

Figure 1 Supplementary Information: Stability of select amino acids after 24 hours of exposure to 80 °C. Results are average of triplicate measurements, with error bars representing +/- one standard deviation.

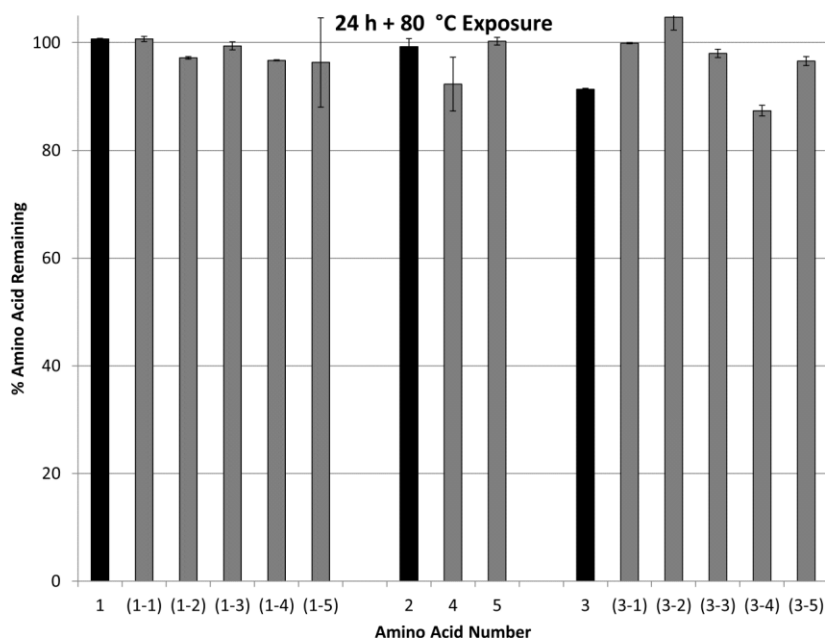
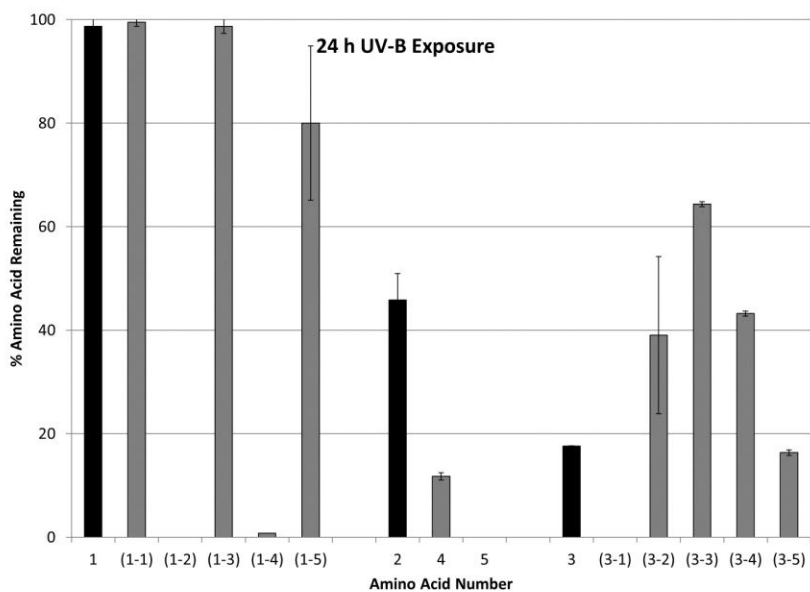
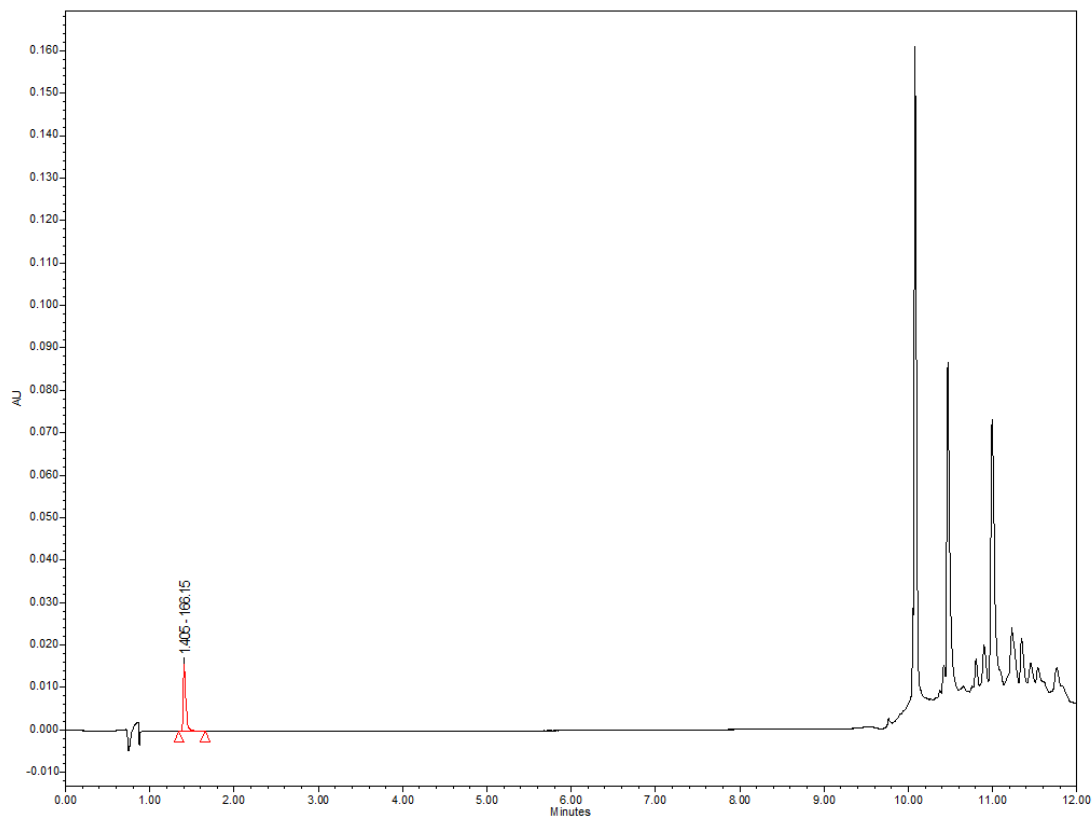


Figure 2 Supplementary Information: Stability of select amino acids after being exposed to UV-B radiation for 24 h. Bars are the average of triplicate measurements, and error bars represent +/- one standard deviation.



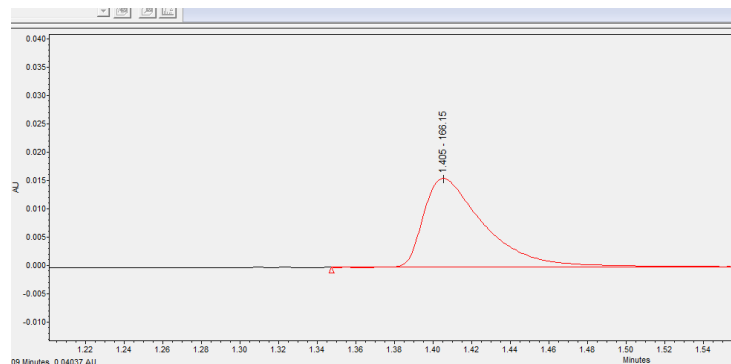
**UV Exposure data** (pg 2-117) of LC absorbance at 254 nm, with integrated area, and corresponding extracted mass spectrum. One sample chromatogram of each control (number not followed by any other identifying information, such as UVA, UVB, UVC), and UVA, UVB, and UVC irradiated samples are shown. Graphs in paper were produced by averaging the results of triplicate measurements. Amino acid identified with number in Tables 1 and 2, 1 for example, is Phe.

#### 1 LC PDA Detector Data with Integrated Peak

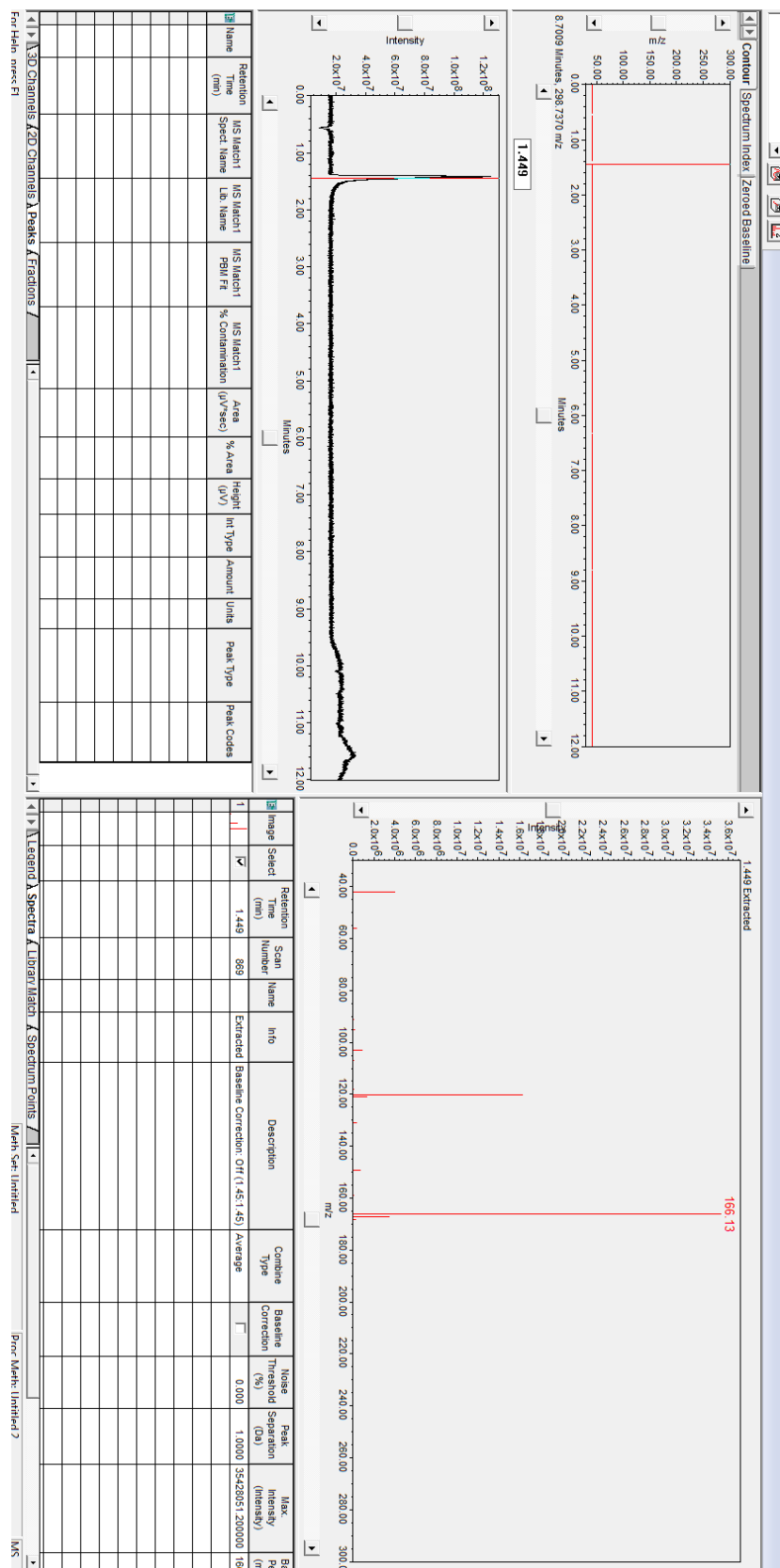


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.405	35021	100.00	15687	bb			Unknown

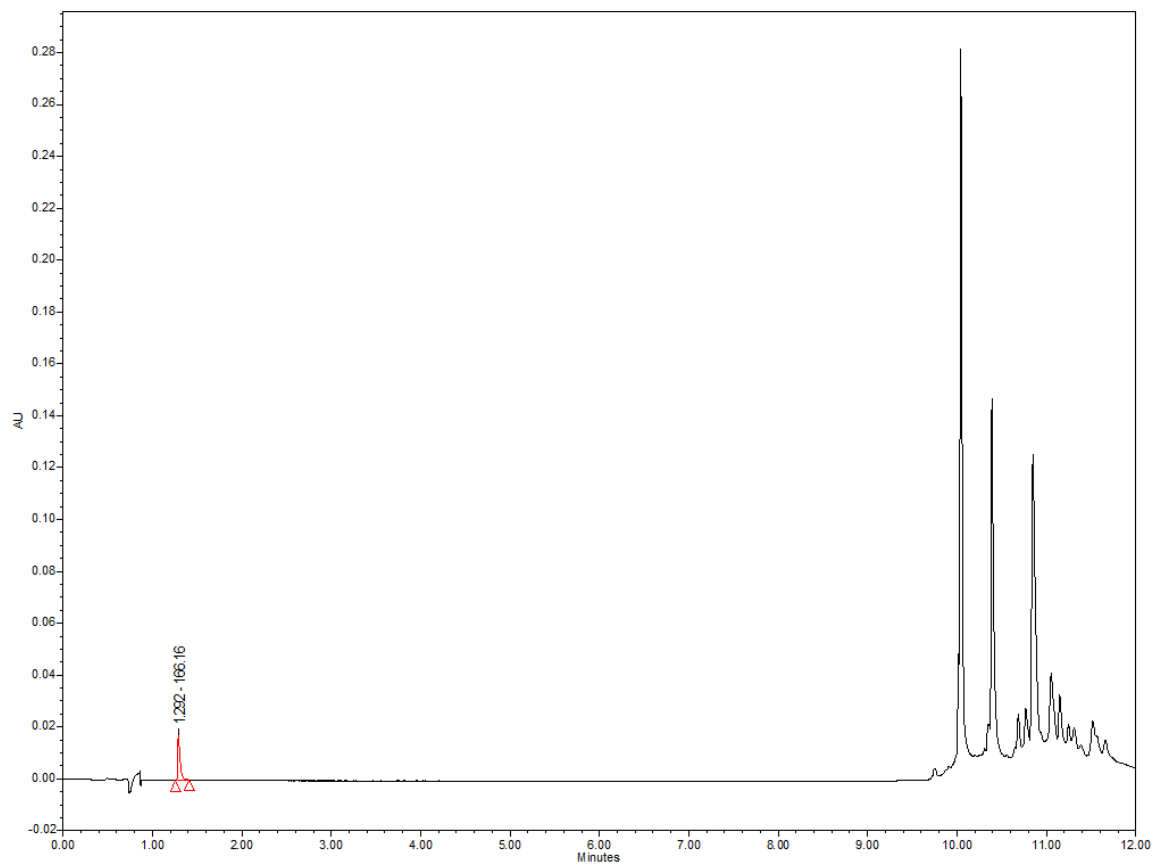
Zoom in of integrated peak shown above:



For Help, press F1

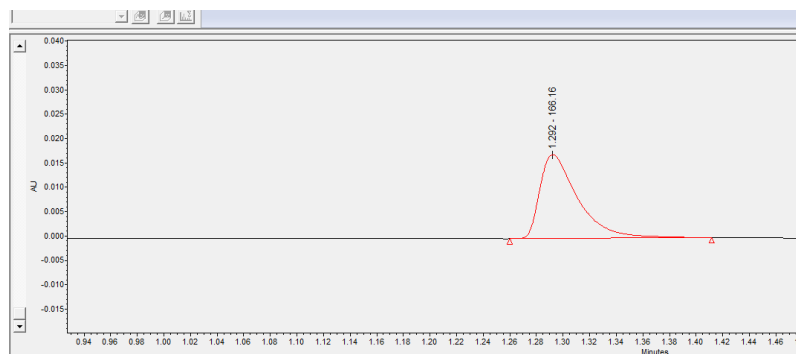


# 1 UV-A LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.292	34000	100.00	17219	bb			Unknown

Zoom in of integrated peak shown above:





**3D Channels / 2D Channels / Peaks / Fractions**

**Contour Spectrum Index / Zeroed Baseline**

**1.316**

**1.316 Extracted**

**Intensity**

**Intensity**

**Retention Time (min)**

**Scan Number**

**Info**

**Description**

**Combine Type**

**Baseline Correction**

**Noise Threshold (%)**

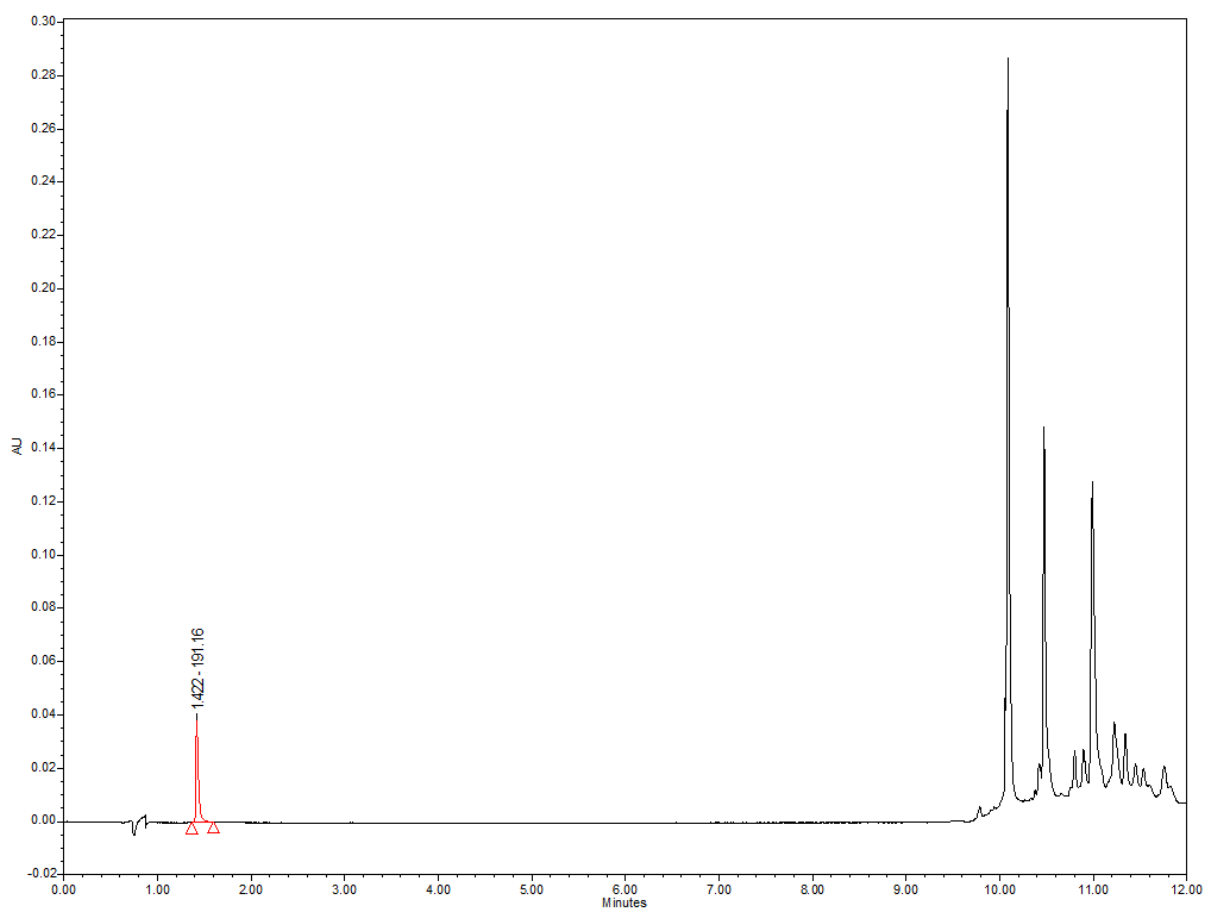
**Peak Separation (m/z)**

**Max Intensity**

**B**

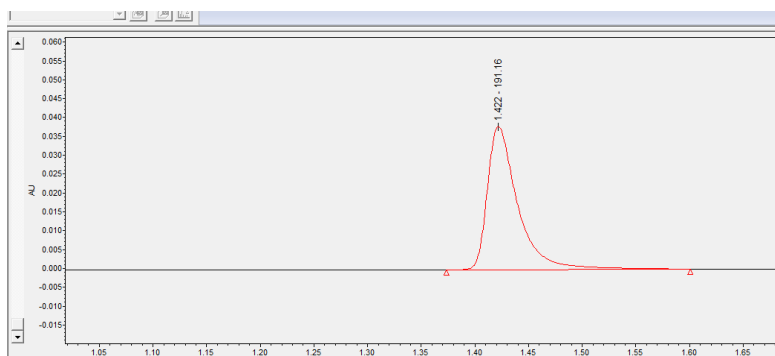
Retention Time (min)	Scan Number	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (m/z)	Max Intensity	B
1.316	789	Extracted	Baseline Correction: Off (1.31:1.32)	Average	1	0.000	1.0000	5414571.200000	16

# 1-1 LC PDA Detector Data with Integrated Peak

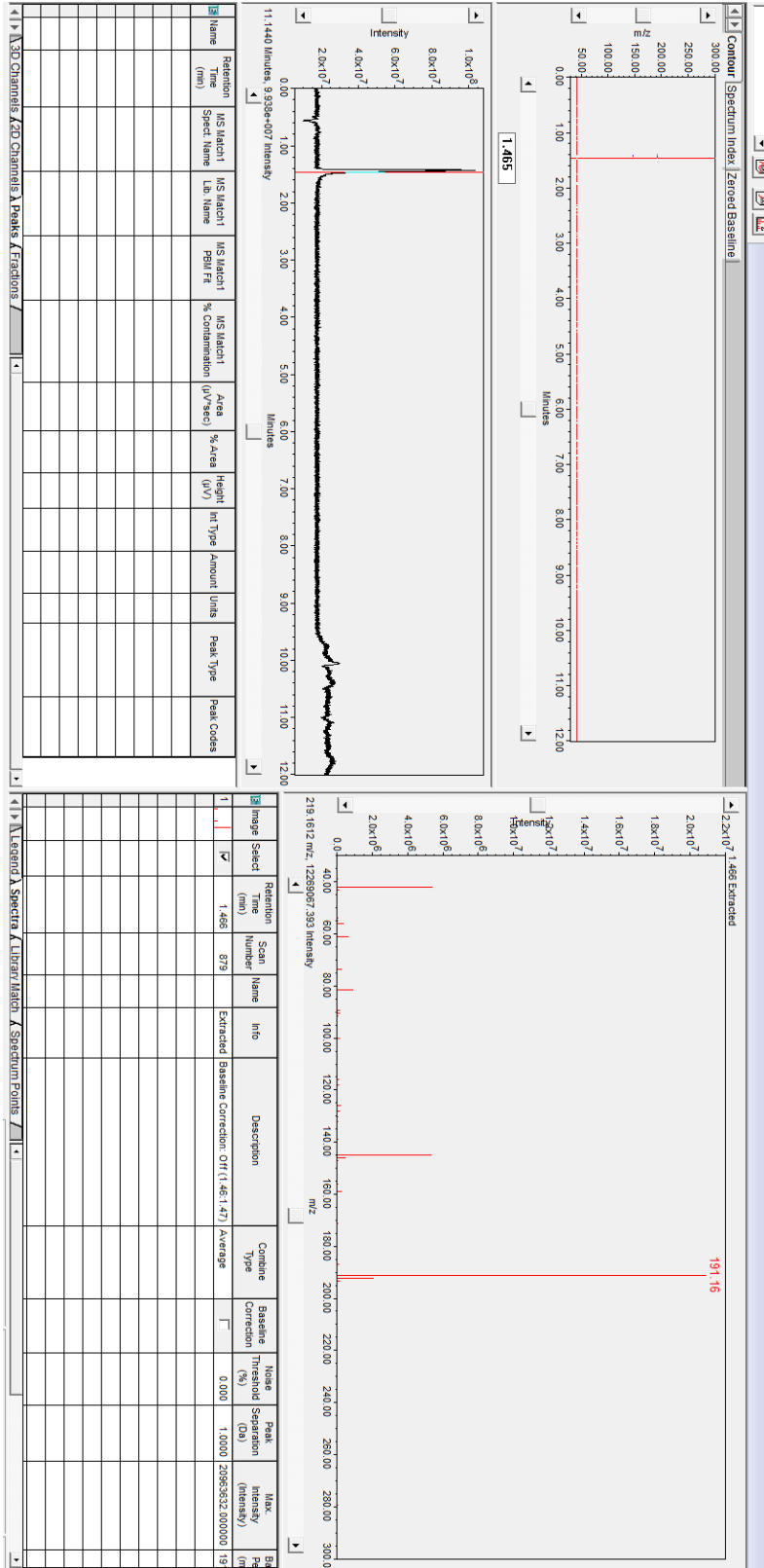


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.422	74917	100.00	37971	bb			Unknown

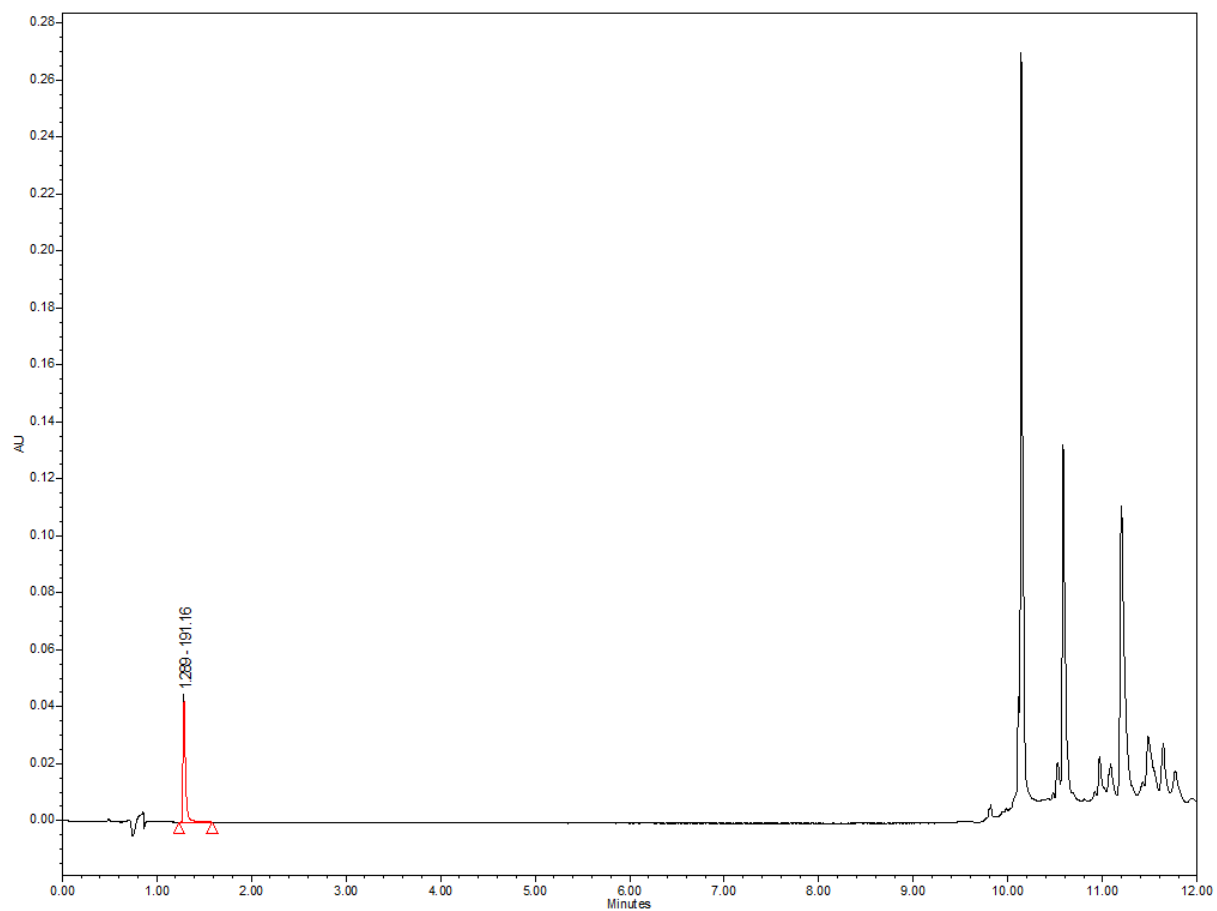
Zoom in of integrated peak shown above:



## 1-1 Mass Spectrum

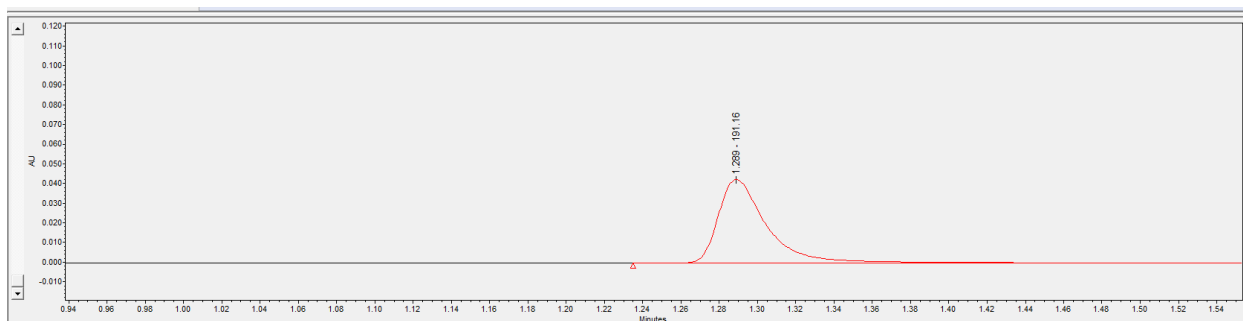


# 1-1 UV-A LC PDA Detector Data with Integrated Peak

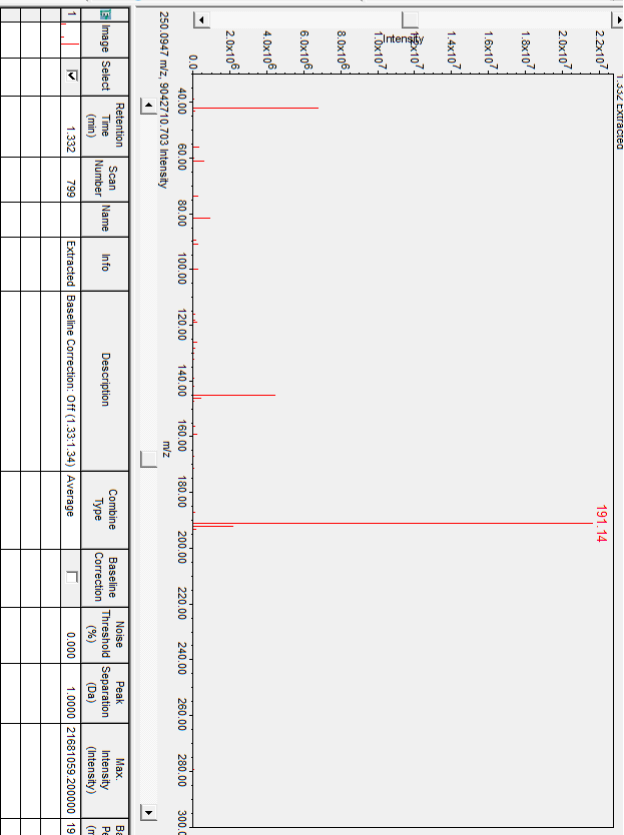
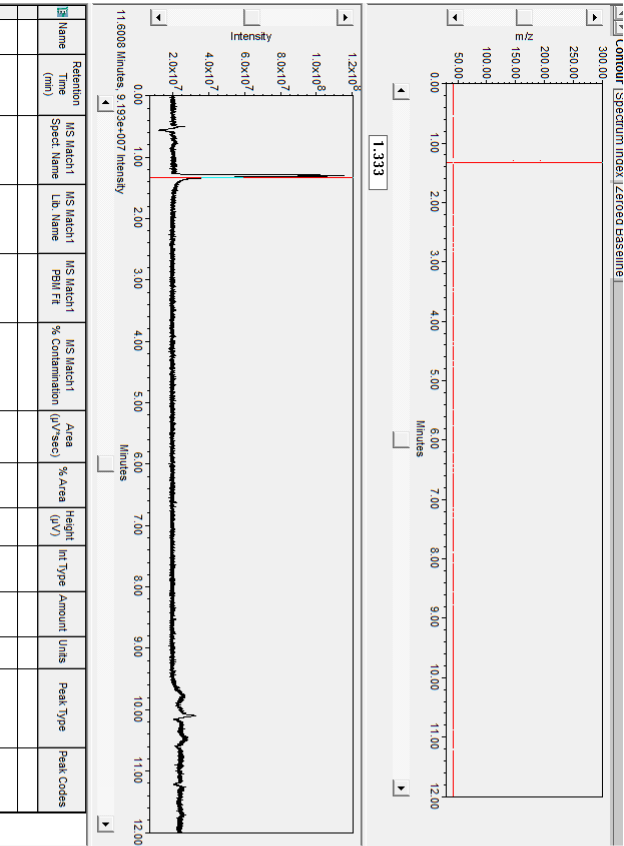


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.289	76760	100.00	42543	bb			Unknown

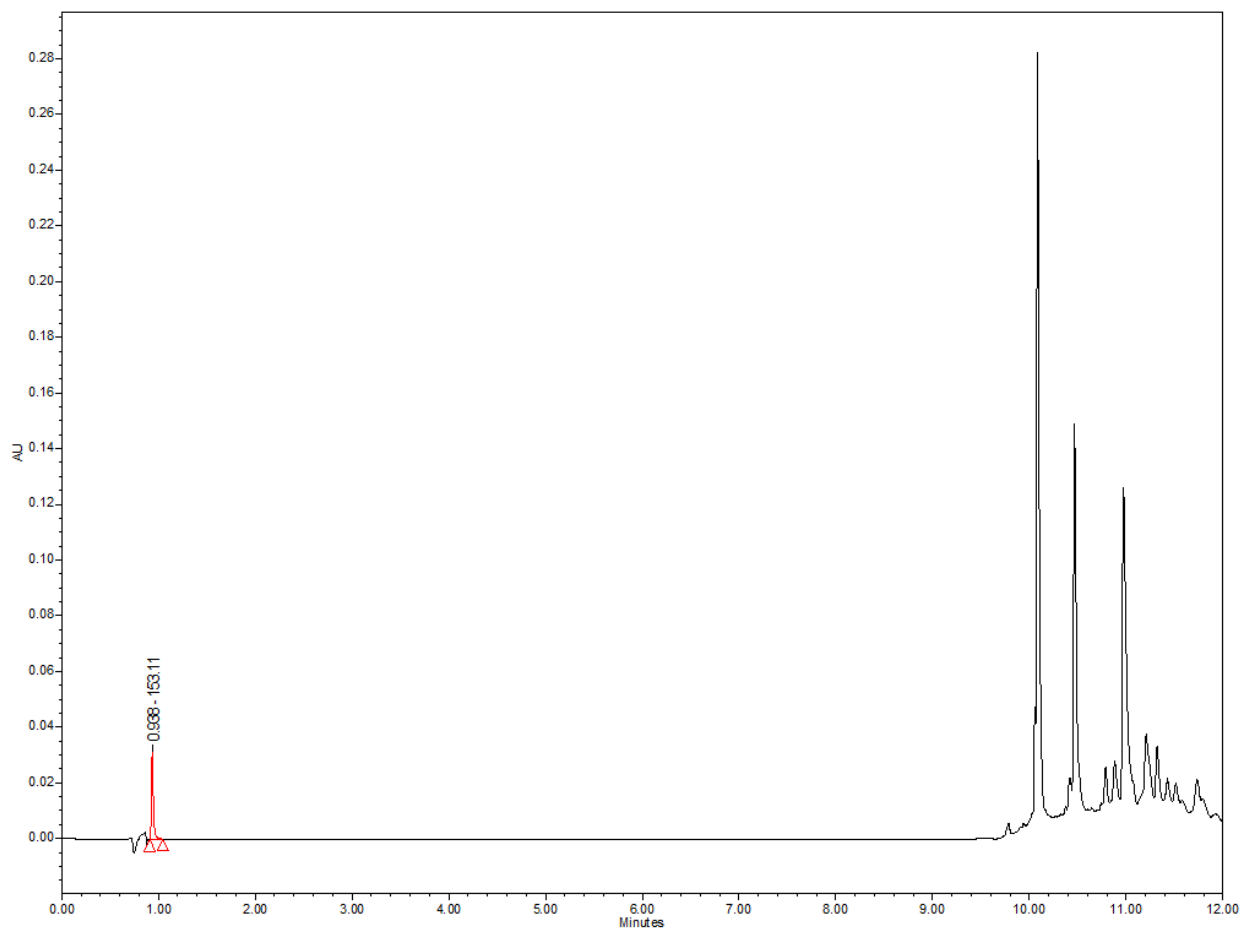
Zoom in of integrated peak shown above:



1-1 UVA Mass Spectrum

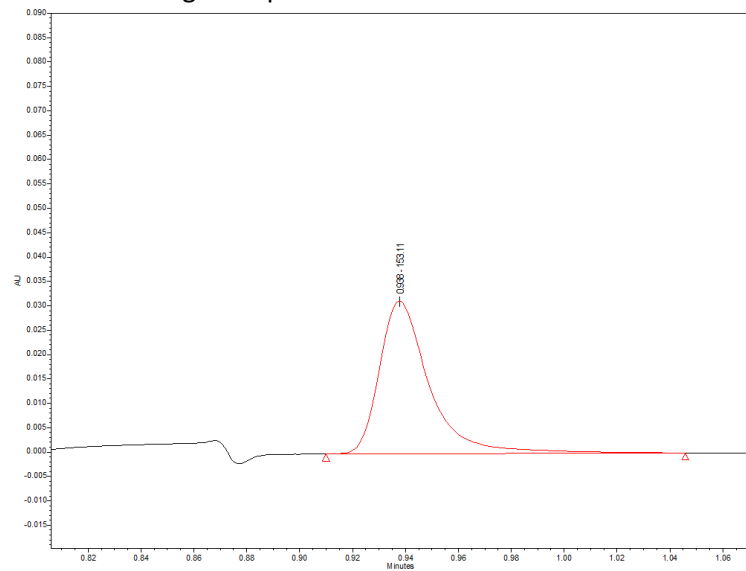


### 1-3 LC PDA Detector Data with Integrated Peak

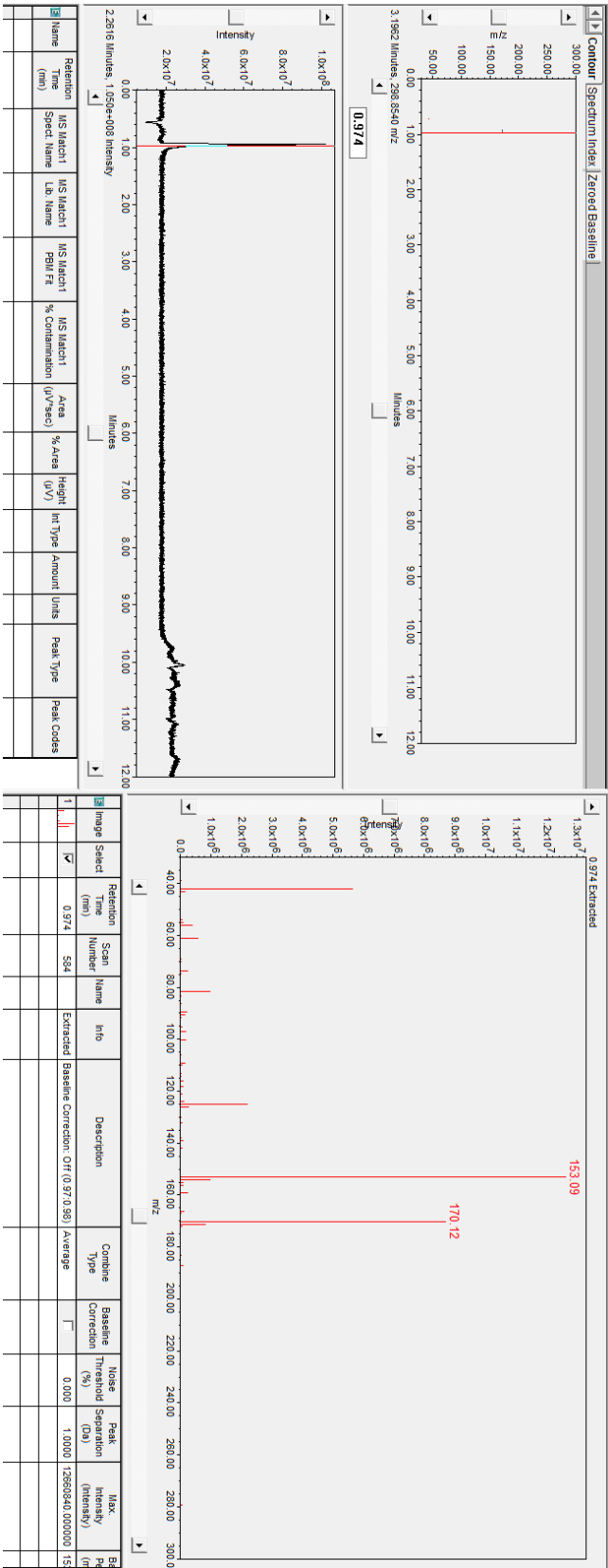


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.938	41162	100.00	31358	bb			Unknown

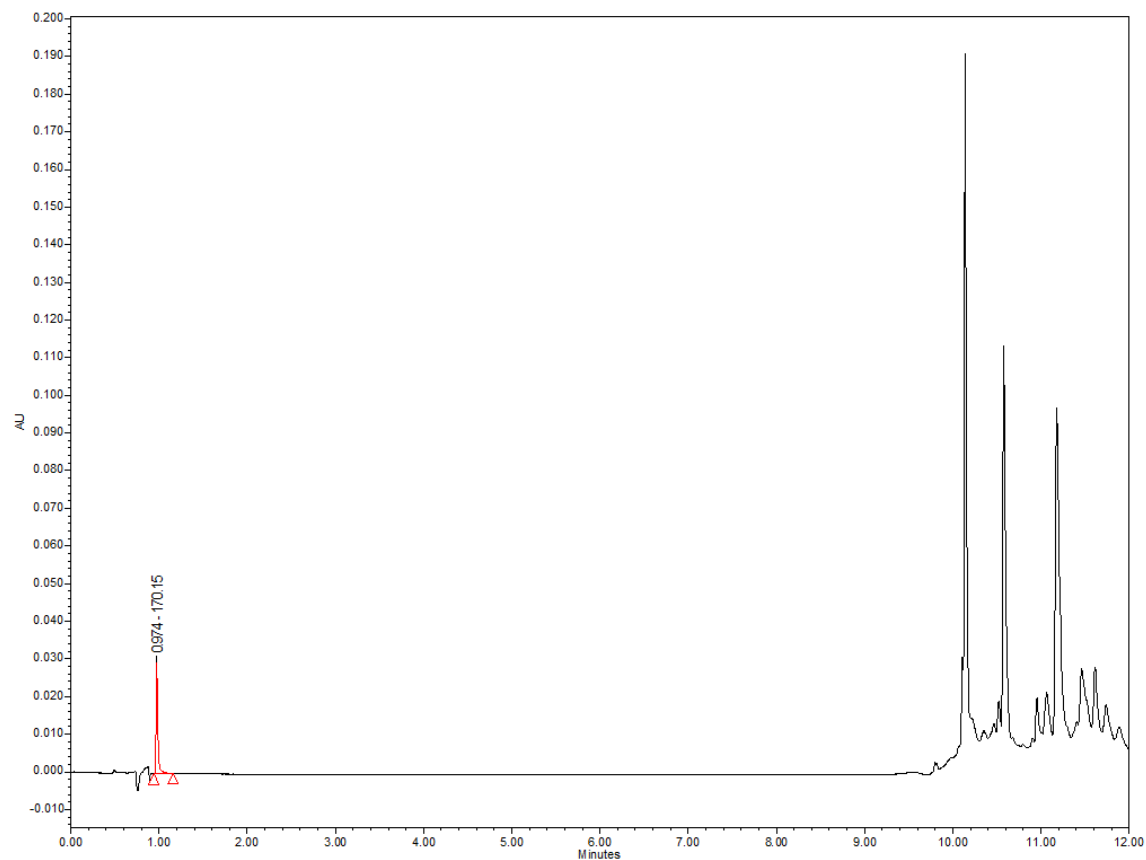
Zoom in of integrated peak shown above:



1-3 Mass Spectrum

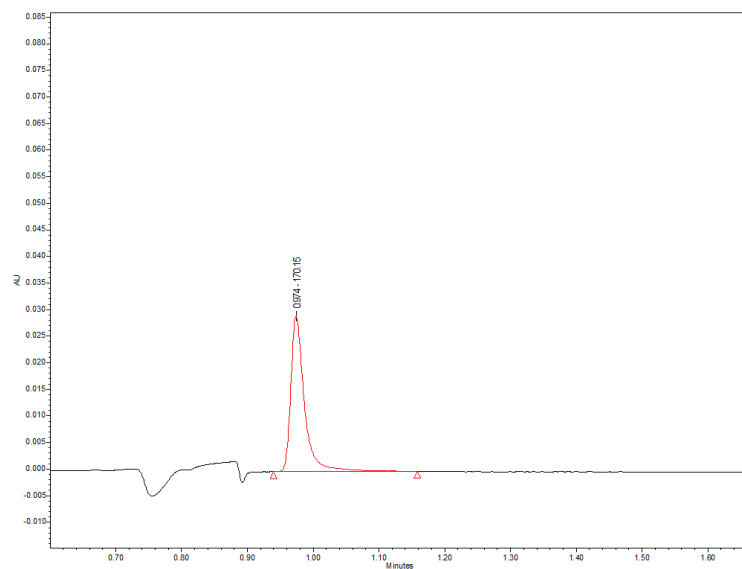


### 1-3 UV-A LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.974	42426	100.00	29318	bb			Unknown

Zoom in of integrated peak shown above:





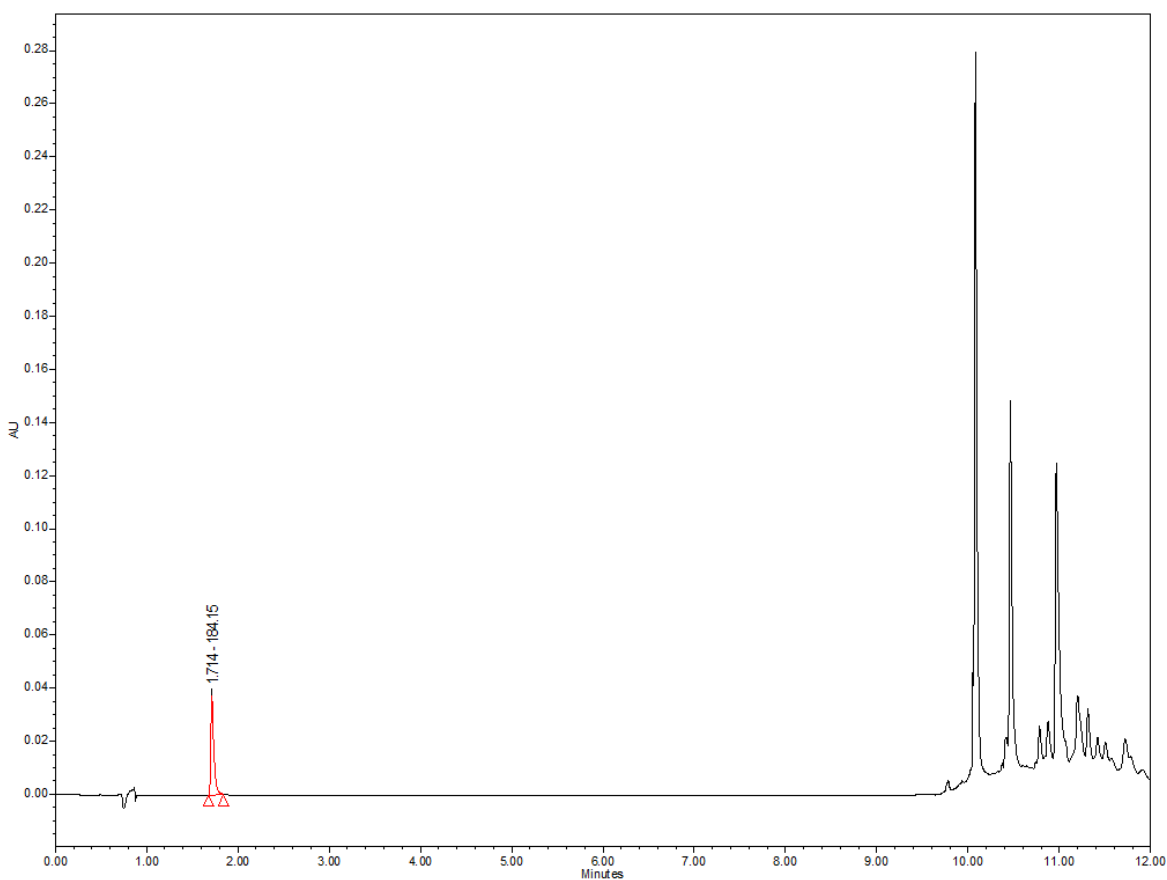
**Contour Spectrum Index, Zeroed Baseline**

1.2683 Minutes, 2.57e+007 Intensity

**0.990 Extracted**

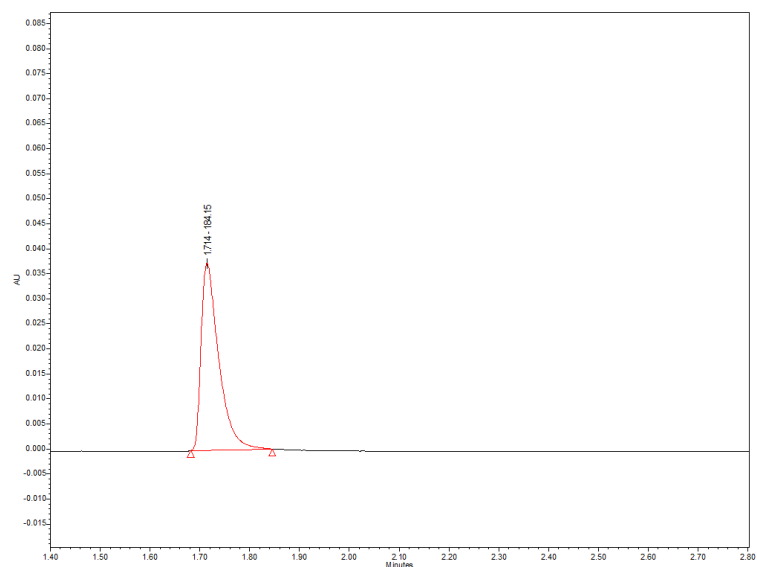
Retention Time (min)	Scan Number	Info	Description	Combine Type	Baseline Correction	Noise Threshold (7s)	Peak Separation (dS)	Max. Intensity (Intensity)
0.990	594	Extracted	Baseline Correction Off (0.990.990)	Average	<input type="checkbox"/>	0.000	1.0000	35.8228.000000

# 1-6 LC PDA Detector Data with Integrated Peak

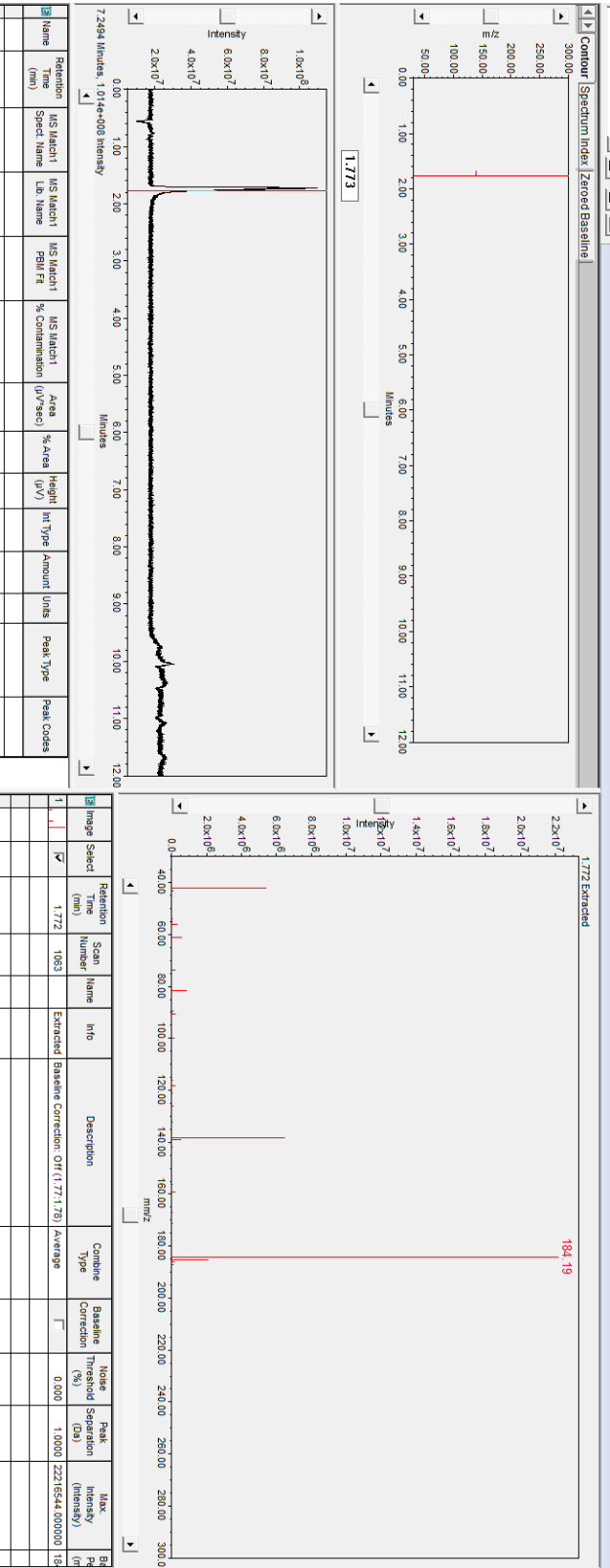


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.714	91863	100.00	37449	bb			Unknown

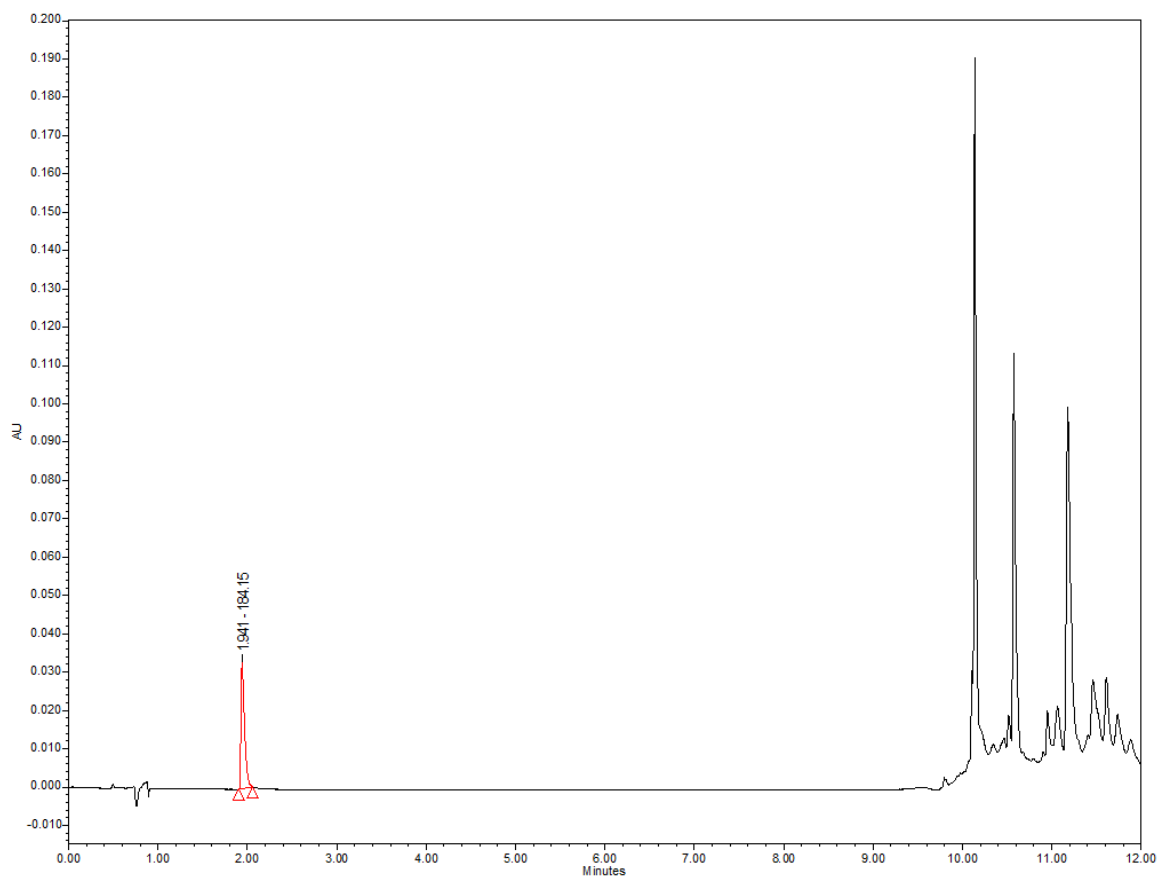
Zoom in of integrated peak shown above:



1-6 Mass Spectrum

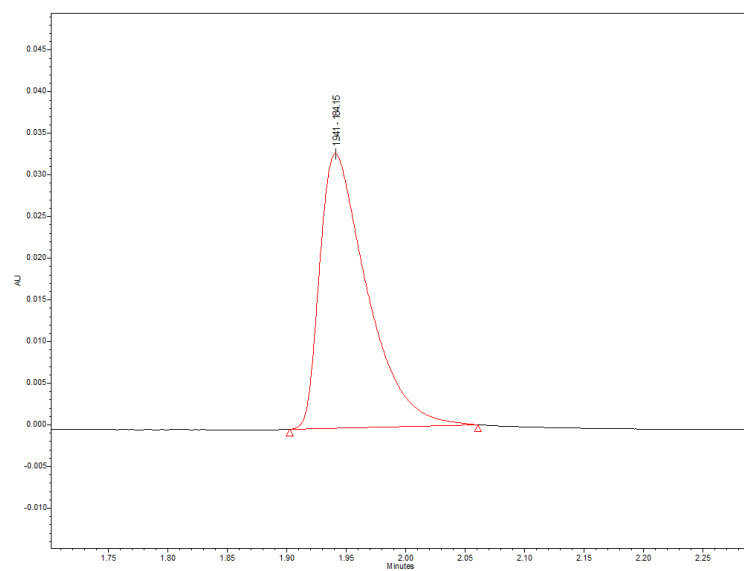


# 1-6 UV-A LC PDA Detector Data with Integrated Peak

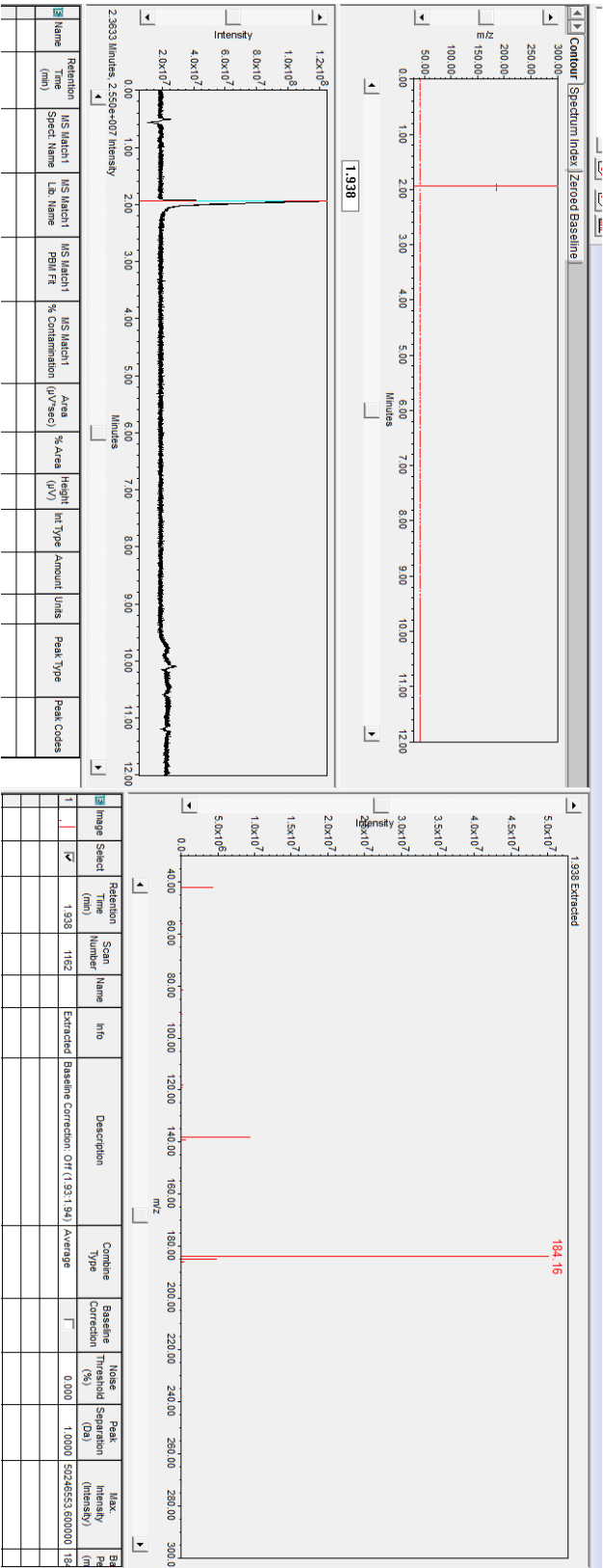


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.941	91434	100.00	32967	bb			Unknown

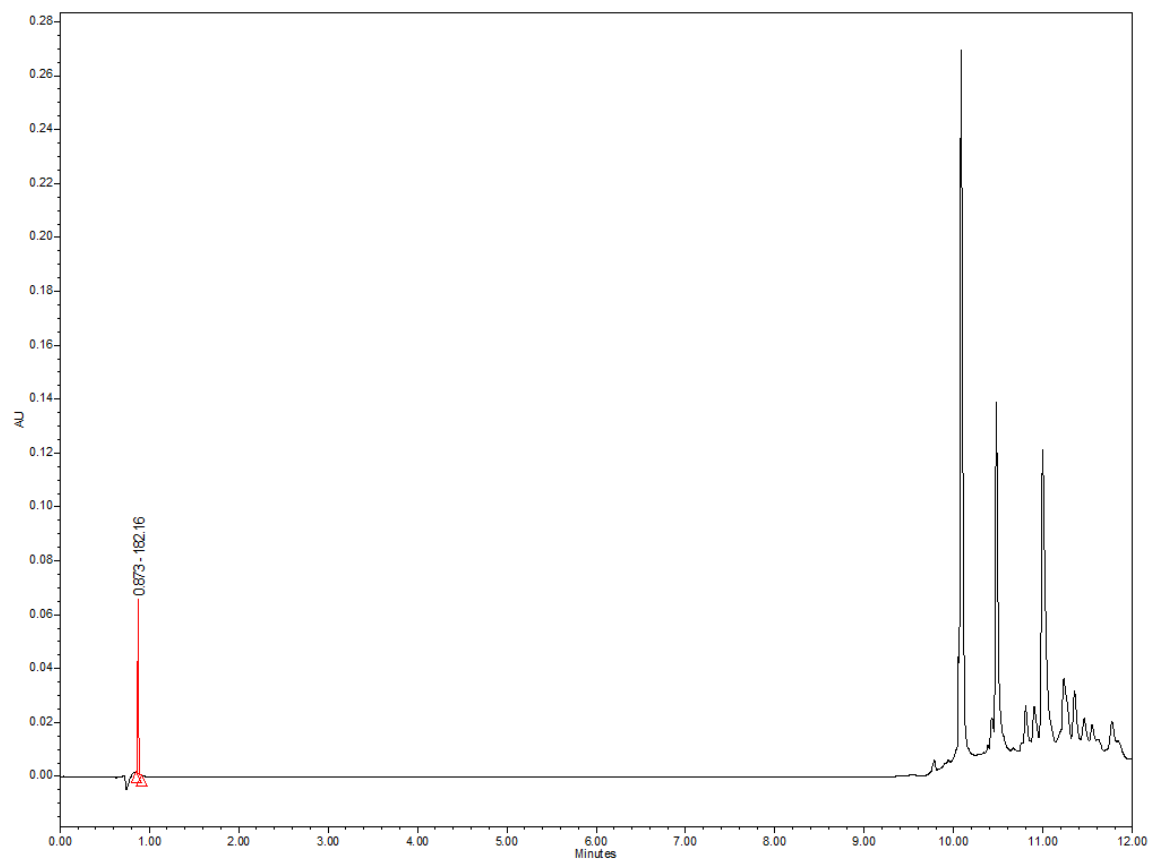
Zoom in of integrated peak shown above:



1-6 UVA Mass Spectrum

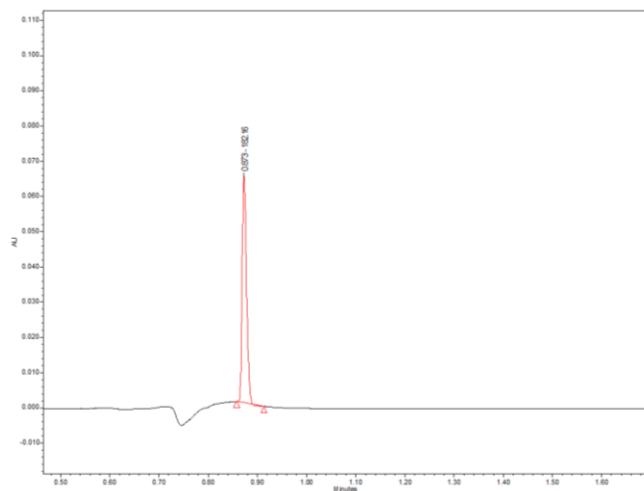


## 2 LC PDA Detector Data with Integrated Peak



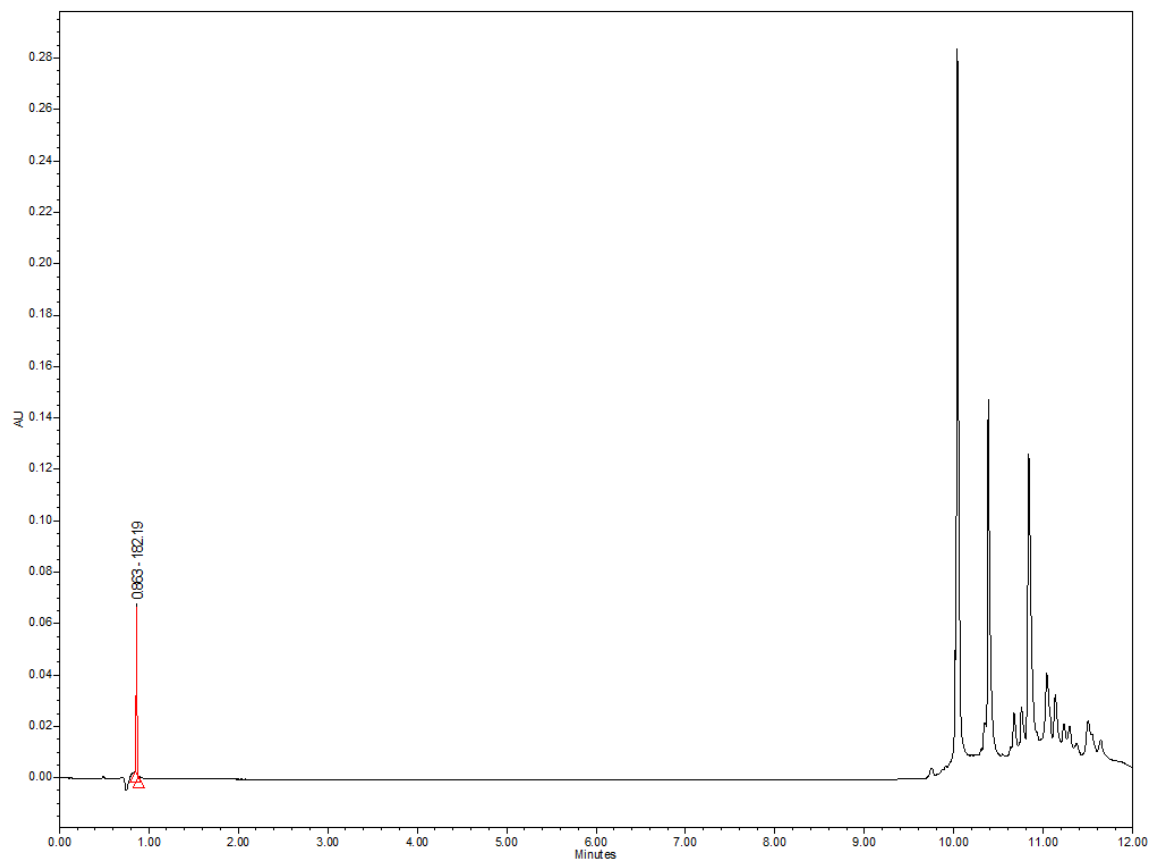
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.873	40039	100.00	64255	bb			Unknown

Zoom in of integrated peak above:



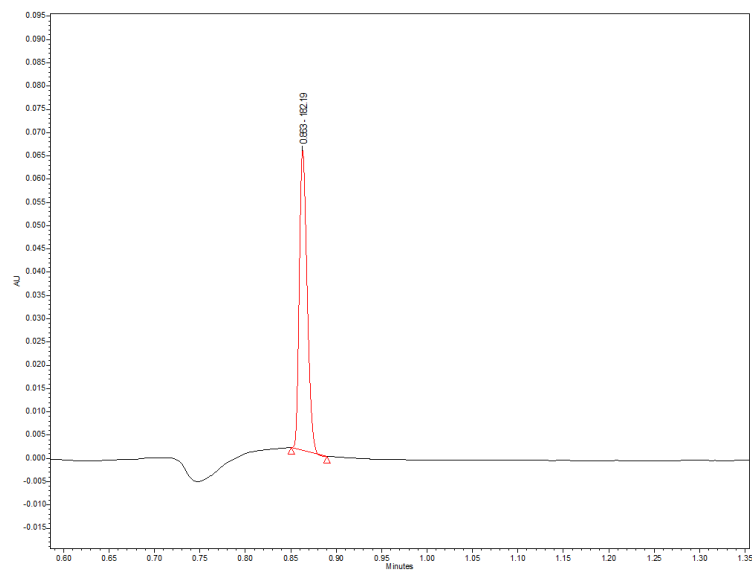
[illegible]

## 2 UV-A LC PDA Detector Data with Integrated Peak



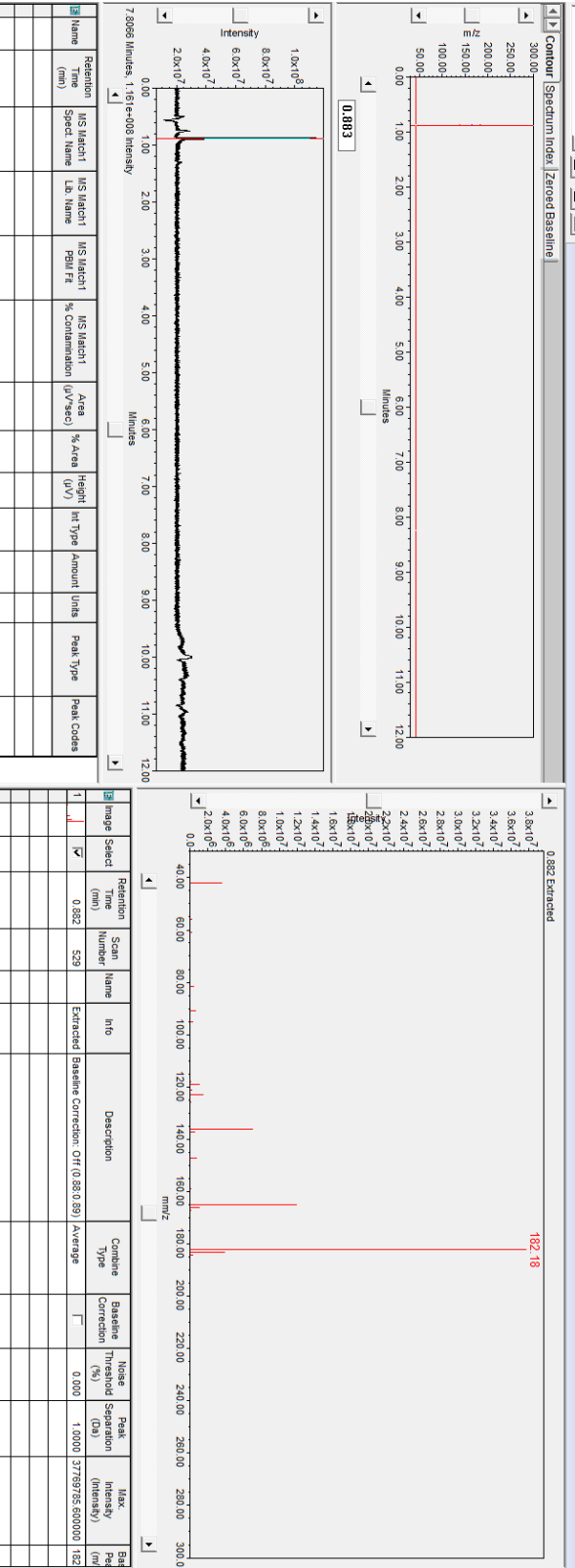
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.863	39363	100.00	64509	bb			Unknown

Zoom in of integrated peak shown above:

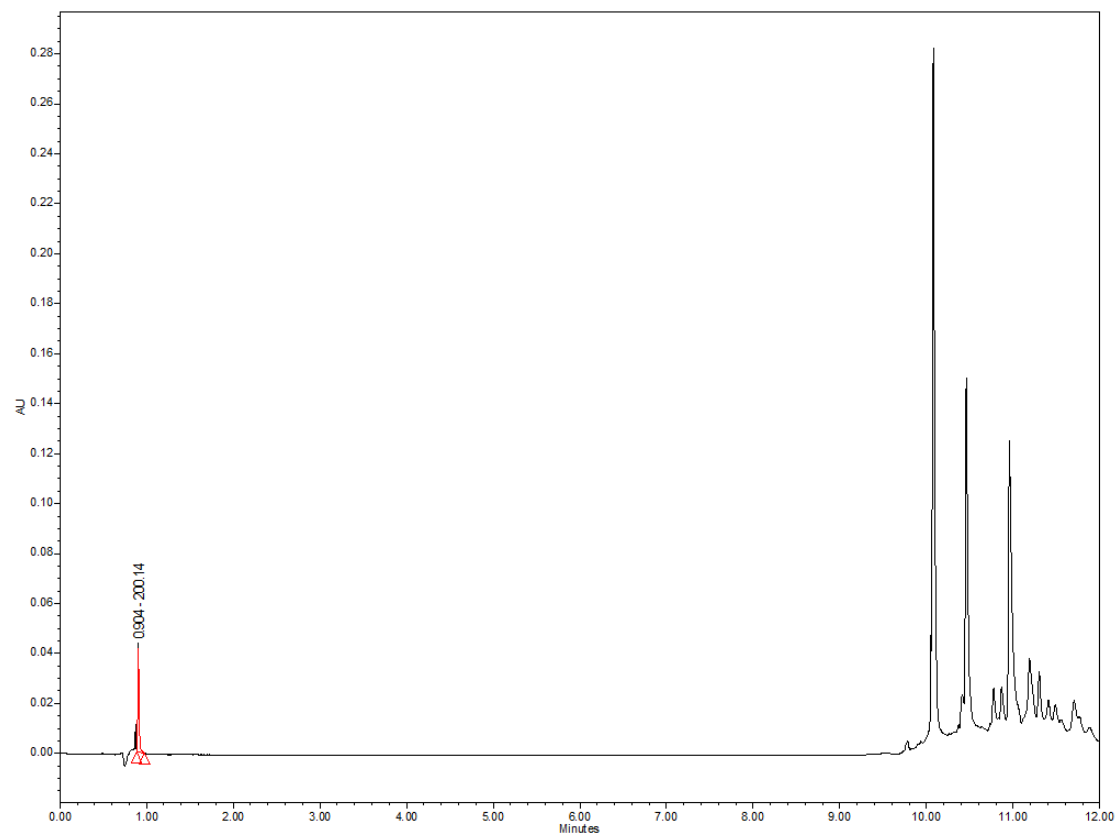




2 UVA Mass Spectrum

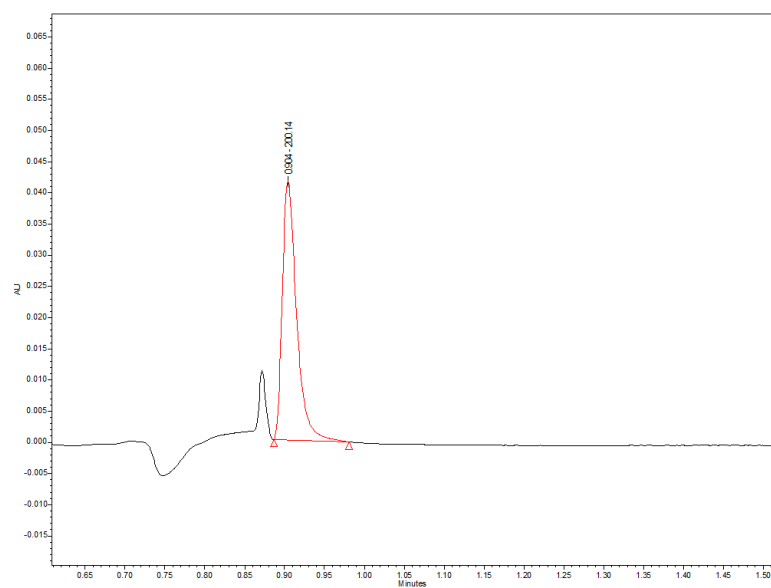


## 2-1 LC PDA Detector Data with Integrated Peak



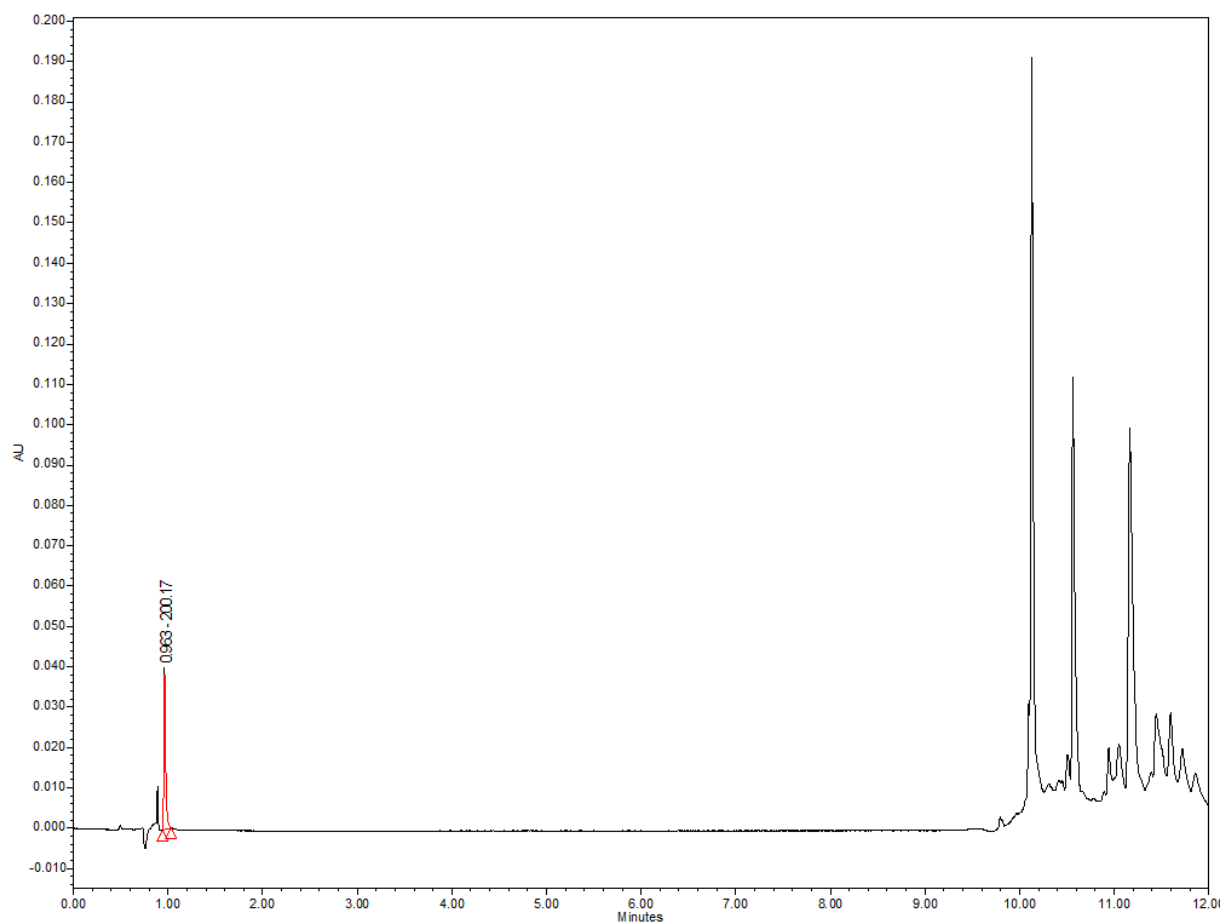
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.904	51406	100.00	41438	bb			Unknown

Zoom in of integrated peak shown above:



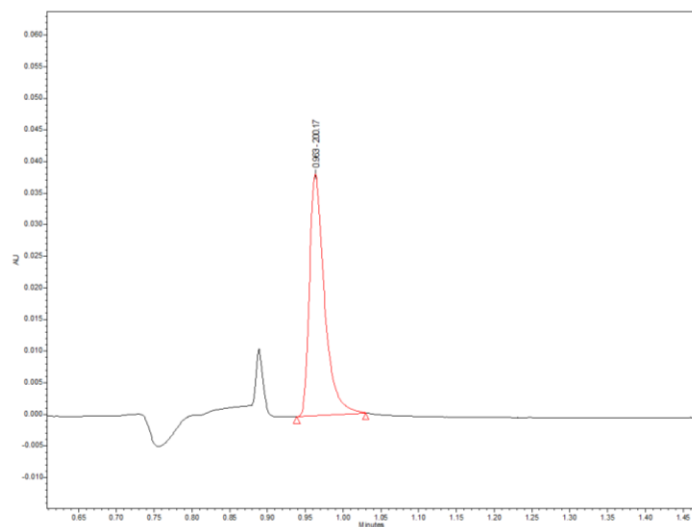
[illegible]

