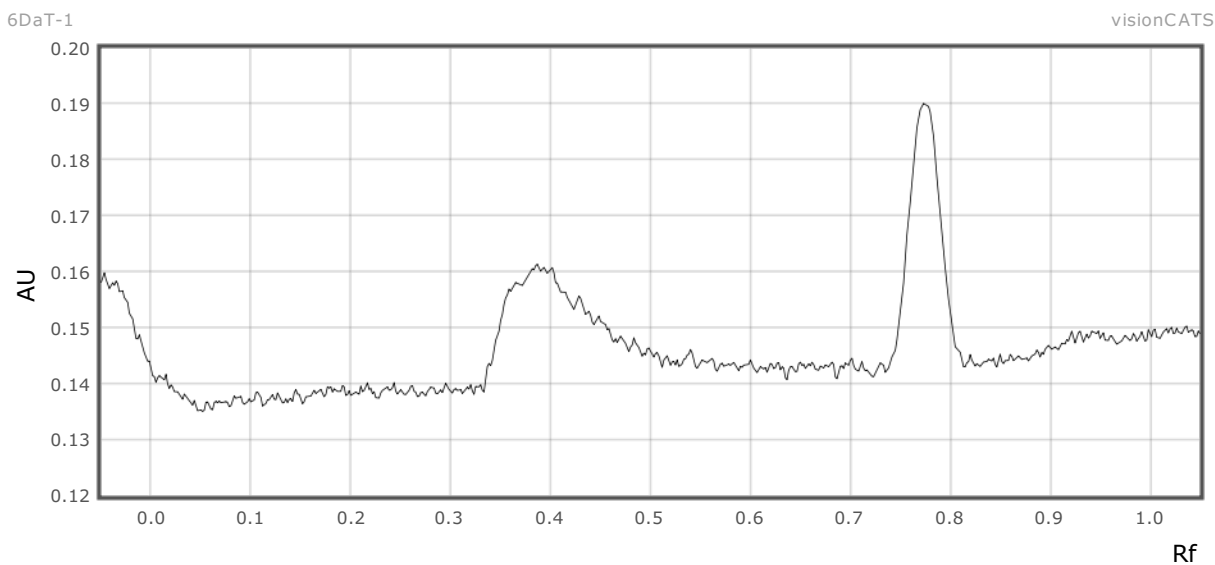
Track 5:

Type Single λ



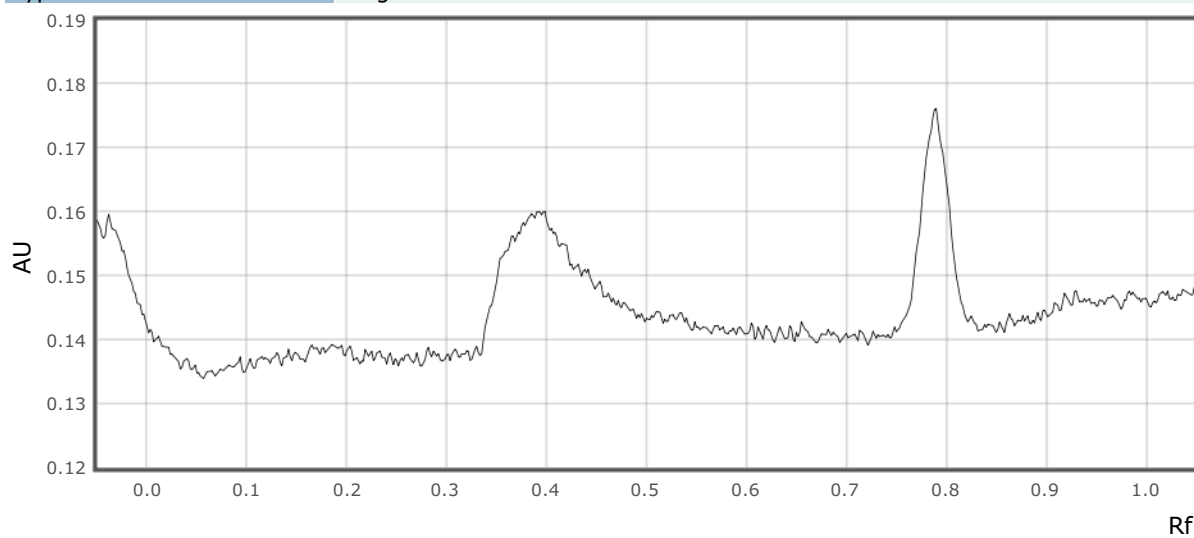
Track 6:

Type Single λ



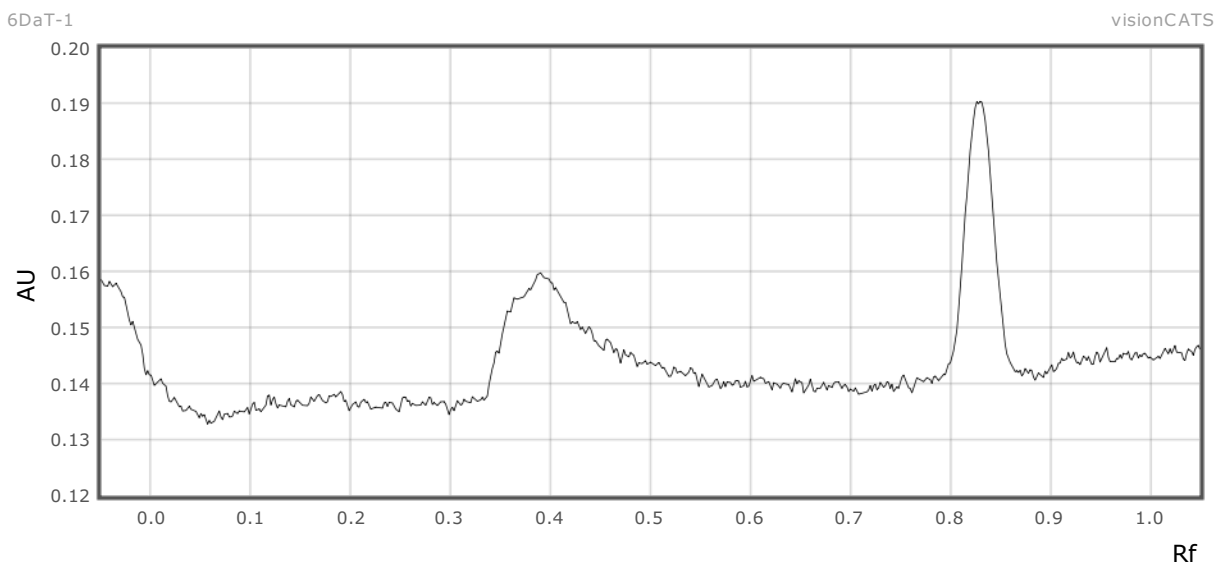
Track 7:

Type Single λ



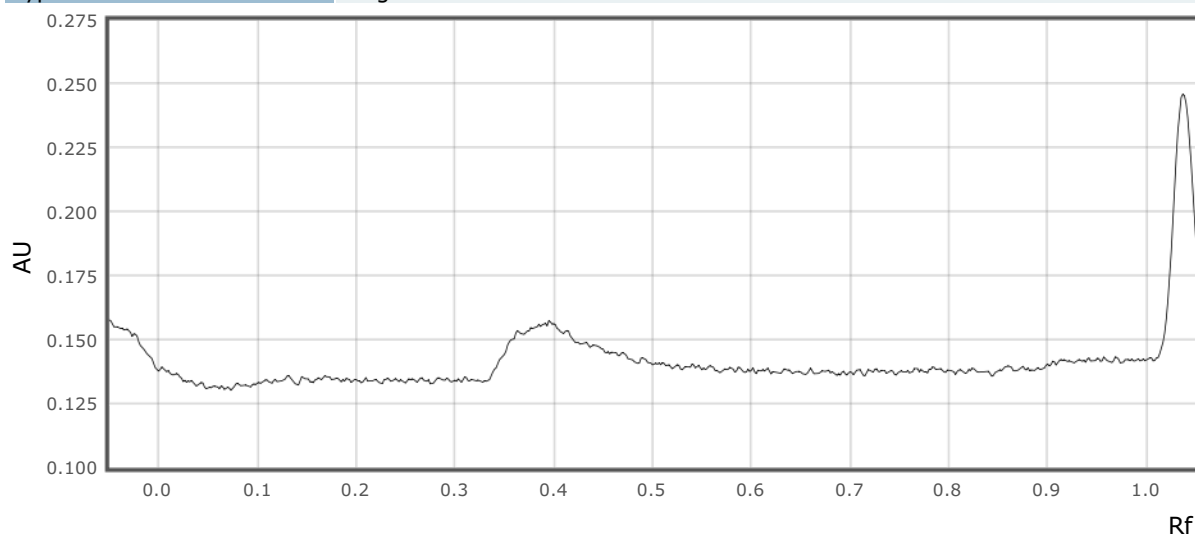
Track 8:

Type Single λ



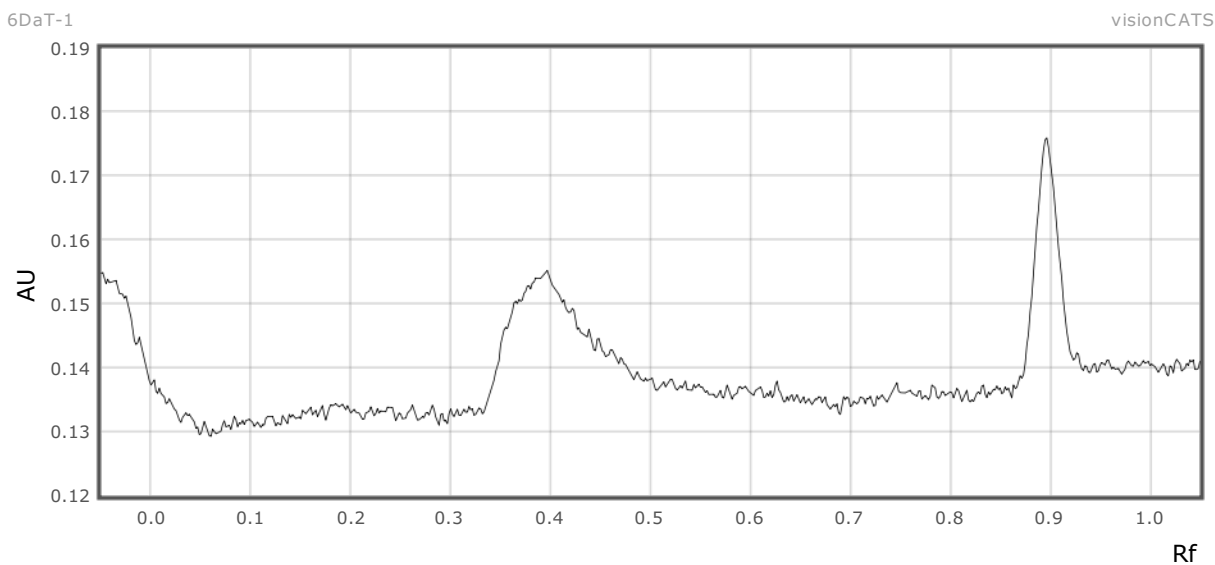
Track 9:

Type Single λ



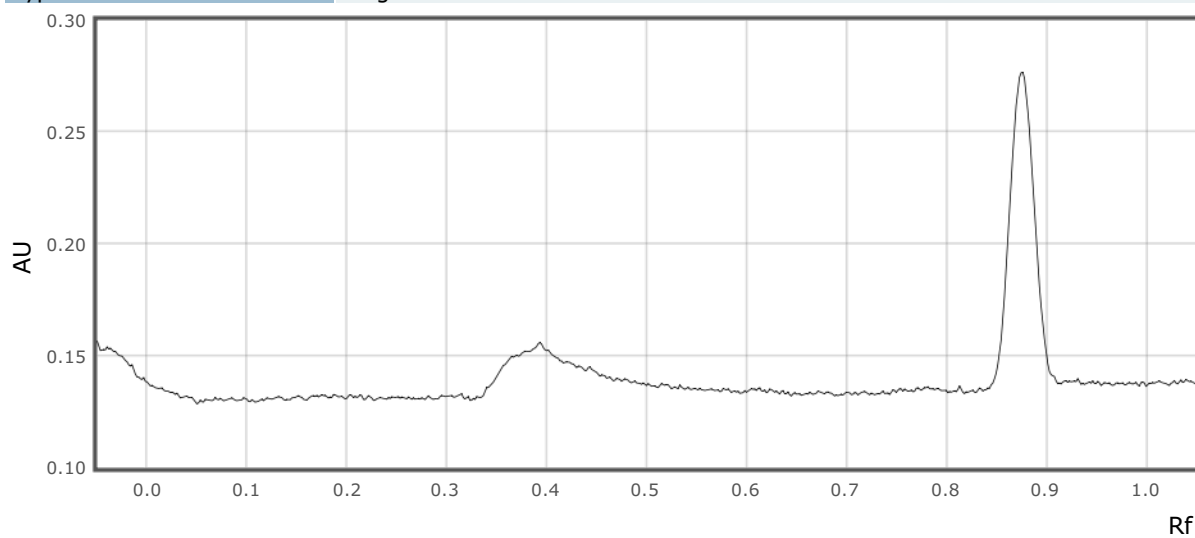
Track 10:

Type Single λ



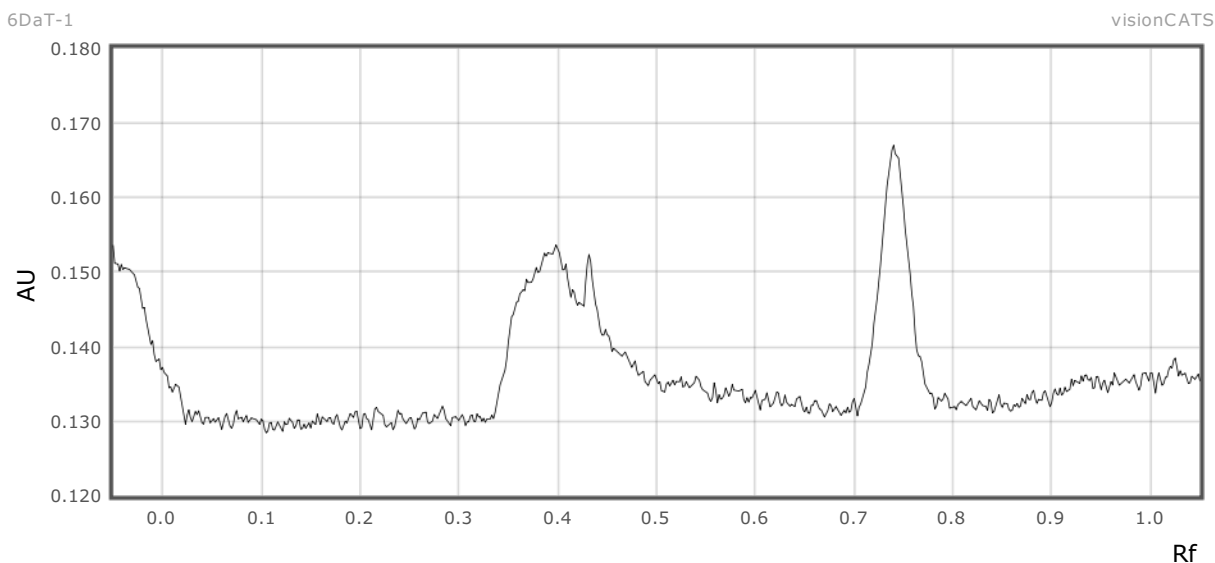
Track 11:

Type Single λ



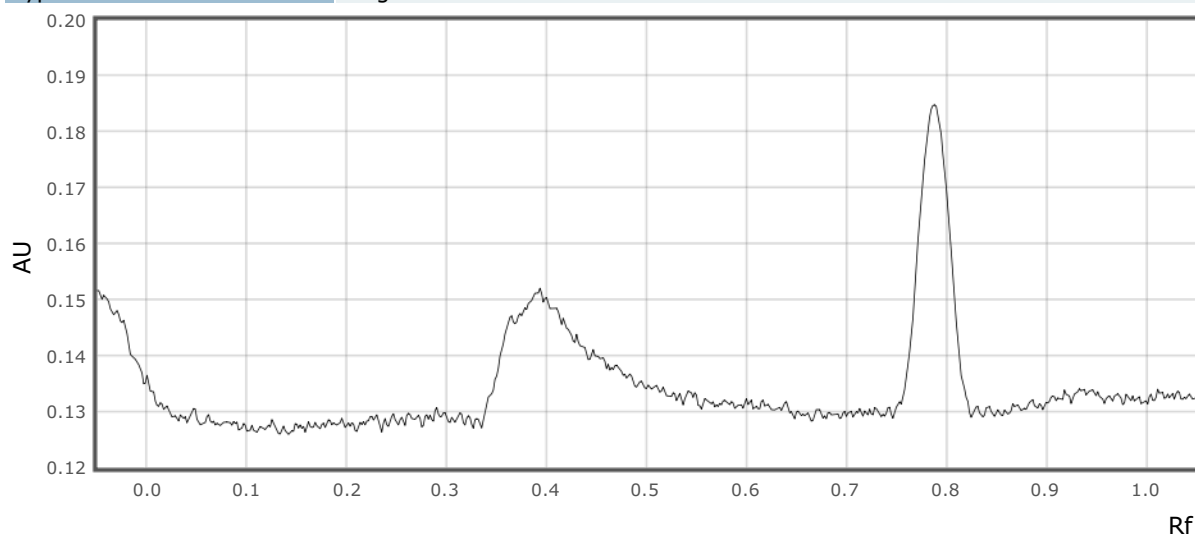
Track 12:

Type Single λ



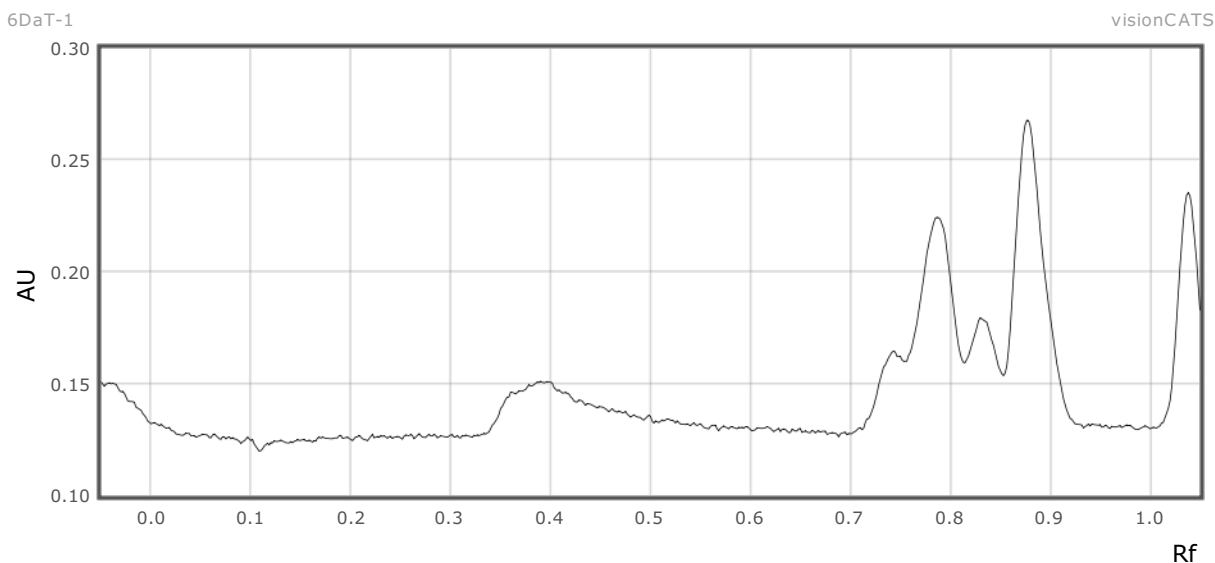
Track 13:

Type Single λ



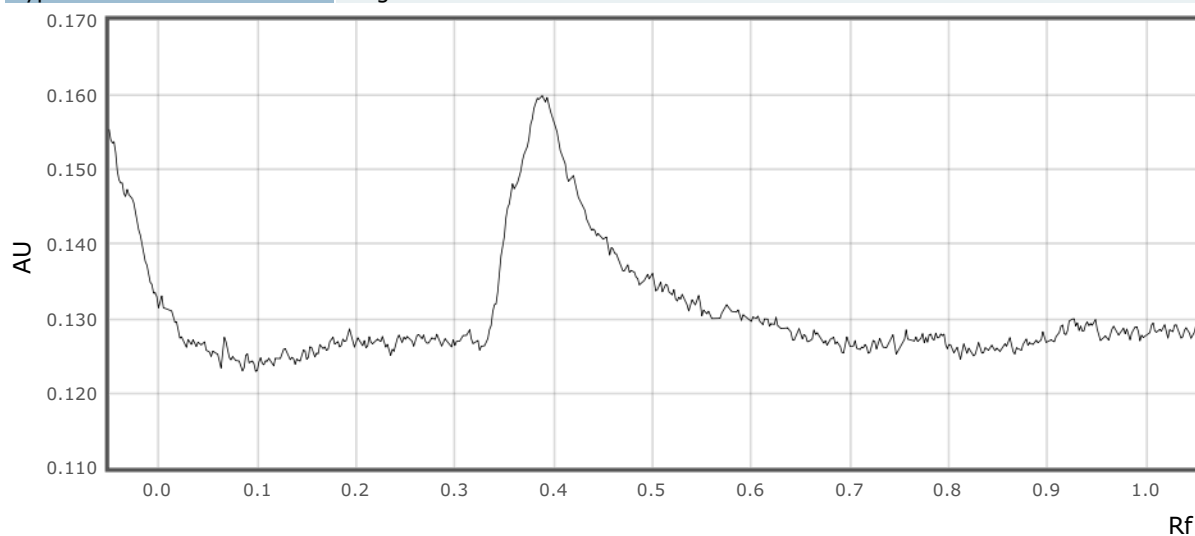
Track 14:

Type Single λ



Track 15:

Type Single λ



Derivatization 1 - dip:

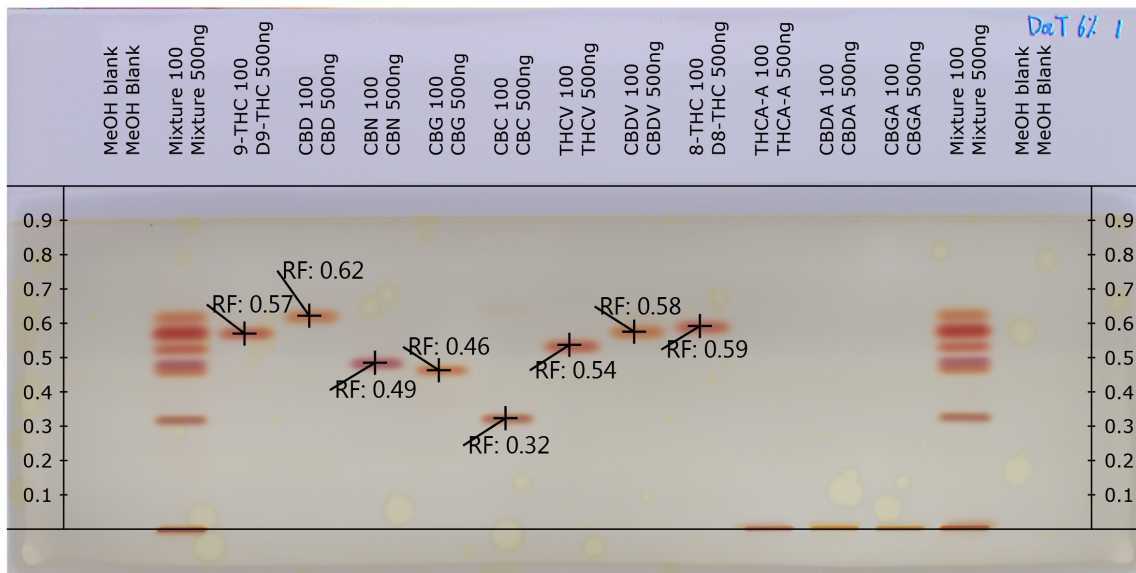
Executed 24-May-2019 15:14:40 visionCATSuser

