

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  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      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  XXXX  XX      X  XX      XX XX  XX
  XXX      XXXXX  XXX  XXX  XX      XXXX  XXXXX X      XXX      XXXXX
                XXXX

```

Sample Name: dimos 23_10 1.1

```

=====
Acq. Operator   : SYSTEM                      Seq. Line :    2
Acq. Instrument : HPLC-OXTLAB                 Location  :   P2-A-02
Injection Date  : 10/23/2023 14:09:58         Inj       :    1
                                           Inj Volume: 20.000 µl
Acq. Method     : C:\Chem32\1\Data\Dimos-Ntina-Xaris-23.10.23 2023-10-23 13-22-49\LACTIC_TEMP
                  .M
Last changed    : 8/28/2023 13:32:52 by SYSTEM
Analysis Method : C:\Chem32\1\Data\Dimos-Ntina-Xaris-23.10.23 2023-10-23 13-22-49\LACTIC_TEMP
                  .M (Sequence Method)
Last changed    : 10/23/2023 15:59:07 by SYSTEM
                  (modified after loading)
=====

```

```

=====
Module                               Type   Firmware rev.   Serial number
-----|-----|-----|-----
RID                                  G7162A D.07.10 [0004] DEAC901169
DAD 2                               G7117C D.07.10 [0004] DEAEK01369
Column Comp. 3                      G7116A D.07.21 [0001] DEAE01239
