

This is a special file, named RPTHEAD.TXT, in the directory of a method which allows you to customize the report header page. It can be used to identify the laboratory which uses the method.

This file is printed on the first page with the report styles:

Header+Short, GLP+Short, GLP+Detail, Short+Spec, Detail+Spec, Full

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      XXXX  XXX
    XX  XX  XX
  XX      XX      XXXXX  XXX XX
  XX      XX XXX  XX    X  XX X XX
  XX    X  XXX XX XXXXXXX  XX X XX
    XX XX  XX XX  XX      XX   XX
      XXXX  XXX  XXX  XXXXX  XXX  XXX
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XXXXXX  X      X      XX
XX  X  XX      XX
XX      XXXXX  XXXXX  XXXXX  XXX      XXXX  XX XXX
  XXXXX  XX      X  XX      XX  XX  XX  XX  XXX XX
    XX  XX      XXXXXX  XX      XX  XX  XX  XX  XX
X  XX  XX XX  X  XX  XX XX  XX  XX  XX  XX  XX
XXXXXX  XXX  XXXXX X  XXX  XXXX  XXXX  XX  XX
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                                     X
XX XXX  XXXXX  XX XXX  XXXX  XX XXX  XXXXX
  XXX XX  XX  X  XX  XX  XX  XX  XXX XX  XX
  XX      XXXXXXX  XX  XX  XX  XX  XX
  XX      XX      XXXXX  XX  XX  XX      XX XX
XXXX      XXXXX  XX      XXXX  XXXX      XXX
                XXXX
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XXX      XXX
  XX      XX
  XX      XXXXX  XXXXX  XX  XXXXX  XX XXX
  XX XXX  XX  X      X  XXXXX  XX  X  XXX XX
  XXX XX  XXXXXXX  XXXXXX  XX  XX  XXXXXXX  XX
  XX  XX  XX      X  XX  XX  XX  XX      XX
  XXX  XXX  XXXXX  XXXXX X  XXXX X  XXXXX  XXXX
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  X      XXX      X
  XX      XX      XX
XXXXX  XXXXX  XXX XX  XX XXX  XX  XXXXX  XXXXX  XXXXX
XX  XX  X  XX X XX  XX  XX  XX      X  XX  XX  X
XX  XXXXXXX  XX X XX  XX  XX  XX  XXXXXXX  XX  XXXXXXX
XX XX  XX      XX  XX  XXXXX  XX  X  XX  XX  XX  XX
  XXX  XXXXX  XXX  XXX  XX      XXXX  XXXXX X  XXX  XXXXX
                XXXX
```

```
=====
Acq. Operator   : user                               Seq. Line :    3
Sample Operator : user
Acq. Instrument : SFC LCMS                           Location  :   D2F-E2
Injection Date  : 11/08/2023 02:24:41                 Inj       :    1
                                                    Inj Volume : 0.200 µl
Different Inj Volume from Sample Entry! Actual Inj Volume : 2.000 µl
Acq. Method     : D:\Data\2023\Yunfei_R0AR\2023-08-10_MBAAd_Calib_2uL 2023-08-11 02-05-23\COL1
                  _5NH4FA_MECN_5T095_1MIN_100-600MS_POS.M
Last changed    : 24/07/2023 15:04:14 by administrator
Analysis Method : D:\Data\2023\Yunfei_R0AR\2023-08-10 baseline corrected\2023-08-11 02-05-23_
                  MBAAd_Calib_2uL\COL1_5NH4FA_MECN_5T095_1MIN_100-600MS_POS.M (Sequence Method
                  )
Last changed    : 24/07/2023 15:04:14 by administrator
Additional Info : Peak(s) manually integrated
=====
```

Module	Type	Firmware rev.	Serial number
Column Comp.	G7116A	D.07.23 [0009]	DEAED08985
Make Up Pump 2	G7110B	D.07.23 [0009]	DEAEH00761
Valve 3	G1170A	D.07.23 [0009]	DEBAD03734
Multisampler 4	G4767A	D.07.24 [0001]	DEAFD00218
LC Pump 5	PumpValveCluster		
Pump 5	G7111B	D.07.24 [0001]	DEAEW03495
SFC Binary Pump 6	G4782A	D.07.23 [0009]	DEAGN00153
DAD 7	G7115A	D.07.23 [0009]	DEAC605436
SFC 8	G4301A	A.03.09 [0005]	SG18067002
ELSD 9	G4260B		GB23230008
Agilent G6125B MSD	G6125B	3.02.50	SG1823N002

Software Revision: Rev. C.01.09 [161] Copyright © Agilent Technologies

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Column(s)

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Column Description : Raptor C18

Serial # : 288

Product# : 9304A52 Batch# : 220519B

Diameter : 2.1 mm Length : 50.0 mm

Particle size : 2.7 µm Void volume : 0.10 ml

Injections : 433

Maximum Pressure : 600.0 bar Maximum pH : 8.0

Minimum pH : 2.0

Maximum Temperature: 60.0 °C

Comment : New 2023-08-03

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Instrument Conditions	:	At Start	At Stop
Column Temp. (left)	:	40.0	40.0 °C
Column Temp. (right)	:	32.4	32.4 °C
Pressure	:	0.0	0.0 bar
Flow	:	0.000	0.000 ml/min

Detector Lamp Burn Times:	Current On-Time	Accumulated On-Time
DAD 1, UV Lamp	: 0.30	849.3 h
DAD 1, Visible Lamp	: 0.00	331.2 h

Solvent Description	:
PMP1, Solvent A	:
PMP2, Solvent A	:
PMP2, Solvent A	:
PMP2, Solvent B	:
PMP2, Solvent B	:

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MSD parameters

Tune file name	:	C:\Users\Public\Documents\ChemStation\1\MStune\6125BTUN\atunes.tun (Wed Aug 2 16:39:21 2023)
Ionization mode	:	ES-API

MSD Instrument Conditions	:	At Start	At Stop
Quad Temp	:	100	100 C
Gas Temp	:	350	350 C
RoughVac	:	2	2 Torr
HighVac	:	5.2E-009	5.2E-009 Torr
CapCur	:	5	655 nA
ChamCur	:	1.1E-001	1.2E-001 µA
DryingGas	:	12	12 l/min
Neb Pres	:	35	35 psi g
Turbo1Spd	:	100	100 %
Turbo1Pwr	:	127	127 W
RF Drive	:	1	15 %
Qd TpDrv	:	16	16 %
Gas TpDrv	:	35	35 %
Neb PrDrv	:	50	49 %
Gas FI Drv	:	62	61 %

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MSD tuning (calibration) parameters

Ionization polarity	:	Positive
Skim1	:	30 V
Skim2	:	
Ion Energy	:	5.0 V
Lens1	:	3.2 V
Lens2	:	
Iris	:	-400 V
HED	:	10000 V
Width Gain	:	-186
Width Offset	:	Variable
