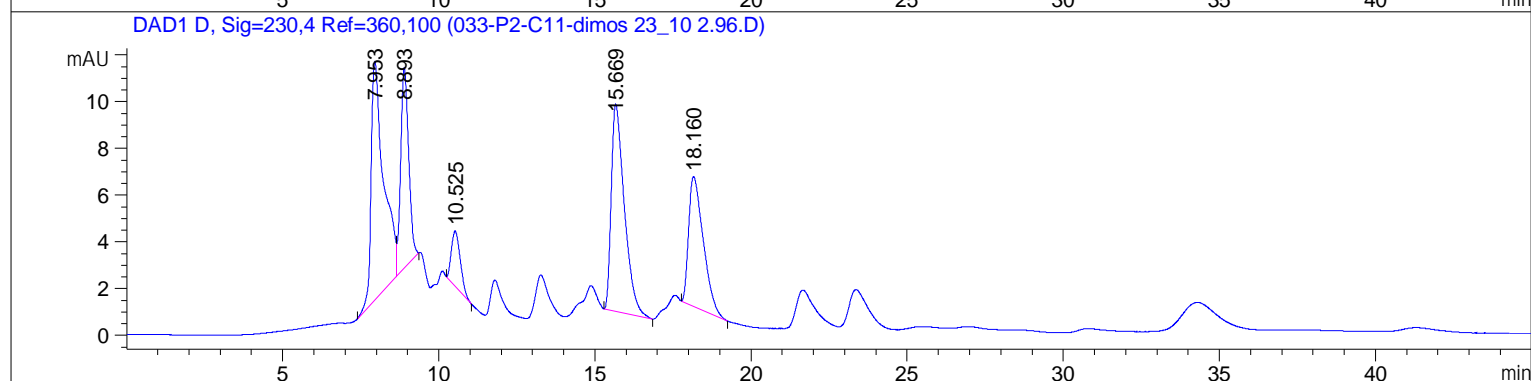
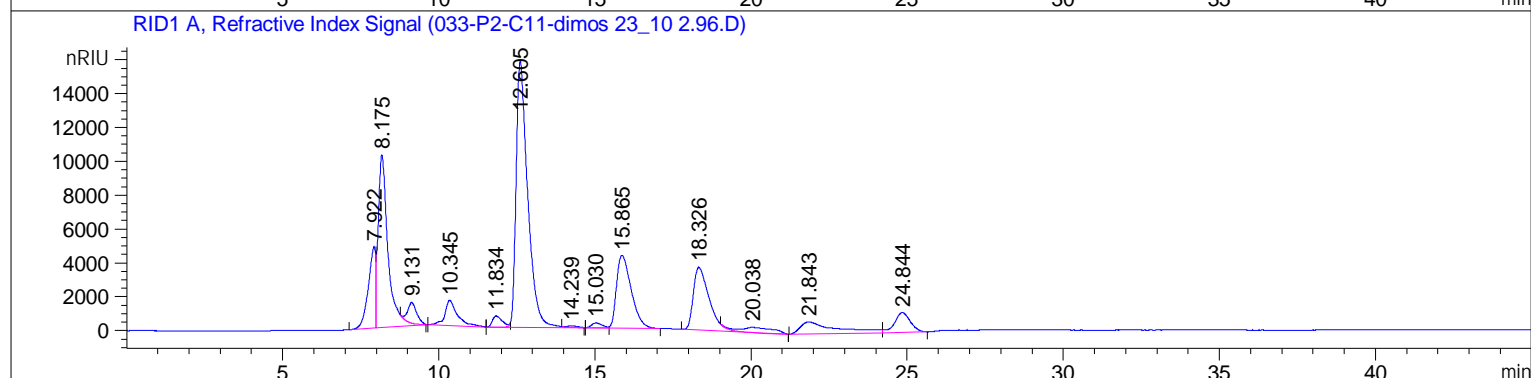
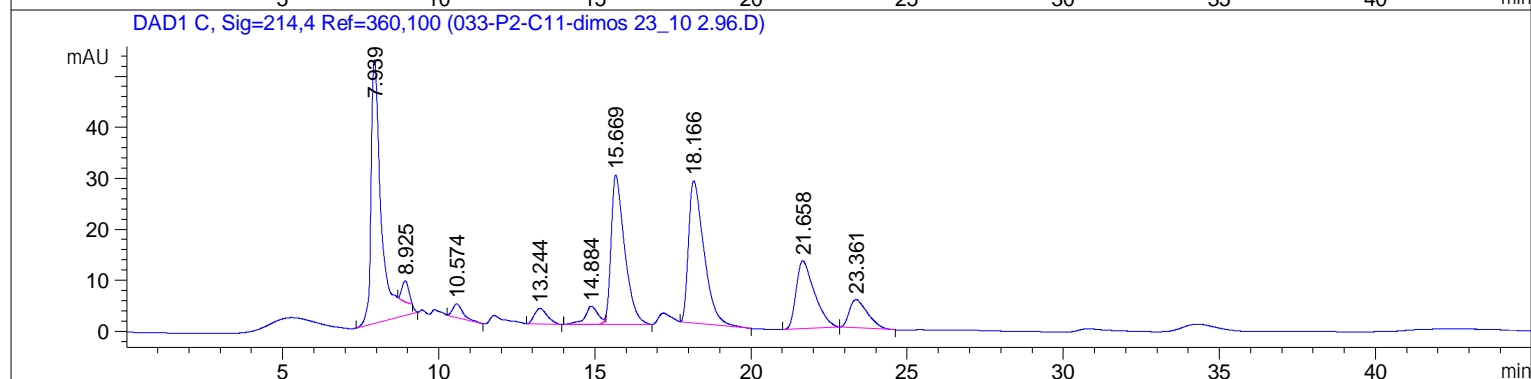