Sampler 4                           G7129A D.07.12 [0002] DEAEQ08976
Quat. Pump 5                        G7111A D.07.21 [0001] DEAEX00817
=====

```

Software Revision: Rev. C.01.07 SR3 [465] Copyright © Agilent Technologies

```

=====
Instrument Conditions :      At Start          At Stop
Column Temp. (left)  :      50.0              50.0 °C
Column Temp. (right) :      50.0              50.0 °C
Pressure              :      41.3              41.7 bar
Flow                  :      0.600             0.600 ml/min
=====

```

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Detector Lamp Burn Times: Current On-Time Accumulated On-Time
DAD 1, UV Lamp          :      2.52           3115.2 h
=====

```

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Solvent Description :
PMP1, Solvent A     :
PMP1, Solvent B     :
PMP1, Solvent C     :
PMP1, Solvent D     :
=====

```

Run Logbook

```

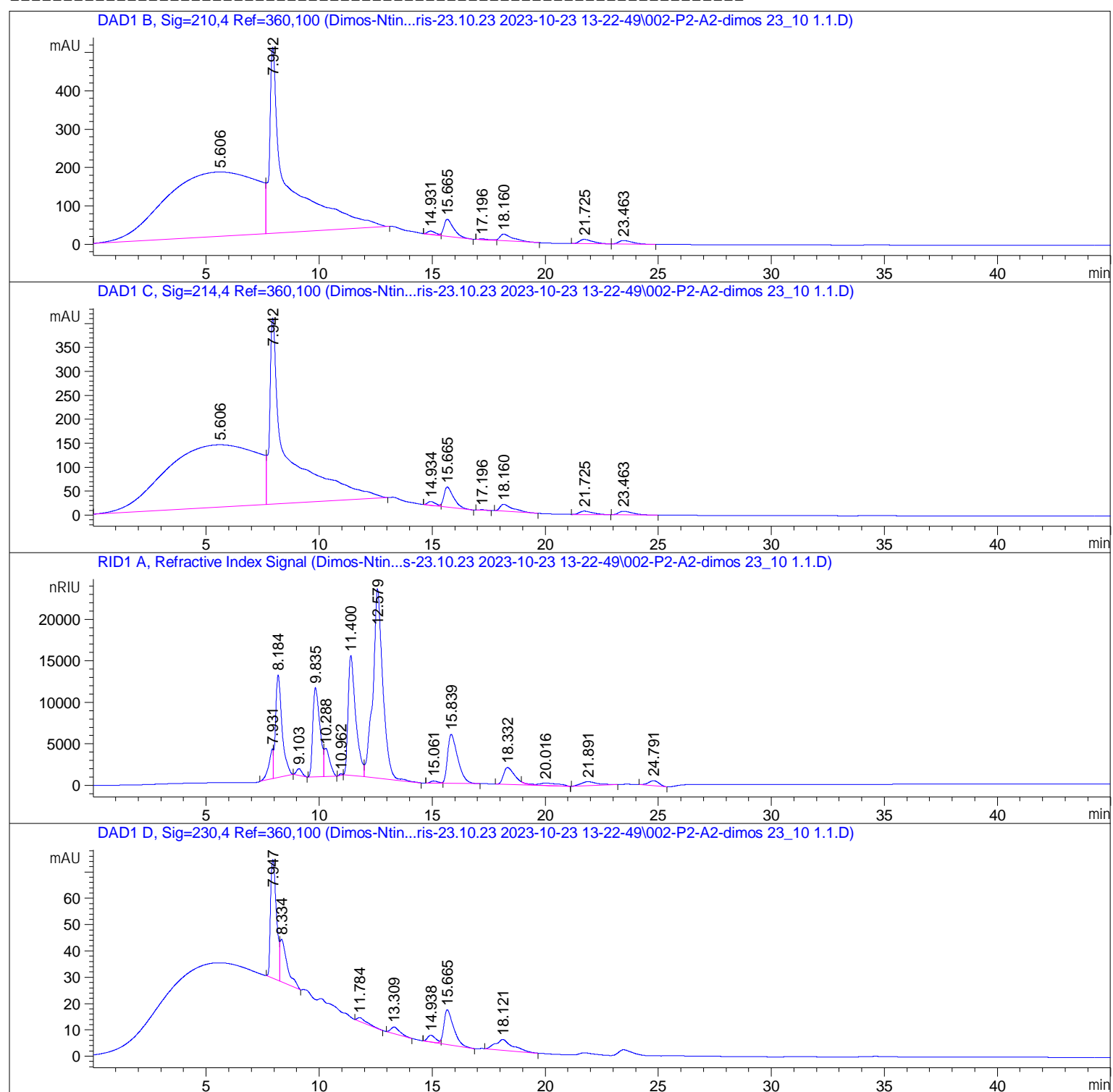
=====
23 Oct 23 03:59 PM
Logbook File: C:\Chem32\...ina-Xaris-23.10.23 2023-10-23 13-22-49\002-P2-A2-dimos 23_10 1.1.D\RUN.LOG
=====

```

Module	# Event Message	Date Time
Method	Method started: line# 2 at location 'P2-A2' > inj# 1	10/23/2023 14:09:00
Method	Instrument running sample P2-A2	10/23/2023 14:09:00