Mass	:	Value

118.08	:	-24
622.03	:	-29

922.01 : -22
1521.97 : -24

Mass Gain : -12.80
Mass Offset : Variabl e
Mass : Val ue

118.08 : 0.752
622.03 : 0.846
922.01 : 0.836
1521.97 : 0.752

Quad DC : 0.00 V
Octopole Peak : 650 V
Octopole Knee :
Lens2DC : Variabl e
Mass : Val ue

50.00 : 0.5
100.00 : 1.0
350.00 : 2.0
1000.00 : 4.0
2000.00 : 6.0

L2RFEn : 1
L2RFPh : 162
L2RF Amp : Variabl e
Mass : Val ue

118.08 : 51
622.03 : 95
922.01 : 105
1521.97 : 145

Mass Filter : Gaussi an
Time Filter : Gaussi an
Time Filter Width : 0.030

Ionization polari ty : Negati ve
Skim1 : 35 V
Skim2 :
Ion Energy : 5.0 V
Lens1 : -3.4 V
Lens2 :
Iris : 400 V
HED : 10000 V
Width Gain : -187
Width Offset : Variabl e
Mass : Val ue

112.99 : -32
601.98 : -76
1033.99 : -74
1633.95 : -32

Mass Gain : -12.85
Mass Offset : Variabl e

Mass	:	Val ue
112.99	:	0.774
601.98	:	0.868
1033.99	:	0.840
1633.95	:	0.774

Quad DC : 0.00 V
Octopole Peak : 650 V
Octopole Knee :
Lens2DC : Vari abl e

Mass	:	Val ue
50.00	:	0.5
100.00	:	1.0
350.00	:	2.0
1000.00	:	4.0
2000.00	:	6.0

L2RFEn : 1
L2RFPh : 162
L2RFamp : Vari abl e

Mass	:	Val ue
112.99	:	70
601.98	:	110
1033.99	:	130
1633.95	:	150

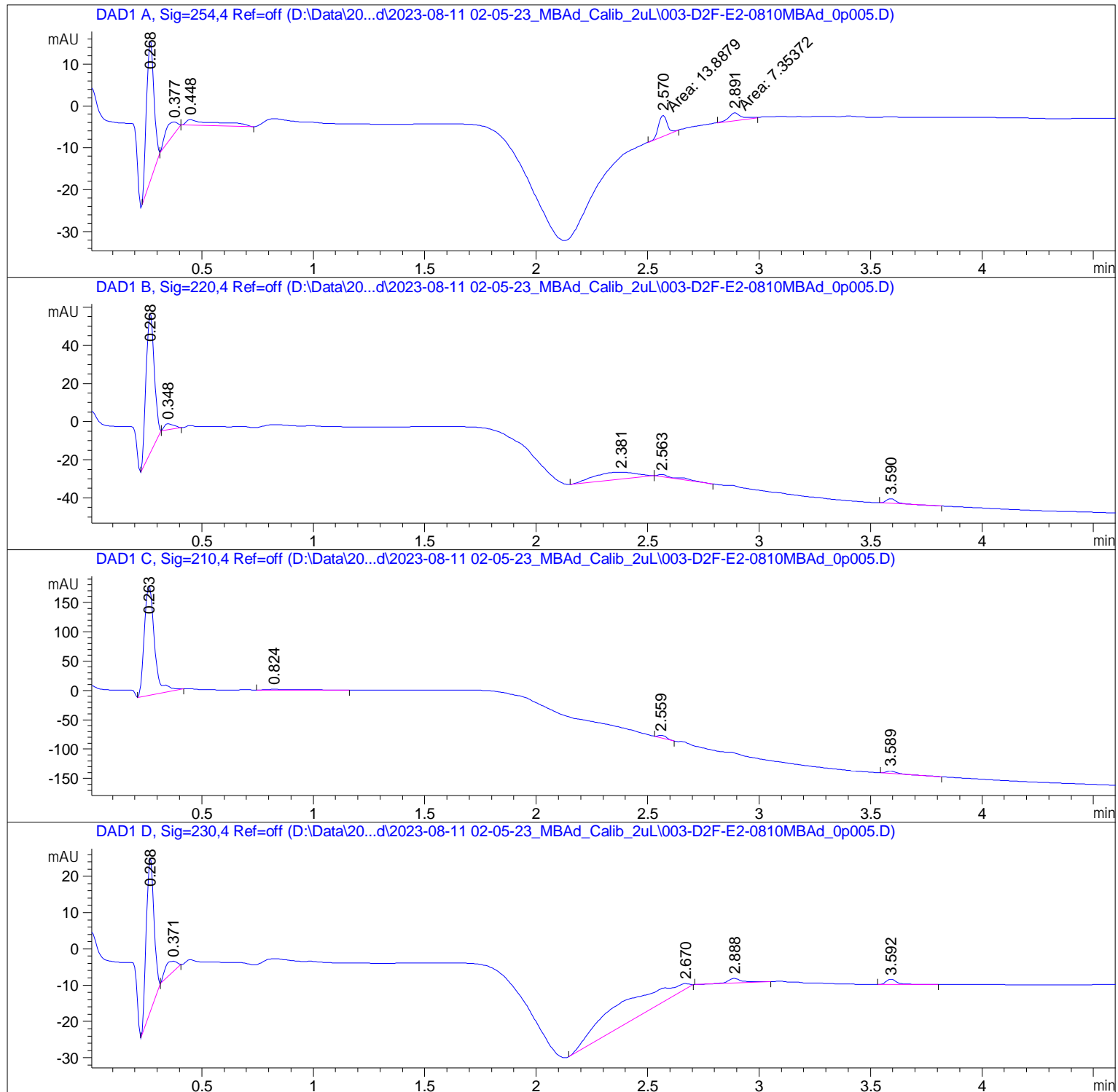
Mass Filter : Gaussi an
Time Filter : Gaussi an
Time Filter Width : 0.030

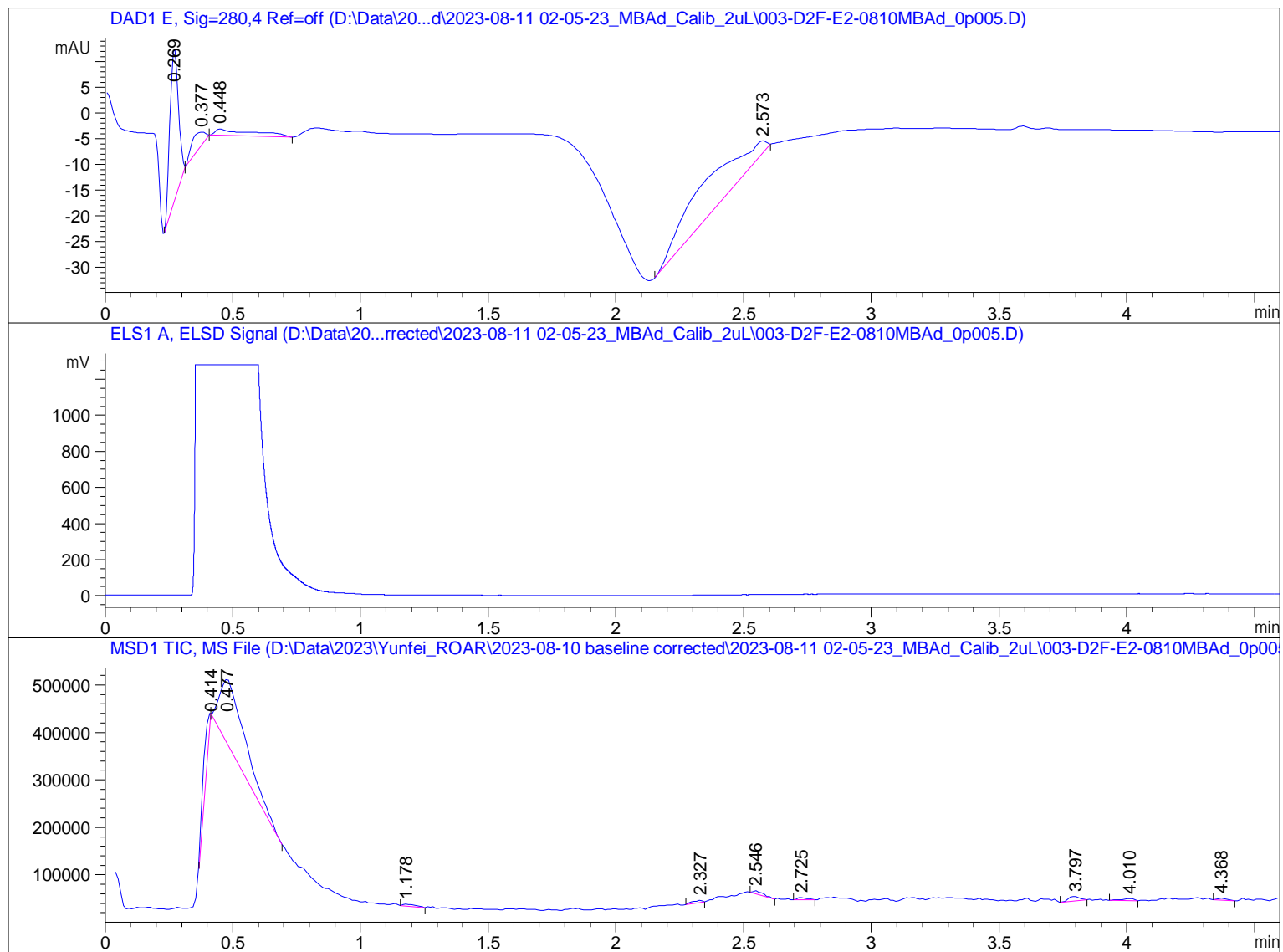
Run Logbook

11 Aug 23 12:14 PM
Logbook File: D:\Data\20...ted\2023-08-11 02-05-23_MBAAd_Cal i b_2uL\003-D2F-E2-0810MBAAd_Op005.D\RUN.LOG

Module	# Event Message	Date Time
Method	Method started: line# 3 at location 'D2F-E2>' ' inj# 1	11/08/2023 02:23:54
CP Macro	PreRun macro: 'LAMPALL ON'	11/08/2023 02:23:55
G4260B	G4260B: ELSD - Autozero	11/08/2023 02:23:55
G4260B	G4260B: ELSD - Al ready swit ched on	11/08/2023 02:23:56
Method	Instrument running sample from location D2F->E2	11/08/2023 02:23:56
G7115A	G7115A: DEAC605436 - Detector: Prepare	11/08/2023 02:24:07
G7115A	G7115A: DEAC605436 - Detector: Idle	11/08/2023 02:24:24
G4767A	G4767A: DEAFD00218 - Draw command finished	11/08/2023 02:24:31
G4767A	G4767A: DEAFD00218 - Sampler wash is active	11/08/2023 02:24:32
G4767A	G4767A: DEAFD00218 - Sampler wash is idle	11/08/2023 02:24:38
G4767A	G4767A: DEAFD00218 - Sample preparation time: >15 sec	11/08/2023 02:24:39
PumpVal ve	G7111B: DEAEW03495 - Run	11/08/2023 02:24:40

PumpValve	G7111B: DEAEW03495 - Postrun	11/08/2023 02: 29: 17
PumpValve	G1170A: DEBAD03715 - Postrun	11/08/2023 02: 29: 17
G4767A	G4767A: DEAFD00218 - Postrun	11/08/2023 02: 29: 17
G7116A	G7116A: DEAED08985 - Postrun	11/08/2023 02: 29: 18
Method	Saving Method COL1_5NH4FA_MECN_5T095_1MIN_10> 0-600MS_POS.M	11/08/2023 02: 31: 28
Method	Instrument run completed	11/08/2023 02: 31: 29
CP Macro	Analyzing rawdata 003-D2F-E2-0810MBAAd_Op005. > D	11/08/2023 02: 31: 30
Method	Saving Method DA.M	11/08/2023 02: 31: 31
Method	Method completed	11/08/2023 02: 31: 43





Area Percent Report

Sorted By : Signal
Multiplier : 1.0000
Dilution : 1.0000
Do not use Multiplier & Dilution Factor with ISTDs

Signal 1: DAD1 A, Sig=254, 4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	0.268	BB	0.0368	78.22271	34.00939	61.6994
2	0.377	BB	0.0748	13.24266	2.81691	10.4454
3	0.448	BB	0.1287	14.07328	1.38339	11.1005
4	2.570	MM	0.0454	13.88794	5.10259	10.9543
5	2.891	MM	0.0646	7.35372	1.89580	5.8004

Totals : 126.78032 45.20808

Signal 2: DAD1 B, Sig=220,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	0.268	BB	0.0396	187.85695	73.99009	70.7172
2	0.348	BB	0.0438	9.14417	3.15230	3.4422
3	2.381	BB	0.2251	51.88420	3.66611	19.5314
4	2.563	BB	0.0943	8.61777	1.22114	3.2441
5	3.590	BBA	0.0510	8.14238	2.43649	3.0651
Totals :				265.64547	84.46613	

Signal 3: DAD1 C, Sig=210,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	0.263	BB	0.0491	597.47943	188.04227	93.8287
2	0.824	BB	0.1659	13.43975	1.15371	2.1106
3	2.559	BB	0.0438	11.71303	4.29497	1.8394
4	3.589	BBA	0.0492	14.14438	4.43715	2.2212
Totals :				636.77659	197.92809	

Signal 4: DAD1 D, Sig=230,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	0.268	BB	0.0376	101.32568	42.84363	39.5172
2	0.371	BB	0.0689	12.16399	2.90158	4.7440
3	2.670	BB	1.0117	131.19020	1.53097	51.1645
4	2.888	BB	0.0694	6.42740	1.35445	2.5067
5	3.592	BB	0.0526	5.30147	1.52273	2.0676
Totals :				256.40874	50.15336	

Signal 5: DAD1 E, Sig=280,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	0.269	BB	0.0370	69.64007	30.05303	31.7610
2	0.377	BB	0.0737	11.82970	2.56950	5.3952
3	0.448	BB	0.1311	13.46113	1.29679	6.1393
4	2.573	BB	0.6070	124.33208	2.43771	56.7045
Totals :				219.26297	36.35703	

Signal 6: ELS1 A, ELSD Signal

Signal 7: MSD1 TIC, MS File

Peak #	RetTime [min]	Type	Width [min]	Area	Height	Area %
1	0.414	BB	0.3435	1.56742e5	5501.26367	12.7718
2	0.477	BB	0.0958	9.54299e5	1.33867e5	77.7590
3	1.178	BB	0.0482	1.45573e4	4050.61646	1.1862
4	2.327	BB	0.0515	1.31342e4	4719.06982	1.0702
5	2.546	BB	0.0506	1.92826e4	6349.19336	1.5712
6	2.725	BB	0.0423	1.39132e4	4953.15820	1.1337
7	3.797	BB	0.0464	2.99019e4	9448.02734	2.4365
8	4.010	BB	0.0522	1.40008e4	4540.66162	1.1408
9	4.368	BB	0.0425	1.14212e4	4037.42236	0.9306

Totals : 1.22725e6 1.77467e5

Summed Peaks Report

Signal 1: DAD1 A, Sig=254,4 Ref=off
Empty table.