## 2-1 UV-A LC PDA Detector Data with Integrated Peak



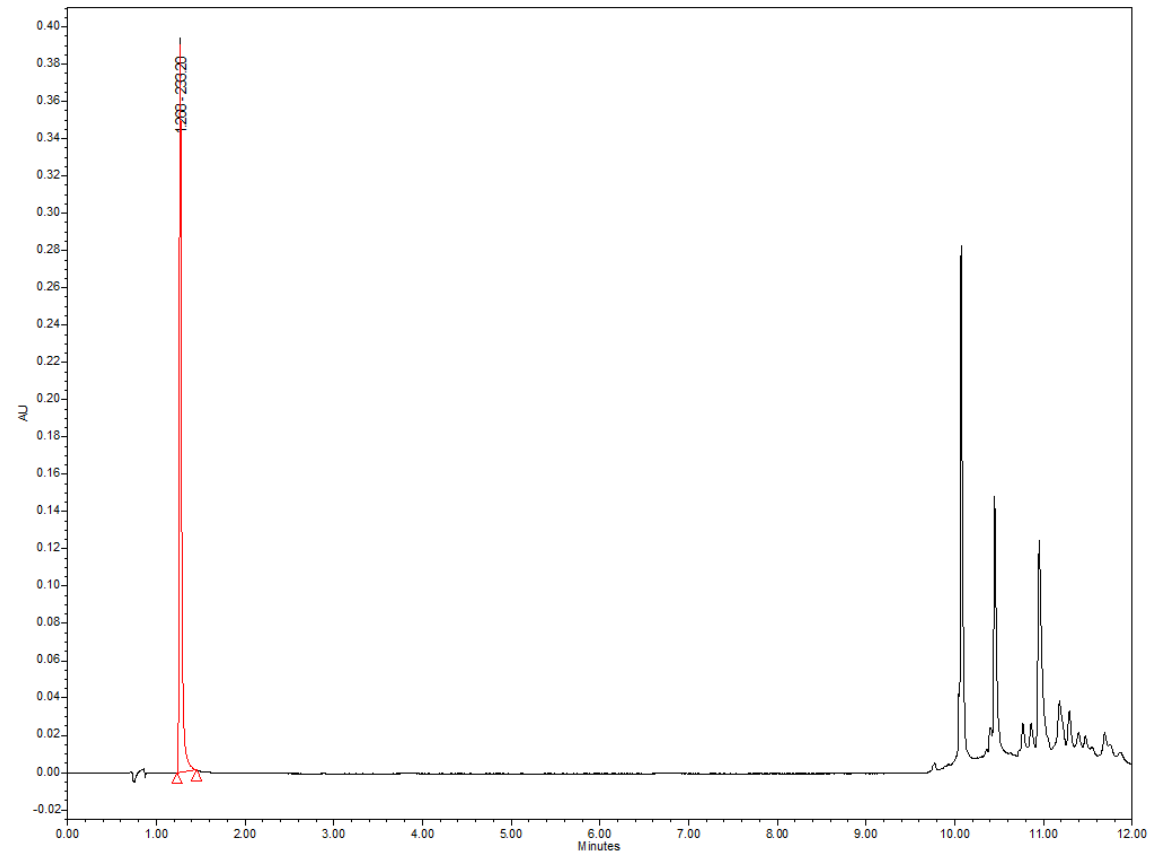
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.963	52738	100.00	38233	bb			Unknown

Zoom in of integrated peak shown above:



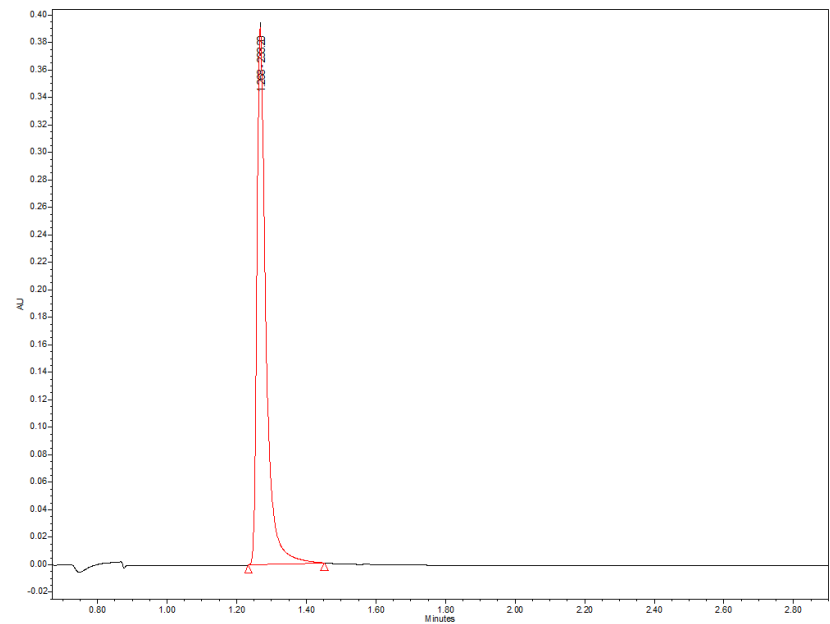
[illegible]

4 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.268	709654	100.00	390718	bb			Unknown

Zoom in of integrated peak shown above:



**Chromatogram Data:**

Retention Time (min)	Intensity
1.316	~8.0 x 10 <sup>7</sup>

**Mass Spectrum Data:**

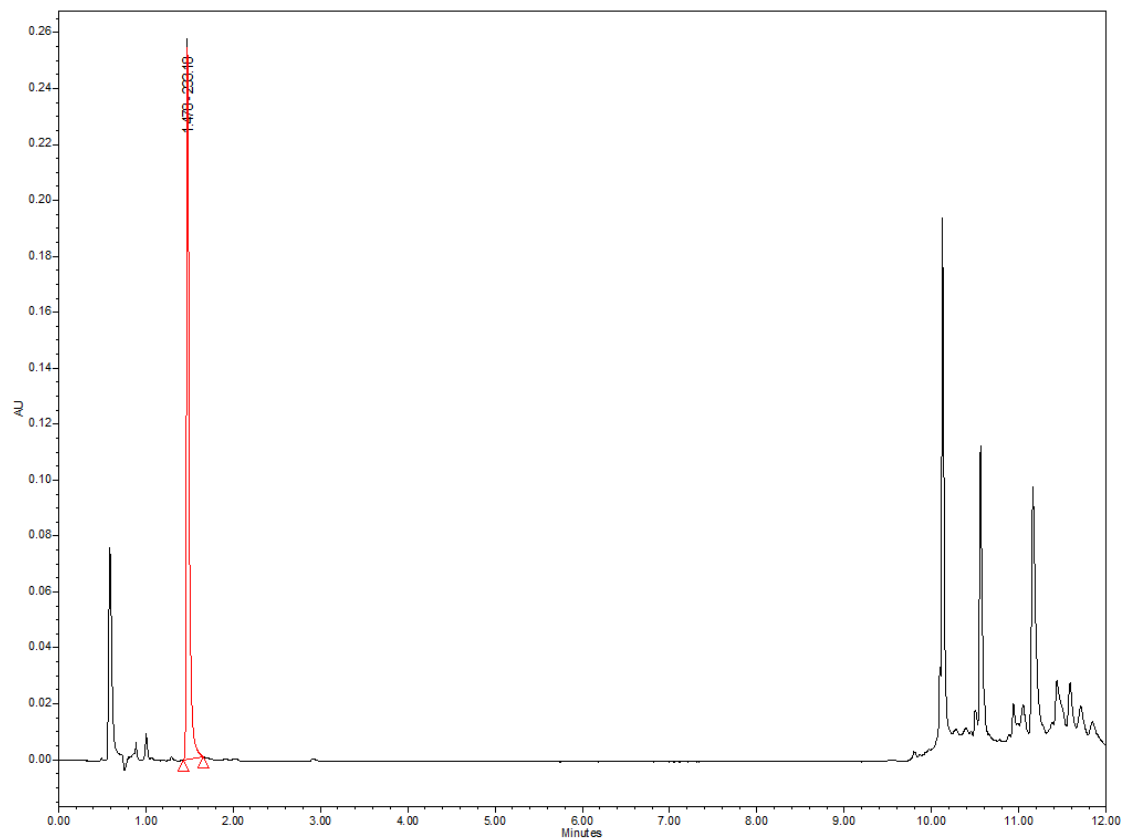
m/z	Intensity
233.19	~2.0 x 10 <sup>8</sup>

**Peak Data Table:**

Peak Number	Retention Time (min)	Scan Number	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Std. Dev. (Intensity)
1	1.316	789	Extracted	Baseline Correction Off (1.311-1.321)	Average	<input type="checkbox"/>	0.000	1.0000	16179530.40000	233.19

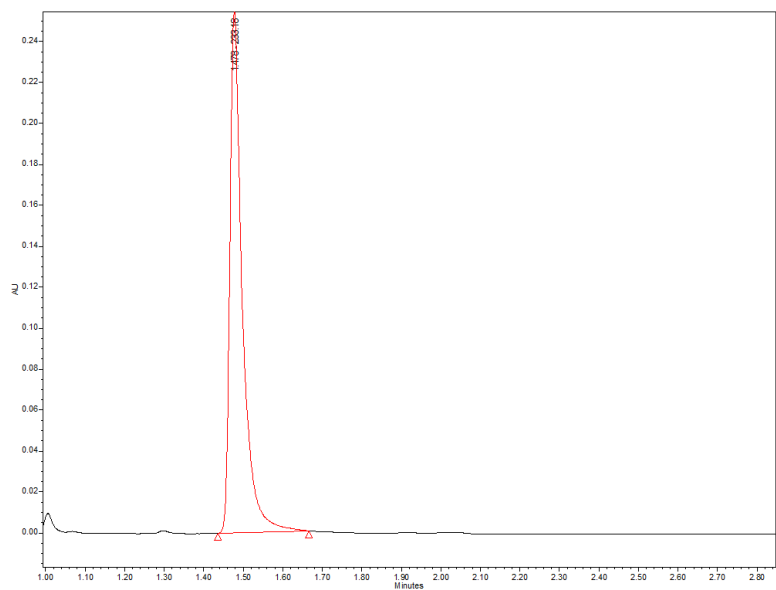
[illegible]

4 UV-A LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.478	559742	100.00	254738	bb			Unknown

Zoom in of integrated peak above:





**Contour** Spectrum Index Zeroed Baseline

Intensity

Minutes

1.496

1.751 Minutes, 1.987e+07 Intensity

Name	Retention Time (min)	MS Inact1 Spect. Name	MS Inact1 Lib. Name	MS Inact1 PDB Ref	MS Inact1 % Contamination	Area (uV*sec)	% Area	Height (uV)	Int. Type	Amount	Units	Peak Type	Peak Codes
1	1.496												

**1.495 Extracted**

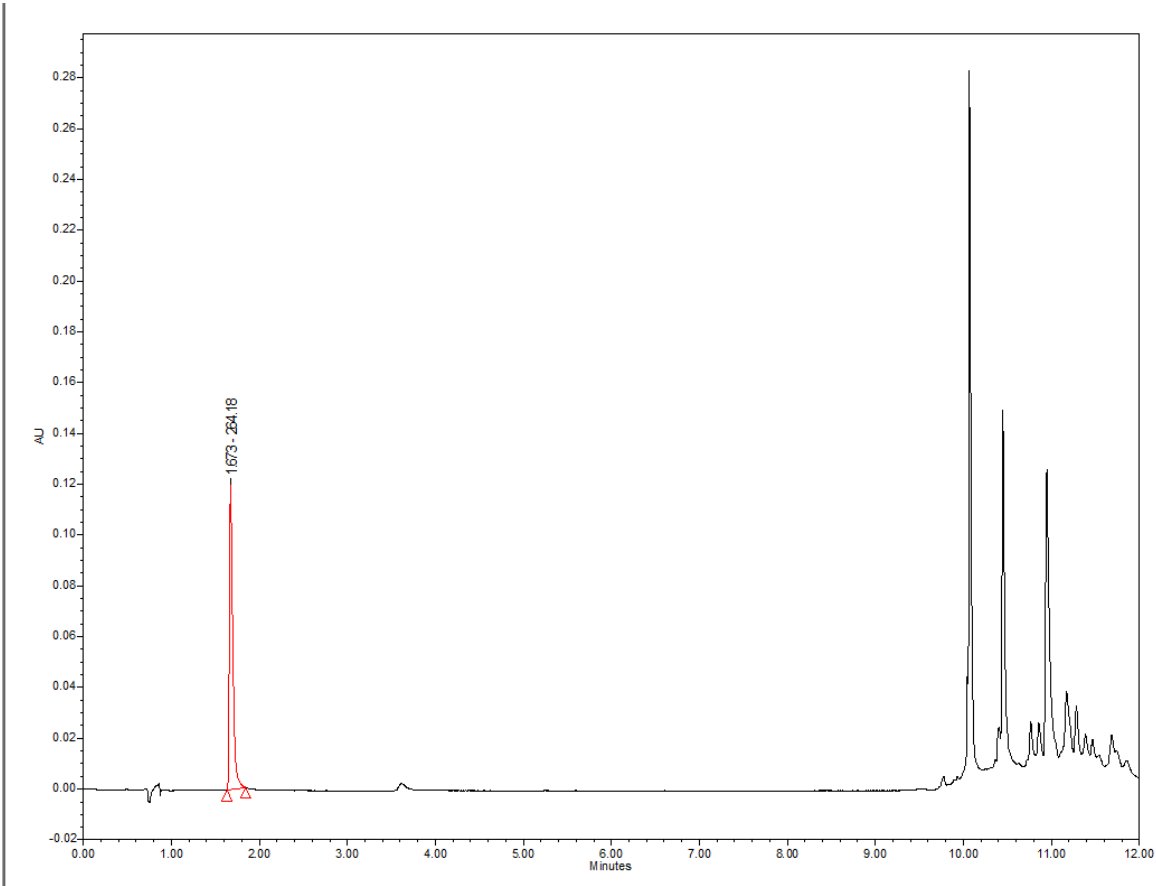
Intensity

m/z

233.19

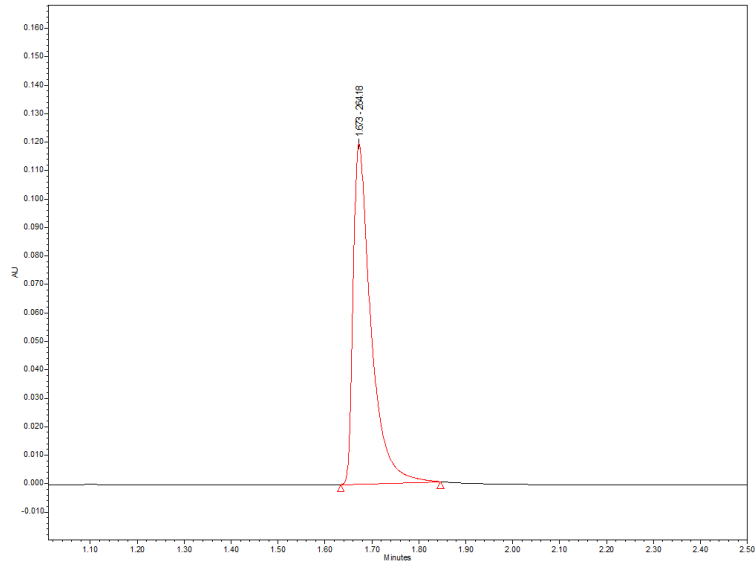
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max Intensity (Intensity)	Base Peak (m/z)
1	1	1.495	897		Extracted	Baseline Correction: Off (1.491, 1.50)	Average		0.000	1.0000	35021382.400000	233

5 LC PDA Detector Data with Integrated Peak

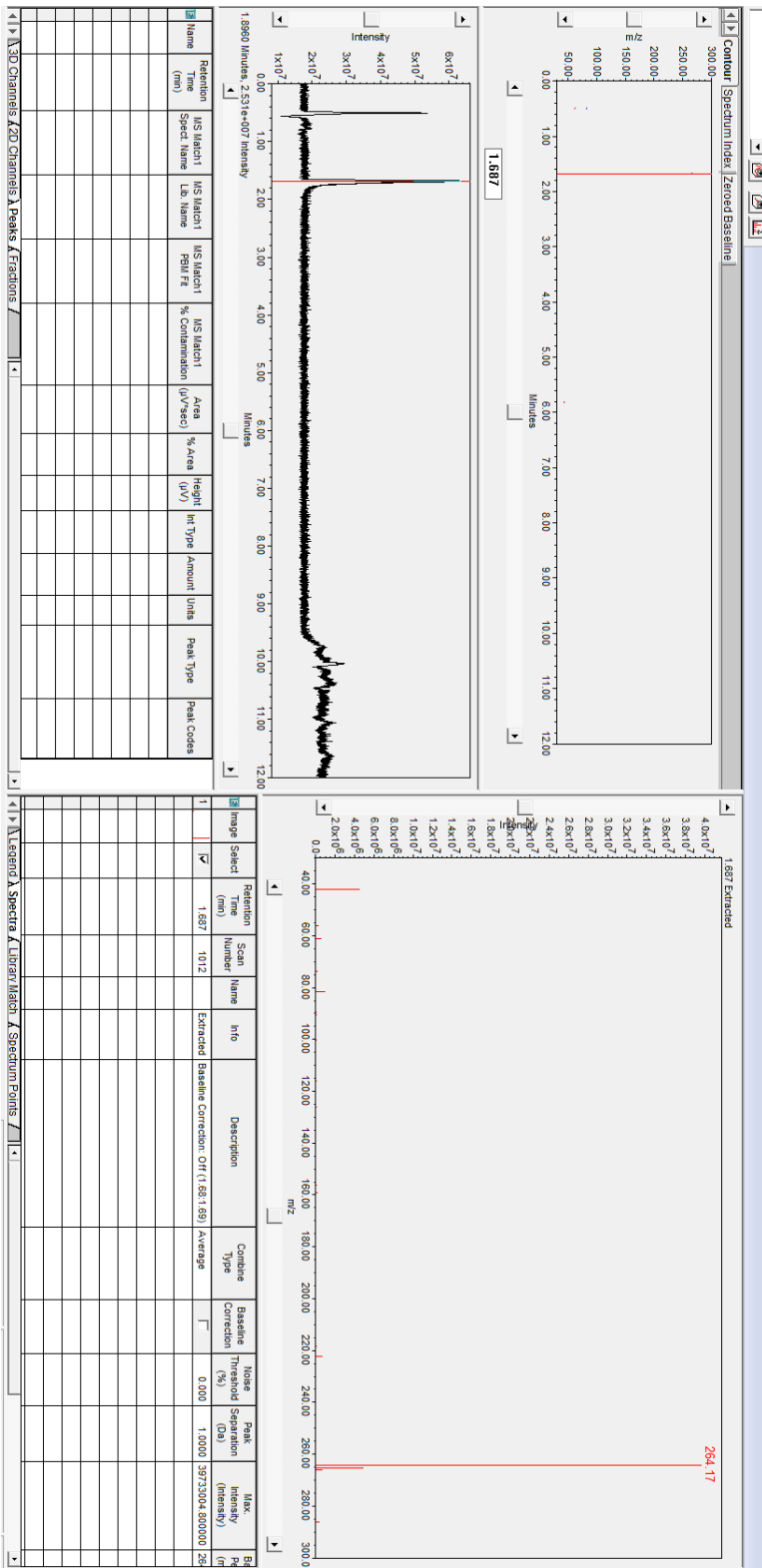


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.673	315272	100.00	119647	bb			Unknown

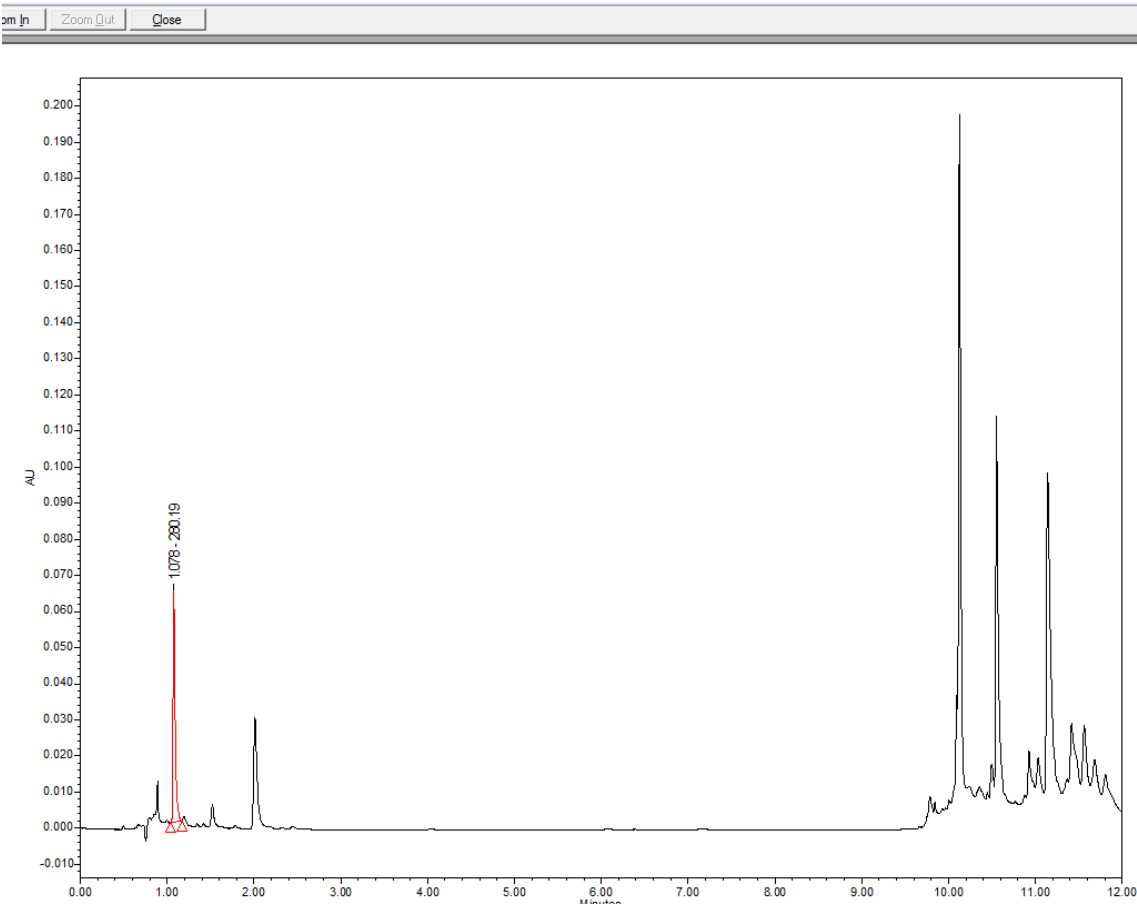
Zoom in of integrated peak shown above:



## 5 Mass Spectrum

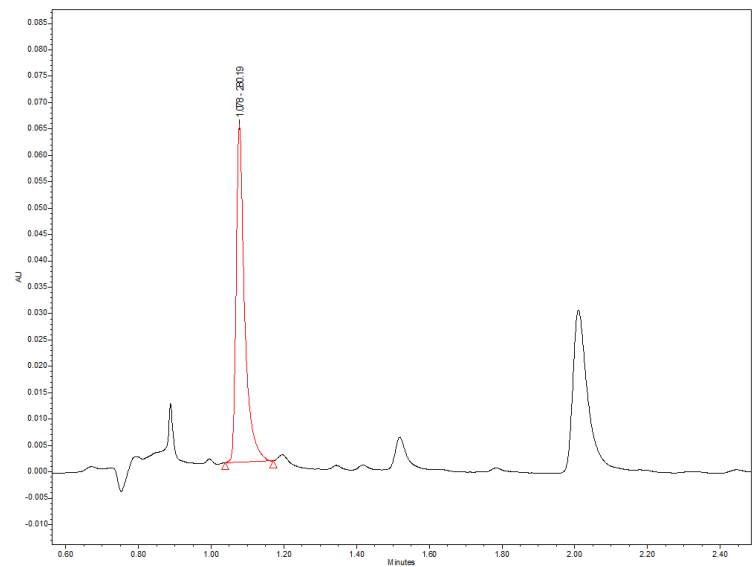


5 UV-A LC PDA Detector Data with Integrated Peak

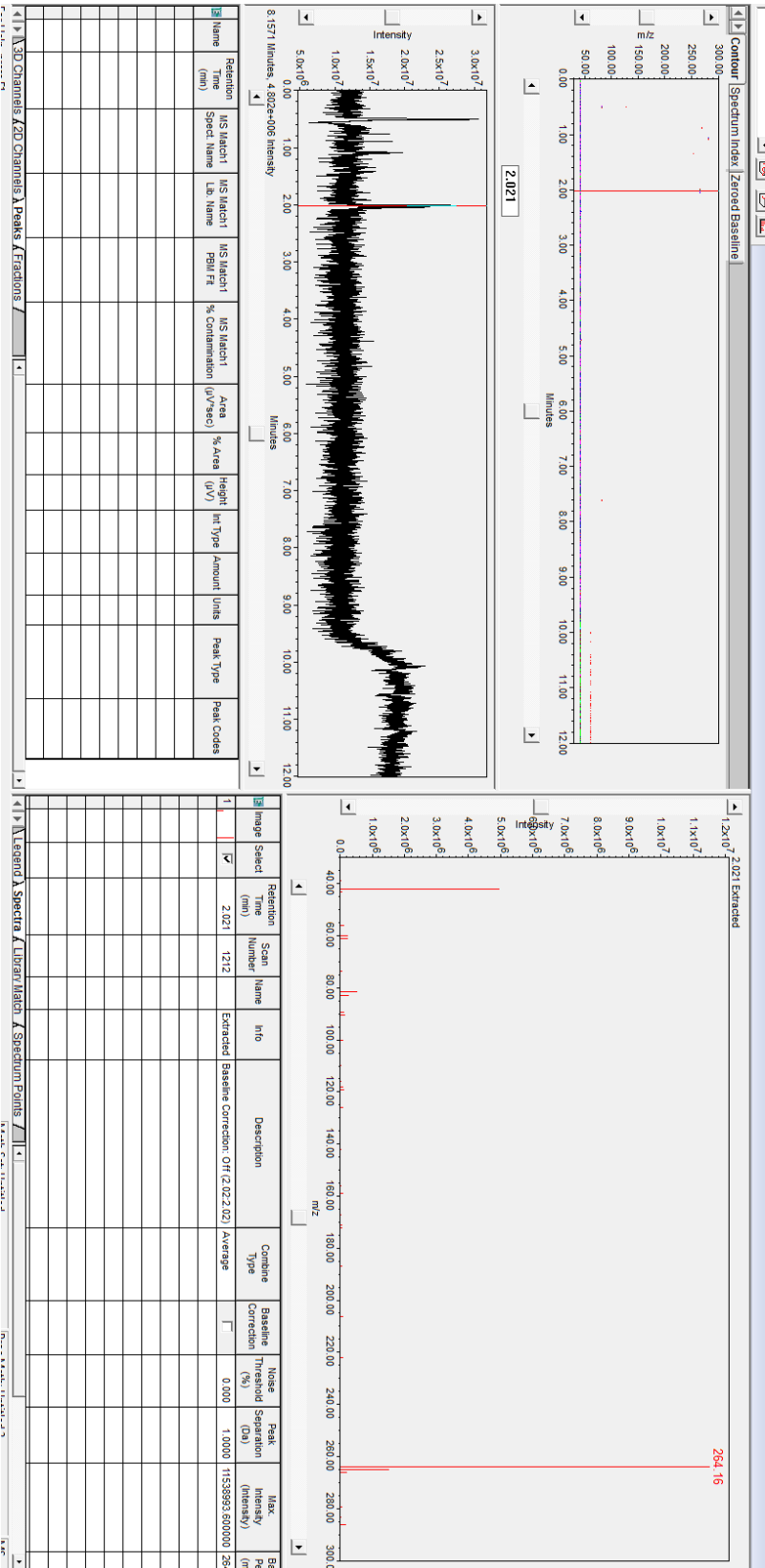


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.078	104654	100.00	64021	bb			Unknown

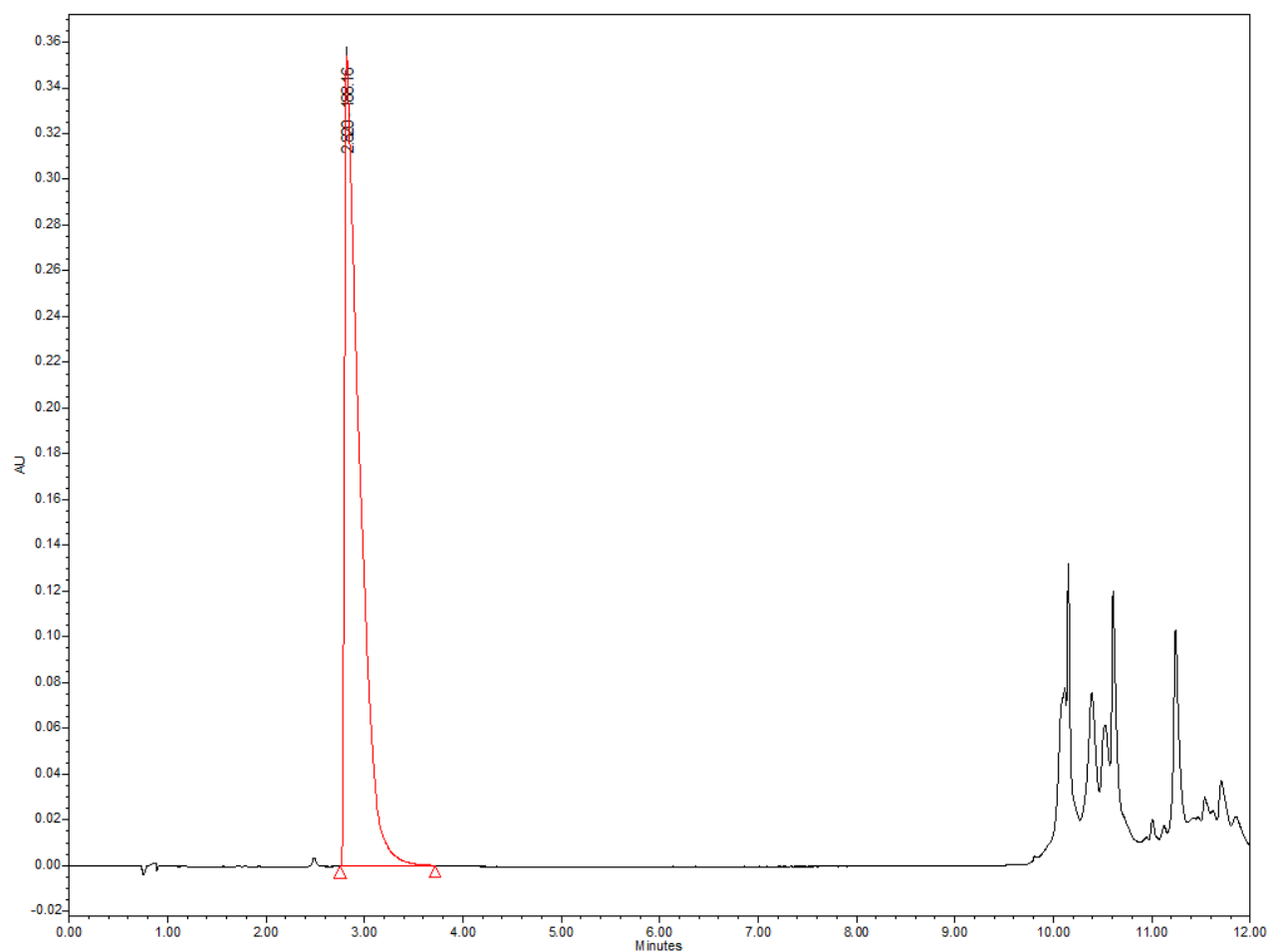
Zoom in on integrated peak shown above:



5 UVA Mass Spectrum



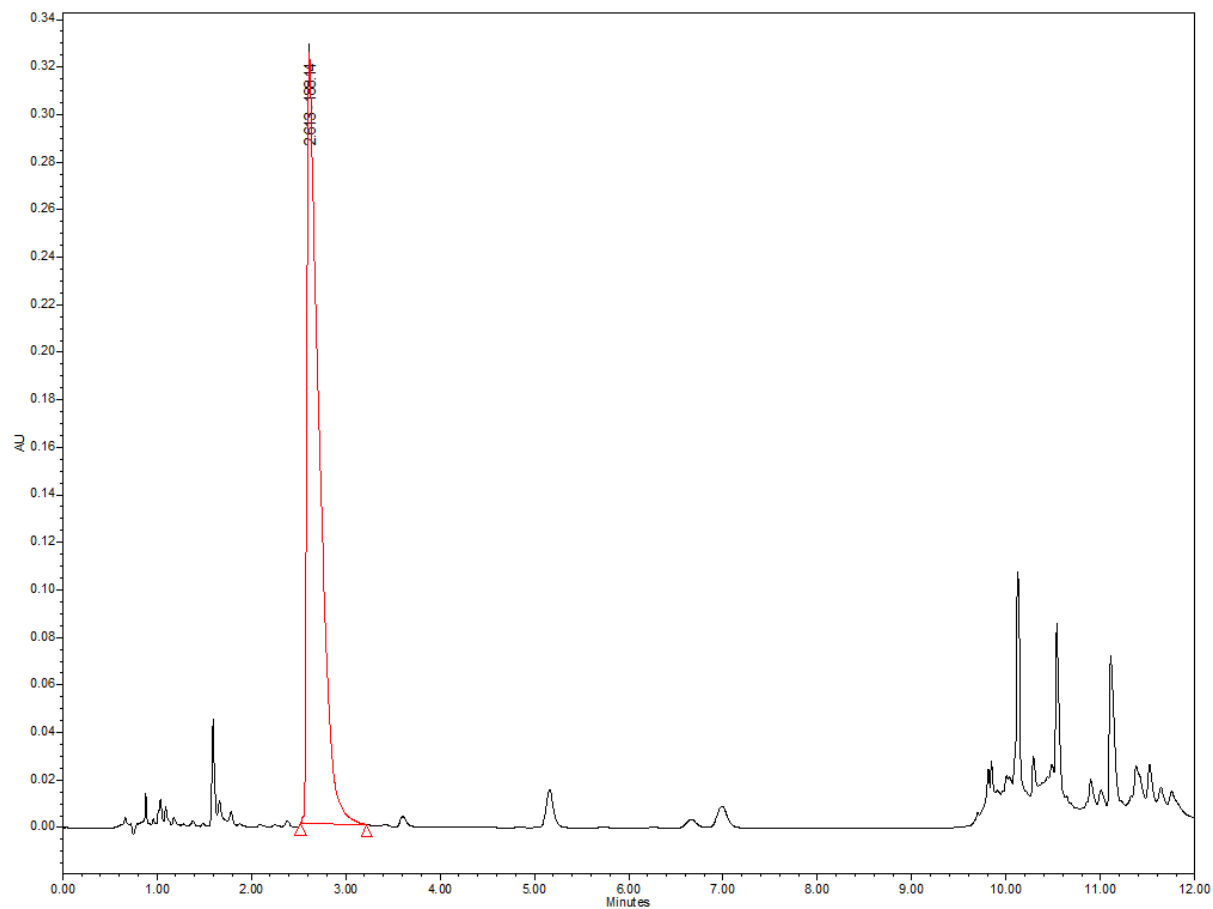
### 3 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.820	3782818	100.00	354418	bb			Unknown

[illegible]

### 3 UV-A LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.613	2938355	100.00	324841	bb			Unknown



**Contour Spectrum Index | Zeroed Baseline**

**Intensity**

**m/z**

**Minutes**

**2.604**

Name	Retention Time (min)	MS Match Spec Name	MS Match Lib Name	MS Match PDB ID	MS Match % Confirmation	Area (μV Sec)	% Area	Height (μV)	Int Type	Amount	Units	Peak Type	Peak Codes

**Legend A Spectra A Library Match A Spectrum Points**

**Relative Intensity**

**m/z**

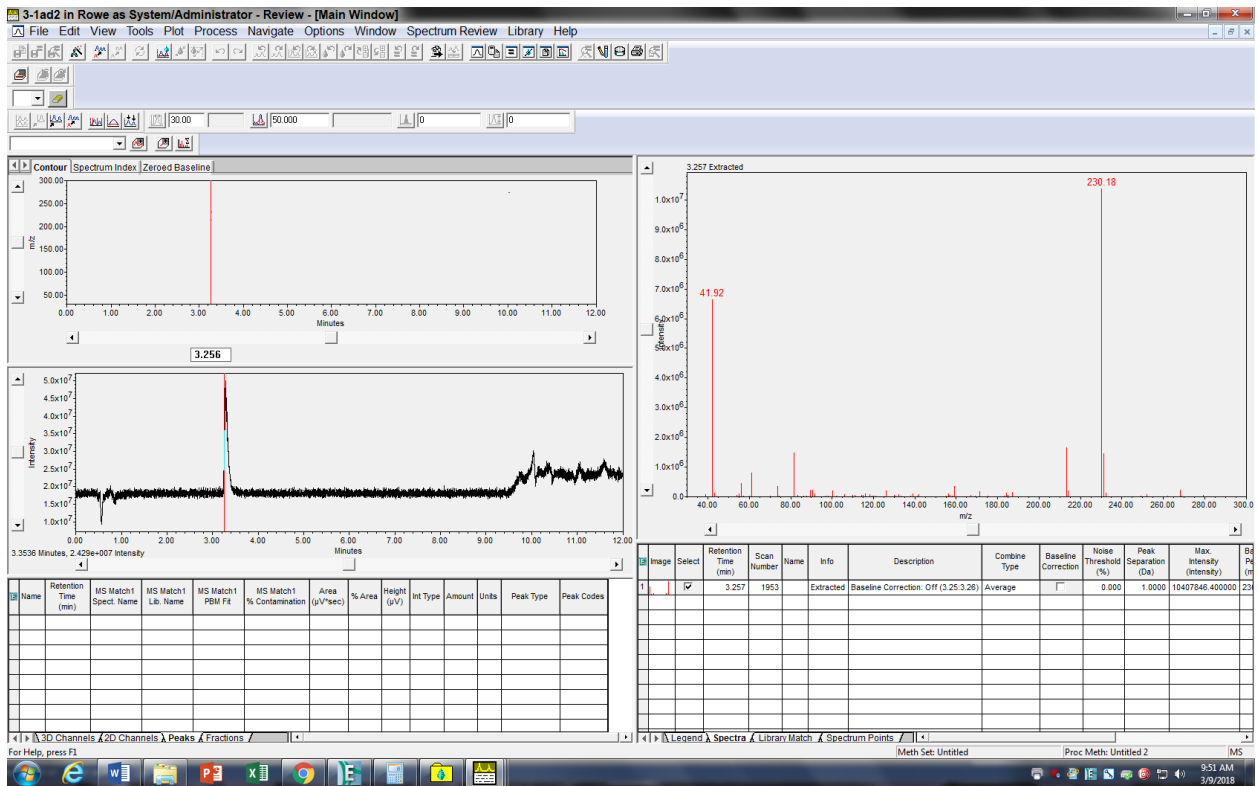
**168.16**

**205.20**

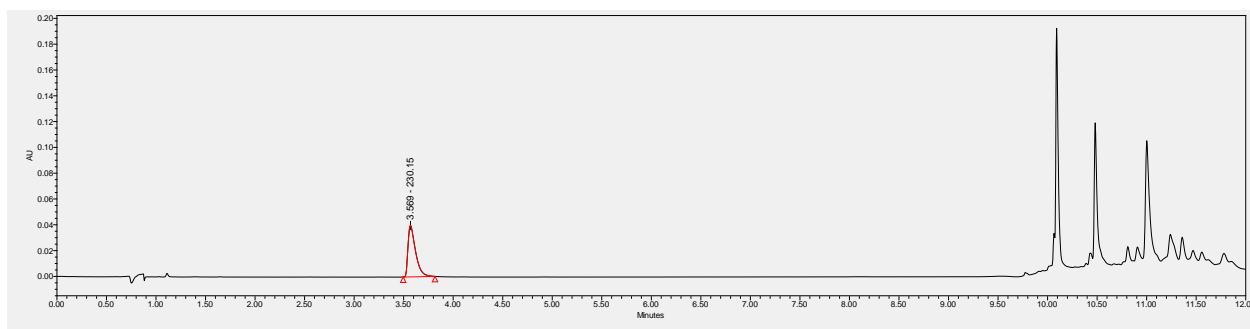
**154.3733**

Image	Spect	Reten Time (min)	Scan Number	Name	Info	Description	Combi Type	BaseLine Correction (%)	Peak Separation (Da)	Max Intensity (Intensity)	B P (f)
1	✓	2.604	1562		Extracted	Baseline Correction Off (2.60 ± 0.1)	Average	0.000	1.0000	40539964.000000	18

### 3-1 Mass Spectrum

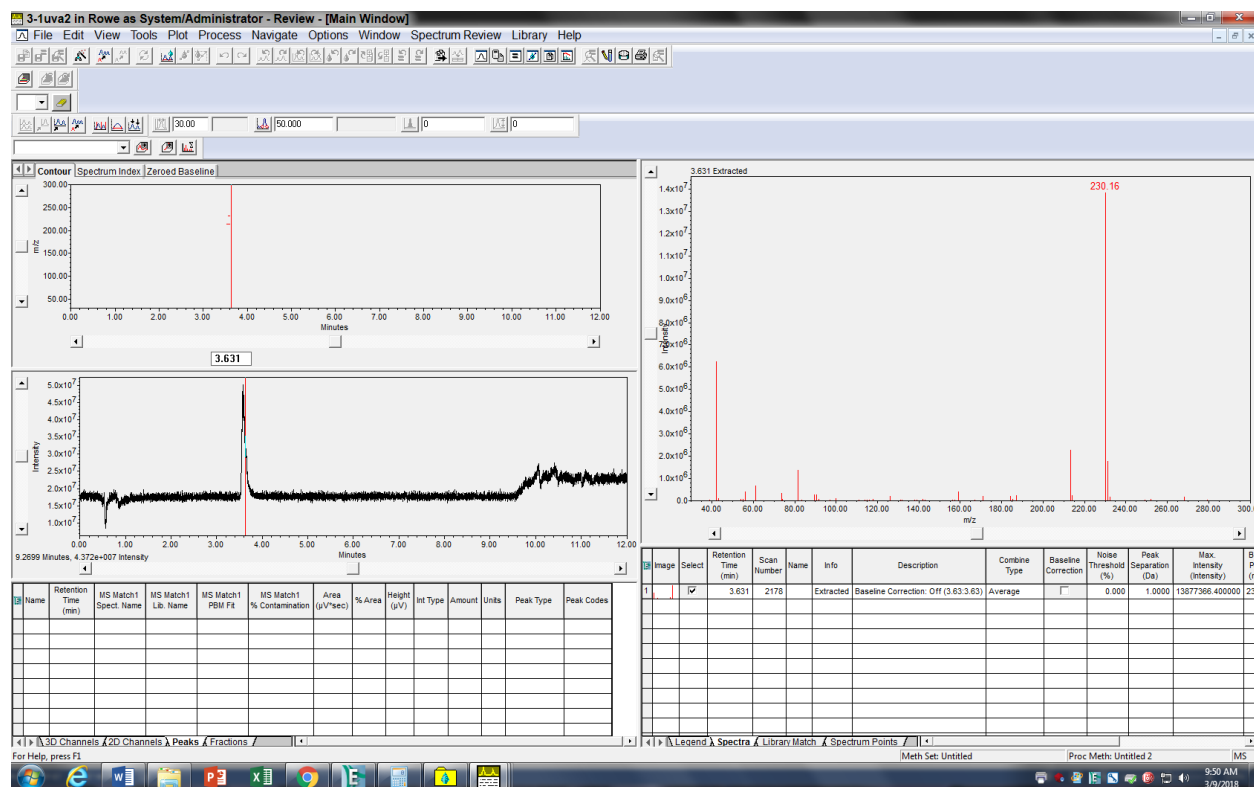


### 3-1 UV-A LC PDA Detector Data with Integrated Peak

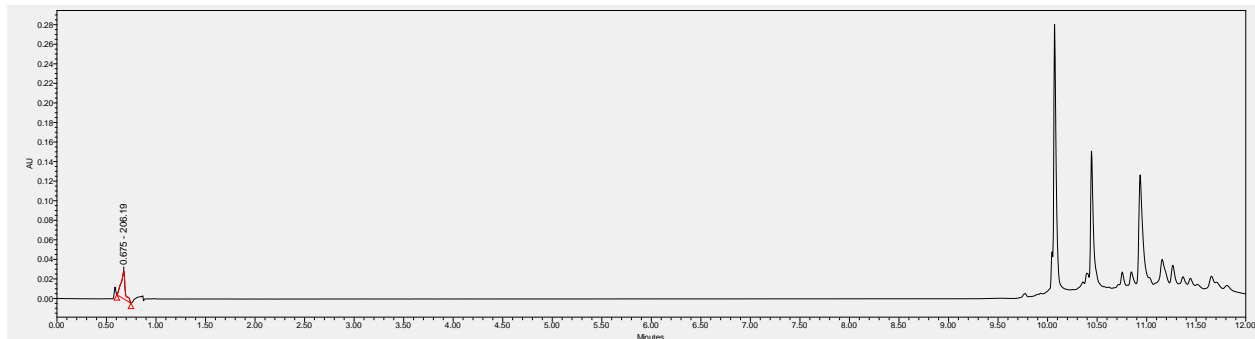


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.569	210426	100.00	39944	bb			Unknown

### 3-1 UVA Mass Spectrum

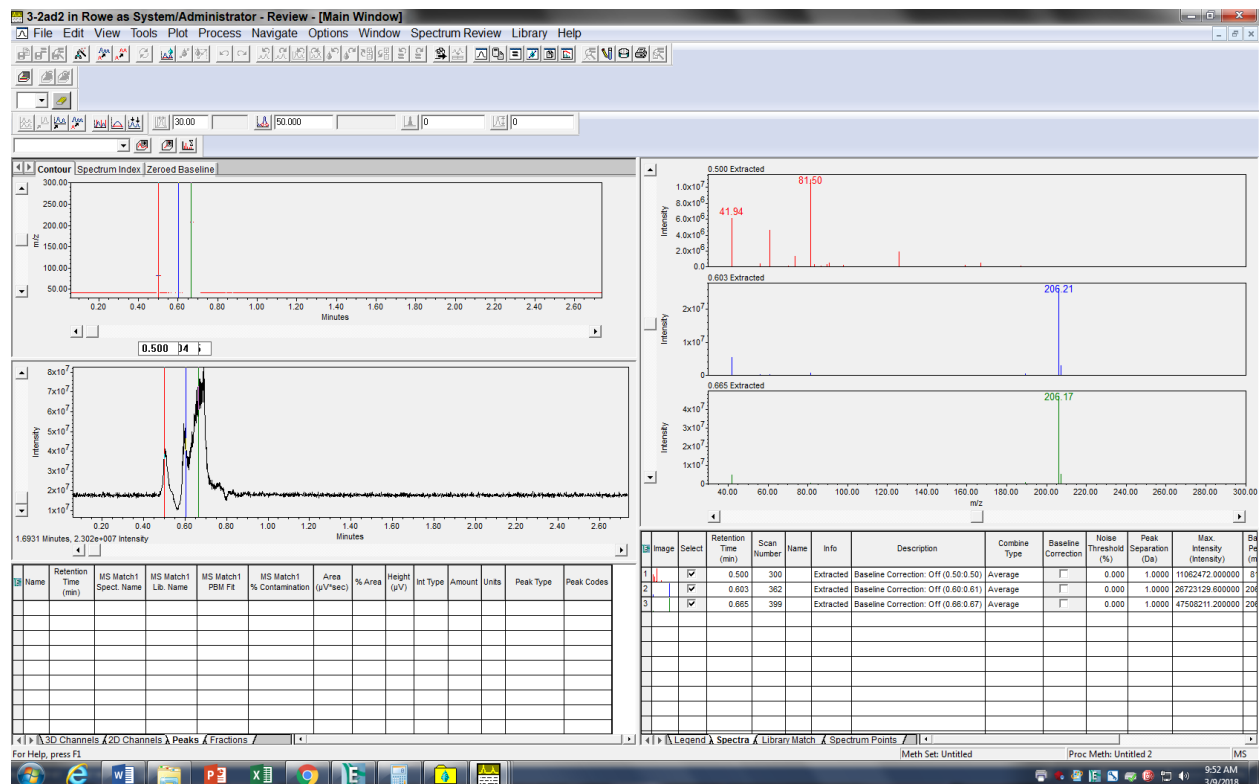


### 3-2 LC PDA Detector Data with Integrated Peak

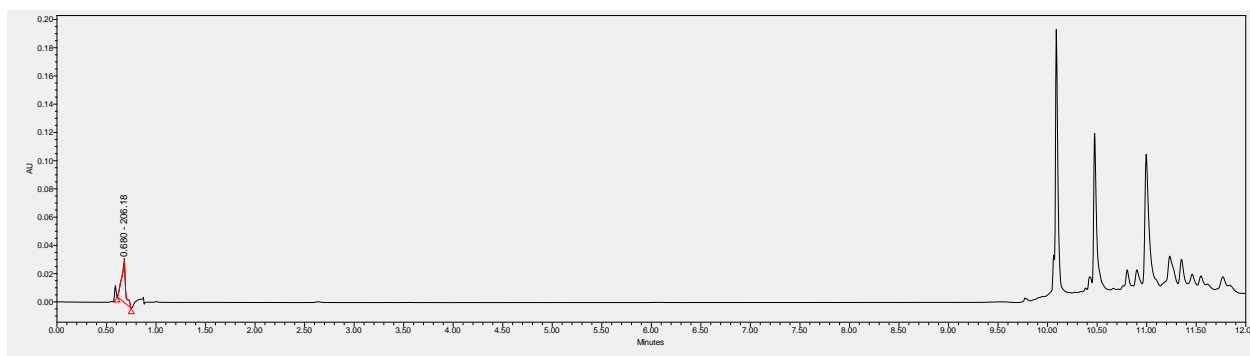


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.675	83012	100.00	28505	bb			Unknown

### 3-2 Mass Spectrum

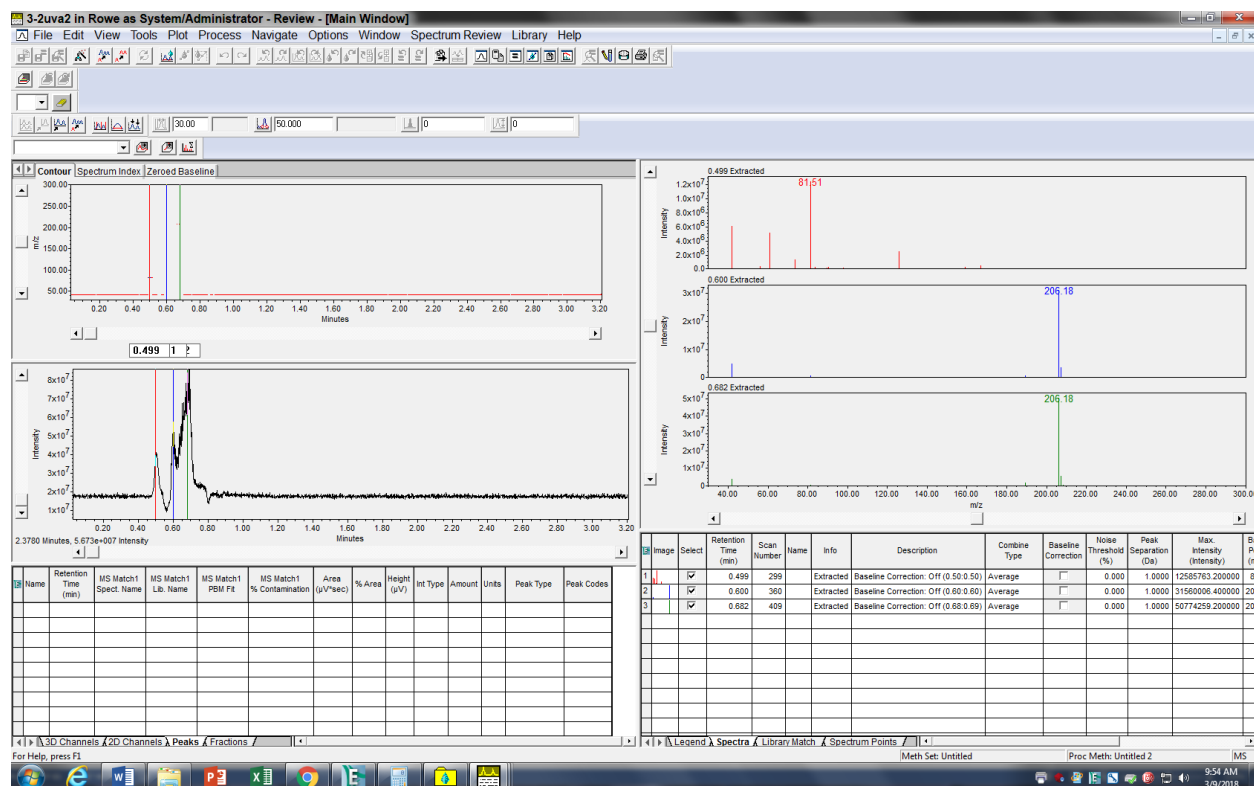


### 3-2 UV-A LC PDA Detector Data with Integrated Peak

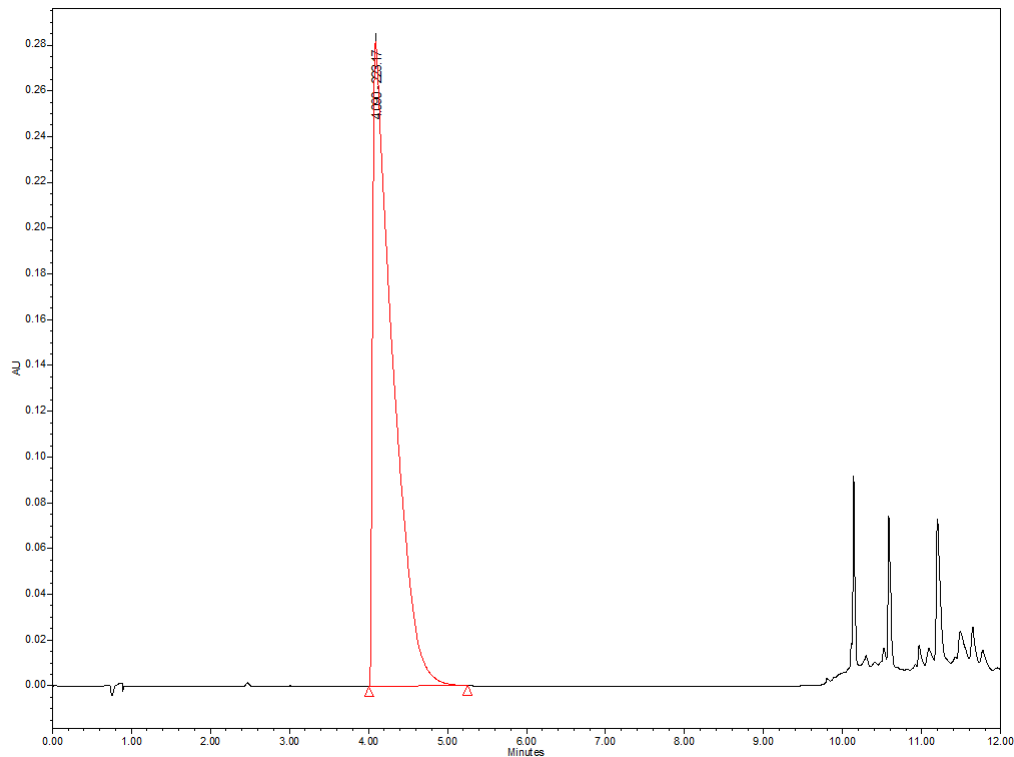


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.680	83099	100.00	28441	bb			Unknown

### 3-2 UVA Mass Spectrum

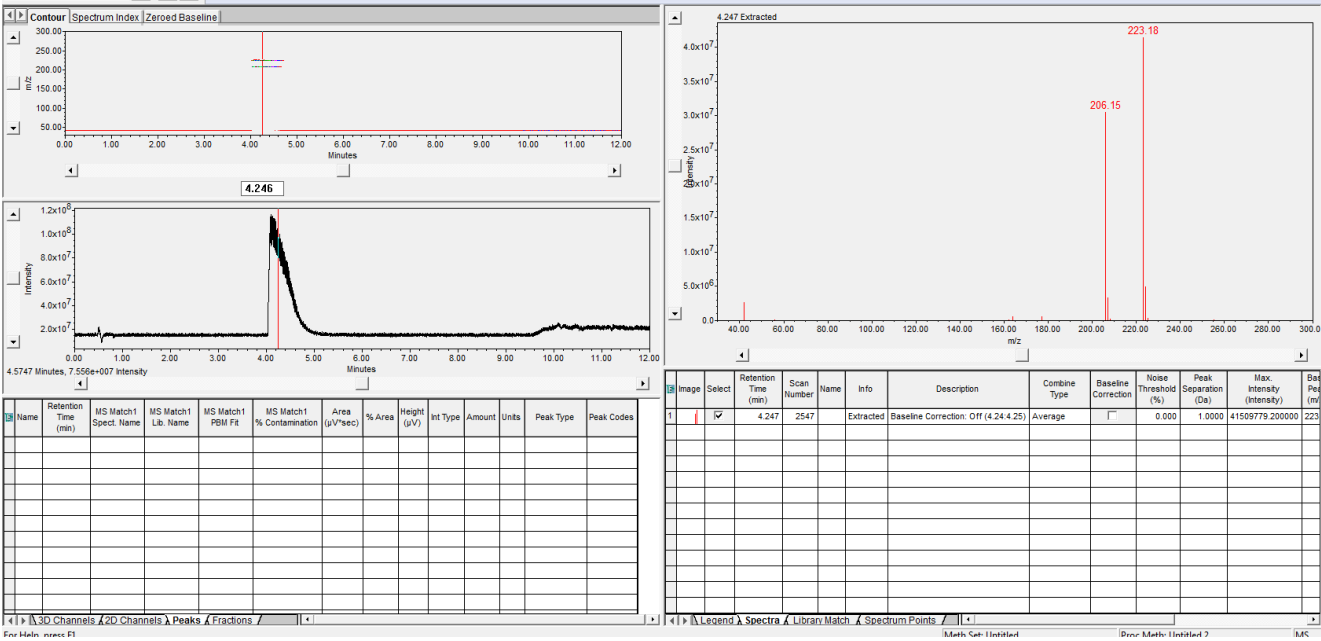


3-3 LC PDA Detector Data with Integrated Peak

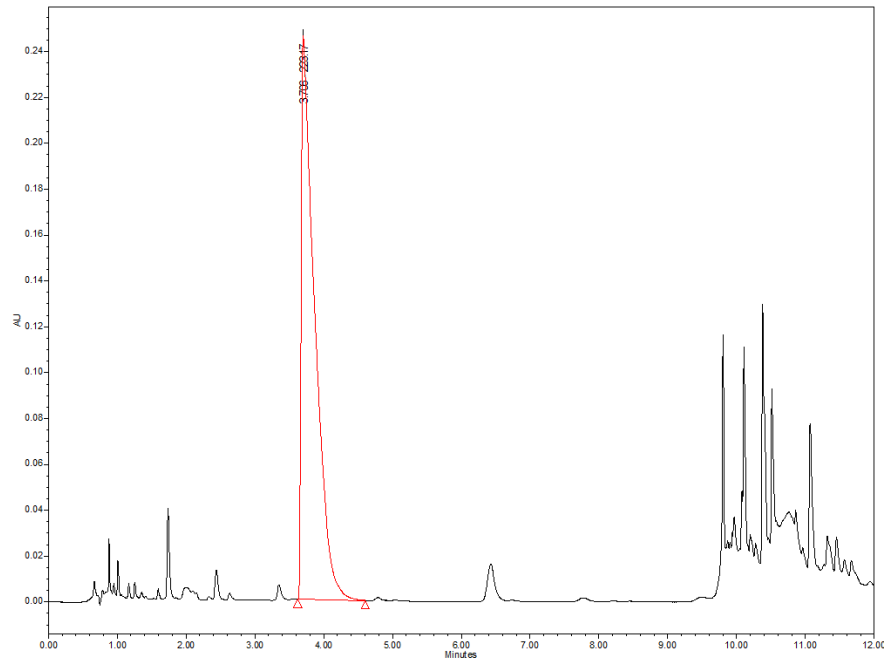


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.090	4951822	100.00	281952	bb			Unknown

3-3 Mass Spectrum

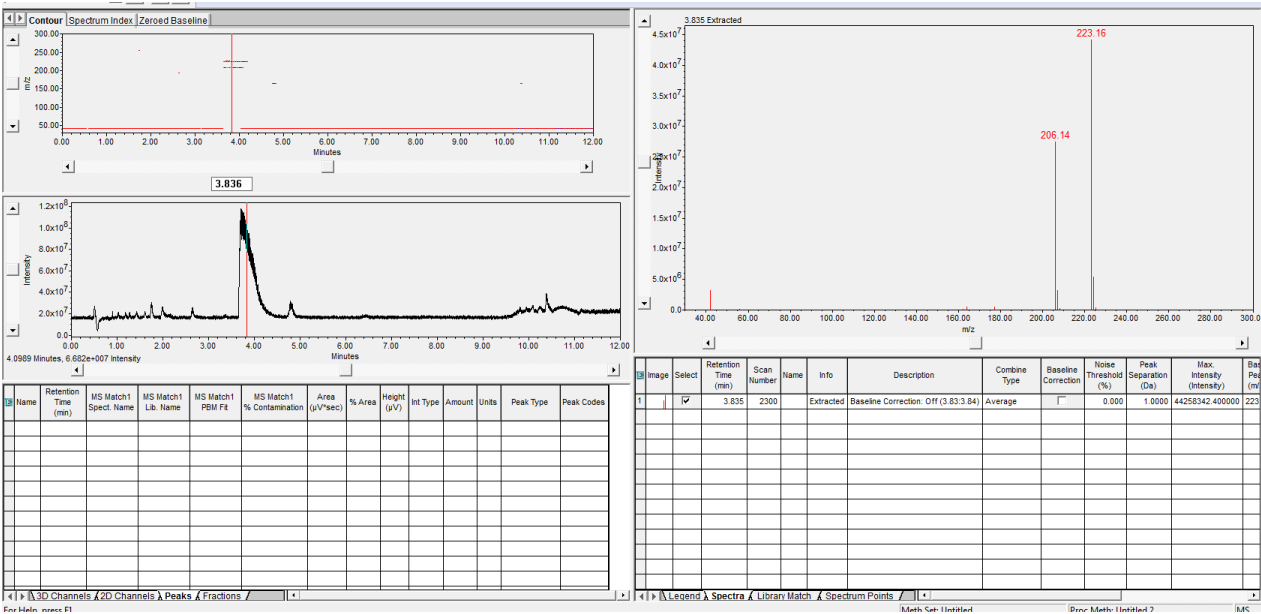


3-3 UV-A LC PDA Detector Data with Integrated Peak

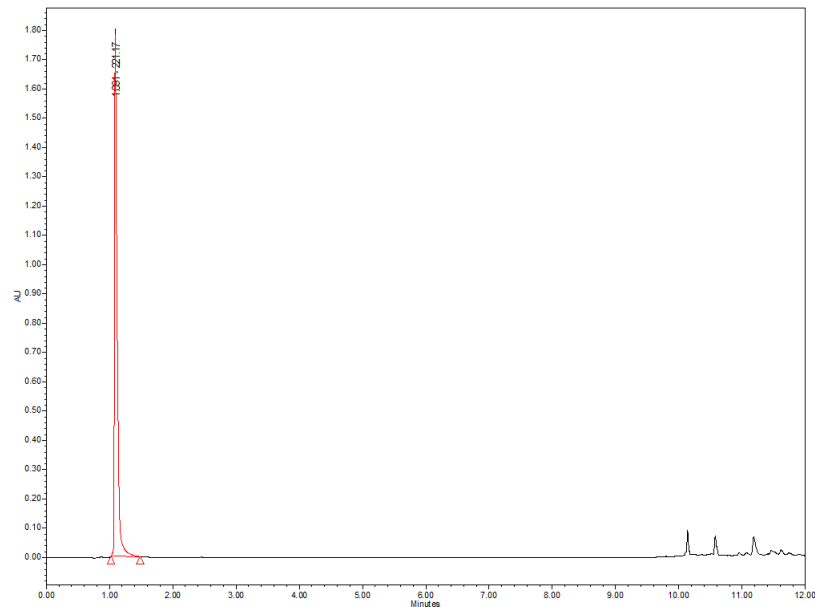


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.706	3346975	100.00	245919	bb			Unknown

3-3 UVA Mass Spectrum

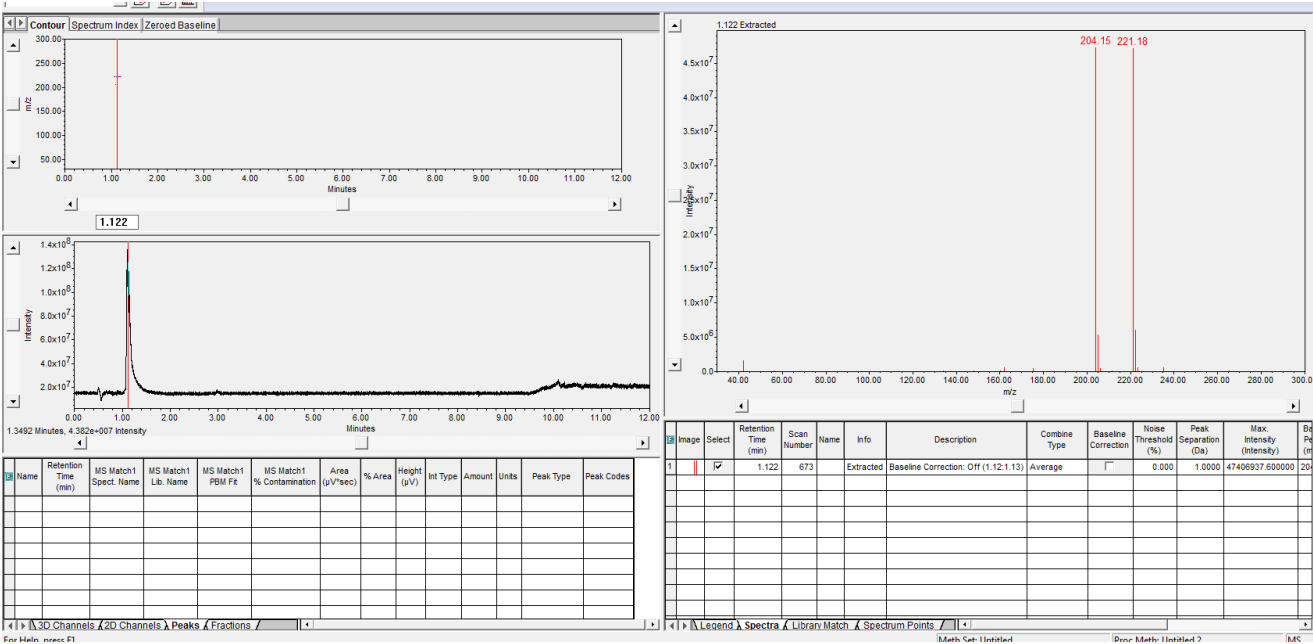


3-4 LC PDA Detector Data with Integrated Peak



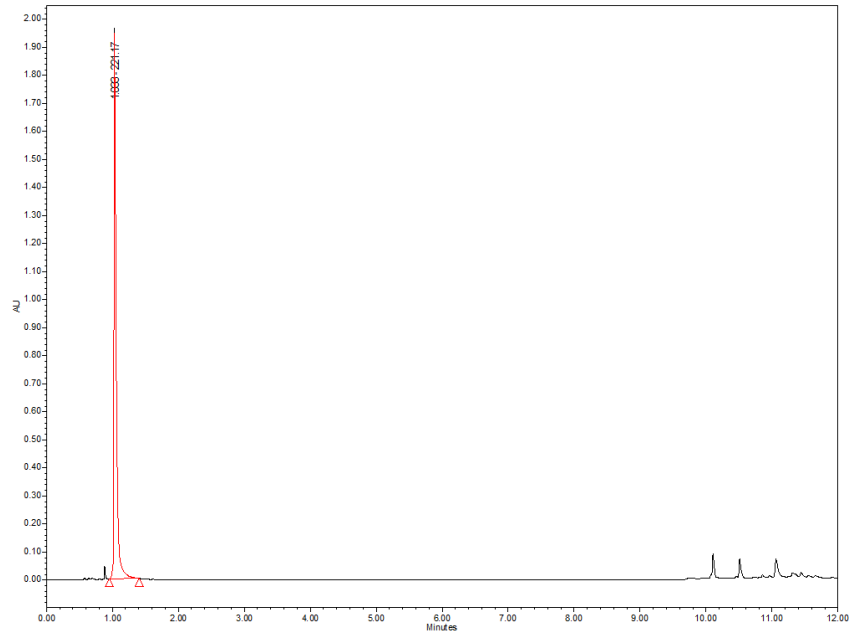
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.091	4974185	100.00	1783005	bb			Unknown

3-4 Mass Spectrum



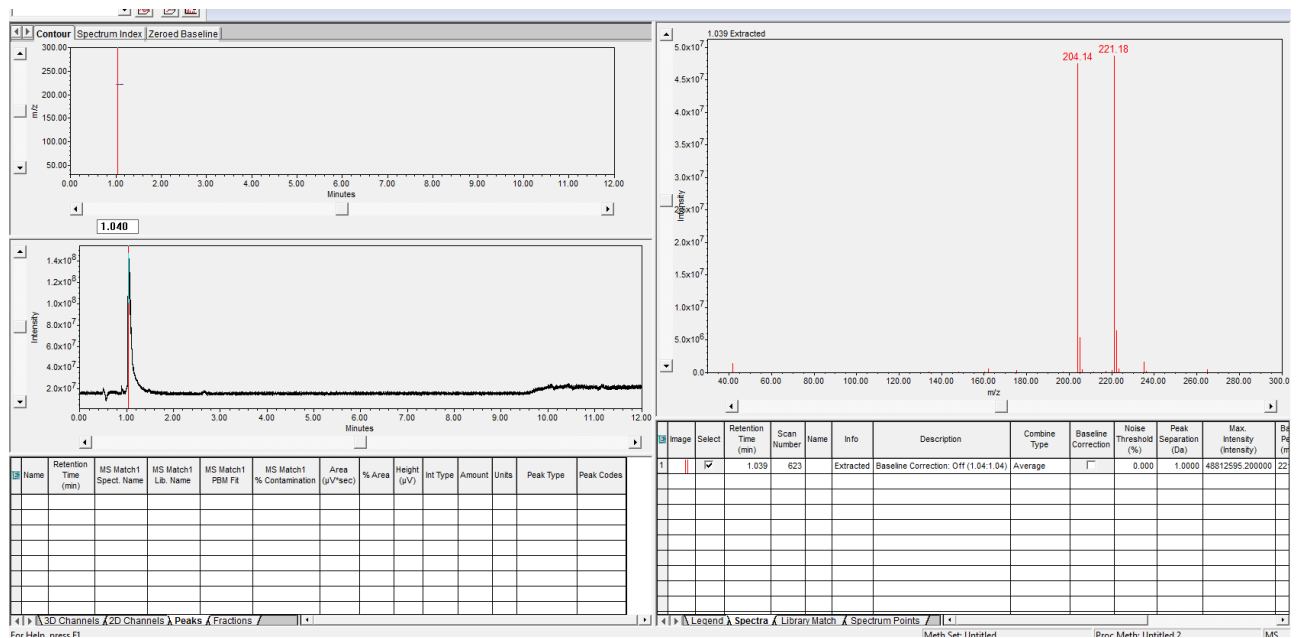


### 3-4 UV-A LC PDA Detector Data with Integrated Peak

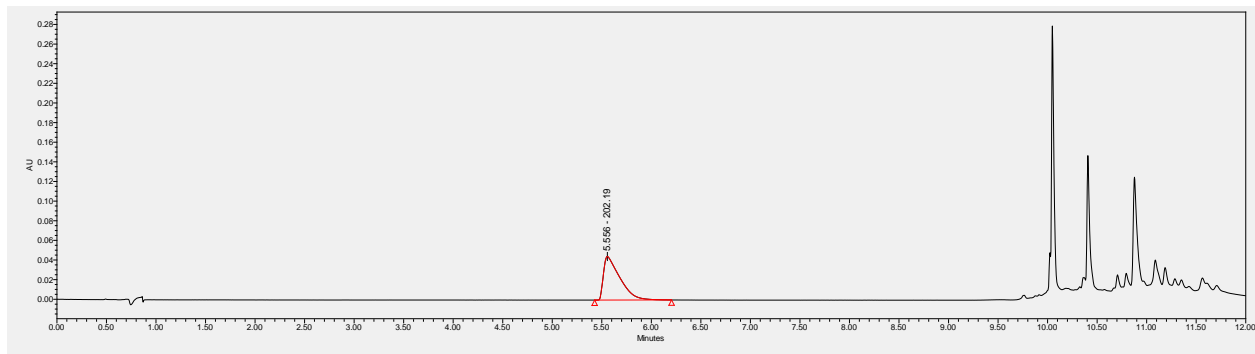


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.033	4802318	100.00	1946698	bb			Unknown

### 3-4 UVA Mass Spectrum

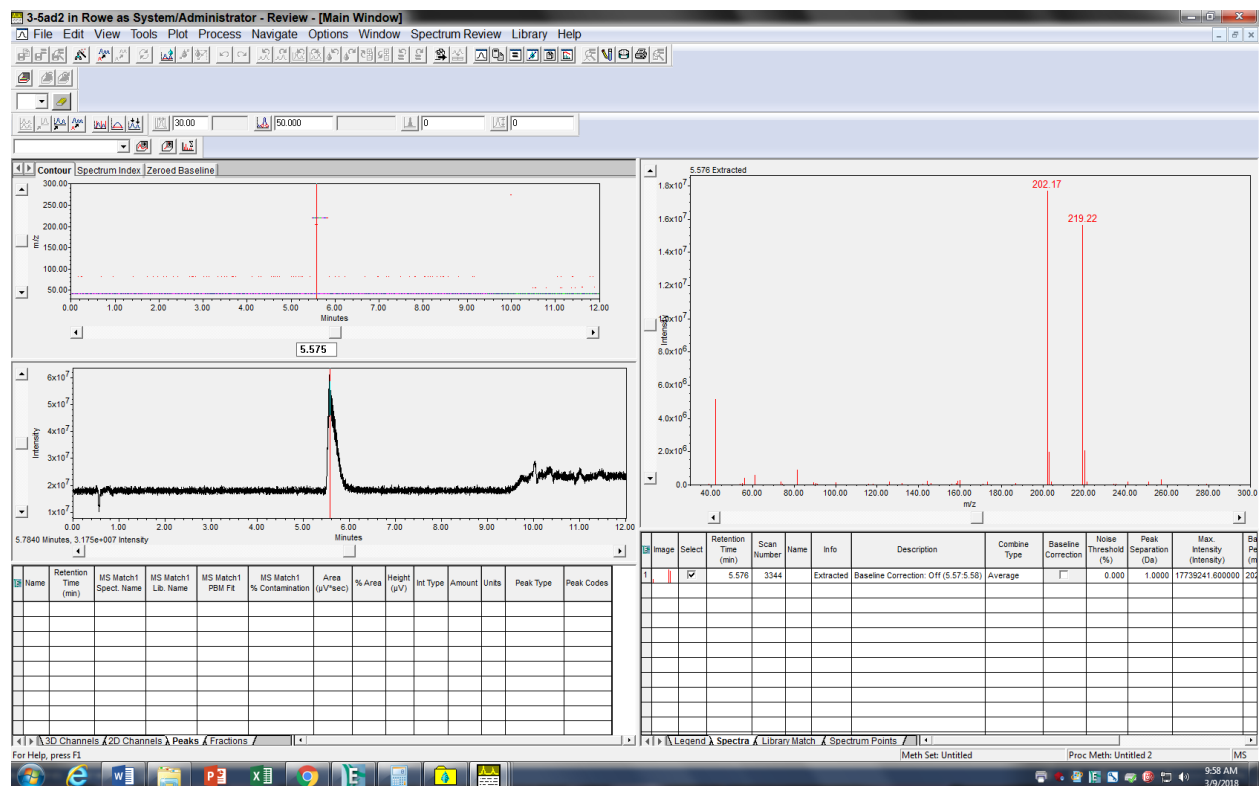


### 3-5 LC PDA Detector Data with Integrated Peak

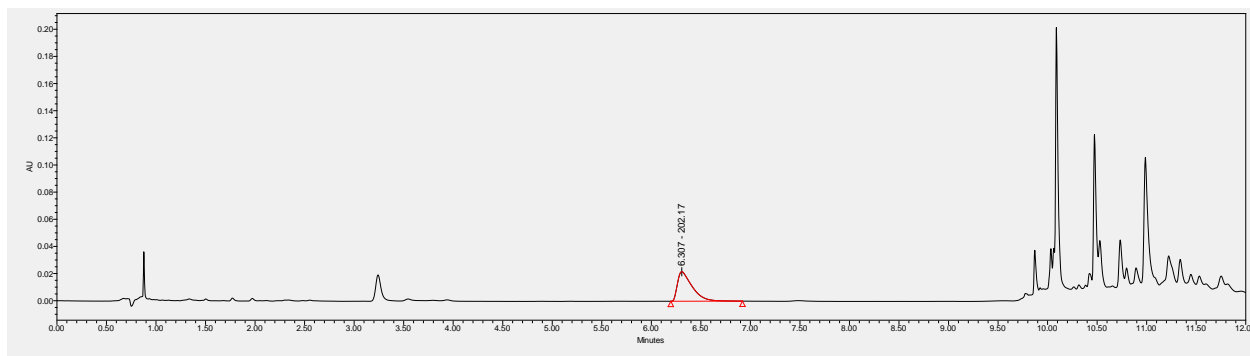


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		5.556	507269	100.00	44401	bb			Unknown

### 3-5 Mass Spectrum

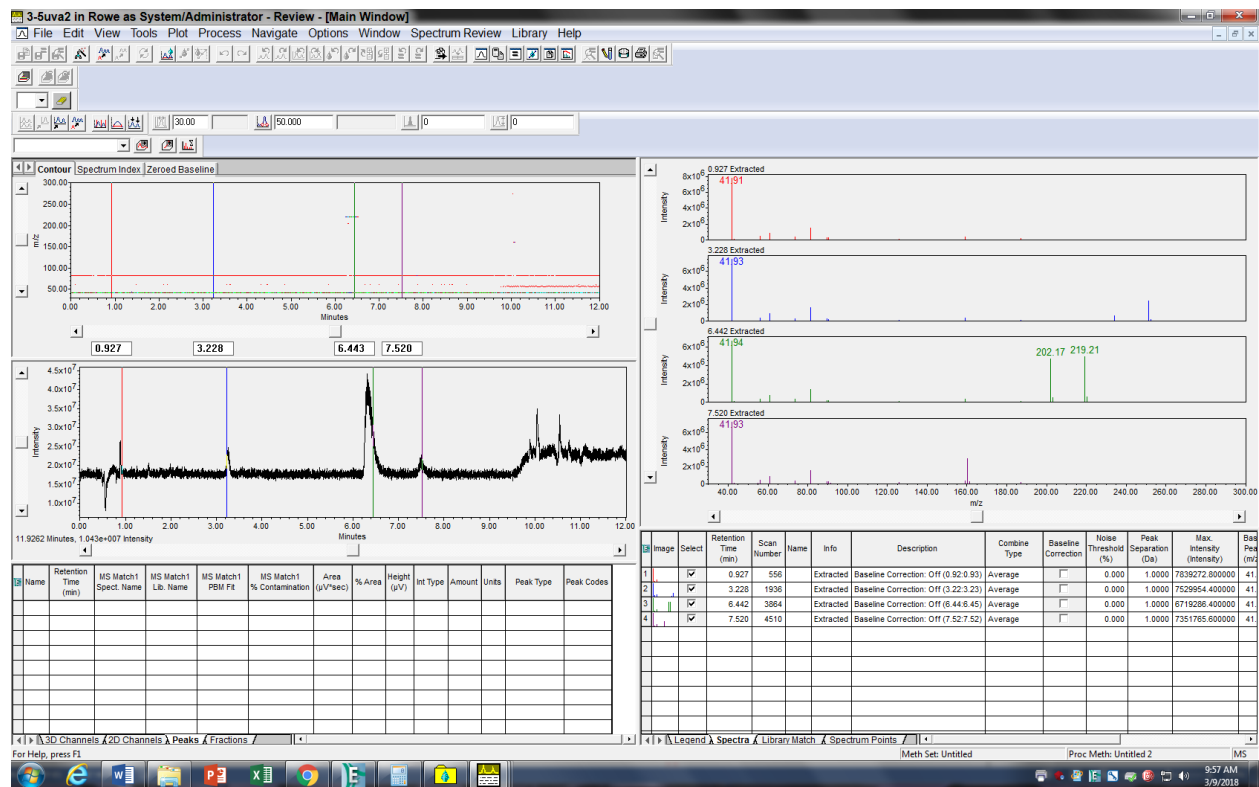


### 3-5 UV-A LC PDA Detector Data with Integrated Peak

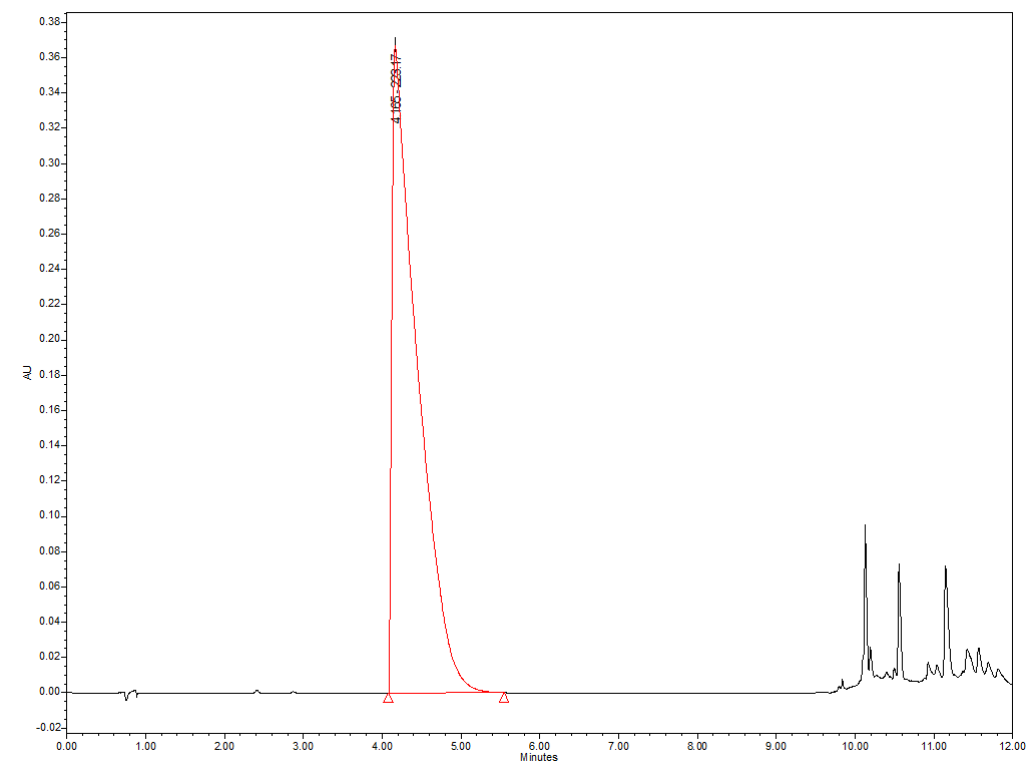


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		6.307	222723	100.00	21758	bb			Unknown