G7117C	G7117C: DEAEK01369 - Detector: Prepare	10/23/2023 14:09:01
G7117C	G7117C: DEAEK01369 - Detector: Idle	10/23/2023 14:09:20
G7129A	G7129A: DEAEQ08976 - Wash procedure active	10/23/2023 14:09:46
G7129A	G7129A: DEAEQ08976 - Wash procedure inactive	10/23/2023 14:09:55
G7129A	G7129A: DEAEQ08976 - Run	10/23/2023 14:09:58
G7162A	G7162A: DEAC901169 - Postrun	10/23/2023 14:54:58
G7129A	G7129A: DEAEQ08976 - Postrun	10/23/2023 14:54:58
G7111A	G7111A: DEAEX00817 - Postrun	10/23/2023 14:54:58

Sample Name: dimos 23_10 1.1

G7116A	G7116A: DEAEM01239 - Postrun	10/23/2023 14:54:58
Method	Instrument run completed	10/23/2023 14:55:05
Method	Saving Method LACTIC_TEMP.M	10/23/2023 14:55:05
CP Macro	Analyzing rawdata 002-P2-A2-dimos 23_10 1.1. > D	10/23/2023 14:55:05
Method	Saving Method DA.M	10/23/2023 14:55:05
Method	Method completed	10/23/2023 14:55:06



External Standard Report

Sorted By : Signal
Calib. Data Modified : 7/1/2021 14:39:58
Multiplier : 1.0000
Dilution : 1.0000
Do not use Multiplier & Dilution Factor with ISTDs

Signal 1: DAD1 B, Sig=210,4 Ref=360,100

RetTime [min]	Type	Area [mAU*s]	Amt/Area	Amount [g/L]	Grp	Name
15.665	VB R	1323.98901	6.56715e-4	8.69484e-1		Lactic acid
18.160	VB R	720.08173	9.16554e-4	6.59994e-1		Acetic acid
21.725	BB	473.34052	0.00000	0.00000		Propionic

Totals : 1.52948