Signal 2: DAD1 B, Sig=220,4 Ref=off
Empty table.

Signal 3: DAD1 C, Sig=210,4 Ref=off
Empty table.

Signal 4: DAD1 D, Sig=230,4 Ref=off
Empty table.

Signal 5: DAD1 E, Sig=280,4 Ref=off
Empty table.

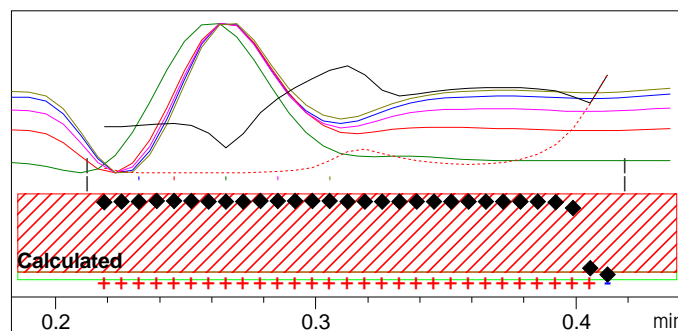
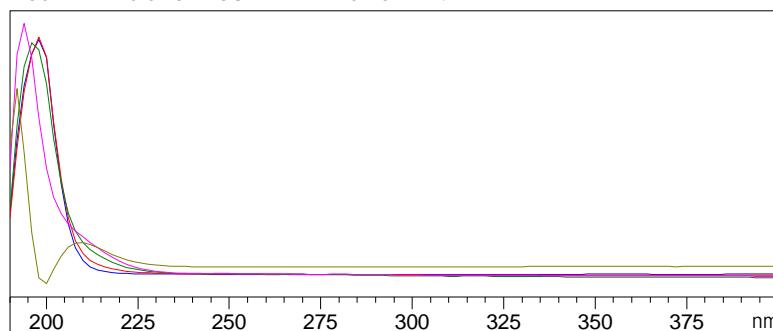
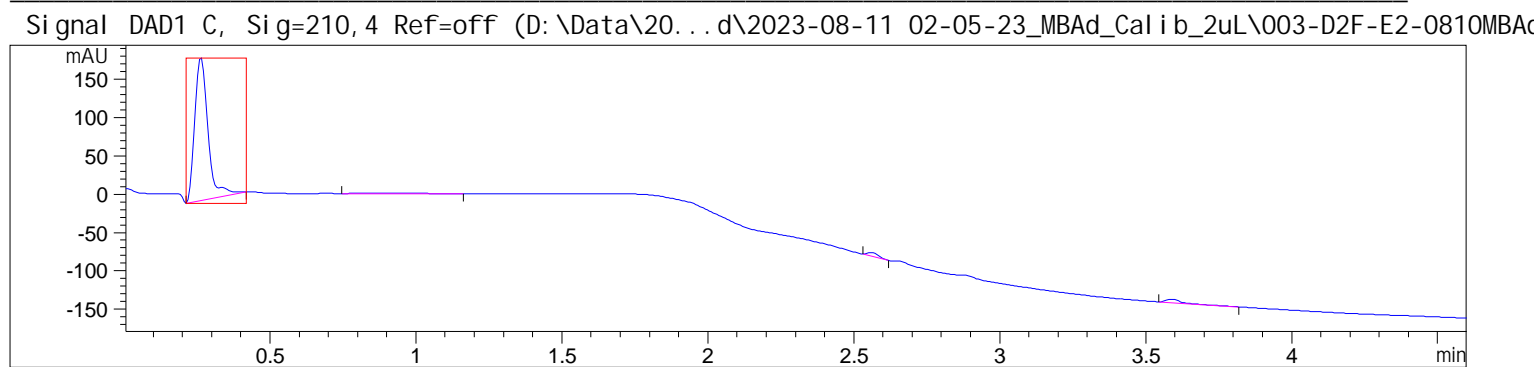
Signal 6: ELS1 A, ELSD Signal
Empty table.

Signal 7: MSD1 TIC, MS File
Empty table.

Final Summed Peaks Report

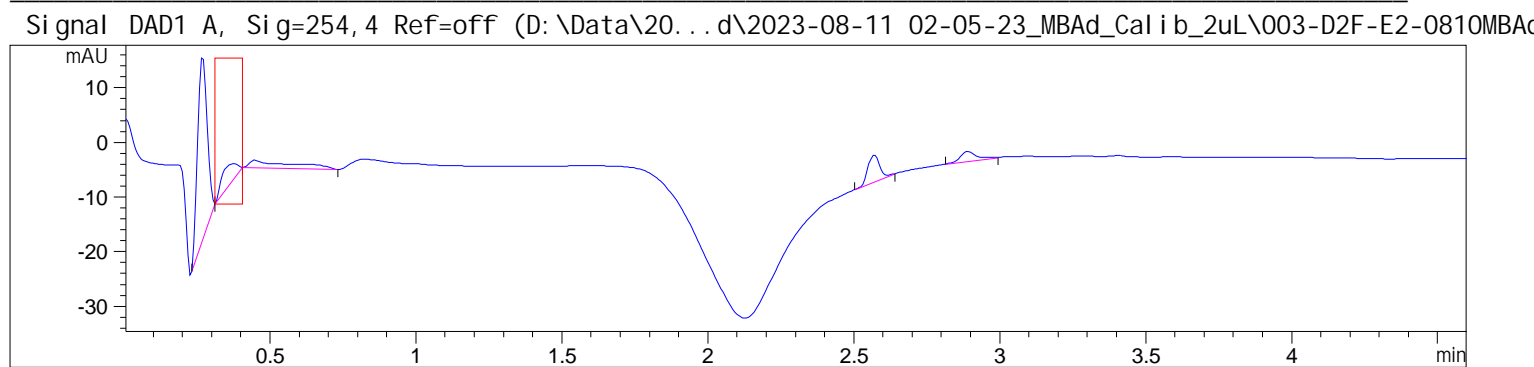
Signal 1: DAD1 A, Sig=254,4 Ref=off

Signal 2: DAD1 B, Sig=220, 4 Ref=off
Signal 3: DAD1 C, Sig=210, 4 Ref=off
Signal 4: DAD1 D, Sig=230, 4 Ref=off
Signal 5: DAD1 E, Sig=280, 4 Ref=off
Signal 6: ELS1 A, ELSD Signal
Signal 7: MSD1 TIC, MS File

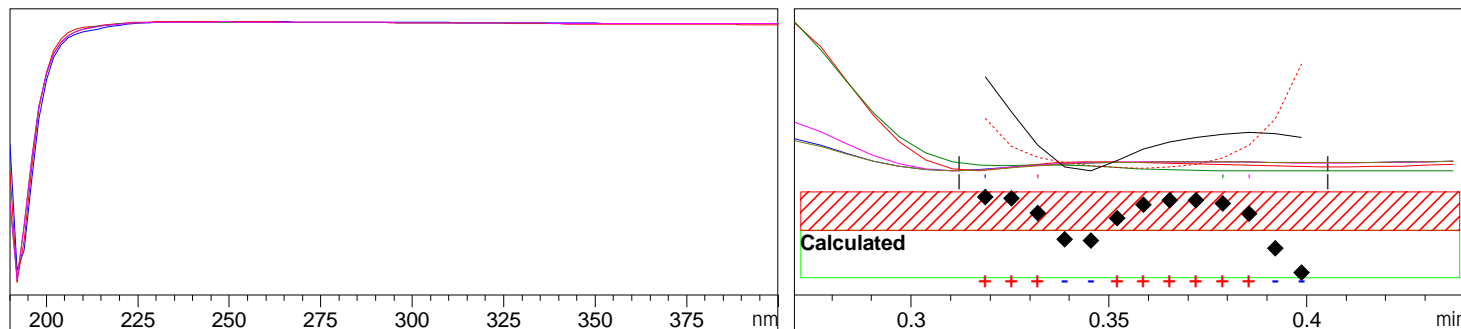


-> The purity factor exceeds the calculated threshold limit. <-

Purity factor : 815.531 (29 of 30 spectra exceed the calculated threshold limit.)
Threshold : 996.663 (Calculated with 29 of 30 spectra)
Reference : Peak start and end spectra (integrated) (0.212 / 0.419)
Spectra : 5 (Selection automatic, 5)
Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)
Warning : Spectral absorbances > 1000 mAU (see help for more information)

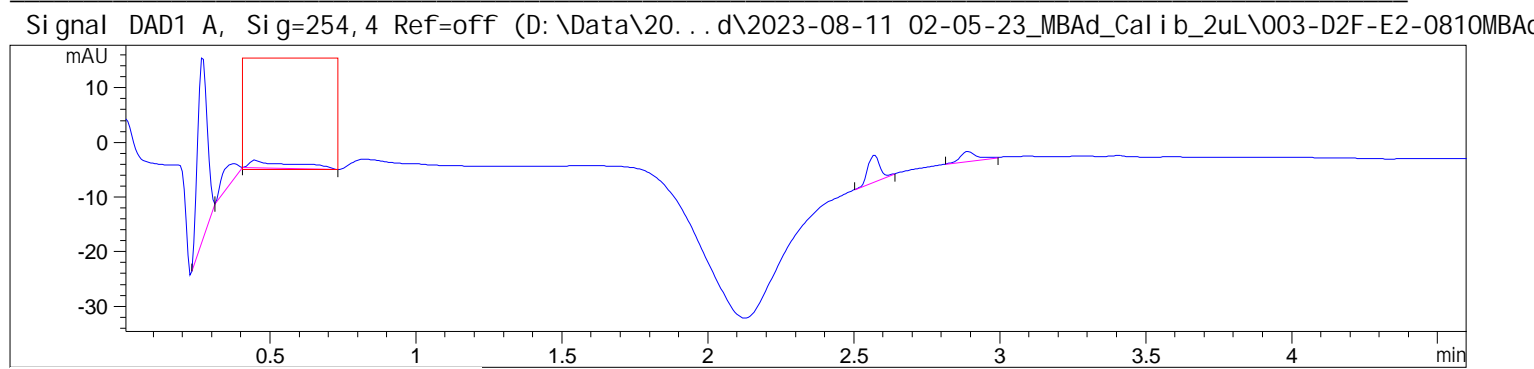


Peak : 2 at 0.377 min Name : ?