### 3-5 UVA Mass Spectrum

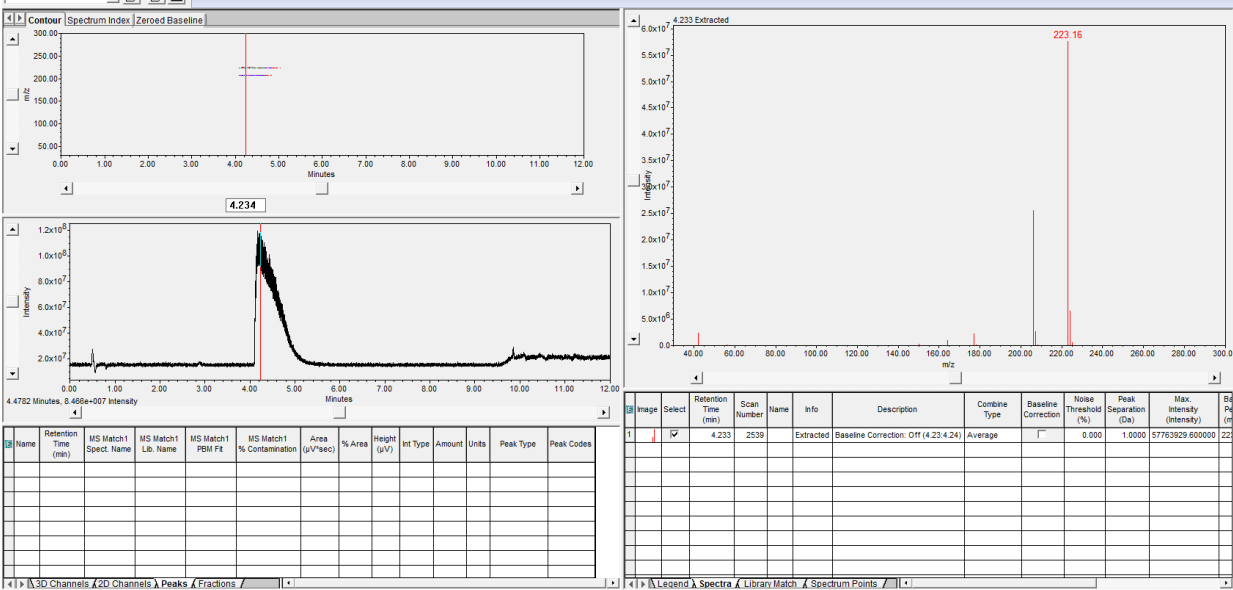


3-6 LC PDA Detector Data with Integrated Peak

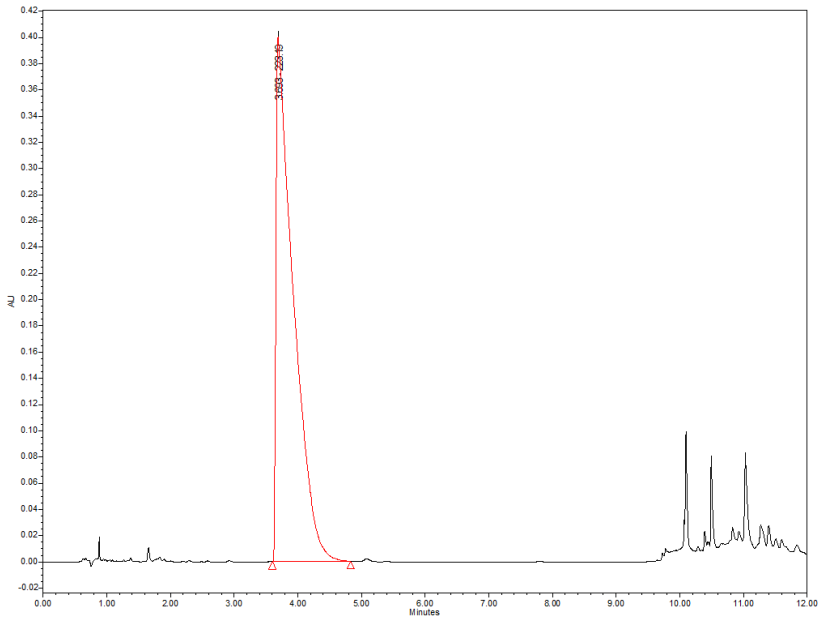


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.165	8174336	100.00	367331	bb			Unknown

3-6 Mass Spectrum

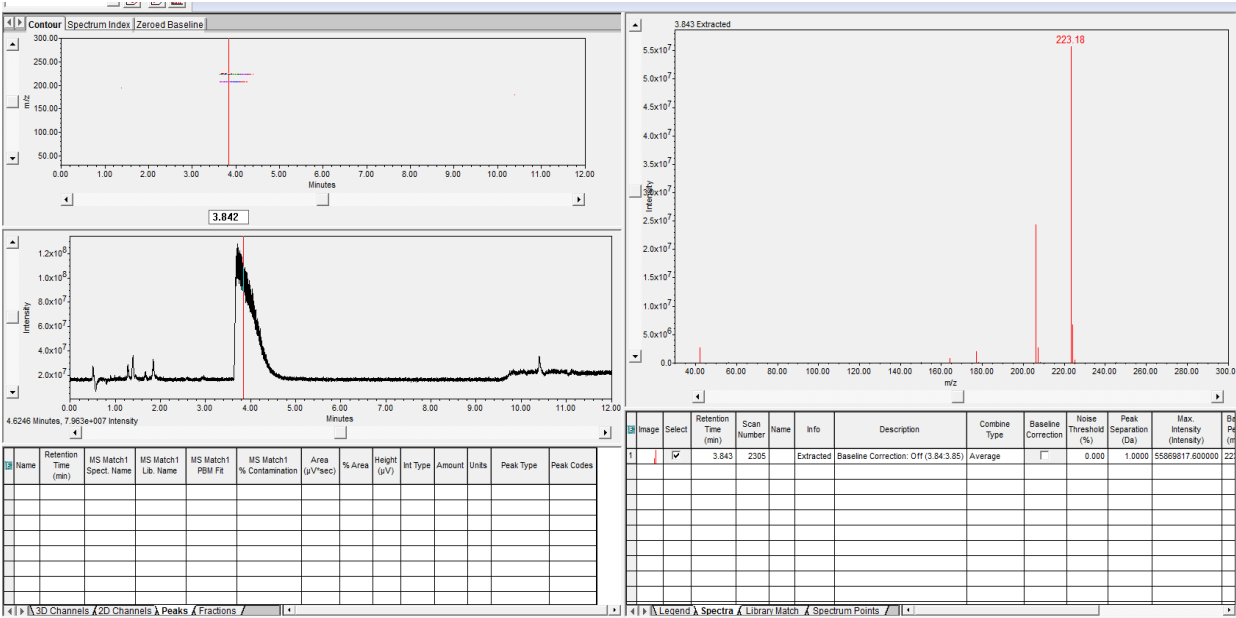


3-6 UV-A LC PDA Detector Data with Integrated Peak

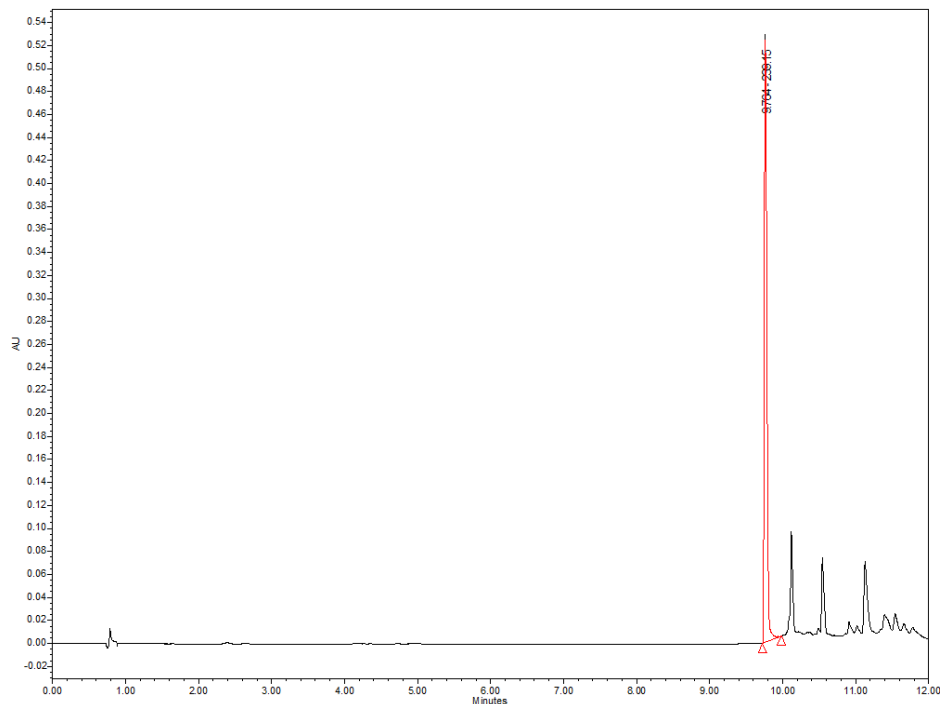


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.693	7561231	100.00	400448	bb			Unknown

3-6 UVA Mass Spectrum

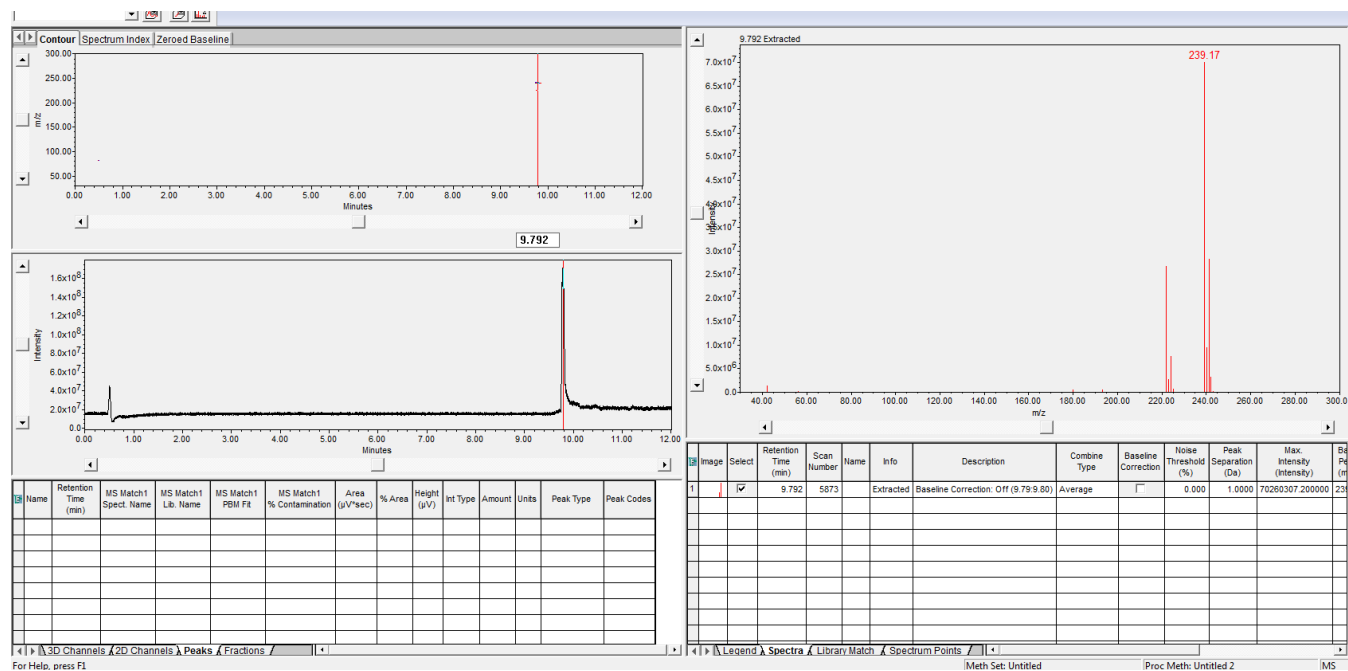


### 3-7 LC PDA Detector Data with Integrated Peak

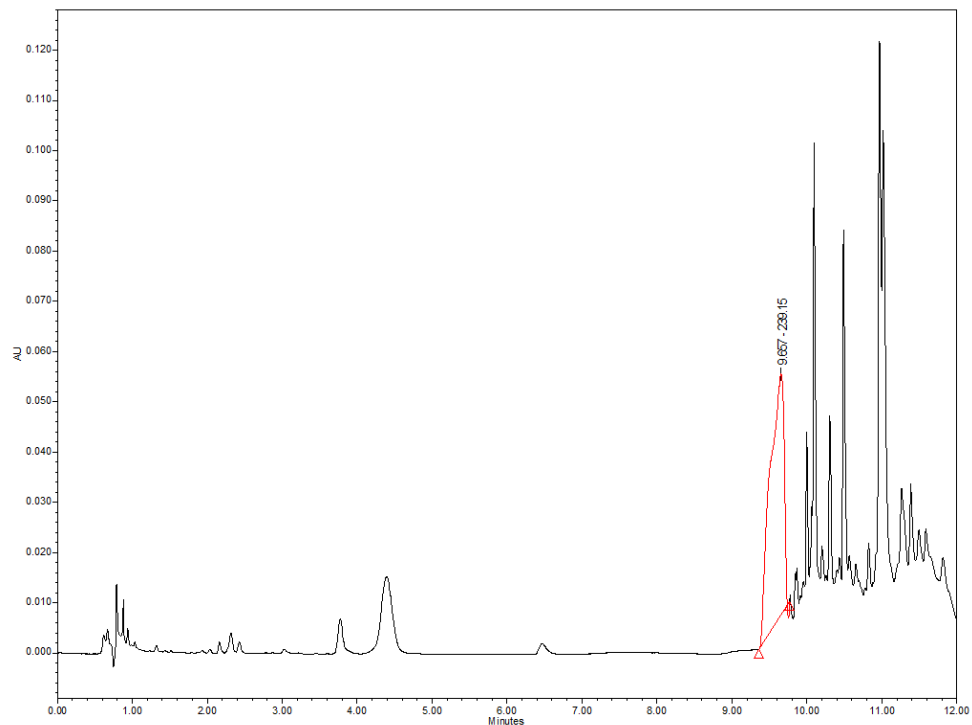


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.764	1219395	100.00	524181	bb			Unknown

### 3-7 Mass Spectrum



### 3-7 UV-A LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.657	635231	100.00	47919	bb			Unknown

### 3-7 UVA Mass Spectrum

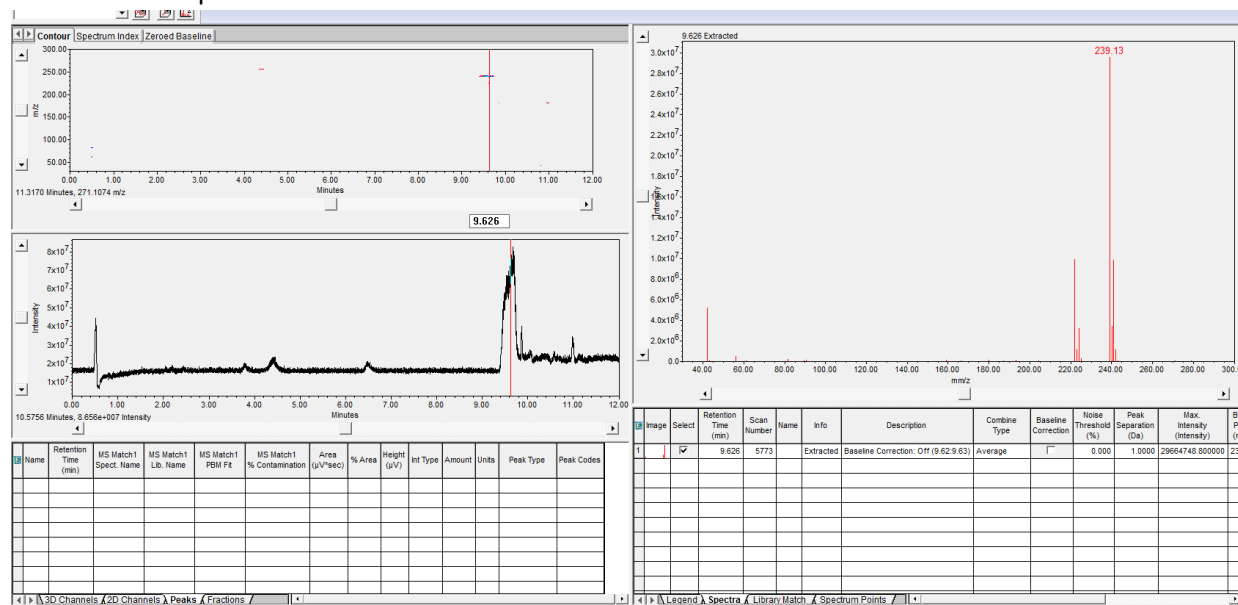
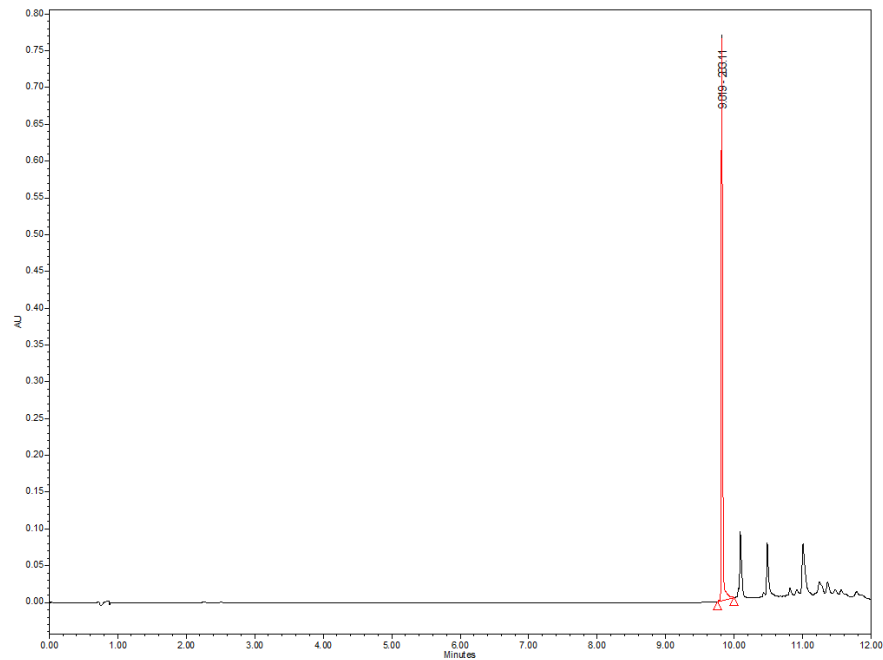


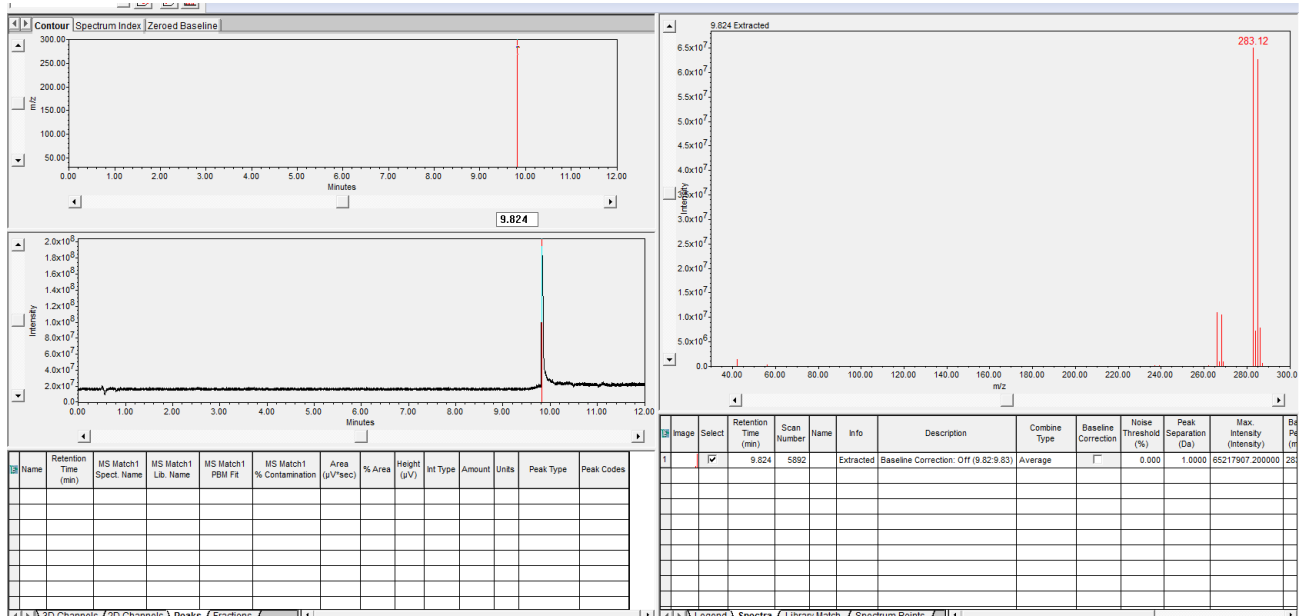
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity	B4 P4 (m)
1	<input checked="" type="checkbox"/>	9.626	5773	Extracted	Baseline Correction: Off (9.62 9.63)	Average			0.000	1.0000	29664746.8000000	23

### 3-8 LC PDA Detector Data with Integrated Peak



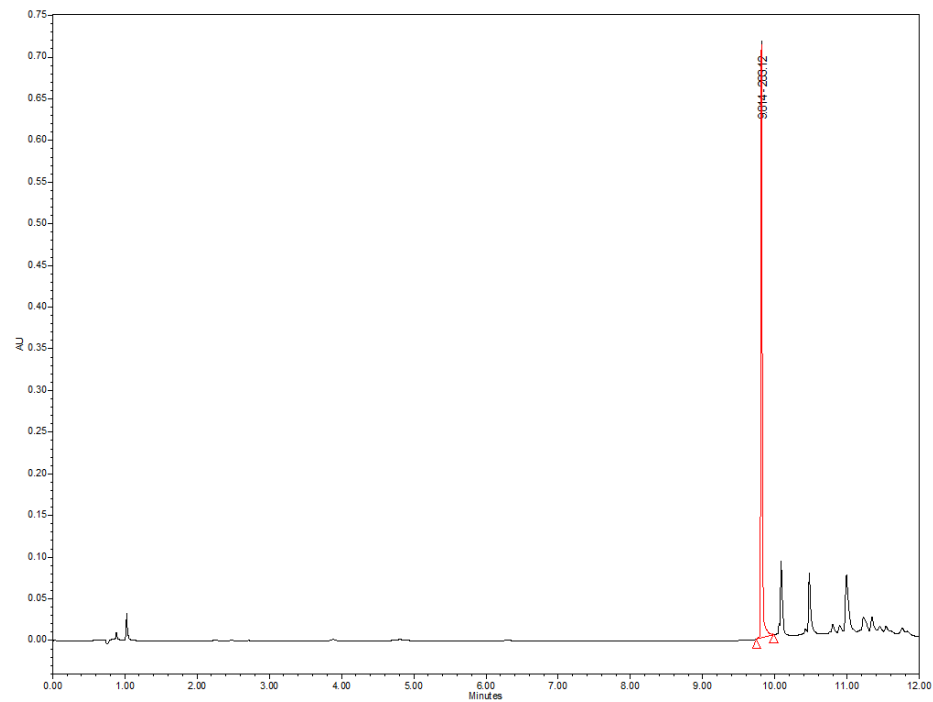
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.819	1146865	100.00	765053	bb			Unknown

### 3-8 Mass Spectrum



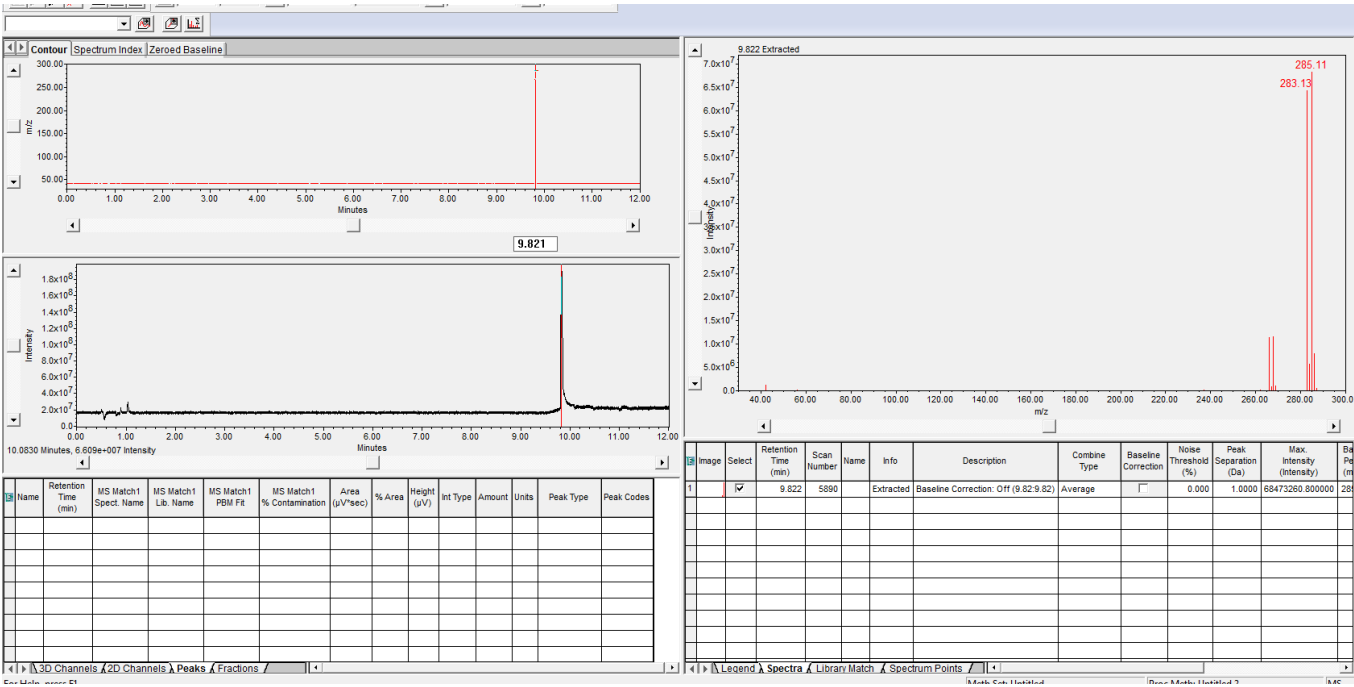


3-8 UV-A LC PDA Detector Data with Integrated Peak

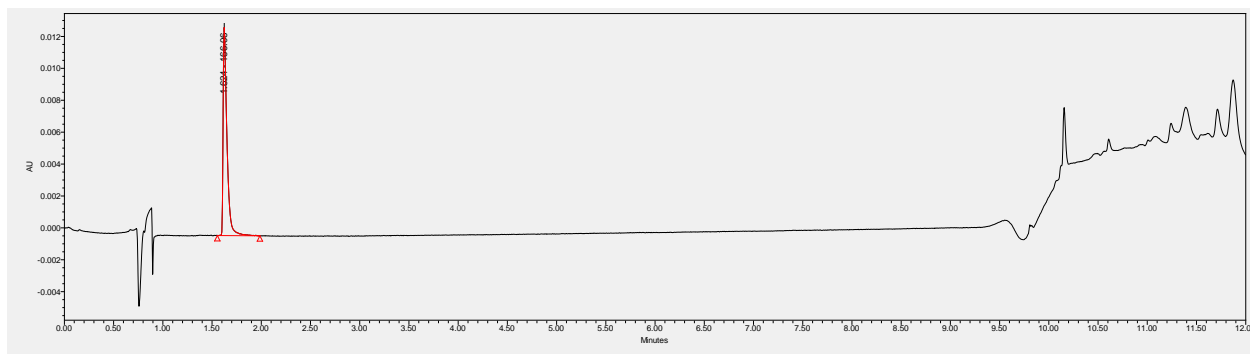


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.814	1071974	100.00	712569	bb			Unknown

3-8 UVA Mass Spectrum

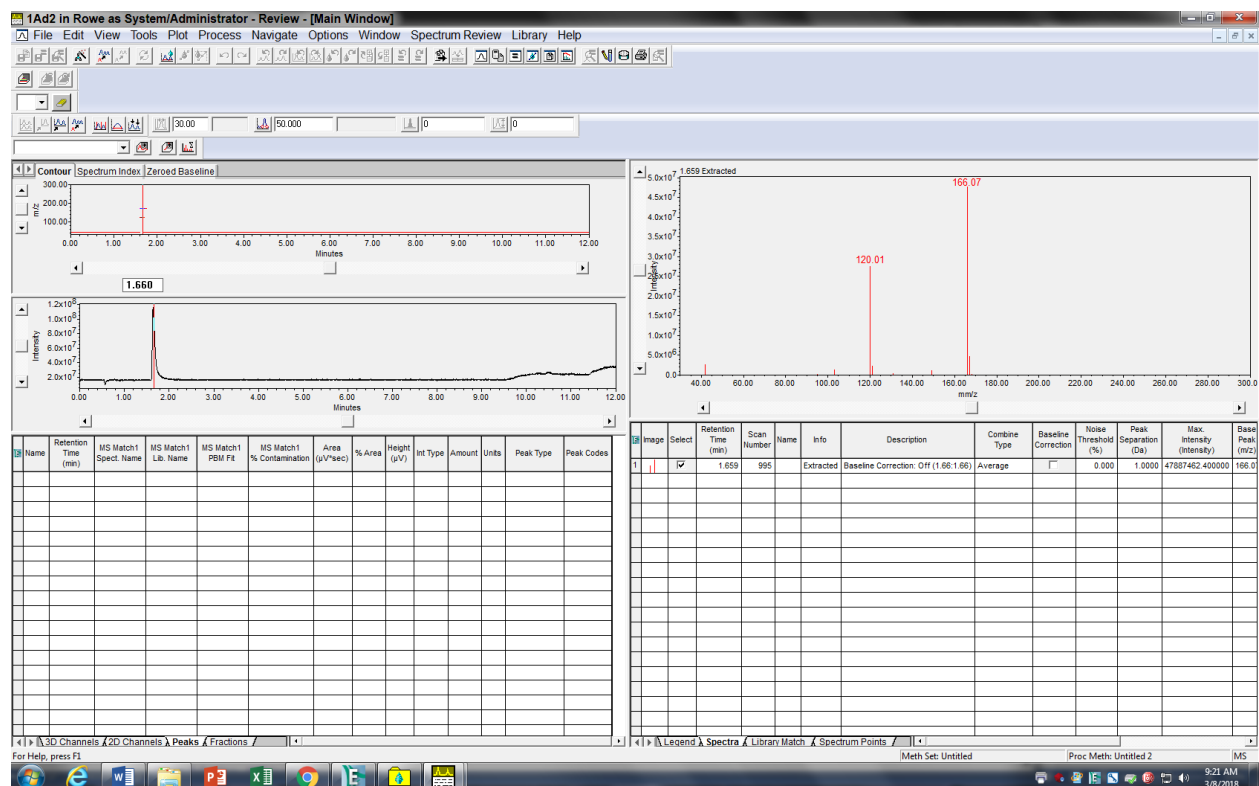


## 1 LC PDA Detector Data with Integrated Peak

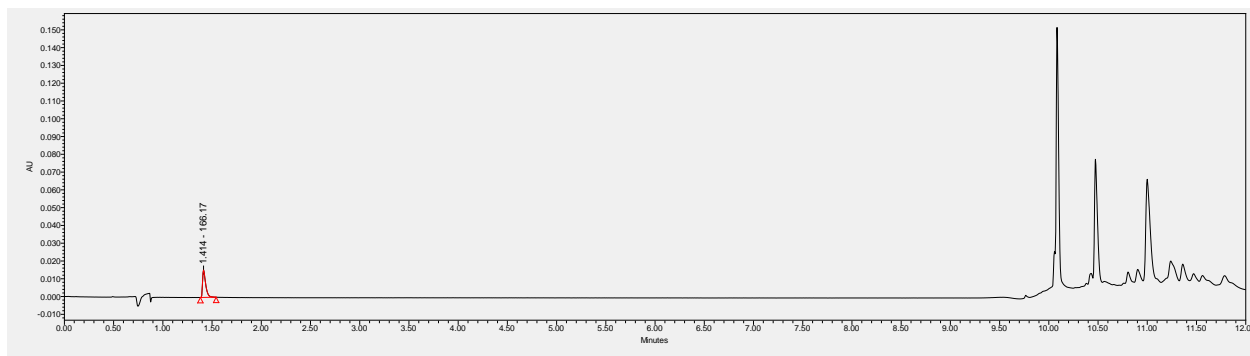


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.624	35449	100.00	13046	bb			Unknown

## 1 Mass Spectrum

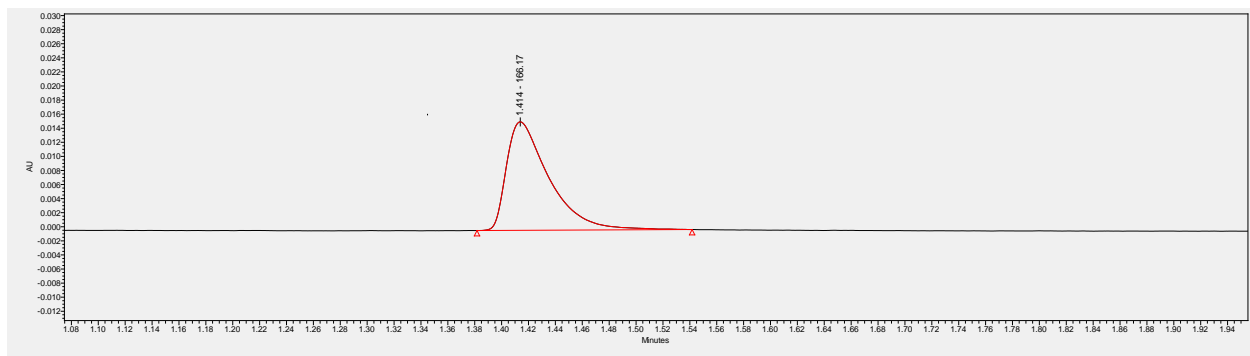


## 1 UV-B LC PDA Detector Data with Integrated Peak

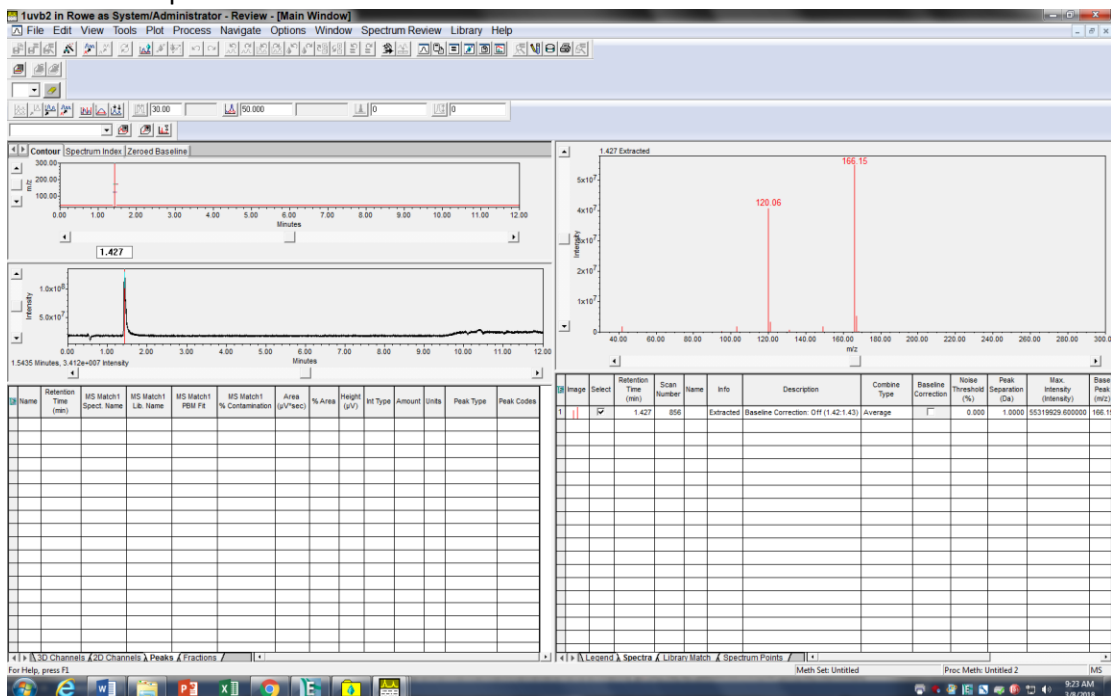


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.414	33814	100.00	15412	bb			Unknown

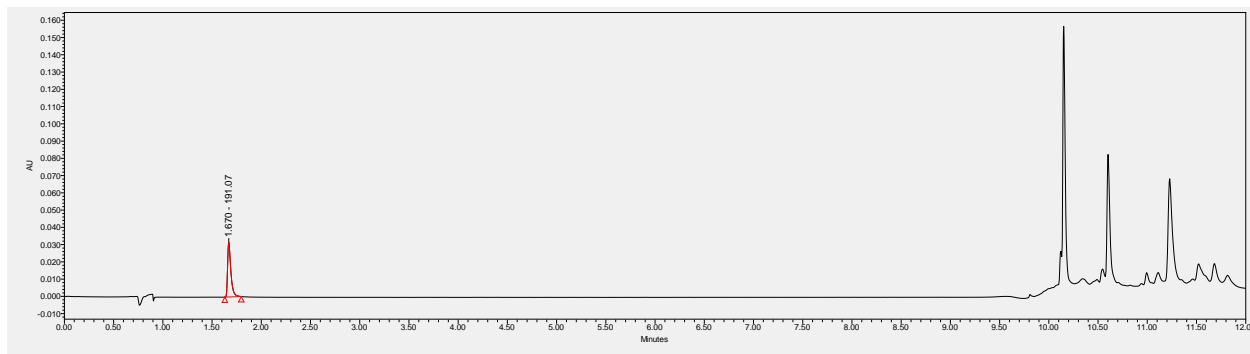
Zoom in of integrated peak shown above:



## 1 UVB Mass Spectrum

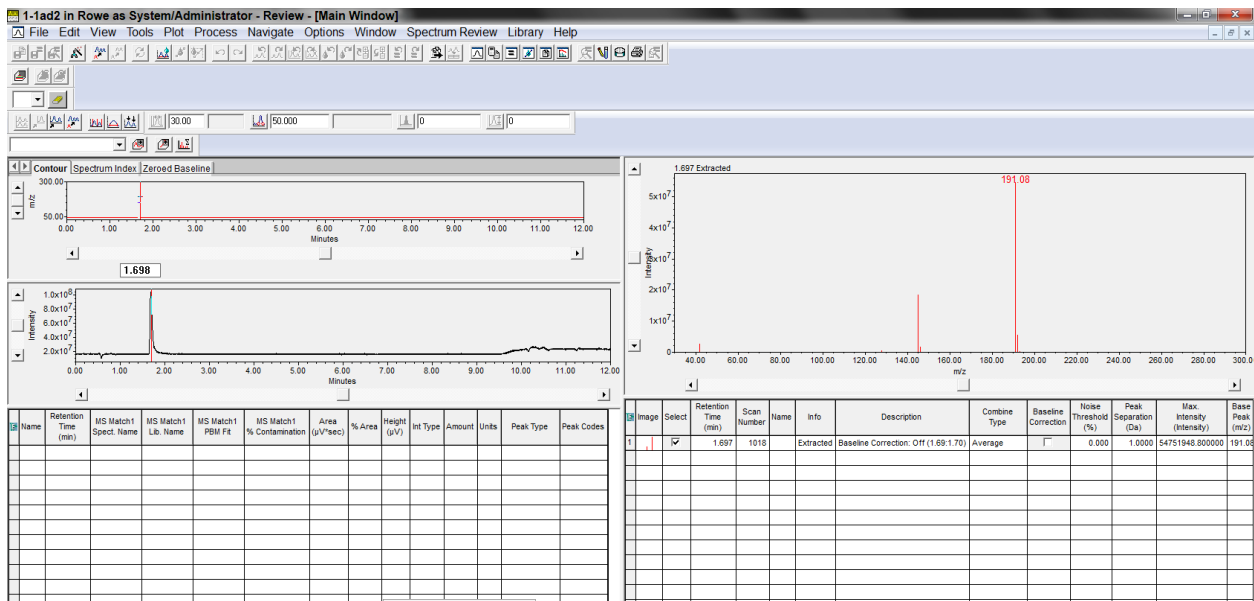


## 1-1 LC PDA Detector Data with Integrated Peak

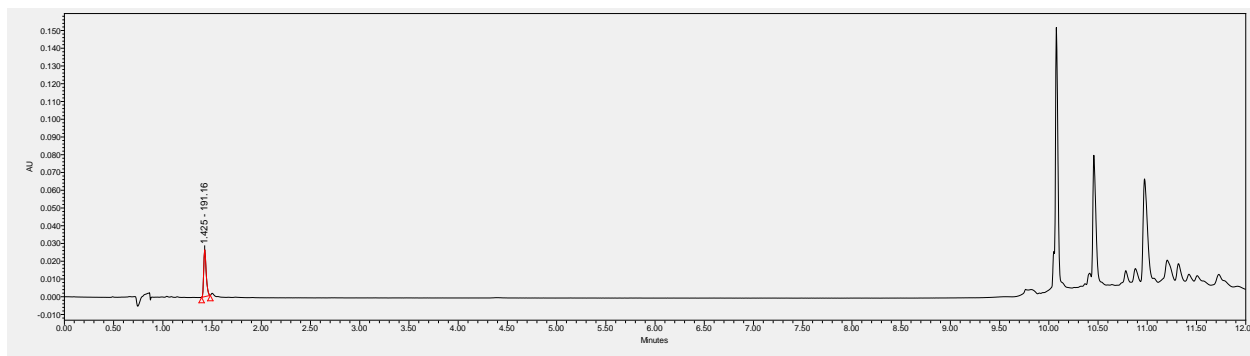


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.670	72903	100.00	31568	bb			Unknown

## 1-1 Mass Spectrum

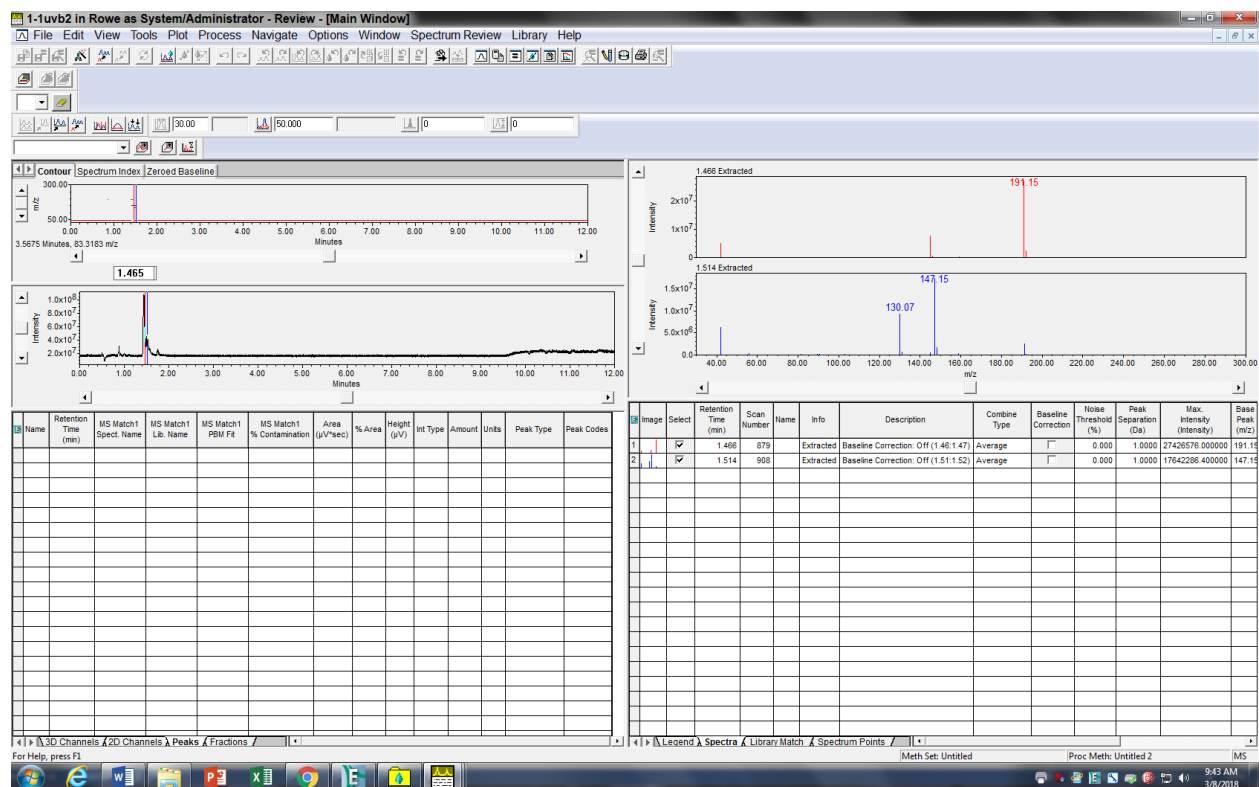


## 1-1UV-B LC PDA Detector Data with Integrated Peak

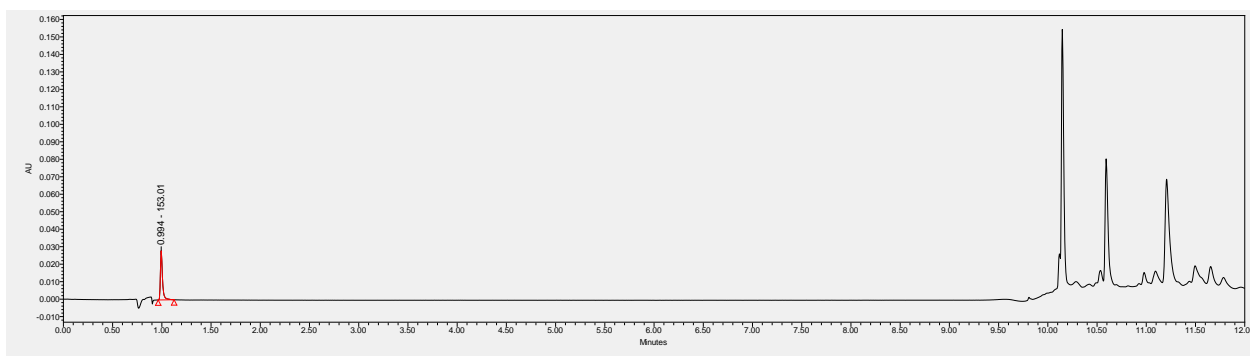


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.425	46285	100.00	26404	bb			Unknown

## 1-1 UVB Mass Spectrum

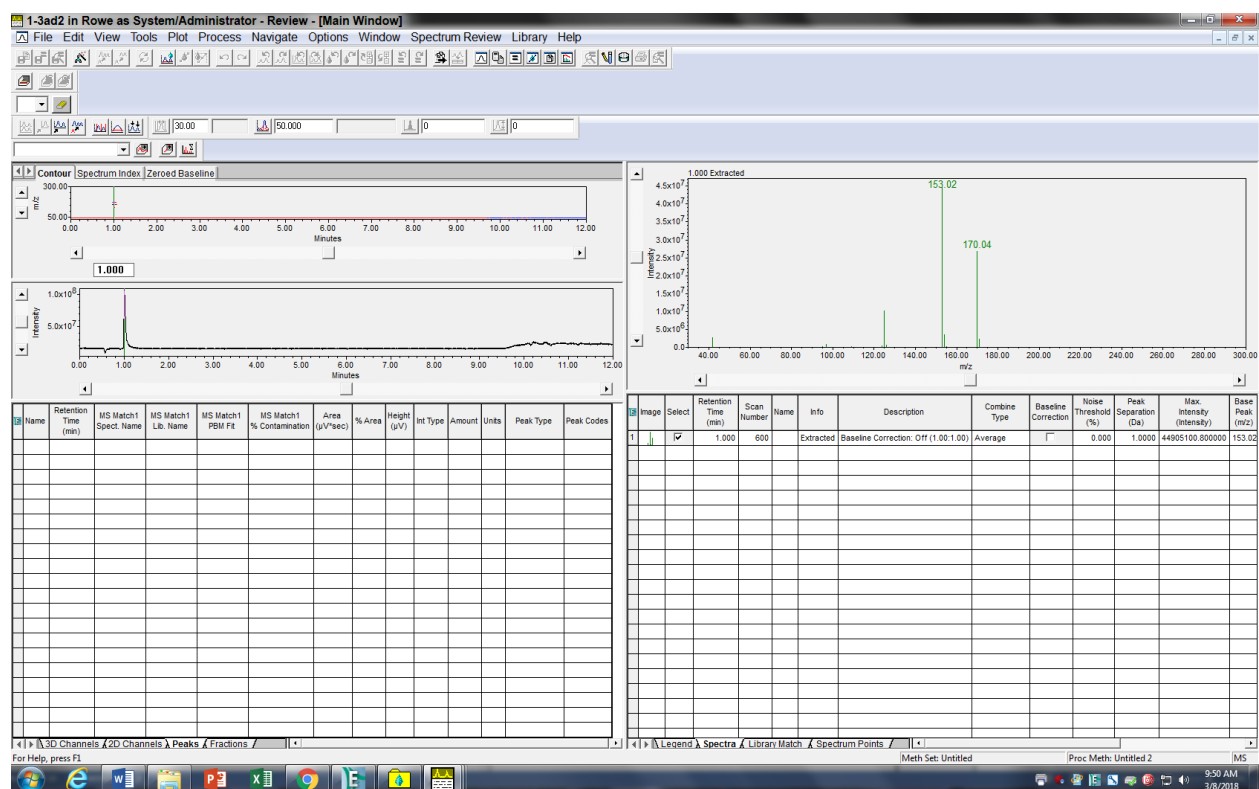


## 1-3 LC PDA Detector Data with Integrated Peak

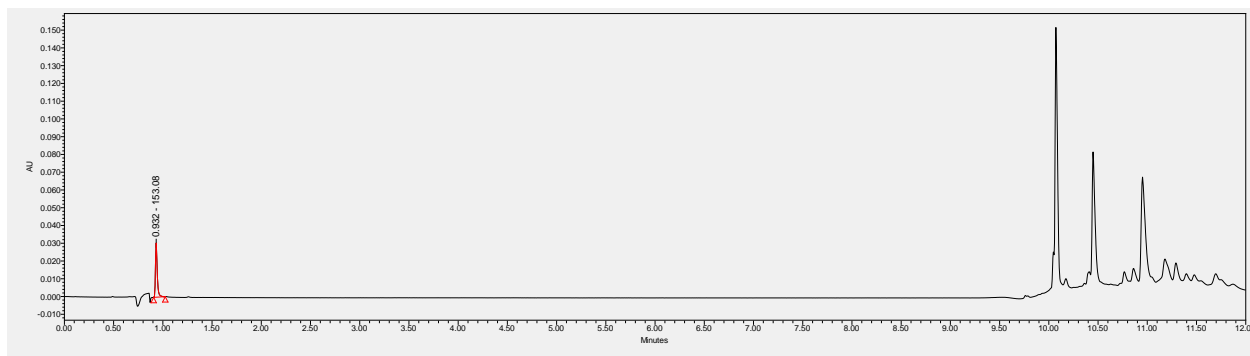


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.994	41530	100.00	28142	bb			Unknown

## 1-3 Mass Spectrum

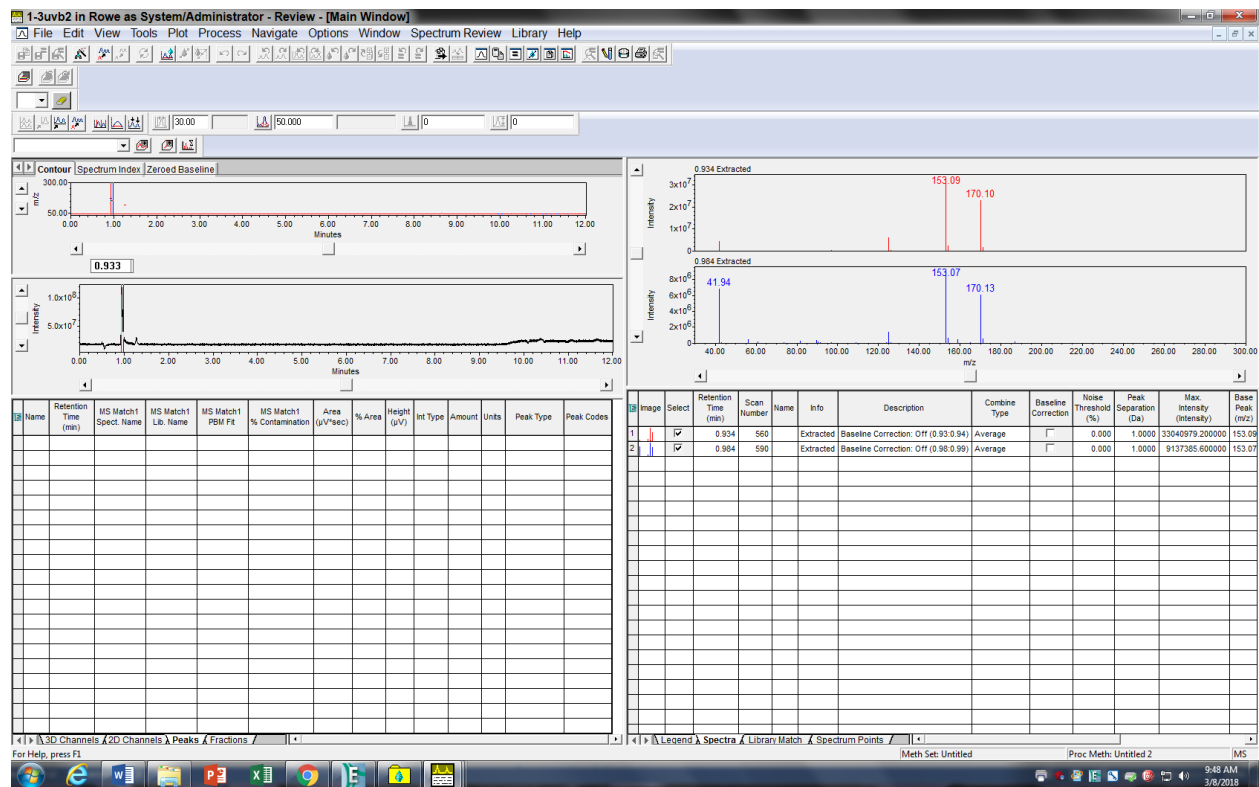


## 1-3 UV-B LC PDA Detector Data with Integrated Peak

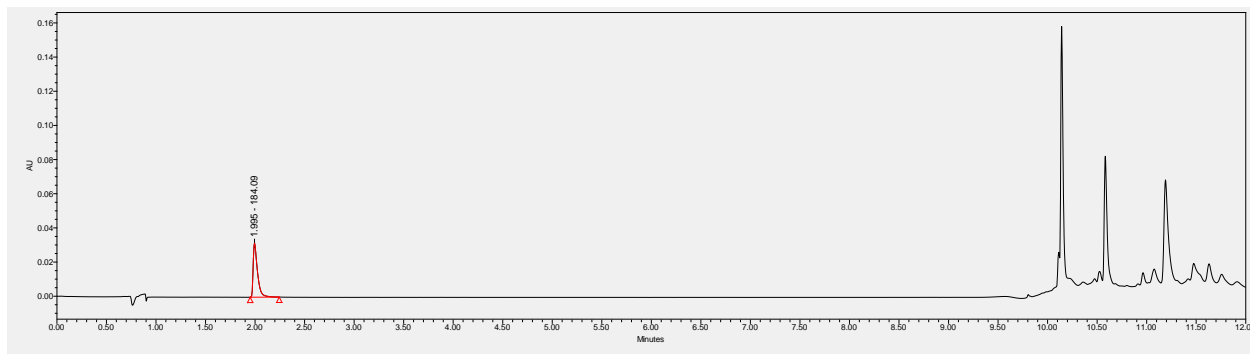


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.932	39169	100.00	30526	bb			Unknown

## 1-3 UVB Mass Spectrum

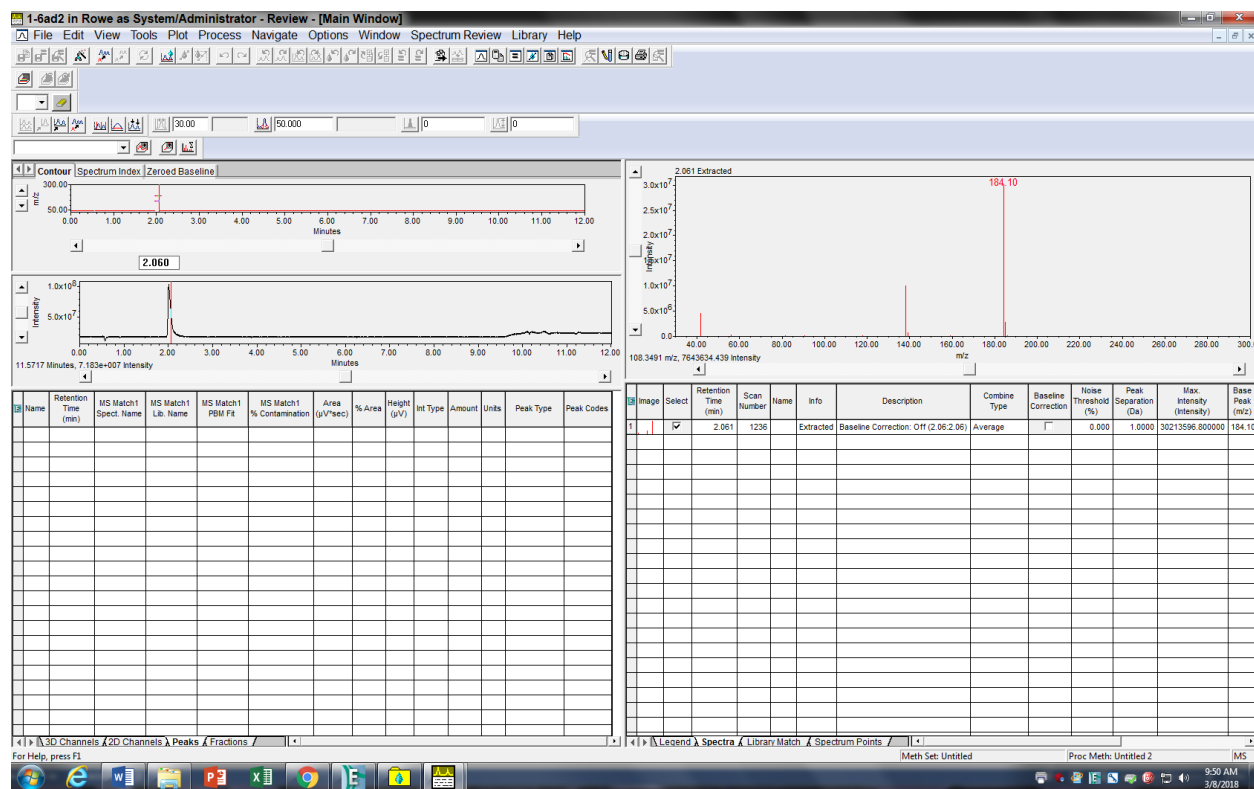


## 1-6 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.995	95629	100.00	31613	bb			Unknown

## 1-6 Mass Spectrum





The chromatogram displays absorbance (AU) on the y-axis (ranging from -0.010 to 0.160) against time in minutes on the x-axis (ranging from 0.00 to 12.00). A small peak is labeled at 1.696 - 1.84.16. A large, sharp peak is visible at approximately 10.0 minutes, followed by several smaller peaks between 10.5 and 11.5 minutes.

	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type	
1		1.696	90617	100.00	37460	bb			Unknown	

The screenshot displays the 1-Subv2 software interface, which is used for data analysis. The main window is titled "1-Subv2 in Rowe as System/Administrator - Review - [Main Window]".

The interface includes a menu bar (File, Edit, View, Tools, Plot, Process, Navigate, Options, Window, Spectrum Review, Library, Help) and a toolbar with various icons for file operations, plotting, and data manipulation.

The central area shows two plots:

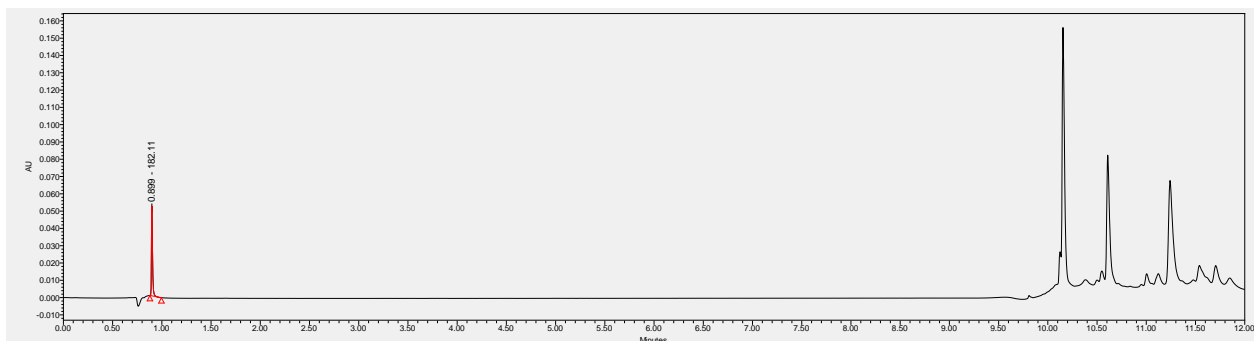
- Chromatogram (Top Left):** A plot of "Intensity" (Y-axis, ranging from 0.0x10<sup>7</sup> to 1.0x10<sup>8</sup>) versus "Minutes" (X-axis, ranging from 0.00 to 12.00). A single prominent peak is labeled "1.724". The plot is titled "Contour Spectrum Index Zeroed Baseline".
- Mass Spectrum (Top Right):** A plot of "Intensity" (Y-axis, ranging from 0 to 6x10<sup>7</sup>) versus "m/z" (X-axis, ranging from 40.00 to 300.00). A single prominent peak is labeled "184.16". The plot is titled "1.724 Extracted".

Below the plots is a table with the following columns:

Name	Retention Time (min)	MS Match1 Spect. Name	MS Match1 Lib. Name	MS Match1 PBM Fit	MS Match1 % Contamination	Area (μV*sec)	% Area	Height (μV)	Int Type	Amount	Units	Peak Type	Peak Codes
1	1.724												

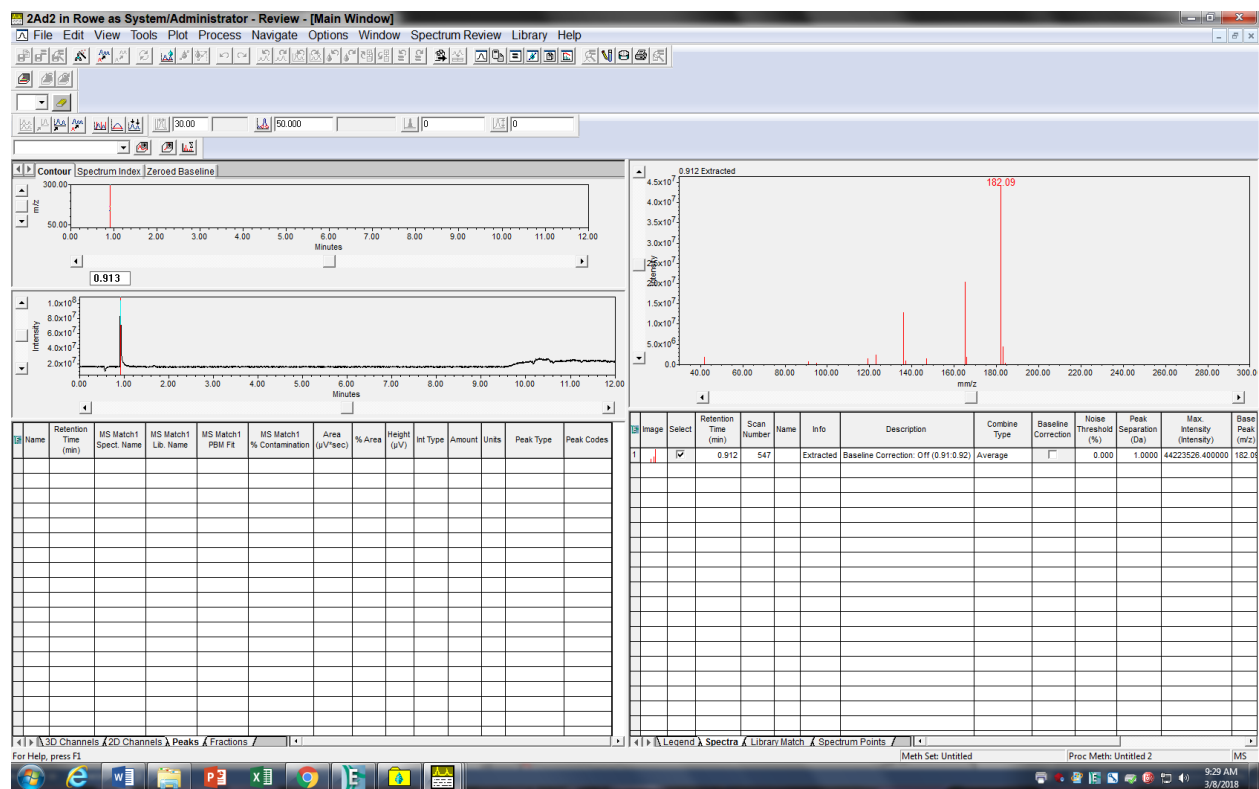
At the bottom of the interface, there is a status bar showing "1-Subv2 Channels 20 Channels 1 Peaks 1 Fractions / 1". The system clock in the bottom right corner indicates the date and time as 3/9/2013, 9:52 AM.

## 2 LC PDA Detector Data with Integrated Peak

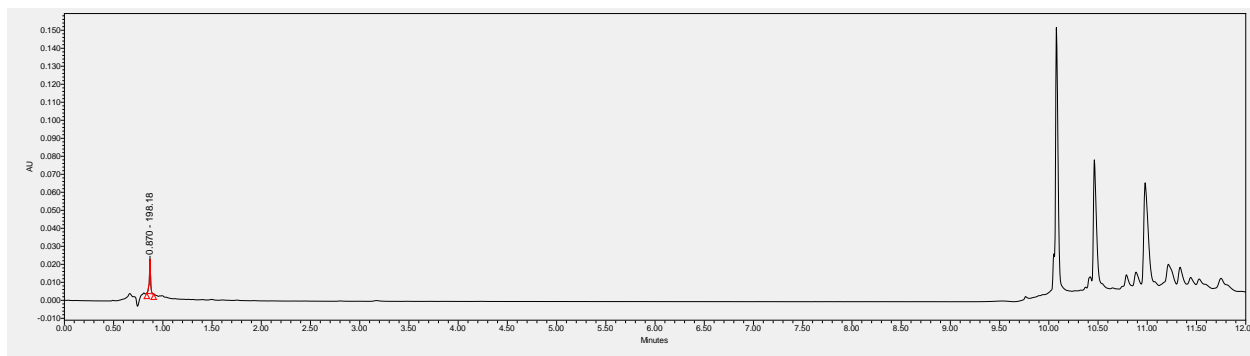


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.899	39379	100.00	52026	bb			Unknown

## 2 Mass Spectrum

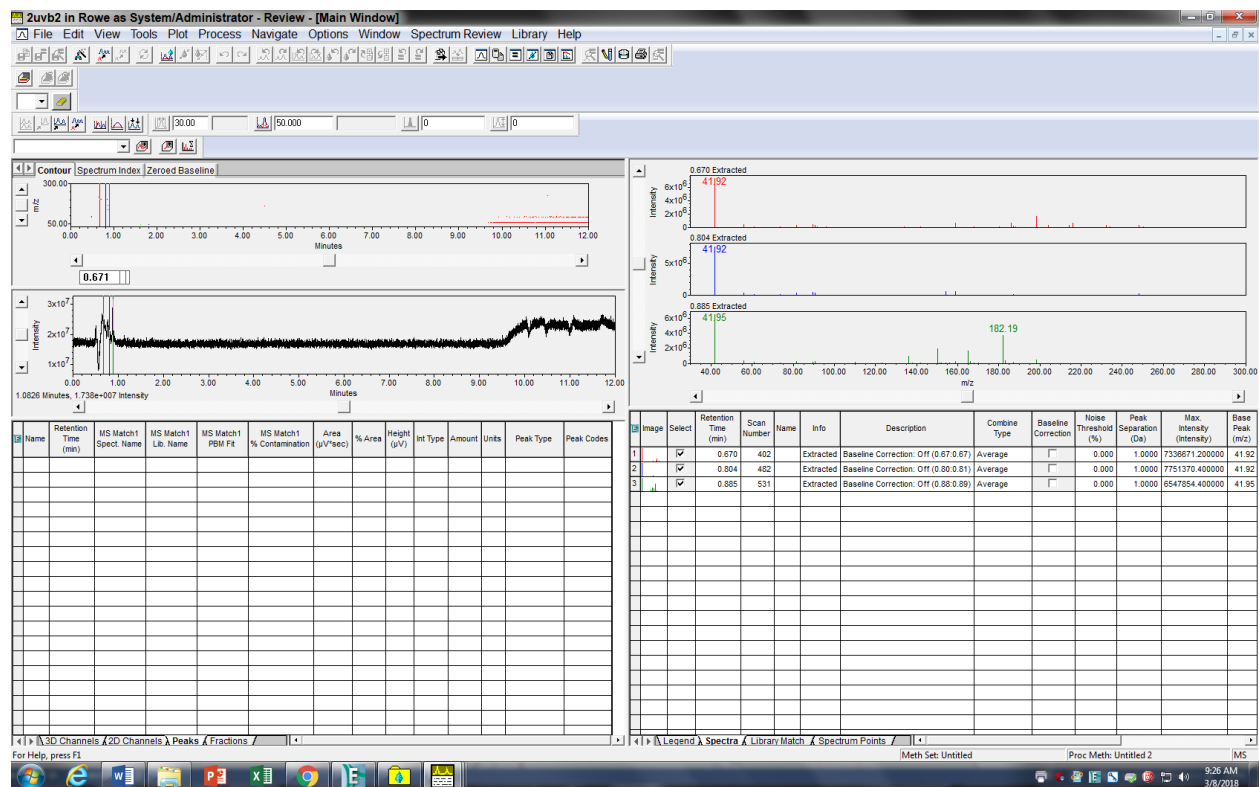


## 2 UV-B LC PDA Detector Data with Integrated Peak

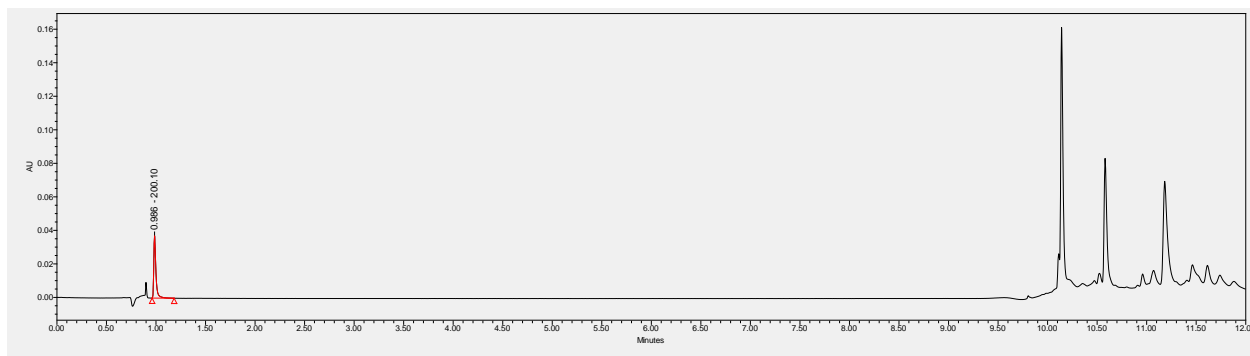


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.870	17015	100.00	18986	bb			Unknown

## 2 UVB Mass Spectrum

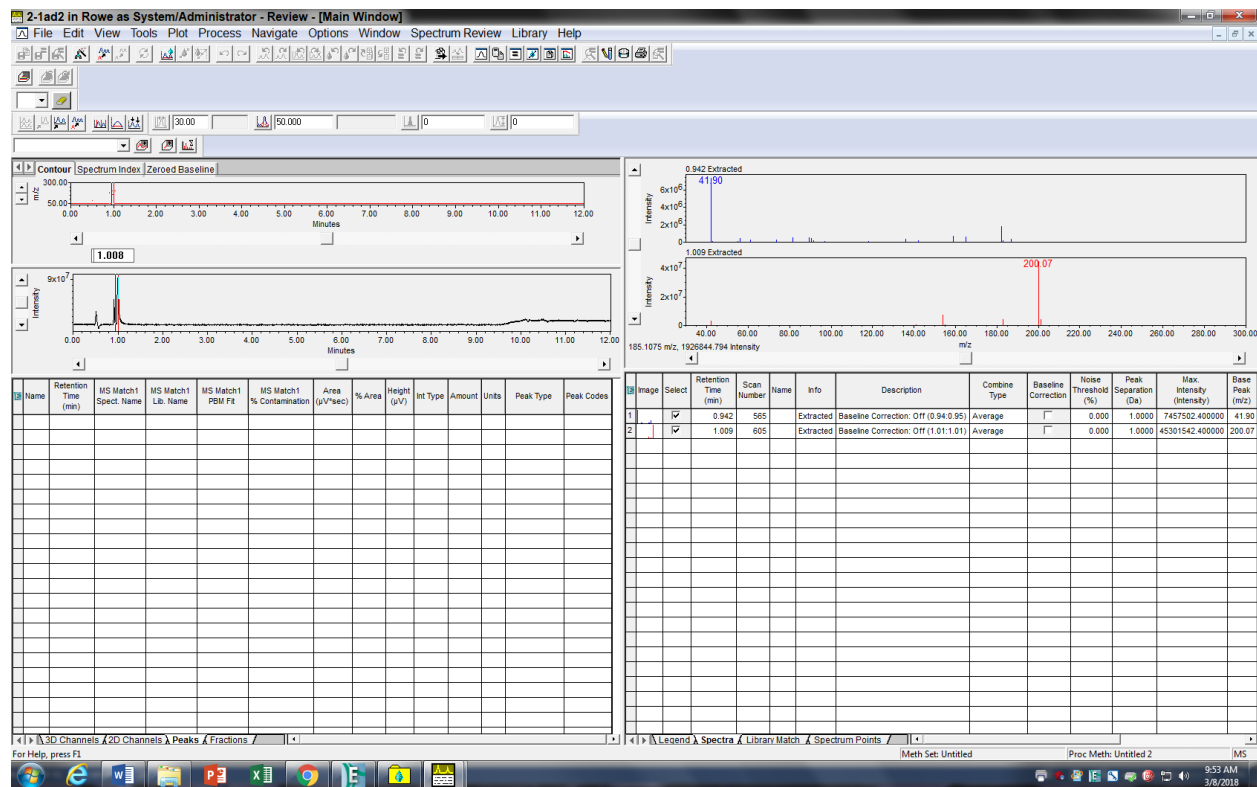


## 2-1 LC PDA Detector Data with Integrated Peak

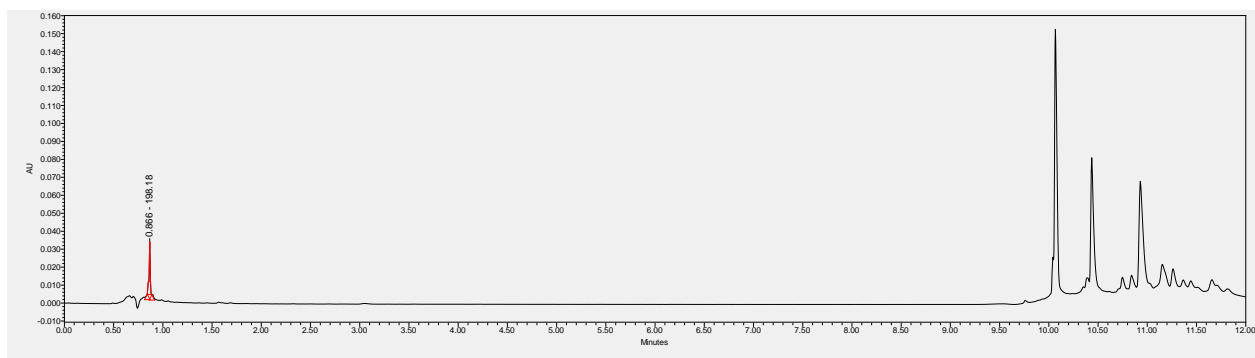


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.986	55681	100.00	37139	bb			Unknown

## 2-1 Mass Spectrum

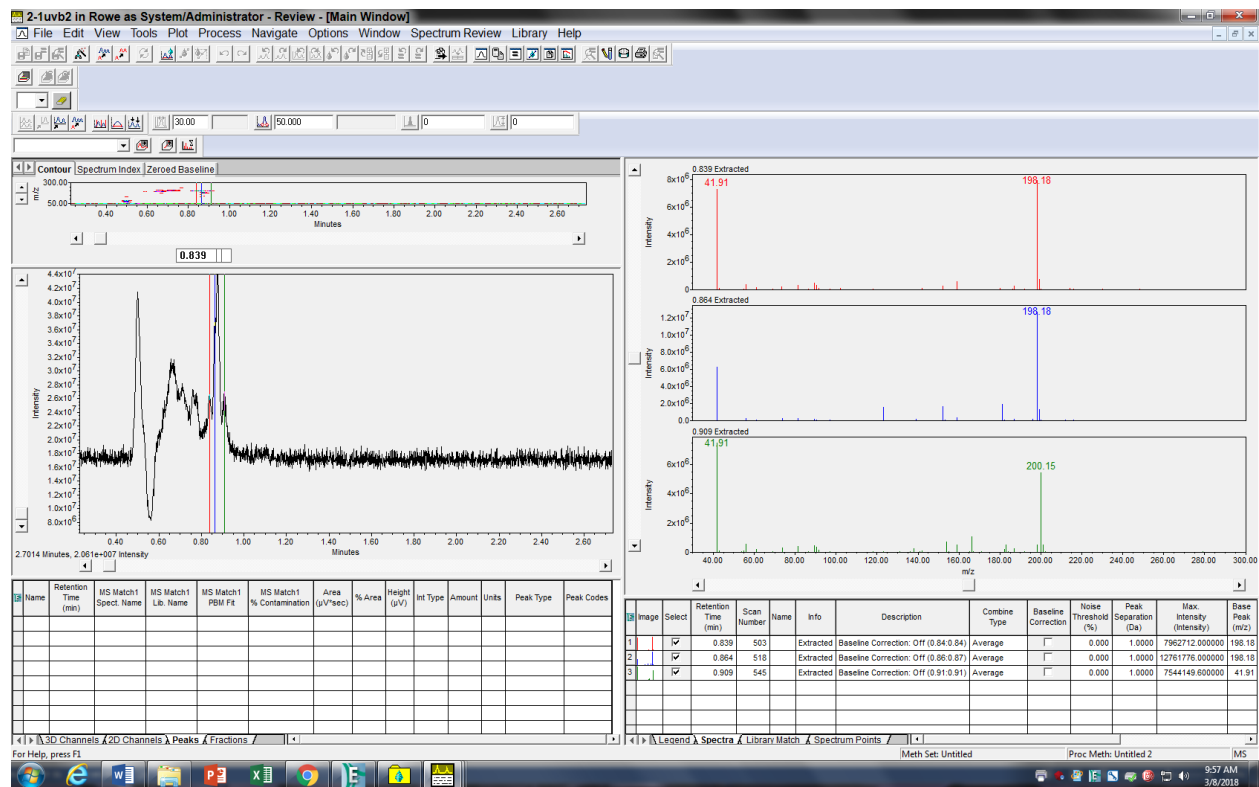


## 2-1 UV-B LC PDA Detector Data with Integrated Peak

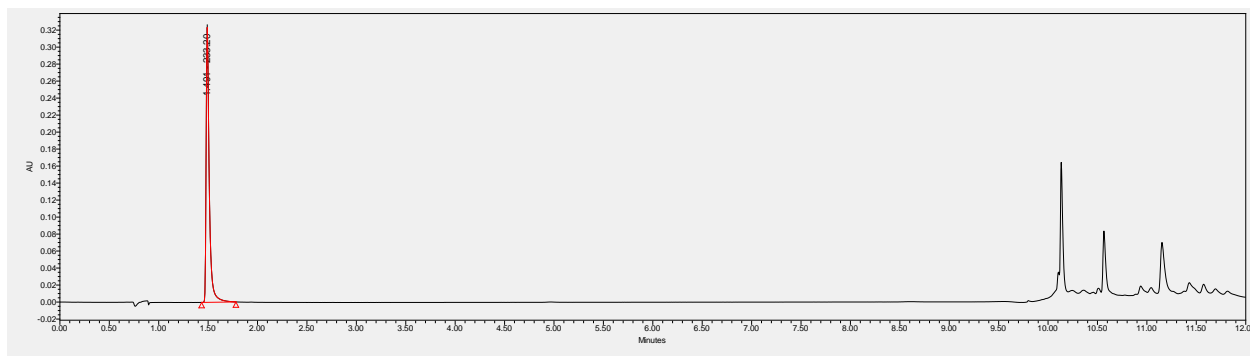


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.866	24197	100.00	29484	bb			Unknown

## 2-1 UVB Mass Spectrum

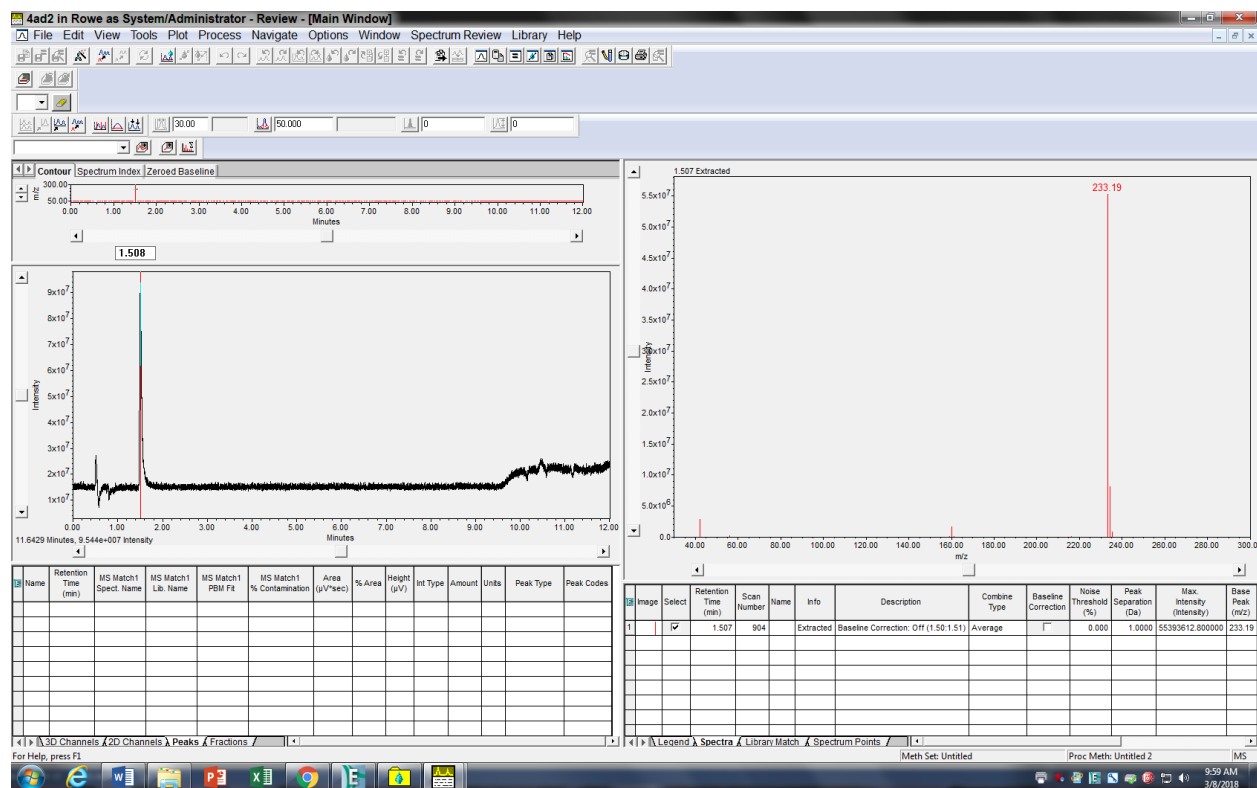


#### 4 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.491	716988	100.00	323502	bb			Unknown

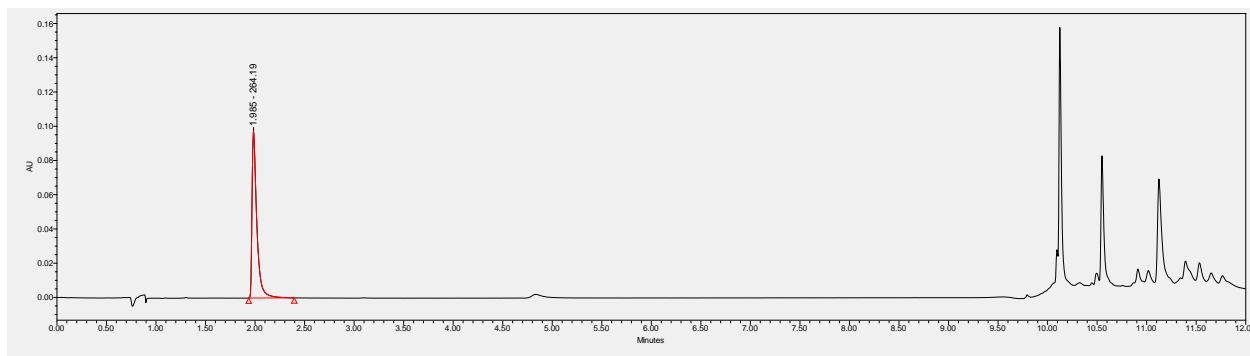
#### 4 Mass Spectrum



The chromatogram displays the detector response (AU) over a 12-minute period. The baseline is stable at approximately 0.005 AU. There are two small peaks at 1.1 and 1.4 minutes. A very large, sharp peak is observed at 10.1 minutes, reaching an AU of approximately 0.155. Two smaller peaks are visible at 10.4 and 10.9 minutes, with AU values of approximately 0.08 and 0.06, respectively. The baseline shows some minor noise and small fluctuations between 10.1 and 11.5 minutes.