DAD1 B, Sig=210,4 Ref=360,100 (033-P2-C11-dimos 23\_10 2.96.D)

The chromatogram displays absorbance (mAU) on the y-axis (0 to 60) against time (min) on the x-axis (0 to 45). The baseline is relatively flat with minor noise. Several distinct peaks are observed and labeled with their retention times. The peak at 7.939 min is the most intense, reaching approximately 60 mAU. Other significant peaks are at 15.669 min and 18.166 min, both around 35 mAU. Smaller peaks are visible at 8.931, 10.587, 13.236, 14.882, 17.190, 21.658, and 23.360 min. A pink line is overlaid on the chromatogram, starting around 7.5 min and ending around 10 min, possibly indicating a specific time range of interest.

Retention Time (min)	Approximate Absorbance (mAU)
7.939	60
8.931	10
10.587	10
13.236	5
14.882	5
15.669	35
17.190	5
18.166	35
21.658	25
23.360	10



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
15.669	VB R	938.74713	6.56523e-4	6.16309e-1		Lactic acid
18.166	VB R	1178.45557	9.16216e-4	1.07972		Acetic acid
21.658	BB	861.96796	0.00000	0.00000		Propionic

Totals : 1.69603

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
9.863		-	-	-		Succrose
11.408		-	-	-		Glucose
11.834	BV E	1.42559e4	3.98341e-7	5.67870e-3		Fructose
24.844	VB	4.11391e4	6.94439e-6	2.85686e-1		Ethanol

Totals : 2.91365e-1

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

3 Warnings or Errors :

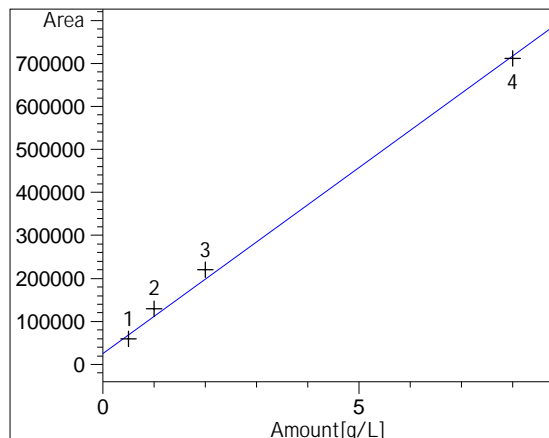
Warning : Calibration warnings (see calibration table listing)  
Warning : Calibrated compound(s) not found  
Warning : Negative results set to zero (cal. curve intercept), (Propionic)

Sample Name: dimos 23\_10 2.96

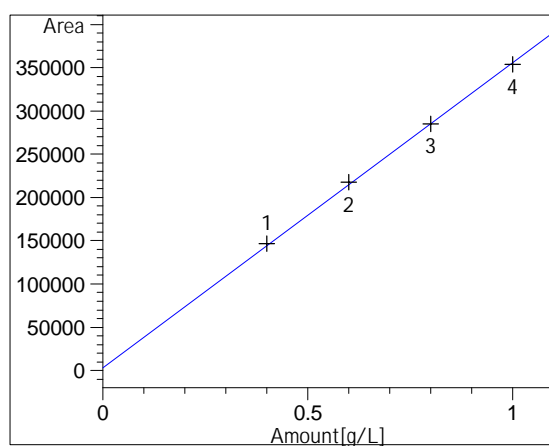
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Calibration Curves