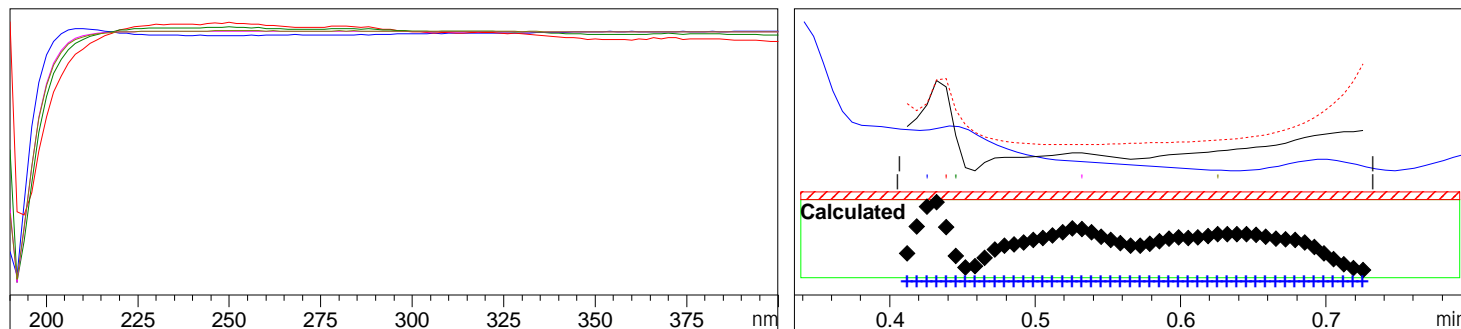


-> The purity factor exceeds the calculated threshold limit. <-

Purity factor : 996.161 (9 of 13 spectra exceed the calculated threshold limit.)
Threshold : 998.551 (Calculated with 9 of 13 spectra)
Reference : Peak start and end spectra (integrated) (0.312 / 0.405)
Spectra : 4 (Selection automatic, 5)
Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)



Peak : 4 at 0.448 min Name : ?



-> The purity factor is within the calculated threshold limit. <-

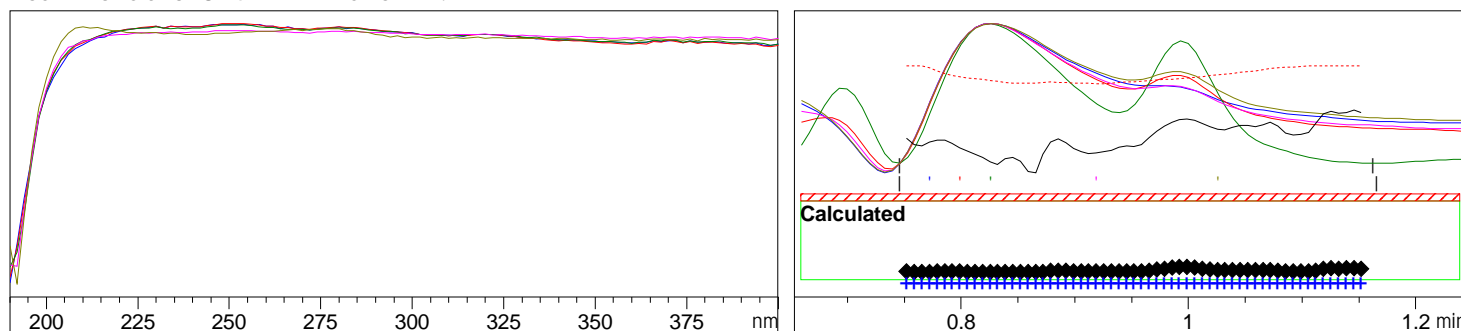
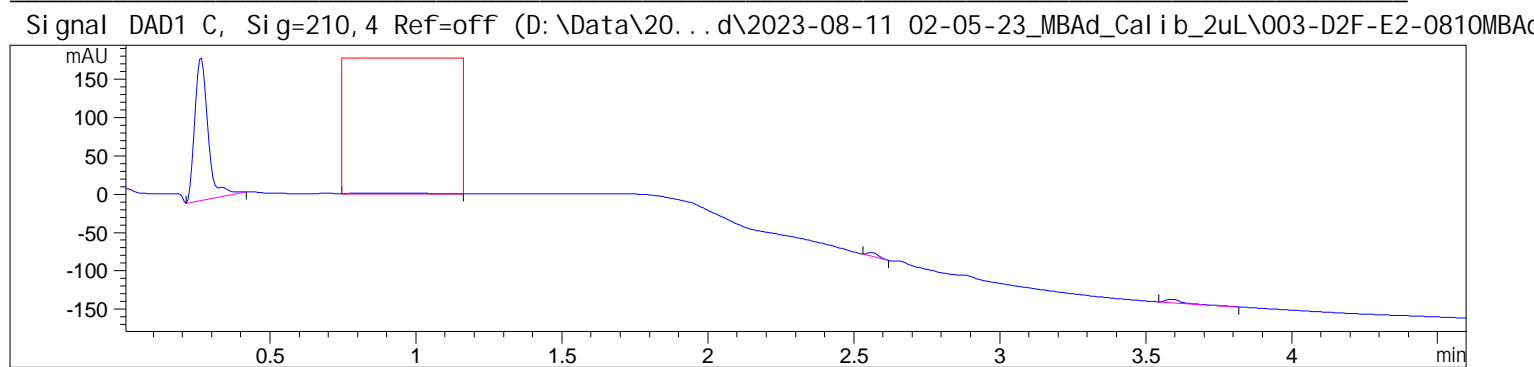
Purity factor : 978.348 (48 of 48 spectra are within the calculated threshold limit.)

Threshold : 933.456 (Calculated with 48 of 48 spectra)

Reference : Peak start and end spectra (integrated) (0.405 / 0.732)

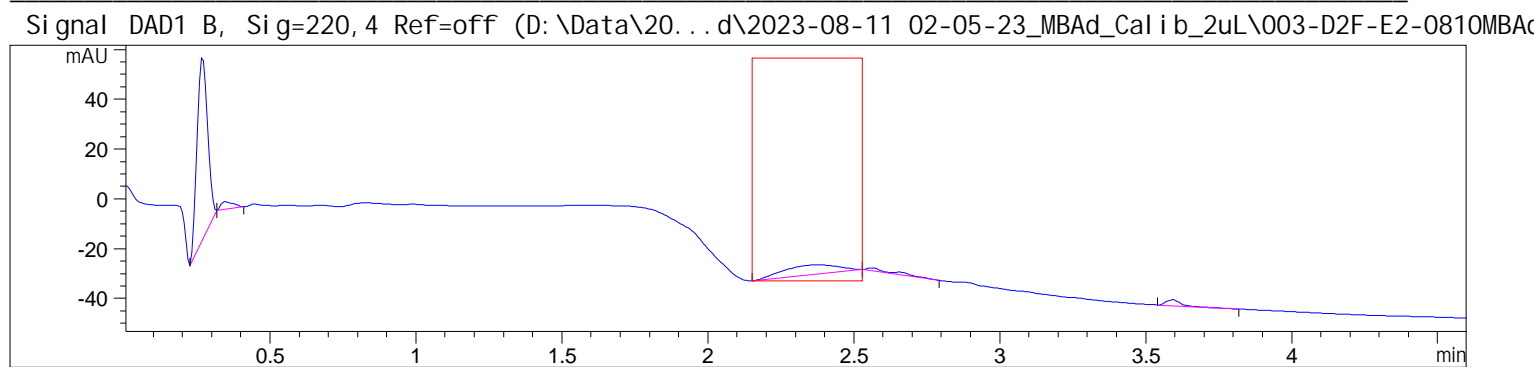
Spectra : 5 (Selection automatic, 5)

Noise Threshold : 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)

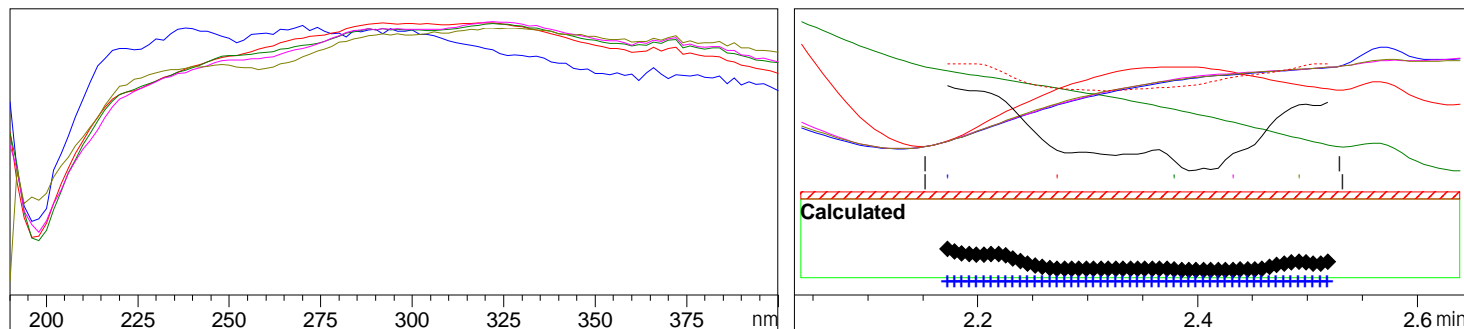


-> The purity factor is within the calculated threshold limit. <-

Purity factor : 983.123 (61 of 61 spectra are within the calculated threshold limit.)
Threshold : 396.799 (Calculated with 61 of 61 spectra)
Reference : Peak start and end spectra (integrated) (0.745 / 1.165)
Spectra : 5 (Selection automatic, 5)
Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)



Peak : 8 at 2.381 min Name : ?