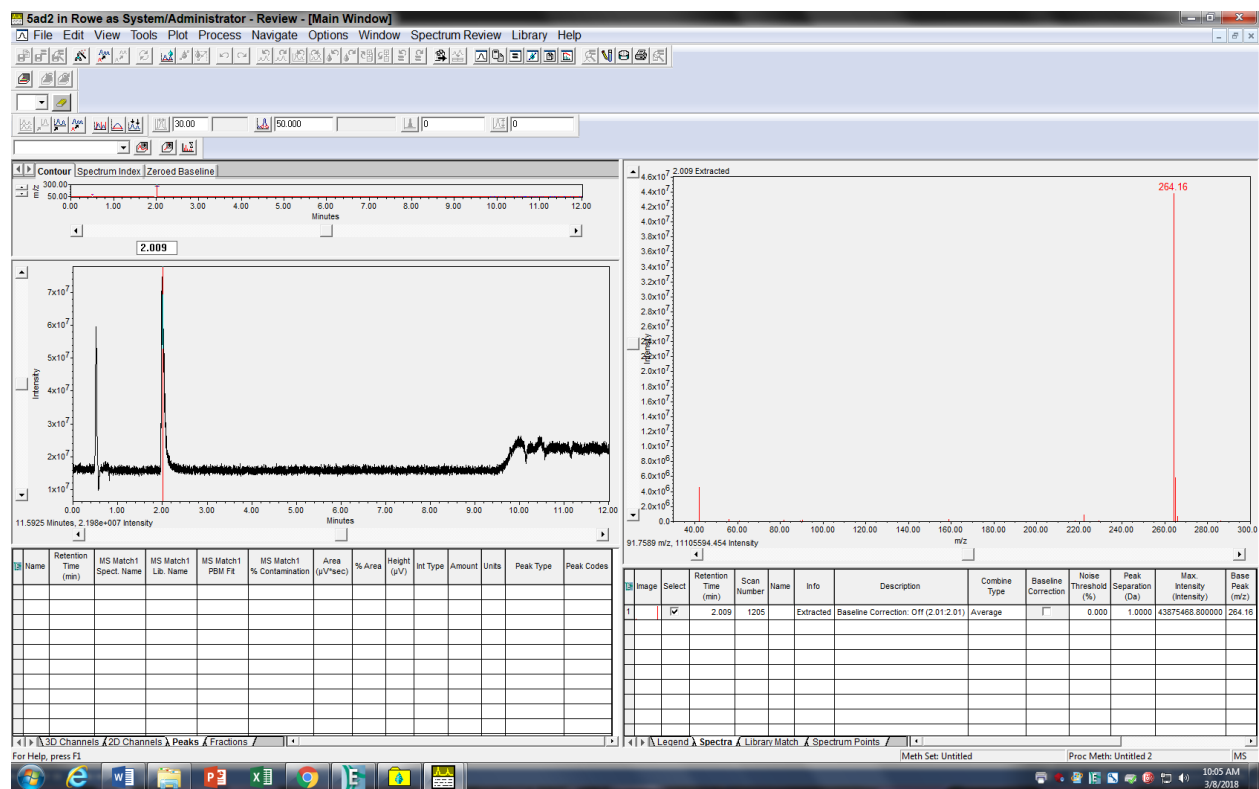
[illegible][illegible]

## 5 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.985	327286	100.00	97203	bb			Unknown

## 5 Mass Spectrum



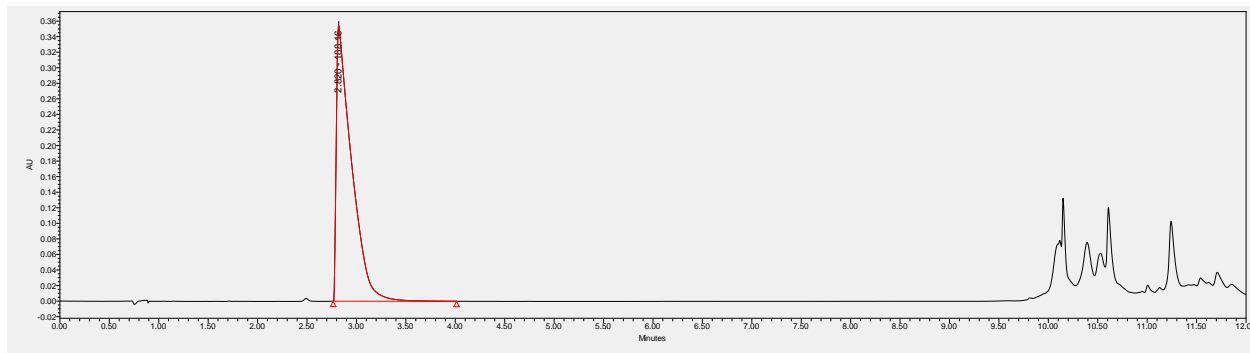


The chromatogram displays absorbance (AU) on the y-axis (ranging from -0.010 to 0.160) against time in minutes on the x-axis (ranging from 0.00 to 12.00). The baseline is relatively flat until approximately 9.50 minutes, where a series of peaks emerge. The most significant peak is at approximately 10.00 minutes, reaching an absorbance of about 0.155 AU. Other notable peaks are observed at approximately 10.30 AU (0.085 AU), 10.70 AU (0.065 AU), and 11.00 AU (0.045 AU). There are also smaller peaks at approximately 0.80 AU (0.065 AU) and 1.00 AU (0.025 AU).

The screenshot displays the Bruker TopSpin software interface. The main window shows a 1D  $^{13}\text{C}$  NMR spectrum with intensity on the y-axis (ranging from  $1.0 \times 10^7$  to  $4.0 \times 10^7$ ) and time in minutes on the x-axis (ranging from 0.20 to 2.60). Several peaks are identified and labeled with their retention times: 0.502, 0.664, 0.70, 0.7, 1.281, 1.475, and 1.980. A zoomed-in view of the spectrum is shown in the top right, highlighting the peaks at 0.502, 0.664, 0.70, 0.7, 1.281, 1.475, and 1.980 minutes. Below the spectrum, a table lists the peak data, including retention time, scan number, name, info, description, combine type, baseline correction, noise threshold, peak separation, max intensity, and peak code.

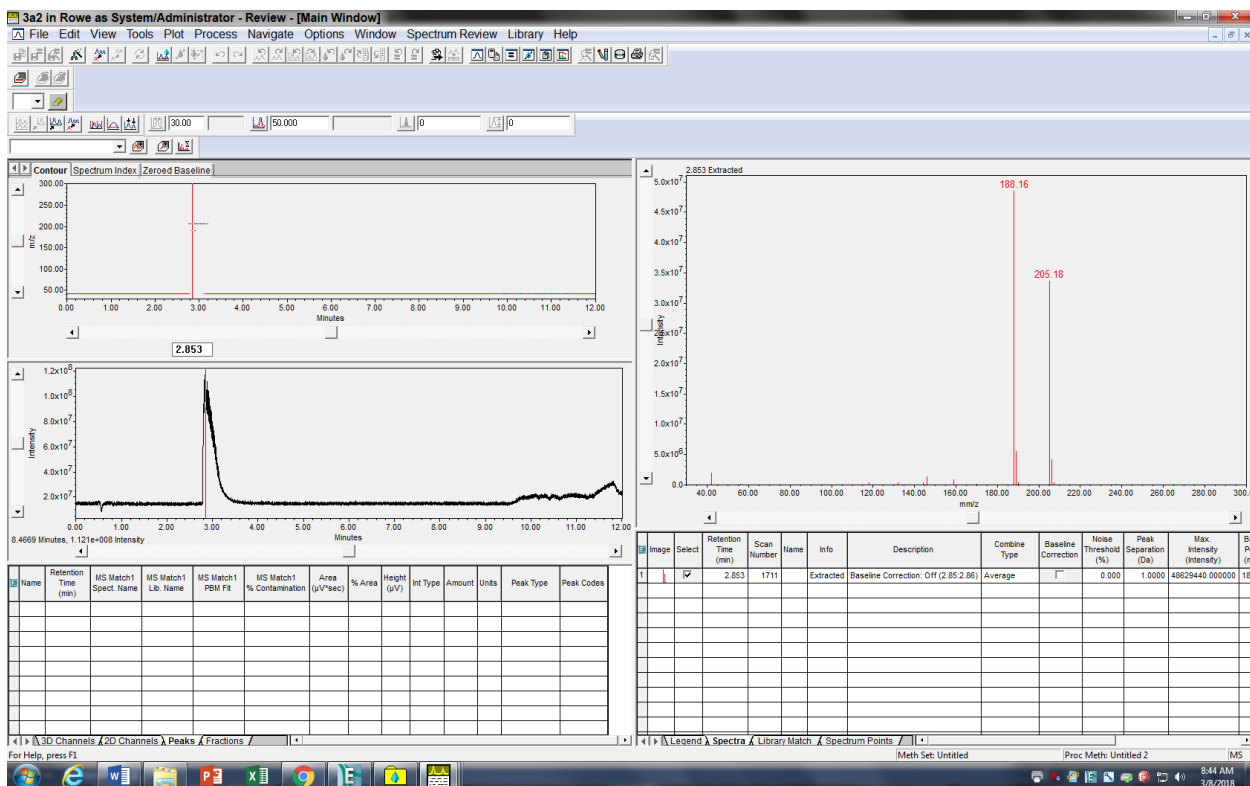
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity	Peak Code
1	<input checked="" type="checkbox"/>	0.502	301	Extracted	Baseline Correction: Off (0.50:0.51)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	20114795.600000	8	
2	<input checked="" type="checkbox"/>	0.664	396	Extracted	Baseline Correction: Off (0.66:0.67)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	7586345.600000	4	
3	<input checked="" type="checkbox"/>	0.759	455	Extracted	Baseline Correction: Off (0.76:0.76)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	7356643.200000	4	
4	<input checked="" type="checkbox"/>	0.870	522	Extracted	Baseline Correction: Off (0.87:0.87)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	6925617.600000	4	
5	<input checked="" type="checkbox"/>	0.884	530	Extracted	Baseline Correction: Off (0.88:0.88)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	7072172.800000	4	
6	<input checked="" type="checkbox"/>	0.967	580	Extracted	Baseline Correction: Off (0.96:0.97)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	8037496.600000	28	
7	<input checked="" type="checkbox"/>	1.281	768	Extracted	Baseline Correction: Off (1.28:1.28)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	7825329.600000	4	
8	<input checked="" type="checkbox"/>	1.299	779	Extracted	Baseline Correction: Off (1.30:1.30)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	8143479.200000	4	
9	<input checked="" type="checkbox"/>	1.476	885	Extracted	Baseline Correction: Off (1.47:1.48)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	8040516.600000	4	

### 3 LC PDA Detector Data with Integrated Peak

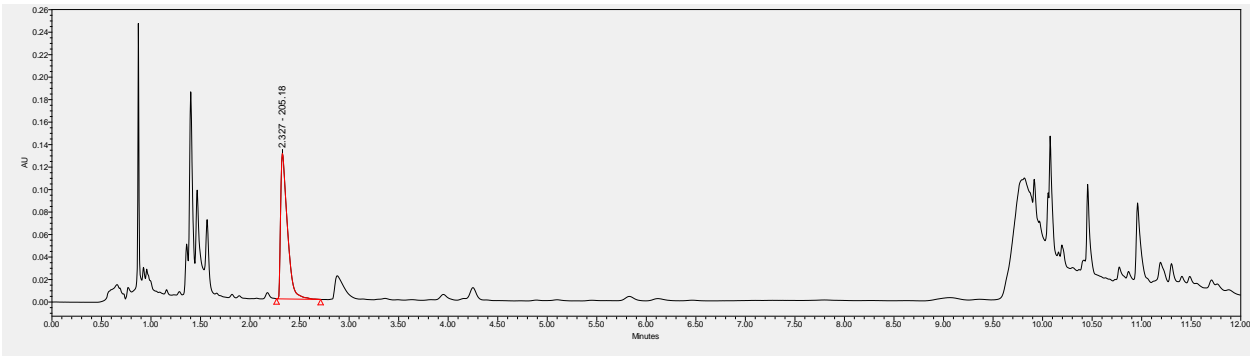


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.820	3792679	100.00	354425	bb			Unknown

### 3 Mass Spectrum

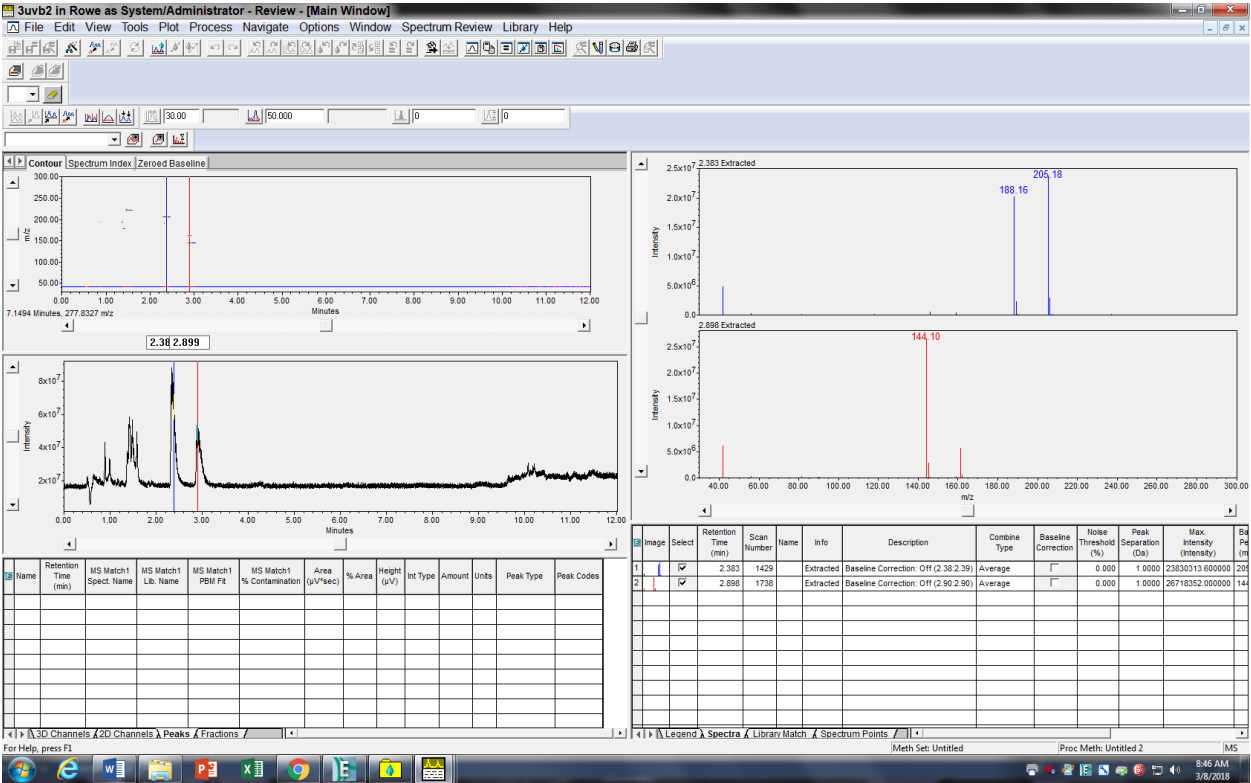


3 UV-B LC PDA Detector Data with Integrated Peak

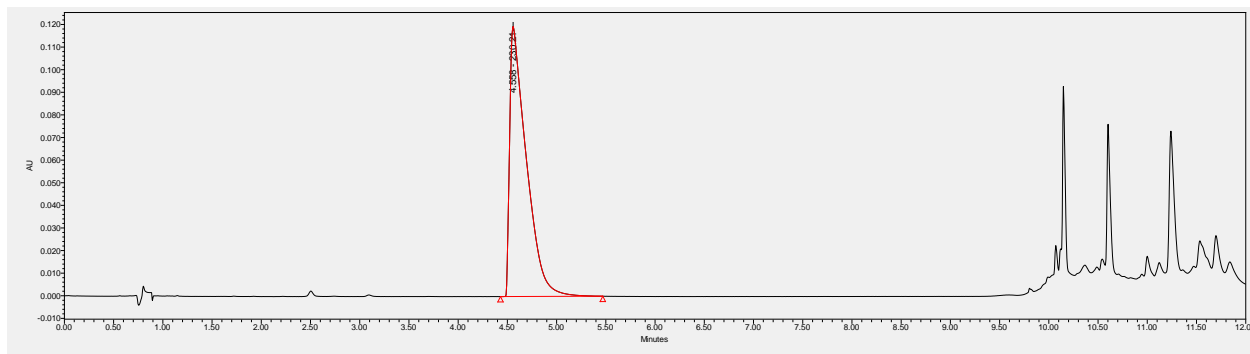


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.327	615698	100.00	129284	bb			Unknown

3 UVB Mass Spectrum

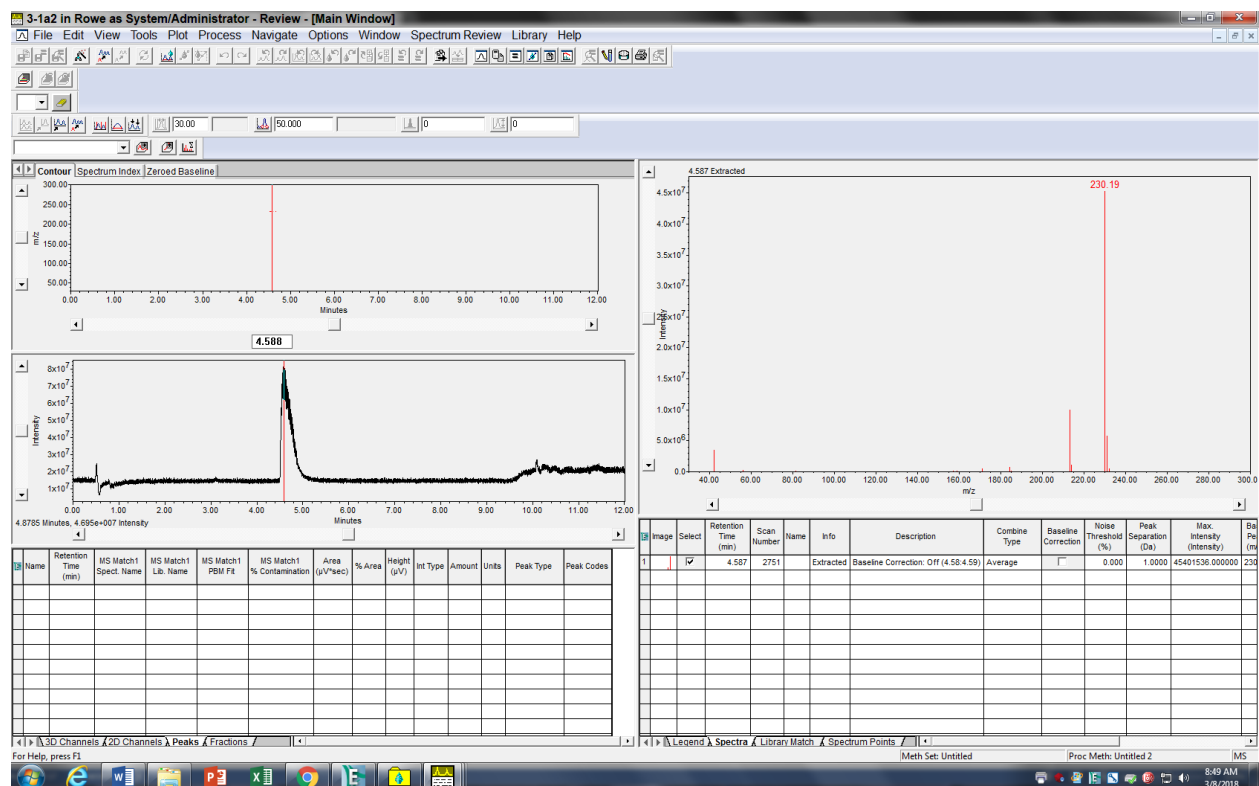


### 3-1 LC PDA Detector Data with Integrated Peak

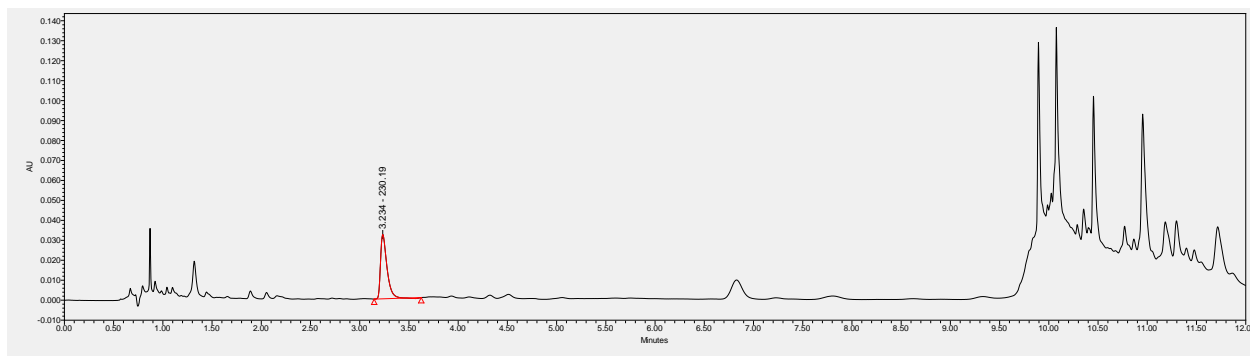


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.558	1402762	100.00	119451	bb			Unknown

### 3-1 Mass Spectrum

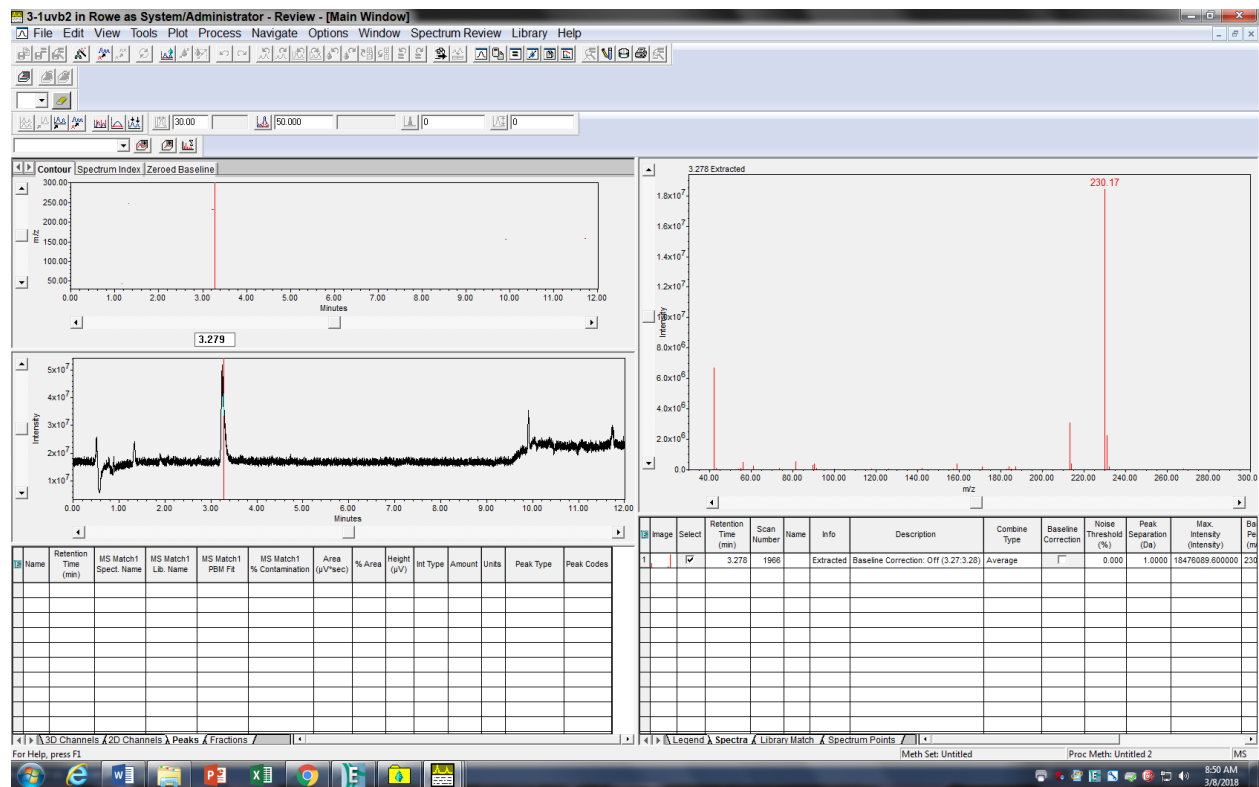


### 3-1 UV-B LC PDA Detector Data with Integrated Peak

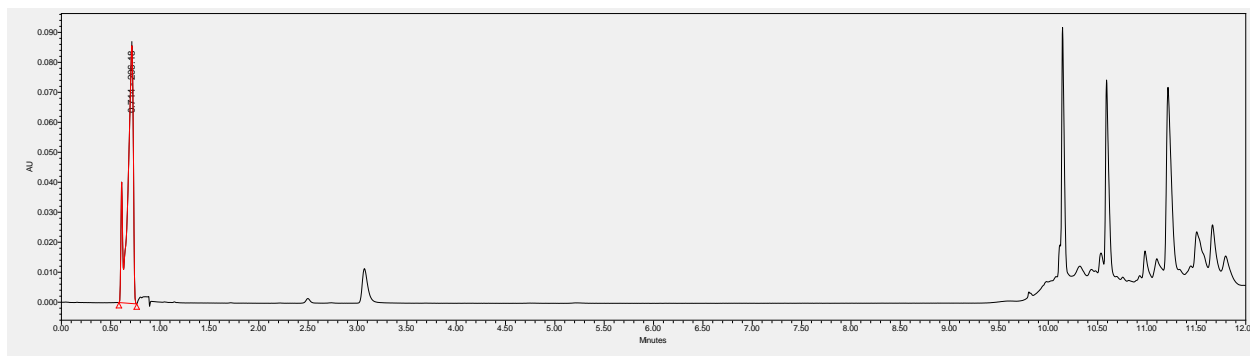


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.234	148564	100.00	32255	bb			Unknown

### 3-1 UVB Mass Spectrum

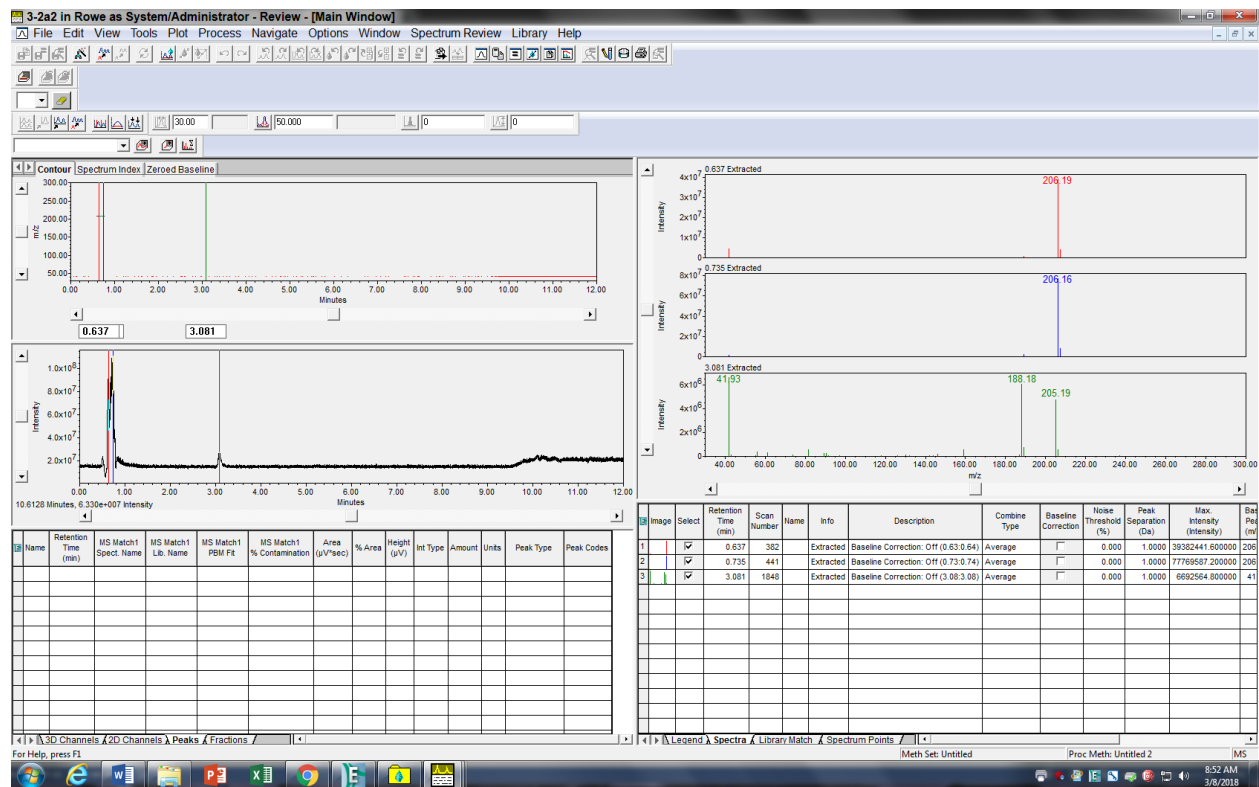


### 3-2 LC PDA Detector Data with Integrated Peak

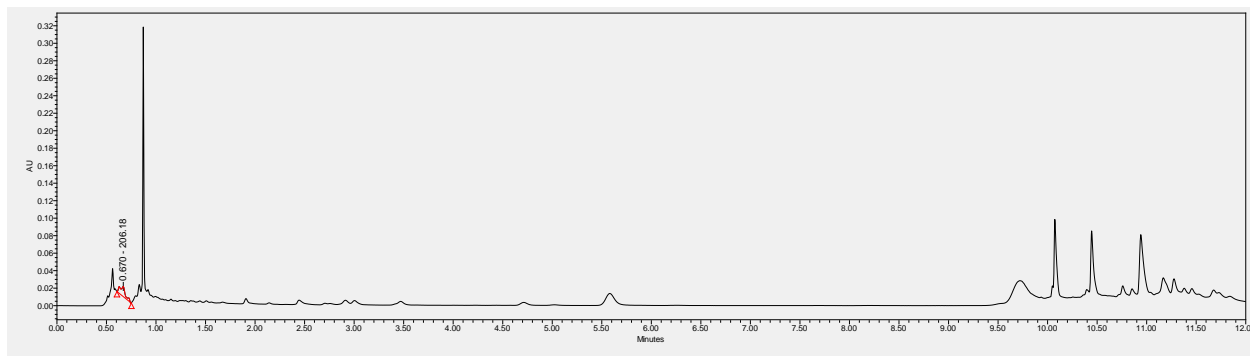


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.714	345619	100.00	85986	bb			Unknown

### 3-2 Mass Spectrum

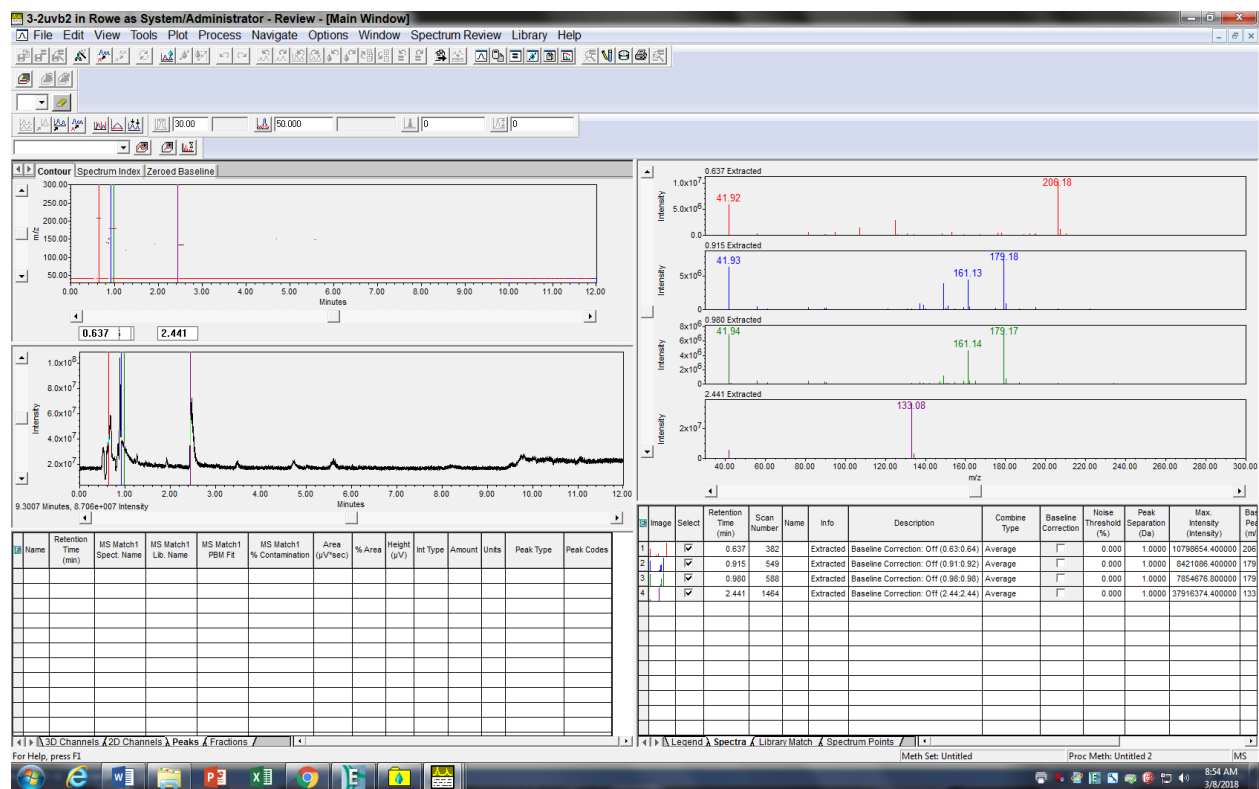


### 3-2 UV-B LC PDA Detector Data with Integrated Peak

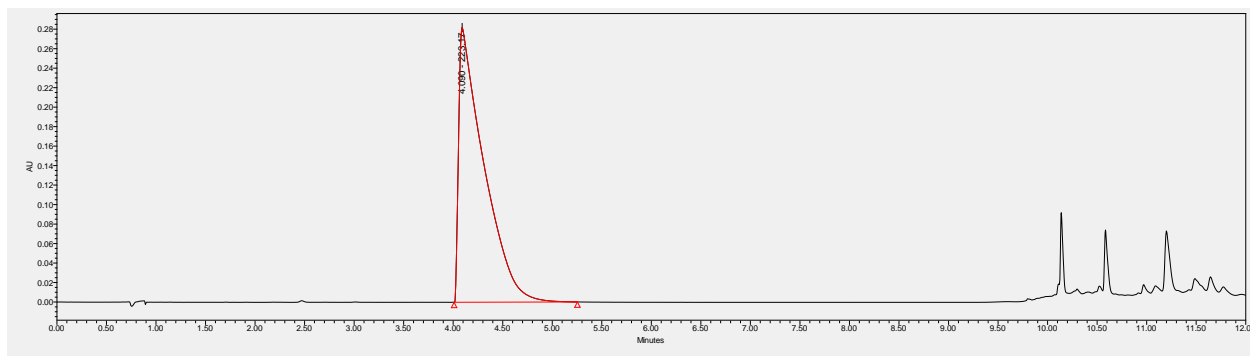


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.670	42294	100.00	11676	bb			Unknown

### 3-2 UVB Mass Spectrum

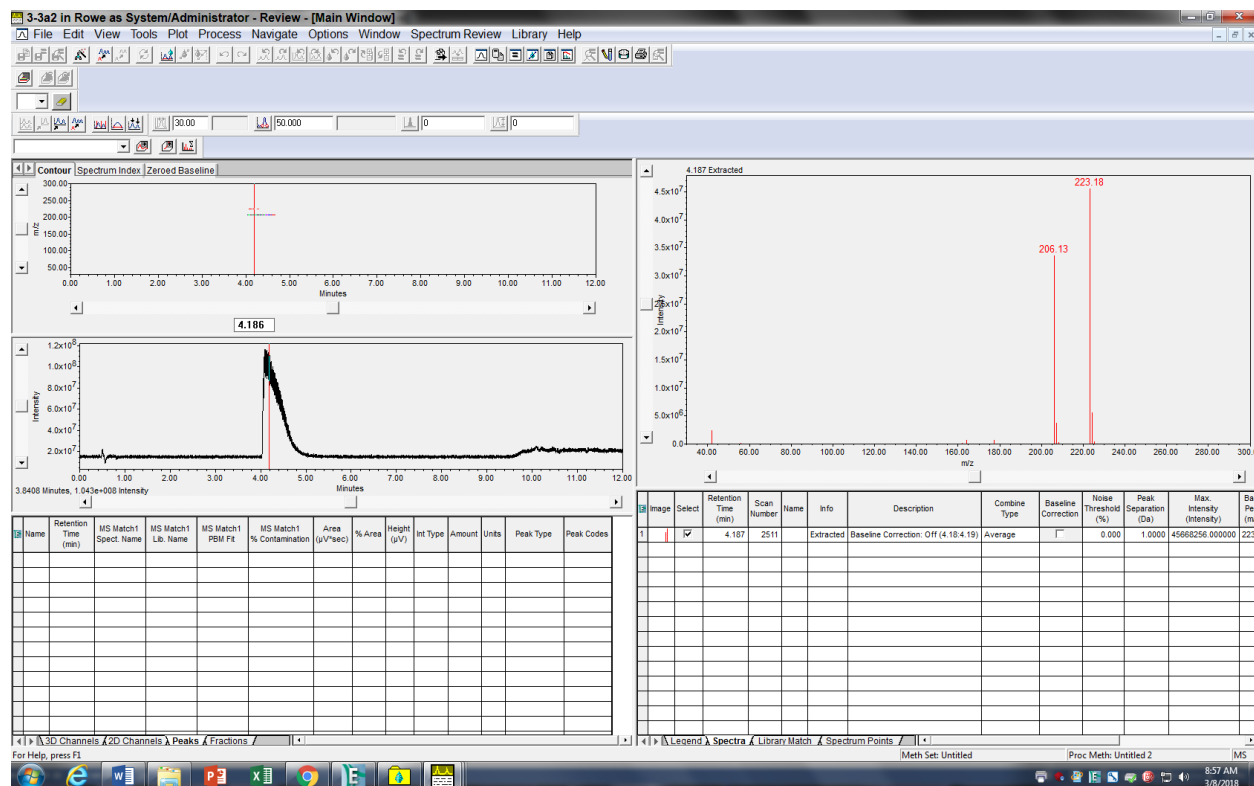


### 3-3 LC PDA Detector Data with Integrated Peak



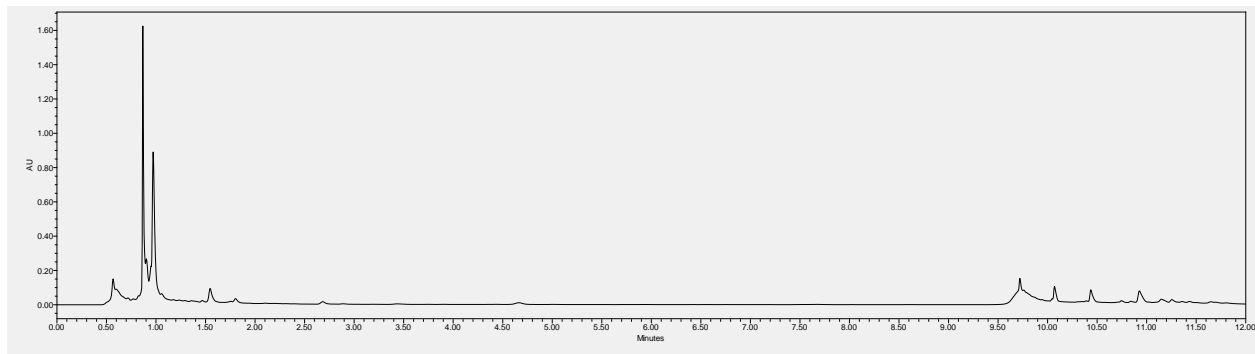
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.090	4951822	100.00	281952	bb			Unknown

### 3-3 Mass Spectrum

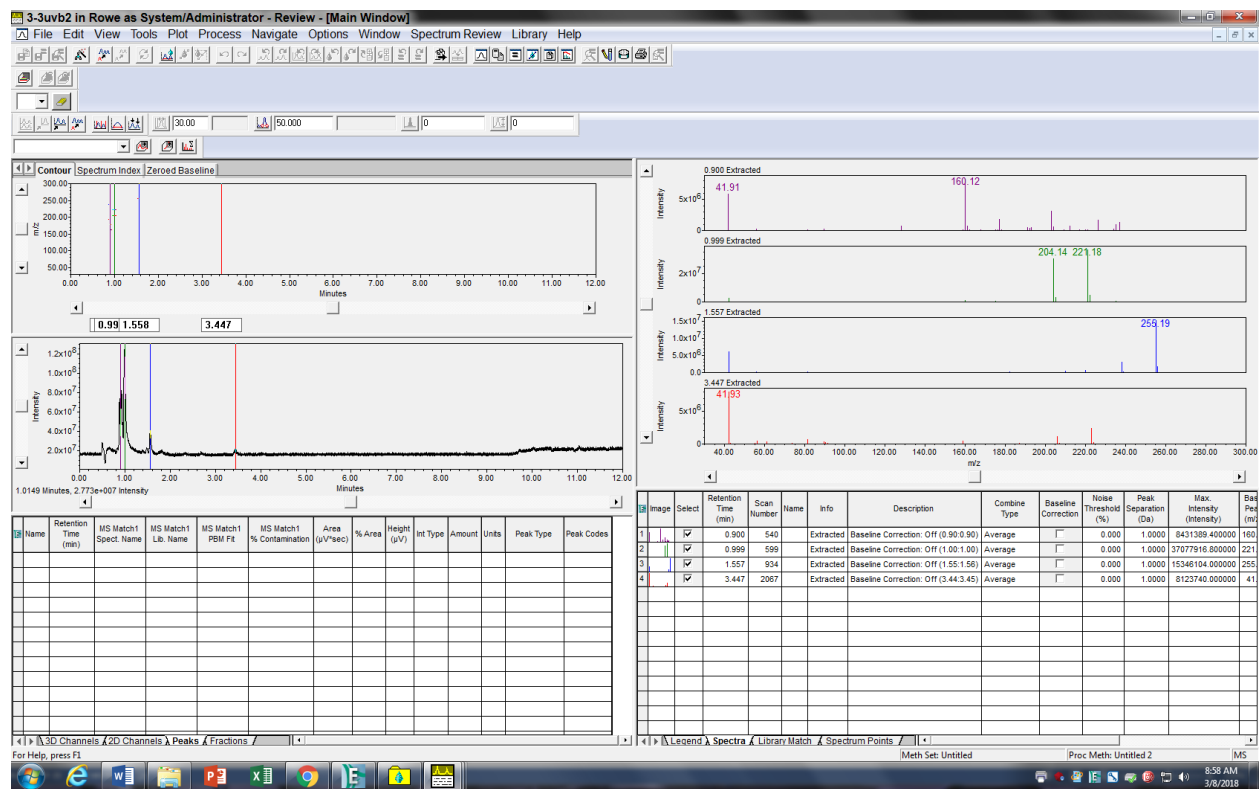




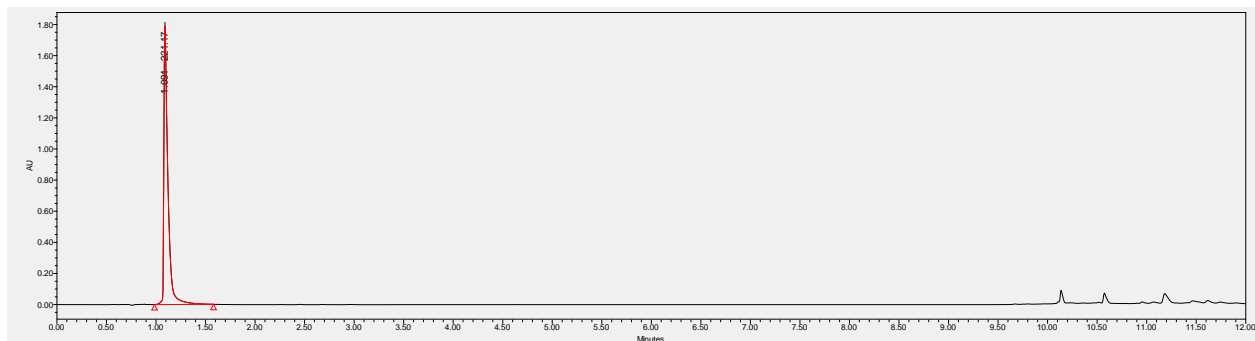
### 3-3 UV-B LC PDA Detector Data with Integrated Peak



### 3-3 UVB Mass Spectrum

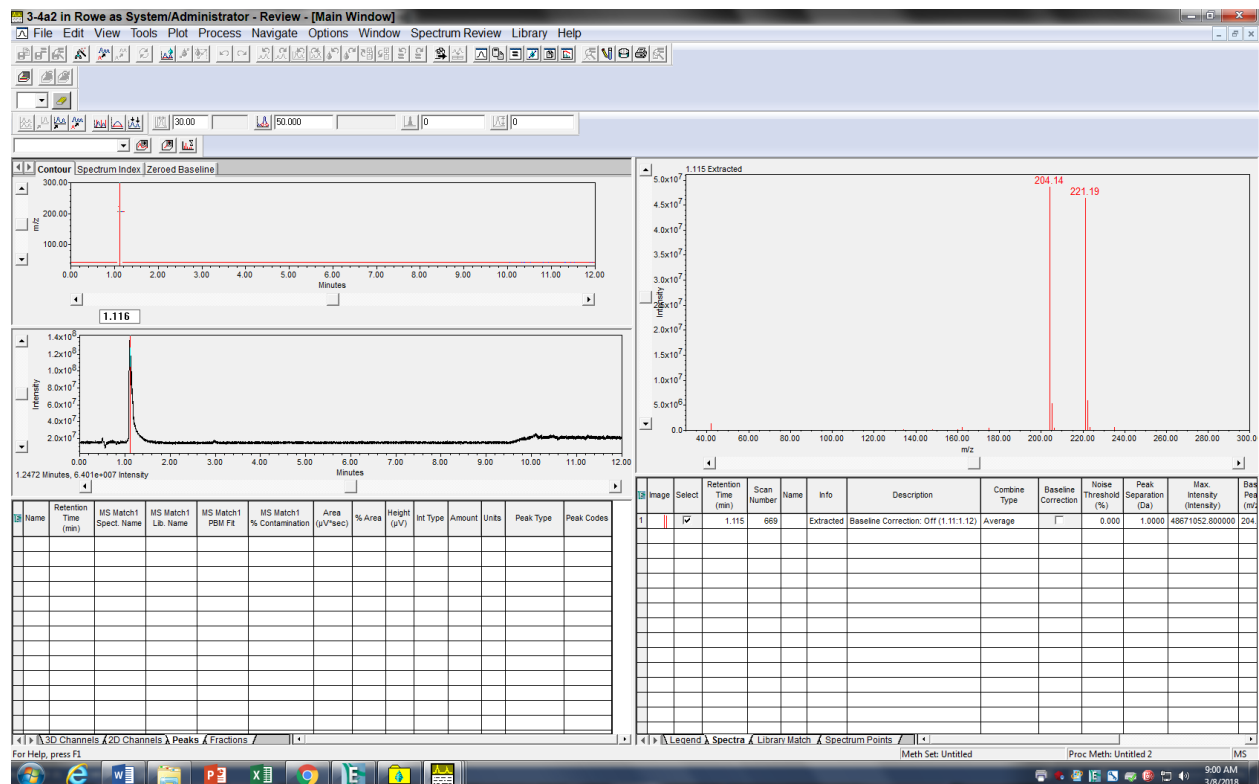


### 3-4 LC PDA Detector Data with Integrated Peak

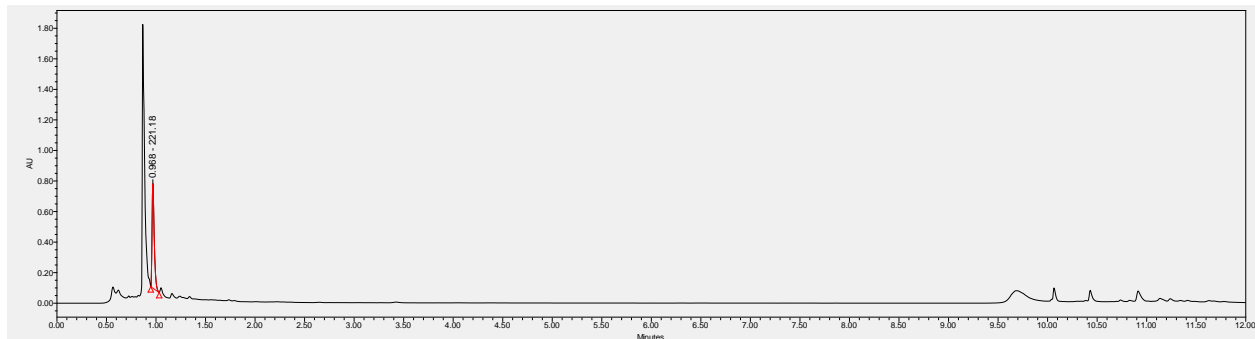


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.091	5071661	100.00	1787026	bb			Unknown

### 3-4 Mass Spectrum

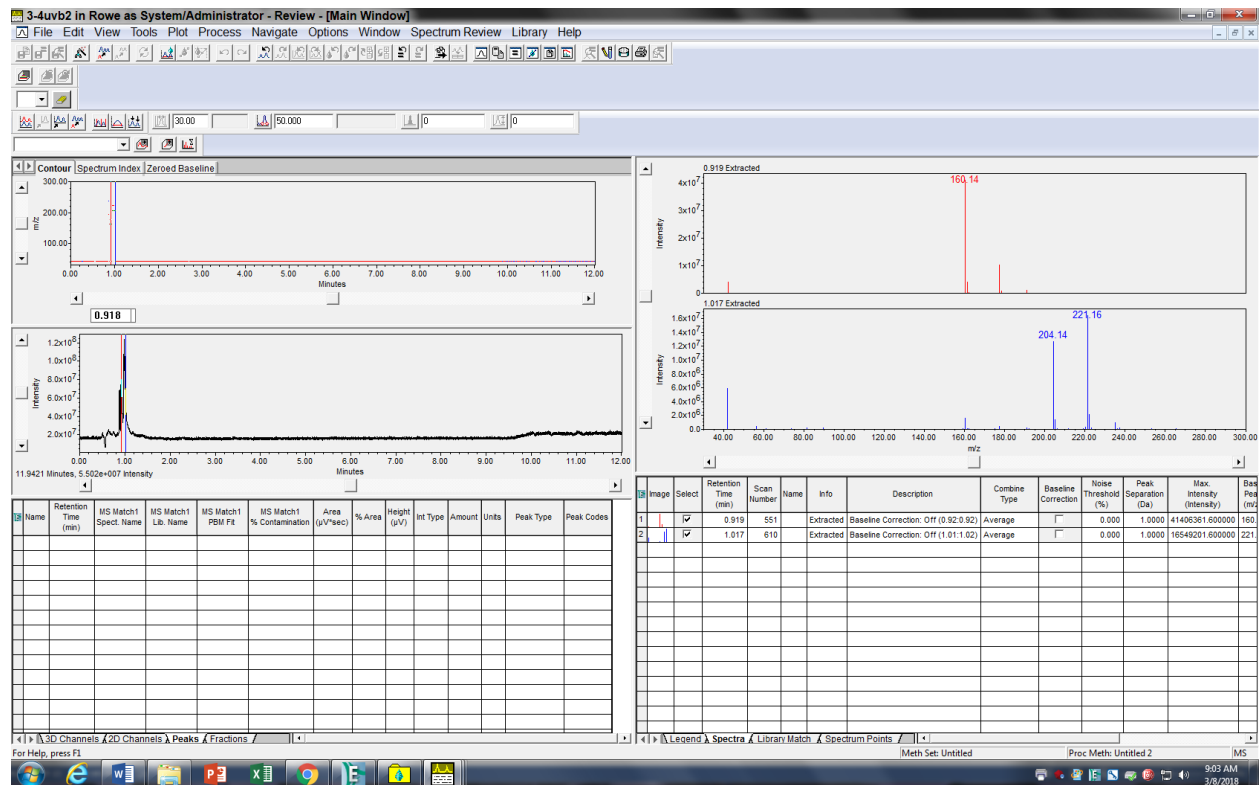


### 3-4 UV-B LC PDA Detector Data with Integrated Peak

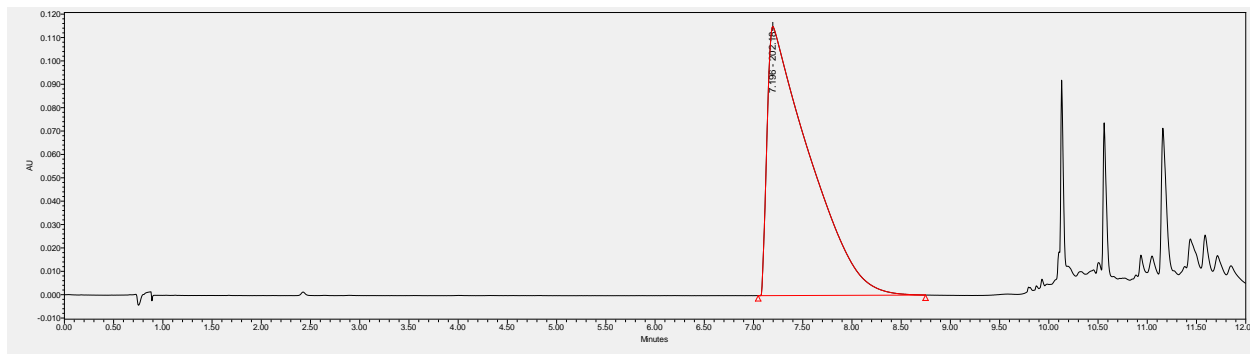


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.968	1043865	100.00	683634	bb			Unknown

### 3-4 UVB Mass Spectrum

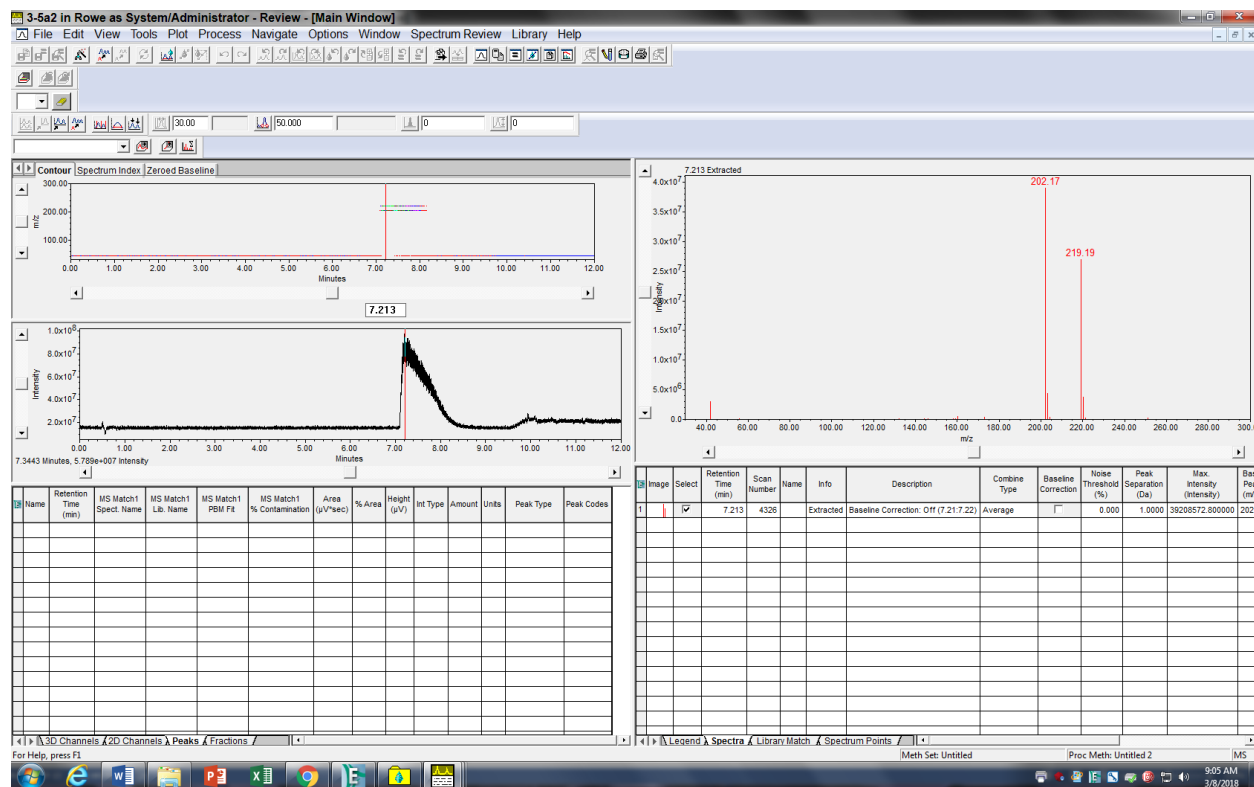


### 3-5 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		7.196	3441249	100.00	115120	bb			Unknown

### 3-5 Mass Spectrum



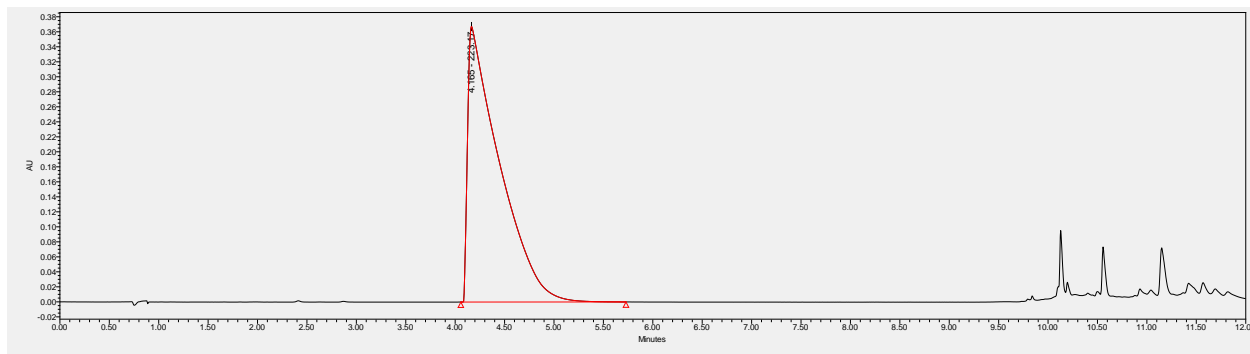
The chromatogram displays absorbance (AU) on the y-axis (ranging from 0.00 to 0.16) against time in minutes on the x-axis (ranging from 0.00 to 12.00). A prominent peak is observed at approximately 5.4 minutes, which is highlighted with a red triangle and labeled with the retention times 5.399 and 219.20. Other significant peaks are visible at approximately 0.9, 2.9, 3.4, and a large cluster of peaks starting around 9.5 minutes, peaking at approximately 9.8 minutes.

3-5uvb2 in Row as System/Administrator - Review - [Main Window]

File Edit View Tools Plot Process Navigate Options Spectrum Review Library Help

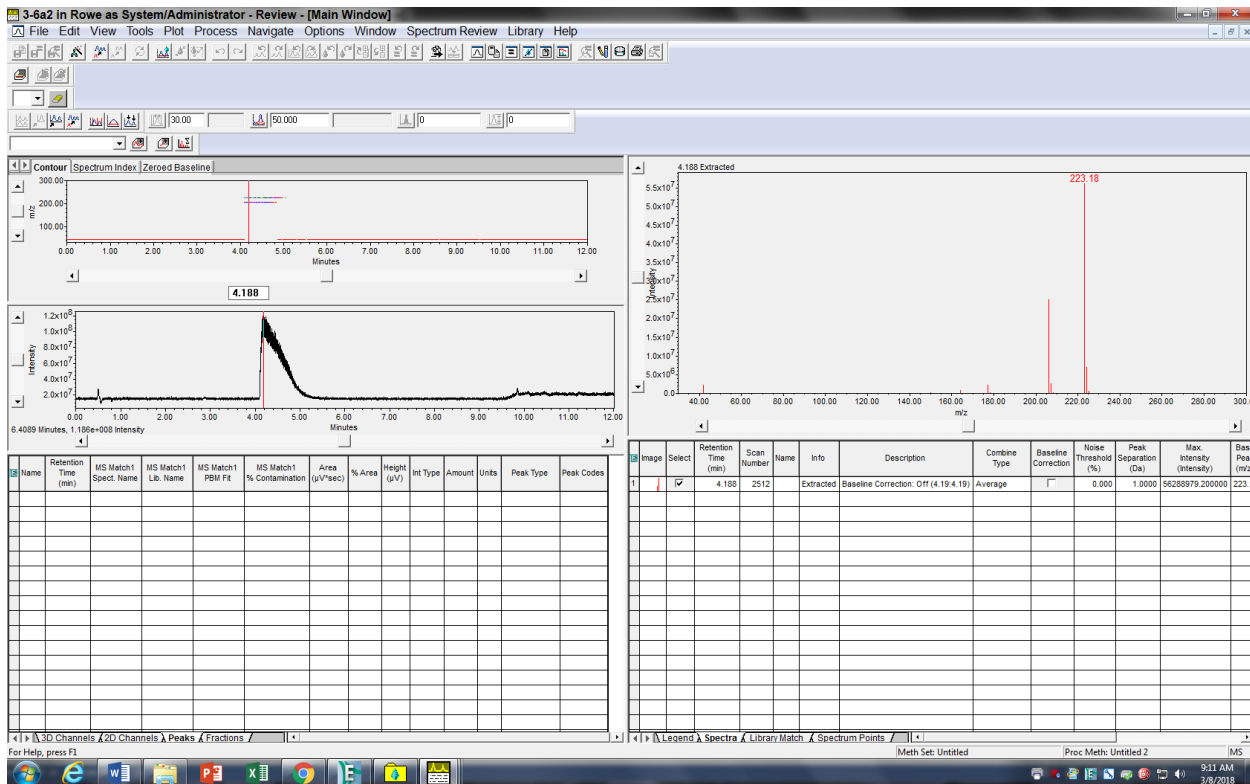
</

### 3-6 LC PDA Detector Data with Integrated Peak

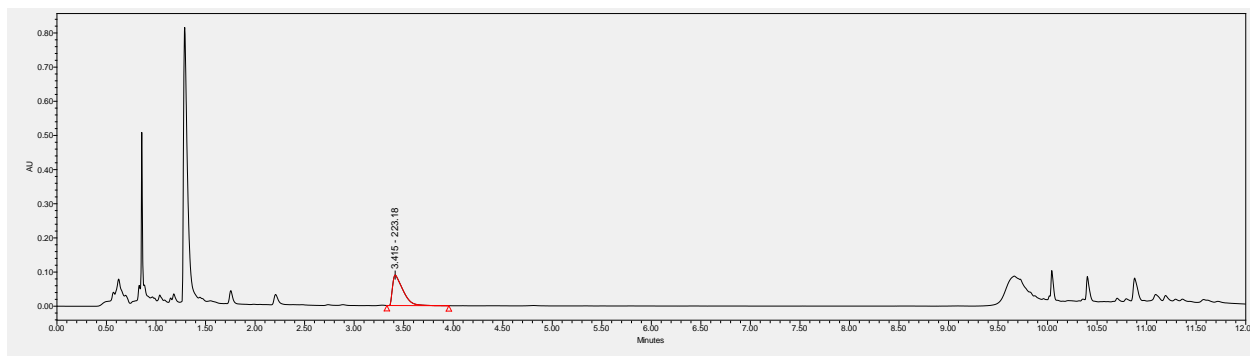


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.165	8184661	100.00	367356	bb			Unknown

### 3-6 Mass Spectrum

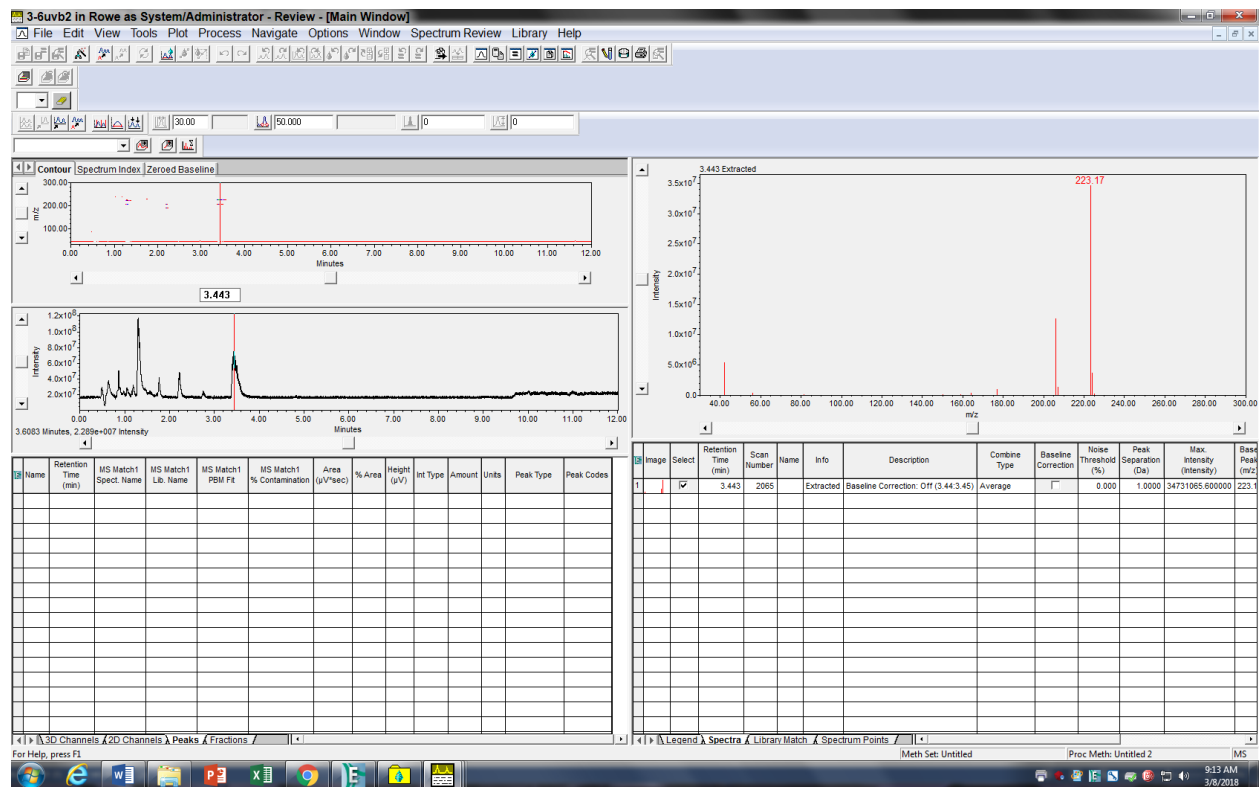


### 3-6 UV-B LC PDA Detector Data with Integrated Peak

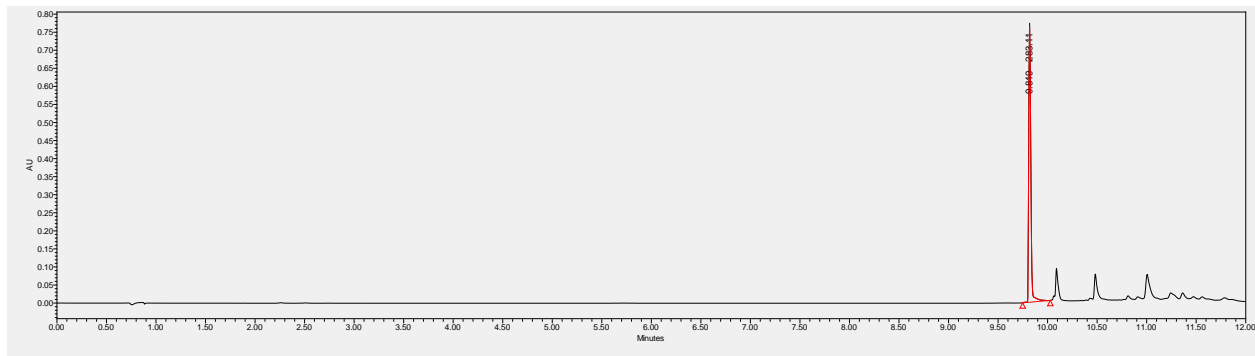


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.415	648197	100.00	89369	bb			Unknown

### 3-6 UVB Mass Spectrum

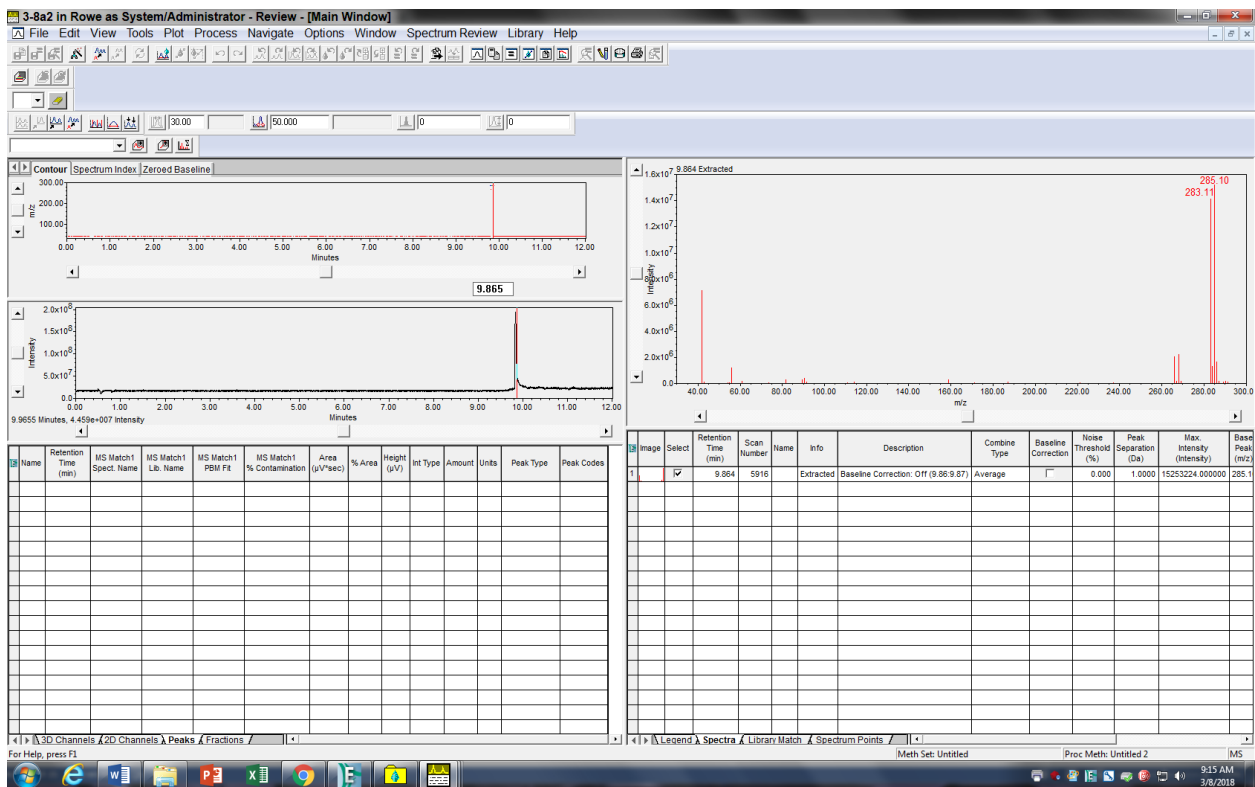


### 3-8 LC PDA Detector Data with Integrated Peak



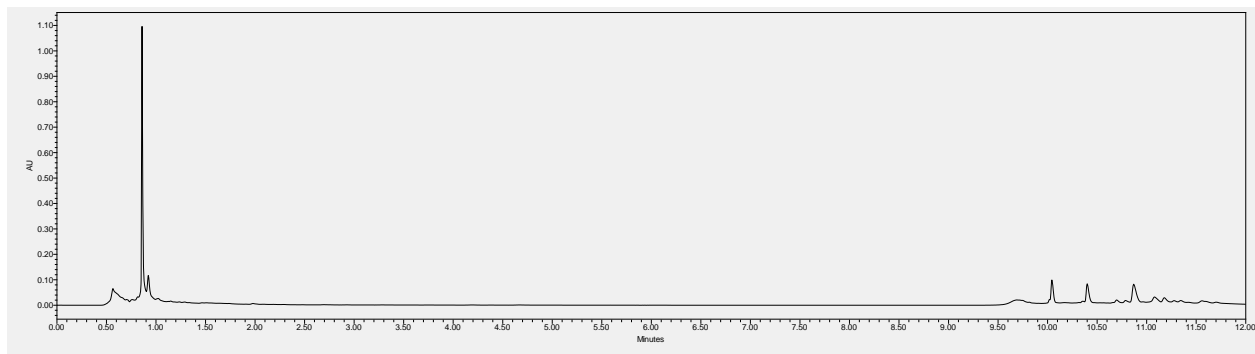
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.819	1148094	100.00	765144	bb			Unknown

### 3-8 Mass Spectrum

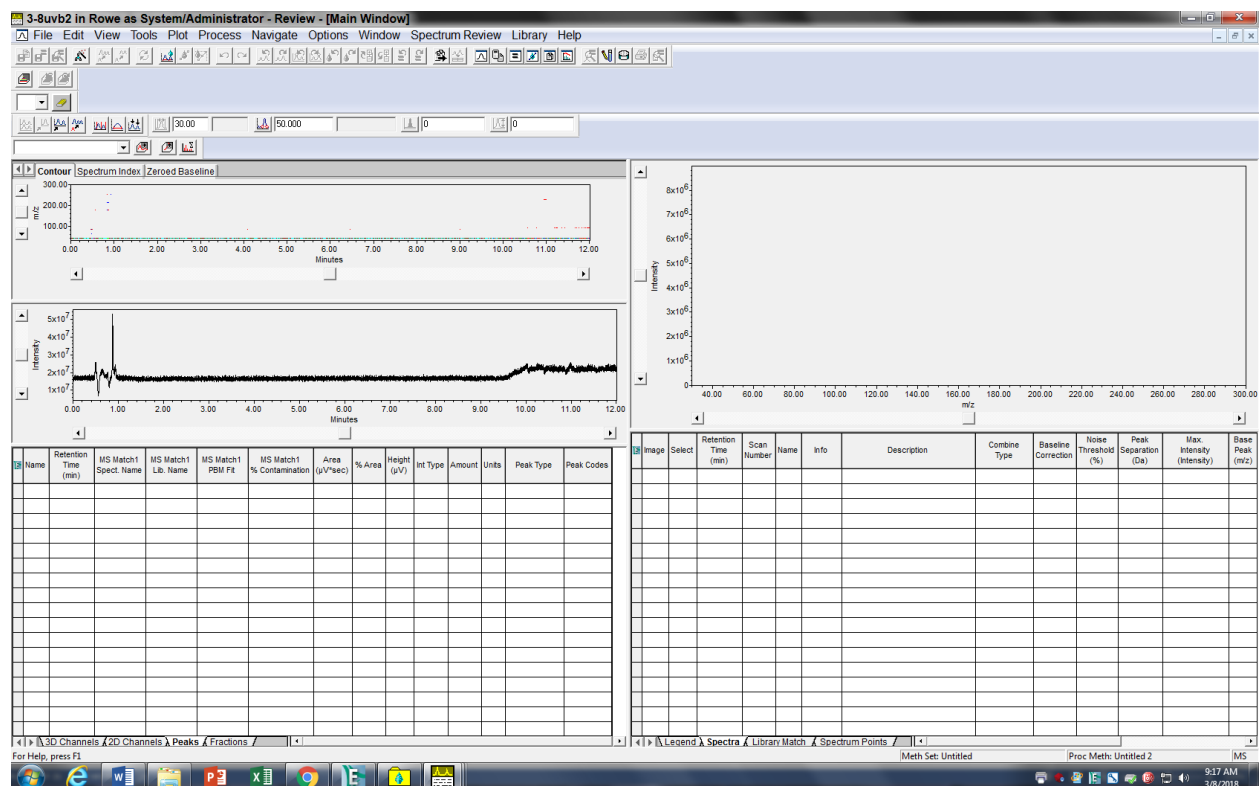




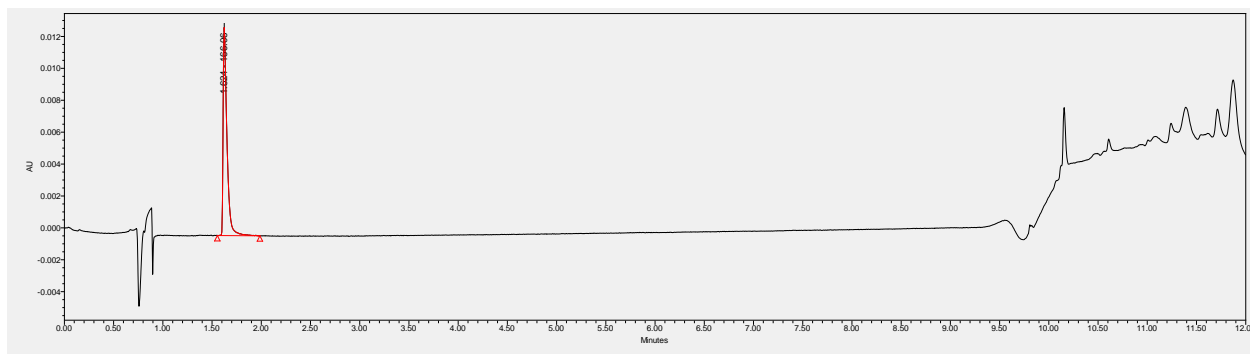
### 3-8 UV-B LC PDA Detector Data with Integrated Peak