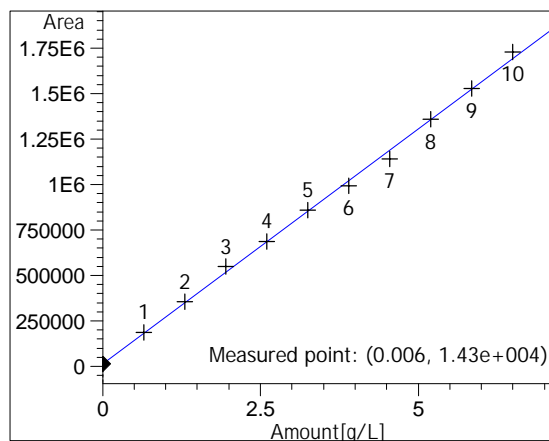
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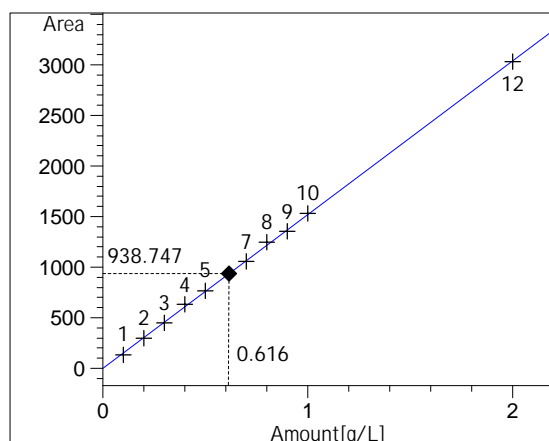
Succrose at exp. RT: 9.863  
 RID1 A, Refractive Index Signal  
 Correlation: 0.99759  
 Residual Std. Dev.: 22755.41185  
 Formula:  $y = mx + b$   
     m: 86671.29673  
     b: 24839.01752  
     x: Amount [g/L]  
     y: Area



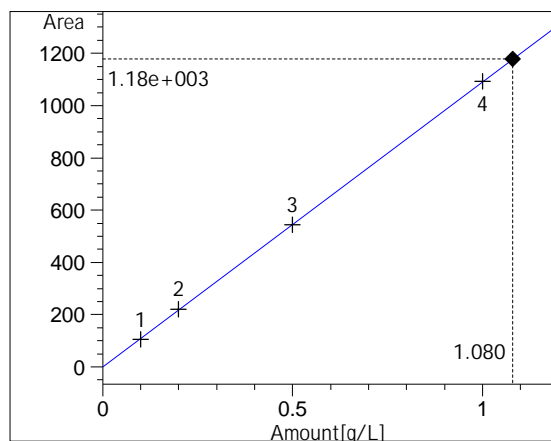
Glucose at exp. RT: 11.408  
 RID1 A, Refractive Index Signal  
 Correlation: 0.99981  
 Residual Std. Dev.: 3023.36360  
 Formula:  $y = mx + b$   
     m: 353737.16216  
     b: 2639.18919  
     x: Amount [g/L]  
     y: Area



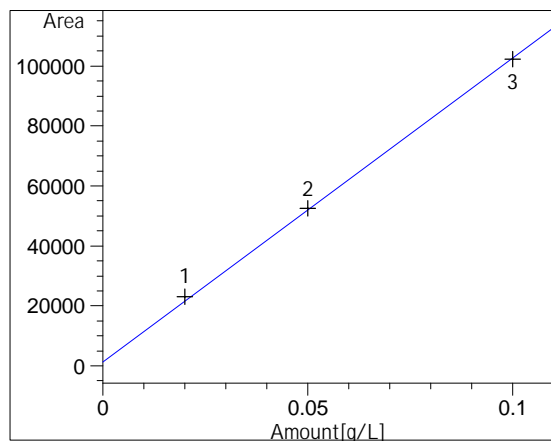
Fructose at exp. RT: 12.100  
 RID1 A, Refractive Index Signal  
 Correlation: 0.99909  
 Residual Std. Dev.: 25081.95145  
 Formula:  $y = mx + b$   
     m: 258721.41259  
     b: 12786.68182  
     x: Amount [g/L]  
     y: Area



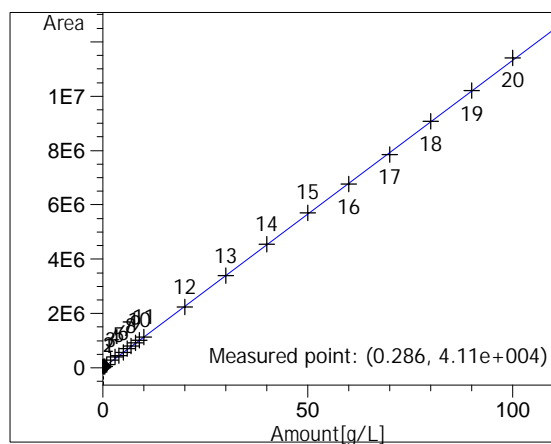
Lactic acid at exp. RT: 15.467  
 DAD1 B, Sig=210, 4 Ref=360, 100  
 Correlation: 0.99983  
 Residual Std. Dev.: 16.65547  
 Formula:  $y = mx + b$   
     m: 1521.64235  
     b: 9.45650e-1  
     x: Amount [g/L]  
     y: Area



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

\*\*\* End of Report \*\*\*