Signal 2: DAD1 C, Sig=214,4 Ref=360,100

Signal 3: RID1 A, Refractive Index Signal

RetTime [min]	Type	Area [nRIU*s]	Amt/Area	Amount [g/L]	Grp	Name
9.835	BV	2.36229e5	1.03247e-5	2.43899		Succrose
11.400	BV	3.37424e5	2.80485e-6	9.46424e-1		Glucose
12.579	VB	7.00292e5	3.79459e-6	2.65732		Fructose
24.791	BB	1.85821e4	4.65862e-6	8.65667e-2		Ethanol

Totals : 6.12930

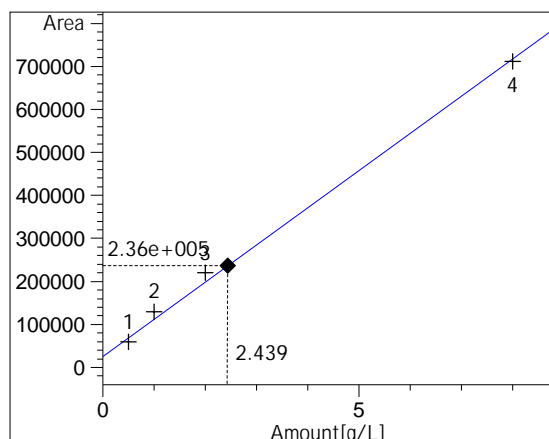
Signal 4: DAD1 D, Sig=230,4 Ref=360,100

2 Warnings or Errors :

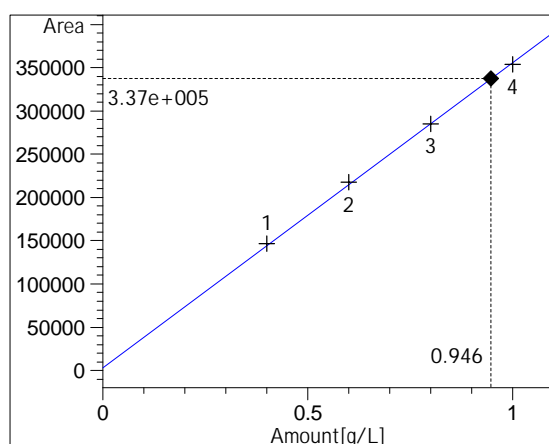
Warning : Calibration warnings (see calibration table listing)
Warning : Negative results set to zero (cal. curve intercept), (Propionic)

Sample Name: di mos 23_10 1.1

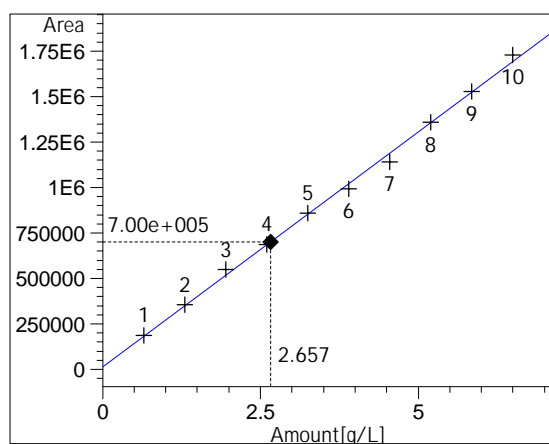
Calibration Curves



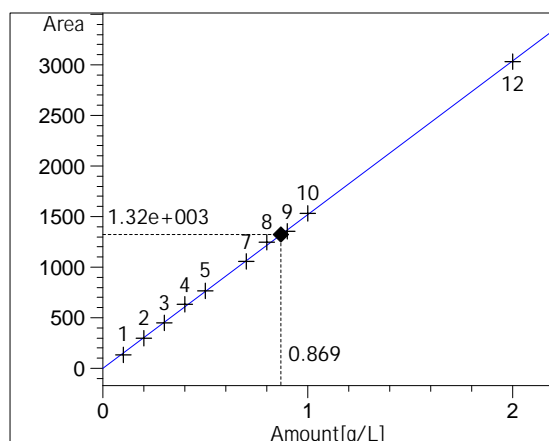
Succrose at exp. RT: 9.863
 RID1 A, Refractive Index Signal
 Correlation: 0.99759
 Residual Std. Dev.: 22755.41185
 Formula: $y = mx + b$
 m: 86671.29673
 b: 24839.01752
 x: Amount[g/L]
 y: Area



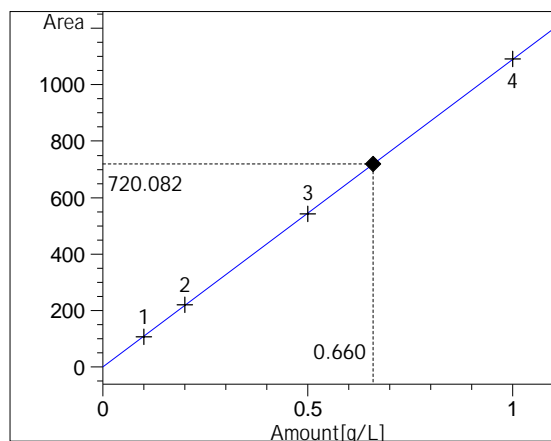
Glucose at exp. RT: 11.408
 RID1 A, Refractive Index Signal
 Correlation: 0.99981
 Residual Std. Dev.: 3023.36360
 Formula: $y = mx + b$
 m: 353737.16216
 b: 2639.18919
 x: Amount[g/L]
 y: Area



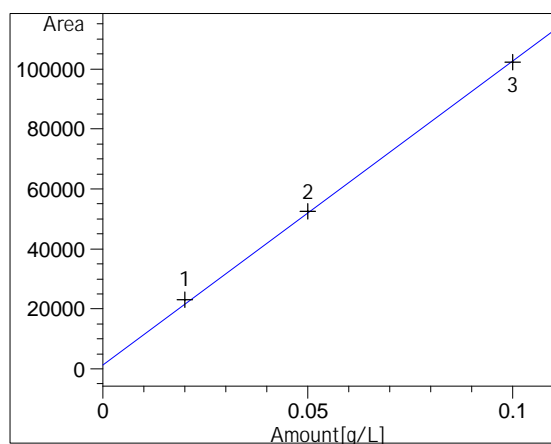
Fructose at exp. RT: 12.100
 RID1 A, Refractive Index Signal