### 3-8 UVB Mass Spectrum

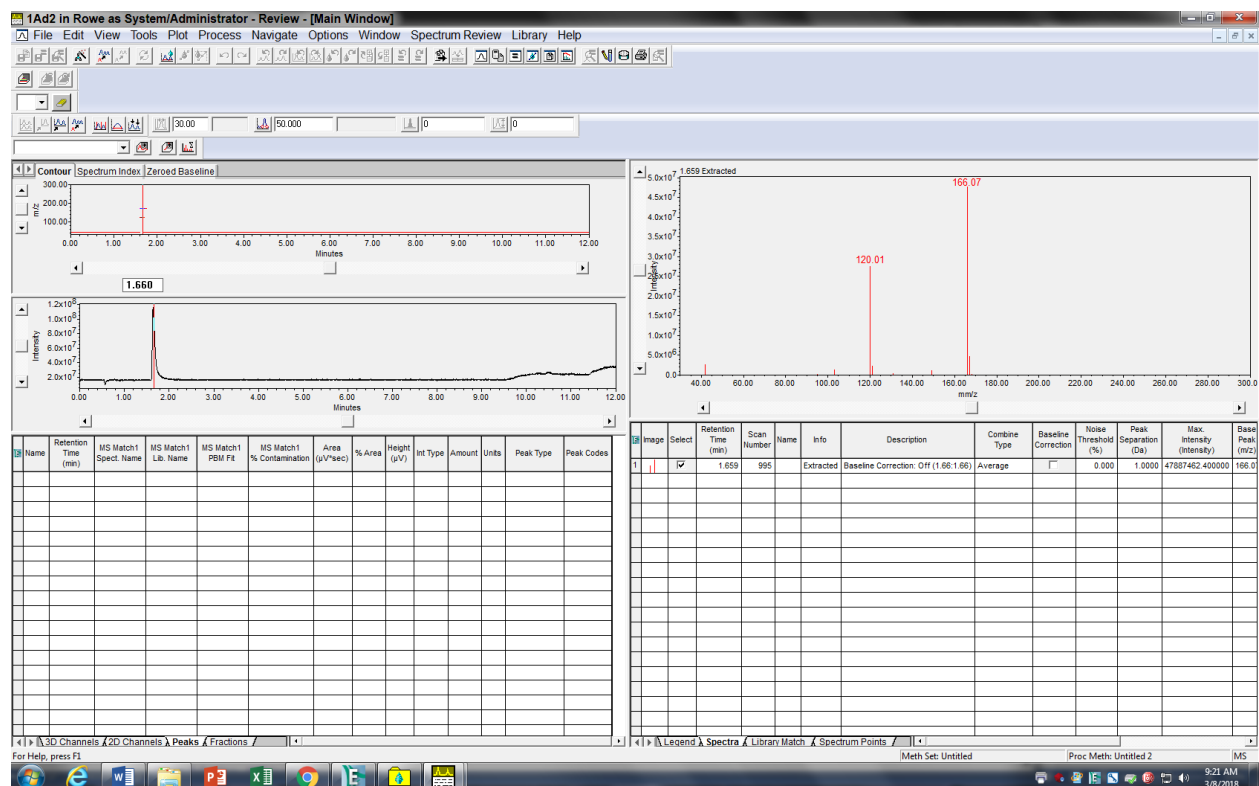


## 1 LC PDA Detector Data with Integrated Peak

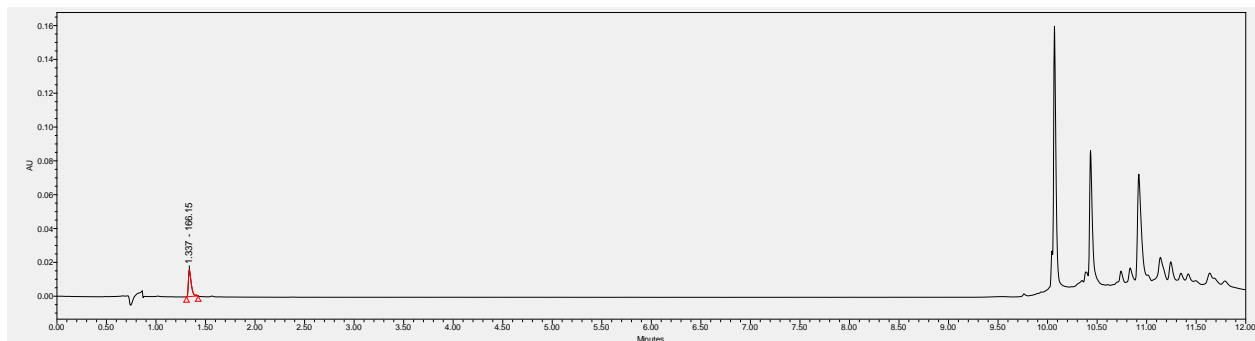


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.624	35449	100.00	13046	bb			Unknown

## 1 Mass Spectrum

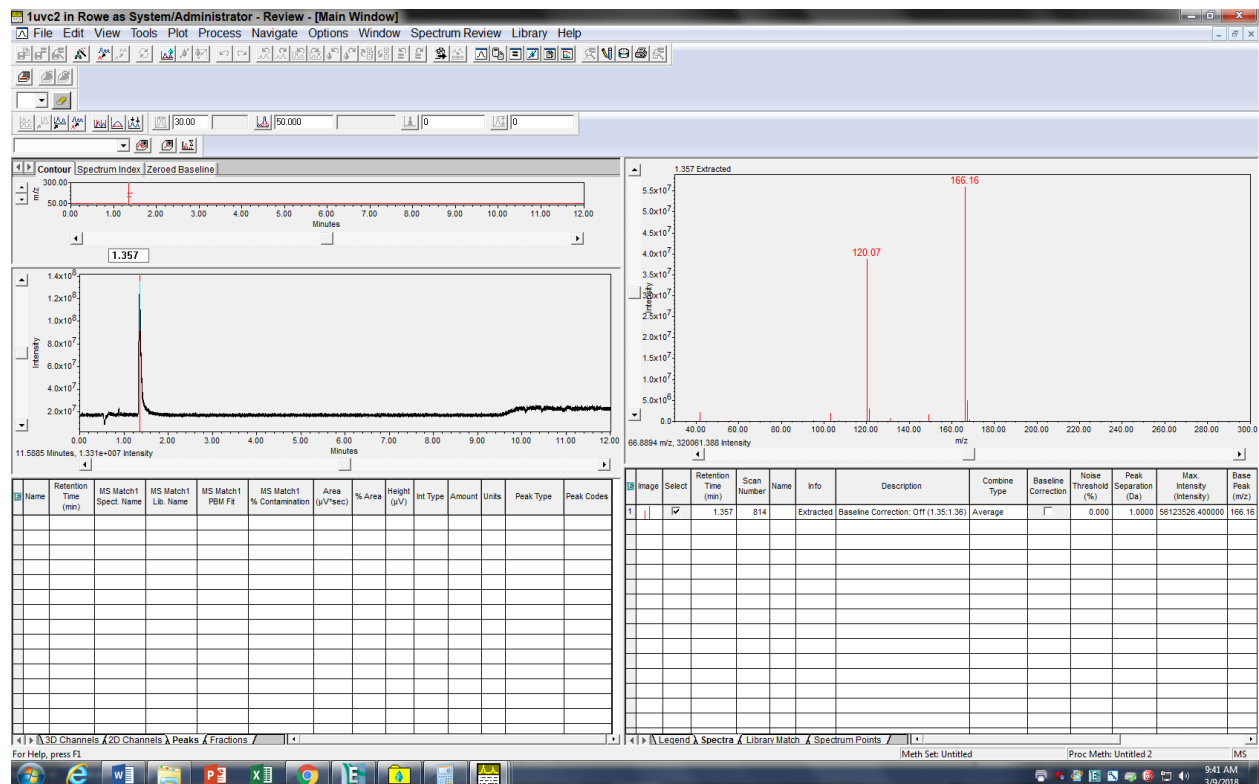


## 1 UV-C LC PDA Detector Data with Integrated Peak

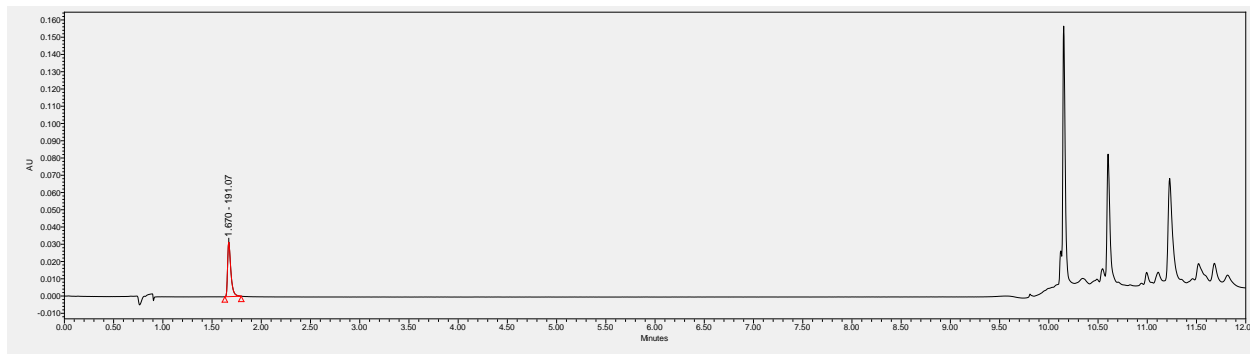


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.337	31687	100.00	15868	bb			Unknown

## 1 UVC Mass Spectrum

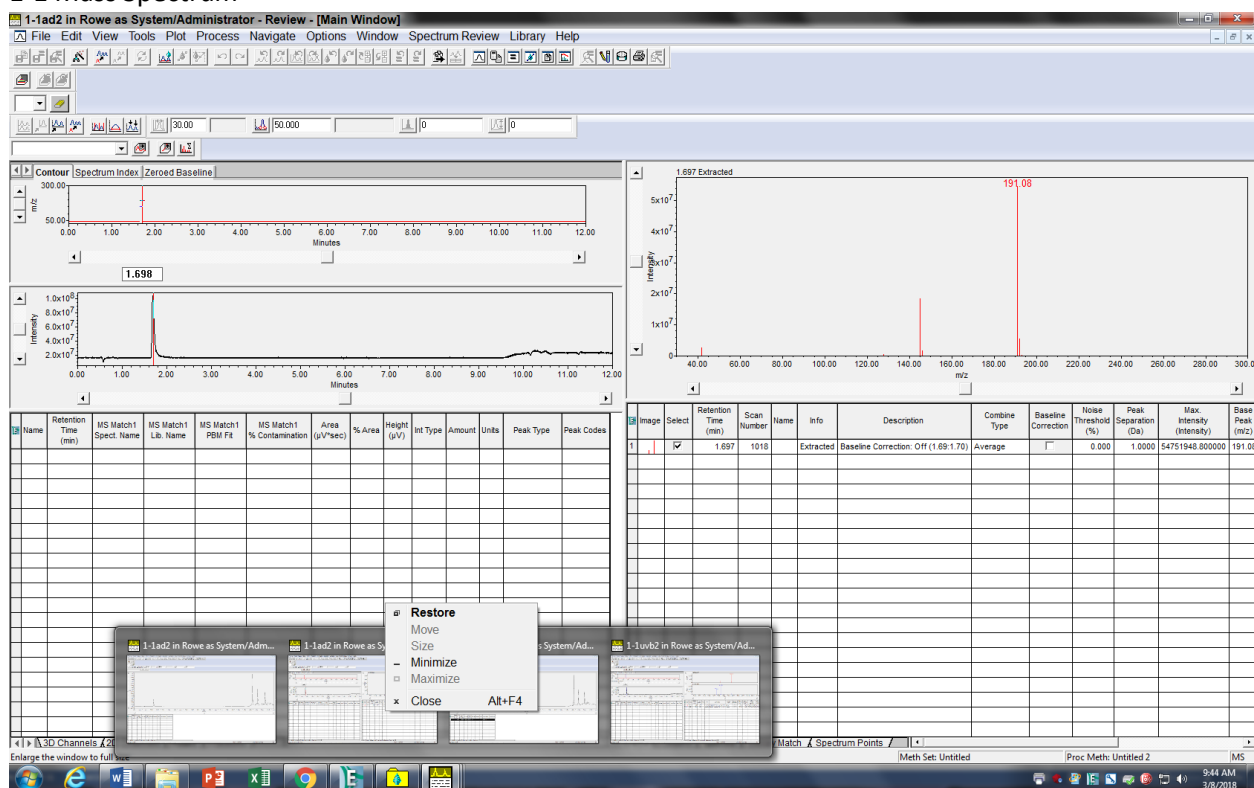


## 1-1 LC PDA Detector Data with Integrated Peak

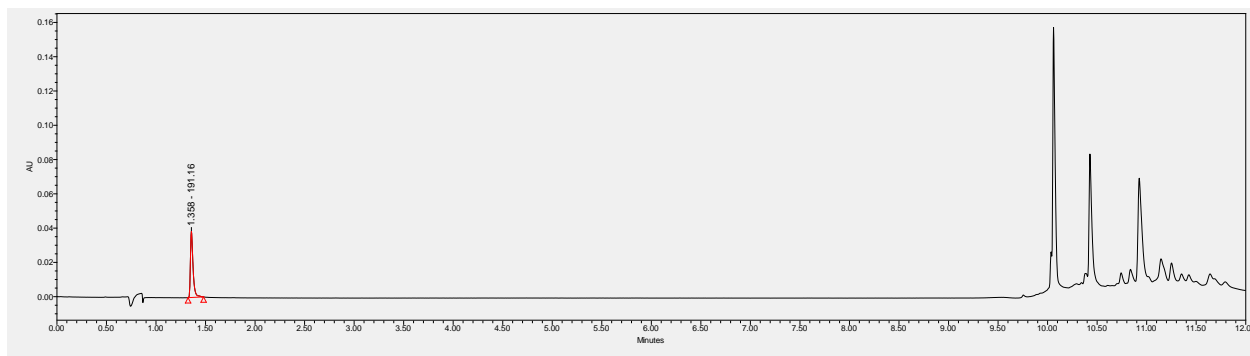


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.670	72903	100.00	31568	bb			Unknown

## 1-1 Mass Spectrum

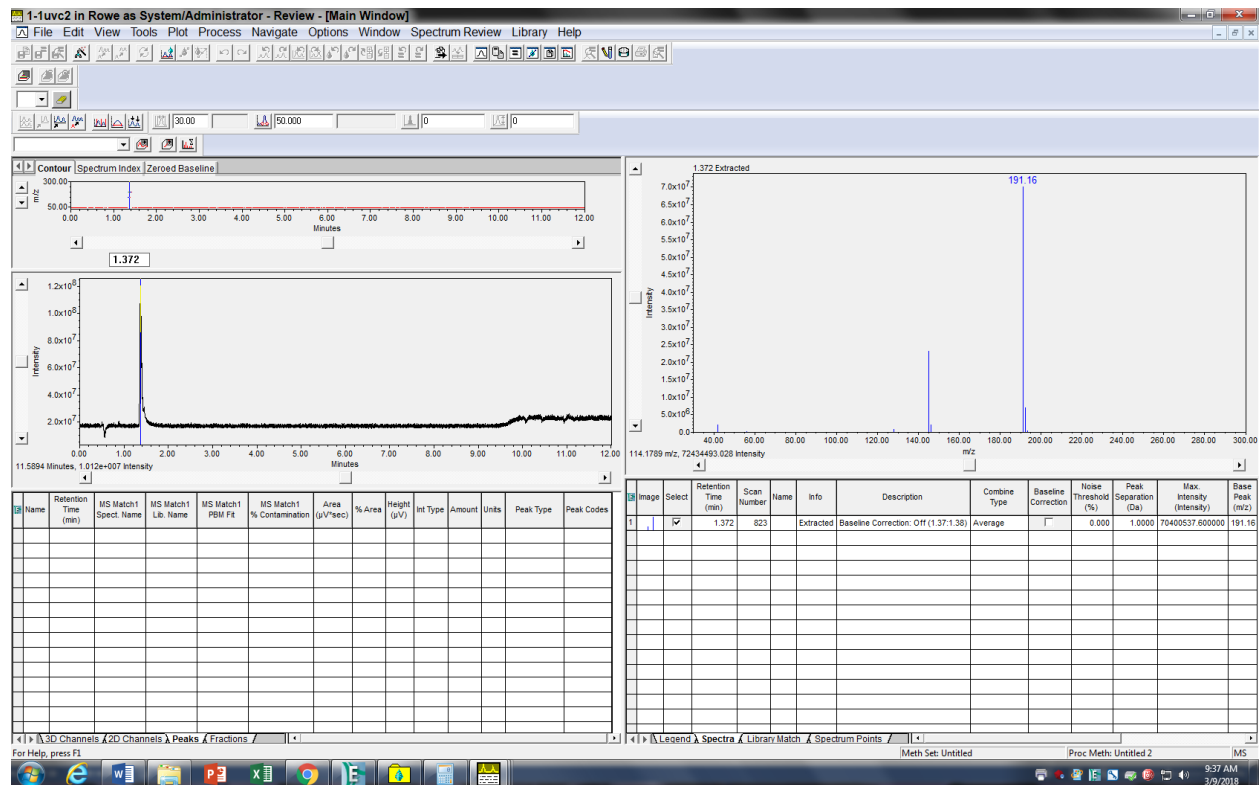


## 1-1UV-C LC PDA Detector Data with Integrated Peak

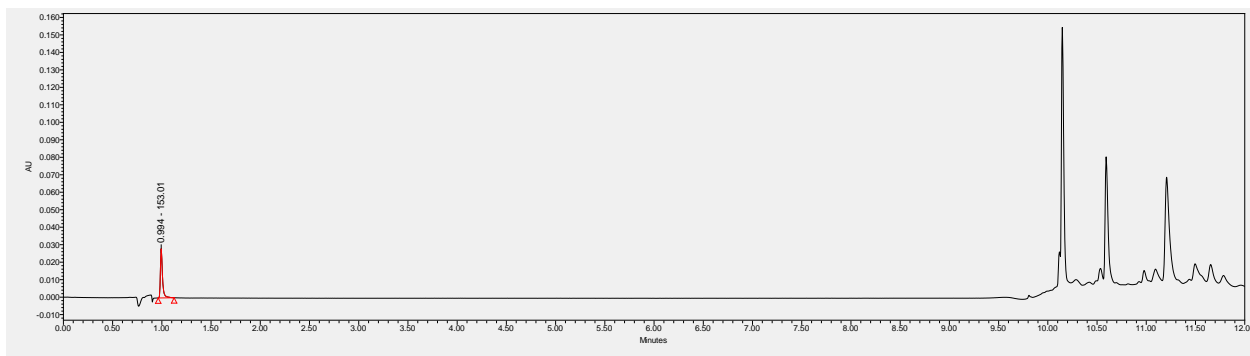


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.358	70672	100.00	38472	bb			Unknown

## 1-1UVC Mass Spectrum

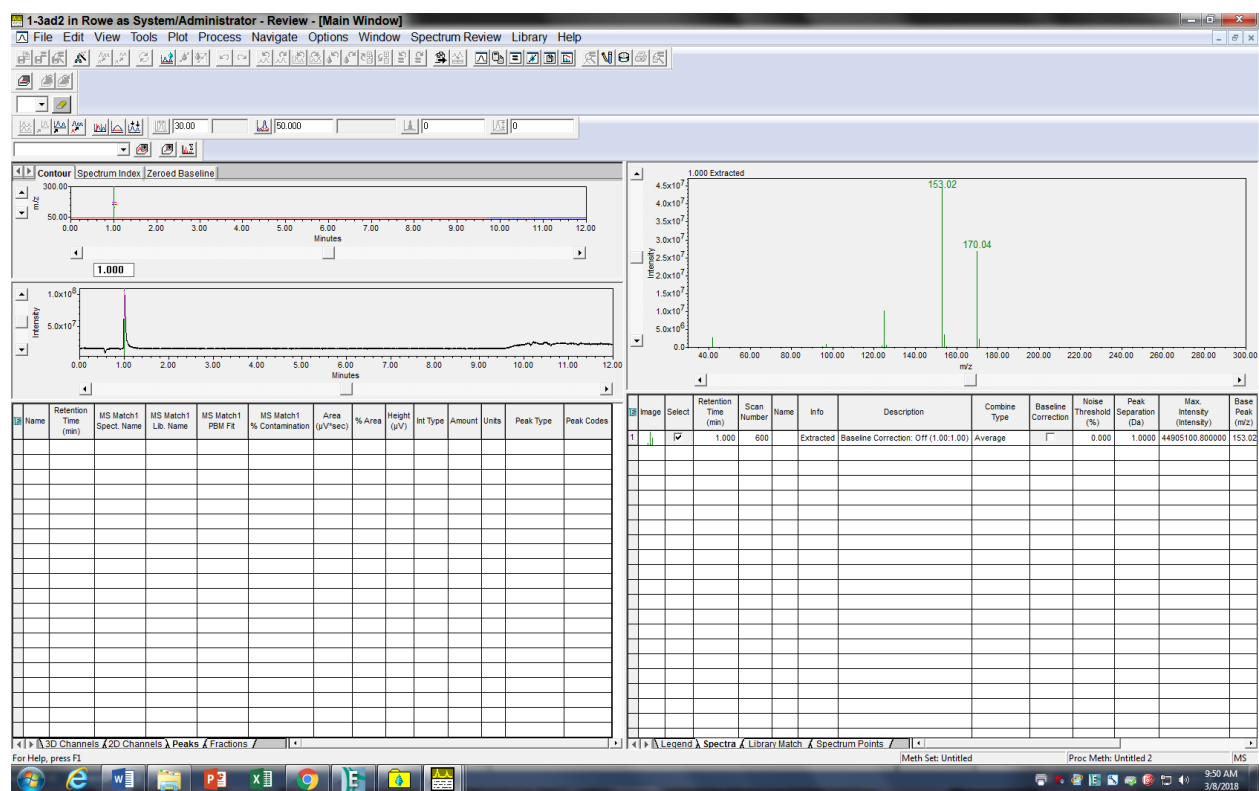


## 1-3 LC PDA Detector Data with Integrated Peak

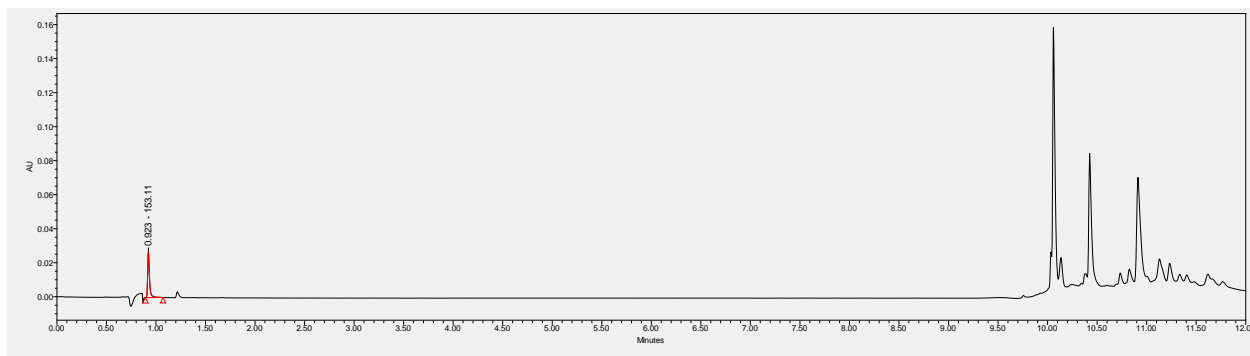


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.994	41530	100.00	28142	bb			Unknown

## 1-3 Mass Spectrum

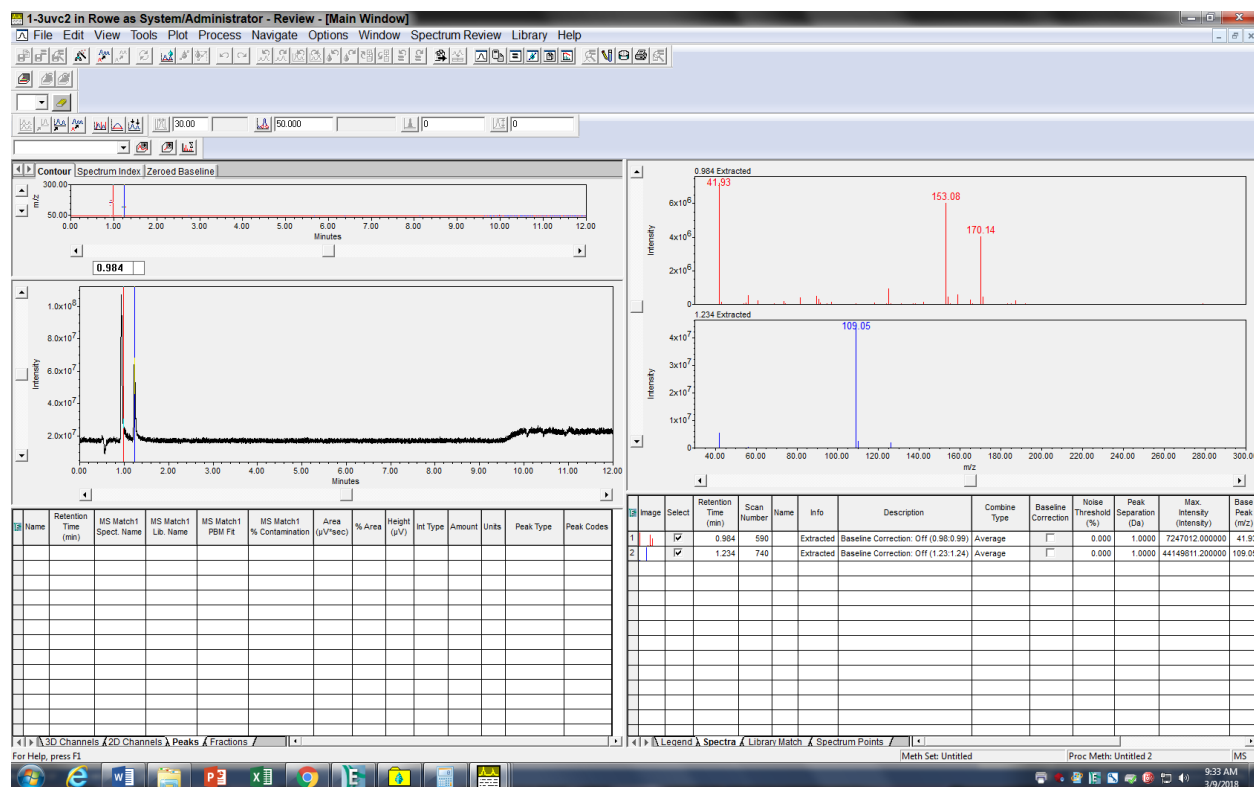


### 1-3 UV-C LC PDA Detector Data with Integrated Peak

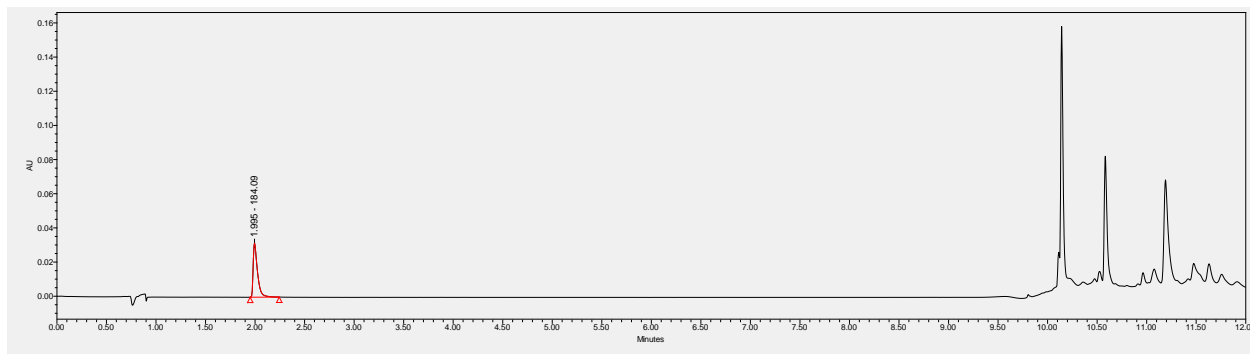


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.923	34905	100.00	26968	bb			Unknown

### 1-3 UVC Mass Spectrum

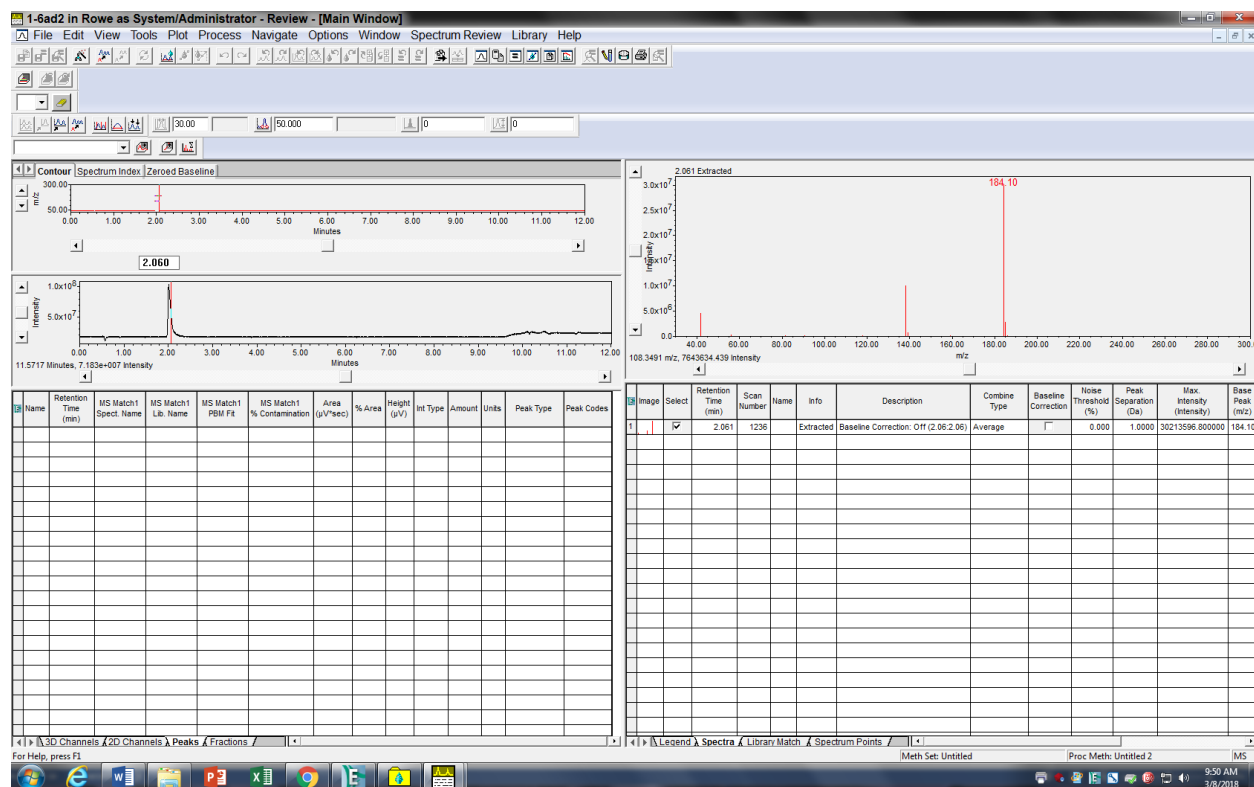


## 1-6 LC PDA Detector Data with Integrated Peak



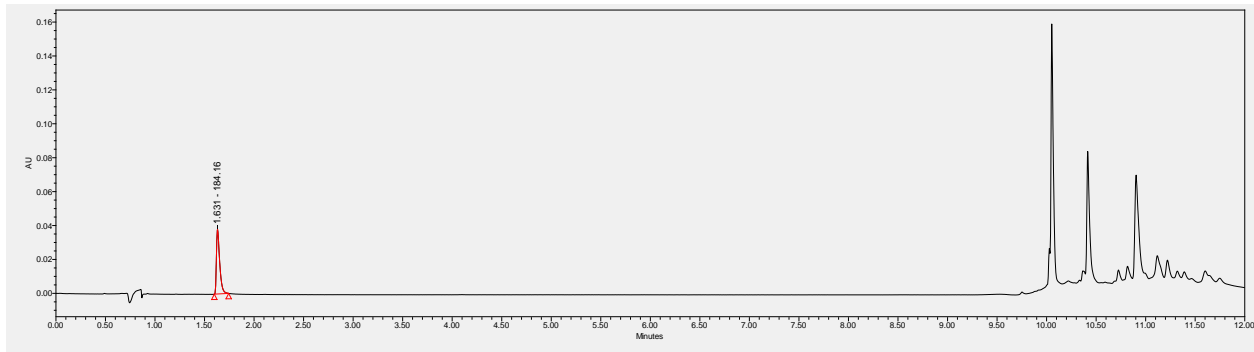
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.995	95629	100.00	31613	bb			Unknown

## 1-6 Mass Spectrum



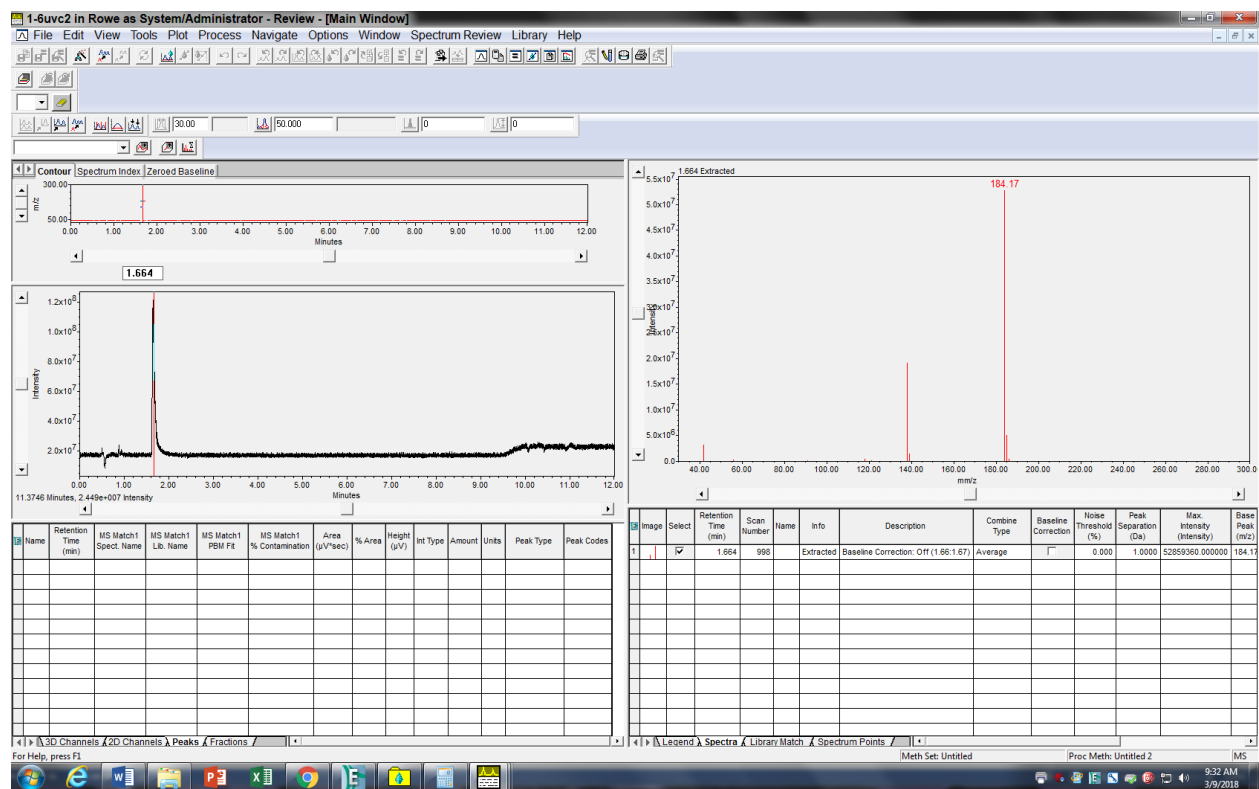


## 1-6 UV-C LC PDA Detector Data with Integrated Peak

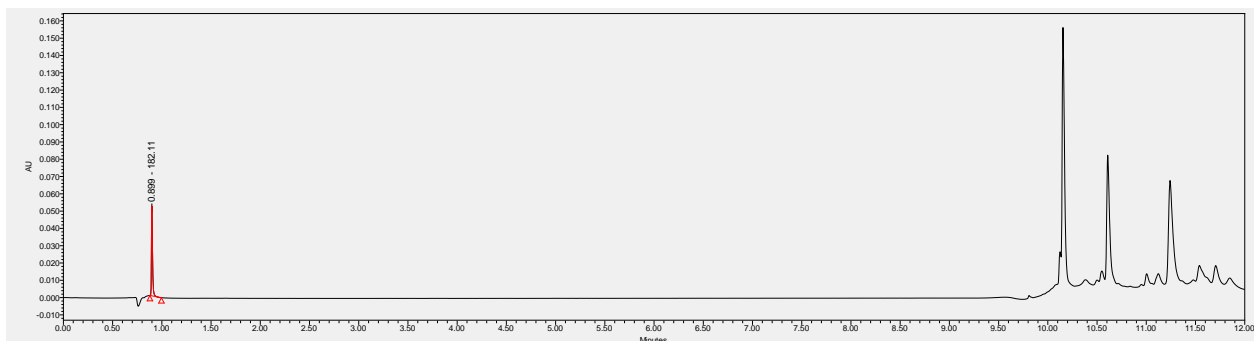


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.631	87632	100.00	38137	bb			Unknown

## 1-6 UVC Mass Spectrum

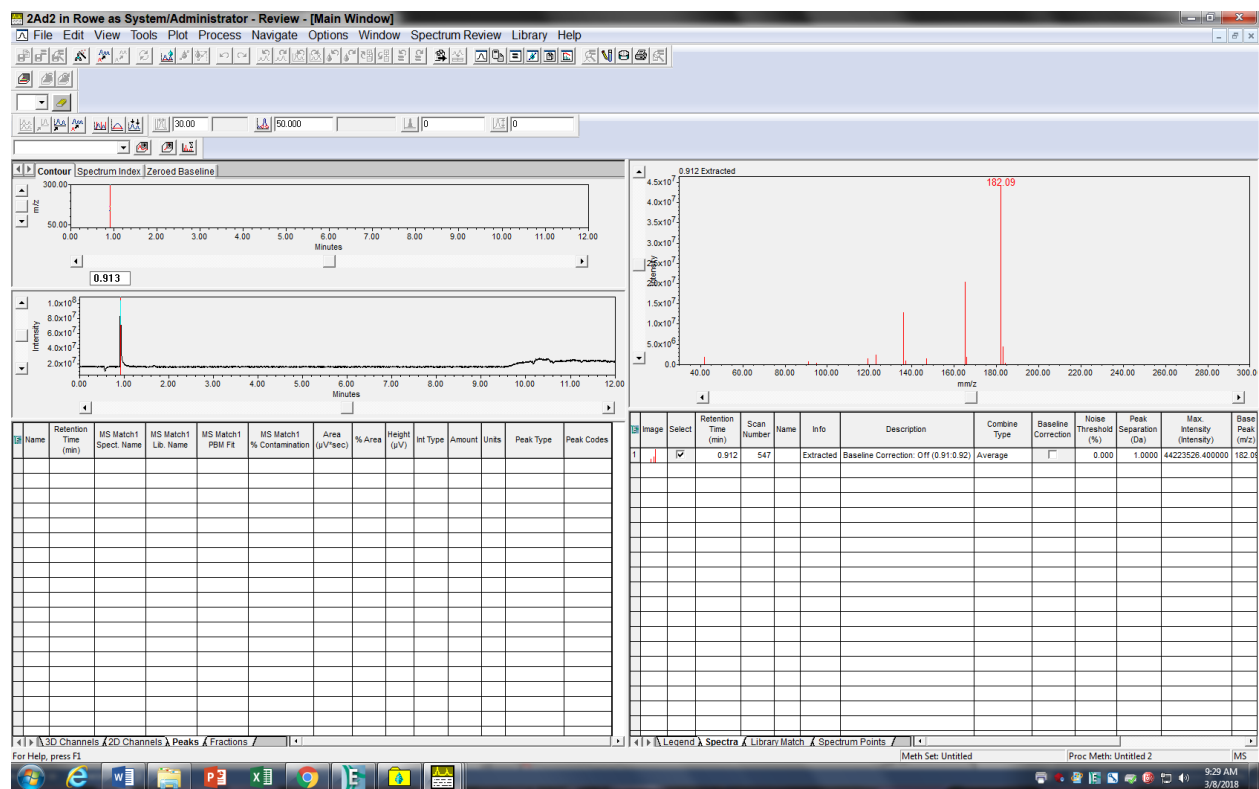


## 2 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.899	39379	100.00	52026	bb			Unknown

## 2 Mass Spectrum



The chromatogram displays the detector response (AU) over a 12-minute period. A significant peak is observed at 10.00 minutes, reaching an AU of approximately 0.16. A smaller peak is visible at 0.965 minutes, reaching an AU of approximately 0.08. The baseline is relatively flat with minor noise and small peaks around 0.8 and 3.0 minutes.

Retention Time (min)	AU
0.965	0.08
10.00	0.16

2uvv2 in Rowe as System/Administrator - Review - [Main Window]

File Edit View Tools Plot Process Navigate Options View Spectrum Review Library Help

30.00 50,000 0 0

Contour Spectrum Index Zeroed Baseline

Intensity vs. Minutes

0.501 0.758 382

Intensity vs. Minutes

Intensity vs. m/z

0.500 Extracted 41.90

0.759 Extracted 182.18

0.882 Extracted 182.19

#	Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Base Peak (m/z)
1		<input checked="" type="checkbox"/>	0.500	300	Extracted	Baseline Correction: Off (0.50:0.50)	Average	<input type="checkbox"/>	0.000	1.0000	8195300.800000	41.90	
2		<input checked="" type="checkbox"/>	0.759	455	Extracted	Baseline Correction: Off (0.76:0.76)	Average	<input type="checkbox"/>	0.000	1.0000	7232840.850000	41.9	
3		<input checked="" type="checkbox"/>	0.882	529	Extracted	Baseline Correction: Off (0.88:0.88)	Average	<input type="checkbox"/>	0.000	1.0000	35294022.400000	182.1	

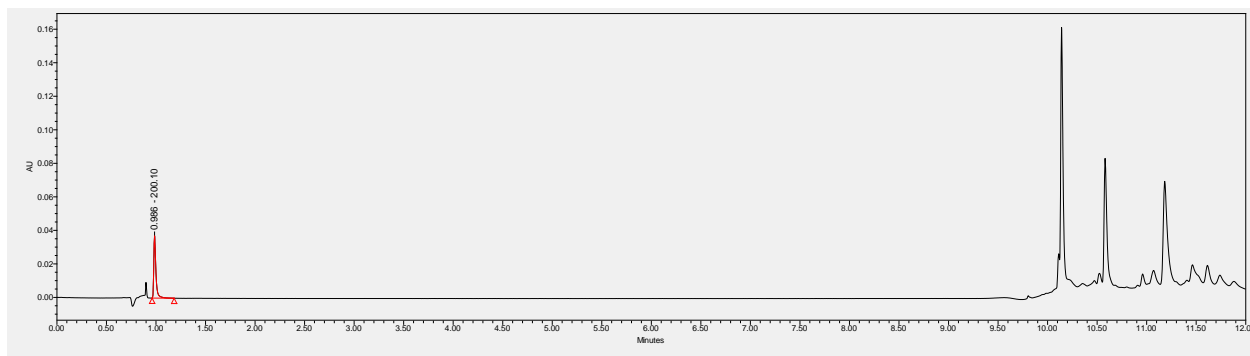
118.7805 m/z, 30859971.932 Intensity

Legend Spectra Library Match Spectrum Points

Meth Set: Untitled Proc Meth: Untitled 2

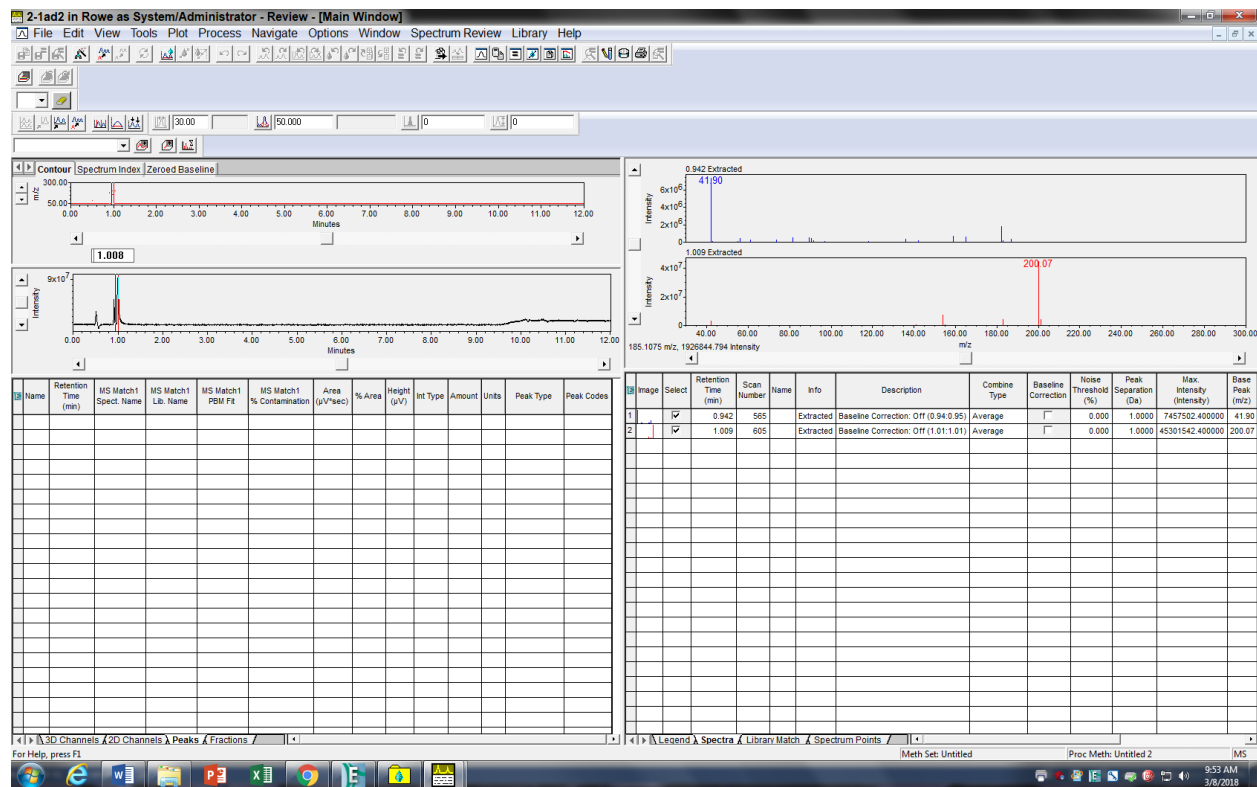
9:29 AM 3/9/2018

## 2-1 LC PDA Detector Data with Integrated Peak

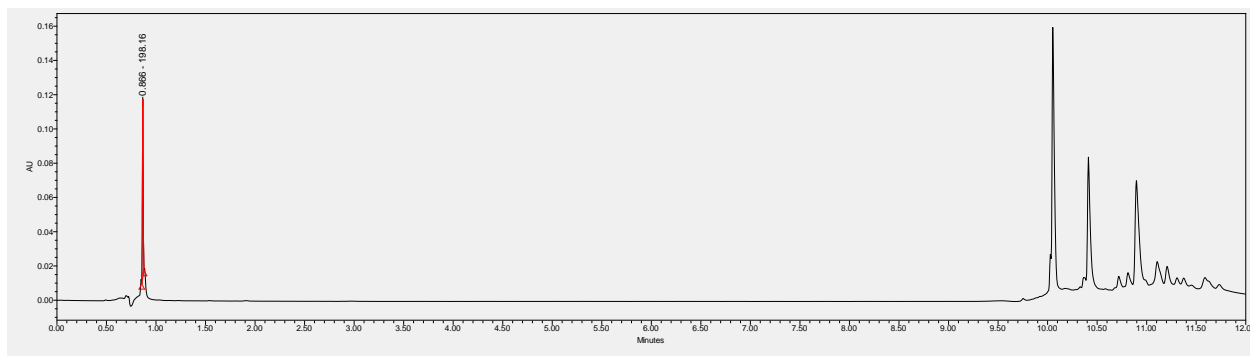


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.986	55681	100.00	37139	bb			Unknown

## 2-1 Mass Spectrum

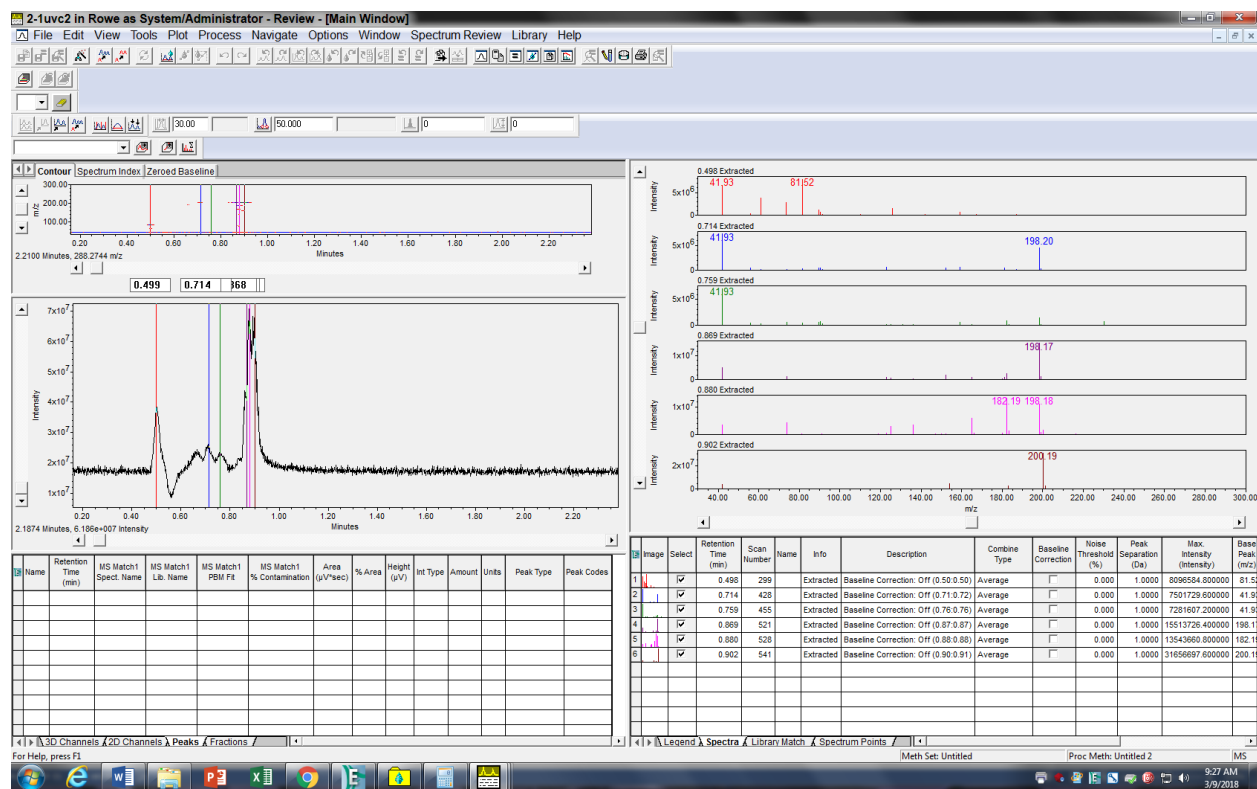


## 2-1 UV-C LC PDA Detector Data with Integrated Peak

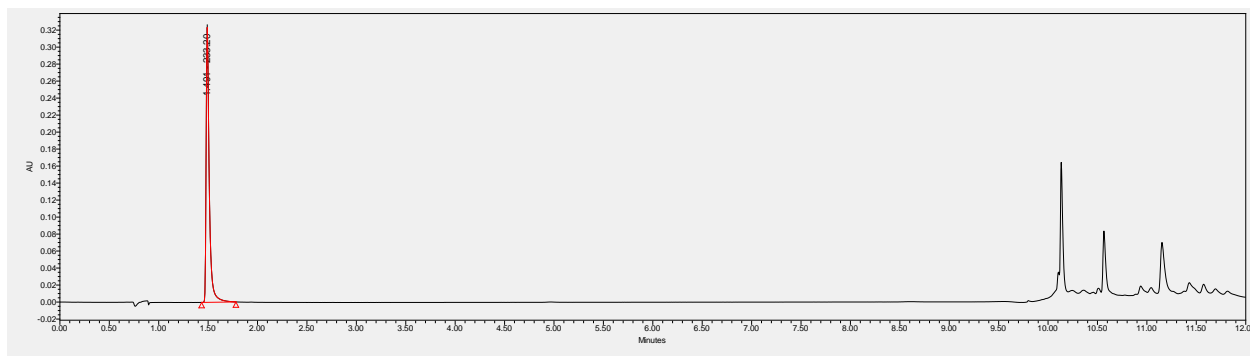


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.866	59089	100.00	105079	bb			Unknown

## 2-1 UVC Mass Spectrum

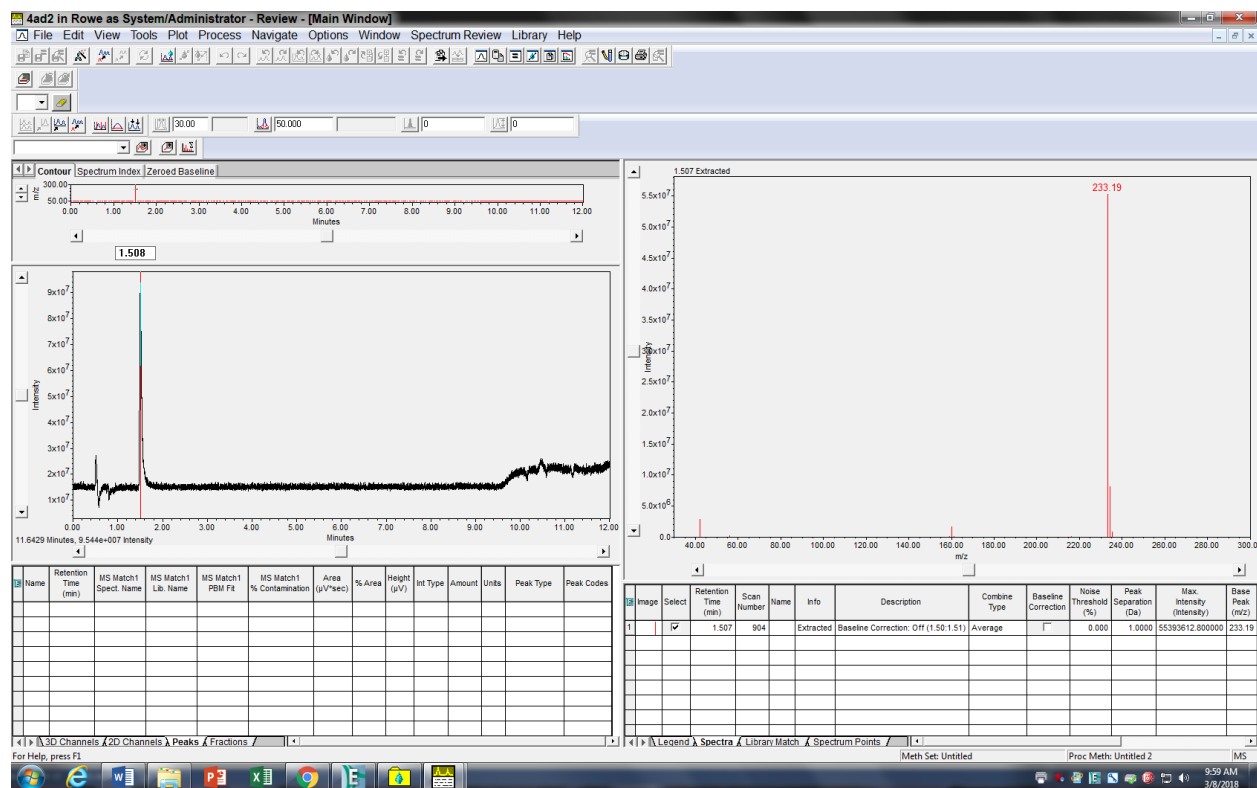


#### 4 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.491	716988	100.00	323502	bb			Unknown

#### 4 Mass Spectrum



The chromatogram displays detector response in AU (Absorbance Units) on the y-axis, ranging from -0.02 to 0.36, against time in minutes on the x-axis, ranging from 0.00 to 12.00. A prominent peak is observed at 1.250 minutes, reaching an AU of approximately 0.35. This peak is highlighted with a red vertical line and a red 'x' at its base. Several smaller peaks are visible in the later time range: a peak at approximately 10.0 minutes (AU ~0.15), a peak at approximately 10.5 minutes (AU ~0.08), and a peak at approximately 11.0 minutes (AU ~0.07). The baseline is relatively flat with minor noise throughout the run.

**Chromatogram Data:**

Peak	Retention Time (min)	Scan Number	Intensity
1	0.495	297	~1.0x10 <sup>8</sup>
2	1.177	705	~1.0x10 <sup>8</sup>
3	1.13	787	~1.0x10 <sup>8</sup>

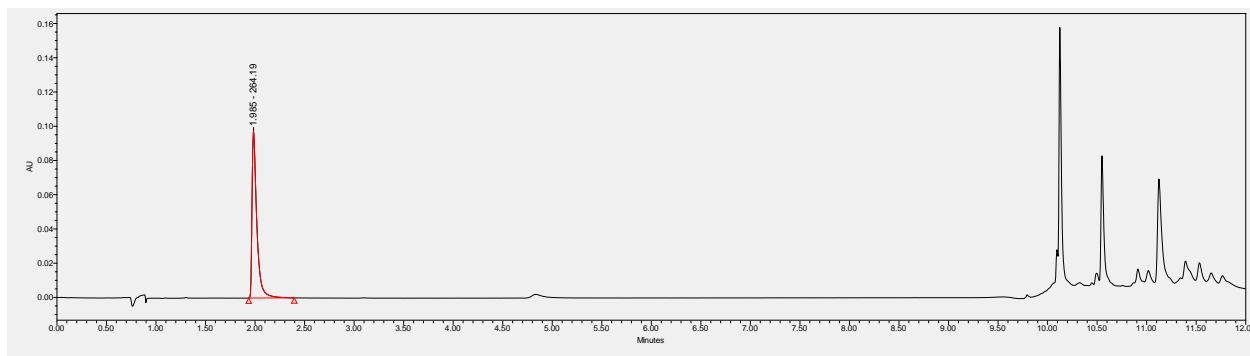
**Extracted Spectra Data:**

Peak	Retention Time (min)	Scan Number	m/z	Intensity
0.495	0.495	297	41.94	~6x10 <sup>6</sup>
0.495	0.495	297	81.52	~6x10 <sup>6</sup>
1.177	1.177	705	233.21	~6x10 <sup>6</sup>
1.13	1.13	787	41.94	~6x10 <sup>6</sup>
1.13	1.13	787	209.19	~6x10 <sup>6</sup>

**Table Data:**

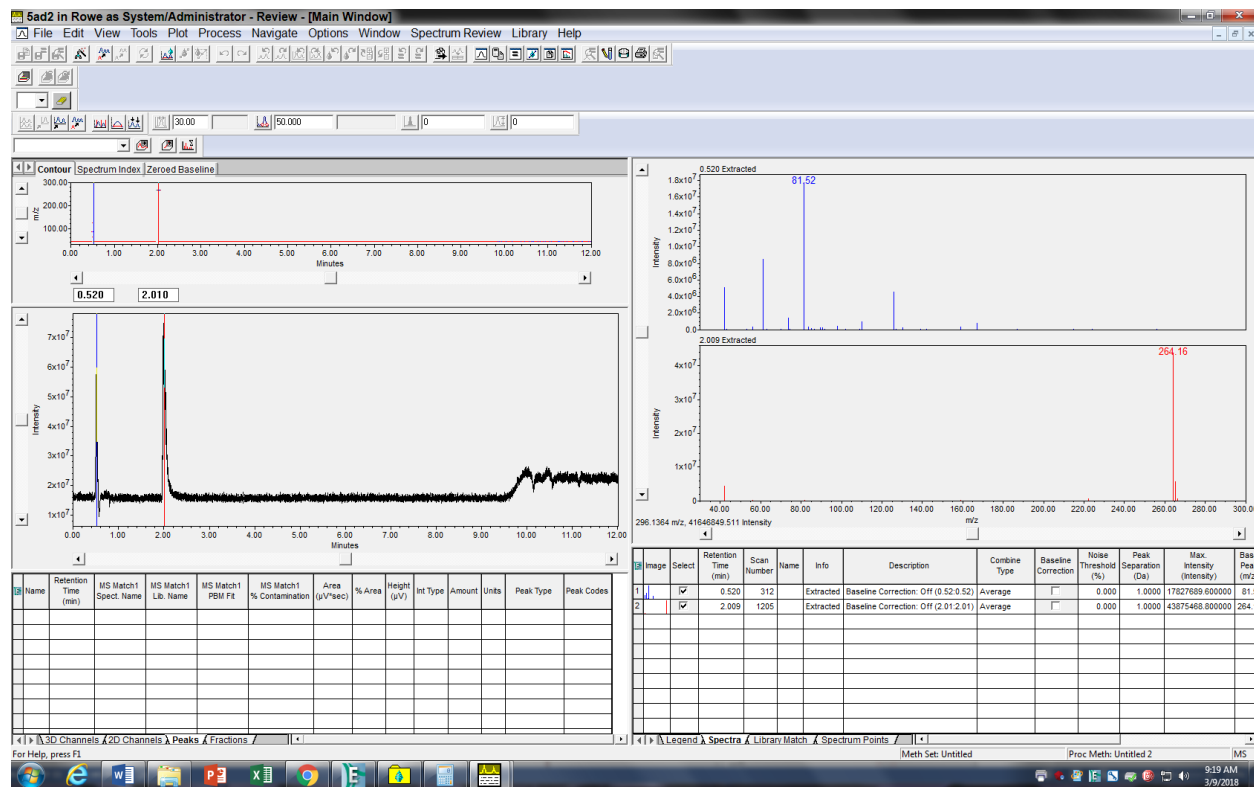
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Base Peak (m/z)
1	<input checked="" type="checkbox"/>	0.495	297		Extracted	Baseline Correction: Off (0.49:0.50)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	6887625.600000	41.9
2	<input checked="" type="checkbox"/>	1.176	705		Extracted	Baseline Correction: Off (1.17:1.18)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	59881968.800000	233.2
3	<input checked="" type="checkbox"/>	1.313	787		Extracted	Baseline Correction: Off (1.31:1.32)	Average	<input checked="" type="checkbox"/>	0.000	1.0000	7004770.400000	41.9

## 5 LC PDA Detector Data with Integrated Peak



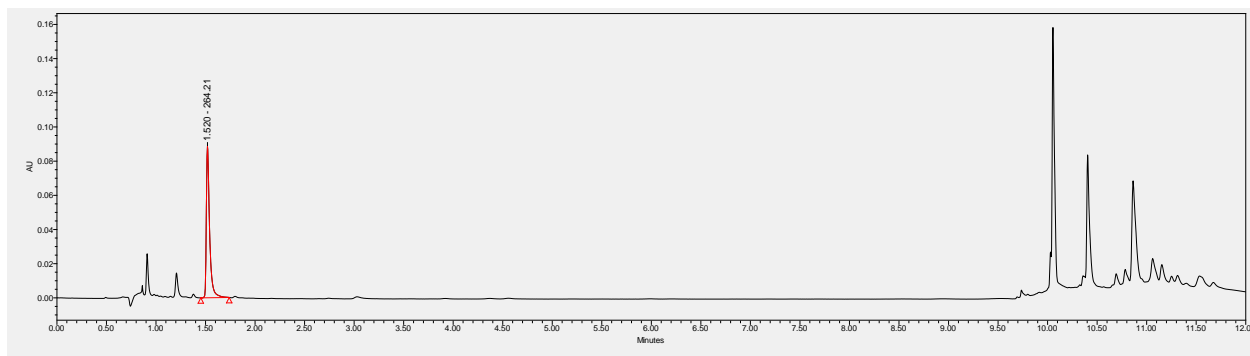
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.985	327286	100.00	97203	bb			Unknown

## 5 Mass Spectrum



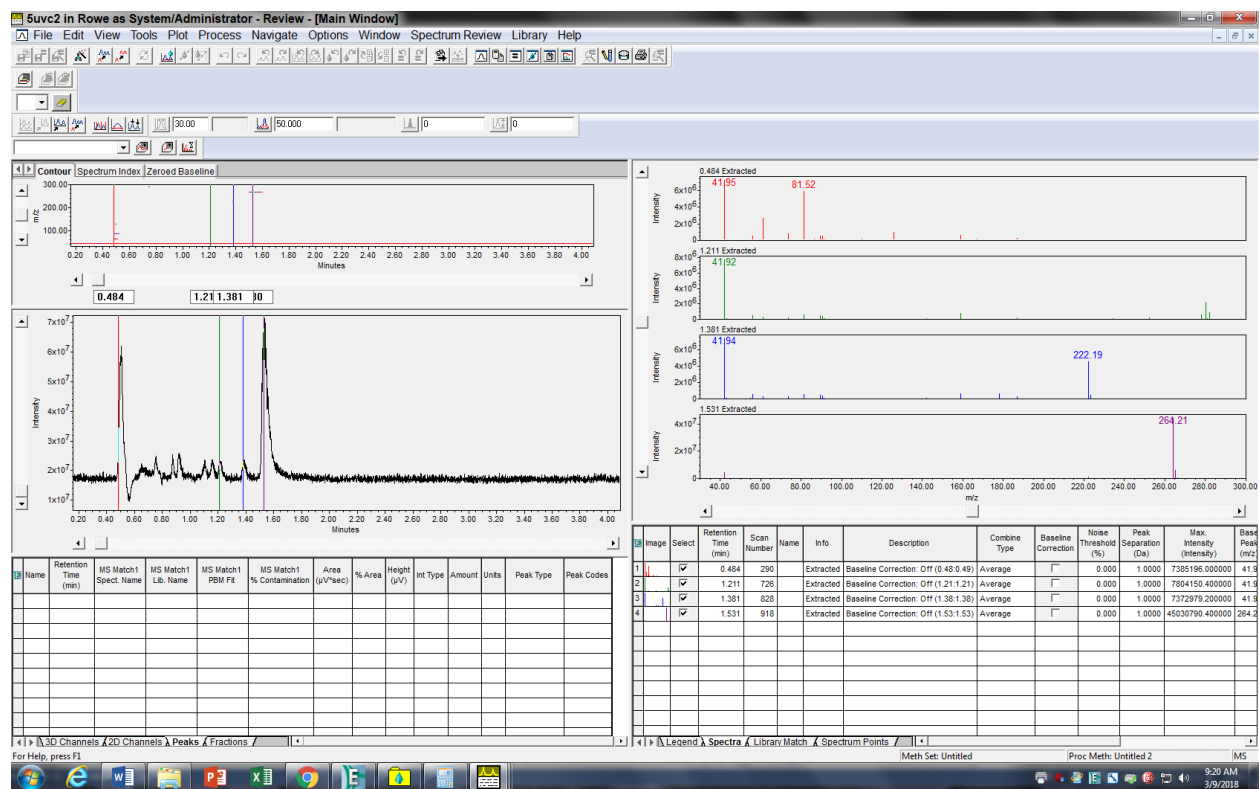


## 5 UV-C LC PDA Detector Data with Integrated Peak

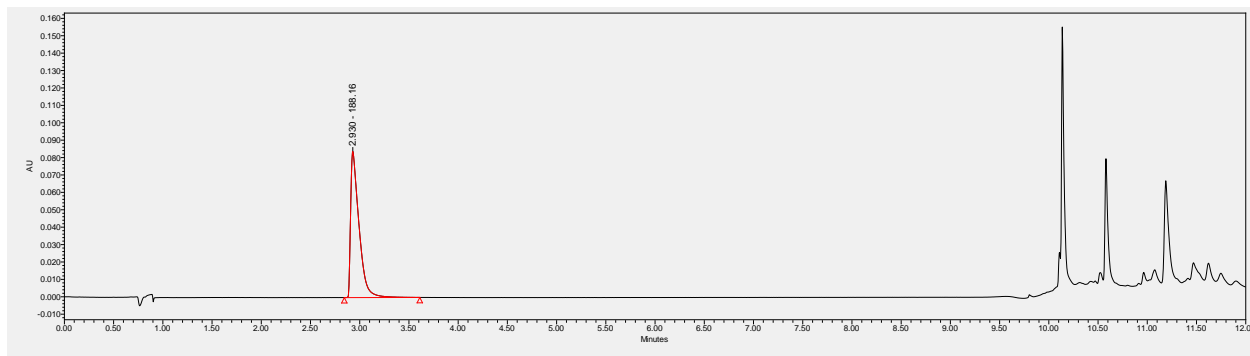


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.520	203226	100.00	88505	bb			Unknown

## 5 UVC Mass Spectrum

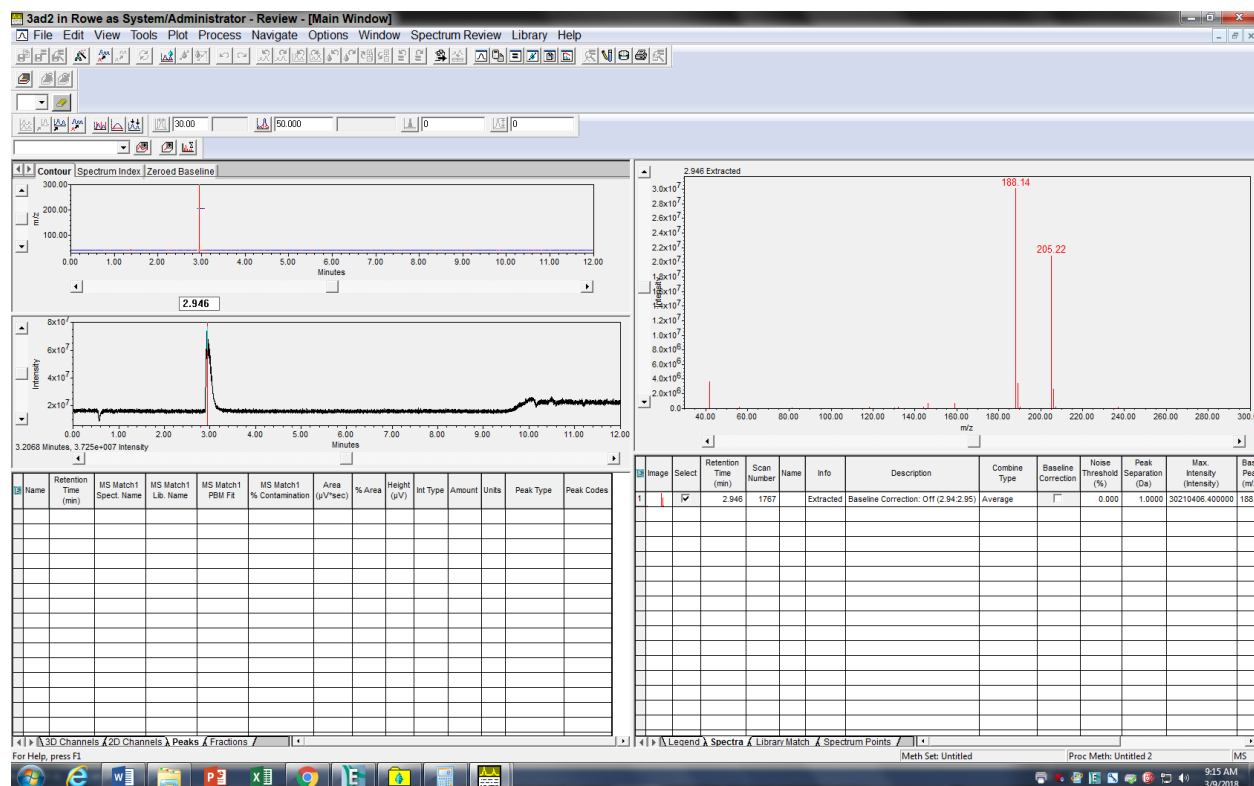


### 3 LC PDA Detector Data with Integrated Peak

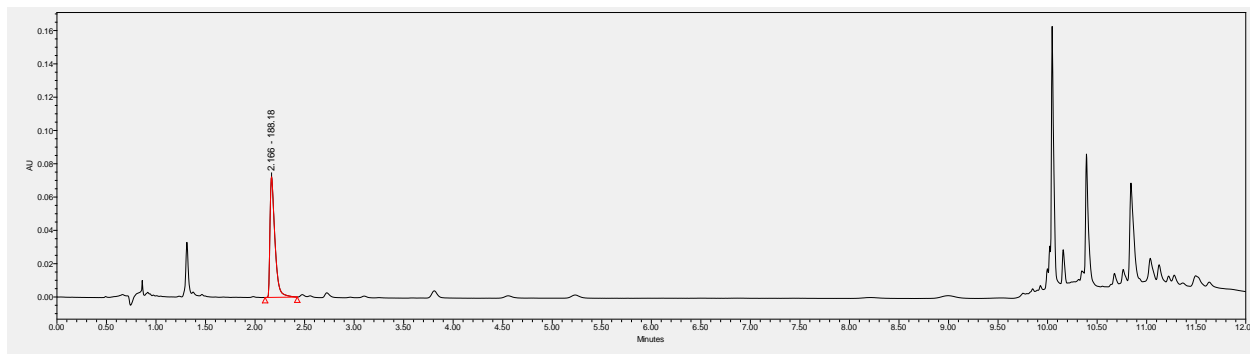


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.930	495936	100.00	83955	bb			Unknown

### 3 Mass Spectrum

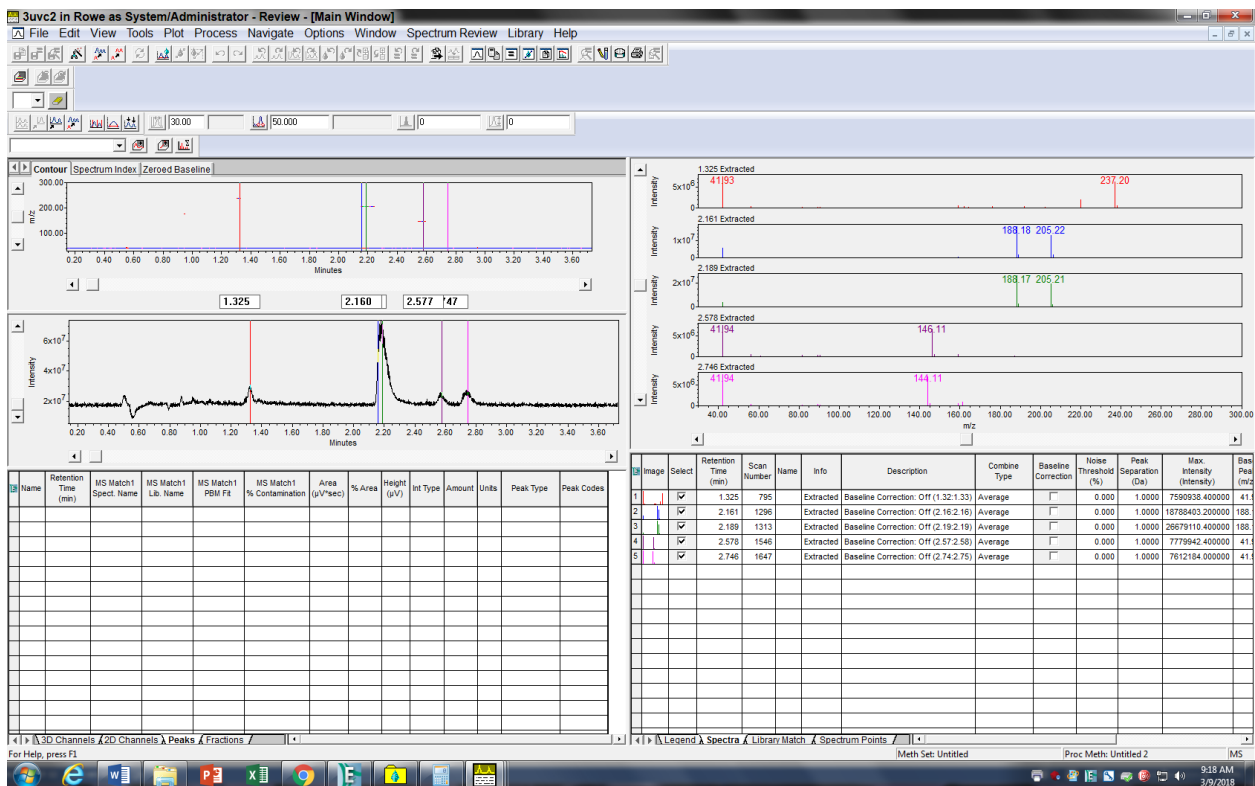


### 3 UV-C LC PDA Detector Data with Integrated Peak

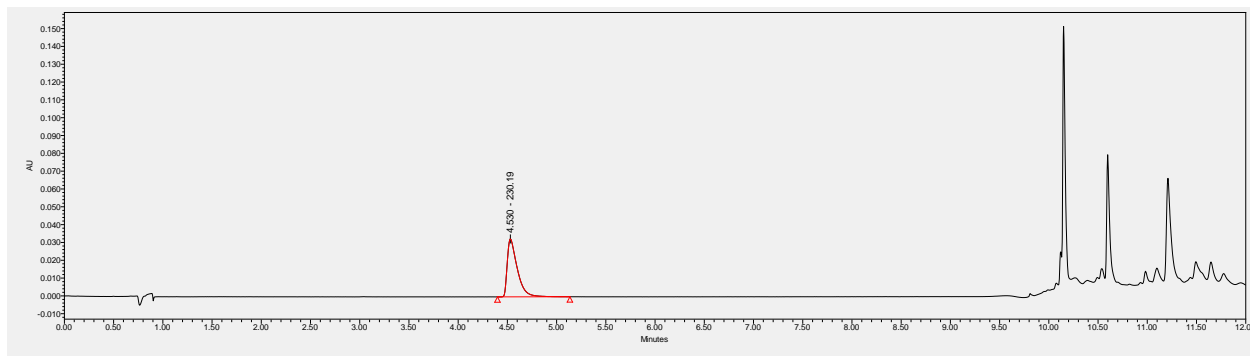


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.930	495936	100.00	83955	bb			Unknown