-> The purity factor is within the calculated threshold limit. <-

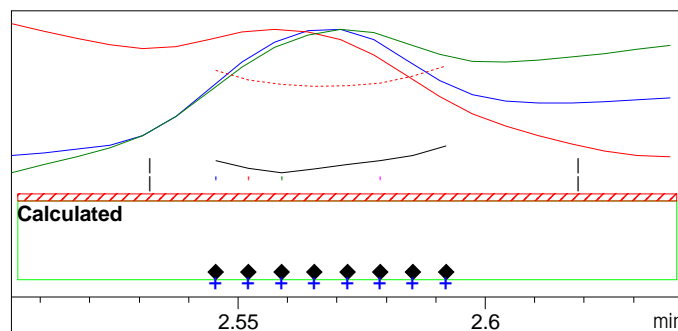
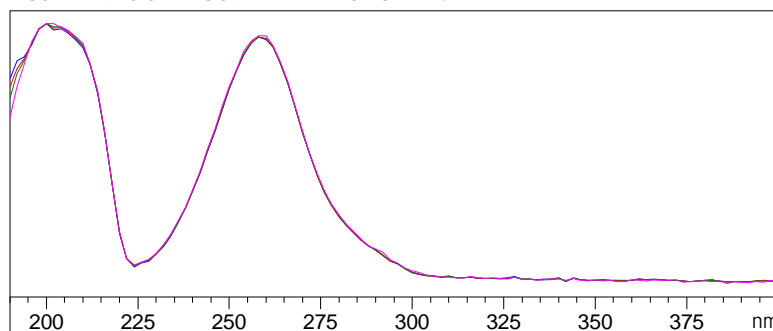
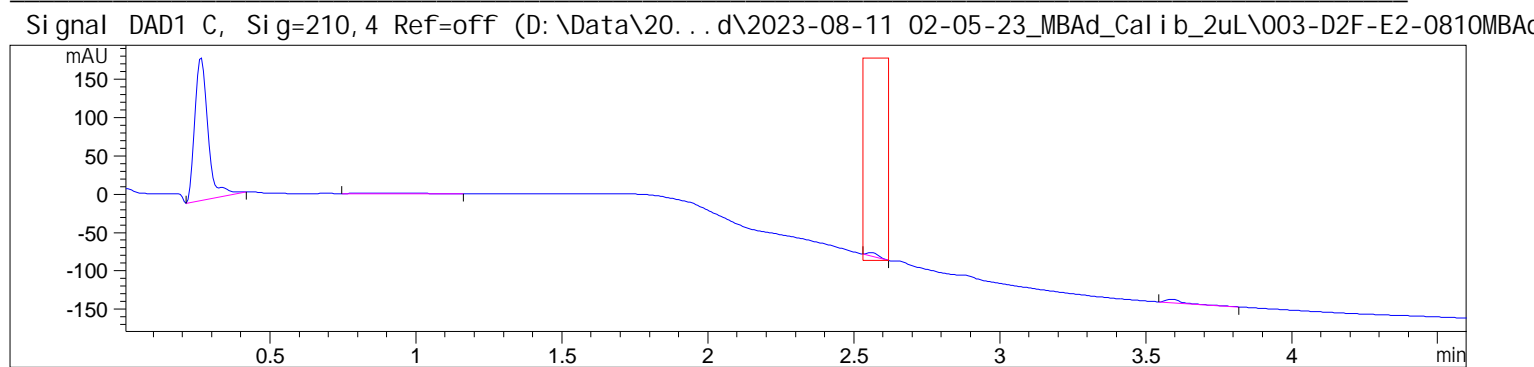
Purity factor : 939.860 (53 of 53 spectra are within the calculated threshold limit.)

Threshold : 485.274 (Calculated with 53 of 53 spectra)

Reference : Peak start and end spectra (integrated) (2.152 / 2.532)

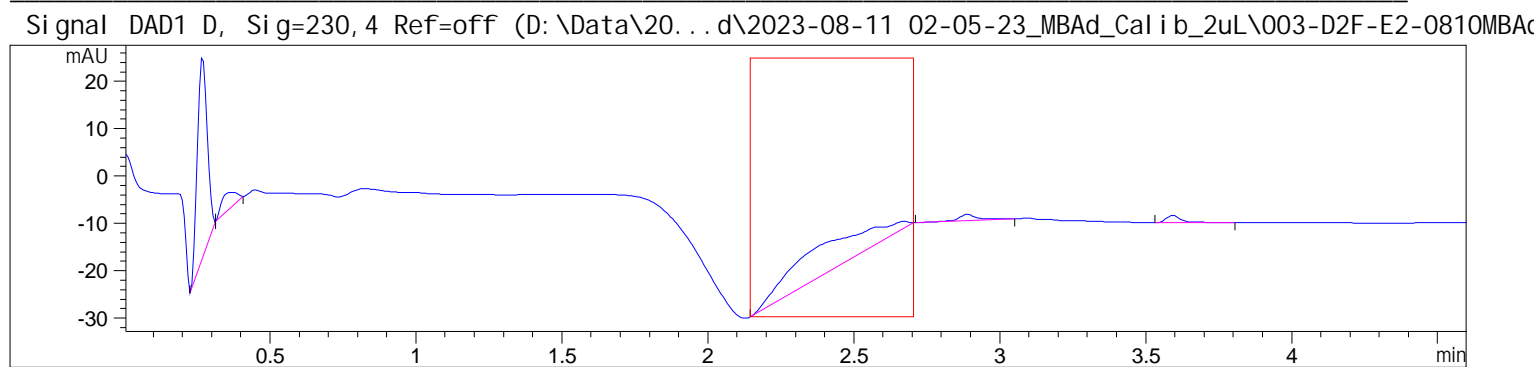
Spectra : 5 (Selection automatic, 5)

Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)

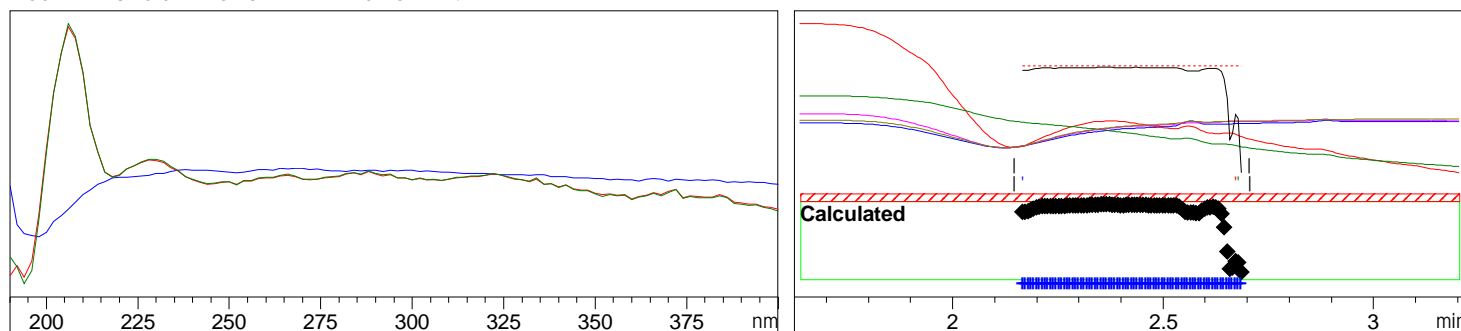


-> The purity factor is within the calculated threshold limit. <-

Purity factor : 999.211 (8 of 8 spectra are within the calculated threshold limit.)
Threshold : 561.669 (Calculated with 8 of 8 spectra)
Reference : Peak start and end spectra (integrated) (2.532 / 2.619)
Spectra : 4 (Selection automatic, 5)
Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)



Peak : 10 at 2.670 min Name : ?



-> The purity factor is within the calculated threshold limit. <-

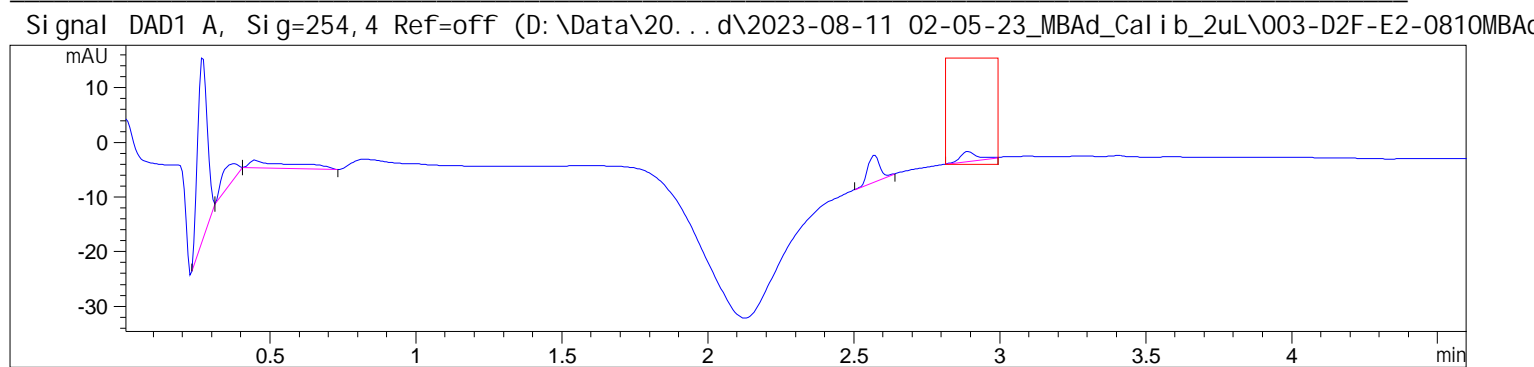
Purity factor : 138.858 (79 of 79 spectra are within the calculated threshold limit.)