 Correlation: 0.99909
 Residual Std. Dev.: 25081.95145
 Formula: $y = mx + b$
 m: 258721.41259
 b: 12786.68182
 x: Amount[g/L]
 y: Area



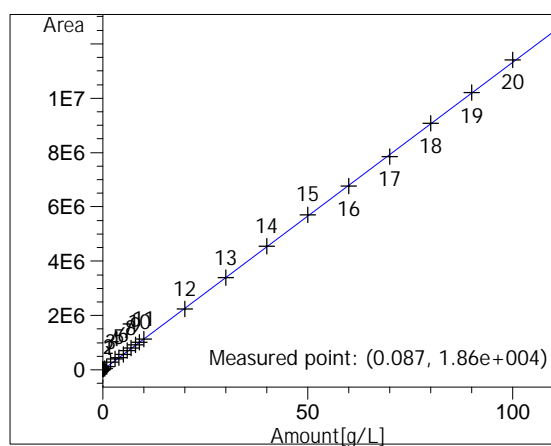
Lactic acid at exp. RT: 15.467
 DAD1 B, Sig=210,4 Ref=360,100
 Correlation: 0.99983
 Residual Std. Dev.: 16.65547
 Formula: $y = mx + b$
 m: 1521.64235
 b: 9.45650e-1
 x: Amount[g/L]
 y: Area



Acetic acid at exp. RT: 18.226
DAD1 B, Sig=210,4 Ref=360,100
Correlation: 0.99999
Residual Std. Dev.: 2.57101
Formula: $y = mx + b$
m: 1092.07931
b: -6.83711e-1
x: Amount[g/L]
y: Area



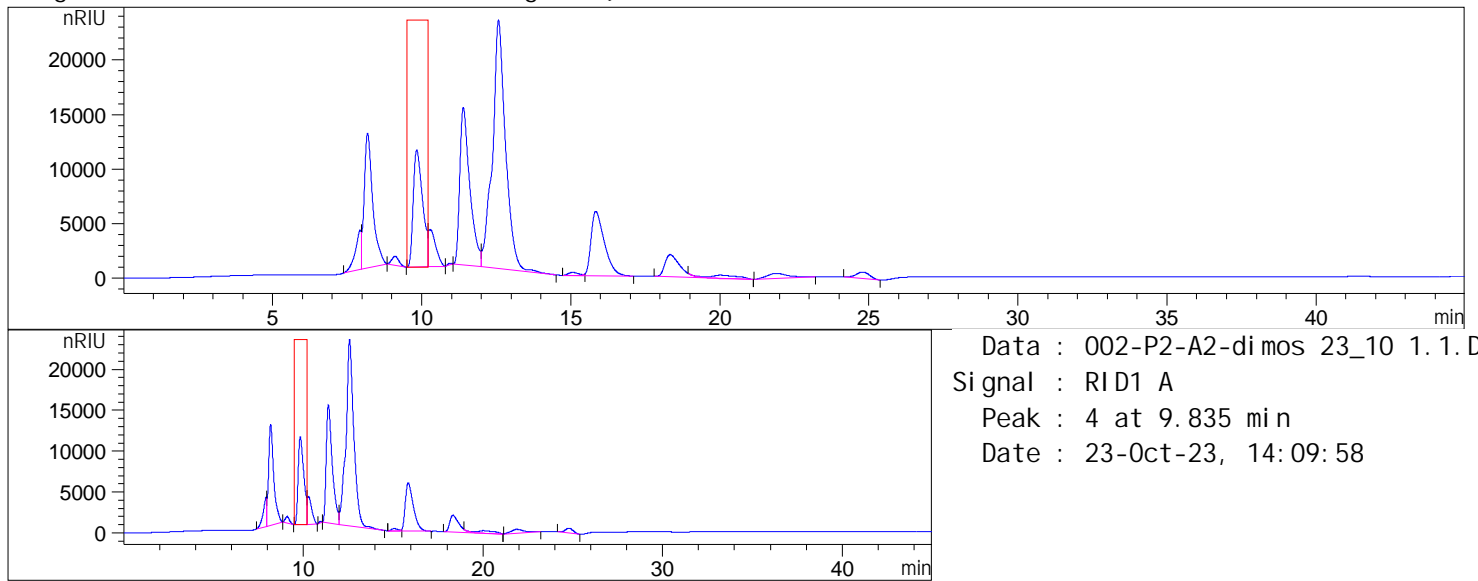
Propionic at exp. RT: 21.787
DAD1 B, Sig=210,4 Ref=360,100
Correlation: 0.99961
Residual Std. Dev.: 1507.31144
Formula: $y = mx + b$
m: 1.01431e6
b: 1395.13216
x: Amount[g/L]
y: Area



Ethanol at exp. RT: 24.978
RID1 A, Refractive Index Signal
Correlation: 0.99996
Residual Std. Dev.: 35437.06772
Formula: $y = mx + b$
m: 113284.07454
b: 8775.42396
x: Amount[g/L]
y: Area

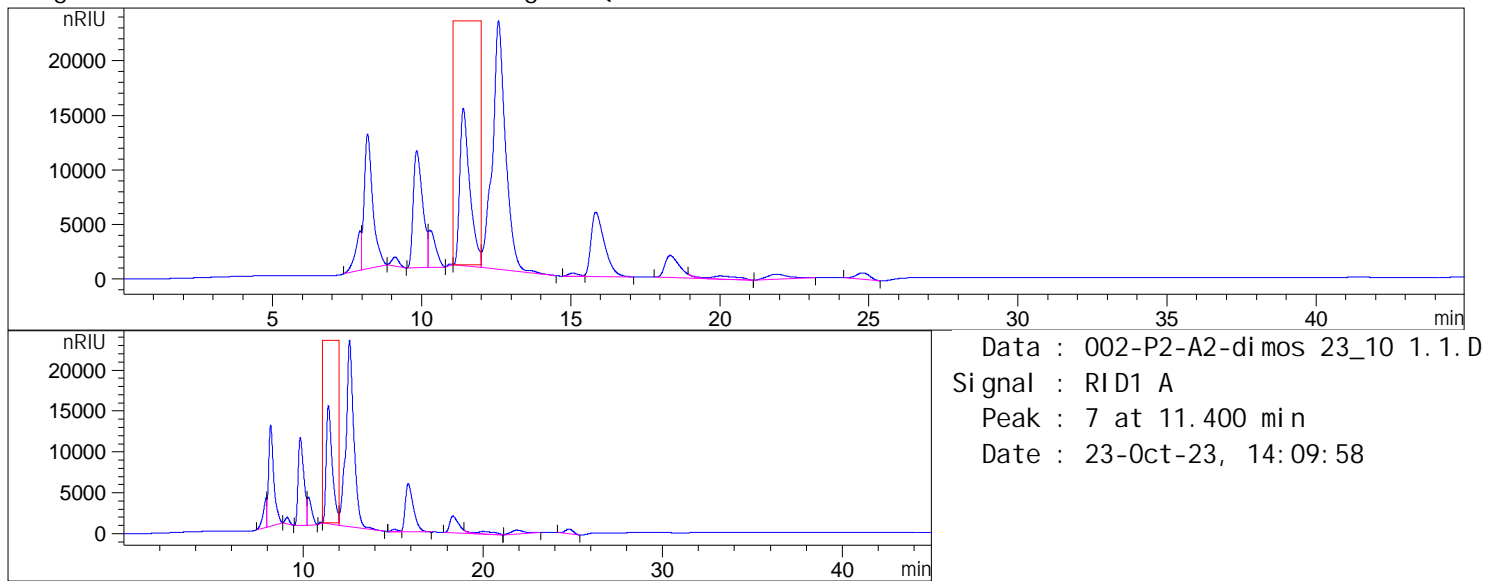
Sample Name: di mos 23_10 1.1

Signal RID1 A, Refractive Index Signal (Dimos-Ntin...s-23.10.23 2023-10-23 13-22-49\002-P2-A2-di mos 23_10 1.1.D



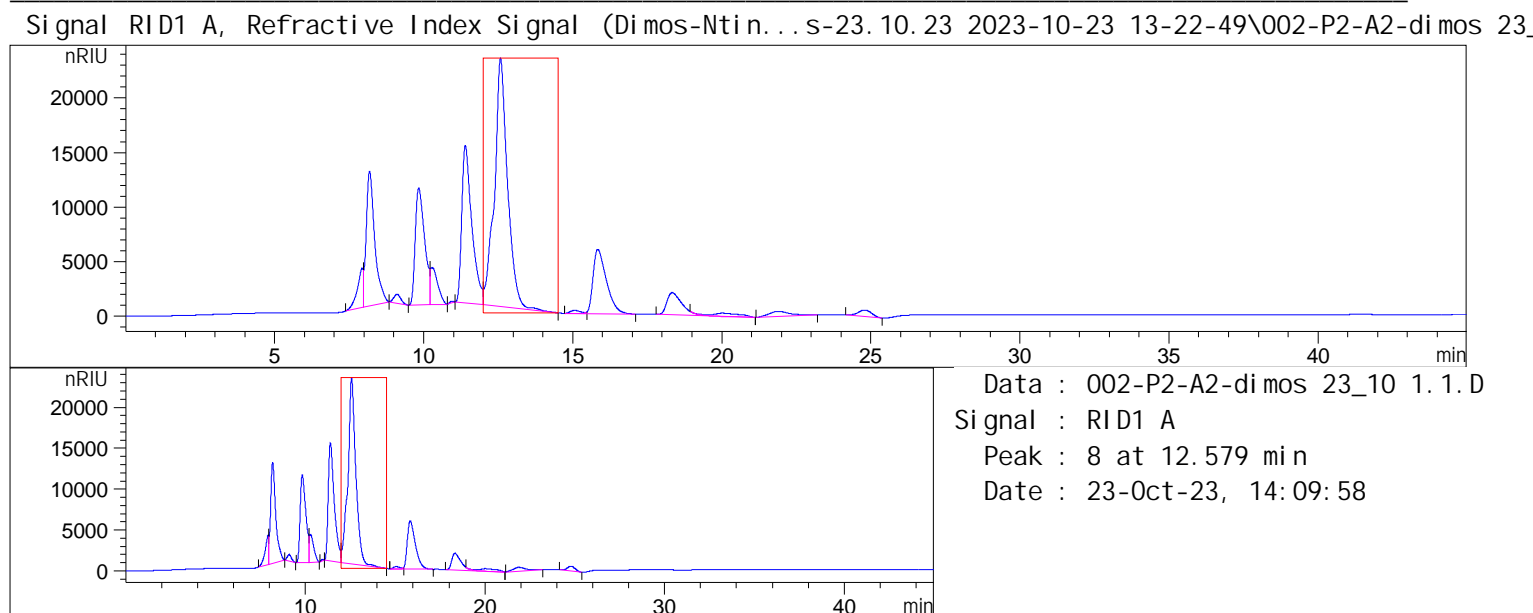
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!! No spectral data found !!