### 3 UVC Mass Spectrum

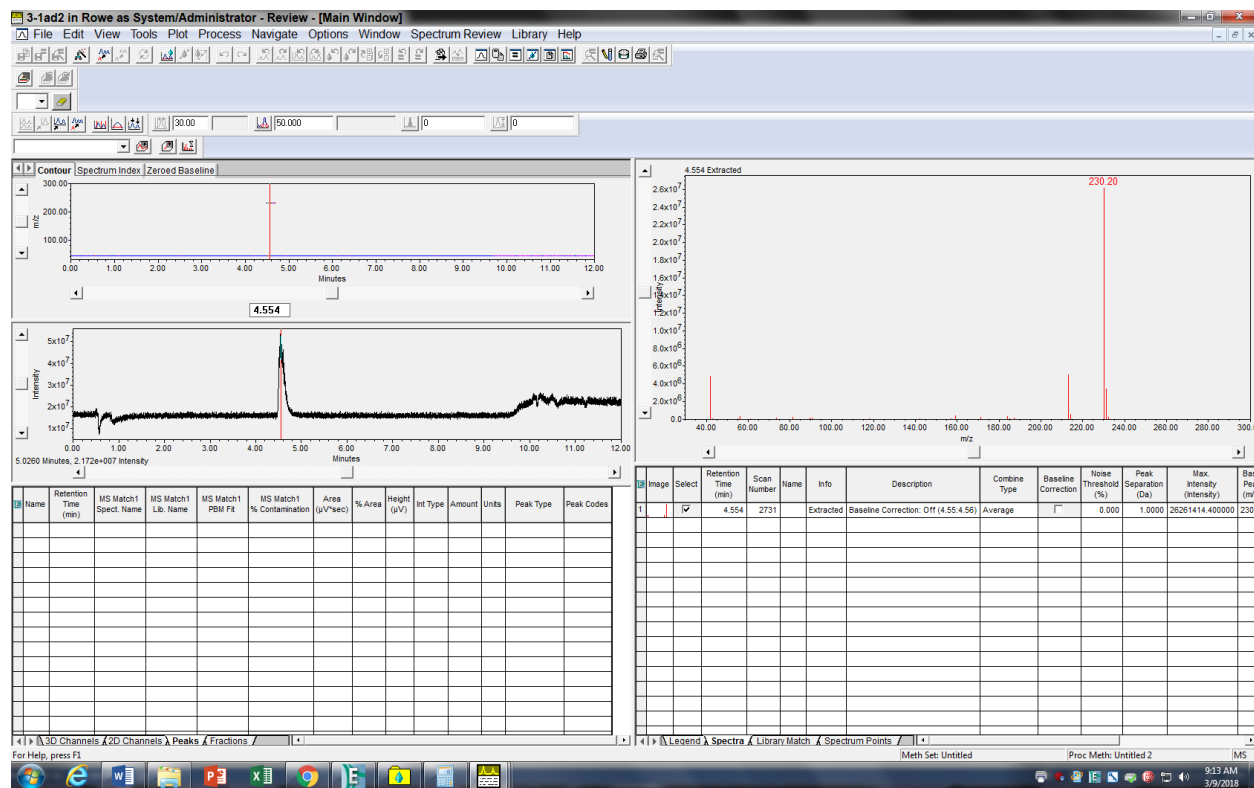


### 3-1 LC PDA Detector Data with Integrated Peak

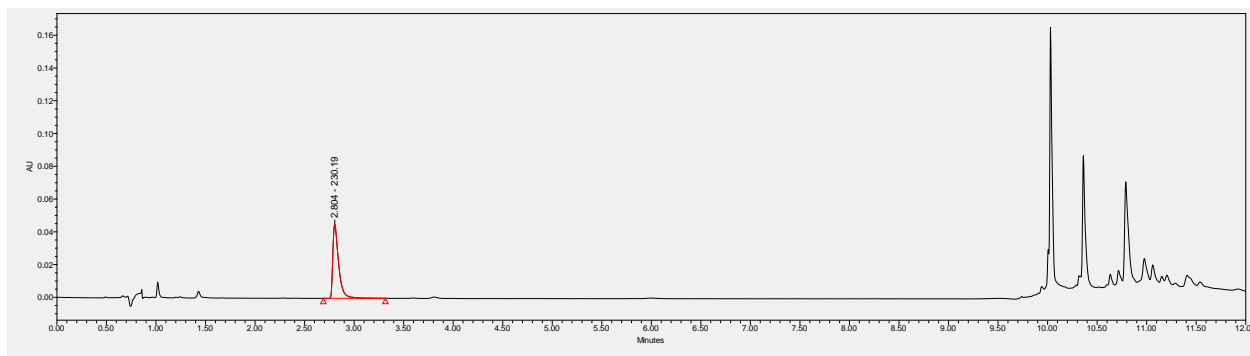


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.530	228635	100.00	32514	bb			Unknown

### 3-1 Mass Spectrum

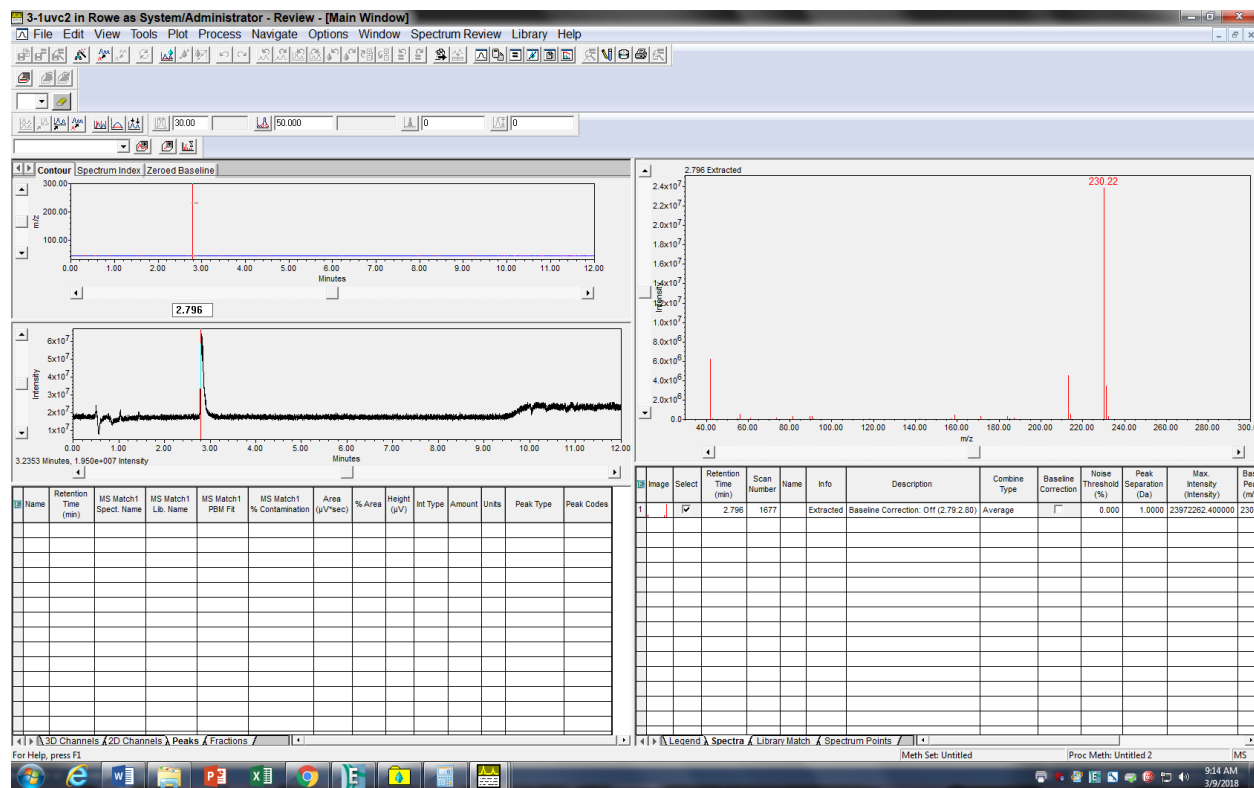


### 3-1 UV-C LC PDA Detector Data with Integrated Peak



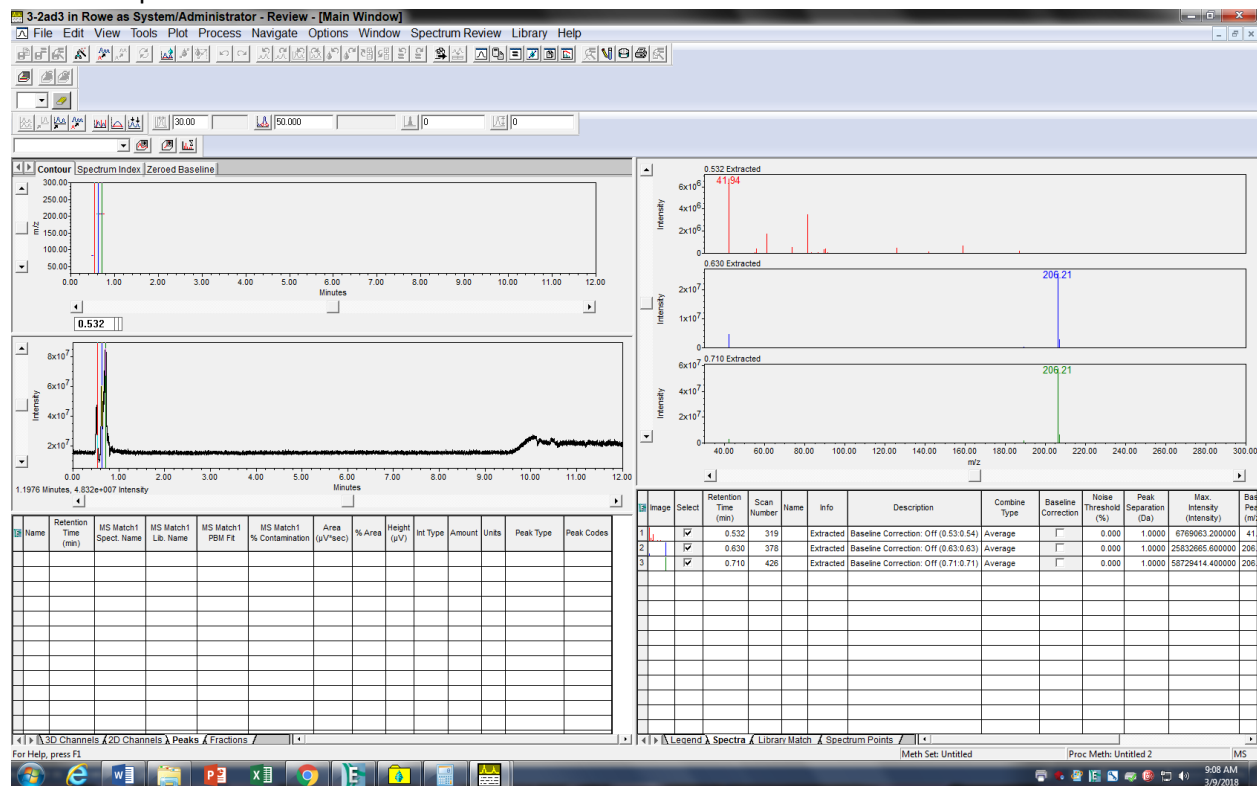
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.804	184039	100.00	45311	bb			Unknown

### 3-1 UVC Mass Spectrum

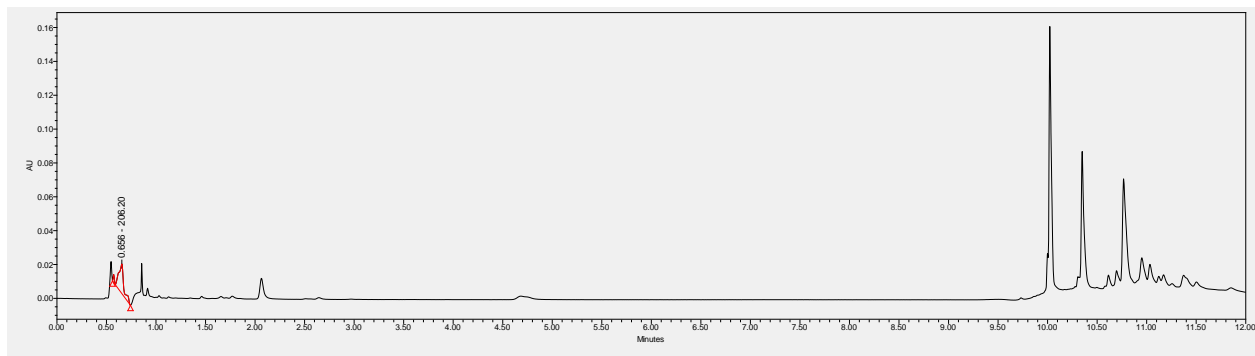


### 3-2 LC PDA Detector Data with Integrated Peak

### 3-2 Mass Spectrum

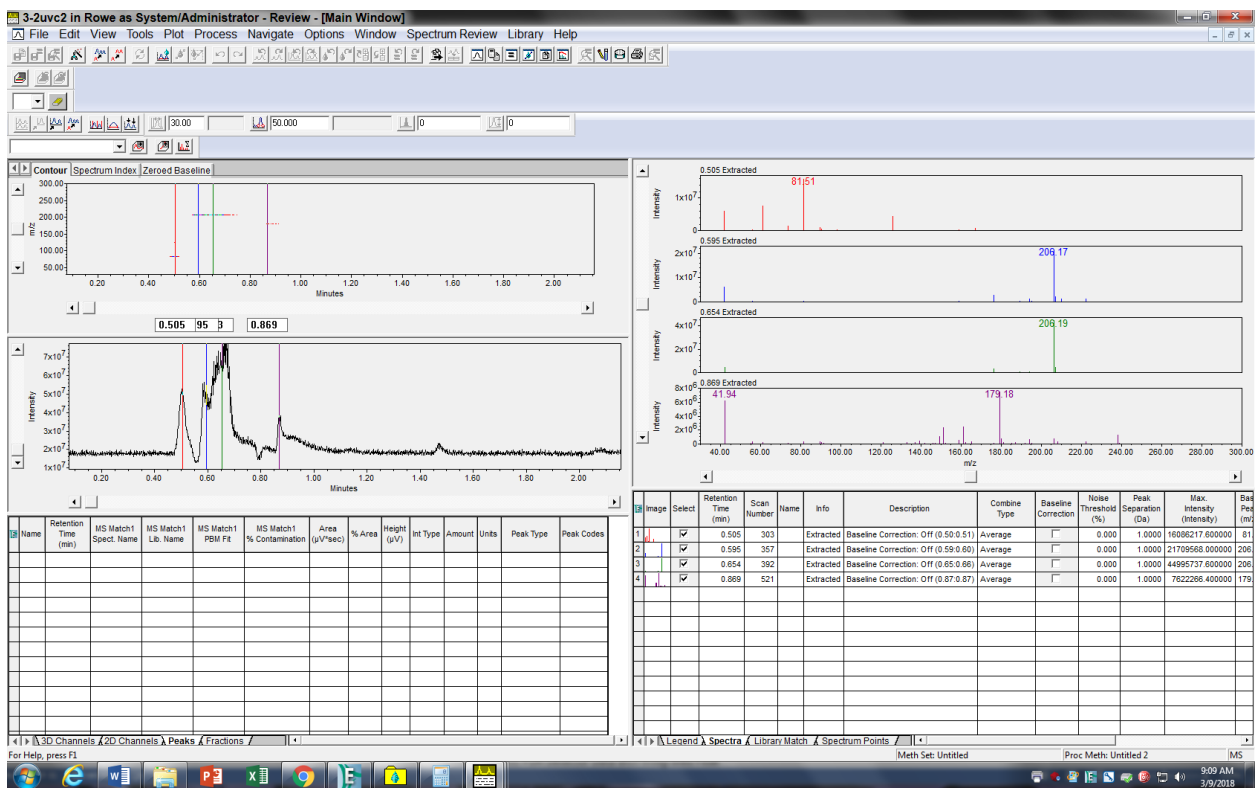


### 3-2 UV-C LC PDA Detector Data with Integrated Peak

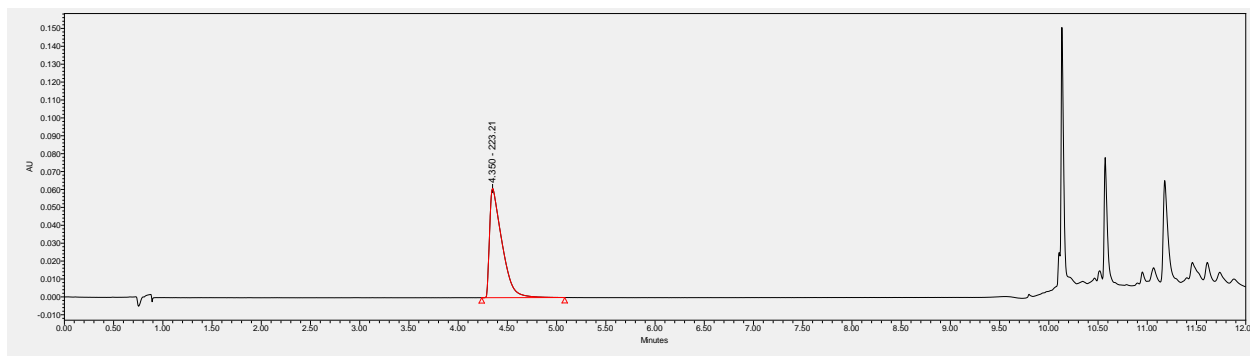


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.656	62342	100.00	17492	bb			Unknown

### 3-2 UVC Mass Spectrum

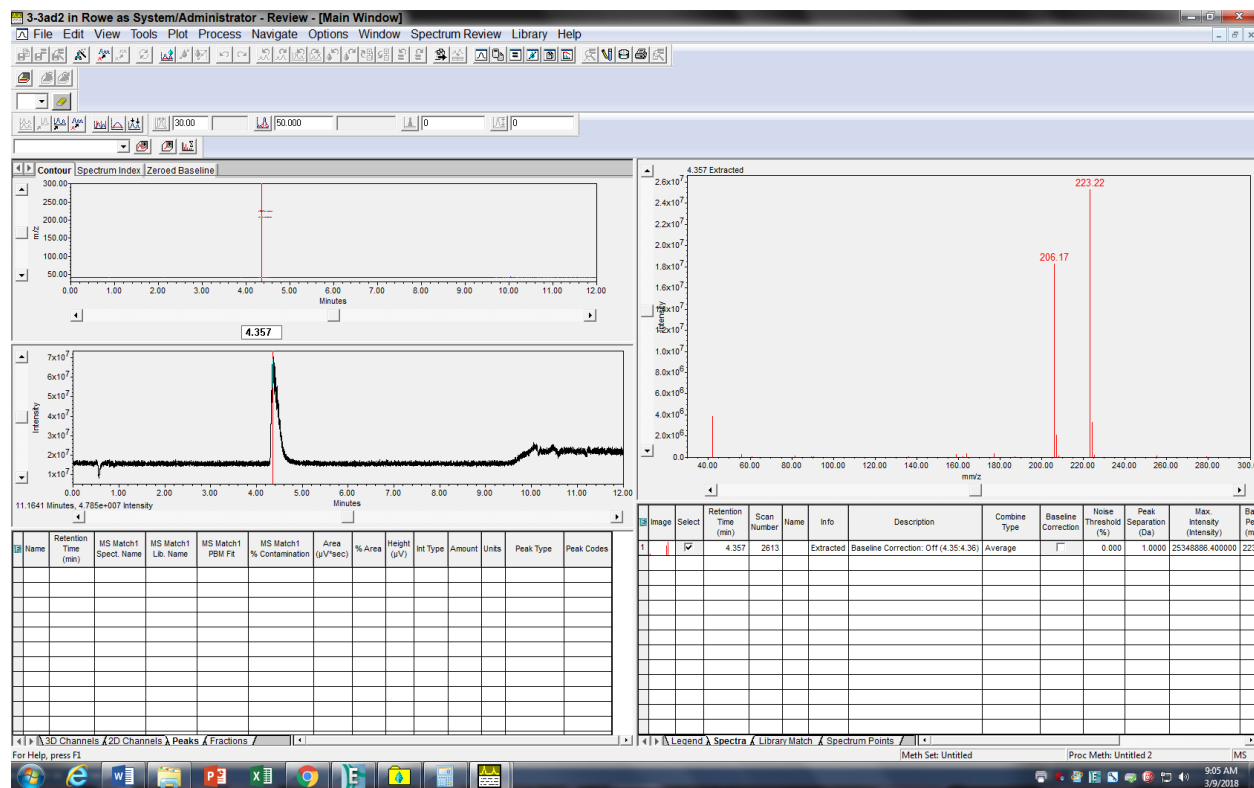


### 3-3 LC PDA Detector Data with Integrated Peak



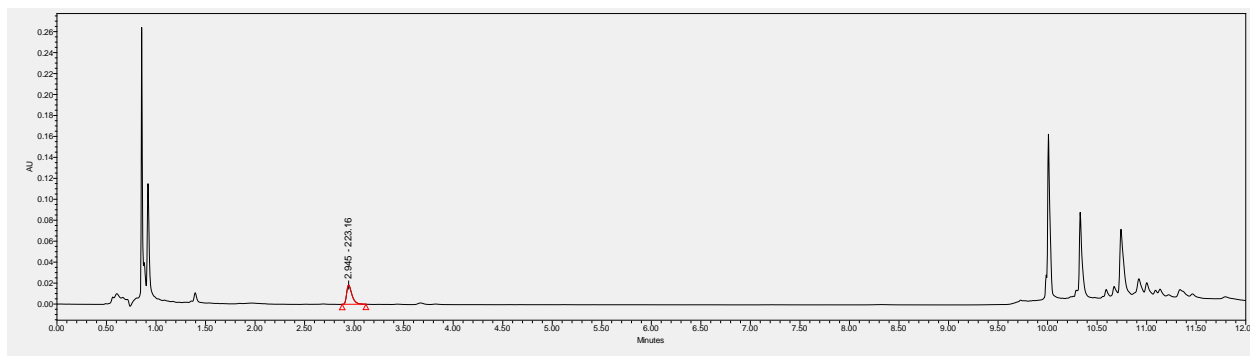
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.350	533792	100.00	61024	bb			Unknown

### 3-3 Mass Spectrum



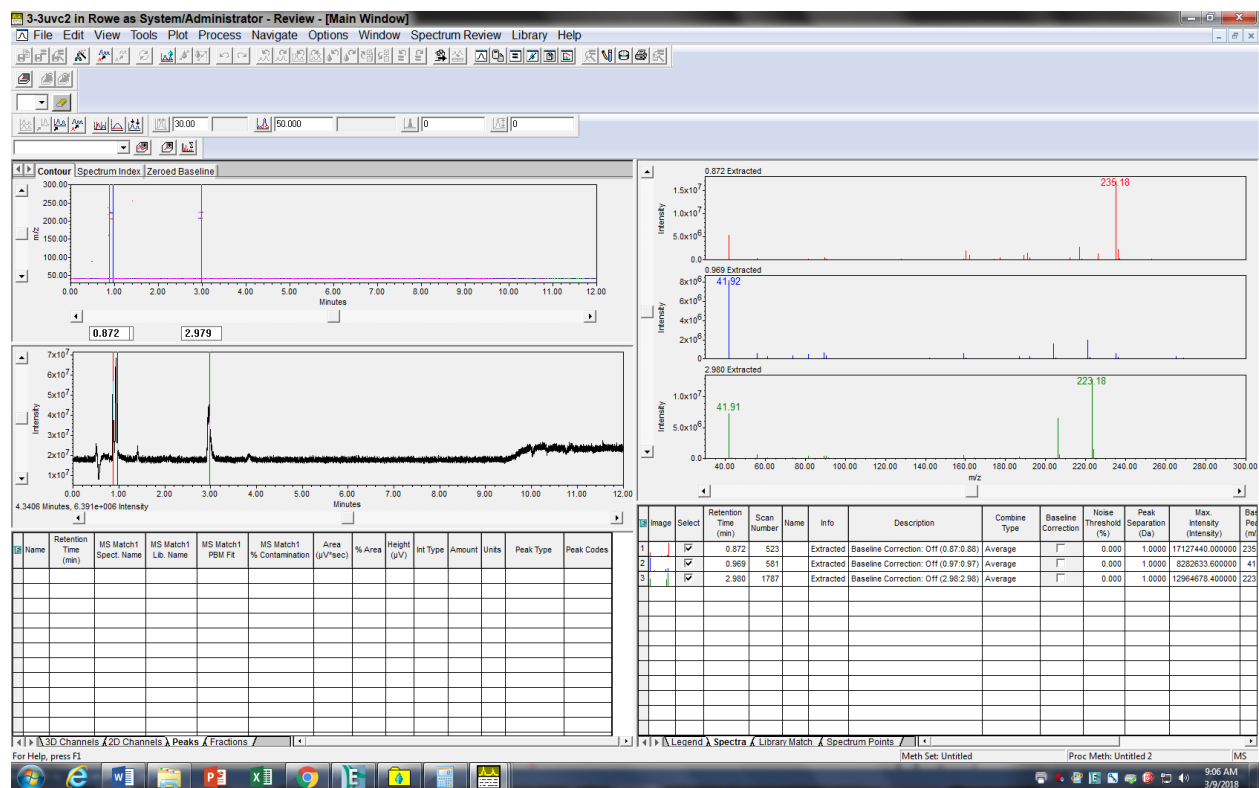


### 3-3 UV-C LC PDA Detector Data with Integrated Peak

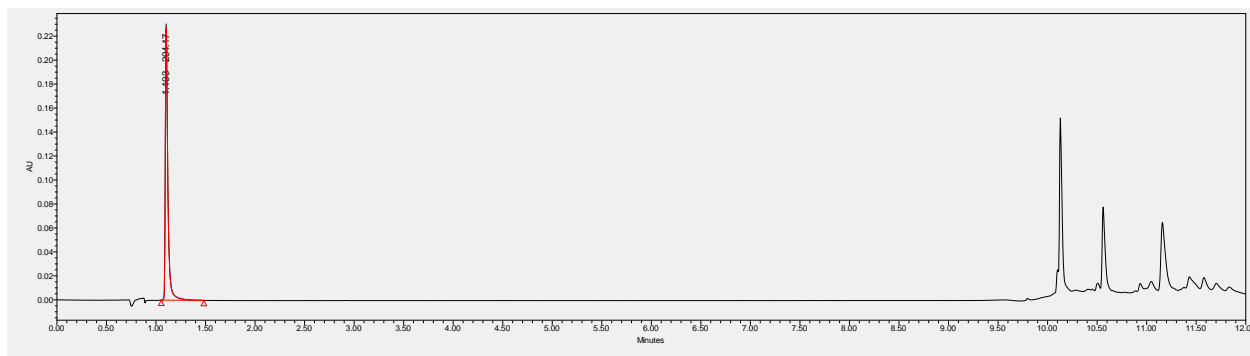


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.945	68888	100.00	18426	bb			Unknown

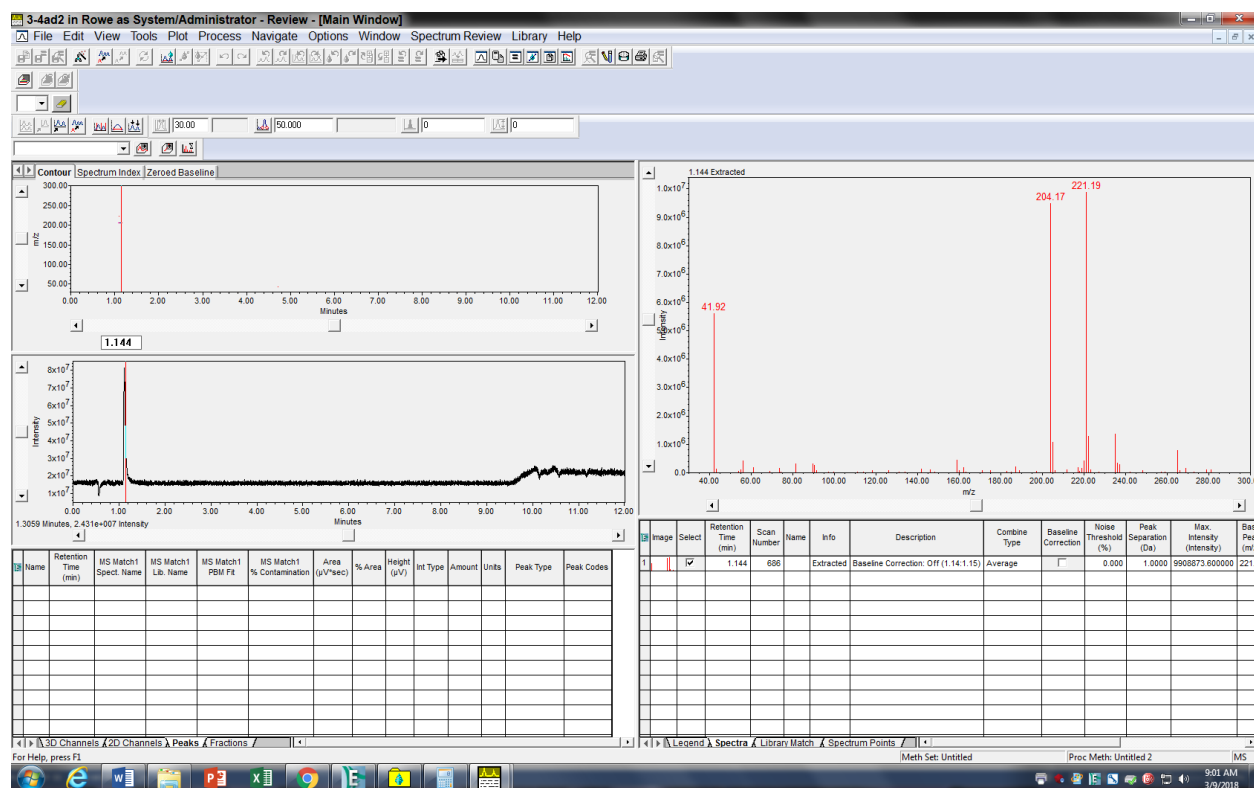
### 3-3 UVC Mass Spectrum



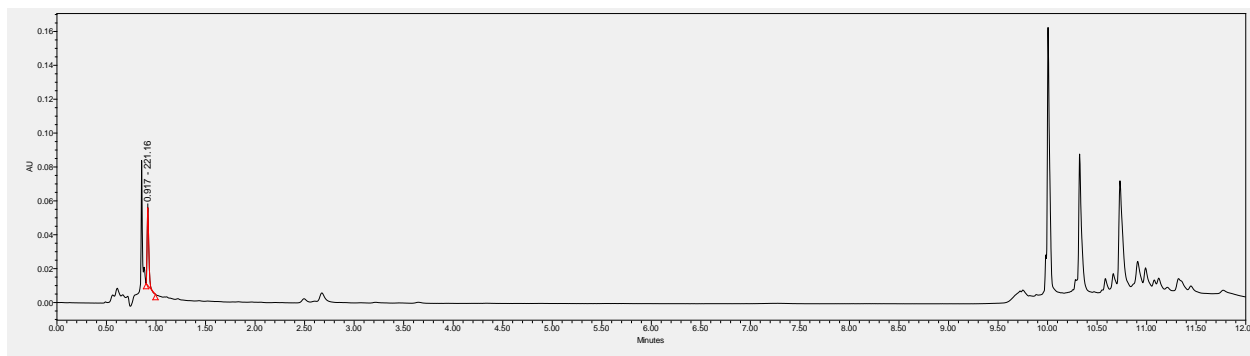
### 3-4 LC PDA Detector Data with Integrated Peak



### 3-4 Mass Spectrum

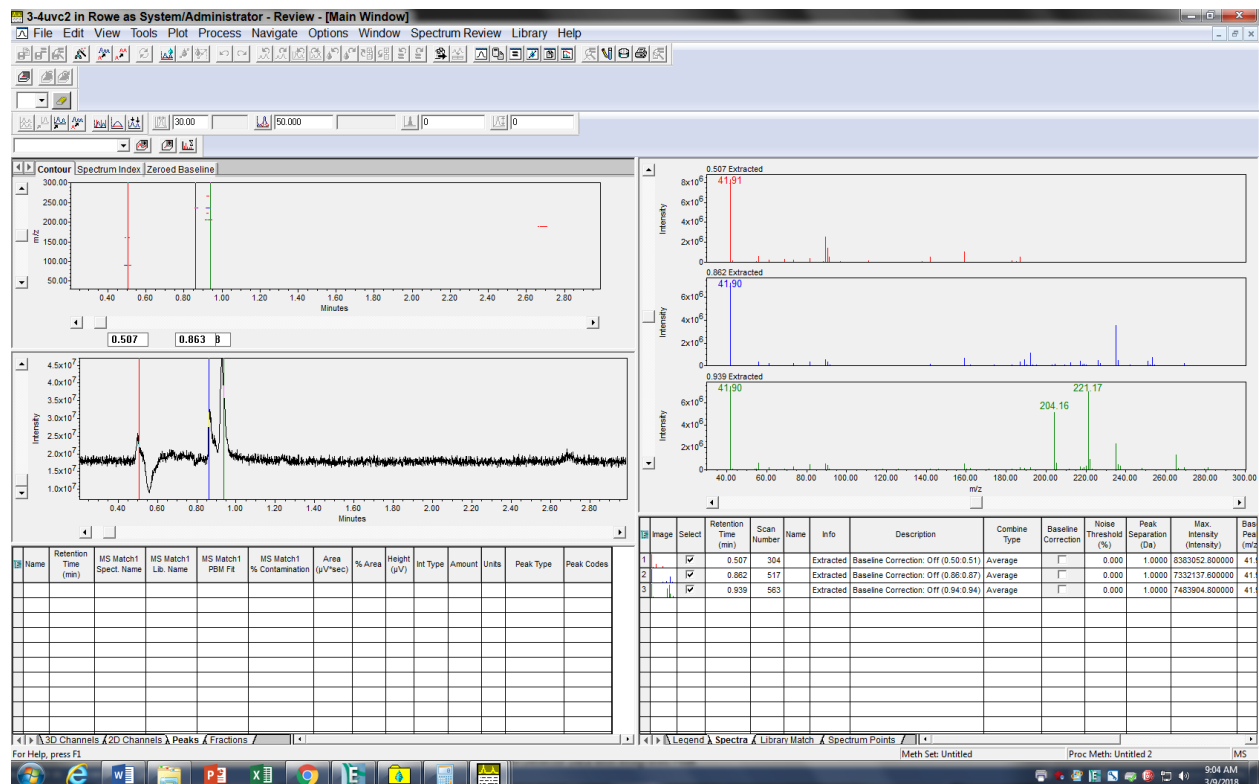


### 3-4 UV-C LC PDA Detector Data with Integrated Peak

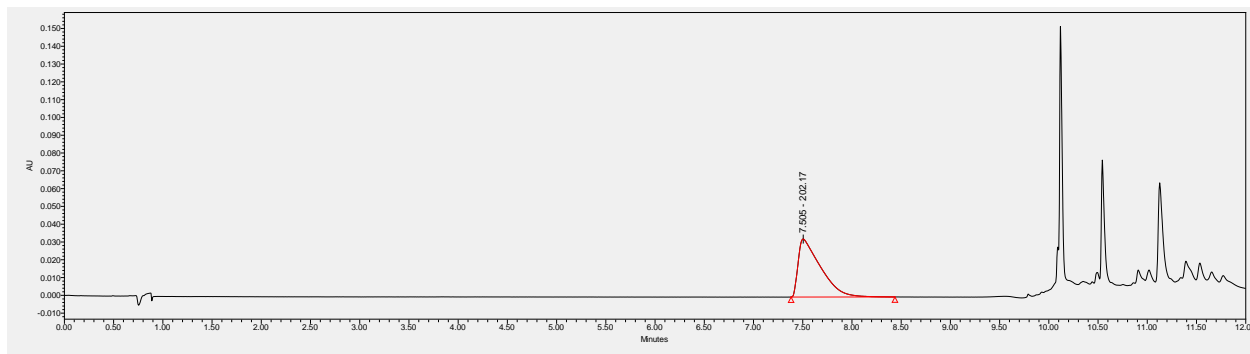


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.917	54423	100.00	45847	bb			Unknown

### 3-4 UVC Mass Spectrum

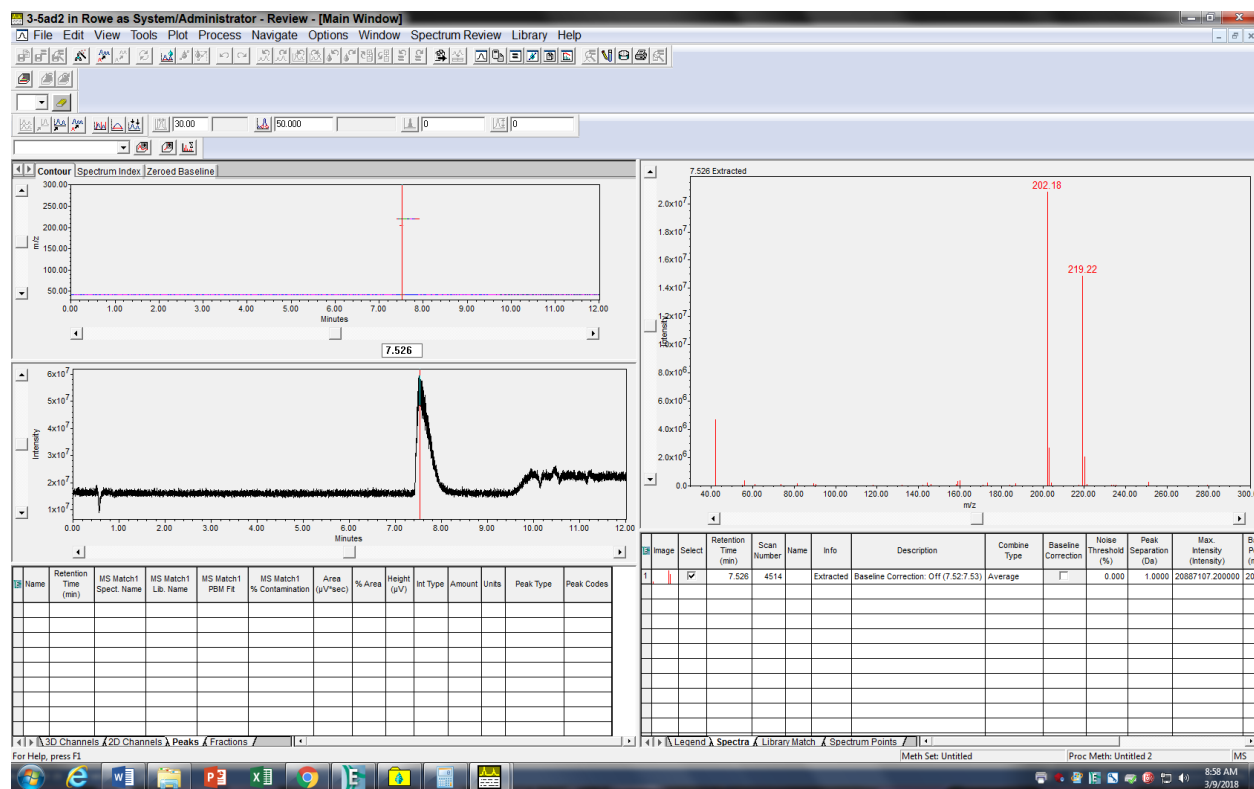


### 3-5 LC PDA Detector Data with Integrated Peak

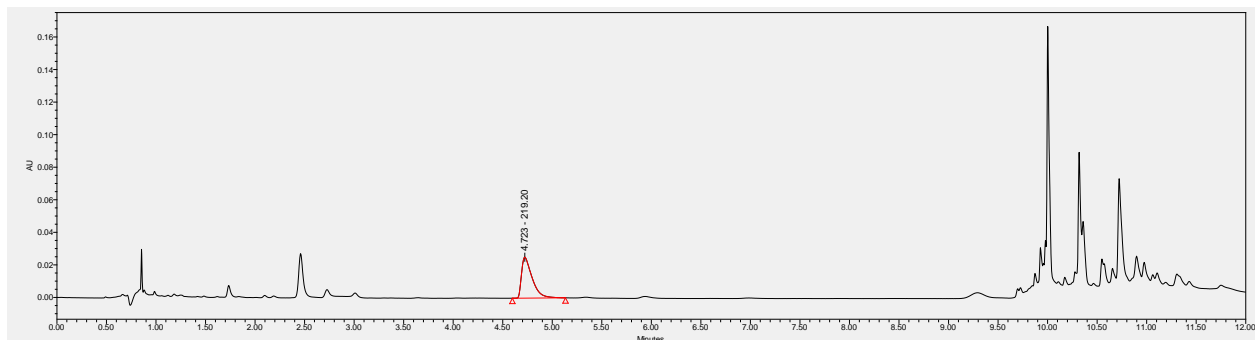


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		7.505	513514	100.00	32606	bb			Unknown

### 3-5 Mass Spectrum

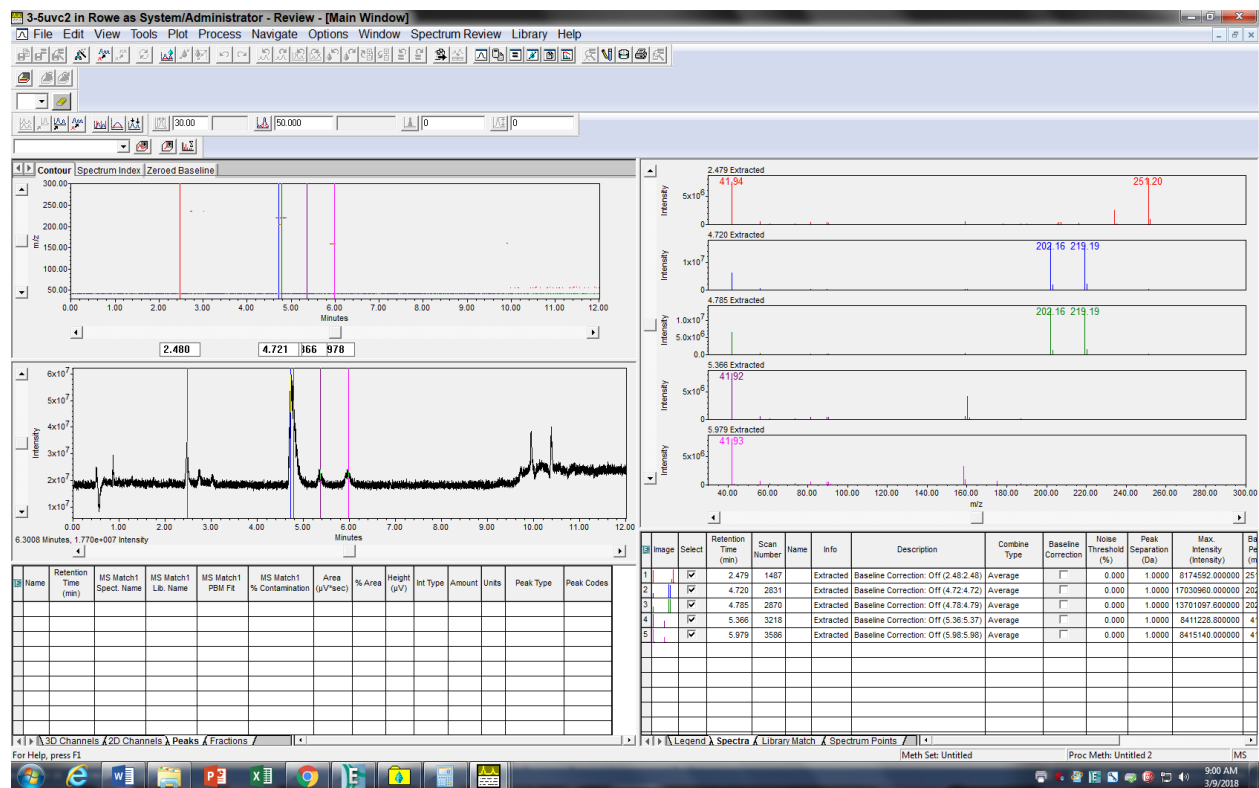


### 3-5 UV-C LC PDA Detector Data with Integrated Peak

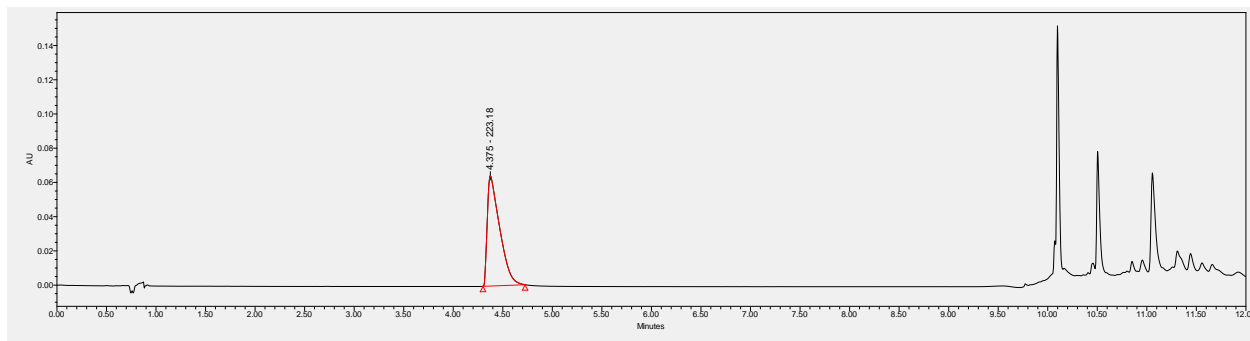


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.723	186241	100.00	25162	bb			Unknown

### 3-5 UVC Mass Spectrum

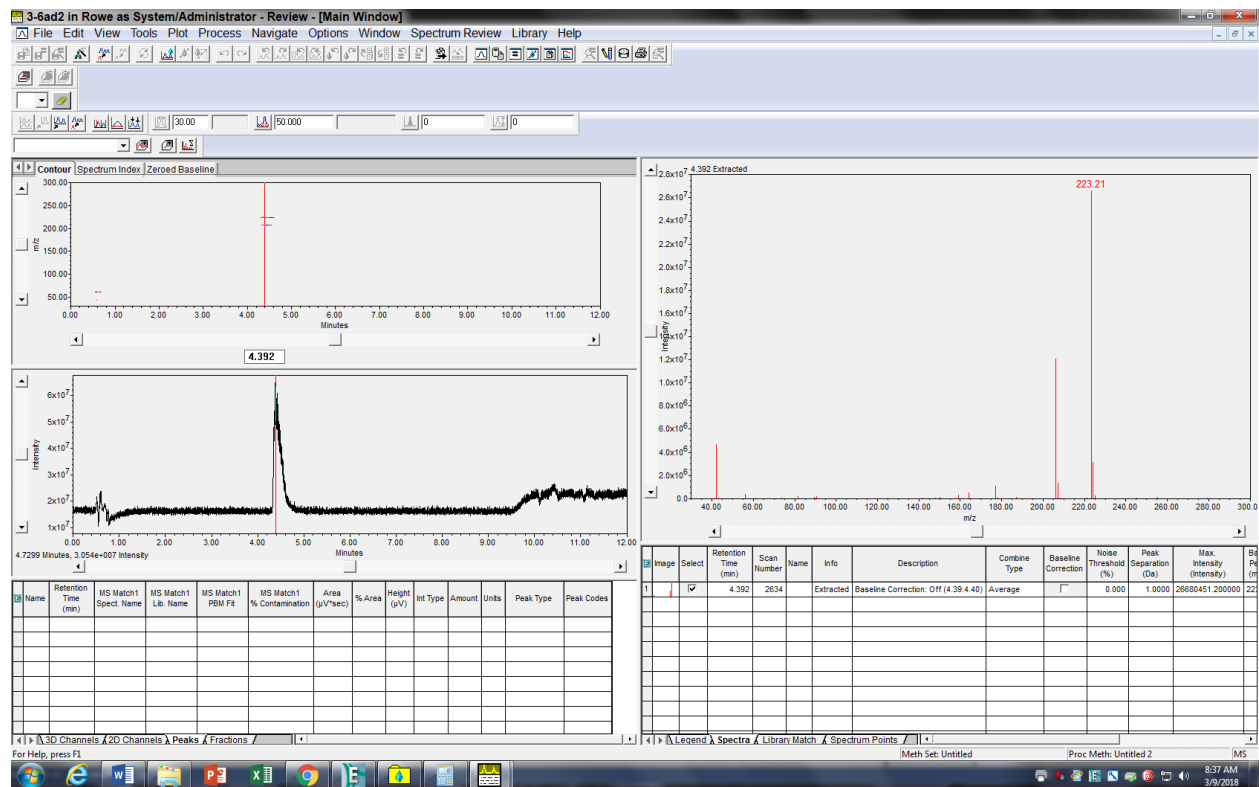


### 3-6 LC PDA Detector Data with Integrated Peak

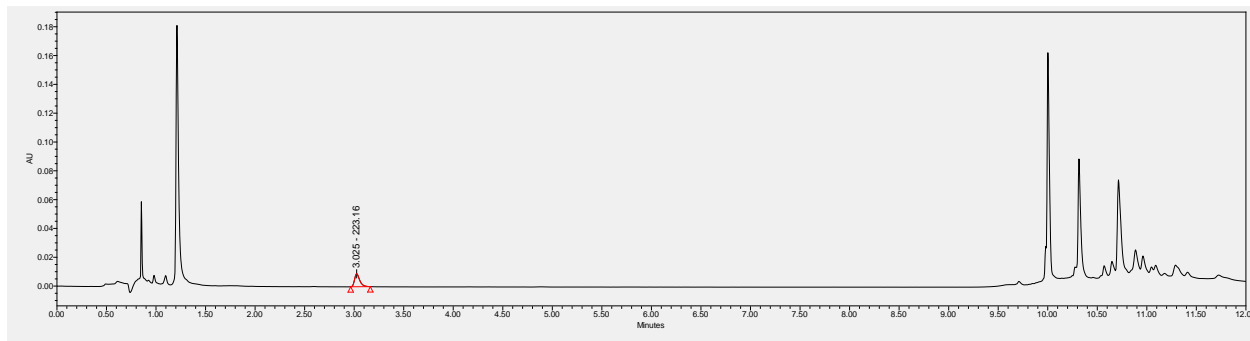


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.375	558204	100.00	64428	bb			Unknown

### 3-6 Mass Spectrum

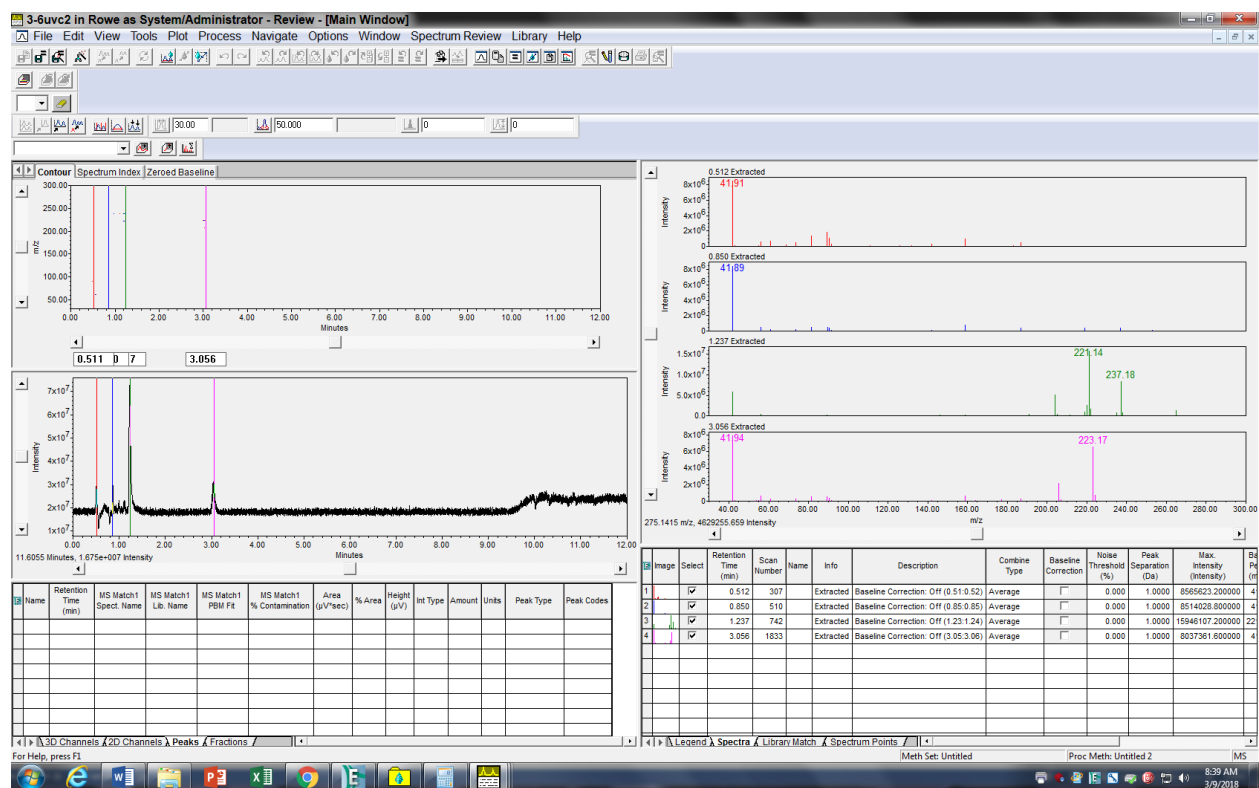


### 3-6 UV-C LC PDA Detector Data with Integrated Peak

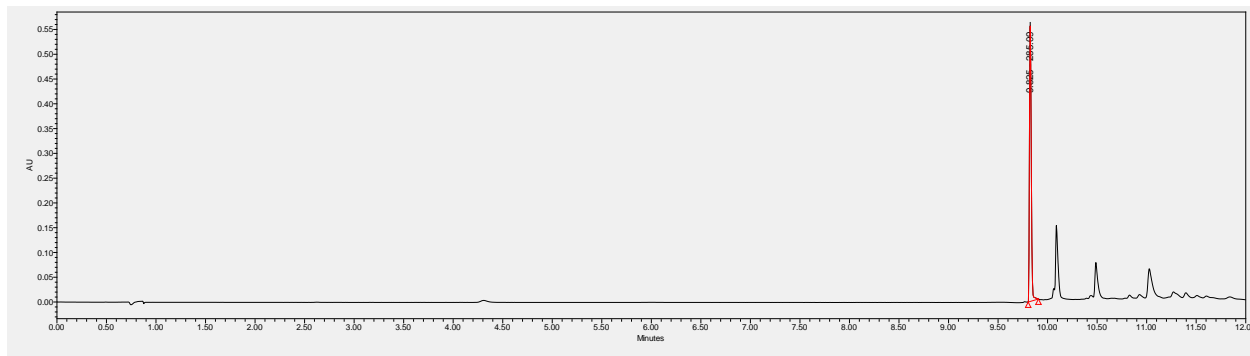


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.025	32696	100.00	8990	bb			Unknown

### 3-6 UVC Mass Spectrum

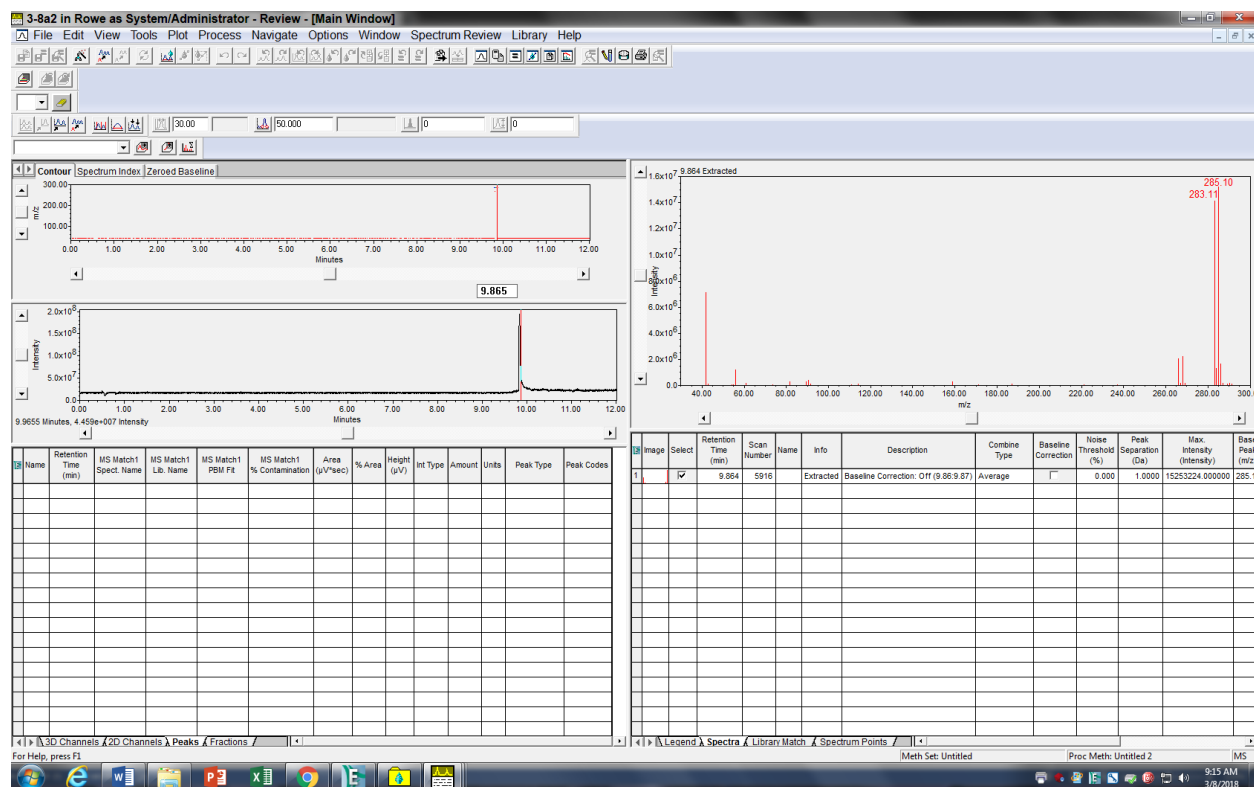


### 3-8 LC PDA Detector Data with Integrated Peak



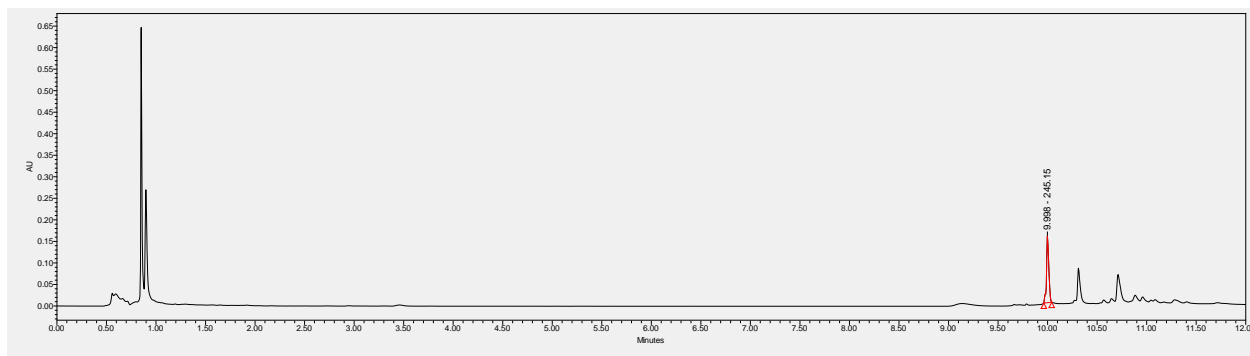
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.825	712780	100.00	556087	bb			Unknown

### 3-8 Mass Spectrum



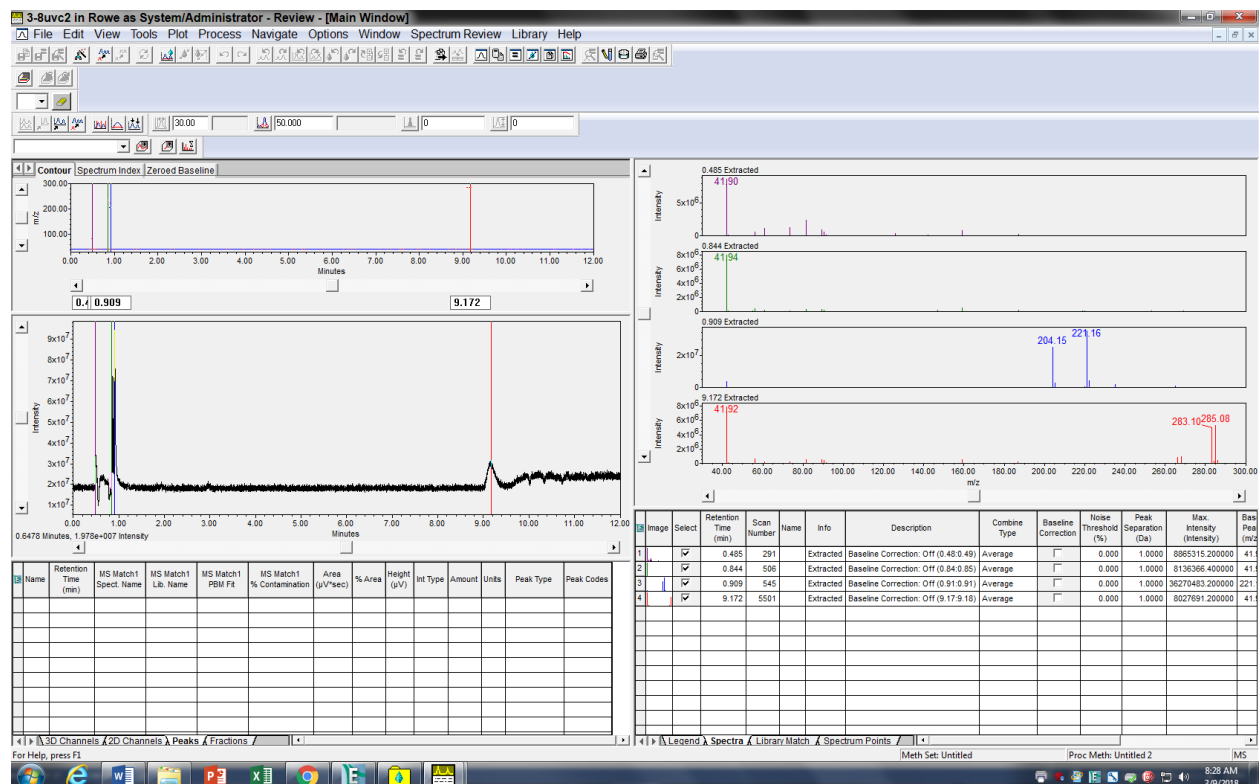


### 3-8 UV-C LC PDA Detector Data with Integrated Peak



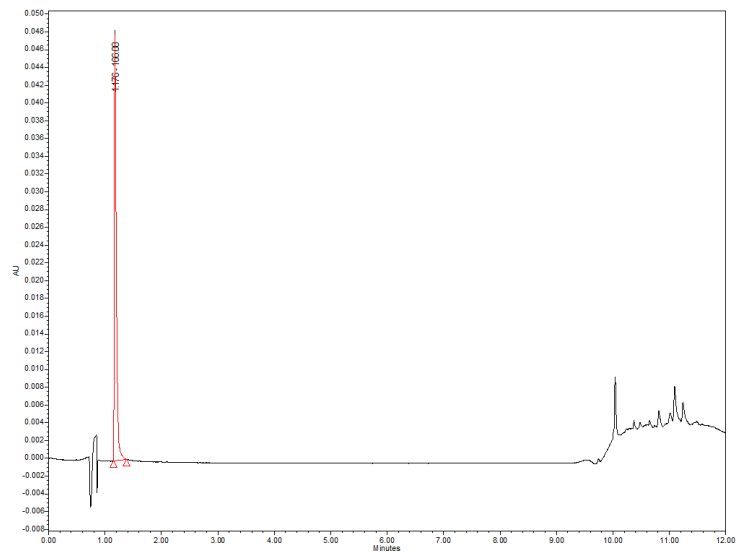
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.998	247473	100.00	154777	bb			Unknown

### 3-8 UVC Mass Spectrum



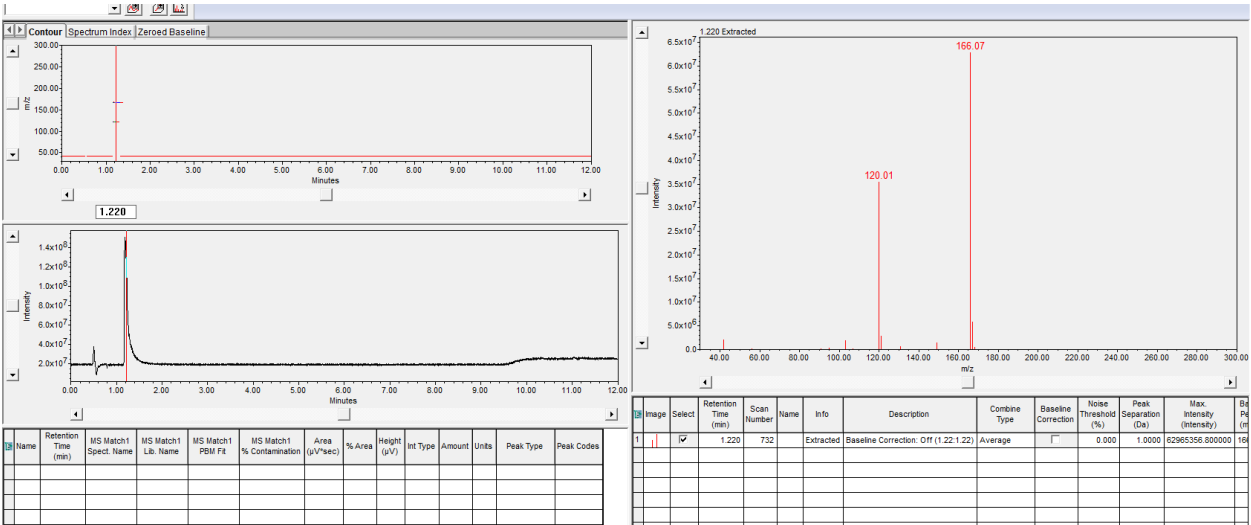
Gamma Exposure data (pg. 118-149) of LC absorbance at 254 nm, with integrated area, and corresponding extracted mass spectrum. One sample chromatogram of each control of all 3.99 krad (low) gamma irradiated samples are shown. Graphs in paper were produced by averaging the results of triplicate measurements. Amino acid identified with number in Tables 1 and 2, 1 for example, is Phe.

1 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.176	112039	100.00	48008	bb			Unknown

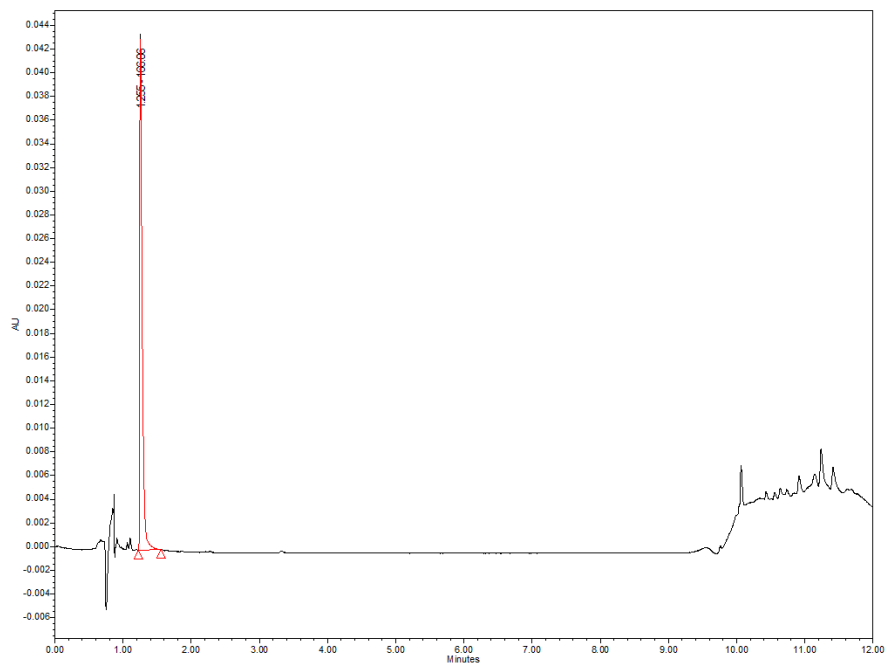
1 Mass Spectrum



	Name	Retention Time (min)	MS Match1 Spect. Name	MS Match1 Lib. Name	MS Match1 PBM File	MS Match1 % Contamination	Area (μV*sec)	% Area	Height (μV)	Int Type	Amount	Units	Peak Type	Peak Codes
1		1.220												

	Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max Intensity (Intensity)	D4 P4 (n)
1		<input checked="" type="checkbox"/>	1.220	732		Extracted	Baseline Correction: Off (1.22:1.22)	Average		0.000	1.0000	62995356.800000	16

## 1 low gamma LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.255	108865	100.00	43177	bb			Unknown

## 1 low gamma Mass Spectrum

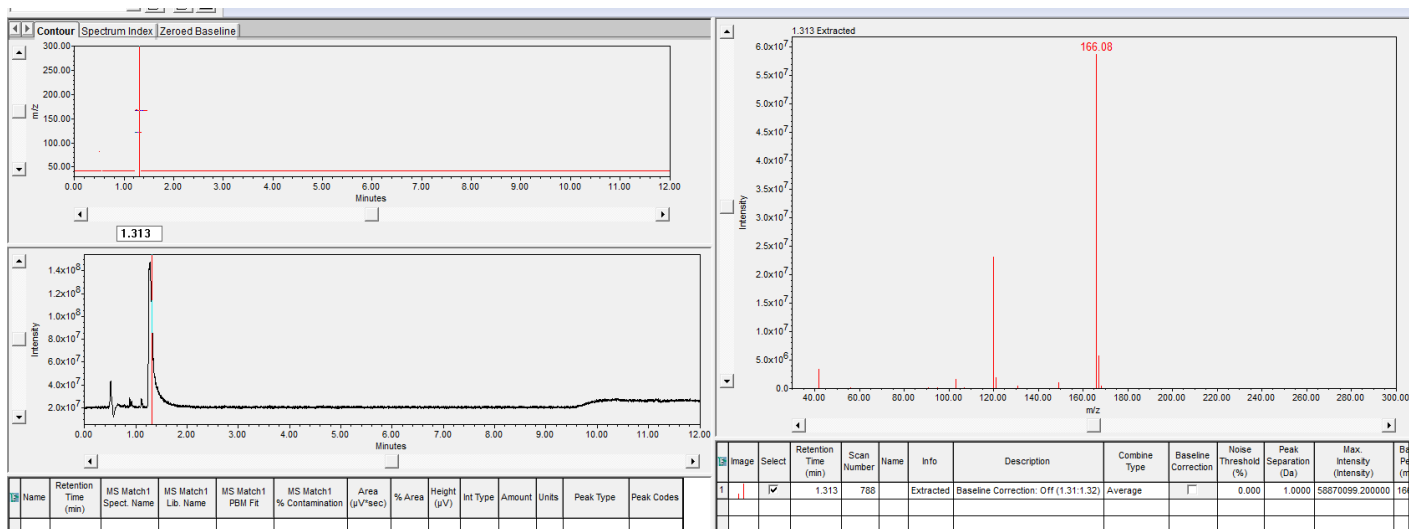
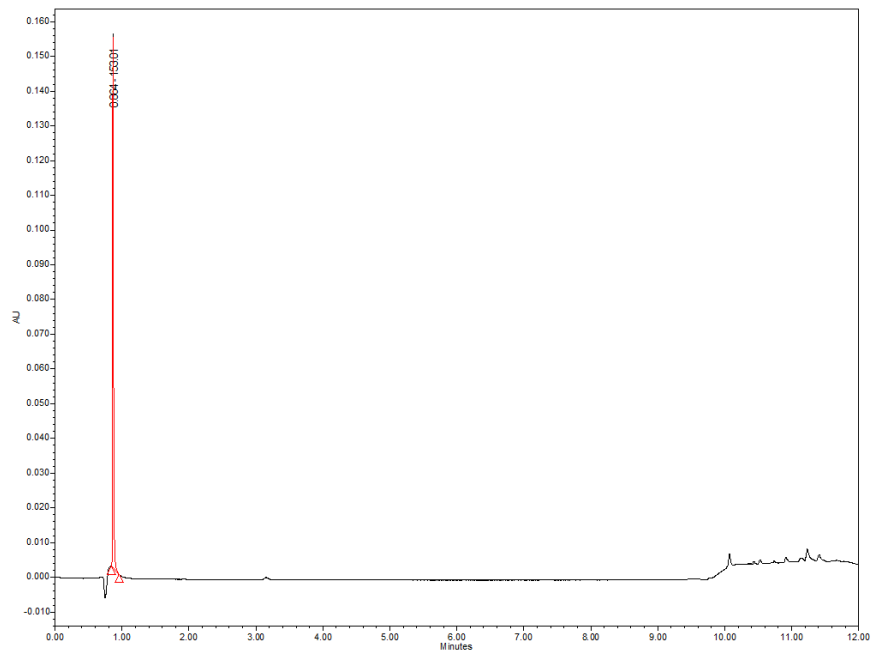


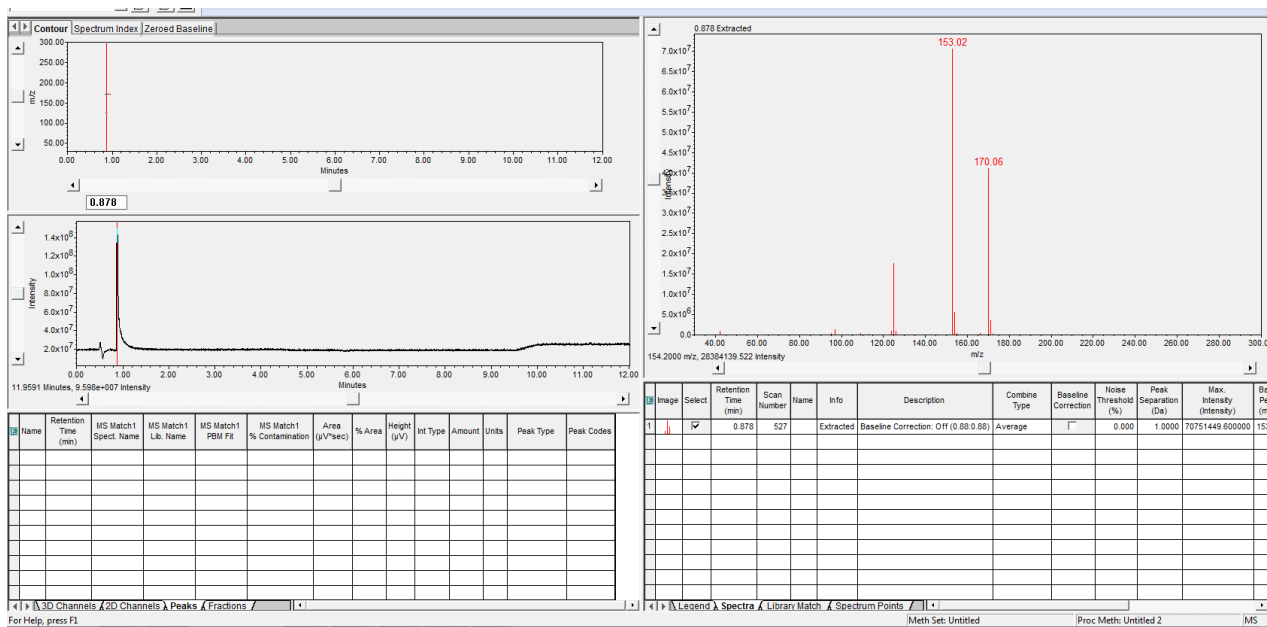
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Bt P4 (m)
1	<input checked="" type="checkbox"/>	1.313	786		Extracted	Baseline Correction: Off (1.31:1.32)	Average	<input type="checkbox"/>	0.000	1.0000	58870099.200000	166.08

## 1-3 LC PDA Detector Data with Integrated Peak

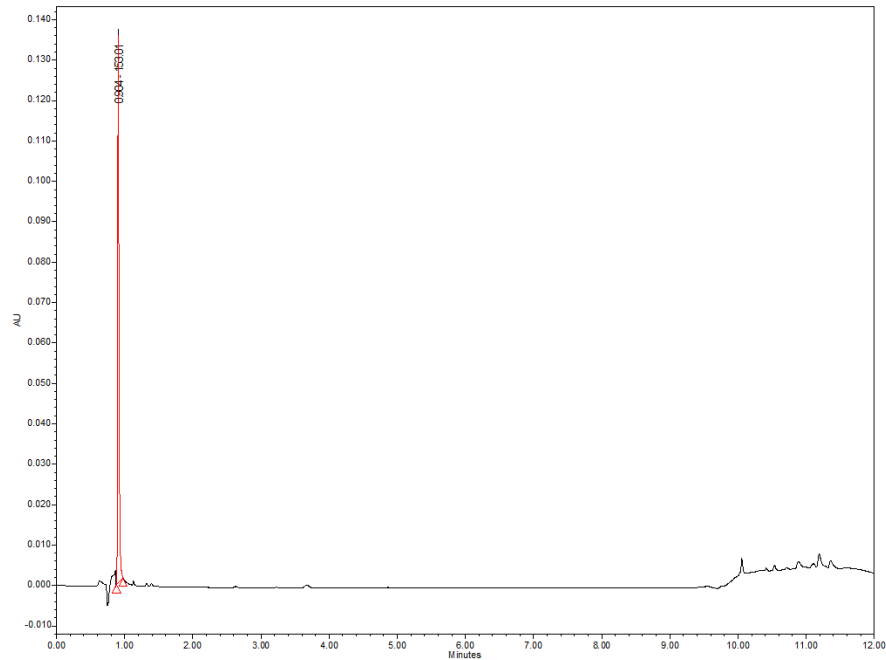


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.864	164670	100.00	152834	bb			Unknown

## 1-3 Mass Spectrum



1-3 low gamma LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.904	169674	100.00	135599	bb			Unknown

1-3 low gamma Mass Spectrum

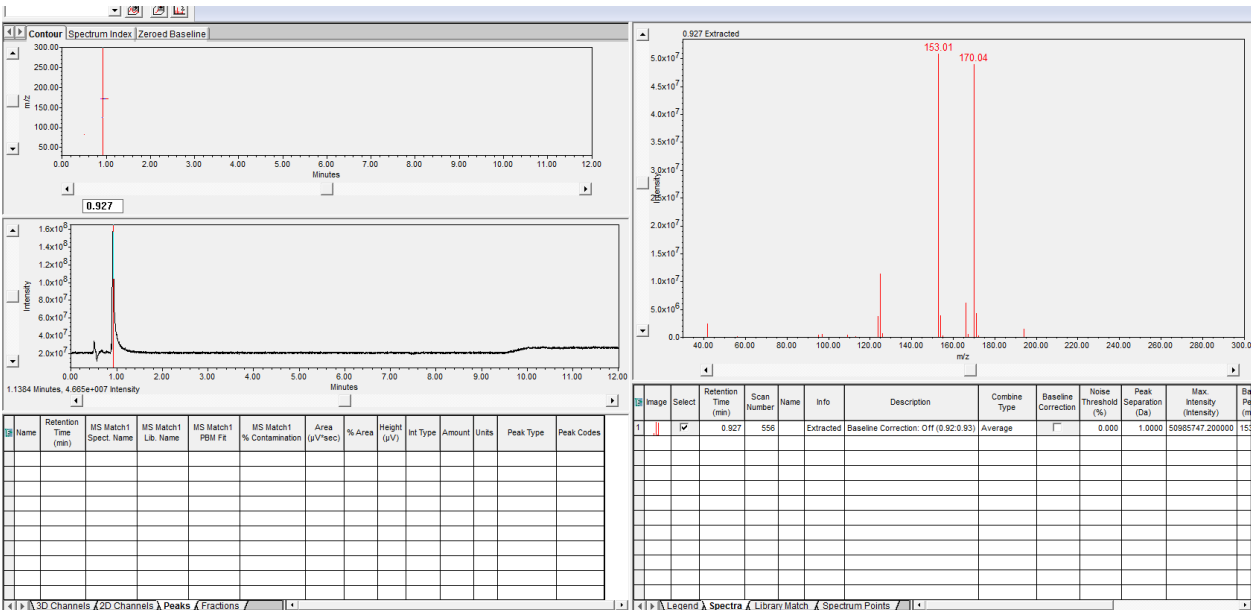
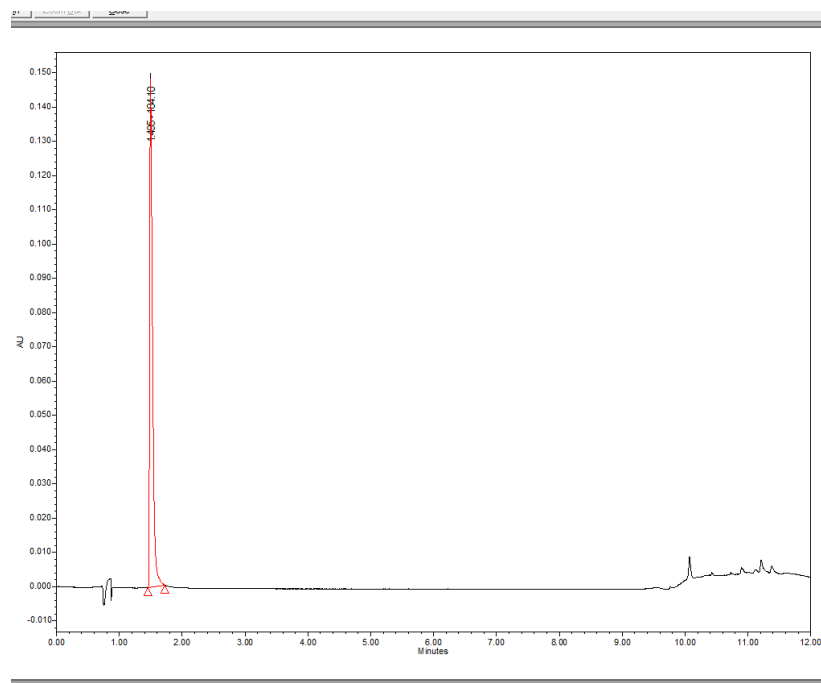


Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Base Pd (m)
1	<input checked="" type="checkbox"/>	0.927	556		Extracted	Baseline Correction: Off (0.92-0.93)	Average	<input type="checkbox"/>	0.000	1.0000	50985747.200000	15