Take image derivatized plate 1a - Visualizer (S/N: 230515):

Executed 24-May-2019 15:14:55 visionCATSuser

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RT White

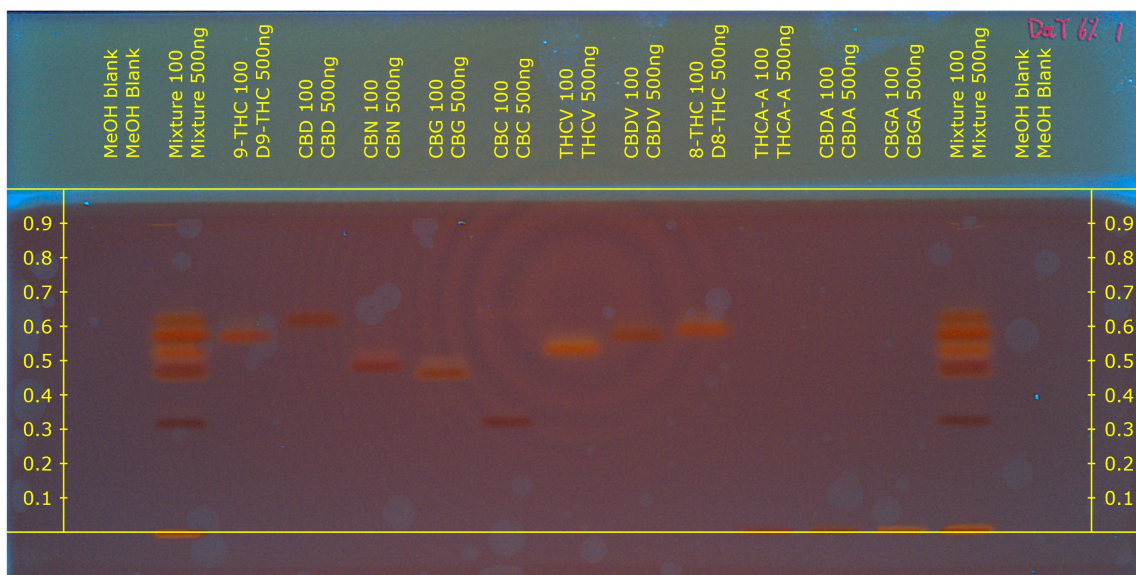
visionCATS
Derivatized, RemTransVis



Exposure	0.073 s
Contrast	1
Normalized exposure	Disabled
Clarify	Disabled
White balance	1.14, 1.09, 0.83

R 366

Derivatized, Remission366



Exposure	9.999 s
Contrast	1
Normalized exposure	Disabled
Clarify	Disabled
White balance	1.00, 1.00, 1.00

Evaluation 1 :

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Validated	false
Step	Take image derivatized plate 1a
Concentration unit type	Mass / volume
Notes	

Definition:

References:

9-THC 100		
Substance Name	Concentration	Purity
9-THC	100.000 µg/ml	100.00 %

CBD 100		
Substance Name	Concentration	Purity
CBD	100.000 µg/ml	100.00 %

CBN 100		
Substance Name	Concentration	Purity
CBN	100.000 µg/ml	100.00 %

CBG 100		
Substance Name	Concentration	Purity
CBG	100.000 µg/ml	100.00 %

CBC 100		
Substance Name	Concentration	Purity
CBC	100.000 µg/ml	100.00 %

THCV 100		
Substance Name	Concentration	Purity
THCV	100.000 µg/ml	100.00 %

CBDV 100		
Substance Name	Concentration	Purity
CBDV	100.000 µg/ml	100.00 %

8-THC 100		
Substance Name	Concentration	Purity
8-THC	100.000 µg/ml	100.00 %

THCA-A 100		
Substance Name	Concentration	Purity
THCA-A	100.000 µg/ml	100.00 %

CBDA 100		
Substance Name	Concentration	Purity
CBDA	100.000 µg/ml	100.00 %

CBGA 100		
Substance Name	Concentration	Purity
CBGA	100.000 µg/ml	100.00 %

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Samples:

Vial ID	Amount	Volume solution	Reference amount	Related to
MeOH blank		0.00 ml		
Mixture 100		0.00 ml		

Integration parameters:

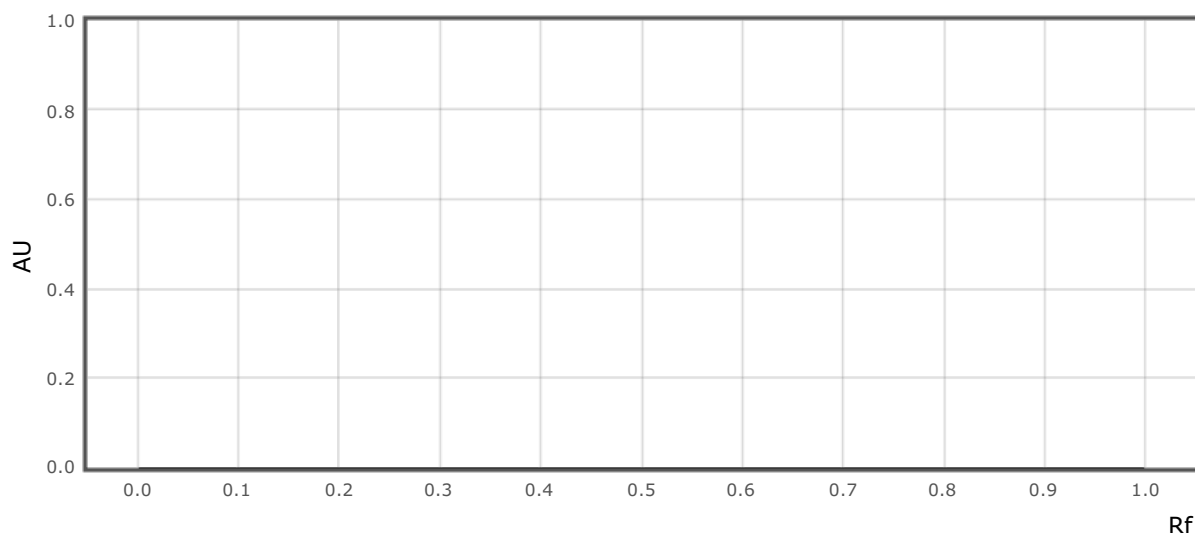
Bounds	[0.000,1.000]
Smoothing	Savitzky-Golay of order 3 and window 7
Baseline correction	Lowest slope with noise 0.05
Profile subtraction	Profile subtraction from track 1
Peaks detection	Gauss (legacy) with sensitivity 0.1, separation 0.1 and threshold 0.1

Scan:

Wavelength	RT White
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Track 1:

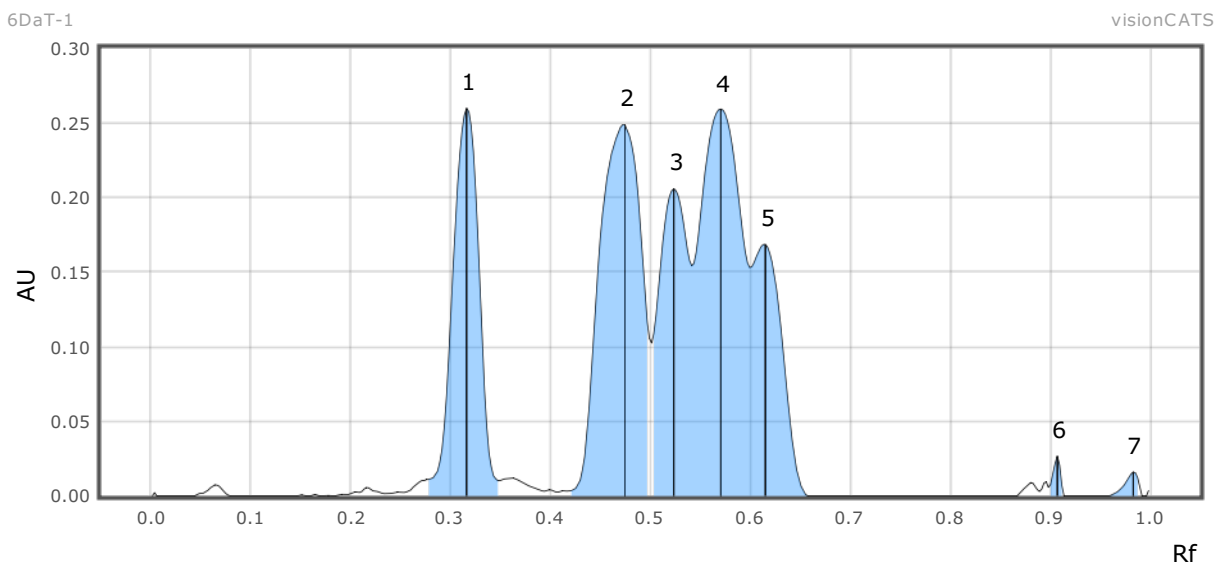
Type	Sample
Vial ID	MeOH blank
Description	MeOH Blank
Volume	2.0 µl



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		

Track 2:

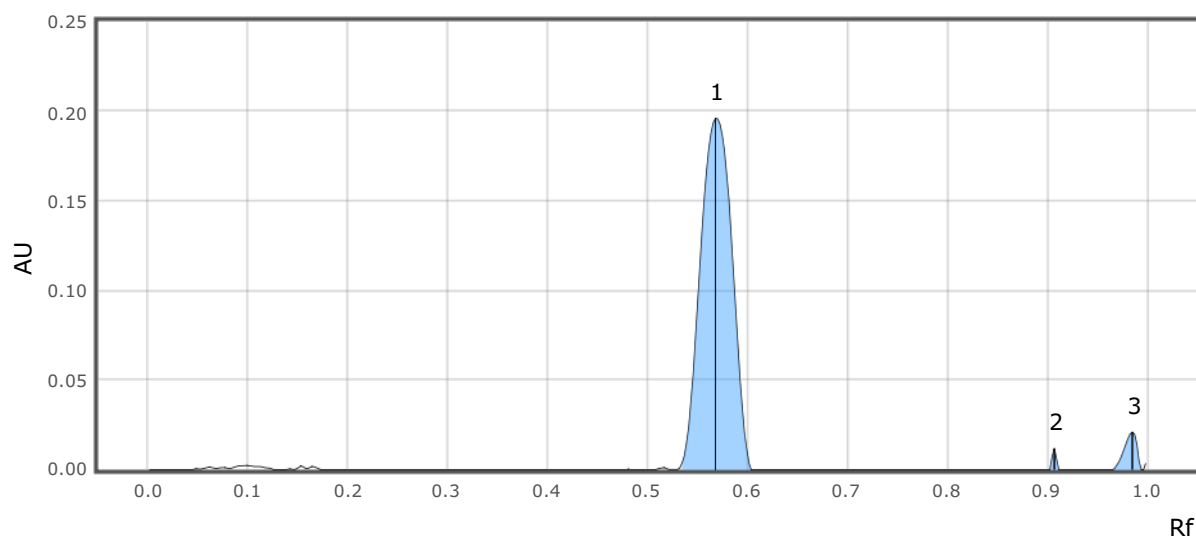
Type	Sample
Vial ID	Mixture 100
Description	Mixture 500ng
Volume	5.0 µl