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

Signal RID1 A, Refractive Index Signal (Dimos-Ntin...s-23.10.23 2023-10-23 13-22-49\002-P2-A2-di mos 23_10 1.1.D

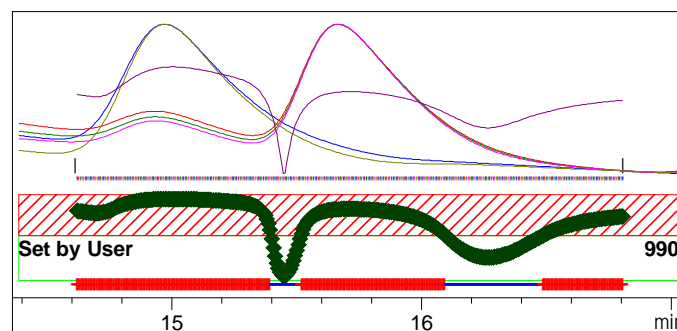
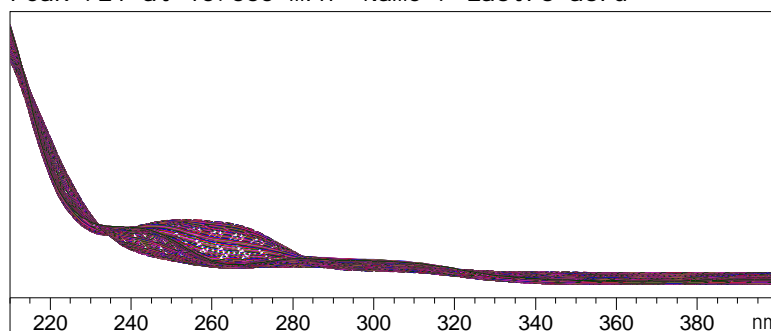
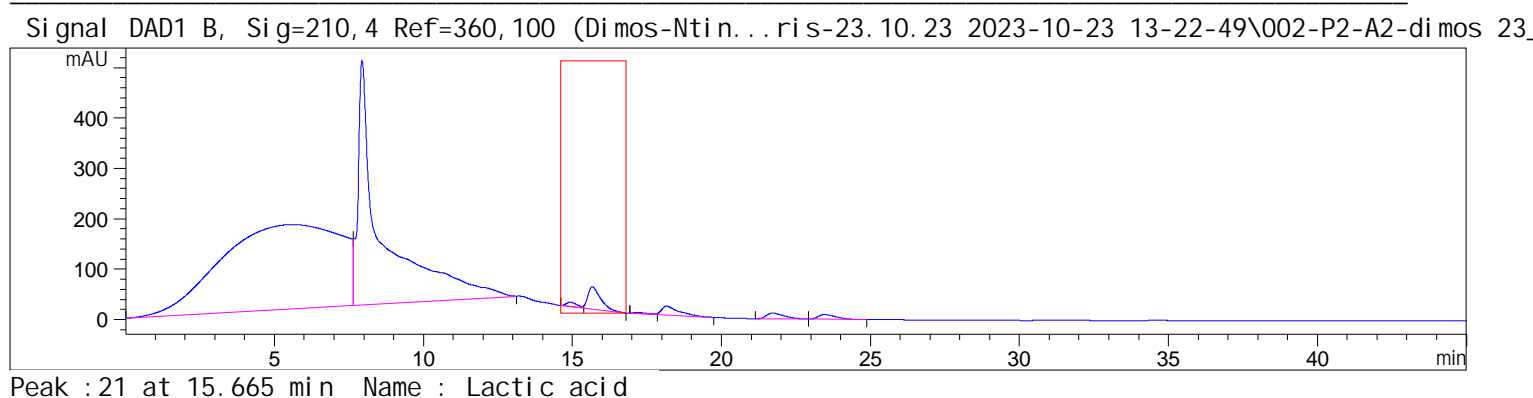


!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!! No spectral data found !!
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

Sample Name: di mos 23_10 1.1



!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!! No spectral data found !!