## 1-6 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.495	454464	100.00	148569	bb			Unknown

## 1-6 Mass Spectrum

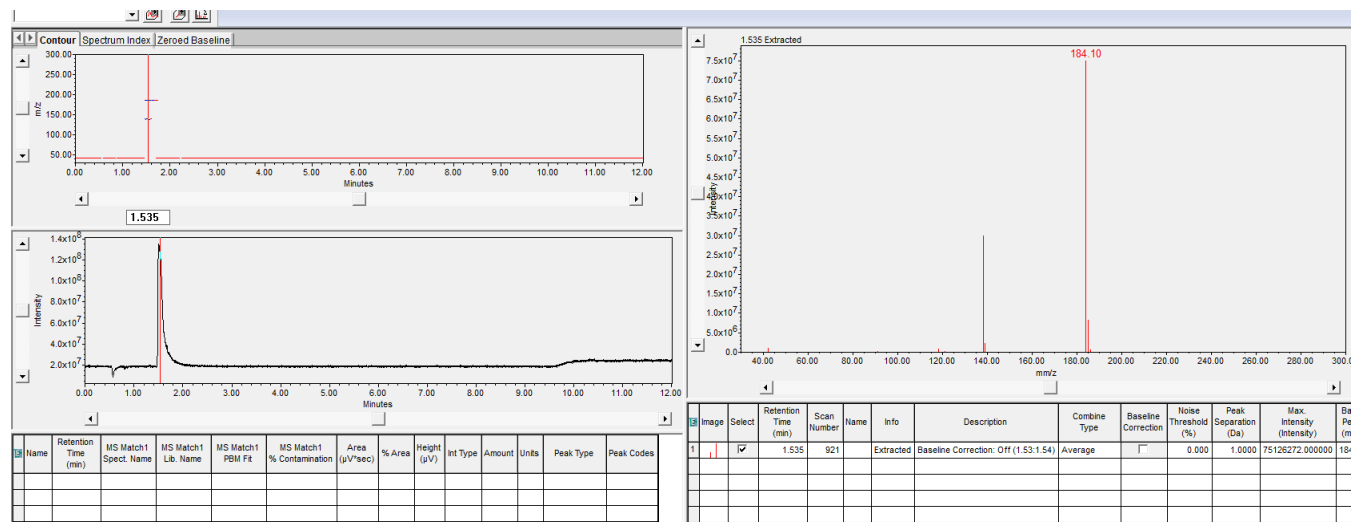
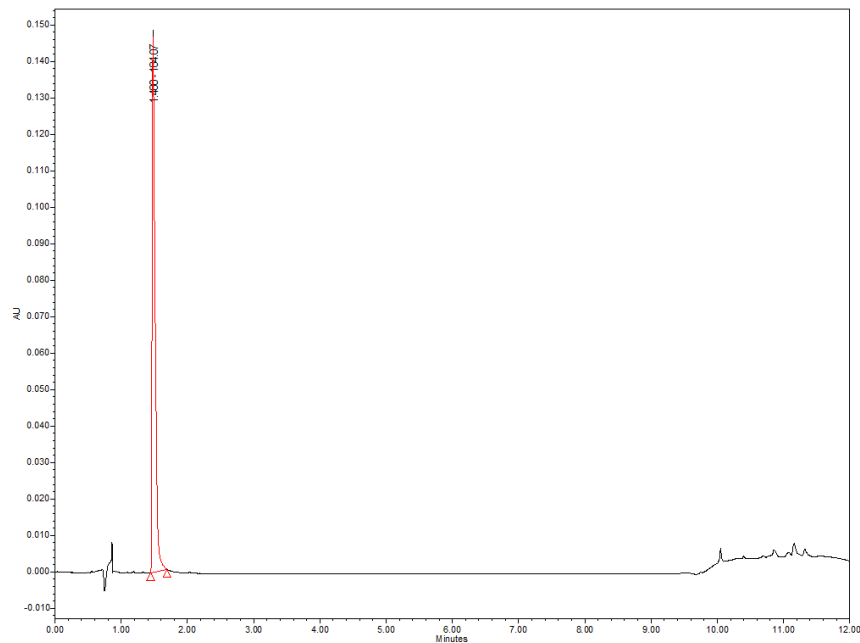


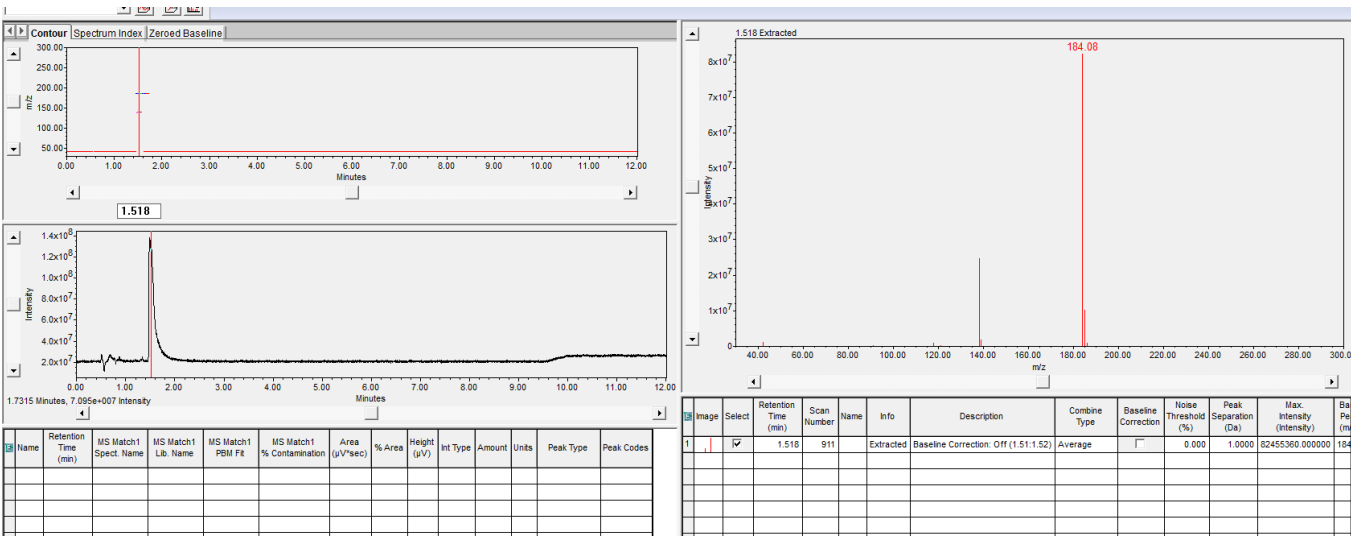
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Ba Pe (m)
1	<input checked="" type="checkbox"/>	1.535	921	Extracted	Baseline Correction: Off (1.53:1.54)	Average			0.000	1.0000	75126272.000000	184

1-6 low gamma LC PDA Detector Data with Integrated Peak

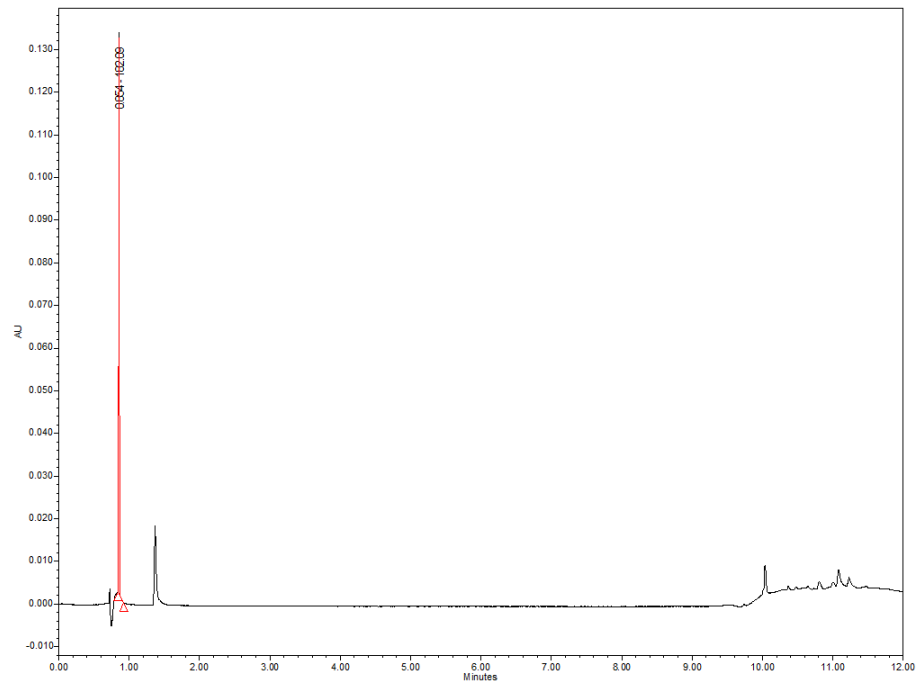


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.480	442237	100.00	146932	bb			Unknown

1-6 low gamma Mass Spectrum

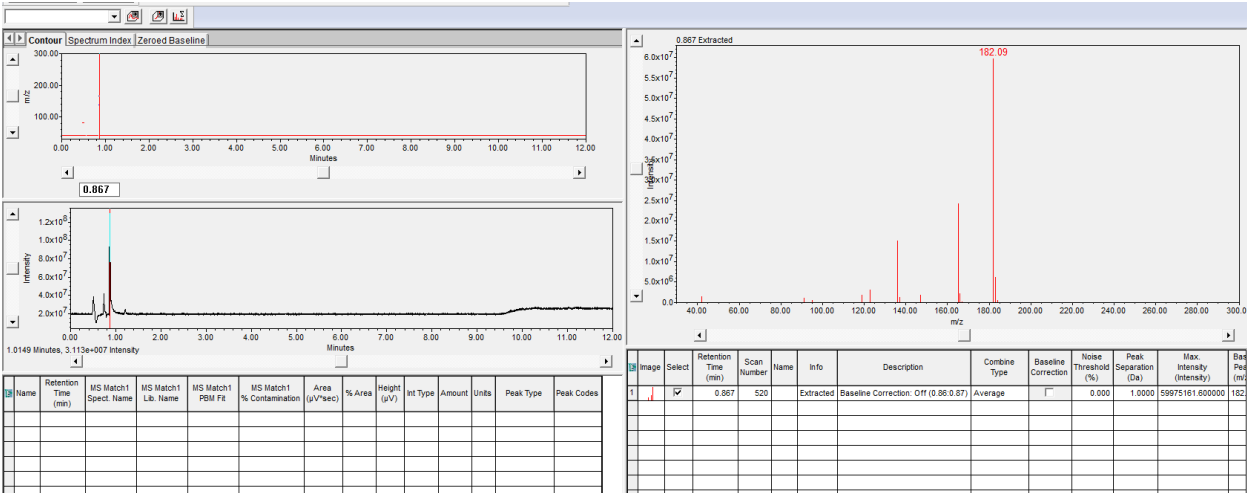


2 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.854	79210	100.00	130402	bb			Unknown

2 Mass Spectrum

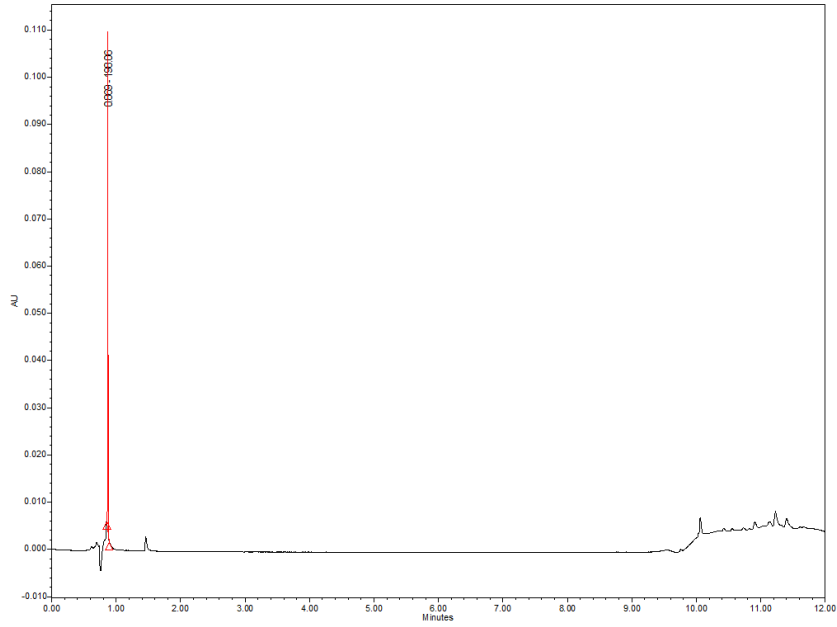


1	Name	Retention Time (min)	MS Match1 Spect. Name	MS Match1 Lib. Name	MS Match1 PSM FA	MS Match1 % Contamination	Area (μV*sec)	% Area	Height (μV)	Int Type	Amount	Units	Peak Type	Peak Codes
1		0.867								Extracted				

1	Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Base Peak (m/z)
1		<input checked="" type="checkbox"/>	0.867	520	Extracted	Baseline Correction: Off (0.86/0.87)	Average			0.000	1.0000	59975161.600000	182

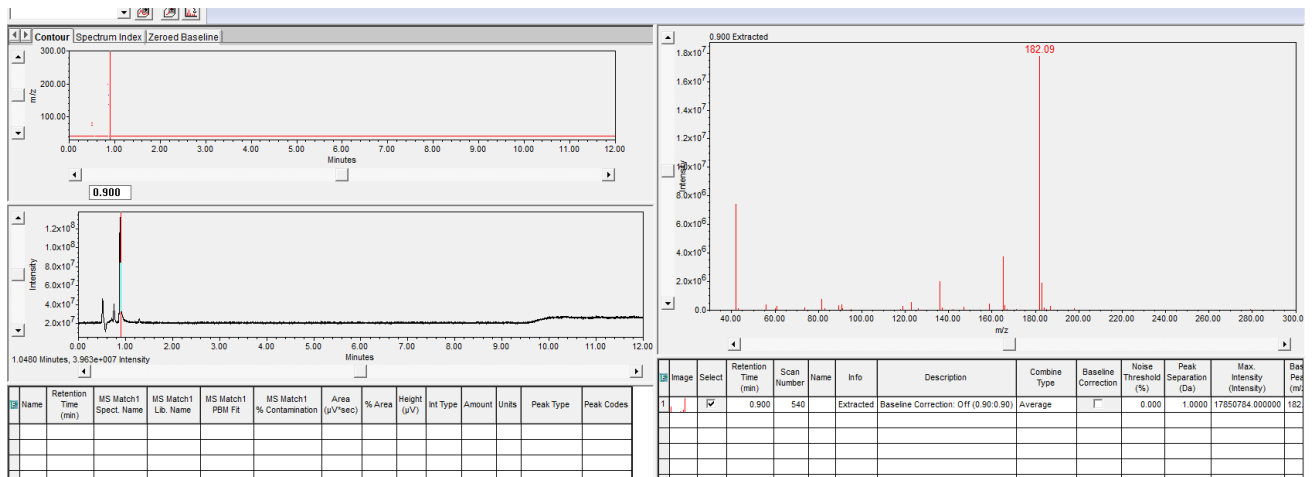


## 2 low gamma LC PDA Detector Data with Integrated Peak

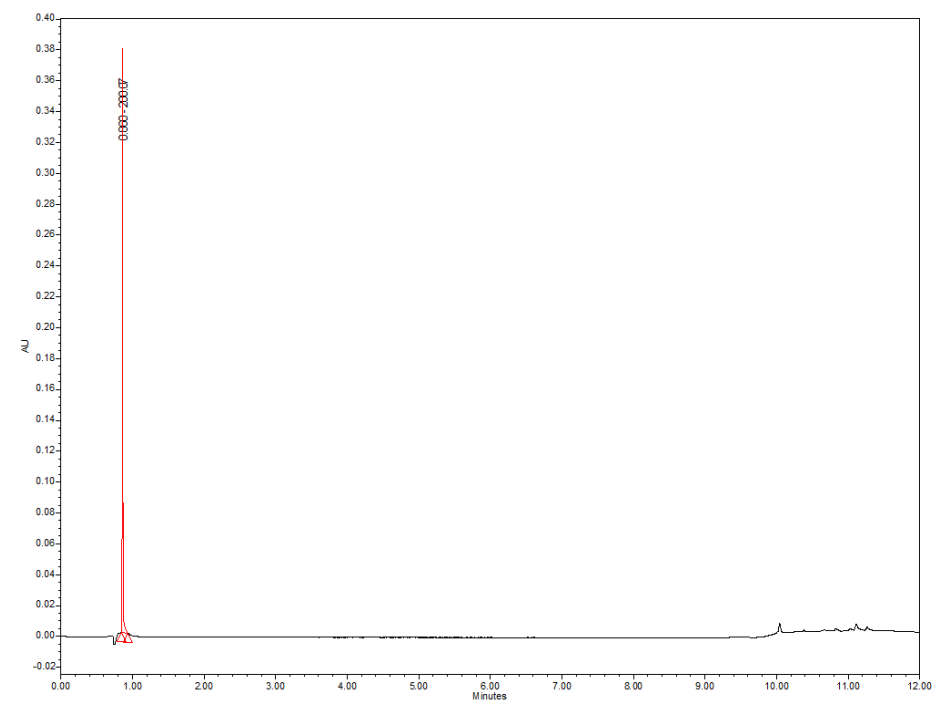


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.869	64135	100.00	105623	bb			Unknown

## 2 low gamma Mass Spectrum

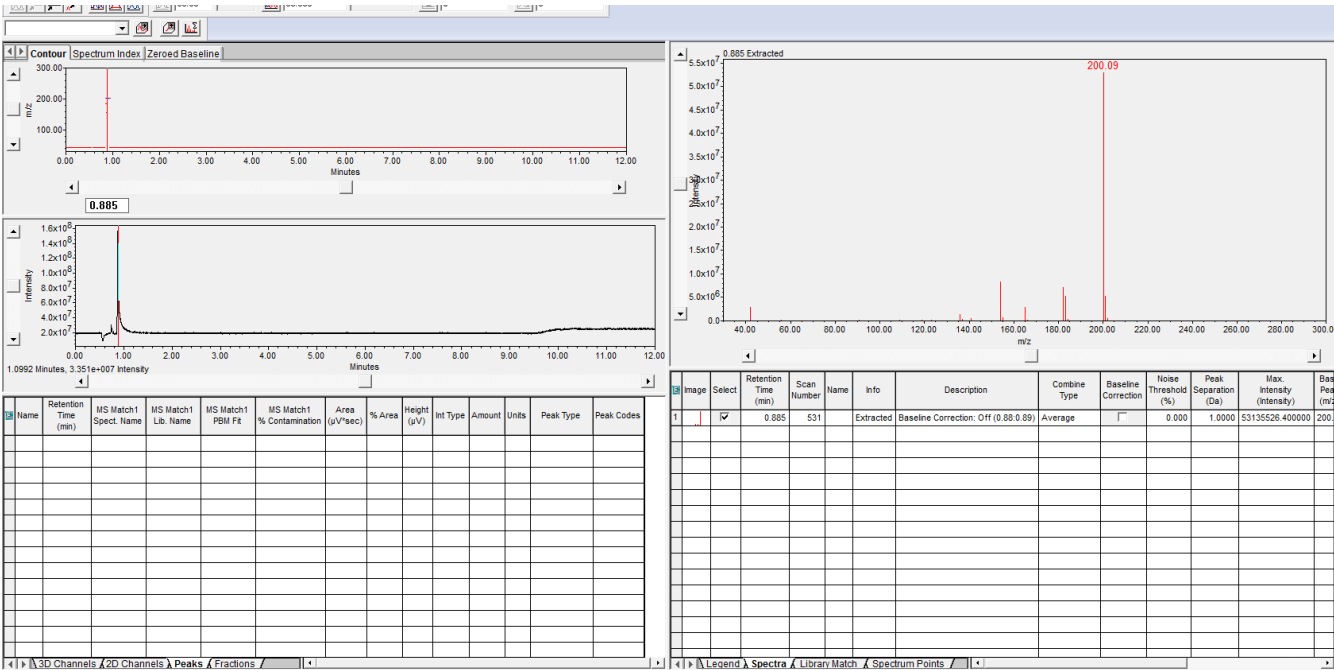


2-1 LC PDA Detector Data with Integrated Peak

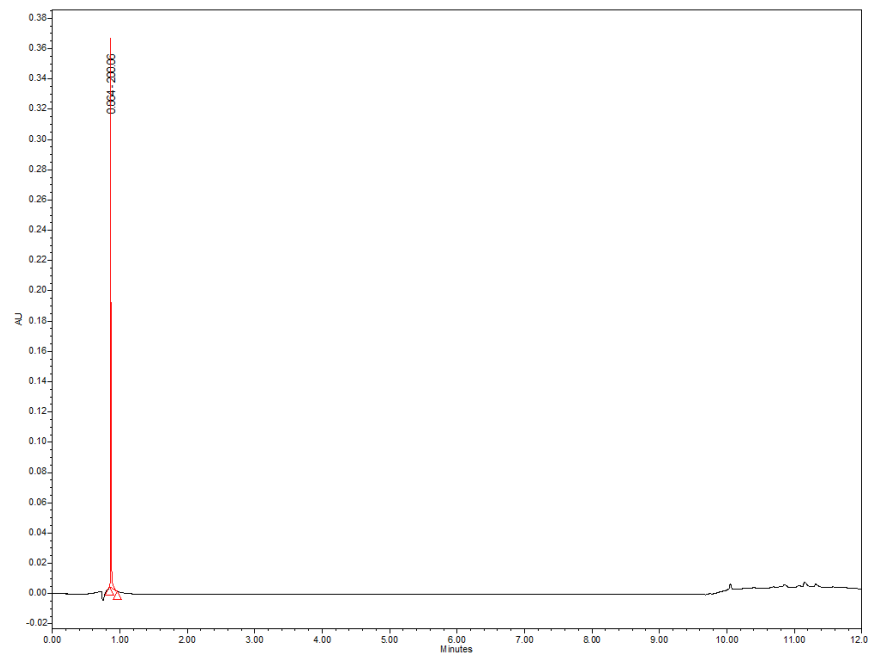


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.860	259974	100.00	379612	bb			Unknown

2-1 Mass Spectrum

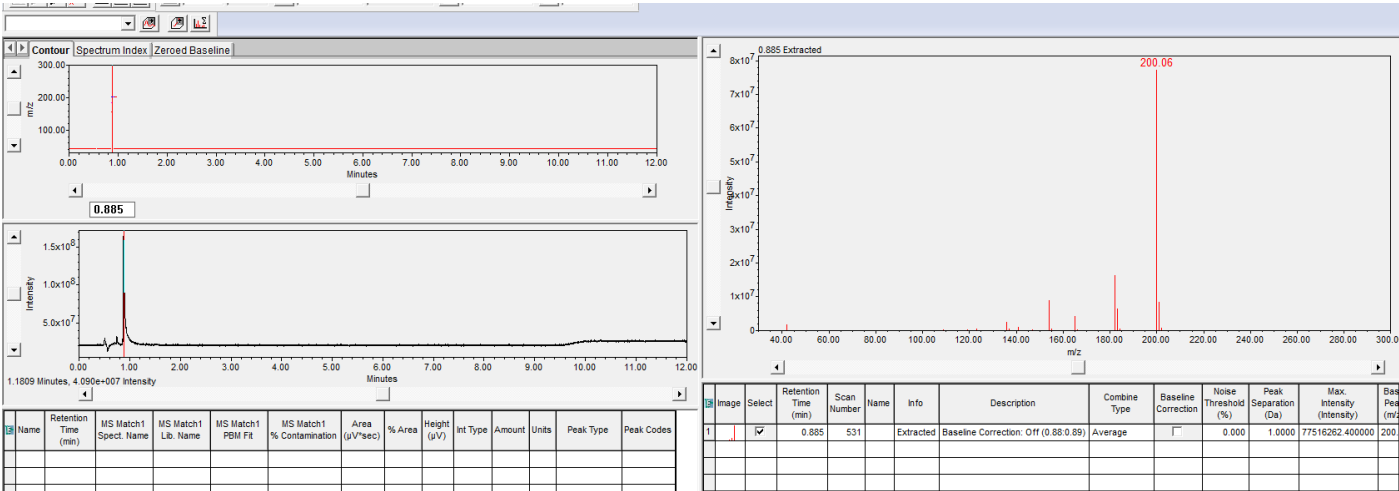


2-1 low gamma LC PDA Detector Data with Integrated Peak

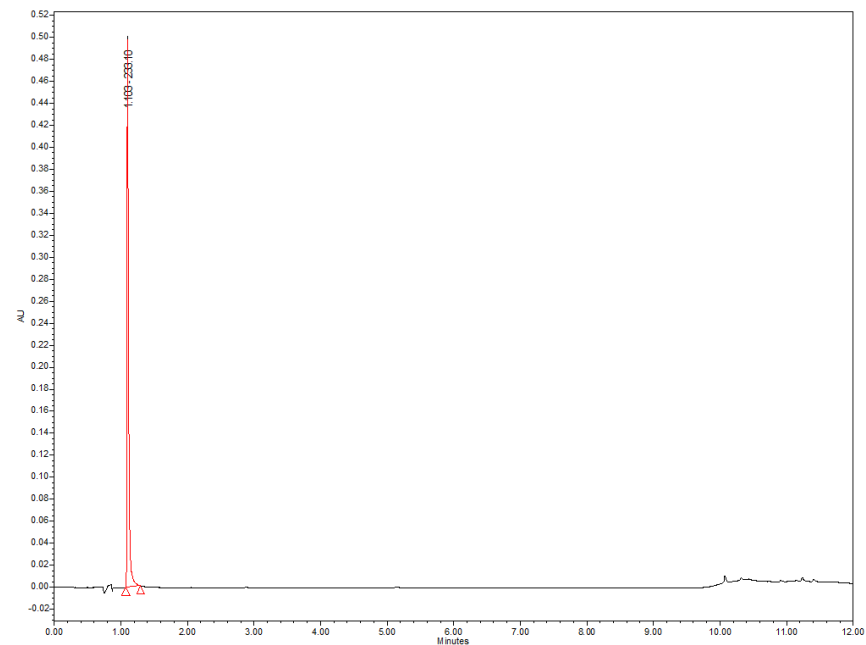


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.864	249960	100.00	363262	bb			Unknown

2-1 low gamma Mass Spectrum



4 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.103	802920	100.00	498268	bb			Unknown

4 Mass Spectrum

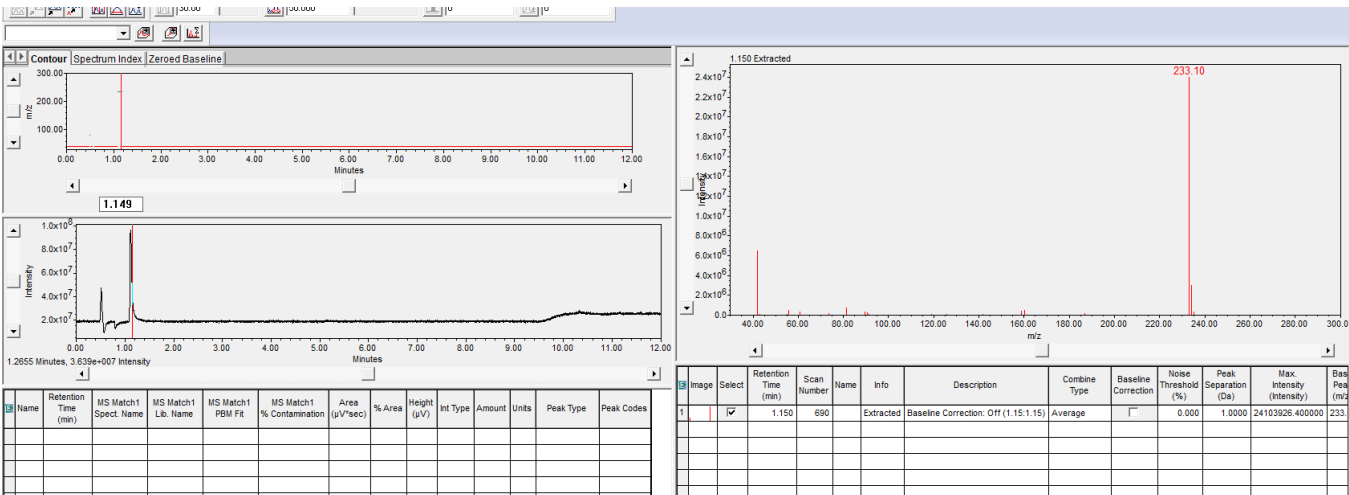
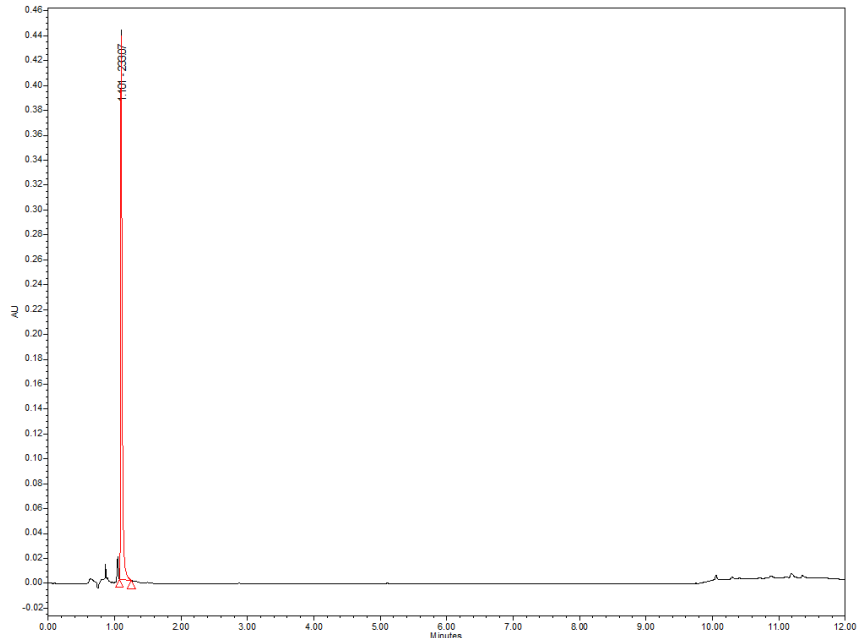


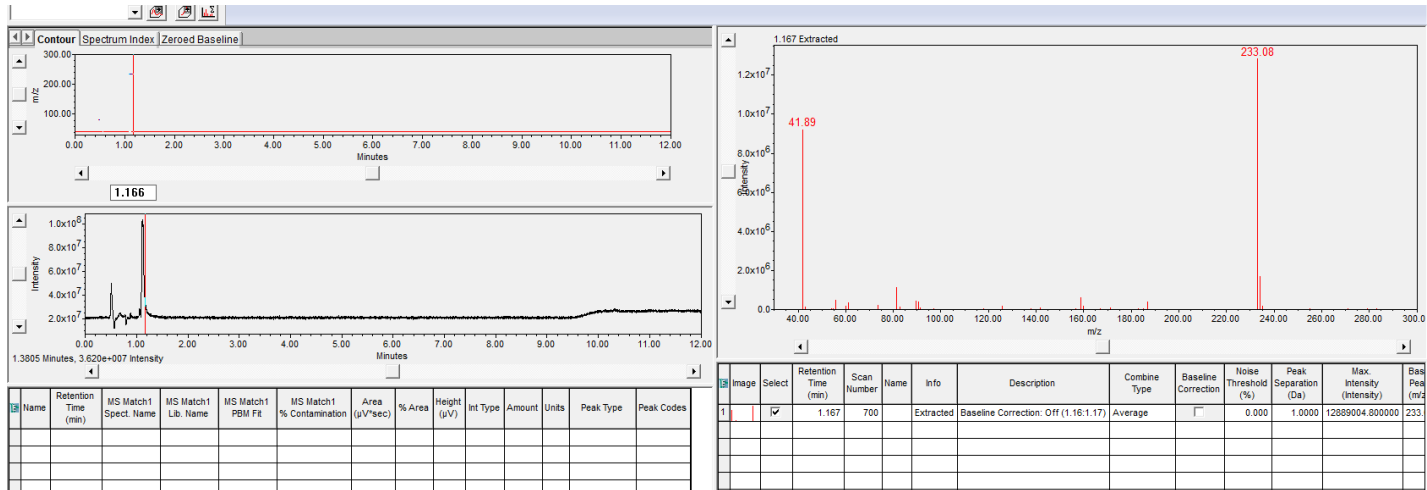
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max Intensity	Bas	Pea
1	<input checked="" type="checkbox"/>	1.150	690	Extracted	Baseline Correction: Off (1.15:1.15)	Average			0.000	1.0000	24103926.400000	233	

#### 4 low gamma LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.101	691384	100.00	437144	bb			Unknown

#### 4 low gamma Mass Spectrum



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.302	571180	100.00	243263	bb			Unknown

**Chromatogram 1 (Top Left):** TIC (Total Ion Chromatogram) showing a single sharp peak at 1.316 minutes. The y-axis is labeled 'm/z' and ranges from 0 to 300.00. The x-axis is labeled 'Minutes' and ranges from 0.00 to 12.00.

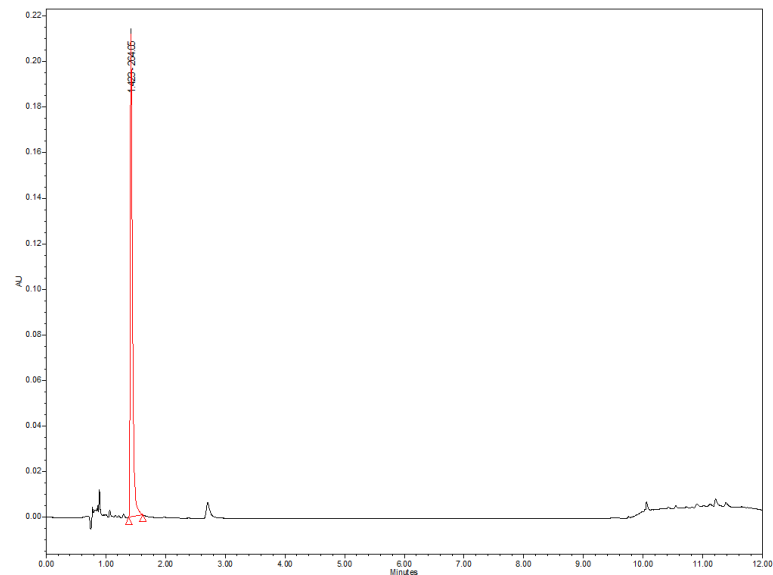
**Chromatogram 2 (Top Right):** Extracted peak at 1.317 minutes. The y-axis is labeled 'Intensity' and ranges from 0 to 7x10<sup>7</sup>. The x-axis is labeled 'm/z' and ranges from 141.1394 to 300.0. A single sharp peak is visible at m/z 264.05.

**Chromatogram 3 (Bottom Left):** Extracted peak at 1.316 minutes. The y-axis is labeled 'Intensity' and ranges from 0 to 8x10<sup>7</sup>. The x-axis is labeled 'Minutes' and ranges from 0.00 to 12.00. A single sharp peak is visible at 1.316 minutes.

**Table 1 (Bottom Right):** Peak table with columns: Image, Select, Retention Time (min), Scan Number, Name, Info, Description, Combine Type, Baseline Correction, Noise Threshold, Peak Separation, Max Intensity, and Bas Peak.

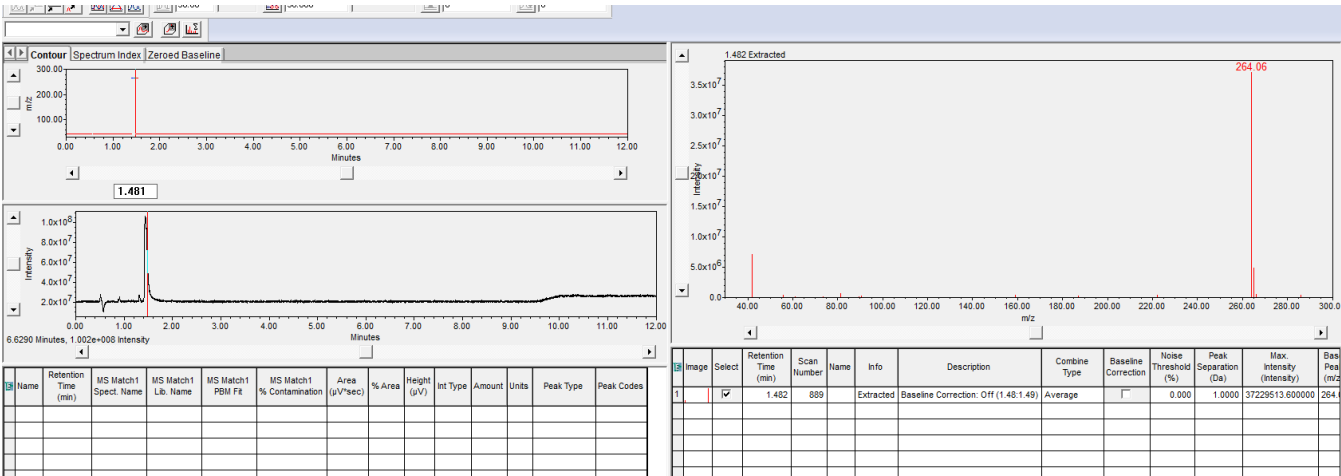
Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold	Peak Separation	Max Intensity	Bas Peak
1	<input checked="" type="checkbox"/>	1.317	790	Extracted		Baseline Correction: Off (1.31:1.32)	Average	<input type="checkbox"/>	0.000	1.0000	67040352.000000	264

5 low gamma LC PDA Detector Data with Integrated Peak

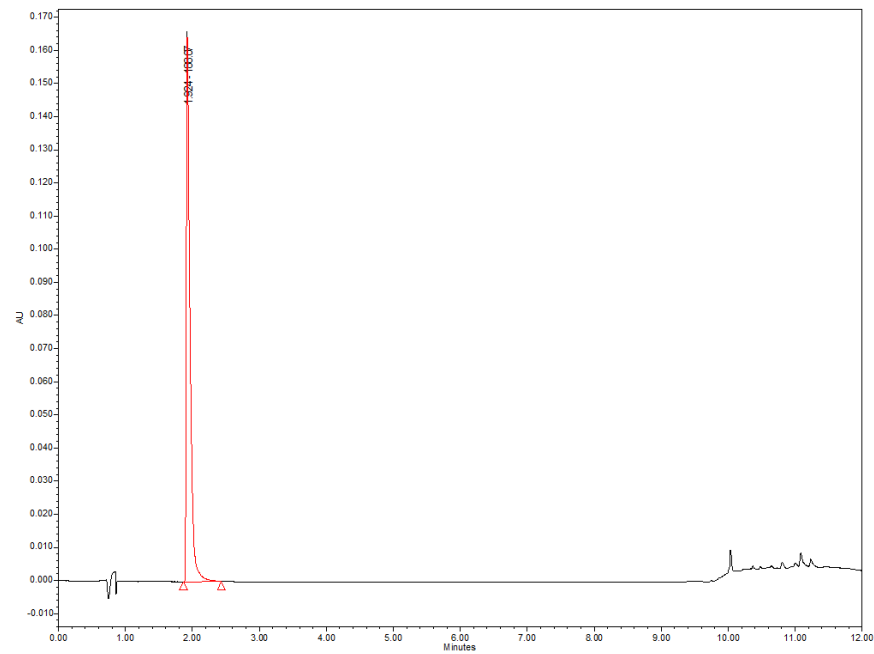


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.423	524622	100.00	212003	bb			Unknown

5 low gamma Mass Spectrum

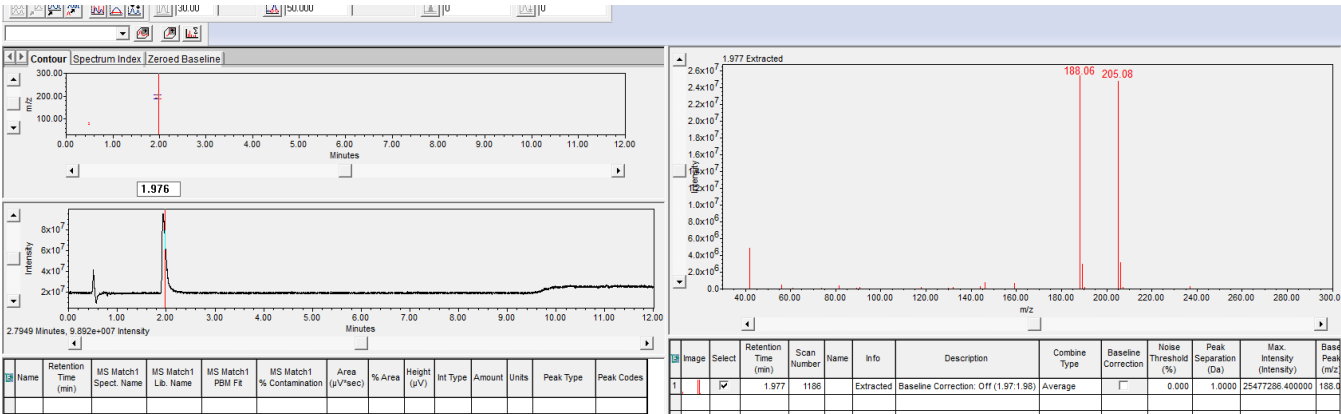


3 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		1.924	648318	100.00	164333	bb			Unknown

3 Mass Spectrum

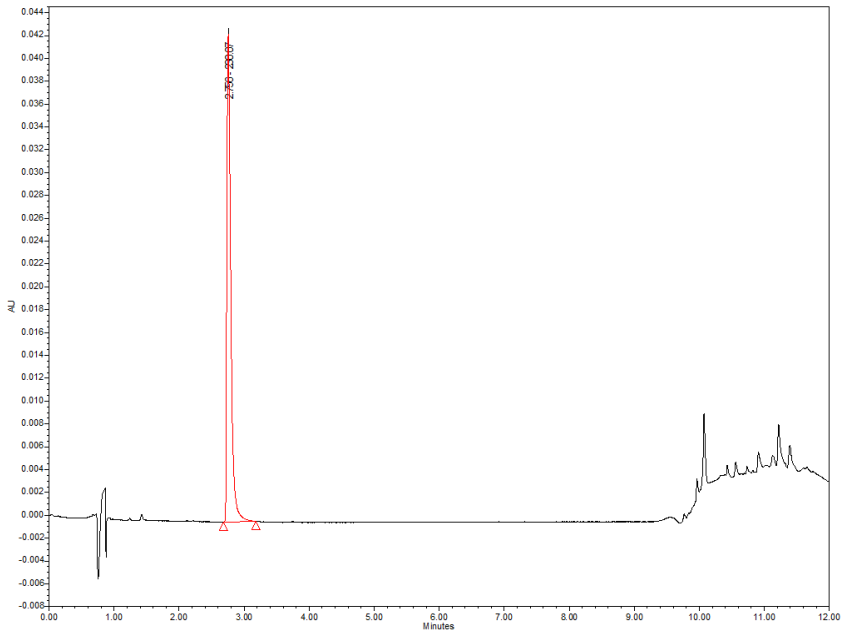




	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.105	570125	100.00	136210	bb			Unknown

Figure 1 displays mass spectrometry data for peak 1. The top panel shows the extracted ion chromatogram (EIC) for peak 1 at 2.133 minutes. The bottom panel shows the mass spectrum of the peak, with the base peak at m/z 205.09. The x-axis is m/z (40-300) and the y-axis is intensity (0.0 to 4.0x10<sup>7</sup>).

3-1 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.756	176427	100.00	42742	bb			Unknown

3-1 Mass Spectrum

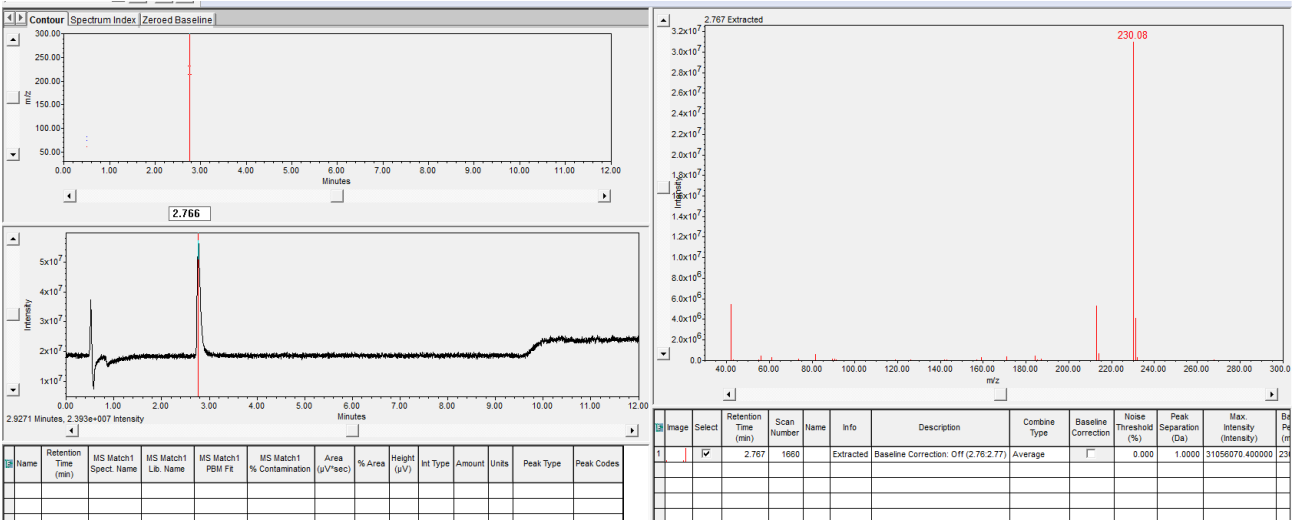
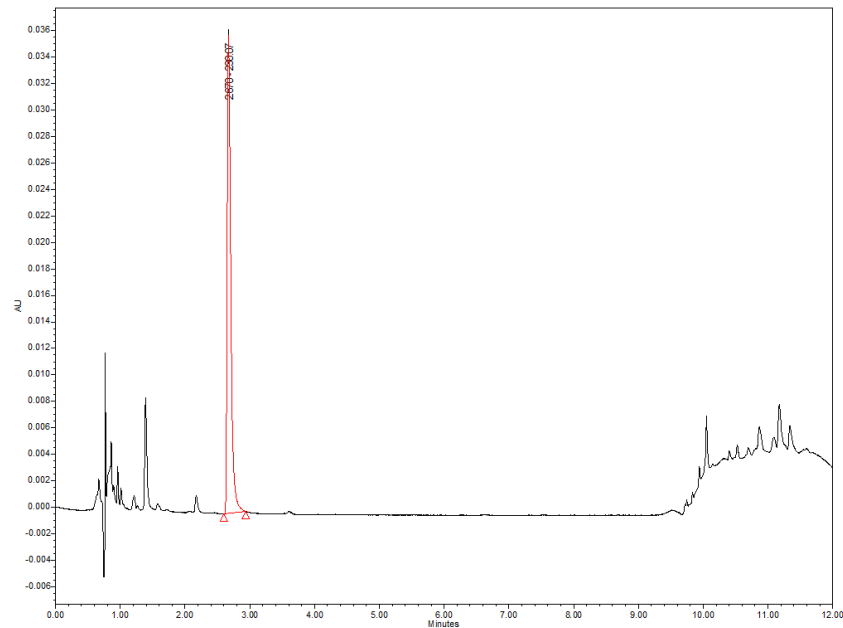


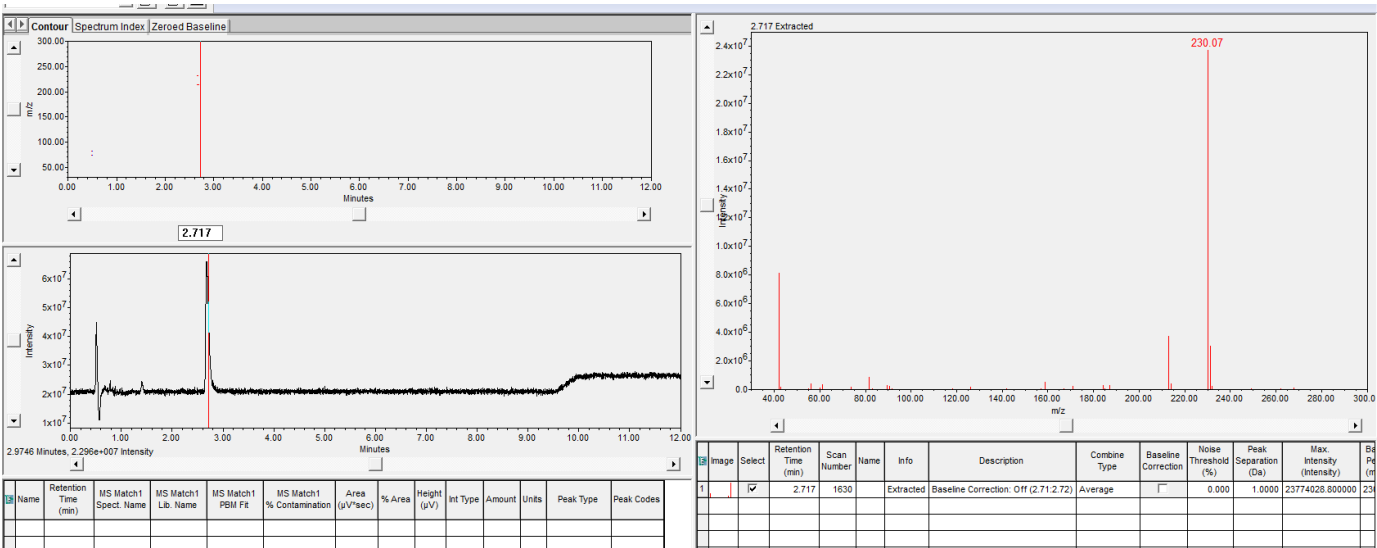
	Image	Select	Retention Time (min)	Scan Number	Name	Info	Description	Combine Type	Baseline Correction	Noise Threshold (%)	Peak Separation (Da)	Max. Intensity (Intensity)	Base Peak (m/z)
1		<input checked="" type="checkbox"/>	2.767	1660	Extracted	Baseline Correction: Off (2.762.77)	Average			0.000	1.0000	31056070.400000	230

3-1 low gamma LC PDA Detector Data with Integrated Peak

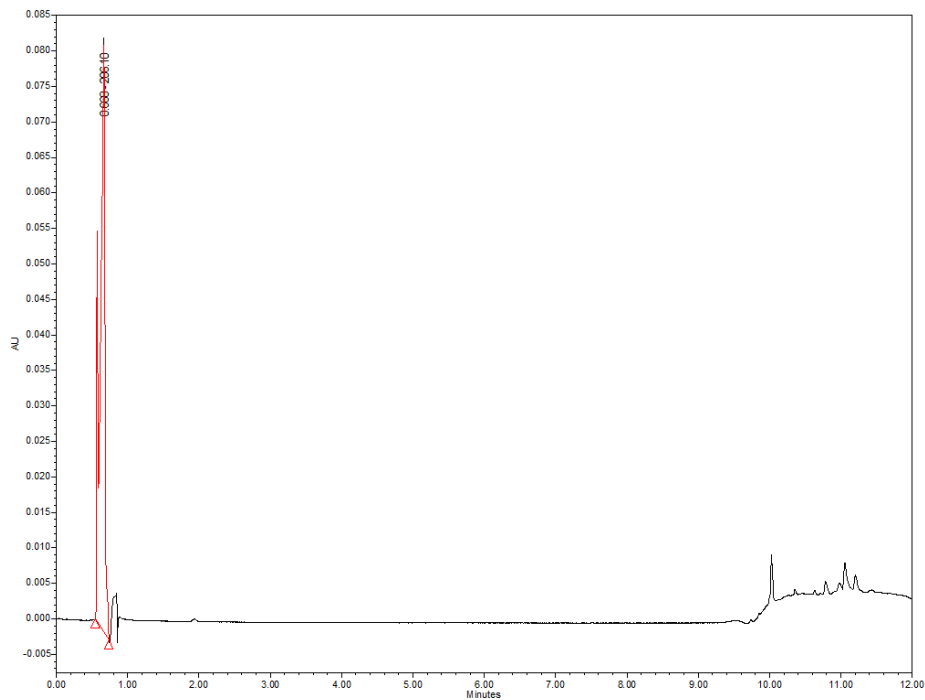


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.670	136579	100.00	36106	bb			Unknown

3-1 low gamma Mass Spectrum

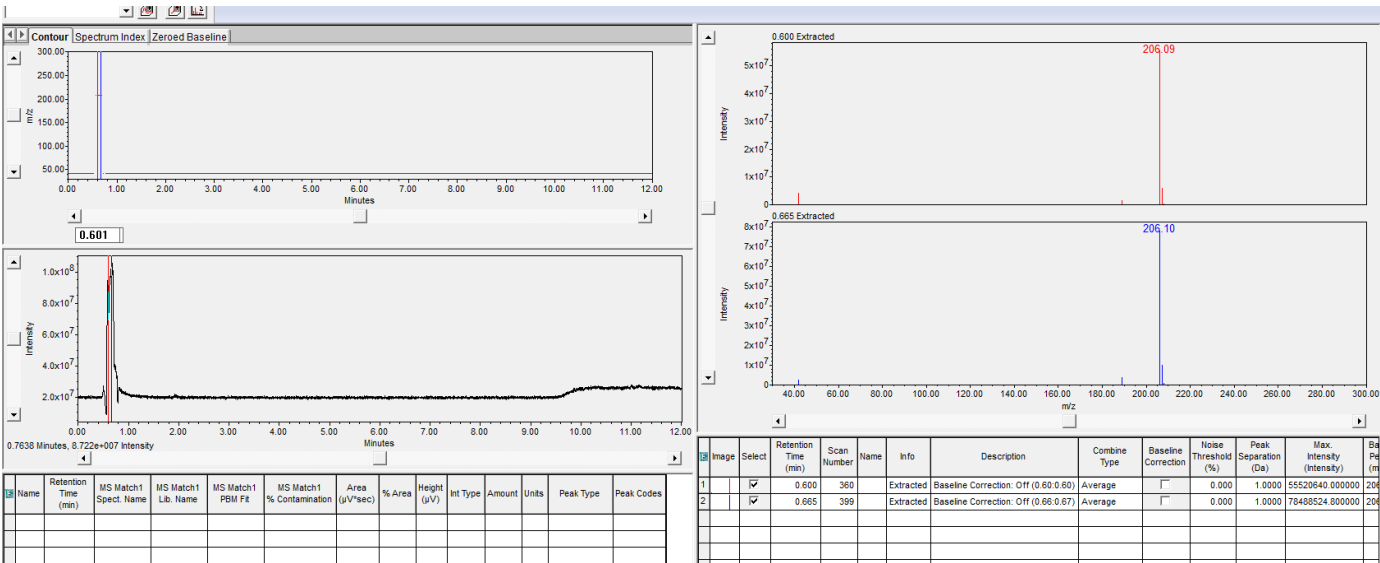


3-2 LC PDA Detector Data with Integrated Peak



	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.668	395712	100.00	82797	bb			Unknown

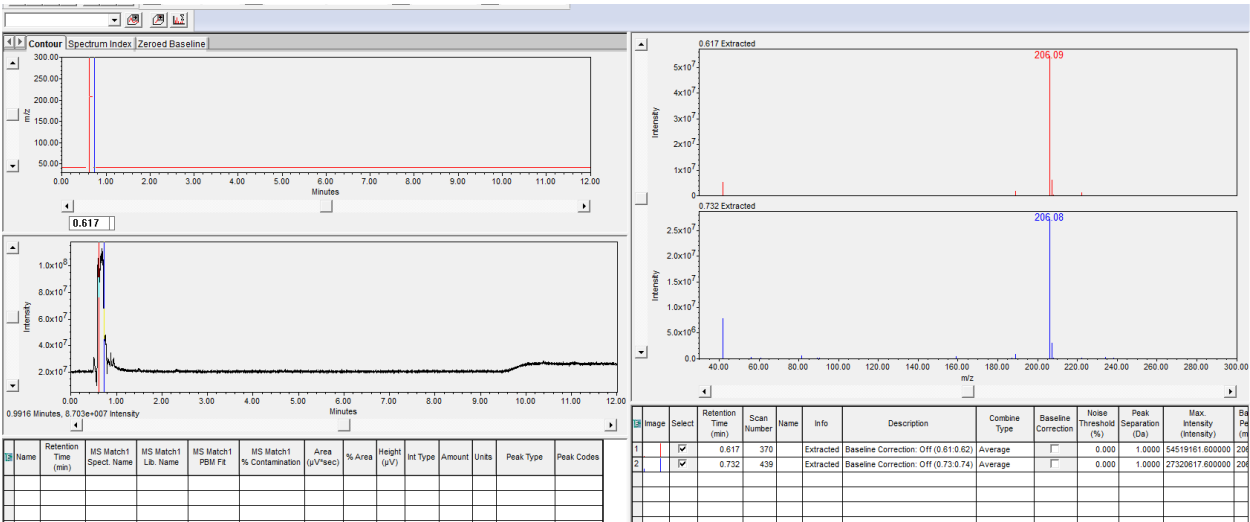
3-2 Mass Spectrum



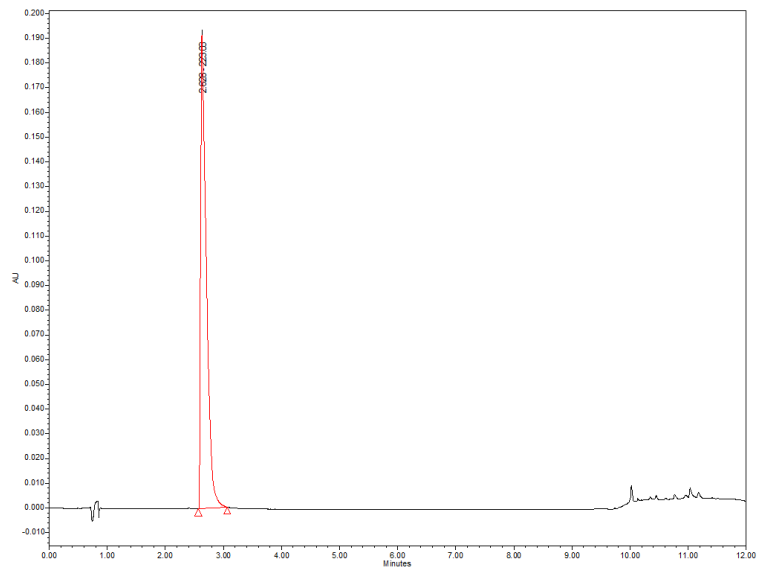
3-2 low gamma LC PDA Detector Data with Integrated Peak

	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.681	389689	100.00	79592	bb			Unknown

3-2 low gamma Mass Spectrum

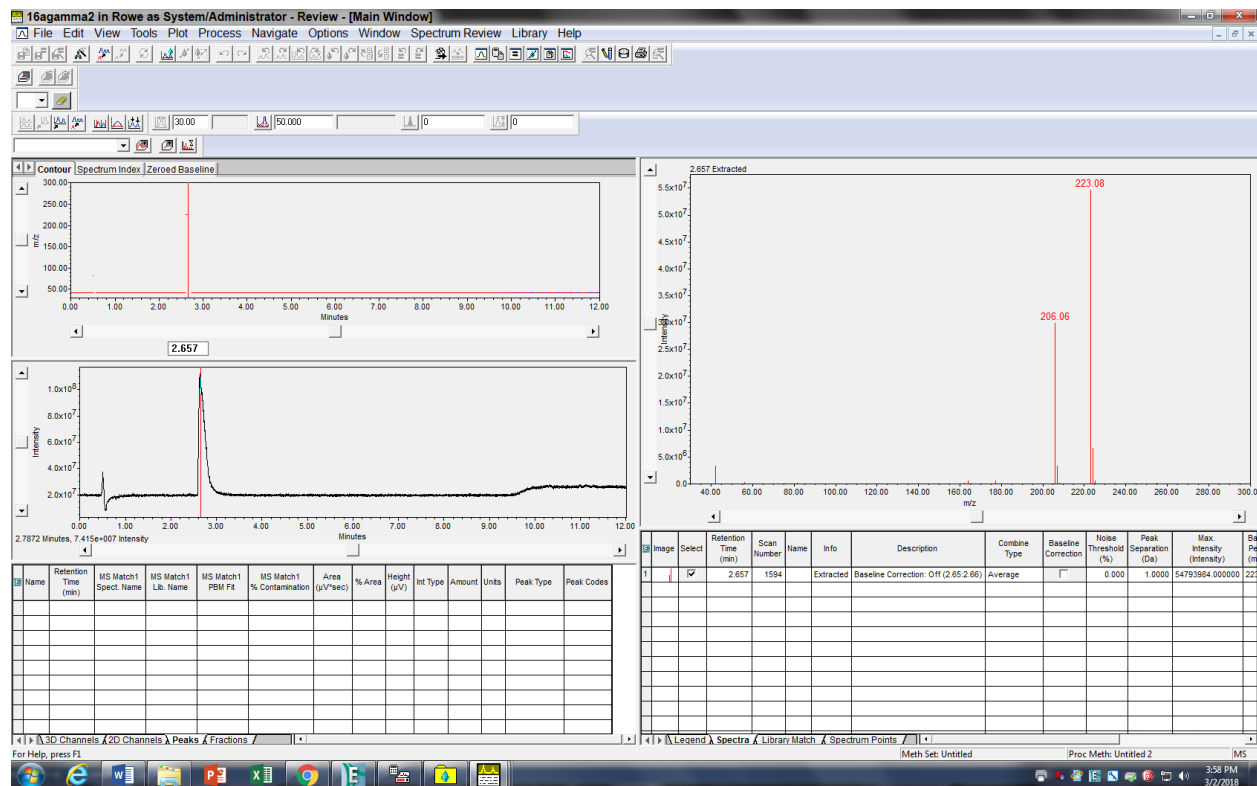


### 3-3 LC PDA Detector Data with Integrated Peak

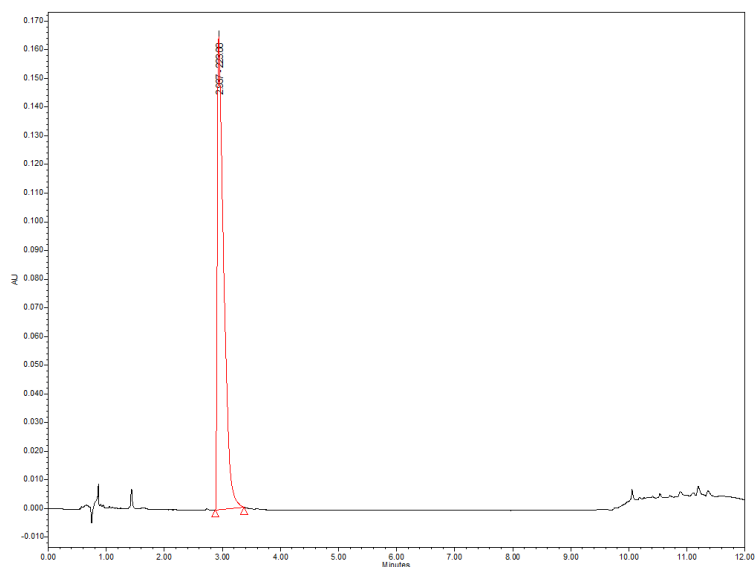


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.628	1325332	100.00	191636	bb			Unknown

### 3-3 Mass Spectrum

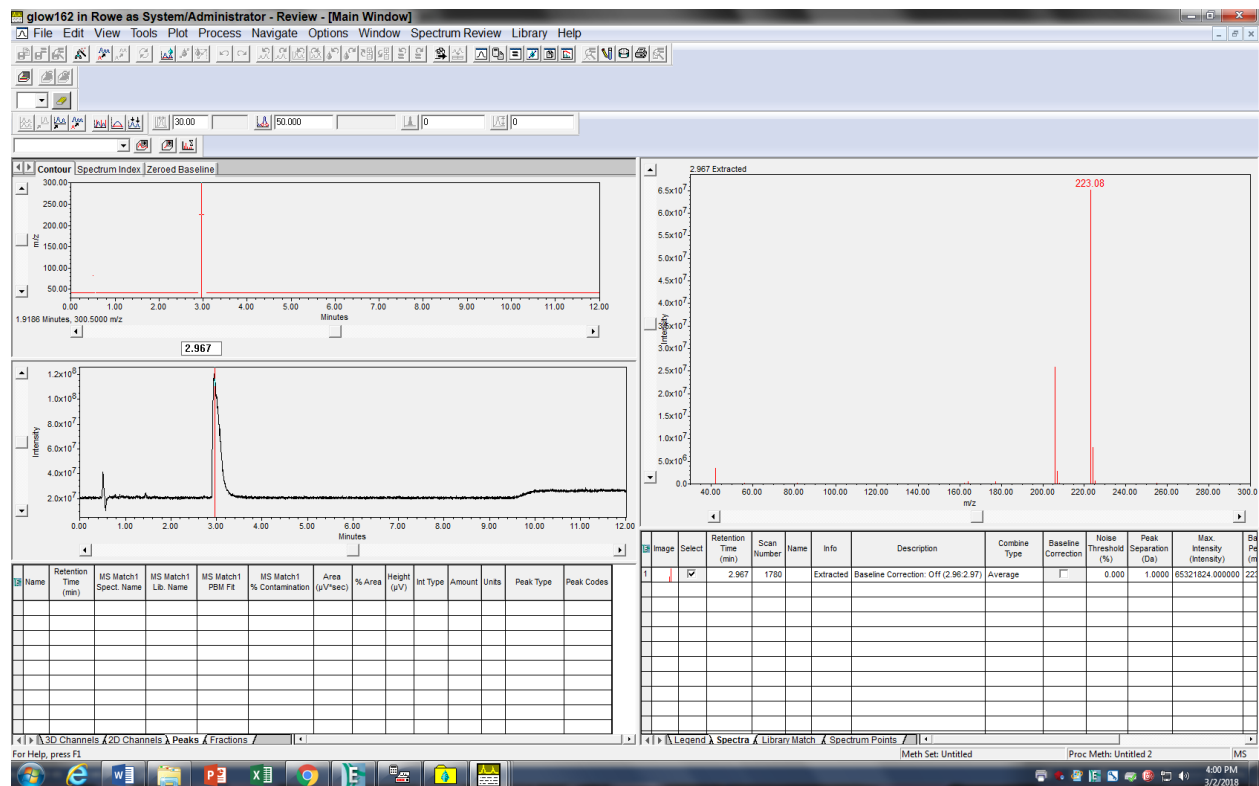


### 3-3 low gamma LC PDA Detector Data with Integrated Peak

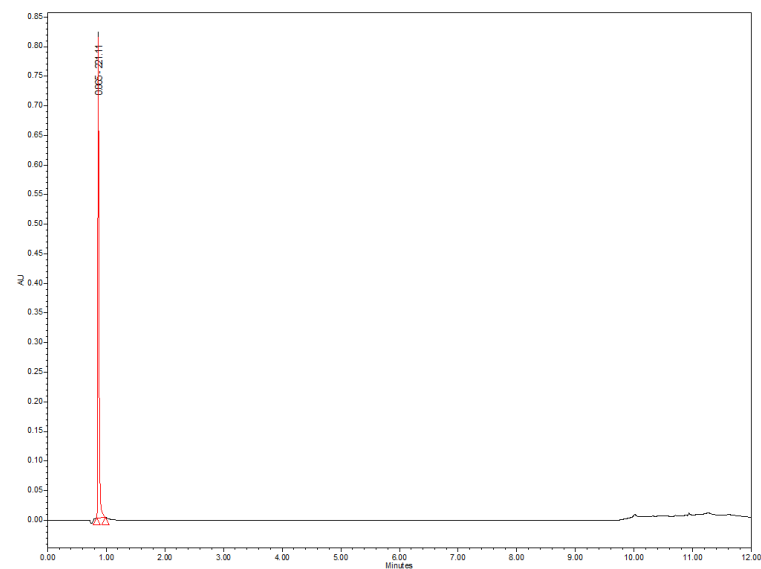


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.937	1273530	100.00	164956	bb			Unknown

### 3-3 low gamma Mass Spectrum

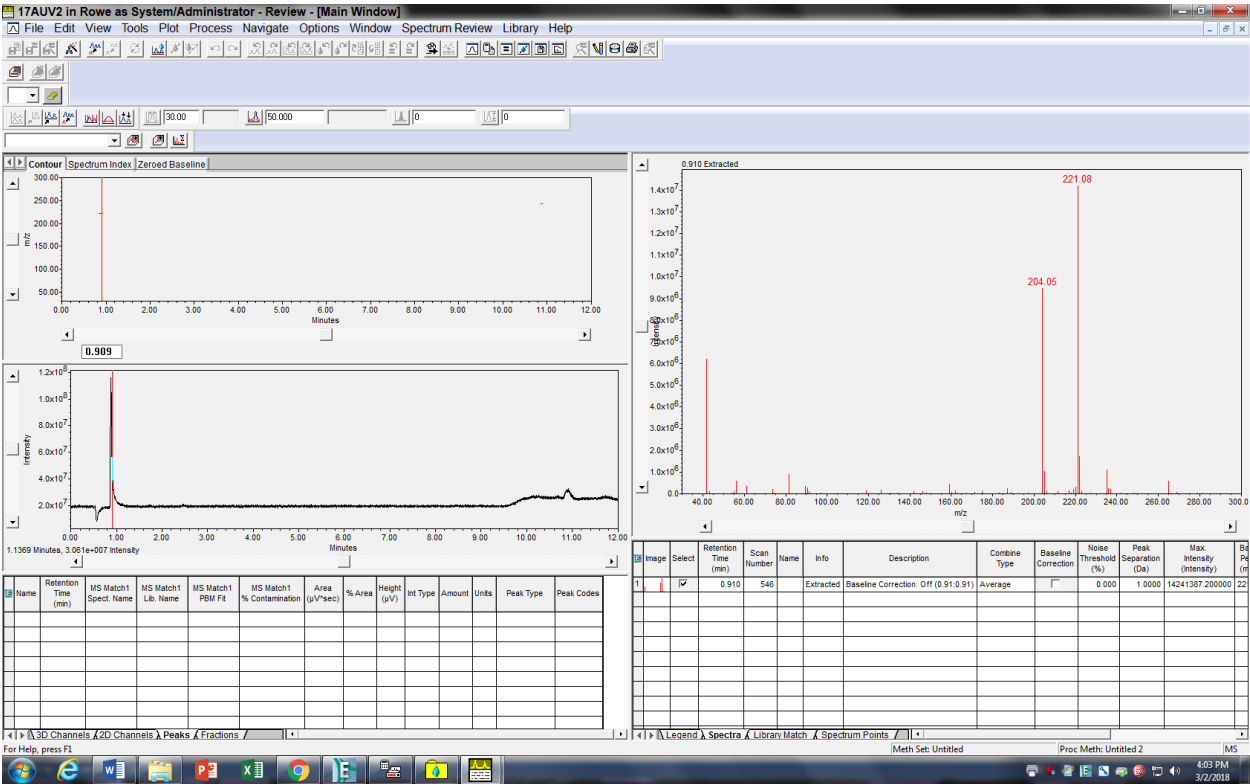


3-4 LC PDA Detector Data with Integrated Peak



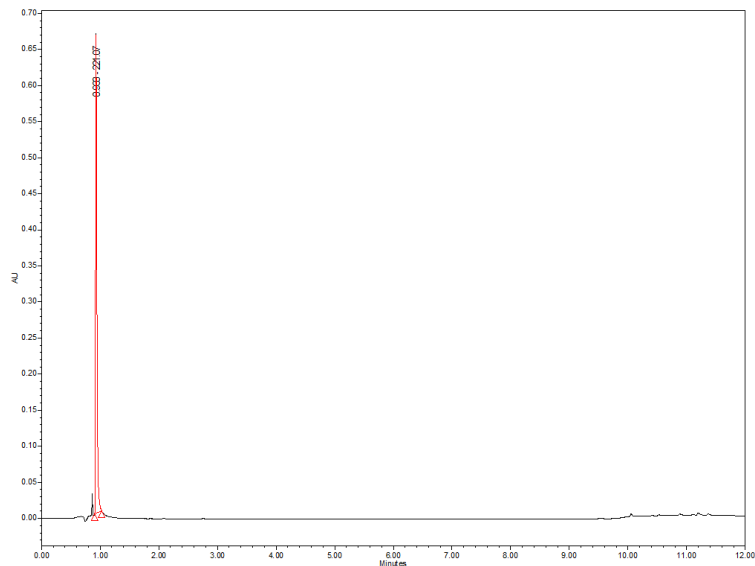
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.865	1082123	100.00	812293	bb			Unknown

3-4 Mass Spectrum



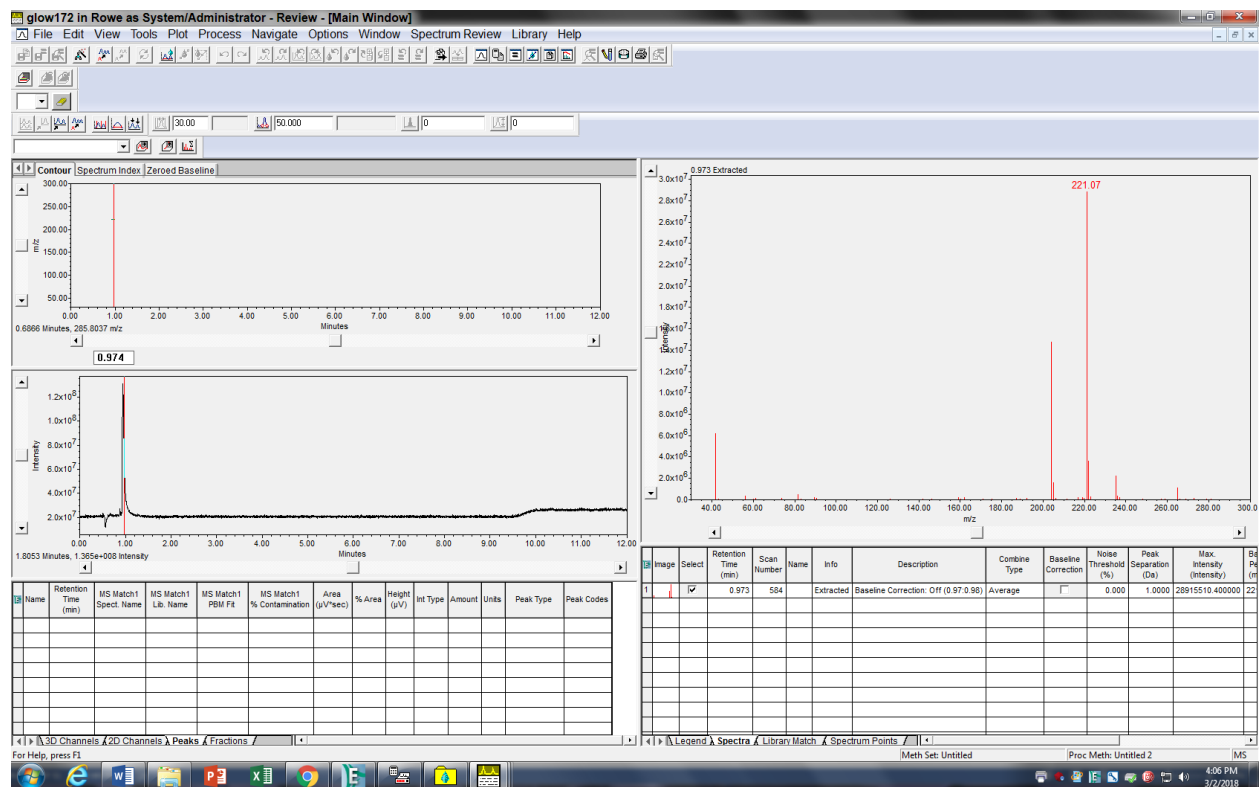


### 3-4 low gamma LC PDA Detector Data with Integrated Peak

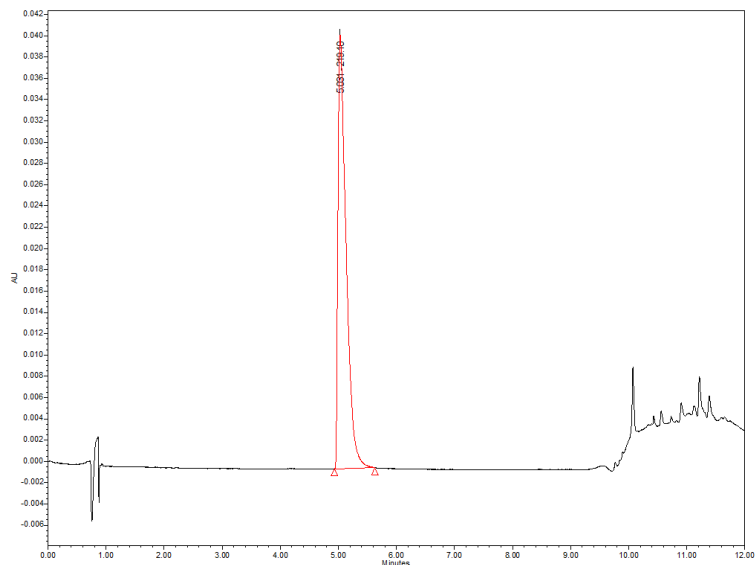


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		0.933	1002079	100.00	662981	bb			Unknown

### 3-4 low gamma Mass Spectrum

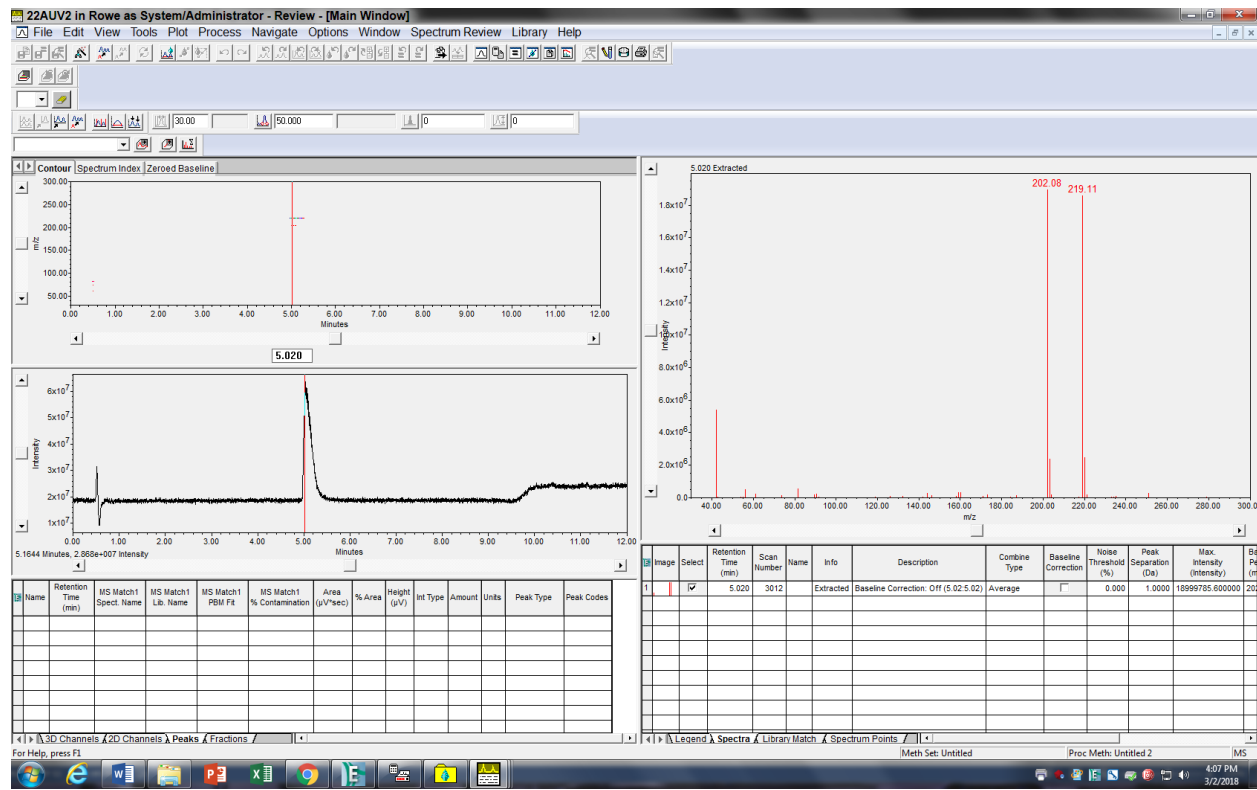


### 3-5 LC PDA Detector Data with Integrated Peak

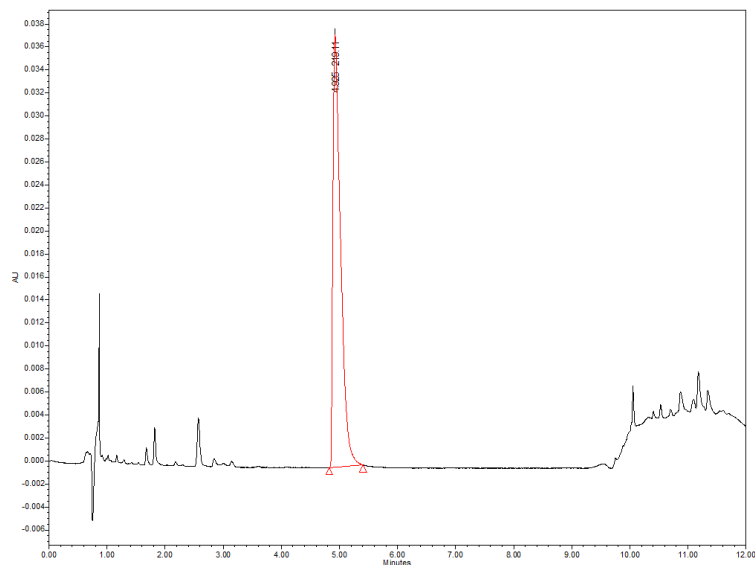


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		5.031	396392	100.00	40761	bb			Unknown

### 3-5 Mass Spectrum

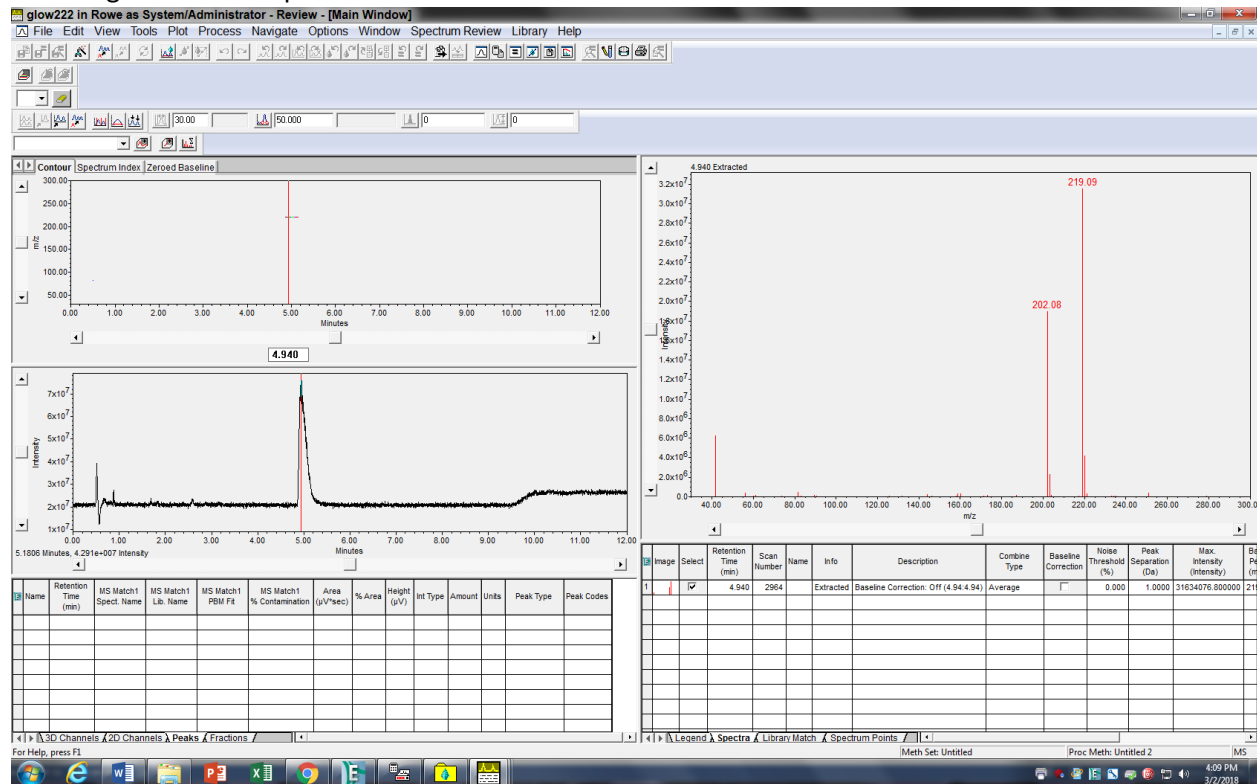


### 3-5 low gamma LC PDA Detector Data with Integrated Peak

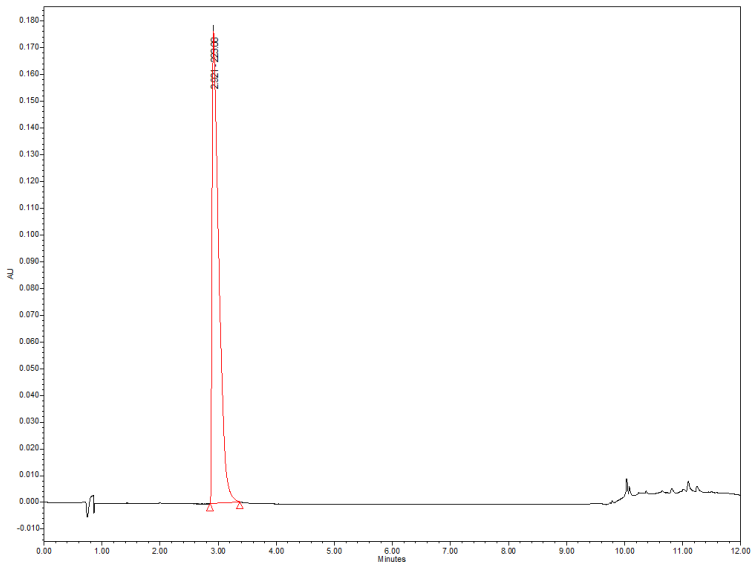


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		4.925	343380	100.00	37621	bb			Unknown

### 3-5 low gamma Mass Spectrum

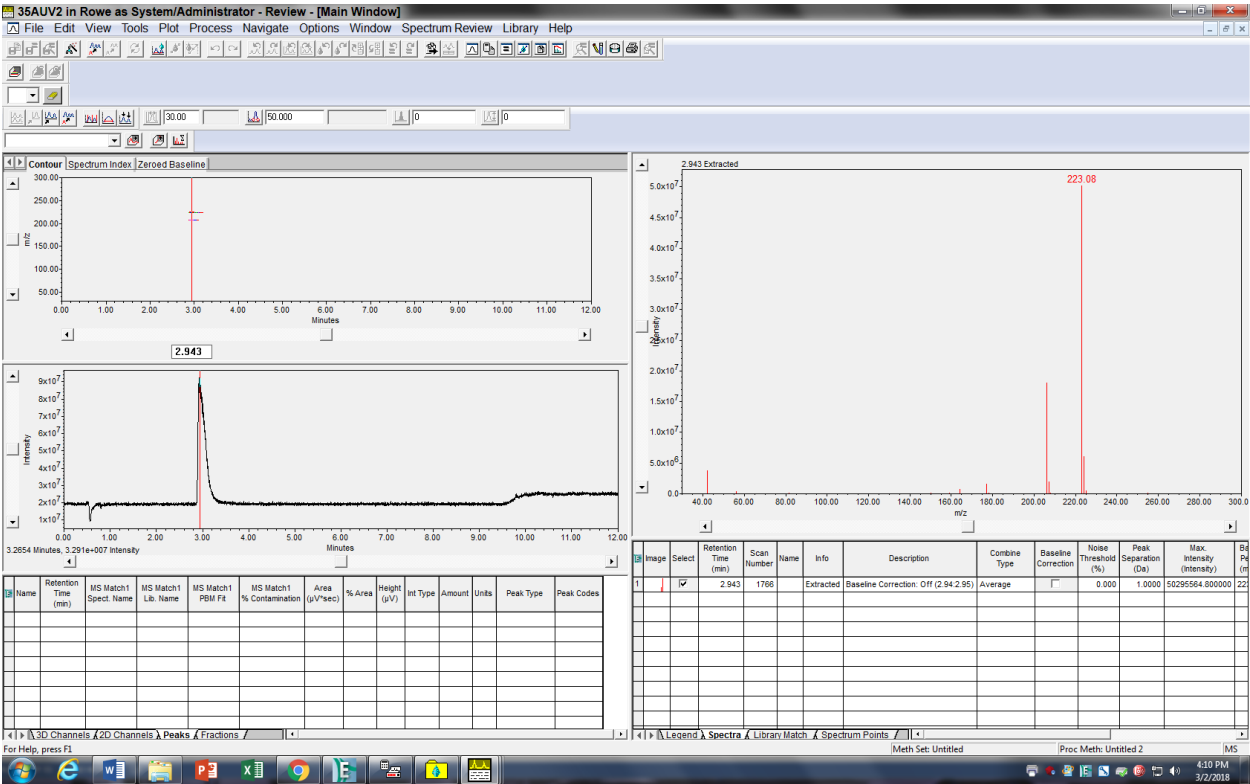


3-6 LC PDA Detector Data with Integrated Peak

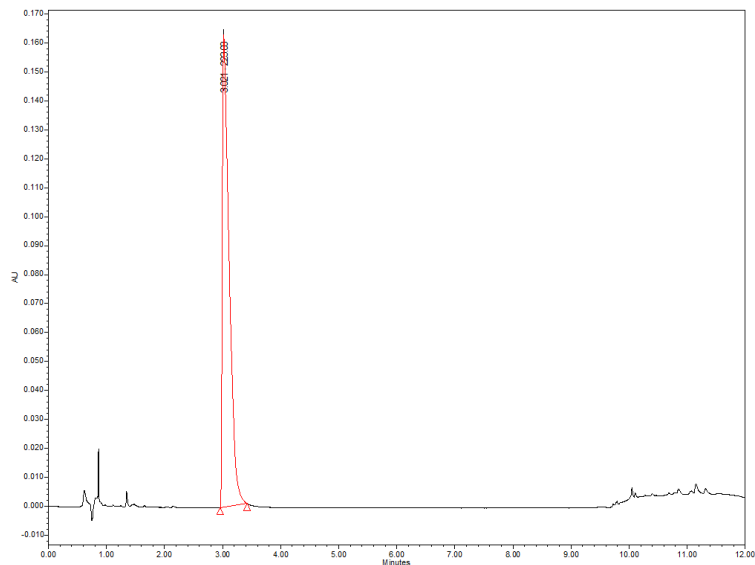


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		2.921	1435309	100.00	176766	bb			Unknown

3-6 Mass Spectrum

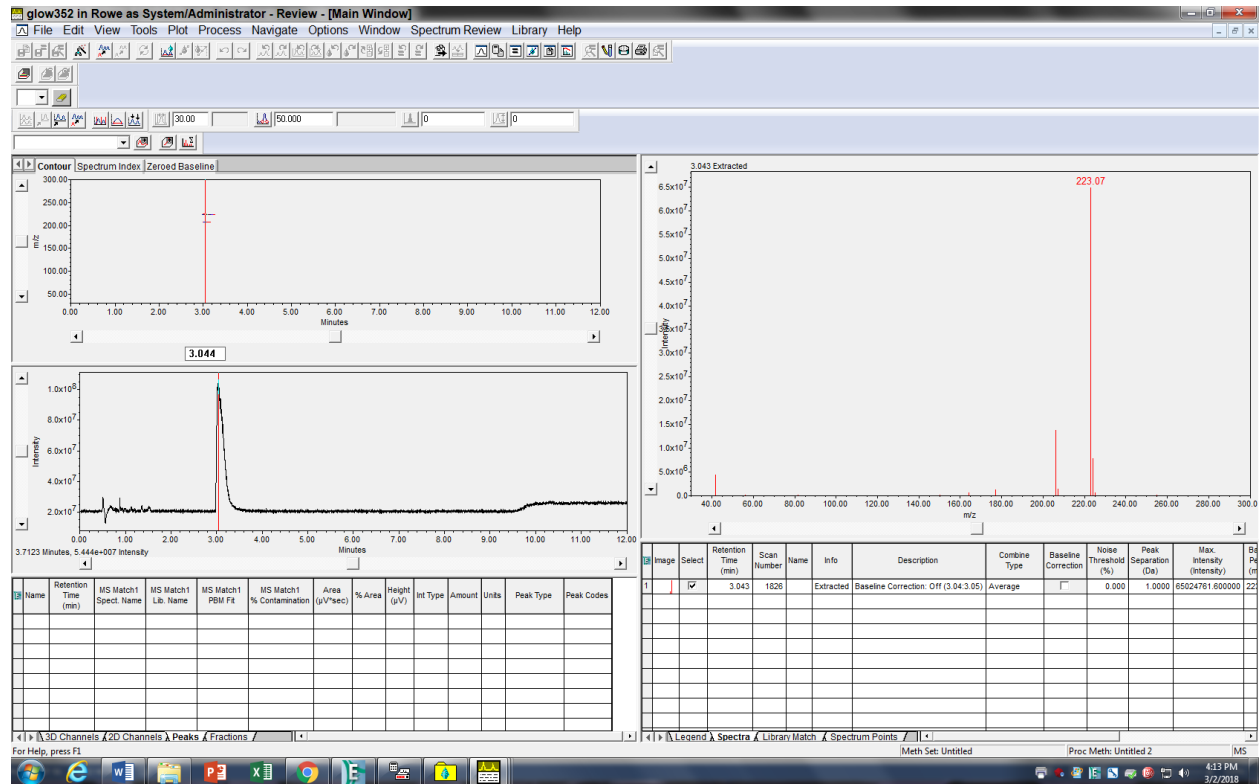


### 3-6 low gamma LC PDA Detector Data with Integrated Peak

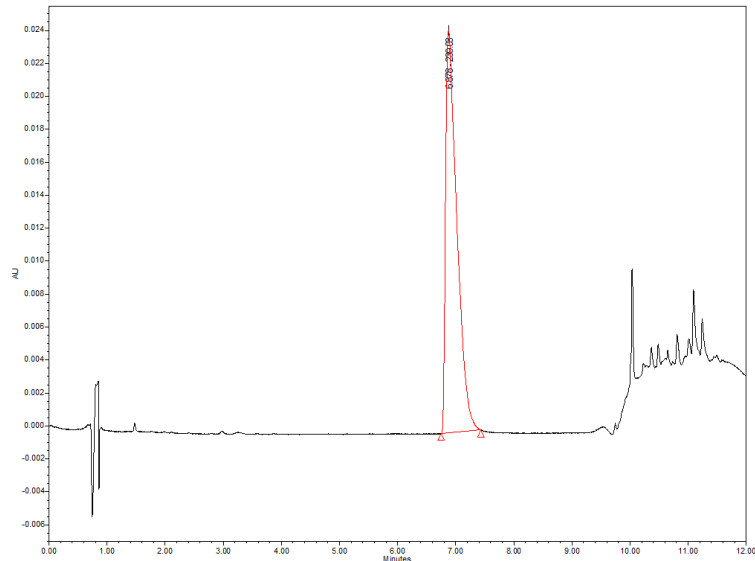


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		3.021	1327610	100.00	163161	bb			Unknown

### 3-6 low gamma Mass Spectrum

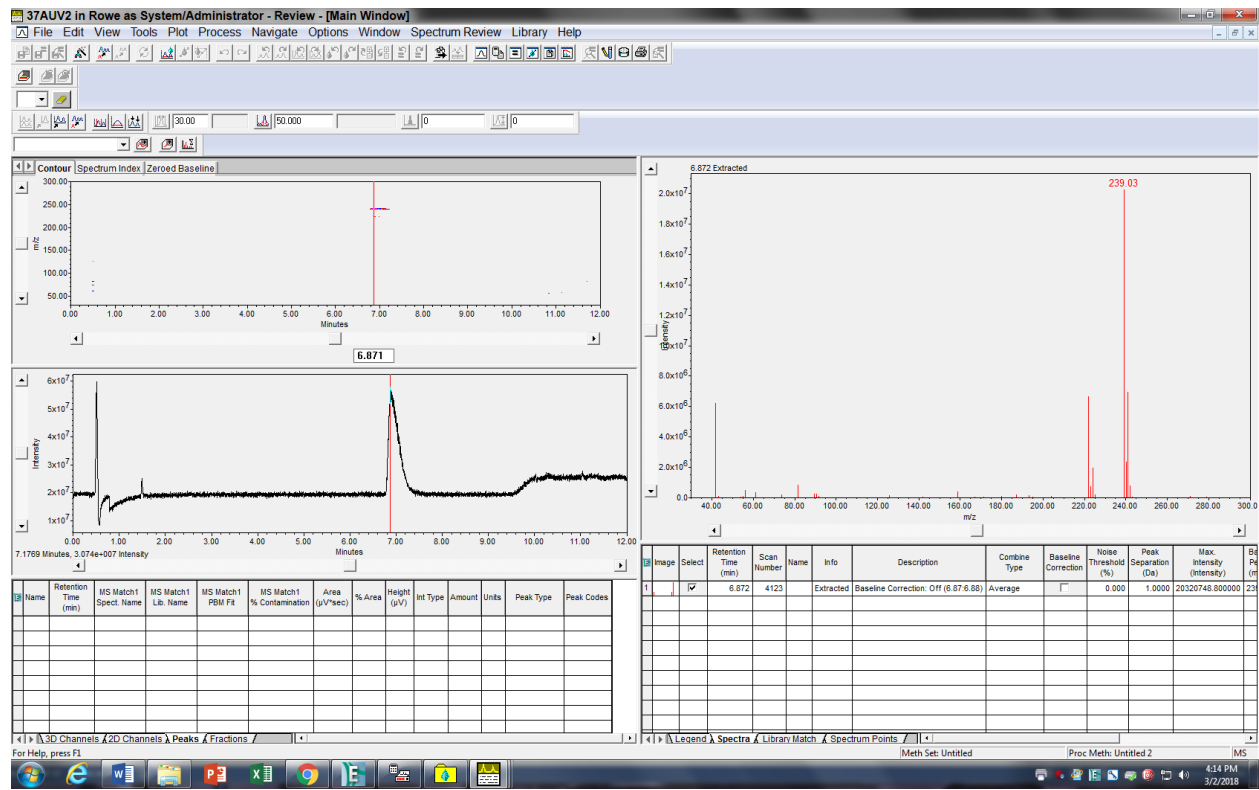


### 3-7 LC PDA Detector Data with Integrated Peak

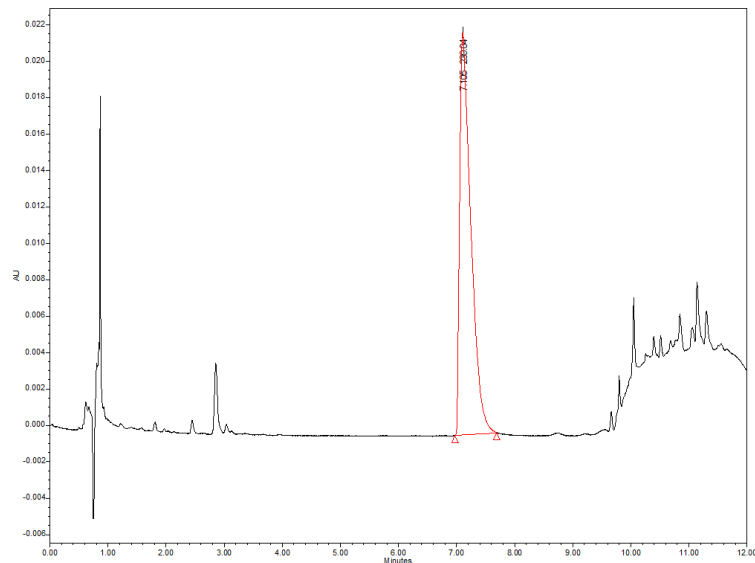


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		6.878	327011	100.00	24411	bb			Unknown

### 3-7 Mass Spectrum

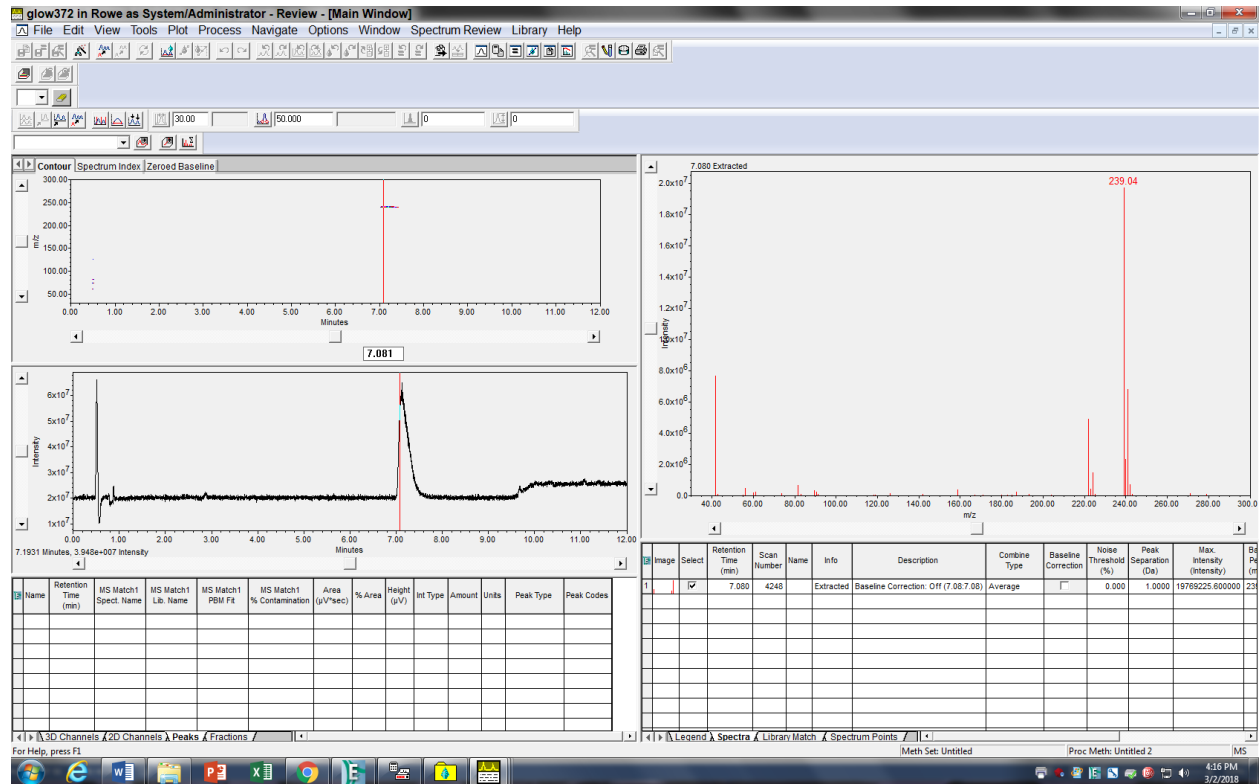


### 3-7 low gamma LC PDA Detector Data with Integrated Peak

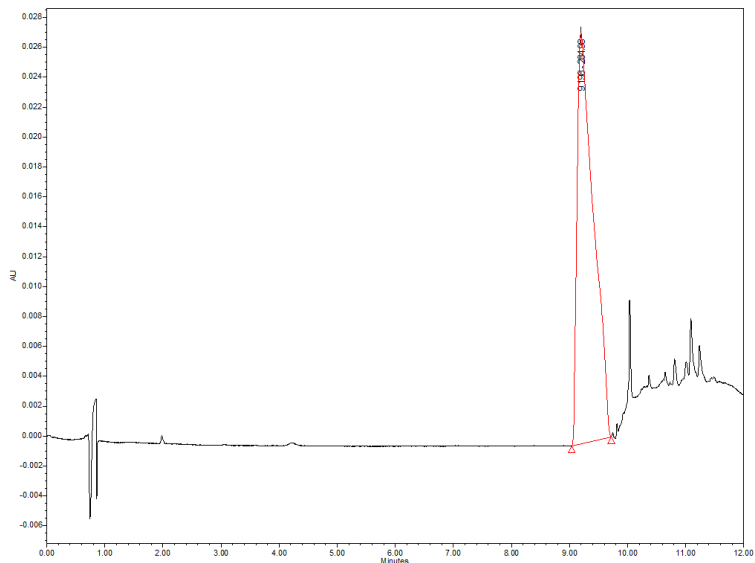


	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		7.105	294309	100.00	22091	bb			Unknown

### 3-7 low gamma Mass Spectrum

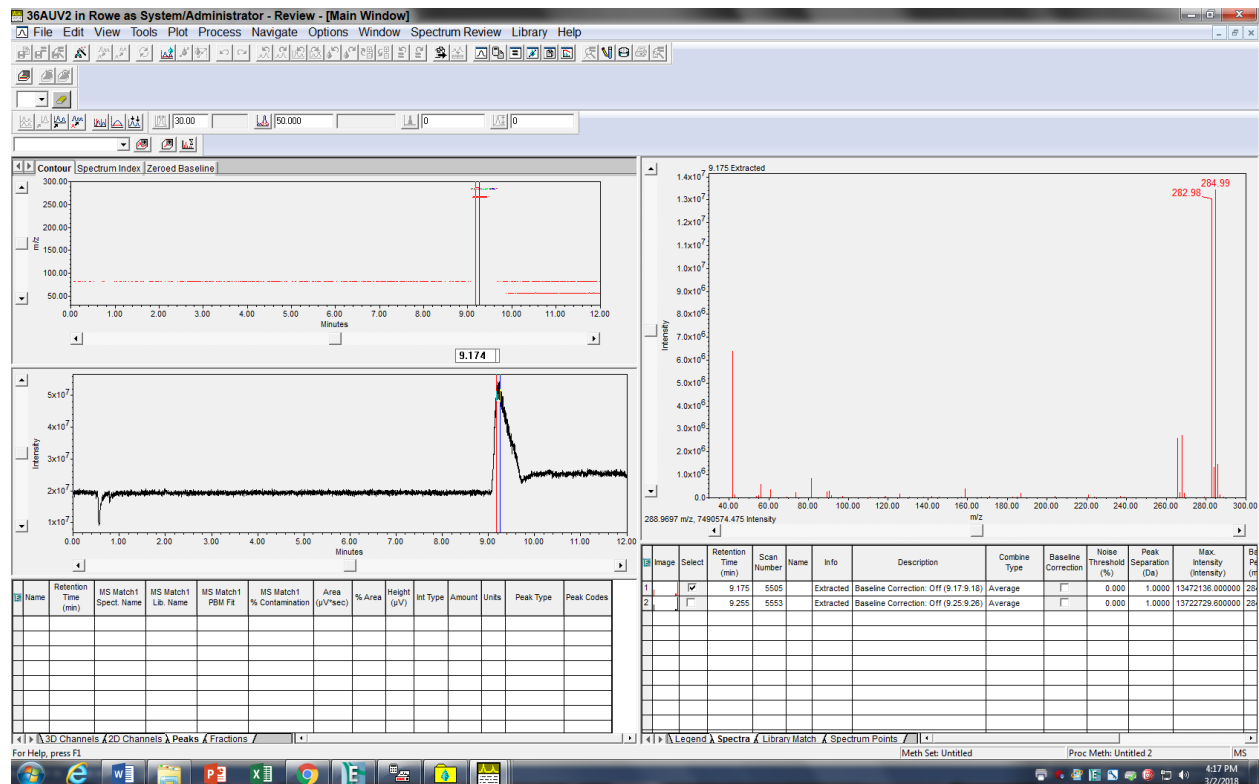


### 3-8 LC PDA Detector Data with Integrated Peak



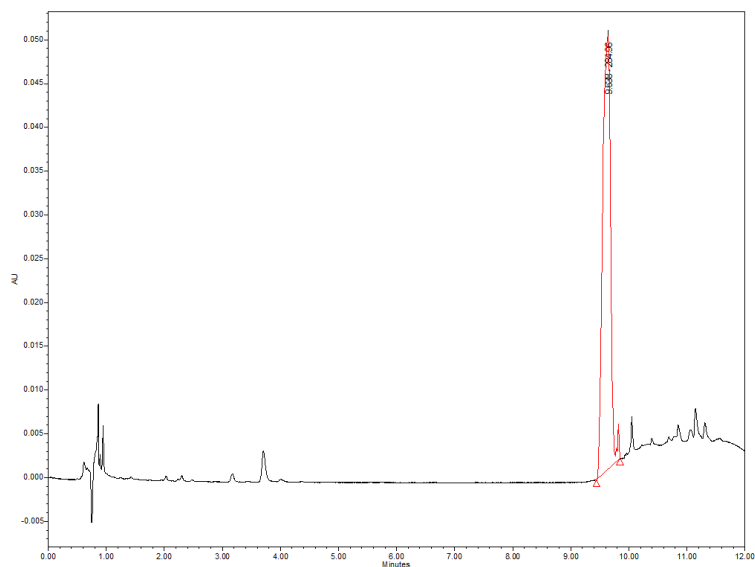
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.198	538110	100.00	27486	bb			Unknown

### 3-8 Mass Spectrum



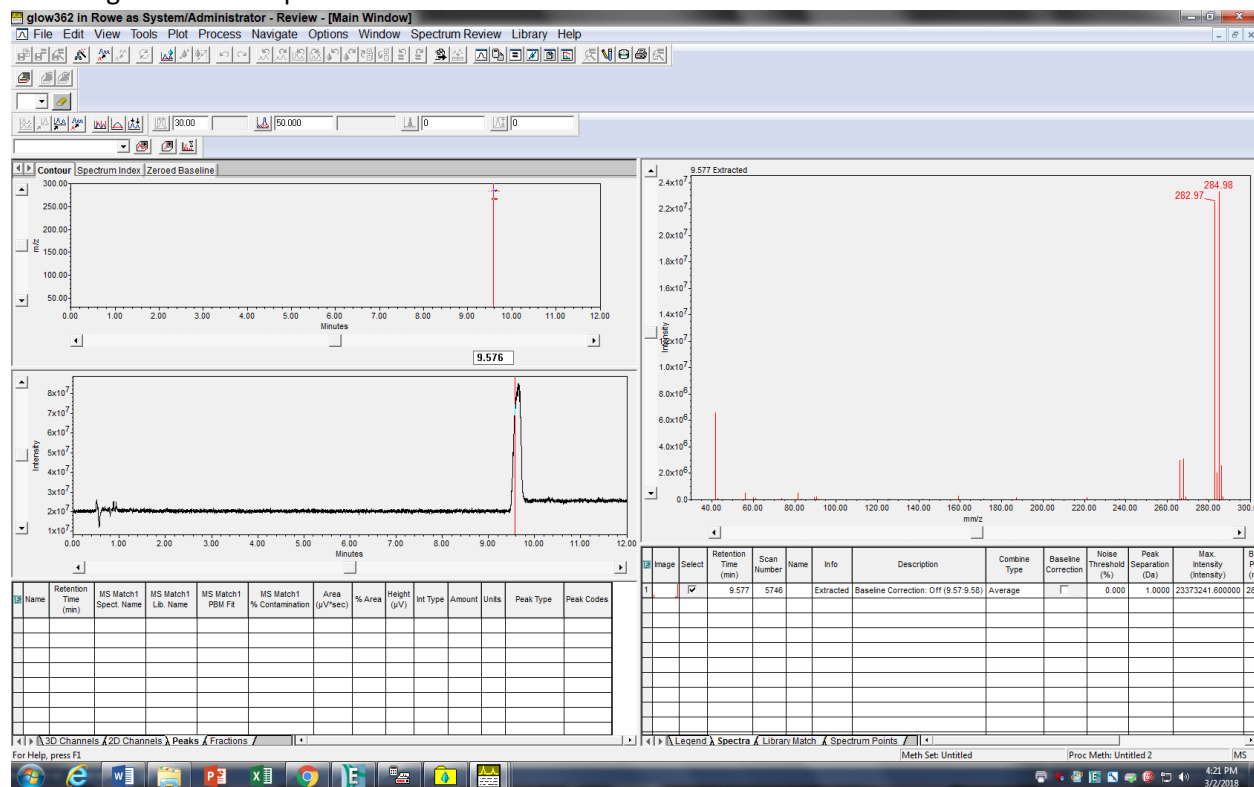


### 3-8 low gamma LC PDA Detector Data with Integrated Peak



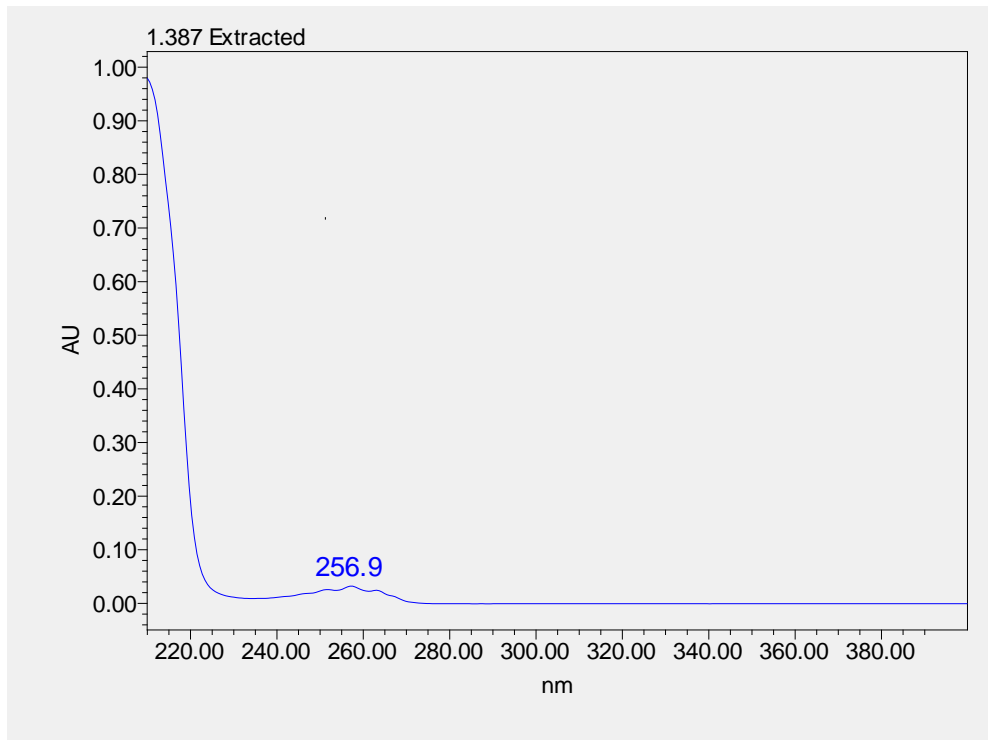
	Name	Retention Time	Area	% Area	Height	Int Type	Amount	Units	Peak Type
1		9.638	472479	100.00	49496	bb			Unknown

### 3-8 low gamma Mass Spectrum

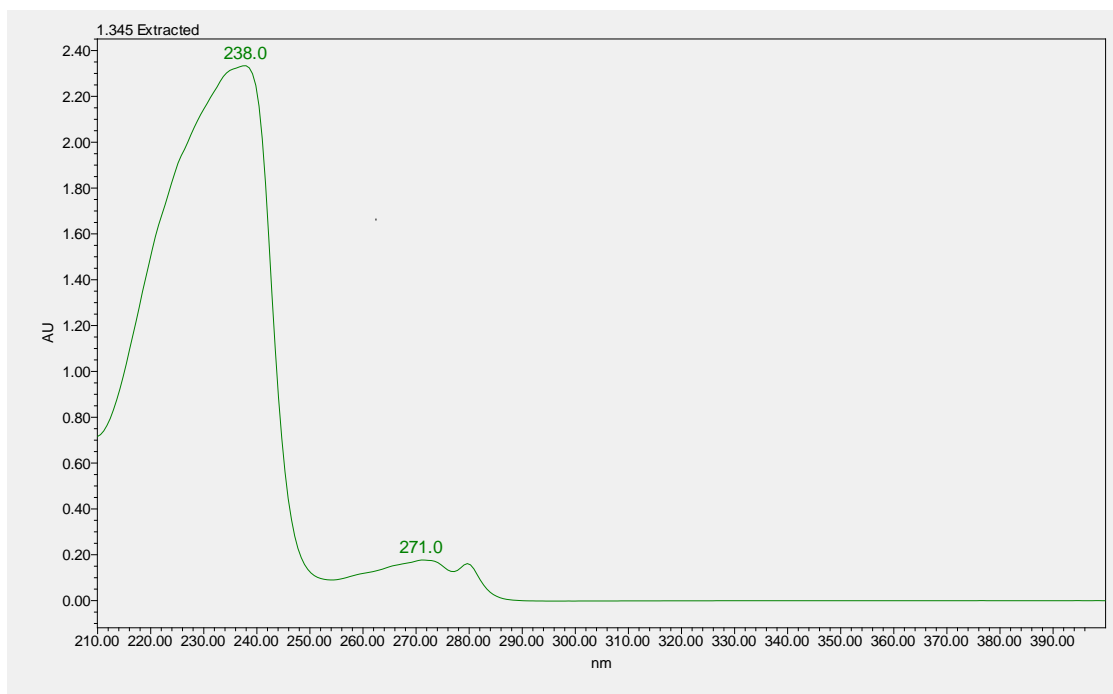


**Absorbance Spectrum of Amino Acids** Absorbance Spectrum of amino acids, resolution = 0.6 nm, range = 209.9nm - 399.9 nm

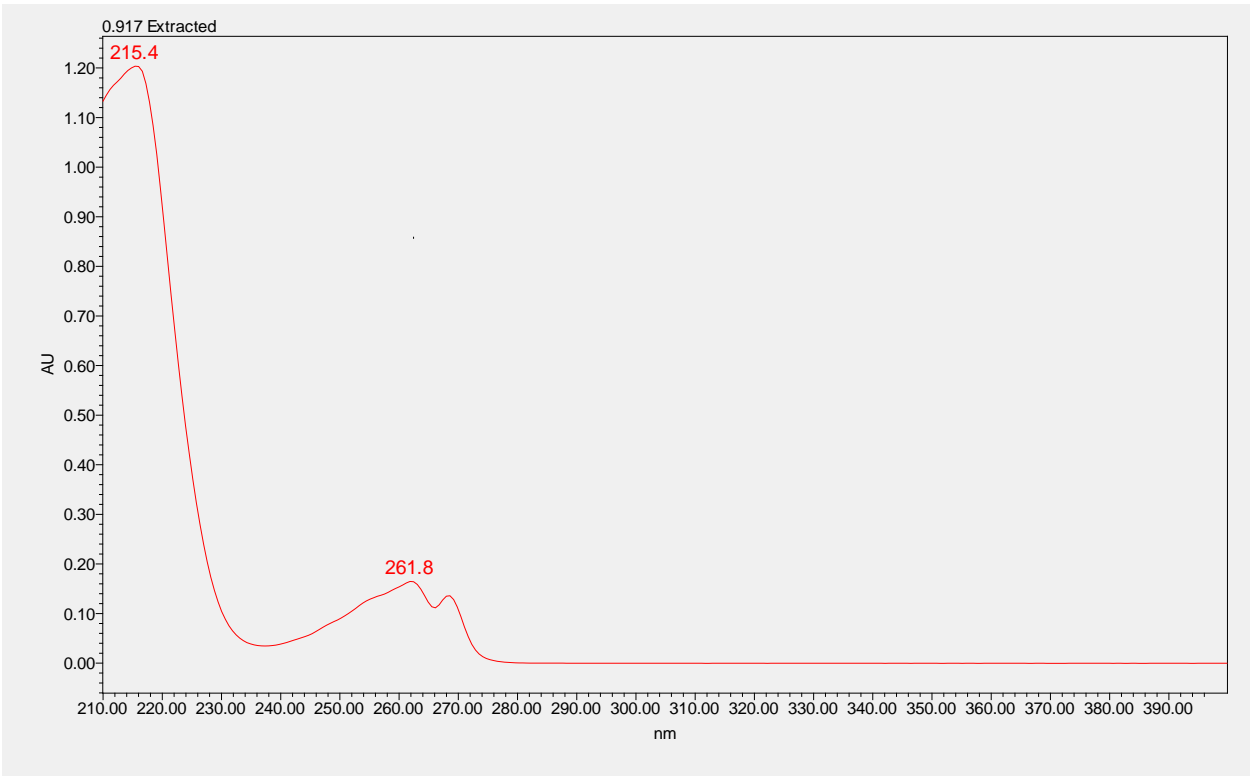
Amino Acid #1



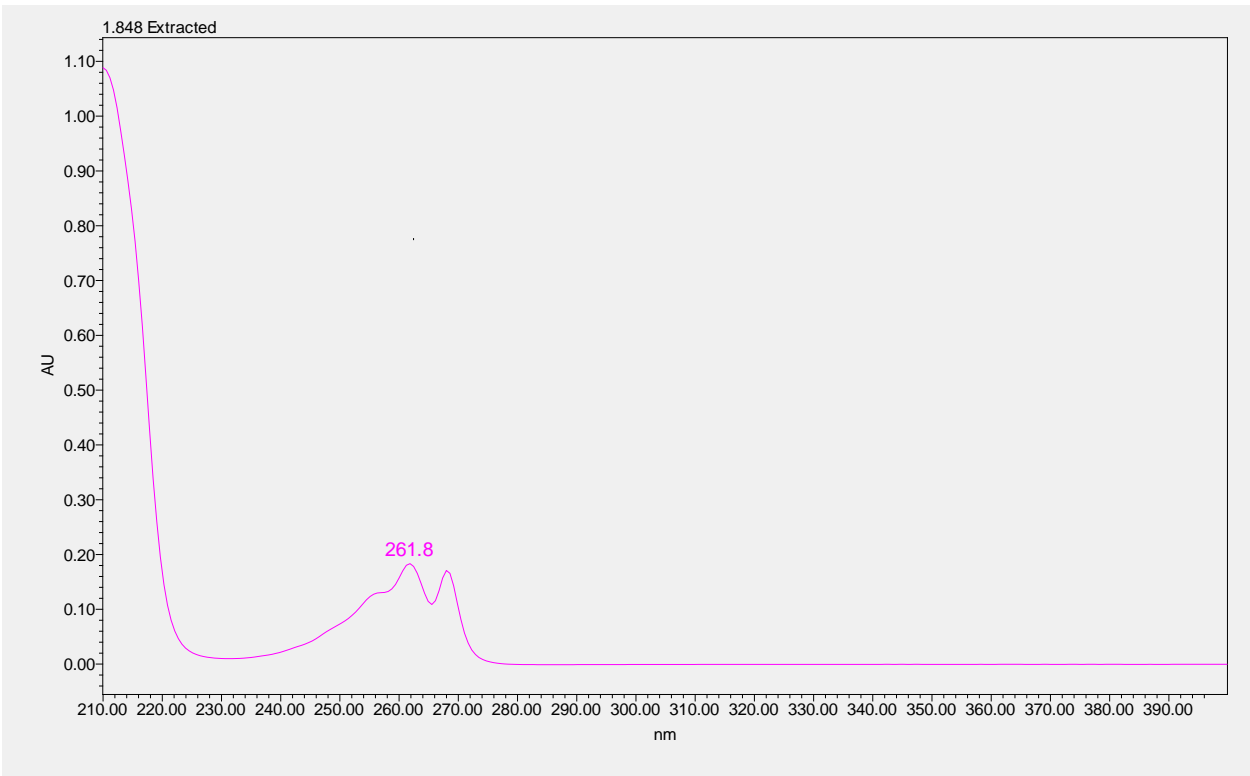
Amino Acid #1-1



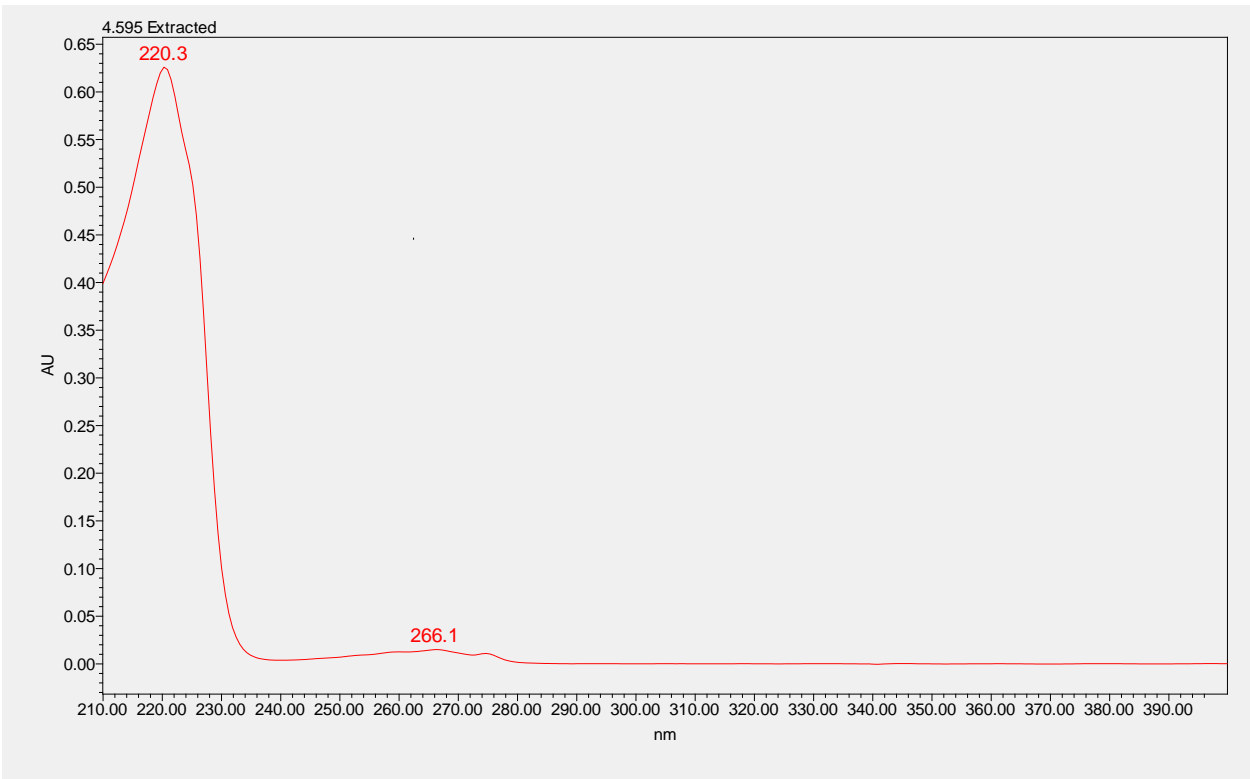
Amino Acid #1-3



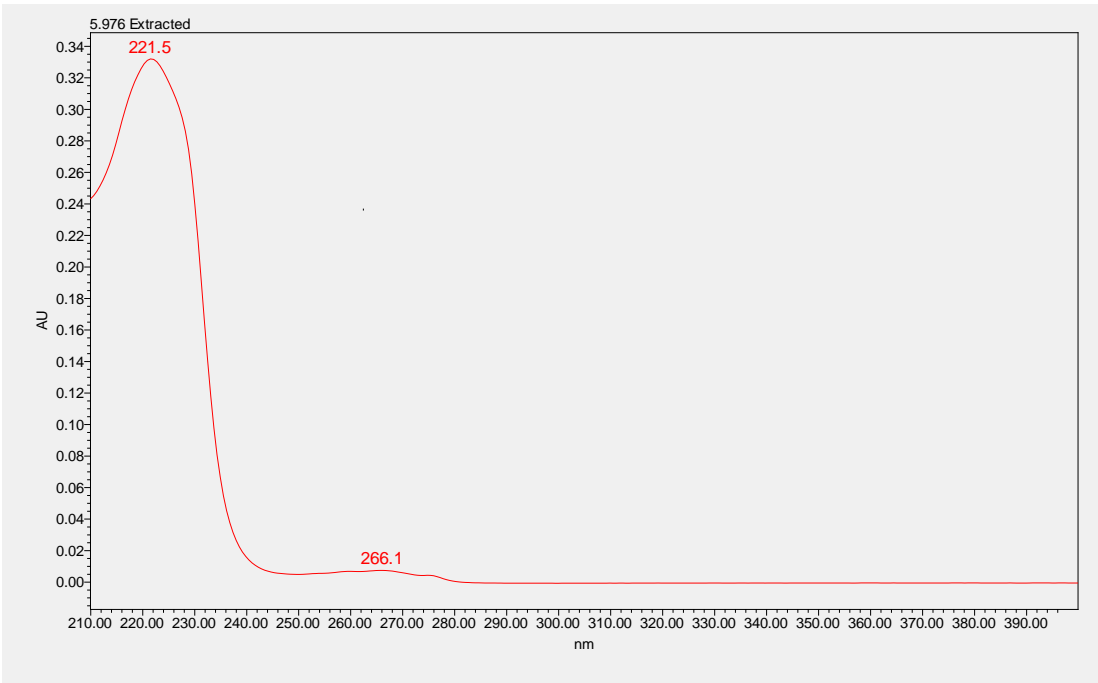
Amino Acid #1-6



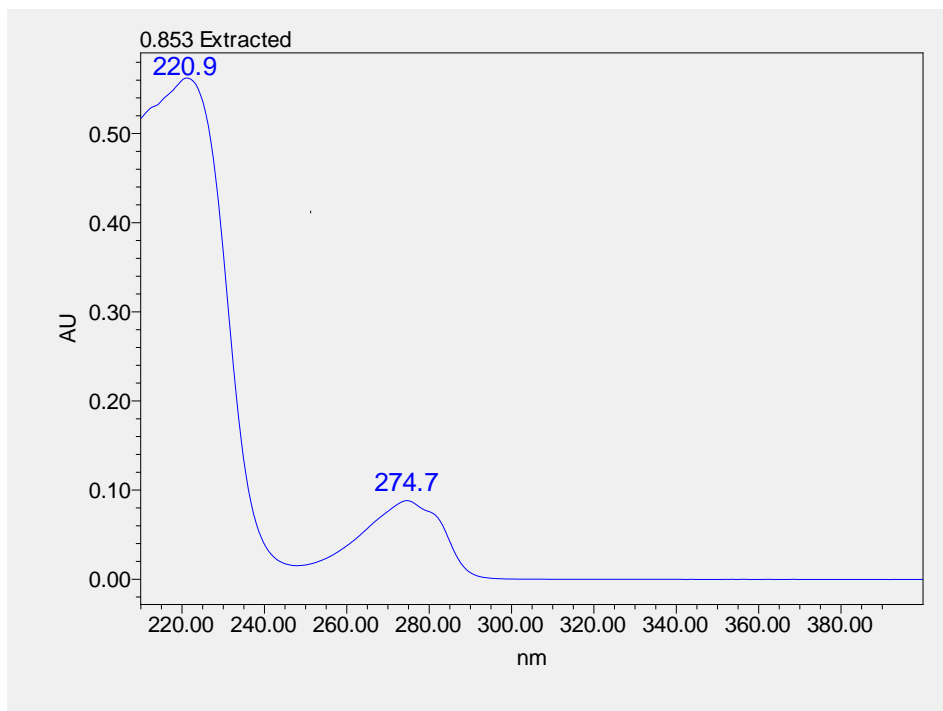
Amino Acid #1-7



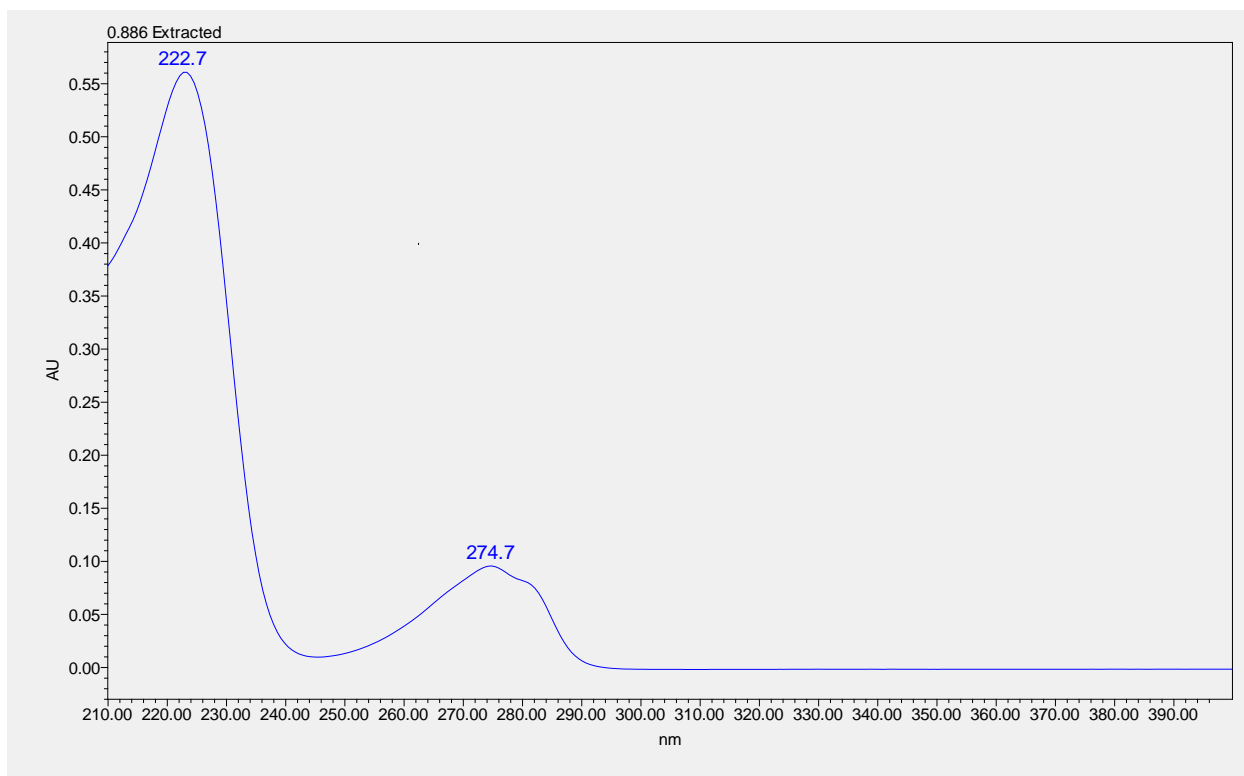
Amino Acid 1-8



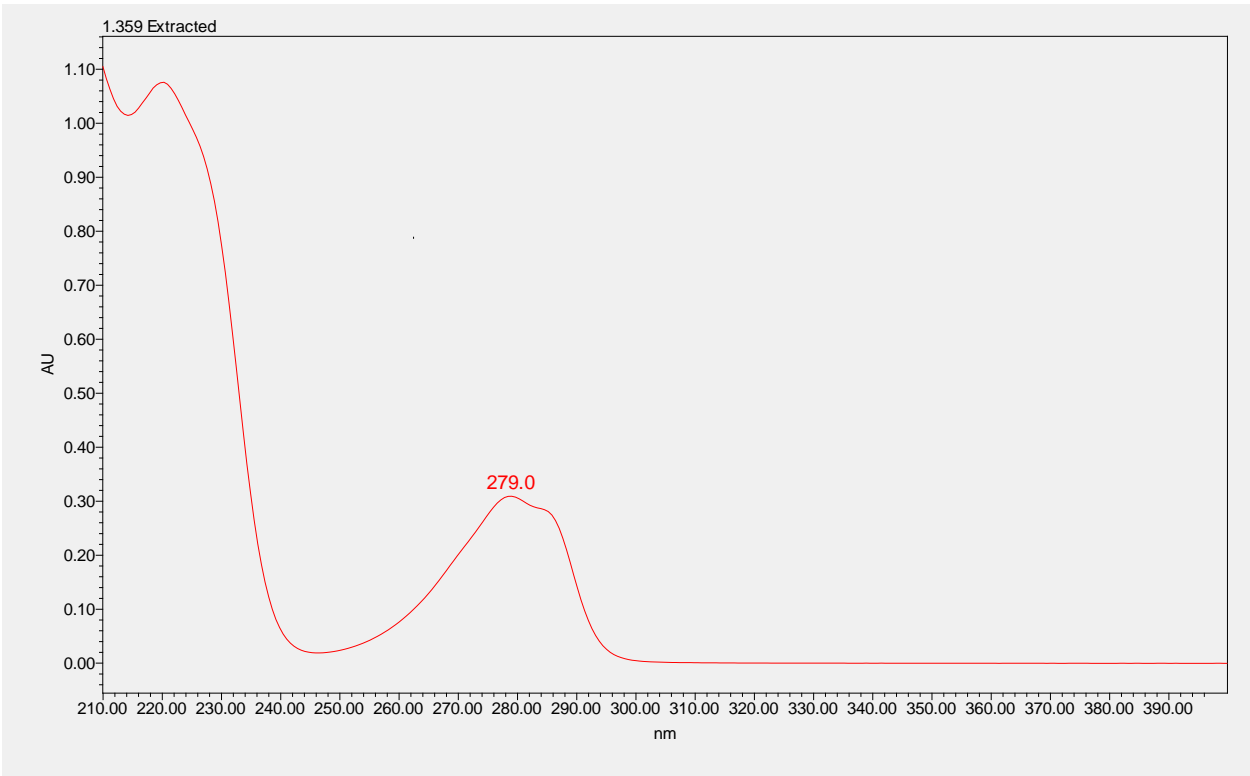
## Amino Acid 2



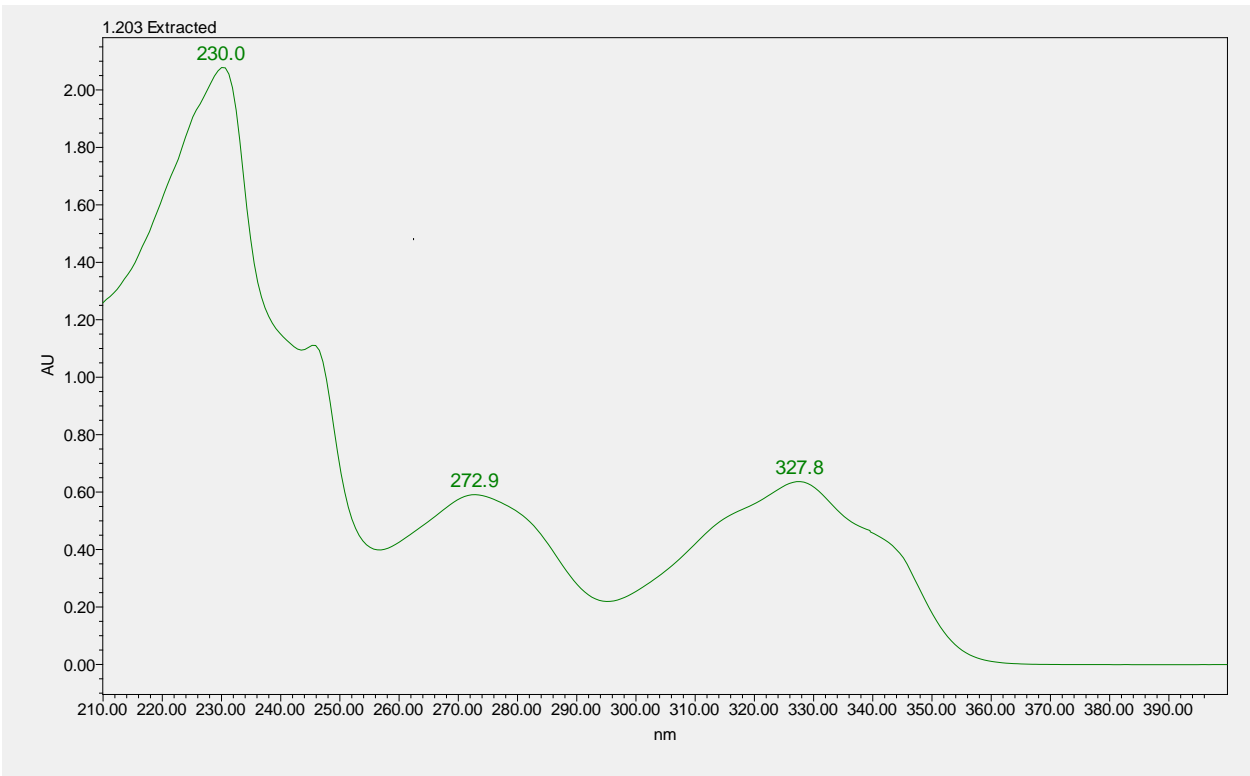
## Amino Acid 2-1



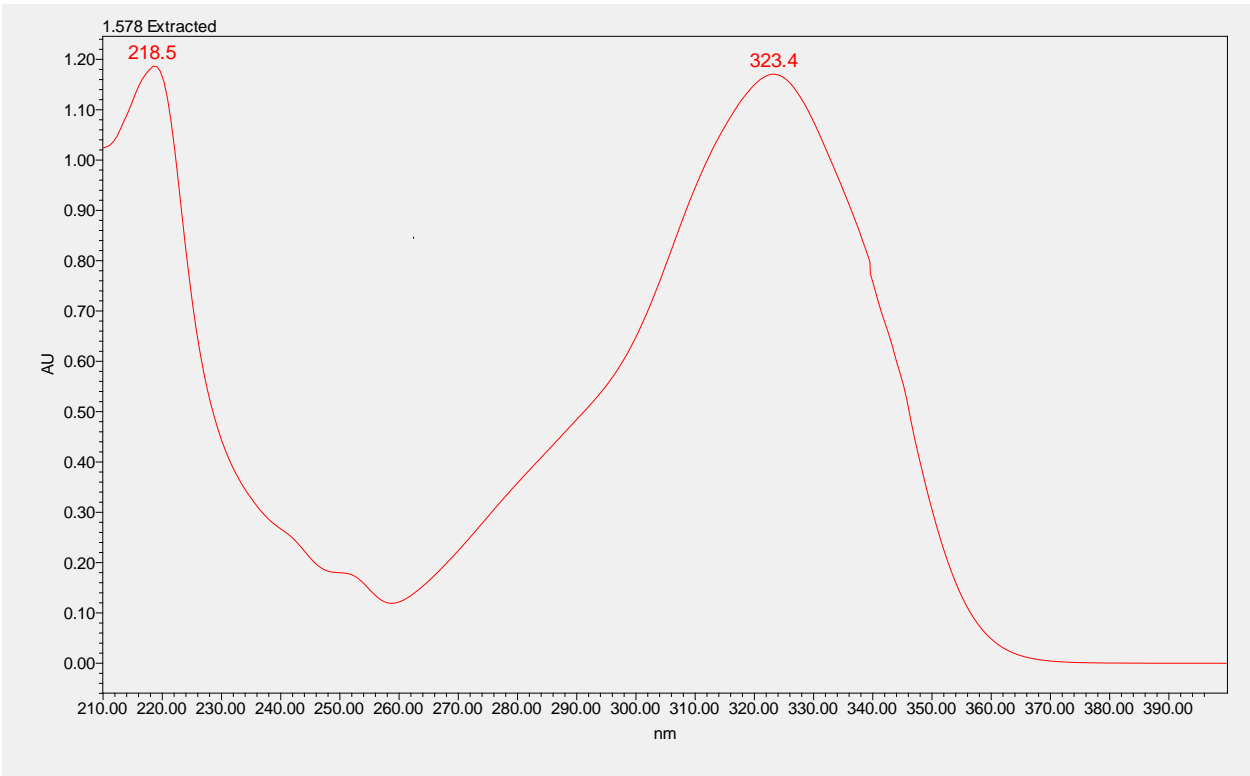
Amino Acid 2-7



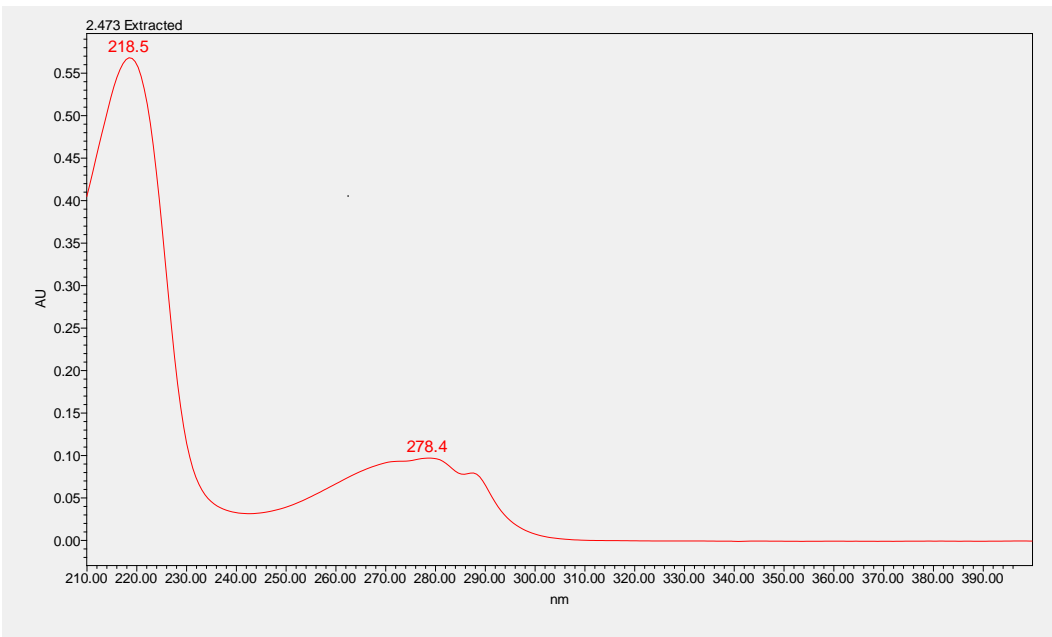
Amino Acid 4



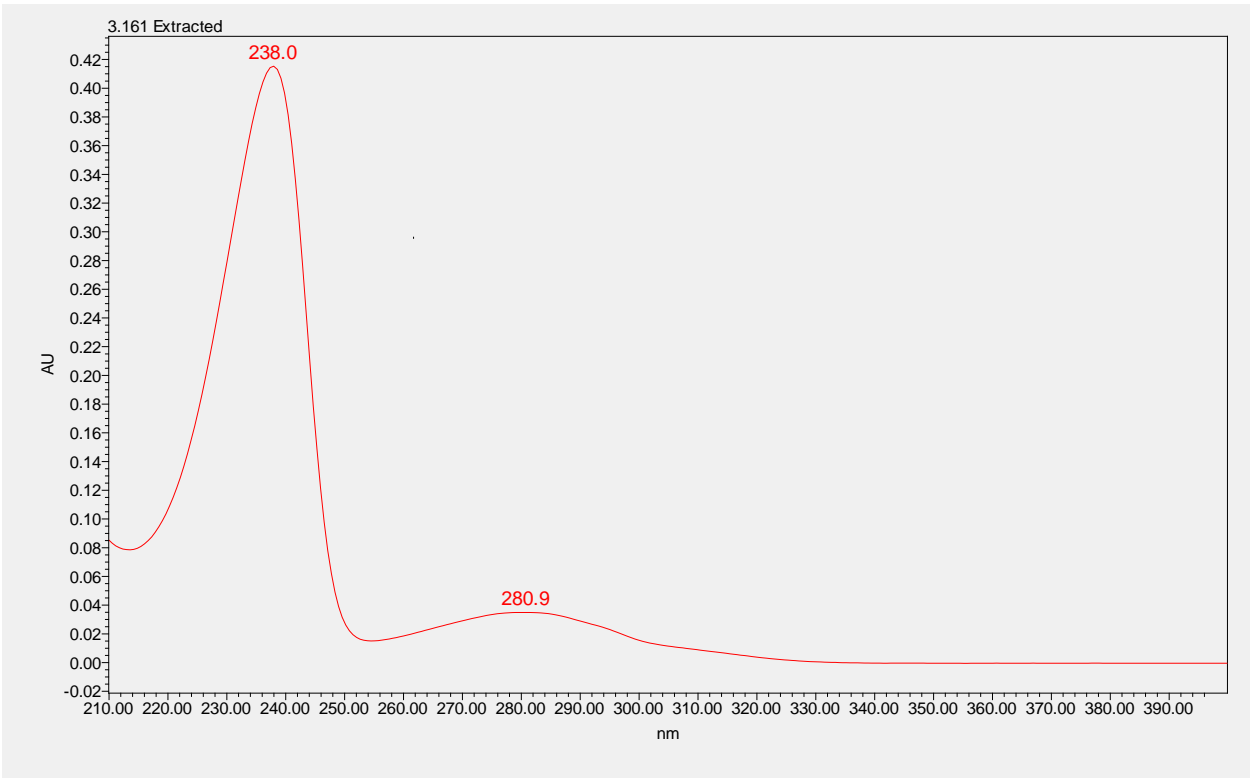
Amino Acid 5



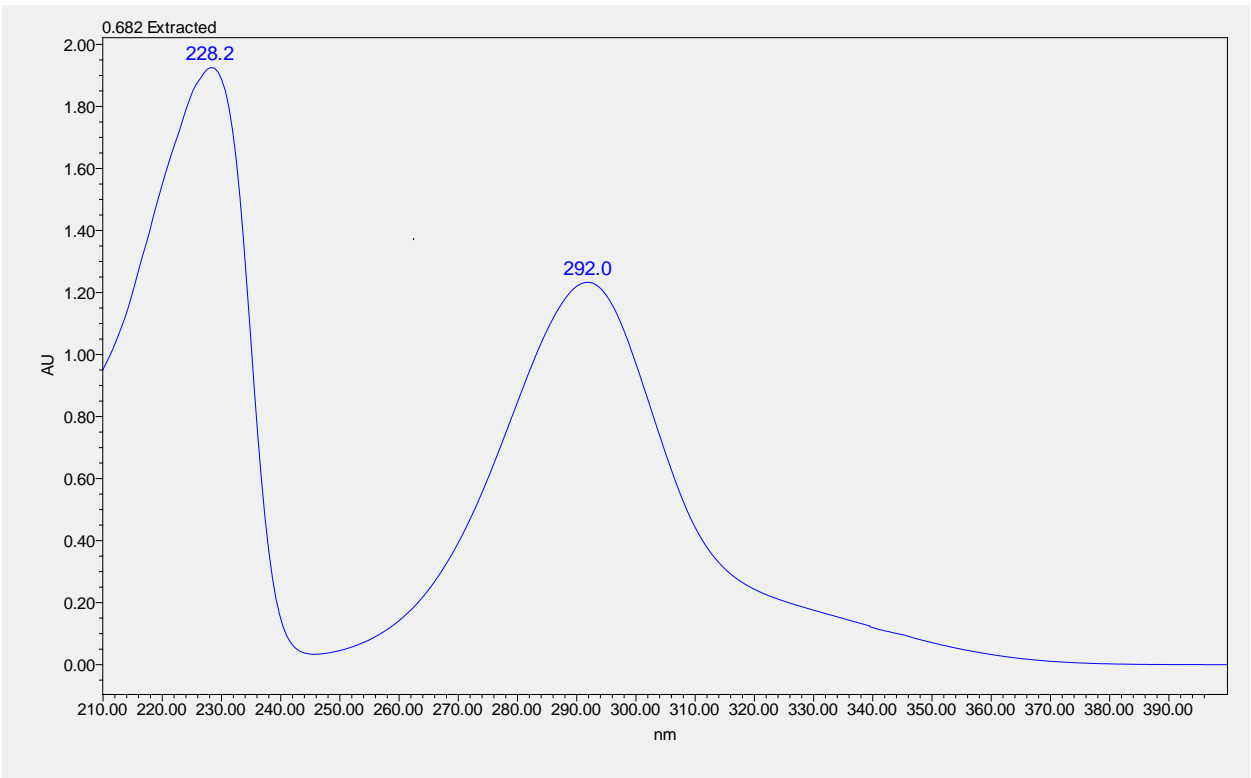
Amino Acid #3



Amino Acid 3-1

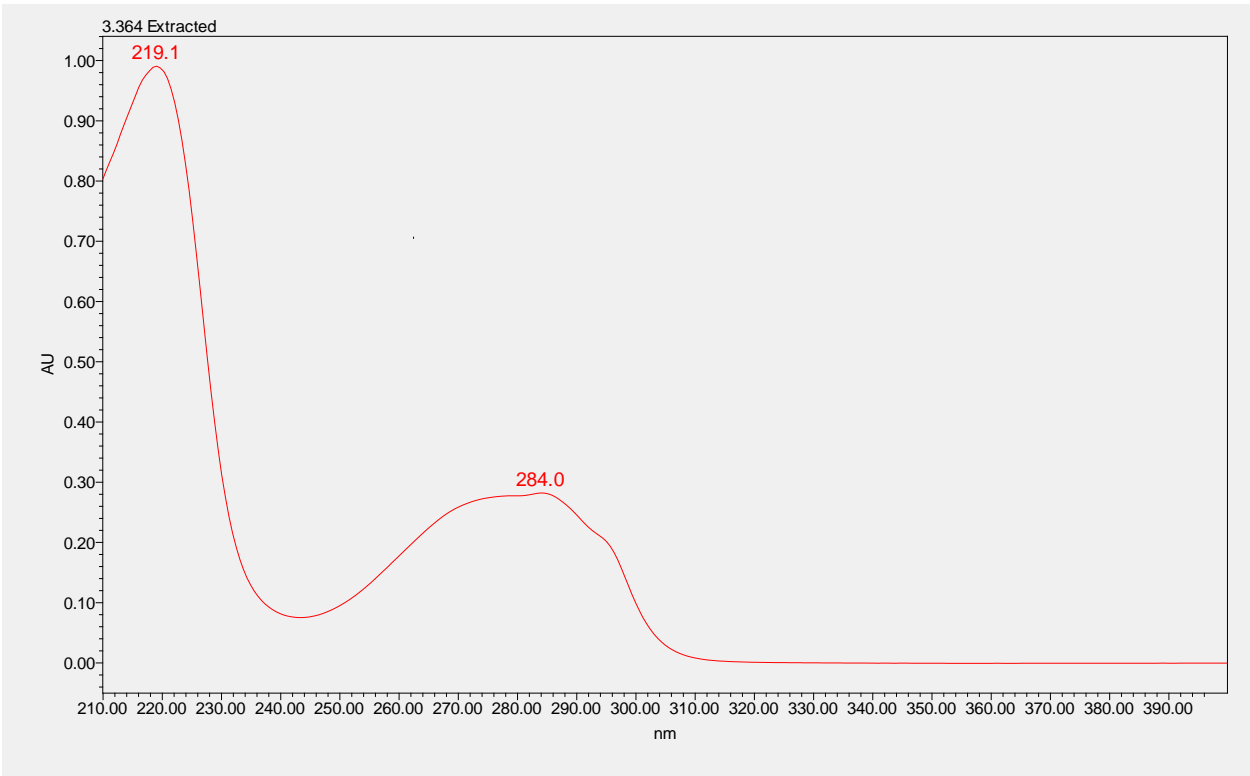


Amino Acid 3-2

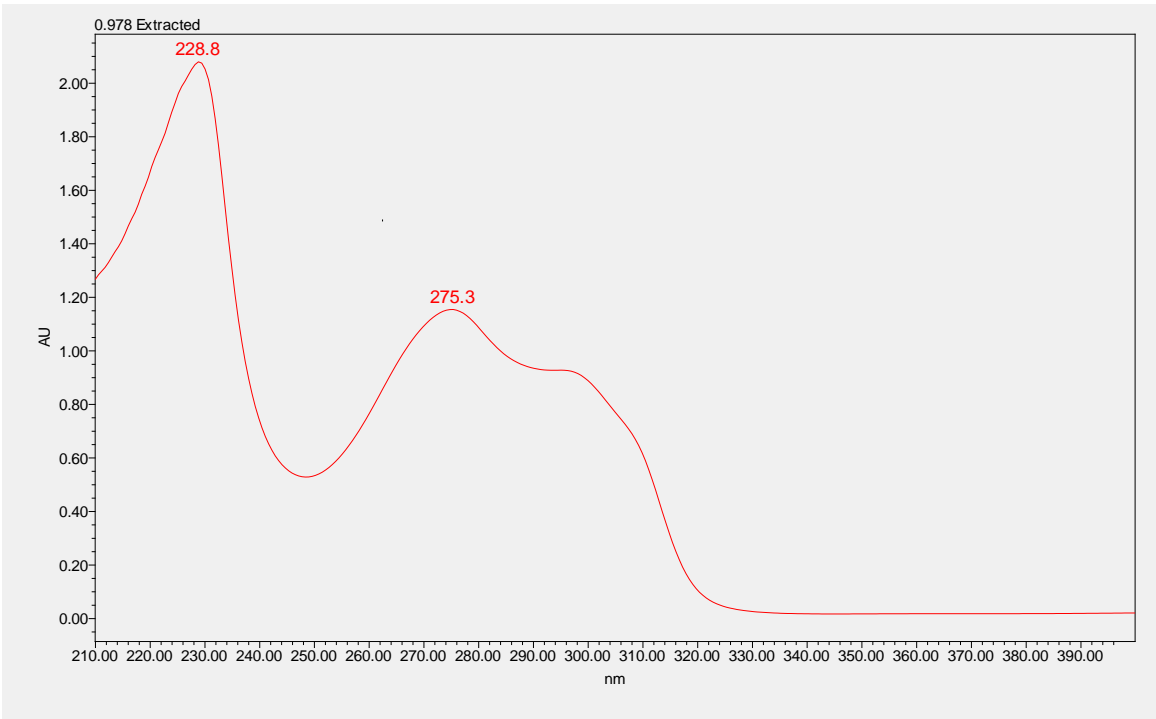




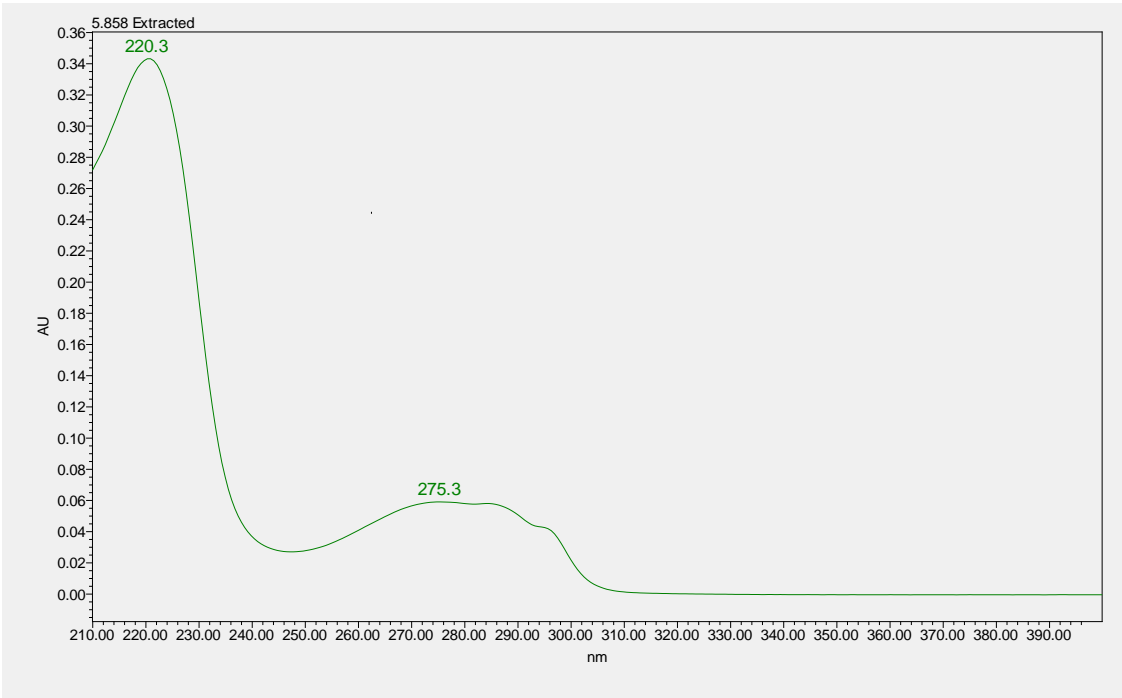
Amino Acid 3-3



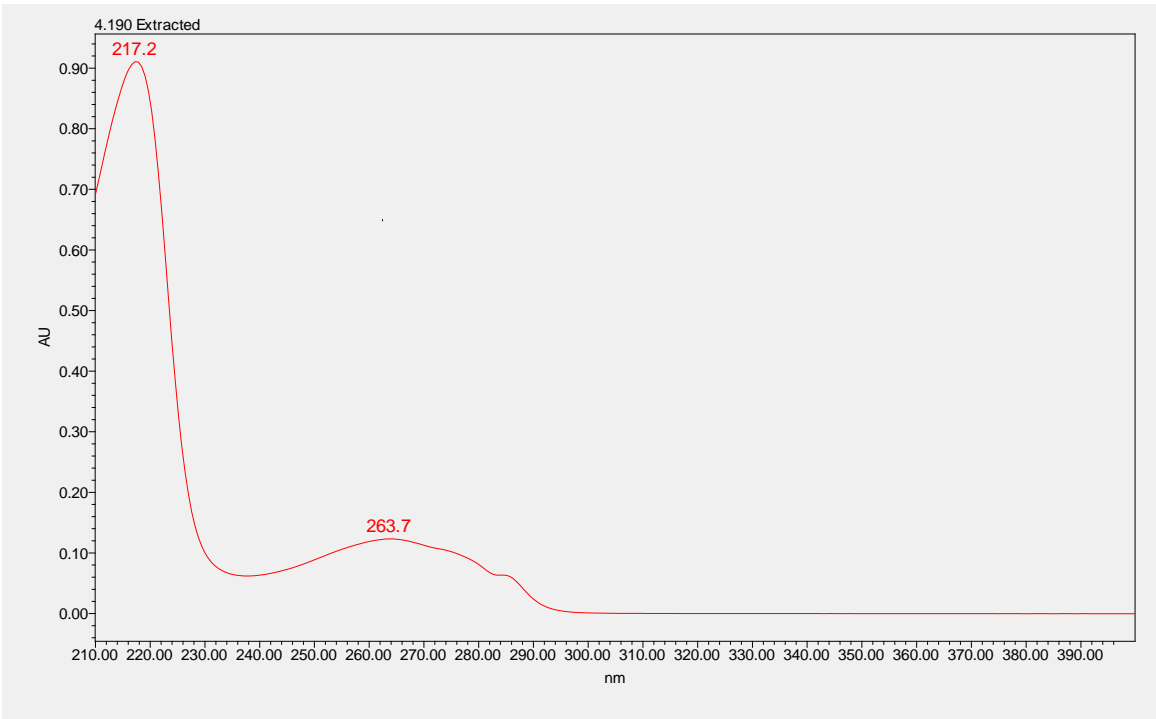
Amino Acid 3-4



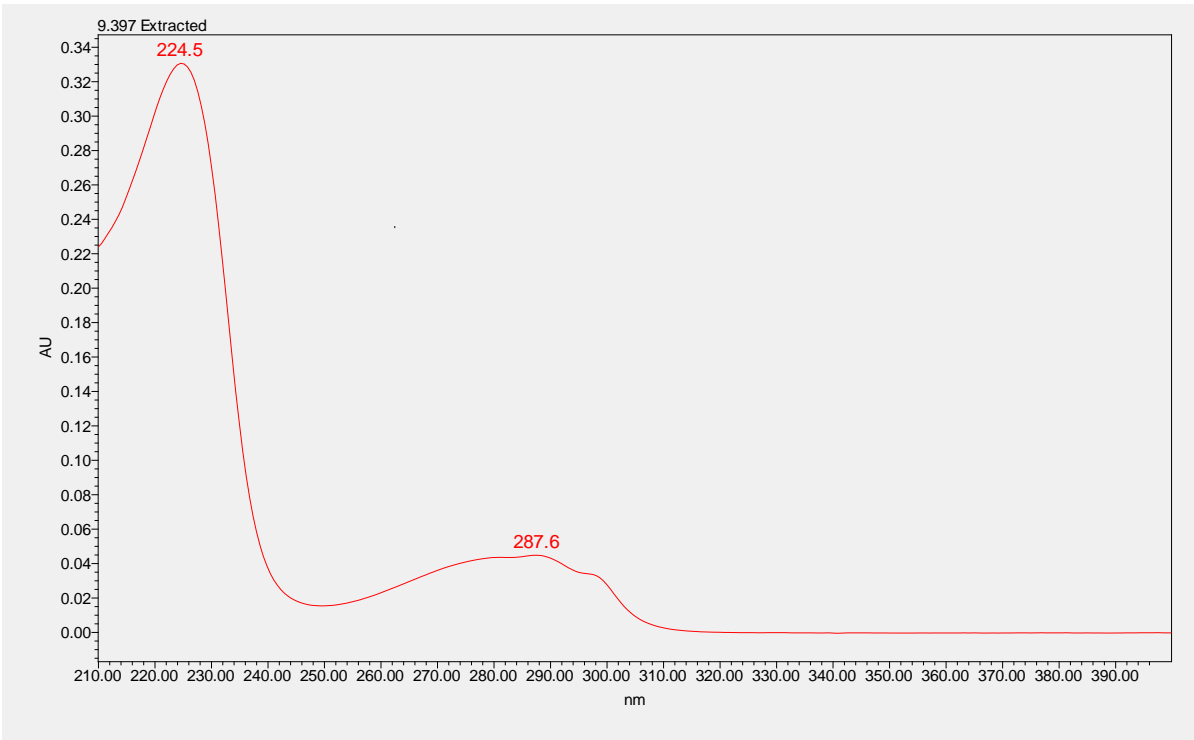
Amino Acid 3-5



Amino Acid #3-6



Amino Acid #3-7



Amino Acid #3-8