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.278	0.0110	0.316	0.2595	21.93	0.347	0.0101	0.00787	17.33	No	
2	0.416	0.0032	0.474	0.2485	21.00	0.499	0.1055	0.01192	26.26	No	
3	0.501	0.1024	0.523	0.2056	17.37	0.541	0.1541	0.00688	15.16	No	
4	0.541	0.1541	0.570	0.2591	21.89	0.599	0.1528	0.01240	27.32	No	
5	0.599	0.1528	0.615	0.1684	14.23	0.657	0.0000	0.00585	12.90	No	
6	0.898	0.0058	0.907	0.0265	2.24	0.914	0.0000	0.00023	0.50	No	
7	0.959	0.0000	0.983	0.0159	1.34	0.992	0.0000	0.00024	0.53	No	

Track 3:

Type	Reference
Vial ID	9-THC 100
Description	D9-THC 500ng
Volume	5.0 µl



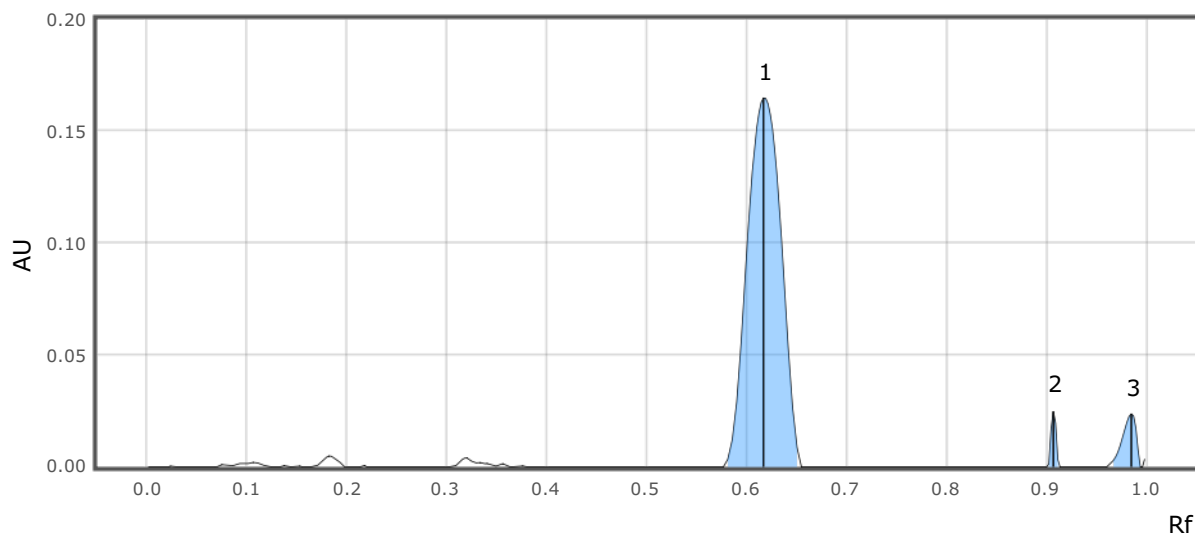
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Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.534	0.0037	0.568	0.1958	85.65	0.604	0.0000	0.00718	95.24	Yes	9-THC
2	0.903	0.0000	0.907	0.0118	5.18	0.912	0.0000	0.00006	0.75	No	
3	0.965	0.0000	0.985	0.0210	9.17	0.994	0.0000	0.00030	4.01	No	

Track 4:

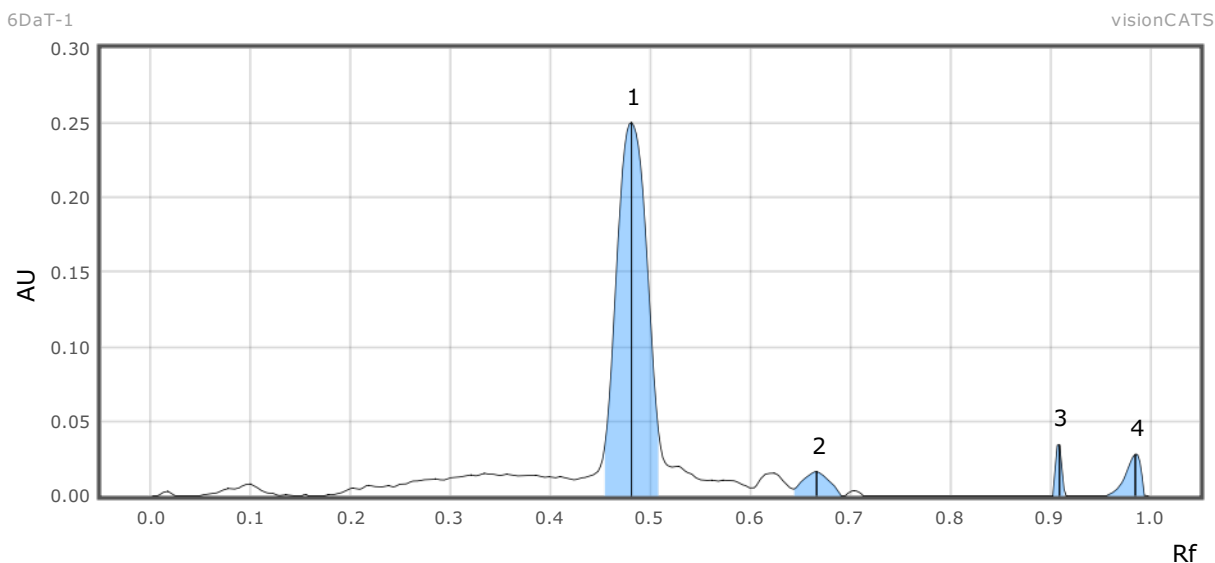
Type	Reference
Vial ID	CBD 100
Description	CBD 500ng
Volume	5.0 µl



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.577	0.0000	0.617	0.1647	77.31	0.655	0.0000	0.00646	92.51	No	CBD
2	0.901	0.0000	0.907	0.0247	11.59	0.914	0.0000	0.00015	2.09	No	
3	0.961	0.0000	0.985	0.0236	11.10	0.994	0.0000	0.00038	5.39	No	

Track 5:

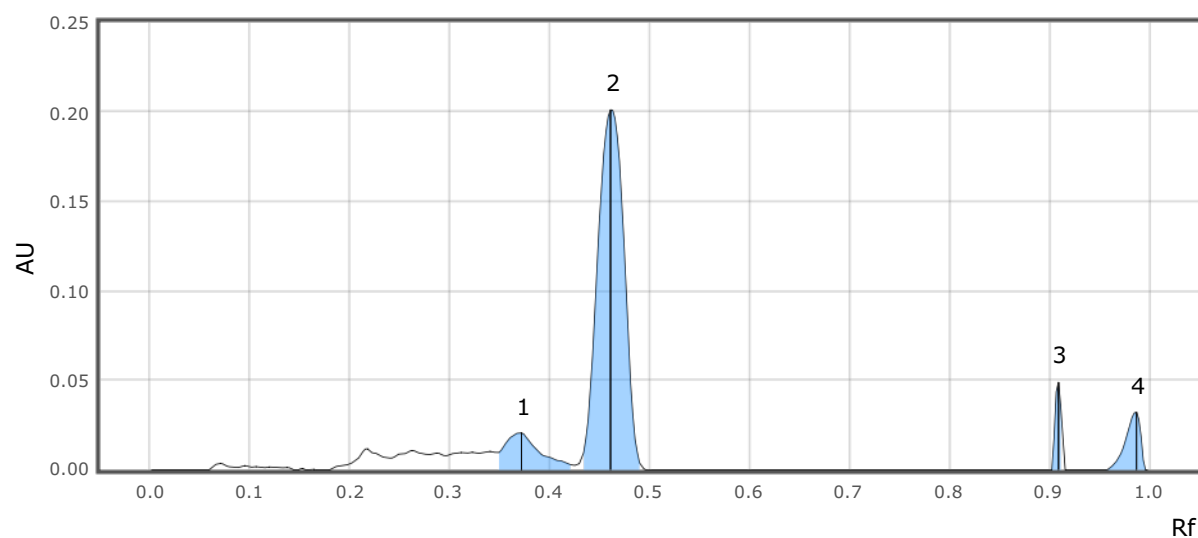
Type	Reference
Vial ID	CBN 100
Description	CBN 500ng
Volume	5.0 µl



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.453	0.0244	0.481	0.2501	76.19	0.510	0.0331	0.00891	88.05	Yes	CBN
2	0.644	0.0045	0.666	0.0162	4.93	0.693	0.0000	0.00049	4.83	No	
3	0.903	0.0000	0.909	0.0341	10.39	0.916	0.0000	0.00024	2.36	No	
4	0.954	0.0000	0.985	0.0279	8.49	0.996	0.0000	0.00048	4.76	No	

Track 6:

Type	Reference
Vial ID	CBG 100
Description	CBG 500ng
Volume	5.0 µl



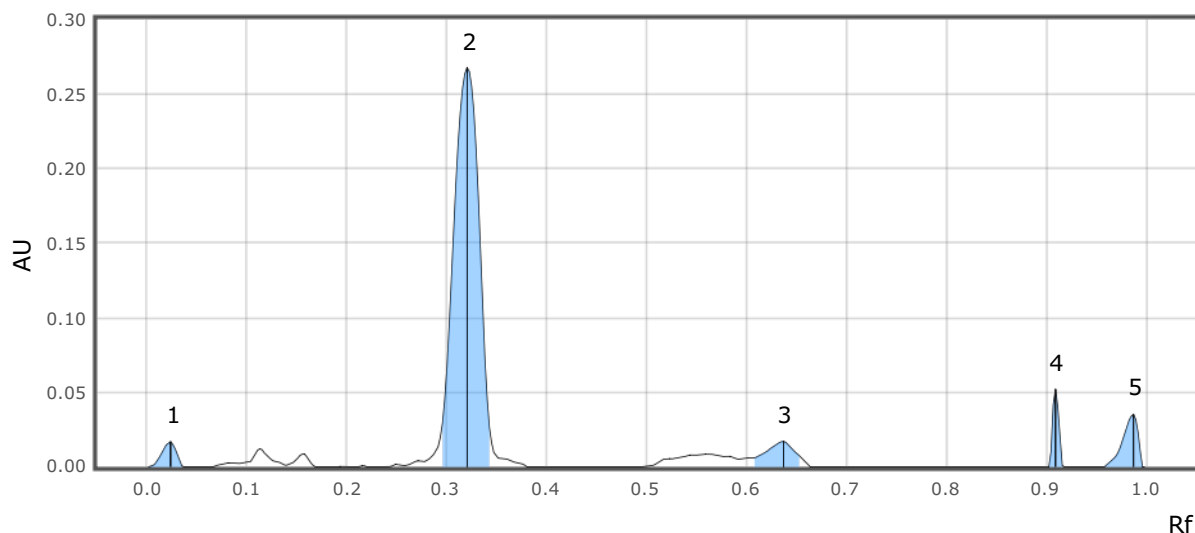
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Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.347	0.0097	0.372	0.0206	6.80	0.425	0.0025	0.00087	10.99	No	
2	0.432	0.0064	0.461	0.2009	66.41	0.490	0.0039	0.00612	77.49	Yes	CBG
3	0.903	0.0000	0.909	0.0489	16.15	0.916	0.0000	0.00036	4.56	No	
4	0.956	0.0000	0.988	0.0322	10.64	0.996	0.0000	0.00055	6.96	No	

Track 7:

Type	Reference
Vial ID	CBC 100
Description	CBC 500ng
Volume	5.0 µl



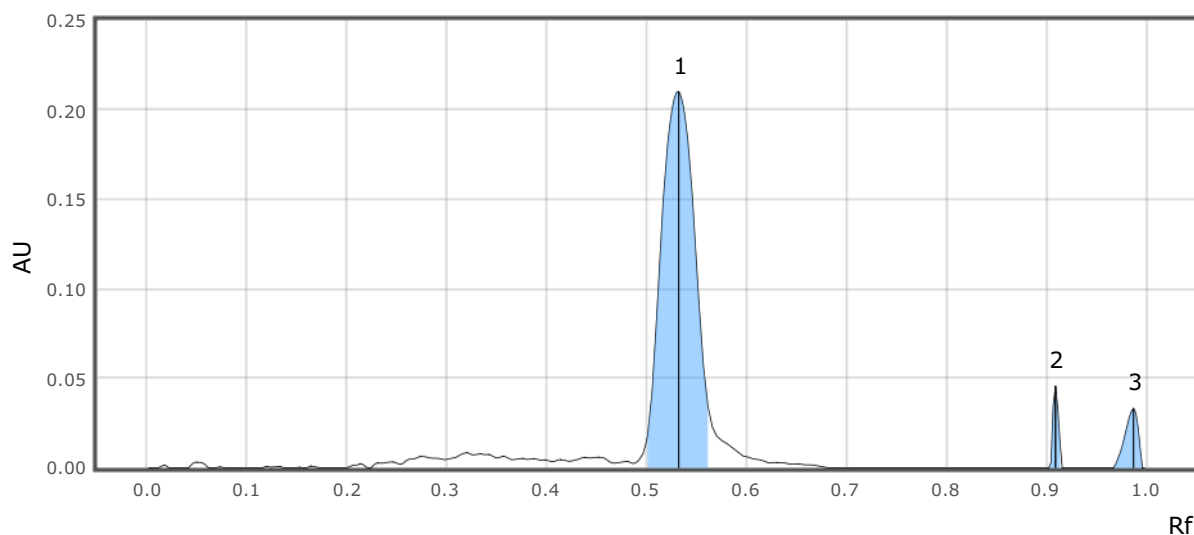
Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.001	0.0000	0.024	0.0169	4.34	0.037	0.0000	0.00027	2.83	No	
2	0.294	0.0199	0.320	0.2674	68.85	0.344	0.0162	0.00784	80.99	Yes	CBC
3	0.606	0.0058	0.637	0.0170	4.38	0.664	0.0000	0.00059	6.05	No	
4	0.903	0.0000	0.909	0.0520	13.39	0.918	0.0000	0.00037	3.82	No	
5	0.956	0.0000	0.988	0.0351	9.03	0.996	0.0000	0.00061	6.31	No	

Track 8:

Type	Reference
Vial ID	THCV 100
Description	THCV 500ng
Volume	5.0 µl

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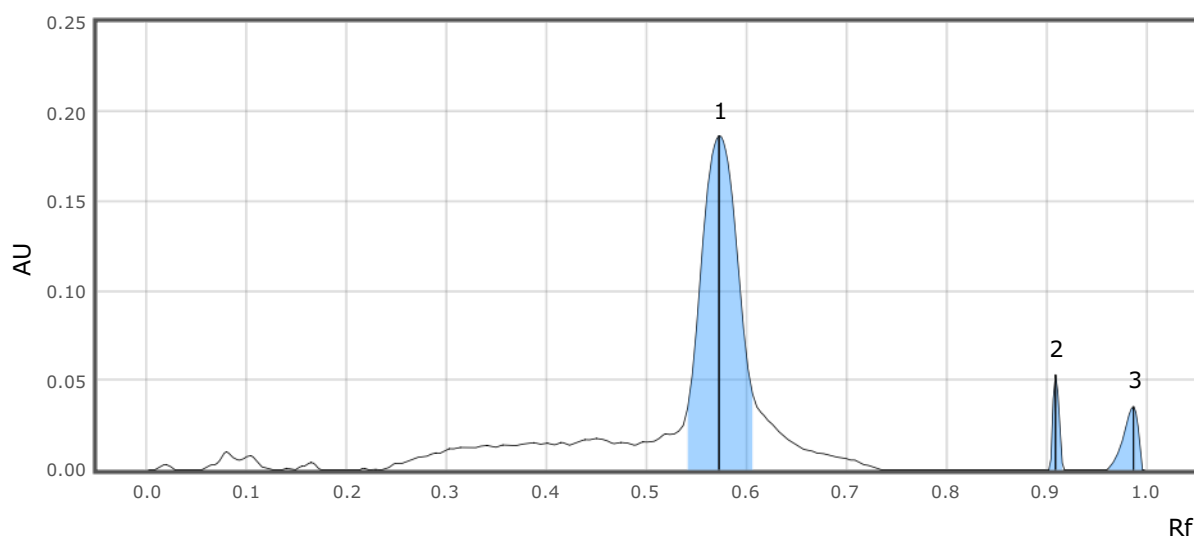
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Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.499	0.0124	0.532	0.2101	72.69	0.566	0.0229	0.00822	91.21	Yes	THCV
2	0.903	0.0000	0.909	0.0456	15.79	0.916	0.0000	0.00029	3.25	No	
3	0.967	0.0000	0.988	0.0333	11.52	0.996	0.0000	0.00050	5.54	No	

Track 9:

Type	Reference
Vial ID	CBDV 100
Description	CBDV 500ng
Volume	5.0 µl

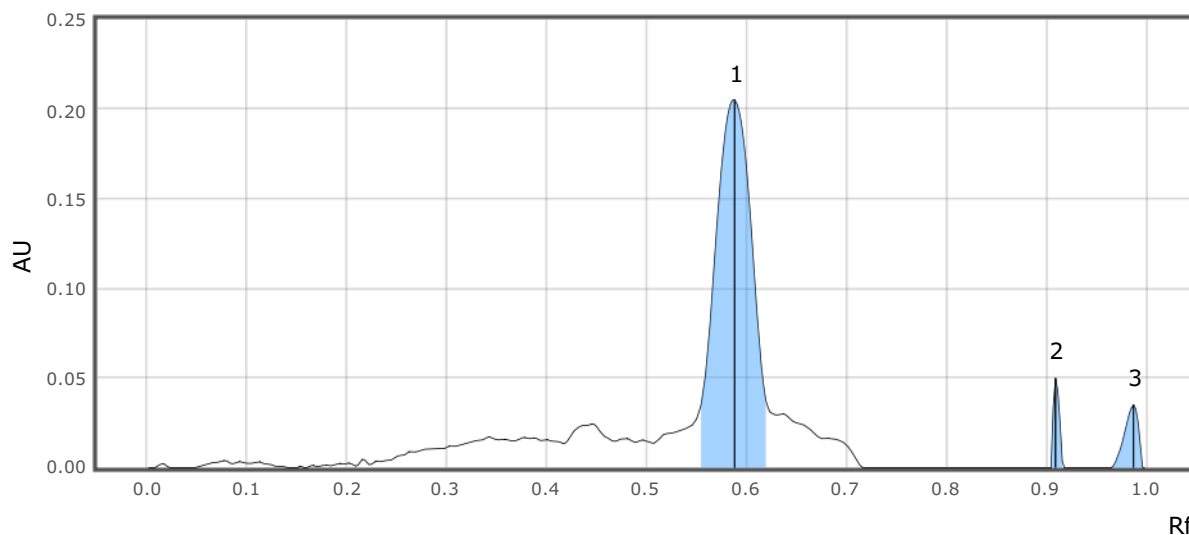


Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.539	0.0294	0.573	0.1865	67.82	0.606	0.0426	0.00803	89.14	Yes	CBDV
2	0.903	0.0000	0.909	0.0531	19.30	0.918	0.0000	0.00038	4.24	No	
3	0.961	0.0000	0.988	0.0354	12.88	0.996	0.0000	0.00060	6.62	No	

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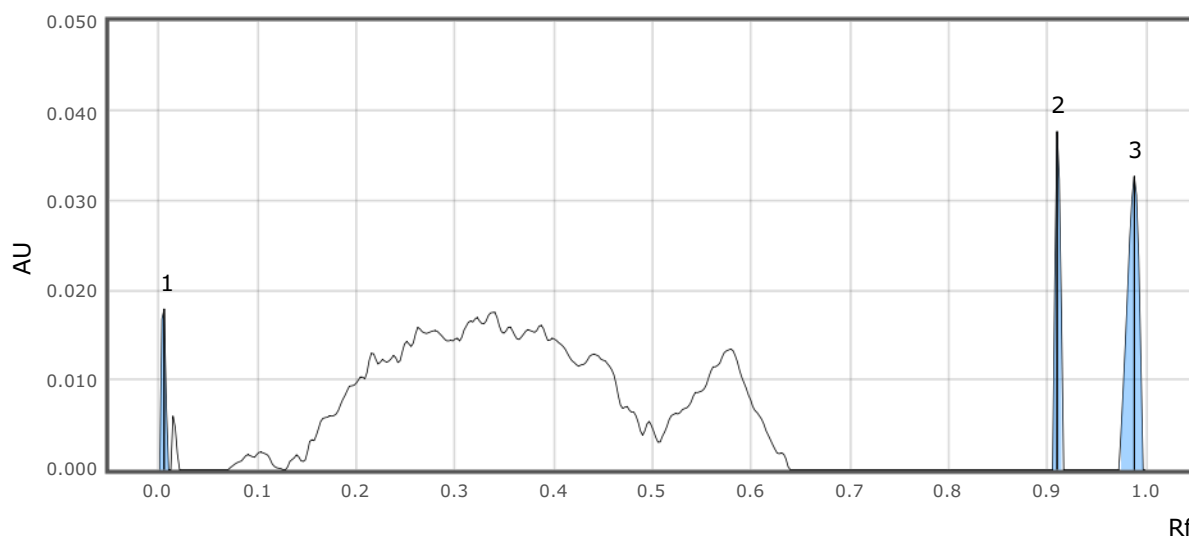
visionCATS

Track 10:											
Type	Reference										
Vial ID	8-THC 100										
Description	D8-THC 500ng										
Volume	5.0 µl										



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.554	0.0349	0.588	0.2050	70.71	0.620	0.0375	0.00859	90.50	Yes	8-THC
2	0.905	0.0000	0.909	0.0499	17.21	0.918	0.0000	0.00035	3.65	No	
3	0.965	0.0000	0.988	0.0350	12.08	0.999	0.0000	0.00055	5.85	No	

Track 11:											
Type	Reference										
Vial ID	THCA-A 100										
Description	THCA-A 500ng										
Volume	5.0 µl										



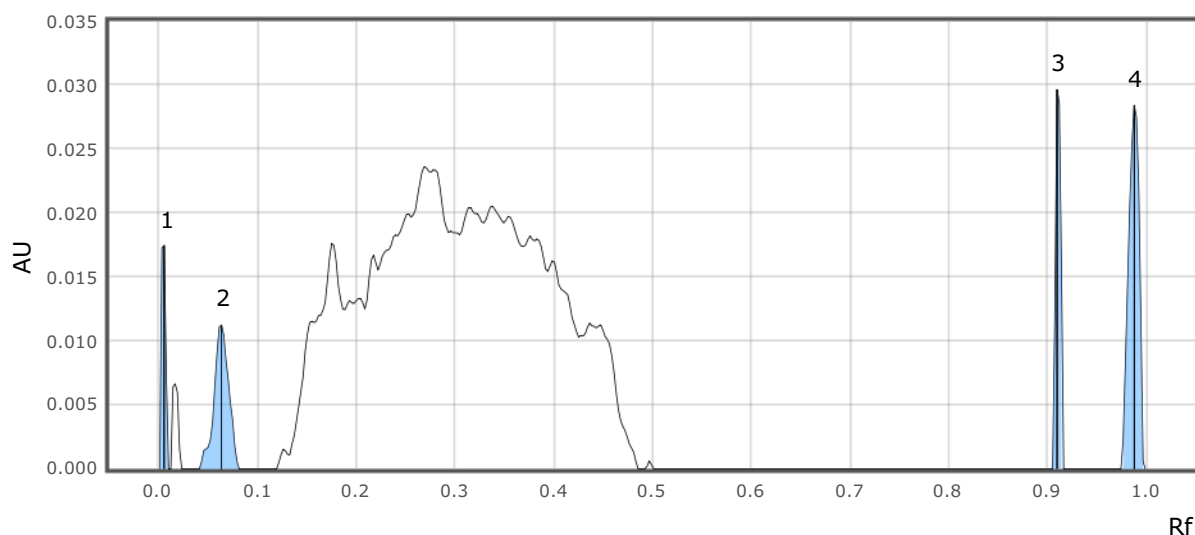
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Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.001	0.0000	0.006	0.0179	20.29	0.010	0.0000	0.00009	11.86	No	THCA-A
2	0.905	0.0000	0.909	0.0377	42.68	0.916	0.0000	0.00024	30.71	No	
3	0.972	0.0000	0.988	0.0327	37.03	0.999	0.0000	0.00045	57.43	No	