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

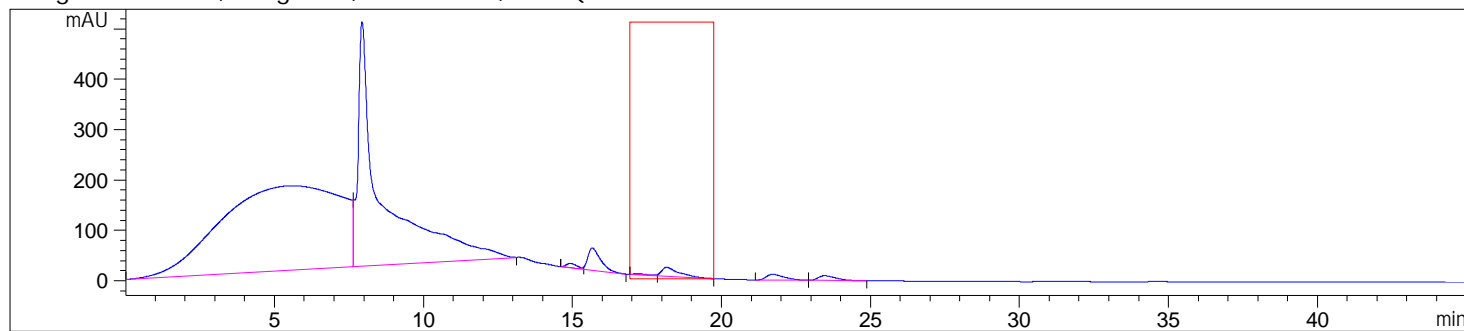


-> The purity factor exceeds the threshold limit. <-

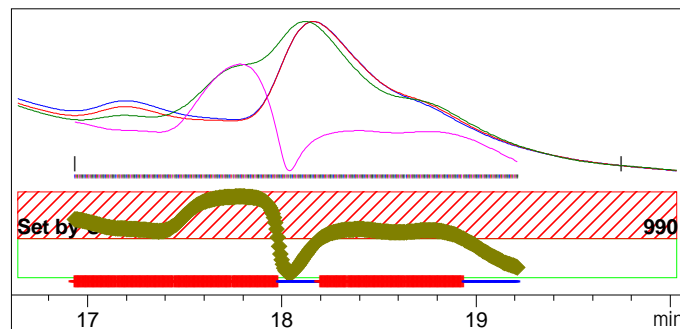
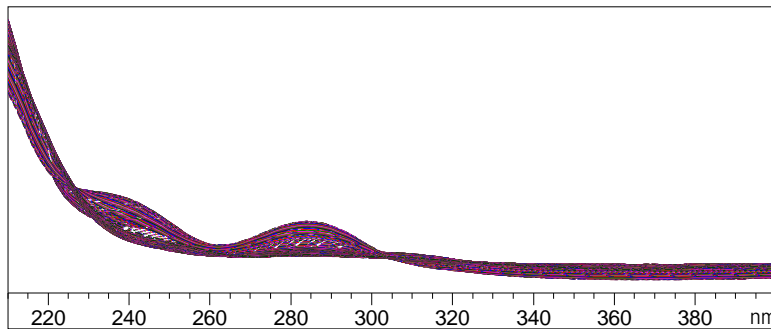
Purity factor : 963.220 (253 of 329 spectra exceed the threshold limit.)
Threshold : 990.000 (Set by user)
Reference : Nearest baseline spectrum (stored) (0.007)
Spectra : 329 (Selection automatic, All)

Sample Name: dimos 23_10 1.1

Signal DAD1 B, Sig=210, 4 Ref=360, 100 (Dimos-Ntin...ris-23.10.23 2023-10-23 13-22-49\002-P2-A2-dimos 23_10 1.1.D)



Peak : 28 at 18.160 min Name : Acetic acid



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

Purity factor : 967.505 (269 of 343 spectra exceed the threshold limit.)

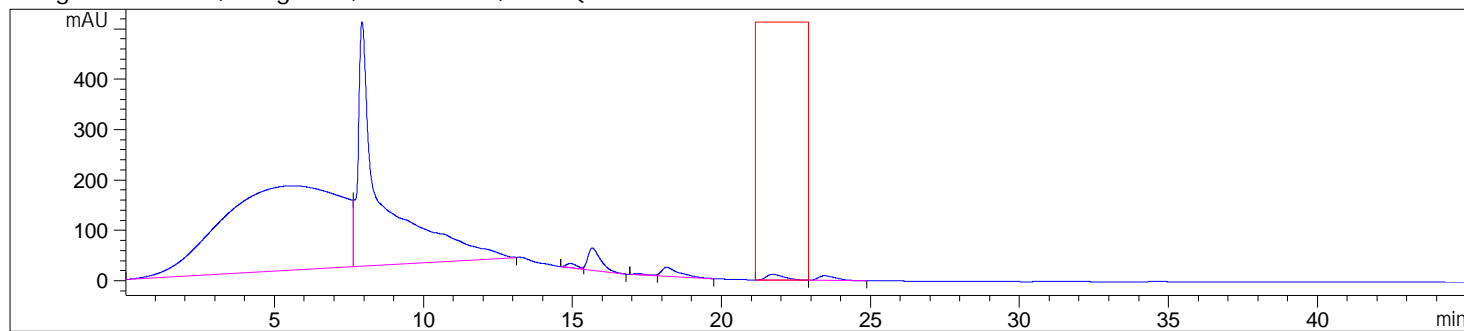
Threshold : 990.000 (Set by user)

Reference : Nearest baseline spectrum (stored) (0.007)

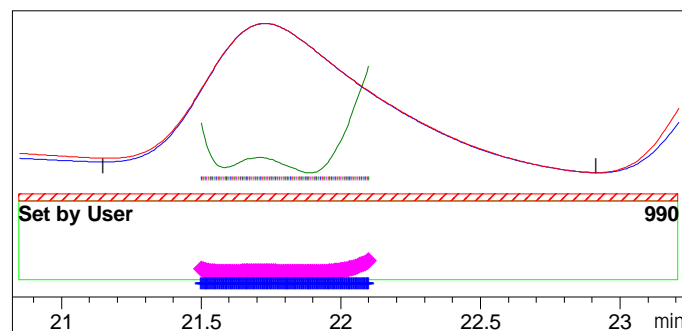
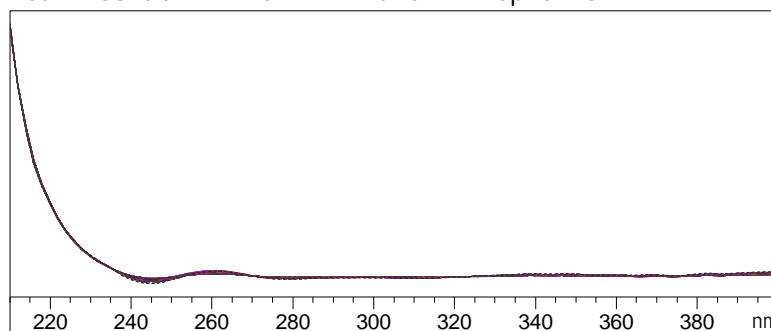