Threshold : 0.000 (Calculated with 79 of 79 spectra)

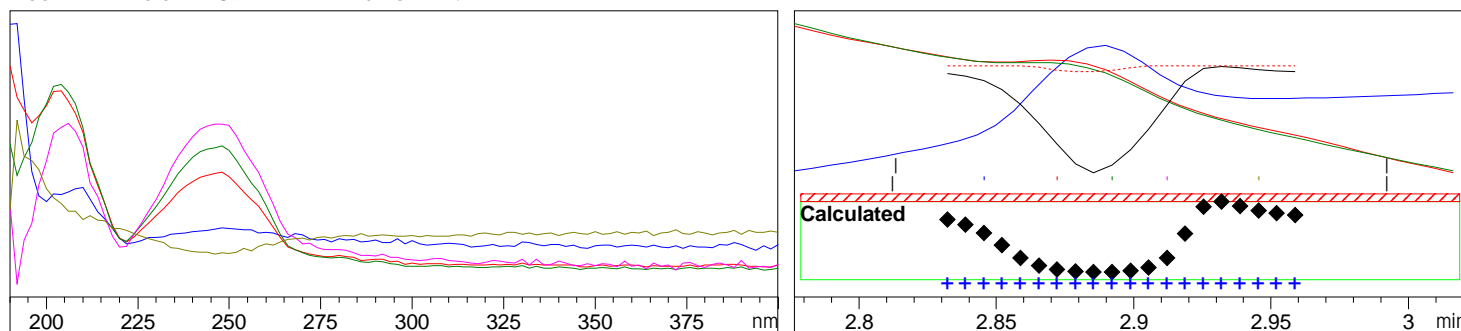
Reference : Peak start and end spectra (integrated) (2.145 / 2.705)

Spectra : 3 (Selection automatic, 5)

Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)

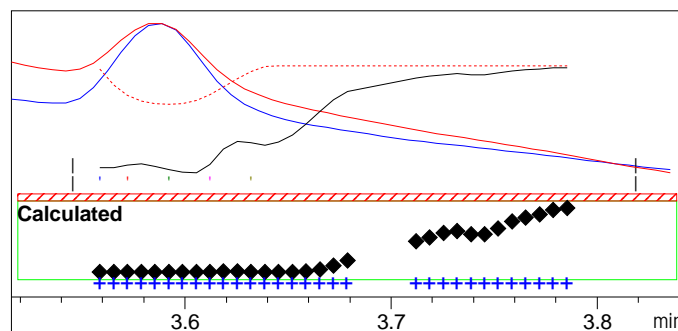
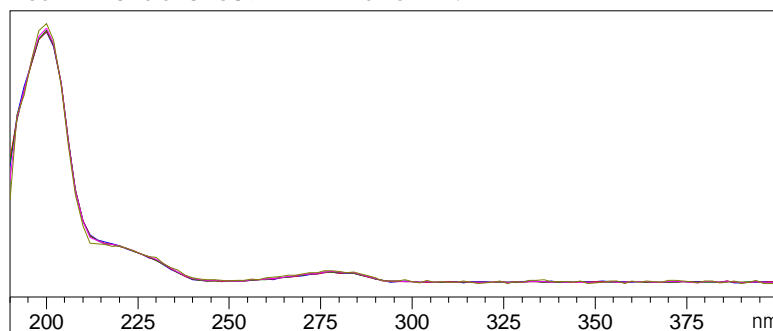
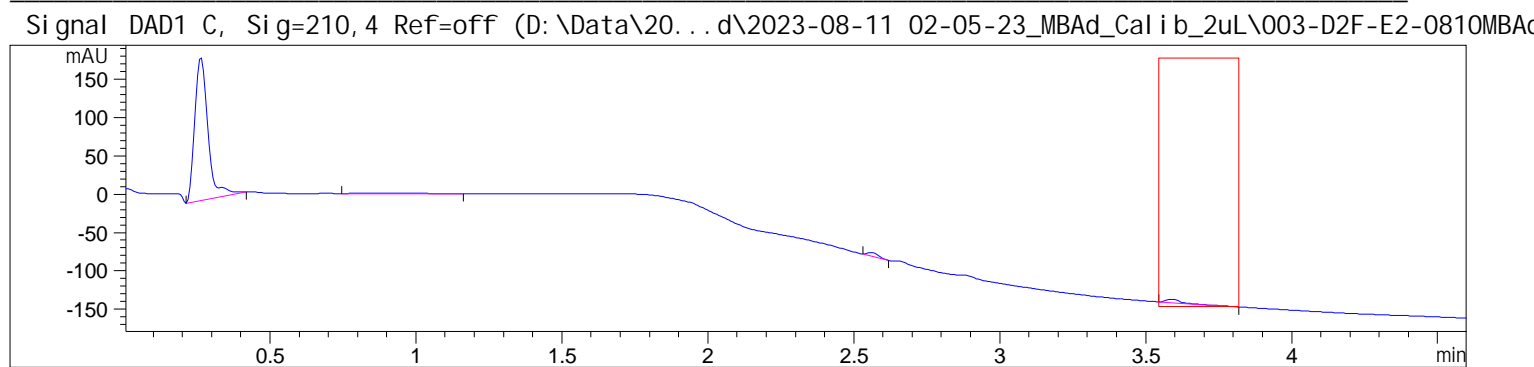


Peak : 12 at 2.891 min Name : ?



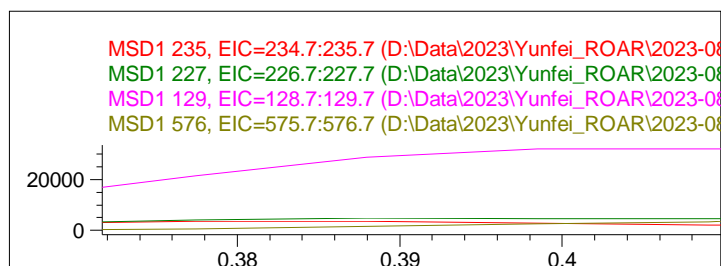
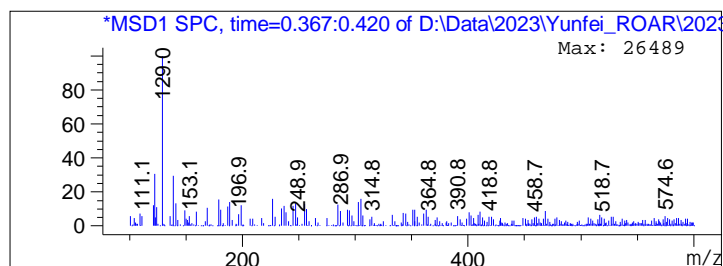
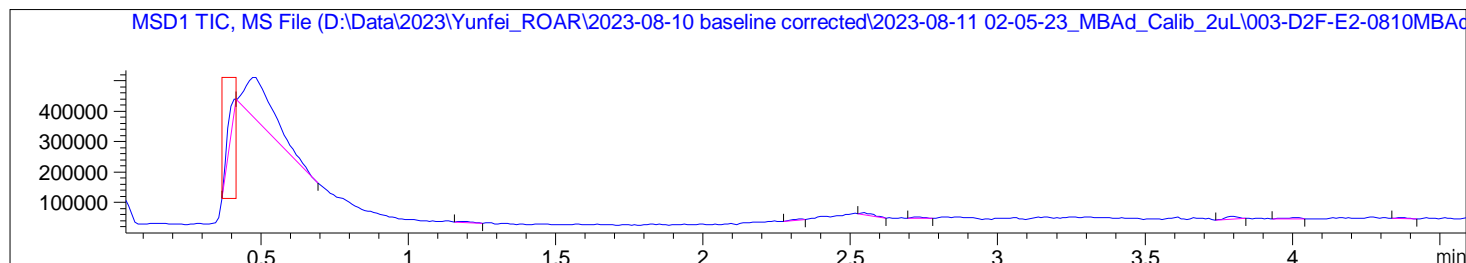
-> The purity factor is within the calculated threshold limit. <-

Purity factor : 547.539 (20 of 20 spectra are within the calculated threshold limit.)
Threshold : 40.816 (Calculated with 20 of 20 spectra)
Reference : Peak start and end spectra (integrated) (2.812 / 2.992)
Spectra : 5 (Selection automatic, 5)
Noise Threshold : 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)



-> The purity factor is within the calculated threshold limit. <-

Purity factor : 736.575 (31 of 31 spectra are within the calculated threshold limit.)
Threshold : 291.235 (Calculated with 31 of 31 spectra)
Reference : Peak start and end spectra (integrated) (3.545 / 3.819)
Spectra : 5 (Selection automatic, 5)
Noise Threshold: 0.745 (12 spectra, St.Dev 0.311 + 3 * 0.1445)

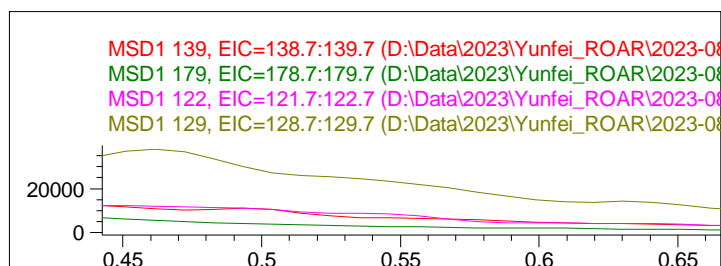
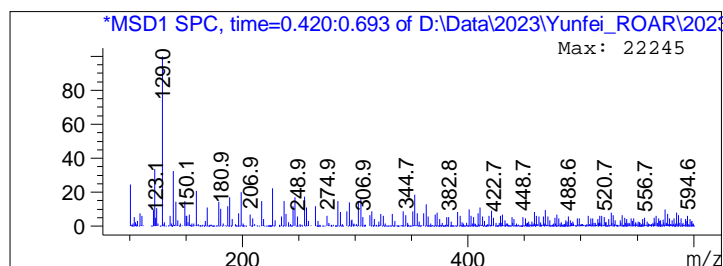
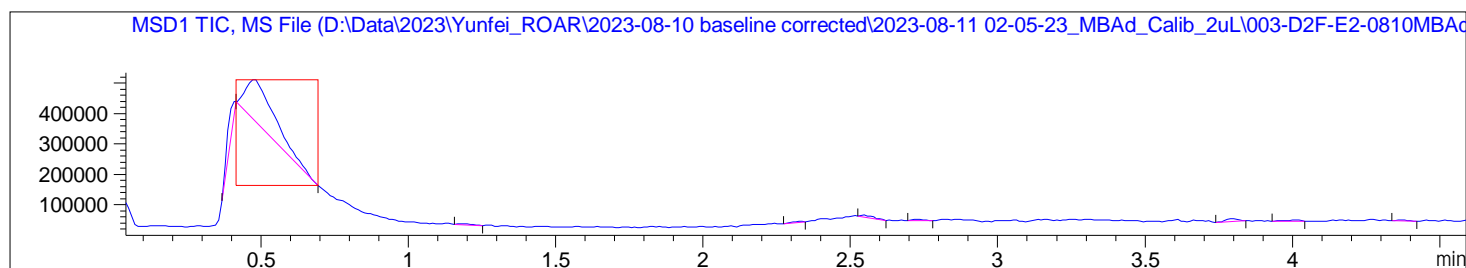


Peak #1 at 0.414 min (0.367 to 0.415 min)

-> The analysis found 2 components, indicating an impure peak. <-

Component 1: Peak at Scan 33.9. Top ions are 235 227 399

Component 2: Peak at Scan 35.0. Top ions are 129 576 255



Peak #2 at 0.477 min (0.415 to 0.693 min)

-> The analysis found 7 components, indicating an impure peak. <-

Component 1: Peak at Scan 38.6. Top ions are 139 179

Component 2: Peak at Scan 39.5. Top ions are 122

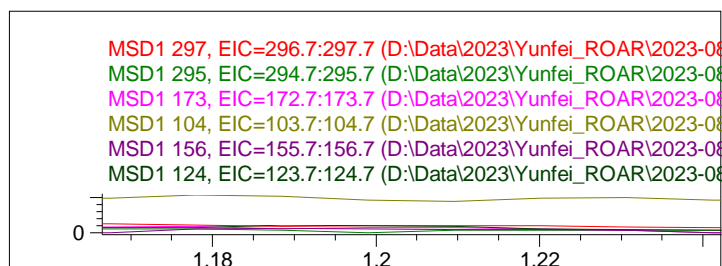
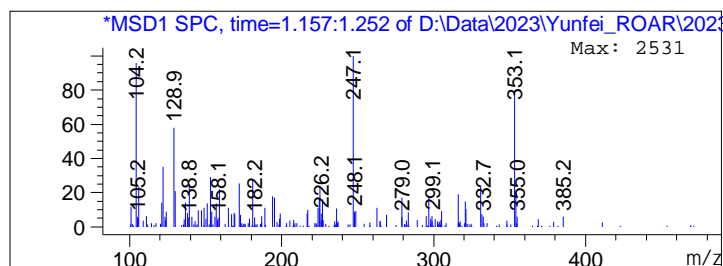
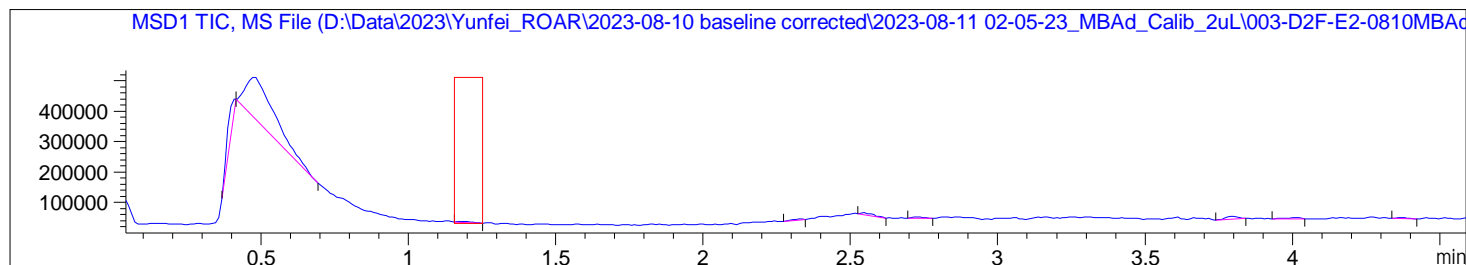
Component 3: Peak at Scan 40.9. Top ions are 129

Component 4: Peak at Scan 42.4. Top ions are 575 585

Component 5: Peak at Scan 43.1. Top ions are 101

Component 6: Peak at Scan 44.8. Top ions are 199 217

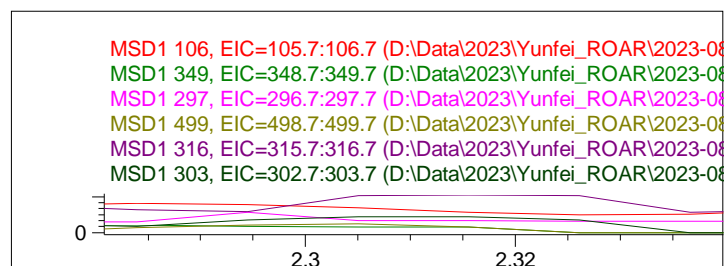
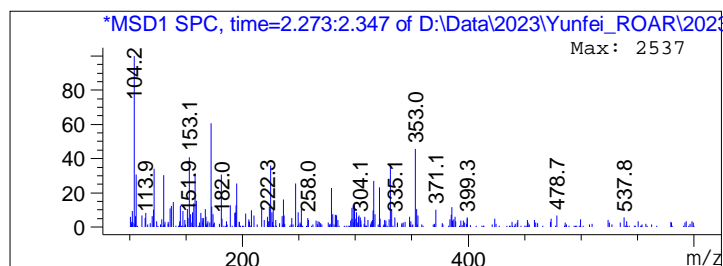
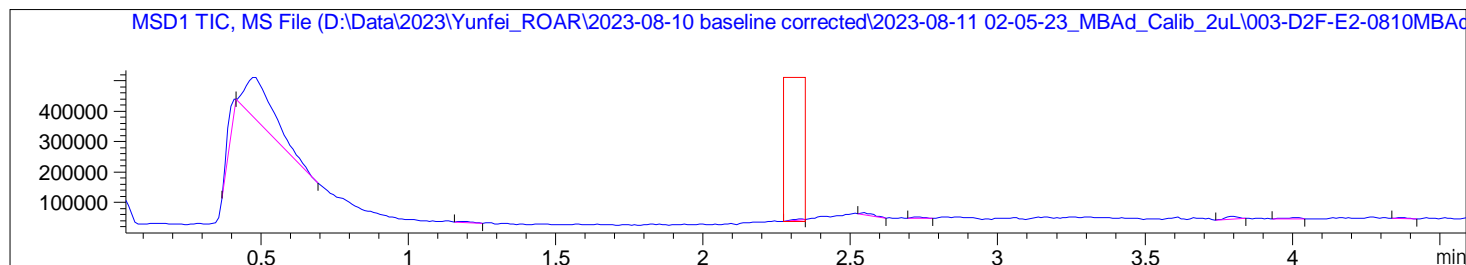
Component 7: Peak at Scan 46.6. Top ions are 159



Peak #3 at 1.178 min (1.157 to 1.252 min)

-> The analysis found 8 components, indicating an impure peak. <-

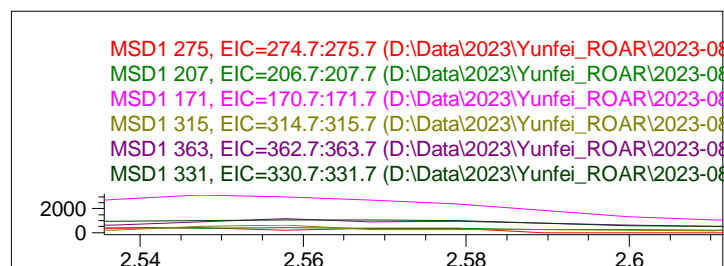
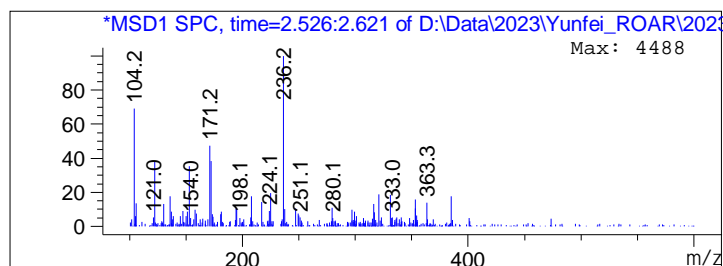
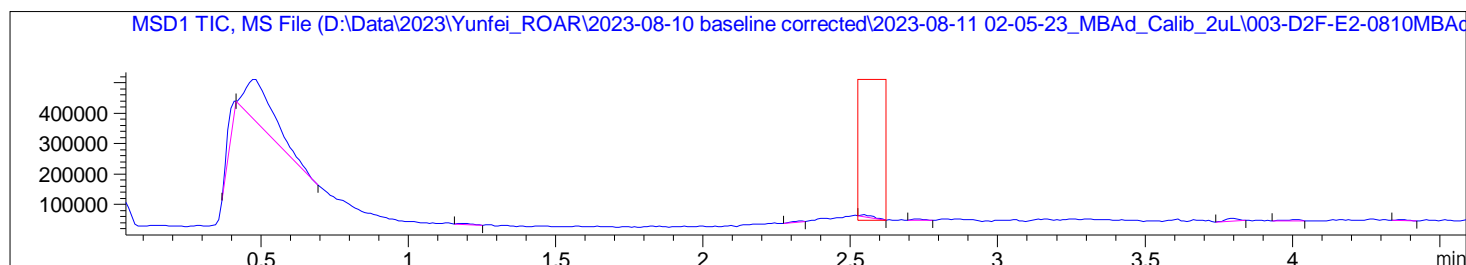
Component 1:	Peak at Scan 108.1.	Top ions are	297 295 269
Component 2:	Peak at Scan 109.2.	Top ions are	173 104 130
Component 3:	Peak at Scan 110.2.	Top ions are	156 124 223
Component 4:	Peak at Scan 111.2.	Top ions are	247 224 333
Component 5:	Peak at Scan 112.0.	Top ions are	199 169 289
Component 6:	Peak at Scan 113.0.	Top ions are	139 303 226
Component 7:	Peak at Scan 113.9.	Top ions are	178 217 331
Component 8:	Peak at Scan 115.0.	Top ions are	353 354 182



Peak #4 at 2.327 min (2.273 to 2.347 min)

-> The analysis found 6 components, indicating an impure peak. <-

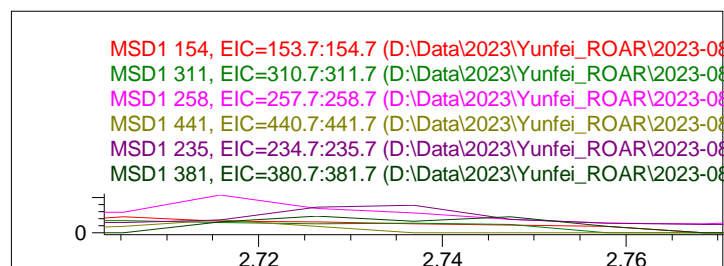
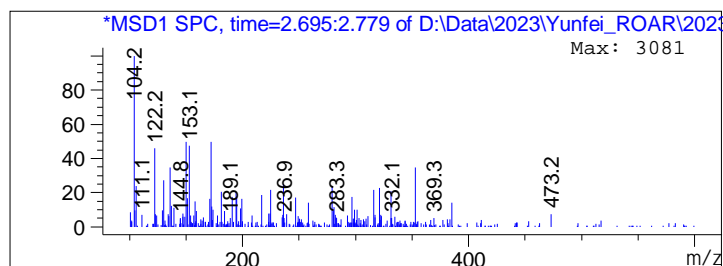
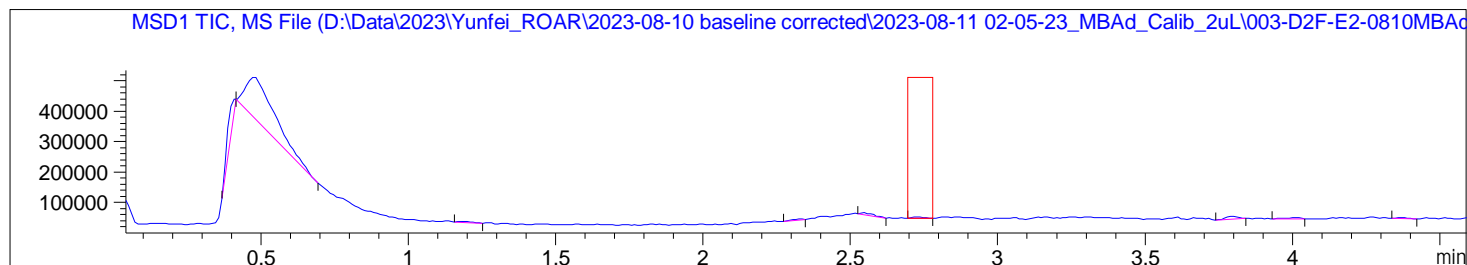
Component 1:	Peak at Scan 214.1.	Top ions are	106 349 122
Component 2:	Peak at Scan 215.3.	Top ions are	297 499 284
Component 3:	Peak at Scan 216.2.	Top ions are	316 303 444
Component 4:	Peak at Scan 217.3.	Top ions are	299 524 156
Component 5:	Peak at Scan 218.4.	Top ions are	282 149 121
Component 6:	Peak at Scan 219.2.	Top ions are	165 399 423



Peak #5 at 2.546 min (2.526 to 2.621 min)

-> The analysis found 7 components, indicating an impure peak. <-

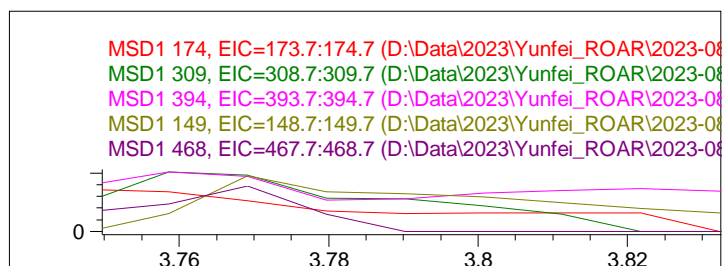
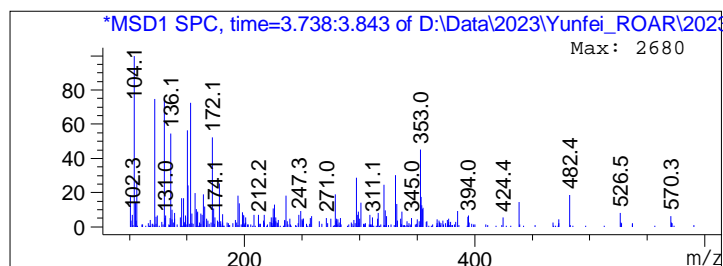
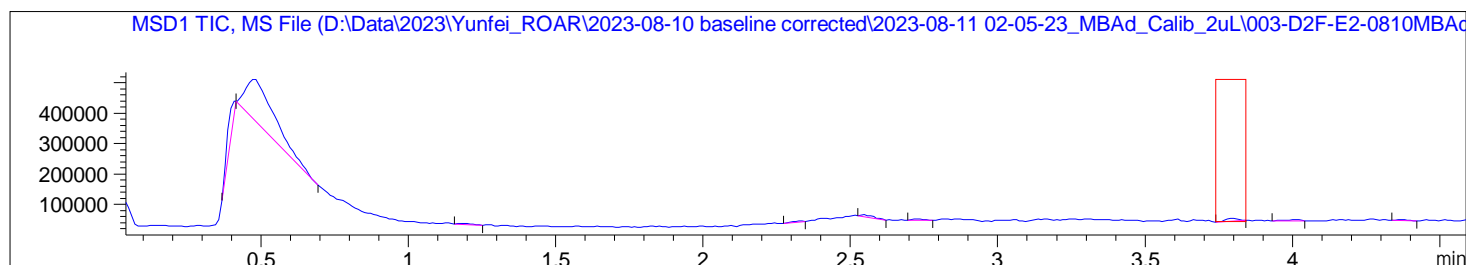
Component 1:	Peak at Scan 238.2.	Top ions are	275 207 517
Component 2:	Peak at Scan 239.2.	Top ions are	171 315 223
Component 3:	Peak at Scan 240.6.	Top ions are	363 331 153
Component 4:	Peak at Scan 241.8.	Top ions are	401 307 333
Component 5:	Peak at Scan 242.9.	Top ions are	385 136 252
Component 6:	Peak at Scan 244.2.	Top ions are	104 240 284
Component 7:	Peak at Scan 245.0.	Top ions are	106 340



Peak #6 at 2.725 min (2.695 to 2.779 min)

-> The analysis found 7 components, indicating an impure peak. <-

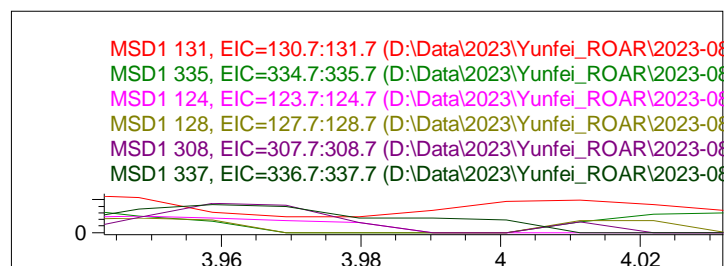
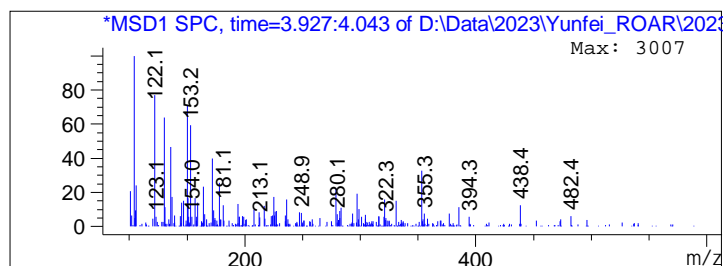
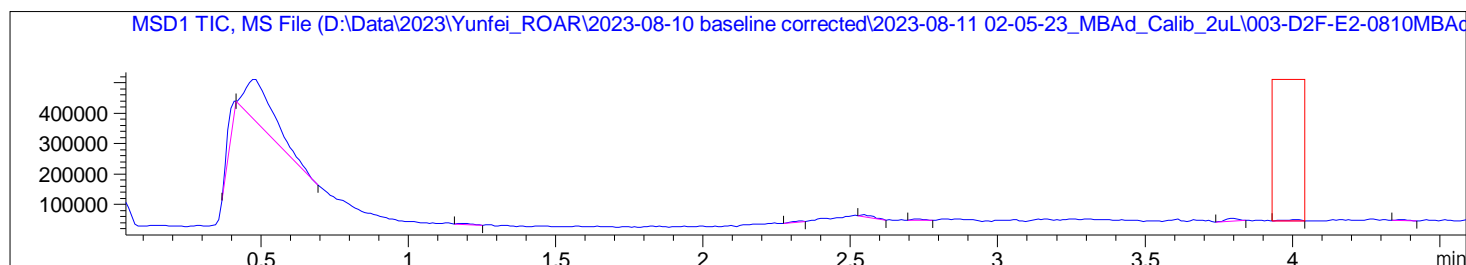
Component 1:	Peak at Scan 254.0.	Top ions are	154 311 121
Component 2:	Peak at Scan 255.0.	Top ions are	258 441 227
Component 3:	Peak at Scan 256.2.	Top ions are	235 381 165
Component 4:	Peak at Scan 257.2.	Top ions are	186 411 104
Component 5:	Peak at Scan 258.1.	Top ions are	150 198 284
Component 6:	Peak at Scan 259.0.	Top ions are	197 383 331
Component 7:	Peak at Scan 260.1.	Top ions are	248 147 282



Peak #7 at 3.797 min (3.739 to 3.843 min)

-> The analysis found 9 components, indicating an impure peak. <-

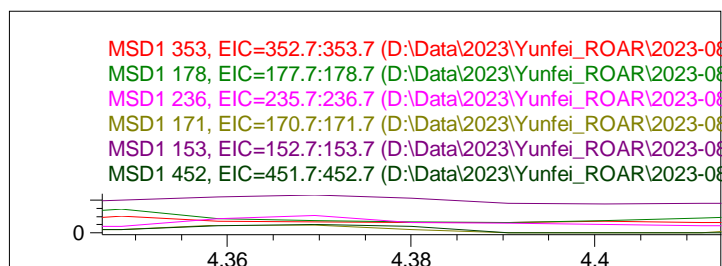
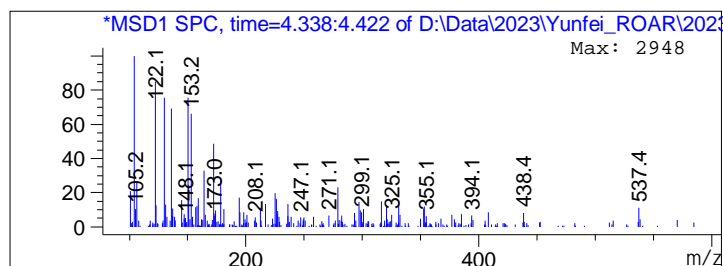
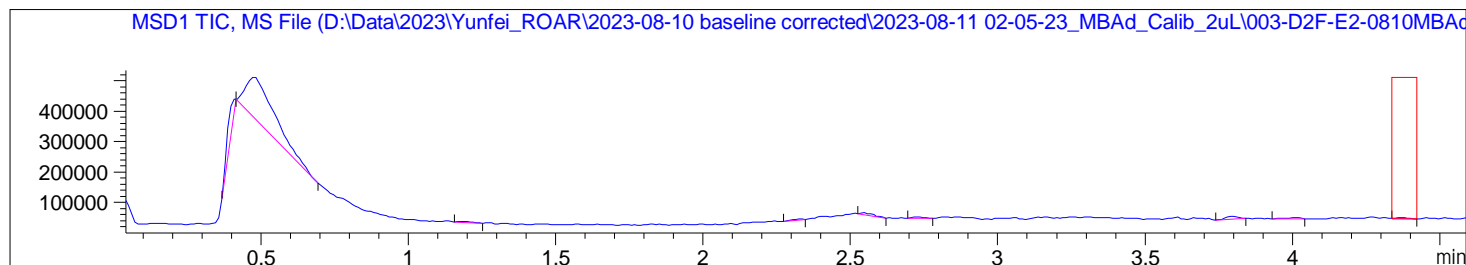
Component 1:	Peak at Scan 352.9.	Top ions are	174
Component 2:	Peak at Scan 354.1.	Top ions are	309 394 265
Component 3:	Peak at Scan 355.2.	Top ions are	149 468 250
Component 4:	Peak at Scan 356.3.	Top ions are	106 298 225
Component 5:	Peak at Scan 357.4.	Top ions are	353 130 213
Component 6:	Peak at Scan 358.4.	Top ions are	331 172 332
Component 7:	Peak at Scan 359.2.	Top ions are	526 379 352
Component 8:	Peak at Scan 360.0.	Top ions are	203 571 336
Component 9:	Peak at Scan 361.2.	Top ions are	354 148 537



Peak #8 at 4.010 min (3.932 to 4.043 min)

-> The analysis found 9 components, indicating an impure peak. <-

Component 1:	Peak at Scan 371.0.	Top ions are	131 335 150
Component 2:	Peak at Scan 372.1.	Top ions are	124 128 537
Component 3:	Peak at Scan 373.1.	Top ions are	308 337 314
Component 4:	Peak at Scan 374.2.	Top ions are	217 165 227
Component 5:	Peak at Scan 375.4.	Top ions are	304 395 297
Component 6:	Peak at Scan 376.2.	Top ions are	282 189 291
Component 7:	Peak at Scan 377.3.	Top ions are	164 105 322
Component 8:	Peak at Scan 378.8.	Top ions are	122 280 136
Component 9:	Peak at Scan 380.0.	Top ions are	293 354 496



Peak #9 at 4.368 min (4.338 to 4.422 min)

-> The analysis found 7 components, indicating an impure peak. <-

Component 1: Peak at Scan 410.0. Top ions are 353 178 439
Component 2: Peak at Scan 411.3. Top ions are 236 171 315
Component 3: Peak at Scan 412.2. Top ions are 153 452 225
Component 4: Peak at Scan 413.4. Top ions are 305 158 438
Component 5: Peak at Scan 414.7. Top ions are 145 537 239
Component 6: Peak at Scan 415.7. Top ions are 159 249 190
Component 7: Peak at Scan 416.4. Top ions are 394

*** End of Report ***