Track 12:

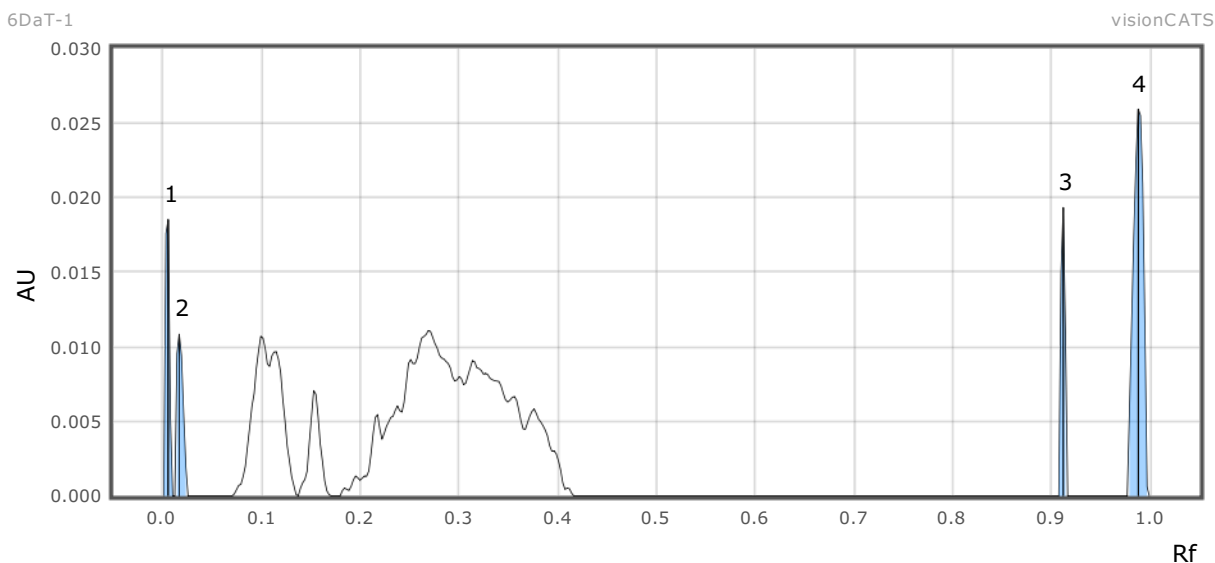
Type	Reference
Vial ID	CBDA 100
Description	CBDA 500ng
Volume	5.0 µl



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.001	0.0000	0.006	0.0175	20.13	0.010	0.0000	0.00009	11.14	No	CBDA
2	0.041	0.0000	0.064	0.0112	12.96	0.082	0.0000	0.00019	23.06	No	
3	0.905	0.0000	0.909	0.0296	34.17	0.916	0.0000	0.00019	22.77	No	
4	0.974	0.0000	0.988	0.0284	32.73	0.999	0.0000	0.00036	43.02	No	

Track 13:

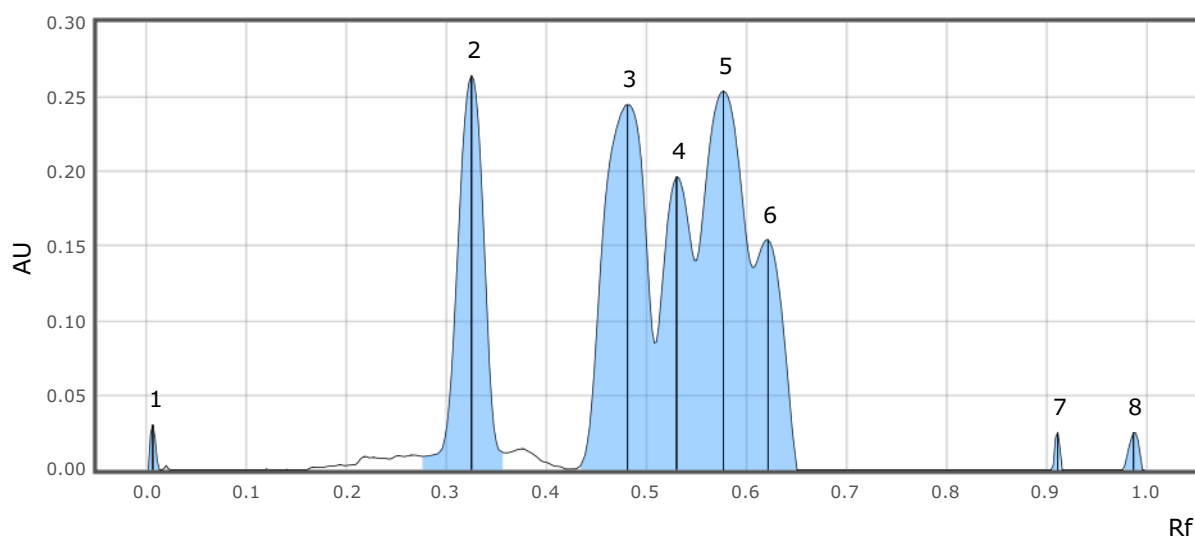
Type	Reference
Vial ID	CBGA 100
Description	CBGA 500ng
Volume	5.0 µl



Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.001	0.0000	0.006	0.0185	24.82	0.010	0.0000	0.00009	15.66	No	CBGA
2	0.012	0.0000	0.017	0.0108	14.53	0.026	0.0000	0.00008	14.07	No	
3	0.907	0.0000	0.912	0.0193	25.90	0.916	0.0000	0.00010	16.96	No	
4	0.976	0.0000	0.988	0.0259	34.75	0.999	0.0000	0.00031	53.30	No	

Track 14:

Type	Sample
Vial ID	Mixture 100
Description	Mixture 500ng
Volume	5.0 µl



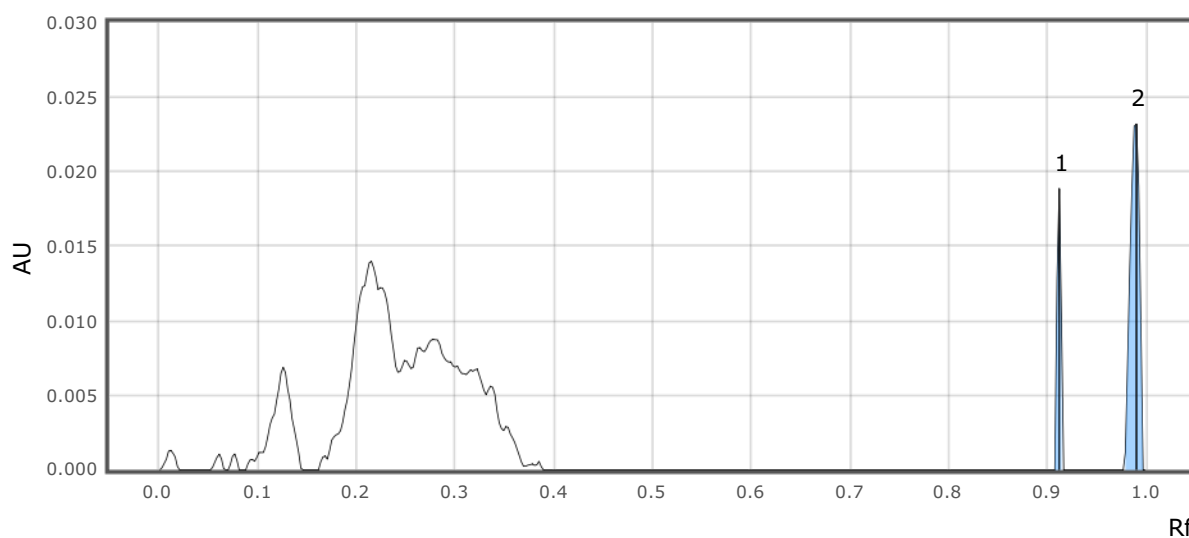
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Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.001	0.0000	0.006	0.0302	2.53	0.012	0.0000	0.00018	0.42	No	
2	0.276	0.0089	0.325	0.2640	22.13	0.358	0.0114	0.00804	18.53	No	
3	0.430	0.0006	0.481	0.2446	20.51	0.508	0.0849	0.01173	27.02	No	
4	0.508	0.0849	0.530	0.1962	16.45	0.548	0.1401	0.00631	14.55	No	
5	0.548	0.1401	0.577	0.2537	21.27	0.606	0.1356	0.01179	27.17	No	
6	0.606	0.1356	0.622	0.1540	12.91	0.651	0.0000	0.00492	11.33	No	
7	0.905	0.0000	0.912	0.0252	2.11	0.916	0.0000	0.00014	0.33	No	
8	0.976	0.0000	0.988	0.0249	2.08	0.999	0.0000	0.00028	0.65	No	

Track 15:

Type	Sample
Vial ID	MeOH blank
Description	MeOH Blank
Volume	2.0 µl



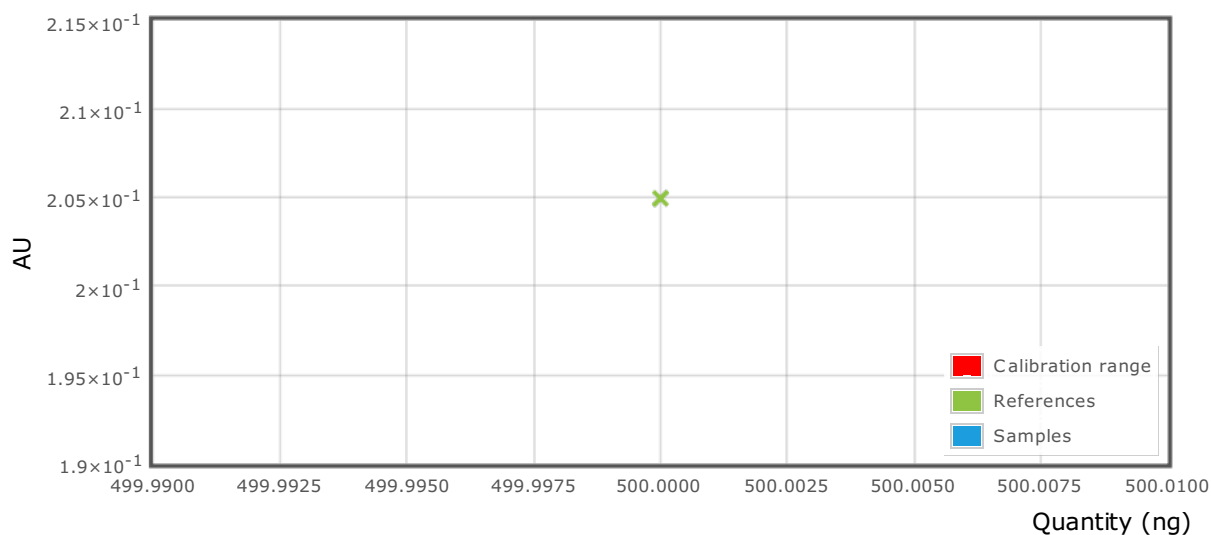
Peak #	Start		Max			End		Area		Manual peak	Substance Name
	Rf	H	Rf	H	%	Rf	H	A	%		
1	0.907	0.0000	0.912	0.0188	44.84	0.916	0.0000	0.00009	26.32	No	
2	0.976	0.0000	0.990	0.0232	55.16	0.996	0.0000	0.00026	73.68	No	