Spectra : 343 (Selection automatic, All)

Sample Name: dimos 23_10 1.1

Signal DAD1 B, Sig=210, 4 Ref=360, 100 (Dimos-Ntin...ris-23.10.23 2023-10-23 13-22-49\002-P2-A2-dimos 23_10 1.1.D)



Peak : 33 at 21.725 min Name : Propionic



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

Purity factor : 999.774 (91 of 91 spectra are within the threshold limit.)

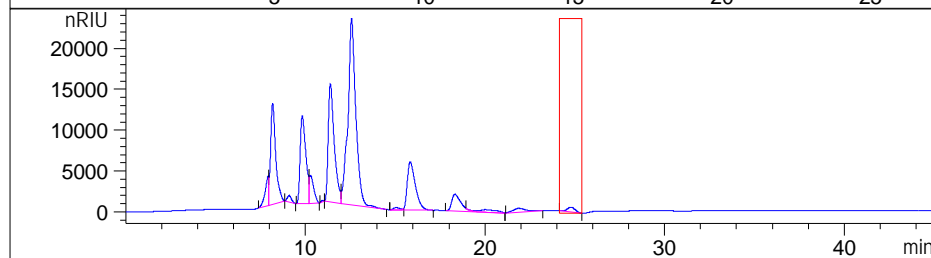
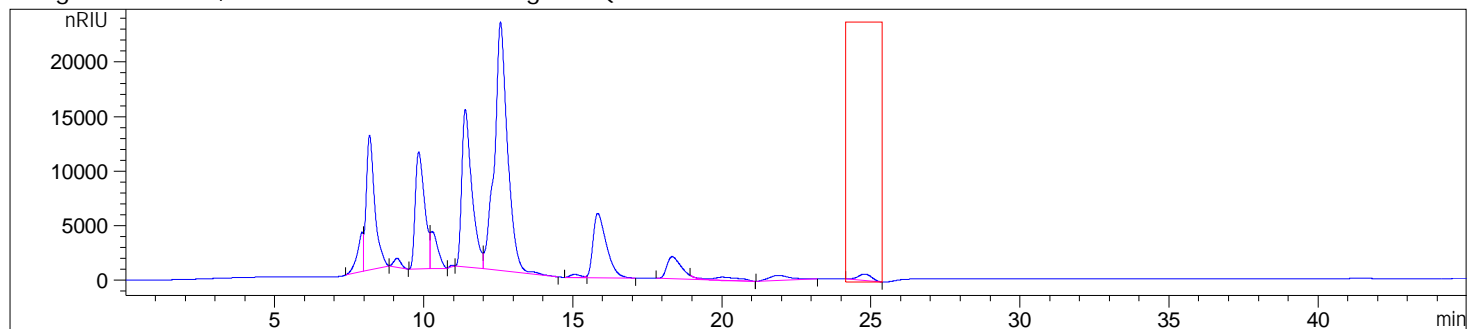
Threshold : 990.000 (Set by user)

Reference : Nearest baseline spectrum (stored) (0.007)

Spectra : 91 (Selection automatic, All)

Sample Name: di mos 23_10 1.1

Signal RID1 A, Refractive Index Signal (Dimos-Ntin...s-23.10.23 2023-10-23 13-22-49\002-P2-A2-di mos 23_10 1.1.D



Data : 002-P2-A2-di mos 23_10 1.1.D
Signal : RID1 A
Peak : 14 at 24.791 min
Date : 23-Oct-23, 14:09:58

!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!! No spectral data found !!

*** End of Report ***