Calibration results:

Height calibration for substance 8-THC @ RT White:

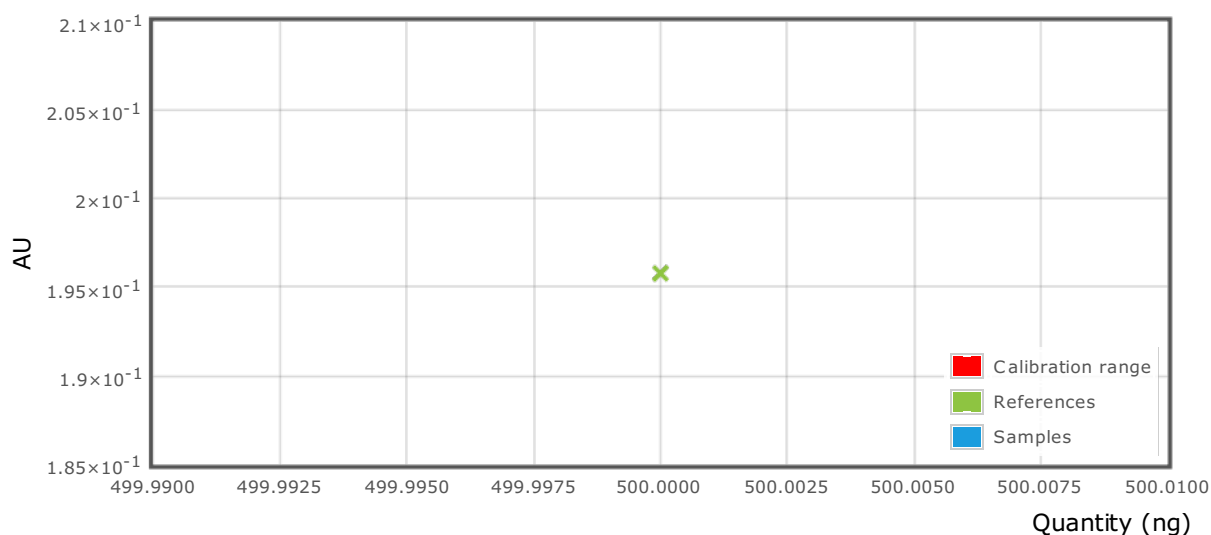
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Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance 9-THC @ RT White:

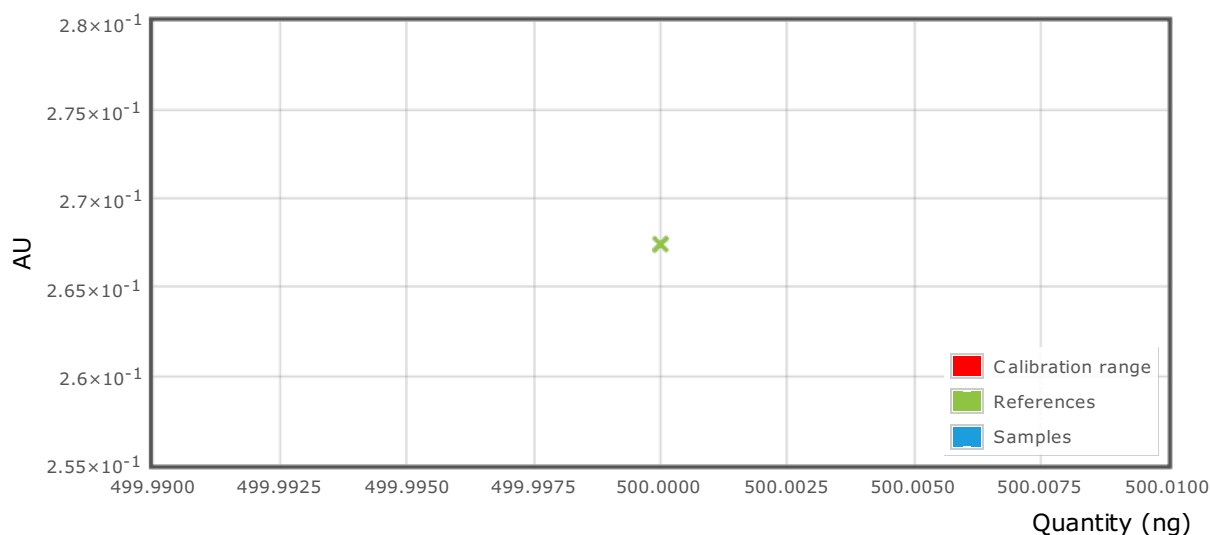


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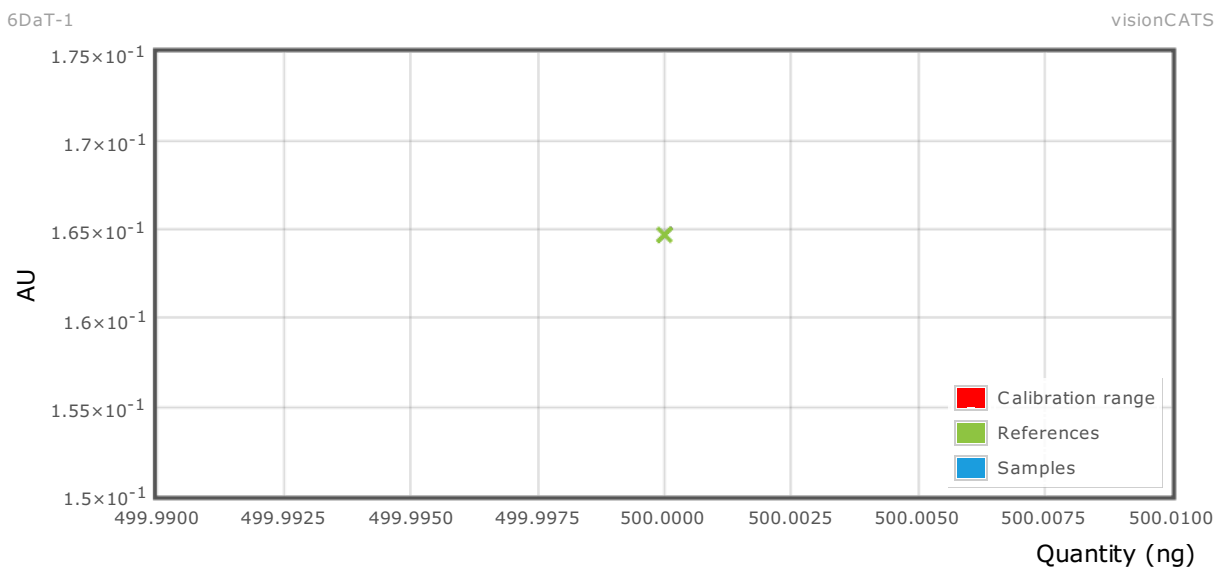
Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBC @ RT White:



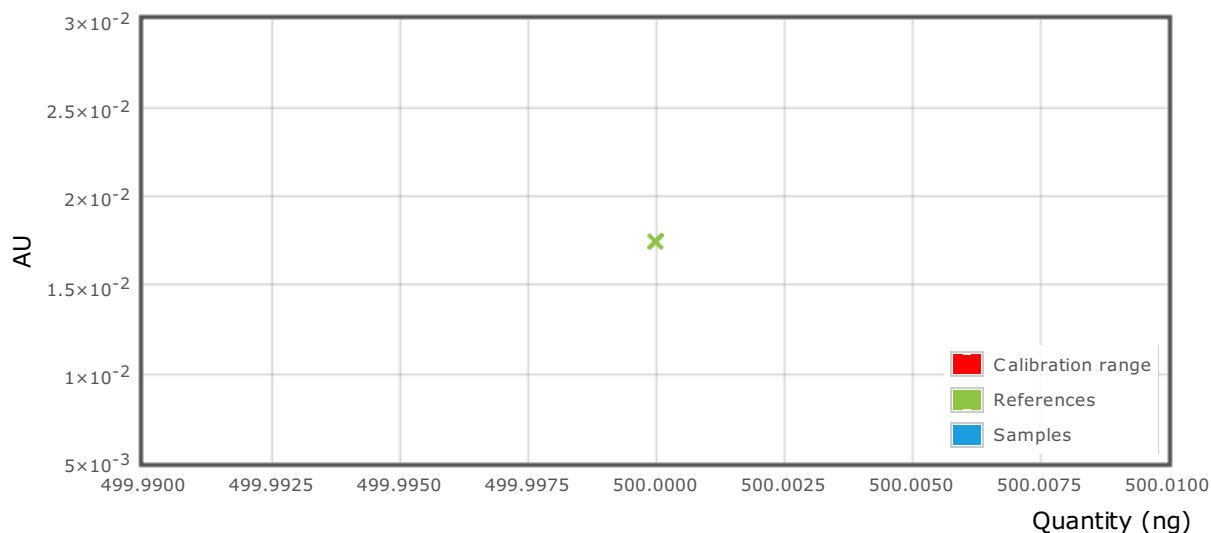
Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBD @ RT White:



Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBDA @ RT White:

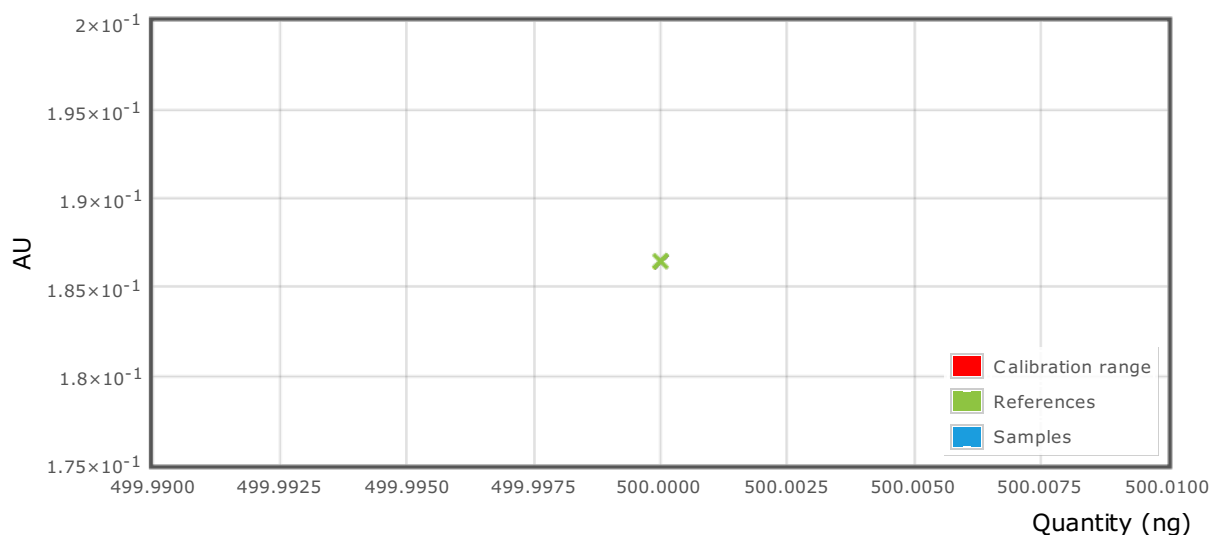


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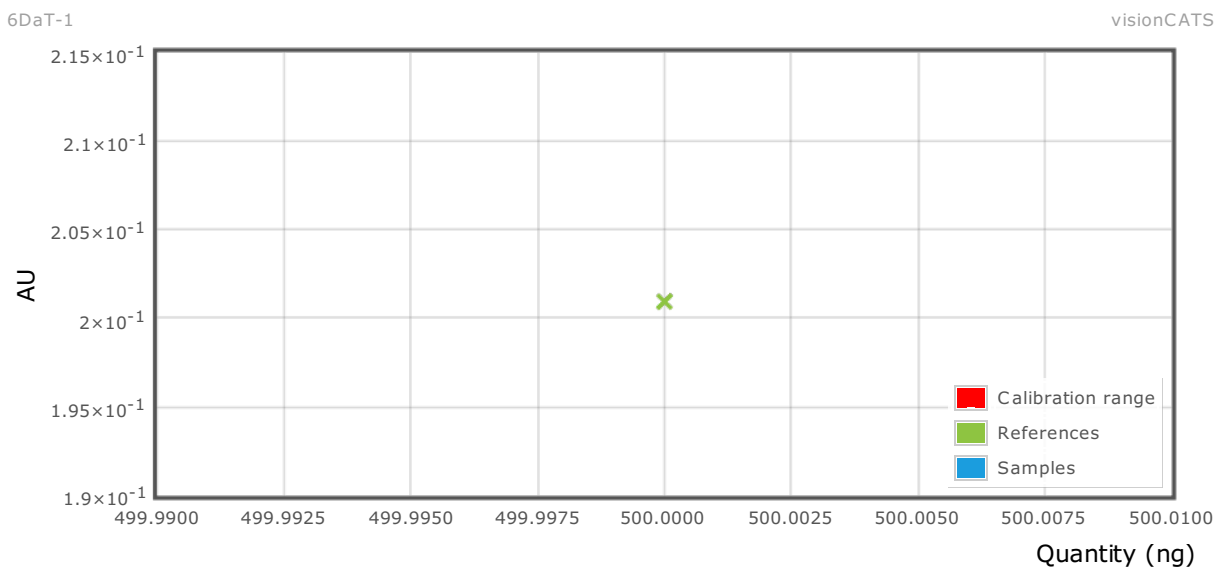
Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBDV @ RT White:



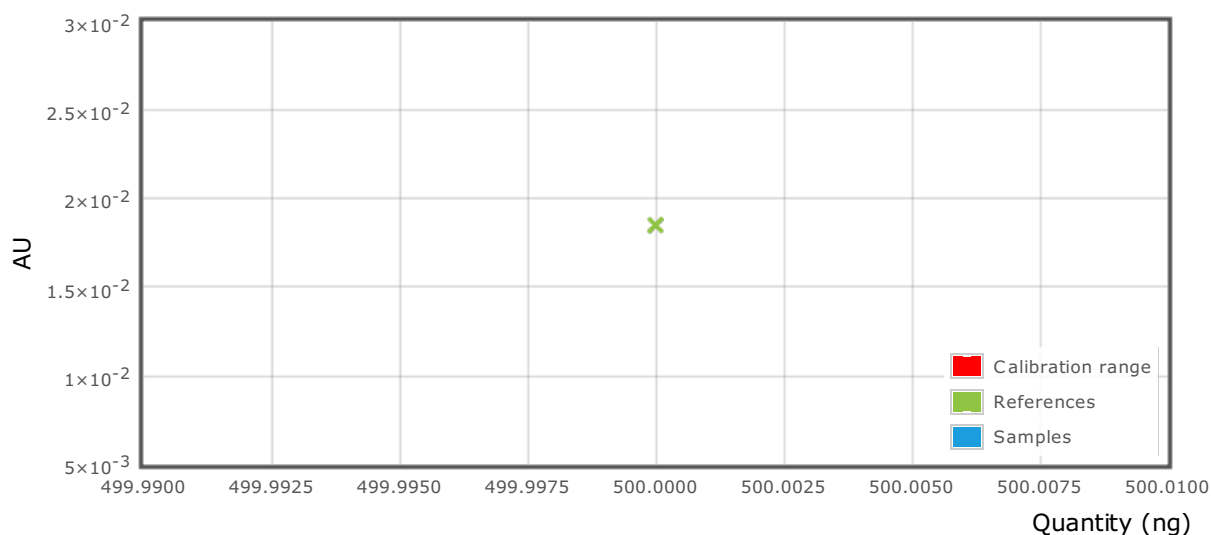
Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBG @ RT White:



Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBGA @ RT White:

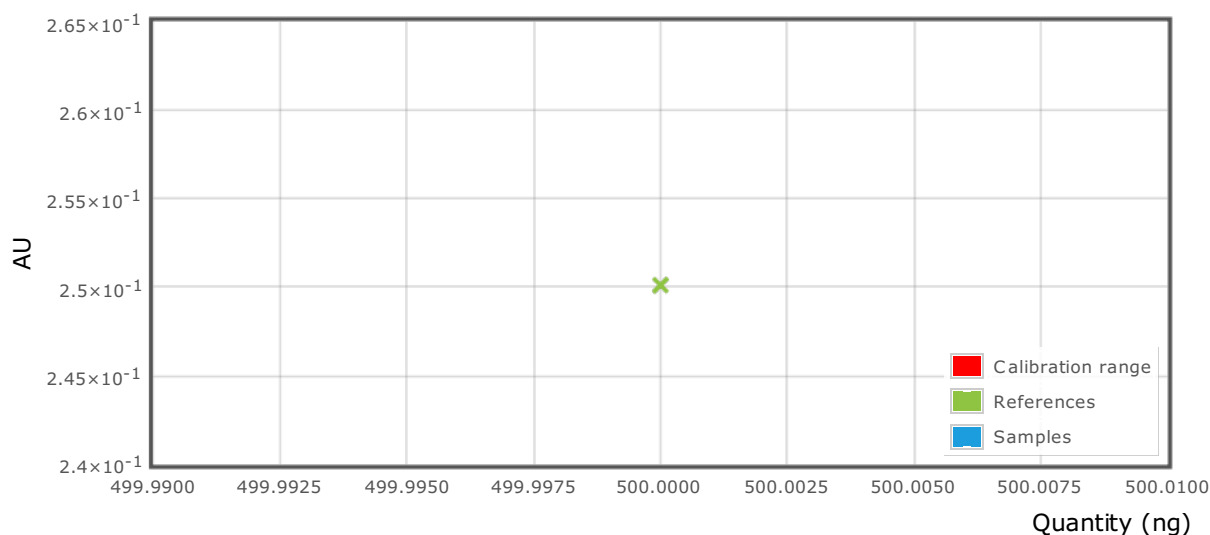


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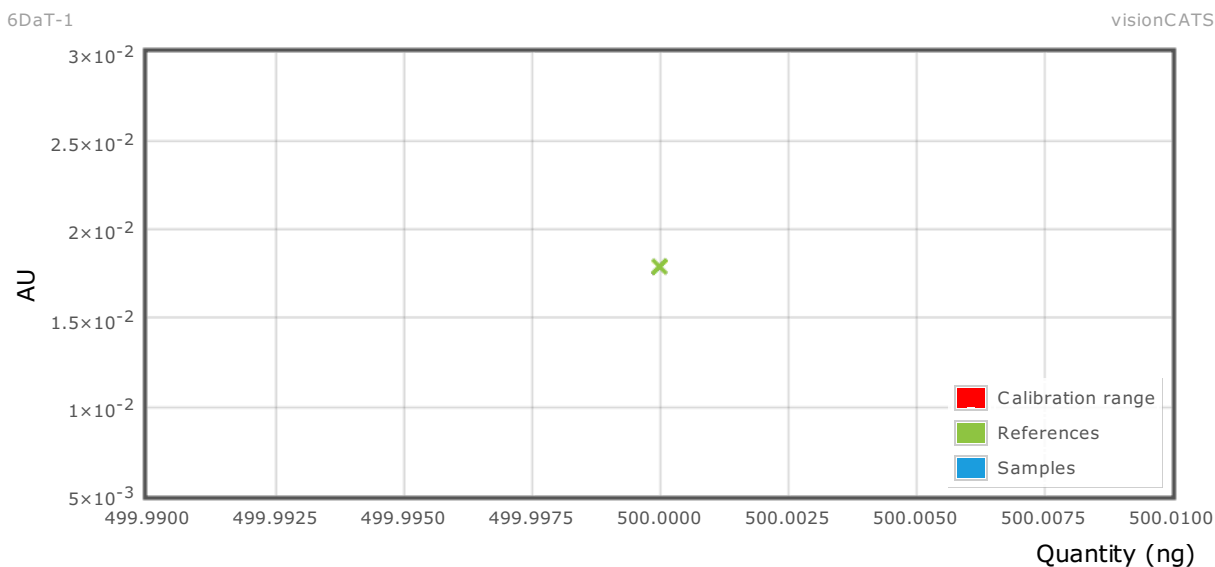
Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance CBN @ RT White:



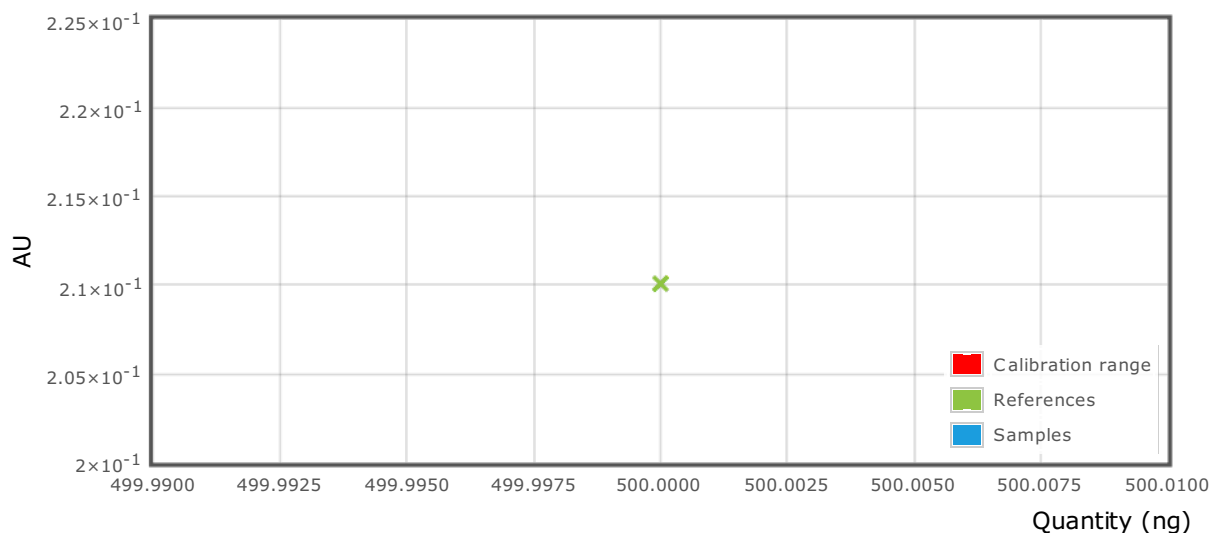
Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance THCA-A @ RT White:



Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Height calibration for substance THCV @ RT White:



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Regression mode	Linear-2
Range deviation	5.00 %
Related substances	Default
Number of references	1
Calibration function	$y=0x$
Coefficient of variation	CV 0.00 %
Correlation coefficient	n/a
	Unable to compute the results for this substance because there wasn't enough groups of references replicas (at least 1 for Linear-1, 2 for Linear2 and Mime-1 and 3 for Polynomial and MiMe-2)

Results:

Substance having no available results		
	CBD	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	CBDA	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	THCV	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	CBN	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	CBC	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	8-THC	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	CBG	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	9-THC	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	THCA-A	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	CBDV	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.
	CBGA	There wasn't any sample application available in the assignments for this substance. Please check that the peaks were correctly detected and assigned for this substance.

A track marked with  means: this result is outside the regression range given by the reference assignments, but is included in the results because it is in the allowed range deviation.

Analyst:

Reviewer: