

**DEPARTMENT OF FORENSIC SCIENCE  
METHOD VALIDATION SUMMARY FORM**

Section: Toxicology


Method: Addition of Xylazine by SPE to the Acid/Base/Neutral Drug Screen

And Quantitation by GC and GCMS

Results recorded?

Procedure documented?

Method fit for use?

Approved by:  Date: 5-1-23

MEMORANDUM FOR RECORD

From: Kimberly Meinweiser

Date: 04/18/2023

Handwritten initials/e-Signature: KM

Subject: Qualitative Validation for Xylazine

DISTRIBUTION:

<input type="checkbox"/> Director	<input type="checkbox"/> FS Lab #:	_____	Approved for SPE base screen, with methapyrilene as the ISTD. Do NOT use for the analysis of liver.  GMA 5/1/23
<input type="checkbox"/> Deputy Director	<input type="checkbox"/> Laboratory Director:	_____	
<input type="checkbox"/> DTS	<input checked="" type="checkbox"/> Program Manager:	Toxicology	
<input type="checkbox"/> HR	<input type="checkbox"/> Supervisor, Section:	_____	
<input type="checkbox"/> Counsel	<input type="checkbox"/> Other:	_____	

The following research for the qualitative determination of xylazine was conducted in the Eastern Laboratory between the dates of 09/23/2021 and 09/28/2021. The proposal was signed on 09/08/2021, which at the time, was a previous revision of the Toxicology SOP. This qualitative validation plan follows what was written in the SOP, revision 19, which was current at the time of the proposal. Data analysis was delayed due to the co-elution of xylazine and tramadol, absence from work and prioritizing casework.

The existing Base Screen GCMS/NPD Solid Phase Extraction (9.6.1) method from the Toxicology SOP was used with a reconstitution of 200uL THIA. The GCMS and NPD parameters are the standard parameters utilized at the Eastern Laboratory for routine base screen analysis.

Interferences from endogenous compounds were evaluated. A total of ten matrix blanks were screened on the GCMS; 1-4 (blood), 6-8 (urine), 9-10 (liver). All were evaluated for the presence of an interfering compound or instrumental response via extracted ion chromatograph for the major indicative ions for xylazine (i.e., m/z 205, 220, 130 and 145). All screened negative for xylazine. Four main matrix blanks were chosen for use in the validation; Matrix 1, Matrix 2, Matrix 6 and Matrix 10.

The analyte was extracted by itself at 2.0 mg/L in four separate matrix blanks. In the same four matrix blanks, the internal standard was extracted by itself. There was no interference documented by the internal standard.

Carryover was evaluated by running three injections of the 0.10 mg/L, 0.50 mg/L, 1.0 mg/L and 2.0 mg/L xylazine standard each followed by a matrix blank. None of the matrix blanks had any carryover observed on either the GCMS or the NPD.

The administrative LOD for the method is 0.005 mg/L. The 0.005 mg/L as well as a 0.010 mg/L were spiked and extracted from four different matrices; two blood, one urine and one liver. When spiked at 0.005 mg/L the retention times and spectral matches were less than acceptable. At 0.010 mg/L the retention times and spectral matches for all matrices were acceptable making 0.010 mg/L an acceptable LOD for xylazine.

Slight retention time shift was observed for Matrix 10 (liver), though still within the +/- 2% window. It would be recommended to run a matrix matched spiked standard for the qualitative determination in liver specimens.

The previously confirmed statewide base control drugs (diphenhydramine, pcp, tramadol, dextromethorphan, amitriptyline, nortriptyline, cyclobenzaprine, citalopram and trazodone) and the base in-house control drugs (doxylamine, chlorpheniramine, methadone, sertraline) were spiked alone at 2.0 mg/L and together with 2.0 mg/L of xylazine. Tramadol and xylazine have a very similar RT on the GCMS; tramadol~ 10.454 min (aafs library match 95) and xylazine~ 10.472 min (aafs library match 97 or >). Although the two drugs coelute, it is clear in all four matrices that there are two compounds present. On the NPD tramadol and xylazine do have slight separation, although if they are seen together, tramadol would need to be quantitated on the Opicoc method. On the NPD tramadol's RT is 11.033 (RRT ~0.9787) and xylazine RT 11.046 (RRT 0.9810) respectively.

Based on these findings, I request approval to qualitatively identify and report xylazine in case work using the modified Base Screen GCMS/NPD Solid Phase Extraction method described above. Data supporting this request is attached.

## MEMORANDUM FOR RECORD

From: Kimberly Meinweiser

Date: 04/18/2023

Handwritten initials/e-Signature: KM

Subject: Qualitative Validation for Xylazine

### DISTRIBUTION:

<input type="checkbox"/> Director	<input type="checkbox"/> FS Lab #:	_____
<input type="checkbox"/> Deputy Director	<input type="checkbox"/> Laboratory Director:	_____
<input type="checkbox"/> DTS	<input checked="" type="checkbox"/> Program Manager:	<u>Toxicology</u>
<input type="checkbox"/> HR	<input type="checkbox"/> Supervisor, Section:	_____ Laboratory: _____
<input type="checkbox"/> Counsel	<input type="checkbox"/> Other:	_____

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MEMORANDUM FOR RECORD

From: Kimberly Meinweiser

Date: 09/08/2021

Handwritten initials/e-Signature: Meinweiser Kimberly Ipu25291 Digitally signed by Meinweiser Kimberly Ipu25291  
Date: 2021.09.08 10:07:38 -04'00'

Subject: Qualitative Xylazine Validation Proposal

**DISTRIBUTION:**

<input type="checkbox"/> Director	<input type="checkbox"/> FS Lab #:	_____	James	Digitally signed by James Hutchings sme19098 Date: 2021.09.08 11:18:08 -04'00'
<input type="checkbox"/> Deputy Director	<input type="checkbox"/> Laboratory Director:	_____	Hutchings	
<input type="checkbox"/> DTS	<input checked="" type="checkbox"/> Program Manager:	James Hutchings, PhD	sme19098	
<input type="checkbox"/> HR	<input checked="" type="checkbox"/> Supervisor, Section:	Connie Luckie, PhD	Laboratory: <u>Eastern Toxicology</u>	
<input type="checkbox"/> Counsel	<input checked="" type="checkbox"/> Other:	Rebecca Wagner, PhD		

I am proposing to perform a validation for the qualitative analysis of xylazine by means of the existing Base Screen GCMD/NPD Solid Phase Extraction (9.6.1) method from the Toxicology SOP. Methapyrilene will be the internal standard. A certified Cerilliant standard will be used to perform the validation.

Xylazine will be studied at three concentrations spread across the linear range of the standard base quantitation curve; 0.010 mg/L, 0.10 mg/L and 2.0 mg/L.

Ten matrix blanks consisting of 5 blood samples, 3 urine samples and 2 liver homogenates will be screened for xylazine. They will also be examined to ensure that they have no endogenous contributions to the analyte signals.

Four matrix blanks will then be chosen for use in the validation.

The analyte will be extracted by itself at the following concentrations; 0.010 mg/L, 0.10 mg/L and 2.0 mg/L in four separate matrix blanks. In these same four matrix blanks, the internal standard will be extracted by itself.

Carryover will be evaluated by running three injections of the 0.10 mg/L xylazine standard followed by a matrix blank. Three more injections of the 2.0 mg/L xylazine standard will be injected followed by another matrix blank.

Common base drugs (diphenhydramine, doxylamine, tramadol, chlorpheniramine, dextromethorphan, methadone, amitriptyline, nortriptyline, cyclobenzaprine, sertraline and trazodone) will be spiked at 2.0 mg/L to evaluate for any interferences in four separate matrix blanks.

The administrative LOD for the current method is 0.0050 mg/L. This concentration will be spiked and extracted from four different matrices; two blood, one urine and one liver.

The predetermined identification criteria are listed below:

Retention time +/- 2%

Identification of peak by library search of the spectrum

Xylazine:

	0.01 RRT	match %	0.1 RRT	0.5 RRT	1.0 RRT	2.0 RRT	LOD RRT	match %
matrix 2	0.979	61	0.979	0.979	0.980	0.981	0.979	34
matrix 1	0.979	70	0.979	0.979	0.980	0.981	0.979	5
matrix 6	0.979	74	0.979	0.979	0.980	0.981	0.979	38
matrix 10	0.985	61	0.984	n/a		split peak	0.985	50

**avg RRT** across matrix 1, 2 and 6 for all conc: **0.980**

Tramadol:

matrix 2	0.978 *
matrix 1	0.976 *
matrix 6	0.978 *
matrix 10	0.978 *

\* Although co-elution of tramadol and xylazine is observed @ 2.0 mg/L, peaks are distinguishable albeit not baseline resolved.

Individualized mass spectral identification with library match is demonstrated in validation data.

# CERTIFICATE of ANALYSIS



## Xylazine

N-(2,6-dimethylphenyl)-5,6-dihydro-4H-1,3-thiazin-2-amine

Item No. 22641 • Batch No. 0508346

Purity Specification:  $\geq 98\%$

Molecular Formula : C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

CAS Number: 7361-61-7

Formula Weight: 220.3

Expiry date: 10JUN2023

Tests	Results
HPLC	Purity: 100.0 %
IR	Conforms
Mass spec	MH+: 221.0
Melting Point	135 - 138 °C
TLC	Purity: 100 %
NMR	Conforms

Reviewed and approved by: Jennifer LaBrecque

#### WARNING

THIS PRODUCT IS FOR RESEARCH USE - NOT FOR HUMAN OR VETERINARY DIAGNOSTIC OR THERAPEUTIC USE. IT IS THE RESPONSIBILITY OF THE PURCHASER TO DETERMINE SUITABILITY FOR OTHER APPLICATIONS.

#### SAFETY DATA

This material should be considered hazardous until further information becomes available. Do not ingest, inhale, get in eyes, on skin, or on clothing. Wash thoroughly after handling. Before use, the user must review the [complete](#) Safety Data Sheet, which has been sent via email to your institution.

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# DRUG STOCK STANDARD PREPARATION LOG

Standard & Concentration Prepared: xylozine working stock - 1mg/mL Assigned Lot#: X41A-20210922

Date Prepared	Assigned Expiration Date	Employee Initials	Drug Name	Vendor Source / Lot # / Expiration Date
9.22.21	6.10.23	KM	xylozine	Cayman 0508346 6.10.23

Method of Preparation: Salt Correction: Most drug standards are prepared as a salt or hydrate, such as phencyclidine hydrochloride, morphine sulfate or benzoyllecgonine - 4H<sub>2</sub>O. The extra weight of the salt or hydrate must be corrected for when preparing liquid drug standards. The following formula is used to determine the amount of drug salt to be weighed.

$$[\text{mg parent drug desired}] \times \left[ \frac{\text{FW drug salt (mg/mmmole)}}{\text{FW parent drug (mg/mmmole)}} \right] = \text{mg drug salt required}$$

FW Drug: 220.33 mg Weighed: 11.65  
 Solvent: MeOH Manuf./Lot: B:J/DZ846-US Volume: 10 mL  
 FW Drug Salt: 256.79

weigh 11.65mg std into a 10mL volumetric + qs to volume w/ MeOH



**ASSAY:**

Base Screen

QC Data File: Xylazine Validation

Analyst(s):

K.Meinweiser

Sample / FS #	Item #	Sample Info
1 MATRIX BLANK 1	BLD	BB0321-1
2 MATRIX BLANK 2	BLD	BB0721-1
3 MATRIX BLANK 3	BLD	BB0121-1
4 MATRIX BLANK 4	BLD	BB0121-2
5 MATRIX BLANK 5	BLD	BB0519-3
6 MATRIX BLANK 6	URN	BU0101017
7 MATRIX BLANK 7	URN	BU06017
8 MATRIX BLANK 8	URN	BU0819
9 MATRIX BLANK 9	LVR	LIV0109
10 MATRIX BLANK 10	LVR	LIV09231 *satisfies 2.8.1.1.4
11		
12		
13		
14		
15		
16		
17		
18		
19		
20		
21		
22		
23		
24		
25		

Specimens removed from TX ADM storage, aliquots removed for analysis, specimens returned to TX ADM storage by, unless otherwise noted:

Analyst: Kimberly Meinweiser

Date: 09.23.21

Time Removed: 0726 Time Returned: 0743

220-F156 Toxicology Barcode Batch Worksheet  
Issued by Toxicology Program Manager  
Issue Date: 02-July-2020

Sample / FS #	Item #	Sample Info
26		
27		
28		
29		
30		
31		
32		
33		
34		
35		
36		
37		
38		
39		
40		
41		
42		
43		
44		
45		
46		
47		
48		
49		
50		

Extraction: LL / SPE SPE lot #: 024170-XD

Control Charts Updated (Initials/Date): N/A

Reviewer / Date: \_\_\_\_\_



## Sequence Table (Front Injector):

Line	Location	SampleName	Method Name	Num Inj	SampleType
1	Vial 131	ETOAC	ALKALI	1	Sample
2	Vial 132	MATRIX BLANK 1 BLD	ALKALI	1	Sample
3	Vial 133	MATRIX BLANK 2 BLD	ALKALI	1	Sample
4	Vial 134	MATRIX BLANK 3 BLD	ALKALI	1	Sample
5	Vial 135	MATRIX BLANK 4 BLD	ALKALI	1	Sample
6	Vial 136	MATRIX BLANK 5 BLD	ALKALI	1	Sample
7	Vial 137	MATRIX BLANK 6 URN	ALKALI	1	Sample
8	Vial 138	MATRIX BLANK 7 URN	ALKALI	1	Sample
9	Vial 139	MATRIX BLANK 8 URN	ALKALI	1	Sample
10	Vial 140	SOLVENT BLANK AFTER MATIX 8	ALKALI	1	Sample
11	Vial 141	MATRIX BLANK 9 LVR	ALKALI	1	Sample
12	Vial 142	SOLVENT BLANK AFTER MATRIX 9	ALKALI	1	Sample
13	Vial 143	MATRIX BLANK 10 LVR	ALKALI	1	Sample
14	Vial 144	SOLVENT BLANK AFTER MATRIX 10	ALKALI	1	Sample
15	Vial 145	OFF	ALKALIOFF	1	Sample

Vial Check  
EL 9.24.21

method: C:\CHEM32\1\METHODS\ALKALI.M  
Modified on: 9/23/2021 at 11:21:17 AM

Method Information

Method: C:\CHEM32\1\METHODS\ALKALI.M  
Modified: 9/23/2021 at 11:21:17 AM

KM/QC DATA IN XYLAZINE VALIDATION

Injection Source and Location

Injection Source: GC Injector

Injection Location: Front

=====

Agilent 7890B

=====

GC  
Oven  
Equilibration Time 0.5 min  
Max Temperature 325 °C  
Slow Fan Disabled  
Oven Program On  
Oven Program 90 °C for 2 min  
Oven#1 then 15 °C/min to 245 °C for 0 min  
Oven#2 then 30 °C/min to 300 °C for 7 min  
Run Time 21.167 min  
Cryo Off

ALS  
Front Injector  
Syringe Size 10 µL  
Syringe A Syringe has not been selected.  
Injection Volume 1 µL  
Solvent A Washes (PreInj) 2  
Solvent A Washes (PostInj) 4  
Solvent A Volume 8 µL  
Solvent B Washes (PreInj) 2  
Solvent B Washes (PostInj) 4  
Solvent B Volume 8 µL  
Sample Washes 0  
Sample Wash Volume 8 µL  
Sample Pumps 2  
Dwell Time (PreInj) 0 min  
Dwell Time (PostInj) 0 min  
Solvent Wash Draw Speed 300 µL/min  
Solvent Wash Dispense Speed 3000 µL/min  
Sample Wash Draw Speed 300 µL/min  
Sample Wash Dispense Speed 3000 µL/min  
Injection Dispense Speed 6000 µL/min  
Viscosity Delay 1 sec  
Sample Depth Disabled  
Injection Type Standard  
L1 Airgap 0.2 µL

Tray  
Barcode heater Disabled  
Barcode mixer Disabled

Sample Overlap  
Mode Sample overlap is not enabled

ALS Errors Pause for user interaction

Front SS Inlet He  
Mode Pulsed Splitless  
Heater On 260 °C  
Pressure On 18.881 psi  
Total Flow On 44.5 mL/min  
Septum Purge Flow On 3 mL/min  
Gas Saver On 20 After 2 min mL/min  
Injection Pulse Pressure 40 psi Until 0.3 min

Purge Flow to Split Vent 40 mL/min at 0.5 min  
Liner Agilent 5190-2293: 900 µL (Splitless, single tape)

Back SS Inlet He

\*\*\*Excluded from Affecting GC's Readiness State\*\*

Mode Splitless  
Heater On 250 °C  
Pressure On 9.801 psi  
Total Flow On 41.7 mL/min  
Septum Purge Flow Off  
Gas Saver Off  
Purge Flow to Split Vent 40 mL/min at 0.5 min  
Liner Agilent 5183-4647; Lot P02-C6823: 870 µL (Split,

Column  
Column #1

RTX-1; SN 1652461  
0 °C-350 °C (350 °C): 30 m x 250 µm x 0.25 µm  
Column lock Unlocked  
In Front SS Inlet He  
Out Front Detector NPD  
(Initial) 90 °C  
Pressure 18.881 psi  
Flow 1.5 mL/min  
Average Velocity 36.082 cm/sec  
Holdup Time 1.3857 min  
Flow Program Off  
Flow Program 1.5 mL/min for 1 min  
Flow then 0.2 mL/min per min to 2.2 mL/min for 0 m  
Run Time 21.167 min

Column #2

10223  
RTX-5  
0 °C-325 °C (325 °C): 30 m x 320 µm x 0.25 µm  
Column lock Unlocked  
In Back SS Inlet He  
Out Back Detector FID  
(Initial) 90 °C  
Pressure 9.801 psi  
Flow 1.7 mL/min  
Average Velocity 31.621 cm/sec  
Holdup Time 1.5812 min  
Flow Program Off  
Flow Program 1.7 mL/min for 1 min  
Flow then 0.2 mL/min per min to 3 mL/min for 0 min  
Run Time 21.167 min

Front Detector NPD

Heater On 310 °C  
H2 Flow On 3 mL/min  
Air Flow On 60 mL/min  
Makeup Flow (Combined) On 10 mL/min  
Carrier Gas Flow Correction Included in Makeup Flow  
Bead On  
Signal 1 Type NPD  
Maximum Bead Voltage 4.095 V  
Bios Bead No

Dry Bead Yes  
Auto Adjust Bead No

Back Detector FID

\*\*\*Excluded from Affecting GC's Readiness State\*\*

Heater On 280 °C  
H2 Flow Off  
Air Flow Off  
Makeup Flow On 25 mL/min  
Carrier Gas Flow Correction Does not affect Makeup or Fuel Flow  
Flame Off  
Electrometer On

Signals

Signal #1: Front Signal  
Description Front Signal  
Details  
Save On  
Data Rate 50 Hz

Signal #2: Back Signal  
Description Back Signal  
Details  
Save Off  
Data Rate 5 Hz

Signal #3: Test Plot  
Description Test Plot  
Details  
Save Off  
Data Rate 50 Hz

Signal #4: Test Plot  
Description Test Plot  
Details  
Save Off  
Data Rate 50 Hz

Run Time Events  
Run Time Events  
Run Time Events Time 0 min  
Run Time Events Event Detector H2 Fuel Flow  
Run Time Events Position 0  
Run Time Events Setpoint Off

Run Time Events Time 2 min  
Run Time Events Event Detector H2 Fuel Flow  
Run Time Events Position 0  
Run Time Events Setpoint On

Sequence Name: C:\MassHunter\GCMS\1\sequence\alkalival.sequence.xml

Comment: Base

Operator: KM/QC DATA IN XYLAZINE VALIDATION

Data Path: C:\MassHunter\GCMS\1\data\BASES\092321\

Instrument Control Pre-Seq Cmd:

Data Analysis Pre-Seq Cmd:

Instrument Control Post-Seq Cmd:

Data Analysis Post-Seq Cmd:

Method Sections To Run

Full Method

Reprocessing Only

Sequence Barcode Options

On Mismatch, Inject Anyway

On Mismatch, Don't Inject

Barcode Disabled

---

Line Type	ALS	File	Method	Sample Name/Misc Info
1) Sample	41	SOL	ALKALI	ETOAC
2) Sample	42	001BLD	ALKALI	MATRIX BLANK 1 BLD
3) Sample	43	002BLD	ALKALI	MATRIX BLANK 2 BLD
4) Sample	44	003BLD	ALKALI	MATRIX BLANK 3 BLD
5) Sample	45	004BLD	ALKALI	MATRIX BLANK 4 BLD
6) Sample	46	005BLD	ALKALI	MATRIX BLANK 5 BLD
7) Sample	47	006URN	ALKALI	MATRIX BLANK 6 URN
8) Sample	48	007URN	ALKALI	MATRIX BLANK 7 URN
9) Sample	49	008URN	ALKALI	MATRIX BLANK 8 URN
10) Sample	50	SOL008	ALKALI	SOLVENT AFTER MATRIX 8
11) Sample	51	009LVR	ALKALI	MATRIX BLANK 9 LVR
12) Sample	52	SOL009	ALKALI	SOLVENT AFTER MATRIX 9
13) Sample	53	010LVR	ALKALI	MATRIX BLANK 10 LVR
14) Sample	54	SOL010	ALKALI	SOLVENT AFTER MATRIX 10

Vial Check  
EL 9.24.21



## Single Quadrupole Acquisition Method - MS Parameters Report

<b>Method file</b>	C:\MassHunter\GCMS\1\methods\ALKALI.m
<b>Tune file</b>	ATUNE.U
<b>Ion source</b>	EI
<b>Source temperature (°C)</b>	230
<b>Quad temperature (°C)</b>	150
<b>Fixed Electron energy (eV)</b>	70.3
<b>Acquisition Type</b>	Scan
<b>Stop time (min)</b>	22.17
<b>Solvent delay (min)</b>	3.00
<b>Trace Ion Detection</b>	False
<b>Gain Factor</b>	1
<b>EM Saver</b>	False
<b>EM Saver Limit</b>	N/A

### Scan Time Segments

Time	Start Mass	End Mass	Threshold	Scan Speed
3.00	40	550	150	1,562 [N=2]

### Timed Events

Time	Type of Event	Parameter
------	---------------	-----------

### Real-Time Plots

Type of Plot	Label	Low Mass	High Mass
Total Ion	N/A	N/A	N/A
Spectrum	N/A	N/A	N/A
Extracted Ion	Scan 1-1	40	550

### Self-Cleaning Ion Source Parameters

Mode	No Cleaning
------	-------------

C:\MassHunter\GCMS\1\methods\ALKALI.m  
 Thu Sep 23 12:02:25 2021

Control Information

-----  
 Sample Inlet : GC  
 Injection Source : GC ALS  
 Injection Location: Front  
 Mass Spectrometer : Enabled

No Sample Prep method has been assigned to this method.

GC  
 GC Summary  
 Run Time 22.167 min  
 Post Run Time 0 min

Oven  
 Temperature  
 Setpoint On  
 (Initial) 90 °C  
 Hold Time 2 min  
 Post Run 0 °C  
 Program  
 #1 Rate 15 °C/min  
 #1 Value 245 °C  
 #1 Hold Time 0 min  
 #2 Rate 30 °C/min  
 #2 Value 300 °C  
 #2 Hold Time 8 min

Equilibration Time 0.5 min  
 Max Temperature 325 °C  
 Maximum Temperature Override Disabled  
 Slow Fan Disabled

ALS  
 Front Injector  
 Syringe Size 10 µL  
 Injection Volume 1 µL  
 Injection Repetitions 1  
 Injection Delay 0 sec  
 Solvent A Washes (PreInj) 4  
 Solvent A Washes (PostInj) 4  
 Solvent A Volume 8 µL  
 Solvent B Washes (PreInj) 4  
 Solvent B Washes (PostInj) 4  
 Solvent B Volume 8 µL  
 Sample Washes 0  
 Sample Wash Volume 8 µL  
 Sample Pumps 2  
 Dwell Time (PreInj) 0 min  
 Dwell Time (PostInj) 0 min  
 Solvent Wash Draw Speed 300 µL/min  
 Solvent Wash Dispense Speed 3000 µL/min  
 Sample Wash Draw Speed 300 µL/min  
 Sample Wash Dispense Speed 3000 µL/min

Injection Dispense Speed	6000 µL/min
Viscosity Delay	0 sec
Sample Depth	Disabled
Tower Fan	On
Solvent Wash Mode	A-A2,B-B2
Sample Overlap Mode	Sample overlap is not enabled
ALS Errors	Pause for user interaction
Front SS Inlet He Mode	Pulsed Splitless
Heater	On 260 °C
Pressure	On 15.177 psi
Total Flow	On 46.869 mL/min
Septum Purge Flow	On 3 mL/min
Septum Purge Flow Mode	Standard
Gas Saver	On 20 After 2 min mL/min
Injection Pulse Pressure	40 psi Until 0.3 min
Purge Flow to Split Vent	42.4 mL/min at 0.5 min
Liner	A Liner has not been selected.
Thermal Aux 2 (MSD Transfer Line) Temperature Setpoint (Initial)	On 280 °C
Column	
Column #1	
Flow	
Setpoint (Initial)	On 1.4687 mL/min
Hold Time	1 min
Post Run	1.2 mL/min
Program	
#1 Rate	0.2 mL/min per min
#1 Value	3 mL/min
#1 Hold Time	0 min
Column Information	Restek 10123
Rtx-1 SN 1652464	
Temperature Range	-60 °C-350 °C (350 °C)
Dimensions	30 m x 250 µm x 0.25 µm
In	Front SS Inlet He
Out	MSD
(Initial)	90 °C
Pressure	15.177 psi
Flow	1.4687 mL/min
Average Velocity	45 cm/sec
Holdup Time	1.1111 min
Control Mode	Ramped Flow
Column Outlet Pressure	0 psi
Front Detector NPD	
Makeup	***Excluded from Affecting GC's Readiness State*** He
Heater	Off
H2 Flow	Off
Air Flow	Off
Makeup Flow	Off

Carrier Gas Flow Correction	Constant Makeup and Fuel Flow
Bead	Off
Maximum Bead Voltage	4.095 V
Bios Bead	No
Dry Bead	Yes
Auto Adjust Bead	No

Signals	
Signal #1:	
Description	None
Signal #2:	
Description	None
Signal #3:	
Description	None
Signal #4:	
Description	None

TUNE PARAMETERS for SN: US71236345

Trace Ion Detection is OFF.

34.610	:	EMISSION	
70.347	:	ENERGY	
28.568	:	REPELLER	
90.157	:	IONFOCUS	
28.500	:	ENTRANCE_LENS	
1870.588	:	EMVOLTS	
			1870.588 : Actual EMV
			0.78 : GAIN FACTOR
2280.000	:	AMUGAIN	
120.063	:	AMUOFFSET	
1.000	:	FILAMENT	
0.000	:	DCPOLARITY	
19.576	:	ENTLENSOFFSET	
-532.000	:	MASSGAIN	
-36.000	:	MASSOFFSET	

END OF TUNE PARAMETERS

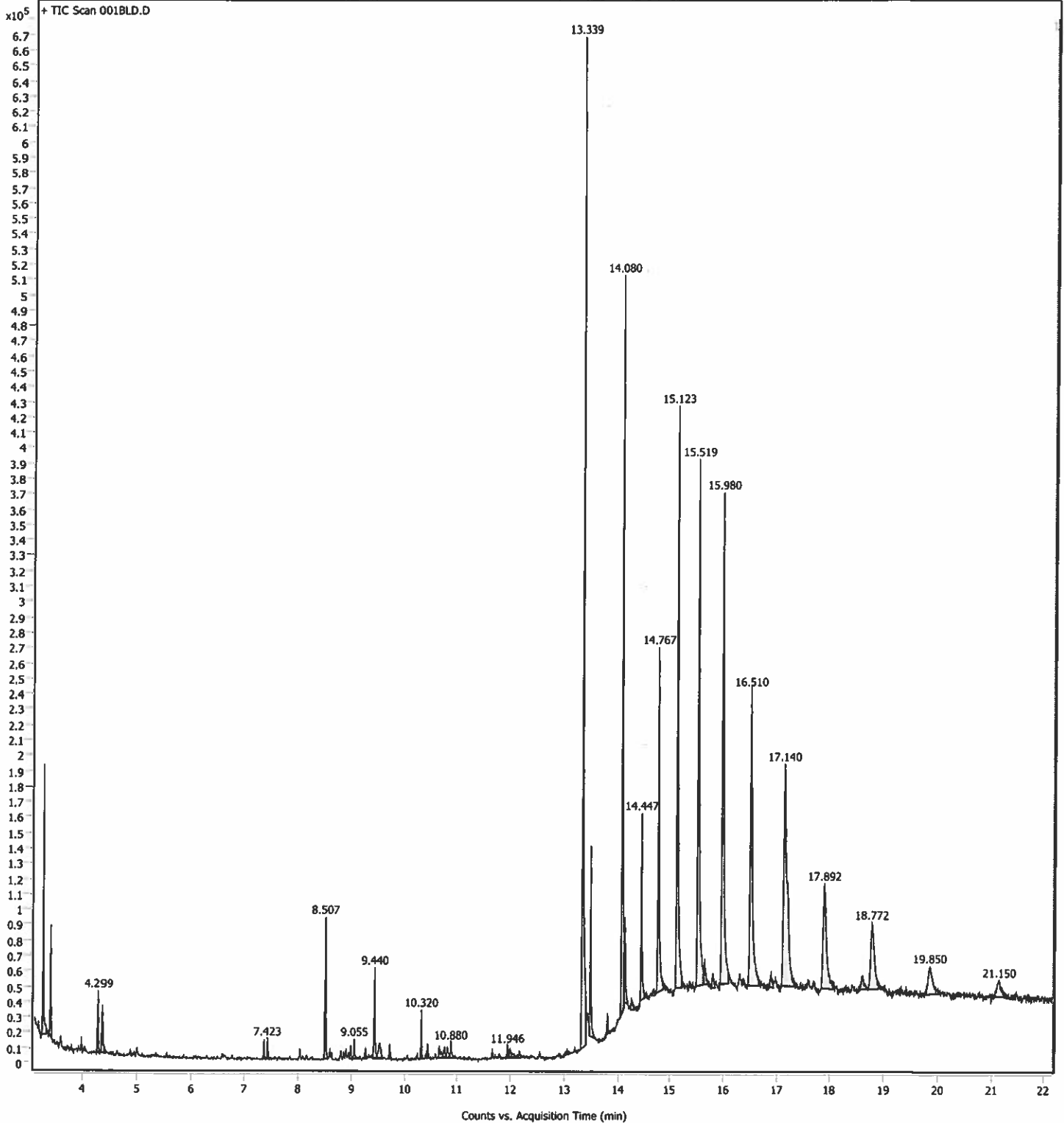
END OF INSTRUMENT CONTROL PARAMETERS

# Analysis Report

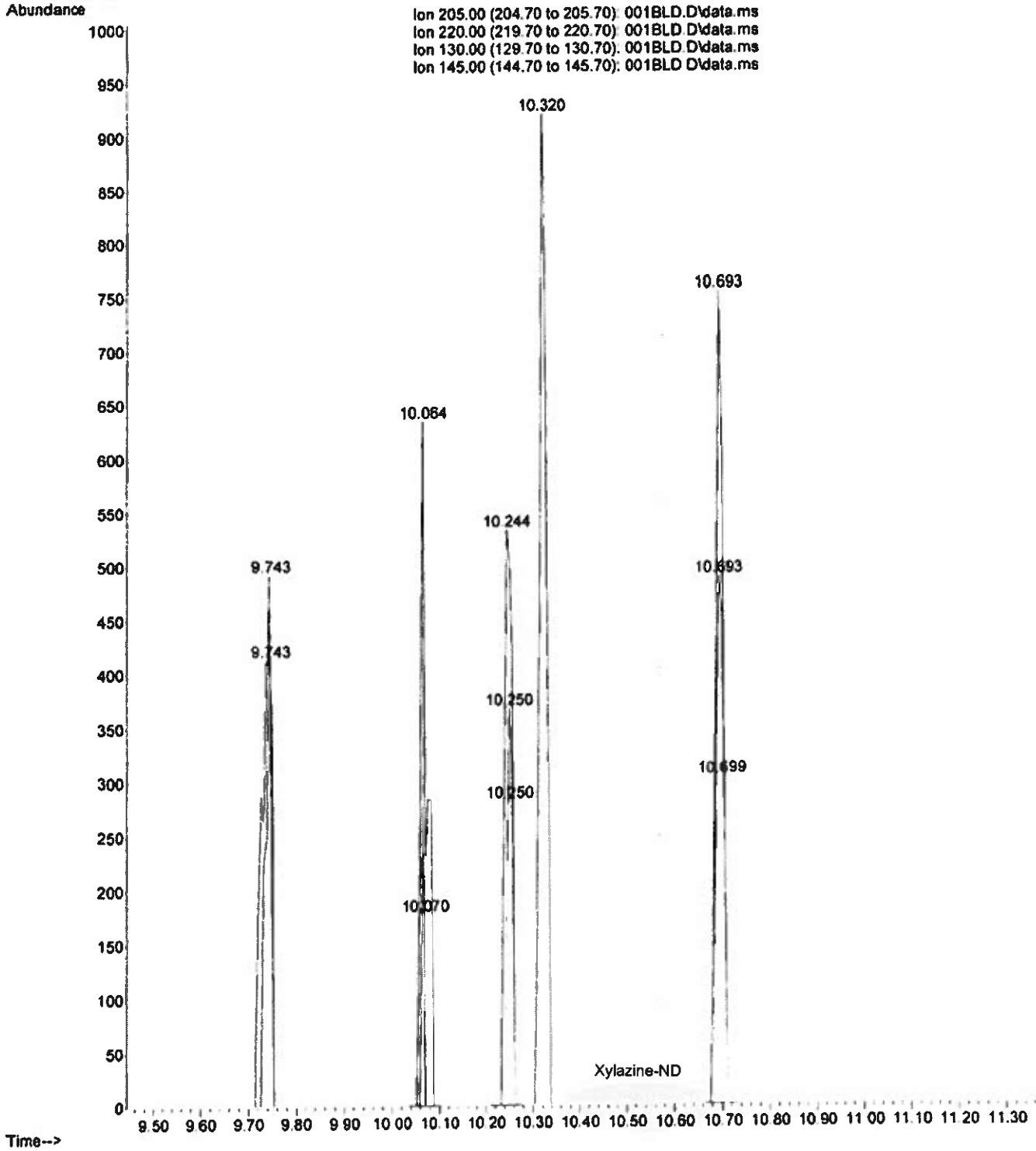
## Sample Information

Sample Name	MATRIX BLANK 1 BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092321\001BLD.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/23/2021 12:34:24 PM (UTC-04:00)
Position	42	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

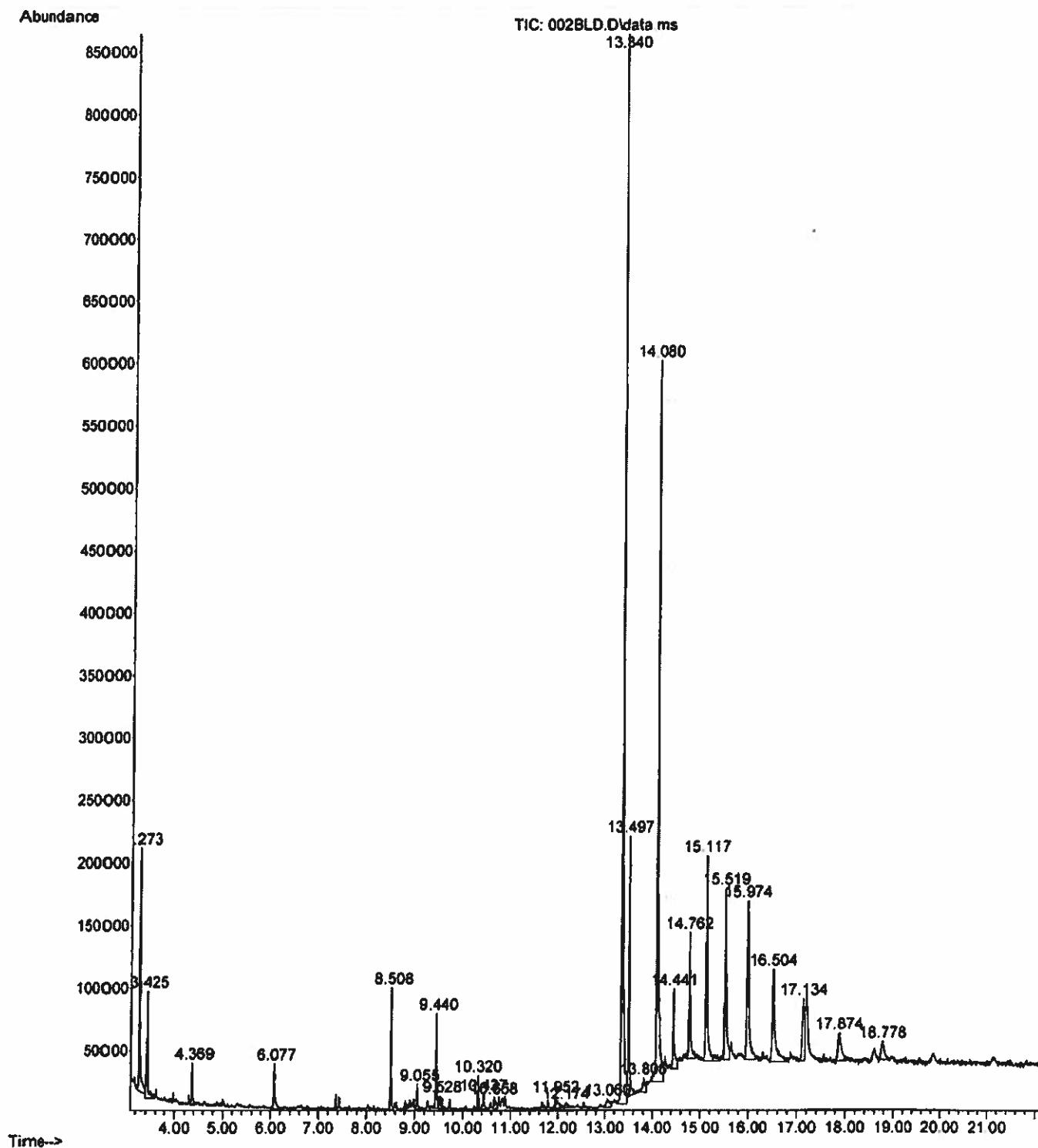
## Sample Chromatograms



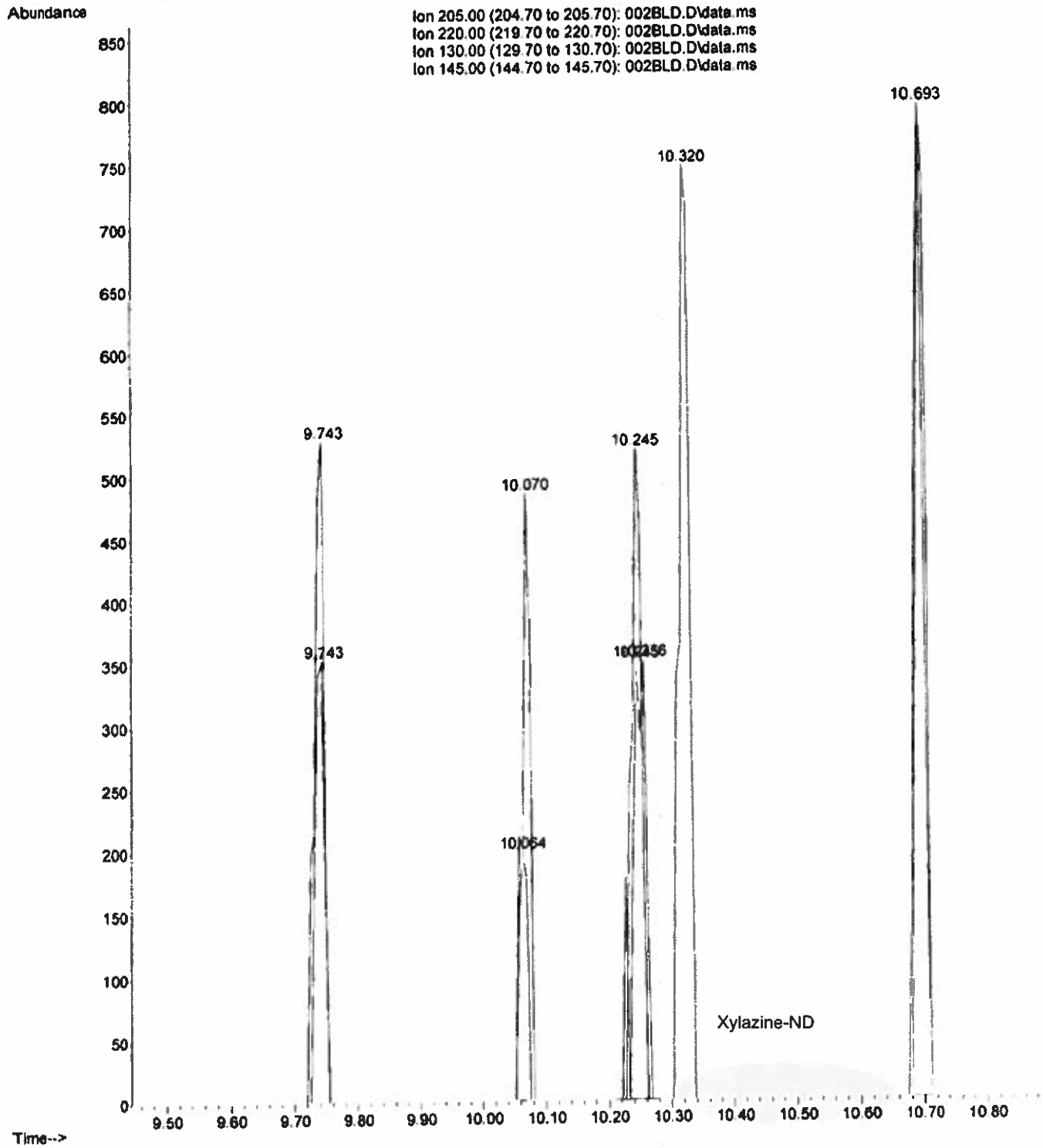
File :C:\Users\TOX\Desktop\092321\001BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 12:34 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 1 BLD  
Misc Info : BB0321-1  
Vial Number: 42



File :C:\Users\TOX\Desktop\092321\002BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 13:00 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 2 BLD  
Misc Info : BB0721-1  
Vial Number: 43

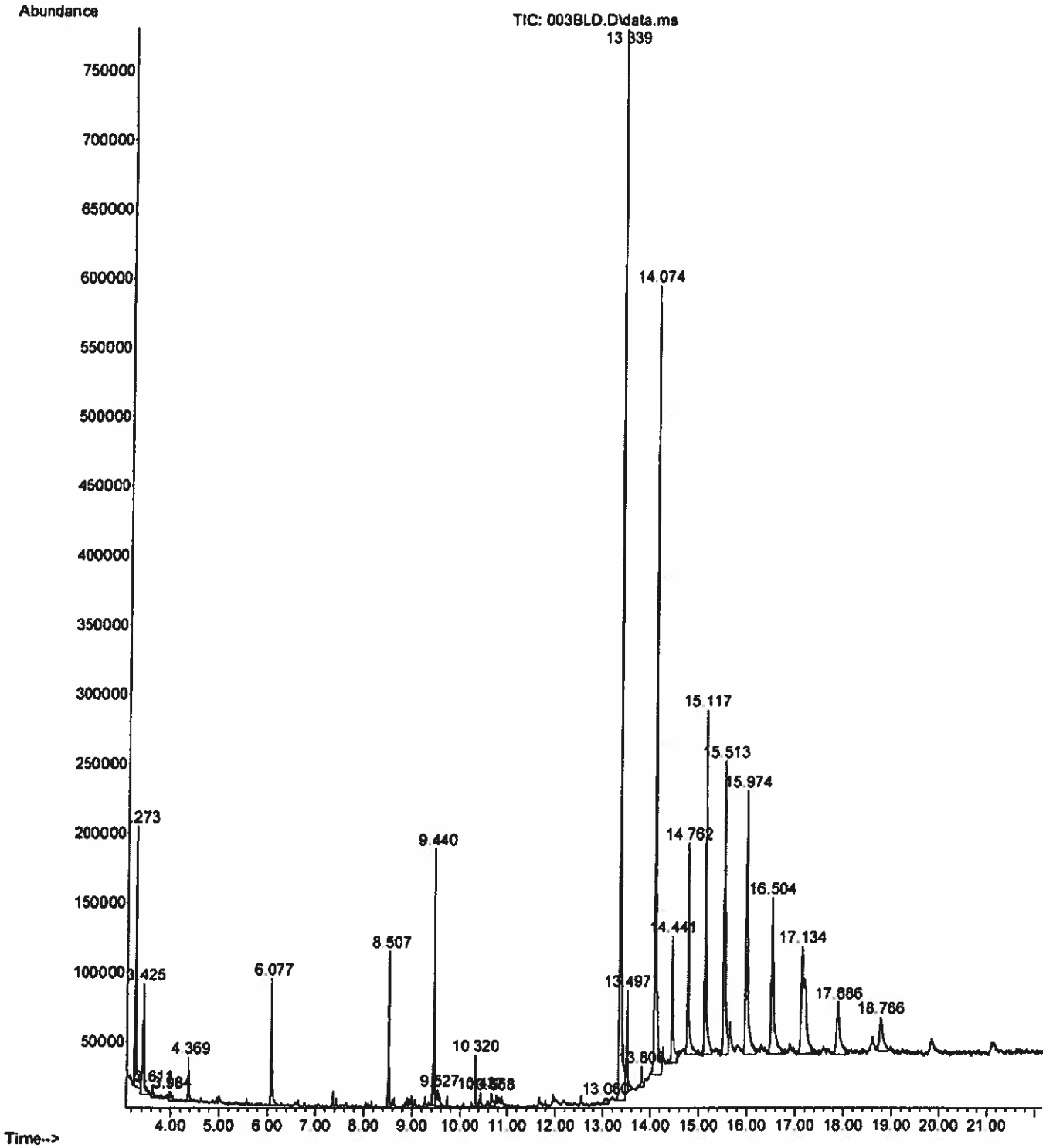


File : C:\Users\TOX\Desktop\092321\002BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 13:00 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 2 BLD  
Misc Info : B80721-1  
Vial Number: 43

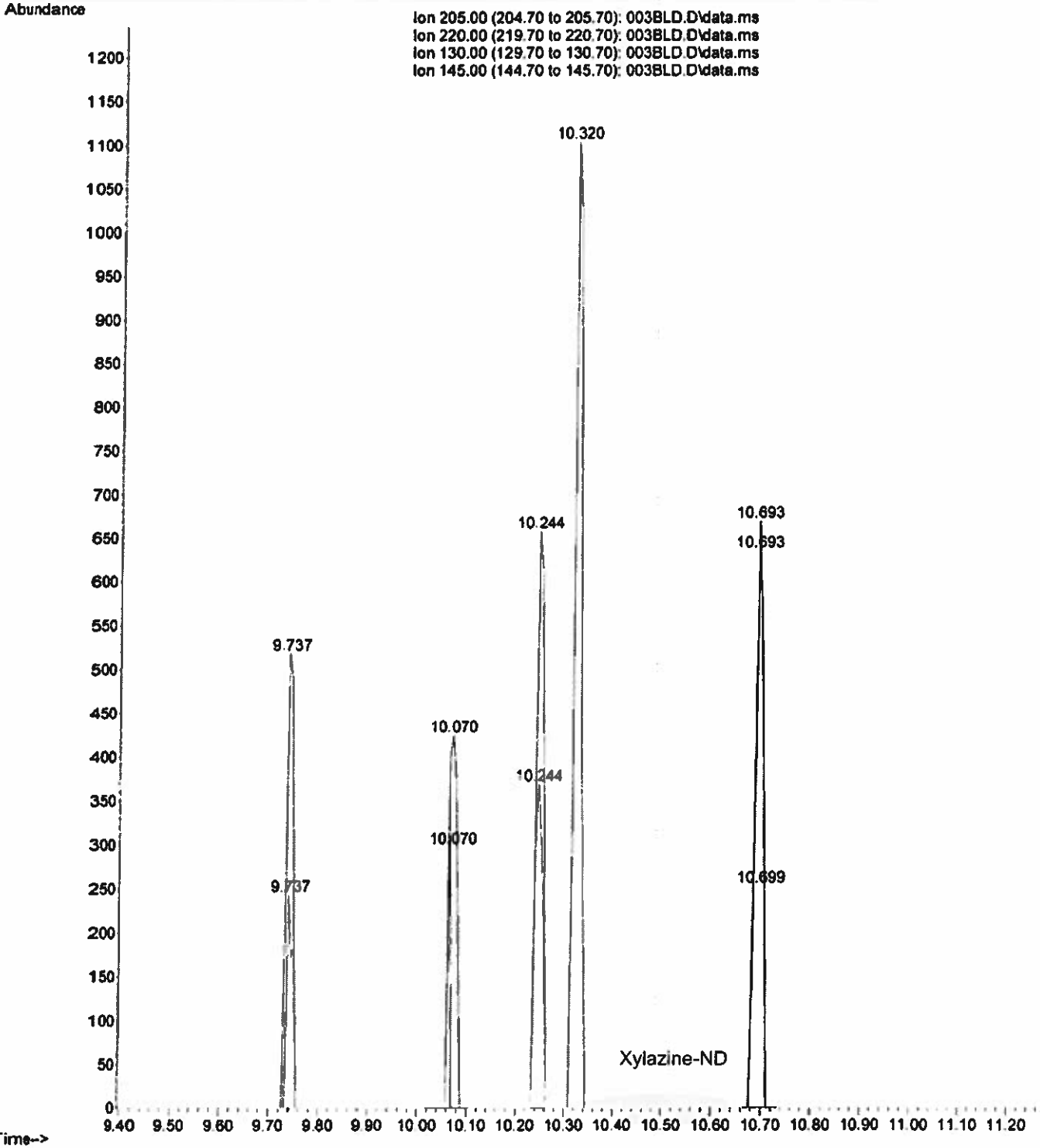




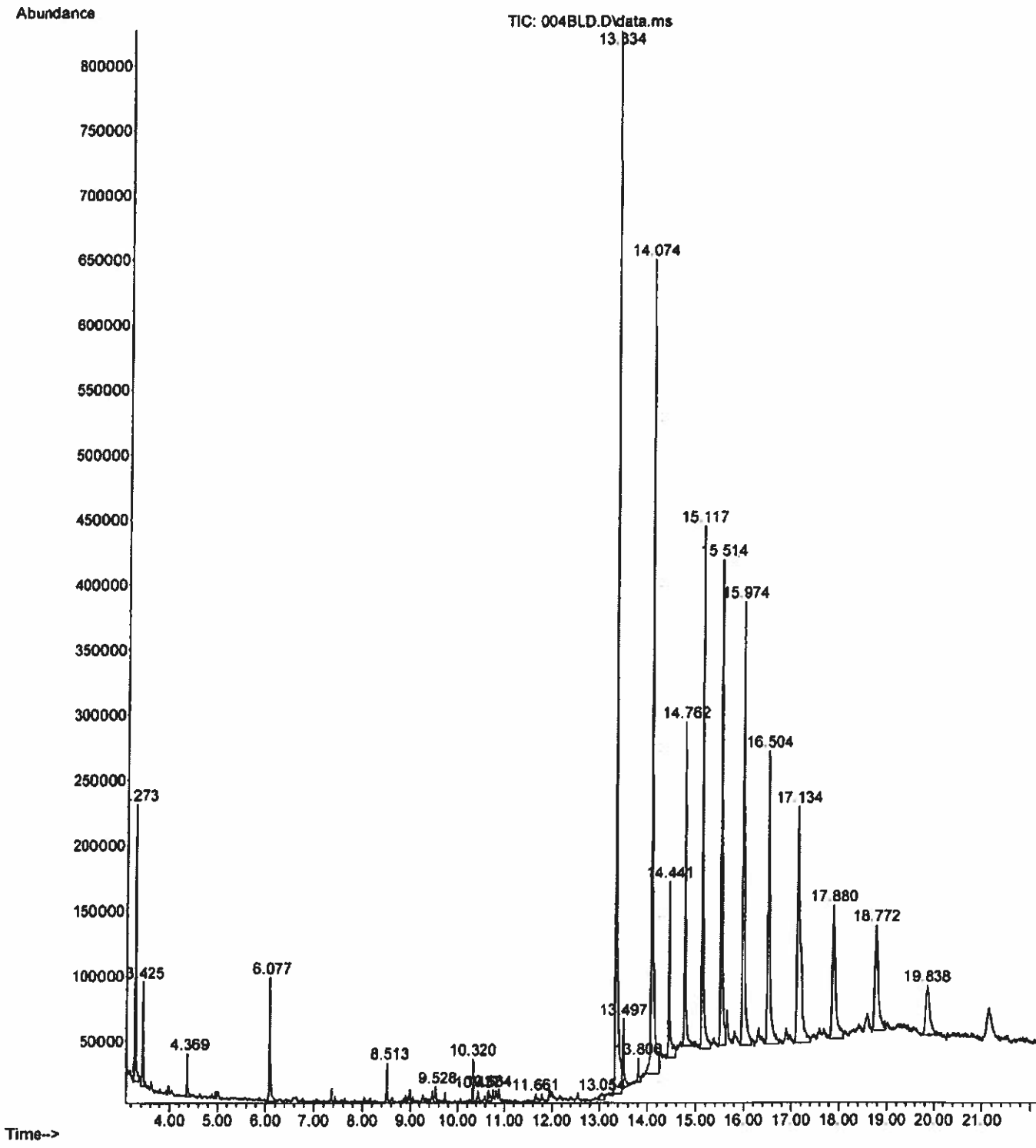
File :C:\Users\TOX\Desktop\092321\003BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 13:26 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 3 BLD  
Misc Info : B80121-1  
Vial Number: 44



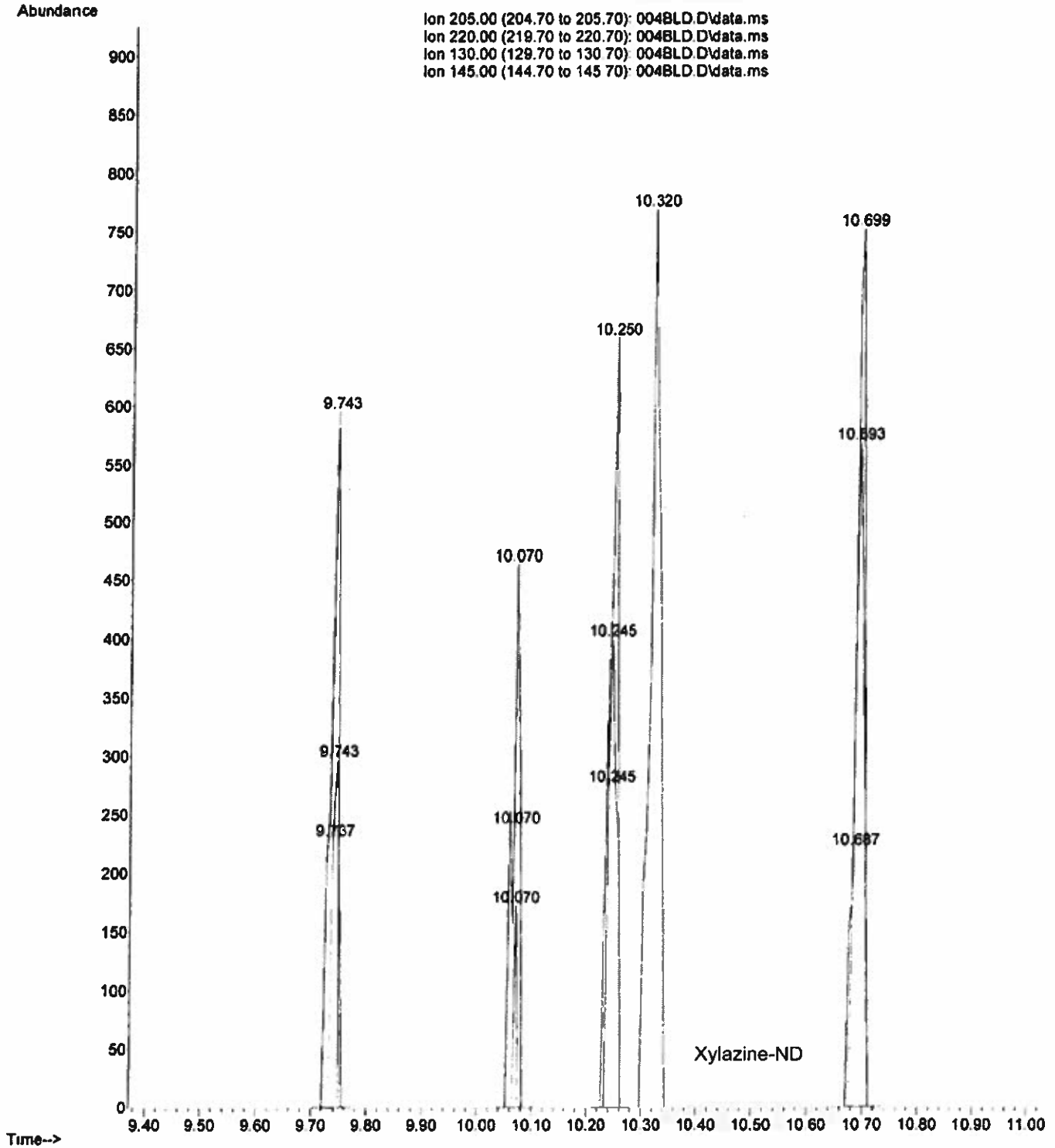
File :C:\Users\TOX\Desktop\092321\003BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 13:26 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 3 BLD  
Misc Info : BB0121-1  
Vial Number: 44



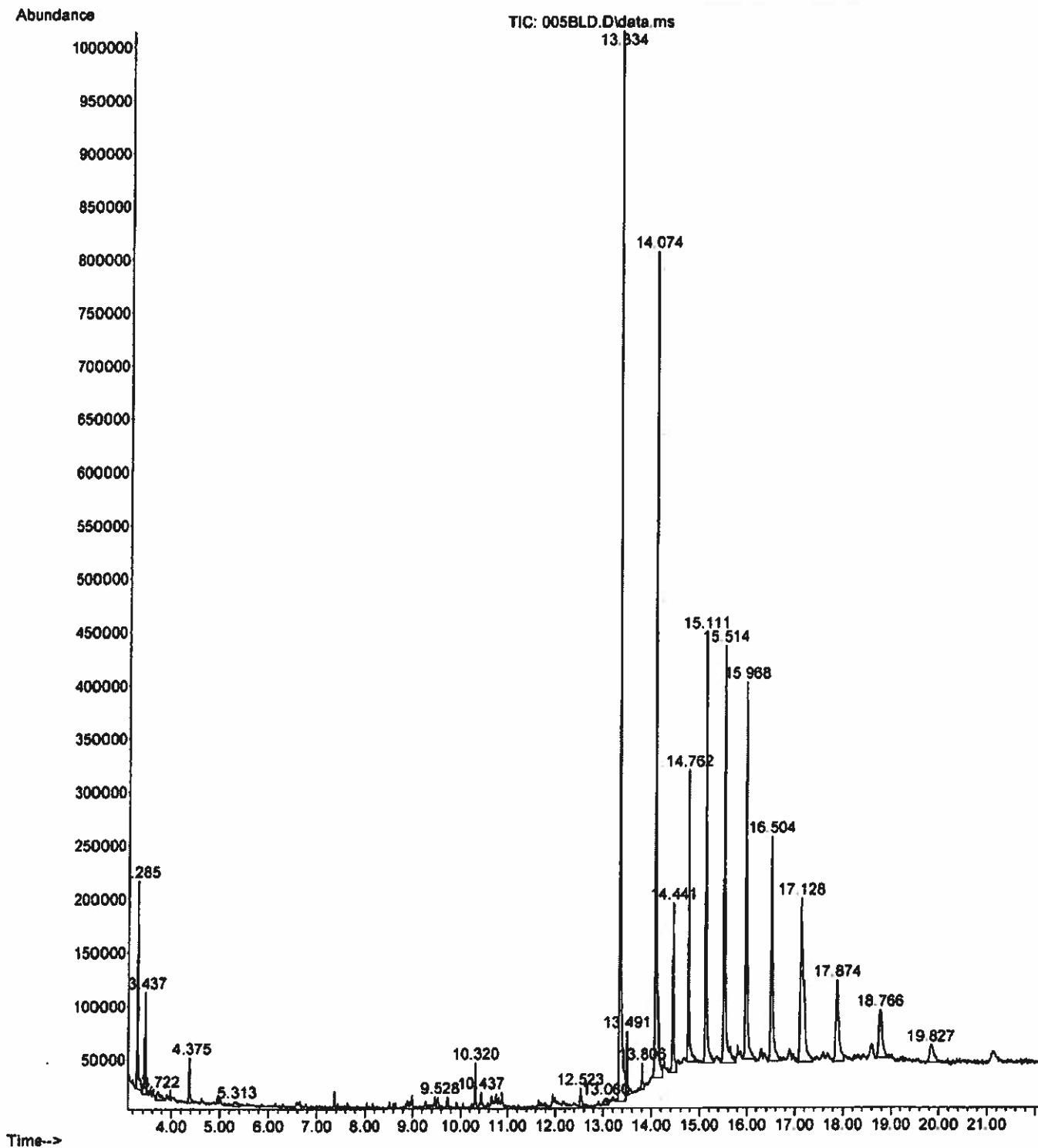
File :C:\Users\TOX\Desktop\092321\004BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 13:51 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 4 BLD  
Misc Info : 880121-2  
Vial Number: 45



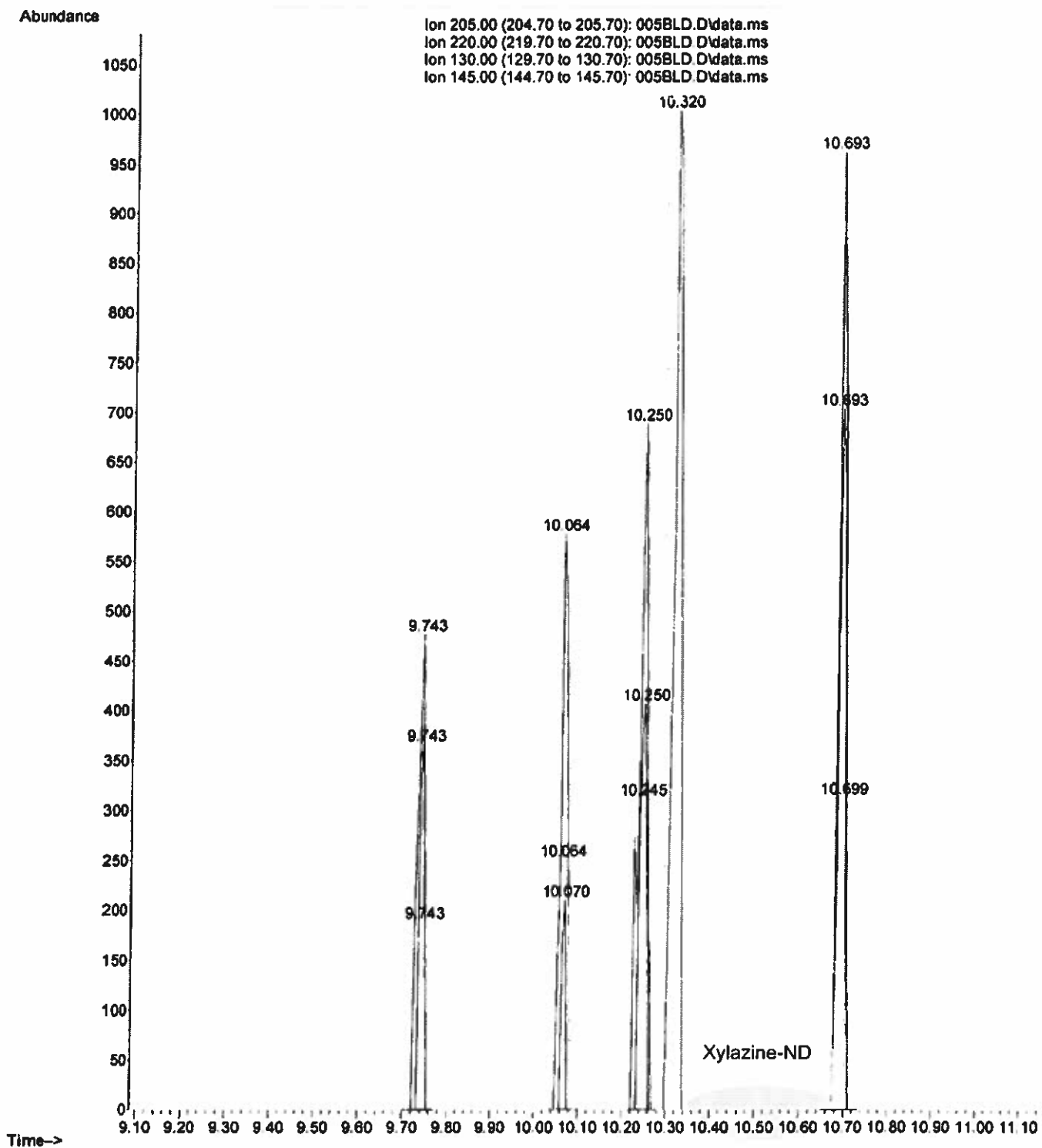
File :C:\Users\TOX\Desktop\092321\004BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 13:51 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 4 BLD  
Misc Info : BB0121-2  
Vial Number: 45



File :C:\Users\TOX\Desktop\092321\005BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 14:17 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 5 BLD  
Misc Info : BB0519-3  
Vial Number: 46



File :C:\Users\TOX\Desktop\092321\005BLD.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 14:17 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 5 BLD  
Misc Info : BB0519-3  
Vial Number: 46

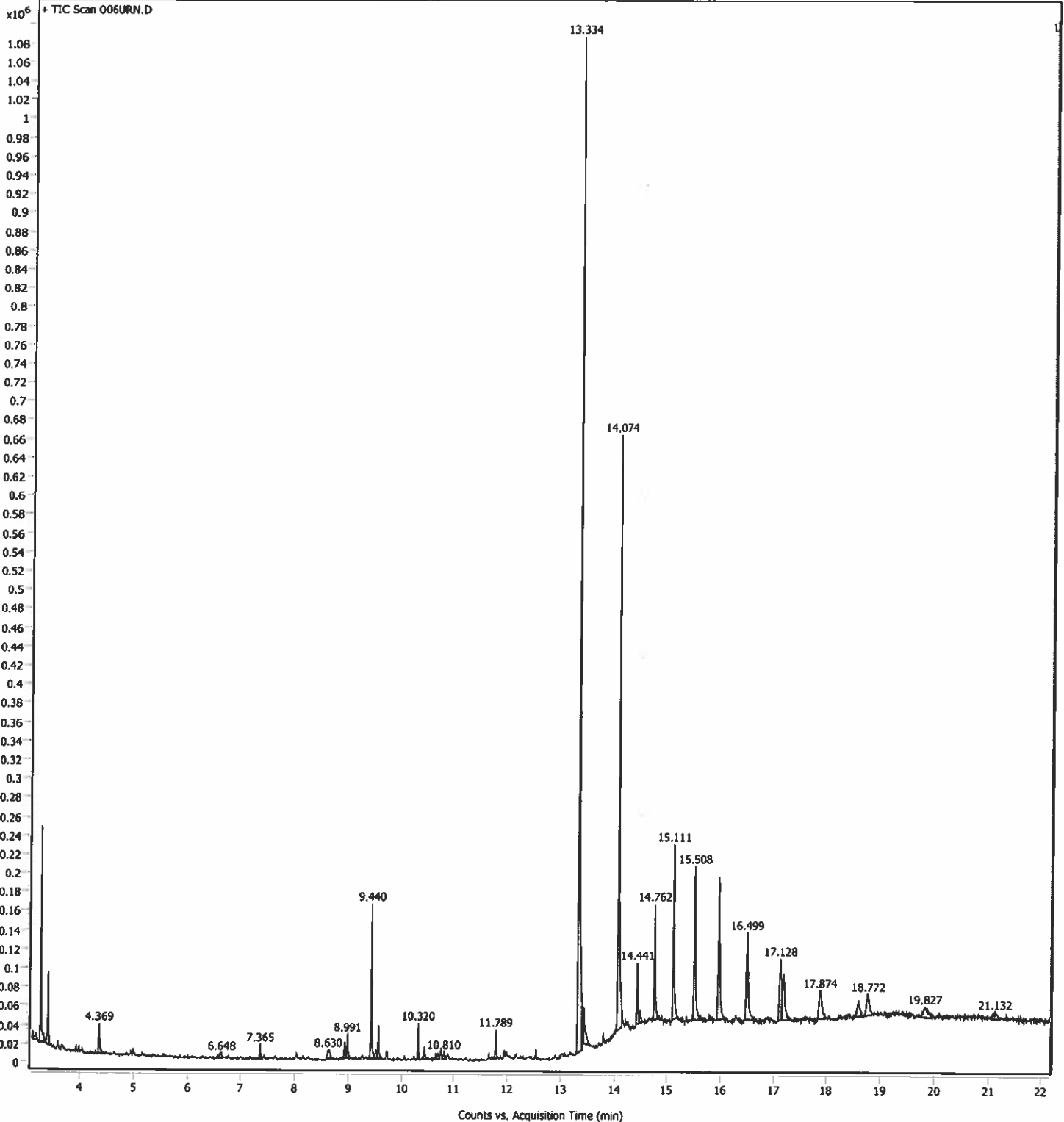


# Analysis Report

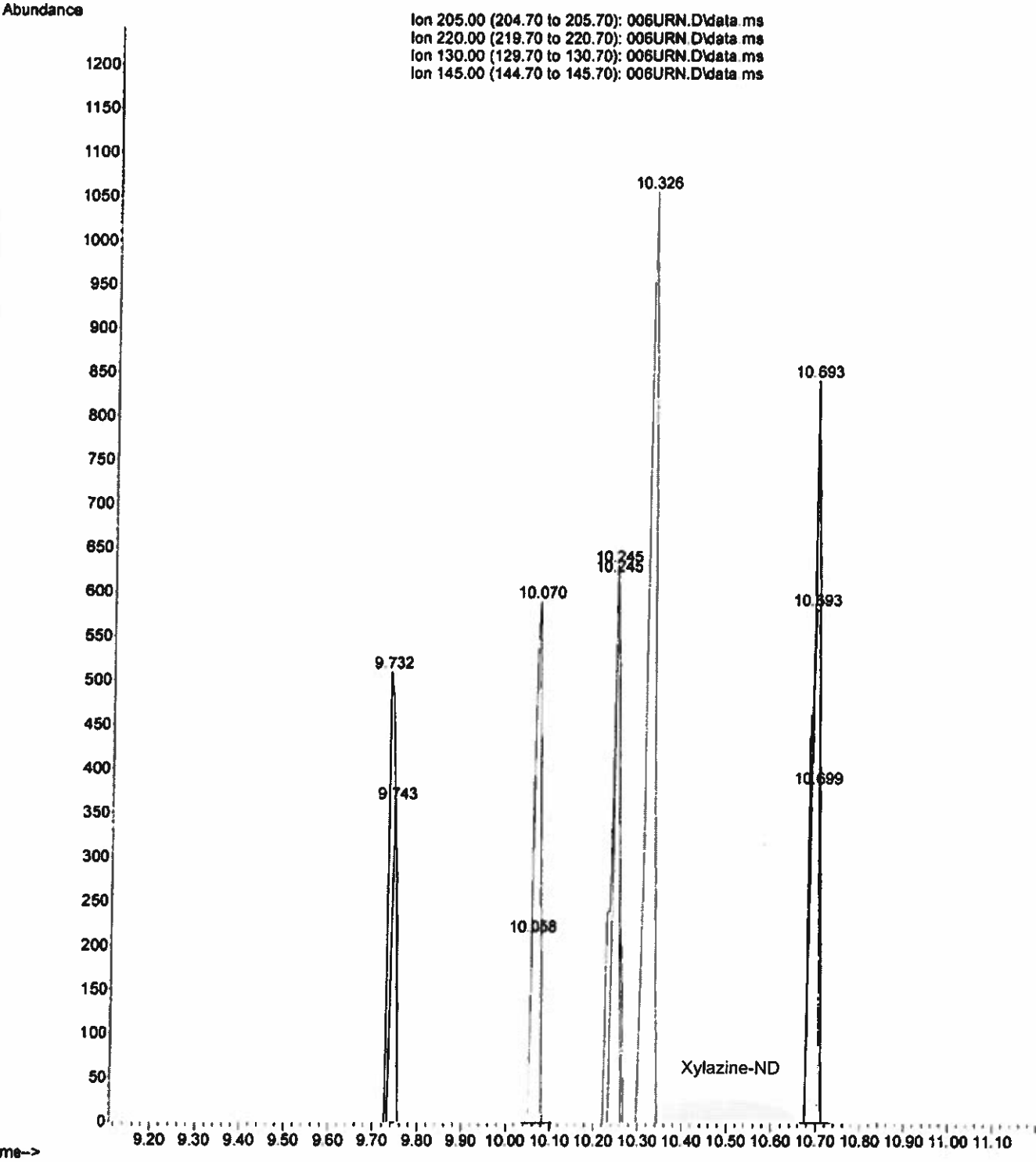
## Sample Information

Sample Name	MATRIX BLANK 6 URN	Data File Path	C:\MassHunter\GCMS\1\data\BASE5\092321\006URN.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/23/2021 2:43:31 PM (UTC-04:00)
Position	47	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms

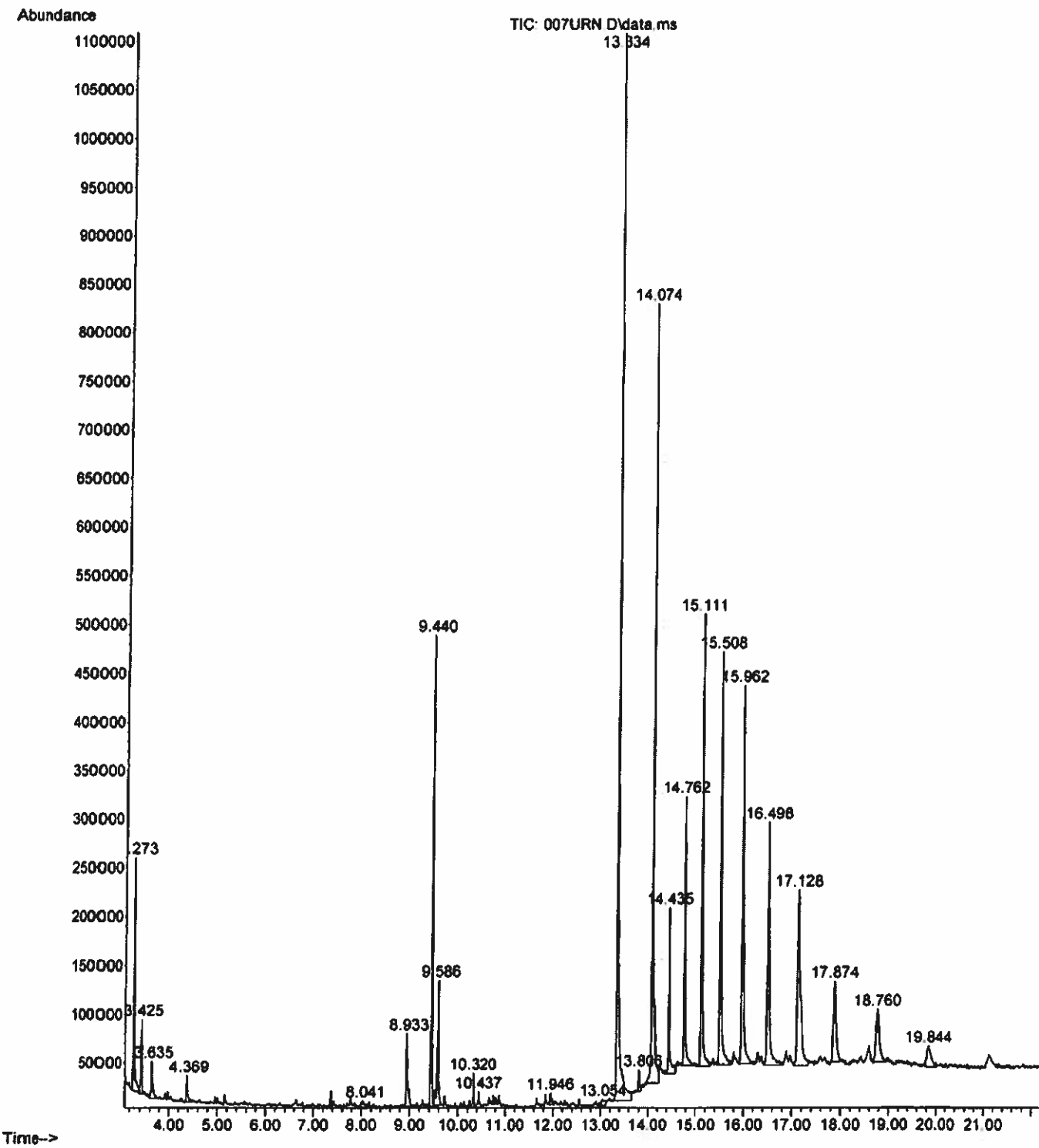


File :C:\Users\TOX\Desktop\092321\006URN.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 14:43 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 6 URN  
Misc Info : BUR101017  
Vial Number: 47

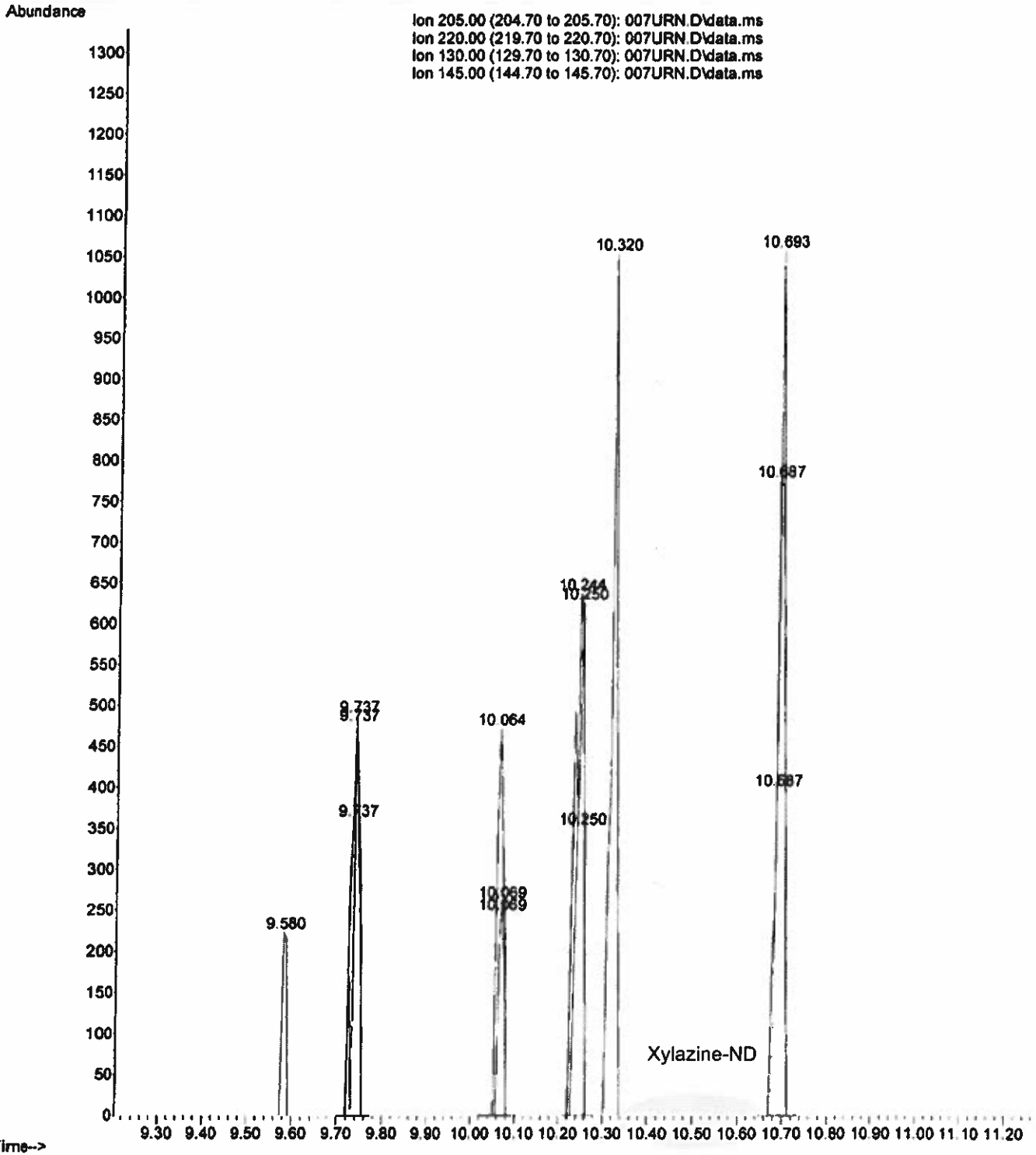




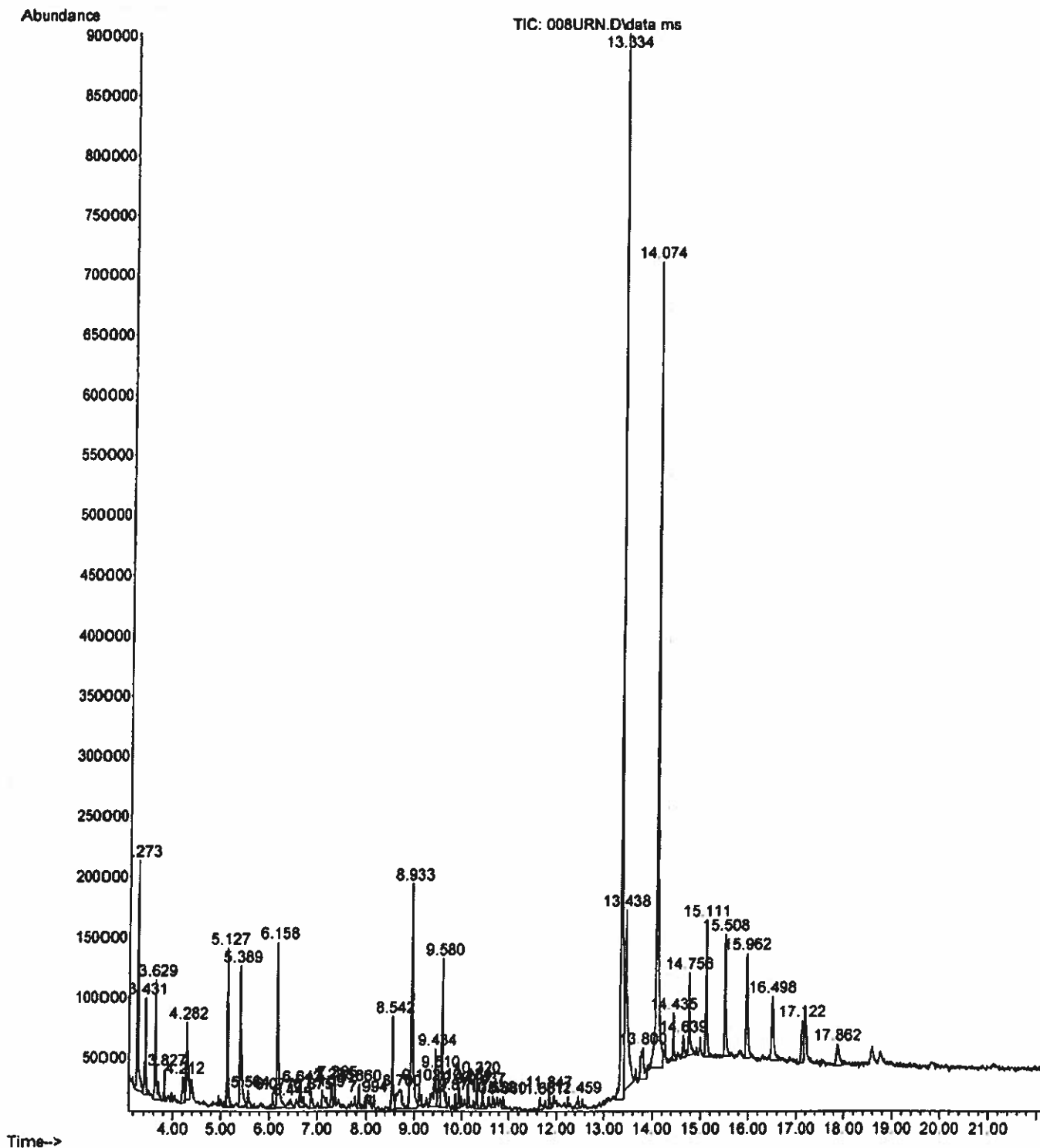
File :C:\Users\TOX\Desktop\092321\007URN.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 15:09 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 7 URN  
Misc Info : BU06017  
Vial Number: 48



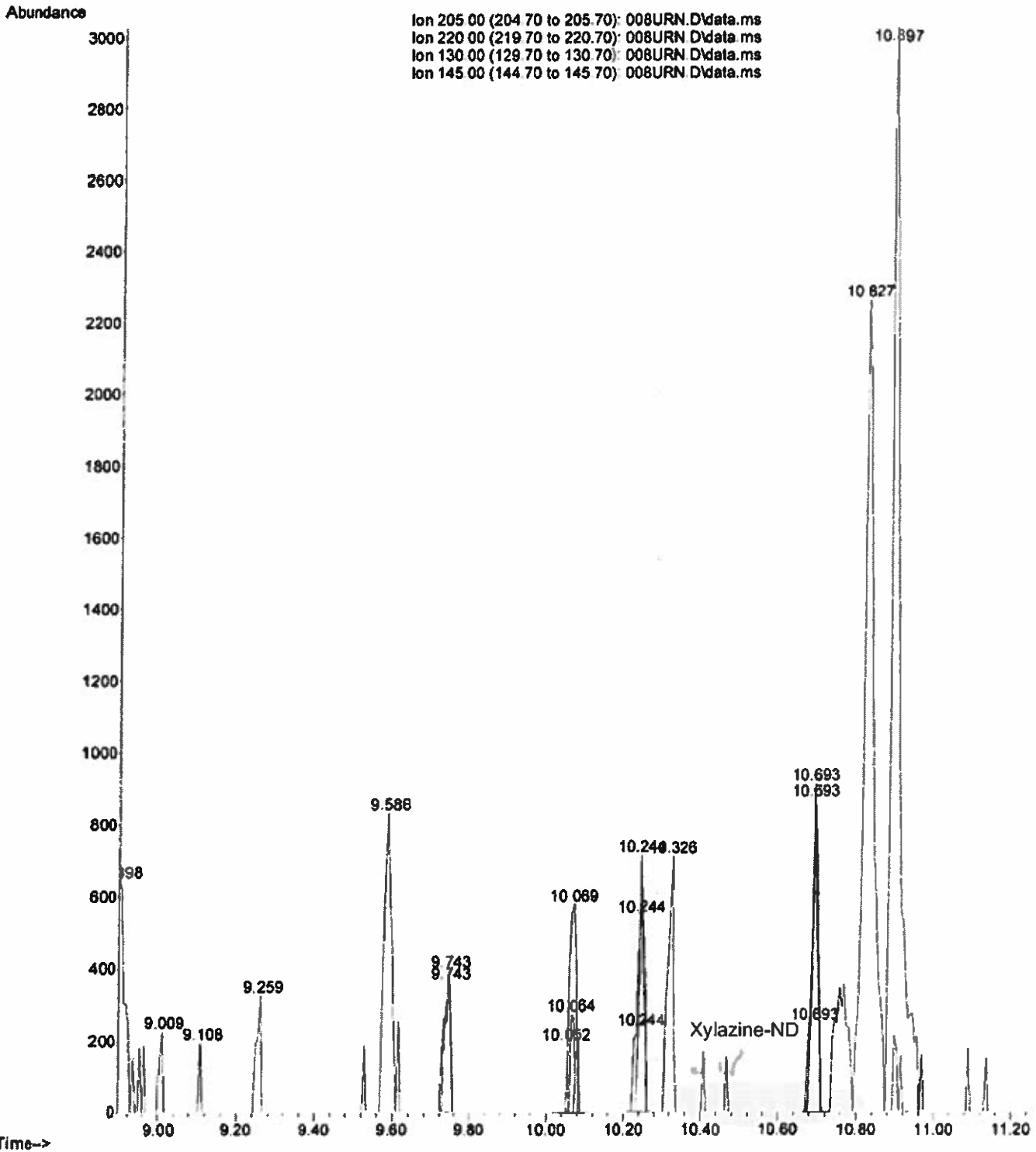
File : C:\Users\TOX\Desktop\092321\007URN.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 15:09 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 7 URN  
Misc Info : BU06017  
Vial Number: 48



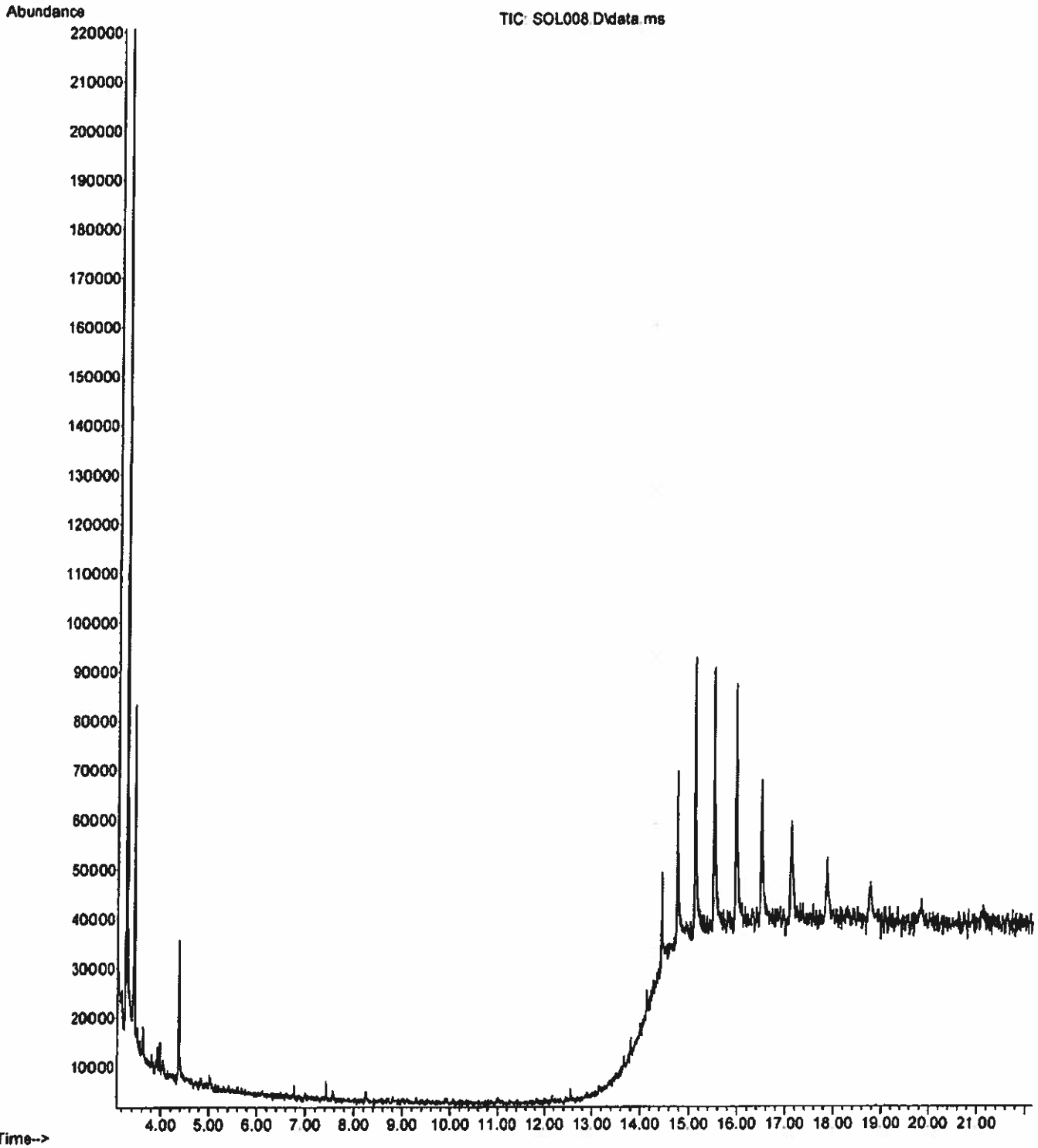
File : C:\Users\TOX\Desktop\092321\008URN.D  
 Operator : KM/QC DATA IN XYLAZINE VALIDATION  
 Acquired : 23 Sep 2021 15:35 using AcqMethod ALKALI.M  
 Instrument : #3 - Enhanced  
 Sample Name: MATRIX BLANK 8 URN  
 Misc Info : BUR0819  
 Vial Number: 49



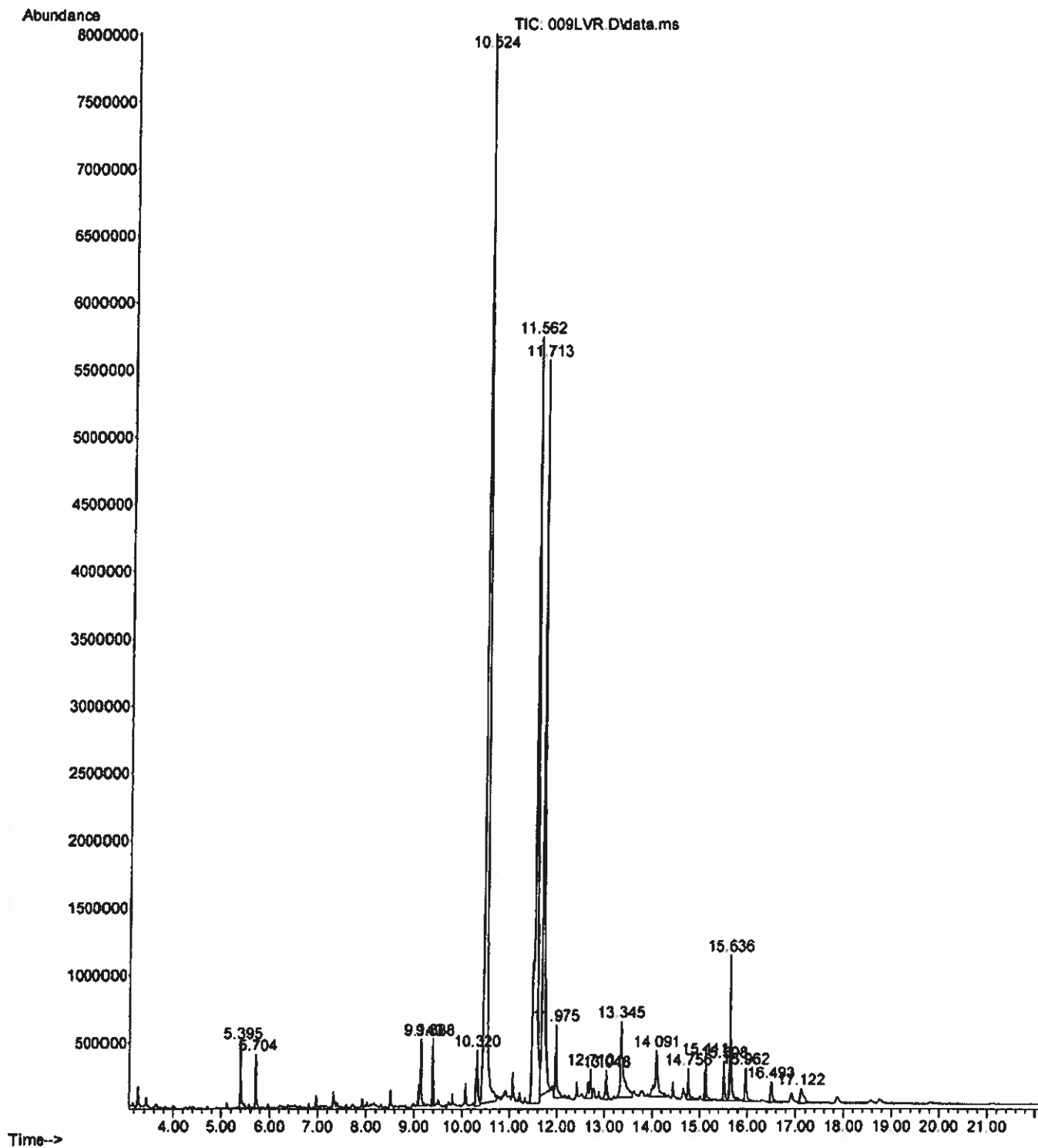
File :C:\Users\TOX\Desktop\092321\008URN.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 15:35 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 8 URN  
Misc Info : BUR0819  
Vial Number: 49



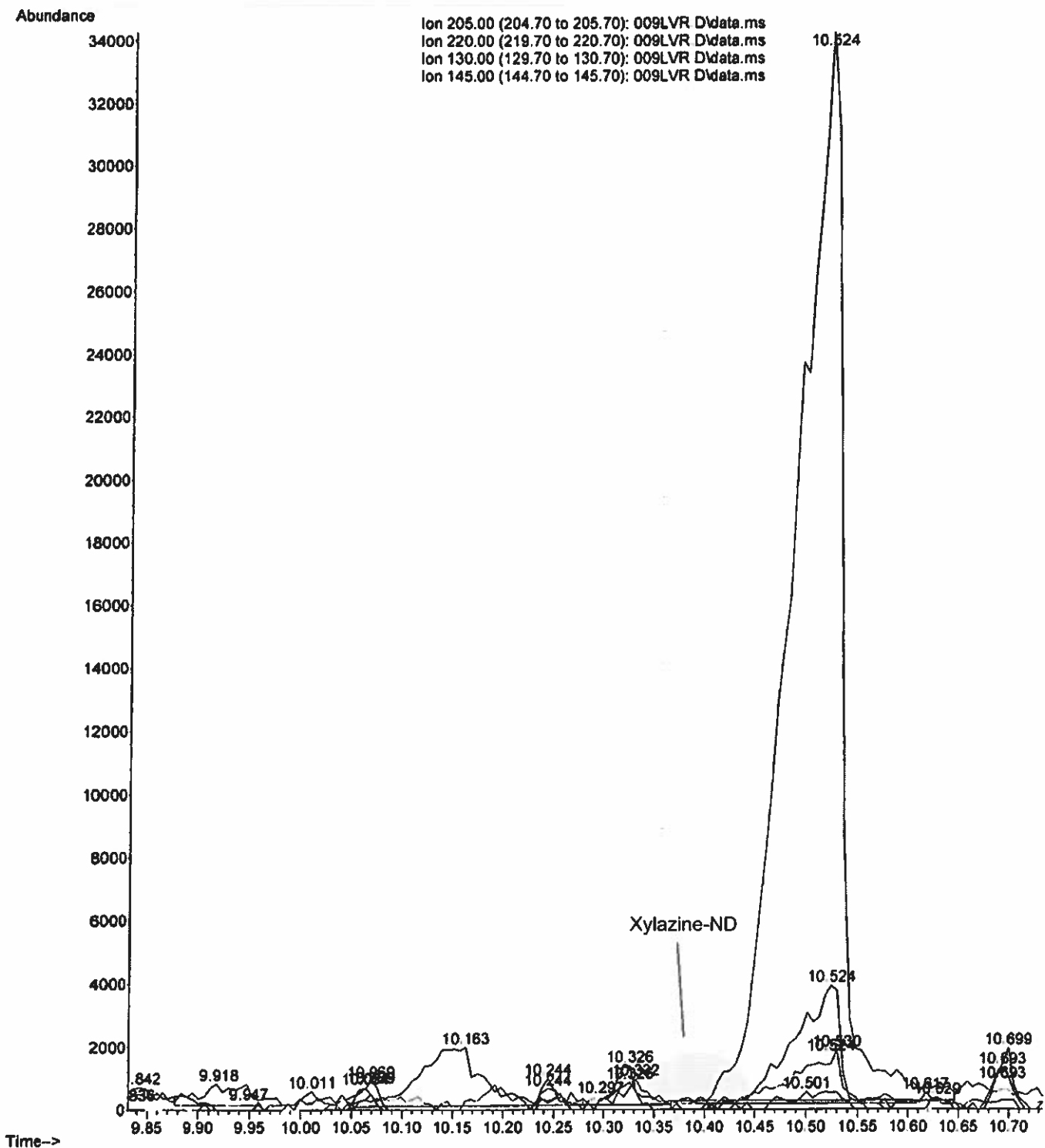
File :C:\Users\TOX\Desktop\092321\SOL008.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 16:01 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: SOLVENT AFTER MATRIX 8  
Misc Info :  
Vial Number: 50



File :C:\Users\TOX\Desktop\092321\009LVR.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 16:26 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 9 LVR  
Misc Info : LIV0109  
Vial Number: 51



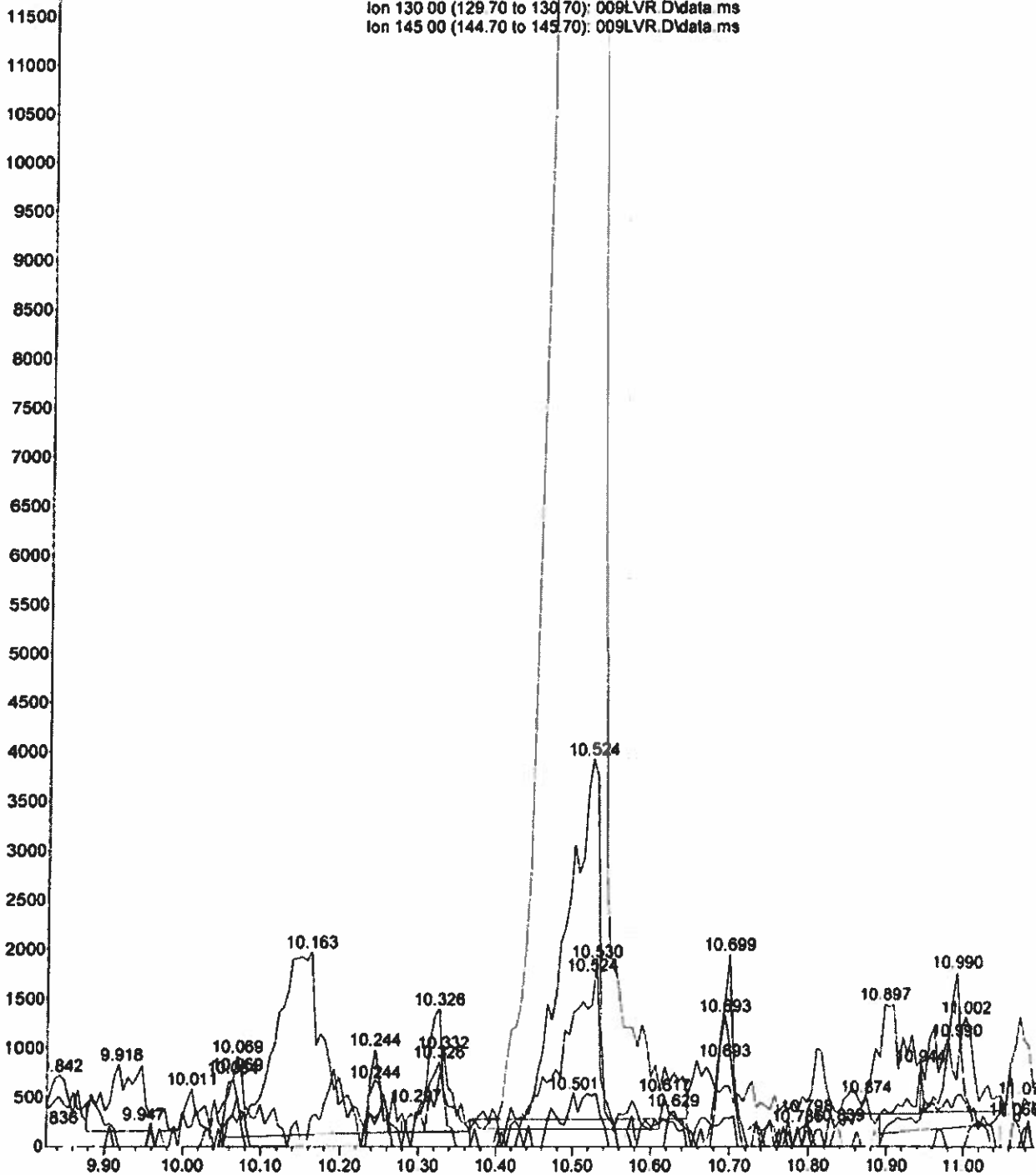
File : C:\Users\TOX\Desktop\092321\009LVR.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 16:26 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 9 LVR  
Misc Info : LIV0109  
Vial Number: 51



File : C:\Users\TOX\Desktop\092321\009LVR.D  
 Operator : KM/QC DATA IN XYLAZINE VALIDATION  
 Acquired : 23 Sep 2021 16:26 using AcqMethod ALKALI.M  
 Instrument : #3 - Enhanced  
 Sample Name: MATRIX BLANK 9 LVR  
 Misc Info : LIV0109  
 Vial Number: 51

Abundance

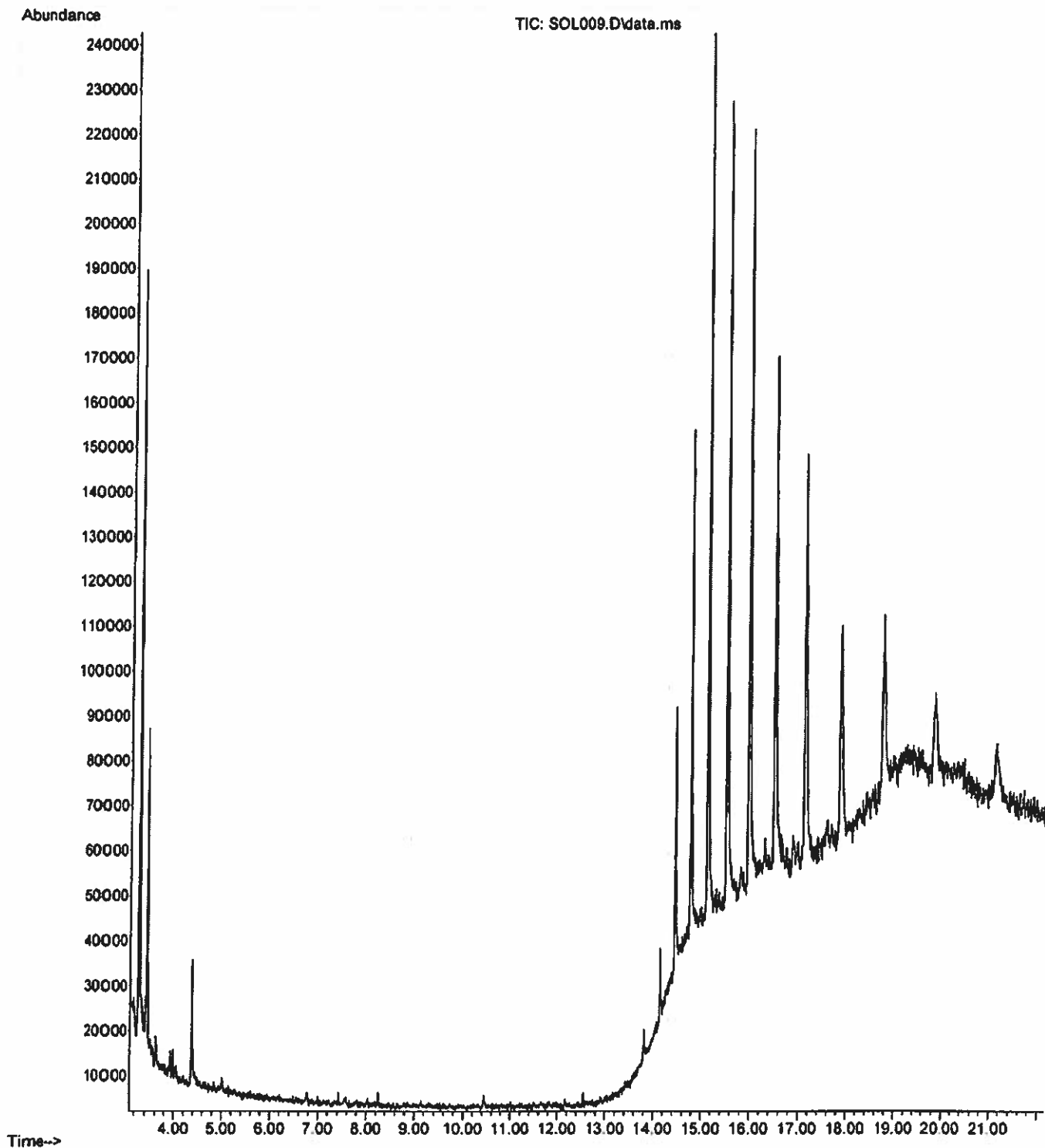
Ion 205.00 (204.70 to 205.70): 009LVR.D\data.ms  
 Ion 220.00 (219.70 to 220.70): 009LVR.D\data.ms  
 Ion 130.00 (129.70 to 130.70): 009LVR.D\data.ms  
 Ion 145.00 (144.70 to 145.70): 009LVR.D\data.ms



Time-->



File :C:\Users\TOX\Desktop\092321\SOL009.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 16:52 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: SOLVENT AFTER MATRIX 9  
Misc Info :  
Vial Number: 52

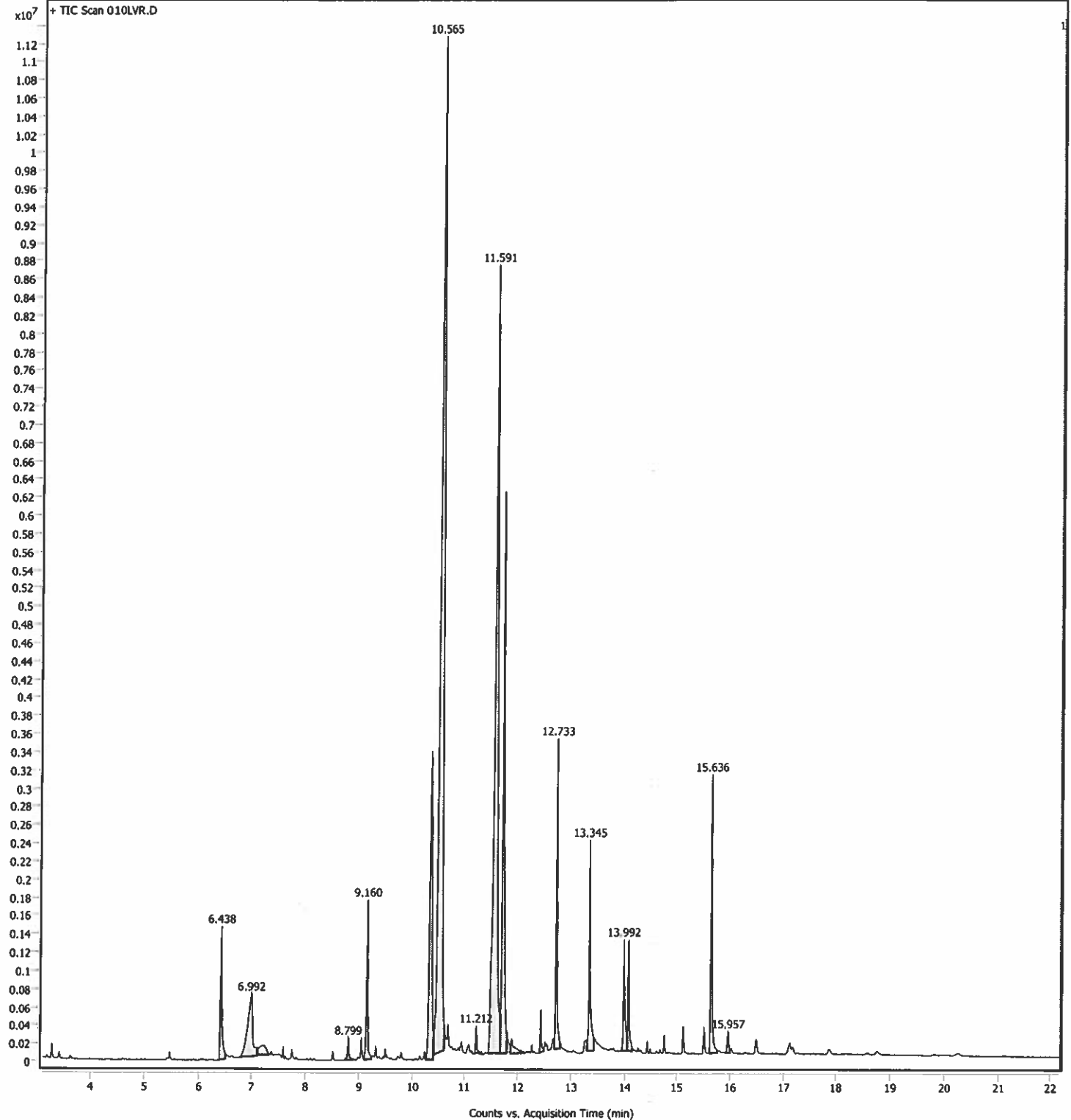


# Analysis Report

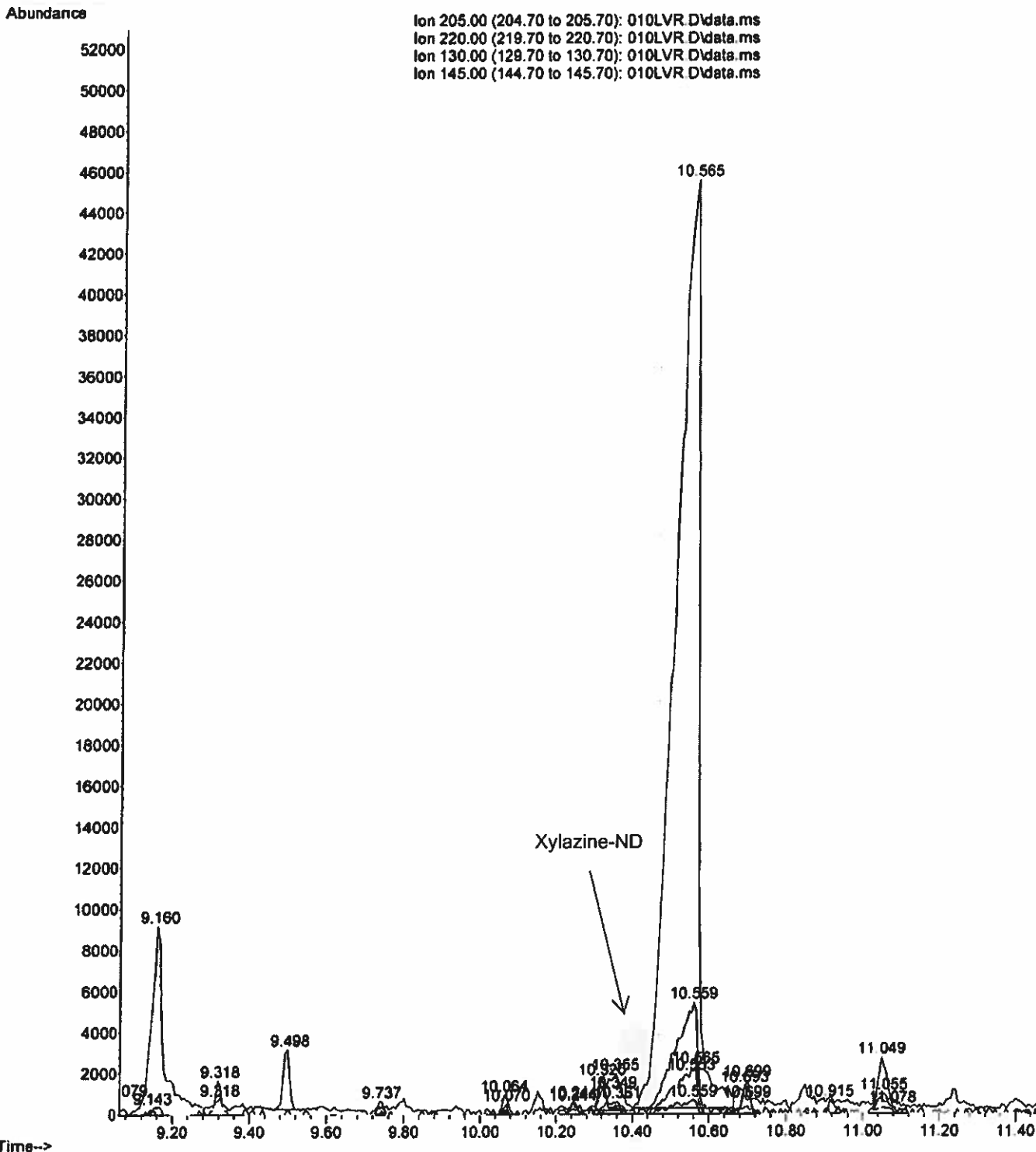
## Sample Information

Sample Name	MATRIX BLANK 10 LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092321\010LVR.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/23/2021 5:18:30 PM (UTC-04:00)
Position	53	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



File : C:\Users\TOX\Desktop\092321\010LVR.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 23 Sep 2021 17:18 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK 10 LVR  
Misc Info : LIV09231  
Vial Number: 53



Ion 205.00 (204.70 to 205.70): 010LVR.D\data.ms  
Ion 220.00 (219.70 to 220.70): 010LVR.D\data.ms  
Ion 130.00 (129.70 to 130.70): 010LVR.D\data.ms  
Ion 145.00 (144.70 to 145.70): 010LVR.D\data.ms

Time-->

**ASSAY:**

Base Screen

QC Data File: Xylazine Validation

**Analyst(s):**

K. Meinweiser

Sample / FS #	Item #	Sample Info
1 MATRIX 2 W/O ISTD	BLD	2.0 mg/L - 400uL of 0.01 mg/mL stock
2 MATRIX 1 W/O ISTD	BLD	
3 MATRIX 6 W/O ISTD	URN	
4 MATRIX 10 W/O ISTD	LVR	*satisfies 2.8.1.1.2
5 MATRIX 2 W/ ISTD ONLY	BLD	
6 MATRIX 1 W/ ISTD ONLY	BLD	
7 MATRIX 6 W/ ISTD ONLY	URN	
8 MATRIX 10 W/ ISTD ONLY	LVR	*satisfies 2.8.1.1.3
9 MATRIX 2 W ISTD	BLD	0.010 mg/L - 20uL of 0.001 mg/mL stock
10 MATRIX 1 W ISTD	BLD	
11 MATRIX 6 W ISTD	URN	
12 MATRIX 10 W ISTD	LVR	
13 MATRIX 2 W ISTD	BLD	0.10 mg/L - 20uL of 0.01 mg/mL stock
14 MATRIX 1 W ISTD	BLD	
15 MATRIX 6 W ISTD	URN	
16 MATRIX 10 W ISTD	LVR	
17 MATRIX 2 W ISTD	BLD	2.0 mg/L - 400uL of 0.01 mg/mL stock
18 MATRIX 1 W ISTD	BLD	
19 MATRIX 6 W ISTD	URN	
20 MATRIX 10 W ISTD	LVR	
21 MATRIX 2 LOD	BLD	0.005 mg/L - 10uL of 0.001 mg/mL stock
22 MATRIX 1 LOD	BLD	
23 MATRIX 6 LOD	URN	
24 MATRIX 10 LOD	LVR	*satisfies 2.8.1.1.6
25		

Specimens removed from TX ADM storage, aliquots removed for analysis, specimens returned to TX ADM storage by, unless otherwise noted:

**Analyst:** Andy Jones

**Date:** 9.28.2021

**Time Removed:** N/A **Time Returned:** N/A

Sample / FS #	Item #	Sample Info
26		
27		
28		
29		
30		
31		
32		
33		
34		
35		
36		
37		
38		
39		
40		
41		
42		
43		
44		
45		
46		
47		
48		
49		
50		

Extraction: LL / SPE SPE lot #: 024170-XD

Control Charts Updated (Initials/Date): N/A

Reviewer / Date: \_\_\_\_\_



## Sequence Table (Front Injector):

vials loaded by AJ

Line	Location	SampleName	Method Name	Num Inj	SampleType
1	Vial 1	ETOAC	ALKALI	1	Sample
2	Vial 2	MATRIX 1 0.10 mg/L - BLD	ALKALI	1	Sample
3	Vial 3	MATRIX 2 0.10 mg/L - BLD	ALKALI	1	Sample
4	Vial 4	MATRIX 6 0.10 mg/L - URN	ALKALI	1	Sample
5	Vial 5	MATRIX BLANK after 0.1 mg/L	ALKALI	1	Sample
6	Vial 6	MATRIX 1 0.50 mg/L - BLD	ALKALI	1	Sample
7	Vial 7	MATRIX 2 0.50 mg/L - BLD	ALKALI	1	Sample
8	Vial 8	MATRIX 6 0.50 mg/L - URN	ALKALI	1	Sample
9	Vial 9	MATRIX BLANK after 0.5 mg/L	ALKALI	1	Sample
10	Vial 10	MATRIX 1 1.0 mg/L - BLD	ALKALI	1	Sample
11	Vial 11	MATRIX 2 1.0 mg/L - BLD	ALKALI	1	Sample
12	Vial 12	MATRIX 6 1.0 mg/L - URN	ALKALI	1	Sample
13	Vial 13	MATRIX BLANK after 1.0 mg/L	ALKALI	1	Sample
14	Vial 14	MATRIX 1 2.0 mg/L - BLD	ALKALI	1	Sample
15	Vial 15	MATRIX 2 2.0 mg/L - BLD	ALKALI	1	Sample
16	Vial 16	MATRIX 6 2.0 mg/L - URN	ALKALI	1	Sample
17	Vial 17	MATRIX BLANK after 2.0 mg/L	ALKALI	1	Sample
18	Vial 19	MATRIX 1 W/ SW CTL - BLD	ALKALI	1	Sample
19	Vial 20	MATRIX 2 W/ SW CTL - BLD	ALKALI	1	Sample
20	Vial 21	MATRIX 6 W/ SW CTL - URN	ALKALI	1	Sample
21	Vial 22	MATRIX 10 W/ 0.1mg/L CAL - ...	ALKALI	1	Sample
22	Vial 23	SOLVENT BLANK	ALKALI	1	Sample
23	Vial 24	MATRIX 1 W/ IH CTL - BLD	ALKALI	1	Sample
24	Vial 25	MATRIX 2 W/ IH CTL - BLD	ALKALI	1	Sample
25	Vial 26	MATRIX 6 W/ IH CTL - URN	ALKALI	1	Sample
26	Vial 27	MATRIX 10 W/ IH CTL - LVR	ALKALI	1	Sample
27	Vial 28	SOLVENT BLANK 1	ALKALI	1	Sample
28	Vial 29	MATRIX 1 W/ SW CTL + XYLAZ...	ALKALI	1	Sample
29	Vial 30	MATRIX 2 W/ SW CTL + XYLAZ...	ALKALI	1	Sample
30	Vial 31	MATRIX 6 W/ SW CTL + XYLAZ...	ALKALI	1	Sample
31	Vial 32	MATRIX 10 W/ 0.1mg/L CAL + ...	ALKALI	1	Sample
32	Vial 33	SOLVENT BLANK 2	ALKALI	1	Sample
33	Vial 34	MATRIX 1 W/ IH CTL + XYLAZ...	ALKALI	1	Sample
34	Vial 35	MATRIX 2 W/ IH CTL + XYLAZ...	ALKALI	1	Sample
35	Vial 36	MATRIX 6 W/ IH CTL + XYLAZ...	ALKALI	1	Sample
36	Vial 37	MATRIX 10 W/ IH CTL + XYLA...	ALKALI	1	Sample
37	Vial 38	SOLVENT BLANK 3	ALKALI	1	Sample
38	Vial 39	SOLVENT BLANK 4	ALKALI	1	Sample
39	Vial 40	MATRIX 1 LOD - BLD	ALKALI	1	Sample
40	Vial 41	MATRIX 2 LOD - BLD	ALKALI	1	Sample
41	Vial 42	MATRIX 6 LOD - URN	ALKALI	1	Sample
42	Vial 43	MATRIX 10 LOD - LVR	ALKALI	1	Sample
43	Vial 44	MATRIX 1 W/O ISTD - BLD	ALKALI	1	Sample
44	Vial 45	MATRIX 2 W/O ISTD - BLD	ALKALI	1	Sample
45	Vial 46	MATRIX 6 W/O ISTD - URN	ALKALI	1	Sample
46	Vial 47	MATRIX 10 W/O ISTD - LVR	ALKALI	1	Sample
47	Vial 48	SOLVENT BLANK 5	ALKALI	1	Sample
48	Vial 49	MATRIX 1 W/ISTD ONLY - BLD	ALKALI	1	Sample
49	Vial 50	MATRIX 2 W/ISTD ONLY - BLD	ALKALI	1	Sample
50	Vial 81	MATRIX 6 W/ISTD ONLY- URN	ALKALI	1	Sample
51	Vial 82	MATRIX 10 W/ISTD ONLY- LVR	ALKALI	1	Sample
52	Vial 83	MATRIX 1 W/ISTD 0.010 mg/L...	ALKALI	1	Sample
53	Vial 84	MATRIX 2 W/ISTD 0.010 mg/L...	ALKALI	1	Sample
54	Vial 85	MATRIX 6 W/ISTD 0.010 mg/L...	ALKALI	1	Sample
55	Vial 86	MATRIX 10 W/ISTD 0.010 mg/...	ALKALI	1	Sample

vial check  
KM 9/29/21

56	Vial 87	MATRIX 1 W/ISTD 0.10 mg/L ...	ALKALI	1	Sample
57	Vial 88	MATRIX 2 W/ISTD 0.10 mg/L ...	ALKALI	1	Sample
58	Vial 89	MATRIX 6 W/ISTD 0.10 mg/L ...	ALKALI	1	Sample
59	Vial 90	MATRIX 10 W/ISTD 0.10 mg/L...	ALKALI	1	Sample
60	Vial 91	MATRIX 1 W/ISTD 0.010 <sup>20</sup> mg/L...	ALKALI	1	Sample
61	Vial 92	MATRIX 2 W/ISTD 2.0 <sup>20</sup> mg/L...	ALKALI	1	Sample
62	Vial 93	MATRIX 6 W/ISTD 2.0 mg/L -...	ALKALI	1	Sample
63	Vial 94	MATRIX 10 W/ISTD 2.0 mg/L-...	ALKALI	1	Sample
64	Vial 95	SOLVENT BLANK 6	ALKALI	1	Sample
65	Vial 96	OFF	ALKALI	1	Sample

vial check 9/29/21

Sequence Name: C:\MassHunter\GCMS\1\sequence\alkalival.sequence.xml

Comment: Base

Operator: KM/QC DATA IN XYLAZINE VALIDATION **KM**

Data Path: C:\MassHunter\GCMS\1\data\BASES\092821\

Instrument Control Pre-Seq Cmd:

Data Analysis Pre-Seq Cmd:

Instrument Control Post-Seq Cmd:

Data Analysis Post-Seq Cmd:

Method Sections To Run

(X) Full Method

( ) Reprocessing Only

Sequence Barcode Options

( ) On Mismatch, Inject Anyway

( ) On Mismatch, Don't Inject

(X) Barcode Disabled

42) Sample	67	036	ALKALI	MATRIX 10 LOD - LVR
43) Sample	68	037	ALKALI	MATRIX 1 W/O ISTD - BLD
44) Sample	69	038	ALKALI	MATRIX 2 W/O ISTD - BLD
45) Sample	70	039	ALKALI	MATRIX 6 W/O ISTD - URN
46) Sample	71	040	ALKALI	MATRIX 10 W/O ISTD - LVR
47) Sample	72	SOL5	ALKALI	Solvent Blank 5
48) Sample	73	041	ALKALI	MATRIX 1 W/ISTD ONLY - BLD
49) Sample	74	042	ALKALI	MATRIX 2 W/ISTD ONLY - BLD
50) Sample	75	043	ALKALI	MATRIX 6 W/ISTD ONLY- URN
51) Sample	76	044	ALKALI	MATRIX 10 W/ISTD ONLY- LVR
52) Sample	77	045	ALKALI	MATRIX 1 W/ISTD 0.010 mg/L -
53) Sample	78	046	ALKALI	MATRIX 2/W ISTD 0.010 mg/L- <b>BLD*</b>
54) Sample	79	047	ALKALI	MATRIX 6 W/ISTD 0.010 mg/L- <b>UR*</b>
55) Sample	80	048	ALKALI	MATRIX 10 W/ISTD 0.010 mg/L-
56) Sample	81	049	ALKALI	MATRIX 1 W/ISTD 0.10 mg/L - <b>BLD*</b>
57) Sample	82	050	ALKALI	MATRIX 2/W ISTD 0.10 mg/L- <b>BLD*</b>
58) Sample	83	051	ALKALI	MATRIX 6 W/ISTD 0.10 mg/L- <b>UR</b>
59) Sample	84	052	ALKALI	MATRIX 10 W/ISTD 0.10 mg/L- <b>LVR*</b>
60) Sample	85	053	ALKALI	MATRIX 1 W/ISTD 2.0 mg/L - <b>BLD*</b>
61) Sample	86	054	ALKALI	MATRIX 2/W ISTD 2.0 mg/L- <b>BLD</b>
62) Sample	87	055	ALKALI	MATRIX 6 W/ISTD 2.0 mg/L- <b>URN</b>
63) Sample	88	056	ALKALI	MATRIX 10 W/ISTD 2.0 mg/L- <b>LVR*</b>
64) Sample	89	SOL6	ALKALI	Solvent Blank 6

**KM 9/29/21**

**vial check KM 9/29/21**



=====  
Calibration Table  
=====

-----  
General Calibration Setting  
-----

Calib. Data Modified : Thursday, November 17, 2022 12:05:52 PM

Rel. Reference Window : 2.000 %  
Abs. Reference Window : 0.000 min  
Rel. Non-ref. Window : 2.000 %  
Abs. Non-ref. Window : 0.000 min  
Uncalibrated Peaks : not reported  
Partial Calibration : Yes, identified peaks are recalibrated  
Correct All Ret. Times: No, only for identified peaks  
  
Curve Type : Linear (some peaks differ, see below)  
Origin : Ignored (some peaks differ, see below)  
Weight : Linear (Amnt) (some peaks differ, see below)

Recalibration Settings:  
Average Response : No Update  
Average Retention Time: No Update

Calibration Report Options :

Printout of recalibrations within a sequence:

Calibration Table after Recalibration

Normal Report after Recalibration

If the sequence is done with bracketing:

Results of first cycle (ending previous bracket)

Default Sample ISTD Information (if not set in sample table):

ISTD #	ISTD Amount [mg/L]	Name
1	1.00000	METHAPYRILENE

-----  
Signal Details  
-----

Signal 1: NPD1 A, Front Signal  
-----

-----  
Overview Table  
-----

RT	Sig	Lvl	Amount [mg/L]	Area	Rsp.Factor	Ref	ISTD #	Compound
11.033	1	1	0.00000	911.87921	0.00000	No	No 1	TRAMADOL
11.046	1	1	0.00000	472.34546	0.00000	No	No 1	XYLAZINE
11.278	1	1	1.00000	820.58740	1.21864e-3	No	Yes 1	METHAPYRILENE

-----  
More compound-specific settings  
-----

Compound: TRAMADOL  
Time Window : From 10.817 min To 11.254 min

Compound: XYLAZINE  
Time Window : From 10.825 min To 11.267 min

Compound: METHAPYRILENE  
Curve Type : Linear  
Origin : Included  
Calibration Level Weights:/  
Level 1 : 1  
-----

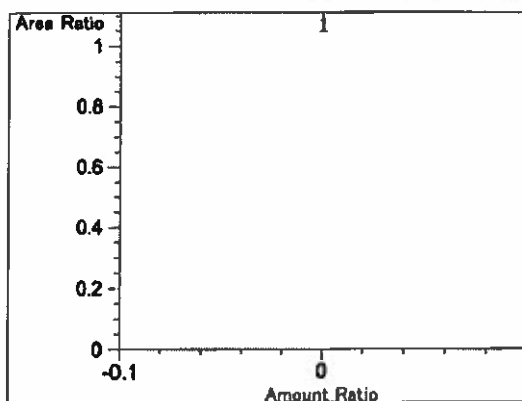
-----  
Peak Sum Table  
-----

\*\*\*No Entries in table\*\*\*  
-----

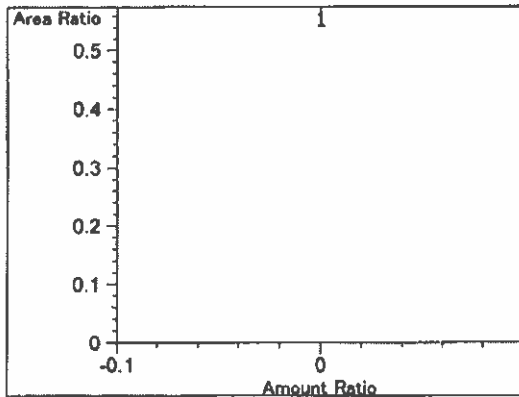
4 Warnings or Errors :

Warning : Curve requires more calibration points., (TRAMADOL)  
Warning : Overlapping peak time windows at 11.033 min, signal 1  
Warning : Overlapping peak time windows at 11.046 min, signal 1  
Warning : Curve requires more calibration points. at 11.033 min, signal 1

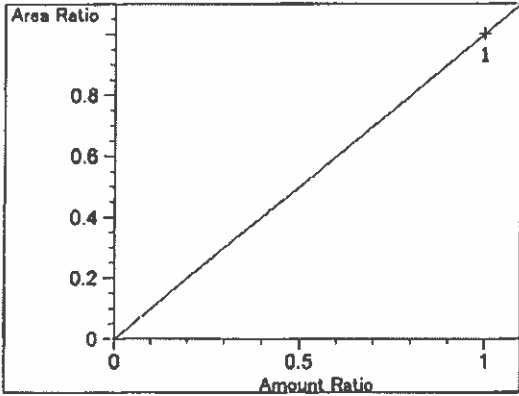
=====  
Calibration Curves  
=====



TRAMADOL at exp. RT: 11.033  
NPD1 A, Front Signal  
Correlation: 1.00000  
Residual Std. Dev.: 0.00000  
Formula:  $y = mx + b$   
m: 0.00000  
b: 0.00000  
x: Amount  
y: Area



XYLAZINE at exp. RT: 11.046  
 NPD1 A, Front Signal  
 Correlation: 1.00000  
 Residual Std. Dev.: 0.00000  
 Formula:  $y = mx + b$   
 m: 0.00000  
 b: 0.00000  
 x: Amount  
 y: Area



METHAPYRILENE at exp. RT: 11.278  
 NPD1 A, Front Signal  
 Correlation: 1.00000  
 Residual Std. Dev.: 0.00000  
 Formula:  $y = mx + b$   
 m: 1.00000  
 b: 0.00000  
 x: Amount  
 y: Area

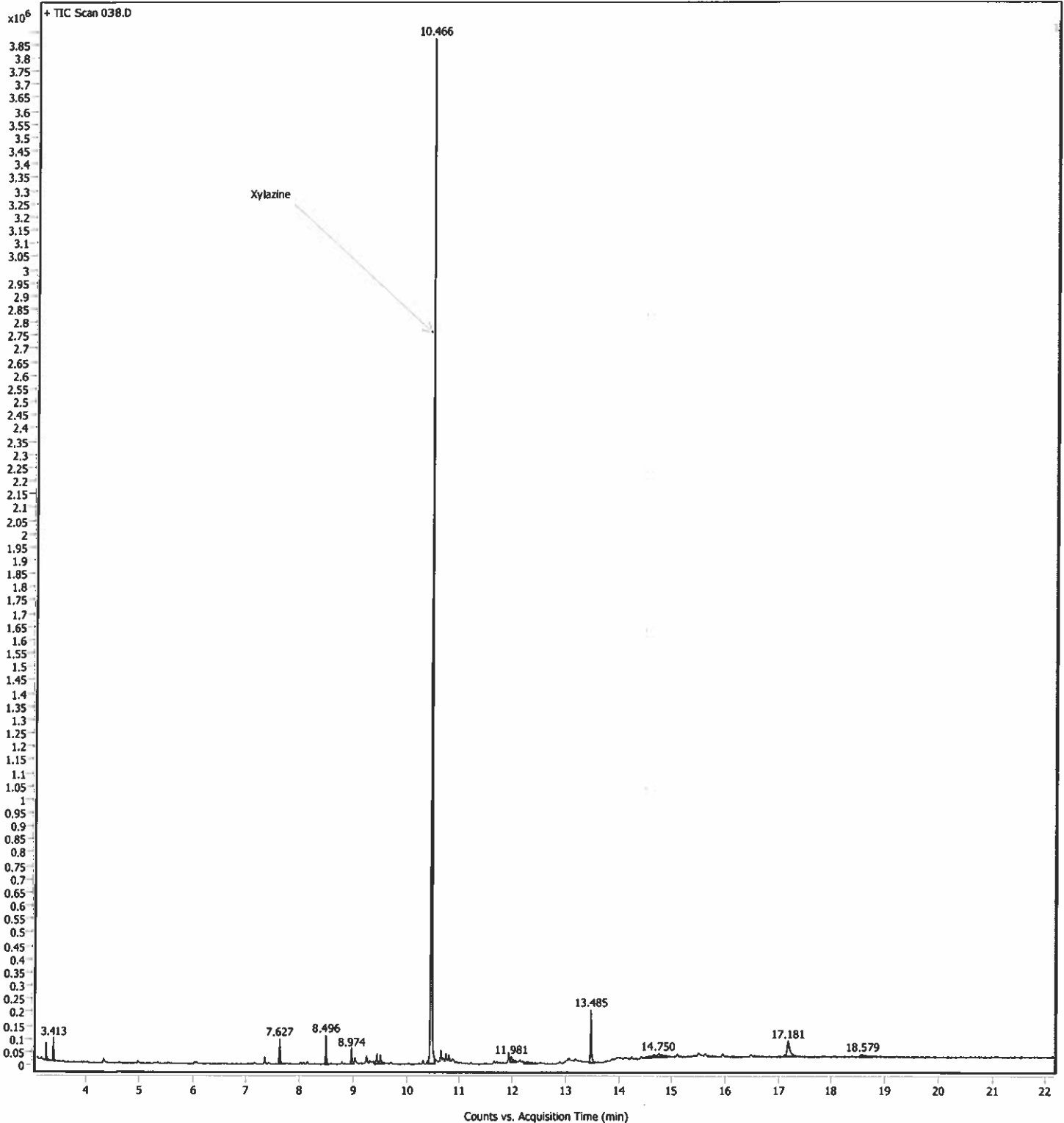


# Analysis Report

## Sample Information

Sample Name	MATRIX 2 W/O ISTD - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\038.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 7:45:35 AM (UTC-04:00)
Position	69	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

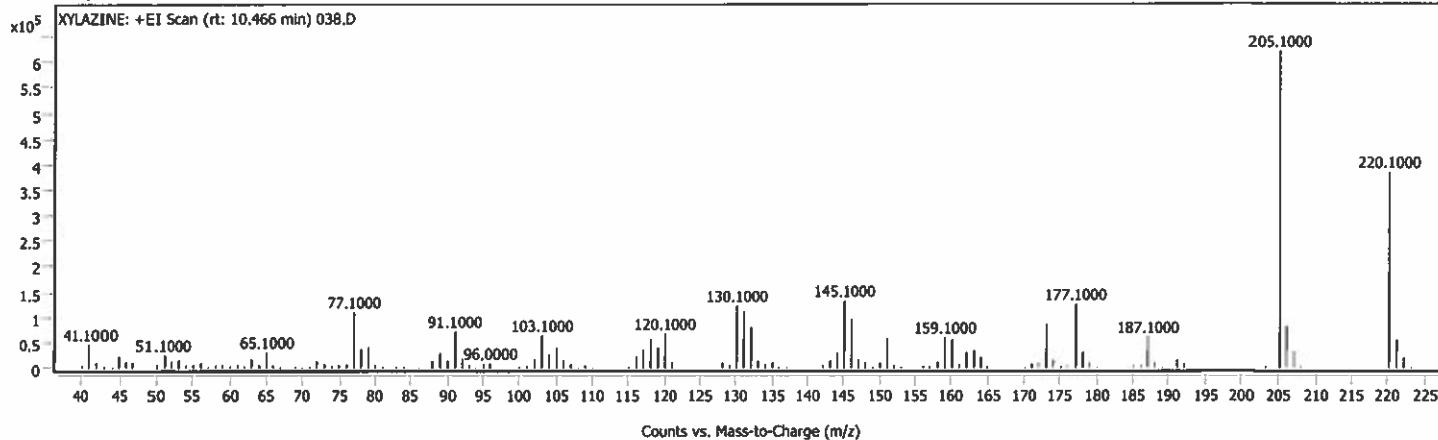
## Sample Spectra

+ Scan (rt: 10.466 min)

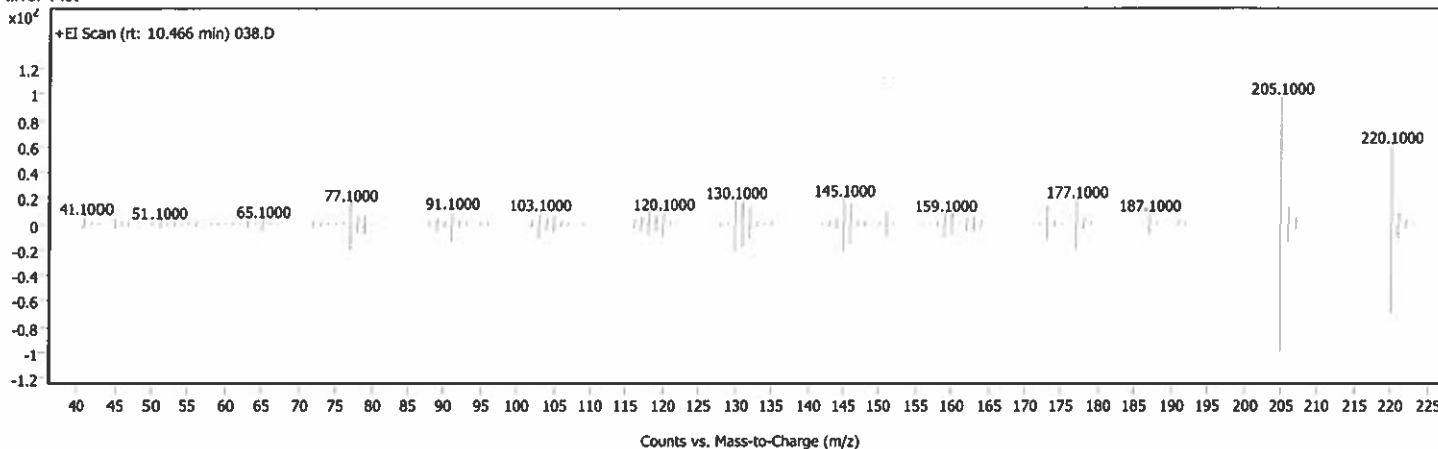
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.58	aafs.l	MATRIX 2 W/O ISTD - BLD	KM/OC DATA IN XYLAZINE VALIDATION

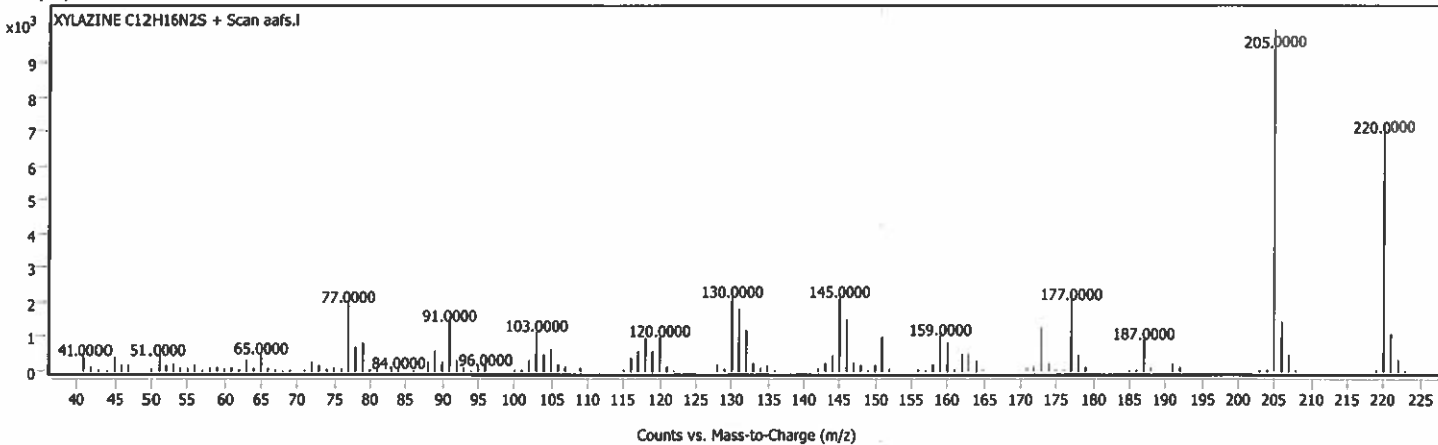
## Observed Spectrum



## Mirror Plot



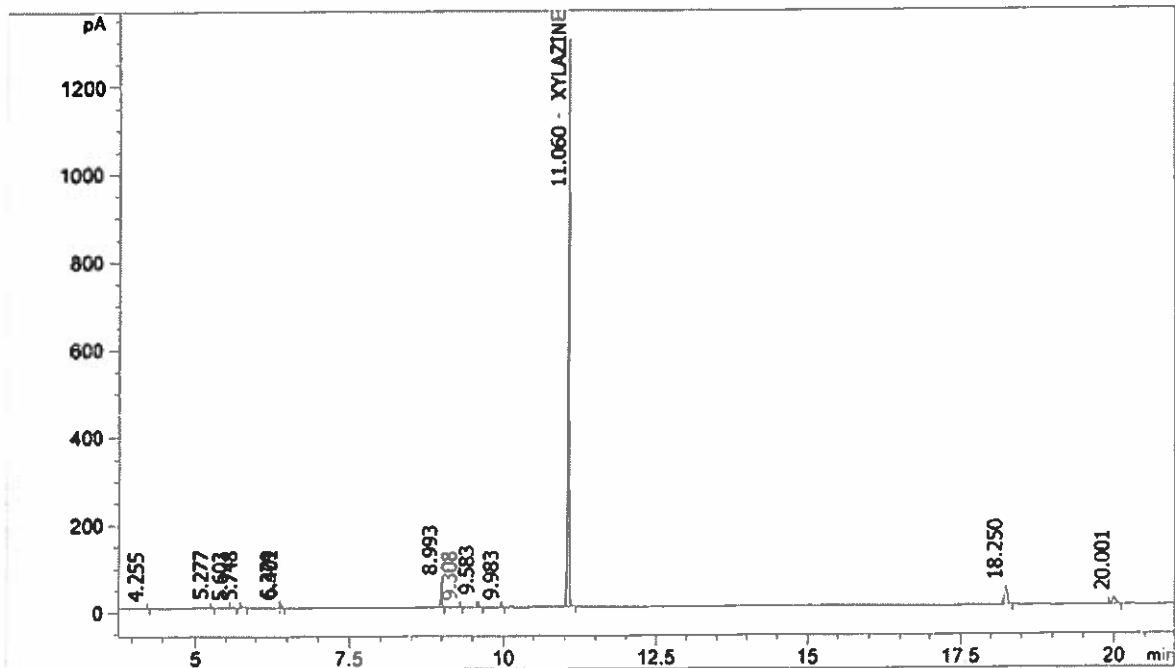
## Library Spectrum



Injection Date: 9/29/2021 7:11:18 AM  
 Sample Name: MATRIX 2 W/O ISTD - BLD  
 Sample Info:

Seq Line: 44  
 Vial 45

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



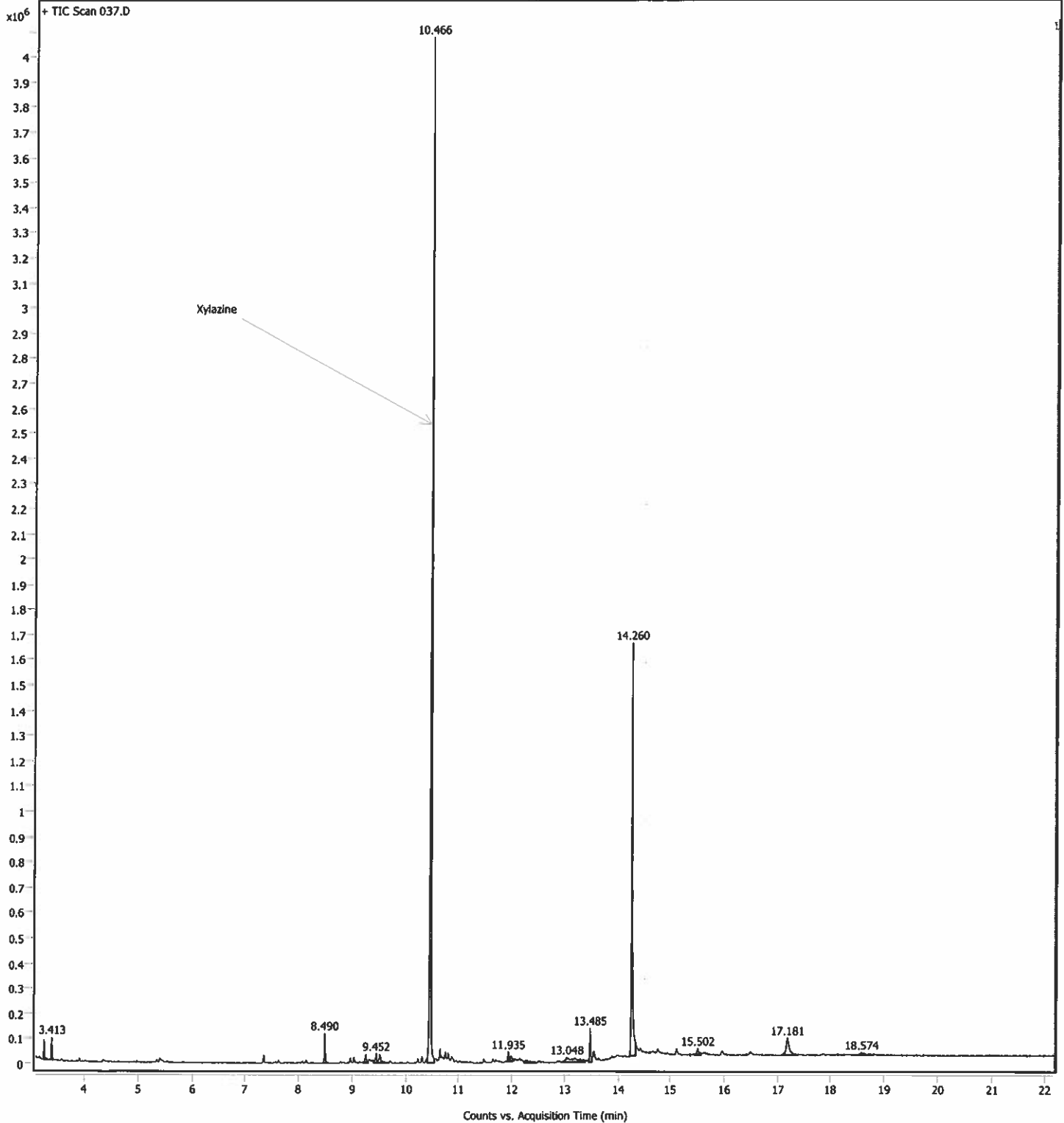
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.060	11.046	1860.743	1293.416	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 W/O ISTD - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\037.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 7:19:48 AM (UTC-04:00)
Position	68	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

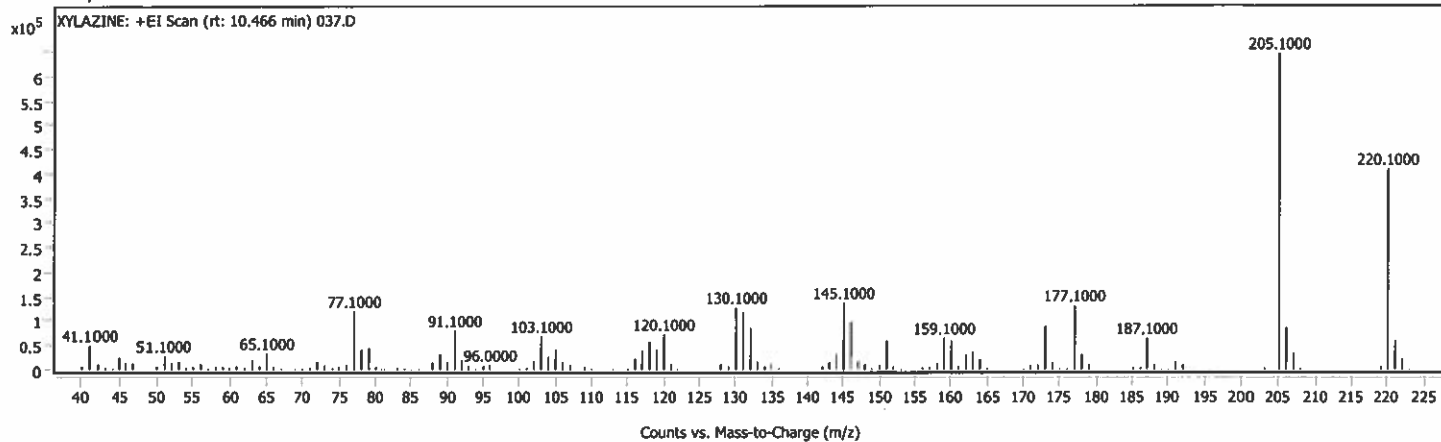
## Sample Spectra

+ Scan (rt: 10.466 min)

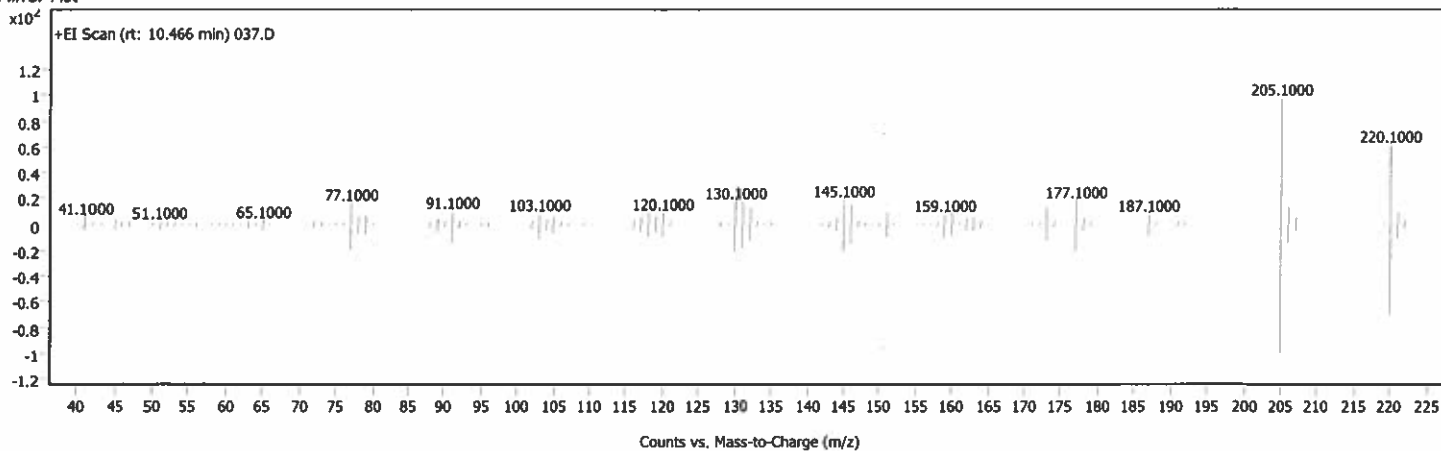
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.66	aafs.1	MATRIX 1 W/O ISTD - BLD	KM/QC DATA IN XYLAZINE VALIDATION

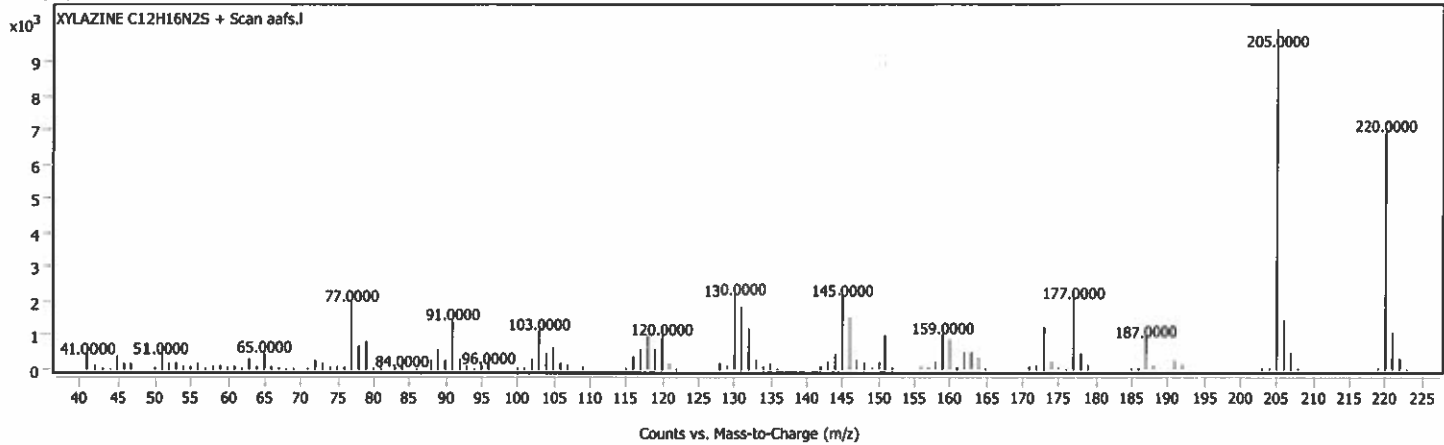
### Observed Spectrum



### Mirror Plot



### Library Spectrum

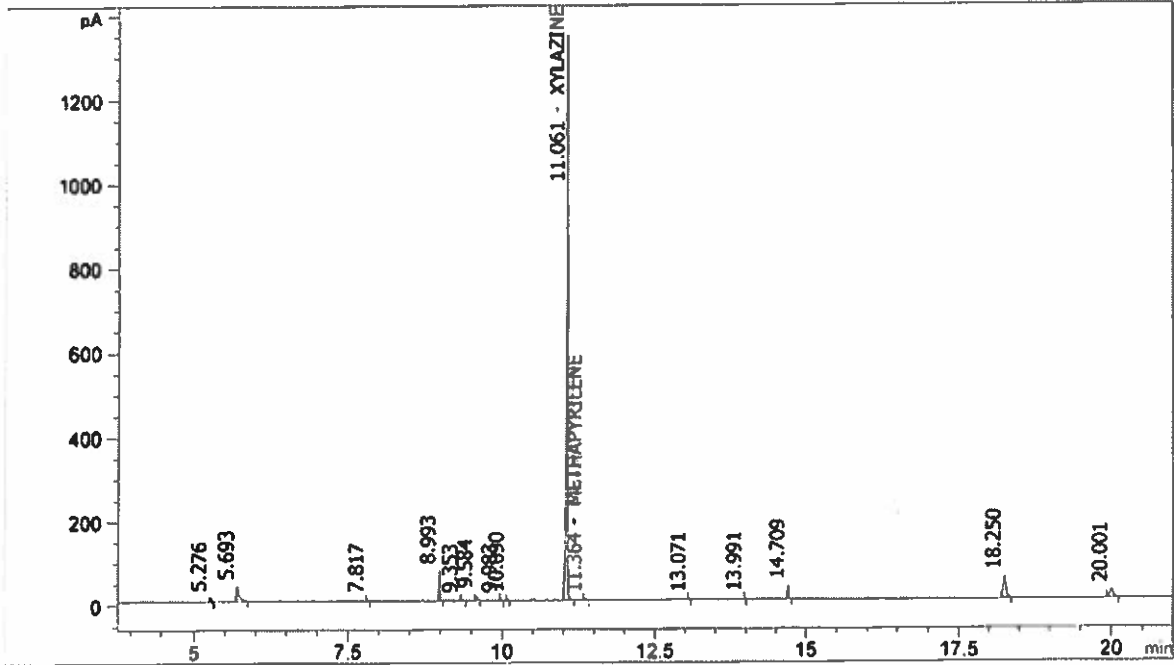




Injection Date: 9/29/2021 6:47:00 AM  
 Sample Name: MATRIX 1 W/O ISTD - BLD  
 Sample Info:

Seq Line: 43  
 Vial 44

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



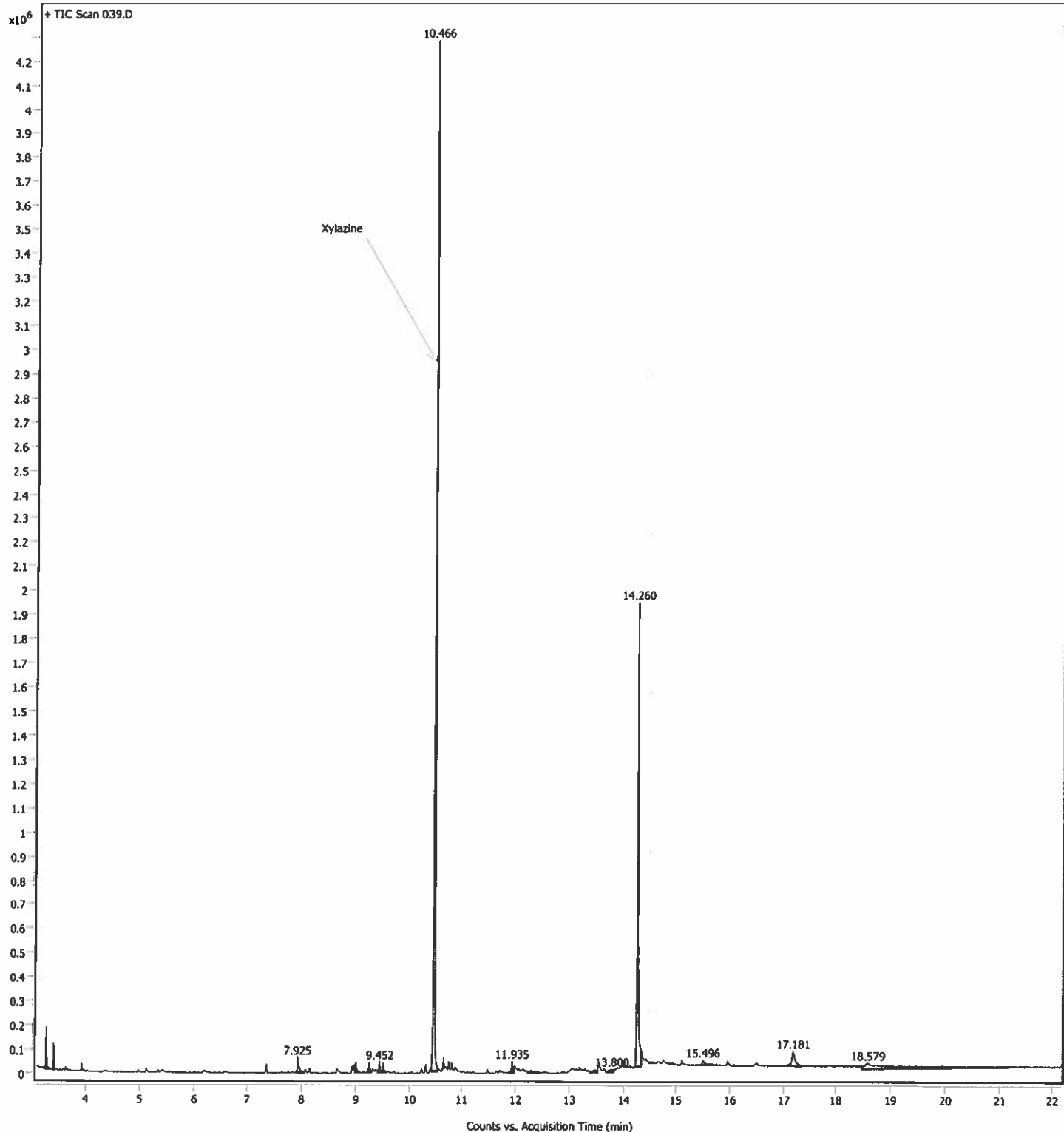
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.061	11.046	2010.912	1337.043	0.000000	XYLAZINE
11.364	11.278	6.070	3.448	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

<b>Sample Name</b>	MATRIX 6 W/O ISTD - URN	<b>Data File Path</b>	C:\MassHunter\GCMS\1\data\BASES\092821\039.D
<b>Instrument</b>	#3 - Enhanced	<b>Acq. Time (Local)</b>	9/29/2021 8:11:23 AM (UTC-04:00)
<b>Position</b>	70	<b>Method Path (Acq)</b>	C:\MassHunter\GCMS\1\methods\ALKALI.M
<b>Operator</b>	KM/QC DATA IN XYLAZINE VALIDATION		2.0 mg/L - 400 uL of 0.01 mg/mL xylazine std

## Sample Chromatograms



# Analysis Report

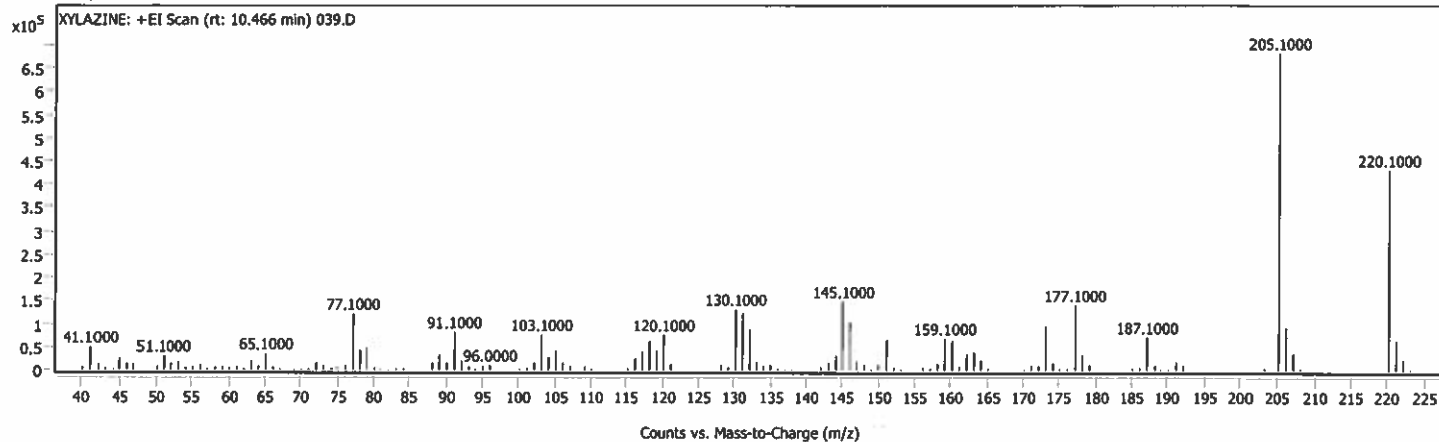
## Sample Spectra

+ Scan (rt: 10.466 min)

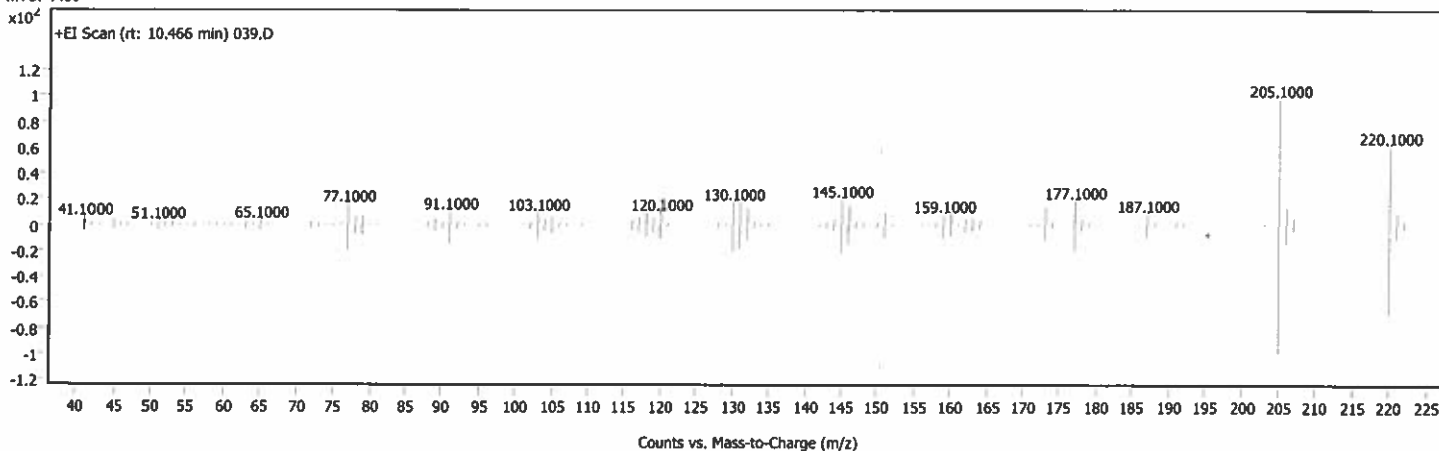
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.62	aafs.l	MATRIX 6 W/O ISTD - URN	KM/QC DATA IN XYLAZINE VALIDATION

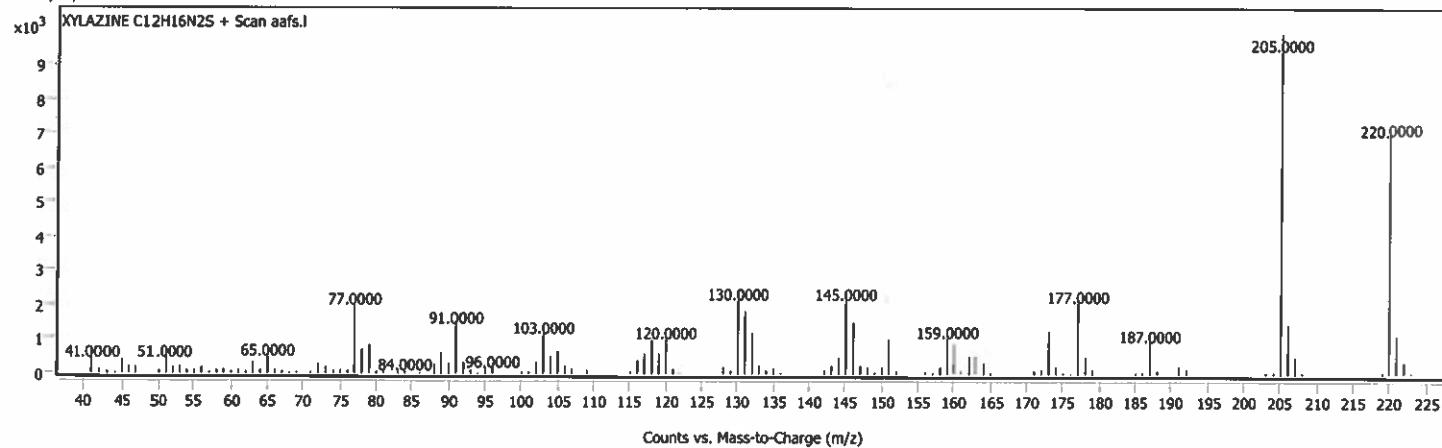
### Observed Spectrum



### Mirror Plot



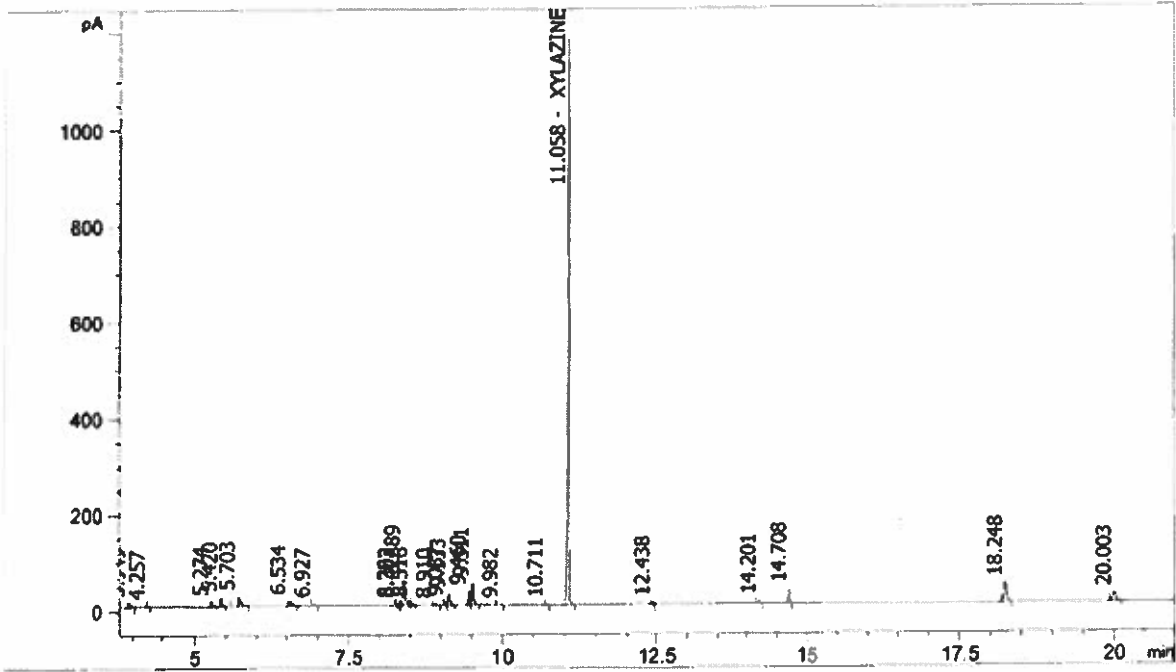
### Library Spectrum



Injection Date: 9/29/2021 7:35:35 AM  
 Sample Name: MATRIX 6 W/O ISTD - URN  
 Sample Info:

Seq Line: 45  
 Vial 46

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



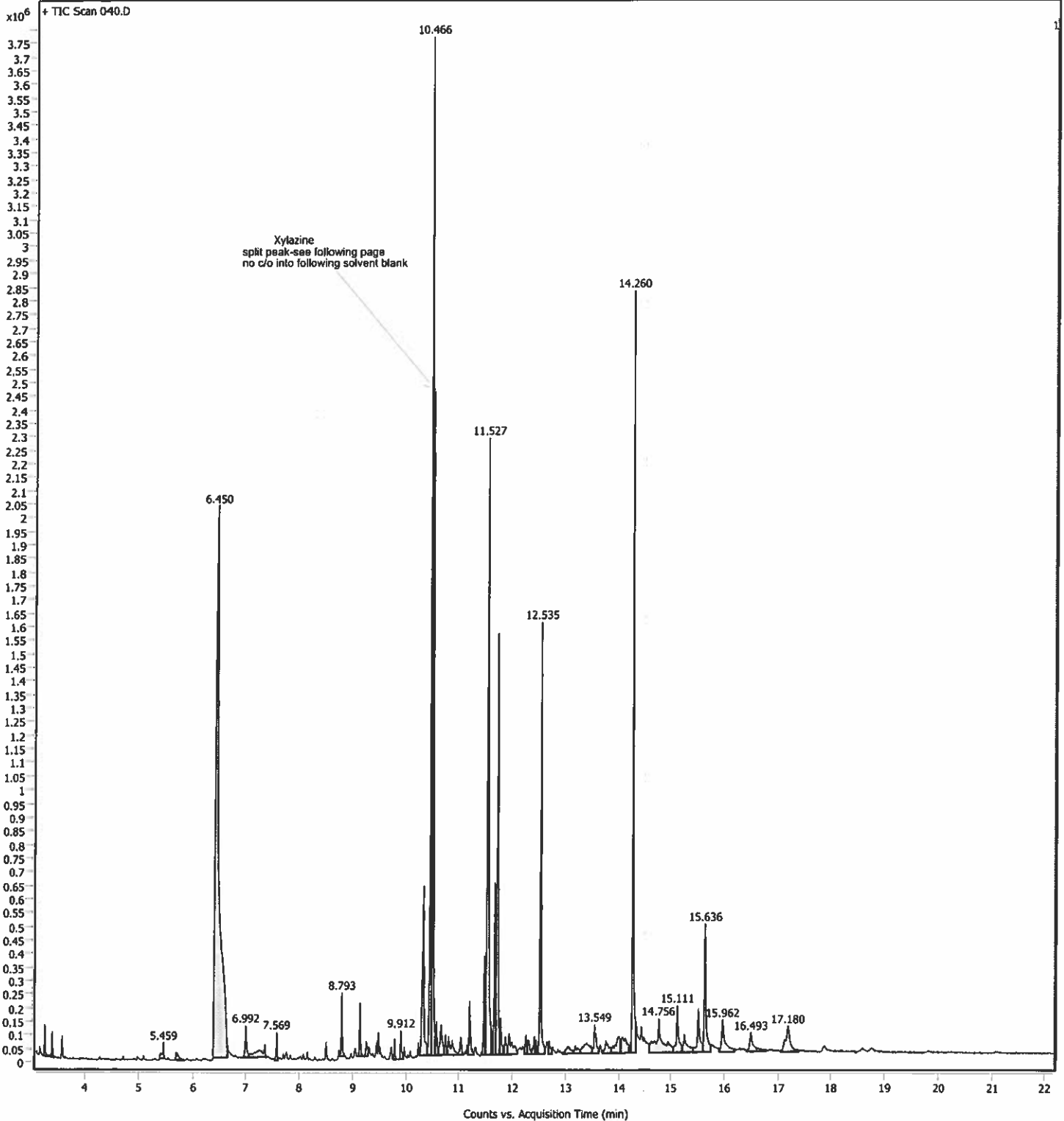
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.058	11.046	1684.577	1168.928	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

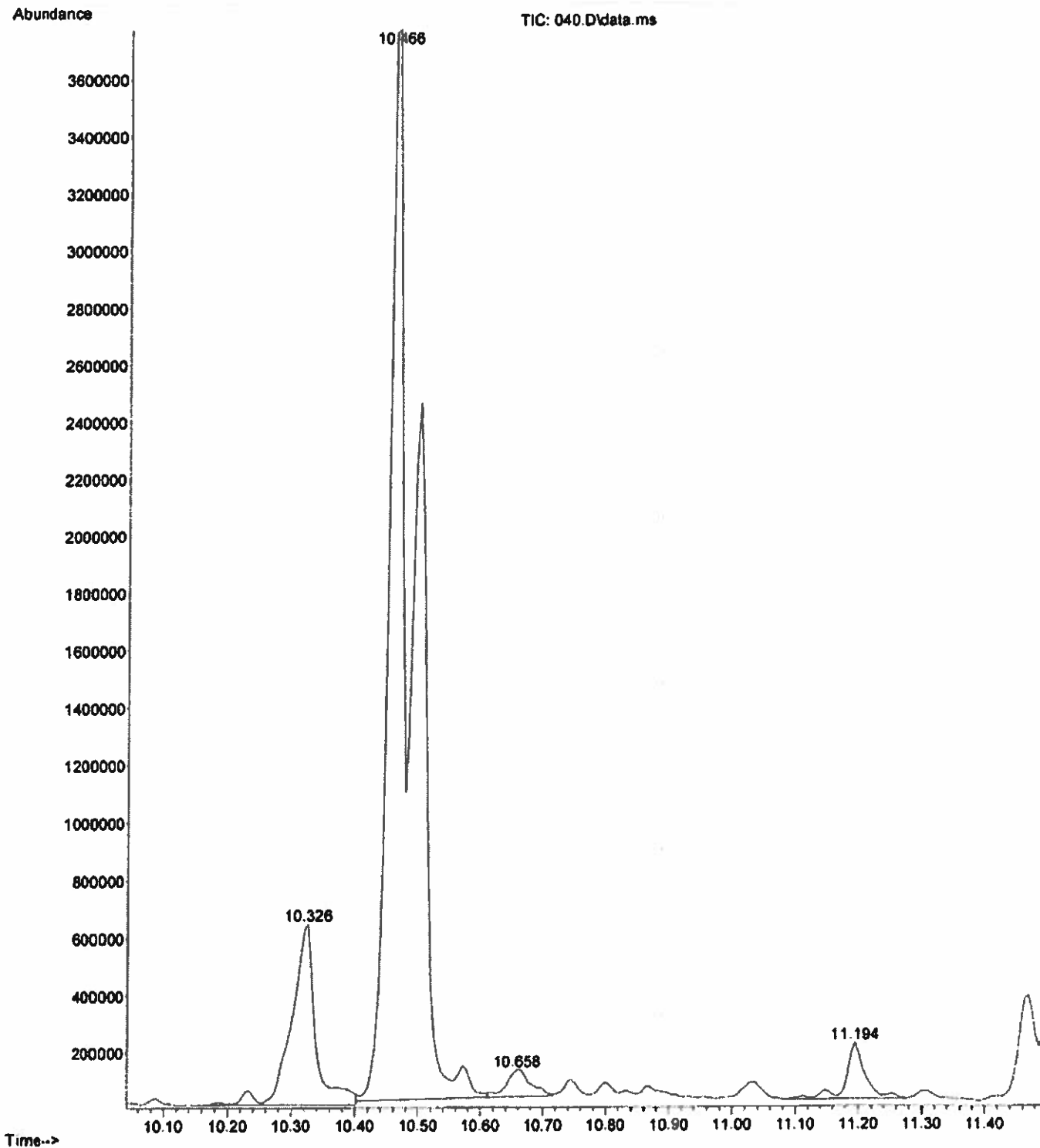
## Sample Information

Sample Name	MATRIX 10 W/O ISTD - LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\040.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 8:37:19 AM (UTC-04:00)
Position	71	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

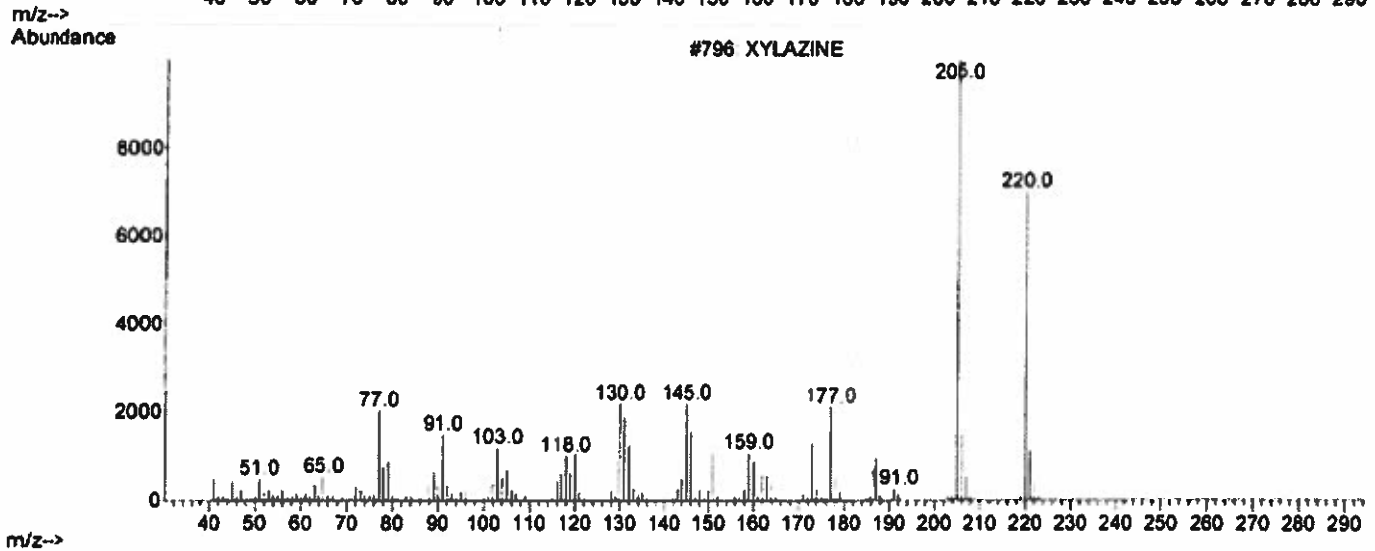
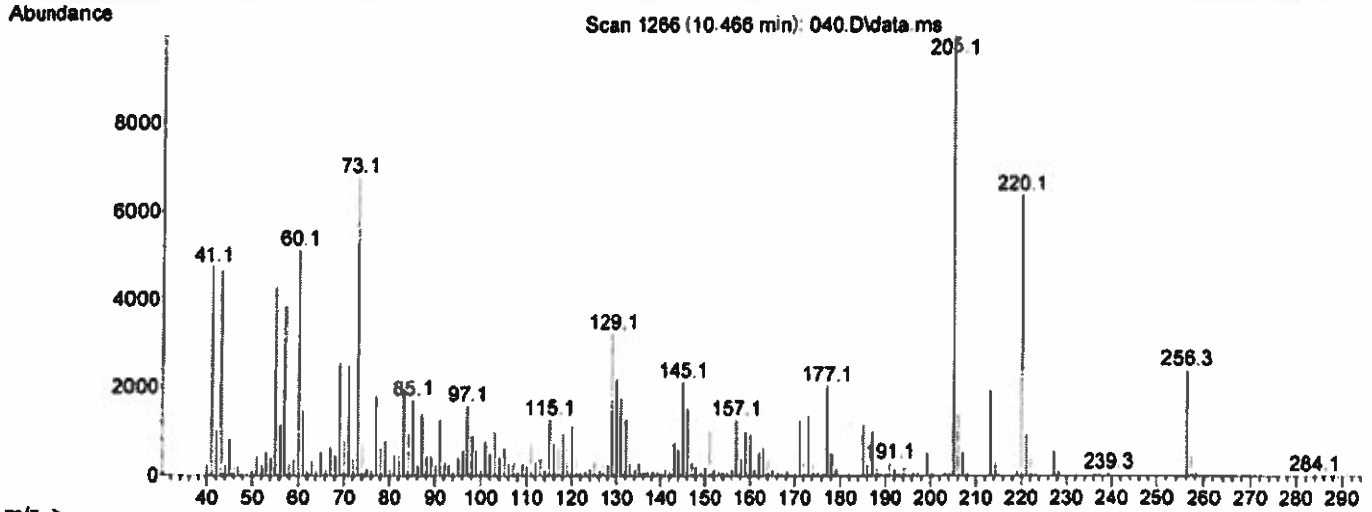
## Sample Chromatograms



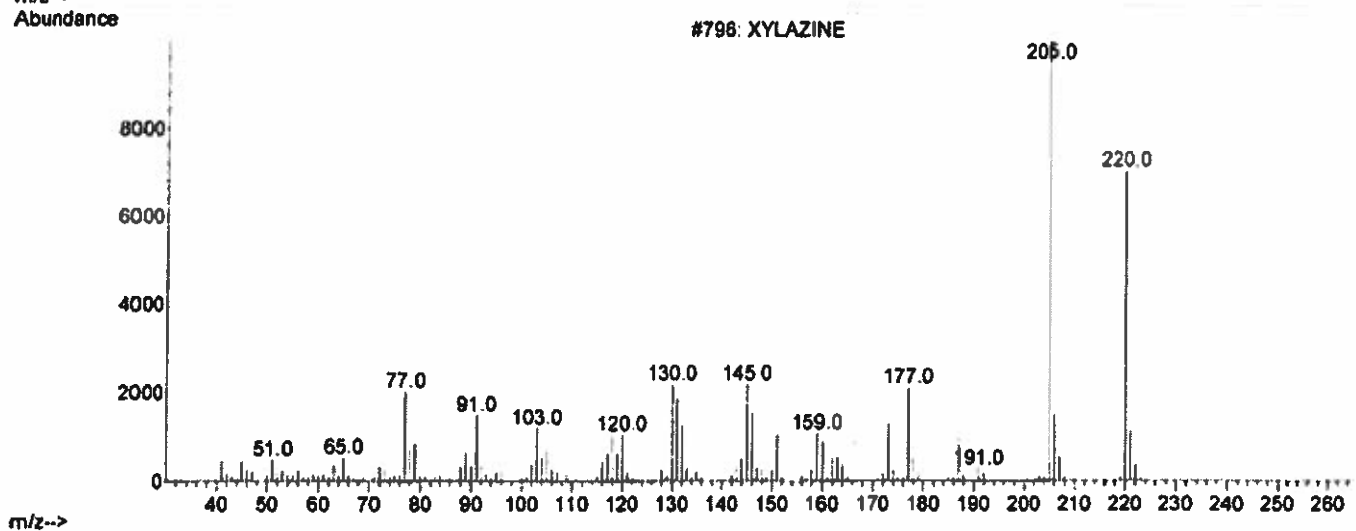
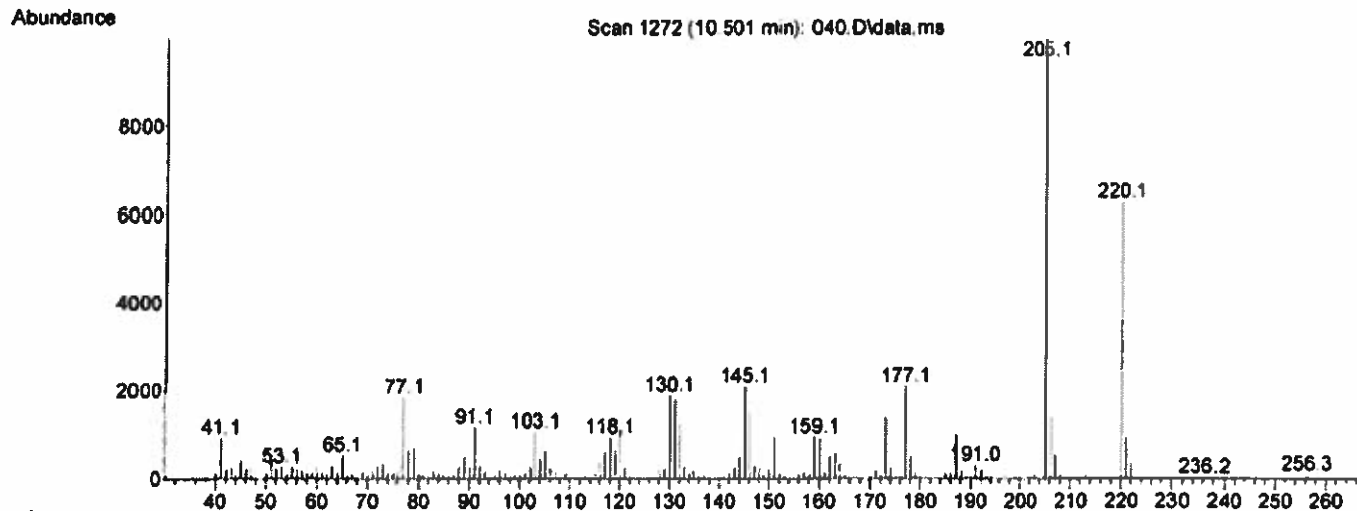
File : C:\Users\TOX\Desktop\092821\040.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 08:37 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 10 W/O ISTD - LVR  
Misc Info : 2.0 mg/L - 400uL of 0.01 mg/mL XYLAZINE WS  
Vial Number: 71



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 99  
ID : XYLAZINE  
Sample Name: MATRIX 10 W/O ISTD - LVR  
File: C:\Users\TOX\Desktop\092821\040.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 08:37  
Vial: 71



Library Searched : C:\MassHunter\Library\aaFs.1  
Quality : 99  
ID : XYLAZINE  
Sample Name: MATRIX 10 W/O ISTD - LVR  
File: C:\Users\TOX\Desktop\092821\040.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 08:37  
Vial: 71



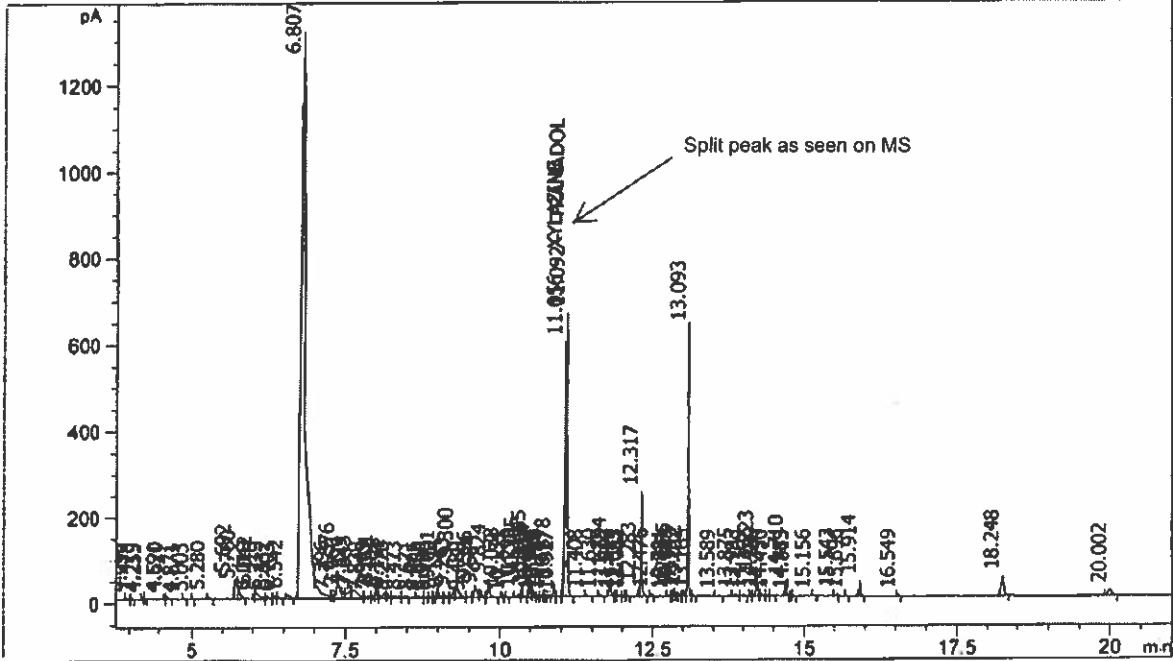


Injection Date: 9/29/2021 7:59:56 AM  
 Sample Name: MATRIX 10 W/O ISTD - ->  
 Sample Info:

Seq Line: 46  
 Vial 47

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

2.0 mg/L - split peak only seen in LVR sample - OK for qual reporting



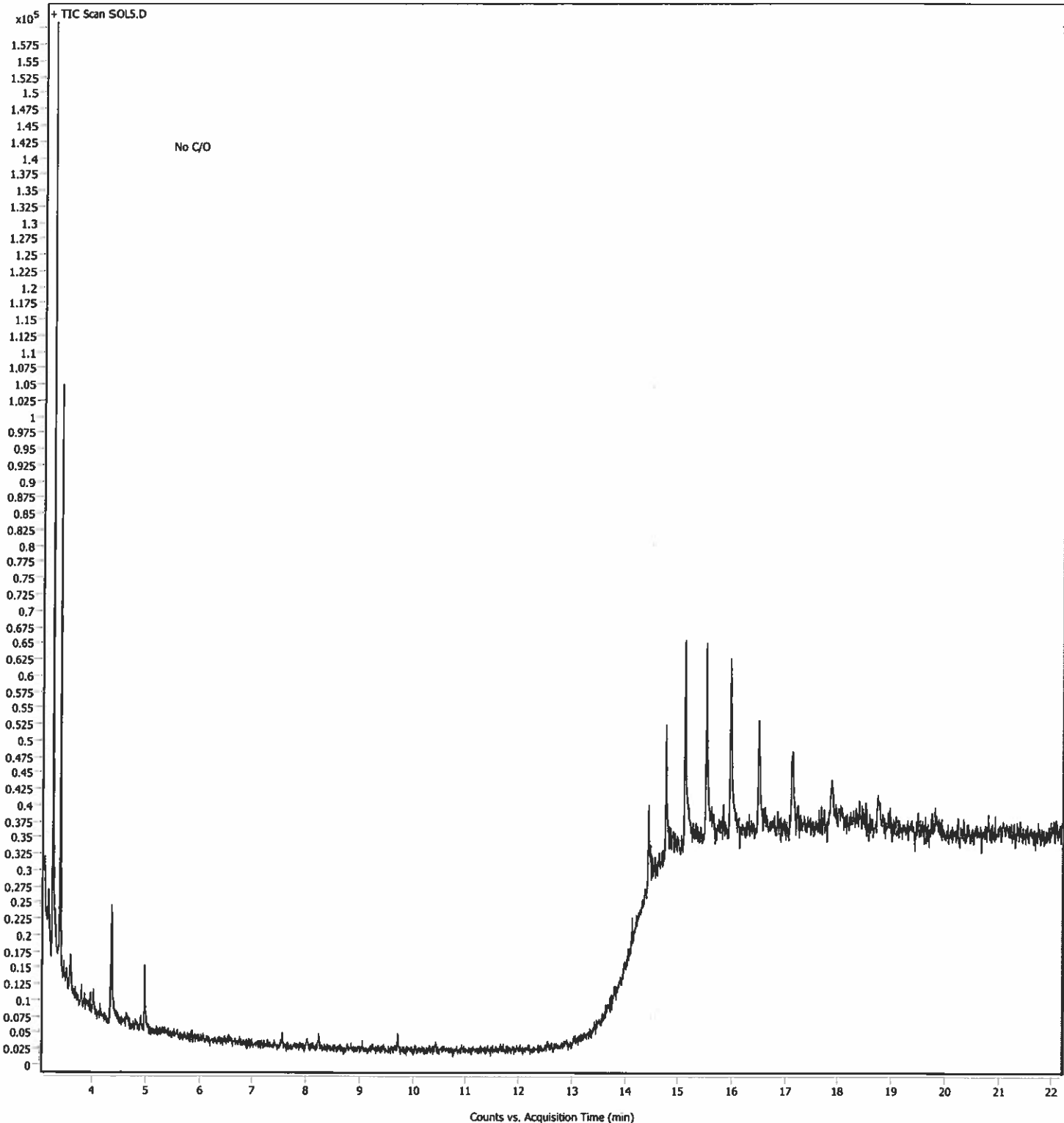
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.056	11.046	806.661	593.069	0.000000	XYLAZINE
11.092	11.033	979.896	658.757	0.000000	TRAMADOL
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	Solvent Blank 5	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\SOL5.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 9:03:08 AM (UTC-04:00)
Position	72	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

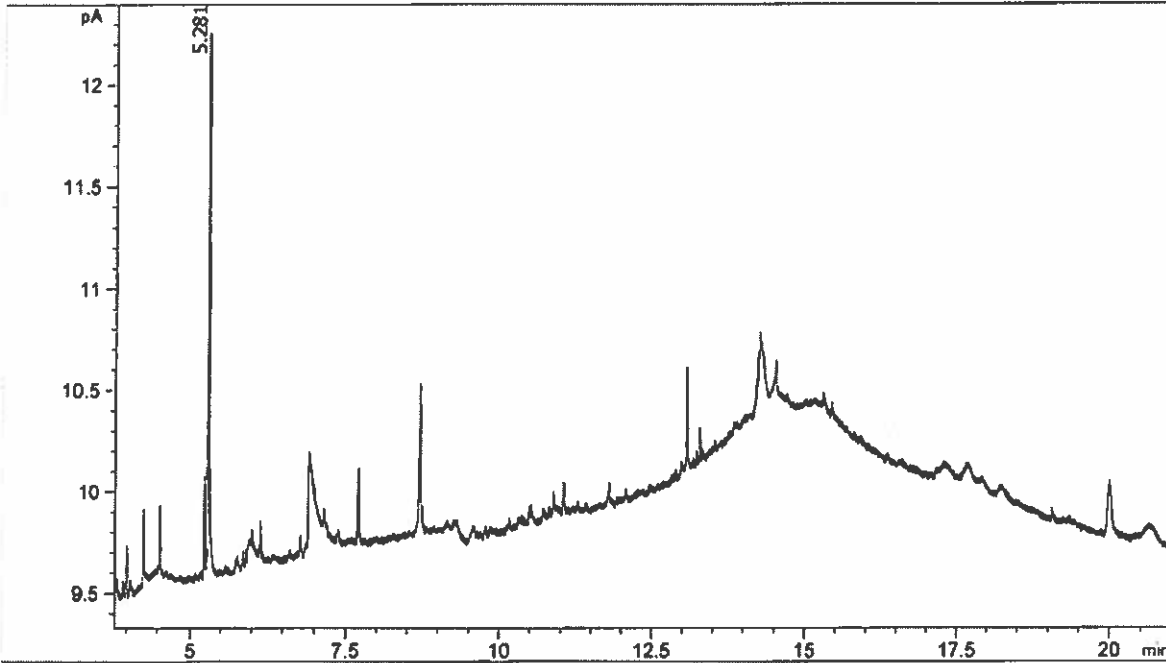
## Sample Chromatograms



Injection Date: 9/29/2021 8:24:14 AM  
Sample Name: SOLVENT BLANK 5  
Sample Info:

Seq Line: 47  
Vial 48

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



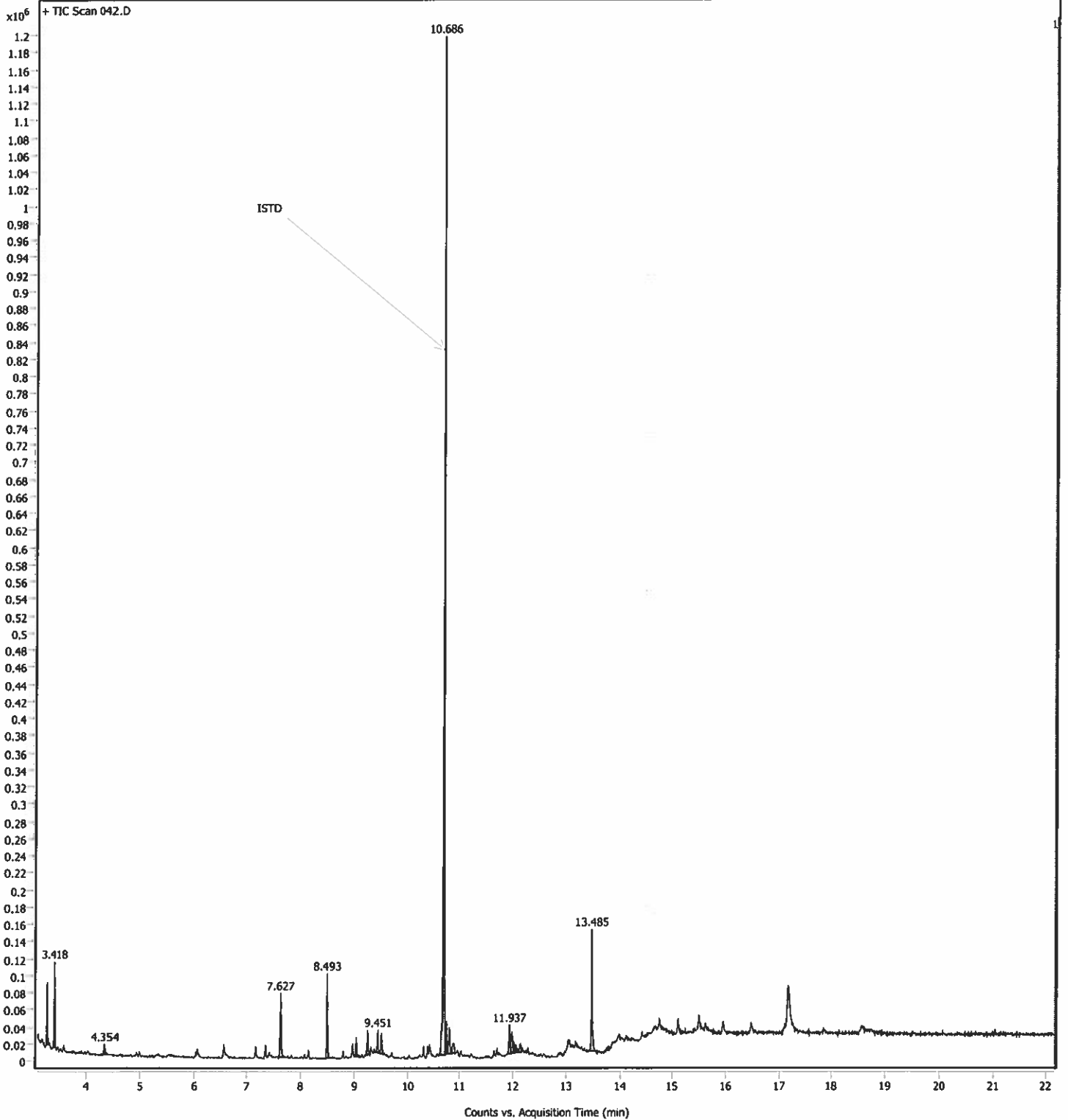
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 2 W/ISTD ONLY - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\042.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 9:54:51 AM (UTC-04:00)
Position	74	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

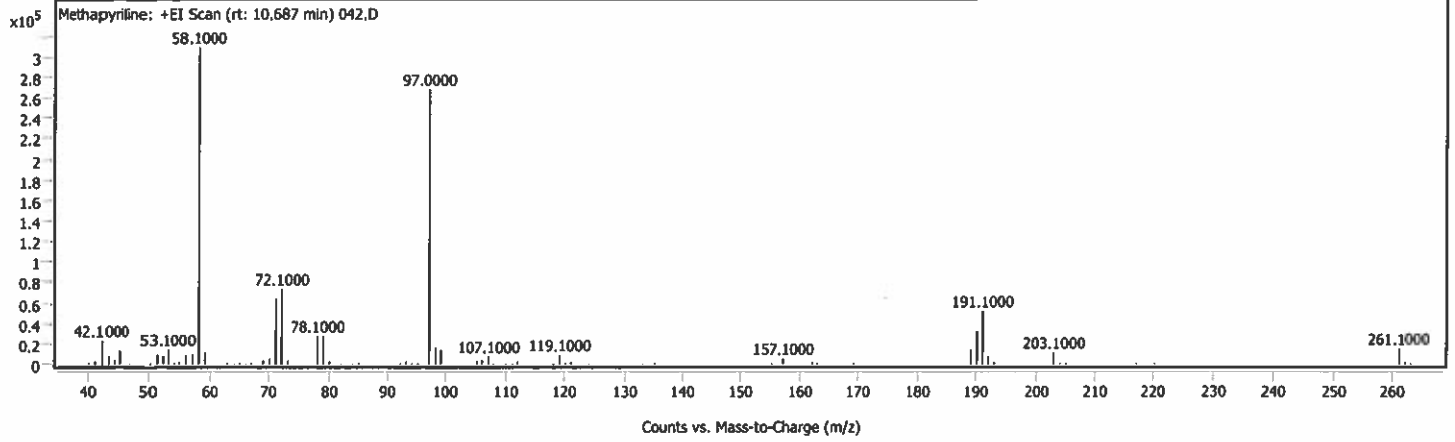
## Sample Spectra

+ Scan (rt: 10.687 min)

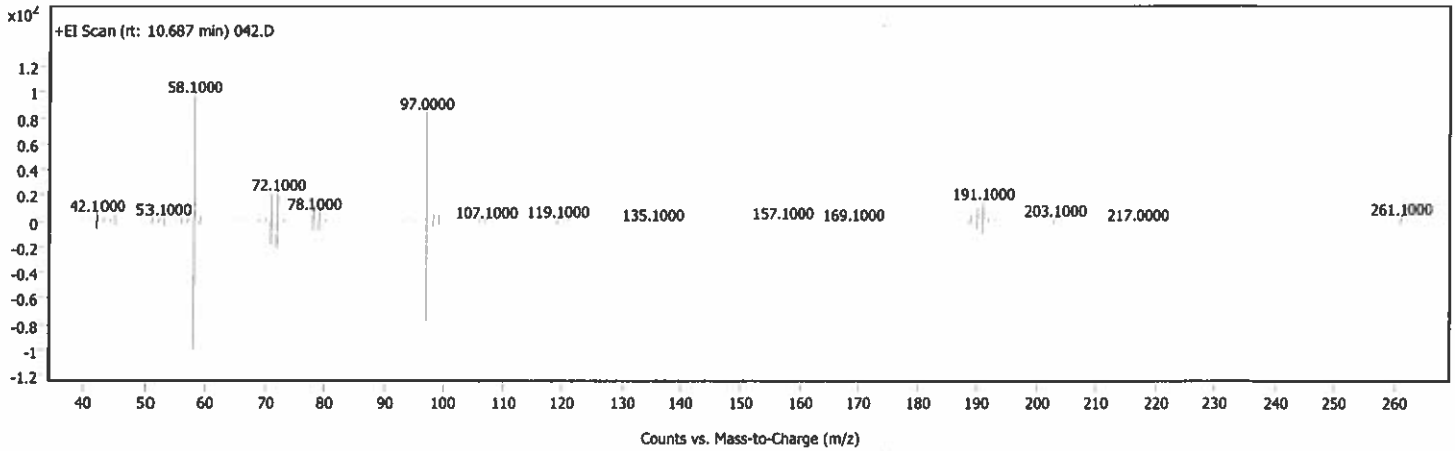
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.56	SWGDRUG 3.3.L	MATRIX 2 W/ISTD ONLY - BLD	KM/QC DATA IN XYLAZINE VALIDATION

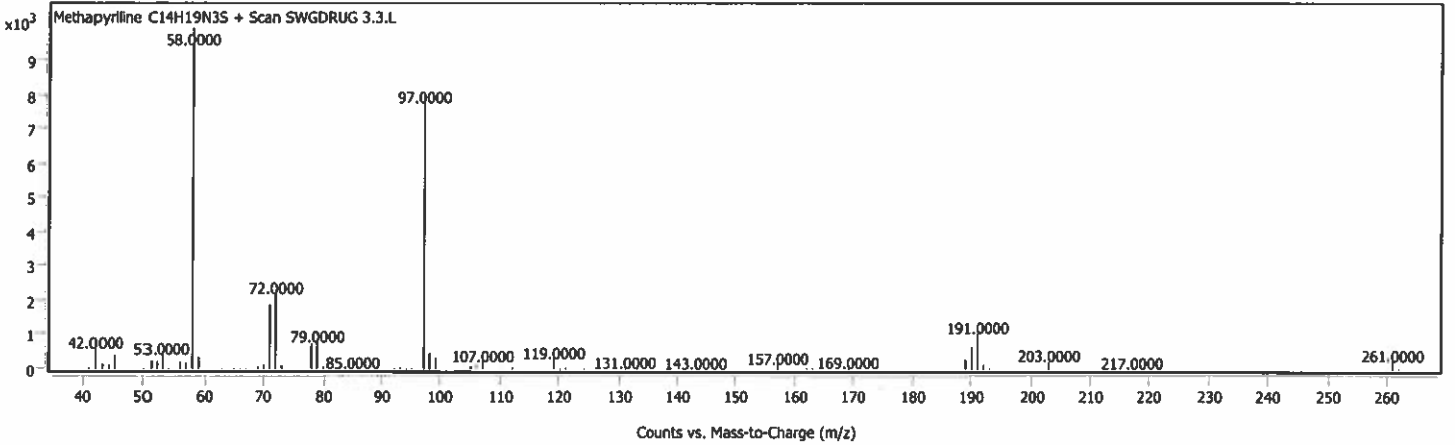
## Observed Spectrum



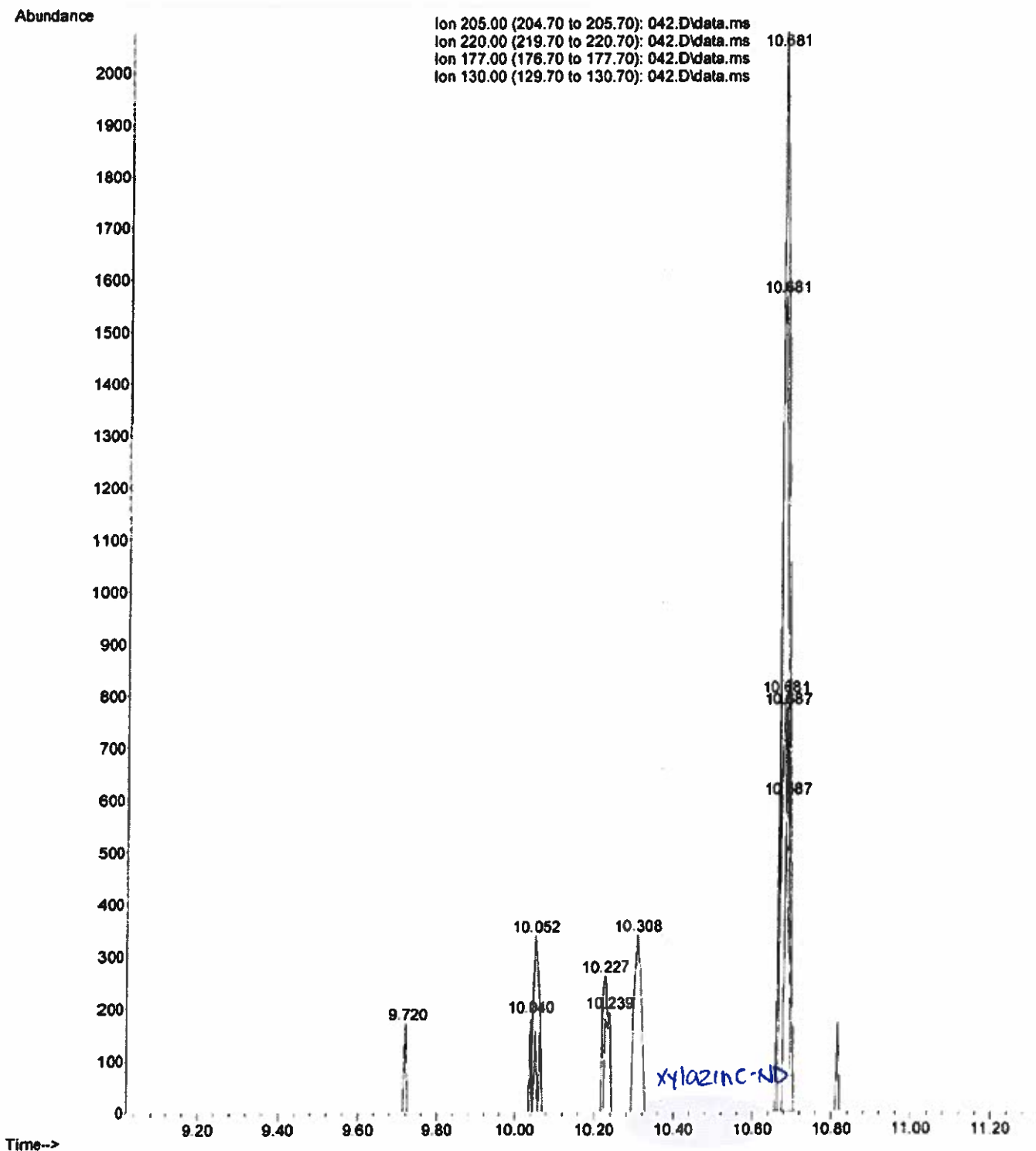
## Mirror Plot



## Library Spectrum



File :C:\Users\TOX\Desktop\092821\042.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 09:54 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 2 W/ISTD ONLY - BLD  
Misc Info :  
Vial Number: 74



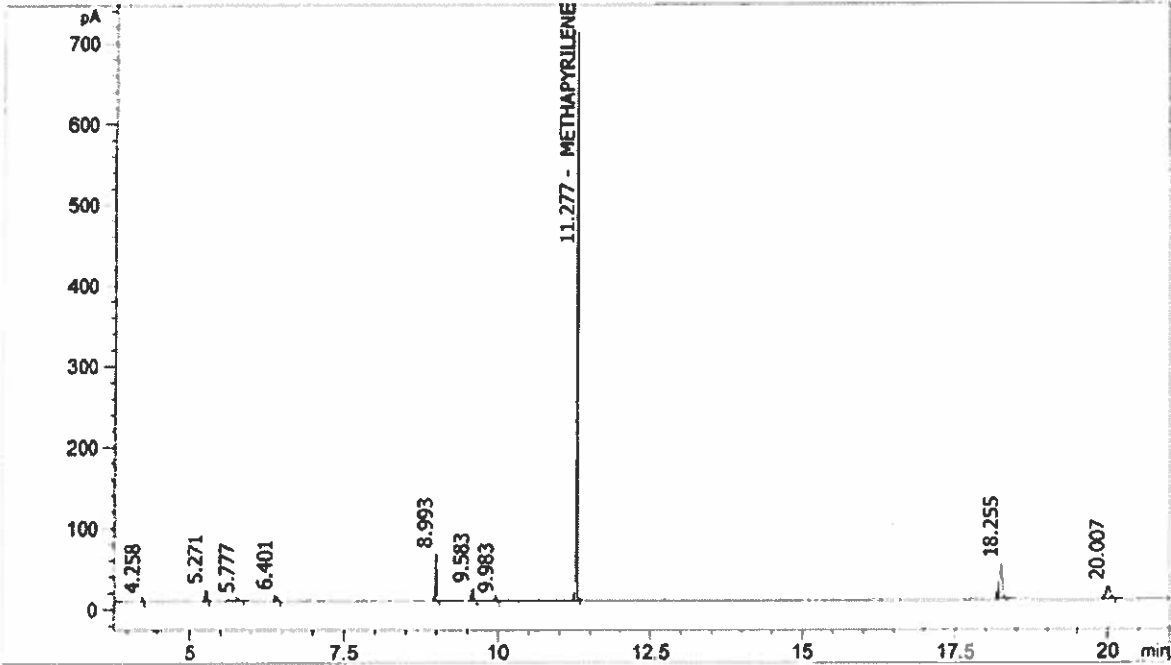
Ion 205.00 (204.70 to 205.70): 042.D\data.ms  
Ion 220.00 (219.70 to 220.70): 042.D\data.ms  
Ion 177.00 (176.70 to 177.70): 042.D\data.ms  
Ion 130.00 (129.70 to 130.70): 042.D\data.ms

xylozinc-ND

Injection Date: 9/29/2021 9:12:52 AM  
Sample Name: MATRIX 2 W/ISTD ONLY ->  
Sample Info:

Seq Line: 49  
Vial 50

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



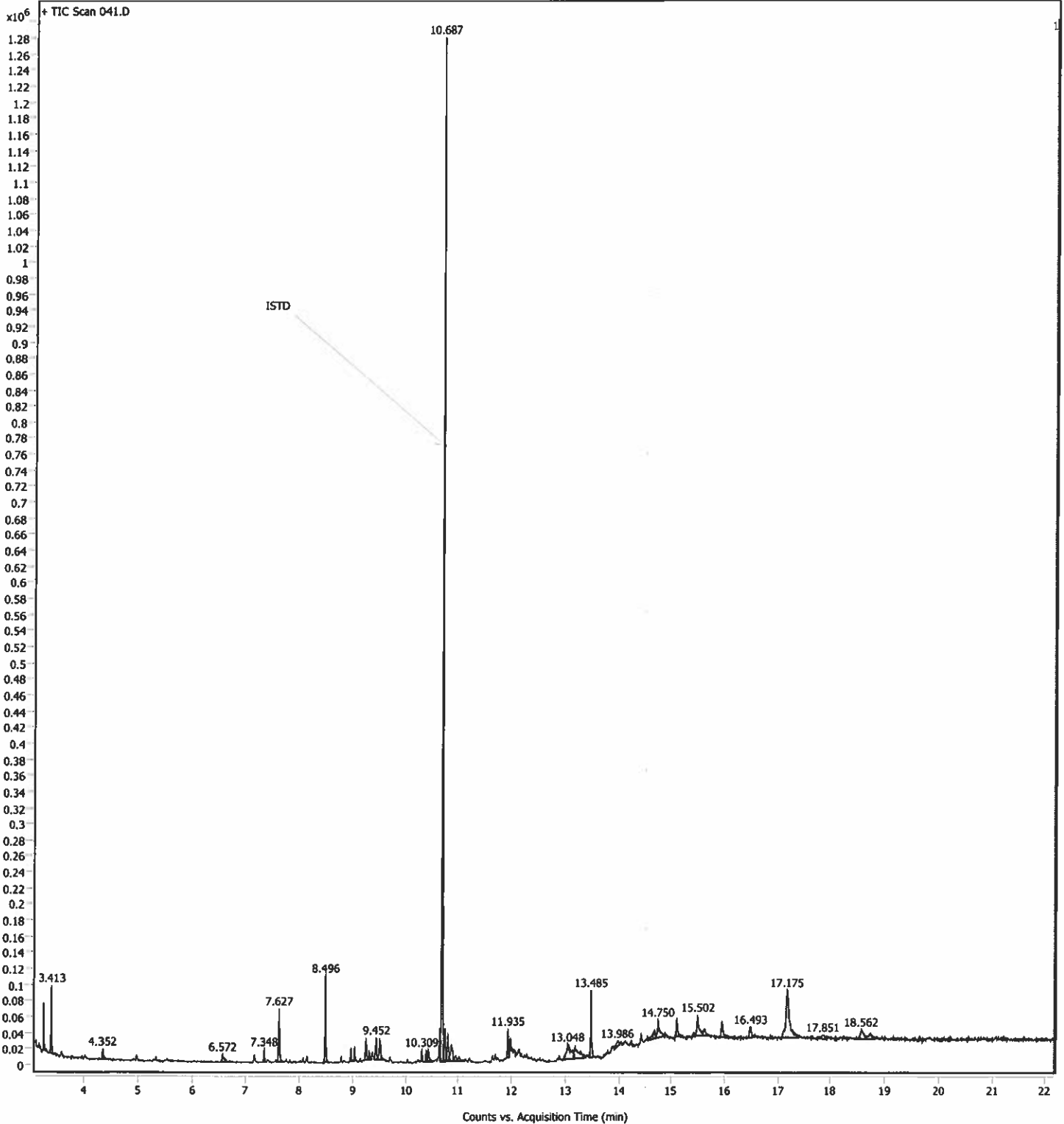
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.277	11.278	805.740	699.676	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

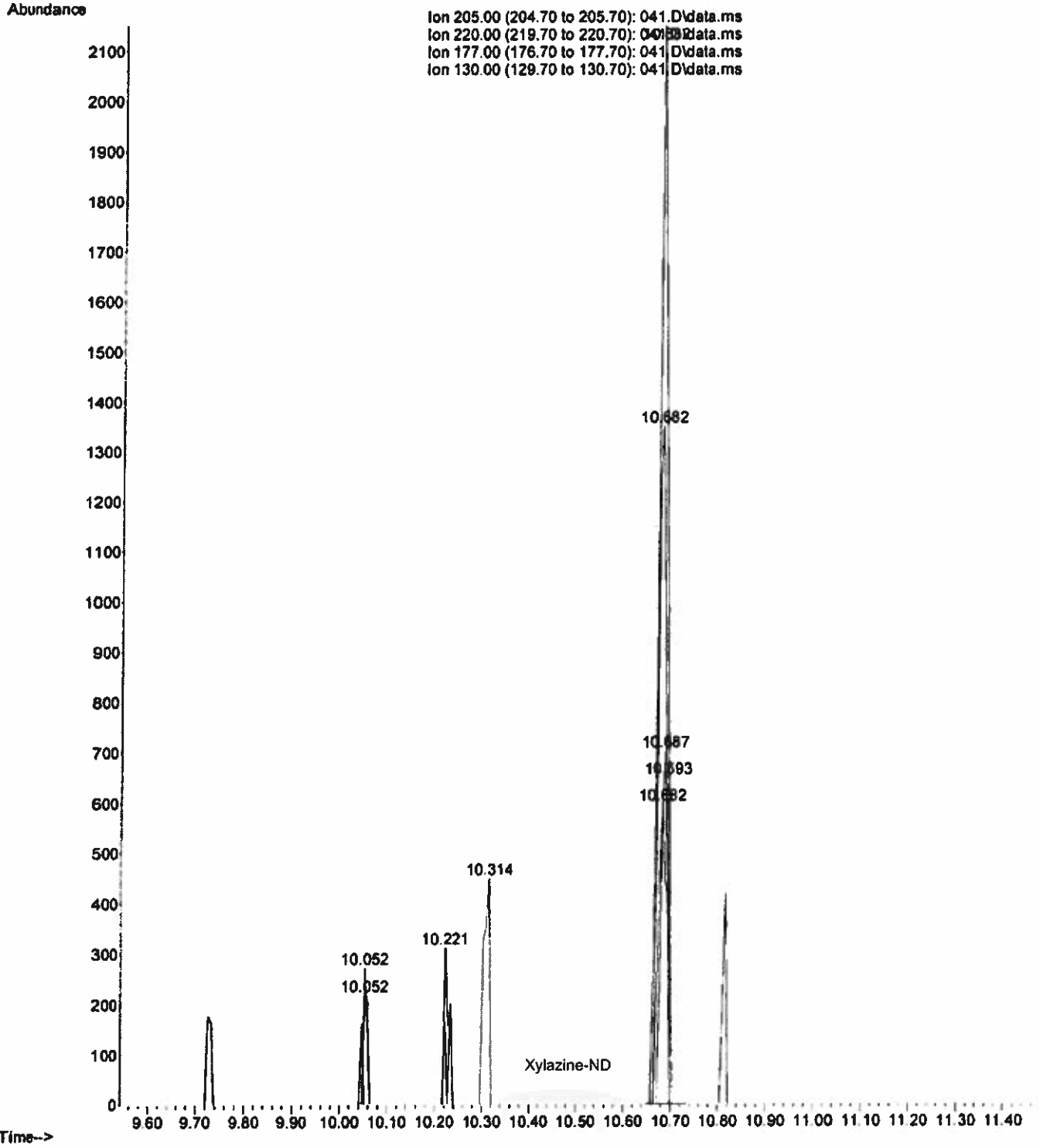
Sample Name	MATRIX 1 W/STD ONLY - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\041.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 9:28:54 AM (UTC-04:00)
Position	73	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





File :C:\Users\TOX\Desktop\092821\041.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 09:28 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 1 W/ISTD ONLY - BLD  
Misc Info :  
Vial Number: 73



# Analysis Report

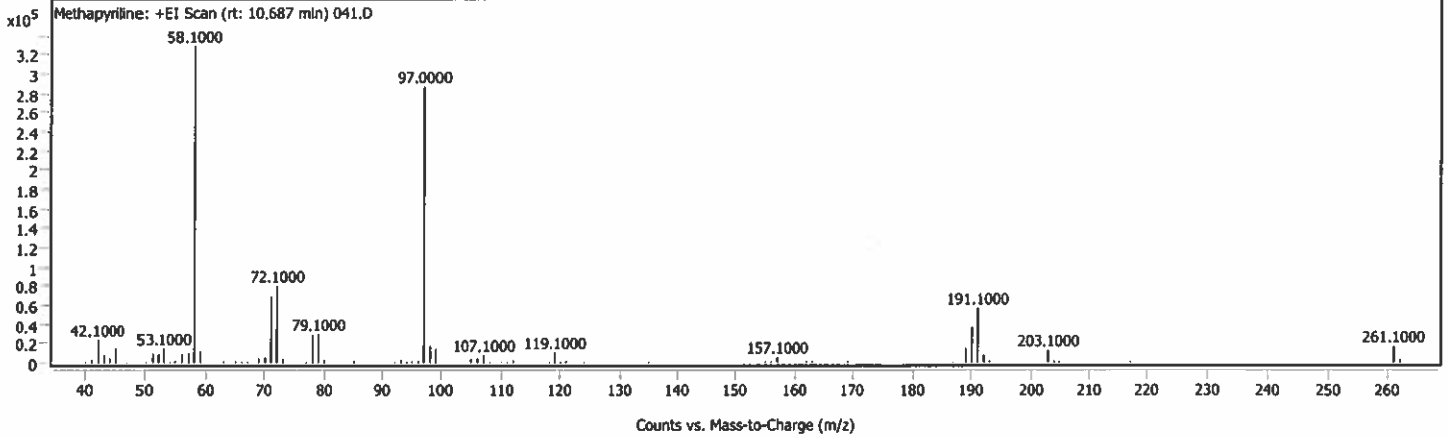
## Sample Spectra

+ Scan (rt: 10.687 min)

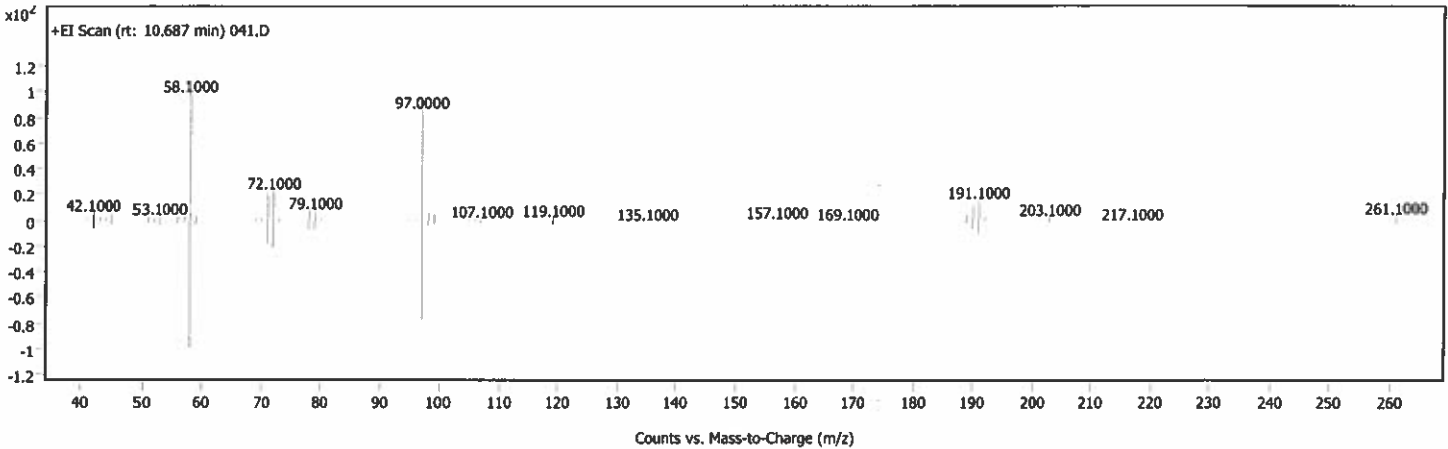
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.19	SWGDRUG 3.3.L	MATRIX 1 W/ISTD ONLY - BLD	KM/QC DATA IN XLAZINE VALIDATION

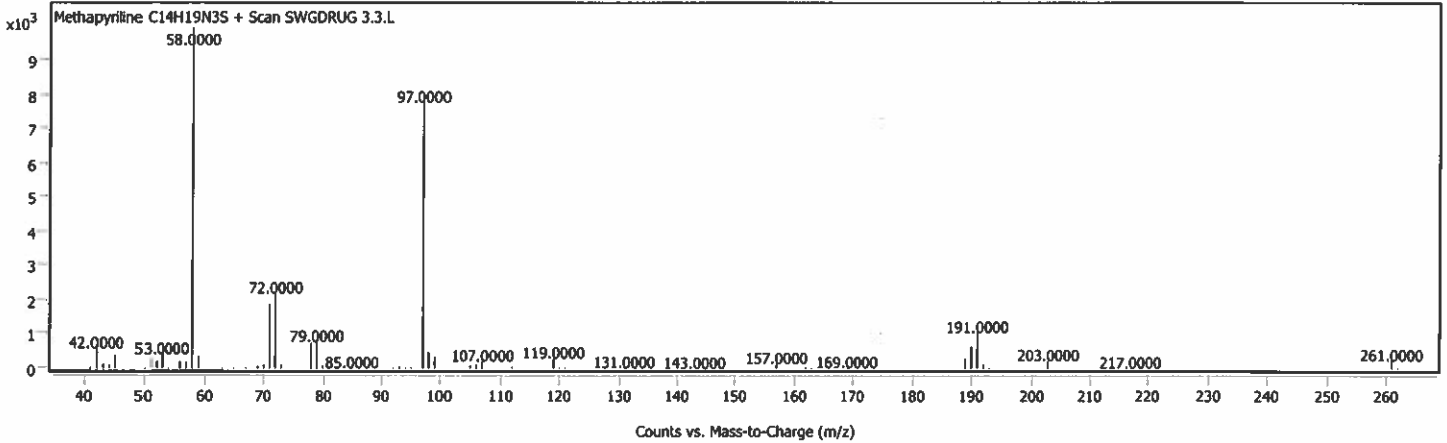
## Observed Spectrum



## Mirror Plot

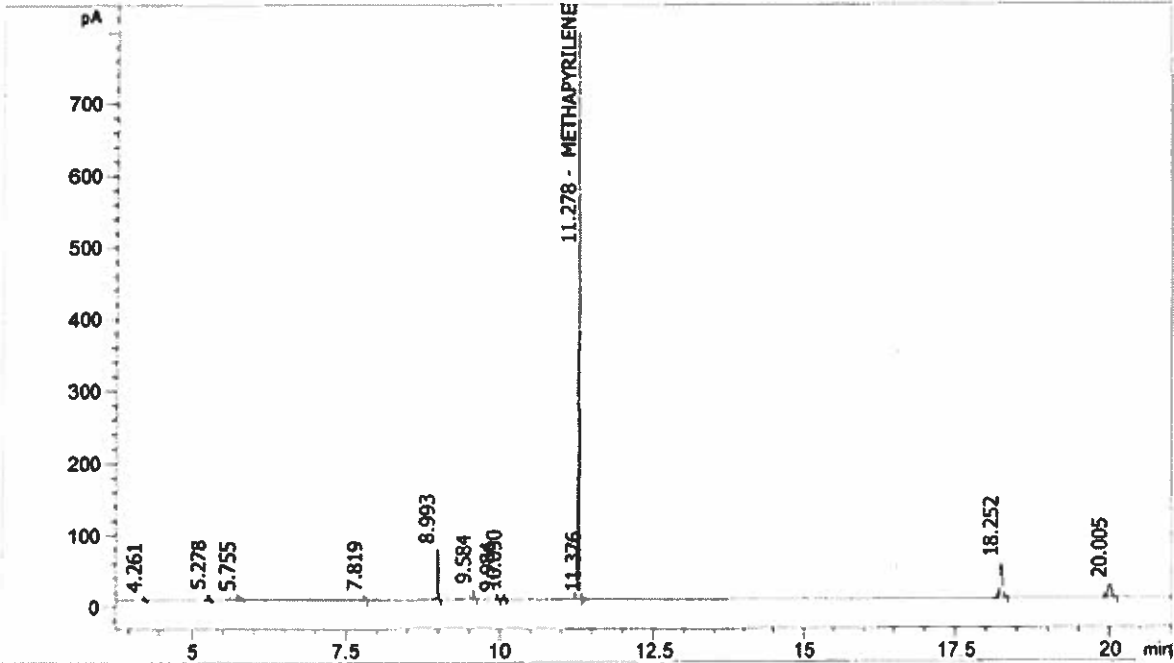


## Library Spectrum



Injection Date: 9/29/2021 8:48:31 AM Seq Line: 48  
 Sample Name: MATRIX 1 W/ISTD ONLY -> Vial 49  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



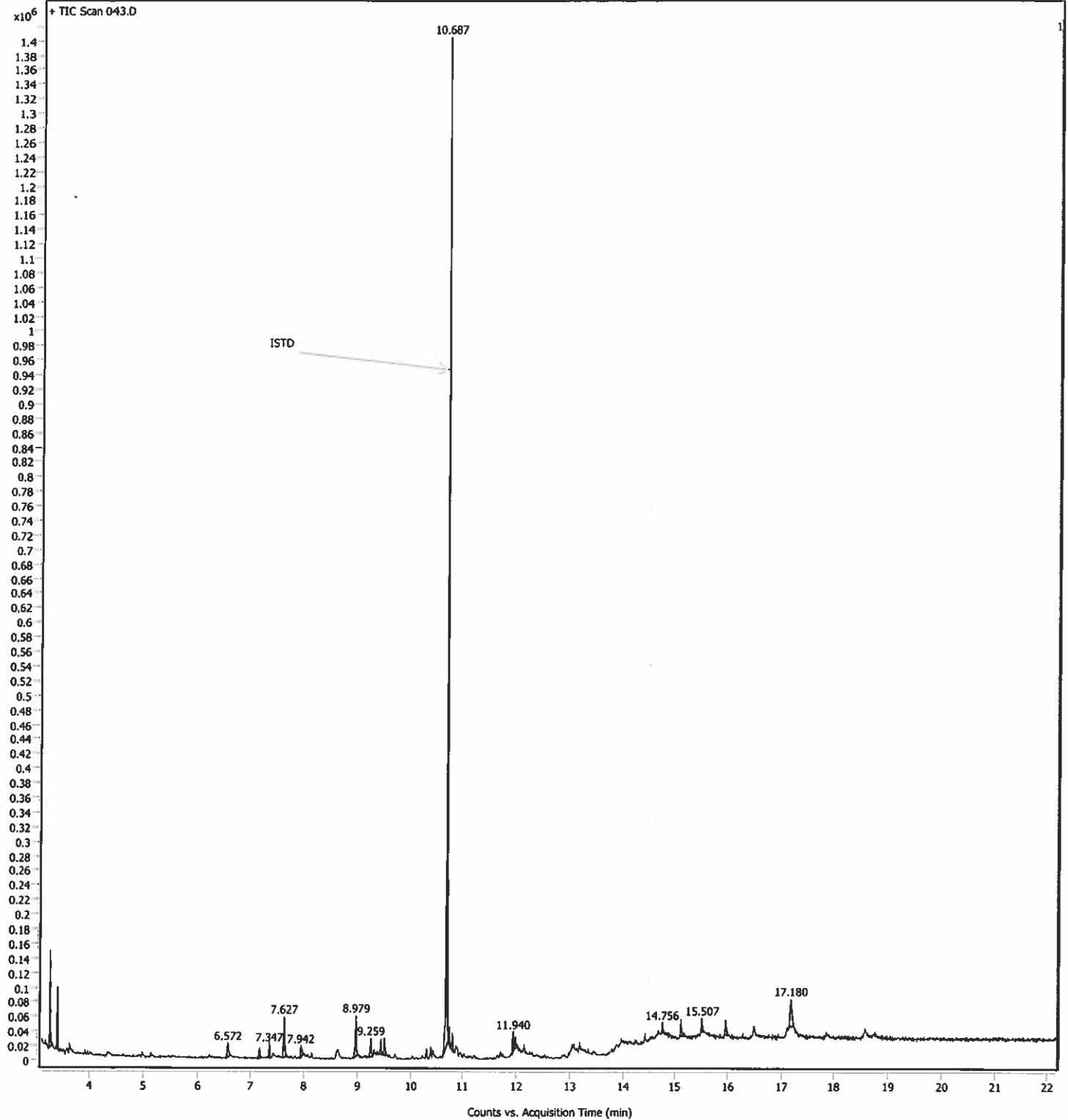
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	917.259	787.908	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 W/ISTD ONLY- URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\043.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 10:20:37 AM (UTC-04:00)
Position	75	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

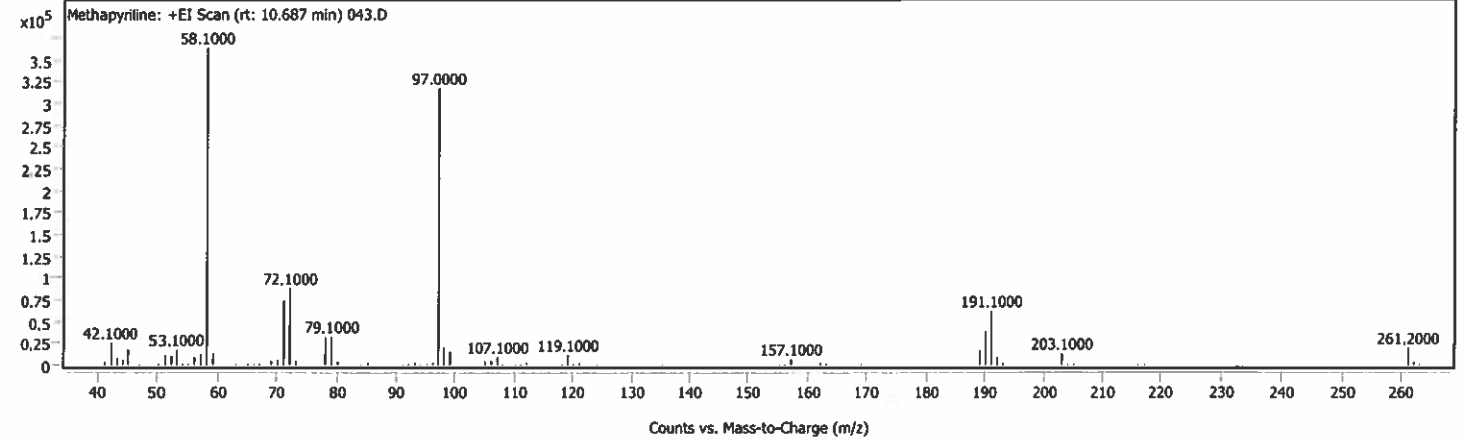
## Sample Spectra

+ Scan (rt: 10.687 min)

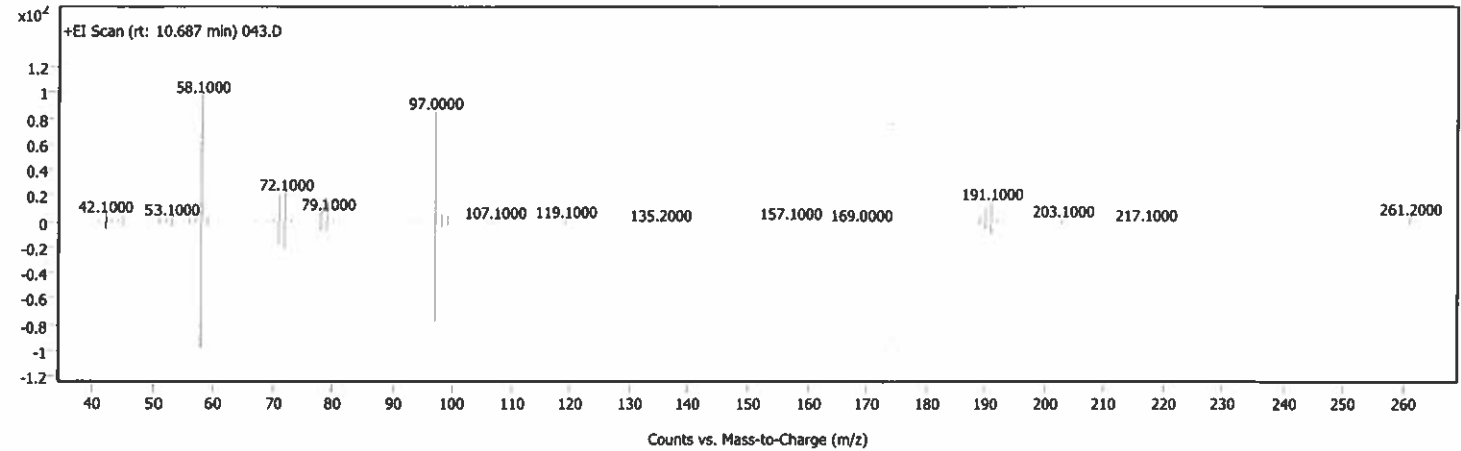
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.29	SWGDRUG 3.3.L	MATRIX 6 W/ISTD ONLY- URN	KM/QC DATA IN XYLAZINE VALIDATION

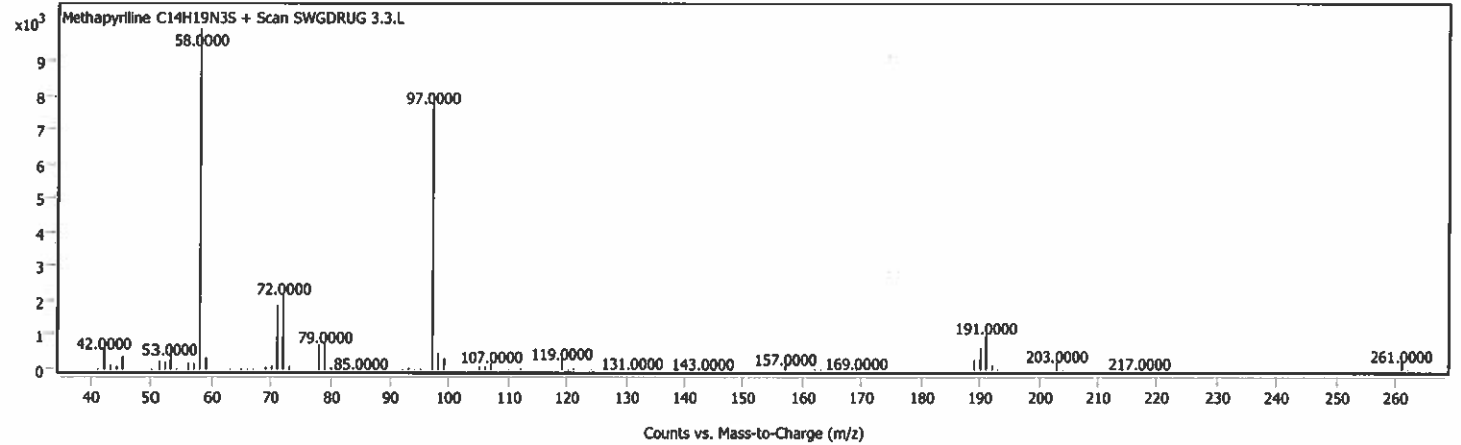
## Observed Spectrum



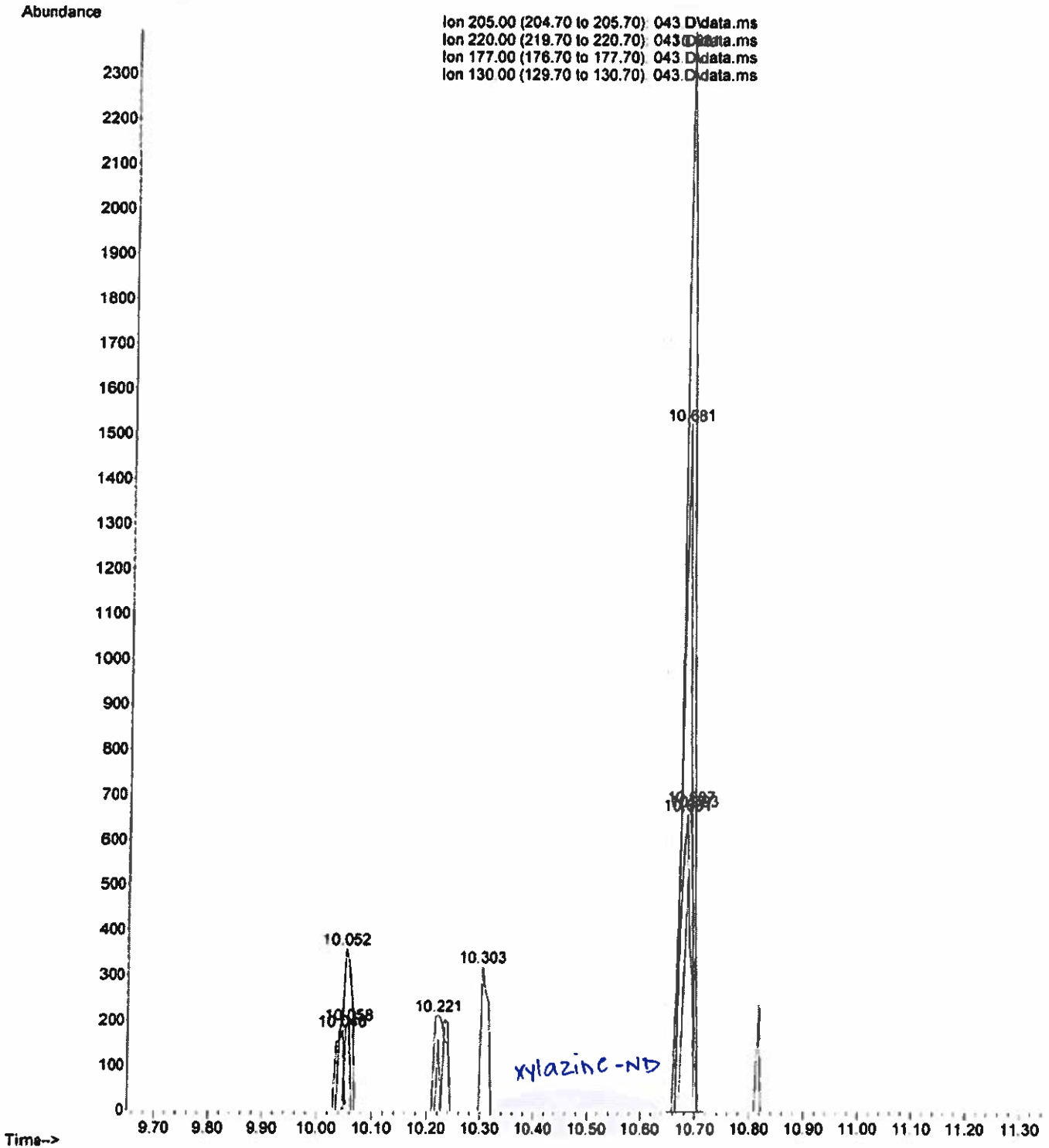
## Mirror Plot



## Library Spectrum

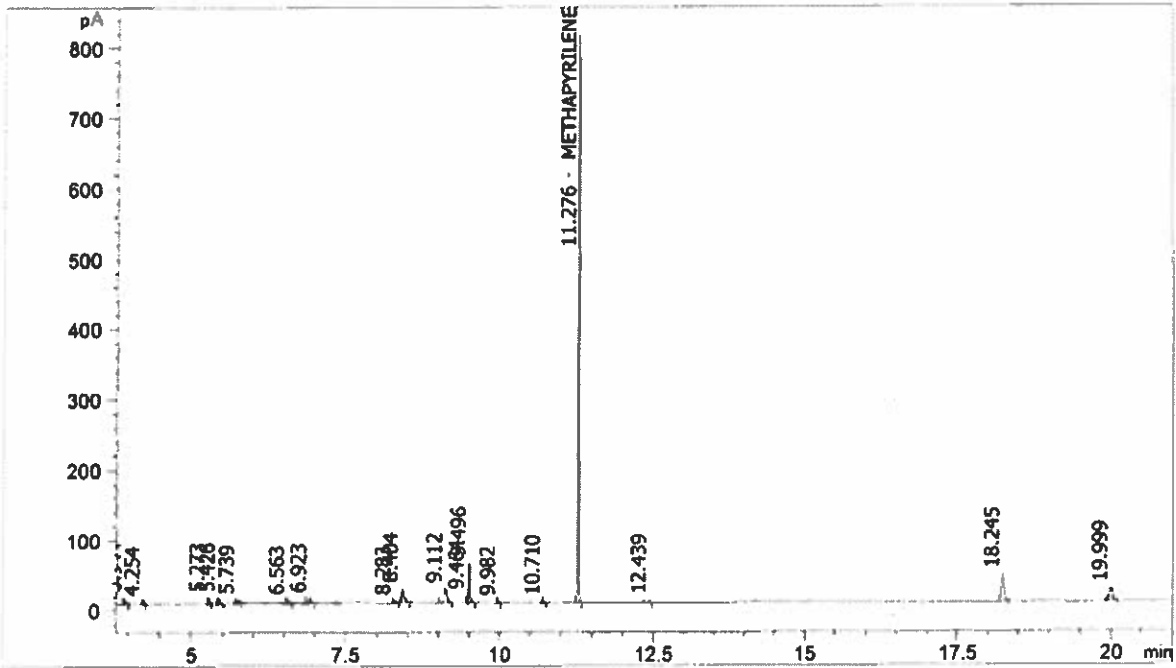


File :C:\Users\TOX\Desktop\092821\043.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 10:20 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 6 W/ISTD ONLY- URN  
Misc Info :  
Vial Number: 75



Injection Date: 9/29/2021 9:37:10 AM Seq Line: 50  
 Sample Name: MATRIX 6 W/ISTD ONLY--> Vial 81  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



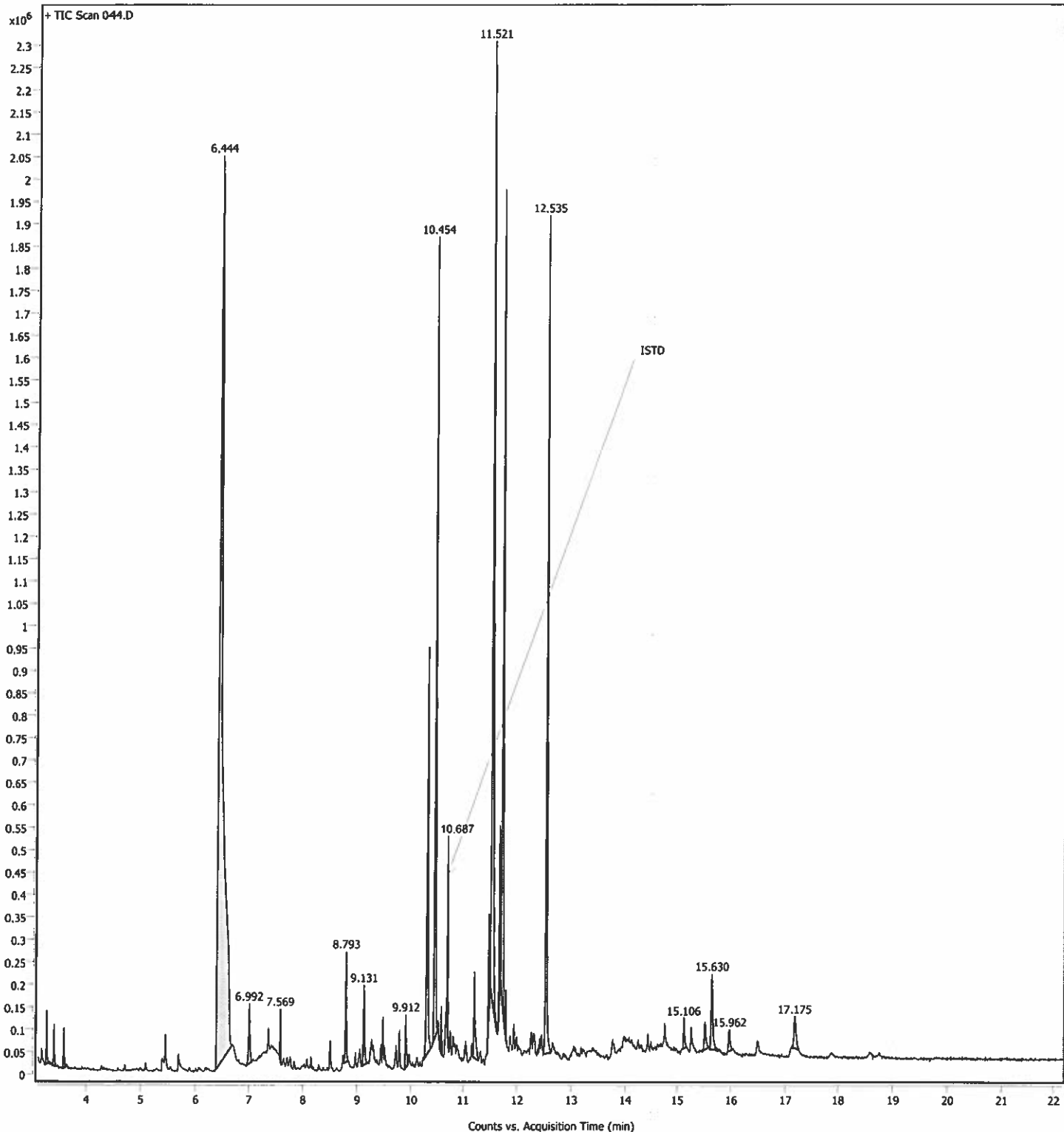
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMACOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.276	11.278	916.839	806.497	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 10 W/ISTD ONLY- LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\044.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 10:46:27 AM (UTC-04:00)
Position	76	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

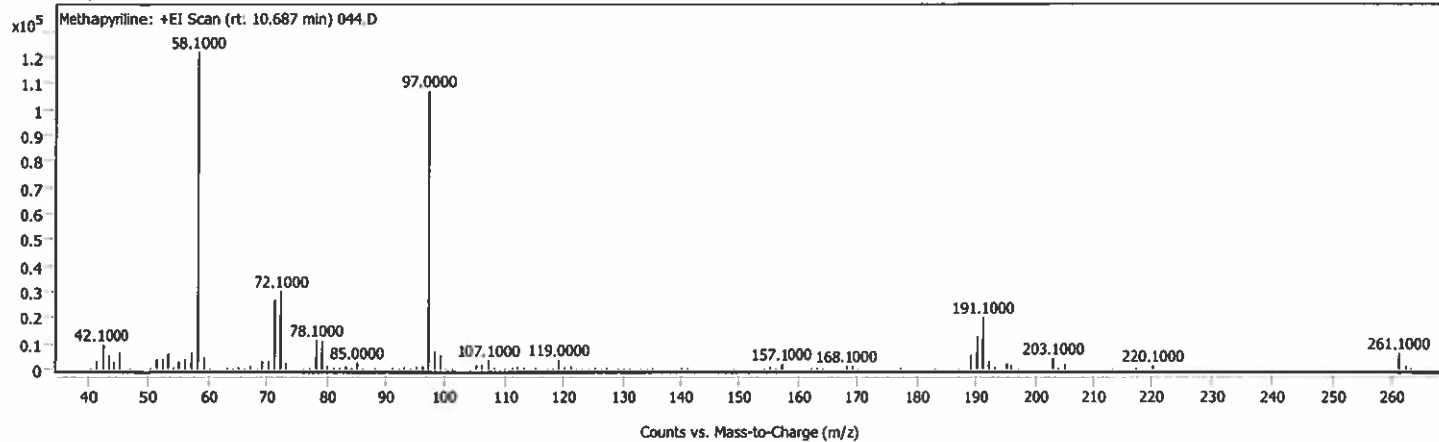
## Sample Spectra

+ Scan (rt: 10.687 min)

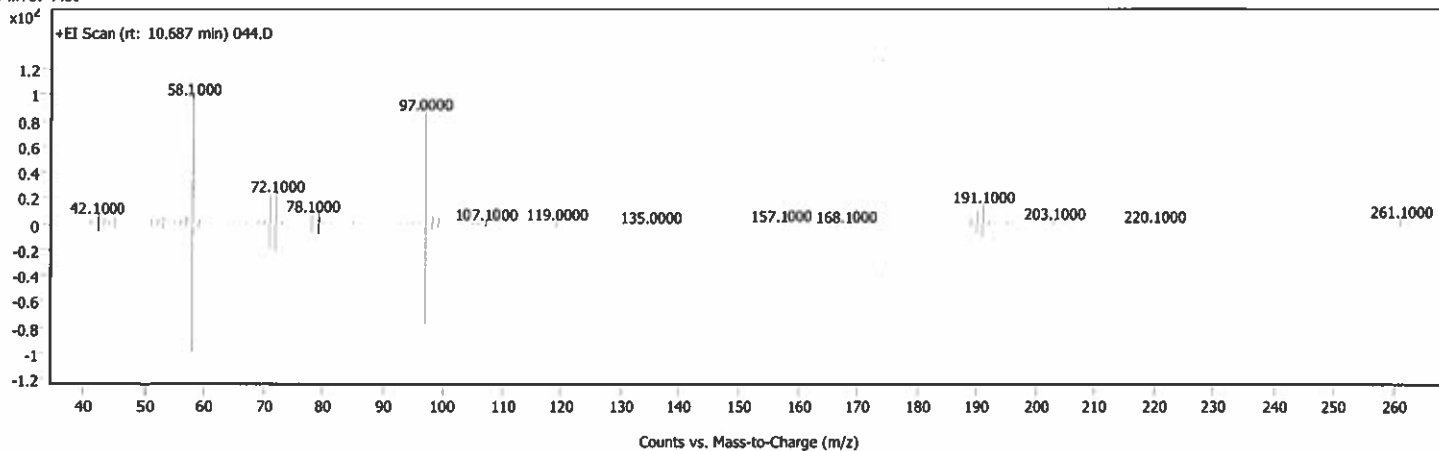
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	92.98	SWGDRUG 3.3.L	MATRIX 10 W/ISTD ONLY- LVR	KM/QC DATA IN XYLAZINE VALIDATION

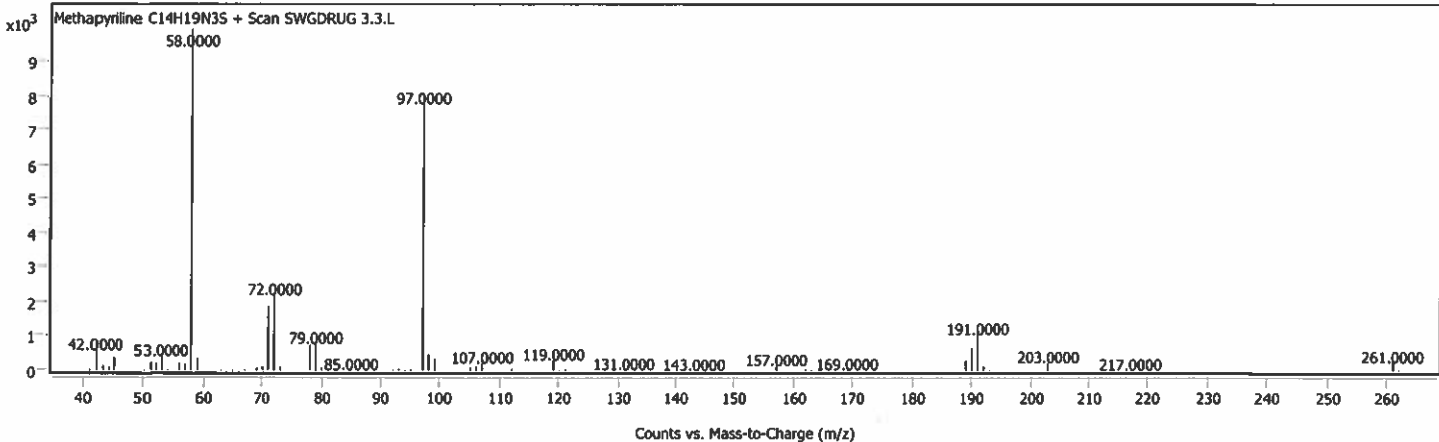
### Observed Spectrum



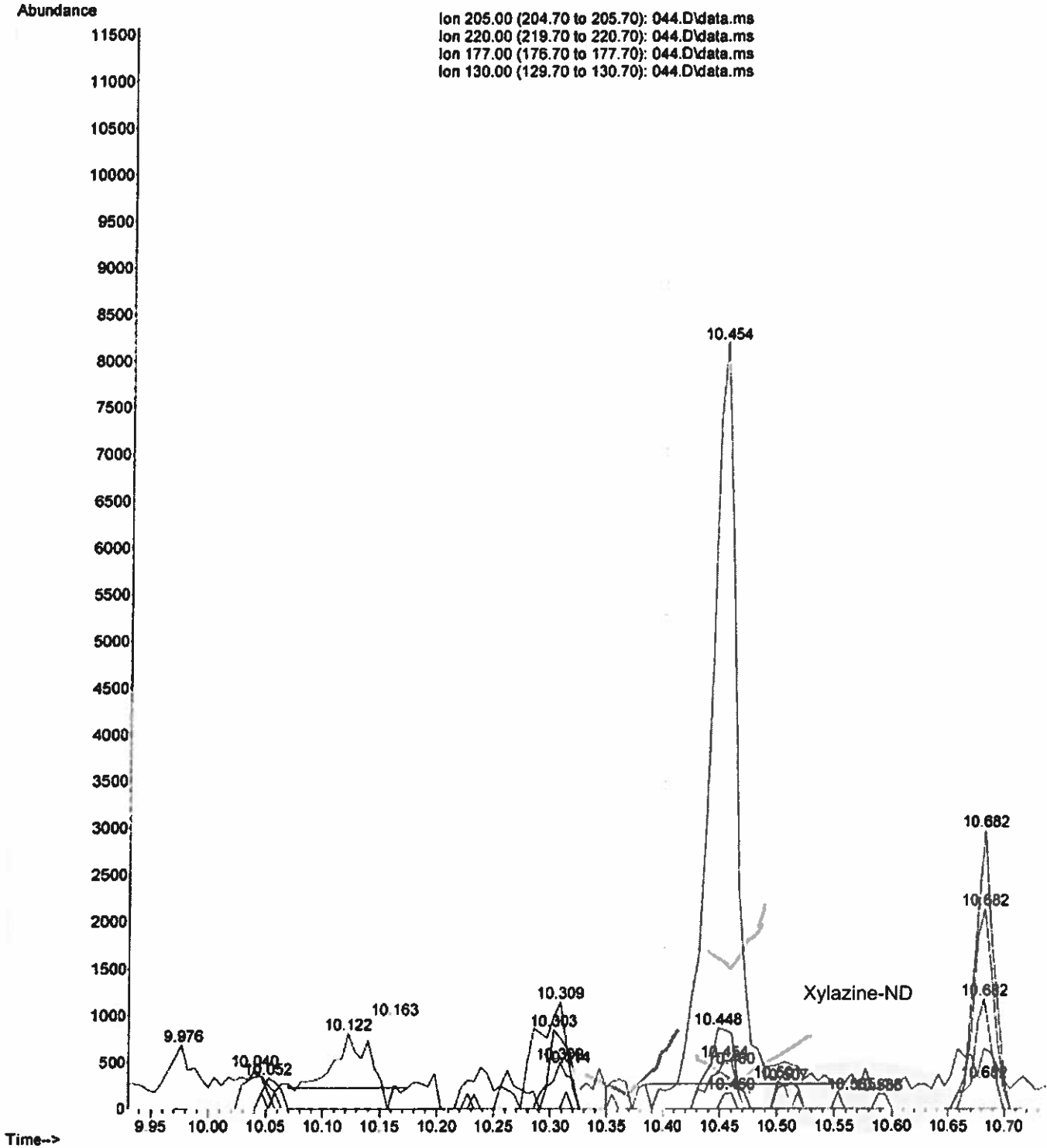
### Mirror Plot



### Library Spectrum



File :C:\Users\TOX\Desktop\092821\044.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 10:46 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 10 W/ISTD ONLY- LVR  
Misc Info :  
Vial Number: 76

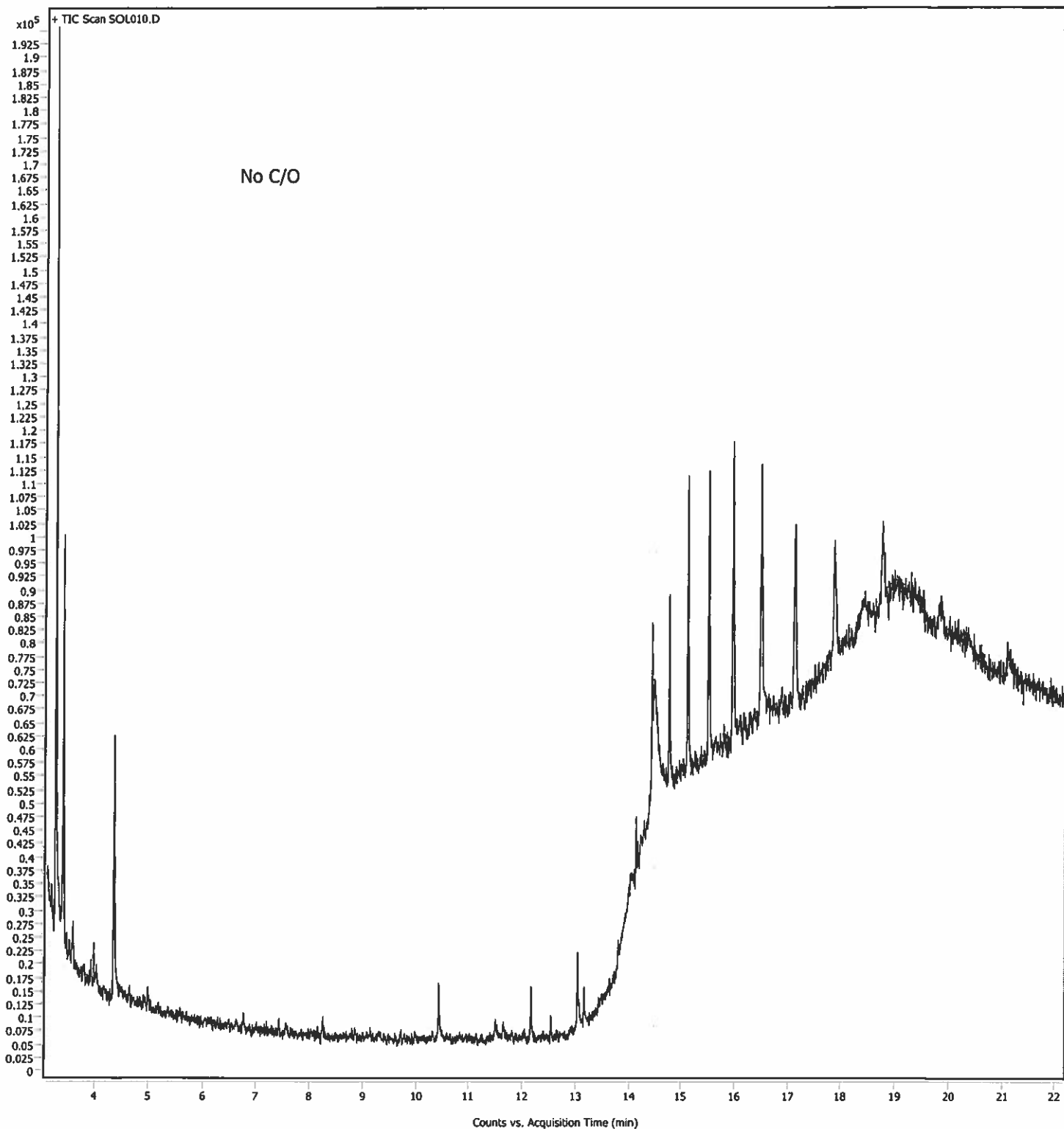


# Analysis Report

## Sample Information

Sample Name	SOLVENT AFTER MATRIX 10	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092321\SOL010.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/23/2021 5:44:18 PM (UTC-04:00)
Position	54	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

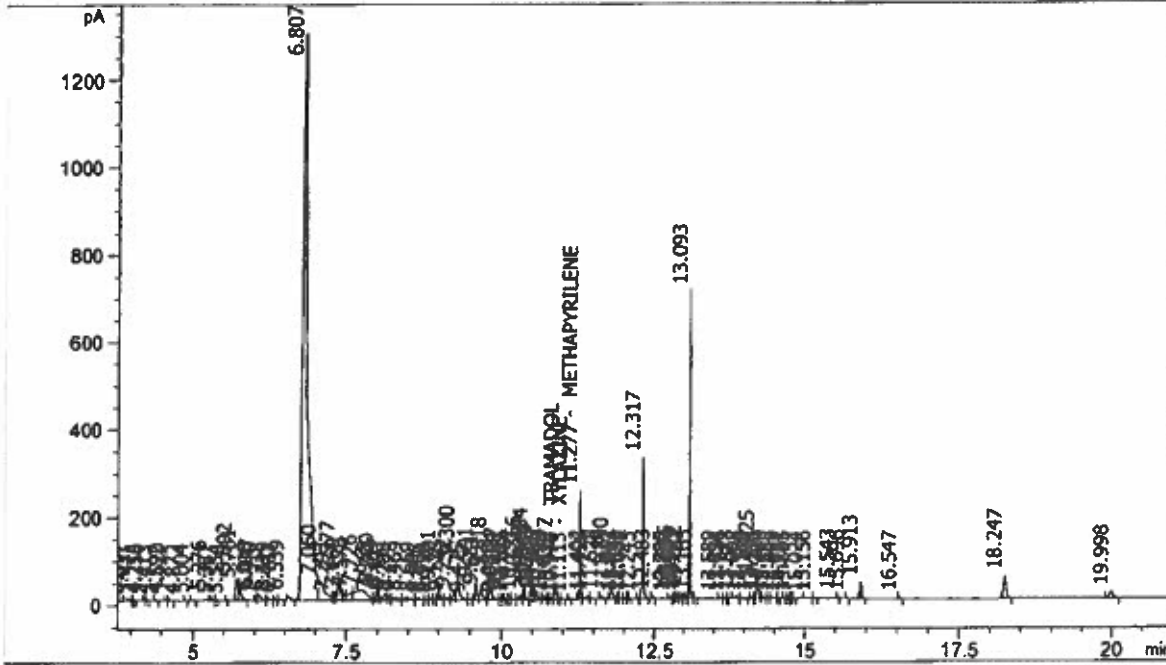
## Sample Chromatograms



Injection Date: 9/29/2021 10:01:28 AM  
 Sample Name: MATRIX 10 W/ISTD ONLY->  
 Sample Info:

Seq Line: 51  
 Vial 82

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

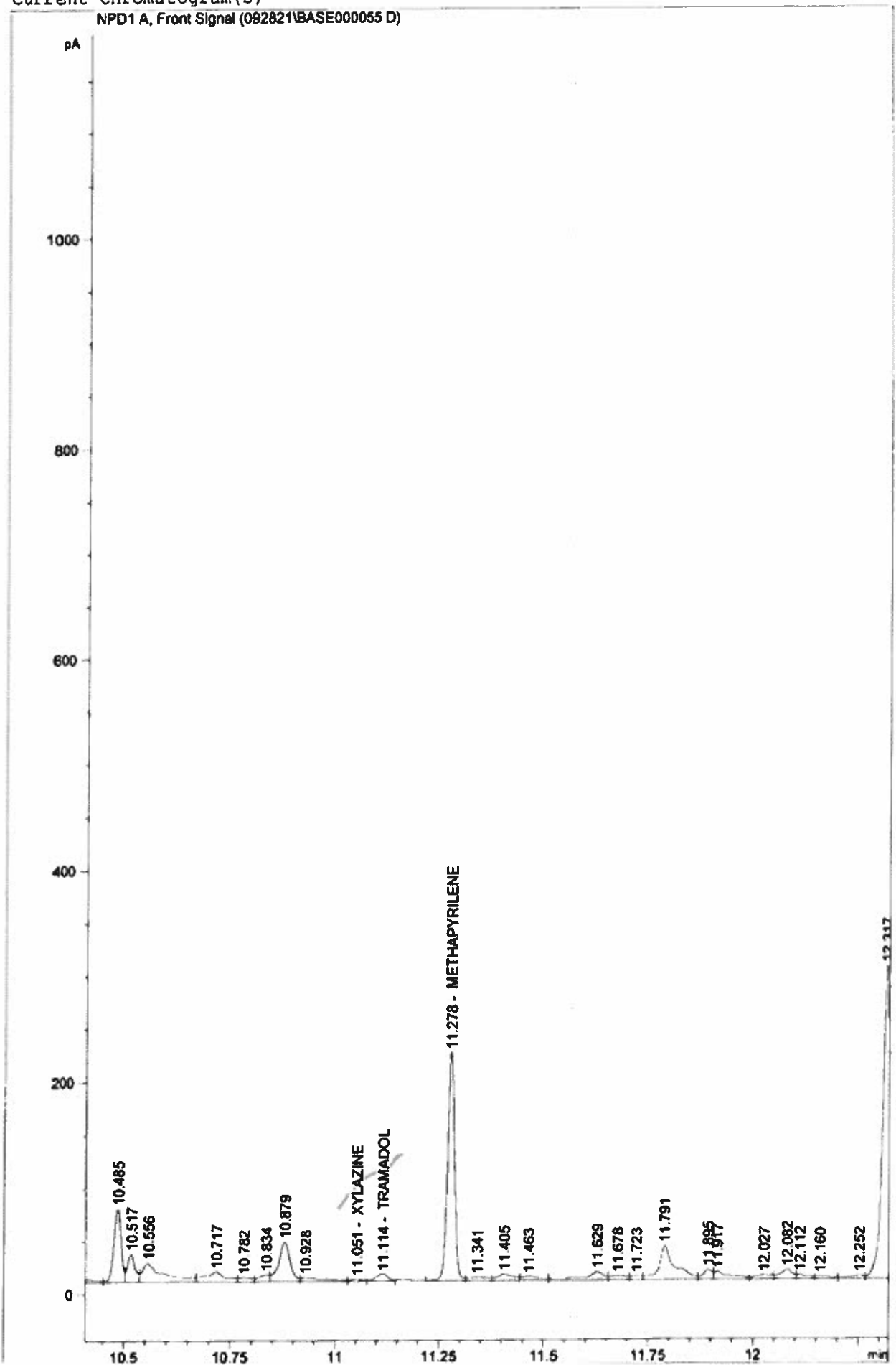


see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
10.978	11.033	5.145	2.034	0.00000	TRAMADOL > ND
11.113	11.046	5.260	2.223	0.00000	XYLAZINE
11.277	11.278	292.603	246.618	1.00000	METHAPYRILENE

Current Chromatogram (s)

NPD1 A, Front Signal (092821\BASE000055 D)

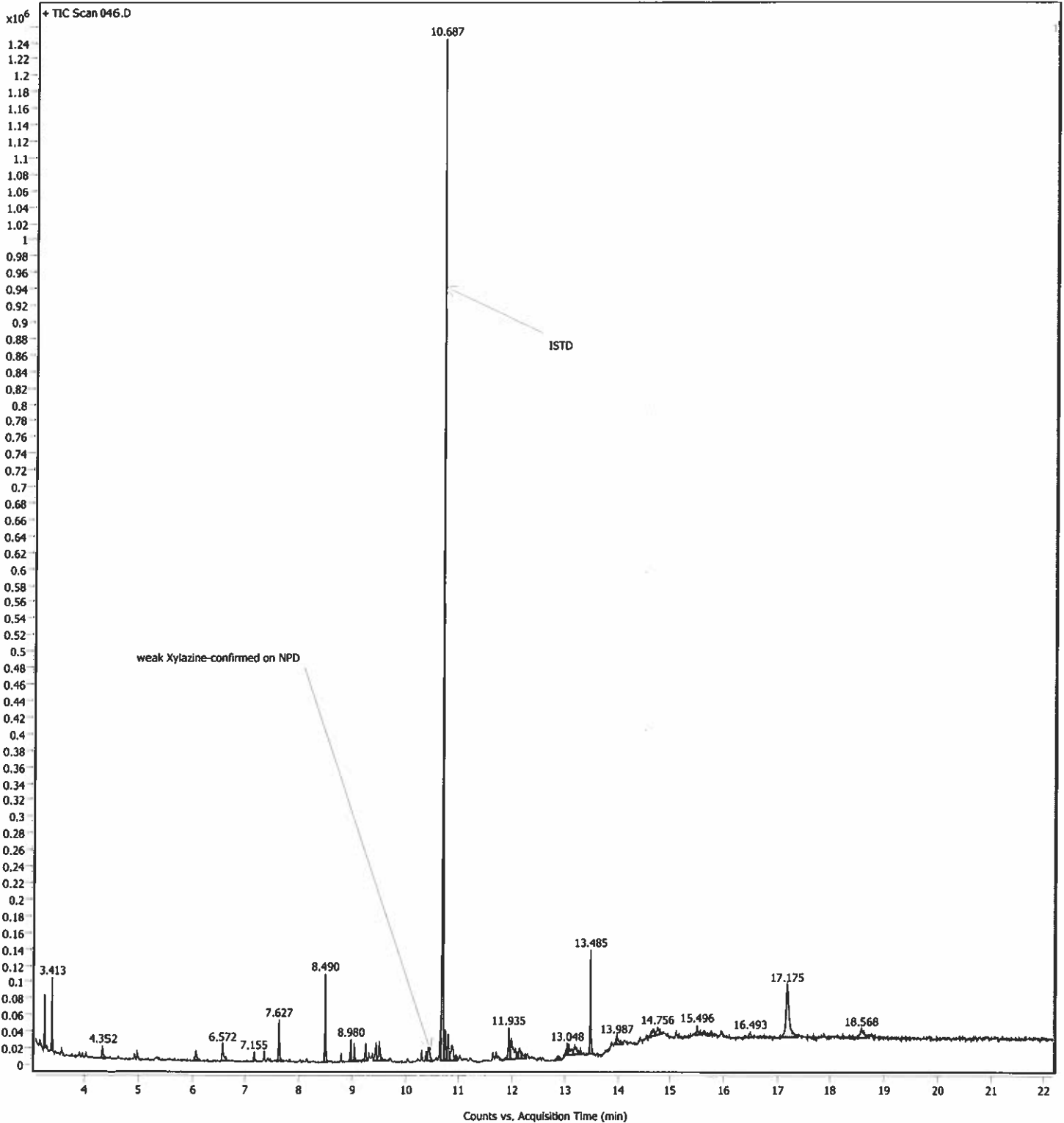


# Analysis Report

## Sample Information

Sample Name	MATRIX 2/W ISTD 0.010 mg/L- BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\046.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 11:38:09 AM (UTC-04:00)
Position	78	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



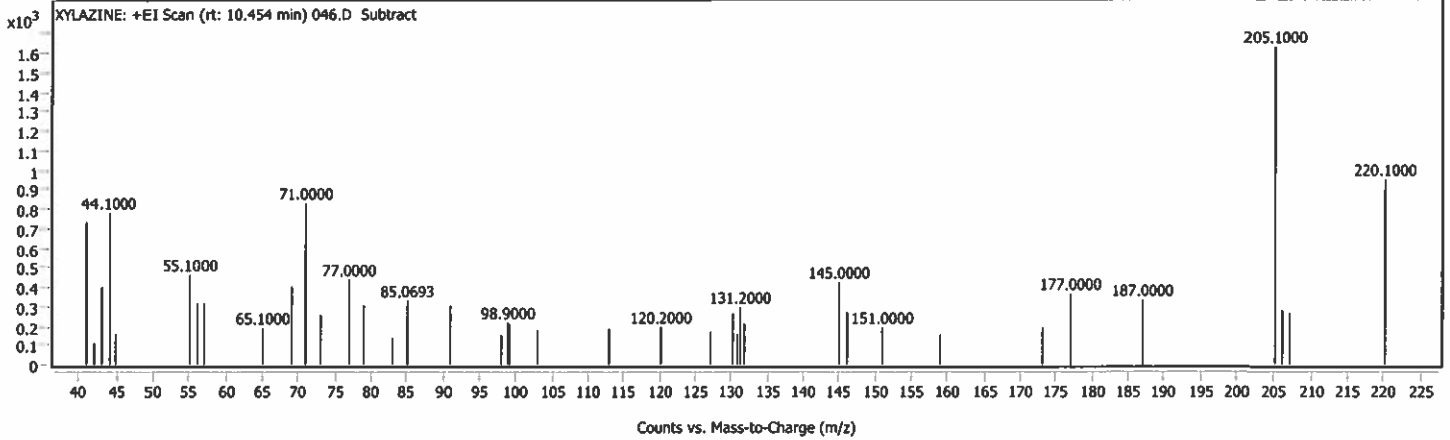
# Analysis Report

+ Scan (rt: 10.454 min) Sub

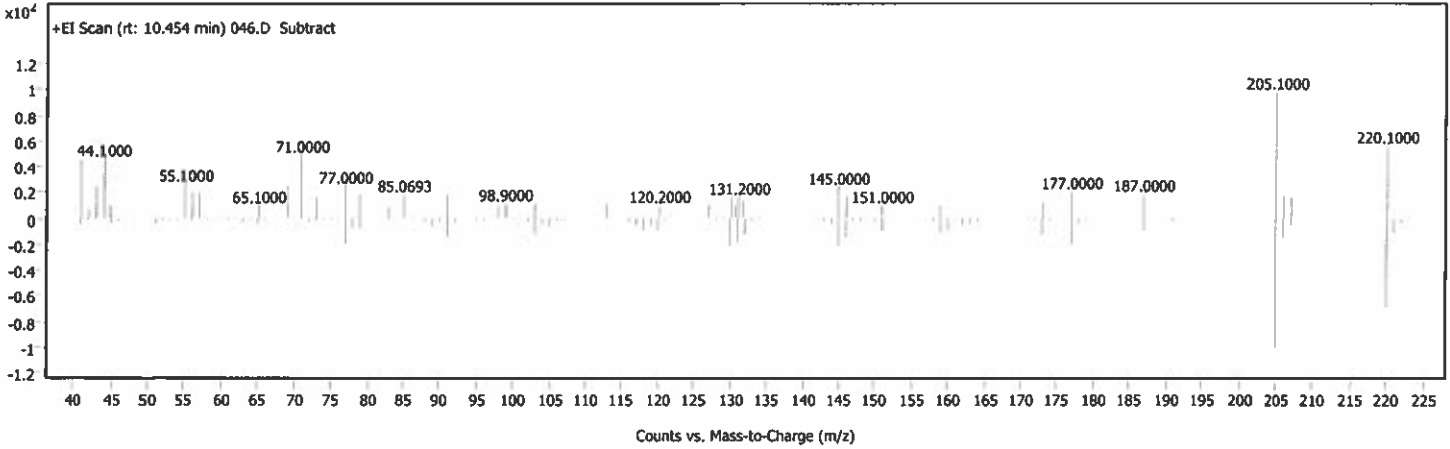
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	61.46	aafs.I	MATRIX 2/W ISTD 0.010 mg/L-BLD	KM/QC DATA IN XYLAZINE VALIDATION

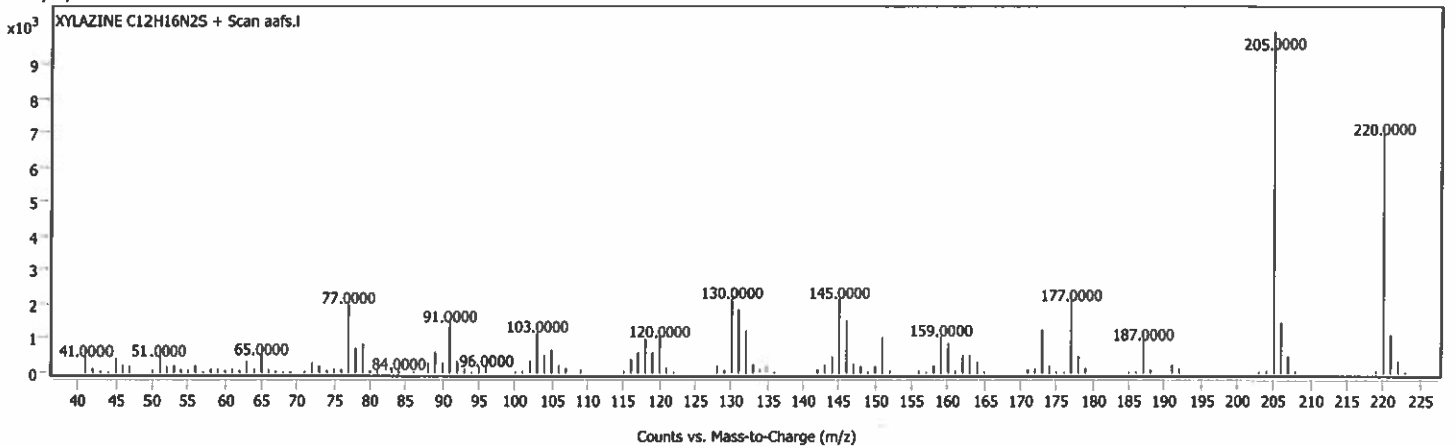
## Observed Spectrum



## Mirror Plot



## Library Spectrum



# Analysis Report

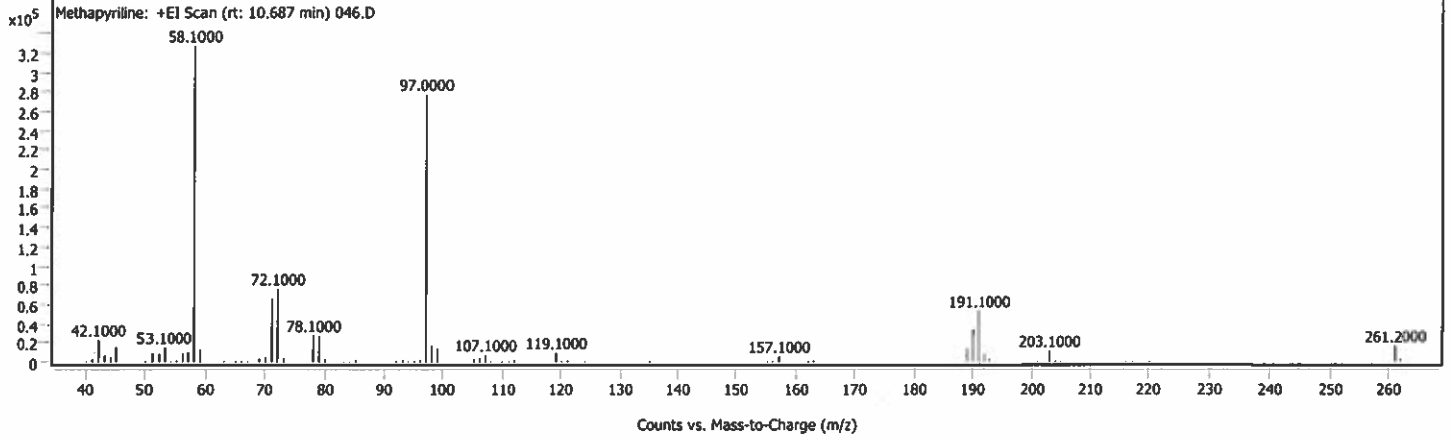
## Sample Spectra

+ Scan (rt: 10.687 min)

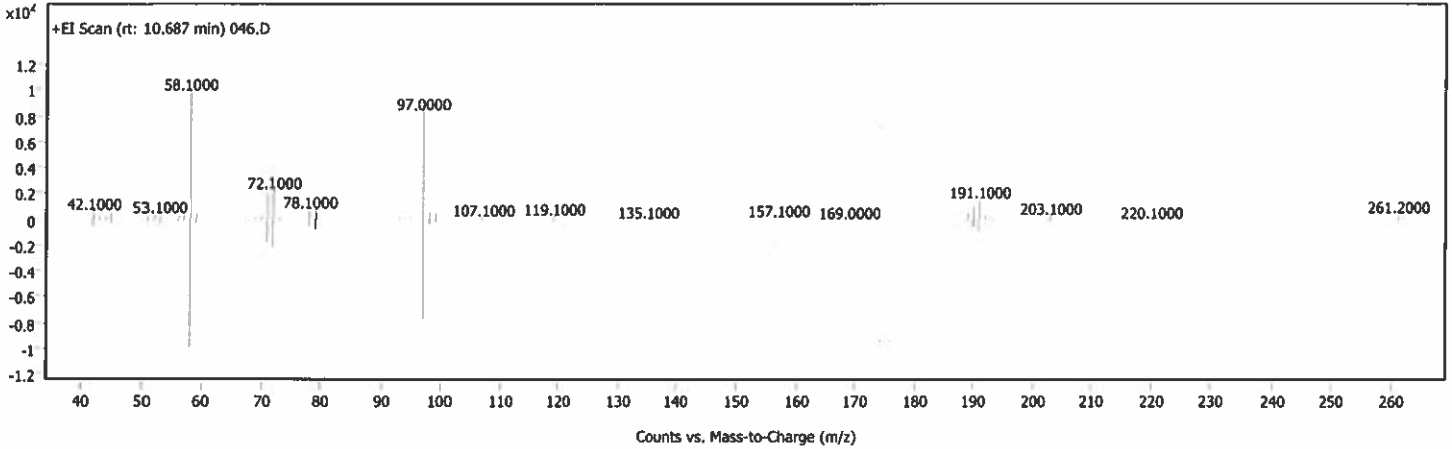
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.23	SWGDRUG 3.3.L	MATRIX 2/W ISTD 0.010 mg/L-BLD	KM/QC DATA IN XYLAZINE VALIDATION

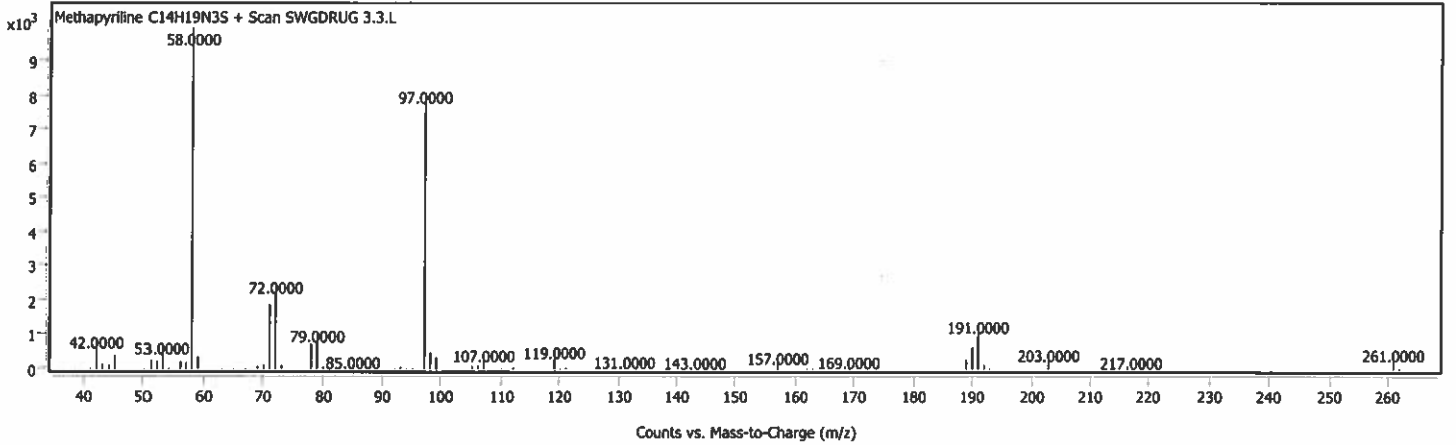
## Observed Spectrum



## Mirror Plot



## Library Spectrum

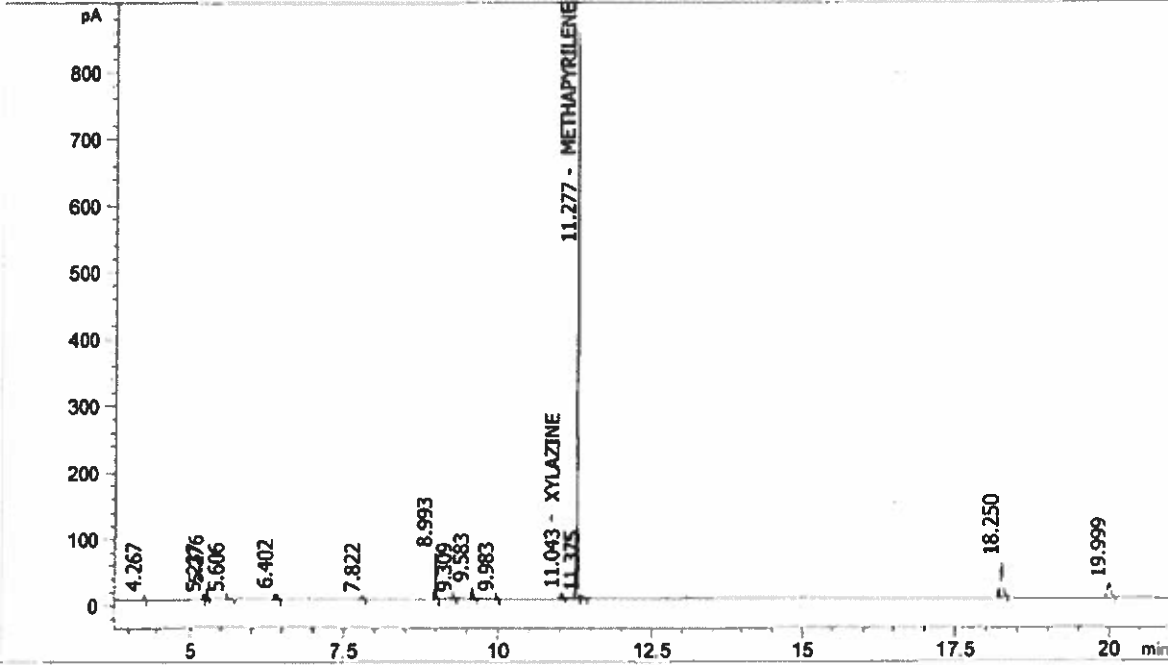




Injection Date: 9/29/2021 10:50:10 AM  
 Sample Name: MATRIX 2 W/ISTD 0.010->  
 Sample Info:

Seq Line: 53  
 Vial 84

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



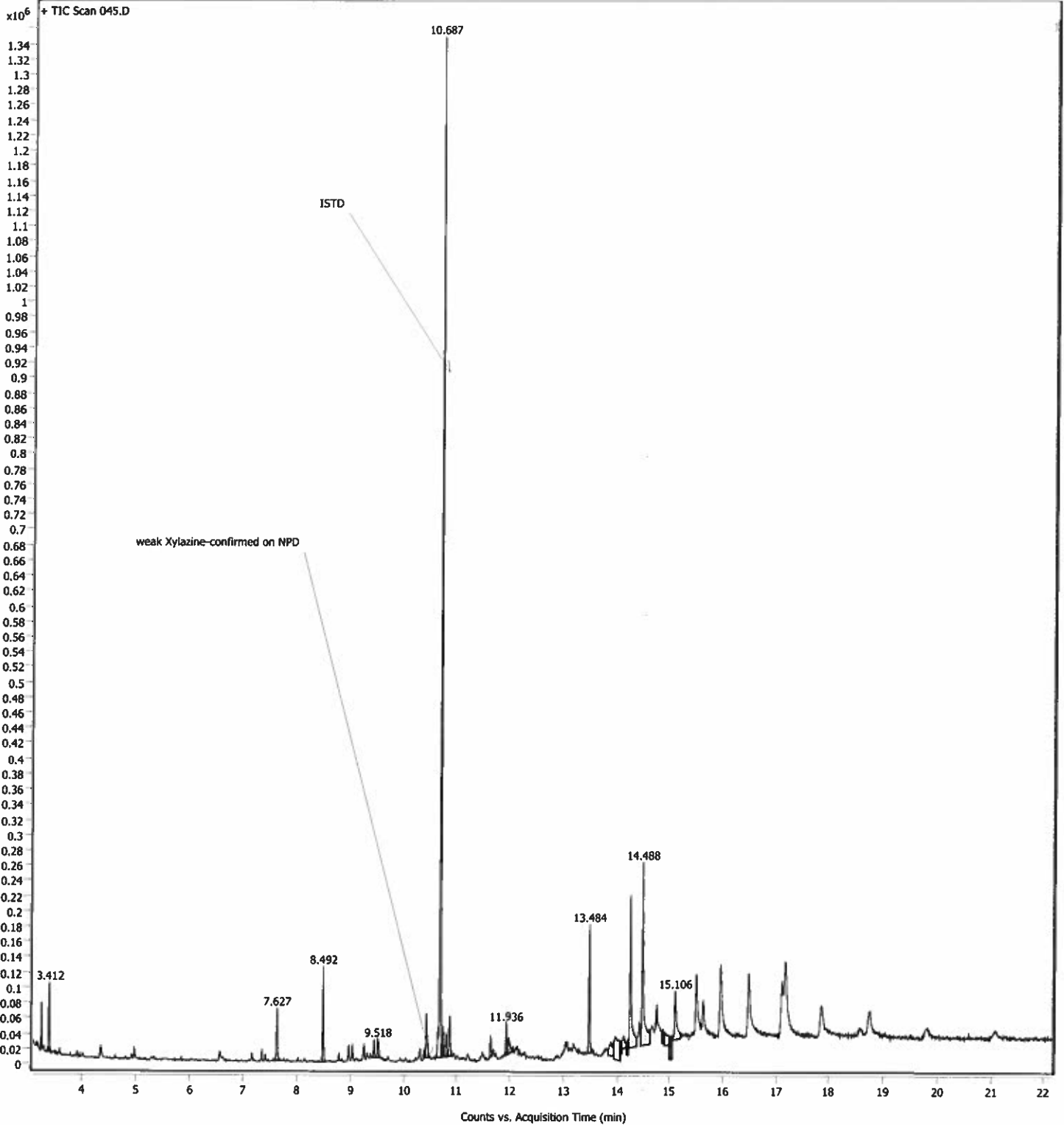
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.043	11.046	8.733	7.203	0.000000	XYLAZINE RRT - 0.9792
11.277	11.278	968.931	852.123	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 W/ISTD 0.010 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\045.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 11:12:22 AM (UTC-04:00)
Position	77	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms

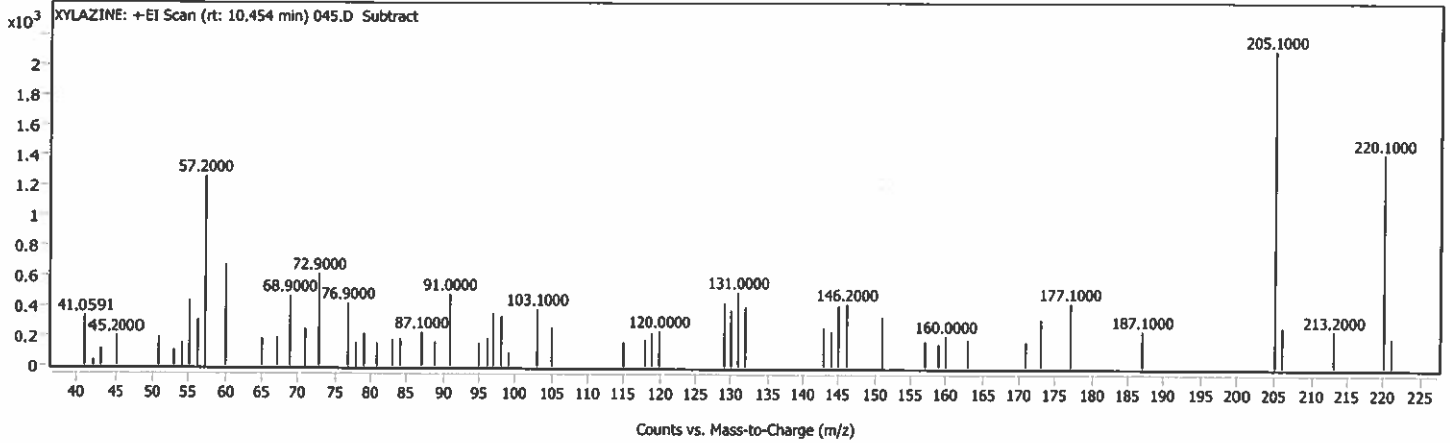


# Analysis Report

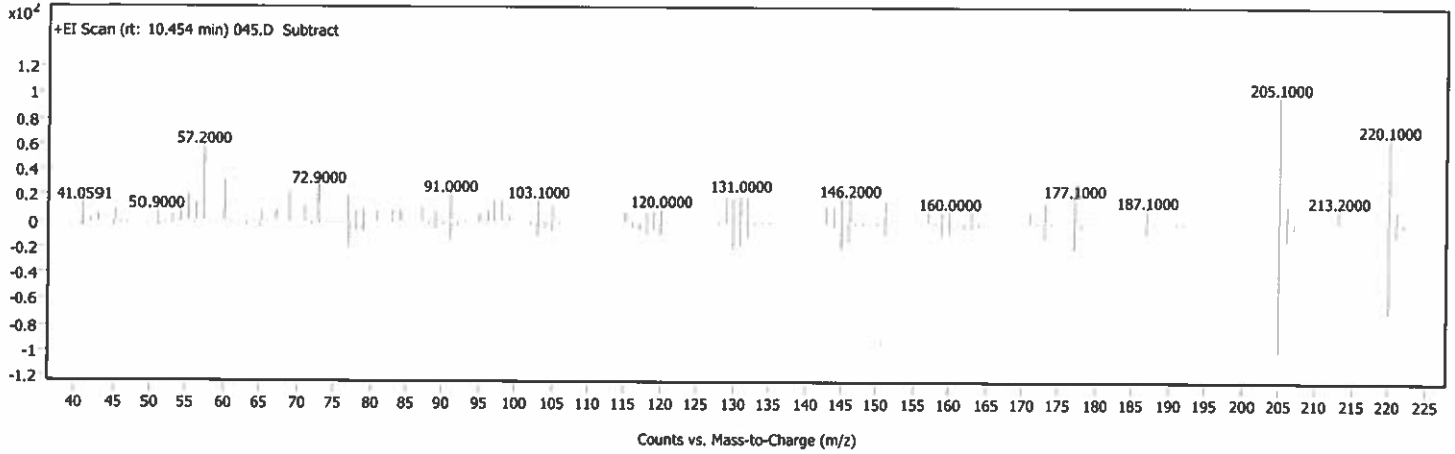
+ Scan (rt: 10.454 min) Sub **XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	70.14	aafs.l	MATRIX 1 W/ISTD 0.010 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

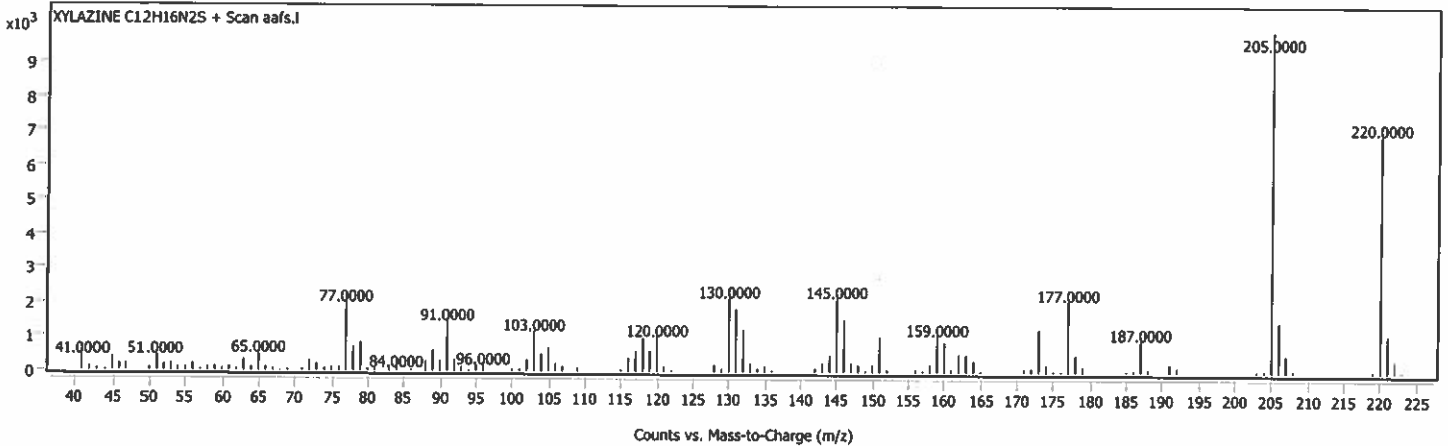
## Observed Spectrum



## Mirror Plot



## Library Spectrum



# Analysis Report

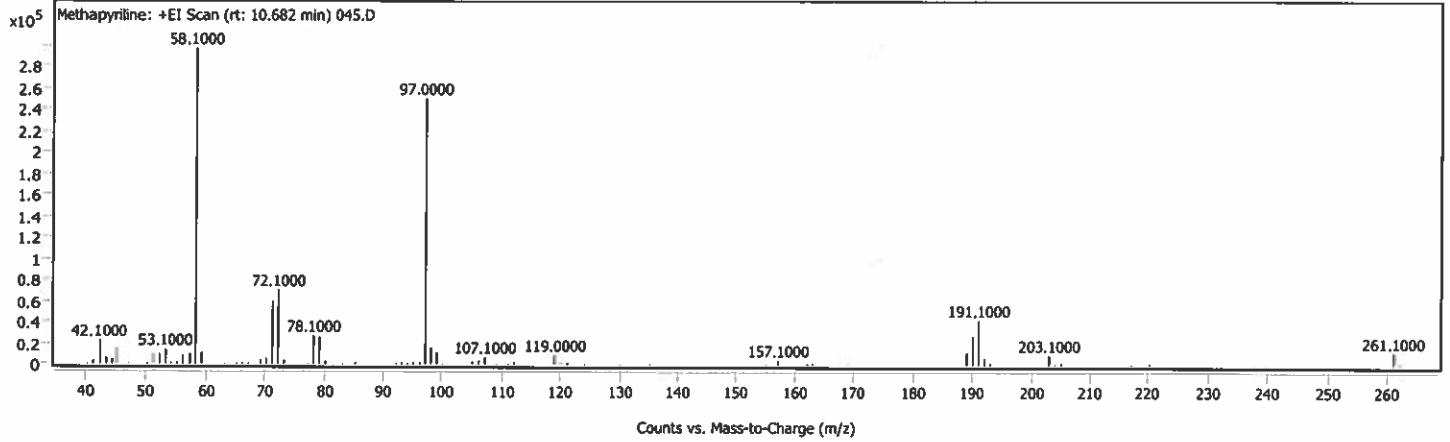
## Sample Spectra

+ Scan (rt: 10.682 min)

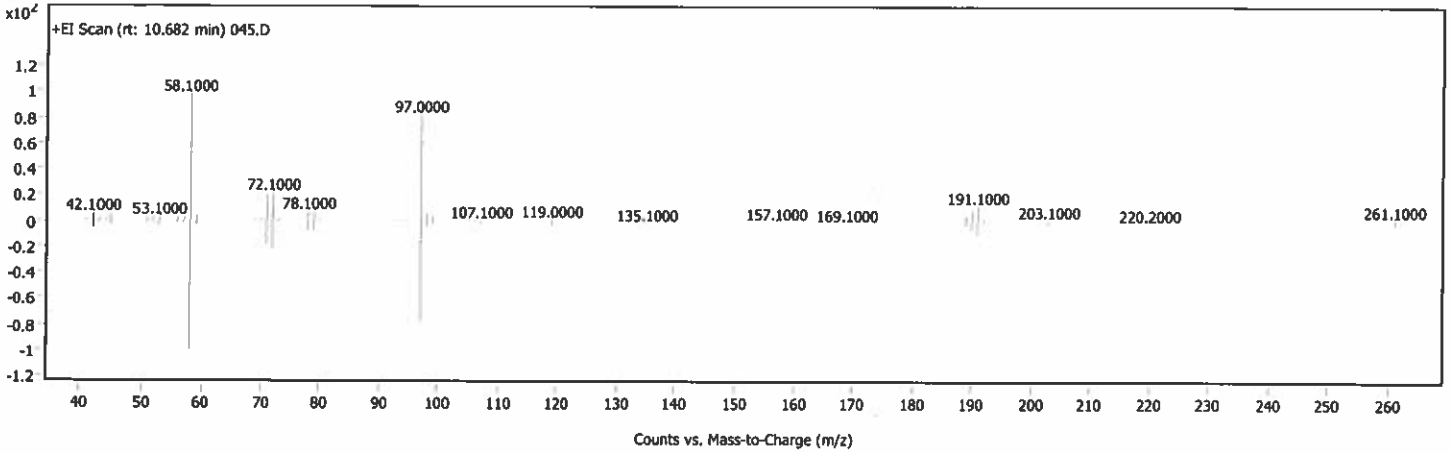
Methapyriline; C<sub>14</sub>H<sub>19</sub>N<sub>3</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C <sub>14</sub> H <sub>19</sub> N <sub>3</sub> S	98.38	SWGDRUG 3.3.L	MATRIX 1 W/ISTD 0.010 mg/L - BLD	KM/QC DATA IN XLAZINE VALIDATION

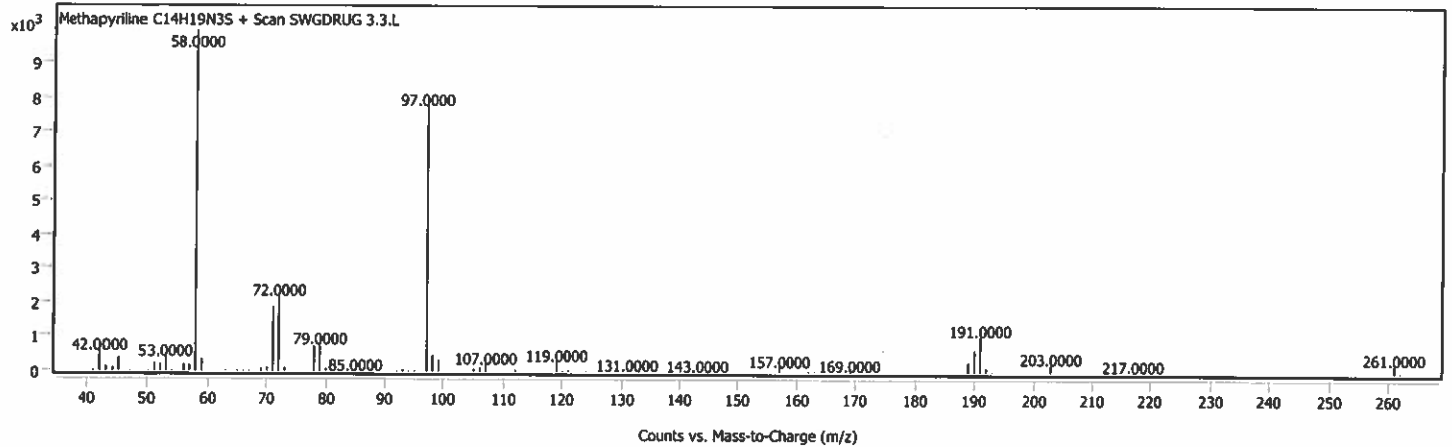
### Observed Spectrum



### Mirror Plot

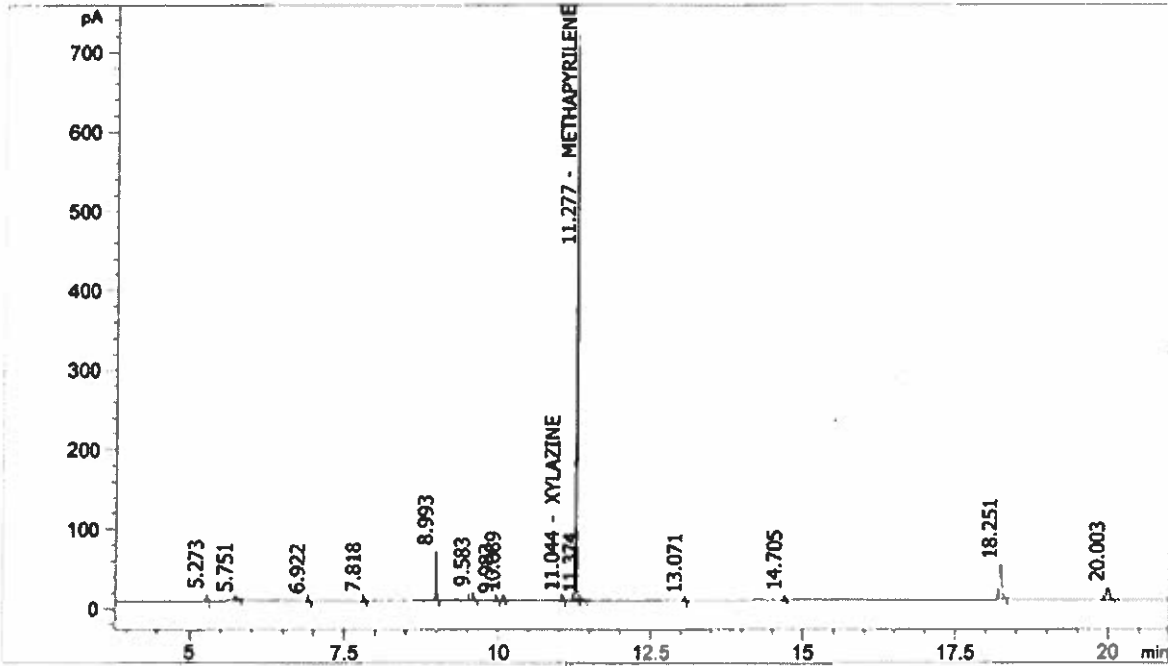


### Library Spectrum



-----  
Injection Date: 9/29/2021 10:25:52 AM Seq Line: 52  
Sample Name: MATRIX 1 W/ISTD 0.010-> Vial 83  
Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION

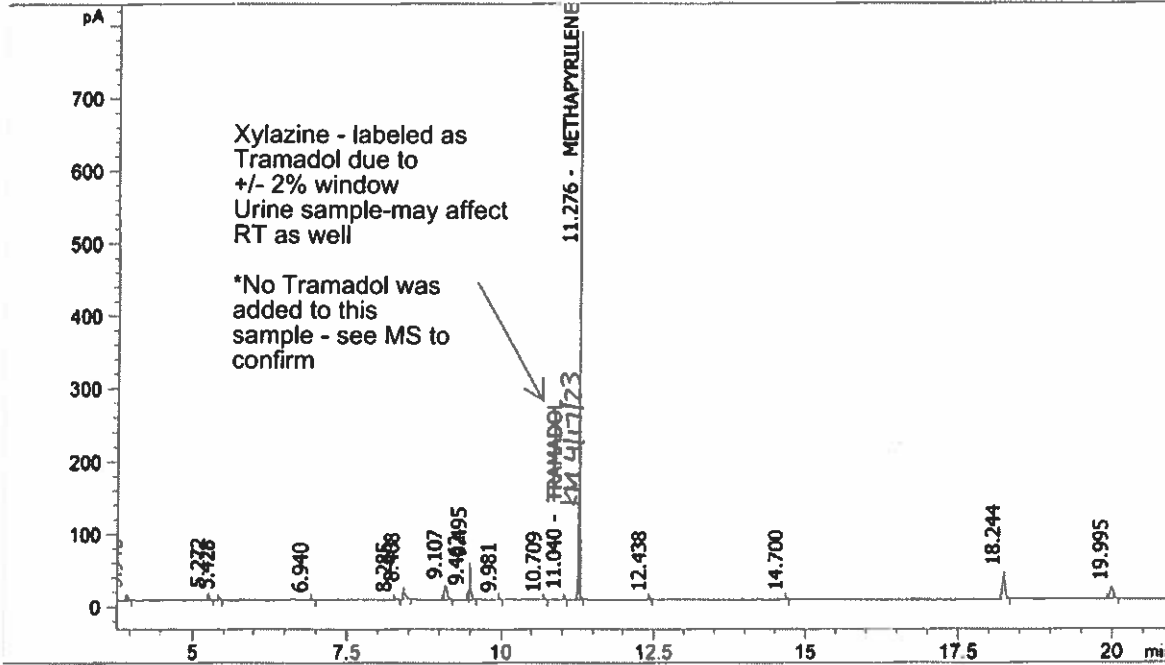


RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.044	11.046	6.392	5.359	0.000000	XYLAZINE RRT - 0.9793
11.277	11.278	802.430	712.589	1.000000	METHAPYRILENE

Injection Date: 9/29/2021 11:14:27 AM  
 Sample Name: MATRIX 6 W/ISTD 0.010->  
 Sample Info:

Seq Line: 54  
 Vial 85

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



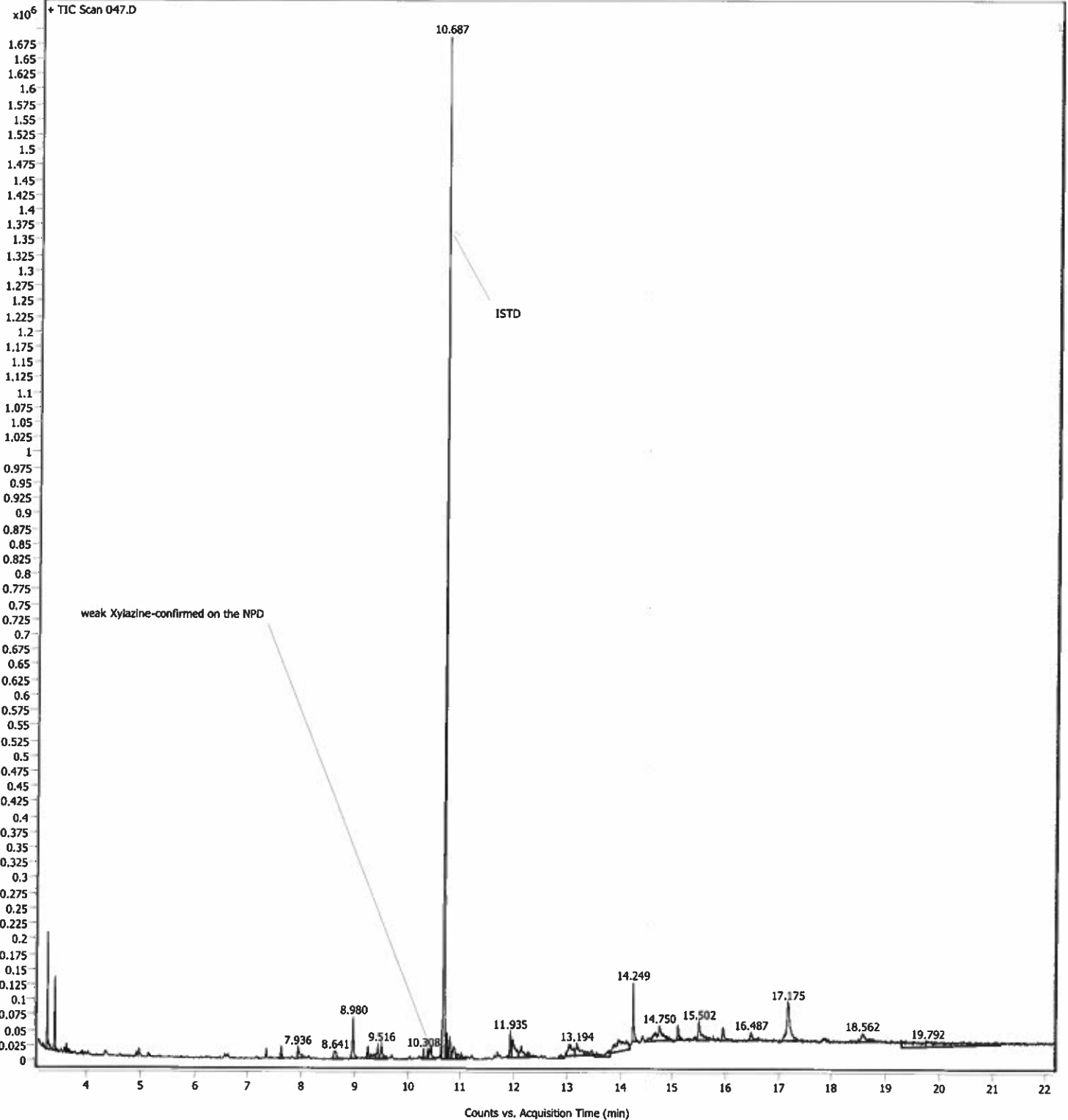
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.040	11.033	7.259	6.226	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE RRT - 0.9791
11.276	11.278	890.075	779.668	1.000000	METHAPYRILENE

# Analysis Report

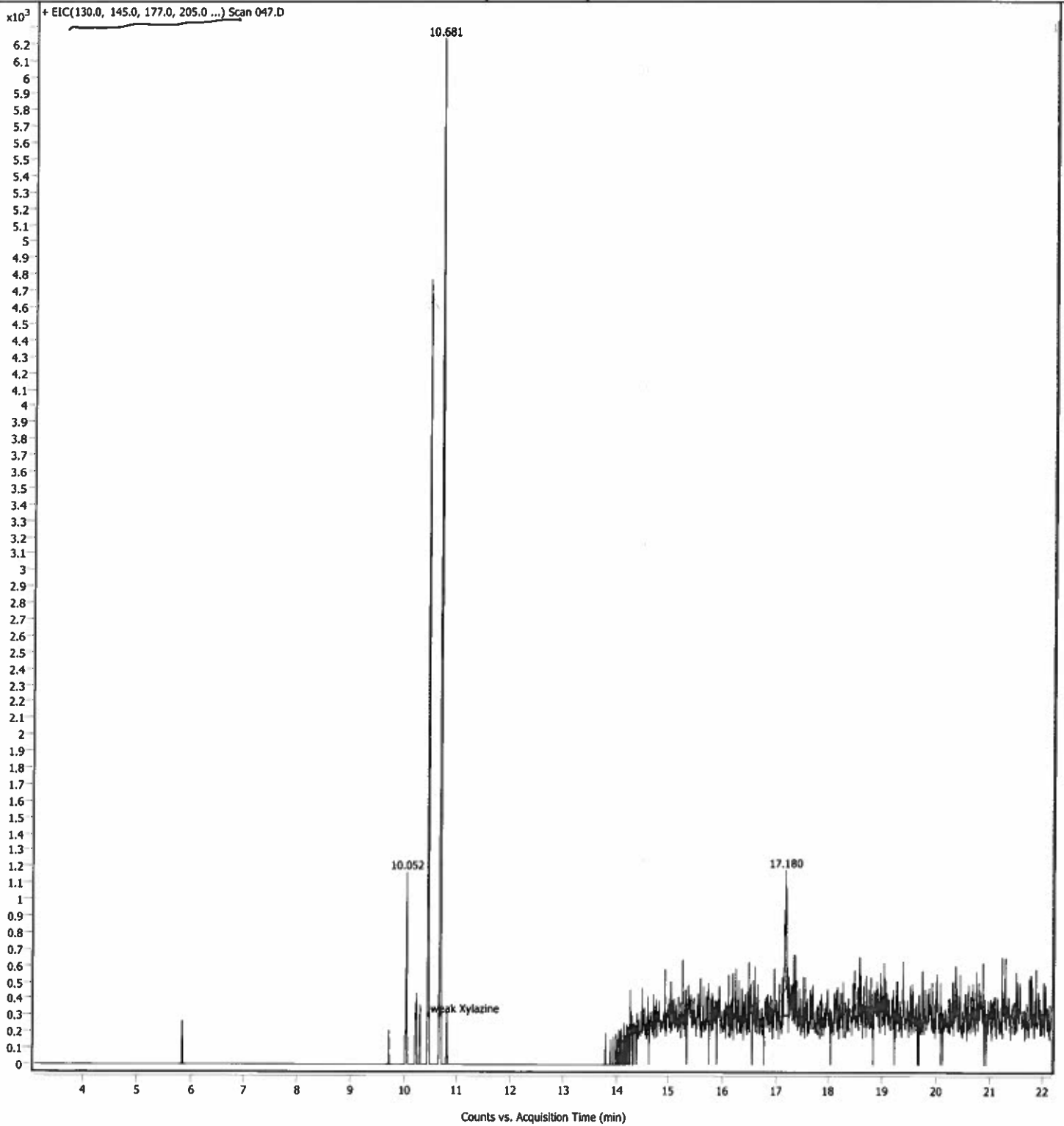
## Sample Information

Sample Name	MATRIX 6 W/ISTD 0.010 mg/L-URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\047.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 12:03:59 PM (UTC-04:00)
Position	79	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report



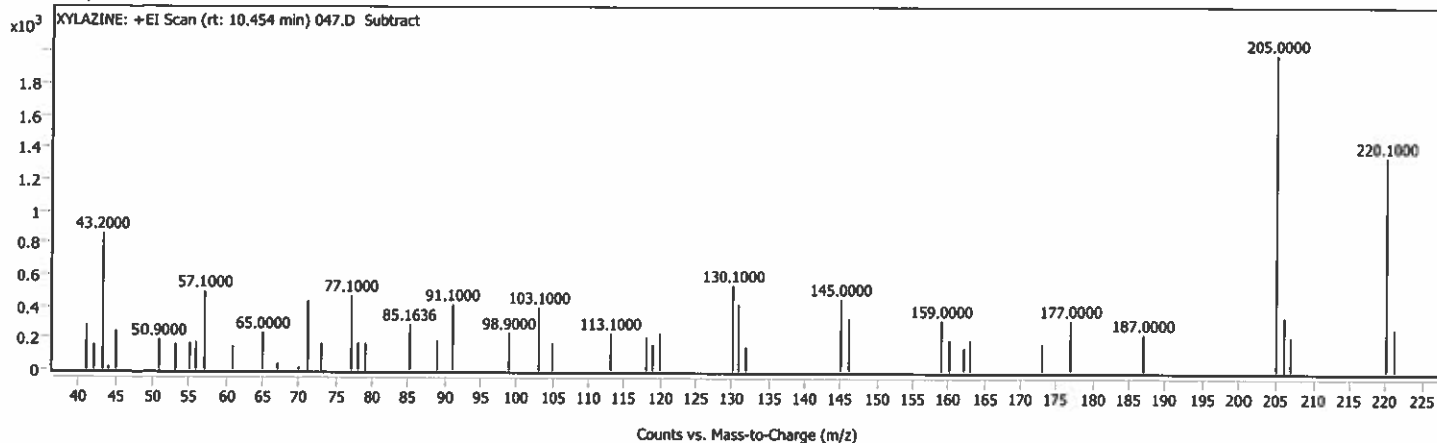


# Analysis Report

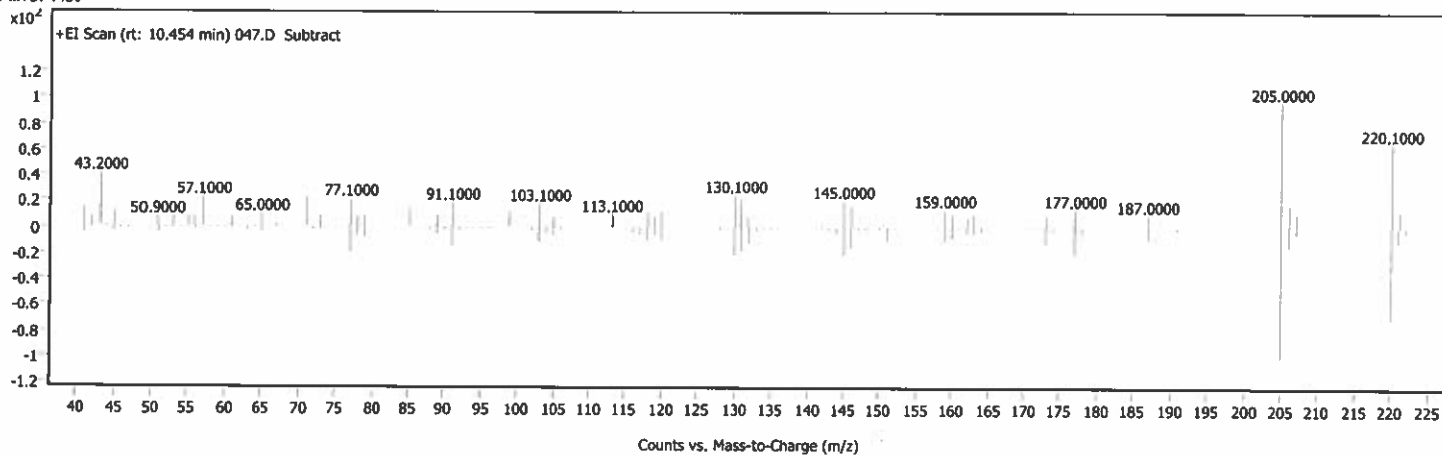
+ Scan (rt: 10.454 min) Sub **XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	74.20	aafs.l	MATRIX 6 W/ISTD 0.010 mg/L-URN	KM/QC DATA IN XYLAZINE VALIDATION

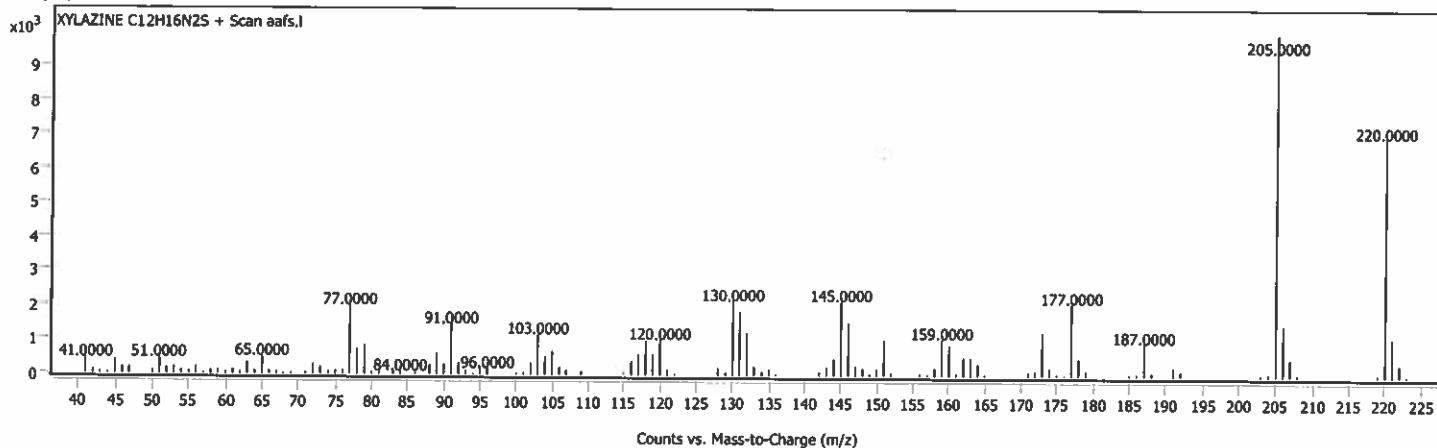
## Observed Spectrum



## Mirror Plot



## Library Spectrum



# Analysis Report

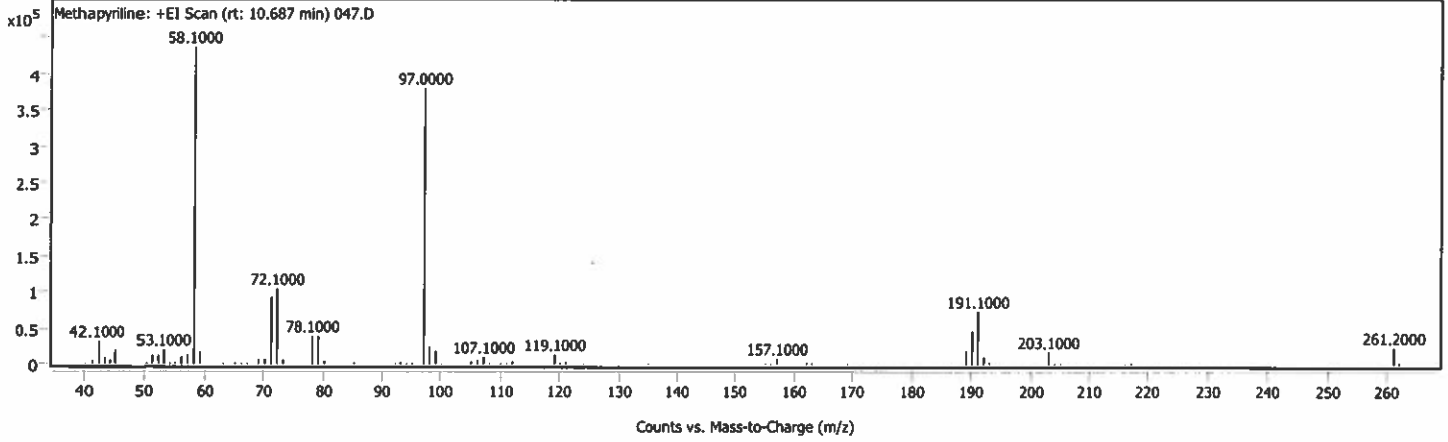
## Sample Spectra

+ Scan (rt: 10.687 min)

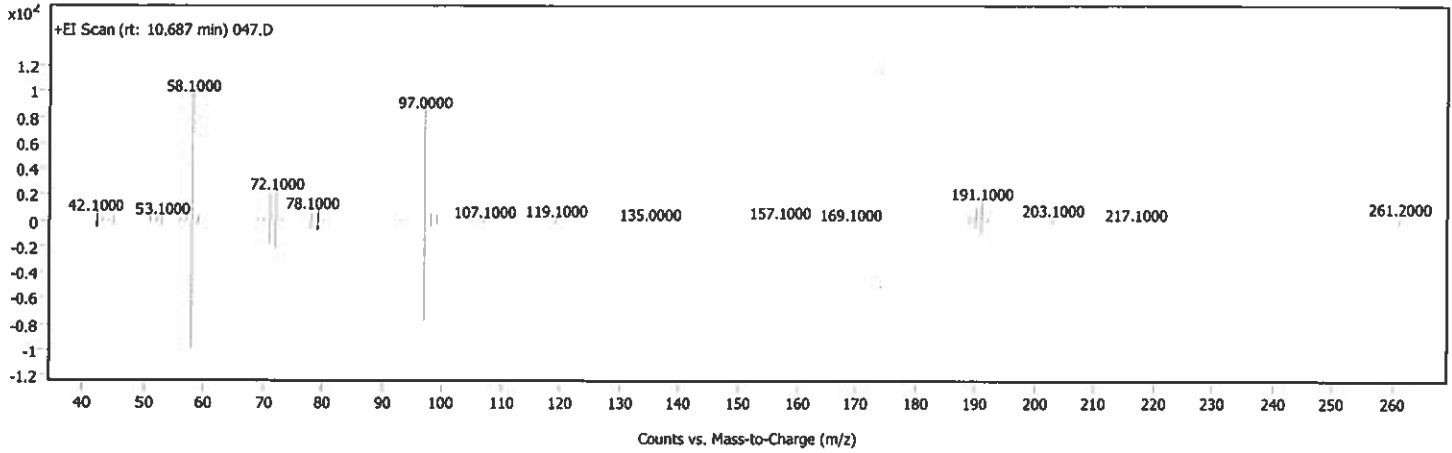
Methapyriline; C<sub>14</sub>H<sub>19</sub>N<sub>3</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C <sub>14</sub> H <sub>19</sub> N <sub>3</sub> S	98.63	SWGDRUG 3.3.L	MATRIX 6 W/ISTD 0.010 mg/L-URN	KM/QC DATA IN XLAZINE VALIDATION

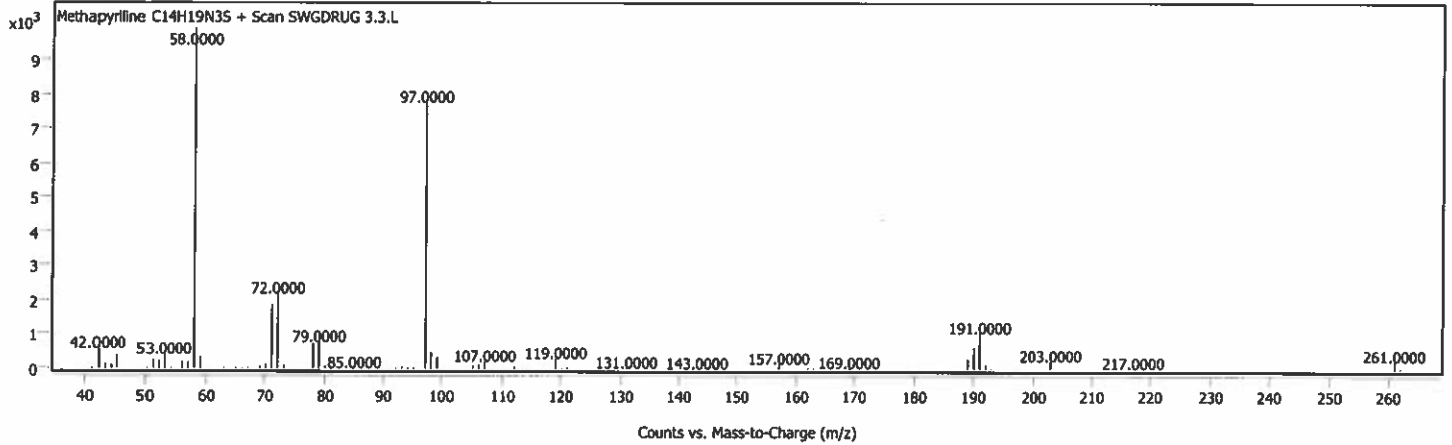
### Observed Spectrum



### Mirror Plot



### Library Spectrum

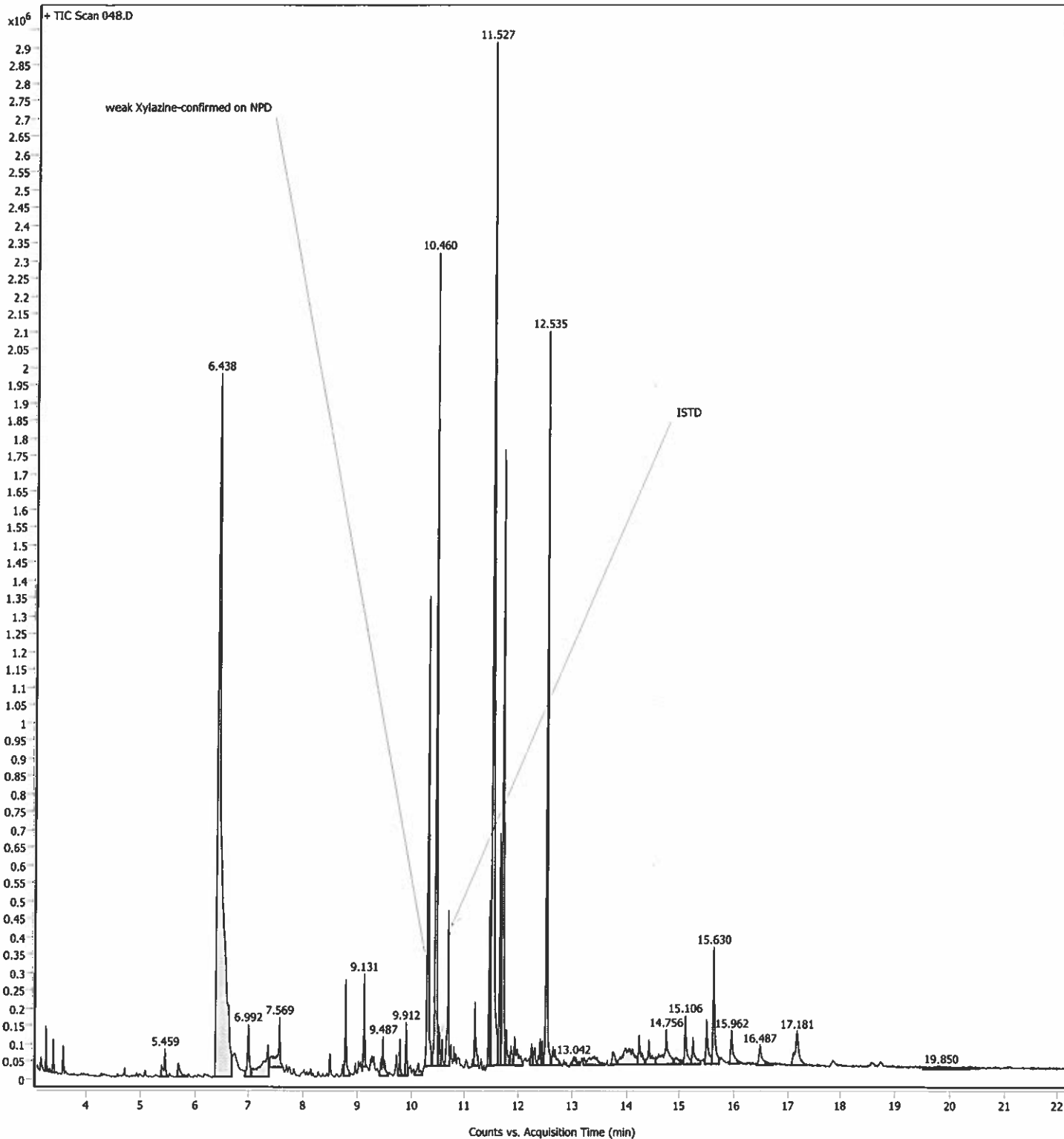


# Analysis Report

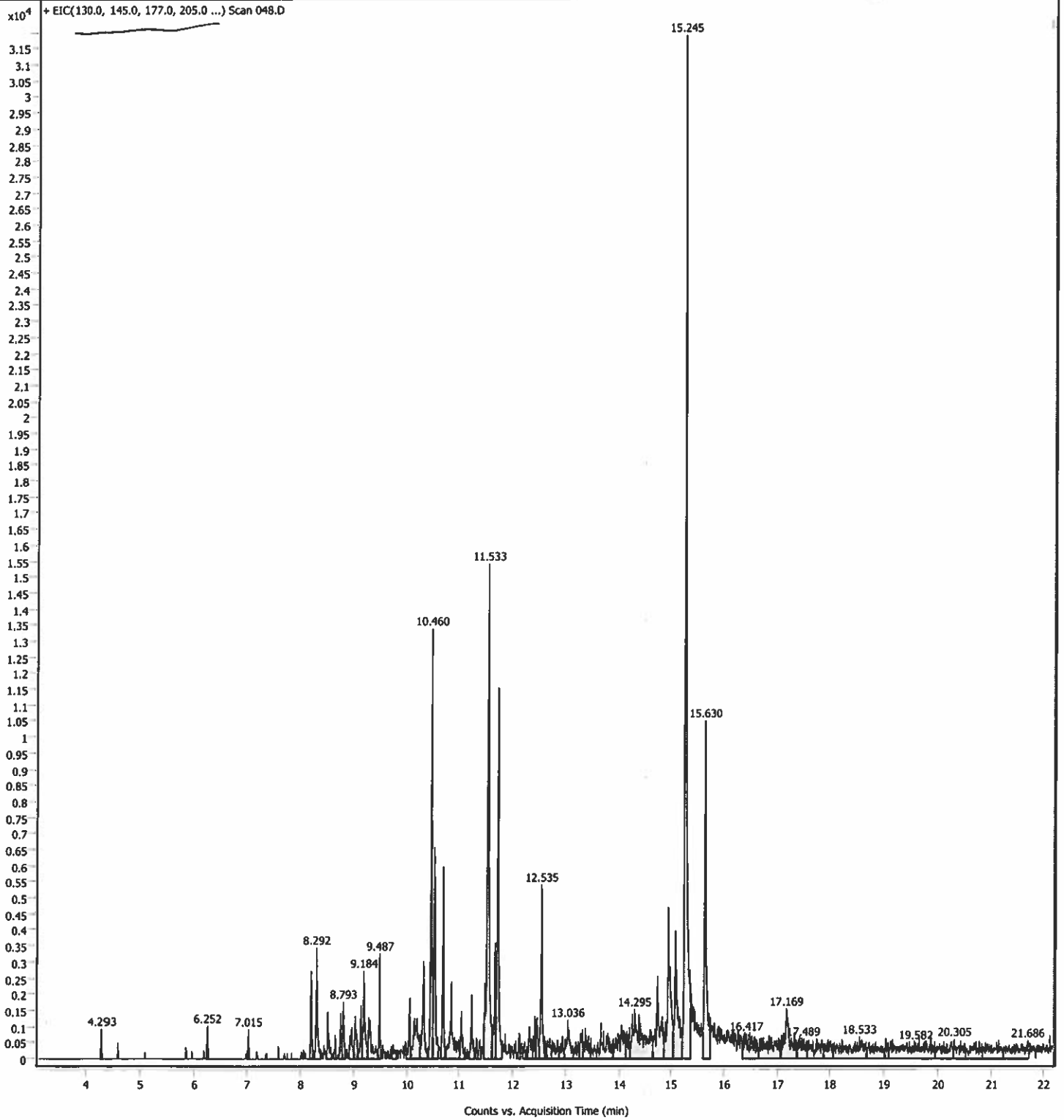
## Sample Information

Sample Name	MATRIX 10 W/ISTD 0.010 mg/L-LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\048.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 12:29:54 PM (UTC-04:00)
Position	80	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report



# Analysis Report

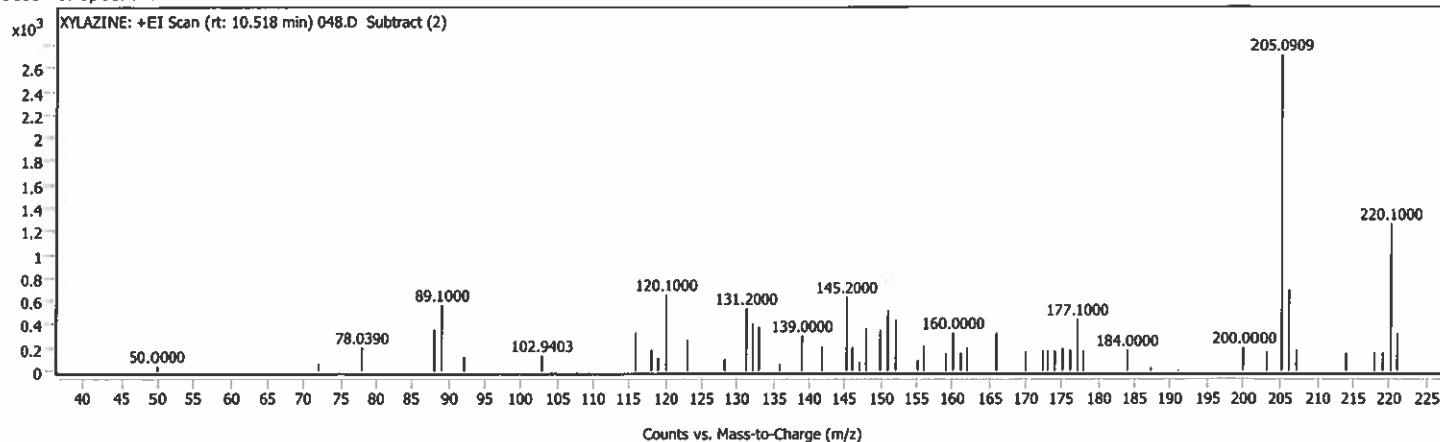
## Sample Spectra

+ Scan (rt: 10.687 min)

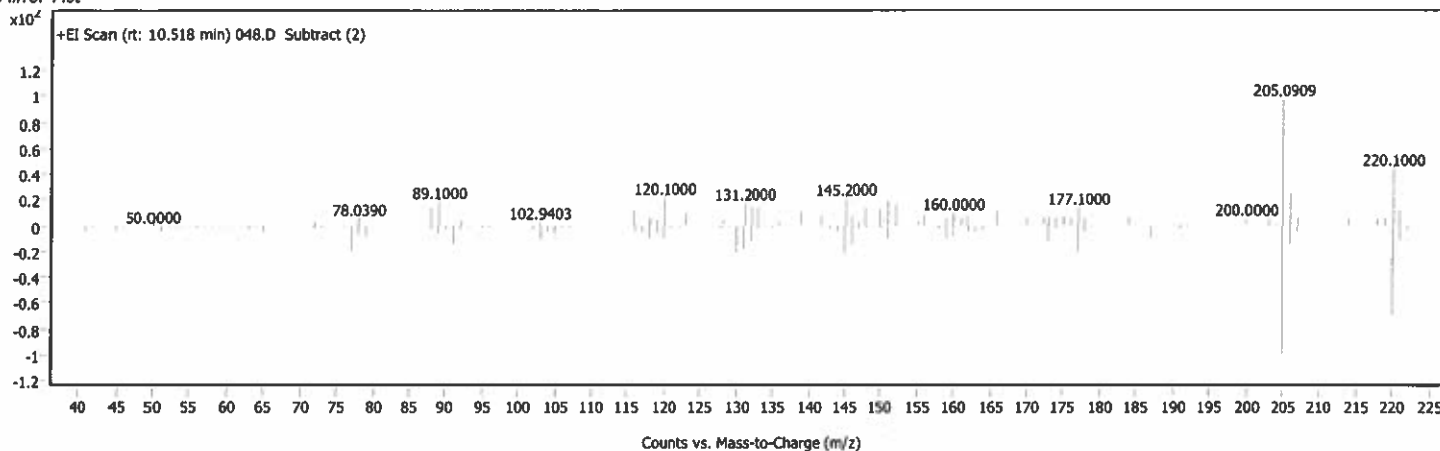
+ Scan (rt: 10.518 min) Sub (2) XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	60.99	aafs.l	MATRIX 10 W/ISTD 0.010 mg/L-LVR	KM/QC DATA IN XYLAZINE VALIDATION

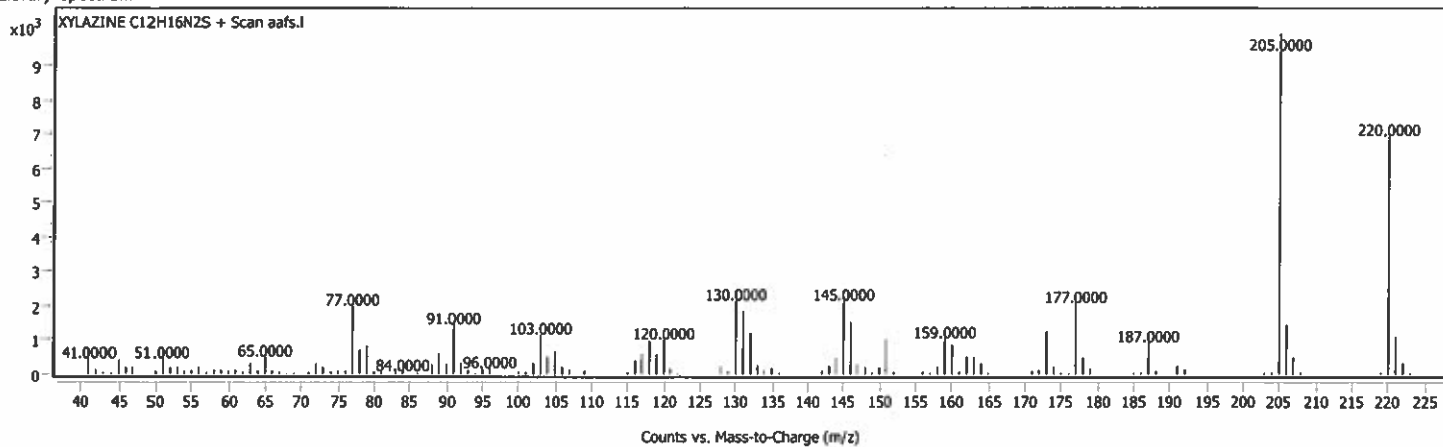
## Observed Spectrum



## Mirror Plot



## Library Spectrum



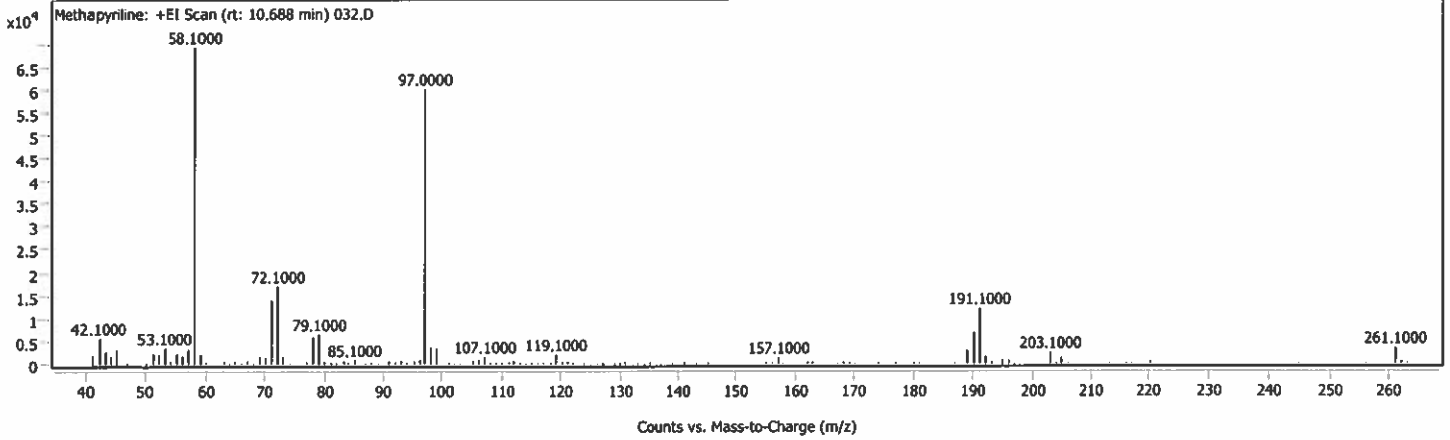
# Analysis Report

+ Scan (rt: 10.688 min)

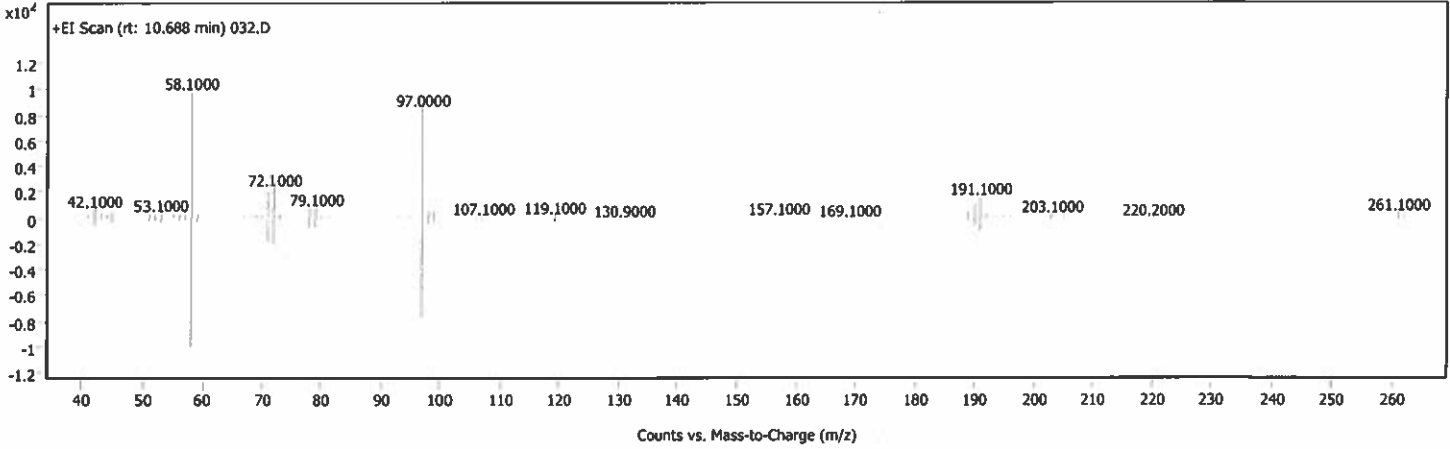
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	93.41	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

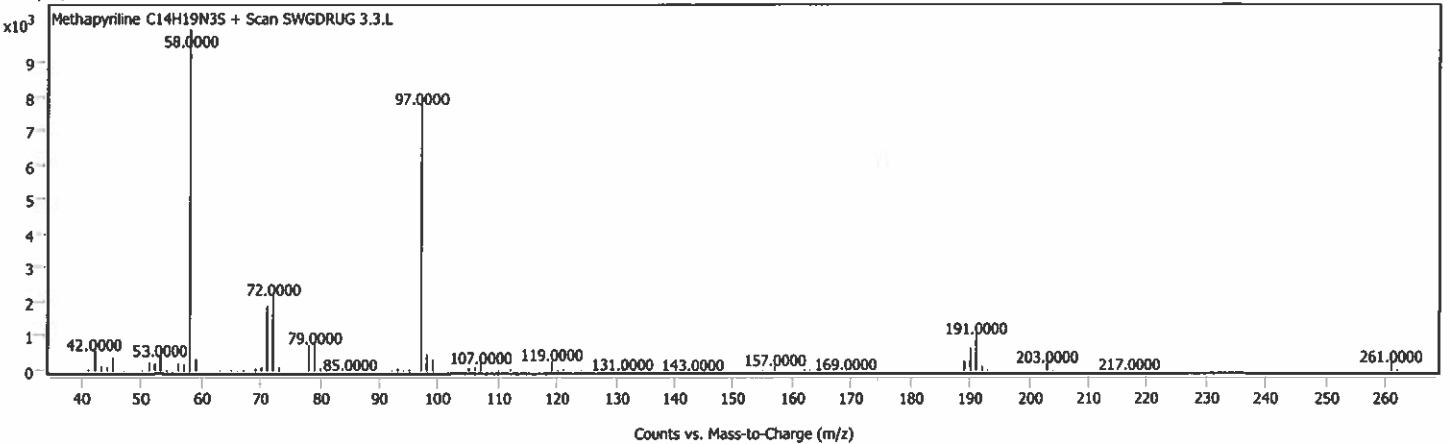
## Observed Spectrum



## Mirror Plot

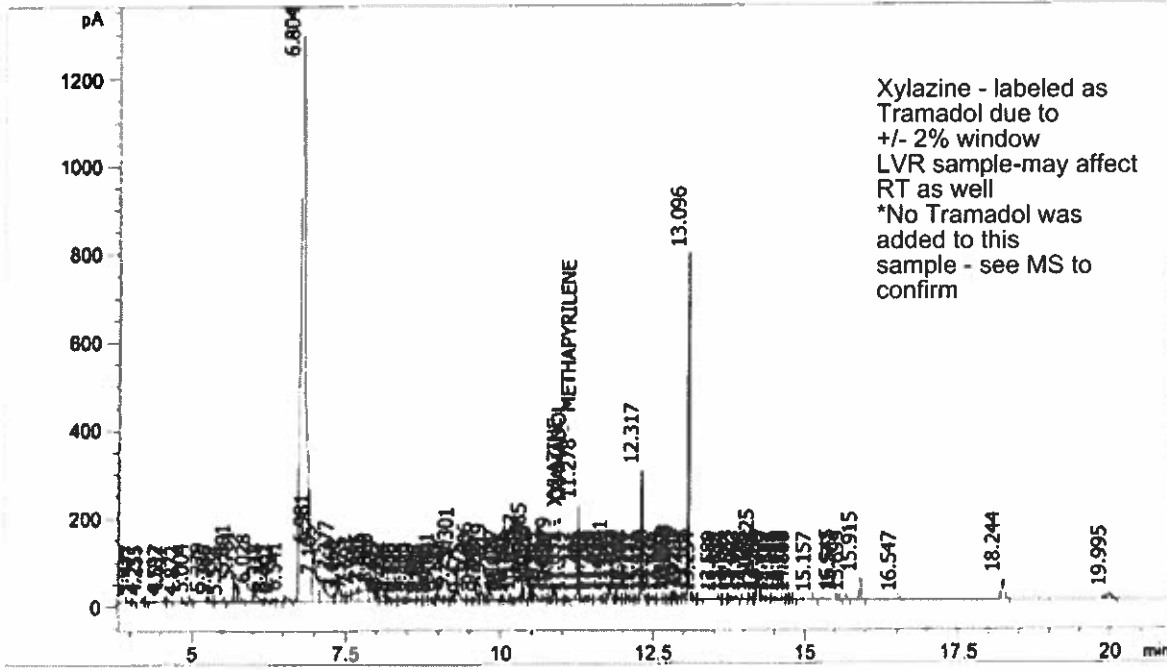


## Library Spectrum



Injection Date: 9/29/2021 11:38:50 AM Seq Line: 55  
 Sample Name: MATRIX 10 W/ISTD 0.01-> Vial 86  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



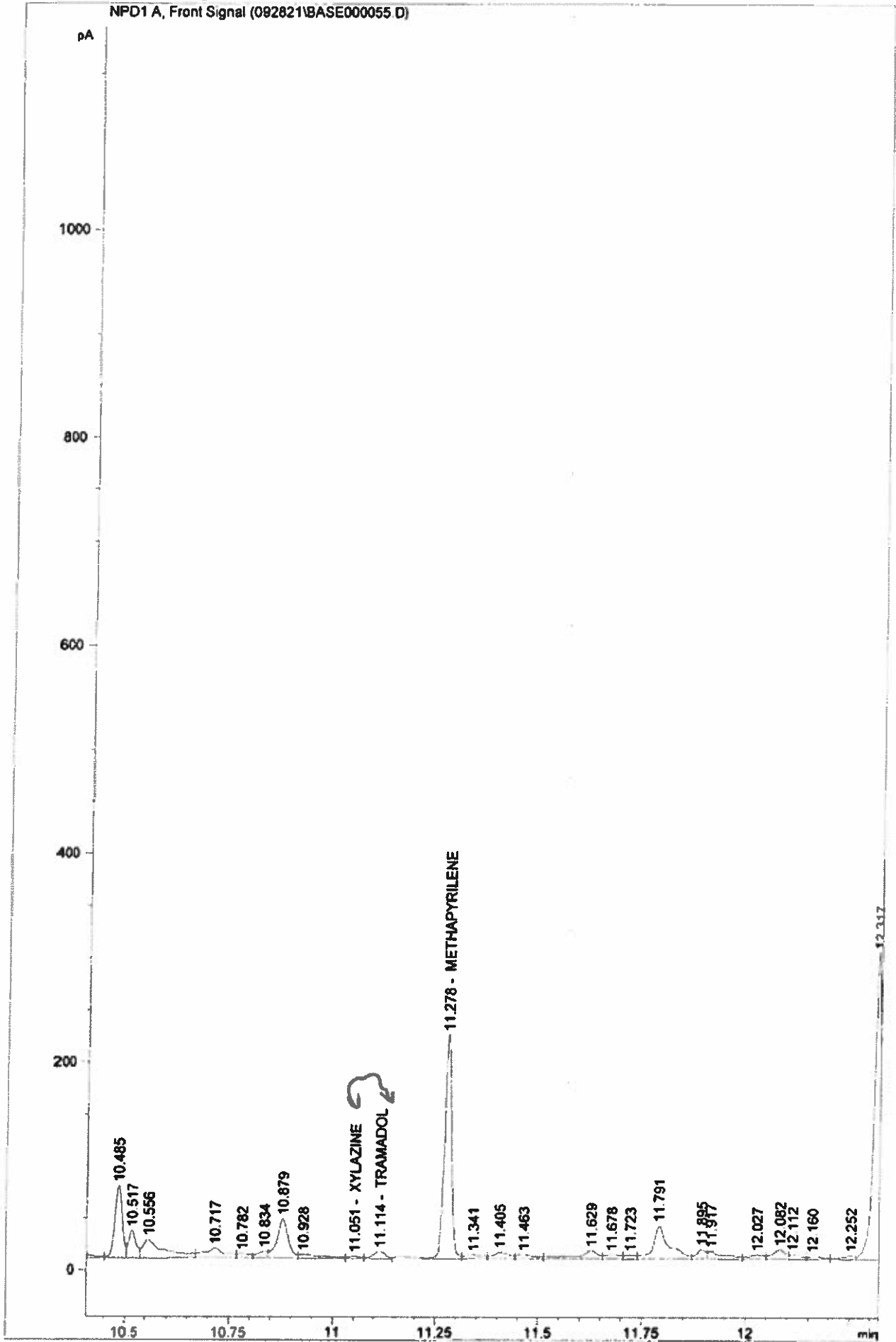
see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.051	11.046	3.632	2.150	0.000000	XYLAZINE
11.114	11.033	13.583	6.802	0.000000	TRAMADOL RRT - 0.9854
11.278	11.278	247.393	215.685	1.000000	METHAPYRILENE

Not used in average of RRTs

Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000055.D)



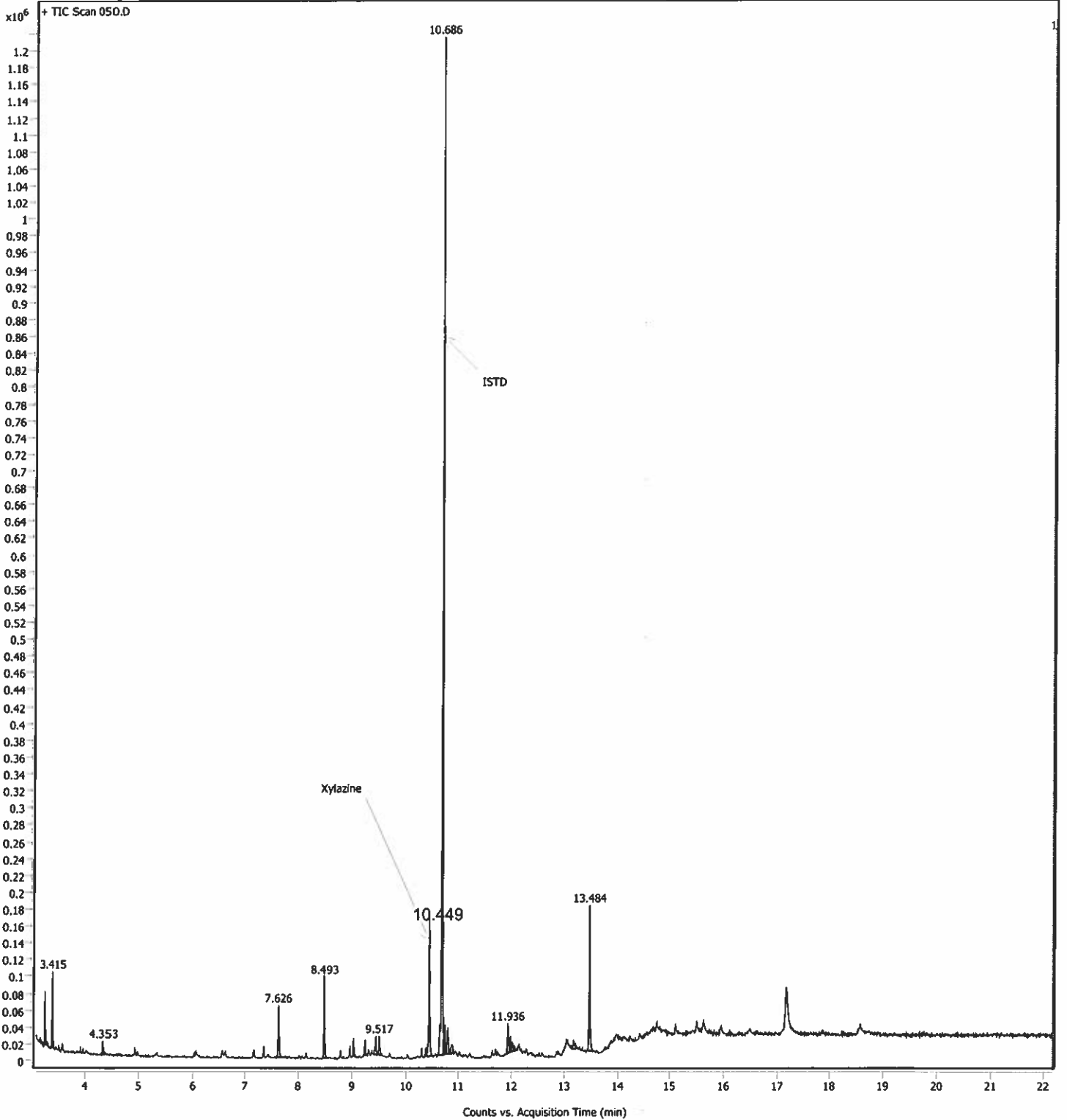


# Analysis Report

## Sample Information

Sample Name	MATRIX 2/W ISTD 0.10 mg/L- BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\050.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 1:21:46 PM (UTC-04:00)
Position	82	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

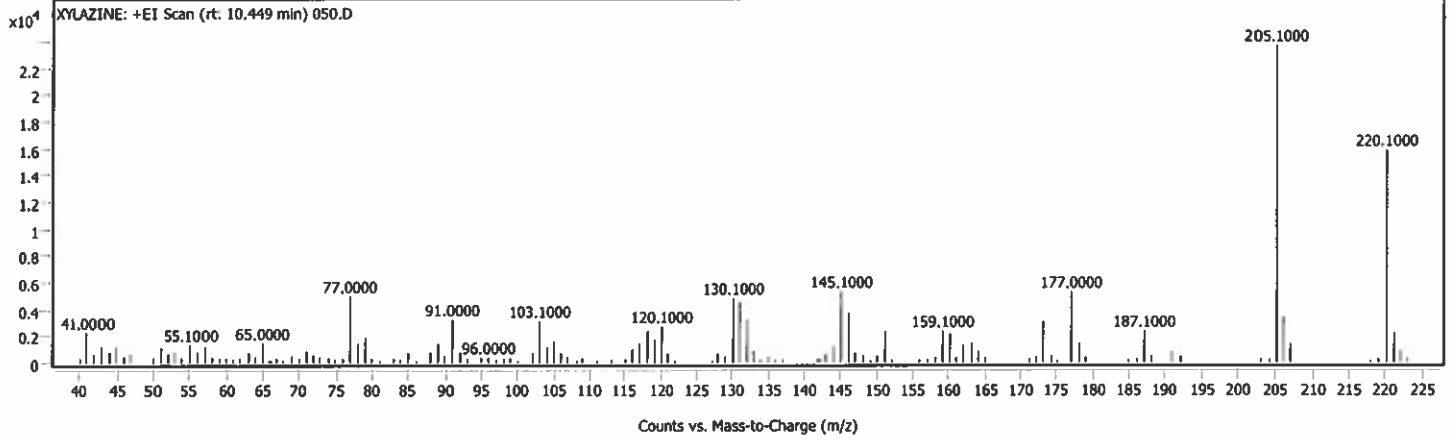
## Sample Spectra

+ Scan (rt: 10.449 min)

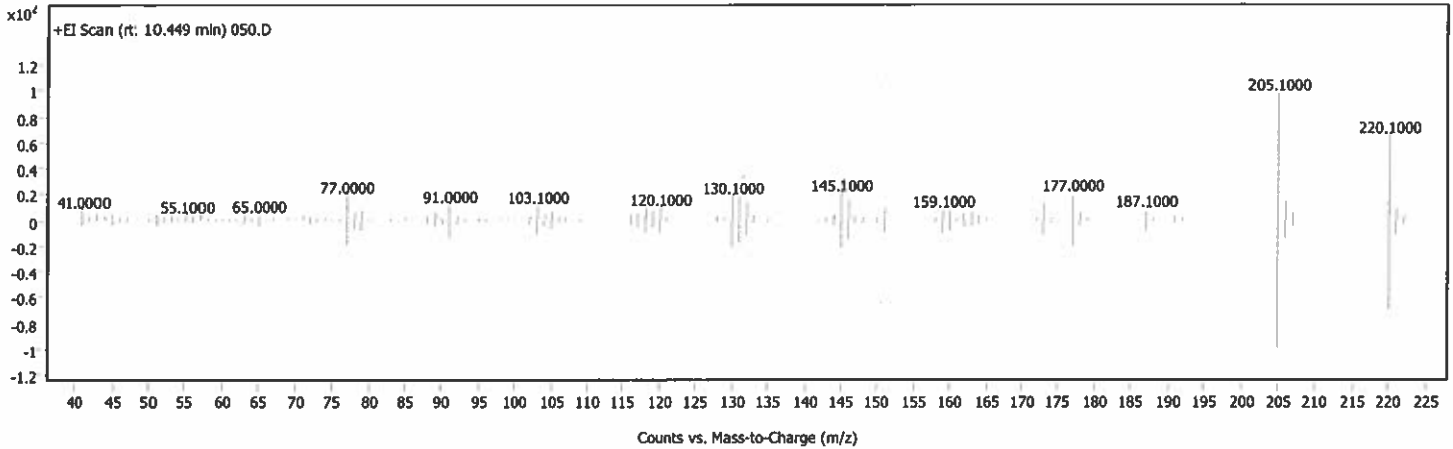
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	95.45	aafs.l	MATRIX 2/W ISTD 0.10 mg/L- BLD	KM/QC DATA IN XYLAZINE VALIDATION

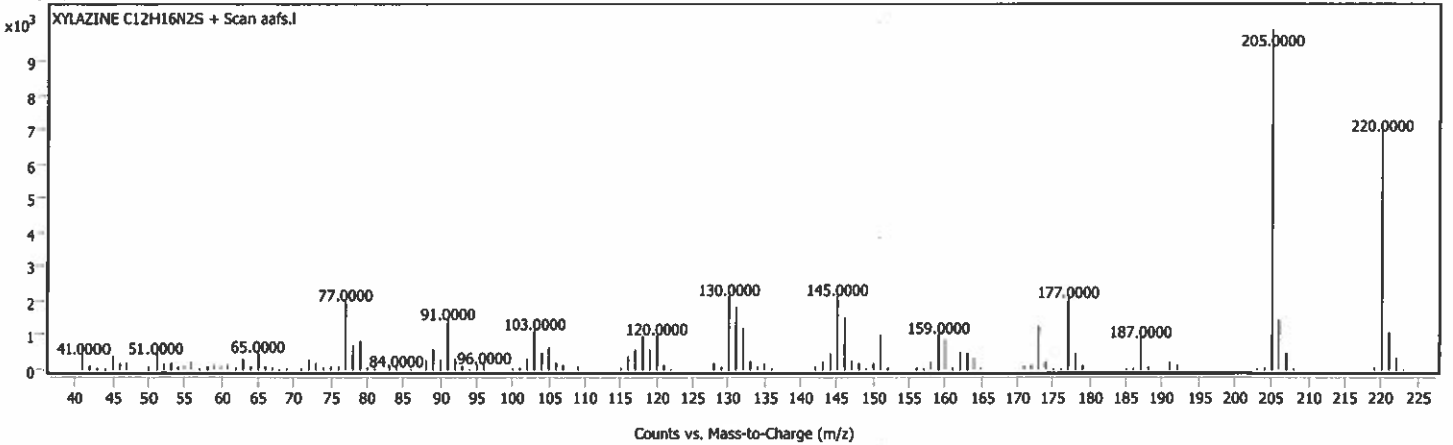
## Observed Spectrum



## Mirror Plot



## Library Spectrum



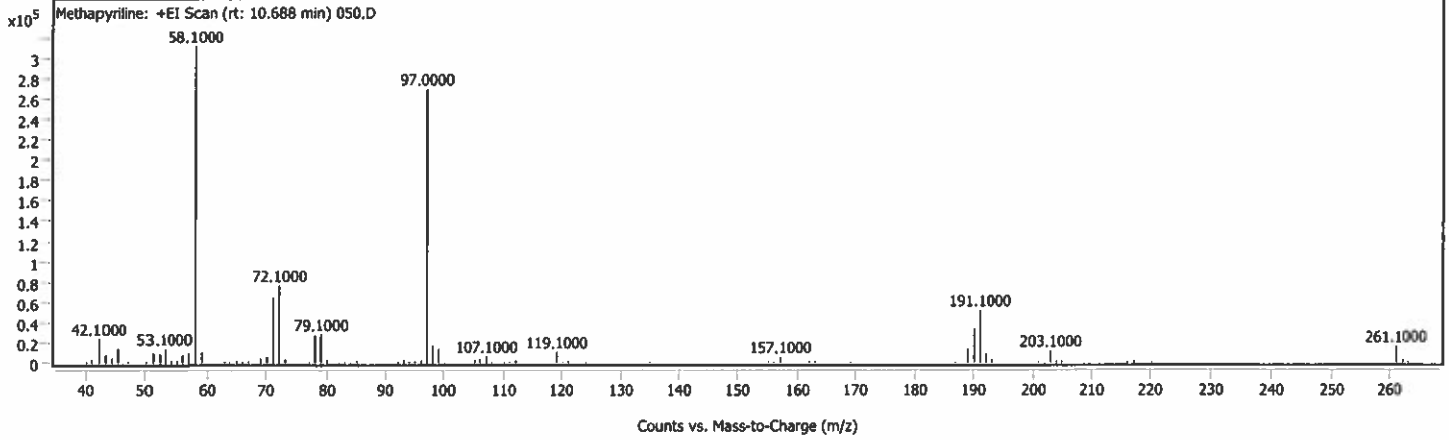
# Analysis Report

+ Scan (rt: 10.688 min)

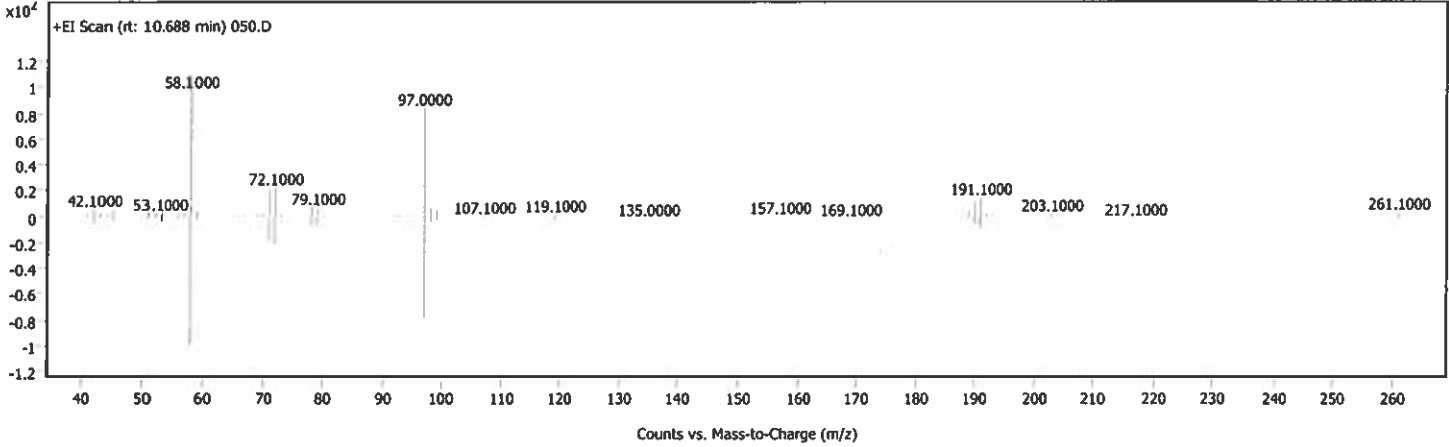
Methapyrine; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyrine	C14H19N3S	98.06	SWGDRUG 3.3.L	MATRIX 2/W ISTD 0.10 mg/L- BLD	KM/QC DATA IN XYLAZINE VALIDATION

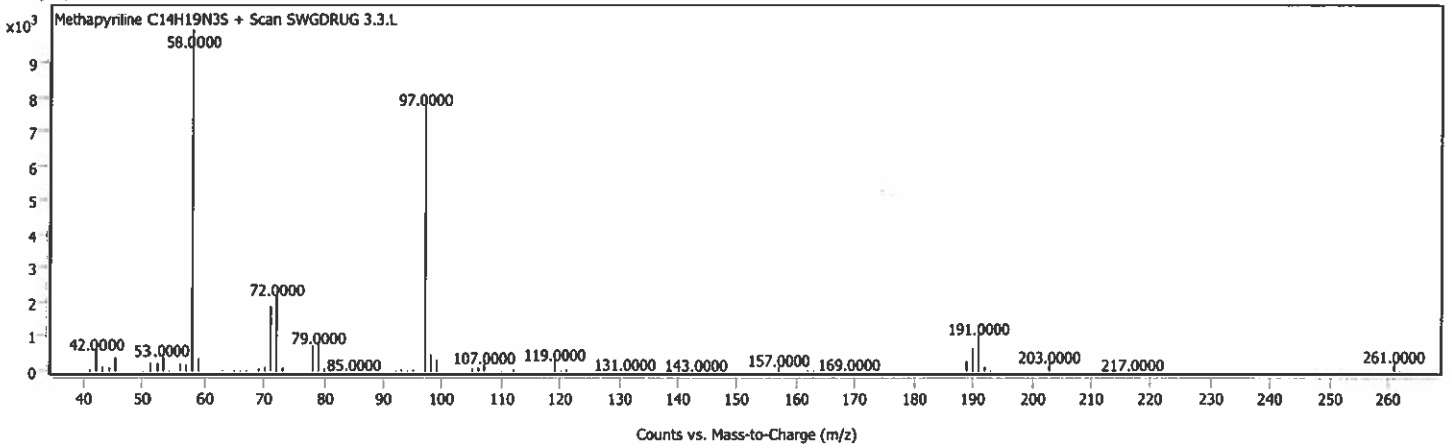
## Observed Spectrum



## Mirror Plot

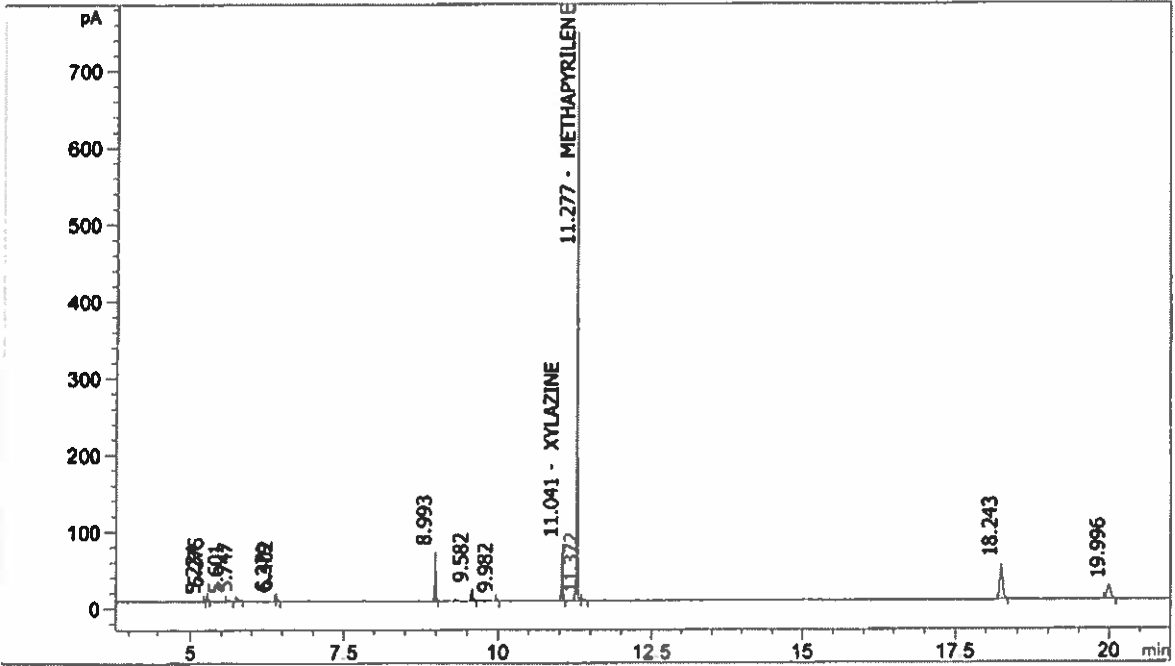


## Library Spectrum



Injection Date: 9/29/2021 12:27:27 PM Seq Line: 57  
 Sample Name: MATRIX 2 W/ISTD 0.10 -> Vial 88  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.041	11.046	85.655	74.466	0.000000	XYLAZINE RRT - 0.9791
11.277	11.278	859.985	734.628	1.000000	METHAPYRILENE

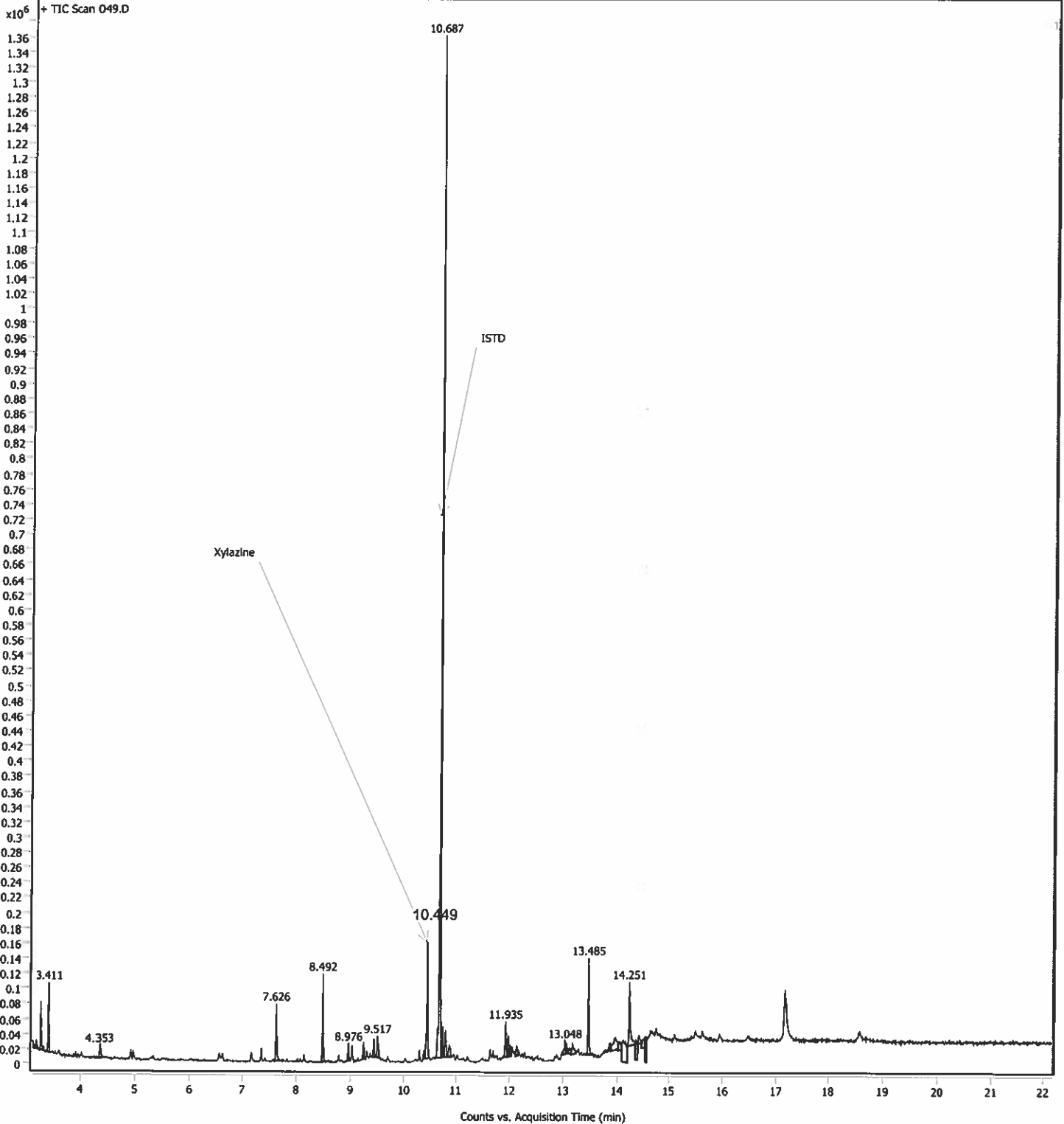
# Analysis Report



## Sample Information

Sample Name	MATRIX 1 W/ISTD 0.10 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\049.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 12:55:43 PM (UTC-04:00)
Position	81	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

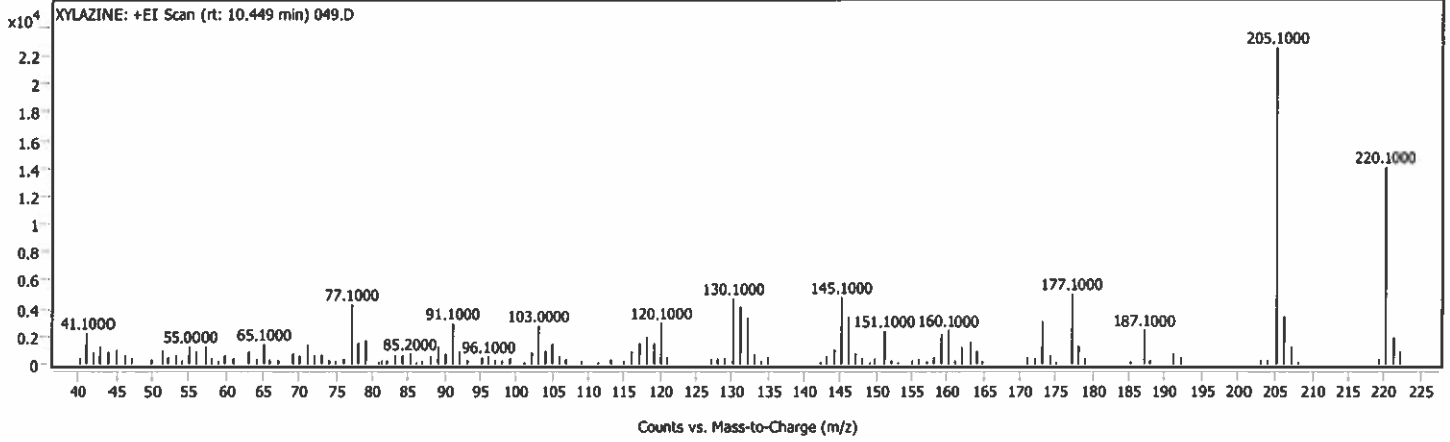
## Sample Spectra

+ Scan (rt: 10.449 min)

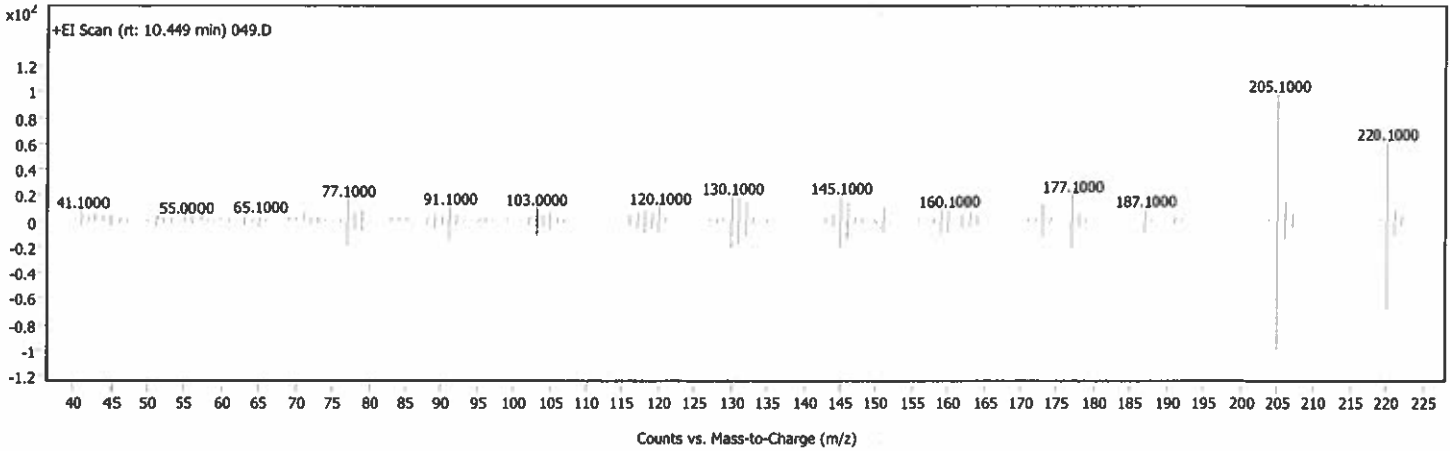
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	94.89	aafs.i	MATRIX 1 W/ISTD 0.10 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

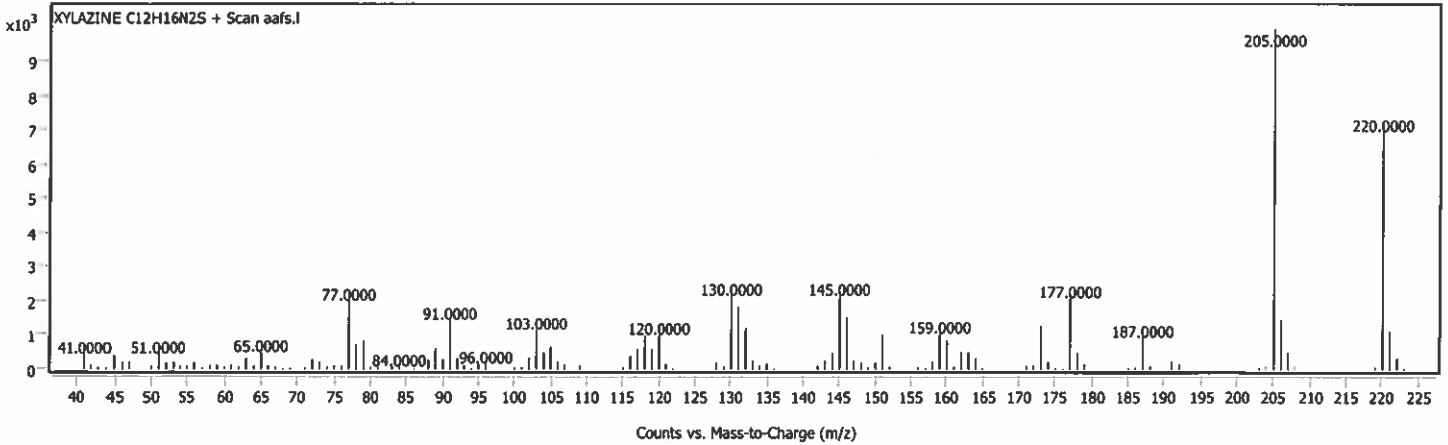
### Observed Spectrum



### Mirror Plot



### Library Spectrum



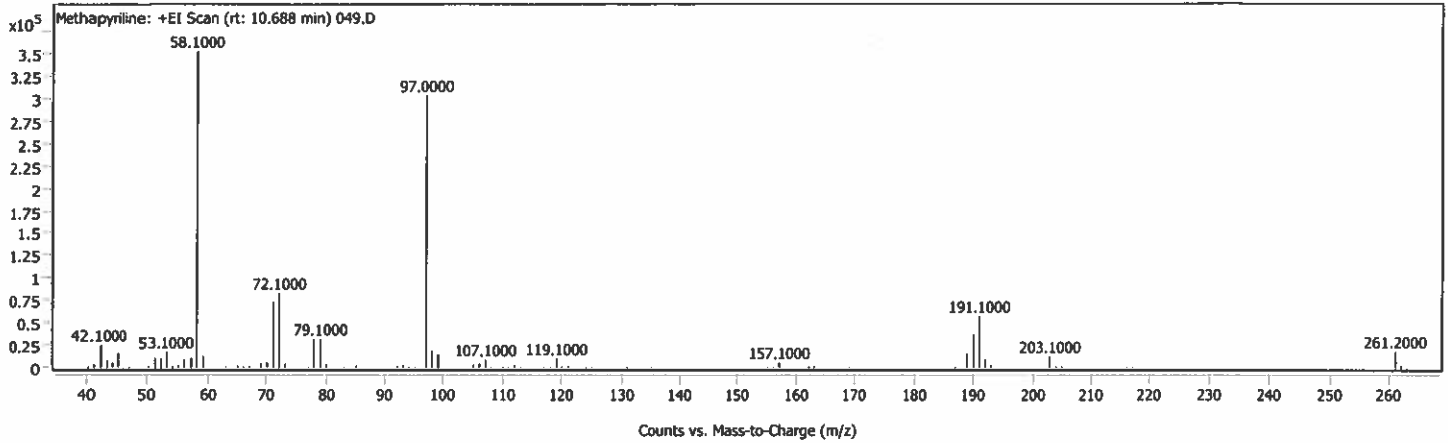
# Analysis Report

+ Scan (rt: 10.688 min)

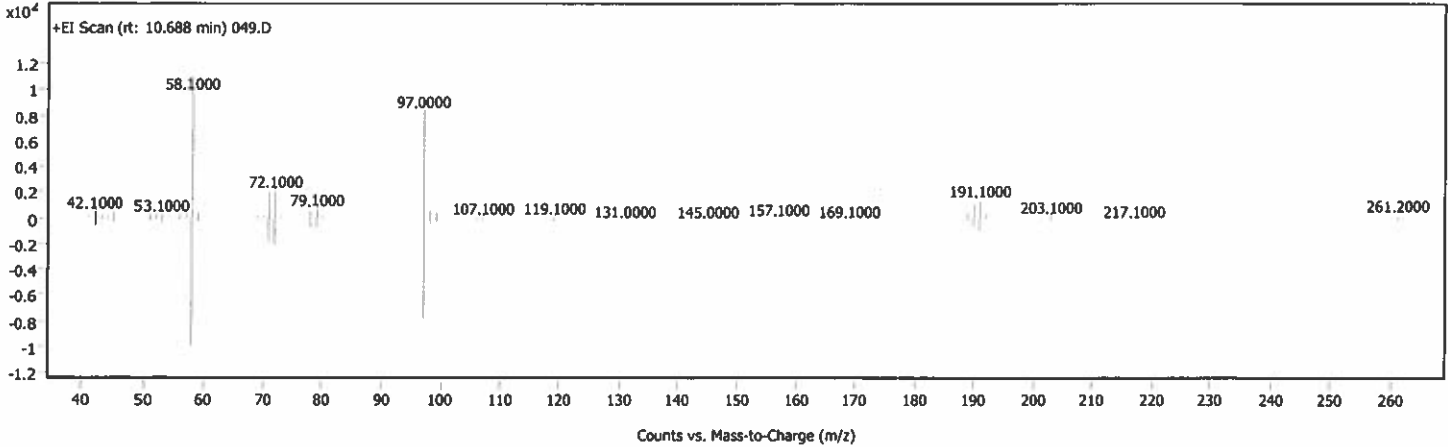
Methapyriline; C<sub>14</sub>H<sub>19</sub>N<sub>3</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C <sub>14</sub> H <sub>19</sub> N <sub>3</sub> S	98.53	SWGDRUG 3.3.L	MATRIX 1 W/ISTD 0.10 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

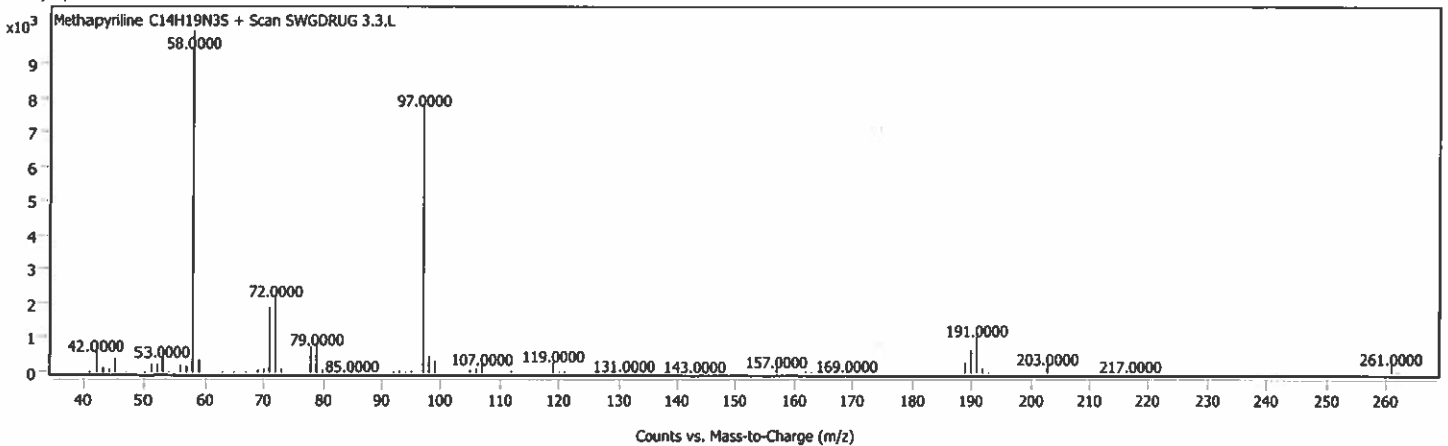
## Observed Spectrum



## Mirror Plot



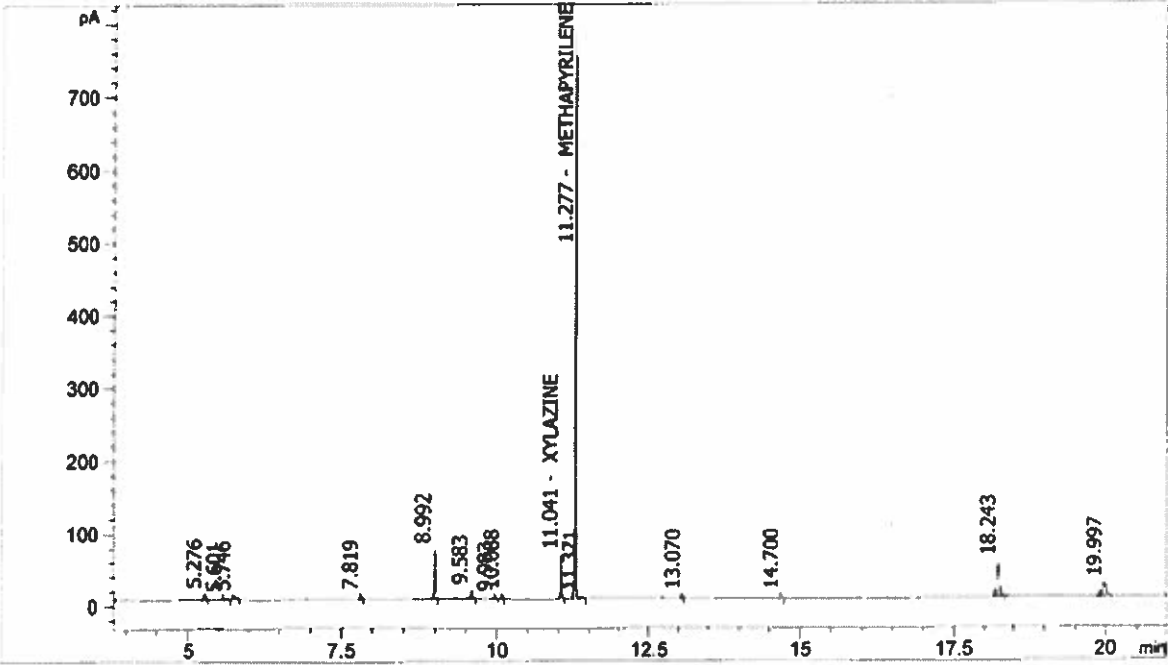
## Library Spectrum



Injection Date: 9/29/2021 12:03:08 PM  
 Sample Name: MATRIX 1 W/ISTD 0.10 ->  
 Sample Info:

Seq Line: 56  
 Vial 87

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.041	11.046	69.536	59.638	0.000000	XYLAZINE RRT - 0.9791
11.277	11.278	883.290	777.130	1.000000	METHAPYRILENE

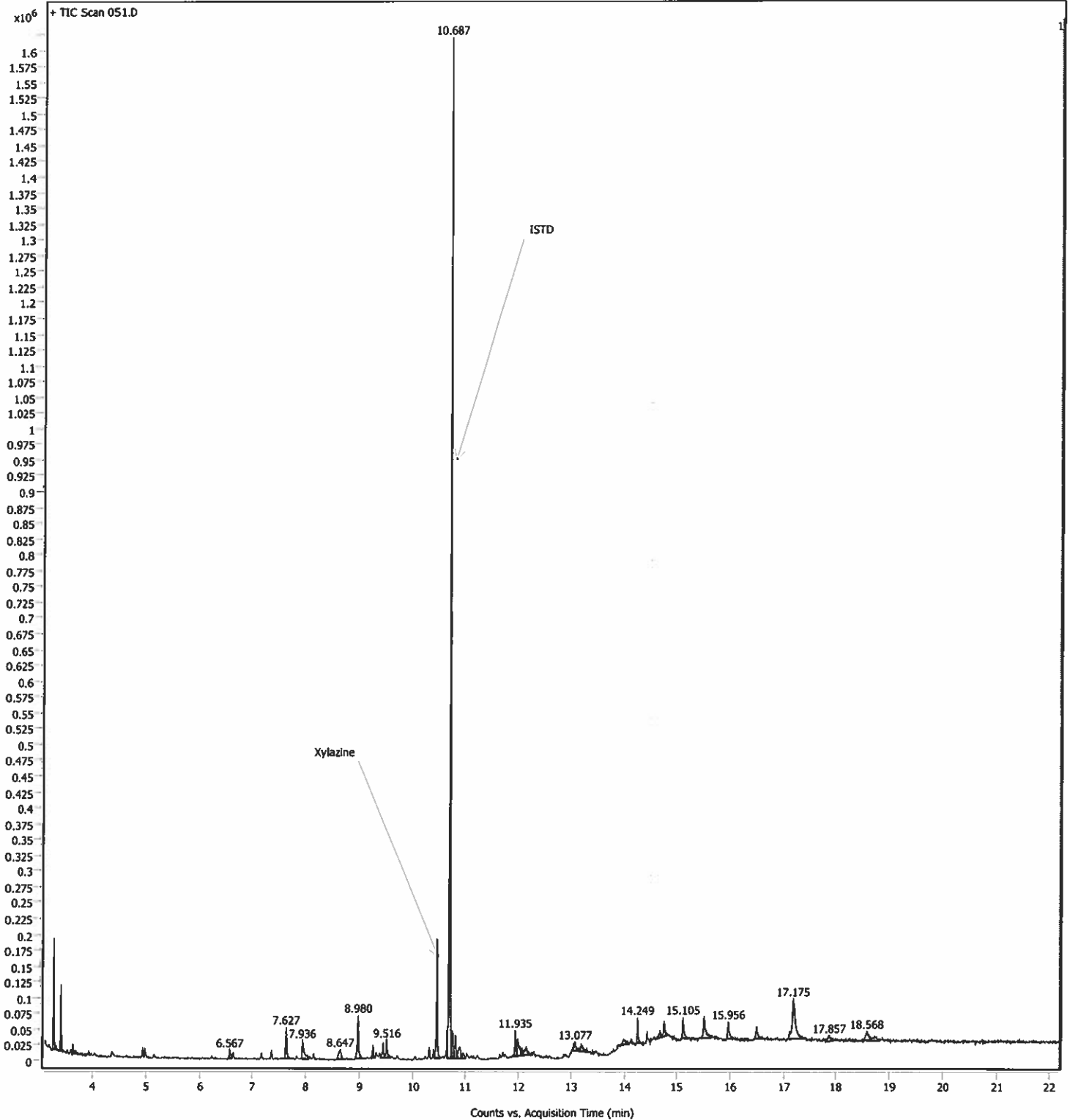


# Analysis Report

## Sample Information

Sample Name	MATRIX 6 W/ISTD 0.10 mg/L- URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\051.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 1:47:38 PM (UTC-04:00)
Position	83	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

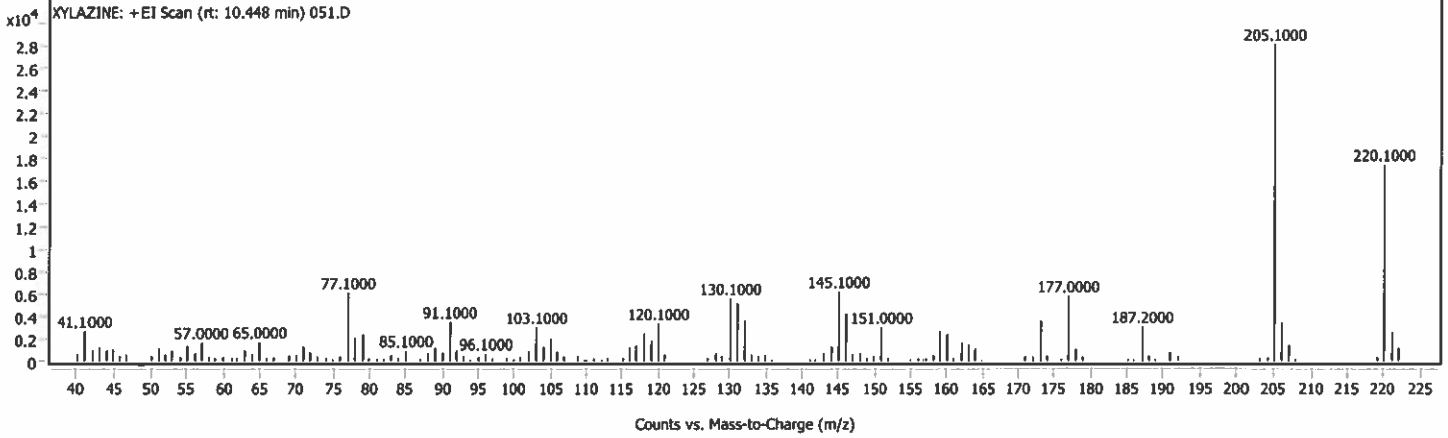
## Sample Spectra

+ Scan (rt: 10.448 min)

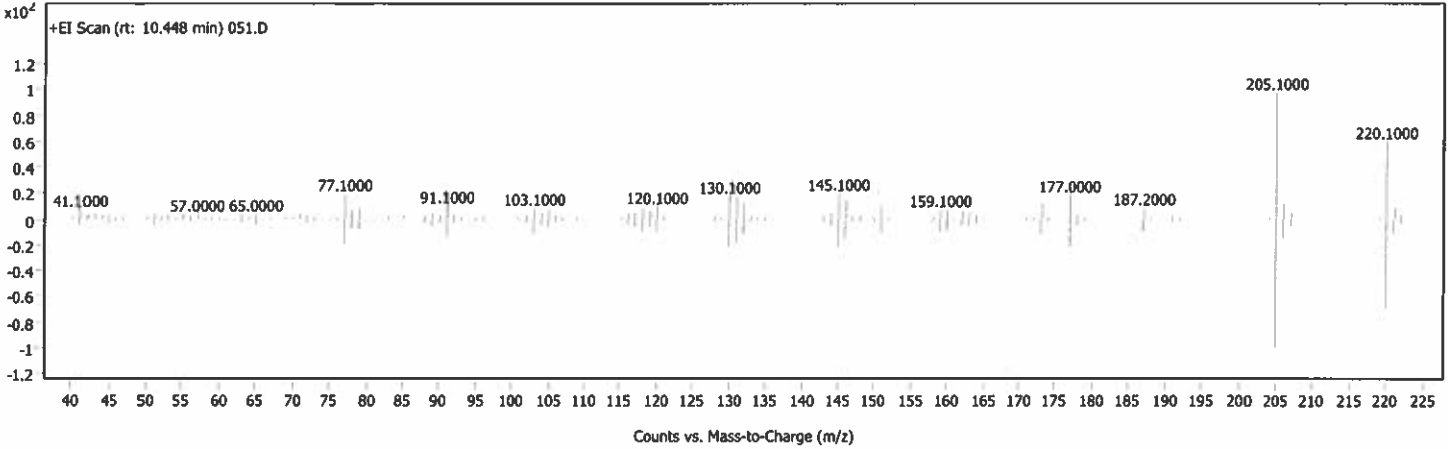
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	95.88	aafs.l	MATRIX 6 W/ISTD 0.10 mg/L- URN	KM/QC DATA IN XYLAZINE VALIDATION

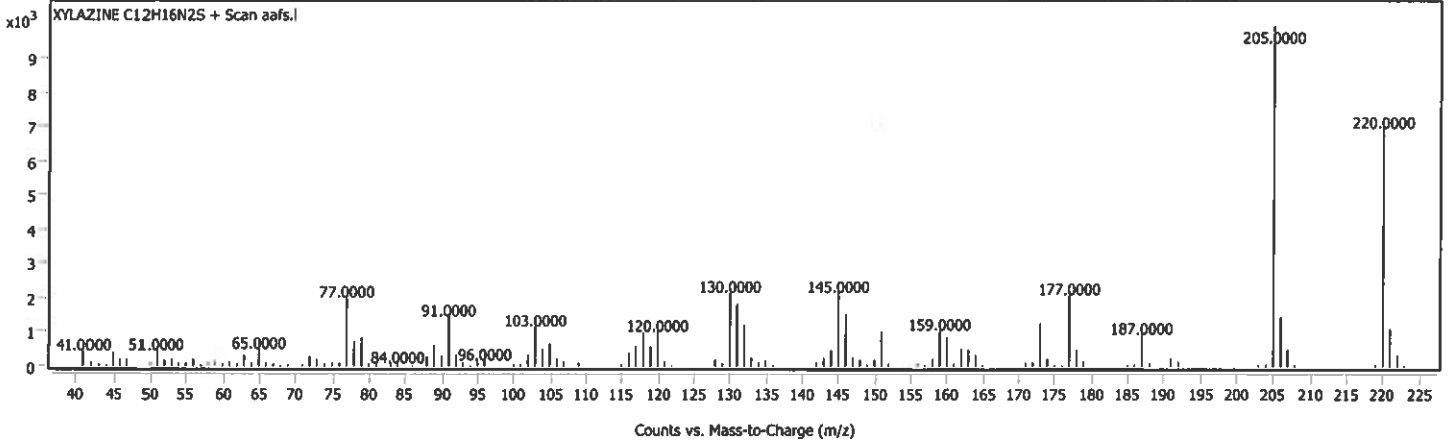
### Observed Spectrum



### Mirror Plot



### Library Spectrum



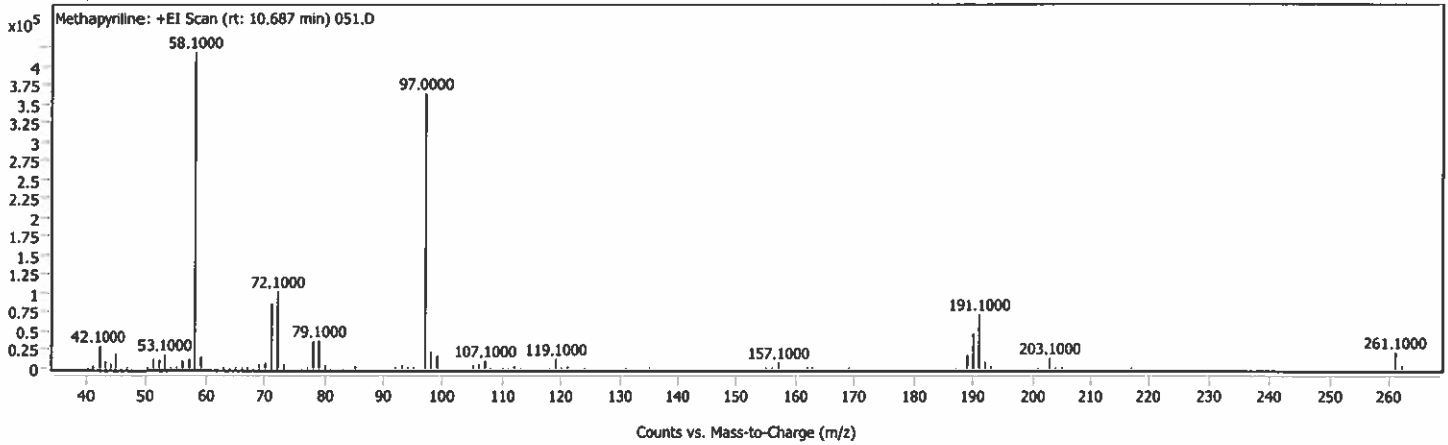
# Analysis Report

+ Scan (rt: 10.687 min)

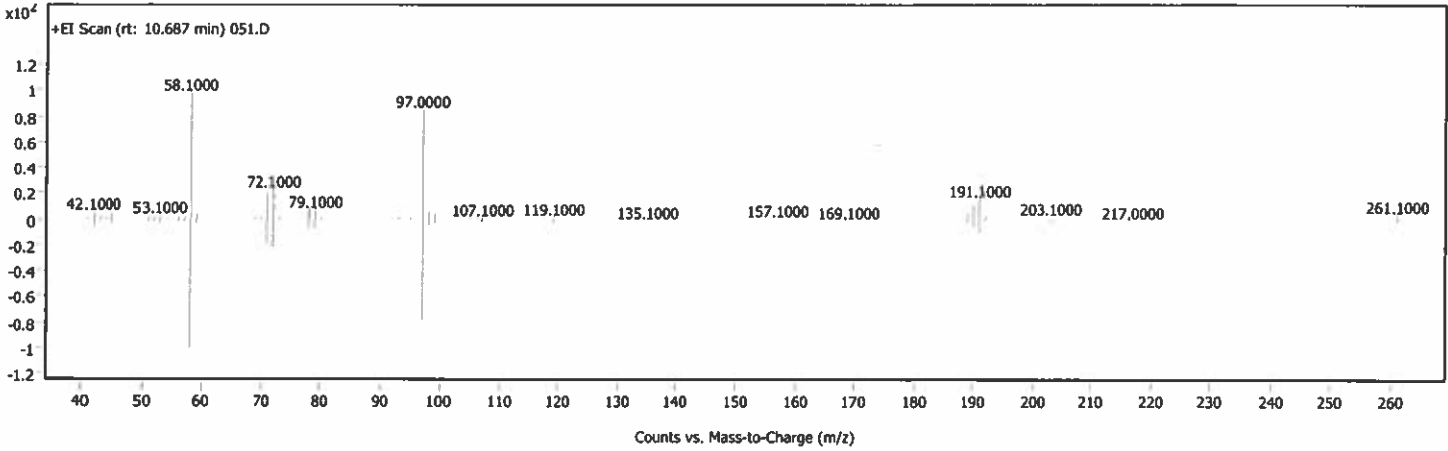
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.47	SWGDRUG 3.3.L	MATRIX 6 W/ISTD 0.10 mg/L- URN	KM/QC DATA IN XYLAZINE VALIDATION

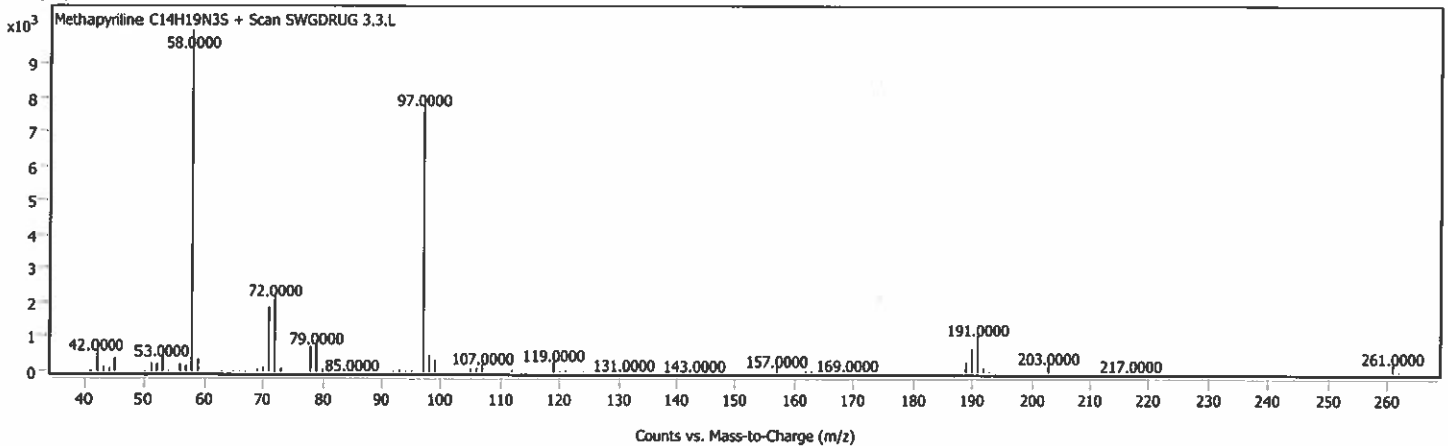
## Observed Spectrum



## Mirror Plot

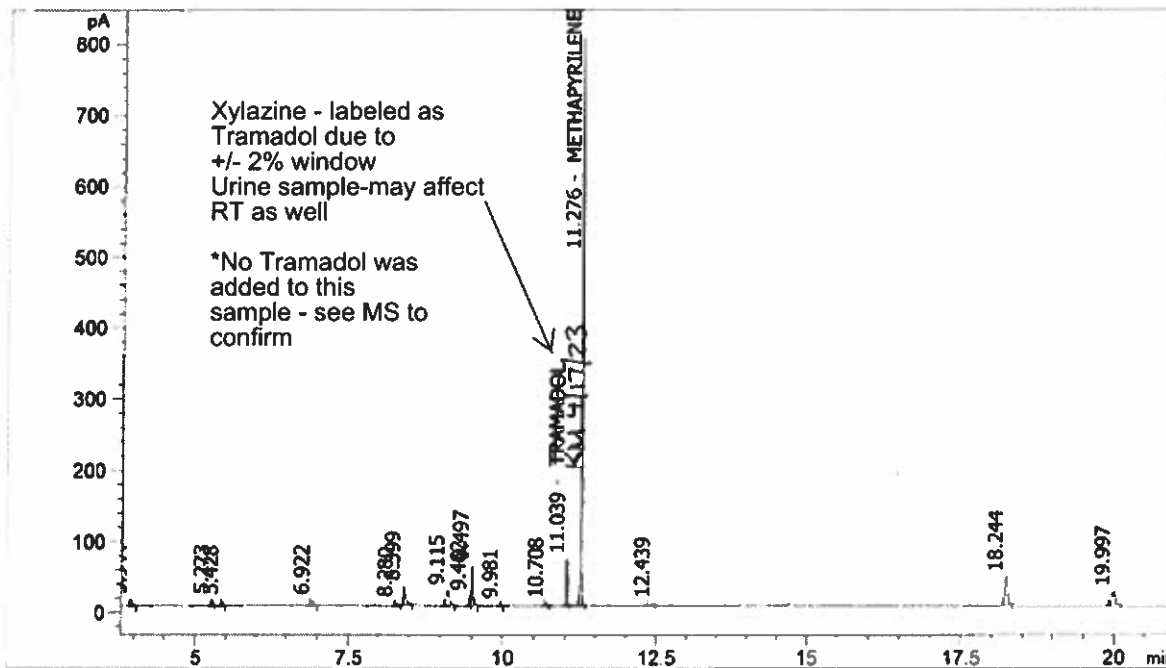


## Library Spectrum



Injection Date: 9/29/2021 12:51:51 PM Seq Line: 58  
 Sample Name: MATRIX 6 W/ISTD 0.10 -> Vial 89  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



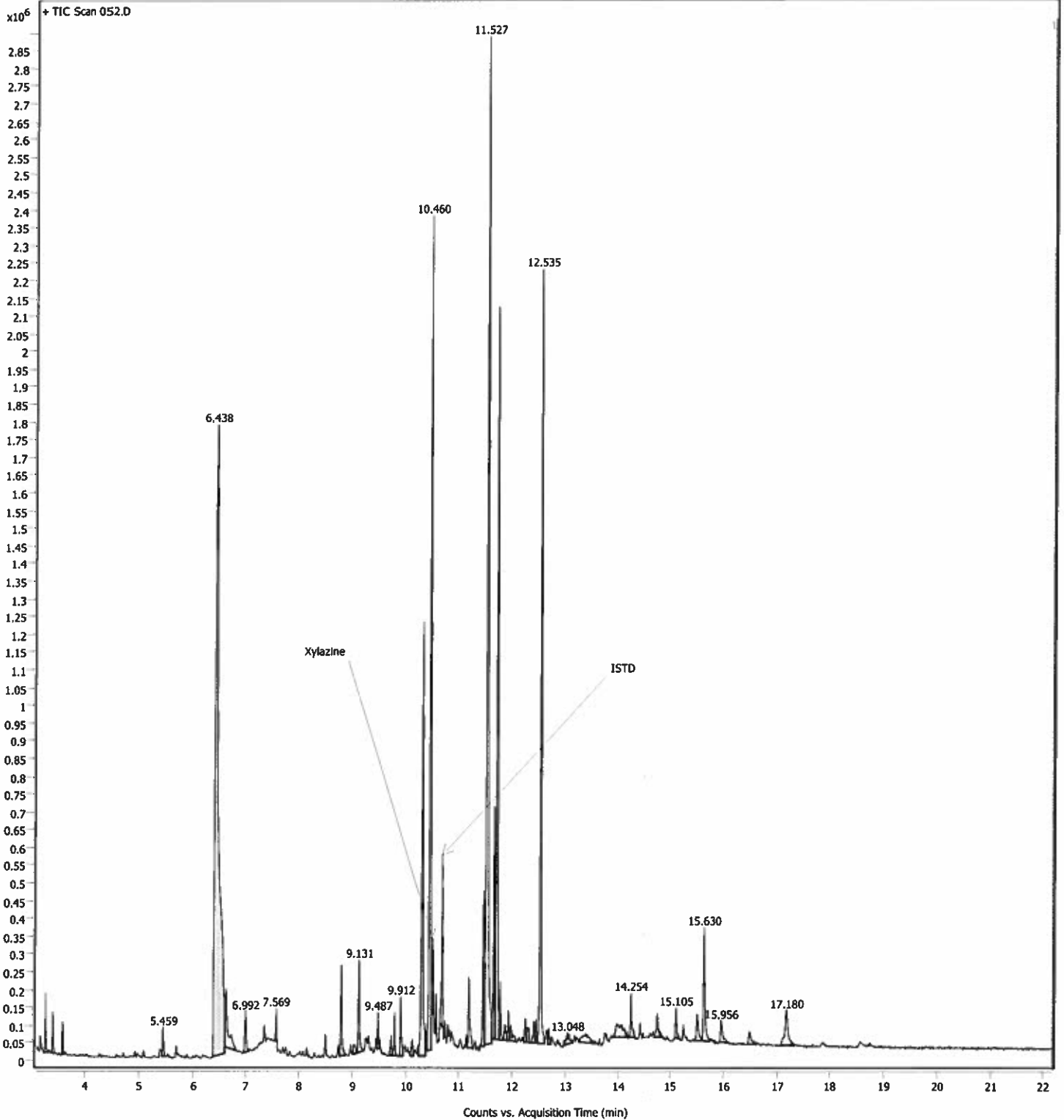
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.039	11.033	74.280	66.514	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE RRT - 0.9790
11.276	11.278	916.292	791.683	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 10 W/ISTD 0.10 mg/L-LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\052.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 2:13:27 PM (UTC-04:00)
Position	84	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



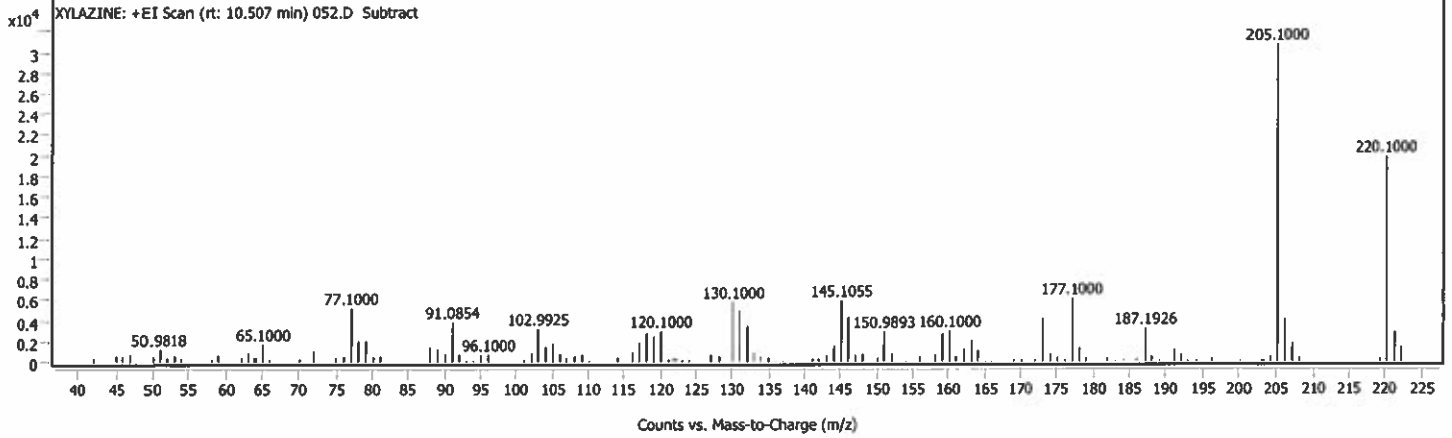
# Analysis Report

## Sample Spectra

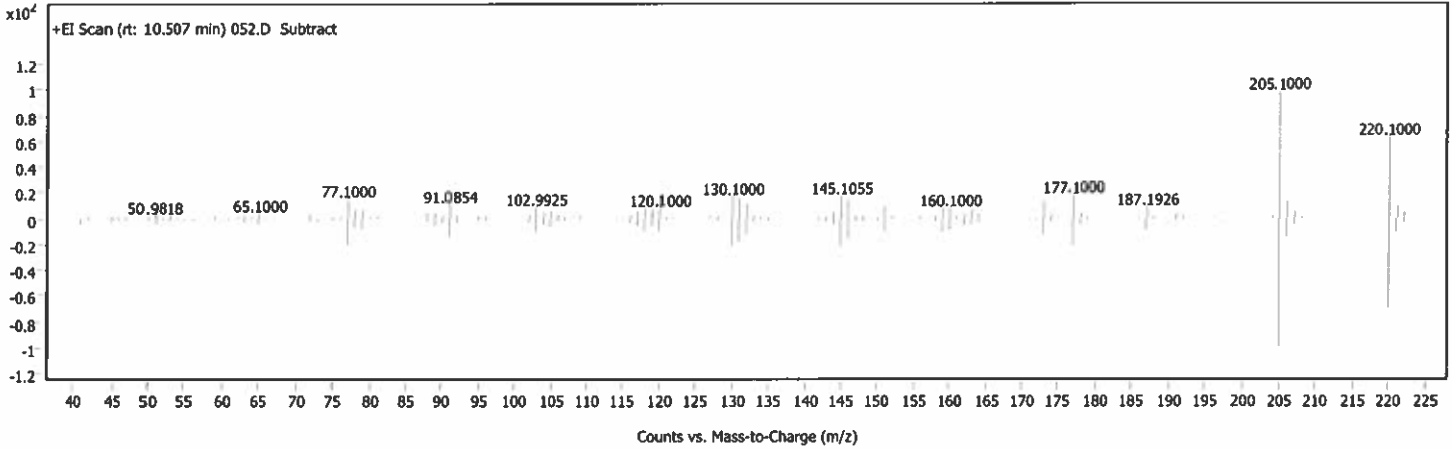
+ Scan (rt: 10.507 min) Sub **XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	94.41	aafs.l	MATRIX 10 W/ISTD 0.10 mg/L-LVR	KM/QC DATA IN XYLAZINE VALIDATION

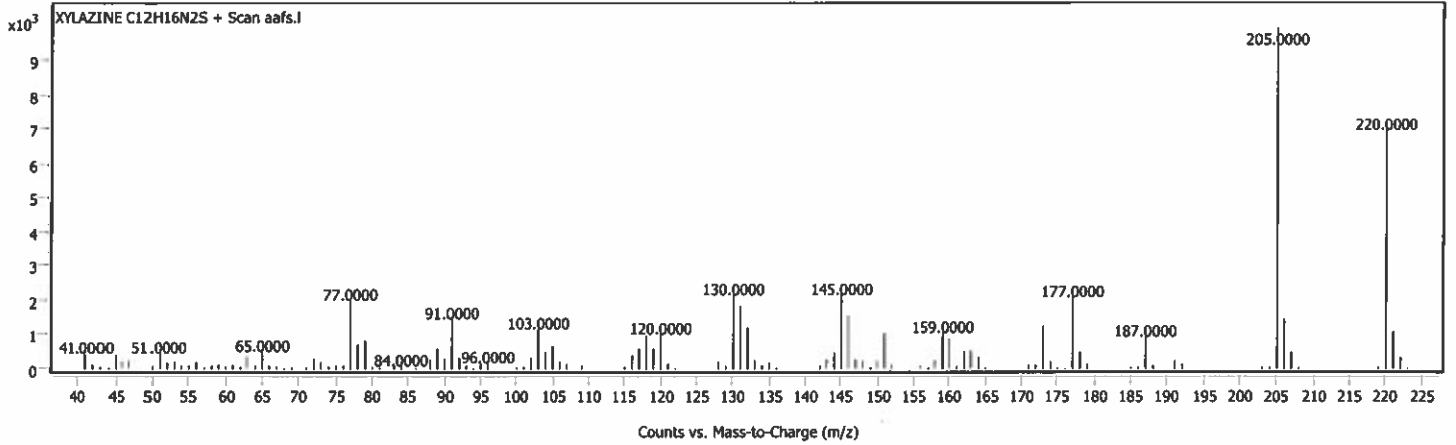
## Observed Spectrum



## Mirror Plot



## Library Spectrum



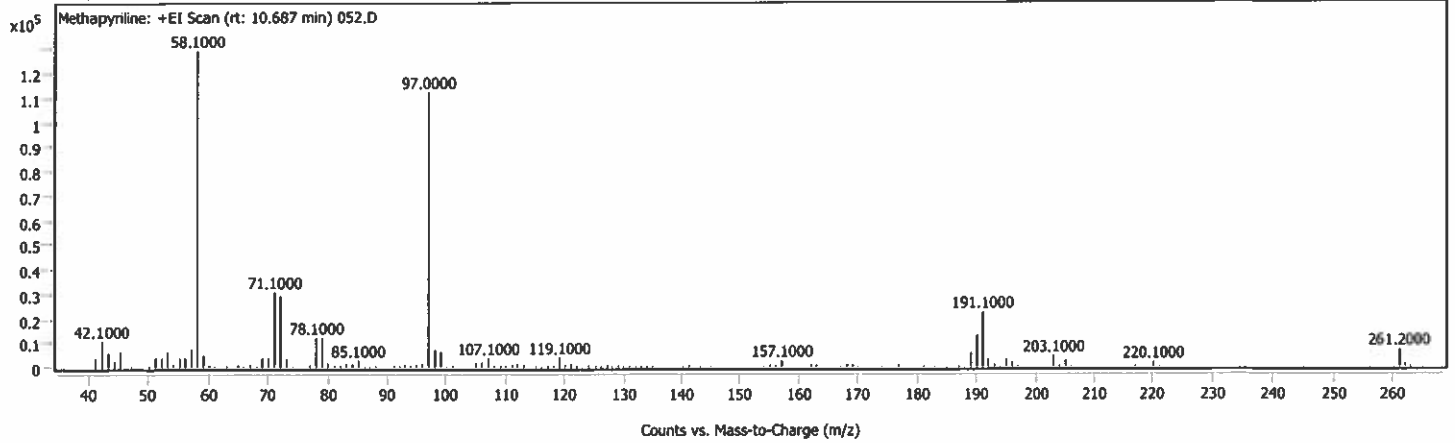
# Analysis Report

+ Scan (rt: 10.687 min)

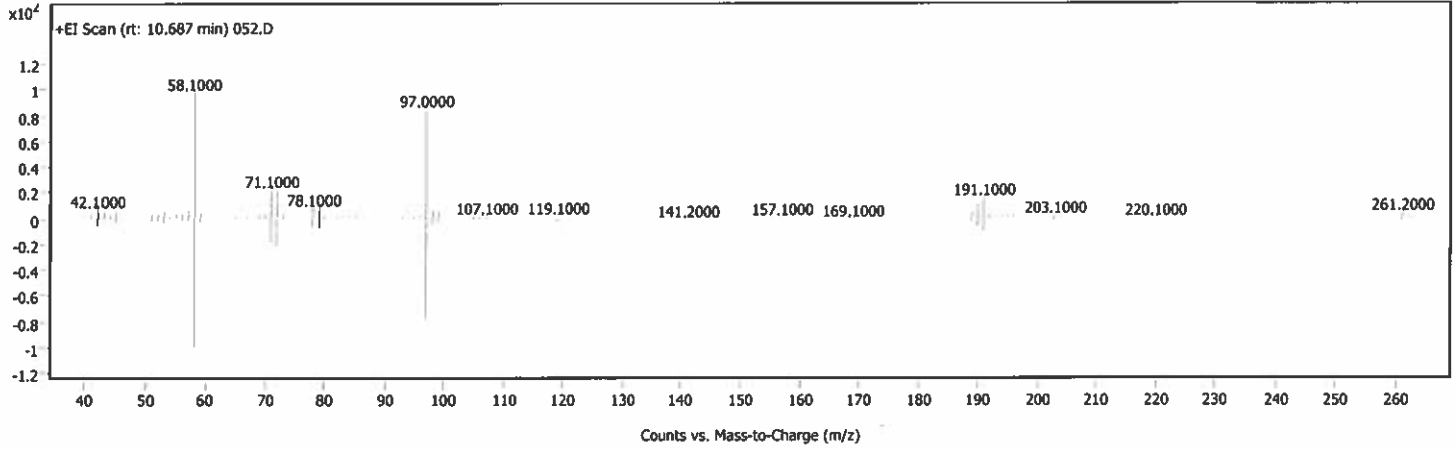
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	92.85	SWGDRUG 3.3.L	MATRIX 10 W/ISTD 0.10 mg/L-LVR	KM/QC DATA IN XYLAZINE VALIDATION

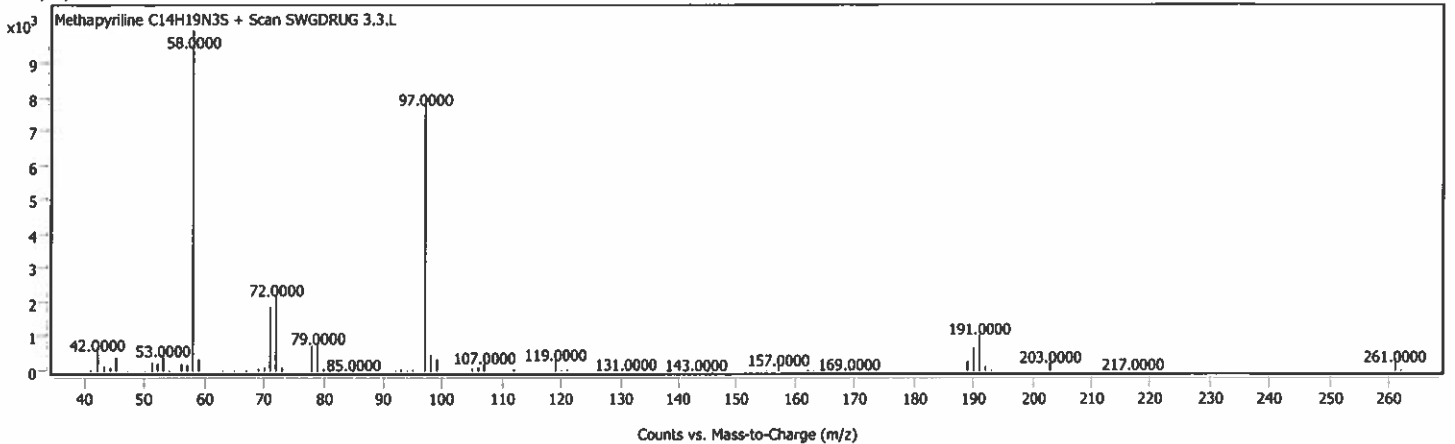
### Observed Spectrum



### Mirror Plot

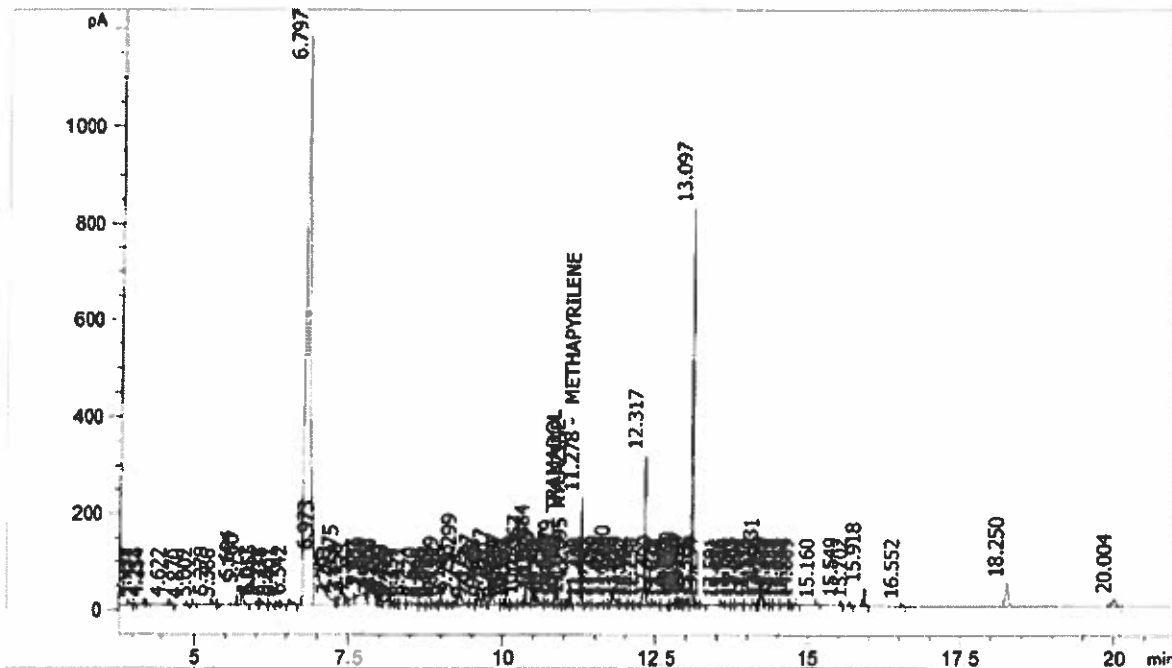


### Library Spectrum



Injection Date: 9/29/2021 1:16:08 PM Seq Line: 59  
 Sample Name: MATRIX 10 W/ISTD 0.10-> Vial 90  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



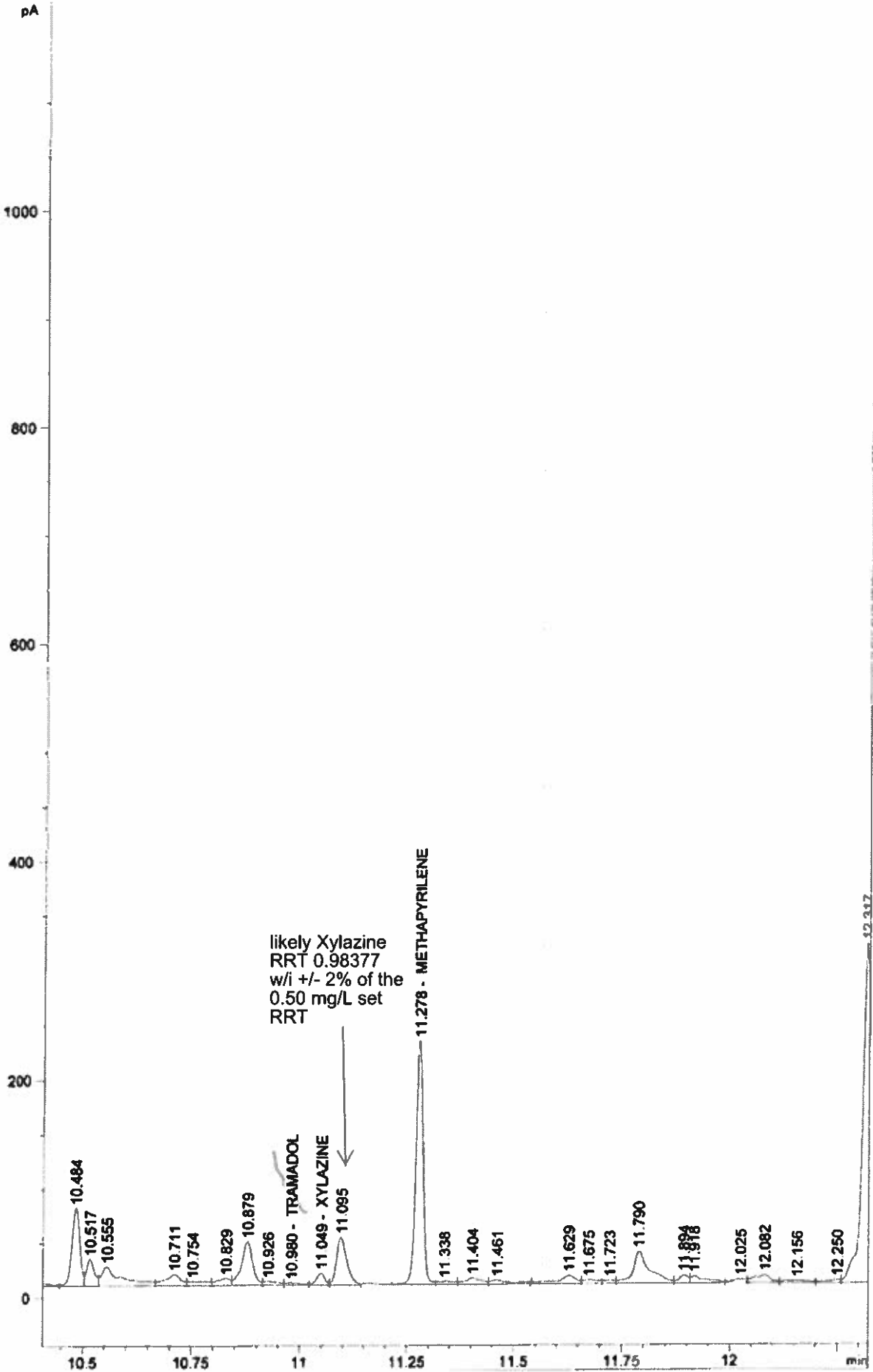
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
10.980	11.033	5.877	2.259	0.000000	TRAMADOL
11.049	11.046	15.449	11.191	0.000000	XYLAZINE see following page
11.278	11.278	267.175	223.779	1.000000	METHAPYRILENE

Not used in average of RRTs



Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000059.D)

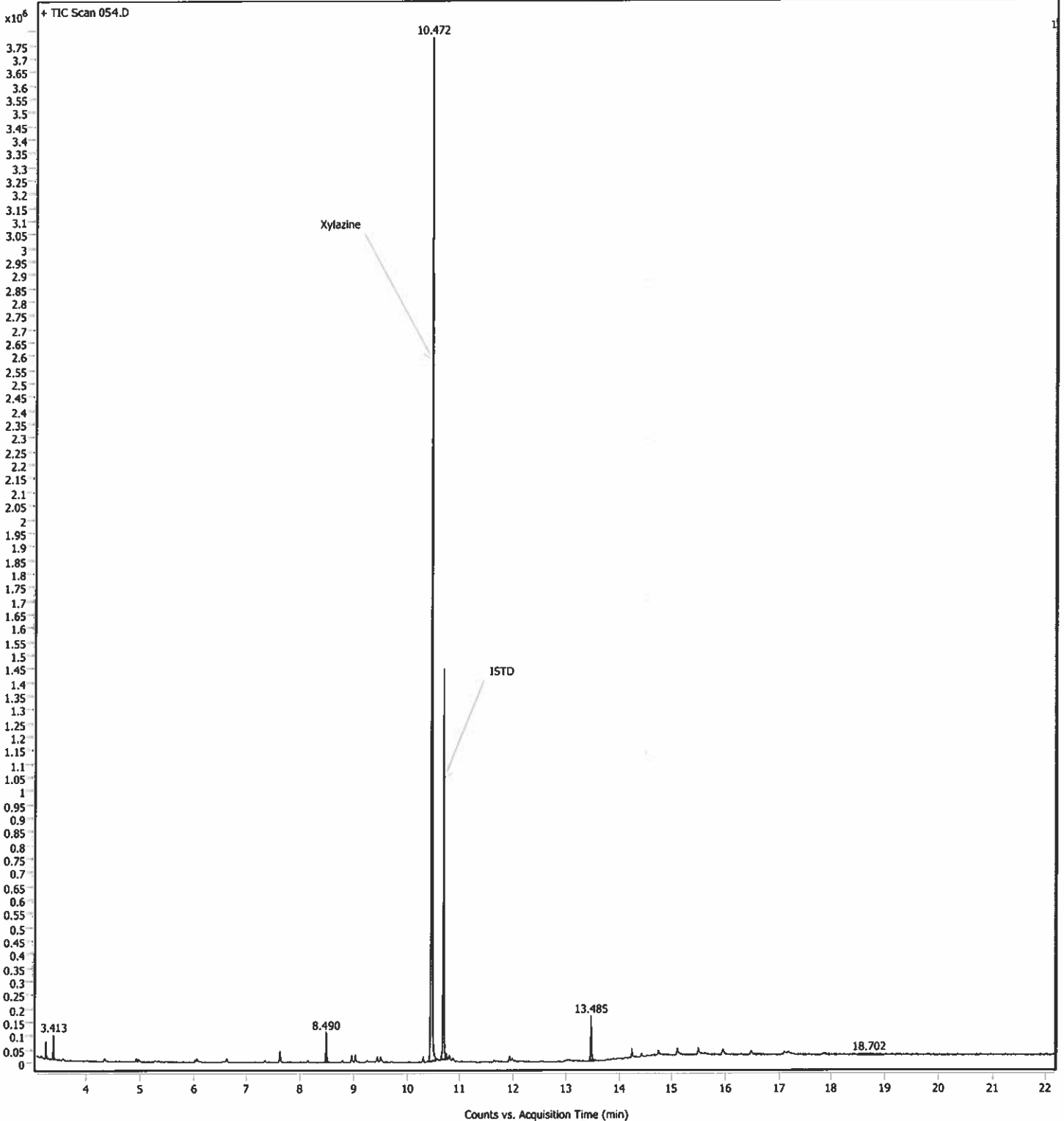


# Analysis Report

## Sample Information

Sample Name	MATRIX 2/W ISTD 2.0 mg/L- BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\054.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 3:05:10 PM (UTC-04:00)
Position	86	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

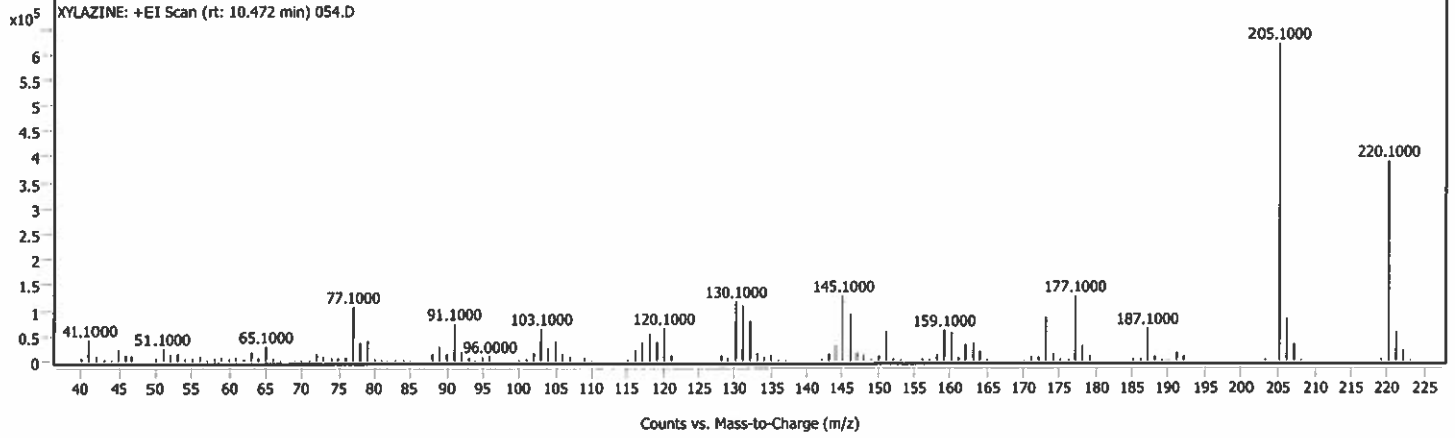
## Sample Spectra

+ Scan (rt: 10.472 min)

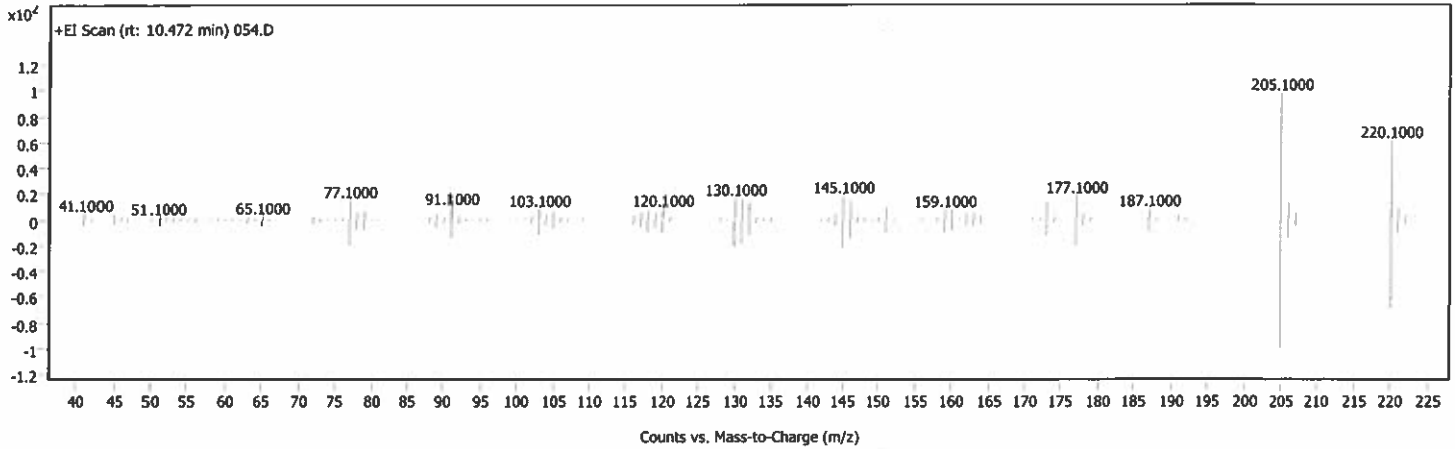
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.72	aafs.I	MATRIX 2/W ISTD 2.0 mg/L- BLD	KM/QC DATA IN XYLAZINE VALIDATION

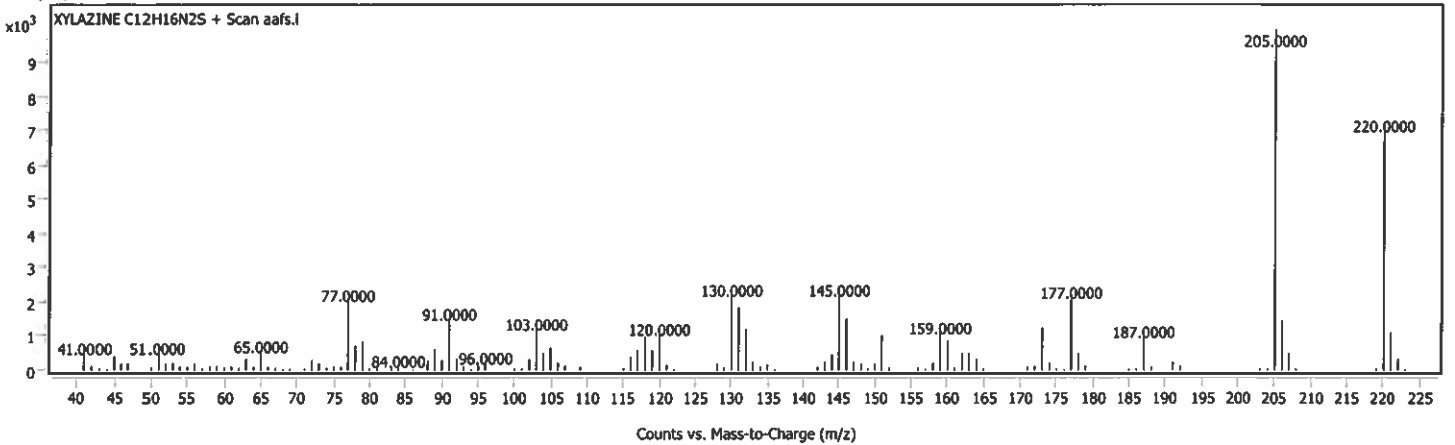
## Observed Spectrum



## Mirror Plot



## Library Spectrum



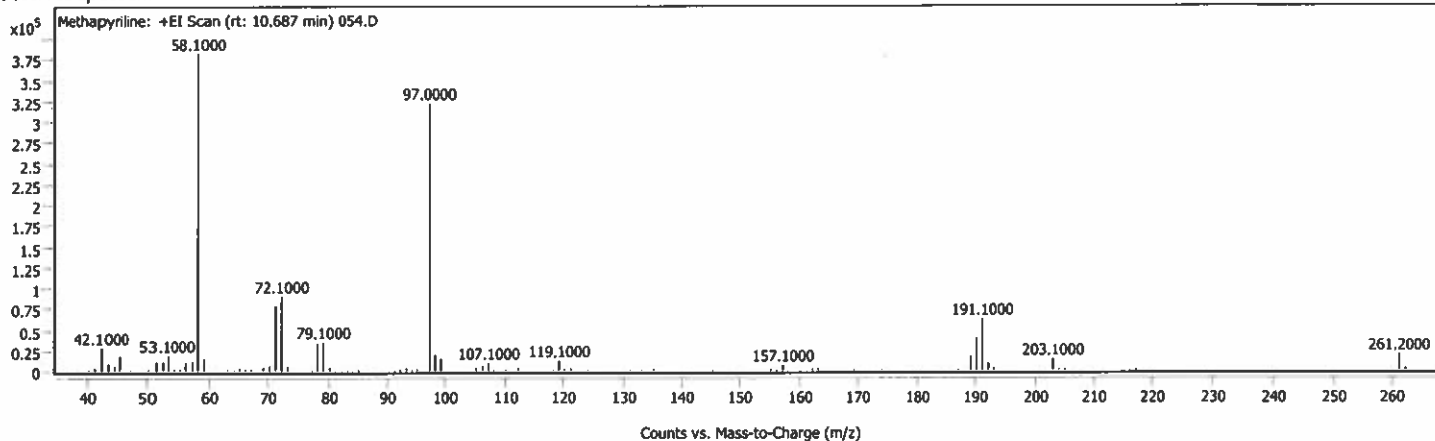
# Analysis Report

+ Scan (rt: 10.687 min)

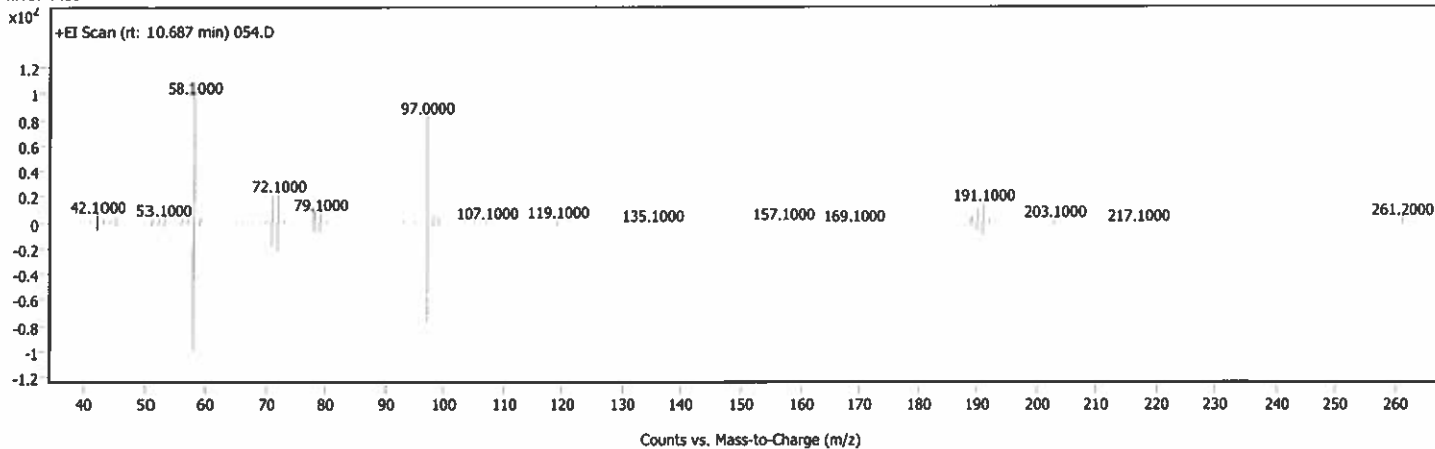
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.61	SWGDRUG 3.3.L	MATRIX 2/W ISTD 2.0 mg/L- BLD	KM/QC DATA IN XYLAZINE VALIDATION

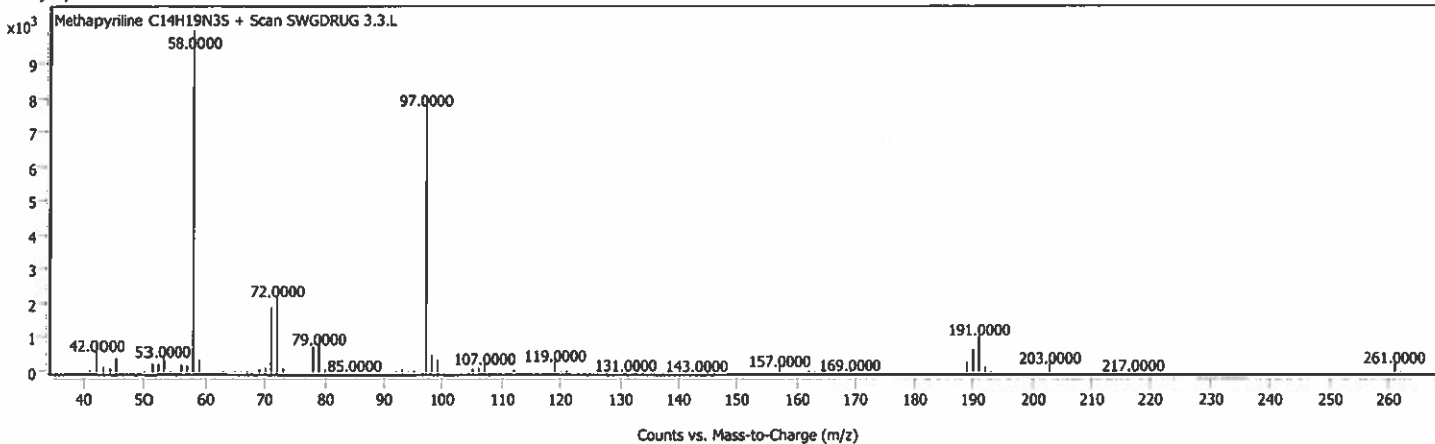
## Observed Spectrum



## Mirror Plot

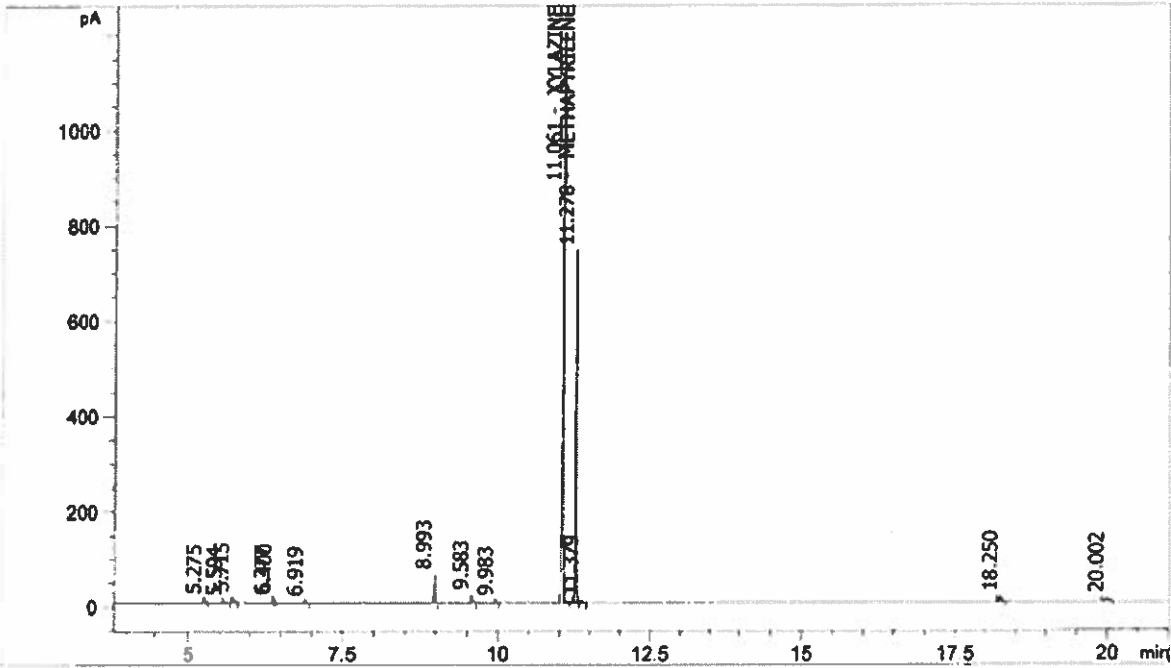


## Library Spectrum



Injection Date: 9/29/2021 2:05:03 PM Seq Line: 61  
 Sample Name: MATRIX 2 W/ISTD 2.0 m-> Vial 92  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



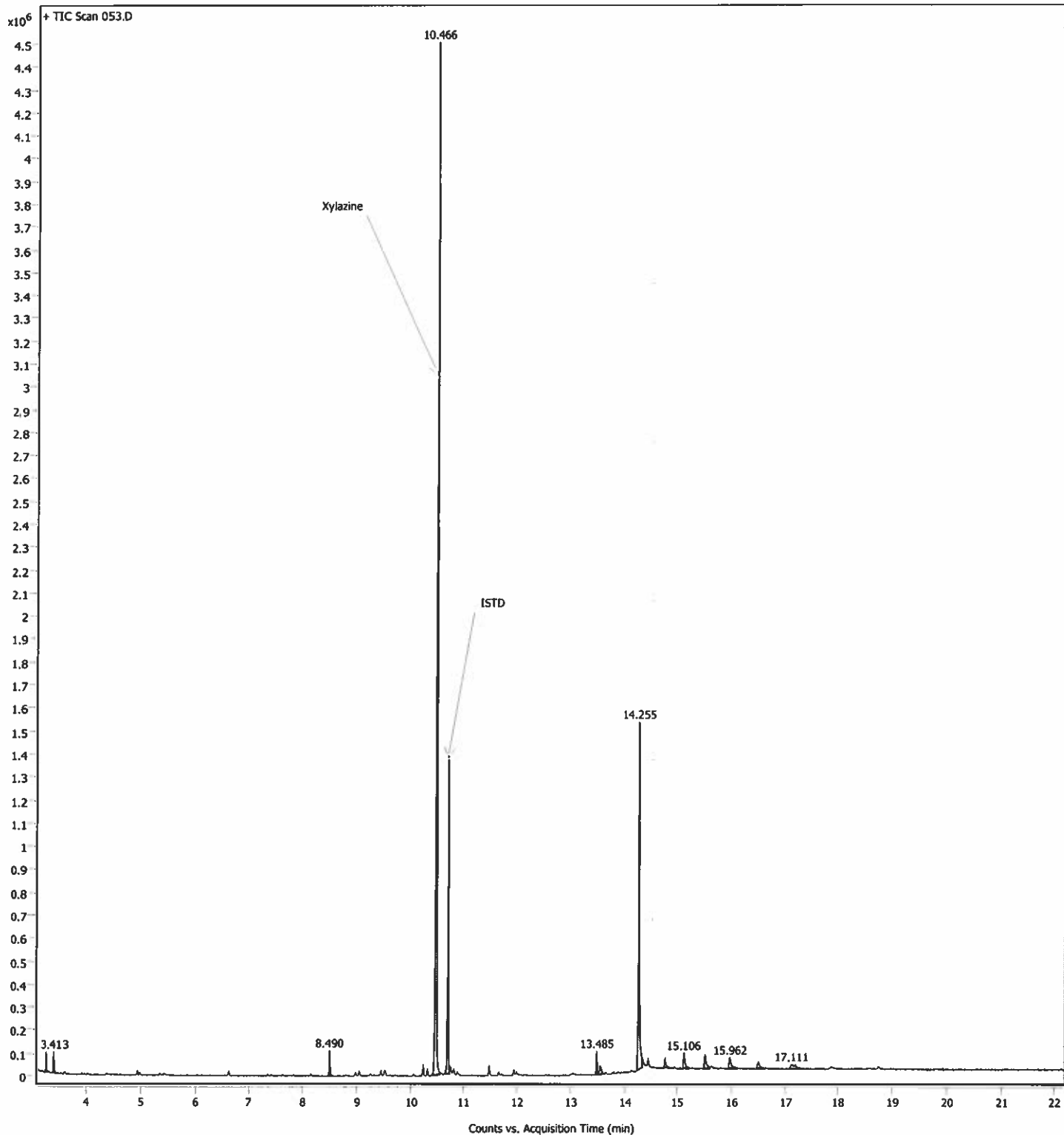
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.061	11.046	1756.485	1180.927	0.000000	XYLAZINE RRT - 0.9808
11.278	11.278	857.154	742.535	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 W/ISTD 2.0 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\053.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 2:39:17 PM (UTC-04:00)
Position	85	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

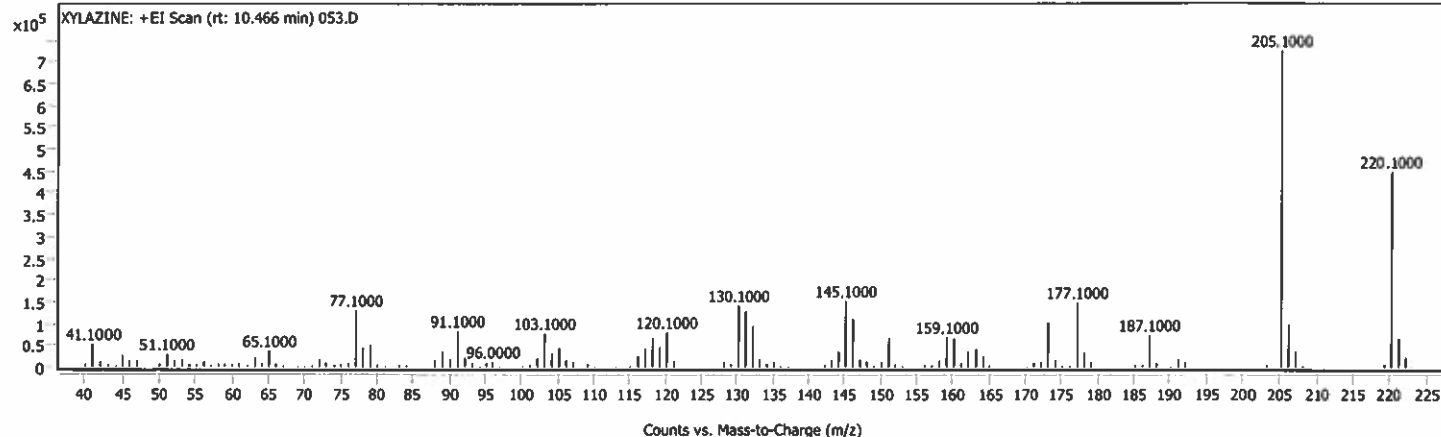
## Sample Spectra

+ Scan (rt: 10.466 min)

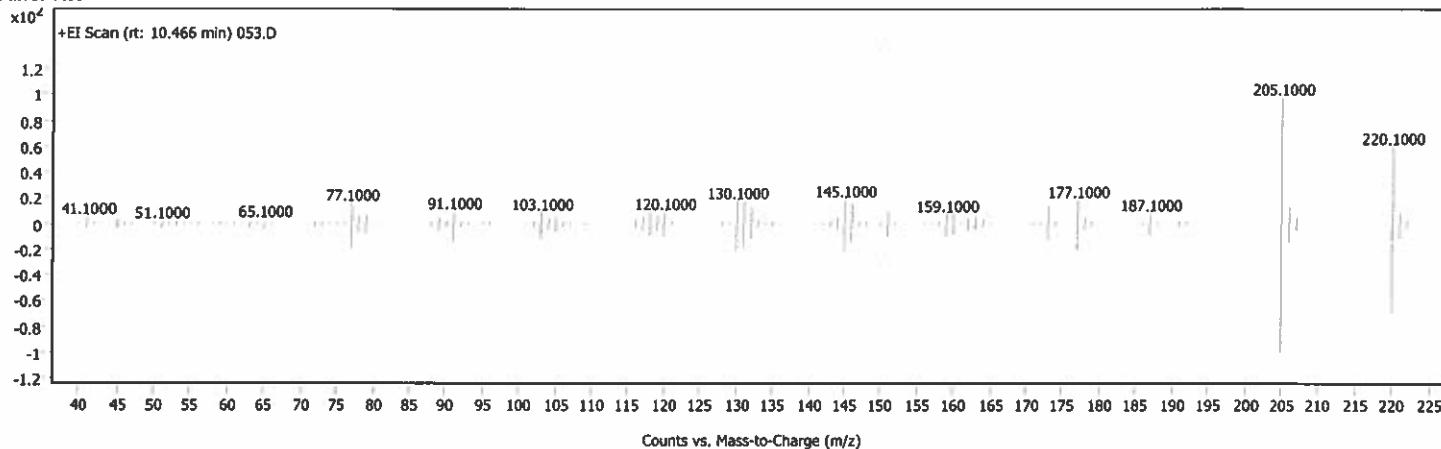
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.75	aafs.l	MATRIX 1 W/ISTD 2.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

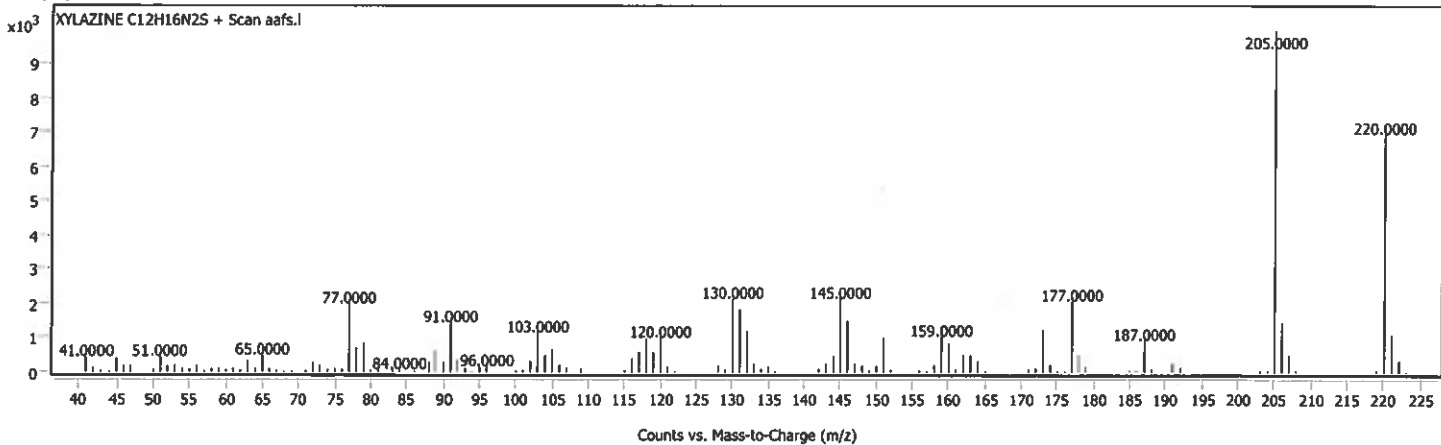
### Observed Spectrum



### Mirror Plot



### Library Spectrum



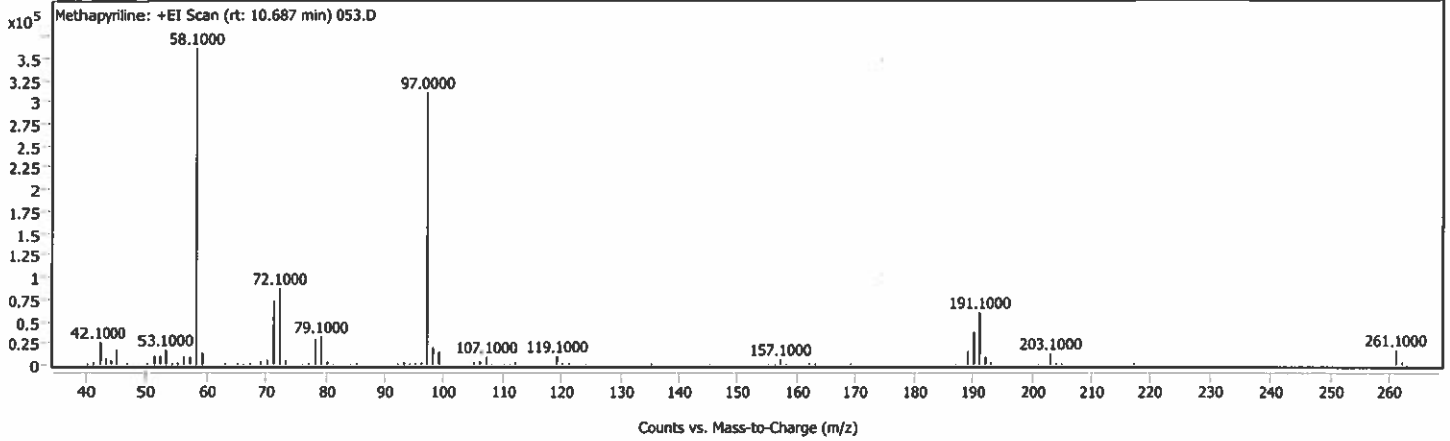
# Analysis Report

+ Scan (rt: 10.687 min)

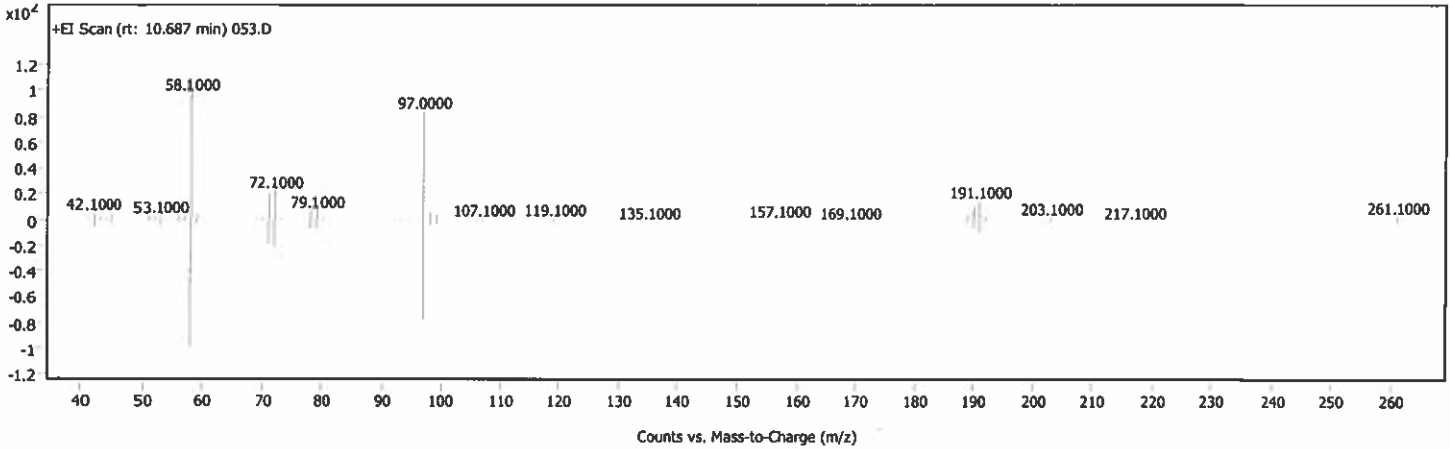
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.44	SWGDRUG 3.3.L	MATRIX 1 W/ISTD 2.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

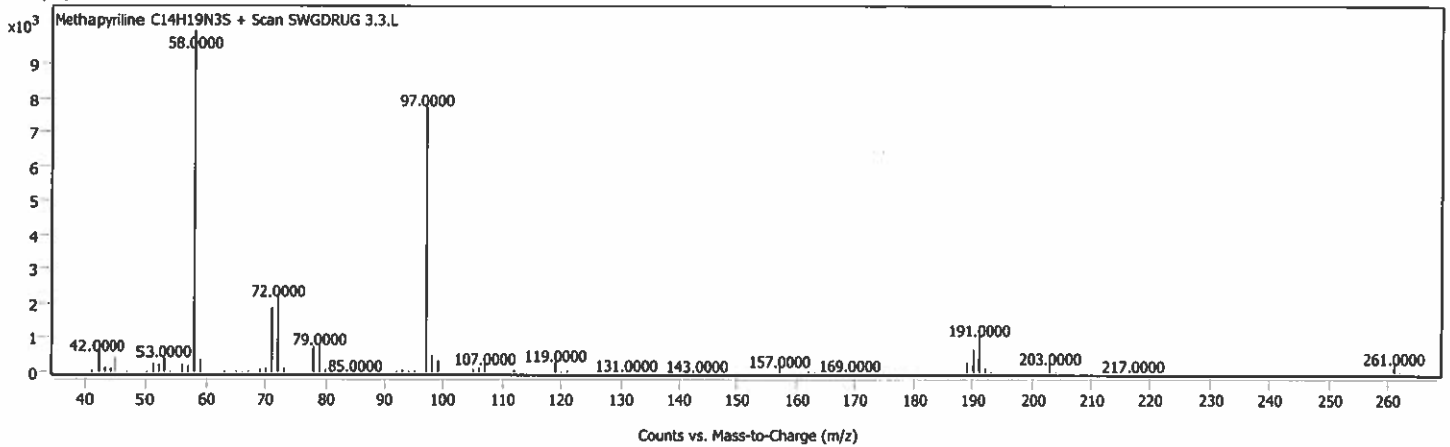
## Observed Spectrum



## Mirror Plot



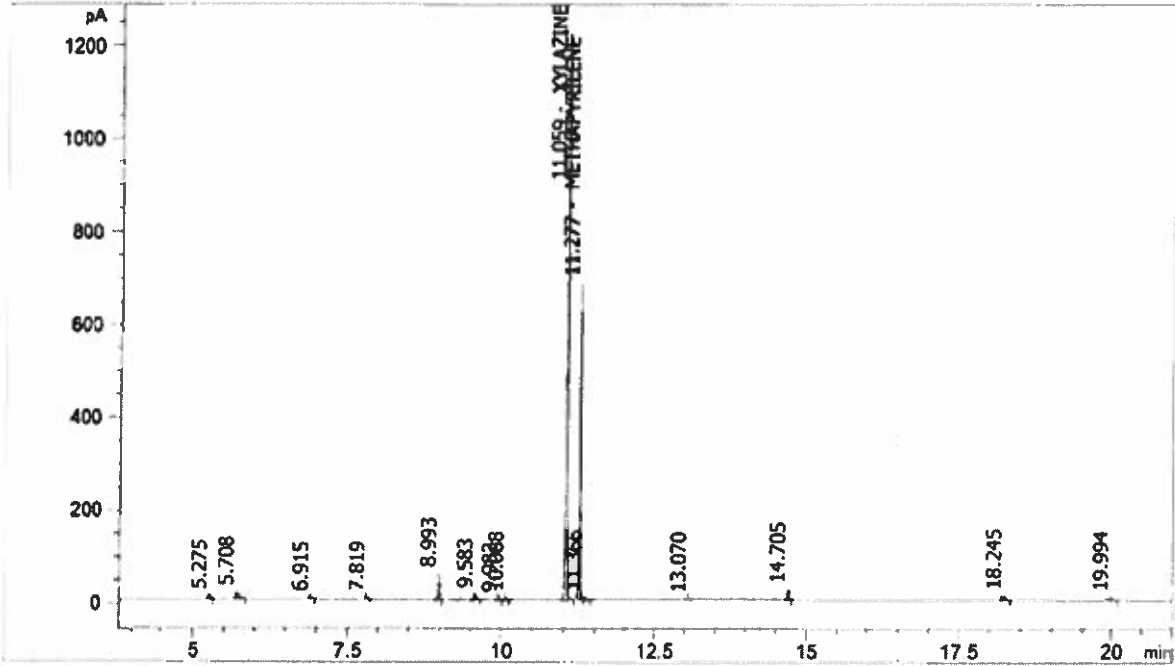
## Library Spectrum





Injection Date: 9/29/2021 1:40:30 PM Seq Line: 60  
 Sample Name: MATRIX 1 W/ISTD 2.0 m-> Vial 91  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



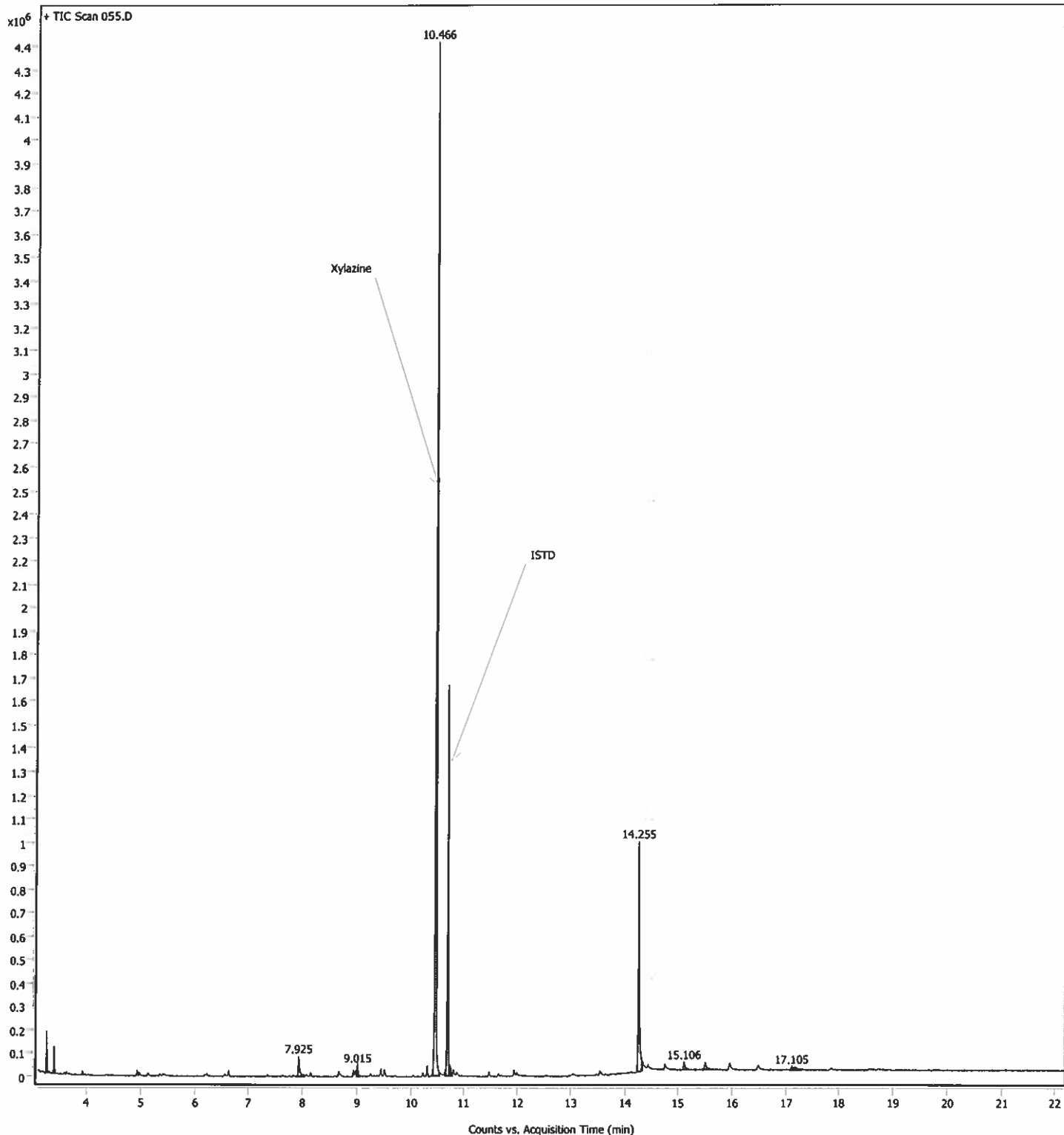
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.059	11.046	1795.546	1218.081	0.000000	XYLAZINE RRT - 0.9807
11.277	11.278	790.096	683.578	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 W/ISTD 2.0 mg/L- URN	Data File Path	C:\MassHunter\GCMS\1\data\BASE5\092821\055.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 3:30:58 PM (UTC-04:00)
Position	87	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

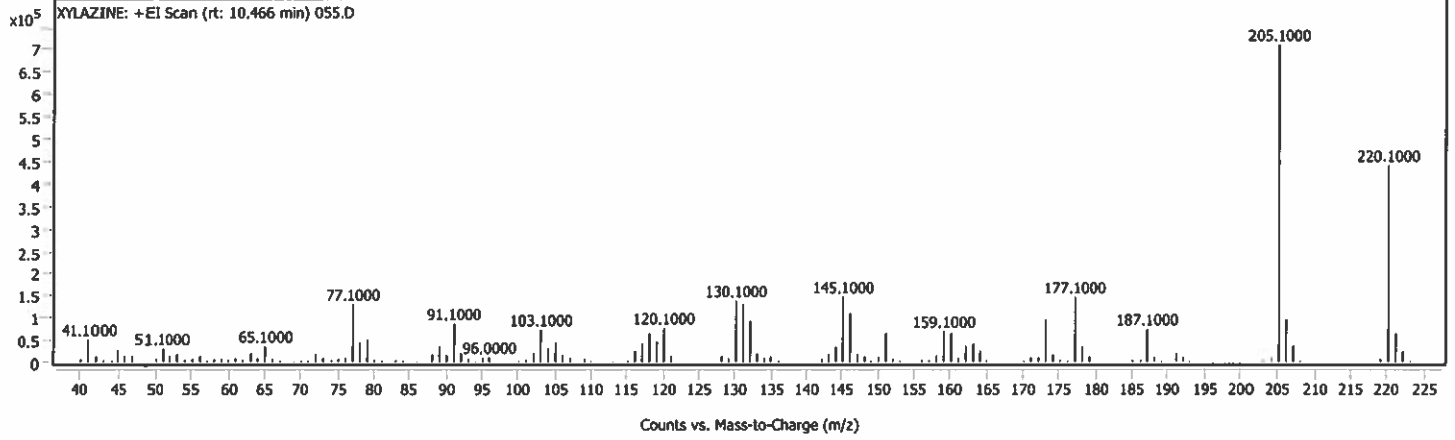
## Sample Spectra

+ Scan (rt: 10.466 min)

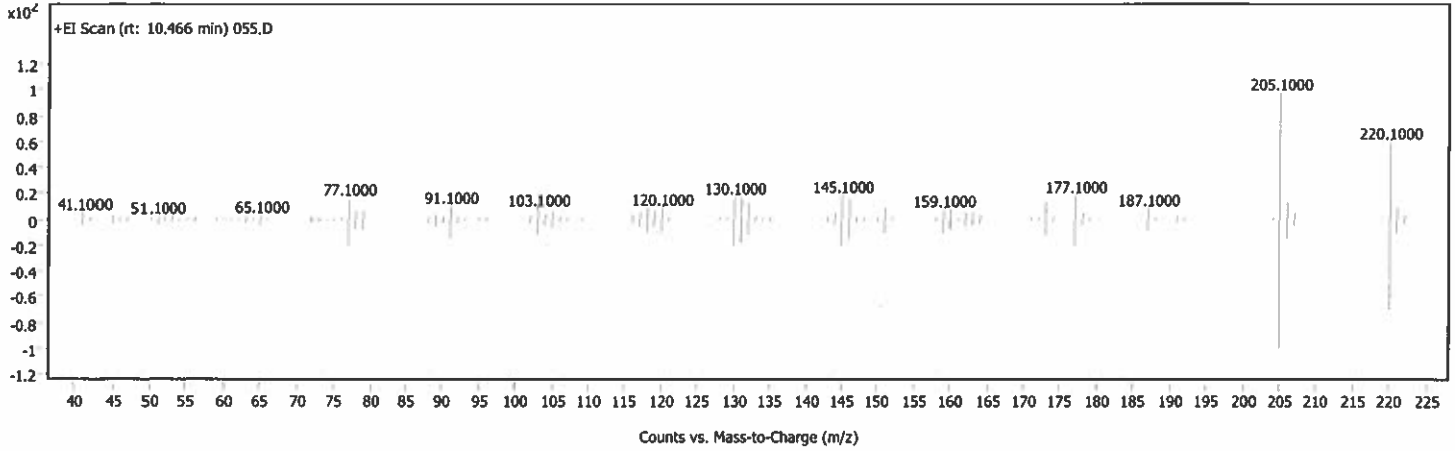
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	97.77	aafs.I	MATRIX 6 W/ISTD 2.0 mg/L- URN	KM/QC DATA IN XYLAZINE VALIDATION

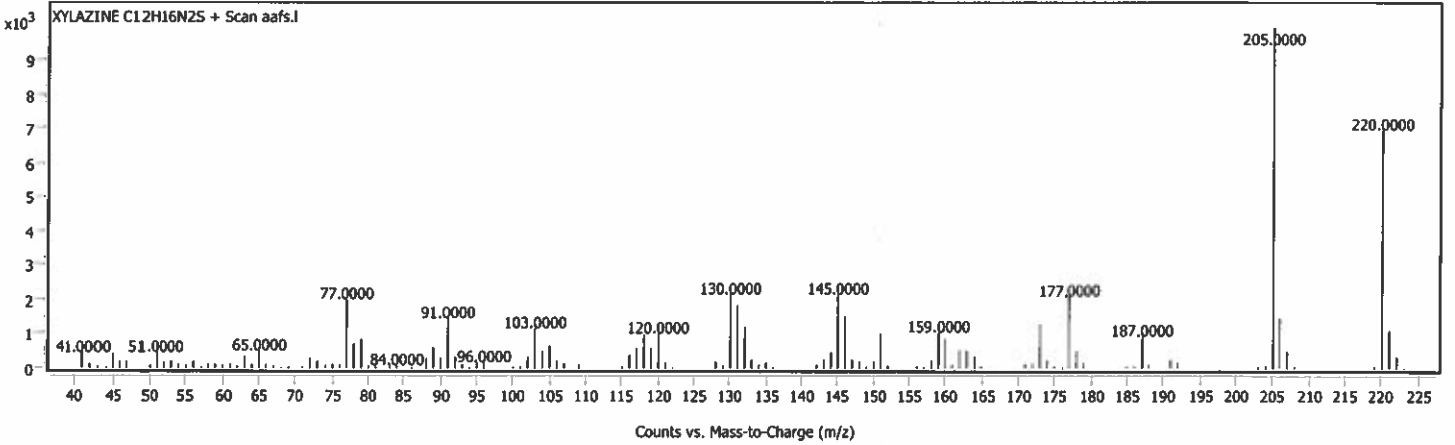
## Observed Spectrum



## Mirror Plot



## Library Spectrum



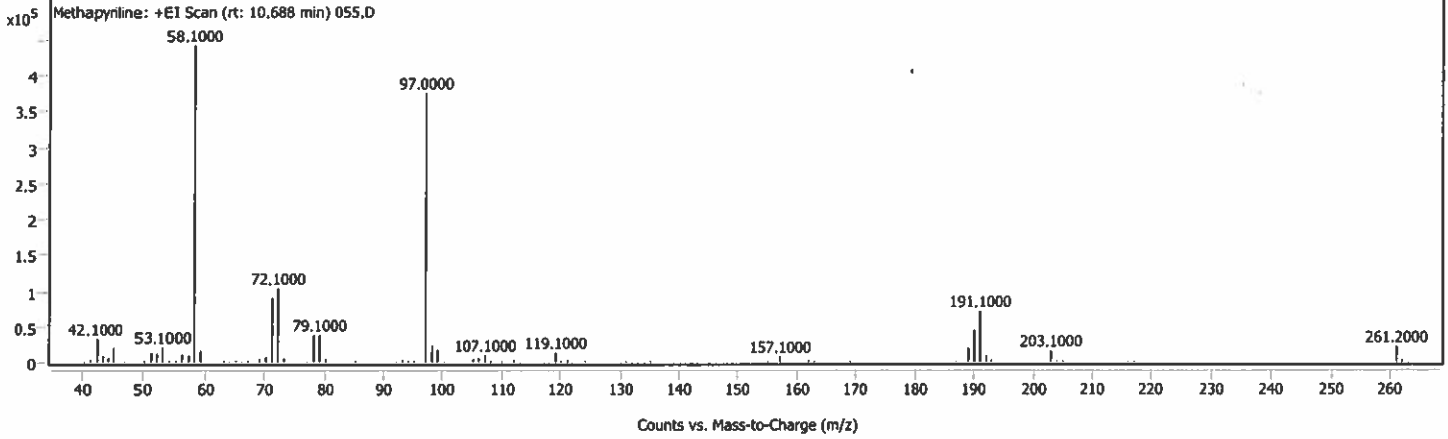
# Analysis Report

+ Scan (rt: 10.688 min)

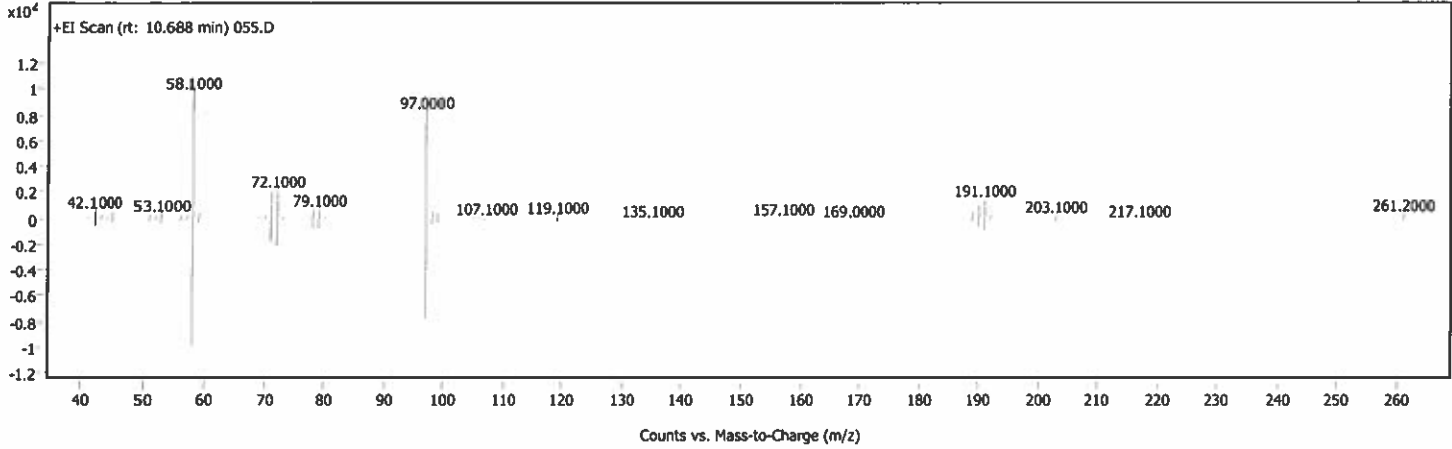
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.81	SWGDRUG 3.3.L	MATRIX 6 W/ISTD 2.0 mg/L- URN	KM/QC DATA IN XLAZINE VALIDATION

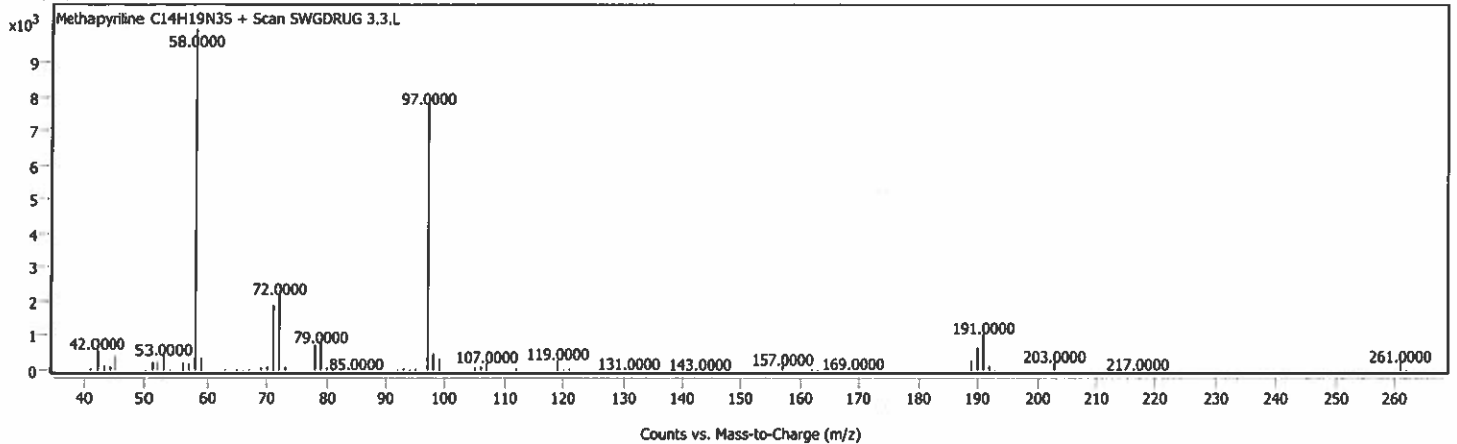
### Observed Spectrum



### Mirror Plot



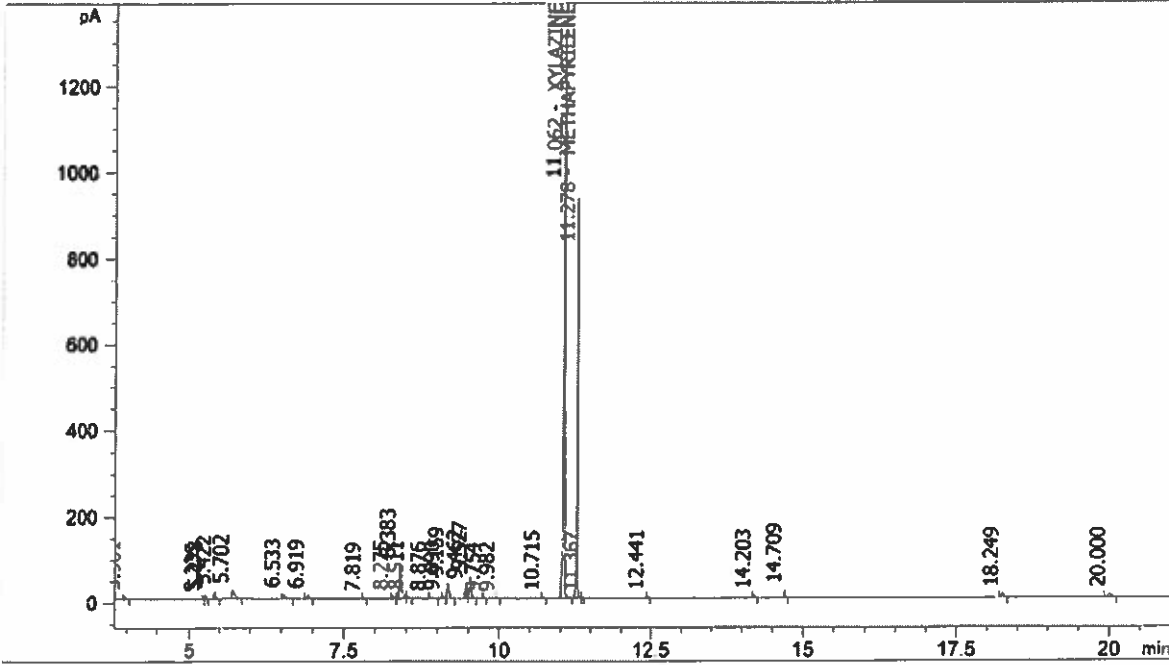
### Library Spectrum



Injection Date: 9/29/2021 2:29:22 PM  
 Sample Name: MATRIX 6 W/ISTD 2.0 m->  
 Sample Info:

Seq Line: 62  
 Vial 93

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



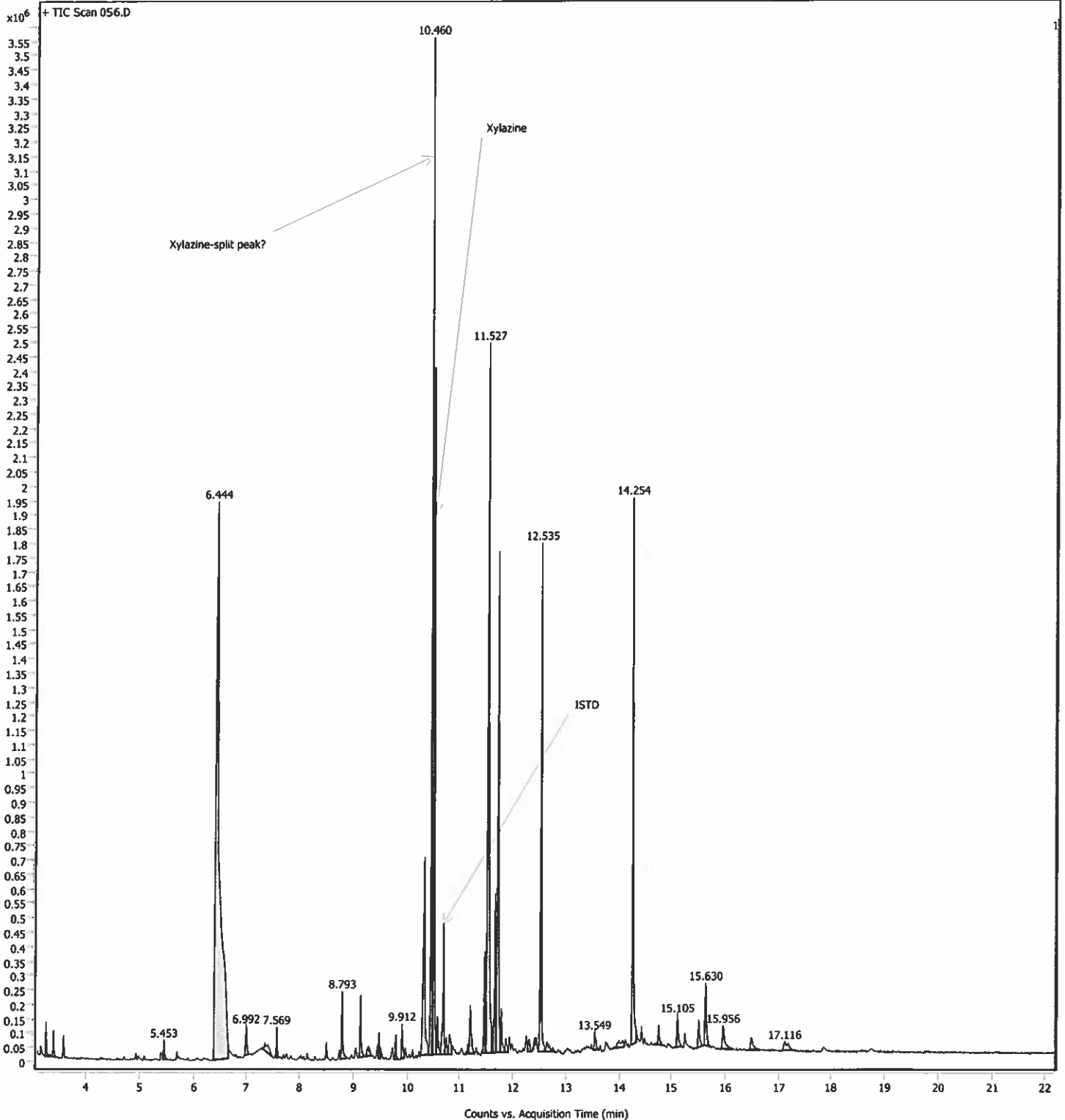
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.062	11.046	2051.870	1314.079	0.000000	XYLAZINE RRT - 0.9808
11.278	11.278	1103.267	929.718	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 10 W/ISTD 2.0 mg/L- LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\056.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 3:56:47 PM (UTC-04:00)
Position	88	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

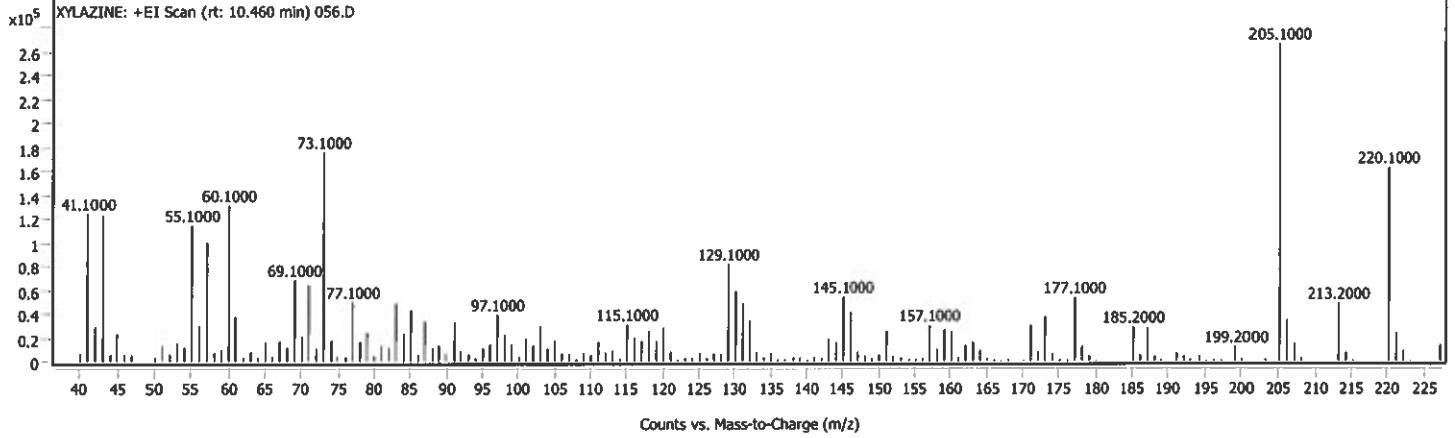
## Sample Spectra

+ Scan (rt: 10.460 min)

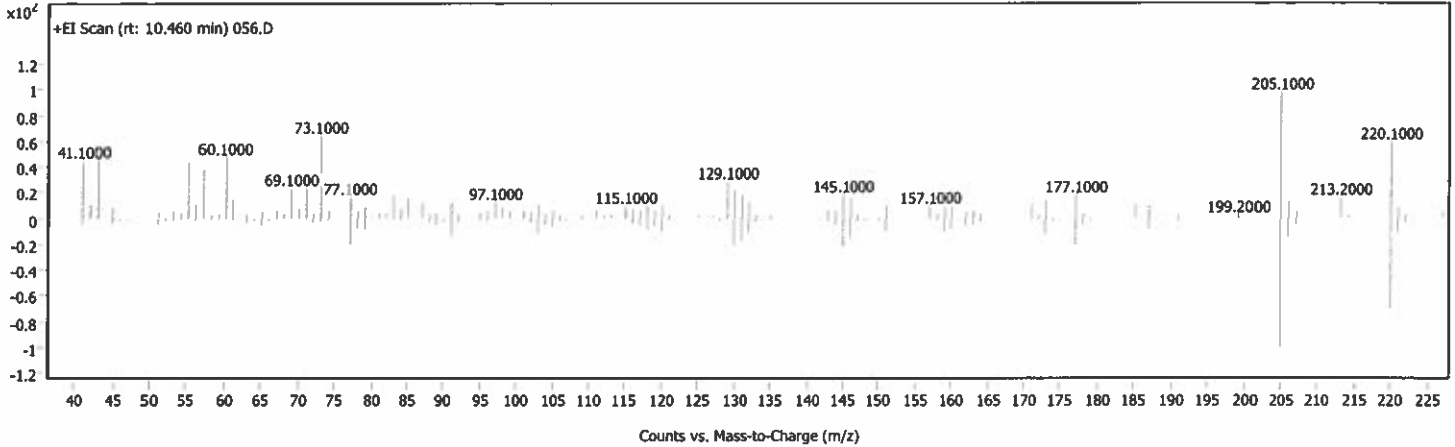
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	73.79	aafs.l	MATRIX 10 W/ISTD 2.0 mg/L- LVR	KM/QC DATA IN XYLAZINE VALIDATION

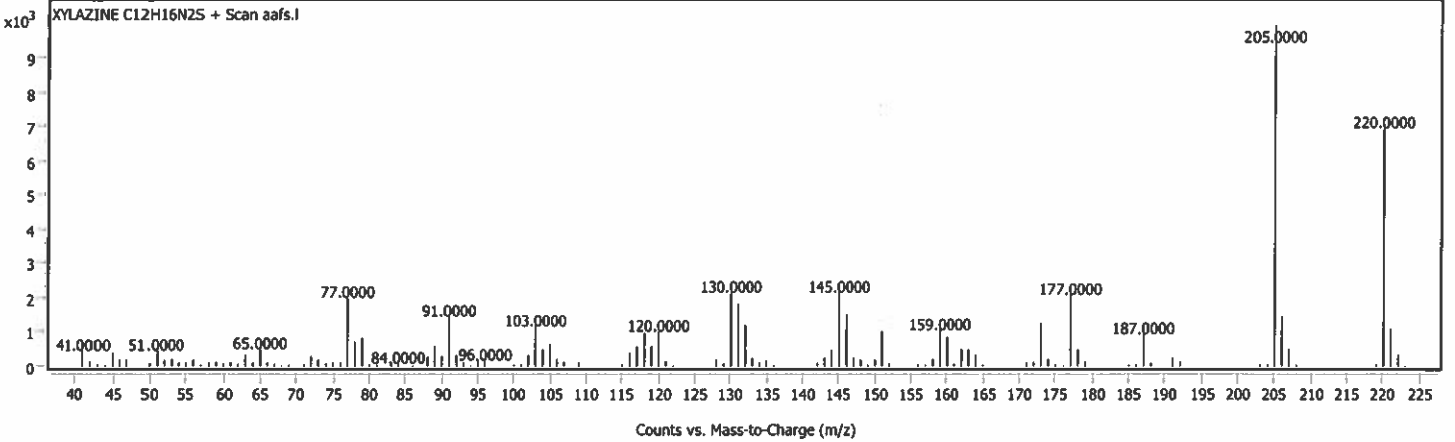
## Observed Spectrum



## Mirror Plot



## Library Spectrum



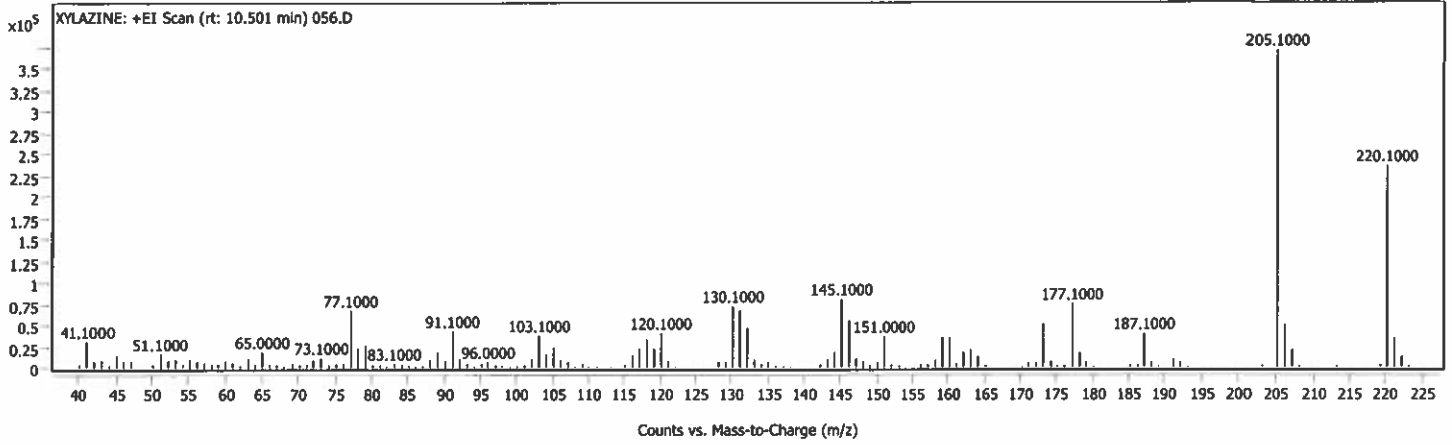
# Analysis Report

+ Scan (rt: 10.501 min)

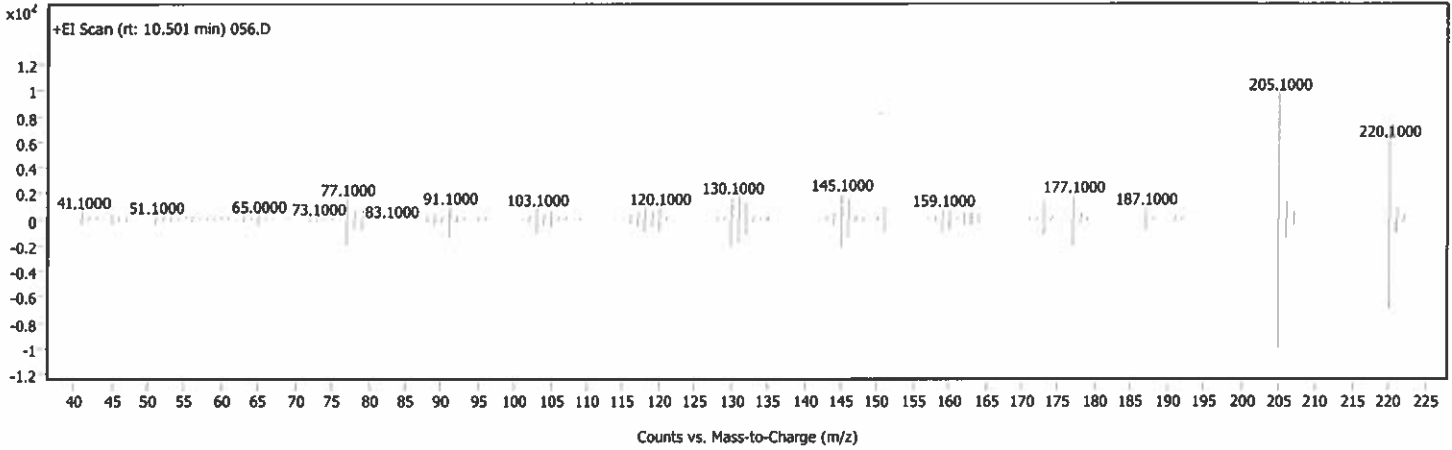
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	96.89	aafs.l	MATRIX 10 W/ISTD 2.0 mg/L- LVR	KM/QC DATA IN XYLAZINE VALIDATION

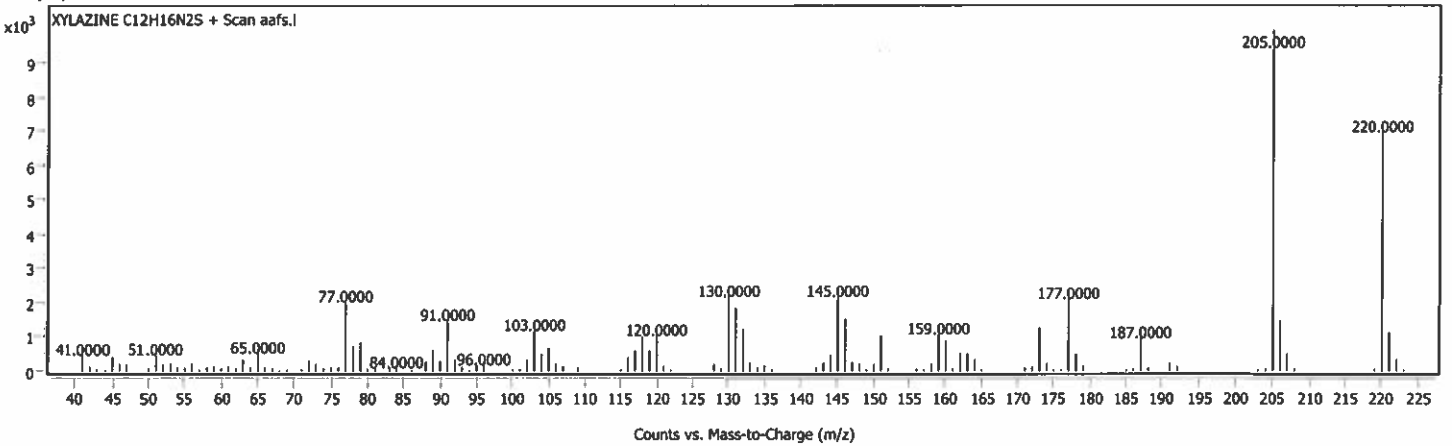
## Observed Spectrum



## Mirror Plot



## Library Spectrum





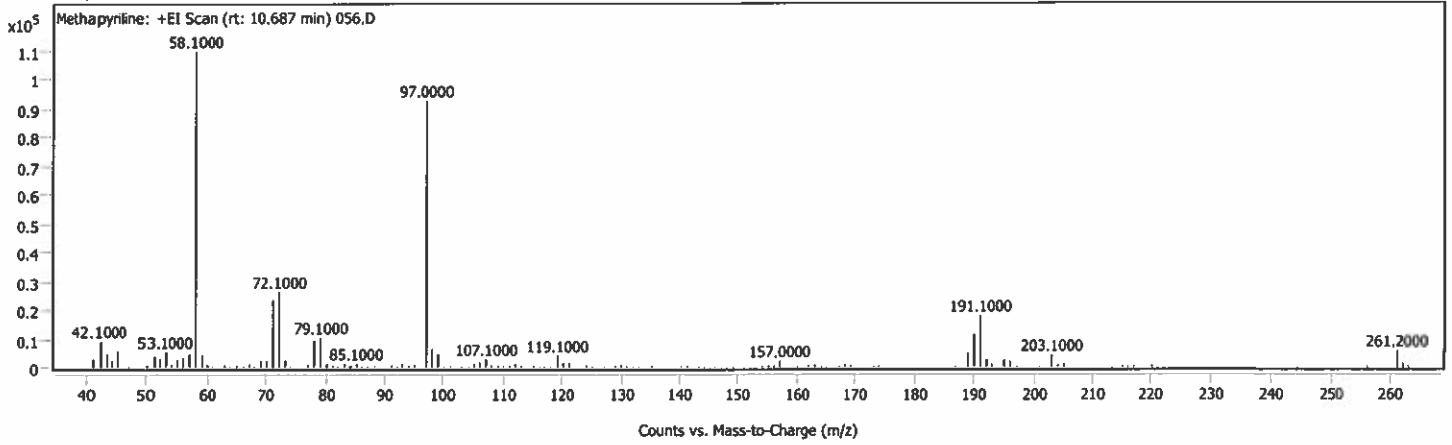
# Analysis Report

+ Scan (rt: 10.687 min)

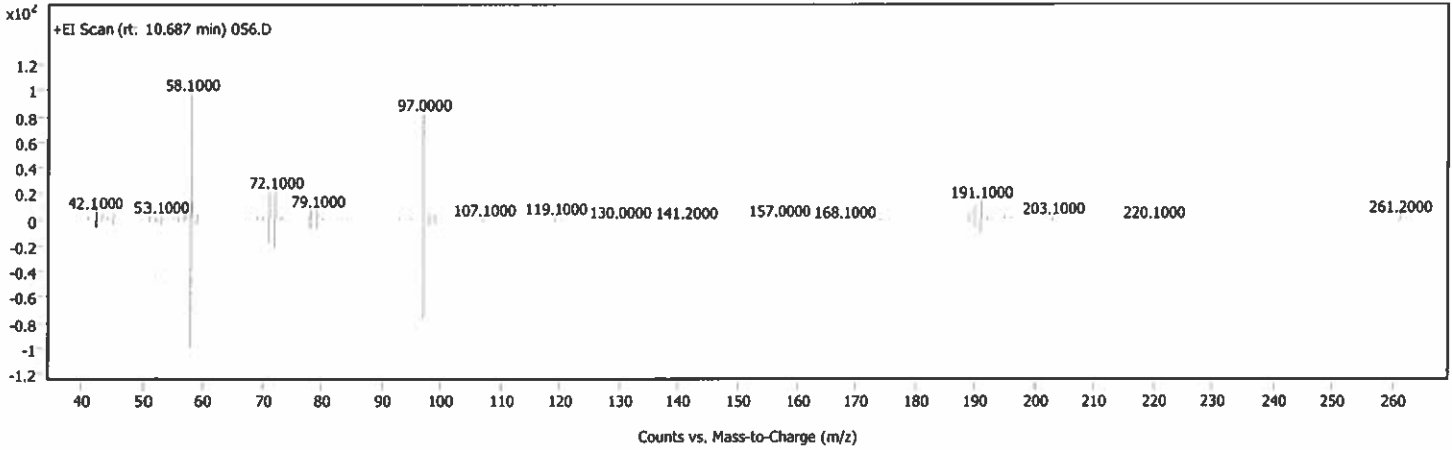
Methapyrine; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyrine	C14H19N3S	93.38	SWGDRUG 3.3.L	MATRIX 10 W/ISTD 2.0 mg/L- LVR	KM/QC DATA IN XYLAZINE VALIDATION

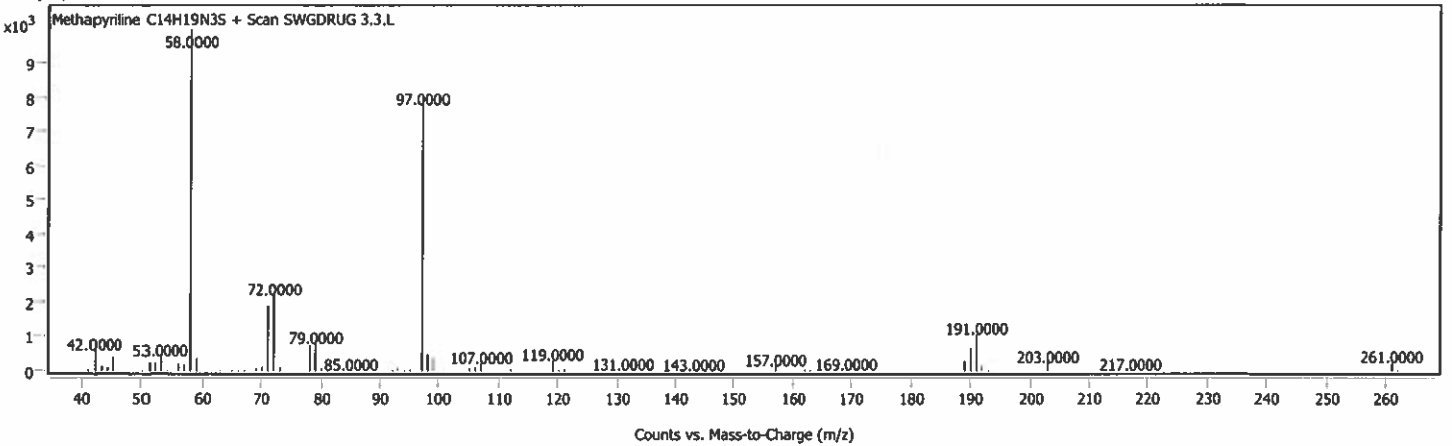
## Observed Spectrum



## Mirror Plot



## Library Spectrum

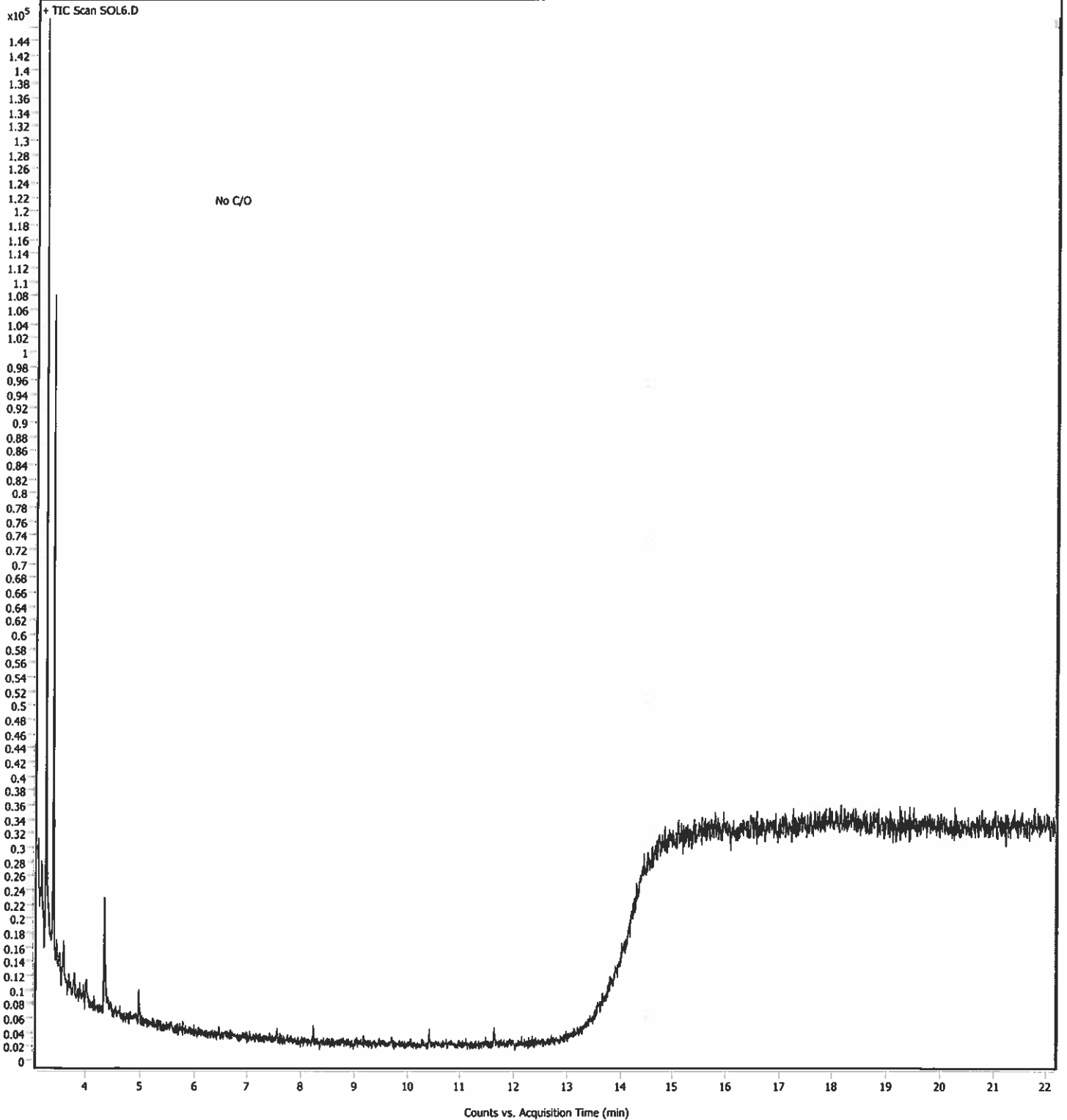


# Analysis Report

## Sample Information

Sample Name	Solvent Blank 6	Data File Path	C:\MassHunter\GCMS\1\data\BASE5\092821\SOL6.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 4:22:39 PM (UTC-04:00)
Position	89	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms

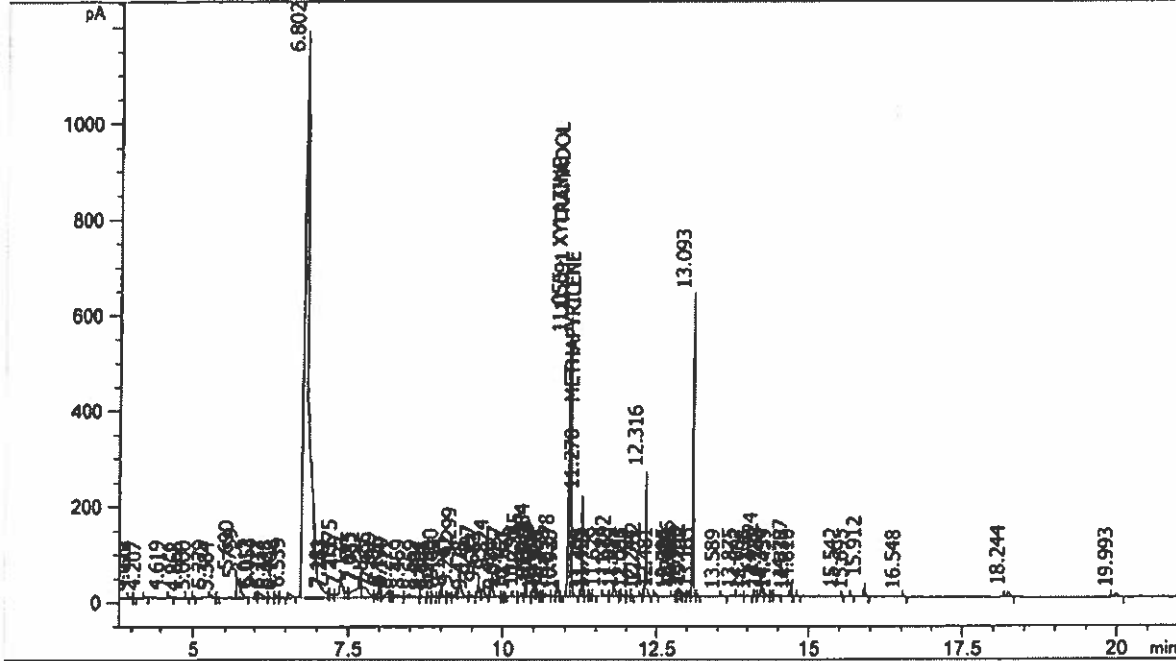


Injection Date: 9/29/2021 2:53:40 PM  
 Sample Name: MATRIX 10 W/ISTD 2.0 ->  
 Sample Info:

Seq Line: 63  
 Vial 94

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

Split peak-OK for qual reporting



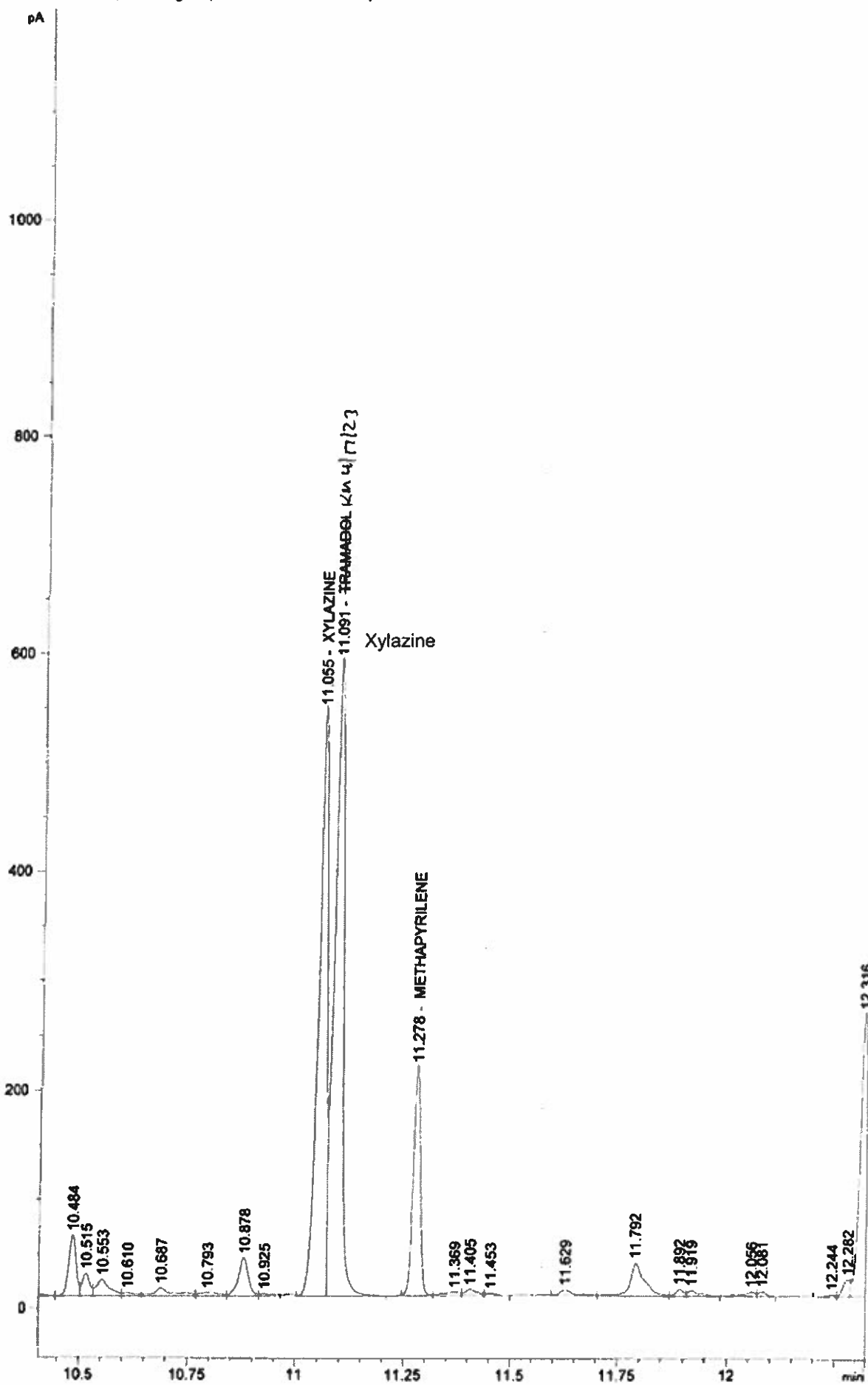
see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.055	11.046	724.084	538.254	0.000000	XYLAZINE
11.091	11.033	857.036	583.173	0.000000	TRAMADOL
11.278	11.278	242.928	210.837	1.000000	METHAPYRILENE

Not used to average RRTs

Current Chromatogram (s)

NPD1 A, Front Signal (092821\BASE000063.D)



Injection Date: 9/29/2021 3:18:03 PM

Seq Line: 64

Sample Name: SOLVENT BLANK 6

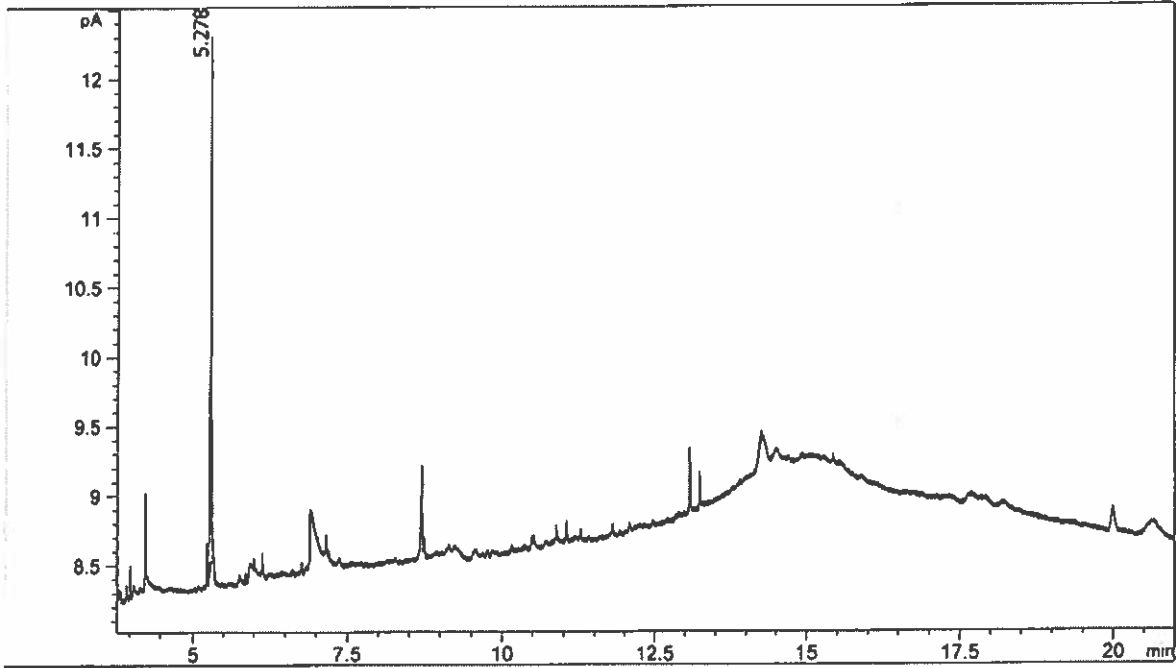
Vial 95

Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M

Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M

KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

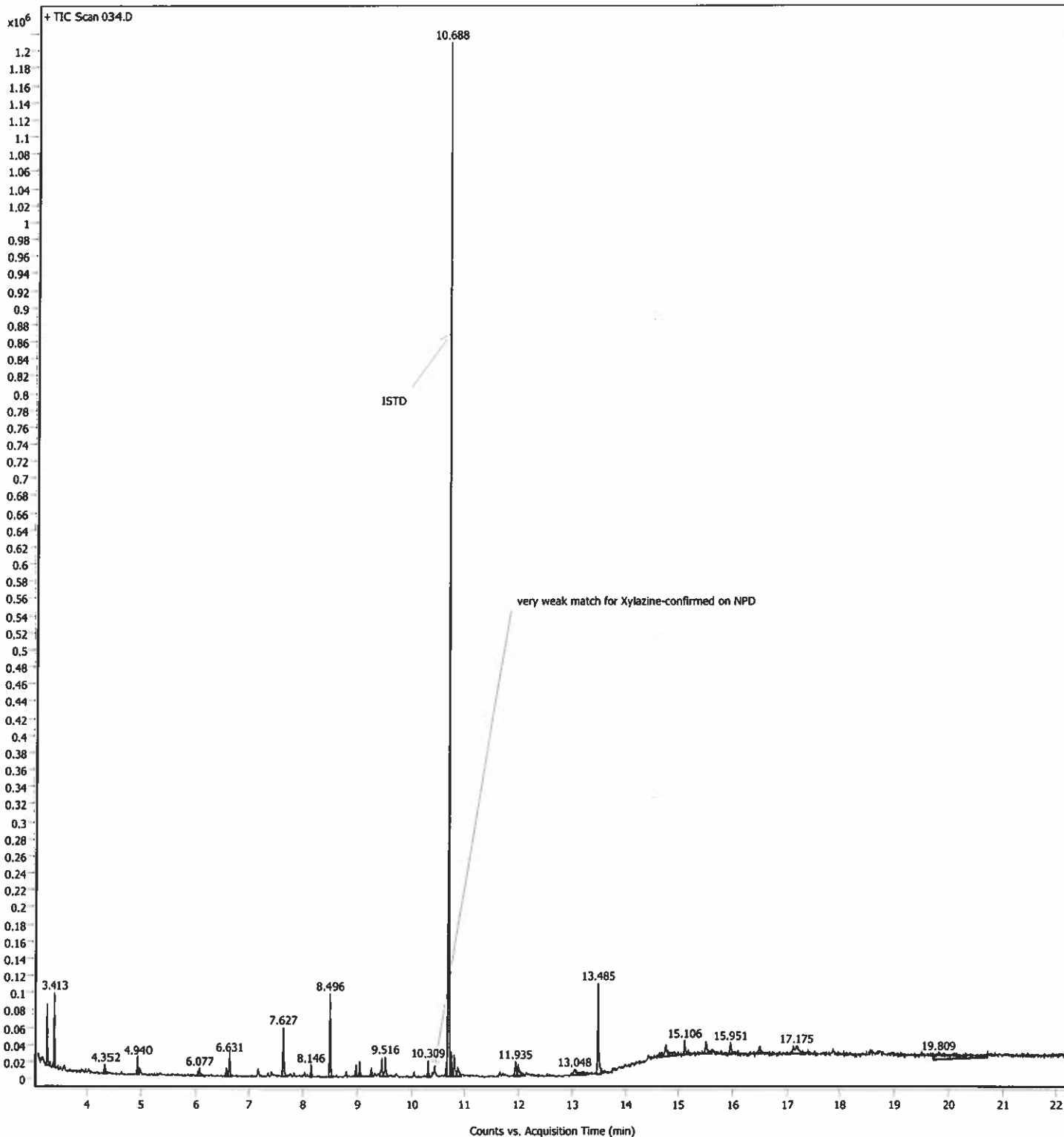
KM/QC DATA IN XYLAZINE VALIDATION

# Analysis Report

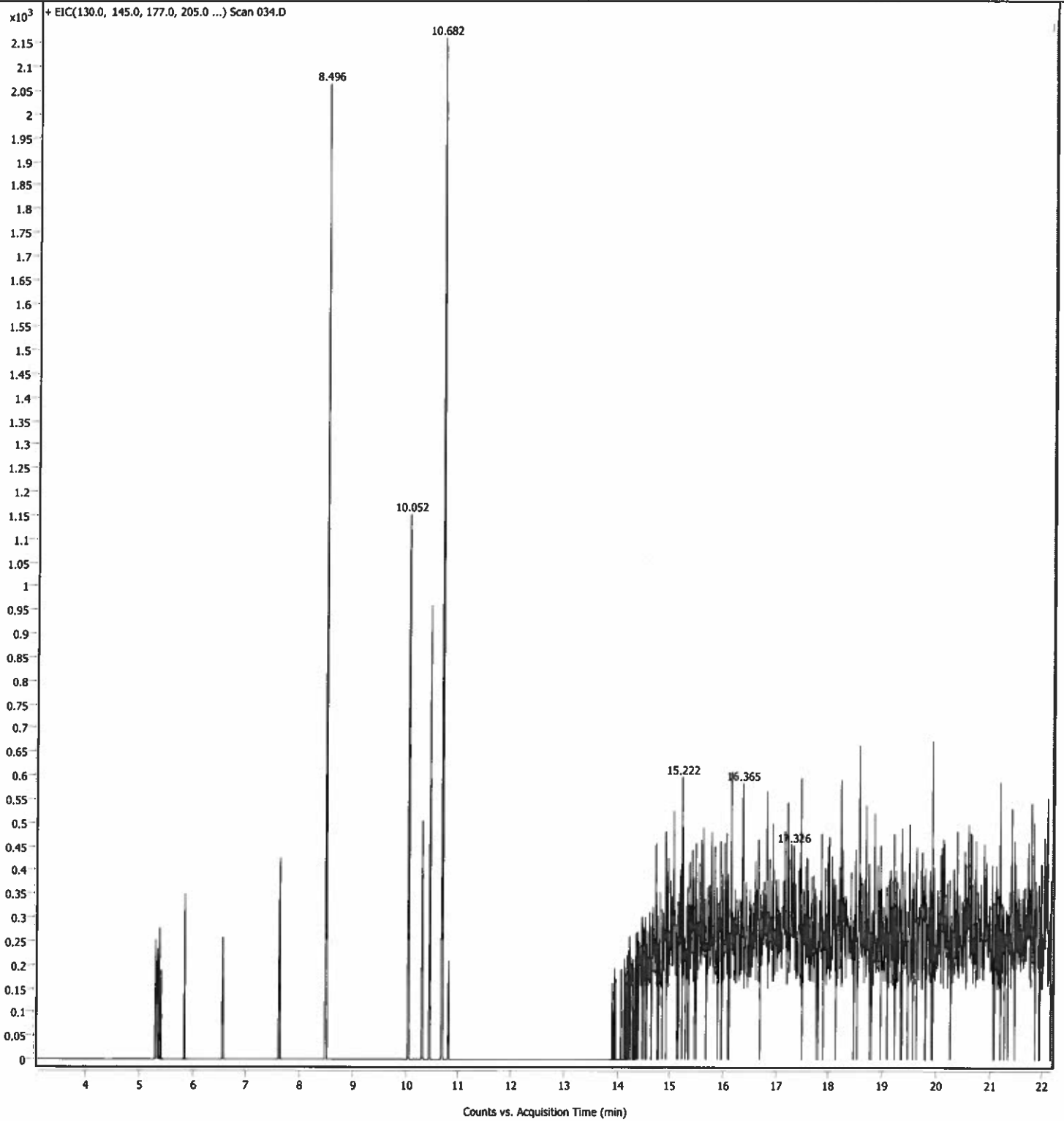
## Sample Information

Sample Name	MATRIX 2 LOD - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\034.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 6:02:19 AM (UTC-04:00)
Position	65	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report



# Analysis Report

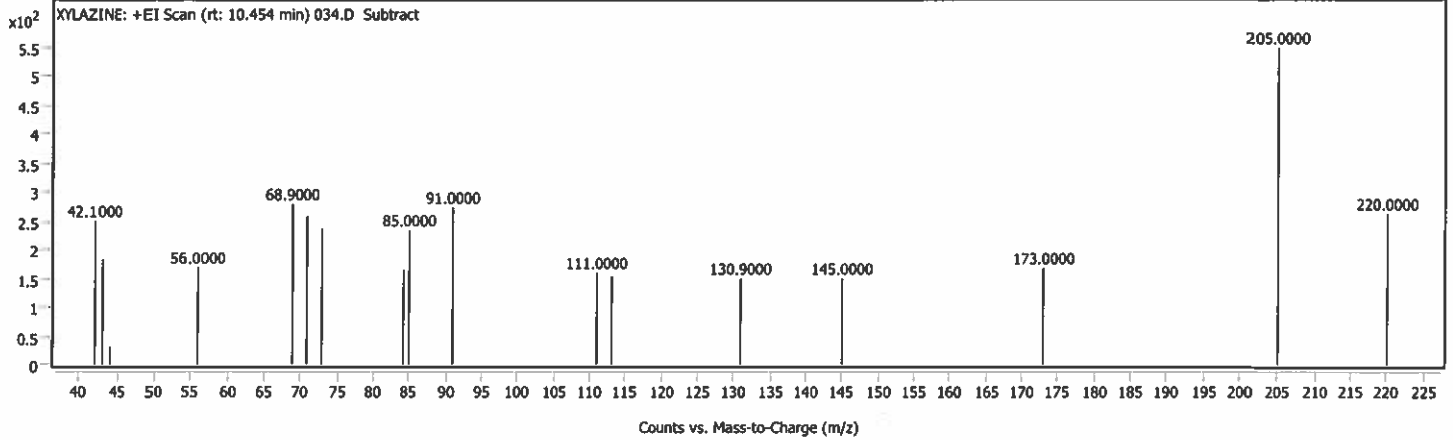
## Sample Spectra

+ Scan (rt: 10.425 min)

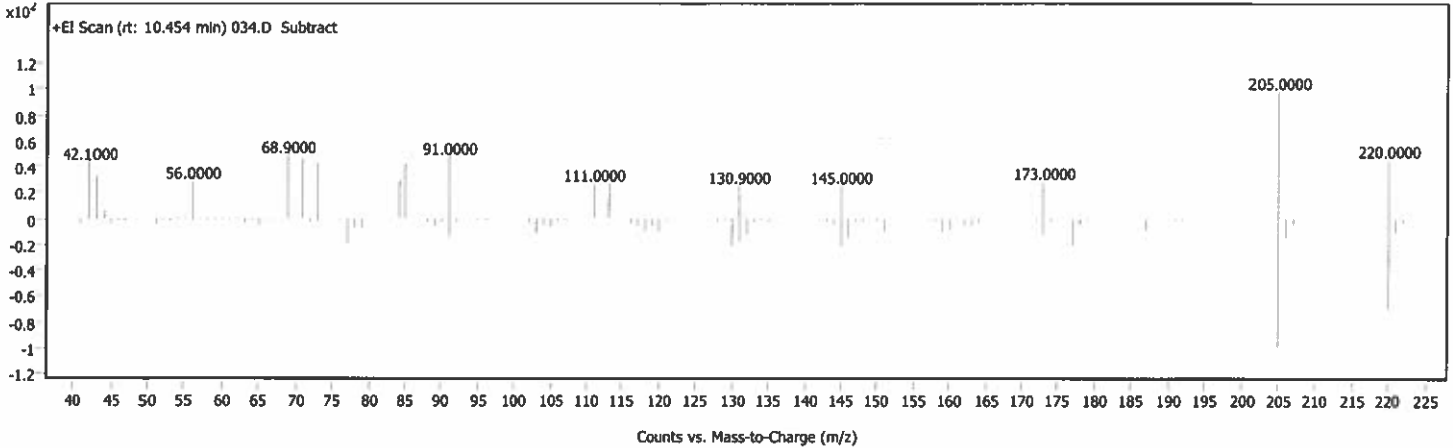
+ Scan (rt: 10.454 min) Sub **XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	34.22	aafs.l	MATRIX 2 LOD - BLD	KM/QC DATA IN XYLAZINE VALIDATION

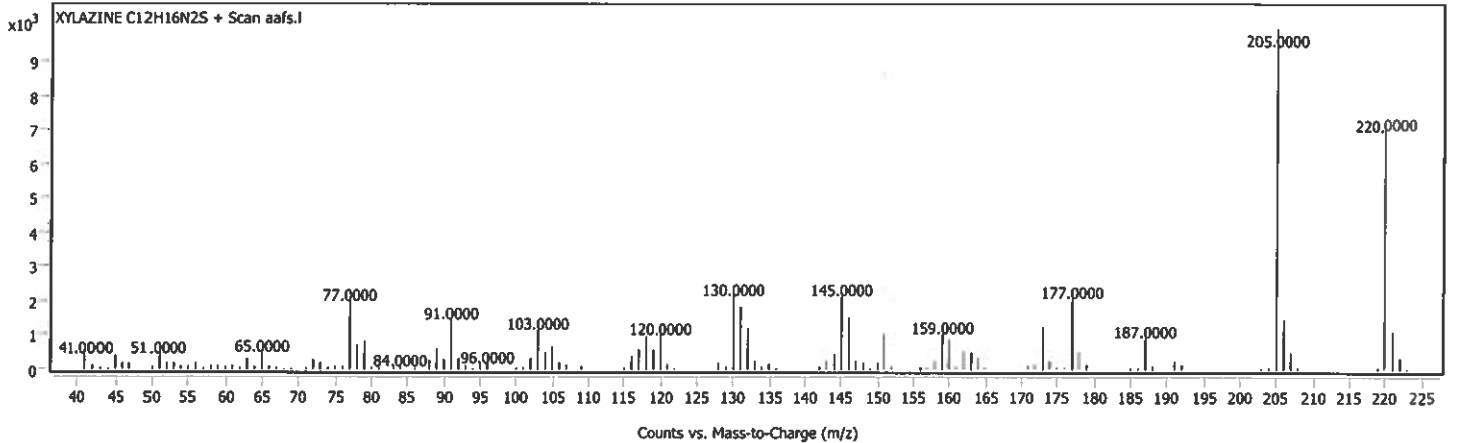
## Observed Spectrum



## Mirror Plot



## Library Spectrum





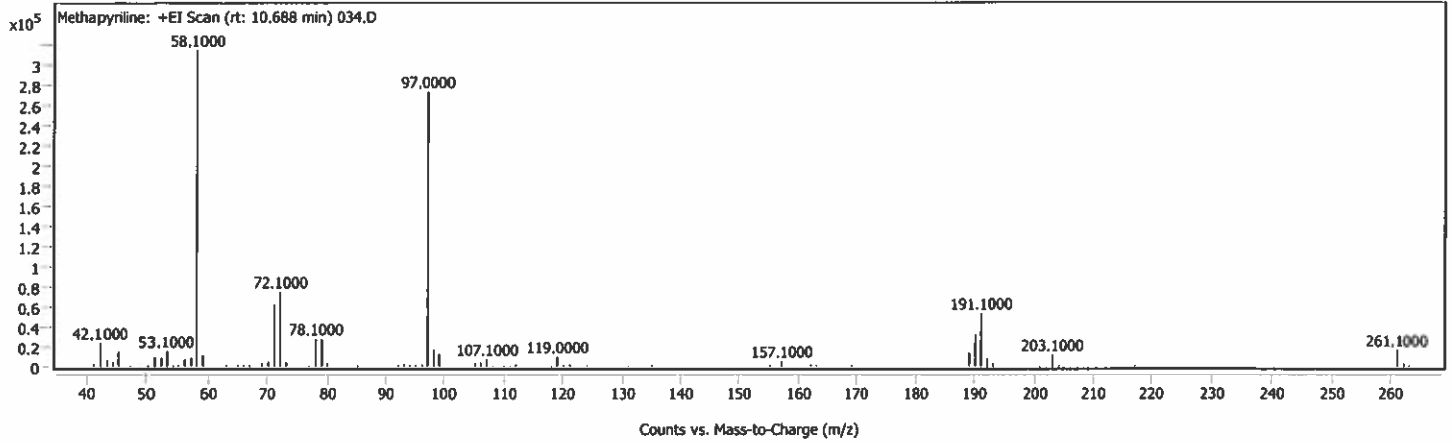
# Analysis Report

+ Scan (rt: 10.688 min)

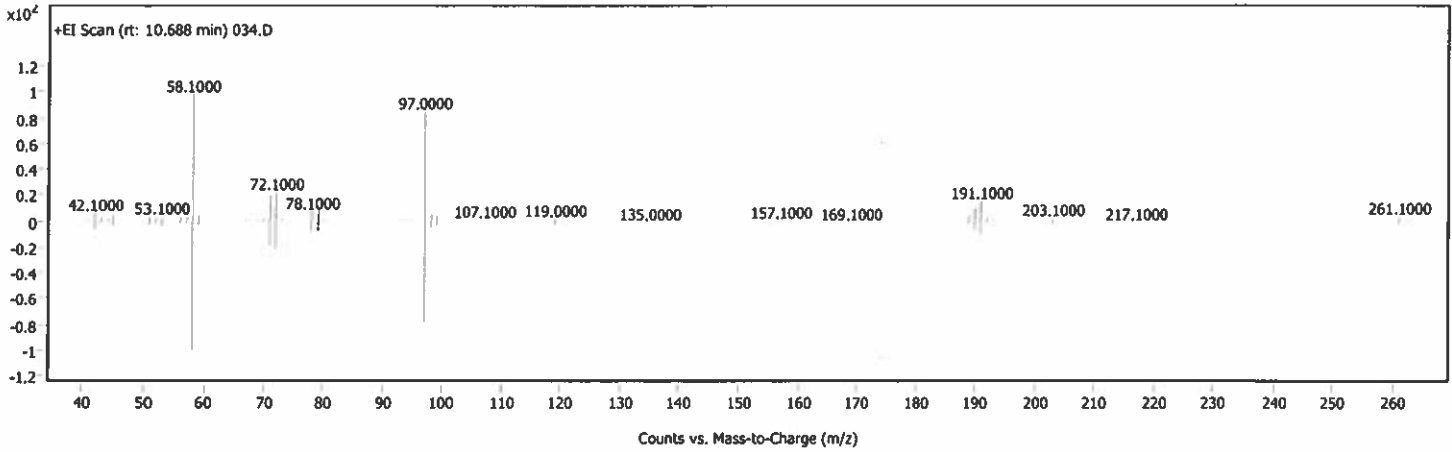
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.54	SWGDRUG 3.3.L	MATRIX 2 LOD - BLD	KM/QC DATA IN XYLAZINE VALIDATION

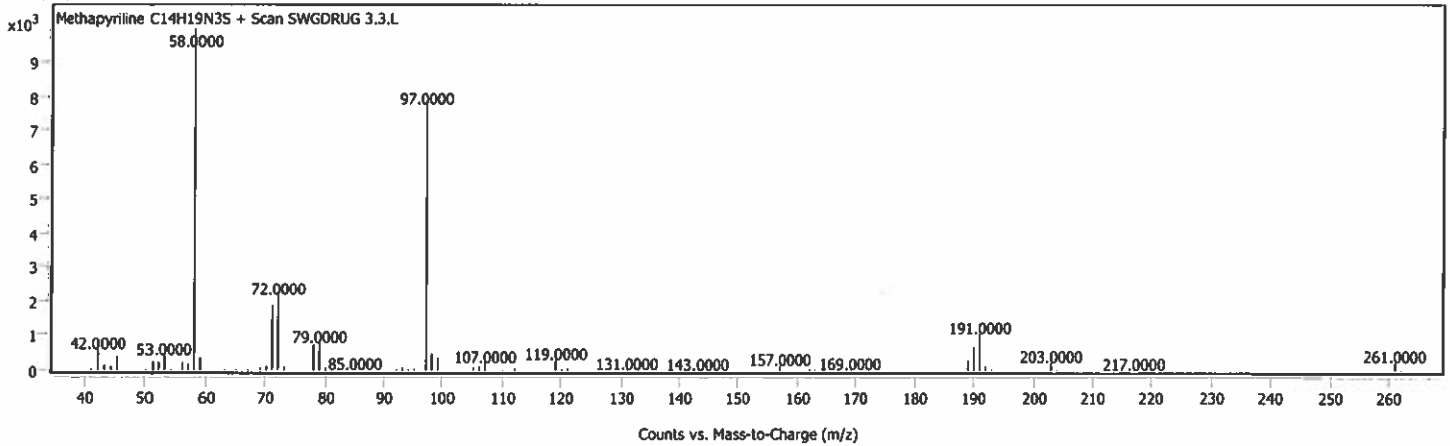
## Observed Spectrum



## Mirror Plot

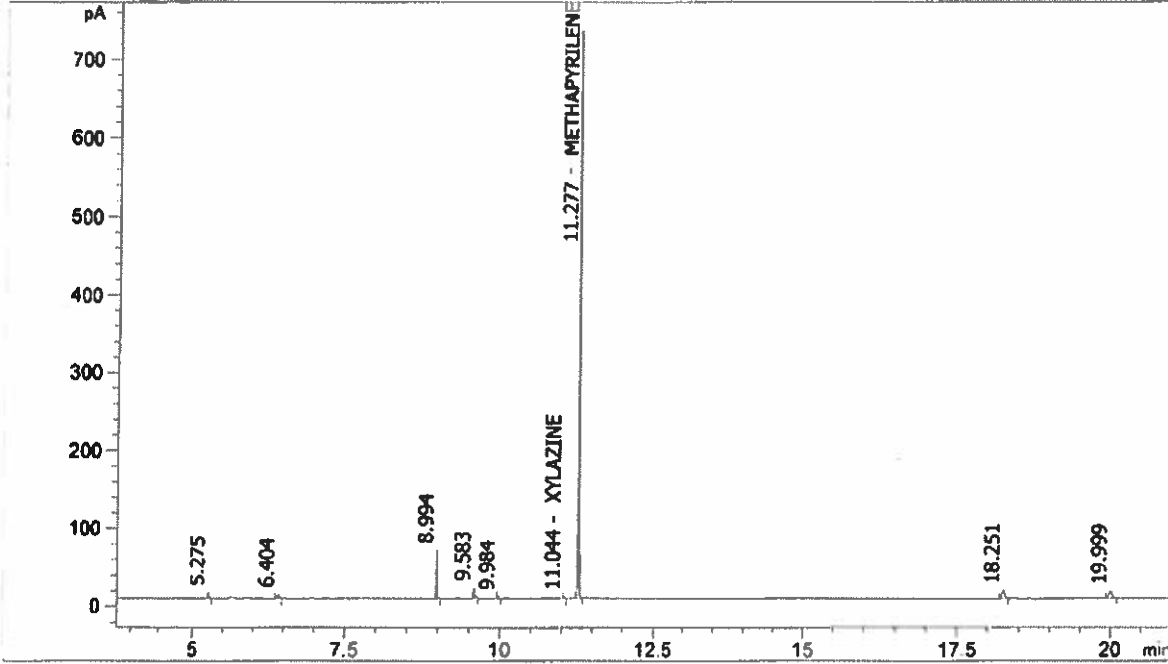


## Library Spectrum



Injection Date: 9/29/2021 5:34:00 AM Seq Line: 40  
 Sample Name: MATRIX 2 LOD - BLD Vial 41  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



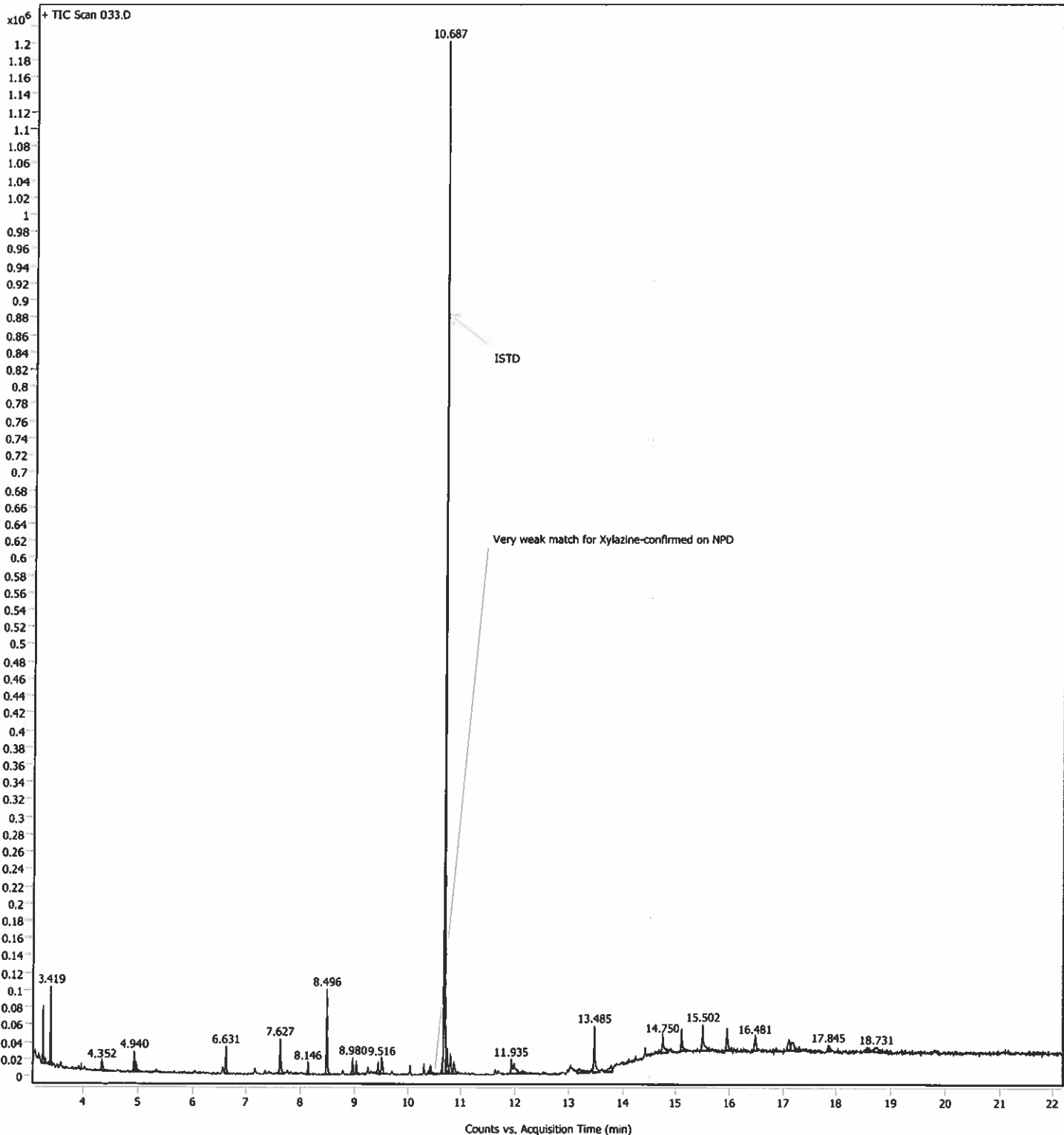
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.044	11.046	3.568	2.702	0.000000	XYLAZINE RRT - 0.9793
11.277	11.278	819.356	726.896	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

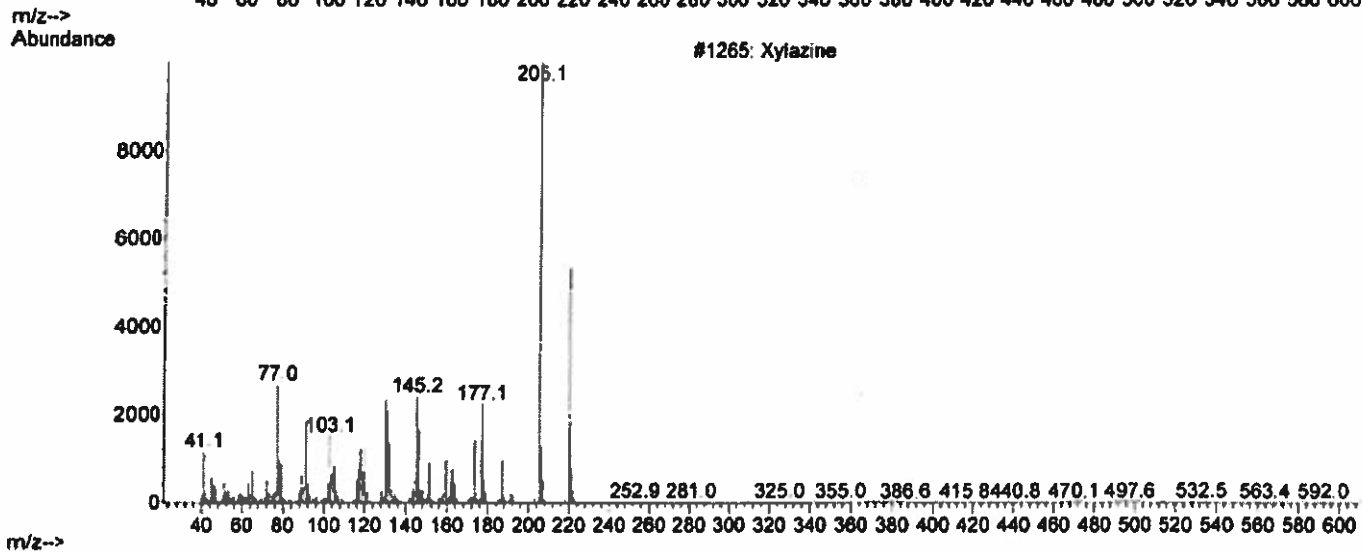
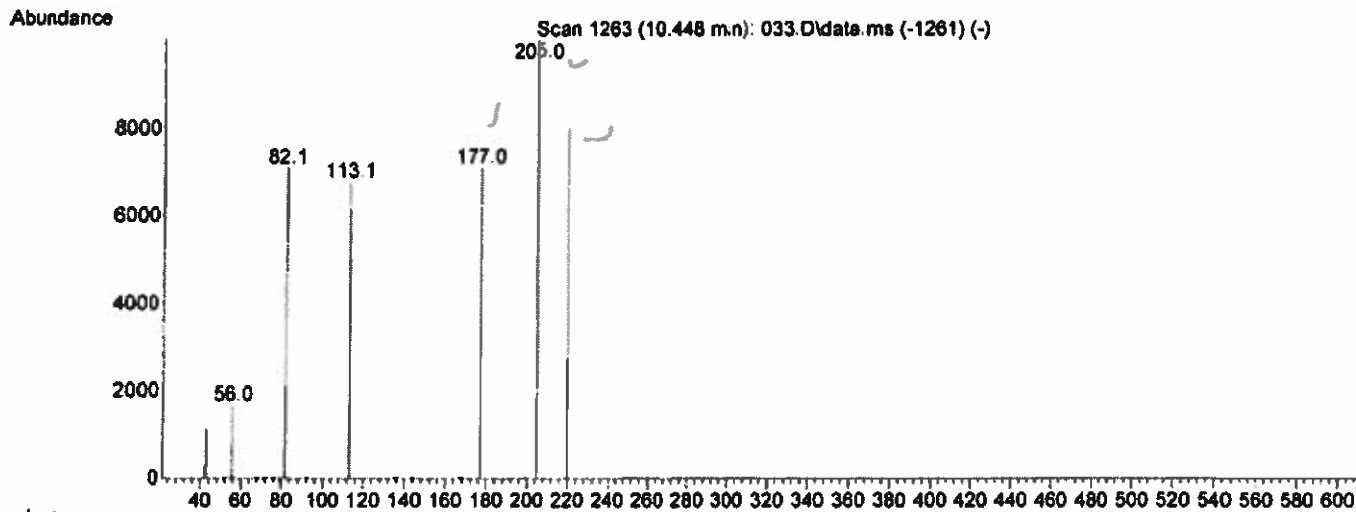
Sample Name	MATRIX 1 LOD - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\033.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 5:36:24 AM (UTC-04:00)
Position	64	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms

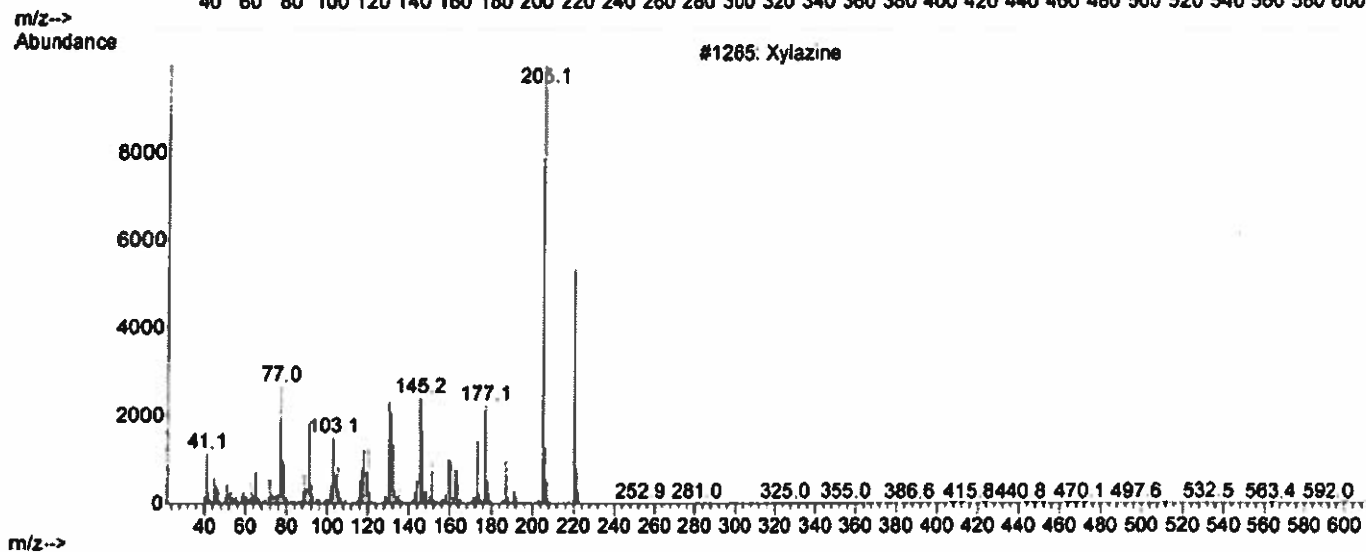
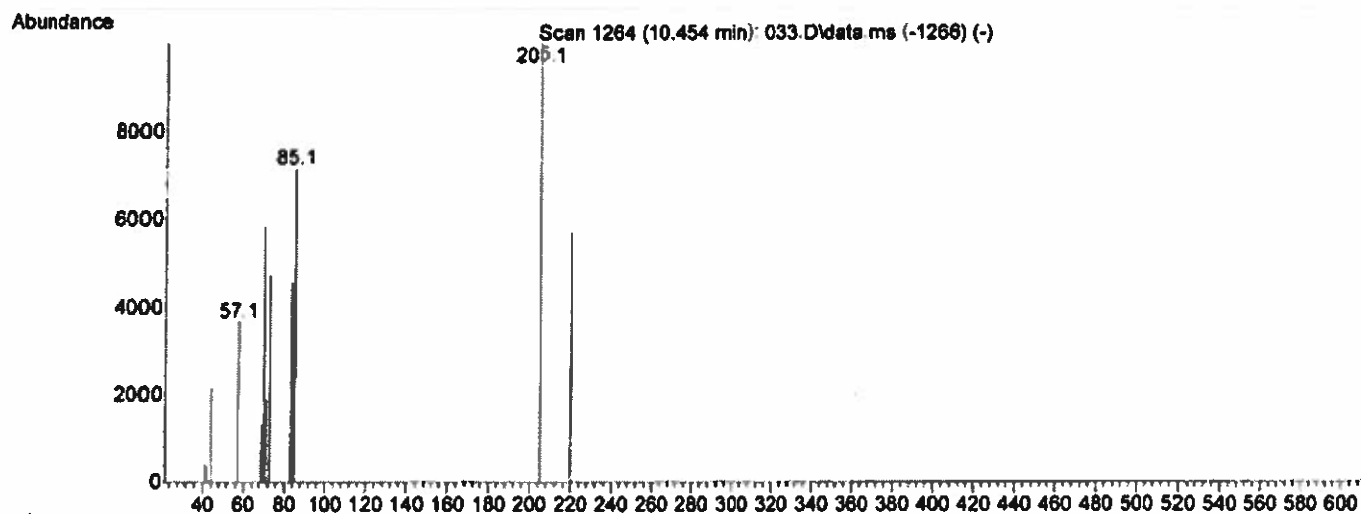




Library Searched : C:\MassHunter\Library\CaymanSpectralLibrary.L  
Quality : 2  
ID : Xylazine  
Sample Name: MATRIX 1 LOD - BLD  
File: C:\Users\TOX\Desktop\092821\033.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 05:36  
Vial: 64



Library Searched : C:\MassHunter\Library\CaymanSpectralLibrary.L  
Quality : 5  
ID : Xylazine  
Sample Name: MATRIX 1 LOD - BLD  
File: C:\Users\TOX\Desktop\092821\033.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 05:36  
Vial: 64



# Analysis Report

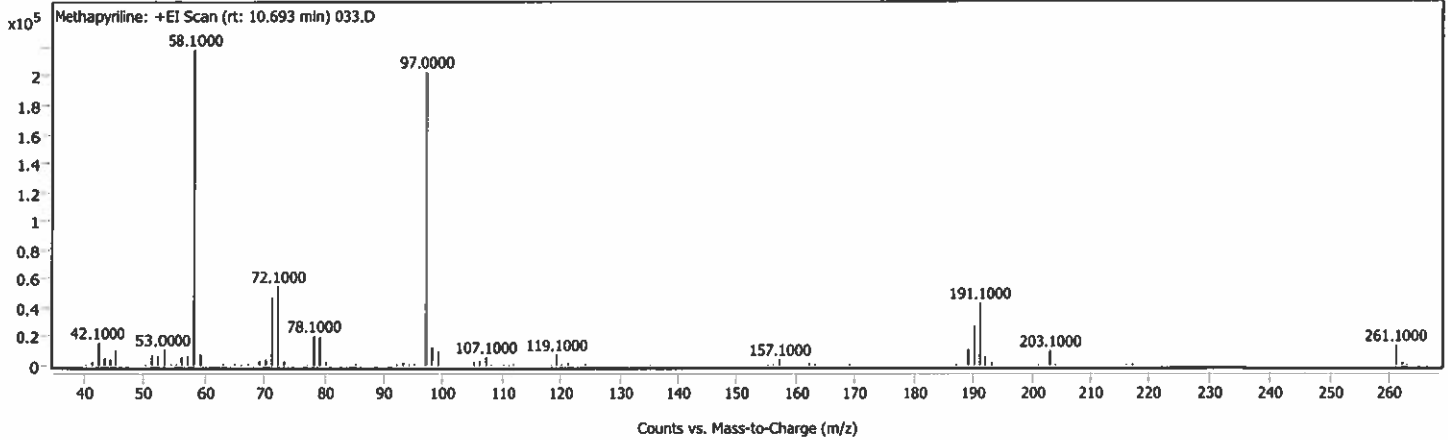
## Sample Spectra

+ Scan (rt: 10.693 min)

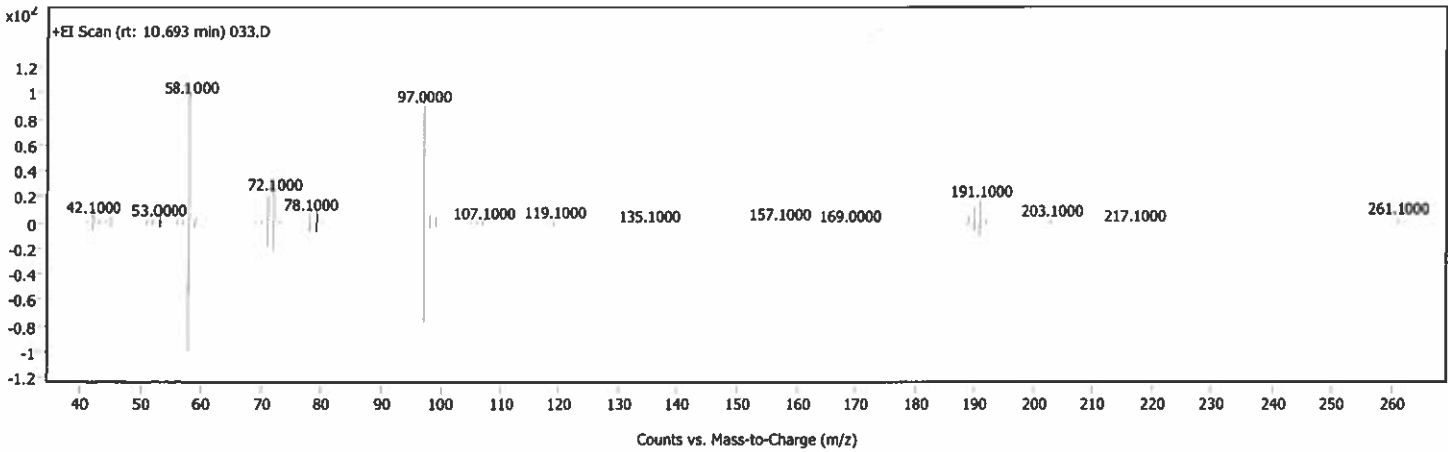
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.20	SWGDRUG 3.3.L	MATRIX 1 LOD - BLD	KM/QC DATA IN XLAZINE VALIDATION

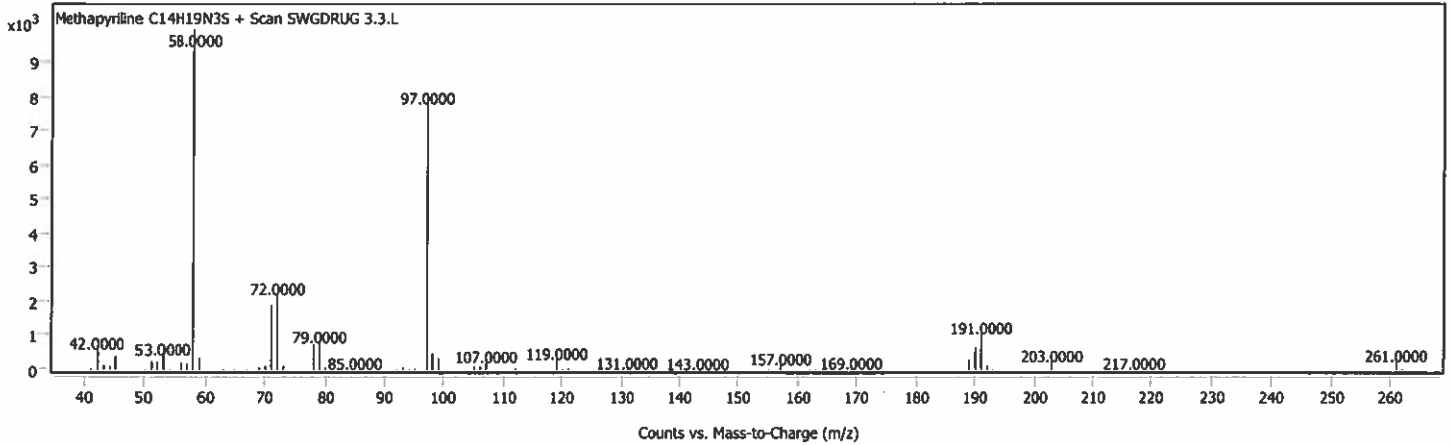
### Observed Spectrum



### Mirror Plot

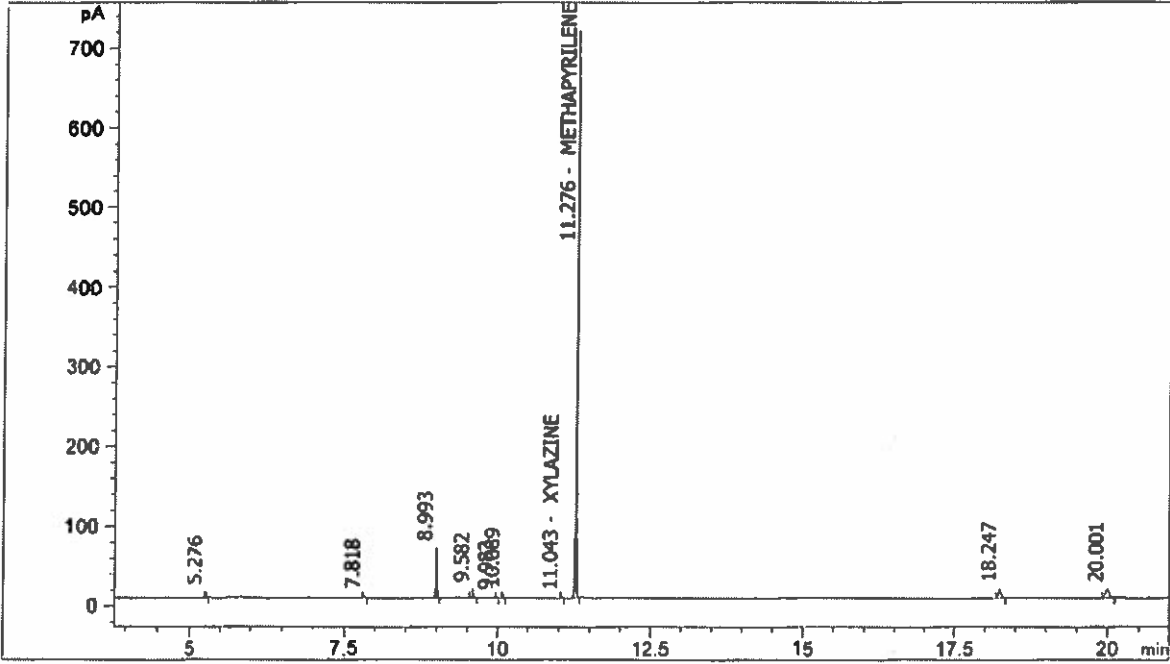


### Library Spectrum



Injection Date: 9/29/2021 5:09:36 AM Seq Line: 39  
 Sample Name: MATRIX 1 LOD - BLD Vial 40  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.043	11.046	2.828	2.134	0.000000	XYLAZINE RRT - 0.9793
11.276	11.278	827.325	711.058	1.000000	METHAPYRILENE

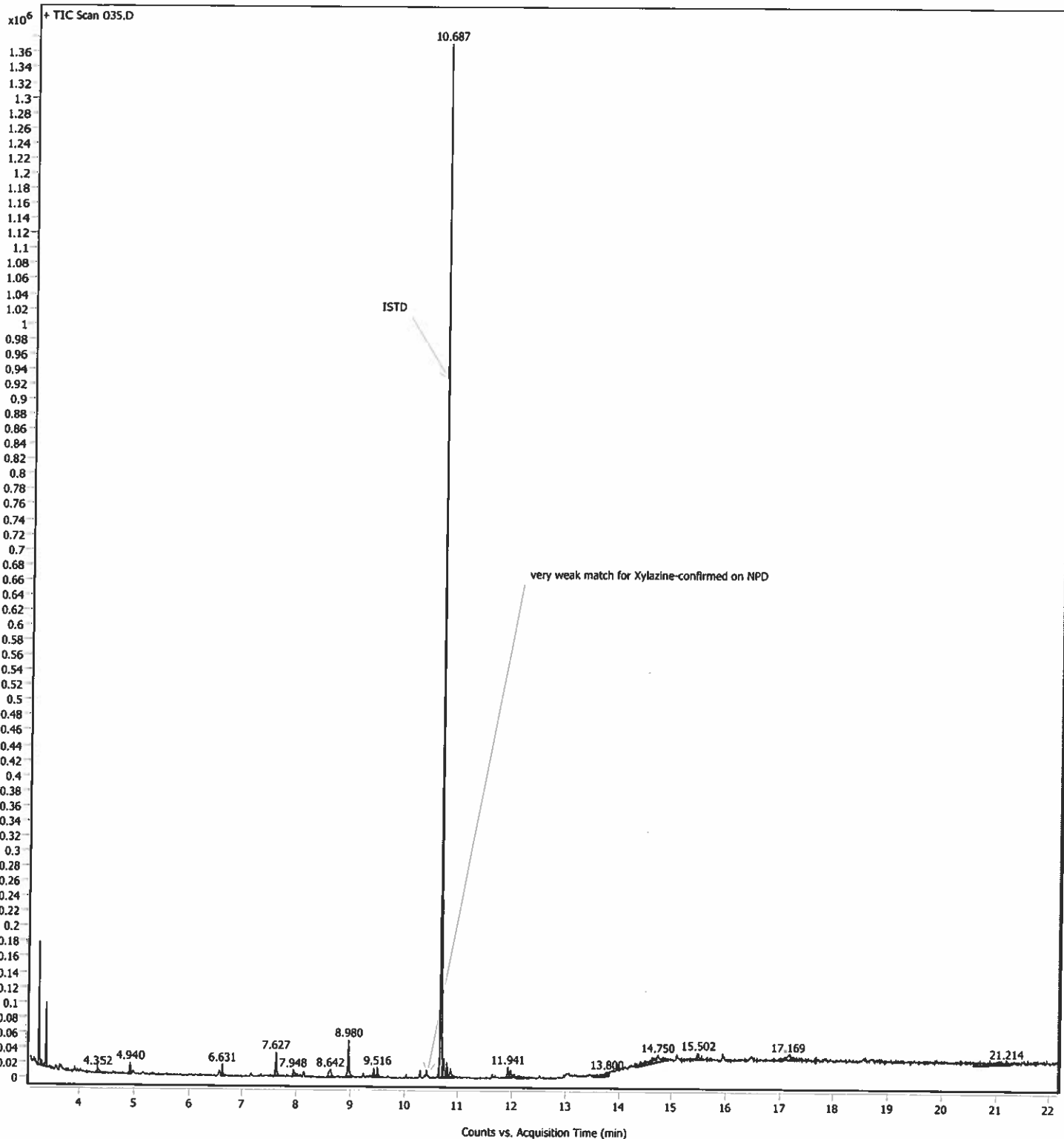


# Analysis Report

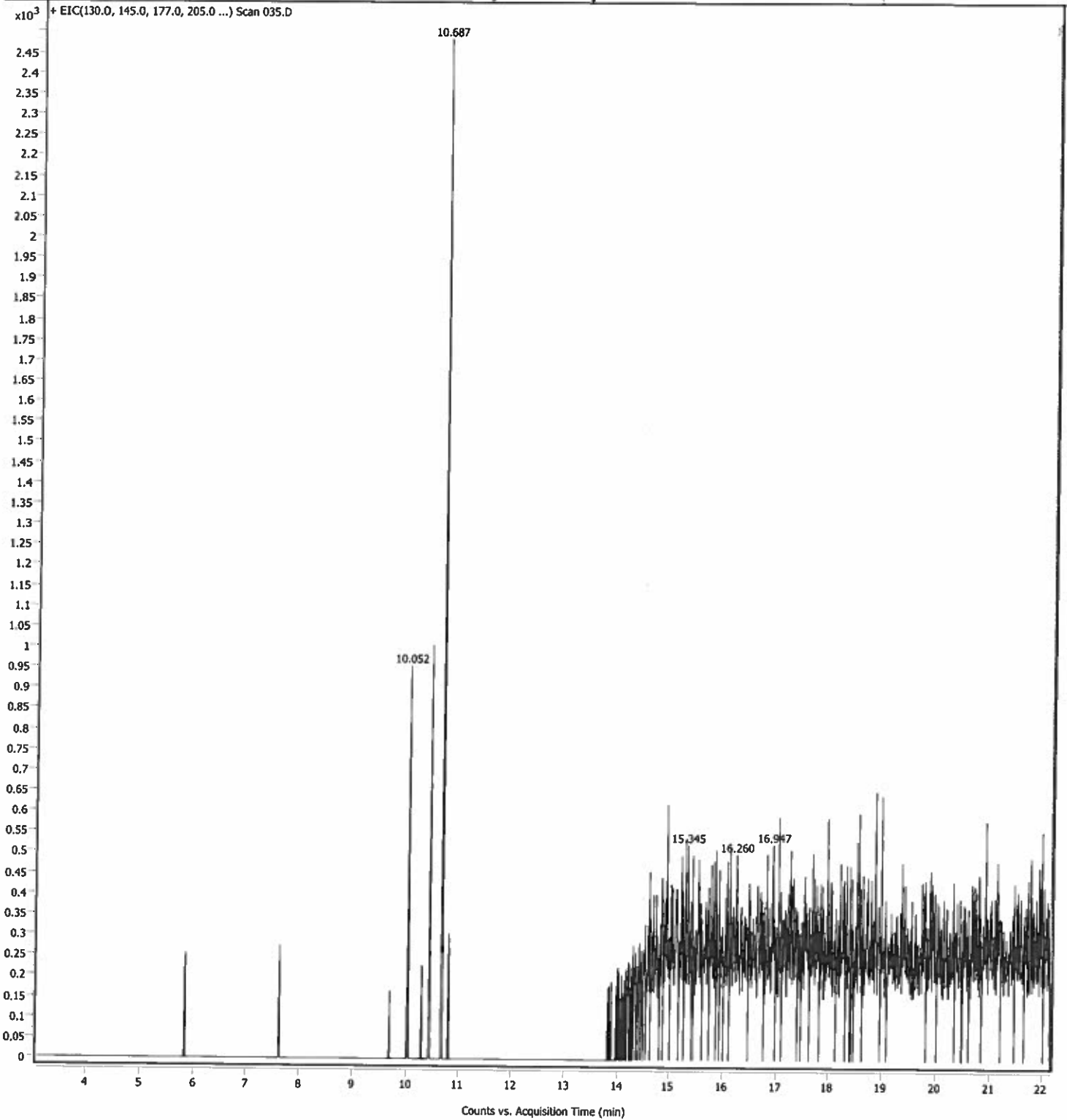
## Sample Information

Sample Name	MATRIX 6 LOD - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\035.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 6:28:06 AM (UTC-04:00)
Position	66	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report



# Analysis Report

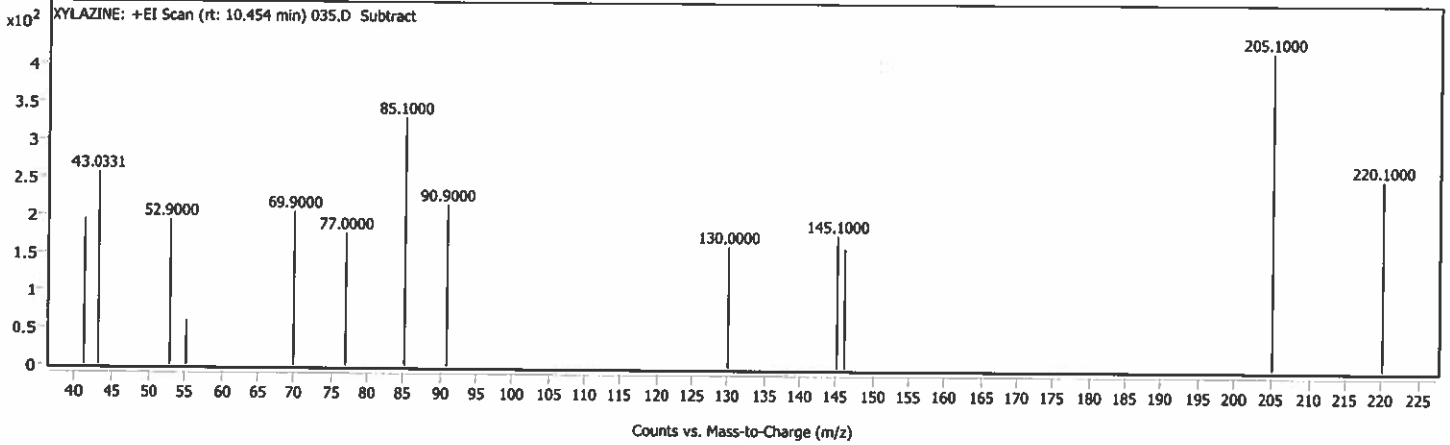
## Sample Spectra

+ Scan (rt: 10.454 min) Sub

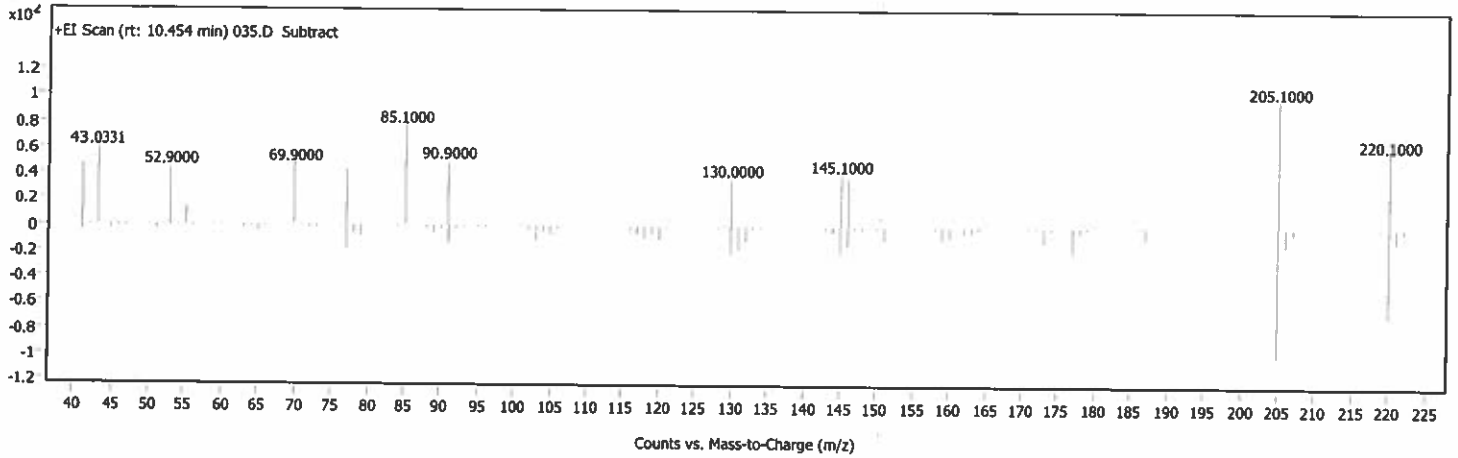
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	38.85	aafs.i	MATRIX 6 LOD - URN	KM/QC DATA IN XYLAZINE VALIDATION

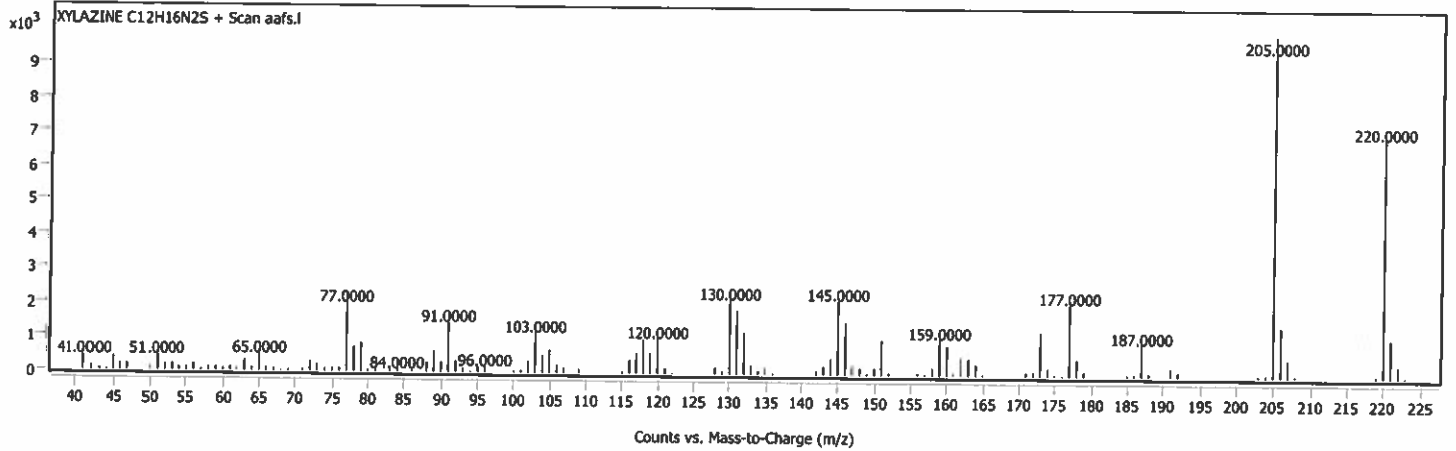
### Observed Spectrum



### Mirror Plot



### Library Spectrum



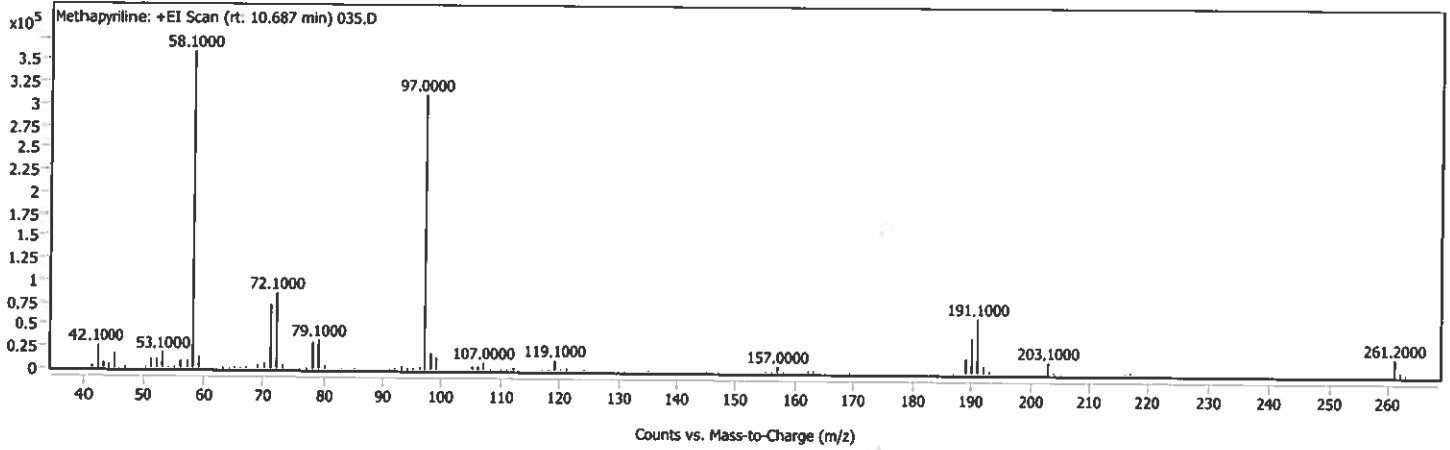
# Analysis Report

+ Scan (rt: 10.687 min)

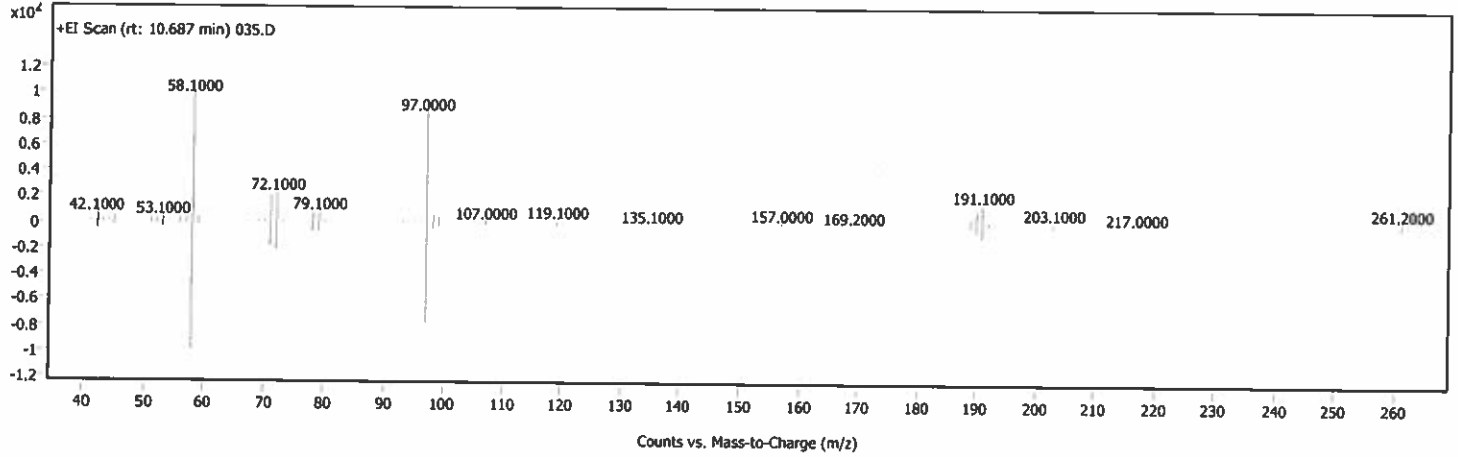
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.64	SWGDRUG 3.3.L	MATRIX 6 LOD - URN	KM/QC DATA IN XYLAZINE VALIDATION

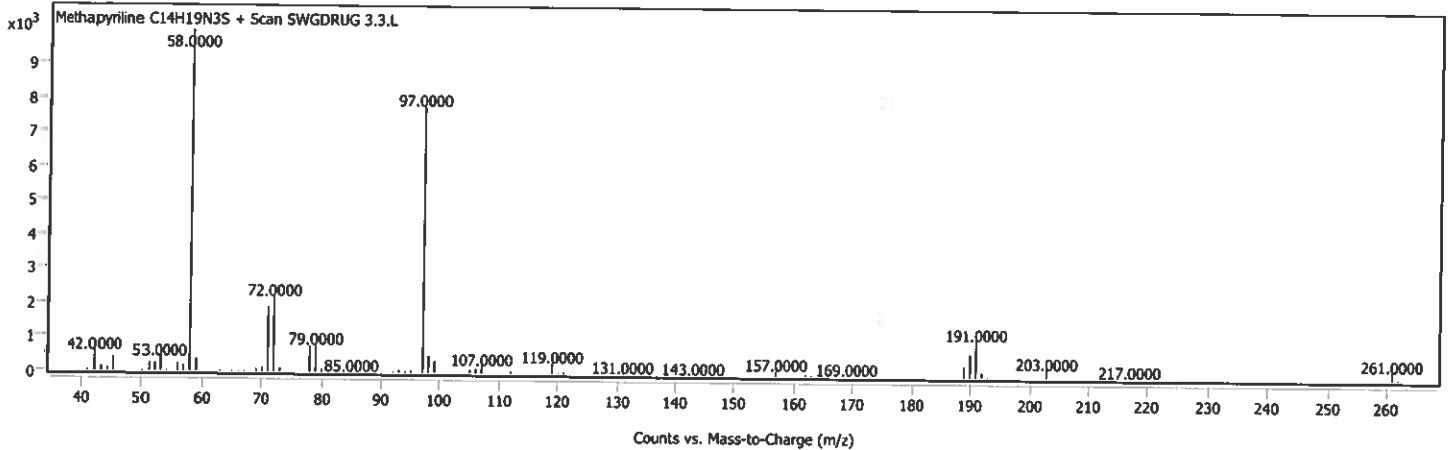
## Observed Spectrum



## Mirror Plot

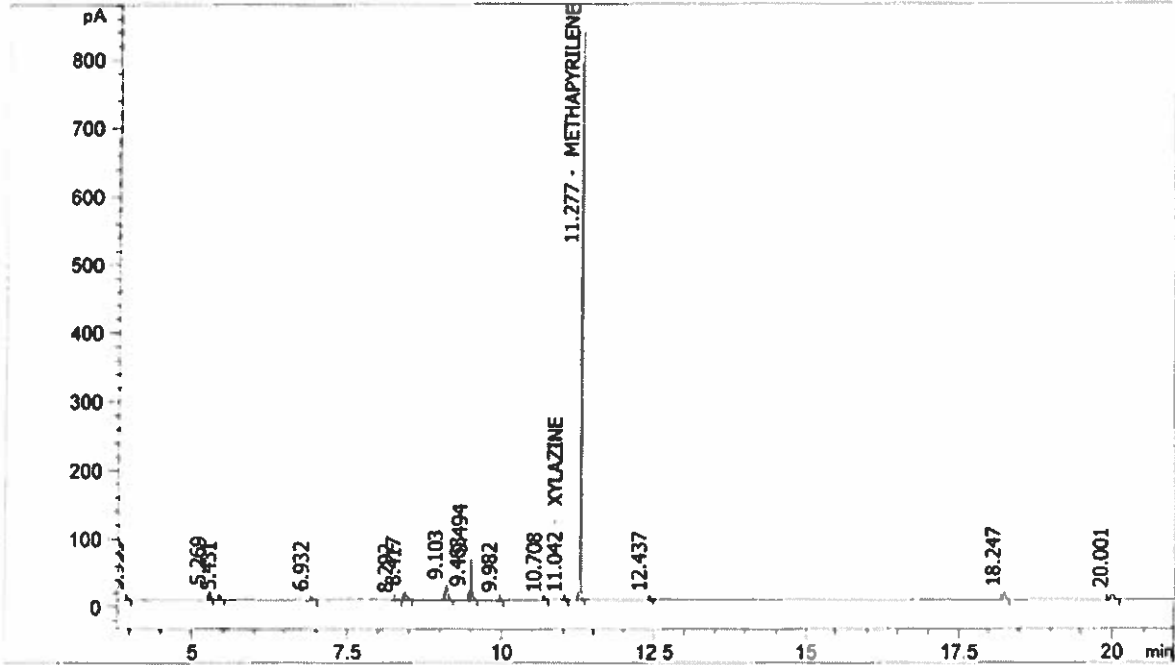


## Library Spectrum



=====  
Injection Date: 9/29/2021 5:58:17 AM Seq Line: 41  
Sample Name: MATRIX 6 LOD - URN Vial 42  
Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



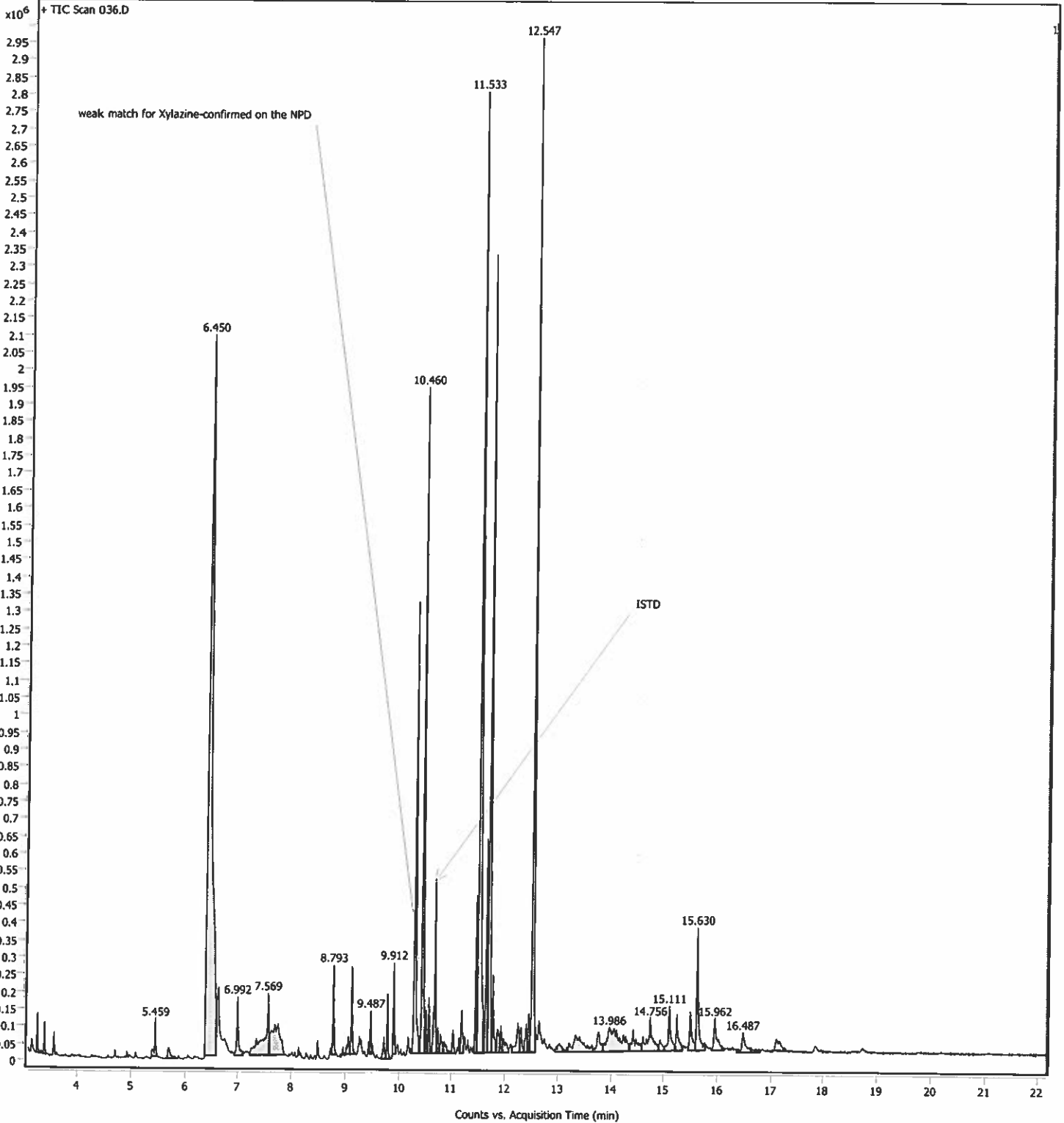
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.042	11.046	3.000	2.467	0.000000	XYLAZINE RRT - 0.9792
11.277	11.278	949.131	825.492	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 10 LOD - LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\036.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 6:53:53 AM (UTC-04:00)
Position	67	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

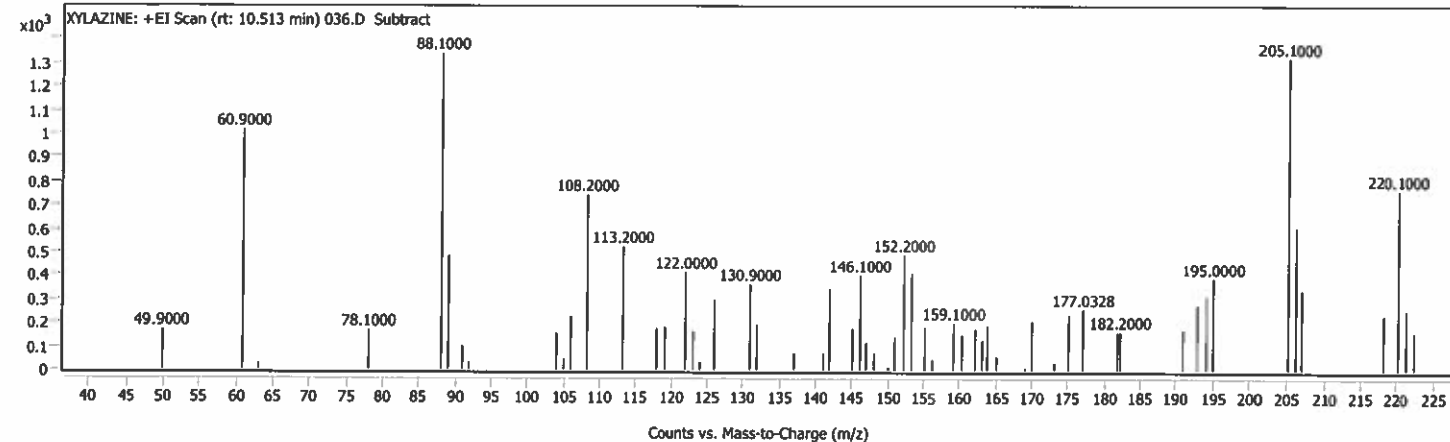
## Sample Spectra

+ Scan (rt: 10.687 min)

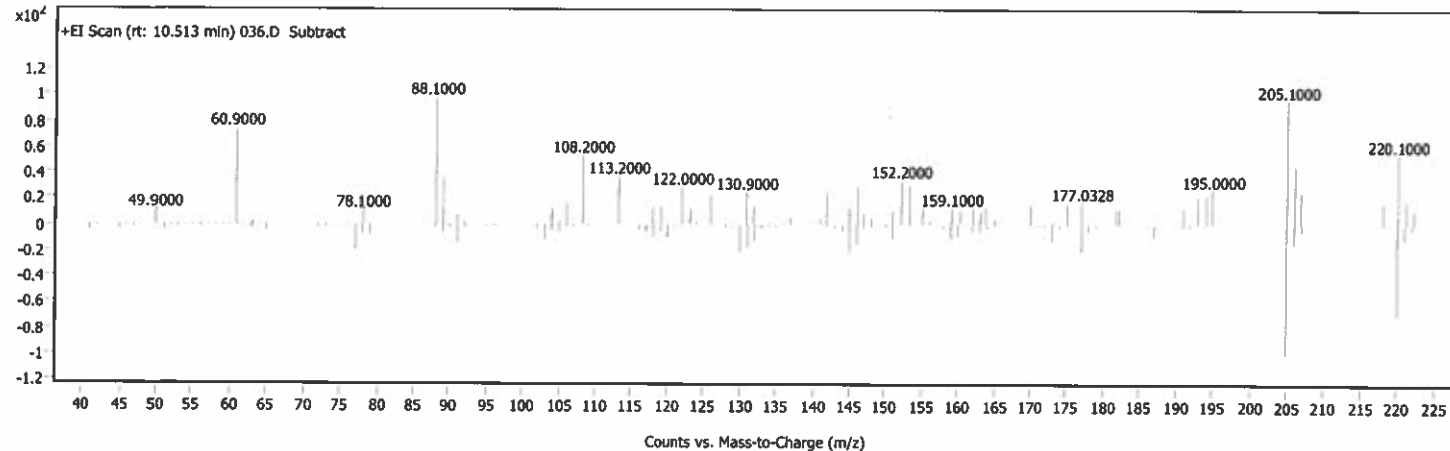
+ Scan (rt: 10.513 min) Sub **XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	50.29	aafs.l	MATRIX 10 LOD - LVR	KM/QC DATA IN XYLAZINE VALIDATION

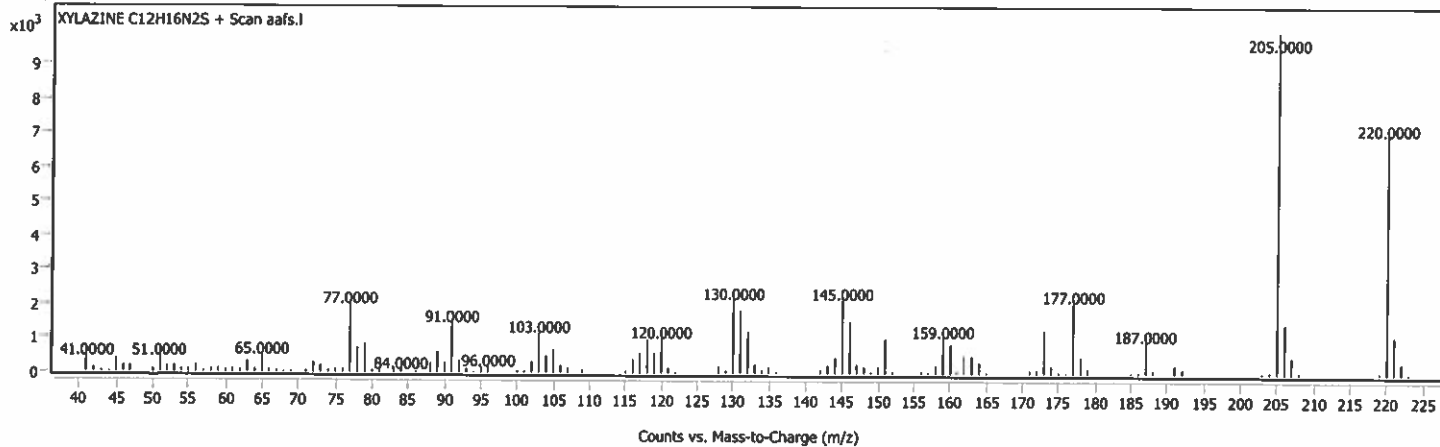
## Observed Spectrum



## Mirror Plot



## Library Spectrum



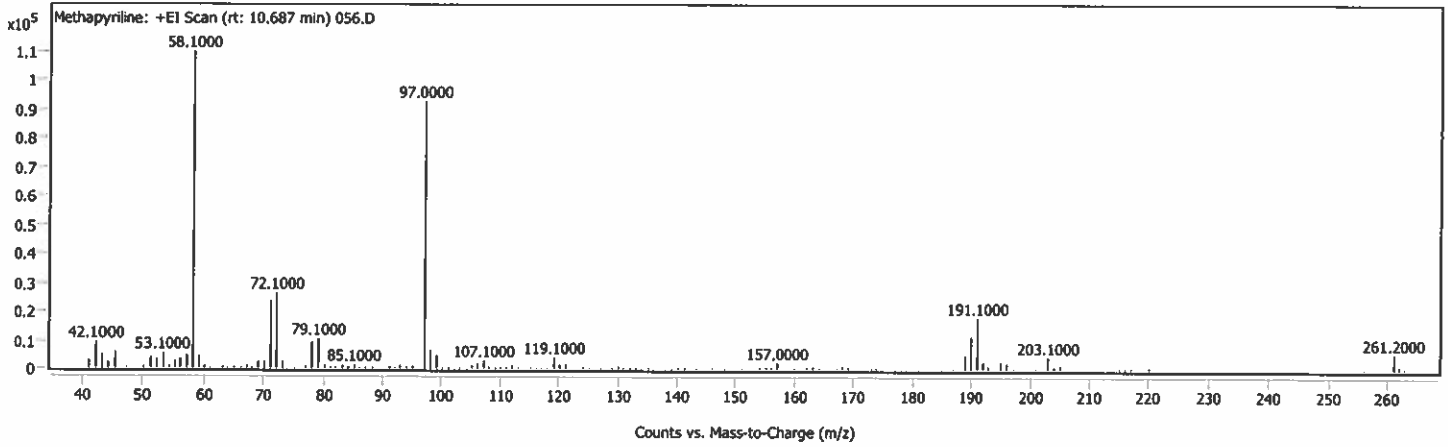
# Analysis Report

+ Scan (rt: 10.687 min)

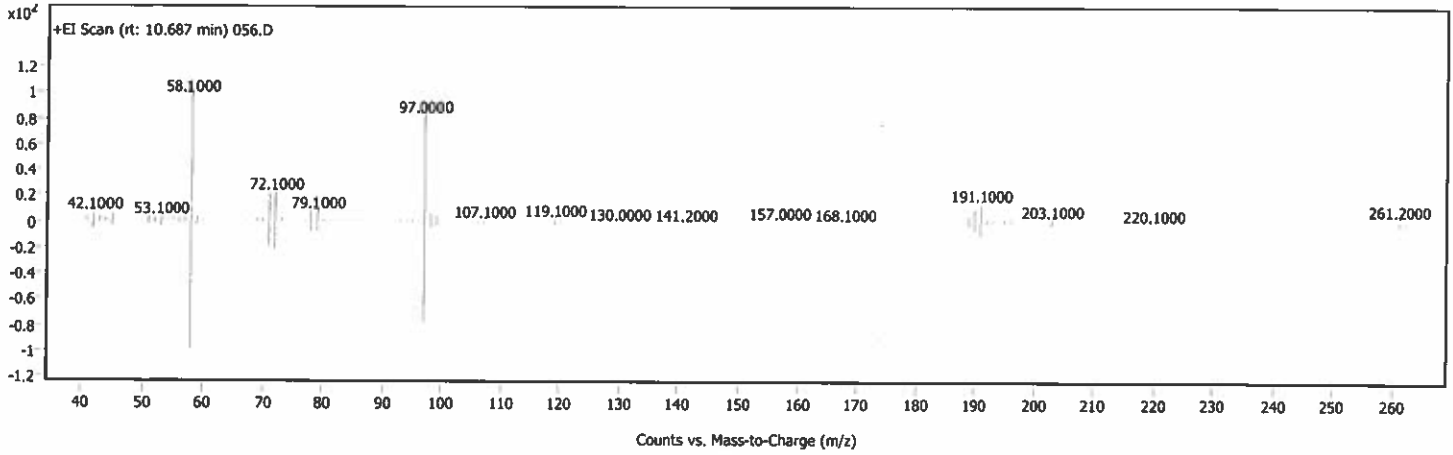
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	93.38	SWGDRUG 3.3.L	MATRIX 10 W/ISTD 2.0 mg/L- LVR	KM/QC DATA IN XYLAZINE VALIDATION

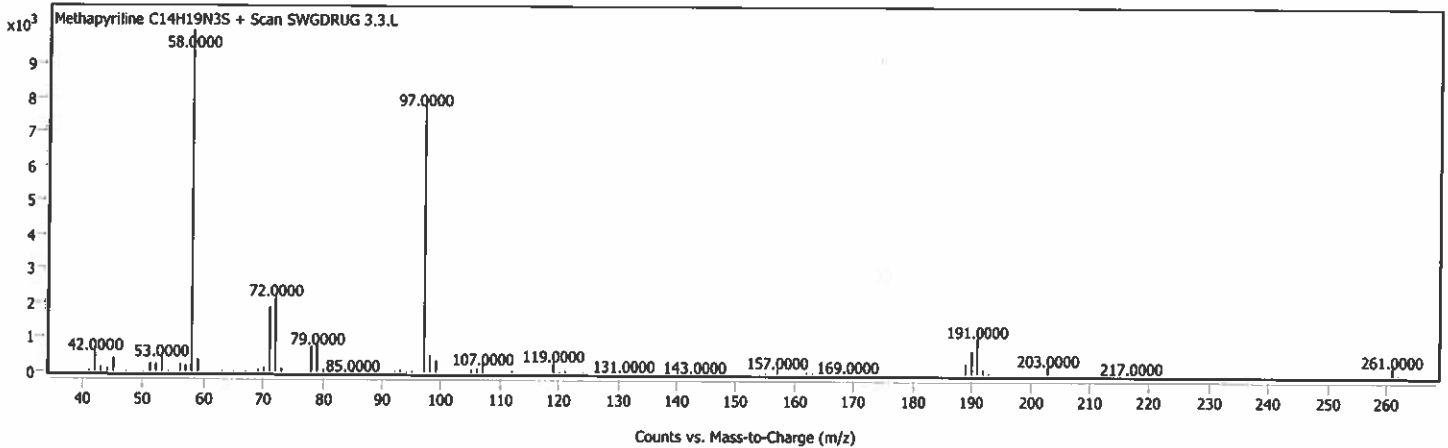
## Observed Spectrum



## Mirror Plot



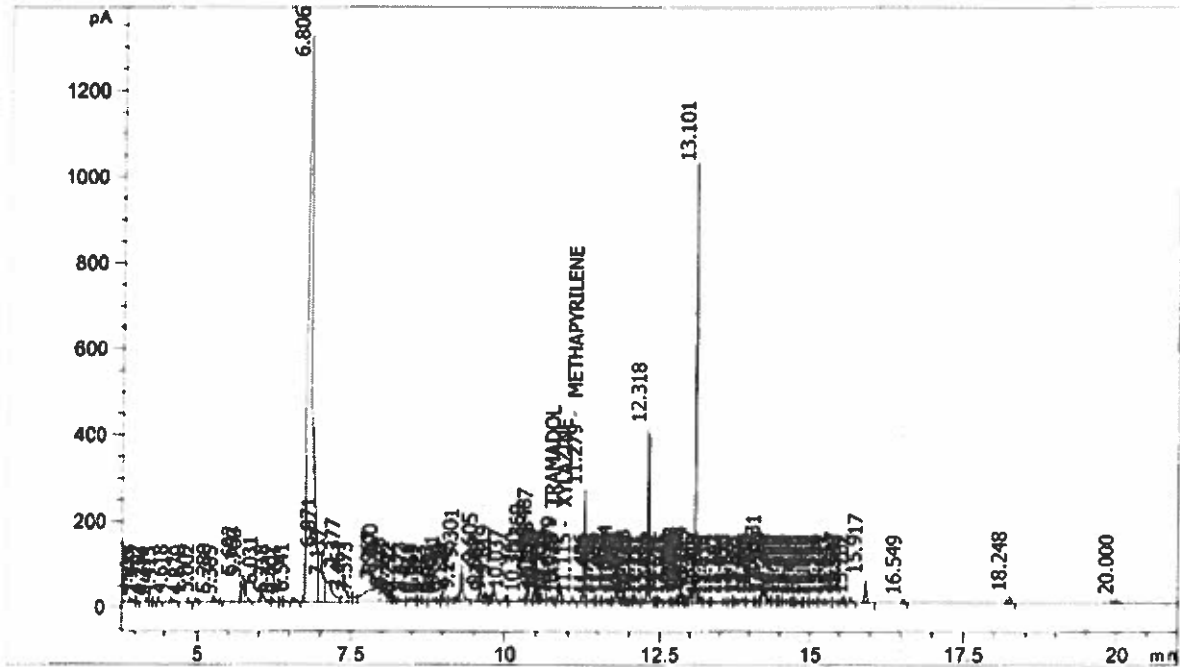
## Library Spectrum





Injection Date: 9/29/2021 6:22:37 AM Seq Line: 42  
 Sample Name: MATRIX 10 LOD - LVR Vial 43  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



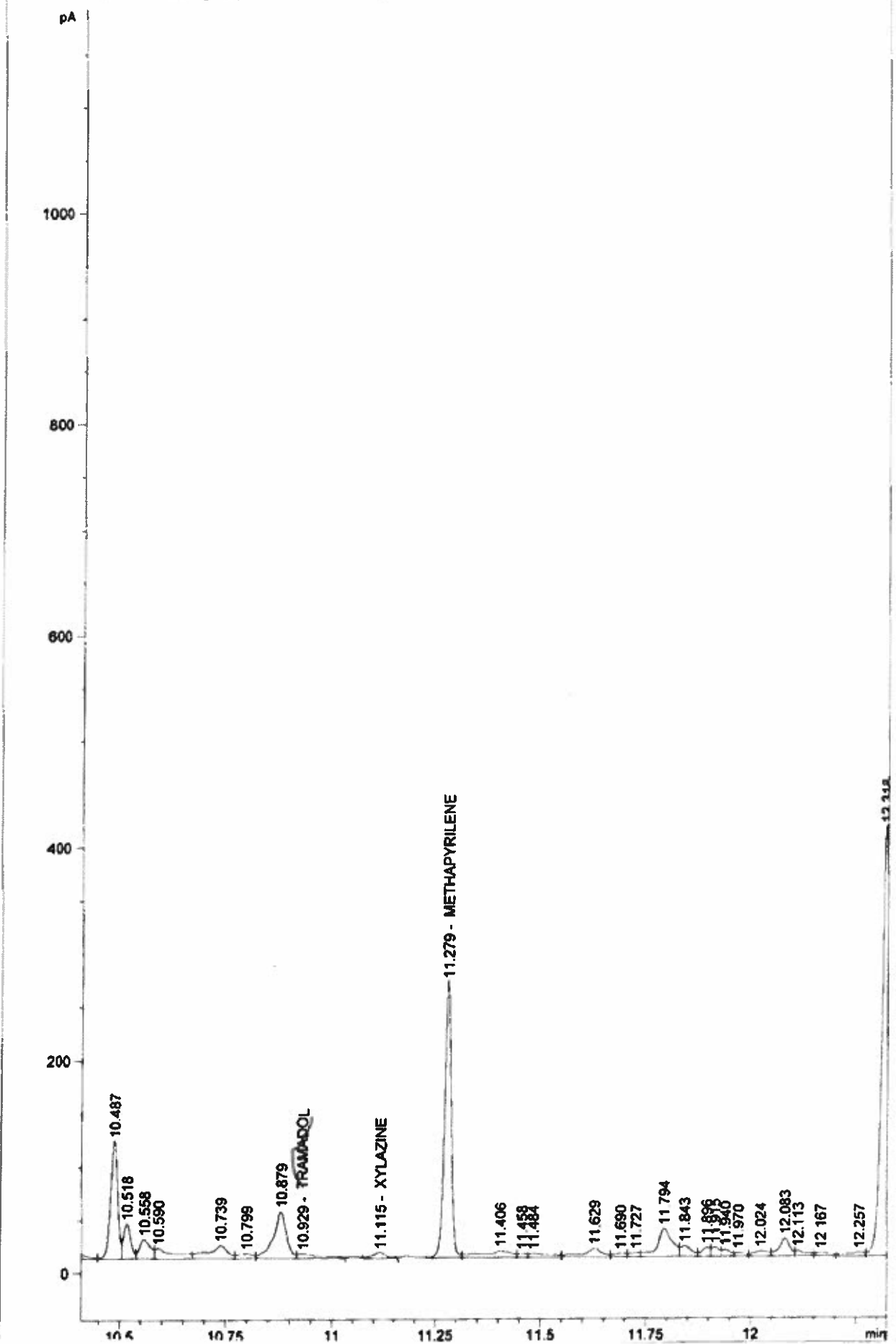
see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
10.929	11.033	15.959	4.366	0.00000	TRAMADOL ND
11.115	11.046	11.746	5.267	0.00000	XYLAZINE RRT - 0.9855
11.279	11.278	298.938	259.493	1.00000	METHAPYRILENE

LVR not used in avg of RRTs

Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000042.D)



**ASSAY:**

Base Screen

QC Data File: Xylazine Validation

**Analyst(s):**

K. Meinweiser

Sample / FS #	Item #	Sample Info
1 MATRIX 2 BASE DRUGS ONLY	BLD	2.0 mg/L - 400uL of 0.01 mg/mL of SWCTL
2 MATRIX 1 BASE DRUGS ONLY	BLD	
3 MATRIX 6 BASE DRUGS ONLY	URN	
4 MATRIX 10 BASE DRUGS ONLY	LVR	2.0 mg/L - 400uL of 0.01 mg/mL of Cal
5 MATRIX 2 BASE DRUGS ONLY	BLD	2.0 mg/L - 400uL of 0.01 mg/mL of IHCTL
6 MATRIX 1 BASE DRUGS ONLY	BLD	
7 MATRIX 6 BASE DRUGS ONLY	URN	
8 MATRIX 10 BASE DRUGS ONLY	LVR	
9 MATRIX 2 BASE DRUGS	BLD	2.0 mg/L - 400uL of 0.01 mg/mL of SWCTL
10 MATRIX 1 BASE DRUGS	BLD	plus 2.0 mg/L - 400uL of 0.01 mg/mL of Xylazine
11 MATRIX 6 BASE DRUGS	URN	
12 MATRIX 10 BASE DRUGS	LVR	2.0 mg/L - 400uL of 0.01 mg/mL of Cal
13 MATRIX 2 BASE DRUGS	BLD	2.0 mg/L - 400uL of 0.01 mg/mL of IHCTL
14 MATRIX 1 BASE DRUGS	BLD	plus 2.0 mg/L - 400uL of 0.01 mg/mL of Xylazine
15 MATRIX 6 BASE DRUGS	URN	
16 MATRIX 10 BASE DRUGS	LVR	*satisfies 2.8.1.1.5
17 MATRIX 2 0.10 mg/L	BLD	20 uL of 0.01 mg/mL std
18 MATRIX 1 0.10 mg/L	BLD	
19 MATRIX 6 0.10 mg/L	URN	
20 MATRIX BLANK		
21 MATRIX 2 0.50 mg/L	BLD	100 uL of 0.01 mg/mL std
22 MATRIX 1 0.50 mg/L	BLD	
23 MATRIX 6 0.50 mg/L	URN	
24 MATRIX BLANK		
25 MATRIX 2 1.0 mg/L	BLD	200 uL of 0.01 mg/mL std

Specimens removed from TX ADM storage, aliquots removed for analysis, specimens returned to TX ADM storage by, unless otherwise noted:

**Analyst:** Kimberly Meinweiser

**Date:** 09.28.21

**Time Removed:** N/A **Time Returned:** N/A

220-F156 Toxicology Barcode Batch Worksheet  
 Issued by Toxicology Program Manager  
 Issue Date: 02-July-2020

Sample / FS #	Item #	Sample Info
26 MATRIX 1 1.0 mg/L	BLD	
27 MATRIX 6 1.0 mg/L	URN	
28 MATRIX BLANK		
29 MATRIX 2 2.0 mg/L	BLD	400 uL of 0.01 mg/mL std
30 MATRIX 1 2.0 mg/L	BLD	
31 MATRIX 6 2.0 mg/L	URN	
32 MATRIX BLANK		*satisfies 2.8.1.1.7
33		
34		
35		
36		
37		
38		
39		
40		
41		
42		
43		
44		
45		
46		
47		
48		
49		
50		

**Extraction:** LL / **SPE** **SPE lot #:** 024170-XD

**Control Charts Updated (Initials/Date):** N/A

**Reviewer / Date:** \_\_\_\_\_



method: C:\CHEM32\1\METHODS\ALKALI.M  
Modified on: 11/2/2022 at 4:25:43 PM

=====  
Agilent 7890B  
=====

GC

Oven  
Equilibration Time 0.5 min  
Max Temperature 325 °C  
Slow Fan Disabled  
Oven Program On  
Oven Program 90 °C for 2 min  
Oven#1 then 15 °C/min to 245 °C for 0 min  
Oven#2 then 30 °C/min to 300 °C for 7 min  
Run Time 21.167 min  
Cryo Off

ALS

Front Injector  
Syringe Size 10 µL  
Syringe A Syringe has not been selected.  
Injection Volume 1 µL  
Solvent A Washes (PreInj) 2  
Solvent A Washes (PostInj) 4  
Solvent A Volume 8 µL  
Solvent B Washes (PreInj) 2  
Solvent B Washes (PostInj) 4  
Solvent B Volume 8 µL  
Sample Washes 0  
Sample Wash Volume 8 µL  
Sample Pumps 2  
Dwell Time (PreInj) 0 min  
Dwell Time (PostInj) 0 min  
Solvent Wash Draw Speed 300 µL/min  
Solvent Wash Dispense Speed 3000 µL/min  
Sample Wash Draw Speed 300 µL/min  
Sample Wash Dispense Speed 3000 µL/min  
Injection Dispense Speed 6000 µL/min  
Viscosity Delay 1 sec  
Sample Depth Disabled  
Injection Type Standard  
L1 Airgap 0.2 µL

Tray

Barcode heater Disabled  
Barcode mixer Disabled

Sample Overlap  
Mode

Sample overlap is not enabled

ALS Errors

Pause for user interaction

Front SS Inlet He  
Mode

Pulsed Splitless

Heater On 260 °C  
Pressure On 18.881 psi  
Total Flow On 44.5 mL/min  
Septum Purge Flow On 3 mL/min  
Gas Saver On 20 After 2 min mL/min  
Injection Pulse Pressure 40 psi Until 0.3 min

method: C:\CHEM32\1\METHODS\ALKALI.M  
Modified on: 11/2/2022 at 4:25:43 PM

Purge Flow to Split Vent  
Liner

40 mL/min at 0.5 min  
Agilent 5190-2793: 900 µL (Splitless, single tape)

Back SS Inlet He

\*\*\*Excluded from Affecting GC's Readiness State\*\*

Mode  
Heater  
Pressure  
Total Flow  
Septum Purge Flow  
Gas Saver  
Purge Flow to Split Vent  
Liner

Splitless  
On 250 °C  
On 9.801 psi  
On 41.7 mL/min  
Off  
Off  
40 mL/min at 0.5 min  
Agilent 5183-4647; Lot P02-C6823: 870 µL (Split,

Column

Column #1  
21080

Restek RTX-1; SN 1652462

0 °C-350 °C (350 °C); 30 m x 250 µm x 0.25 µm

Column lock

Unlocked

In

Front SS Inlet He

Out

Front Detector NPD

(Initial)

90 °C

Pressure

18.881 psi

Flow

1.5 mL/min

Average Velocity

36.082 cm/sec

Holdup Time

1.3857 min

Flow Program

On

Flow Program

1.5 mL/min for 1 min

Flow

then 0.2 mL/min per min to 2.2 mL/min for 0 m

Run Time

21.167 min

Column #2

10223

RTX-5

0 °C-325 °C (325 °C); 30 m x 320 µm x 0.25 µm

Column lock

Unlocked

In

Back SS Inlet He

Out

Back Detector FID

(Initial)

90 °C

Pressure

9.801 psi

Flow

1.7 mL/min

Average Velocity

31.621 cm/sec

Holdup Time

1.5812 min

Flow Program

On

Flow Program

1.7 mL/min for 1 min

Flow

then 0.2 mL/min per min to 3 mL/min for 0 min

Run Time

21.167 min

Front Detector NPD

Heater

On 310 °C

H2 Flow

On 3 mL/min

Air Flow

On 60 mL/min

Makeup Flow (Combined)

On 10 mL/min

Carrier Gas Flow Correction

Included in Makeup Flow

Bead

On

Signal 1 Type

NPD

Maximum Bead Voltage

4.095 V

Bios Bead

No

method: C:\CHEM32\1\METHODS\ALKALI.M  
Modified on: 11/2/2022 at 4:25:43 PM

Dry Bead Yes  
Auto Adjust Bead No

Back Detector FID

\*\*\*Excluded from Affecting GC's Readiness State\*\*  
Heater On 280 °C  
H2 Flow Off  
Air Flow Off  
Makeup Flow On 25 mL/min  
Carrier Gas Flow Correction Does not affect Makeup or Fuel Flow  
Flame Off  
Electrometer On

Signals

Signal #1: Front Signal  
Description Front Signal  
Details  
Save On  
Data Rate 50 Hz

Signal #2: Back Signal  
Description Back Signal  
Details  
Save Off  
Data Rate 5 Hz

Signal #3: Test Plot  
Description Test Plot  
Details  
Save Off  
Data Rate 50 Hz

Signal #4: Test Plot  
Description Test Plot  
Details  
Save Off  
Data Rate 50 Hz

Run Time Events

Run Time Events  
Run Time Events Time 0 min  
Run Time Events Event Detector H2 Fuel Flow  
Run Time Events Position 0  
Run Time Events Setpoint Off

Run Time Events Time 2 min  
Run Time Events Event Detector H2 Fuel Flow  
Run Time Events Position 0  
Run Time Events Setpoint On

Sequence Name: C:\MassHunter\GCMS\1\sequence\alkalival.sequence.xml

Comment: Base

Operator: KM/QC DATA IN XYLAZINE VALIDATION KM

Data Path: C:\MassHunter\GCMS\1\data\BASES\092821\

Instrument Control Pre-Seq Cmd:

Data Analysis Pre-Seq Cmd:

Instrument Control Post-Seq Cmd:

Data Analysis Post-Seq Cmd:

Method Sections To Run

(X) Full Method

( ) Reprocessing Only

Sequence Barcode Options

( ) On Mismatch, Inject Anyway

( ) On Mismatch, Don't Inject

(X) Barcode Disabled

vials loaded by AJ

Line	Type	ALS	File	Method	Sample Name/Misc Info
1)	Sample	26	ETOAC	ALKALI	ETOAC
2)	Sample	27	001	ALKALI	MATRIX 1 0.10 mg/L - BLD
3)	Sample	28	002	ALKALI	MATRIX 2 0.10 mg/L - BLD
4)	Sample	29	003	ALKALI	MATRIX 6 0.10 mg/L - URN
5)	Sample	30	004	ALKALI	MATRIX BLANK after 0.1 mg/L
6)	Sample	31	005	ALKALI	MATRIX 1 0.50 mg/L - BLD
7)	Sample	32	006	ALKALI	MATRIX 2 0.50 mg/L - BLD
8)	Sample	33	007	ALKALI	MATRIX 6 0.50 mg/L - URN
9)	Sample	34	008	ALKALI	MATRIX BLANK after 0.5 mg/L
10)	Sample	35	009	ALKALI	MATRIX 1 1.0 mg/L - BLD
11)	Sample	36	010	ALKALI	MATRIX 2 1.0 mg/L - BLD
12)	Sample	37	011	ALKALI	MATRIX 6 1.0 mg/L - URN
13)	Sample	38	012	ALKALI	MATRIX BLANK after 1.0 mg/L
14)	Sample	39	013	ALKALI	MATRIX 1 2.0 mg/L - BLD
15)	Sample	40	014	ALKALI	MATRIX 2 2.0 mg/L - BLD
16)	Sample	41	015	ALKALI	MATRIX 6 2.0 mg/L - URN
17)	Sample	42	016	ALKALI	MATRIX BLANK after 2.0 mg/L
18)	Sample	43	0147	ALKALI	MATRIX 1 W/ SW CTL - BLD
19)	Sample	44	018	ALKALI	MATRIX 2 W/ SW CTL - BLD
20)	Sample	45	019	ALKALI	MATRIX 6 W/ SW CTL - URN
21)	Sample	46	020	ALKALI	MATRIX 10 W/ 0.01 mg/L CAL -
22)	Sample	47	SOL	ALKALI	Solvent Blank
23)	Sample	48	021	ALKALI	MATRIX 1 W/ IH CTL - BLD
24)	Sample	49	022	ALKALI	MATRIX 2 W/ IH CTL - BLD
25)	Sample	50	023	ALKALI	MATRIX 6 W/ IH CTL - URN
26)	Sample	51	024	ALKALI	MATRIX 10 W/ IH CTL - LVR
27)	Sample	52	SOL1	ALKALI	Solvent Blank 1
28)	Sample	53	025	ALKALI	MATRIX 1 W/ SW CTL + XYLAZINE
29)	Sample	54	026	ALKALI	MATRIX 2 W/ SW CTL + XYLAZINE
30)	Sample	55	027	ALKALI	MATRIX 6 W/ SW CTL + XYLAZINE
31)	Sample	56	028	ALKALI	MATRIX 10 W/ 0.01 mg/L CAL +
32)	Sample	57	SOL2	ALKALI	Solvent Blank 2
33)	Sample	58	029	ALKALI	MATRIX 1 W/ IH CTL + XYLAZINE
34)	Sample	59	030	ALKALI	MATRIX 2 W/ IH CTL + XYLAZINE
35)	Sample	60	031	ALKALI	MATRIX 6 W/ IH CTL + XYLAZINE
36)	Sample	61	032	ALKALI	MATRIX 10 W/ IH CTL + XYLAZINE KM 9/29/21
37)	Sample	62	SOL3	ALKALI	Solvent Blank 3
38)	Sample	63	SOL4	ALKALI	Solvent Blank 4
39)	Sample	64	033	ALKALI	MATRIX 1 LOD - BLD
40)	Sample	65	034	ALKALI	MATRIX 2 LOD - BLD
41)	Sample	66	035	ALKALI	MATRIX 6 LOD - URN

vial check 9/29/21 KM



## Single Quadrupole Acquisition Method - MS Parameters Report

<b>Method file</b>	C:\MassHunter\GCMS\1\methods\ALKALI.m
<b>Tune file</b>	ATUNE.U
<b>Ion source</b>	EI
<b>Source temperature (°C)</b>	230
<b>Quad temperature (°C)</b>	150
<b>Fixed Electron energy (eV)</b>	70.3
<b>Acquisition Type</b>	Scan
<b>Stop time (min)</b>	22.17
<b>Solvent delay (min)</b>	3.00
<b>Trace Ion Detection</b>	False
<b>Gain Factor</b>	1
<b>EM Saver</b>	False
<b>EM Saver Limit</b>	N/A

### Scan Time Segments

Time	Start Mass	End Mass	Threshold	Scan Speed
3.00	40	550	150	1,562 [N=2]

### Timed Events

Time	Type of Event	Parameter
------	---------------	-----------

### Real-Time Plots

Type of Plot	Label	Low Mass	High Mass
Total Ion	N/A	N/A	N/A
Spectrum	N/A	N/A	N/A
Extracted Ion	Scan 1-1	40	550

### Self-Cleaning Ion Source Parameters

Mode	No Cleaning
------	-------------

C:\MassHunter\GCMS\1\methods\ALKALI.m  
 Tue Sep 28 11:40:23 2021

Control Information

-----  
 Sample Inlet : GC  
 Injection Source : GC ALS  
 Injection Location: Front  
 Mass Spectrometer : Enabled

No Sample Prep method has been assigned to this method.

GC  
 GC Summary  
 Run Time 22.167 min  
 Post Run Time 0 min

Oven  
 Temperature  
 Setpoint On  
 (Initial) 90 °C  
 Hold Time 2 min  
 Post Run 0 °C  
 Program  
 #1 Rate 15 °C/min  
 #1 Value 245 °C  
 #1 Hold Time 0 min  
 #2 Rate 30 °C/min  
 #2 Value 300 °C  
 #2 Hold Time 8 min

Equilibration Time 0.5 min  
 Max Temperature 325 °C  
 Maximum Temperature Override Disabled  
 Slow Fan Disabled

ALS  
 Front Injector  
 Syringe Size 10 µL  
 Injection Volume 1 µL  
 Injection Repetitions 1  
 Injection Delay 0 sec  
 Solvent A Washes (PreInj) 4  
 Solvent A Washes (PostInj) 4  
 Solvent A Volume 8 µL  
 Solvent B Washes (PreInj) 4  
 Solvent B Washes (PostInj) 4  
 Solvent B Volume 8 µL  
 Sample Washes 0  
 Sample Wash Volume 8 µL  
 Sample Pumps 2  
 Dwell Time (PreInj) 0 min  
 Dwell Time (PostInj) 0 min  
 Solvent Wash Draw Speed 300 µL/min  
 Solvent Wash Dispense Speed 3000 µL/min  
 Sample Wash Draw Speed 300 µL/min  
 Sample Wash Dispense Speed 3000 µL/min

Injection Dispense Speed	6000 µL/min
Viscosity Delay	0 sec
Sample Depth	Disabled
Tower Fan	On
Solvent Wash Mode	A-A2,B-B2
Sample Overlap Mode	Sample overlap is not enabled
ALS Errors	Pause for user interaction
Front SS Inlet He Mode	Pulsed Splitless
Heater	On 260 °C
Pressure	On 15.177 psi
Total Flow	On 46.869 mL/min
Septum Purge Flow	On 3 mL/min
Septum Purge Flow Mode	Standard
Gas Saver	On 20 After 2 min mL/min
Injection Pulse Pressure	40 psi Until 0.3 min
Purge Flow to Split Vent	42.4 mL/min at 0.5 min
Liner	A Liner has not been selected.
Thermal Aux 2 (MSD Transfer Line) Temperature	
Setpoint (Initial)	On 280 °C
Column	
Column #1	
Flow	
Setpoint (Initial)	On 1.4687 mL/min
Hold Time	1 min
Post Run	1.2 mL/min
Program	
#1 Rate	0.2 mL/min per min
#1 Value	3 mL/min
#1 Hold Time	0 min
Column Information	Restek 10123
Rtx-1 SN 1652464	
Temperature Range	-60 °C-350 °C (350 °C)
Dimensions	30 m x 250 µm x 0.25 µm
In	Front SS Inlet He
Out	MSD
(Initial)	90 °C
Pressure	15.177 psi
Flow	1.4687 mL/min
Average Velocity	45 cm/sec
Holdup Time	1.1111 min
Control Mode	Ramped Flow
Column Outlet Pressure	0 psi
Front Detector NPD	
Makeup	***Excluded from Affecting GC's Readiness State***
Heater	He
H2 Flow	Off
Air Flow	Off
Makeup Flow	Off

Carrier Gas Flow Correction	Constant Makeup and Fuel Flow
Bead	Off
Maximum Bead Voltage	4.095 V
Blos Bead	No
Dry Bead	Yes
Auto Adjust Bead	No

Signals	
Signal #1:	
Description	None
Signal #2:	
Description	None
Signal #3:	
Description	None
Signal #4:	
Description	None

TUNE PARAMETERS for SN: US71236345

-----

Trace Ion Detection is OFF.

34.610	:	EMISSION	
70.347	:	ENERGY	
28.232	:	REPELLER	
90.157	:	IONFOCUS	
32.000	:	ENTRANCE_LENS	
1894.118	:	EMVOLTS	
			1894.118 : Actual EMV
			0.81 : GAIN FACTOR
2284.000	:	AMUGAIN	
119.688	:	AMUOFFSET	
1.000	:	FILAMENT	
0.000	:	DCPOLARITY	
18.573	:	ENTLENSOFFSET	
-530.000	:	MASSGAIN	
-36.000	:	MASSOFFSET	

END OF TUNE PARAMETERS

-----

END OF INSTRUMENT CONTROL PARAMETERS

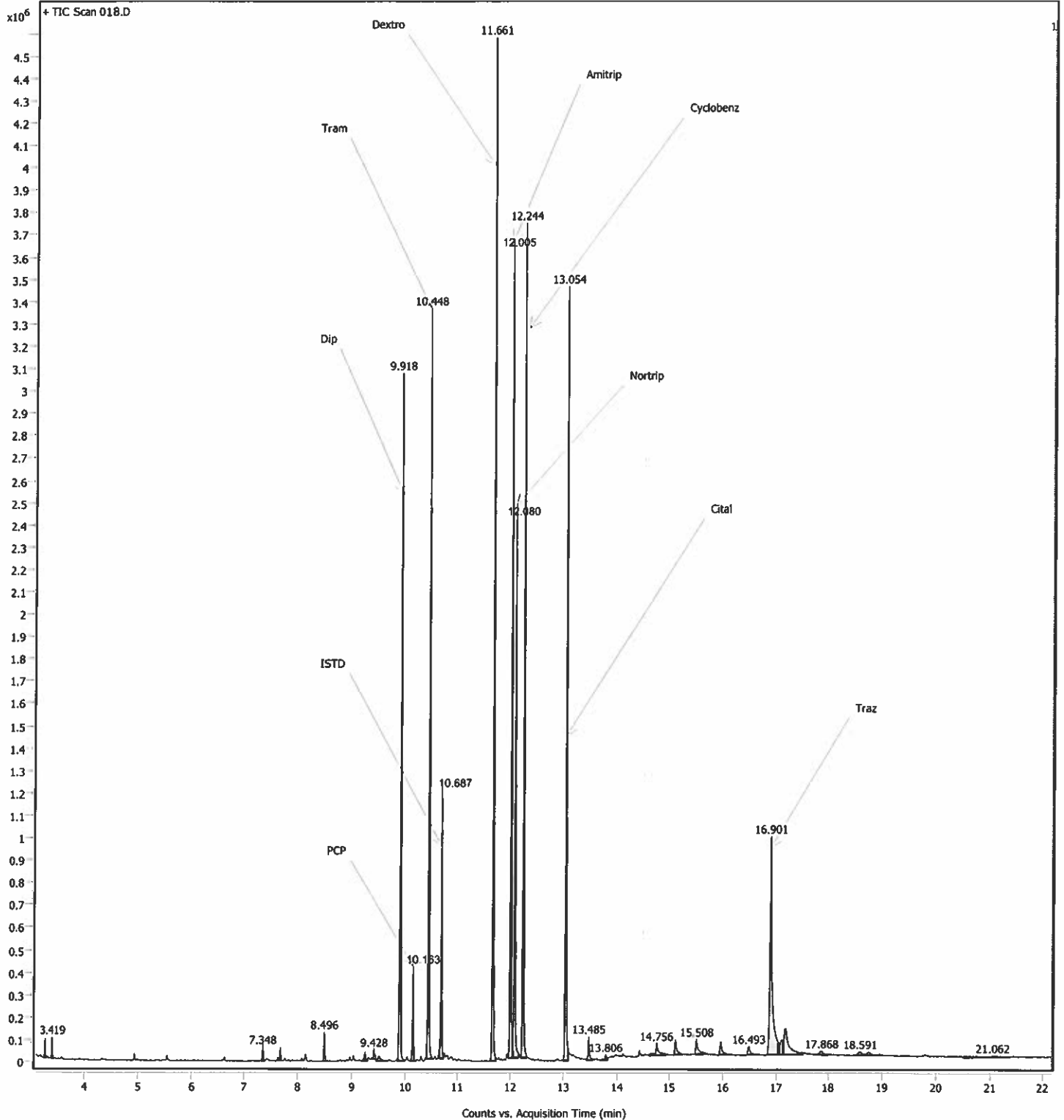
-----

# Analysis Report

## Sample Information

Sample Name	MATRIX 2 W/ SW CTL - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\018.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 8:59:46 PM (UTC-04:00)
Position	44	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

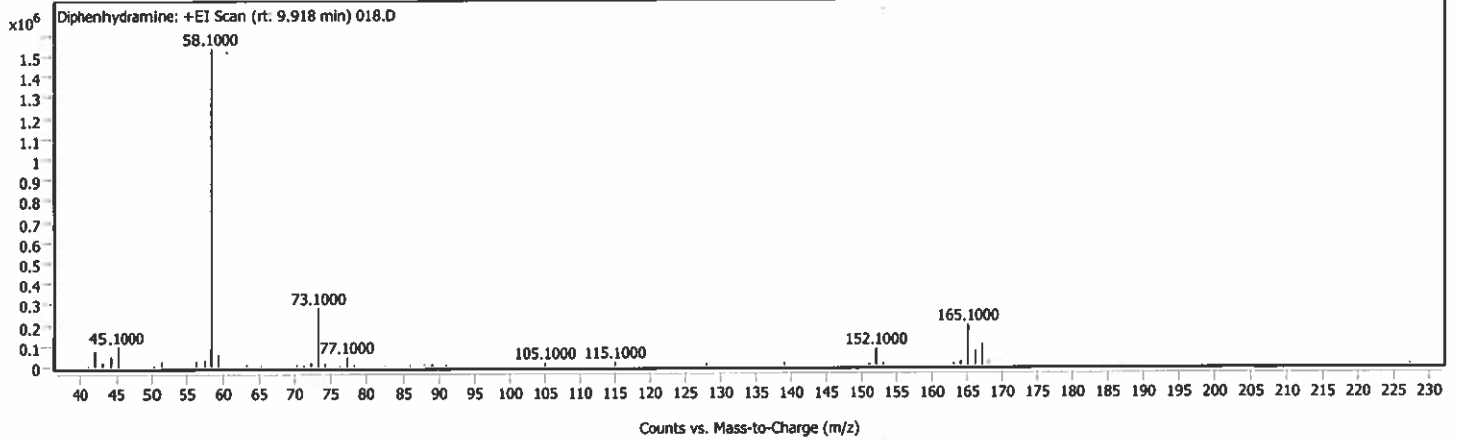
## Sample Spectra

+ Scan (rt: 9.918 min)

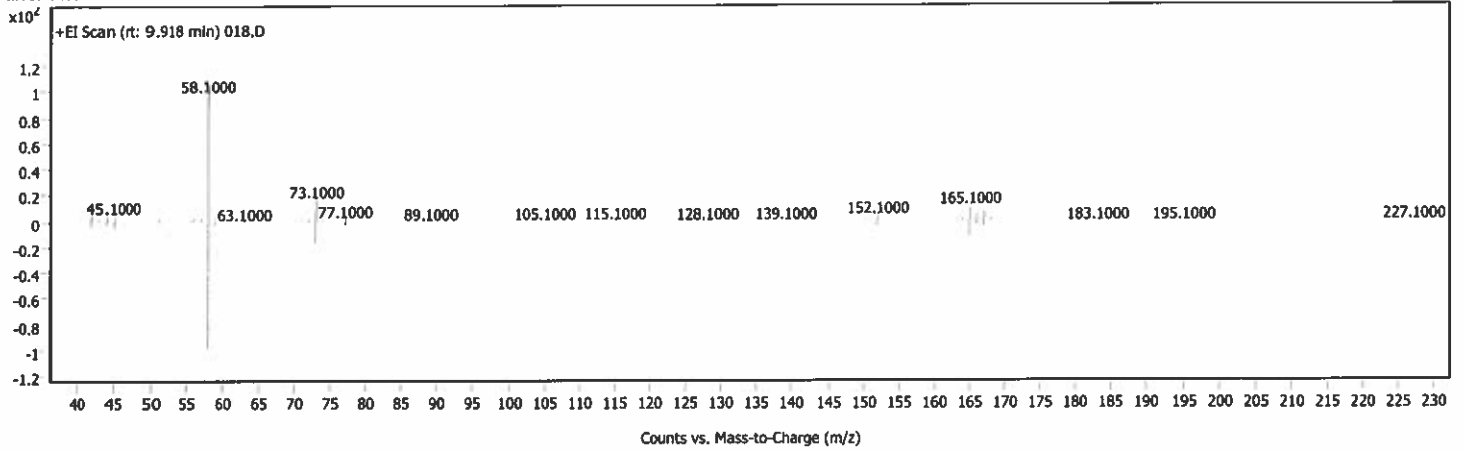
Diphenhydramine; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Diphenhydramine	C17H21NO	98.29	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

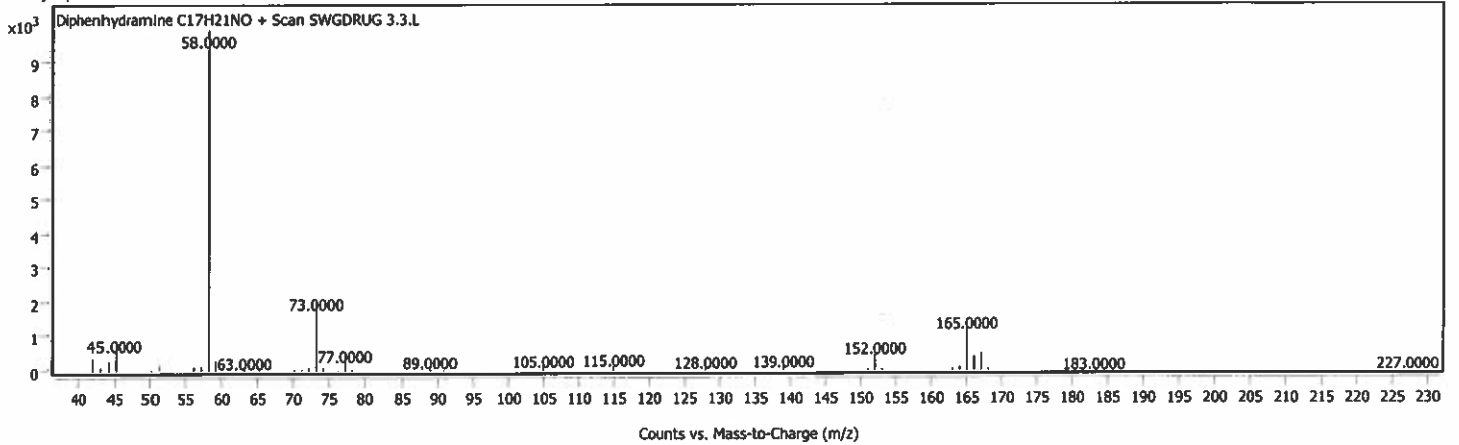
## Observed Spectrum



## Mirror Plot



## Library Spectrum



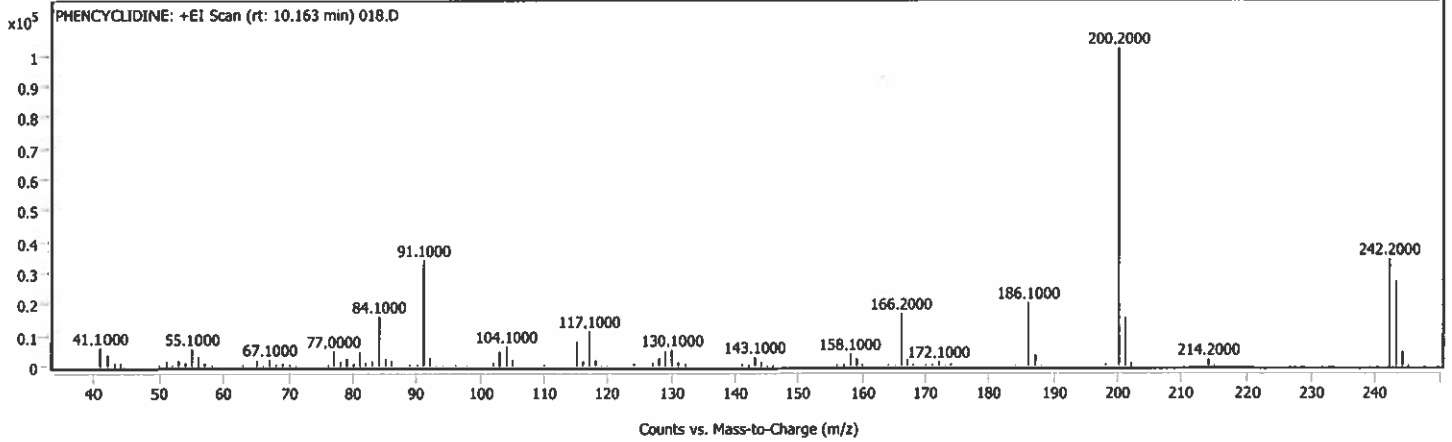
# Analysis Report

+ Scan (rt: 10.163 min)

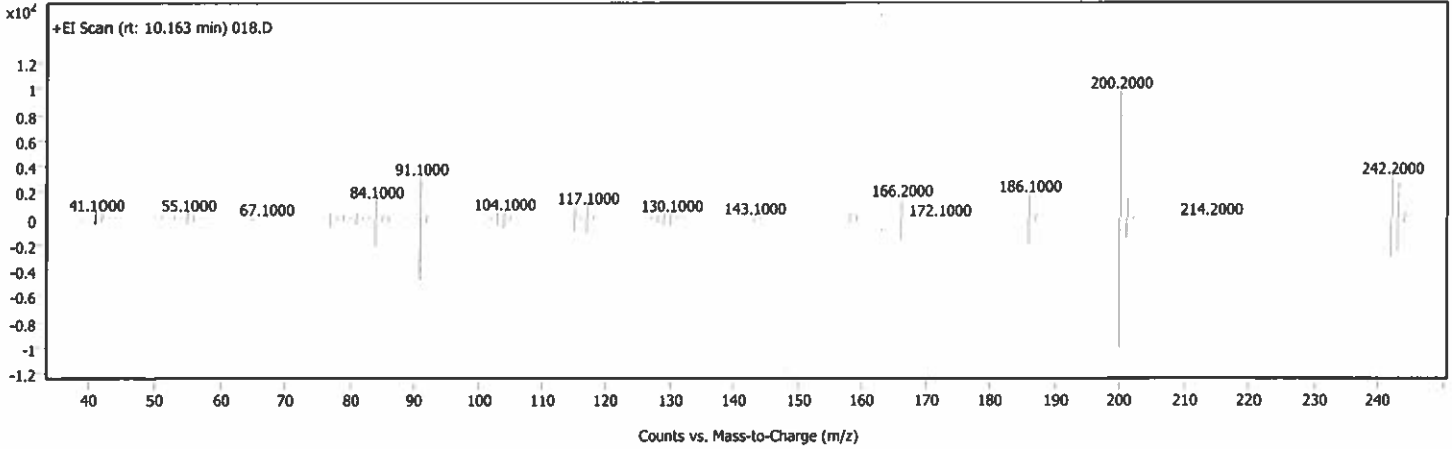
PHENCYCLIDINE; C17H25N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
PHENCYCLIDINE	C17H25N	98.84	aafs.I	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

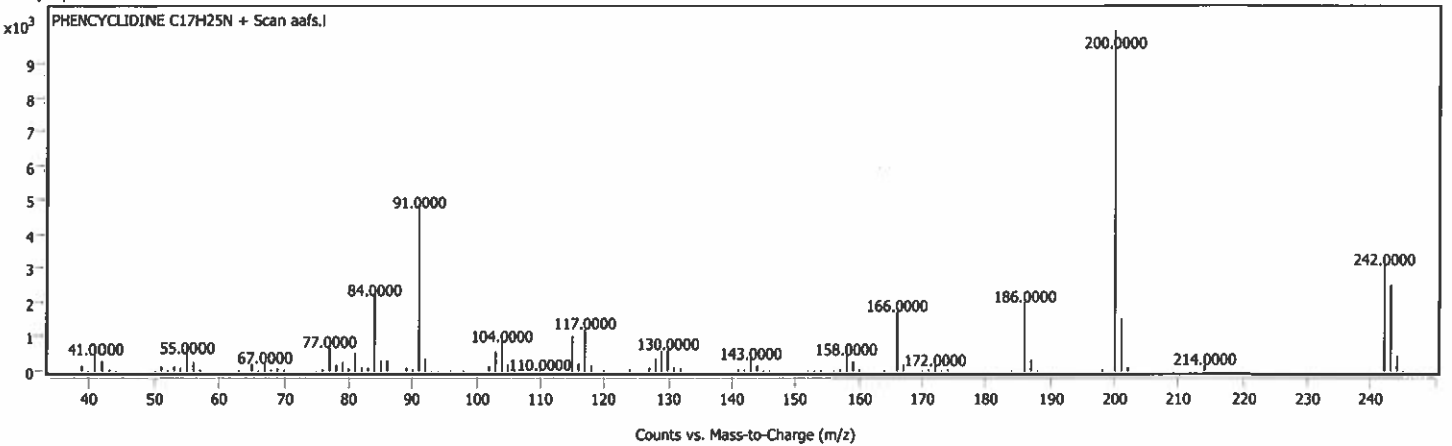
## Observed Spectrum



## Mirror Plot



## Library Spectrum



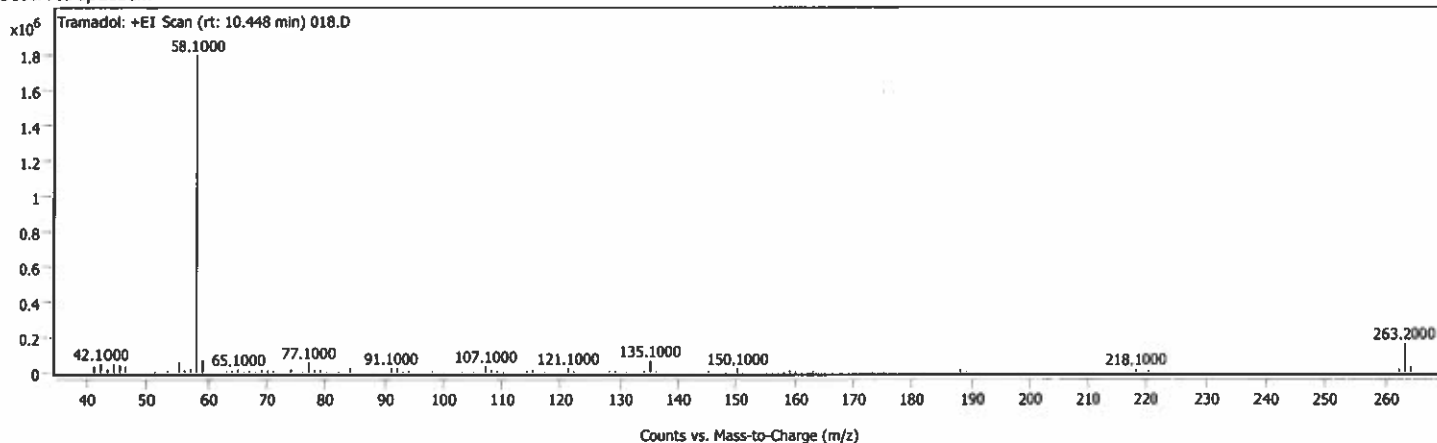
# Analysis Report

+ Scan (rt: 10.448 min)

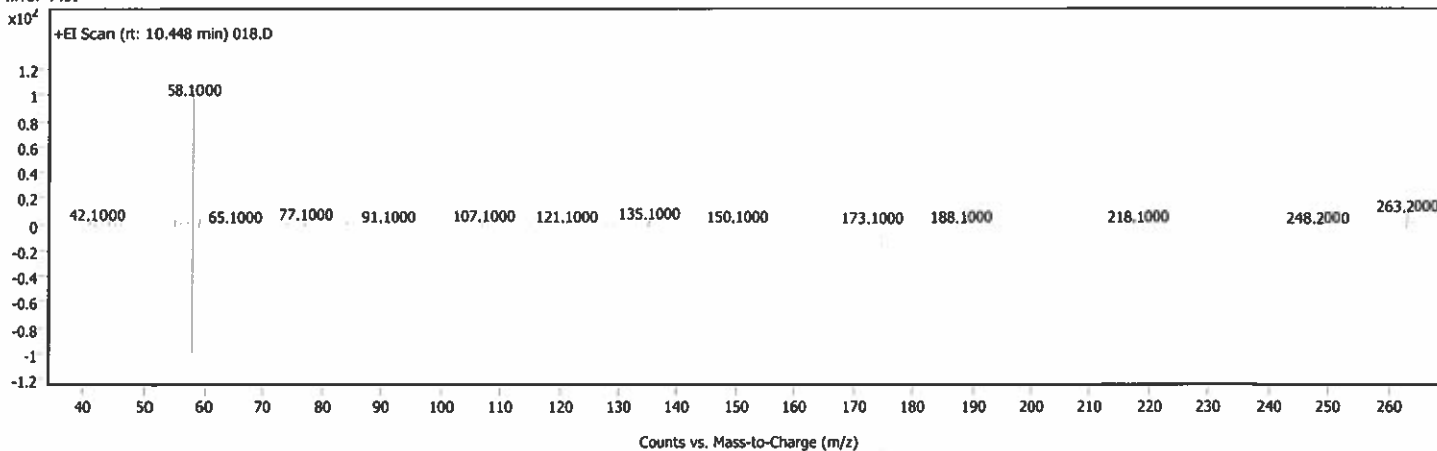
Tramadol; C16H25NO2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Tramadol	C16H25NO2	96.16	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

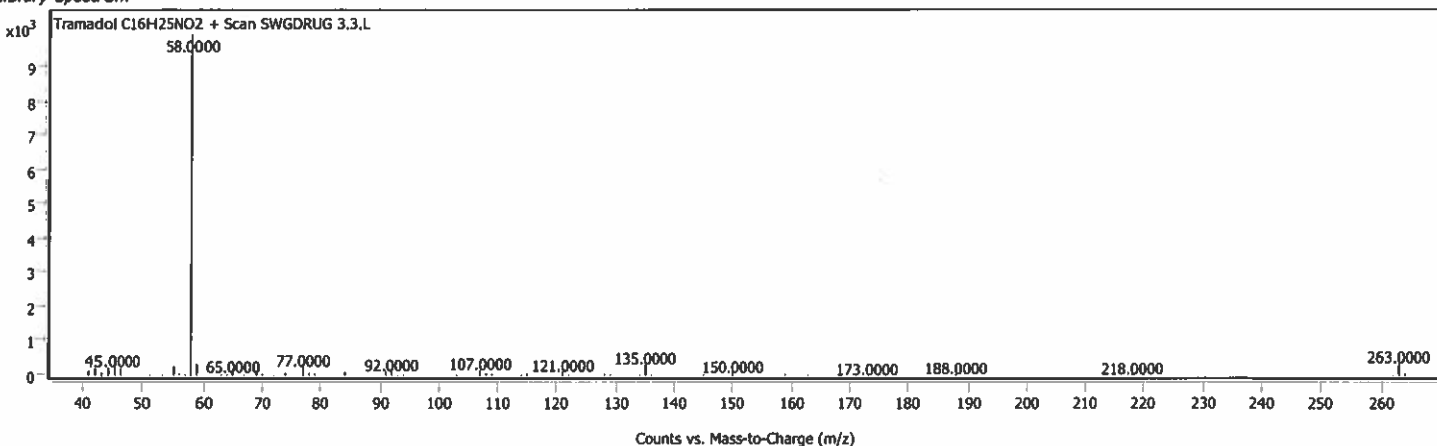
## Observed Spectrum



## Mirror Plot



## Library Spectrum





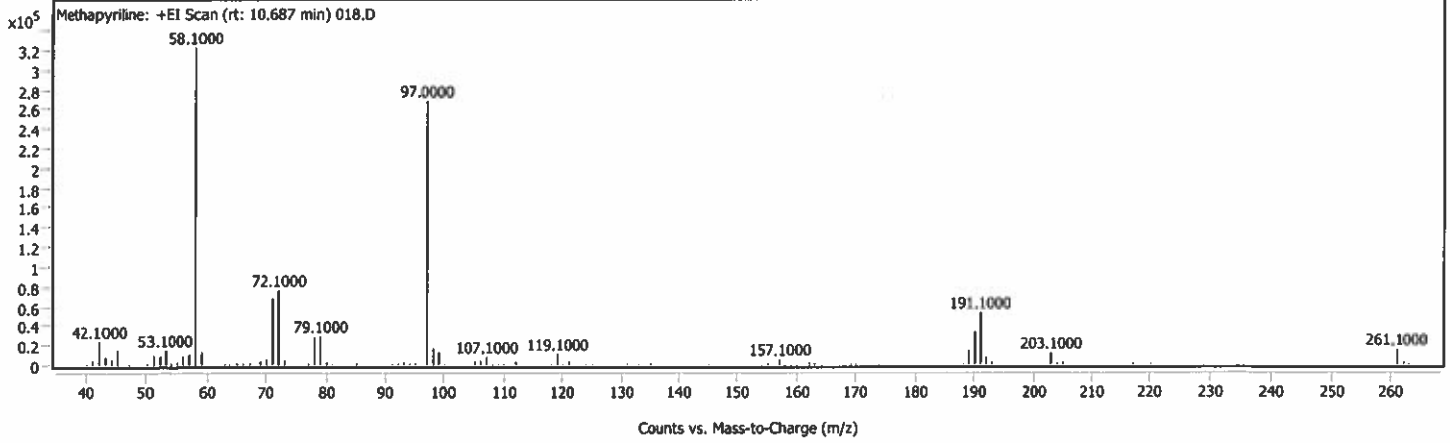
# Analysis Report

+ Scan (rt: 10.687 min)

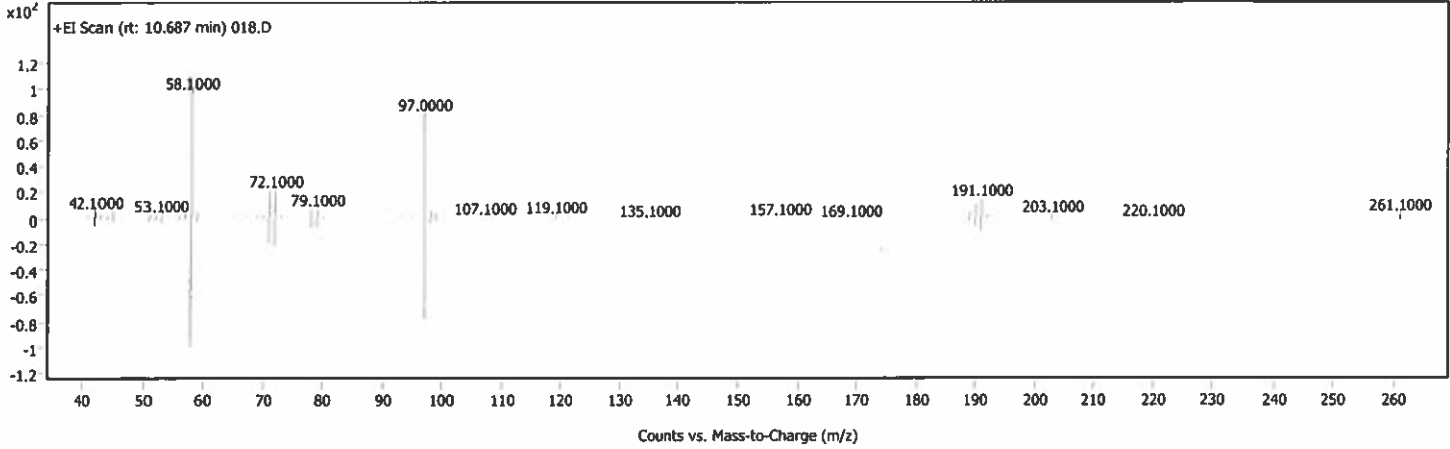
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.08	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

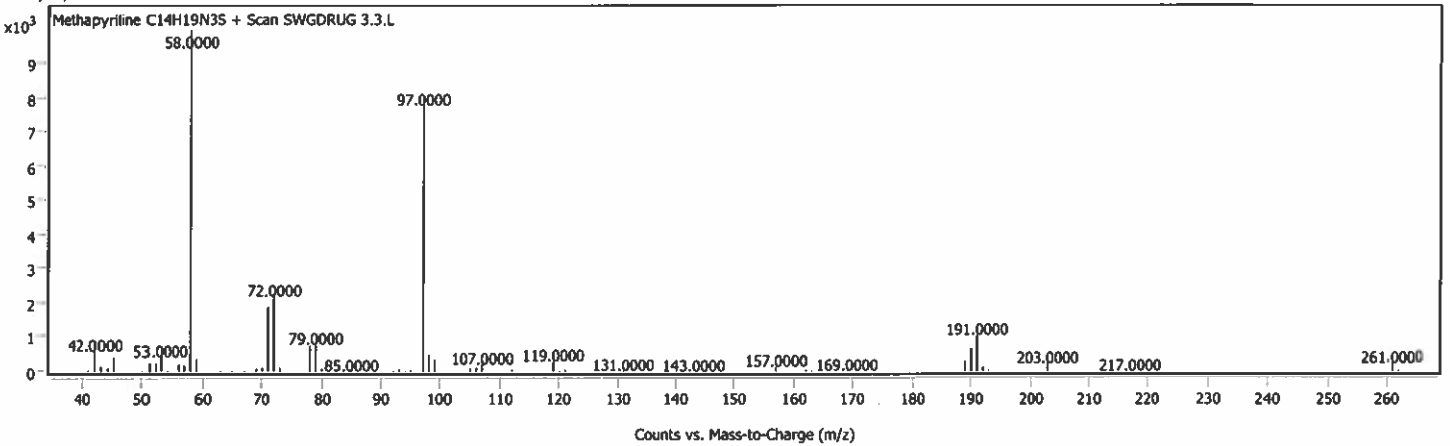
## Observed Spectrum



## Mirror Plot



## Library Spectrum



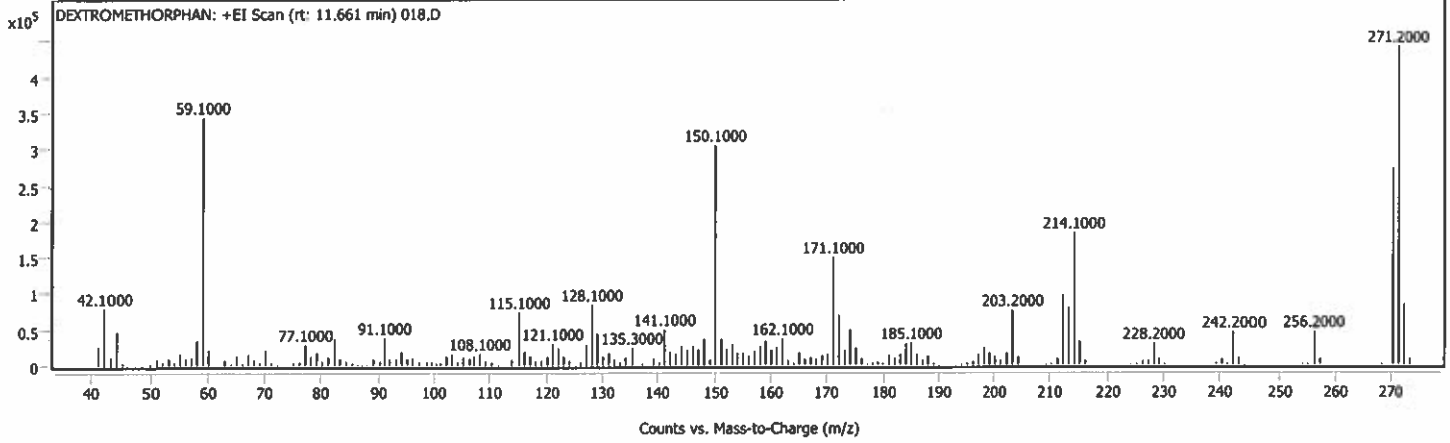
# Analysis Report

+ Scan (rt: 11.661 min)

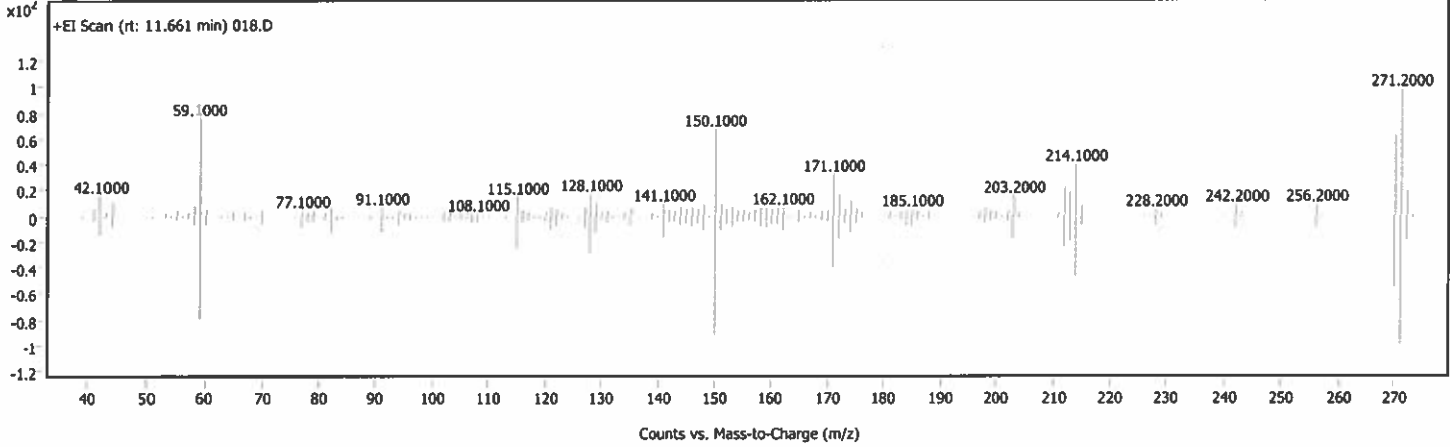
DEXTROMETHORPHAN; C<sub>18</sub>H<sub>25</sub>NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C <sub>18</sub> H <sub>25</sub> NO	92.43	aafs.I	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

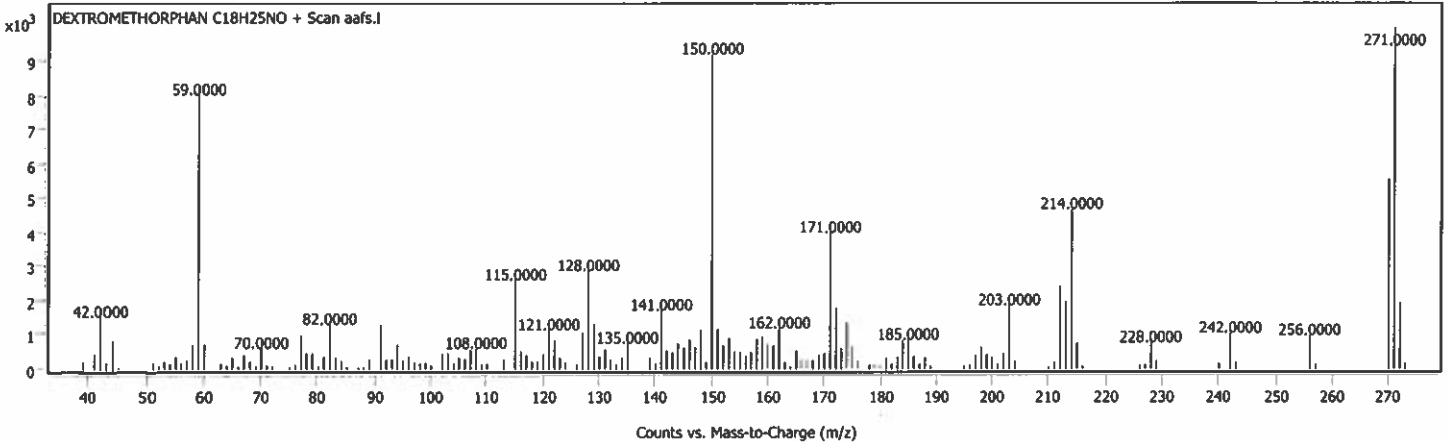
## Observed Spectrum



## Mirror Plot



## Library Spectrum



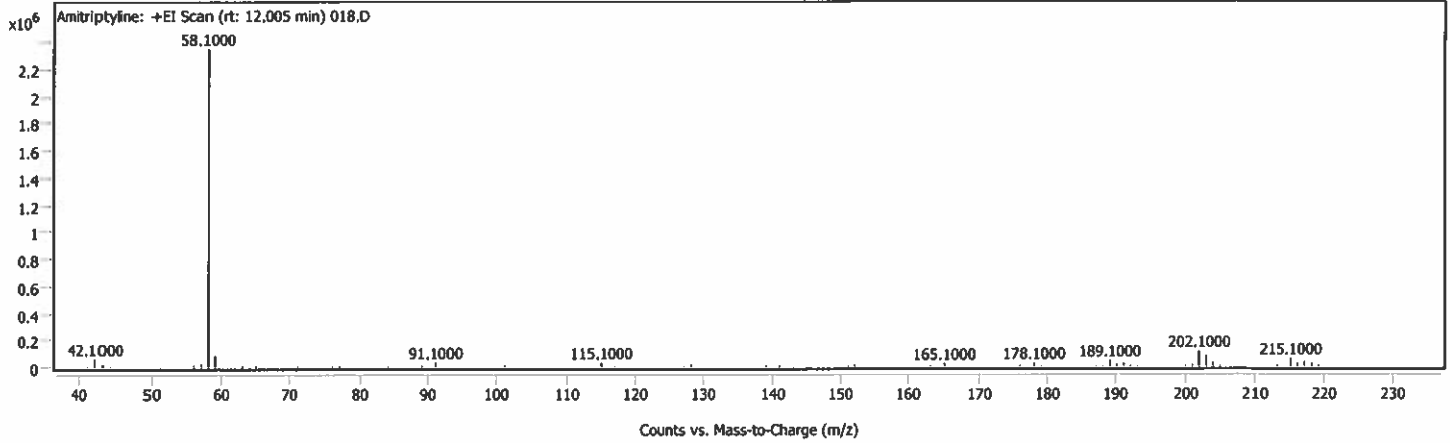
# Analysis Report

+ Scan (rt: 12.005 min)

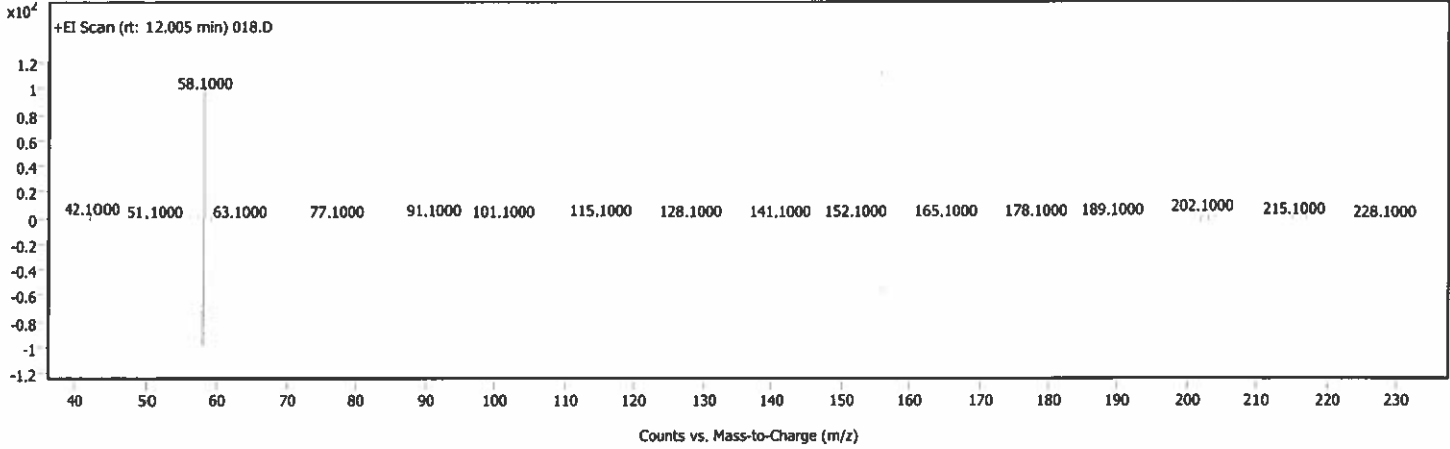
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	97.94	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

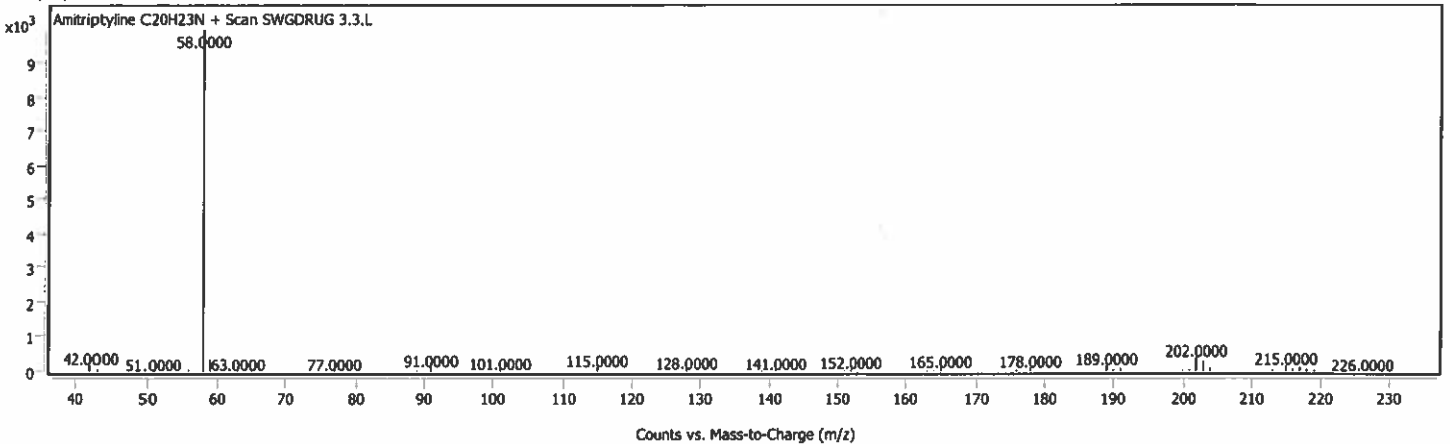
## Observed Spectrum



## Mirror Plot



## Library Spectrum



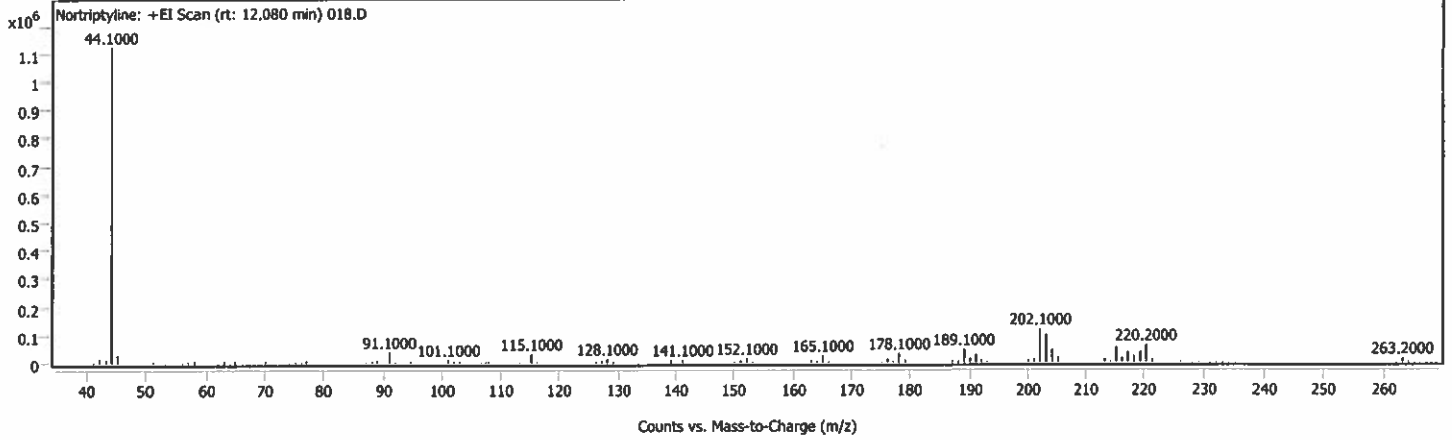
# Analysis Report

+ Scan (rt: 12.080 min)

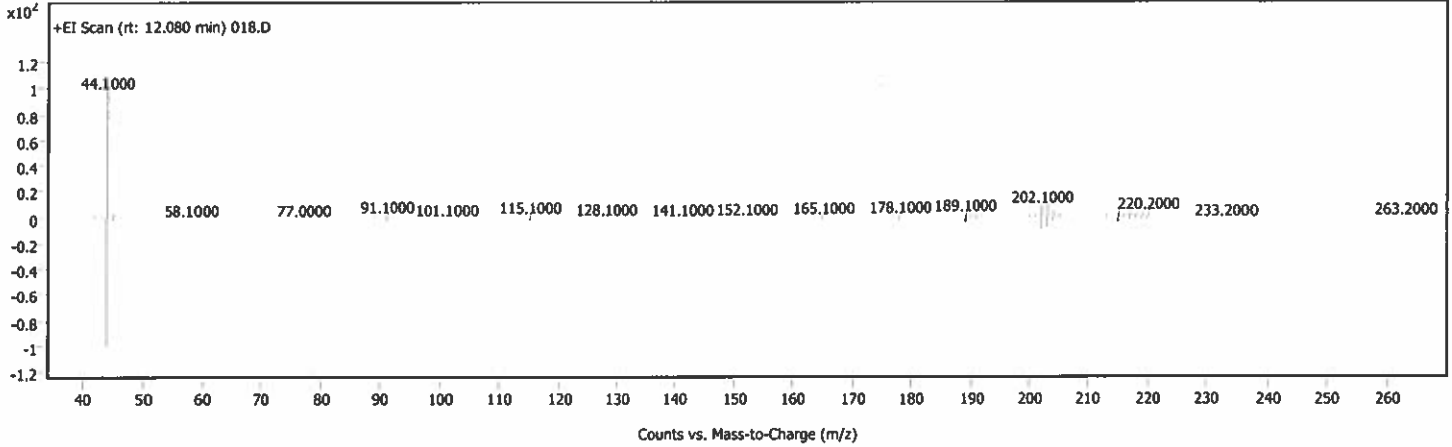
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	97.77	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

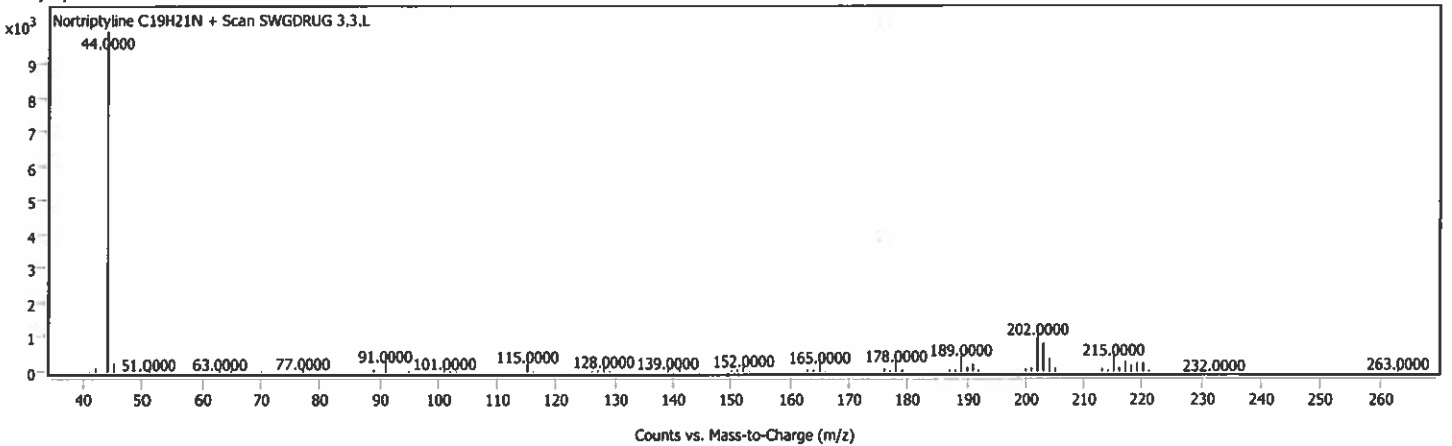
## Observed Spectrum



## Mirror Plot



## Library Spectrum



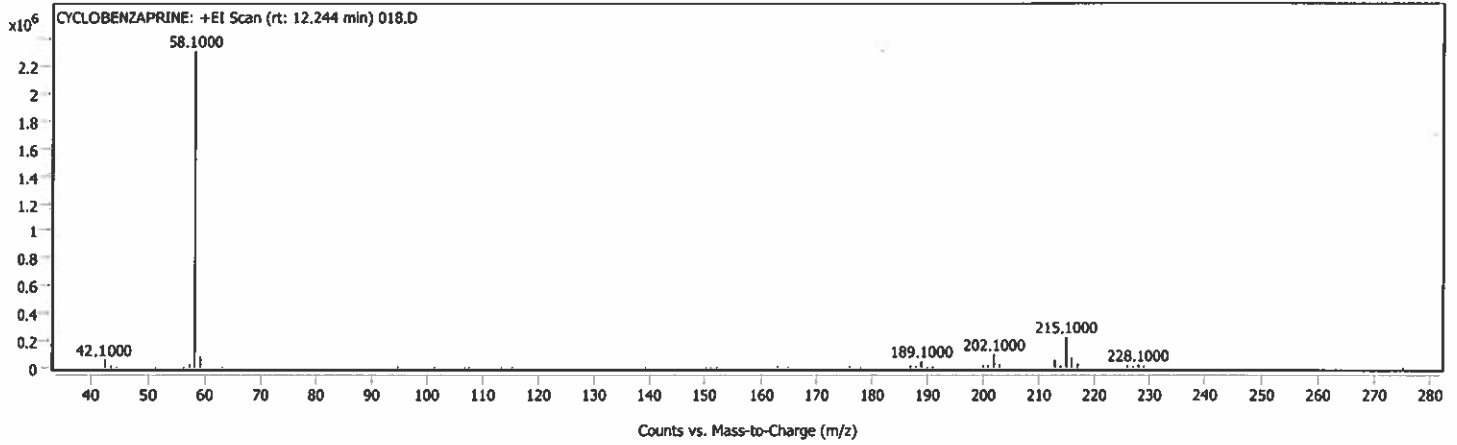
# Analysis Report

+ Scan (rt: 12.244 min)

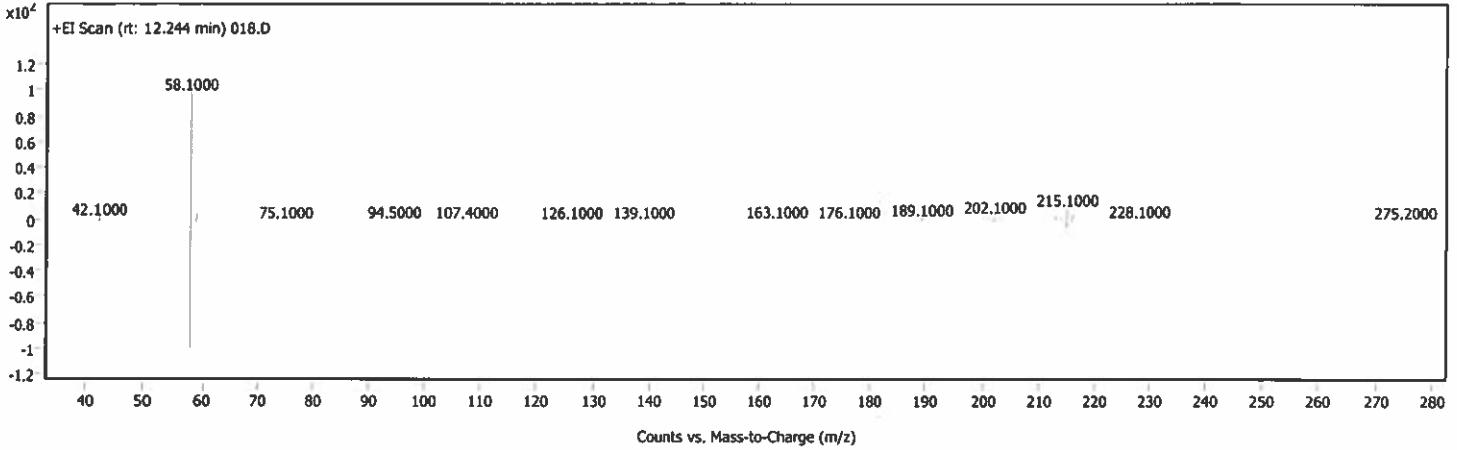
CYCLOBENZAPRINE; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CYCLOBENZAPRINE	C20H21N	96.83	aafs.l	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

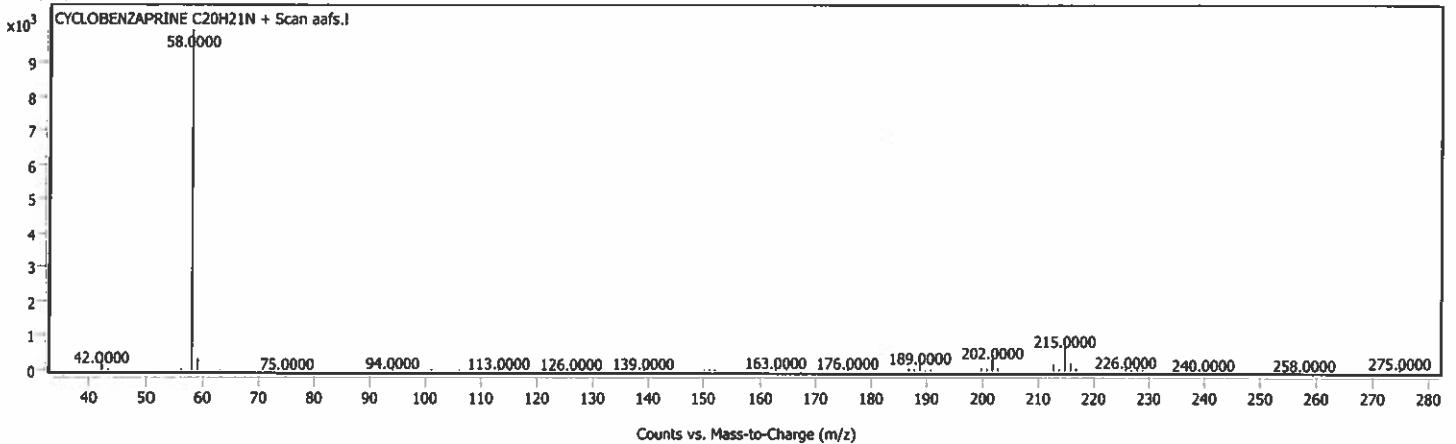
### Observed Spectrum



### Mirror Plot



### Library Spectrum



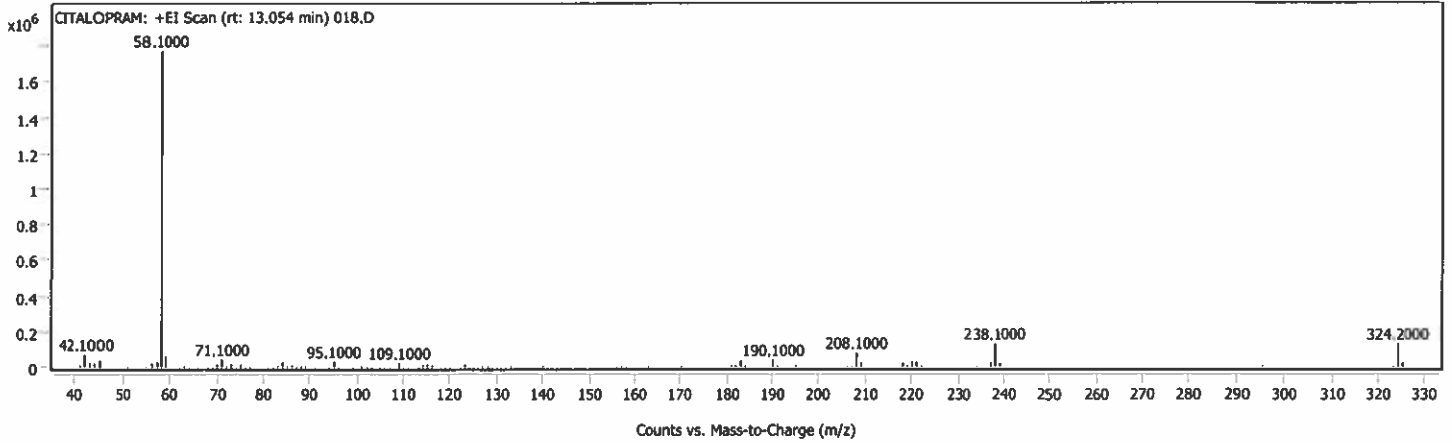
# Analysis Report

+ Scan (rt: 13.054 min)

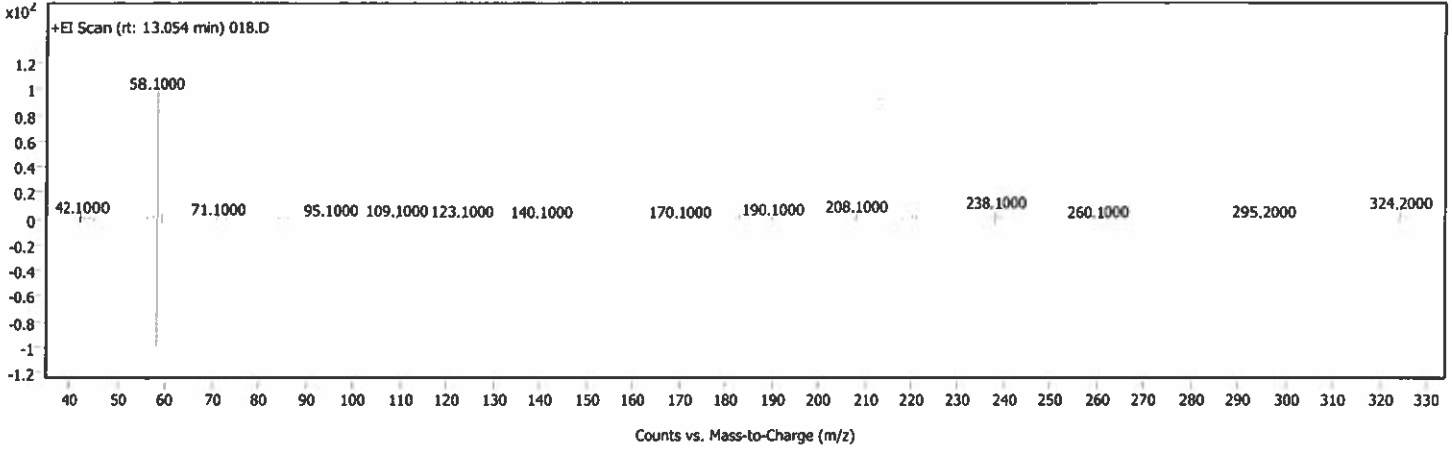
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	96.09	aafs.I	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

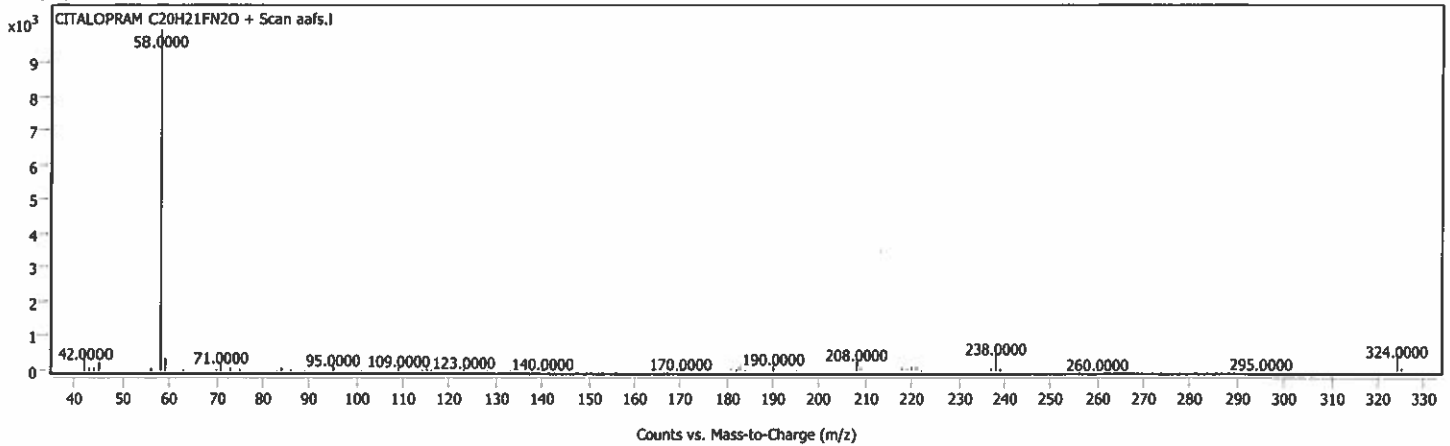
### Observed Spectrum



### Mirror Plot



### Library Spectrum



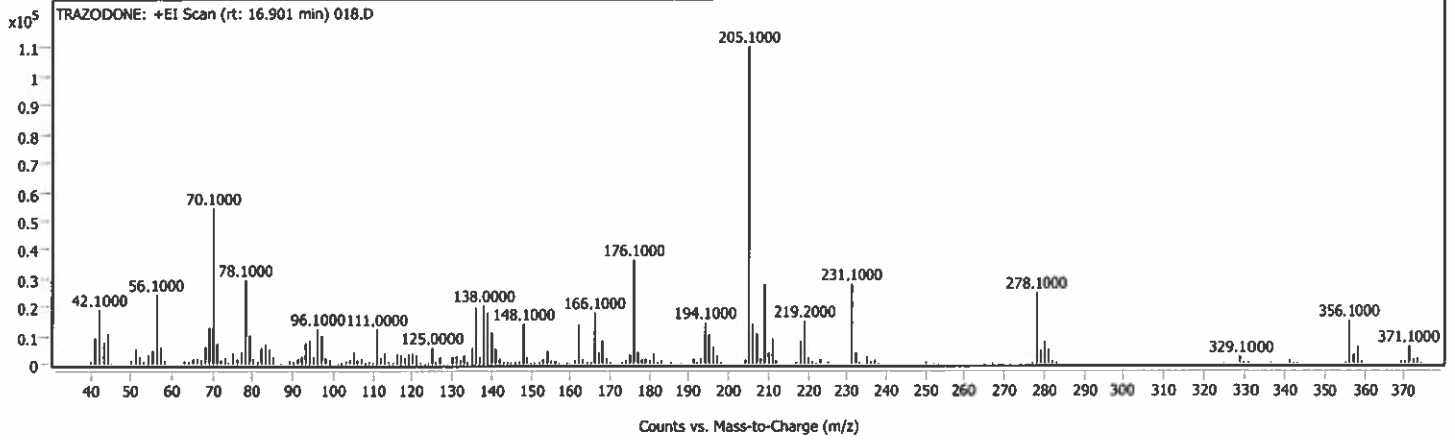
# Analysis Report

+ Scan (rt: 16.901 min)

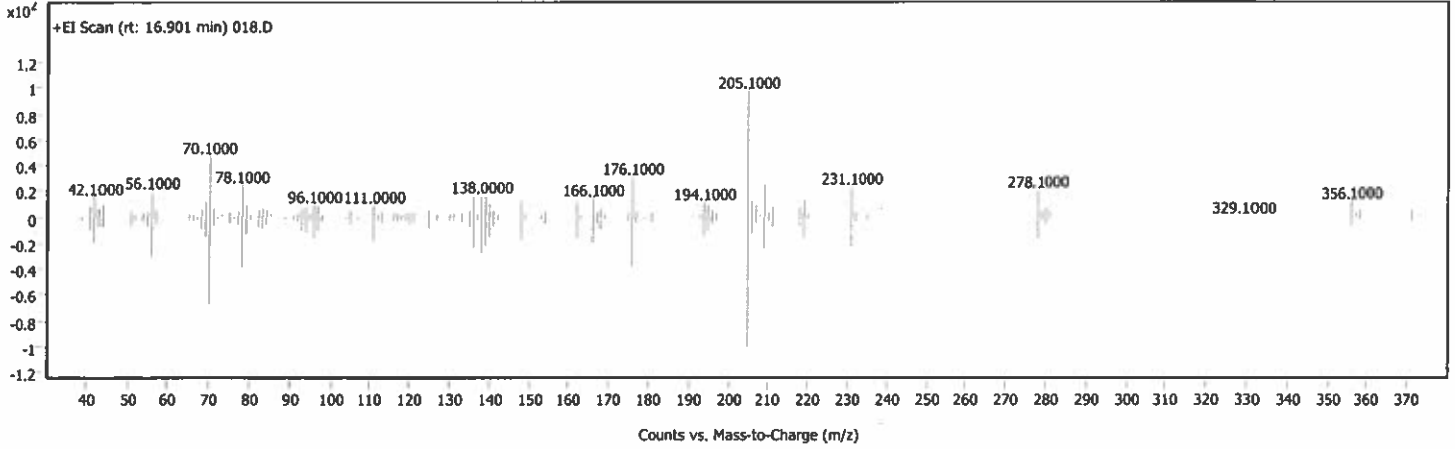
TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		95.19	aafs.1	MATRIX 2 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

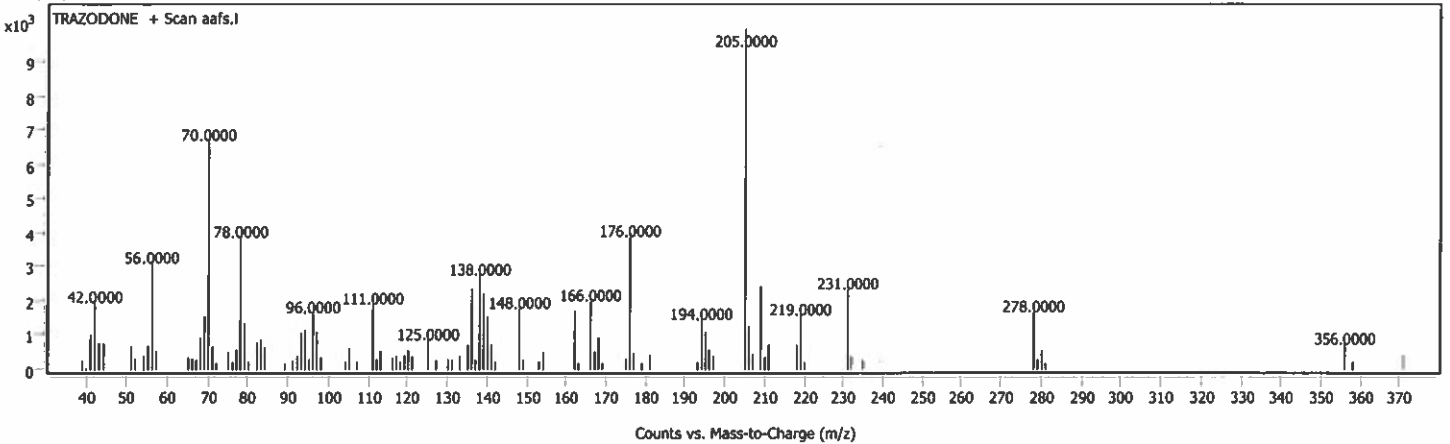
### Observed Spectrum



### Mirror Plot



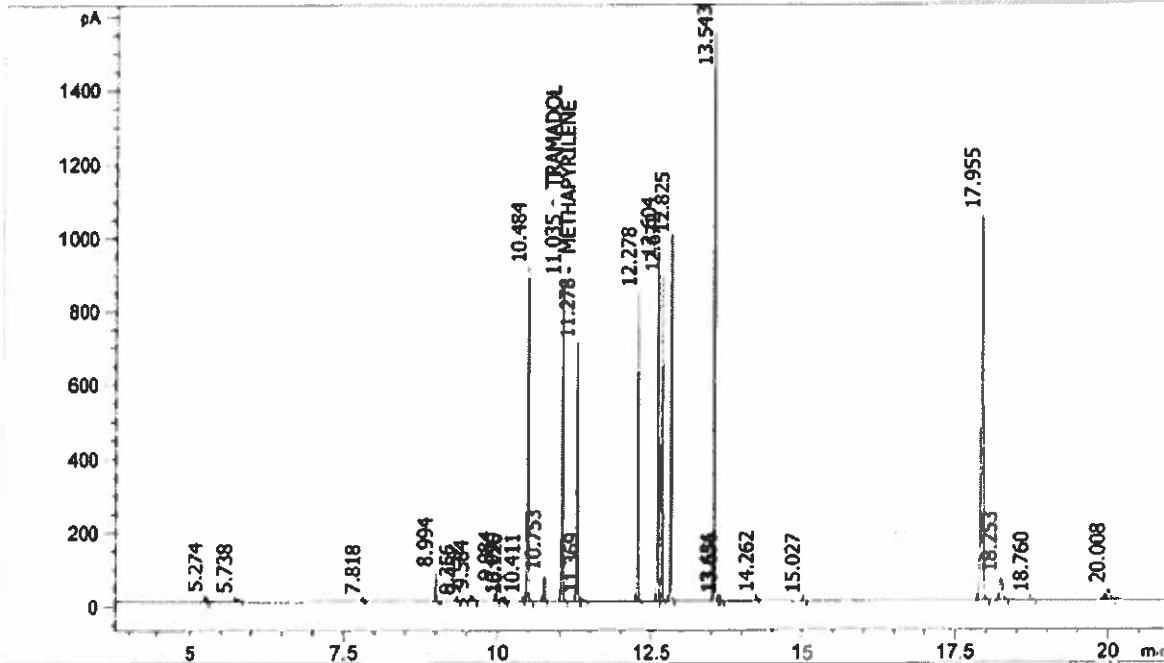
### Library Spectrum



Injection Date: 9/28/2021 9:03:11 PM  
 Sample Name: MATRIX 2 W/ SW CTL - ->  
 Sample Info:

Seq Line: 19  
 Vial 20

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.035	11.033	1079.018	871.382	0.000000	TRAMADOL RRT - 0.9785
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	799.039	694.321	1.000000	METHAPYRILENE



# Analysis Report

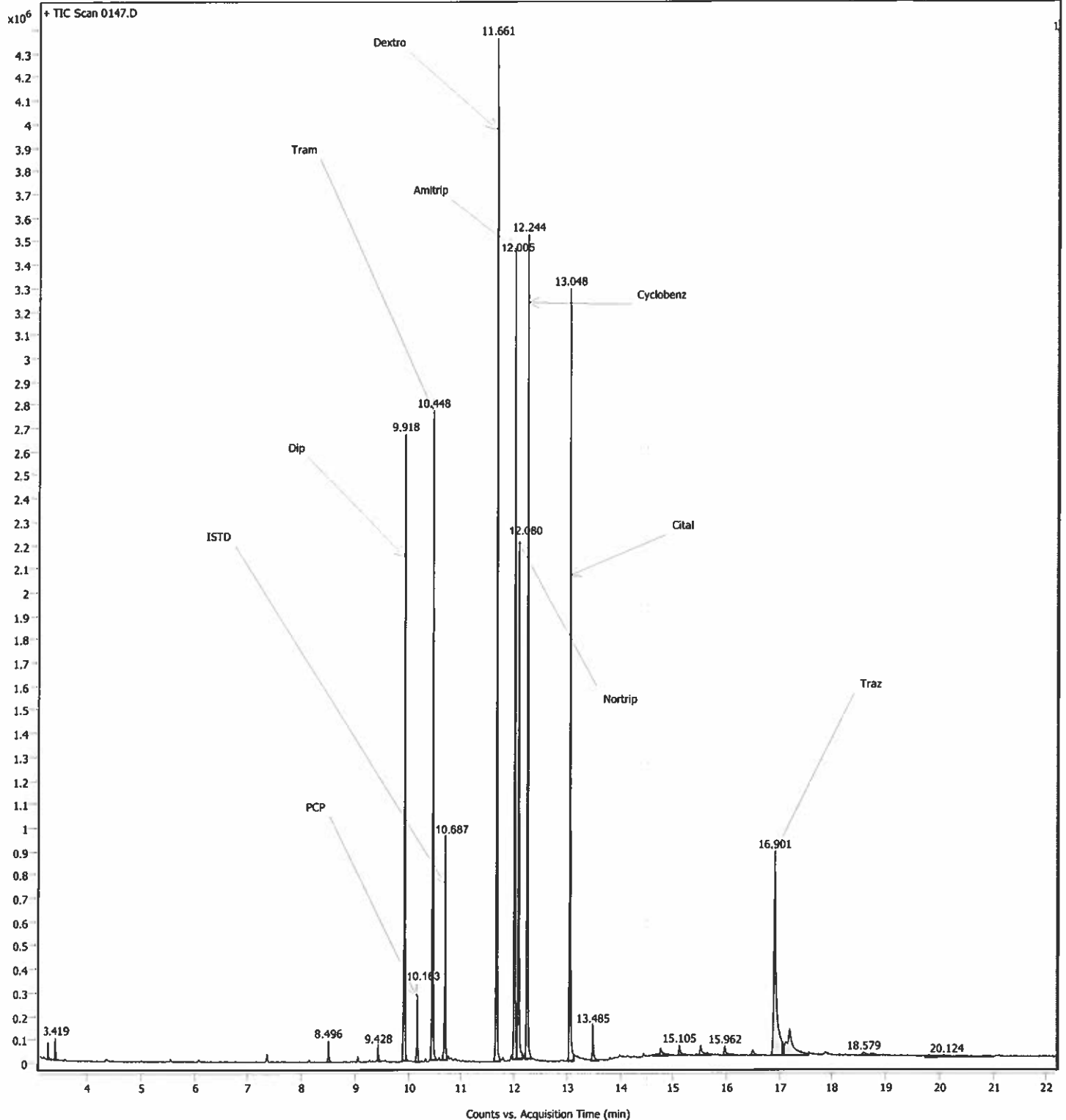
## Sample Information

**Sample Name** MATRIX 1 W/ SW CTL - BLD  
**Instrument** #3 - Enhanced  
**Position** 43  
**Operator** KM/QC DATA IN XYLAZINE VALIDATION

**Data File Path**  
**Acq. Time (Local)**  
**Method Path (Acq)**

C:\MassHunter\GCMS\1\data\BASES\092821\0147.D  
9/28/2021 8:33:54 PM (UTC-04:00)  
C:\MassHunter\GCMS\1\methods\ALKALI.M

## Sample Chromatograms



# Analysis Report

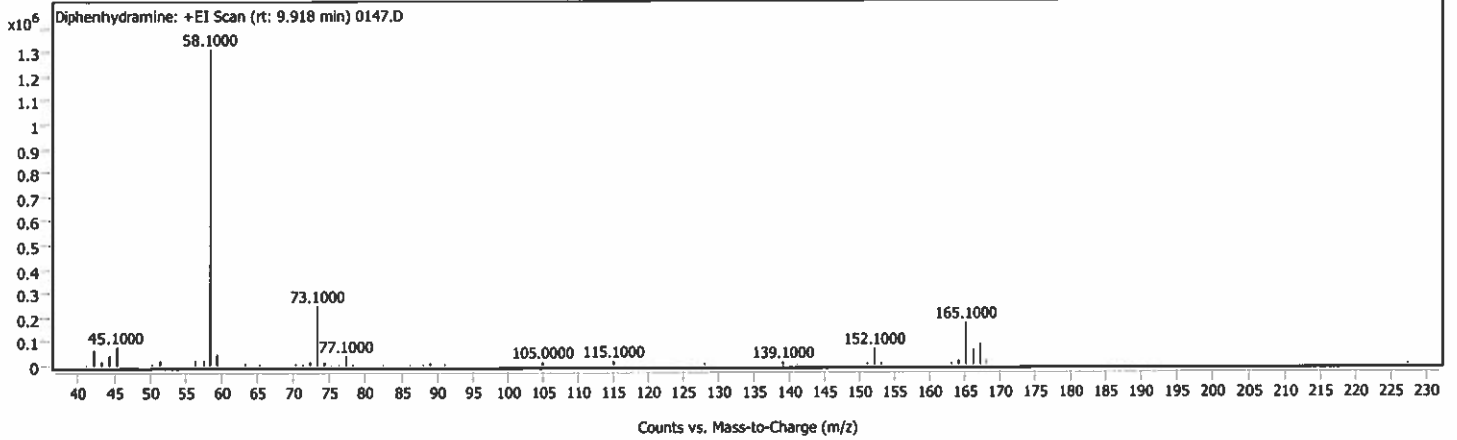
## Sample Spectra

+ Scan (rt: 9.918 min)

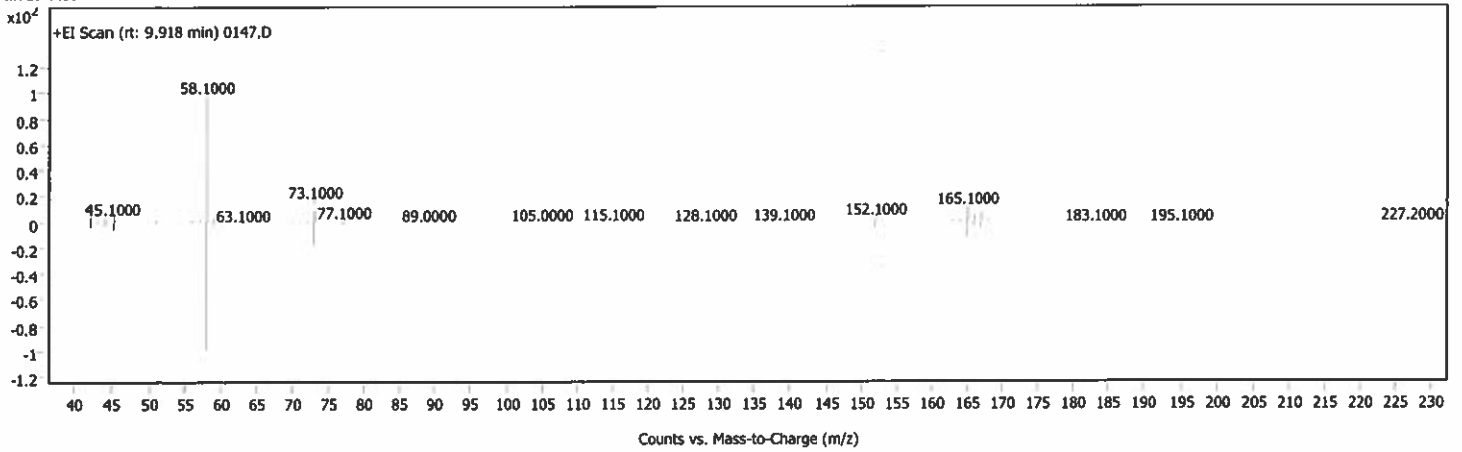
Diphenhydramine; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Diphenhydramine	C17H21NO	98.42	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

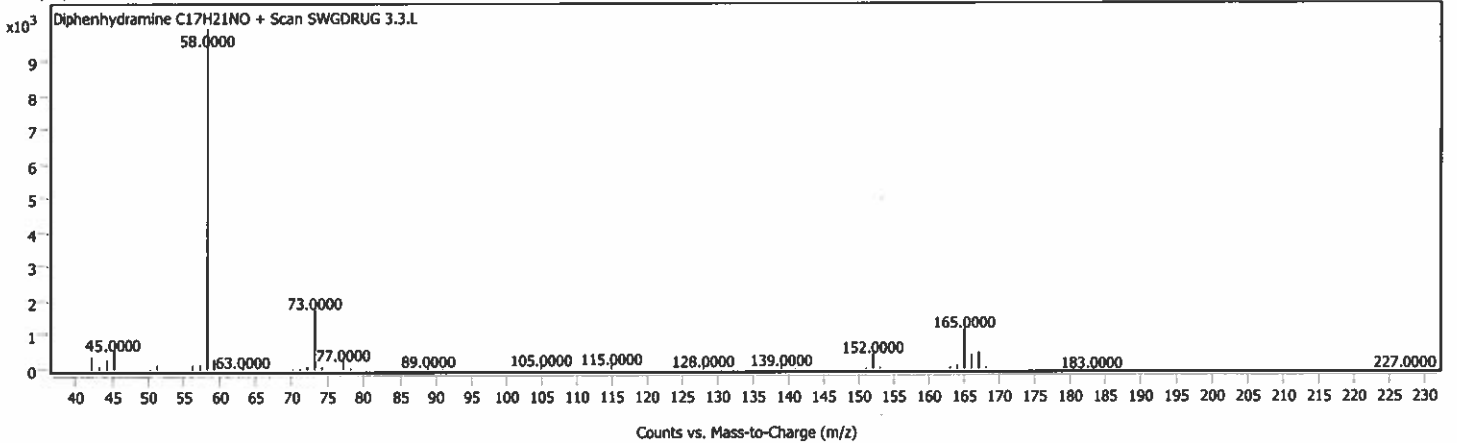
## Observed Spectrum



## Mirror Plot



## Library Spectrum



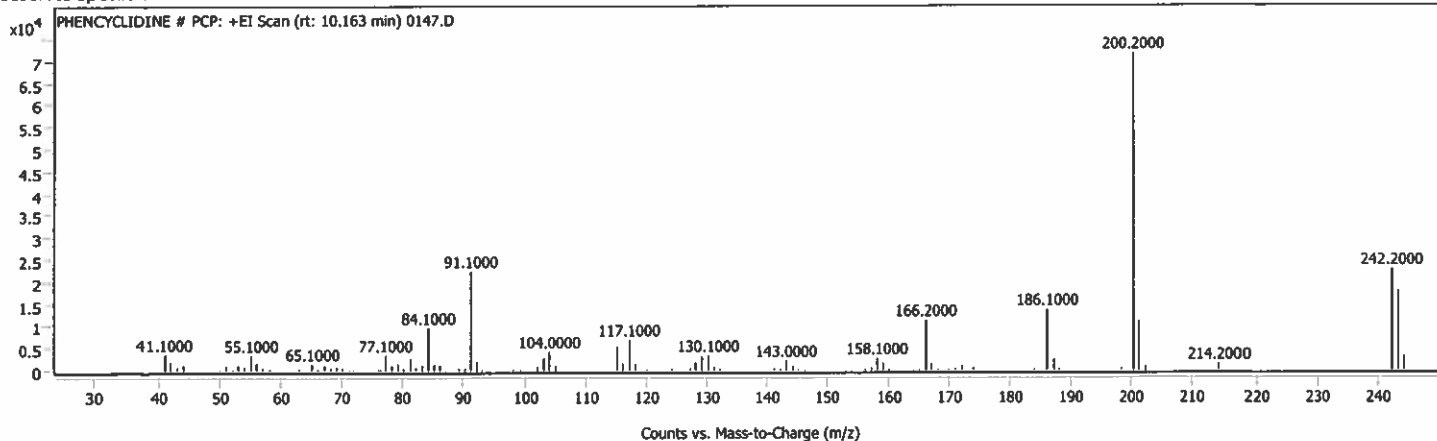
# Analysis Report

+ Scan (rt: 10.163 min)

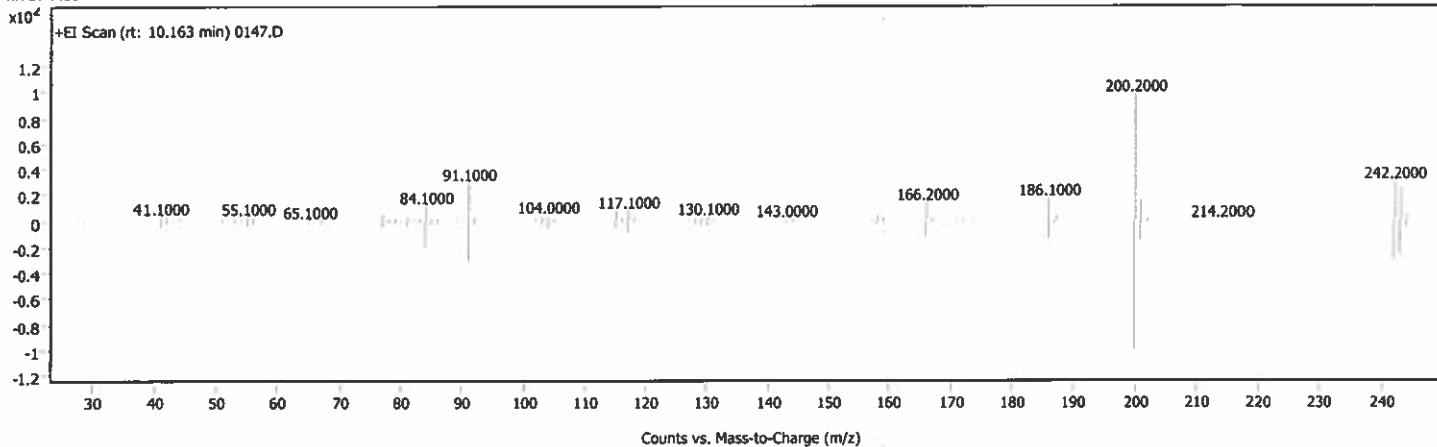
PHENCYCLIDINE # PCP; C17H25N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
PHENCYCLIDINE # PCP	C17H25N	98.27	aafs.I	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

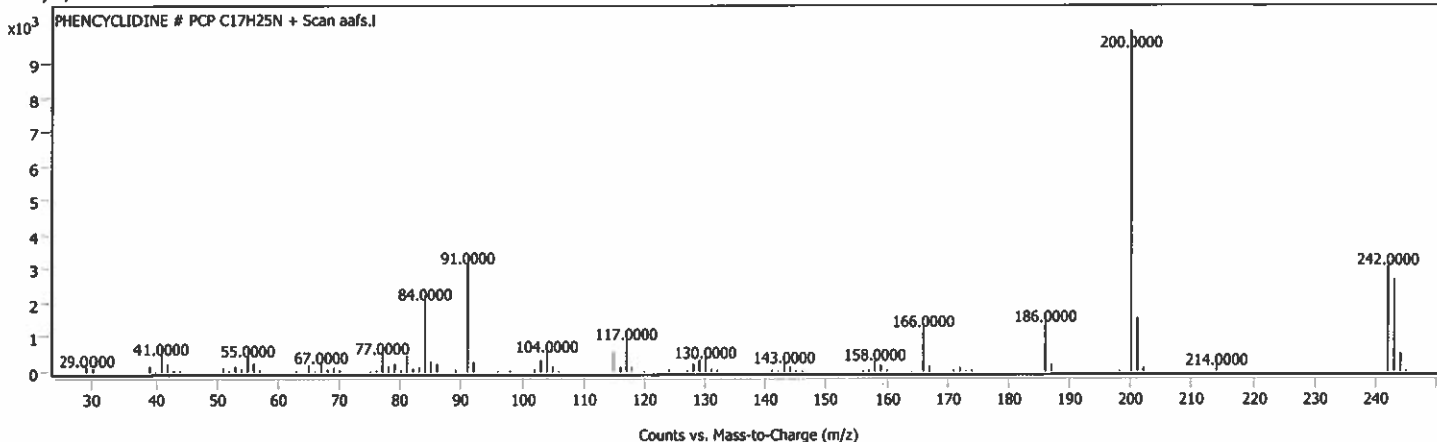
## Observed Spectrum



## Mirror Plot



## Library Spectrum



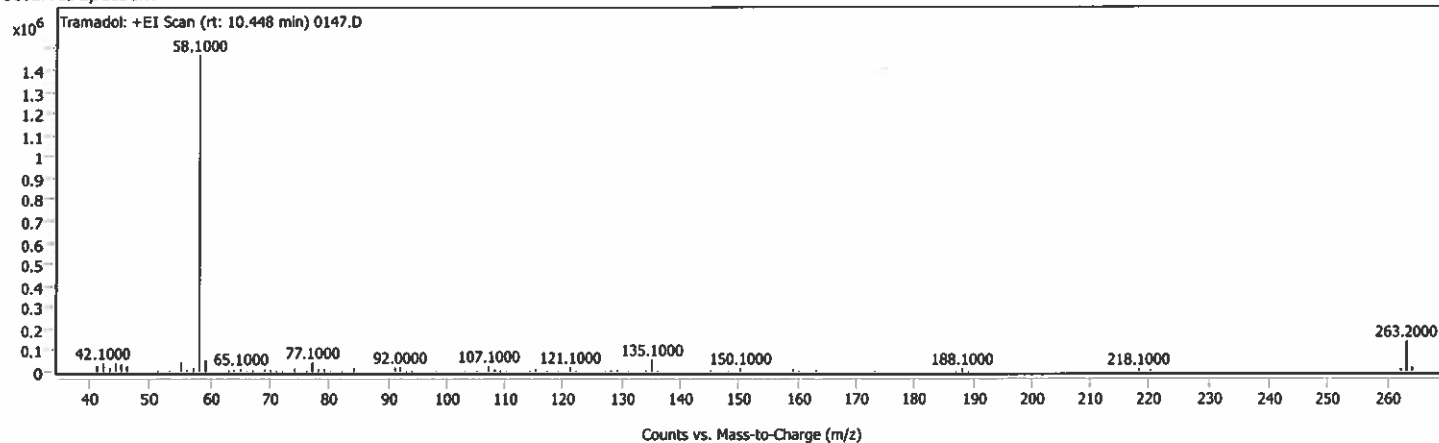
# Analysis Report

+ Scan (rt: 10.448 min)

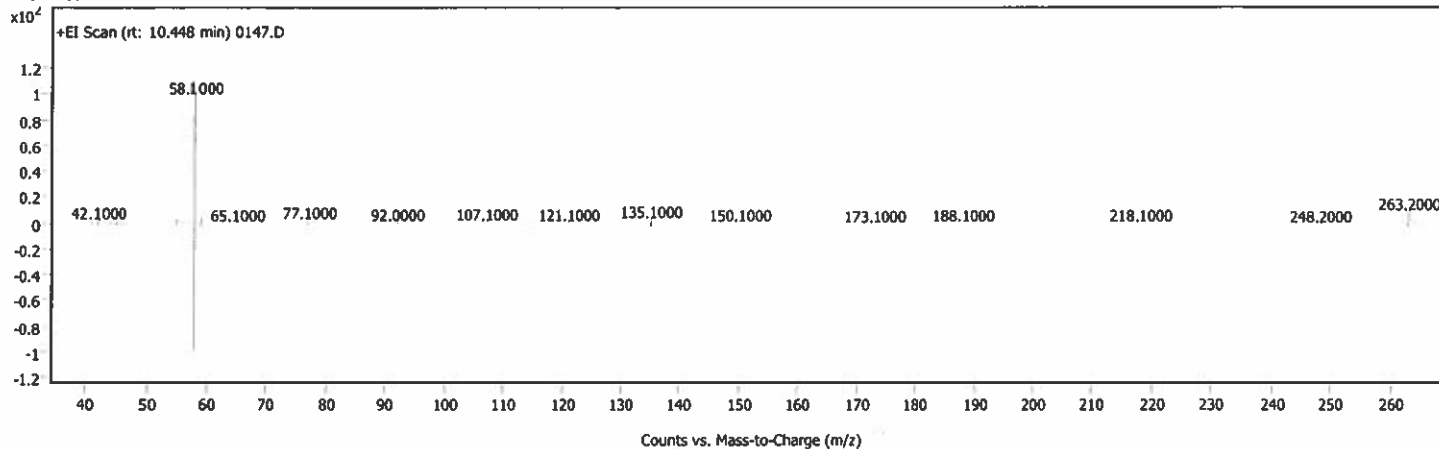
Tramadol; C16H25NO2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Tramadol	C16H25NO2	95.46	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

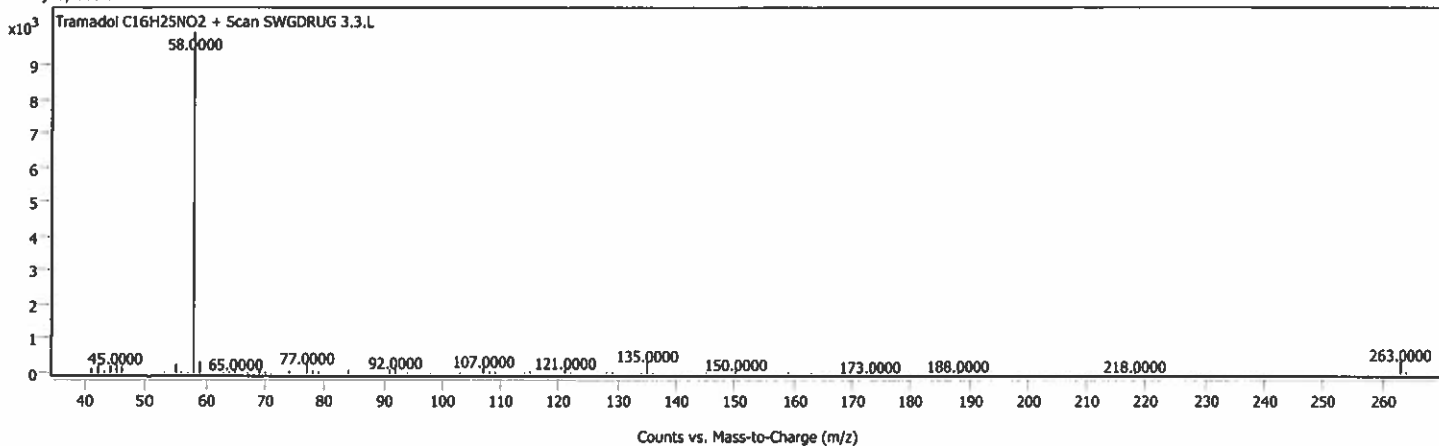
## Observed Spectrum



## Mirror Plot



## Library Spectrum



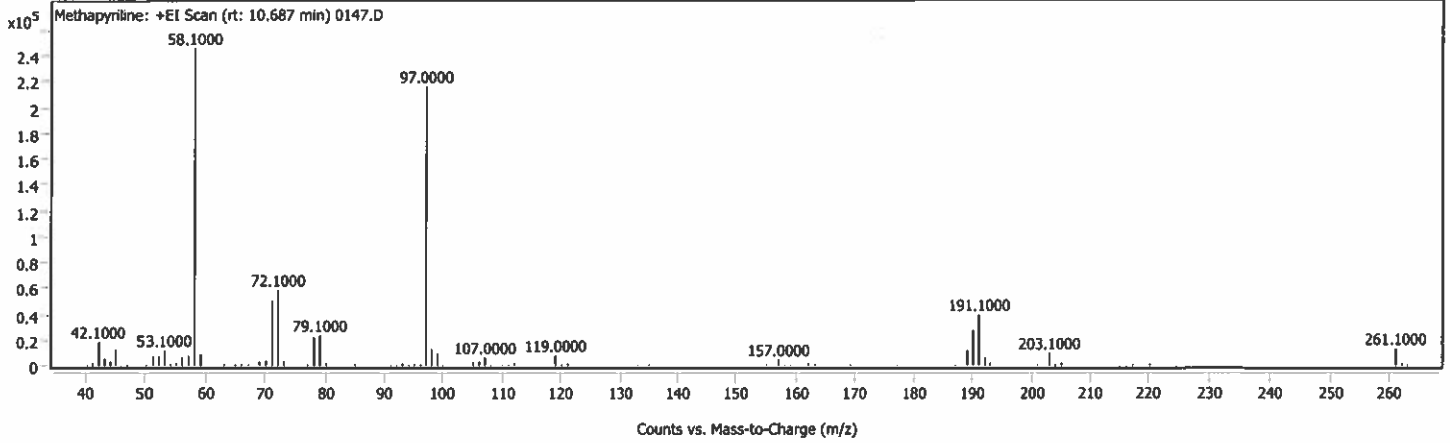
# Analysis Report

+ Scan (rt: 10.687 min)

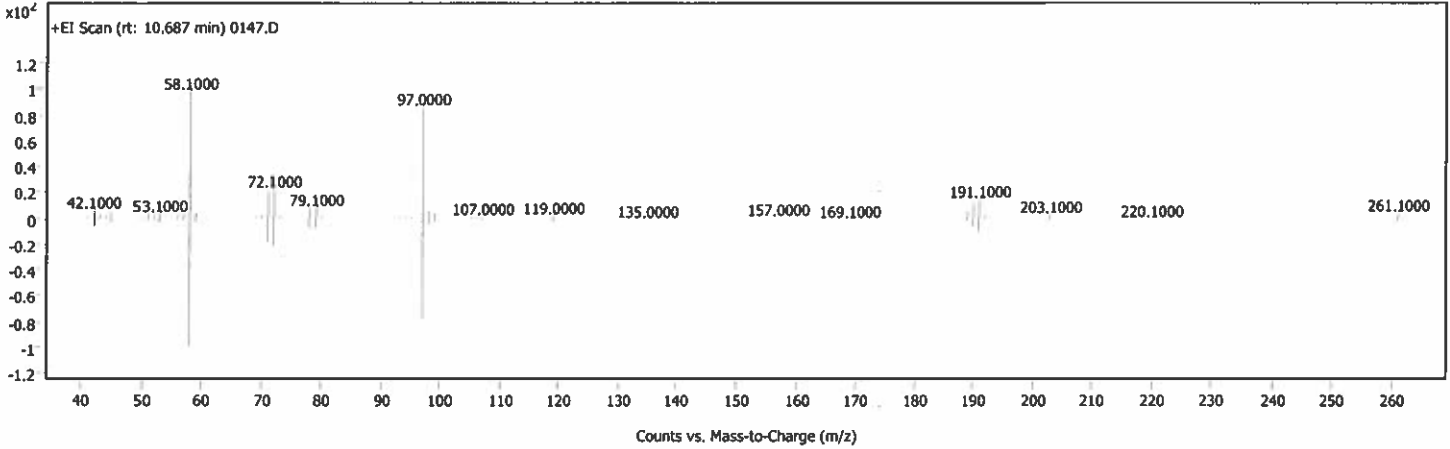
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	97.55	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XLAZINE VALIDATION

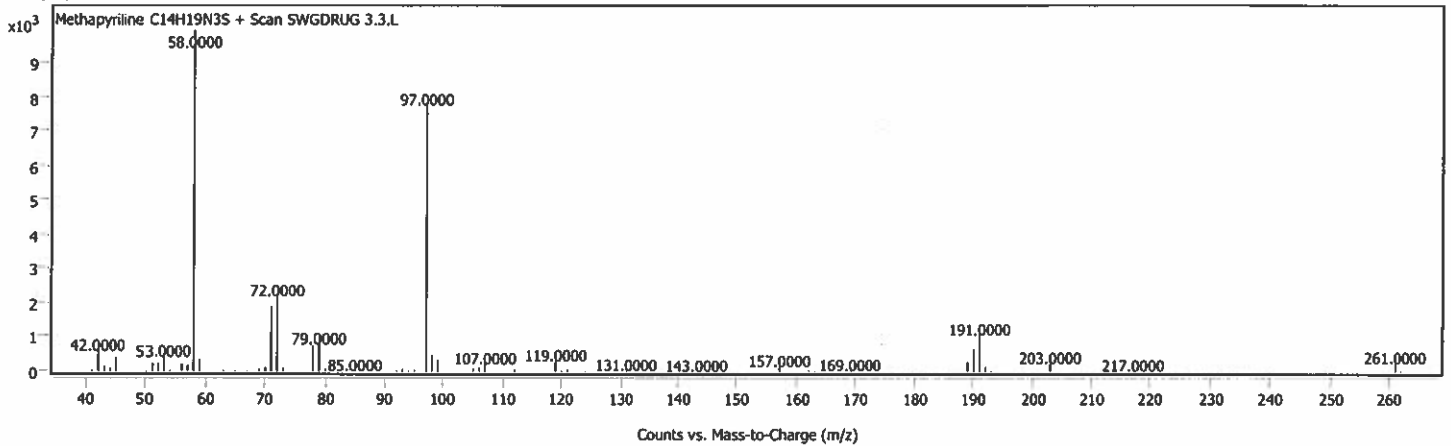
### Observed Spectrum



### Mirror Plot



### Library Spectrum



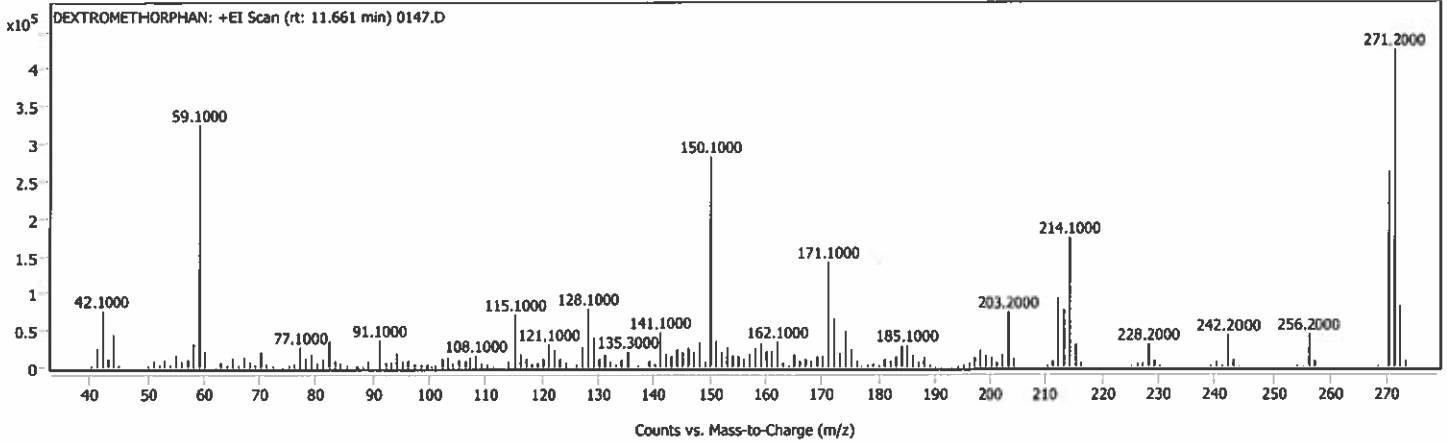
# Analysis Report

+ Scan (rt: 11.661 min)

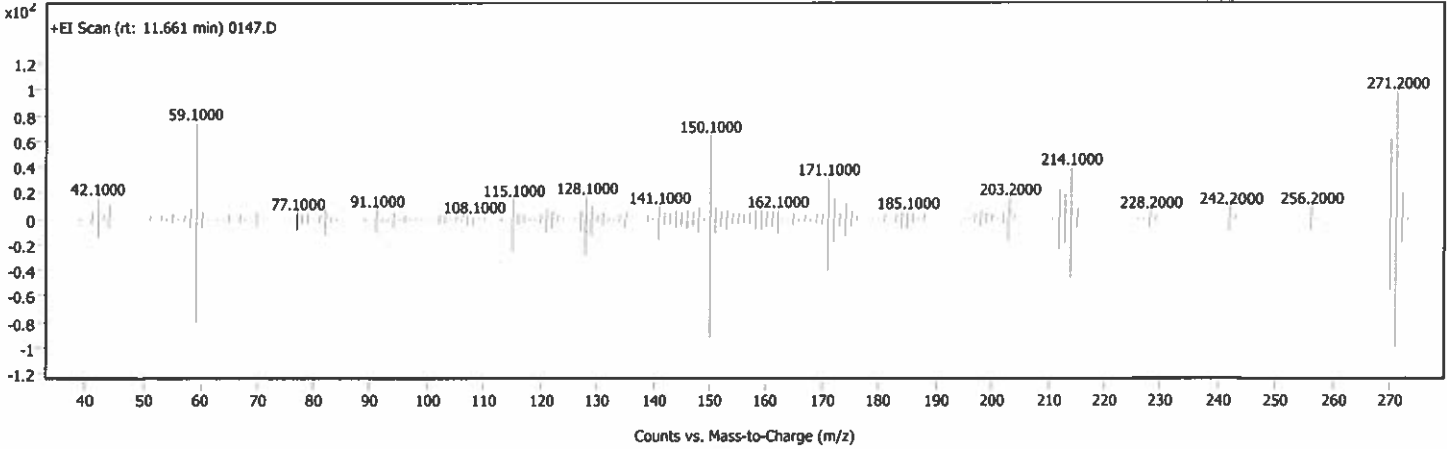
DEXTROMETHORPHAN; C18H25NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	92.19	aafs.I	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

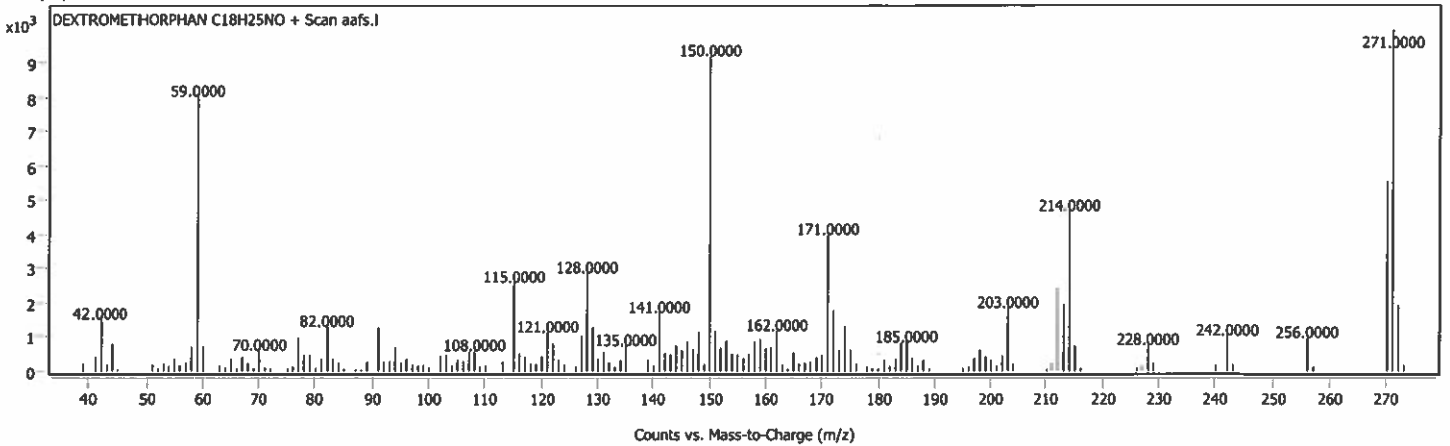
### Observed Spectrum



### Mirror Plot



### Library Spectrum



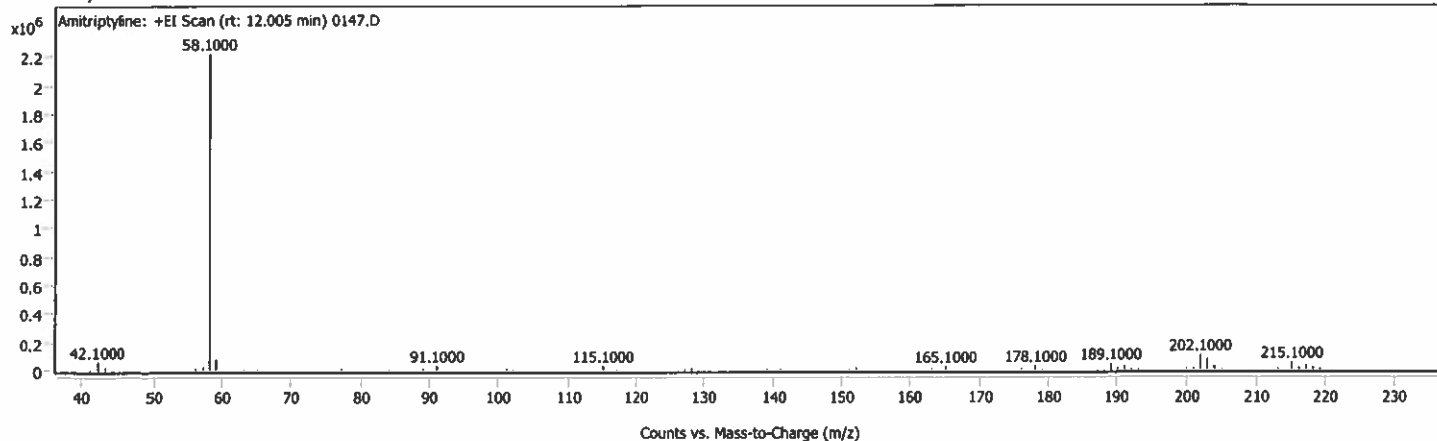
# Analysis Report

+ Scan (rt: 12.005 min)

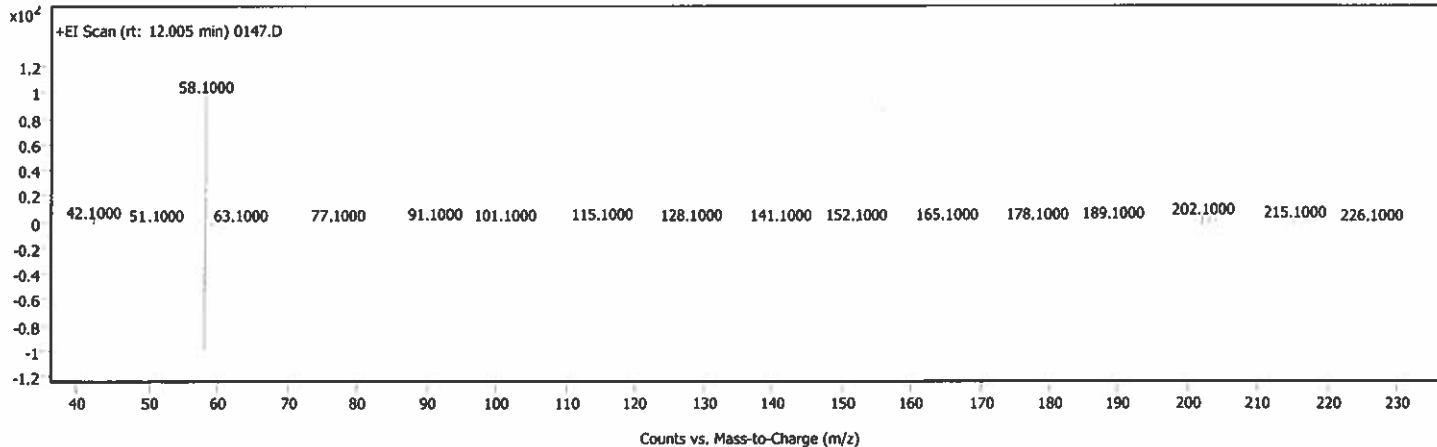
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	97.96	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

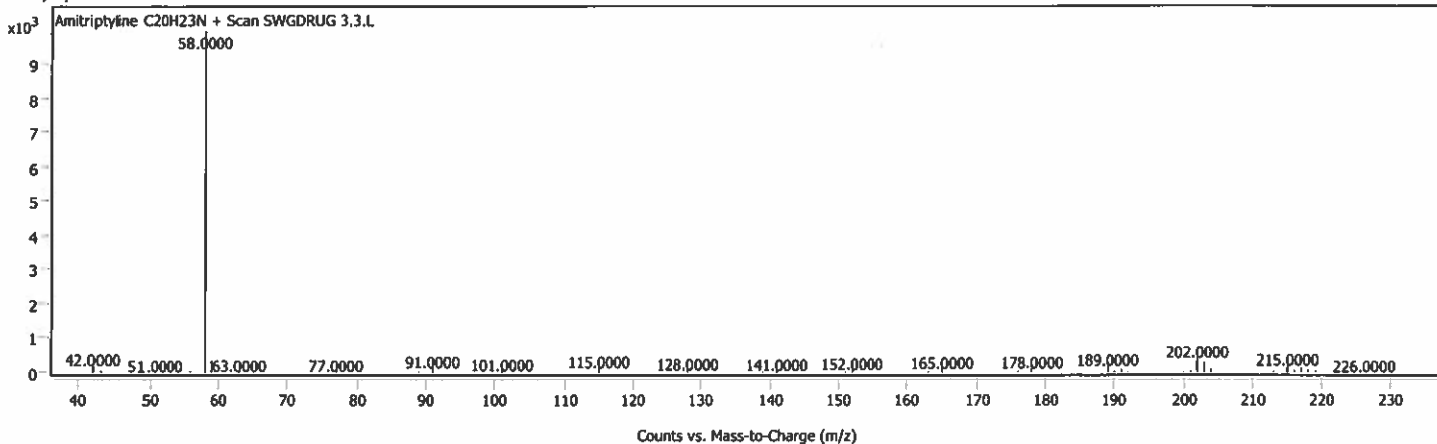
## Observed Spectrum



## Mirror Plot



## Library Spectrum



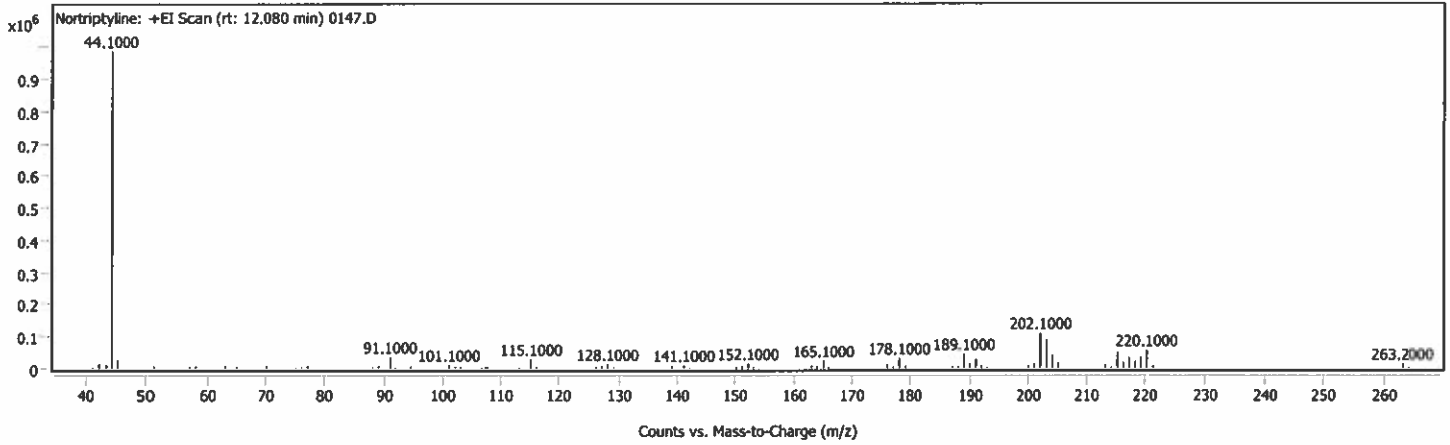
# Analysis Report

+ Scan (rt: 12.080 min)

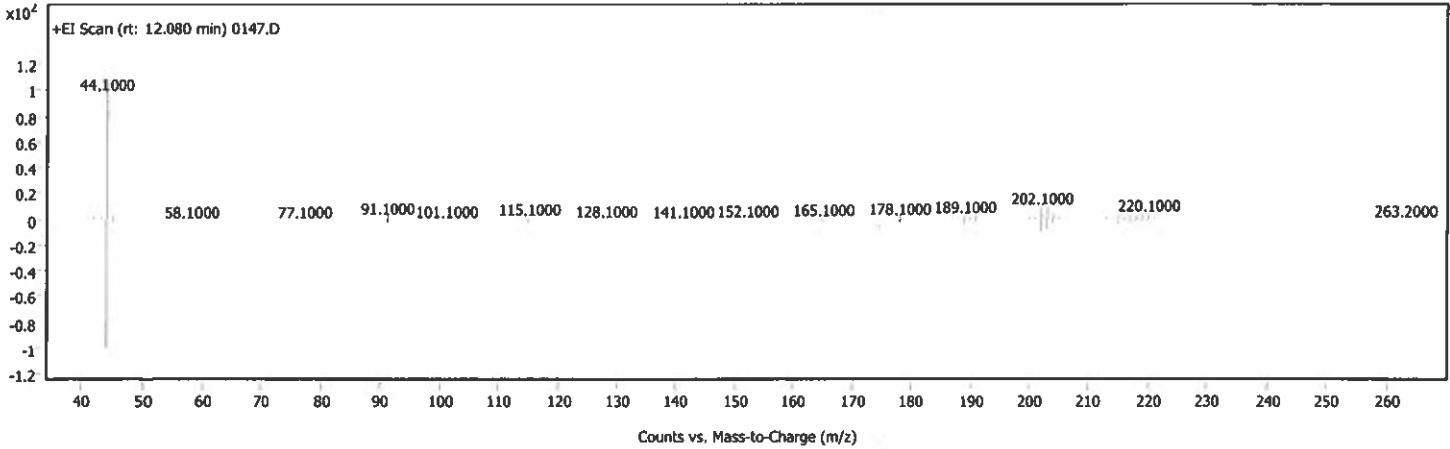
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	97.83	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

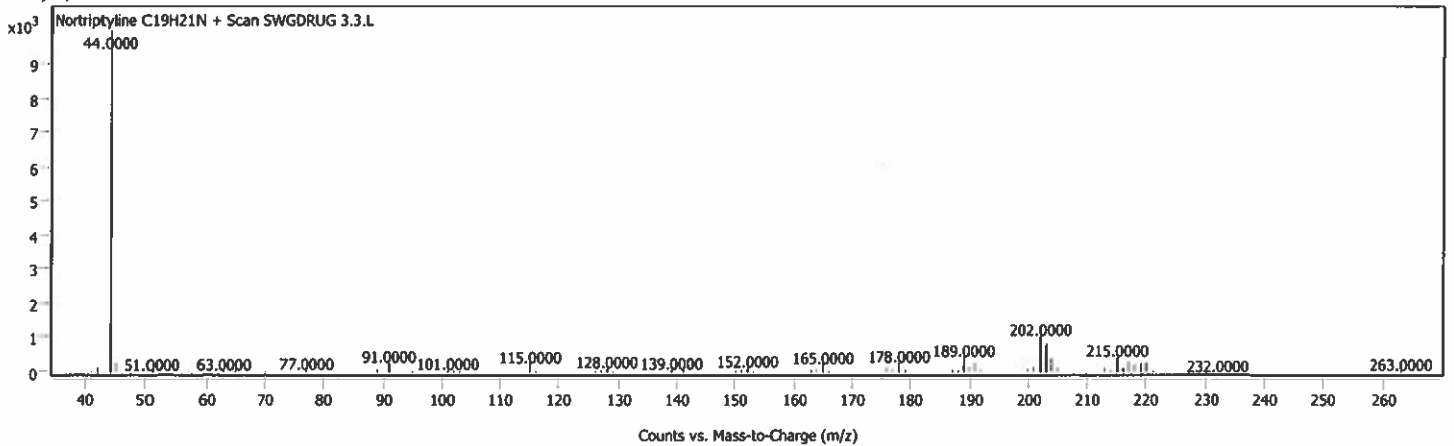
### Observed Spectrum



### Mirror Plot



### Library Spectrum





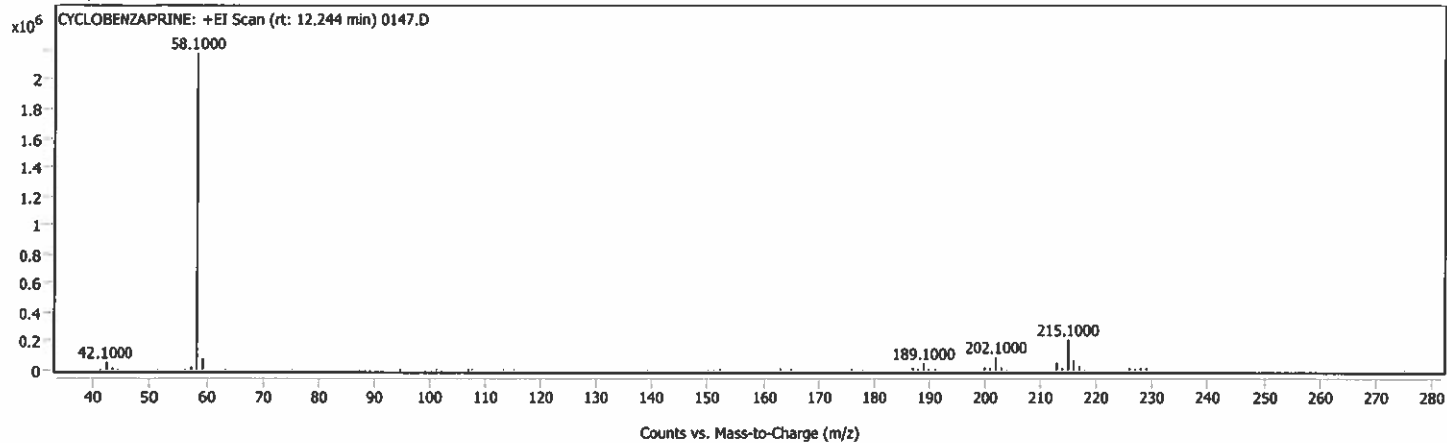
# Analysis Report

+ Scan (rt: 12.244 min)

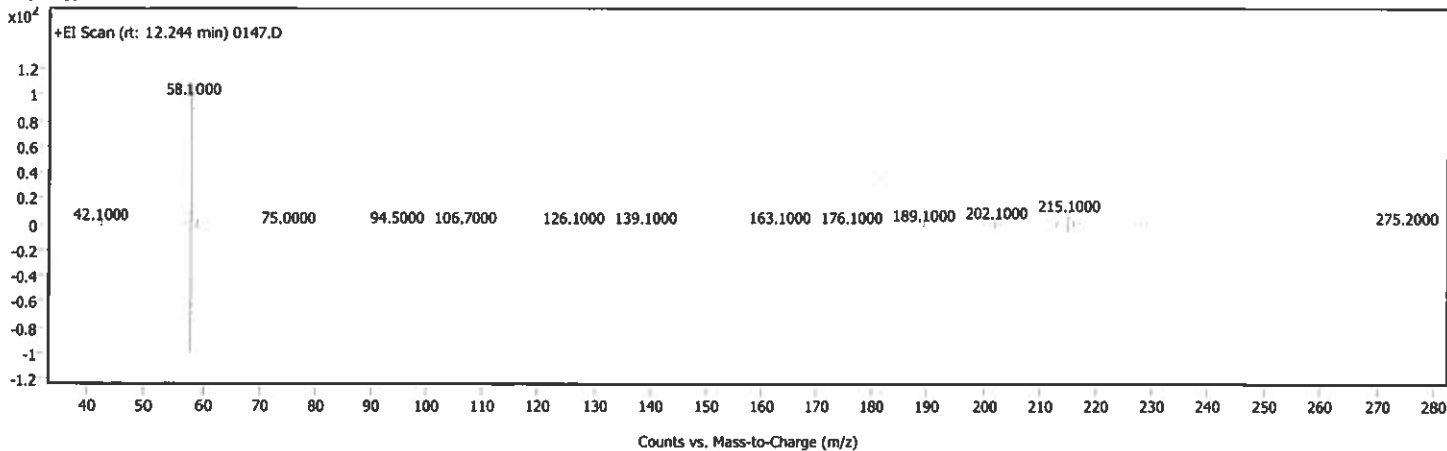
CYCLOBENZAPRINE; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CYCLOBENZAPRINE	C20H21N	96.65	aafs.1	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

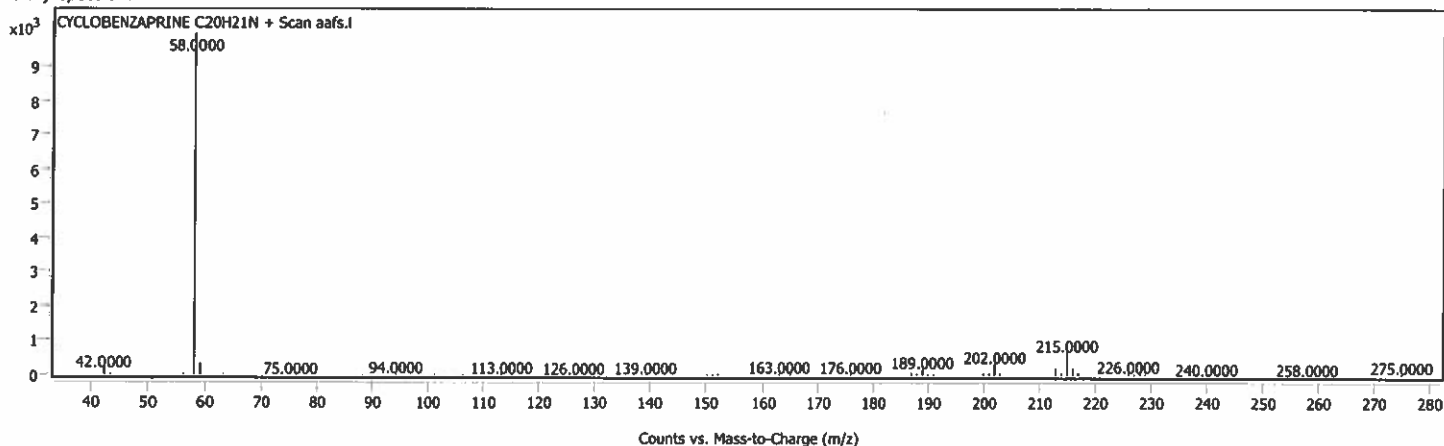
### Observed Spectrum



### Mirror Plot



### Library Spectrum



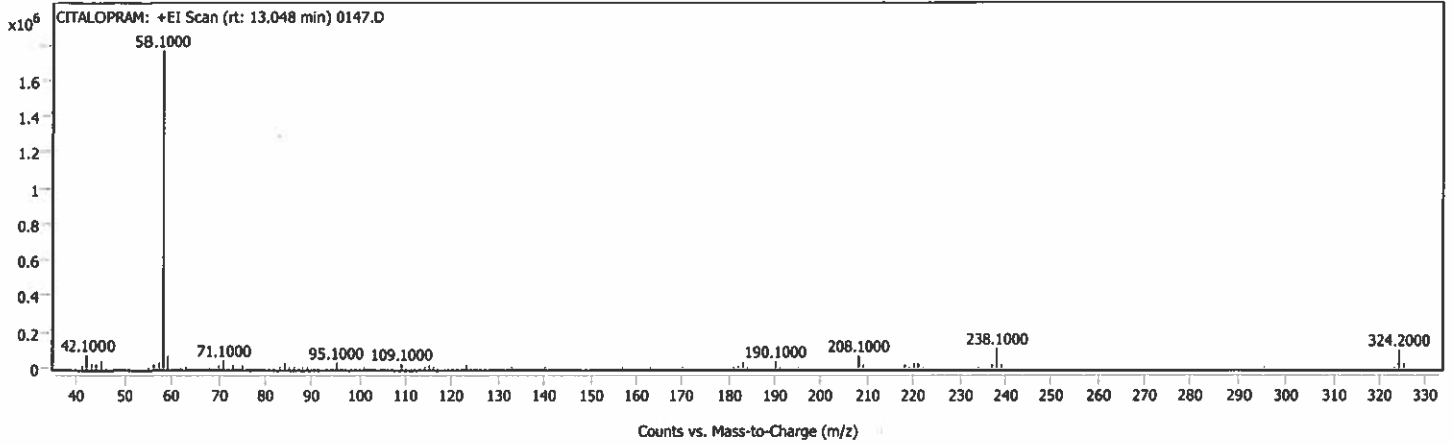
# Analysis Report

+ Scan (rt: 13.048 min)

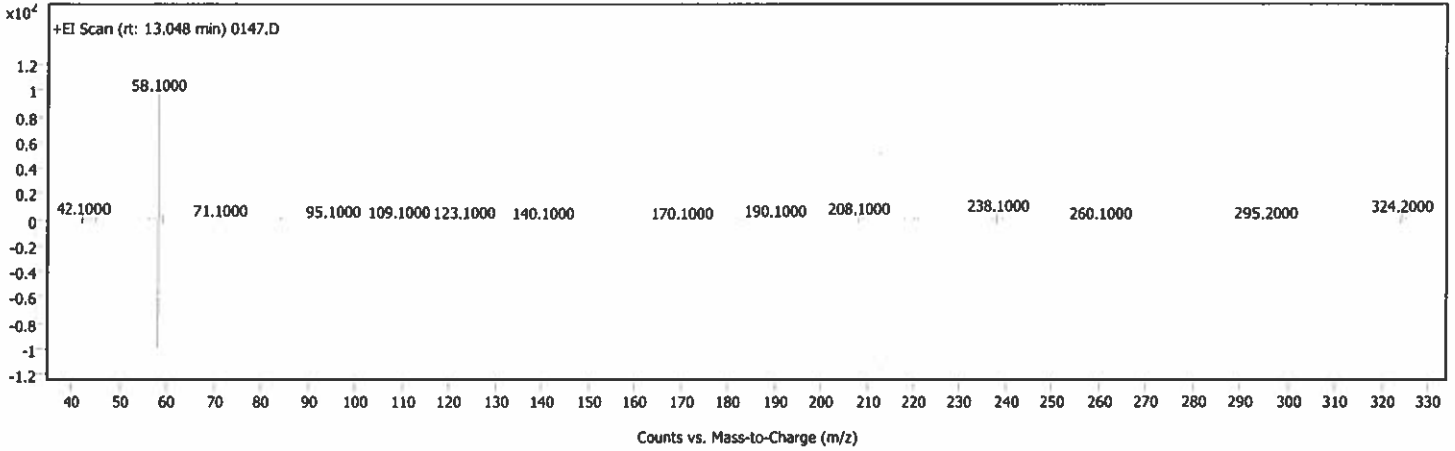
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	96.68	aafs.l	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

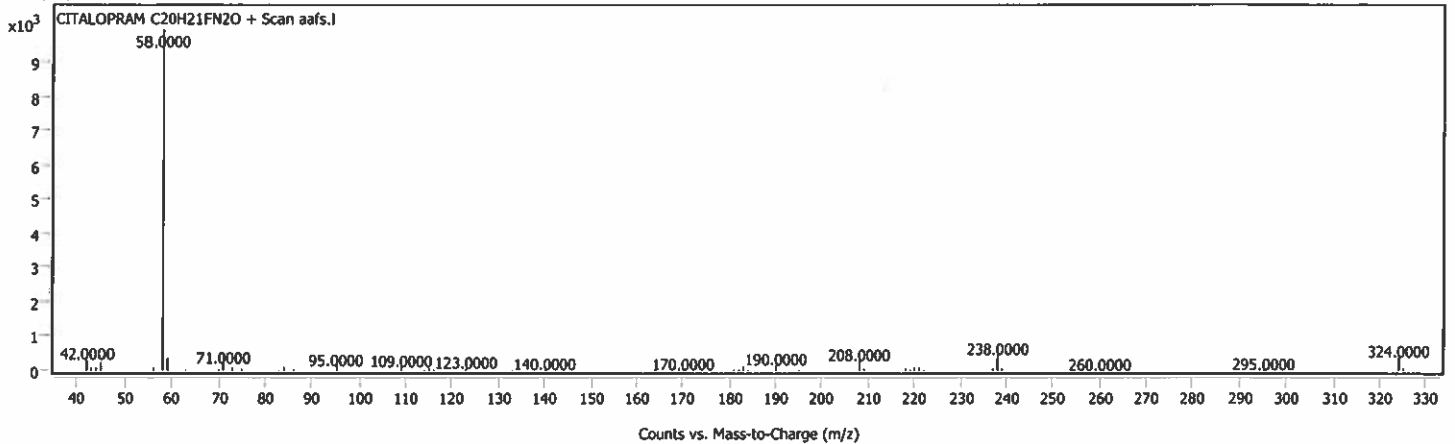
## Observed Spectrum



## Mirror Plot



## Library Spectrum



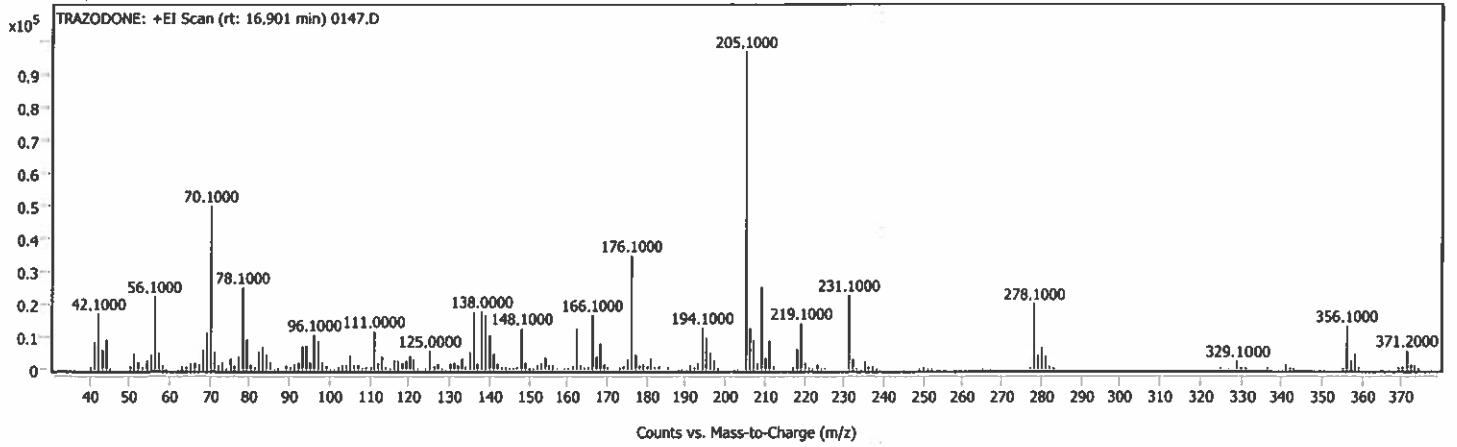
# Analysis Report

+ Scan (rt: 16.901 min)

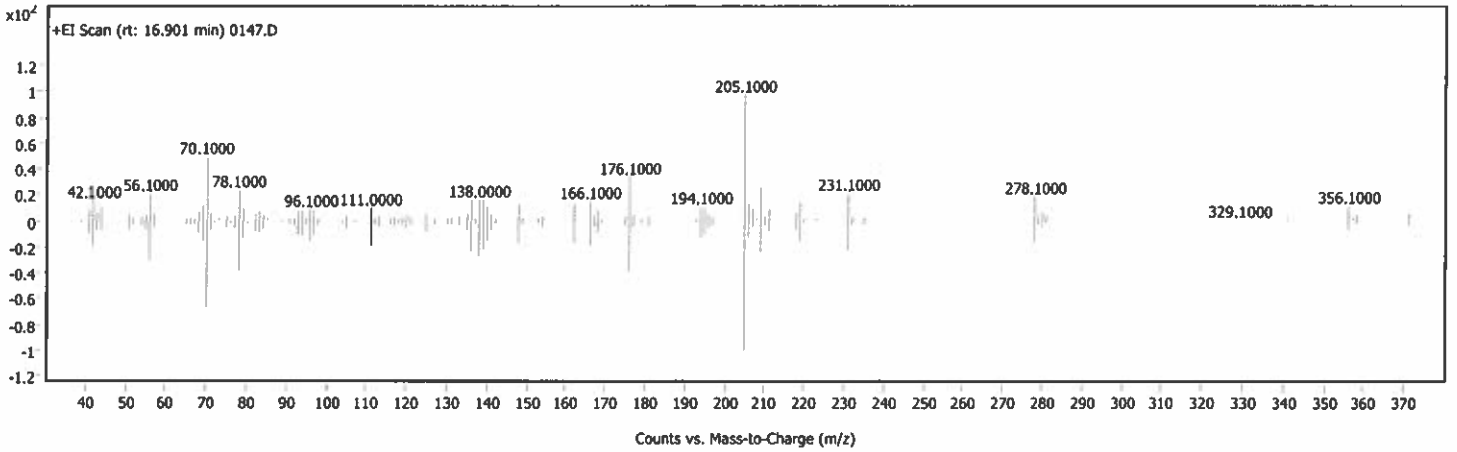
TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		95.40	aafs.I	MATRIX 1 W/ SW CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

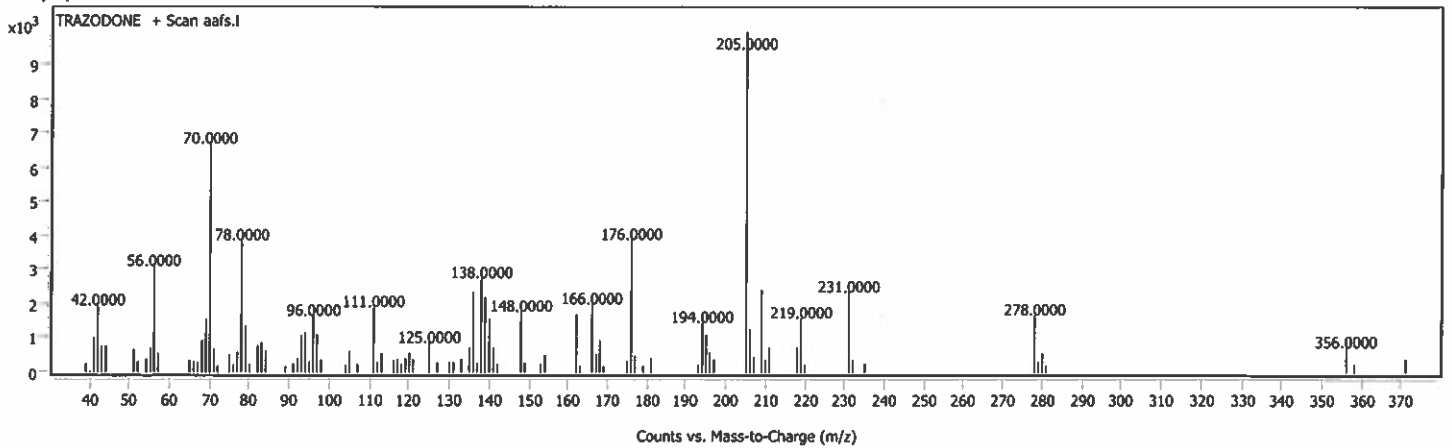
### Observed Spectrum



### Mirror Plot

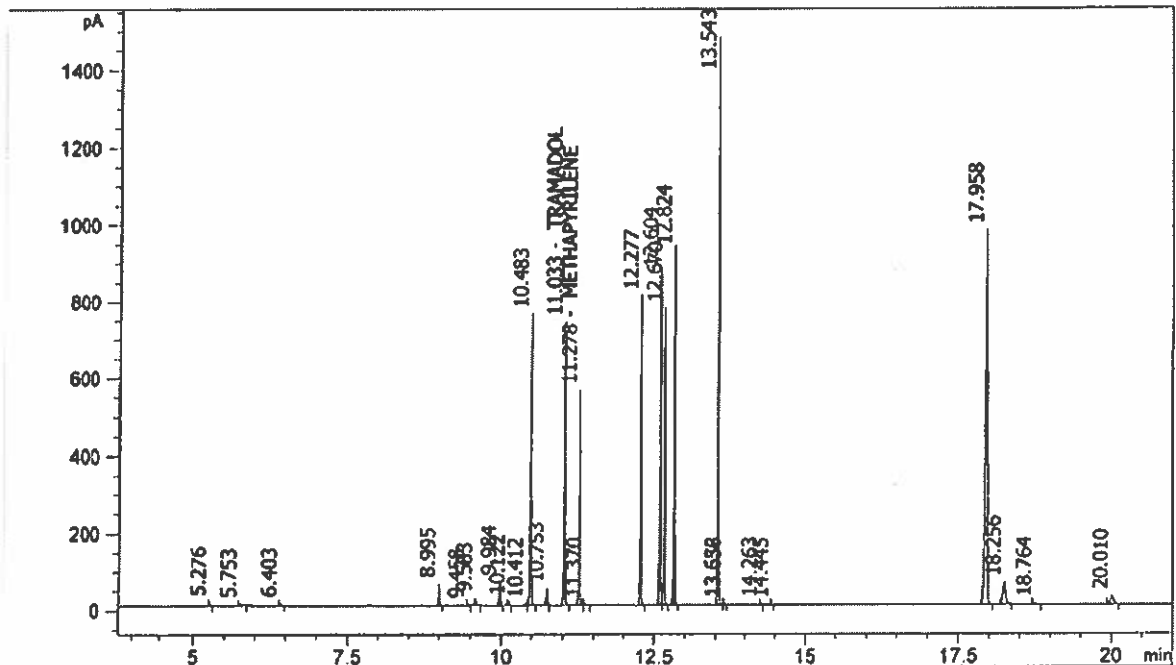


### Library Spectrum



Injection Date: 9/28/2021 8:38:50 PM Seq Line: 18  
 Sample Name: MATRIX 1 W/ SW CTL - -> Vial 19  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.033	11.033	911.867	733.780	0.000000	TRAMADOL RRT - 0.9783
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	648.027	555.823	1.000000	METHAPYRILENE

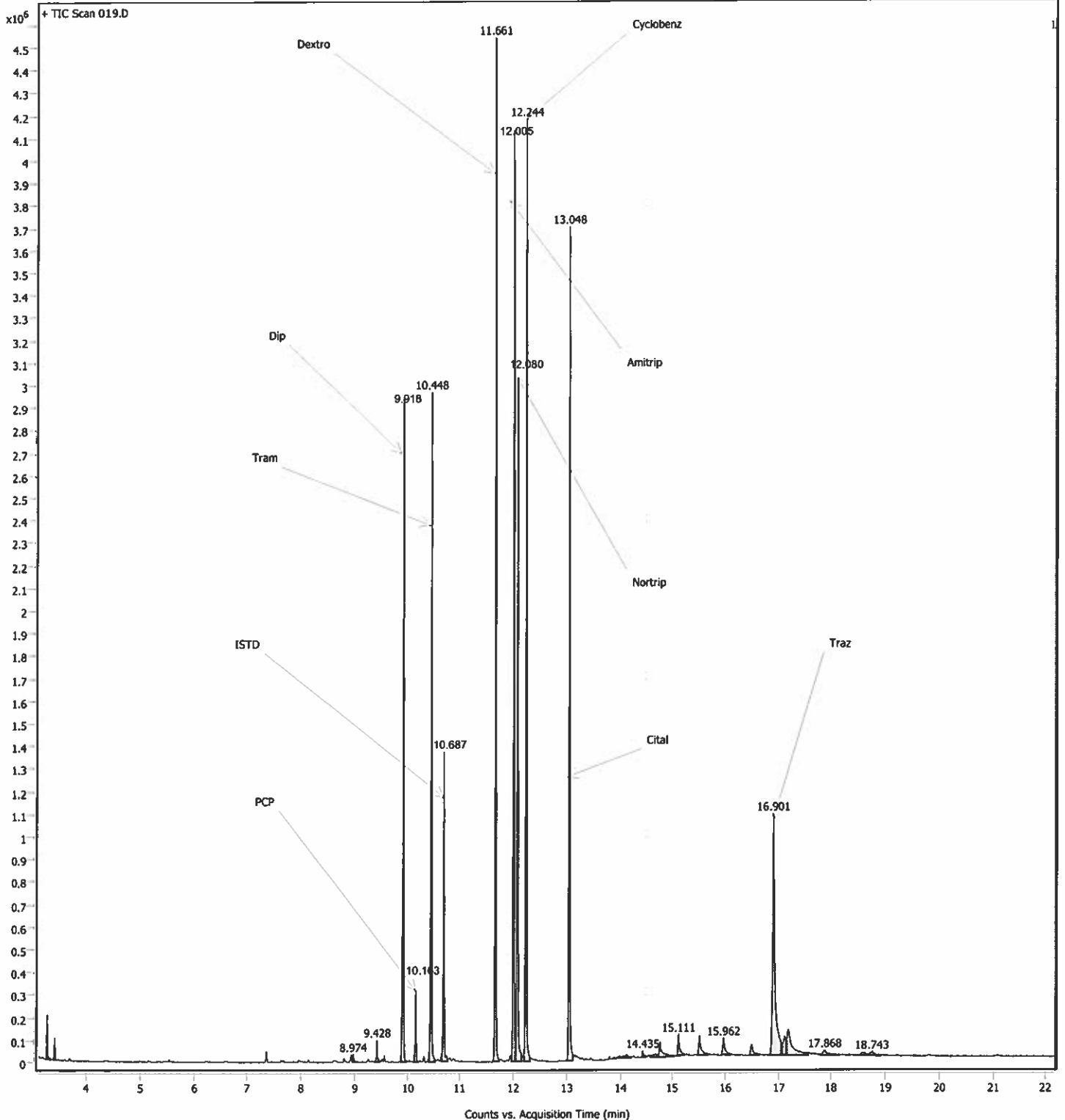
\*RT for Tramadol set from sample

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 W/ SW CTL - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASE5\092821\019.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 9:25:34 PM (UTC-04:00)
Position	45	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

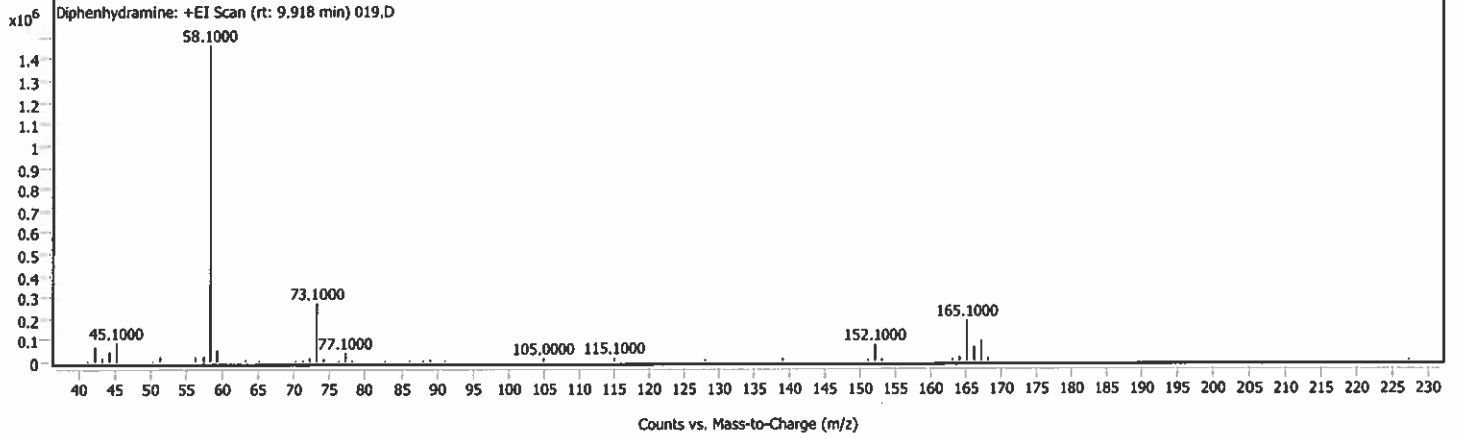
## Sample Spectra

+ Scan (rt: 9.918 min)

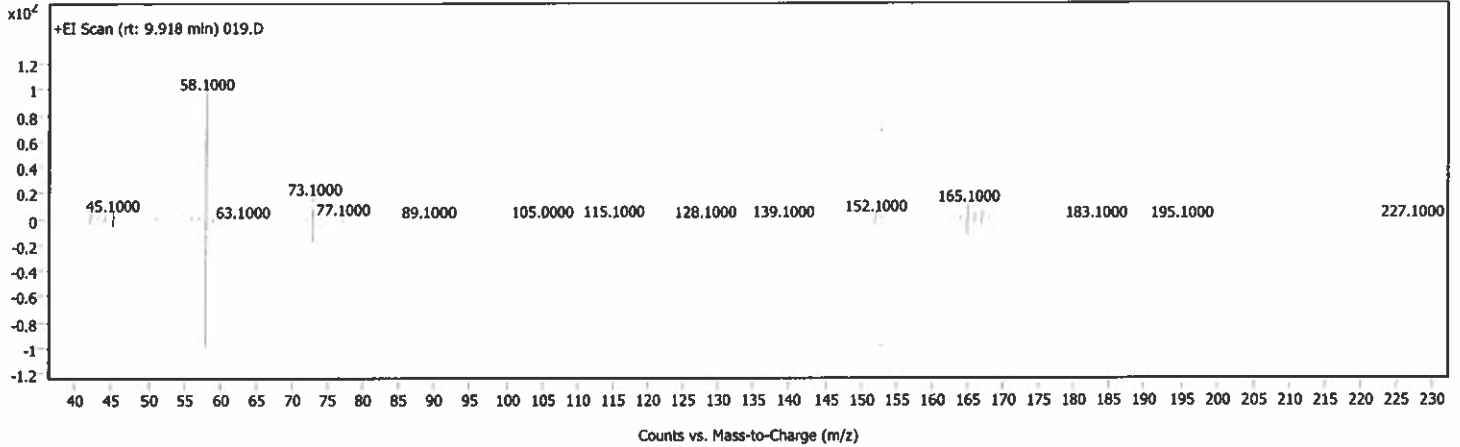
Diphenhydramine; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Diphenhydramine	C17H21NO	98.31	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

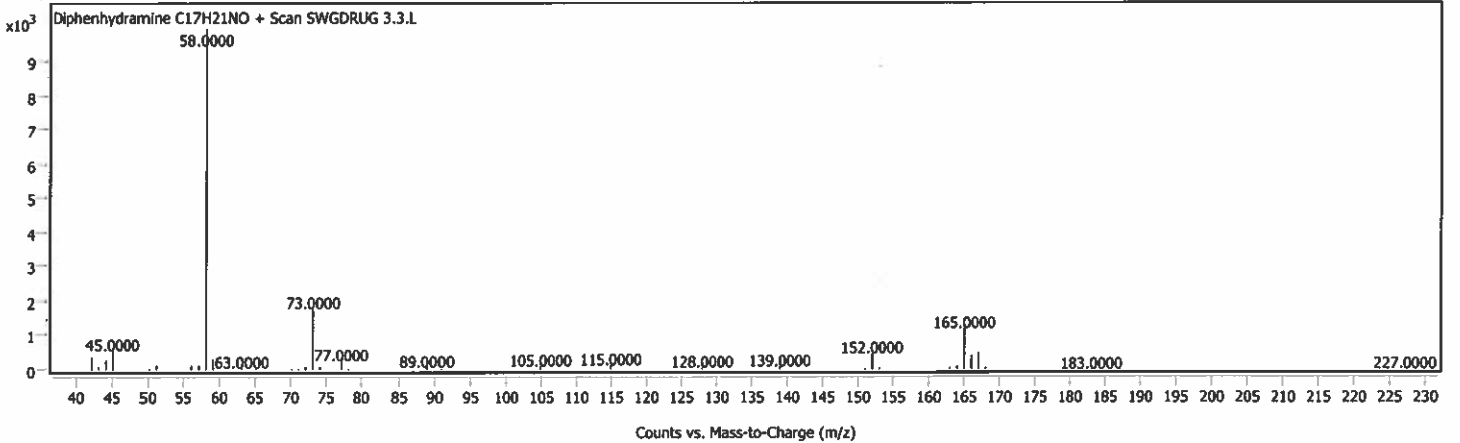
## Observed Spectrum



## Mirror Plot



## Library Spectrum



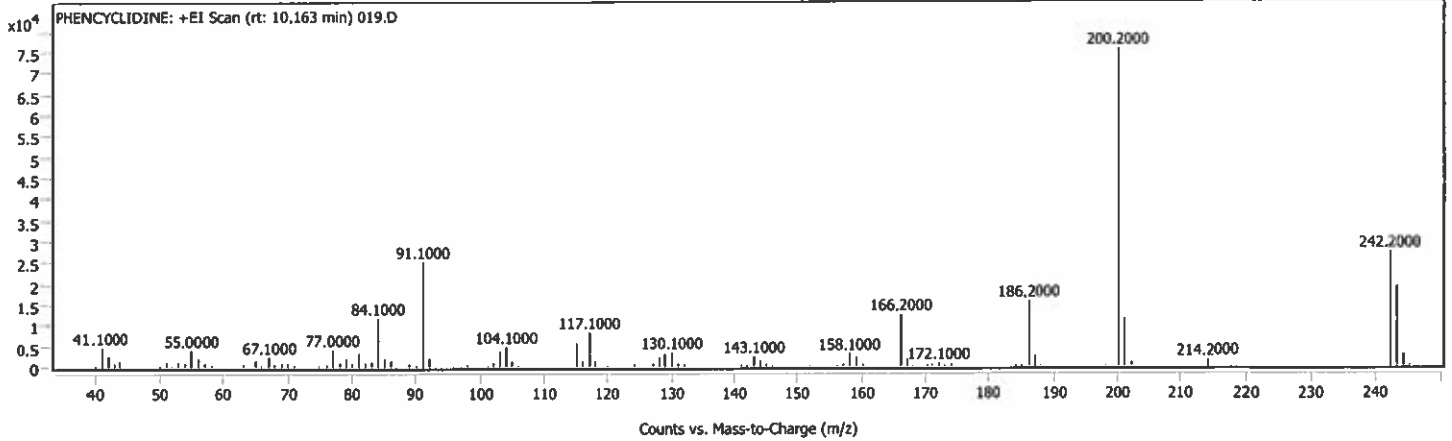
# Analysis Report

+ Scan (rt: 10.163 min)

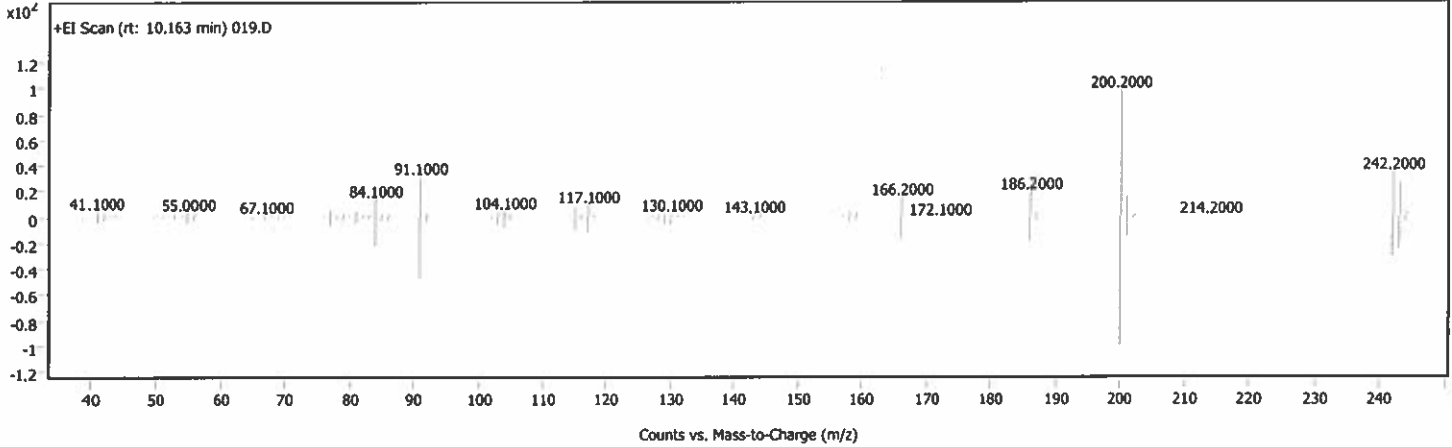
PHENCYCLIDINE; C17H25N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
PHENCYCLIDINE	C17H25N	98.50	aafs.I	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

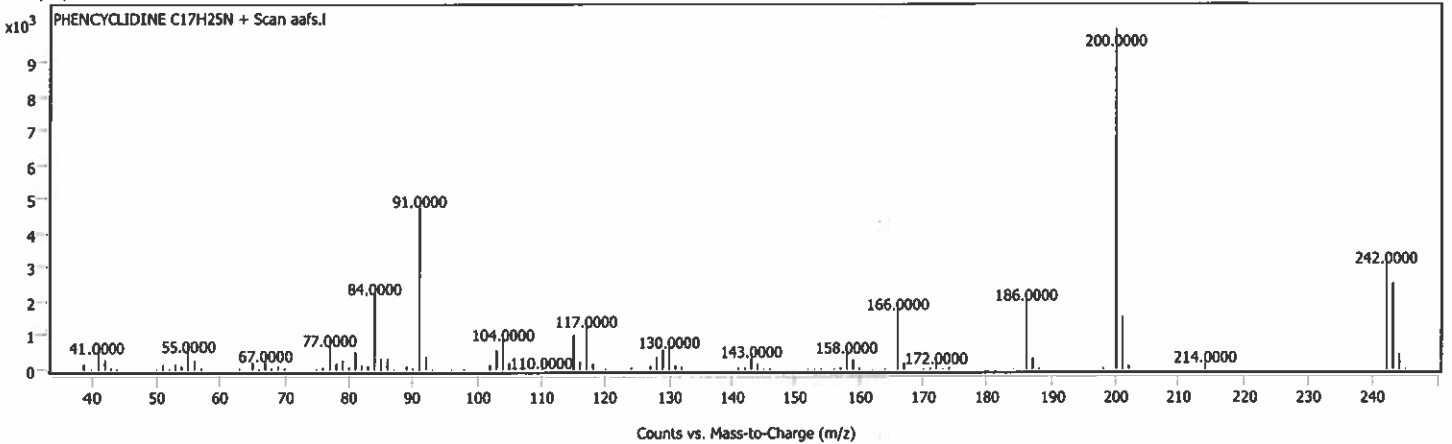
## Observed Spectrum



## Mirror Plot



## Library Spectrum



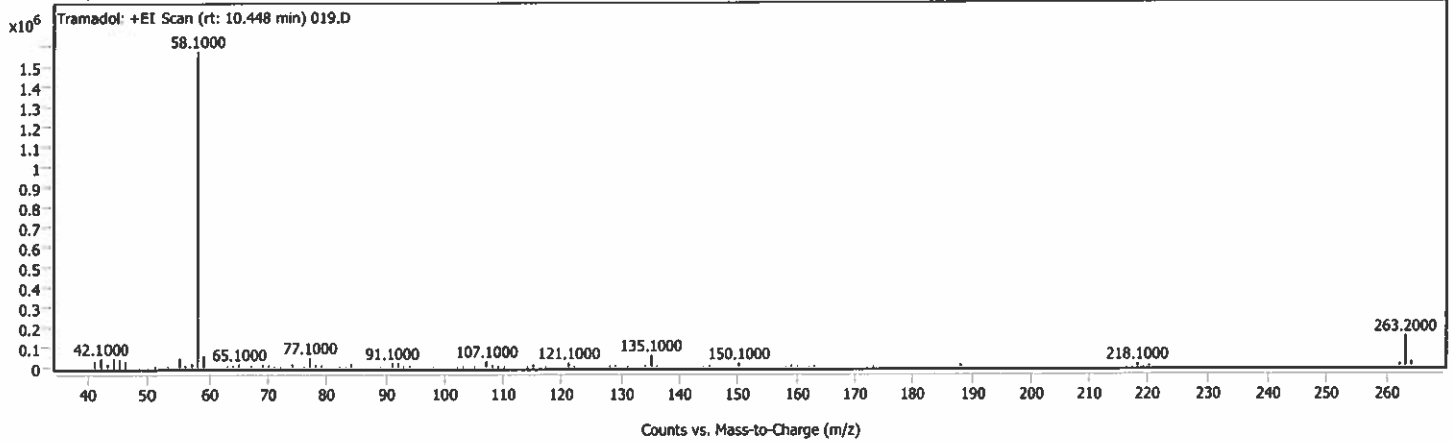
# Analysis Report

+ Scan (rt: 10.448 min)

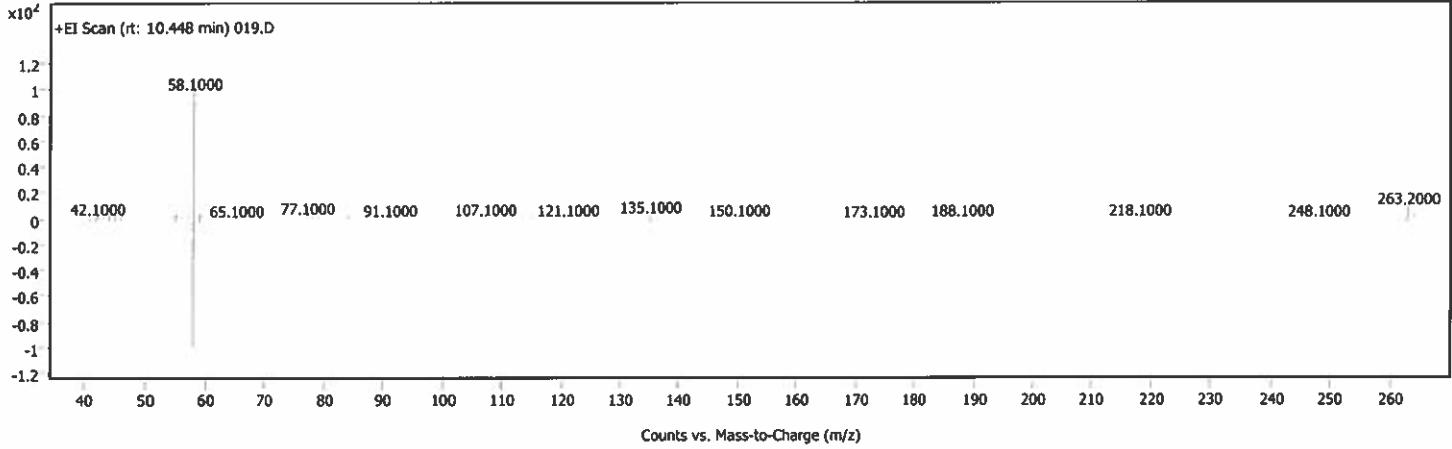
Tramadol; C16H25NO2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Tramadol	C16H25NO2	95.71	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

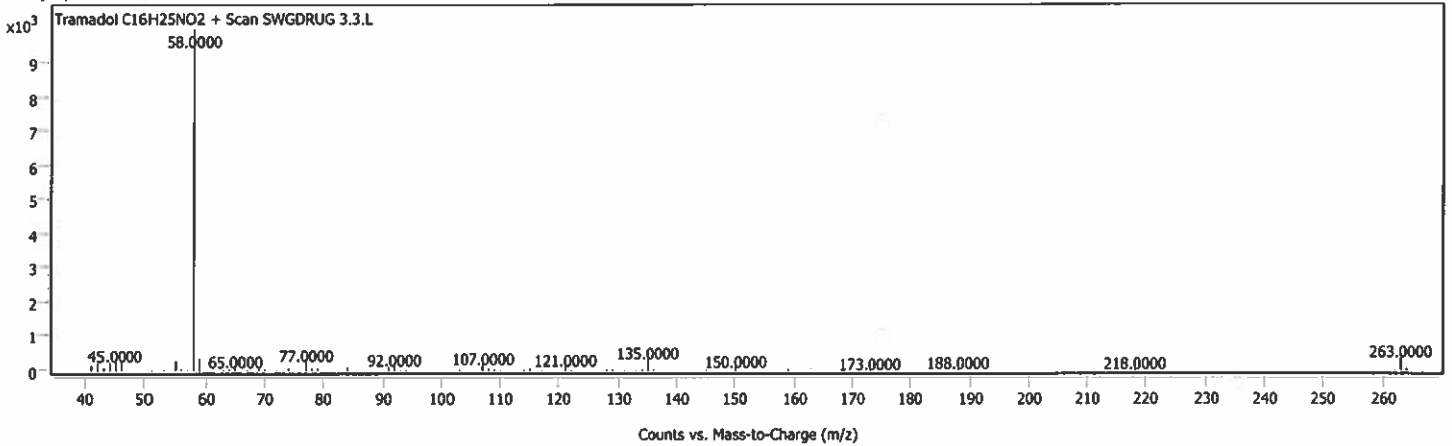
## Observed Spectrum



## Mirror Plot



## Library Spectrum





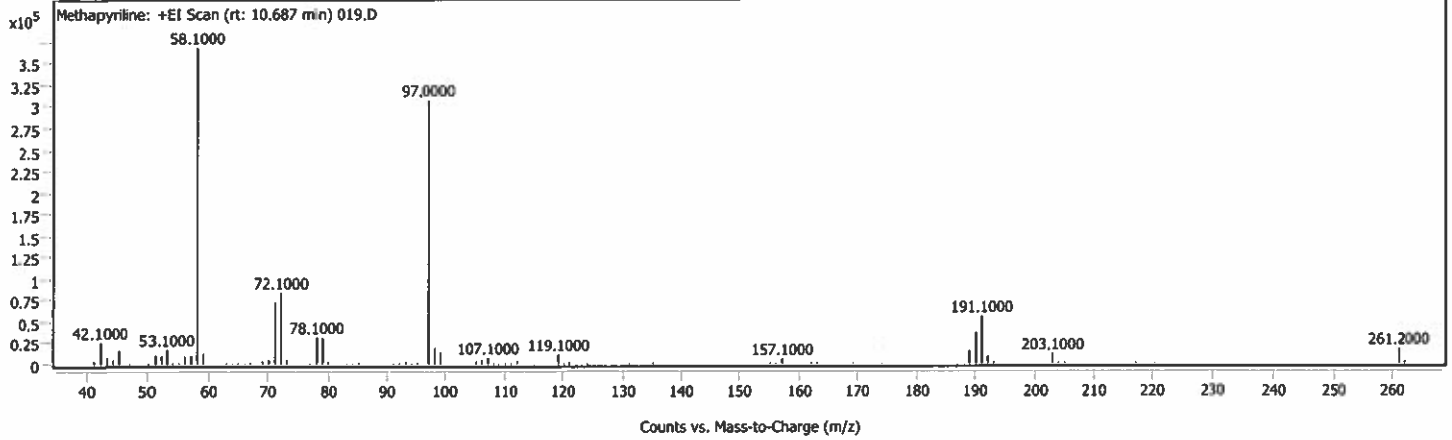
# Analysis Report

+ Scan (rt: 10.687 min)

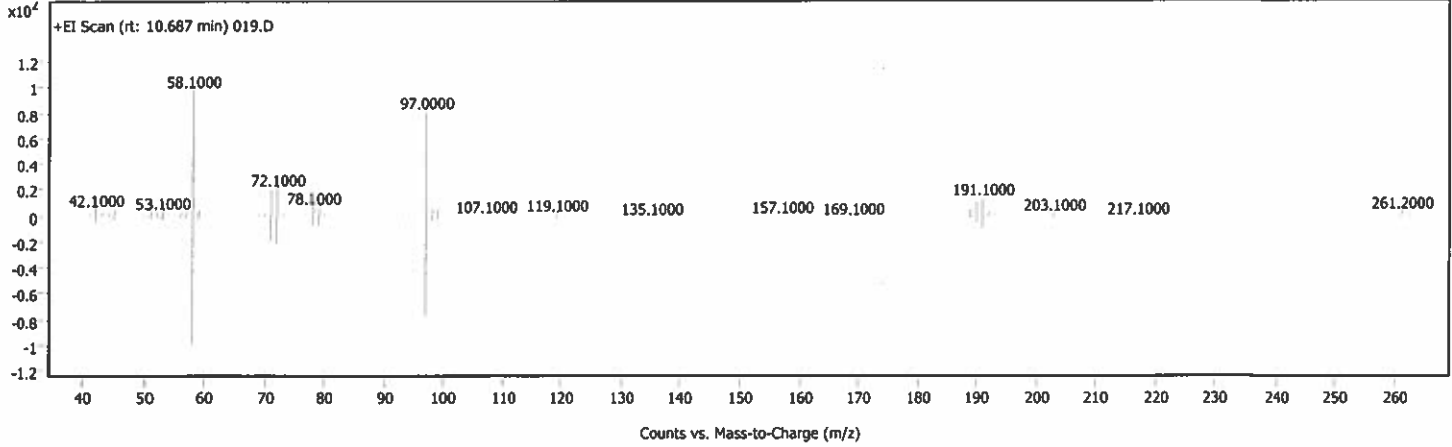
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.37	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XLAZINE VALIDATION

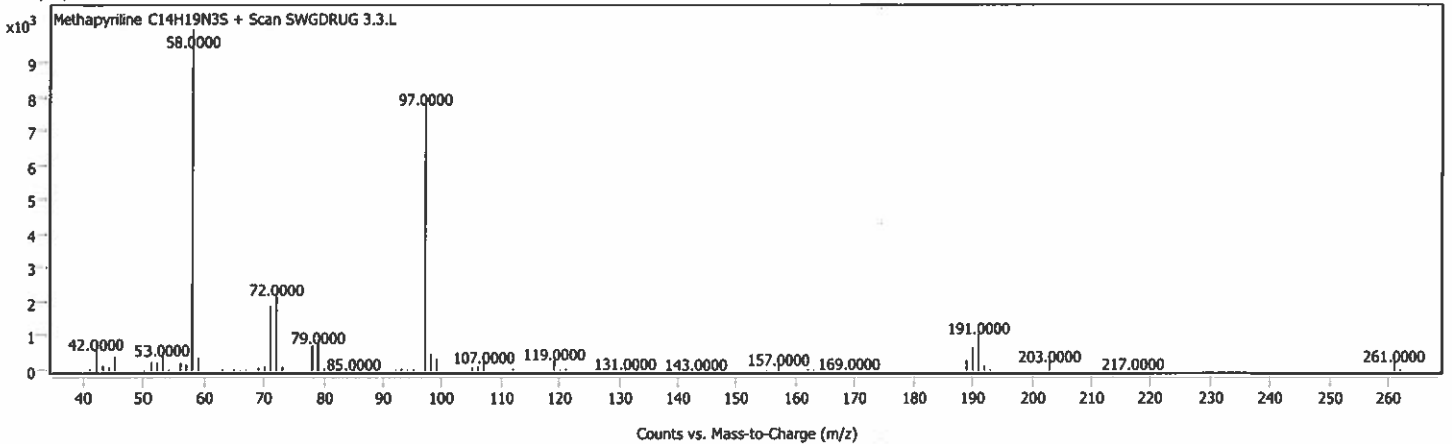
## Observed Spectrum



## Mirror Plot



## Library Spectrum



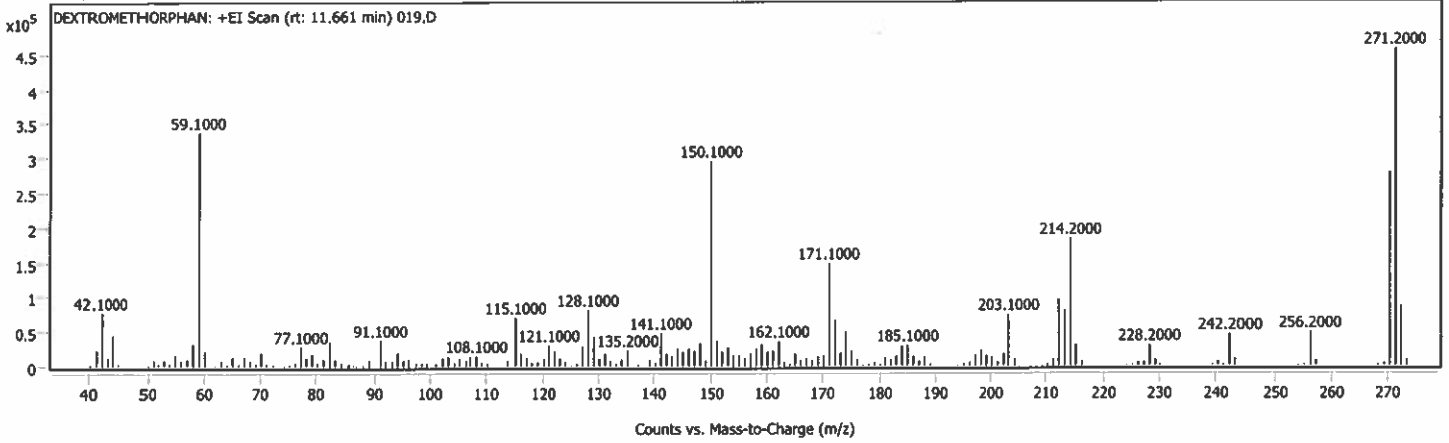
# Analysis Report

+ Scan (rt: 11.661 min)

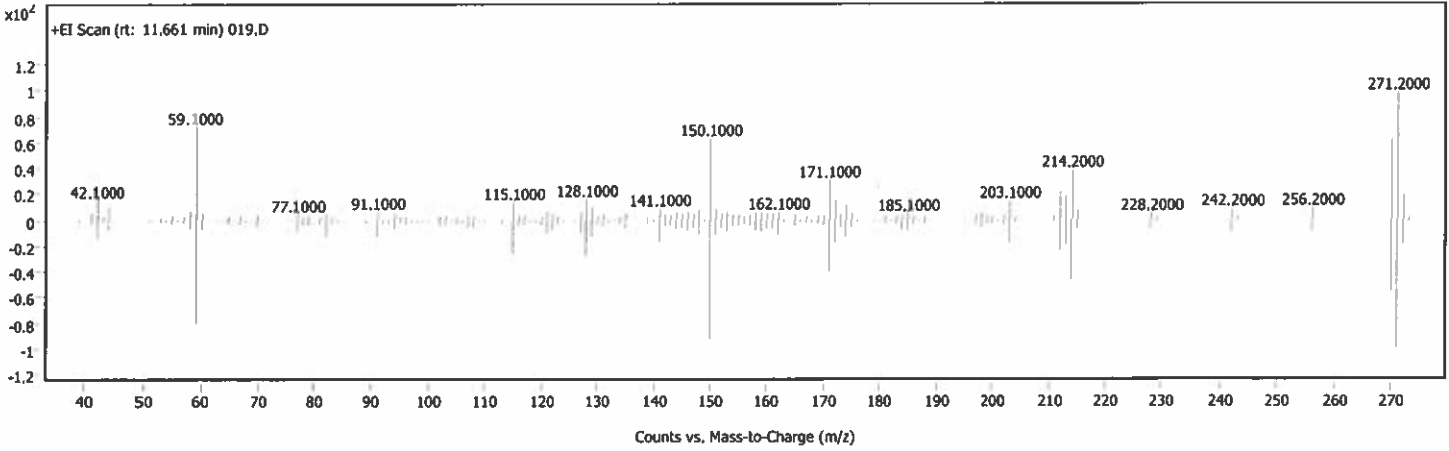
**DEXTROMETHORPHAN; C18H25NO**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	92.32	aafs.I	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

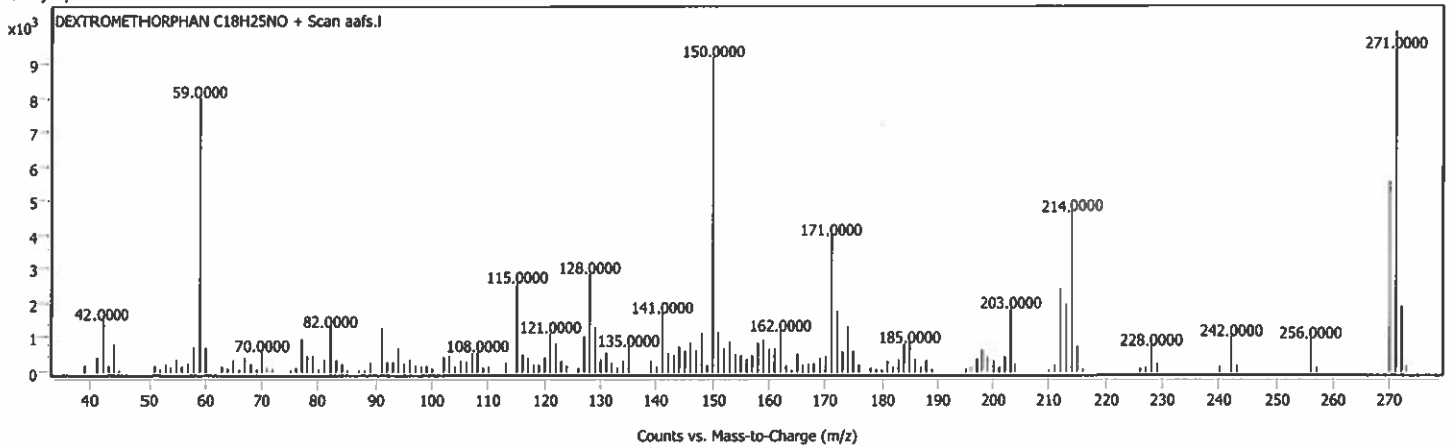
### Observed Spectrum



### Mirror Plot



### Library Spectrum



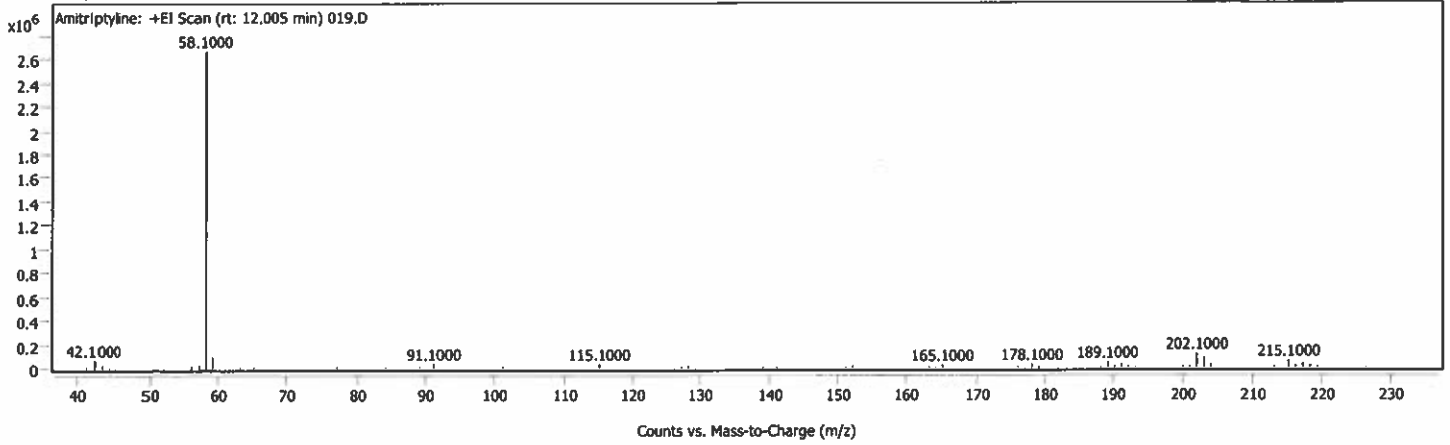
# Analysis Report

+ Scan (rt: 12.005 min)

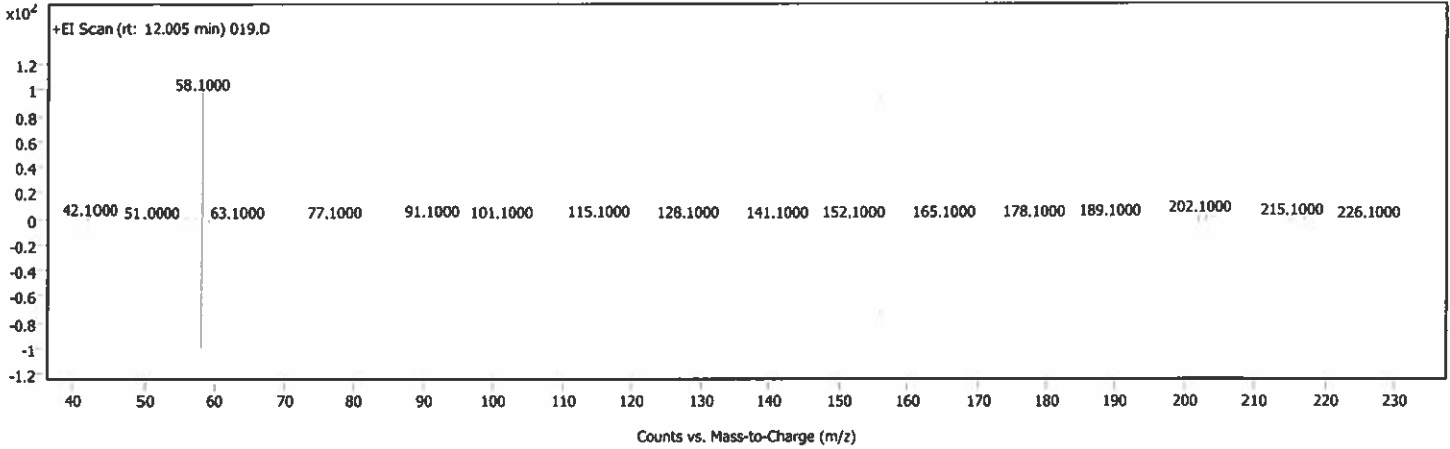
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	98.02	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

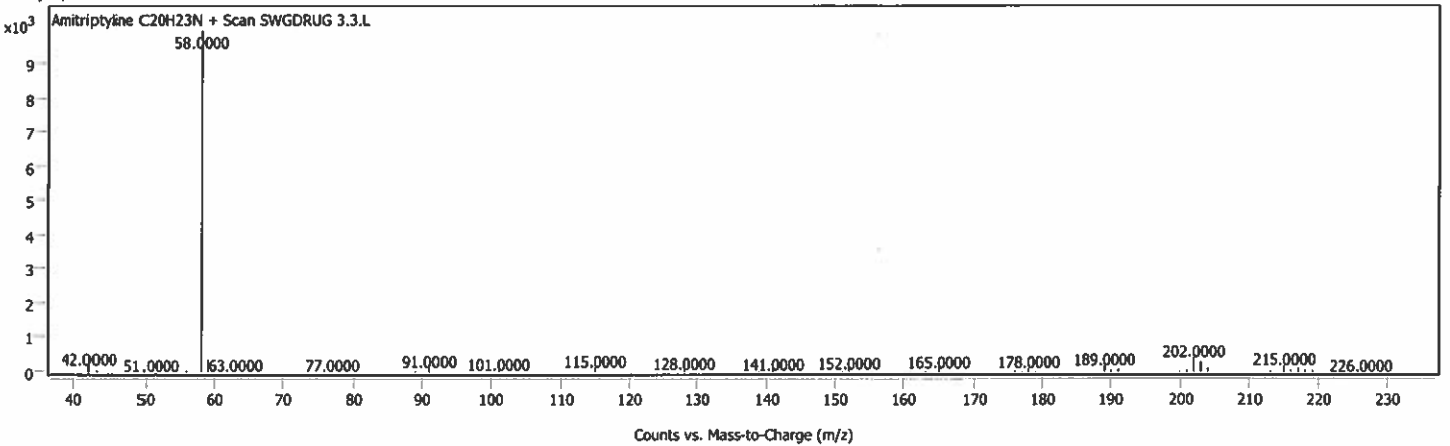
### Observed Spectrum



### Mirror Plot



### Library Spectrum



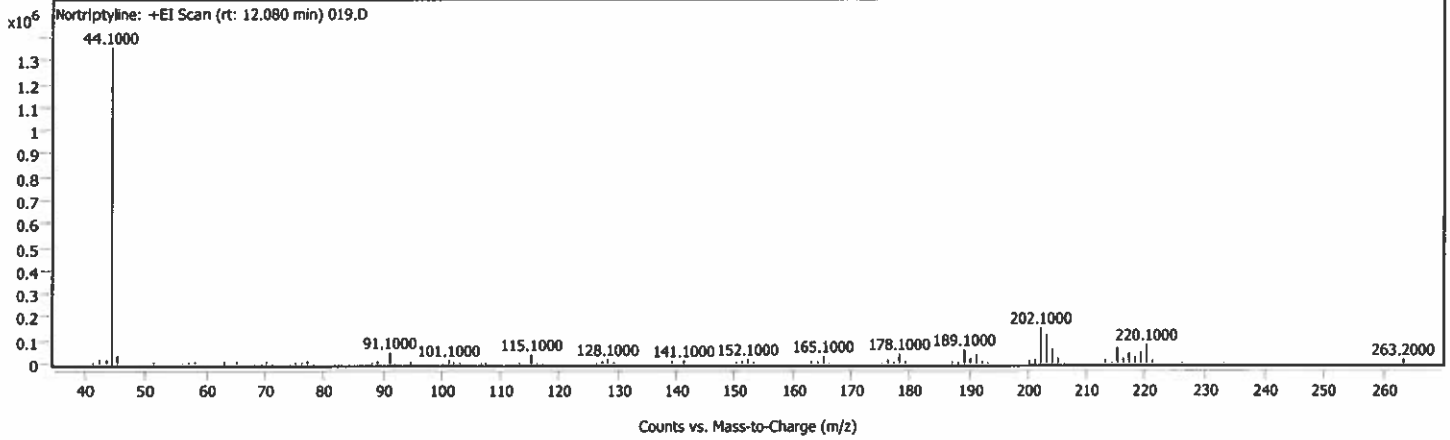
# Analysis Report

+ Scan (rt: 12.080 min)

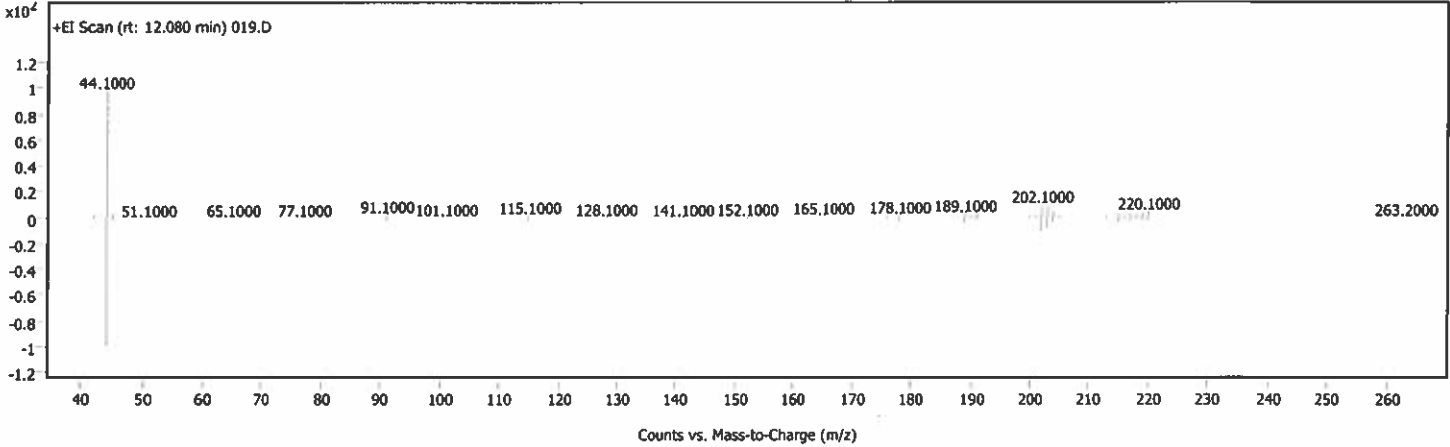
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	97.74	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

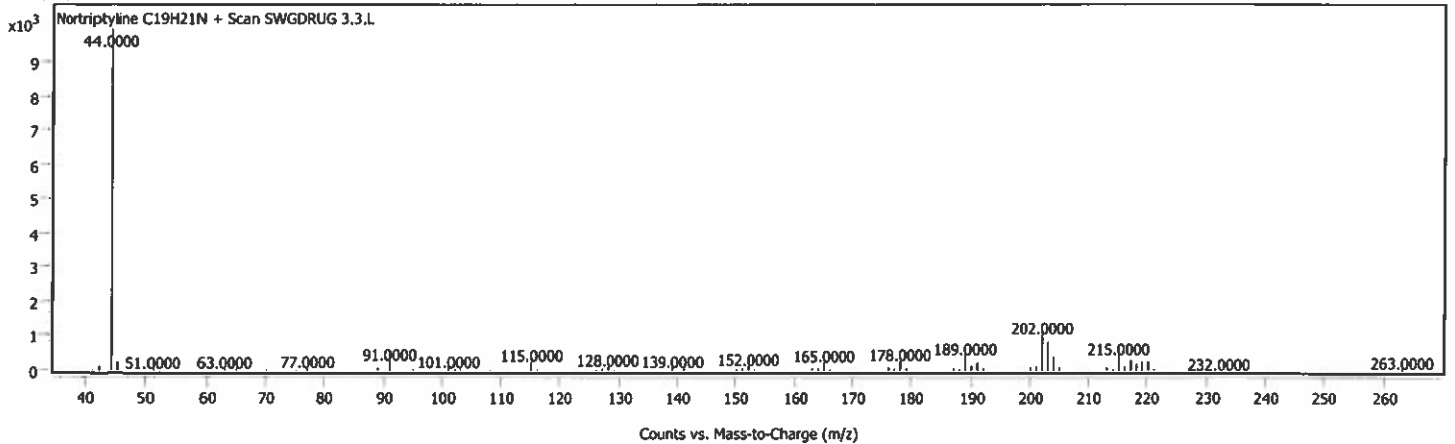
## Observed Spectrum



## Mirror Plot



## Library Spectrum



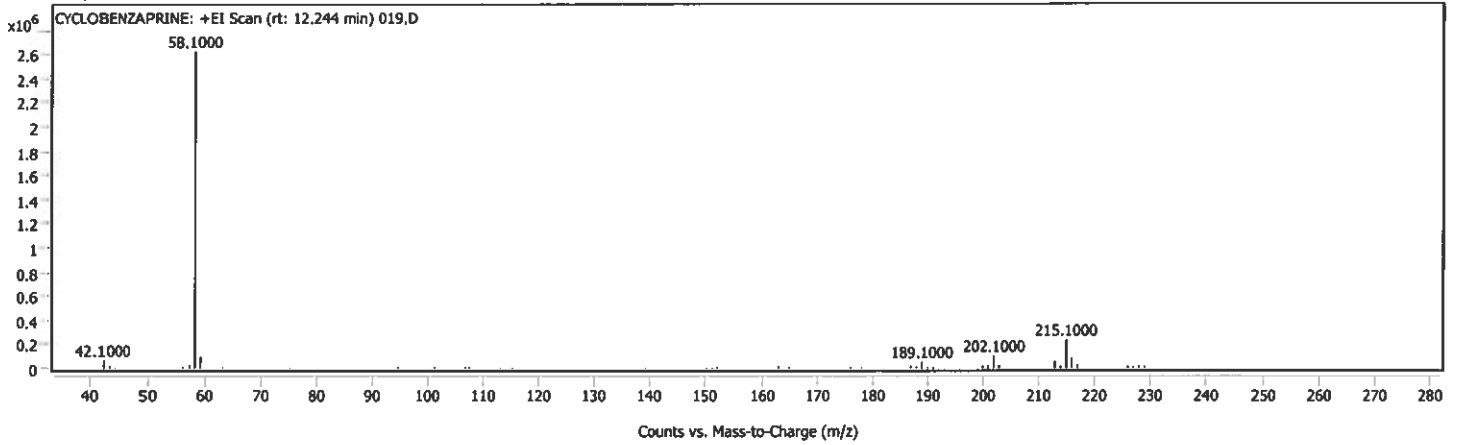
# Analysis Report

+ Scan (rt: 12.244 min)

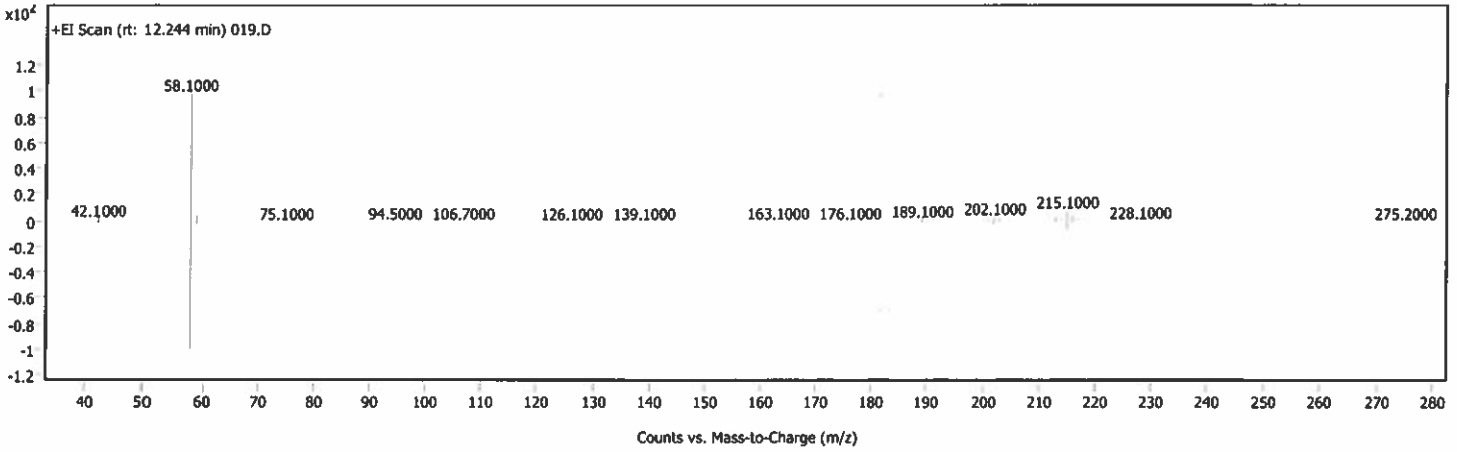
CYCLOBENZAPRINE; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CYCLOBENZAPRINE	C20H21N	96.38	aafs.I	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

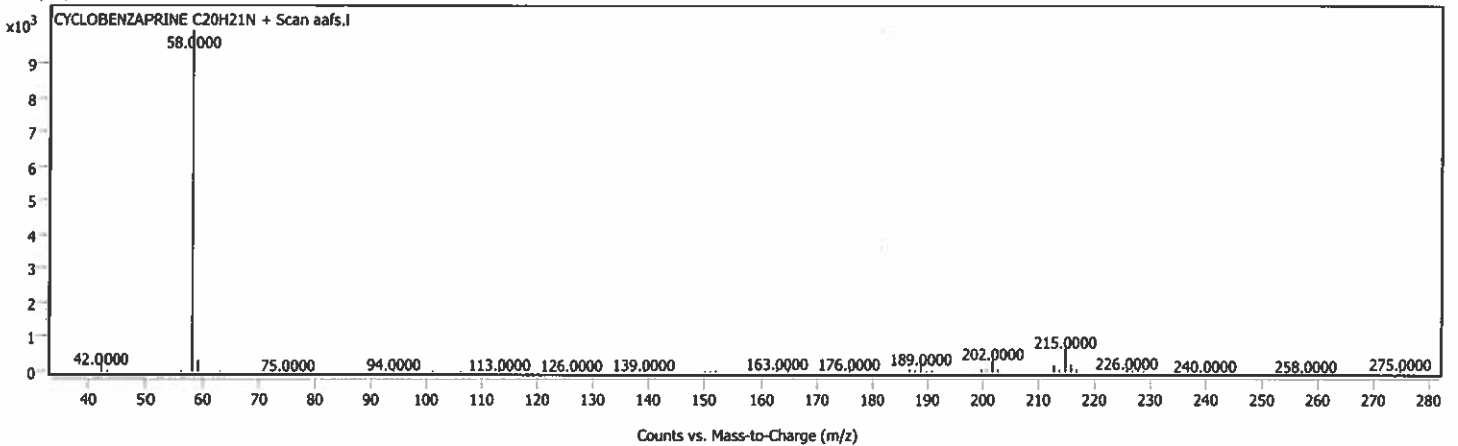
## Observed Spectrum



## Mirror Plot



## Library Spectrum



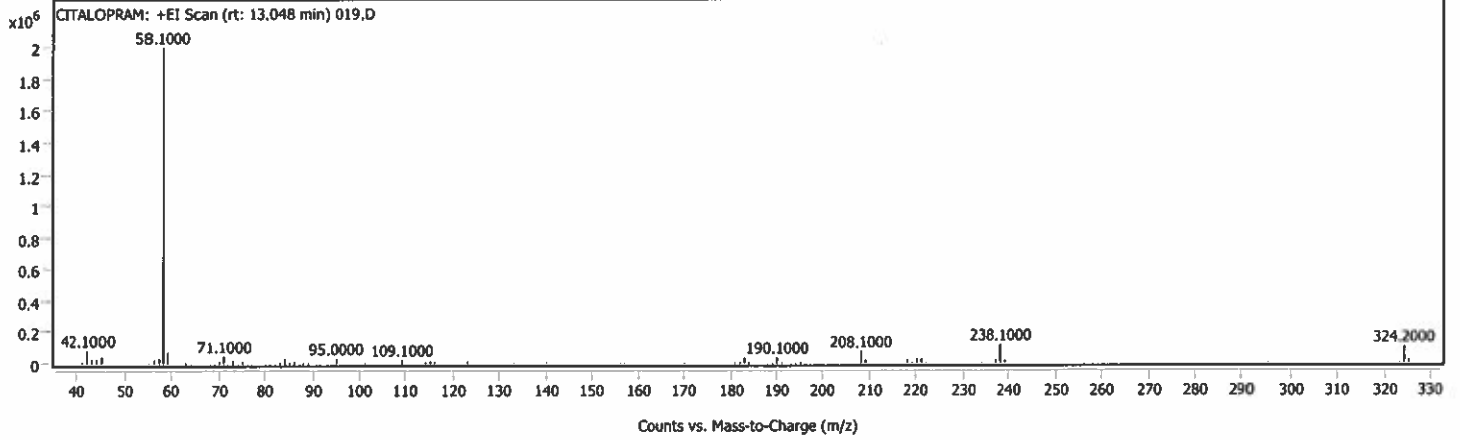
# Analysis Report

+ Scan (rt: 13.048 min)

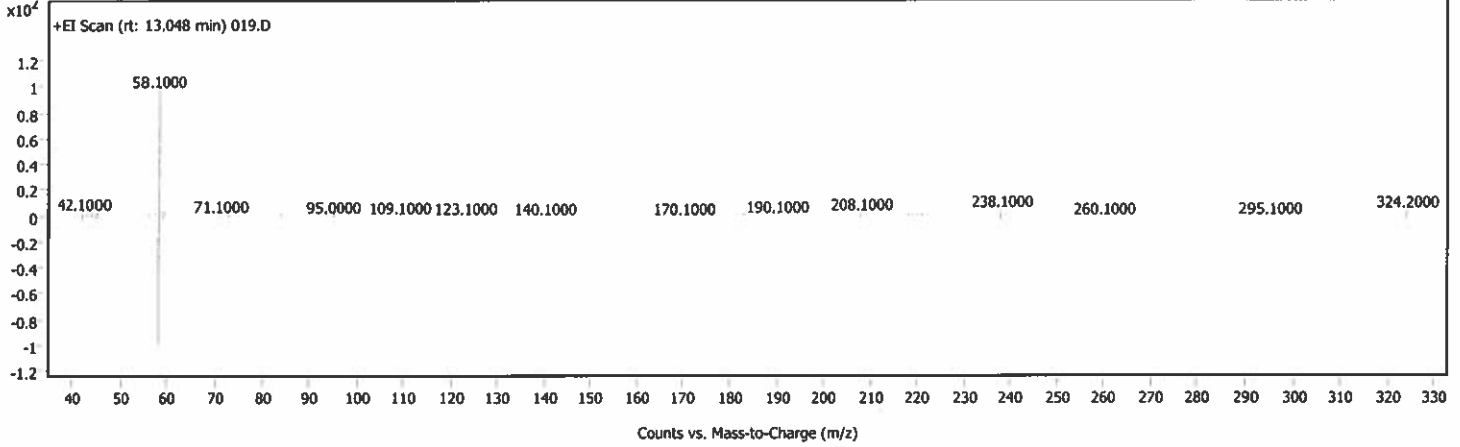
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	96.72	aafs.l	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XLAZINE VALIDATION

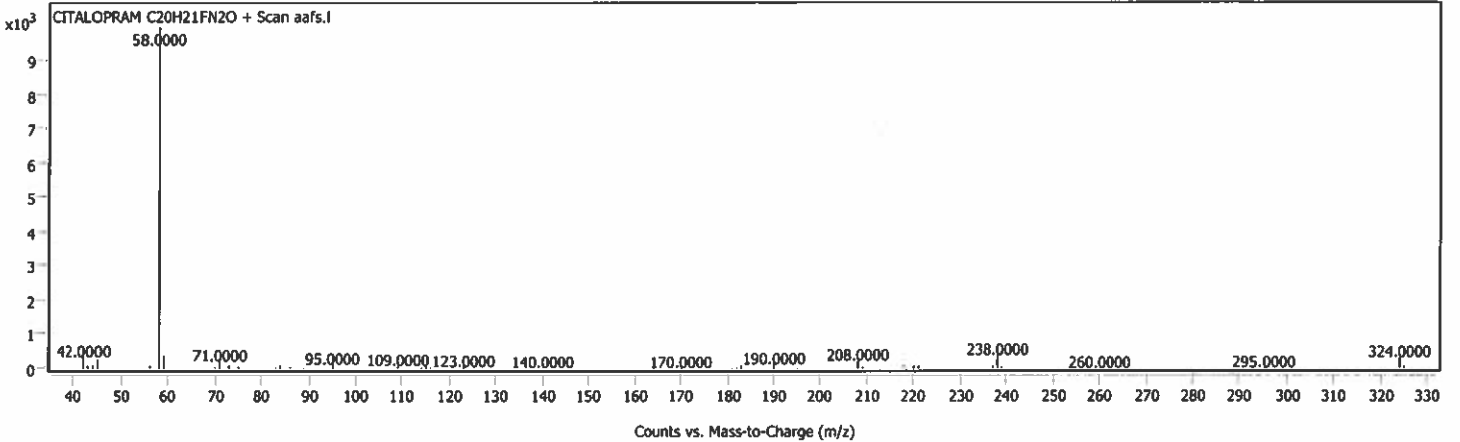
### Observed Spectrum



### Mirror Plot



### Library Spectrum



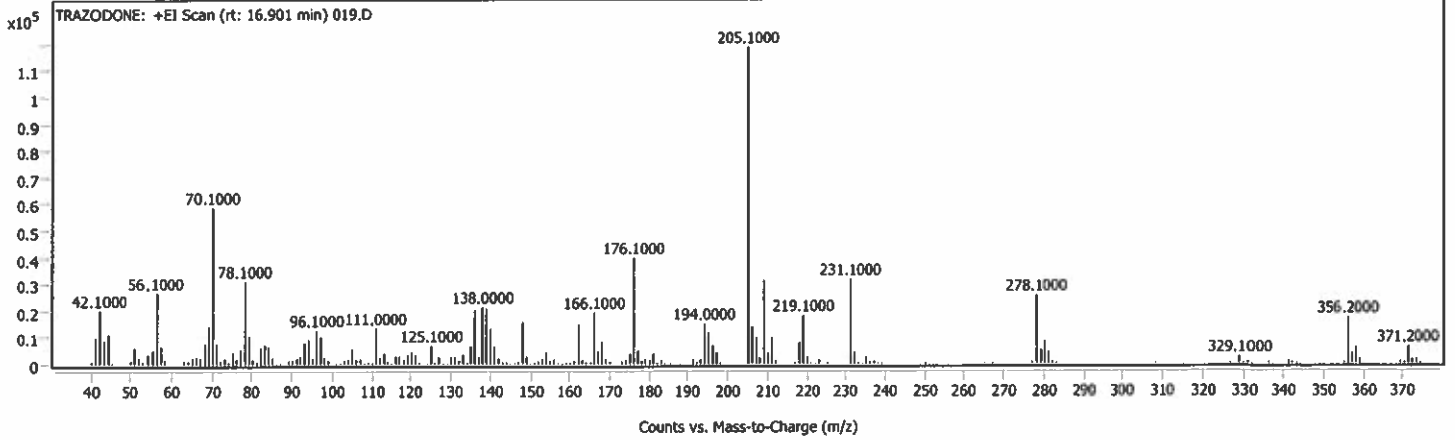
# Analysis Report

+ Scan (rt: 16.901 min)

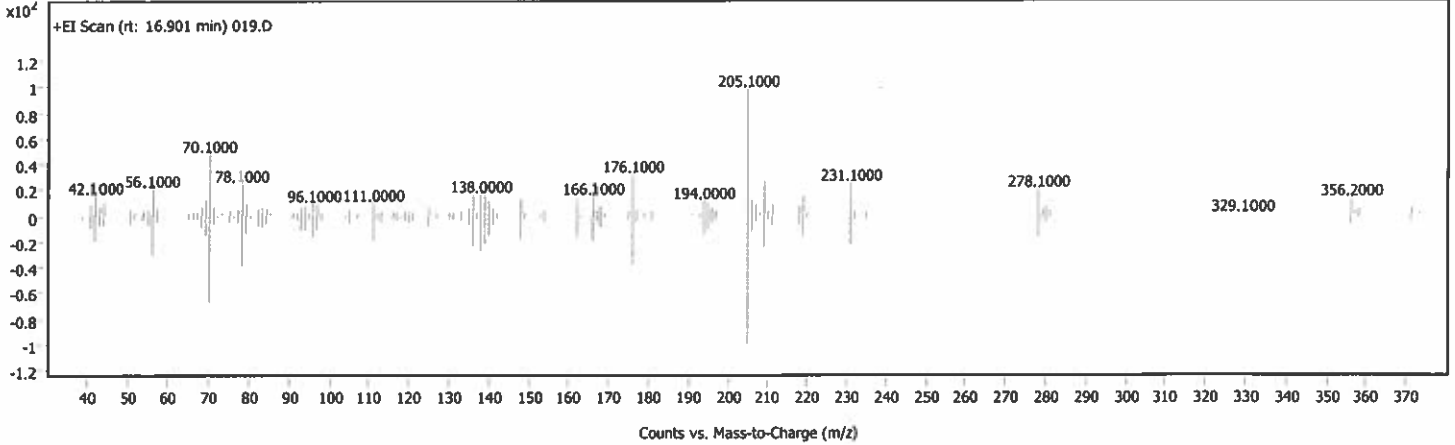
TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		95.29	aafs.I	MATRIX 6 W/ SW CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

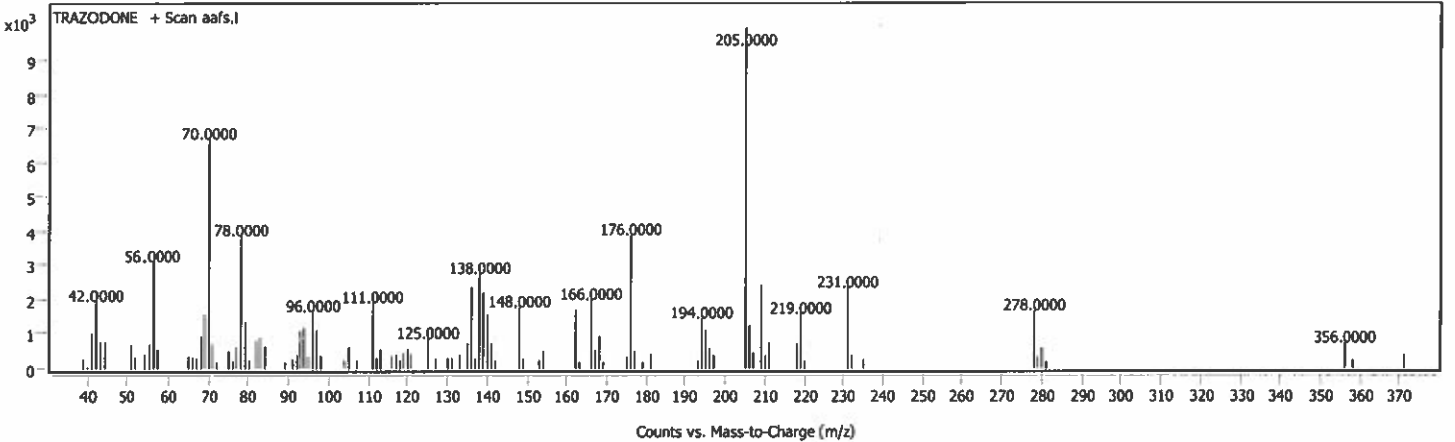
### Observed Spectrum



### Mirror Plot

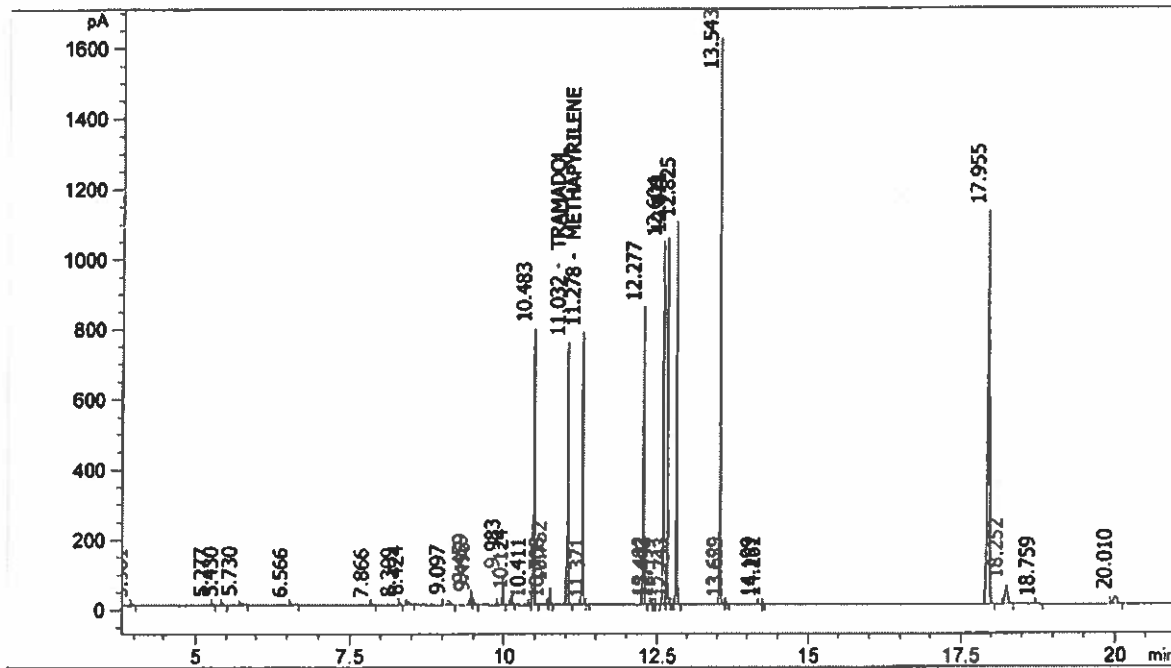


### Library Spectrum



Injection Date: 9/28/2021 9:27:29 PM Seq Line: 20  
 Sample Name: MATRIX 6 W/ SW CTL - -> Vial 21  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.032	11.033	949.633	748.275	0.000000	TRAMADOL RRT - 0.9782
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	901.578	775.774	1.000000	METHAPYRILENE



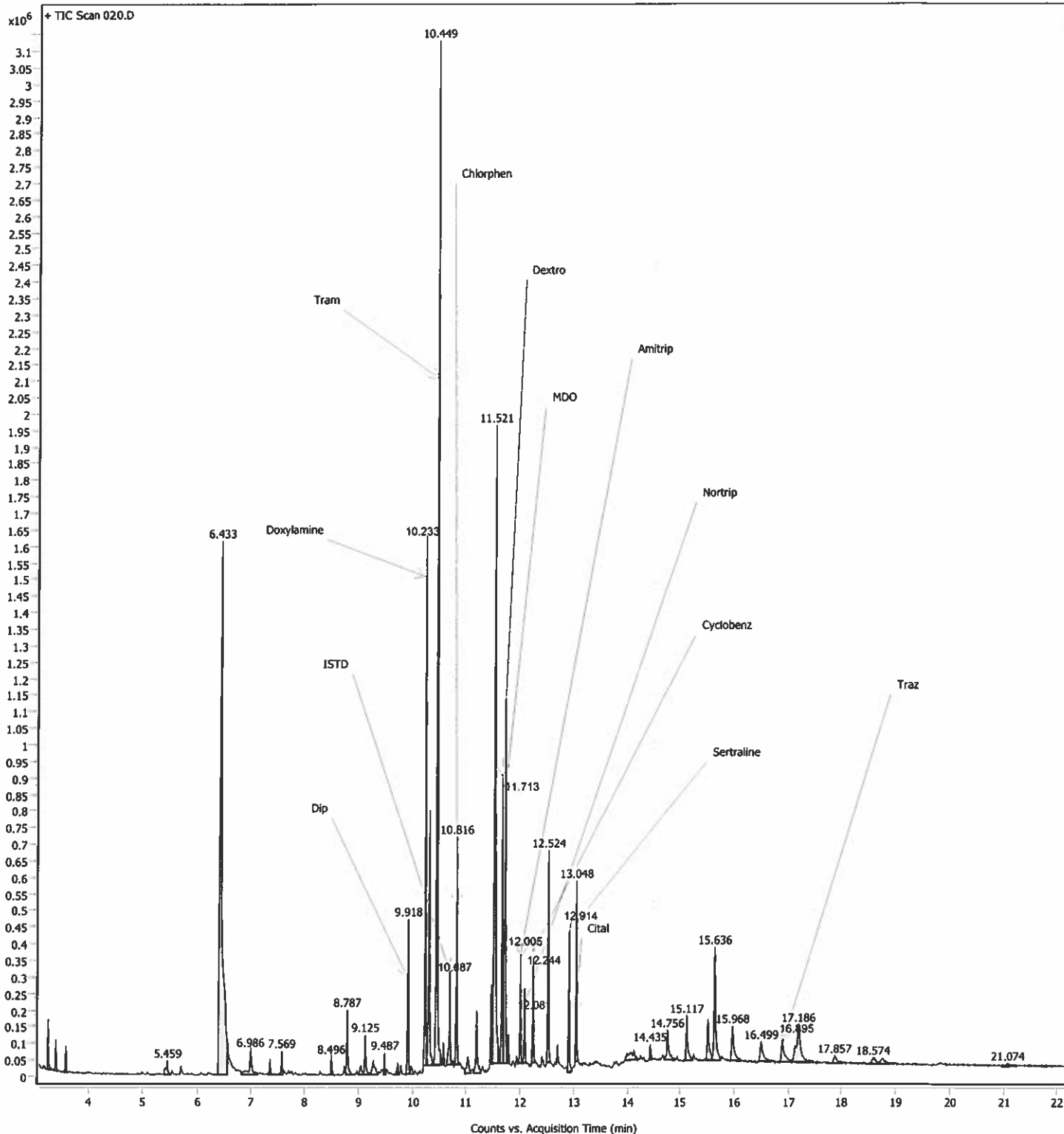
# Analysis Report

## Sample Information

Sample Name	MATRIX 10 W/ 0.01 mg/L CAL - LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\020.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 9:51:21 PM (UTC-04:00)
Position	46	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

2.0 mg/L - 400 ul of 0.01 mg/L KM 4/17/23

## Sample Chromatograms



# Analysis Report

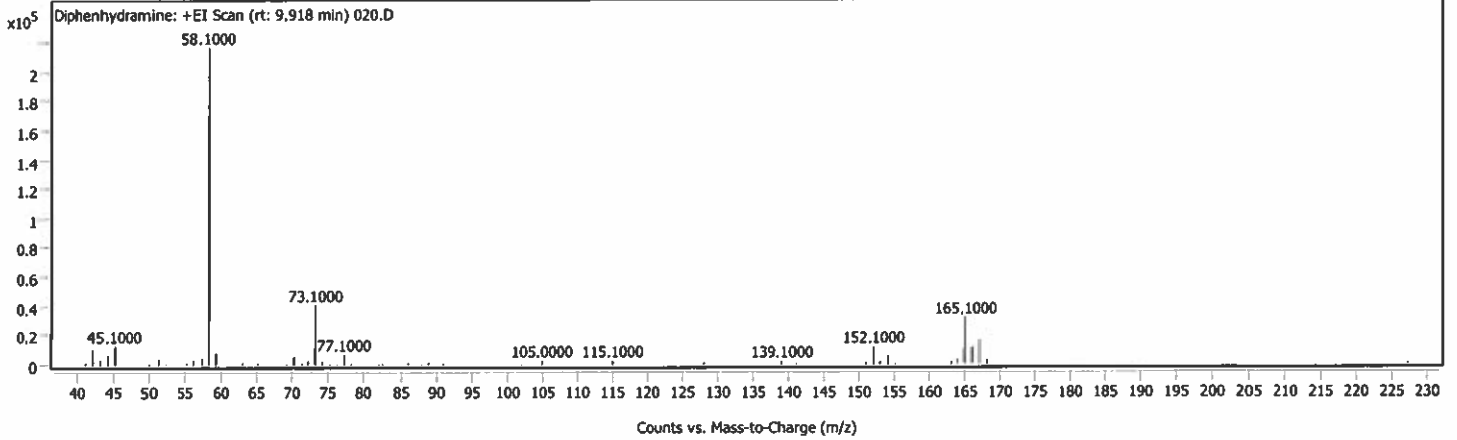
## Sample Spectra

+ Scan (rt: 9.918 min)

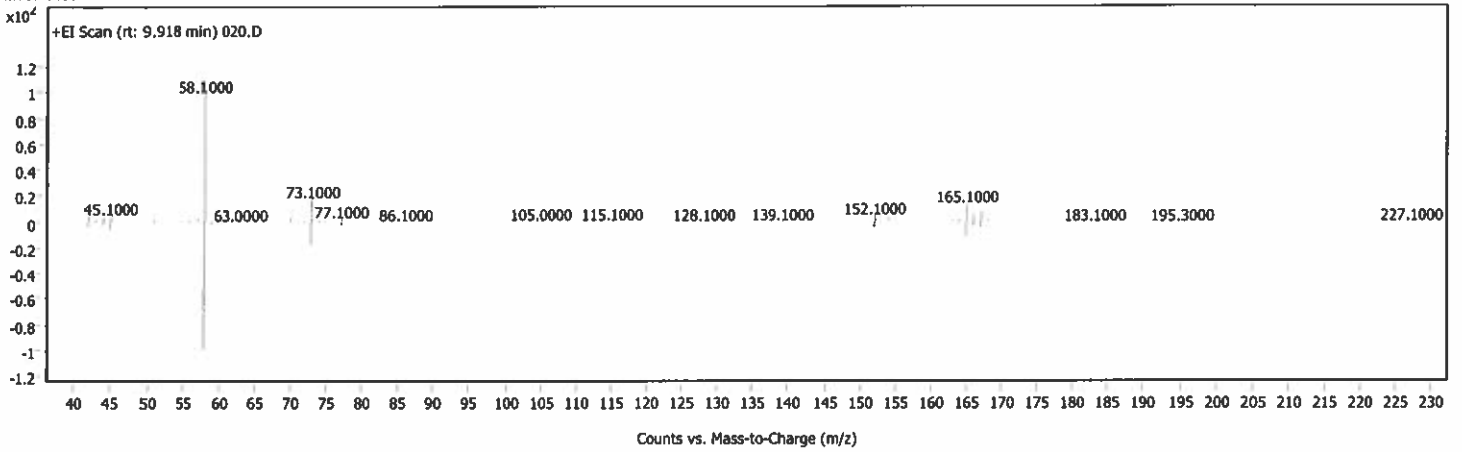
Diphenhydramine; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Diphenhydramine	C17H21NO	95.61	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XLAZINE VALIDATION

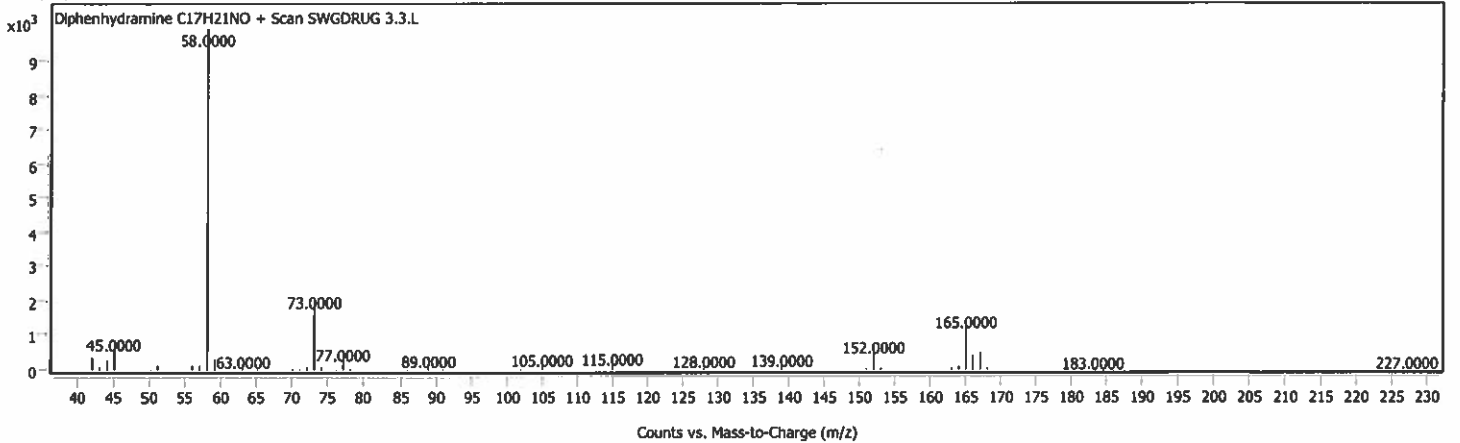
## Observed Spectrum



## Mirror Plot



## Library Spectrum



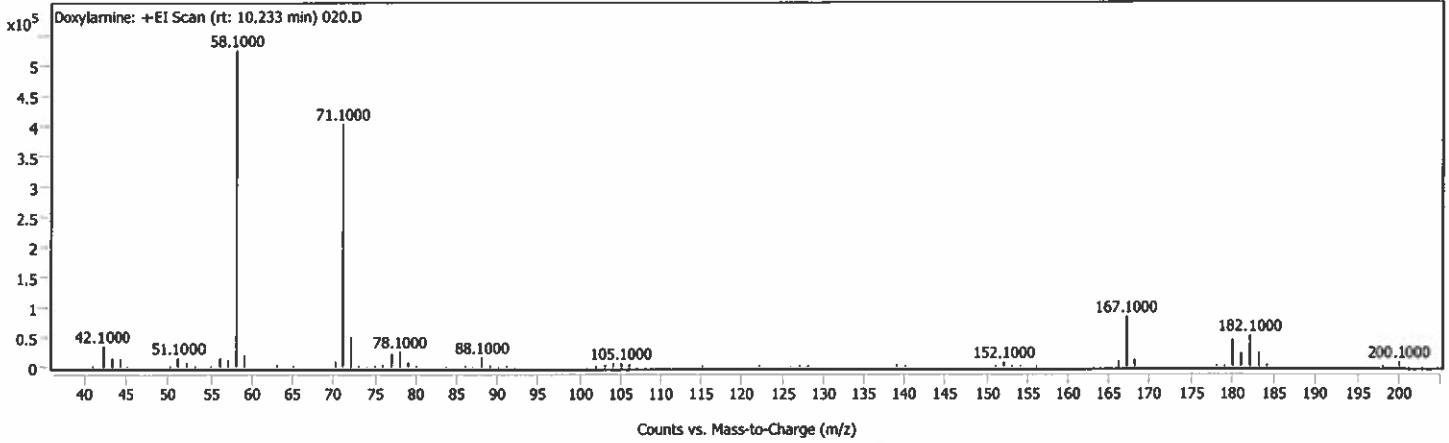
# Analysis Report

+ Scan (rt: 10.233 min)

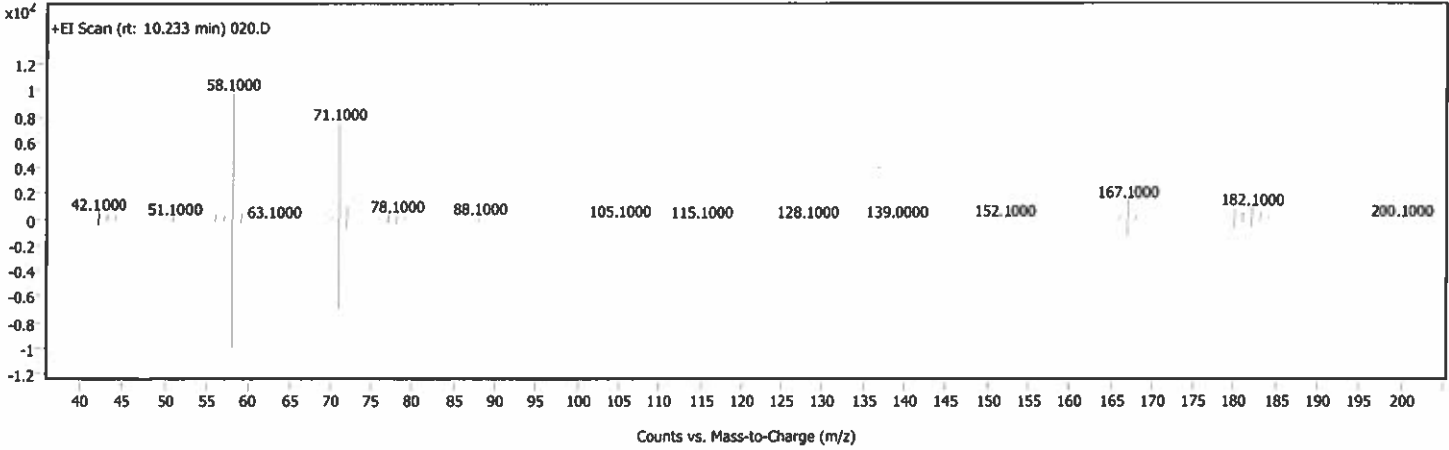
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.83	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

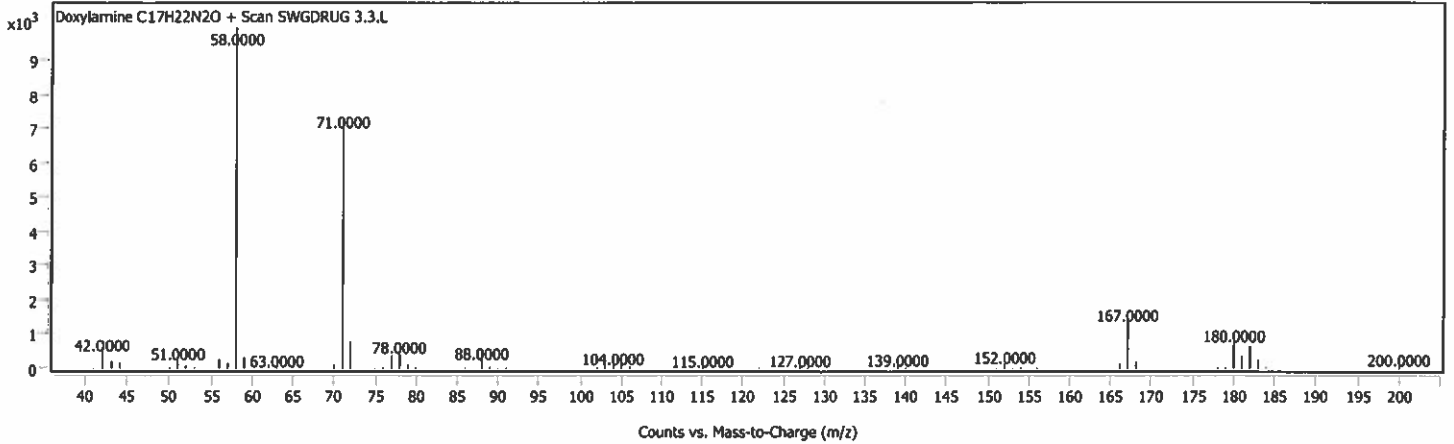
### Observed Spectrum



### Mirror Plot



### Library Spectrum



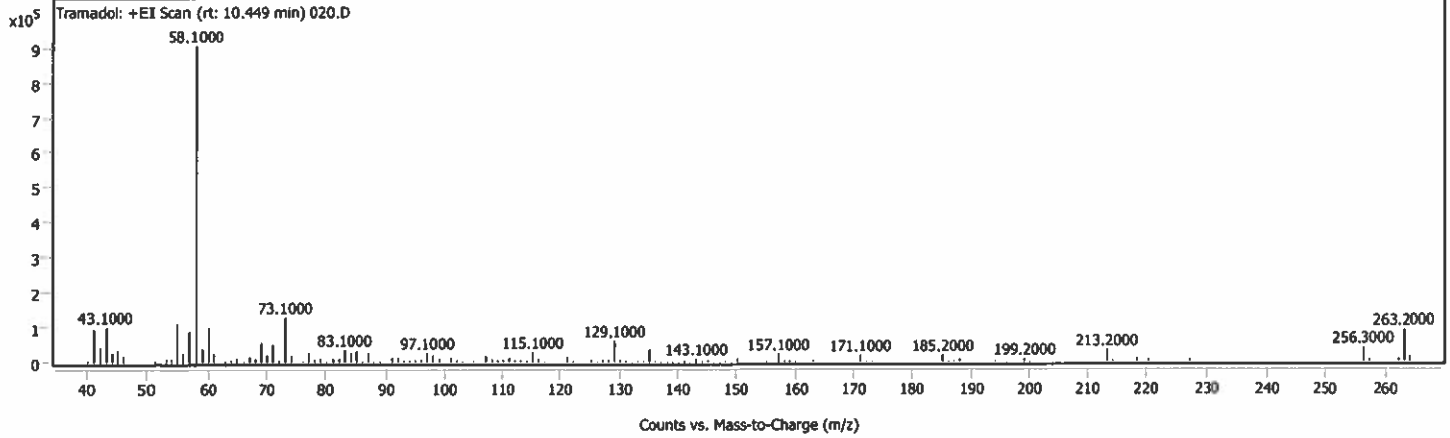
# Analysis Report

+ Scan (rt: 10.449 min)

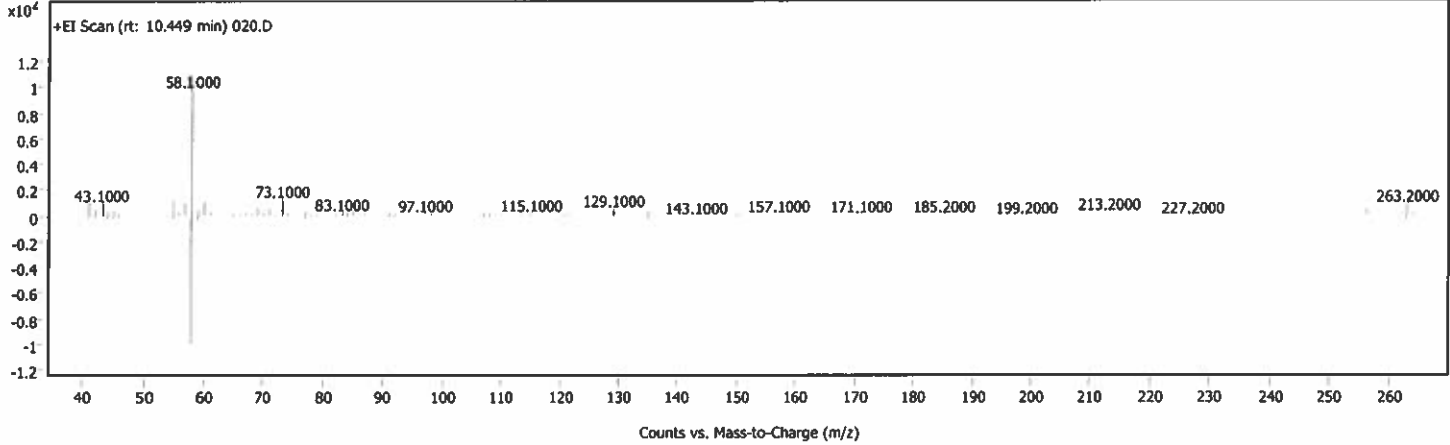
Tramadol; C16H25NO2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Tramadol	C16H25NO2	71.28	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

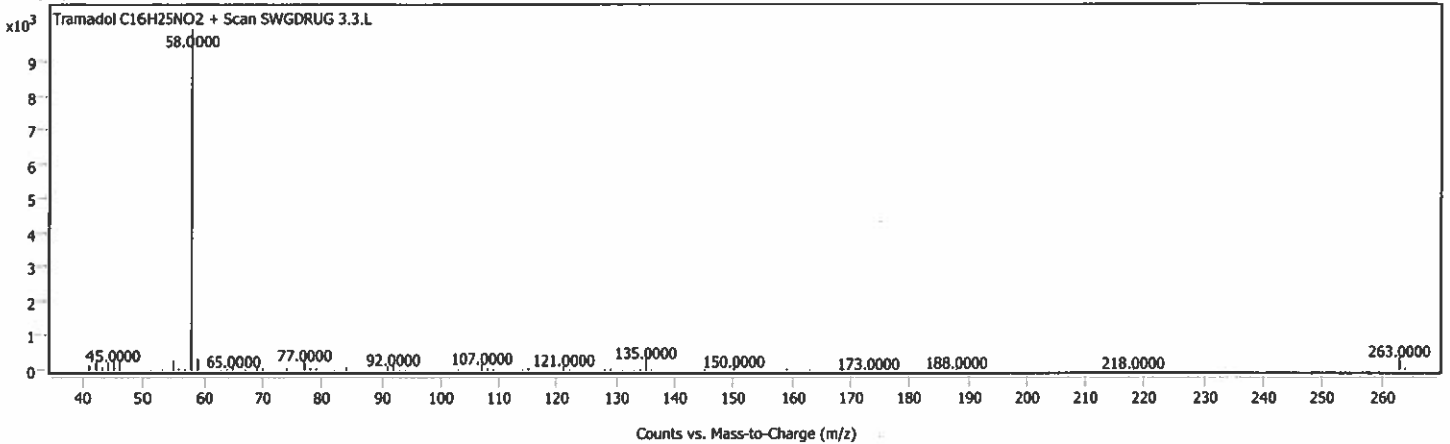
## Observed Spectrum



## Mirror Plot



## Library Spectrum



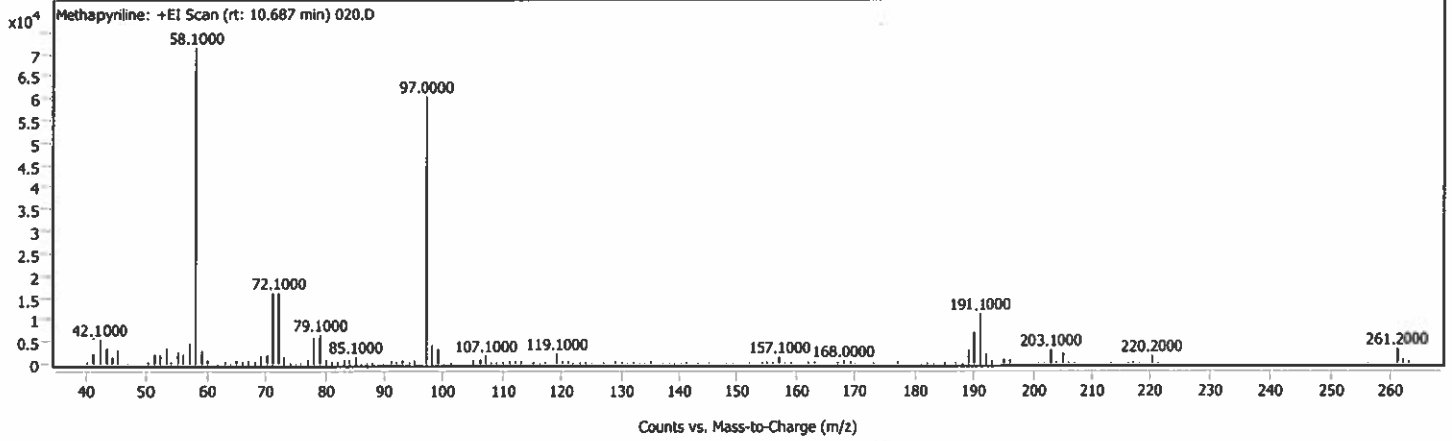
# Analysis Report

+ Scan (rt: 10.687 min)

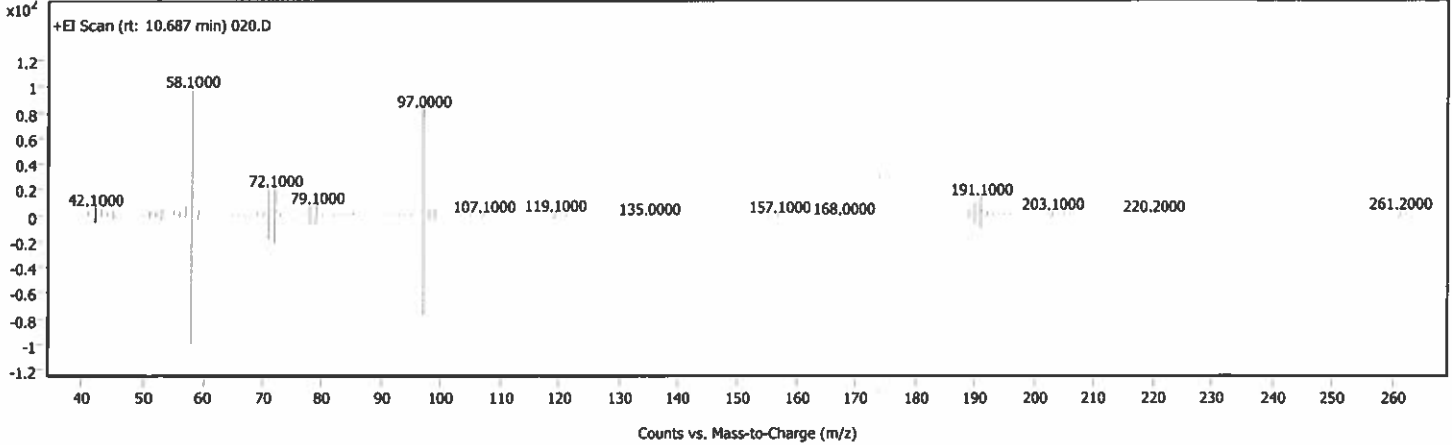
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	91.11	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

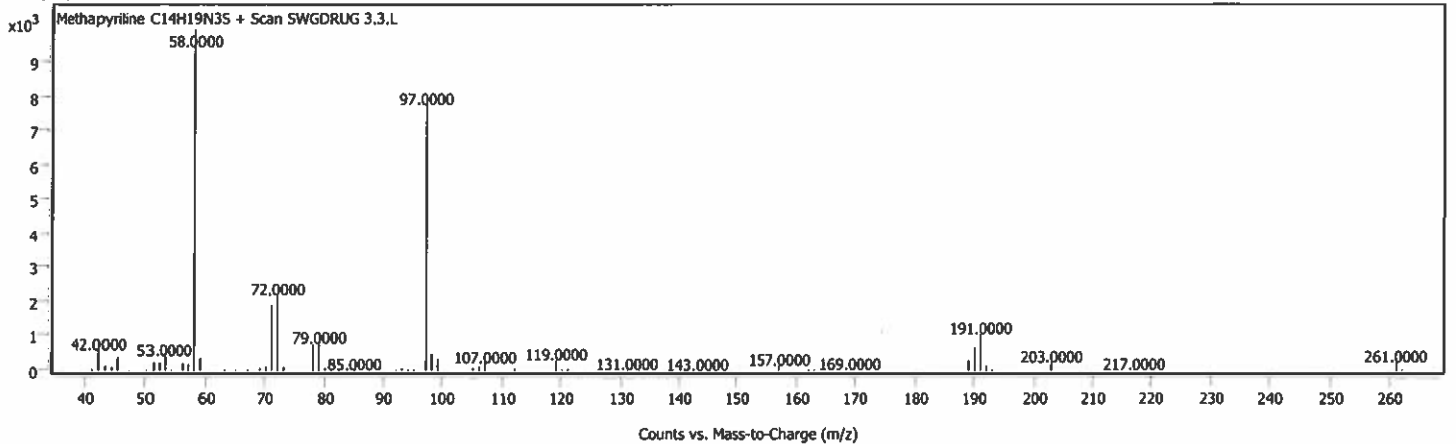
## Observed Spectrum



## Mirror Plot



## Library Spectrum



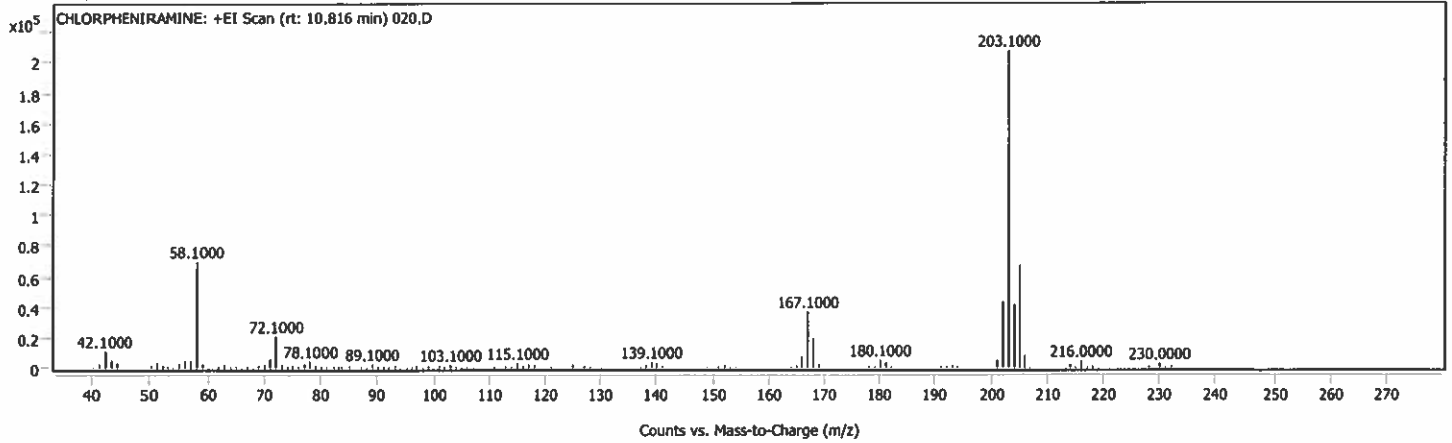
# Analysis Report

+ Scan (rt: 10.816 min)

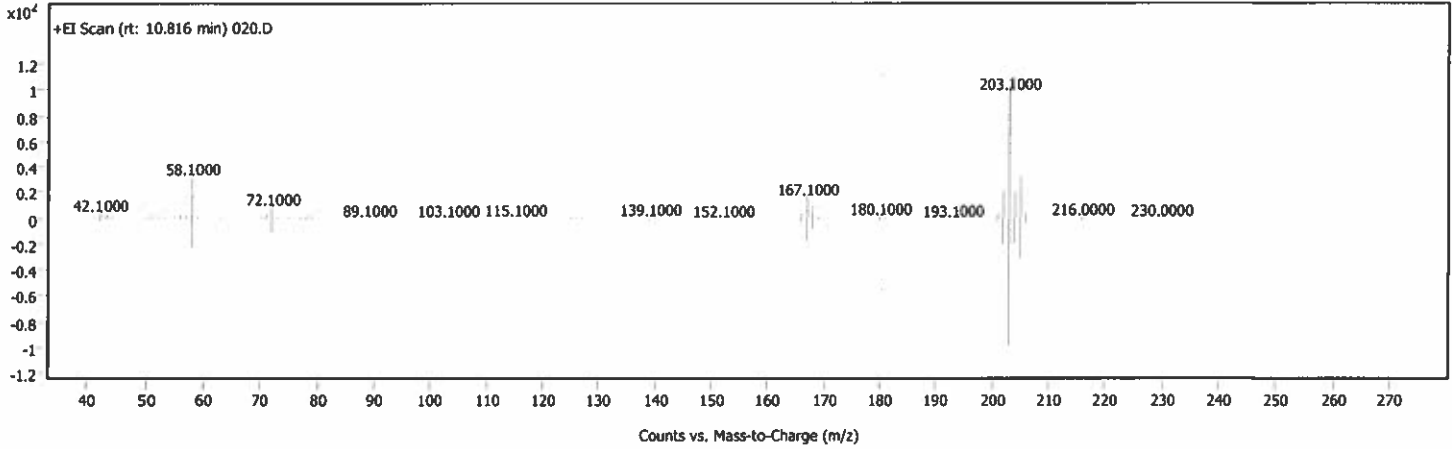
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	98.30	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

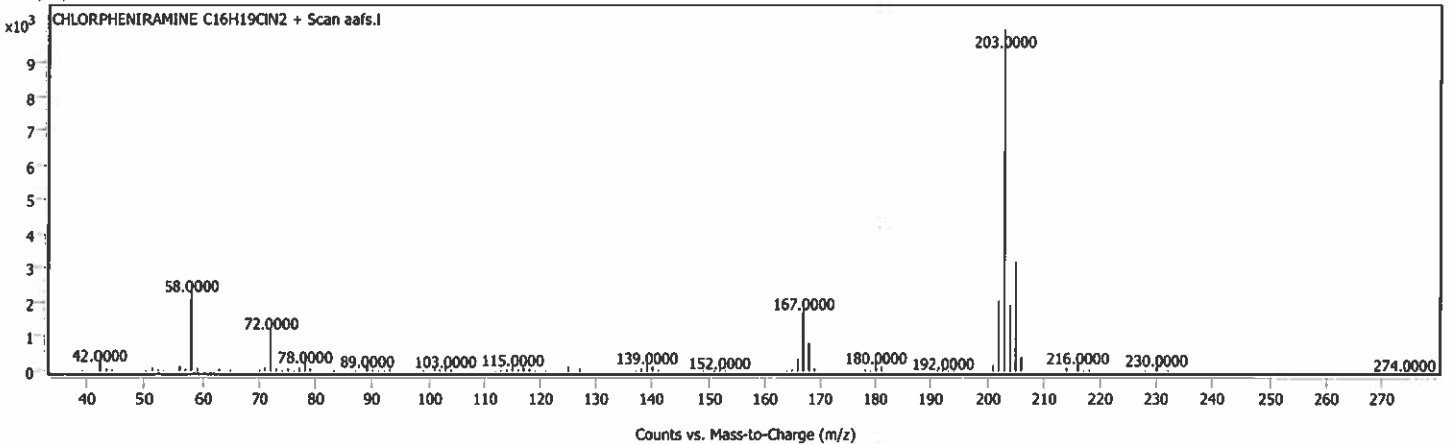
### Observed Spectrum



### Mirror Plot



### Library Spectrum

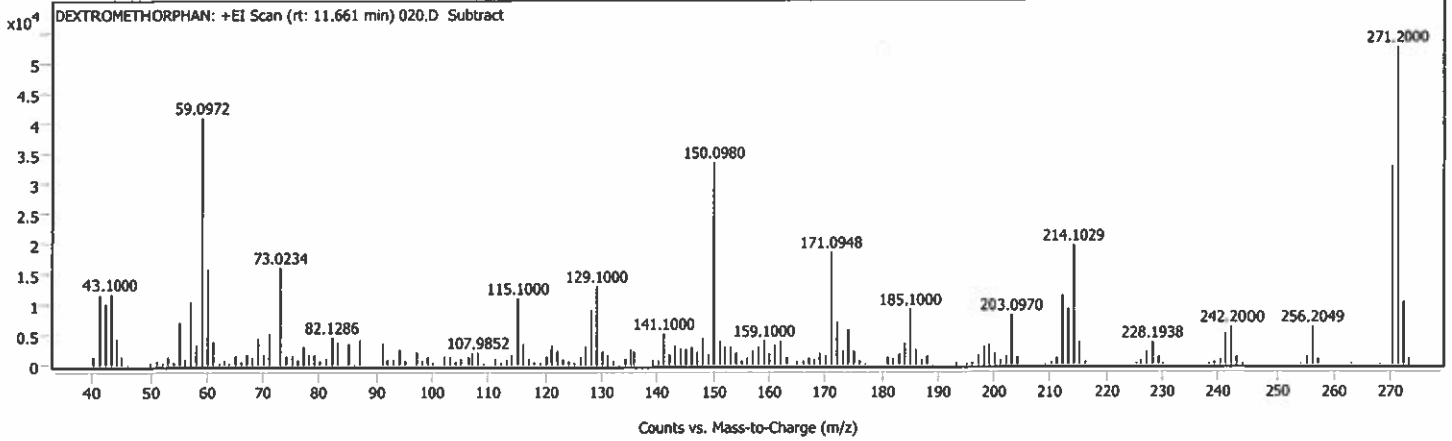


# Analysis Report

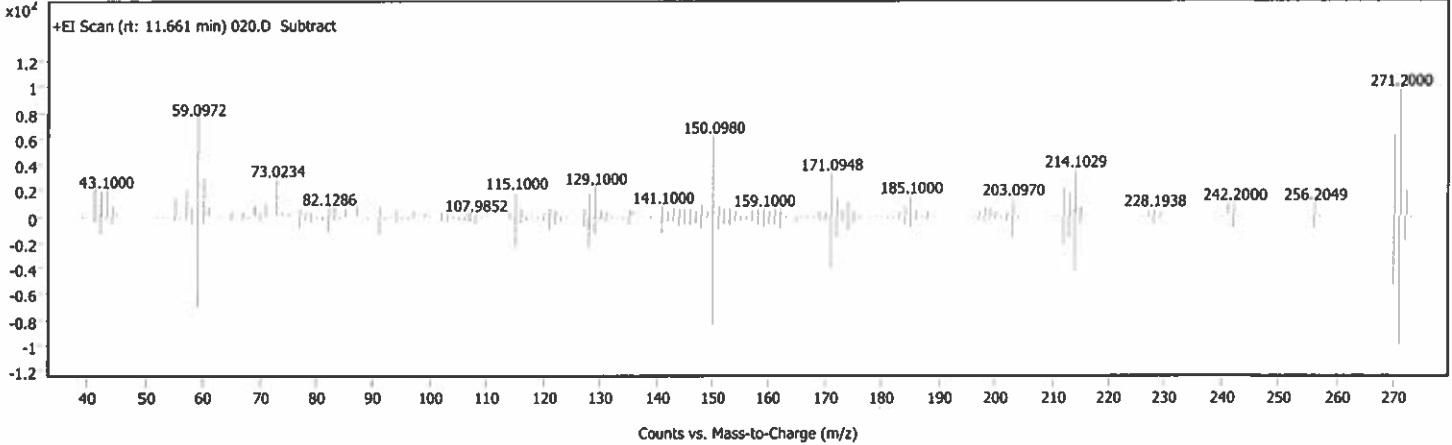
+ Scan (rt: 11.661 min) Sub DEXTROMETHORPHAN; C18H25NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	83.26	aafs.l	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

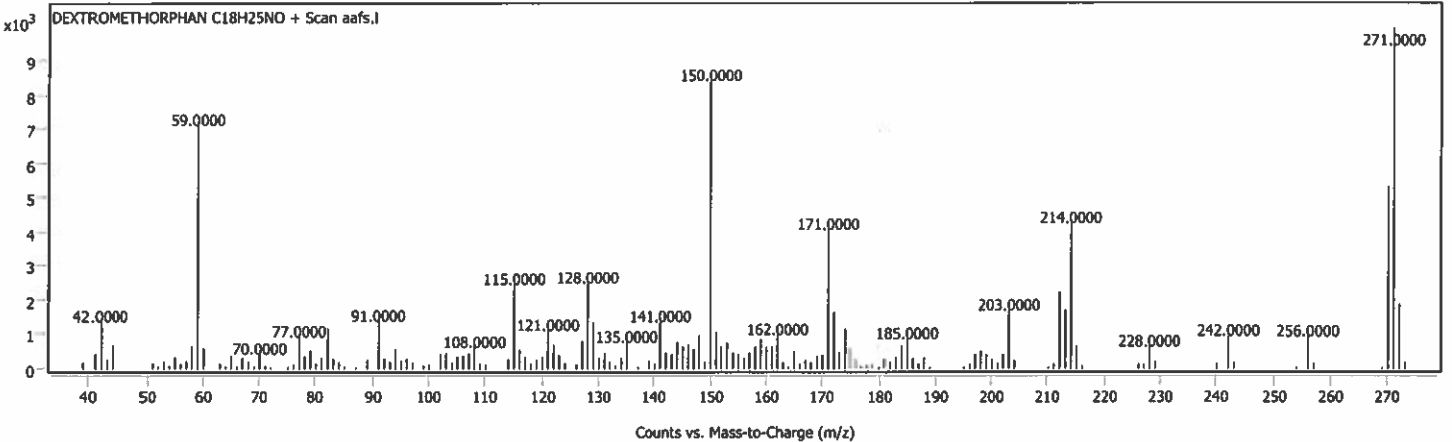
## Observed Spectrum



## Mirror Plot



## Library Spectrum



# Analysis Report

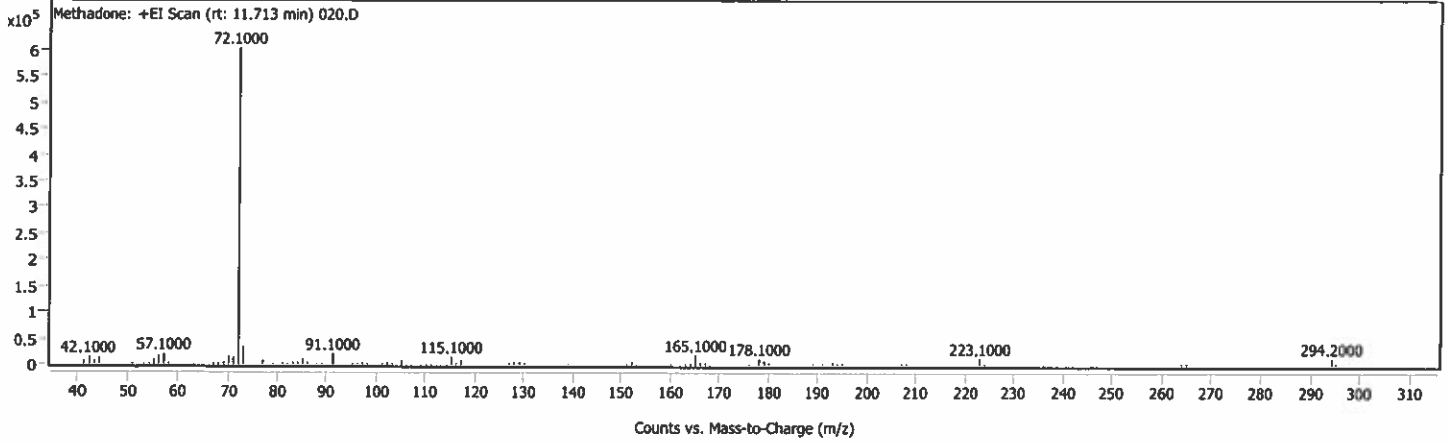


+ Scan (rt: 11.713 min)

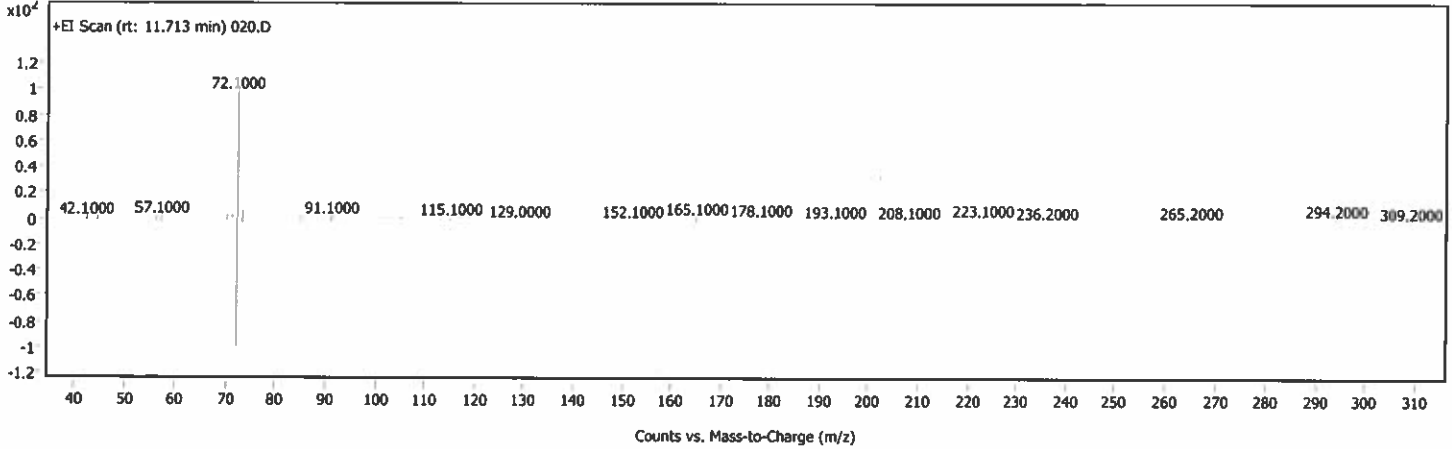
Methadone; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C21H27NO	92.81	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

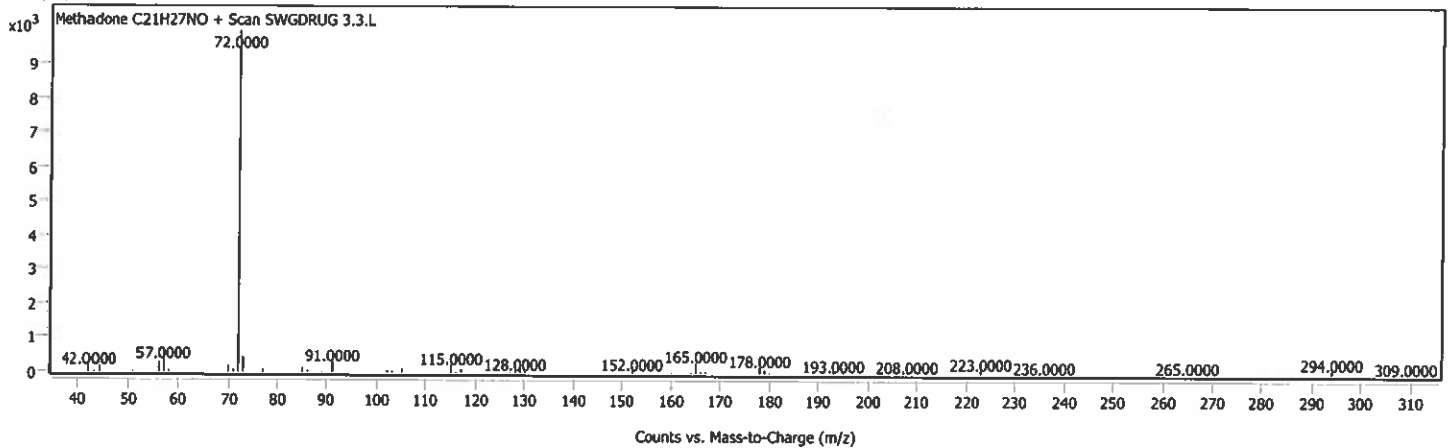
## Observed Spectrum



## Mirror Plot



## Library Spectrum





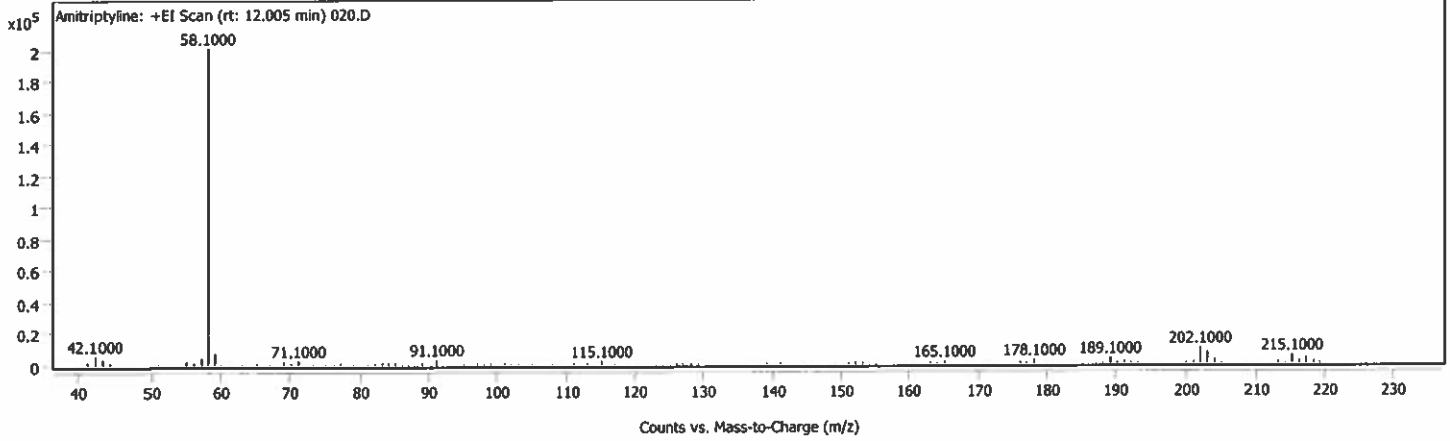
# Analysis Report

+ Scan (rt: 12.005 min)

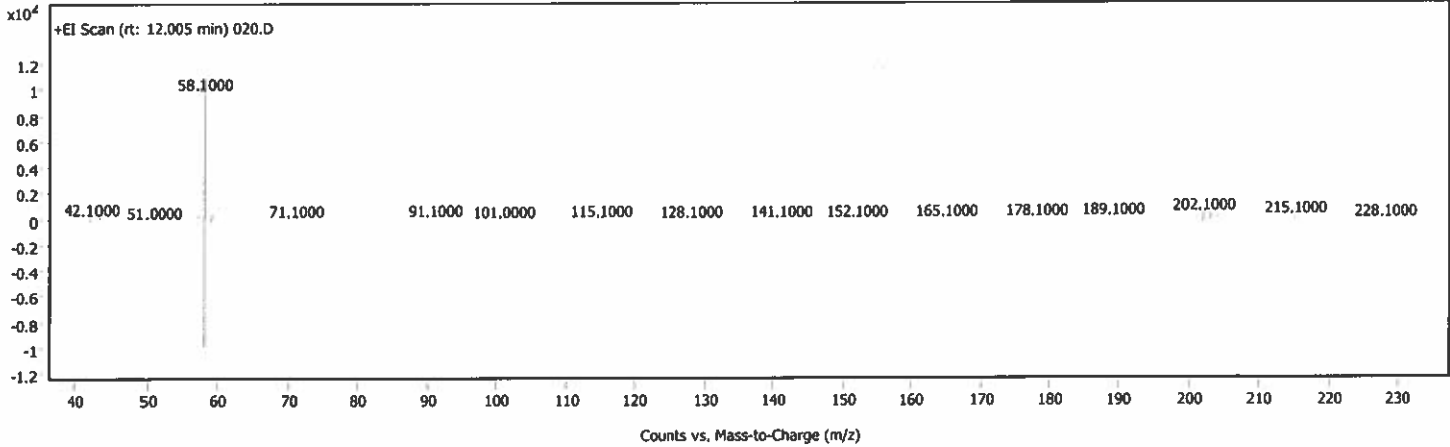
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	91.90	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

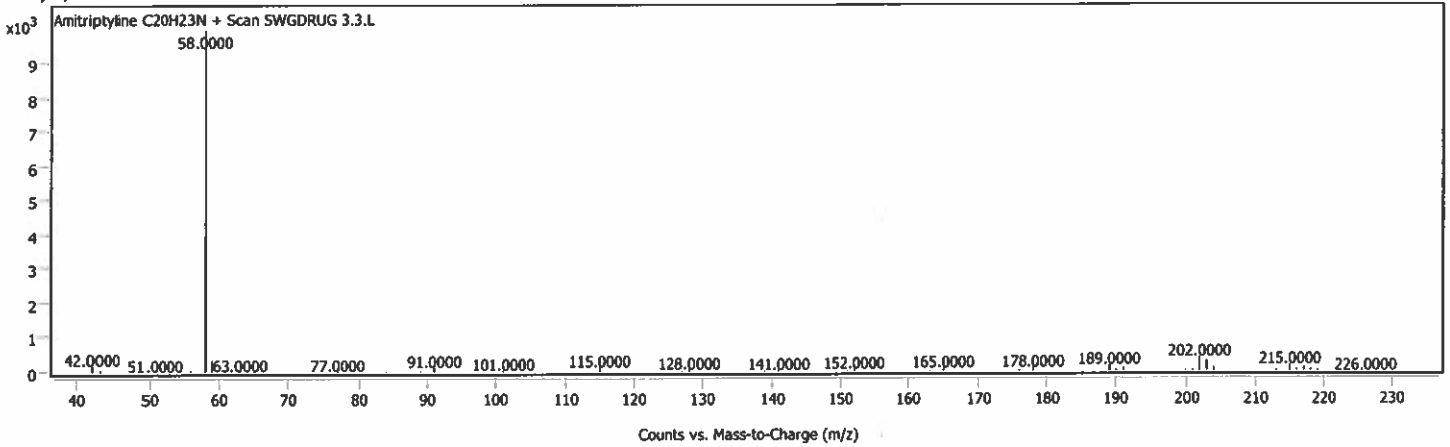
## Observed Spectrum



## Mirror Plot



## Library Spectrum



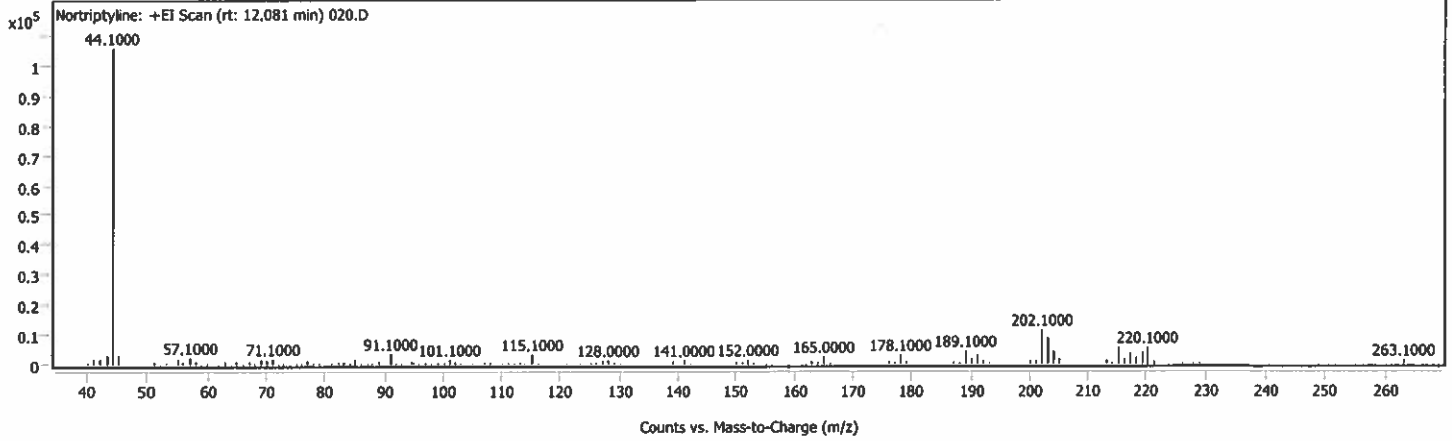
# Analysis Report

+ Scan (rt: 12.081 min)

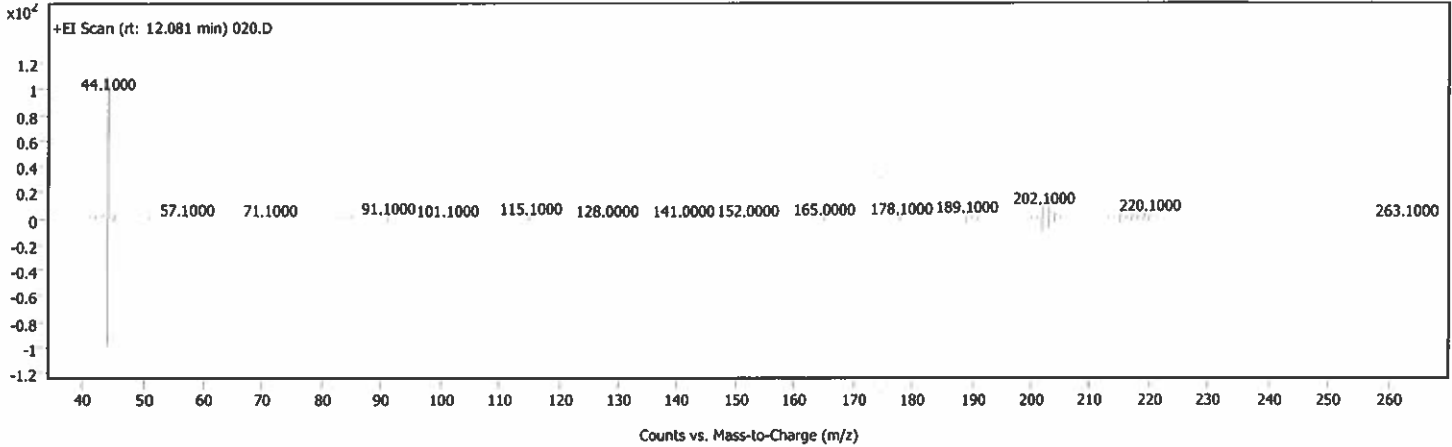
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	92.68	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

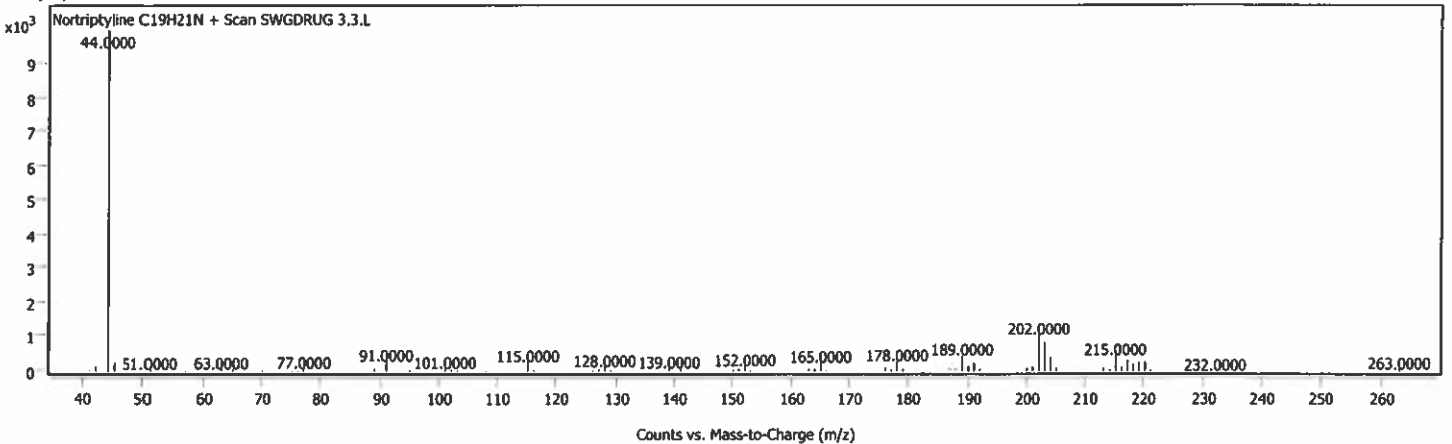
### Observed Spectrum



### Mirror Plot



### Library Spectrum



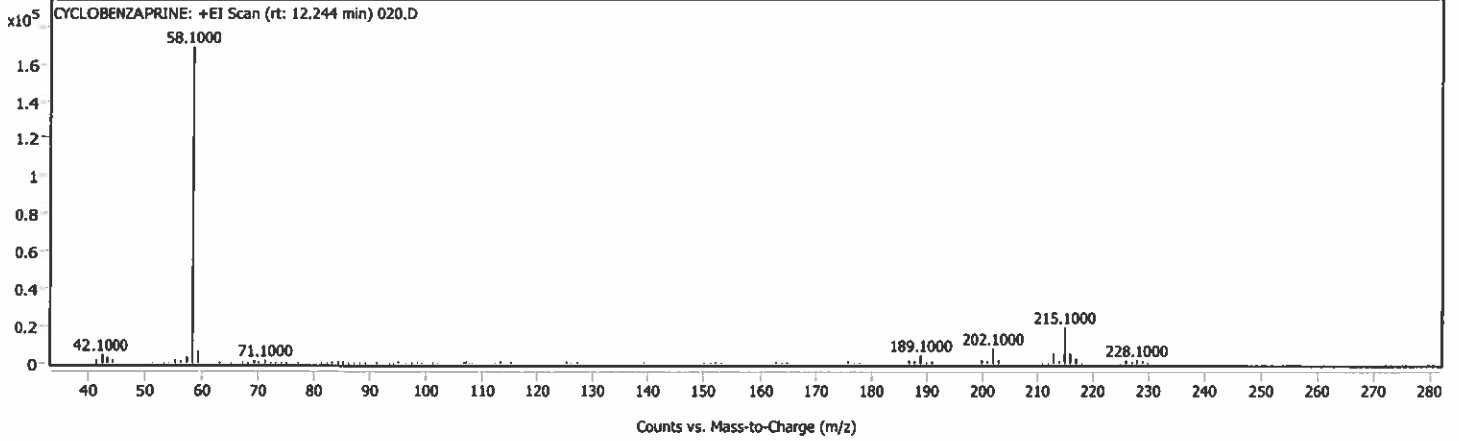
# Analysis Report

+ Scan (rt: 12.244 min)

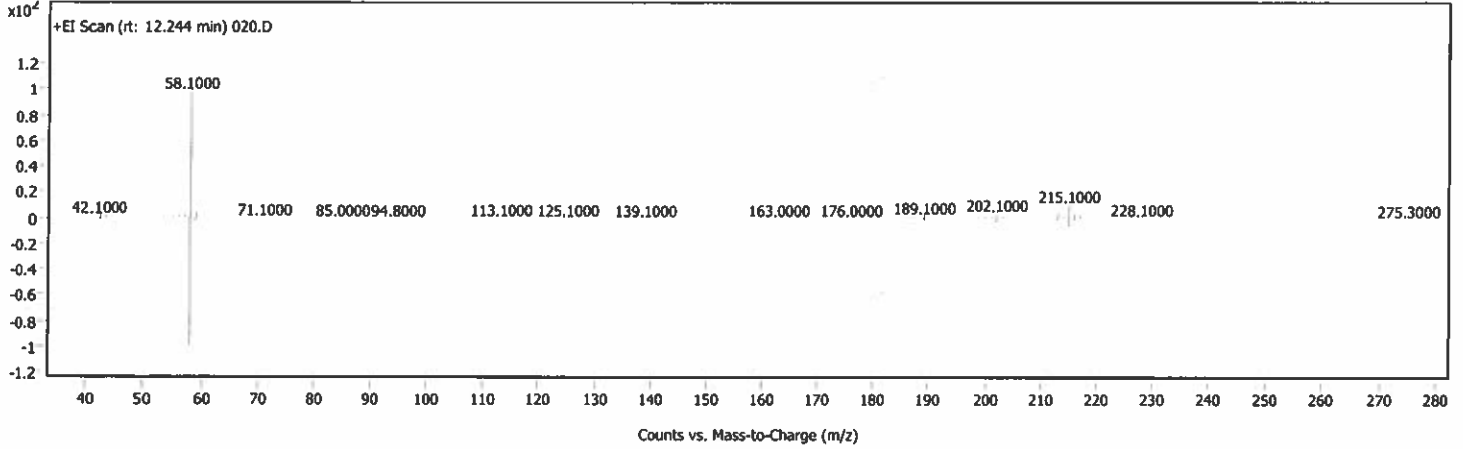
CYCLOBENZAPRINE; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CYCLOBENZAPRINE	C20H21N	91.06	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XLAZINE VALIDATION

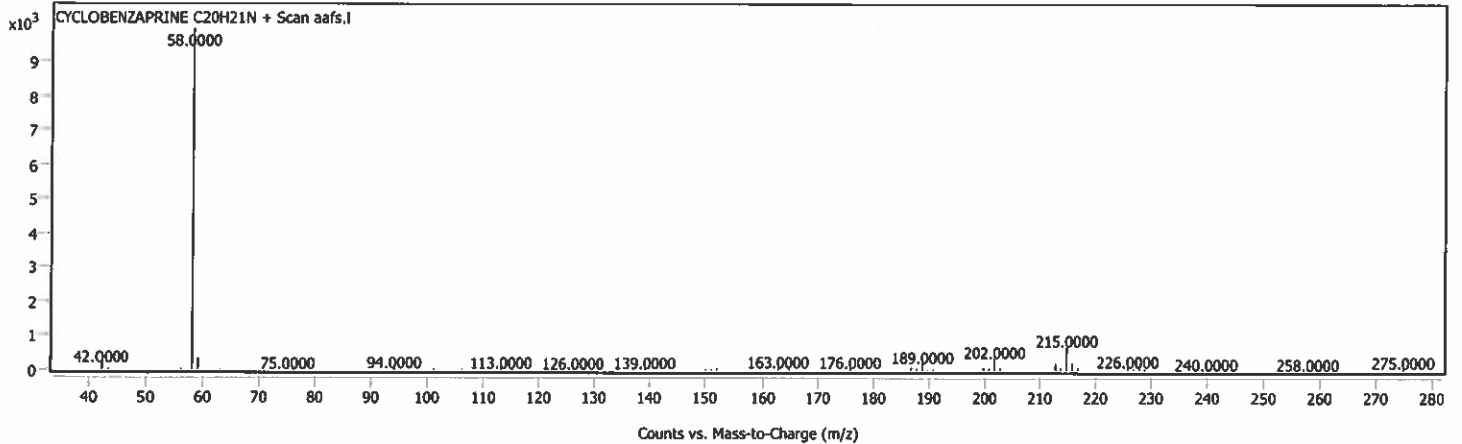
### Observed Spectrum



### Mirror Plot



### Library Spectrum



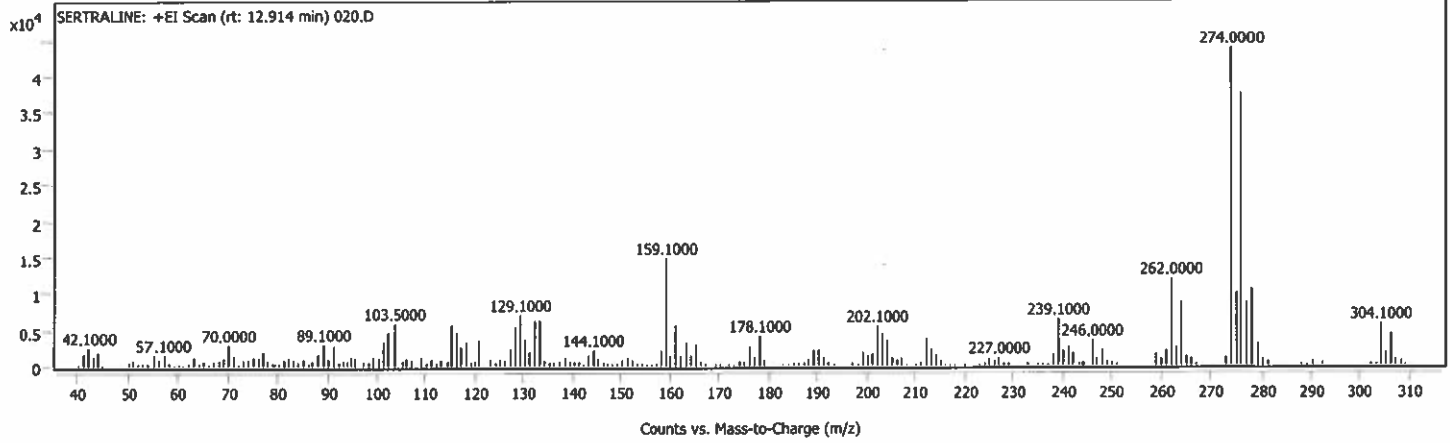
# Analysis Report

+ Scan (rt: 12.914 min)

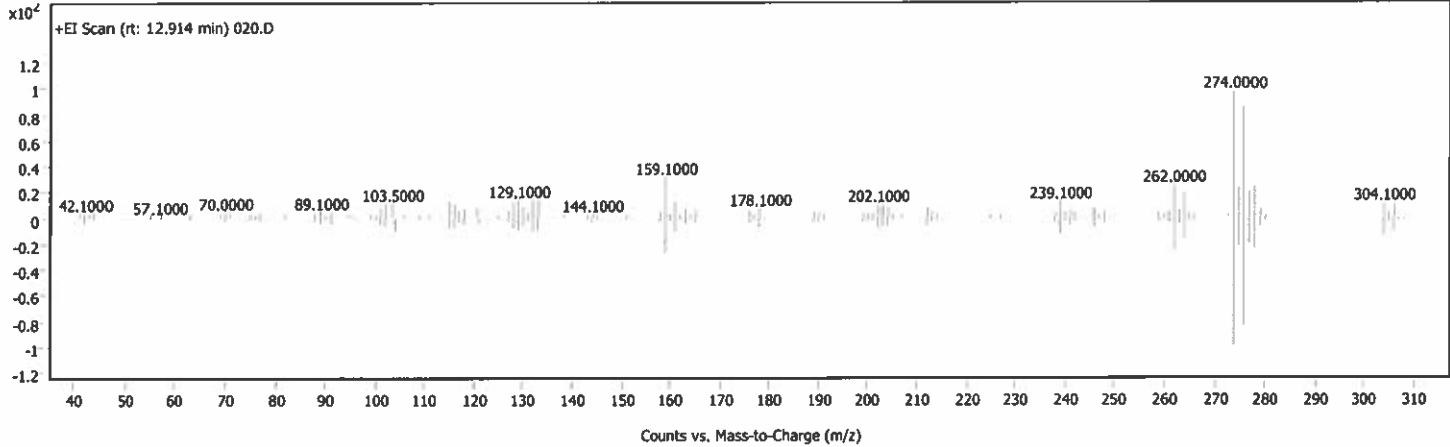
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	90.27	aafs.l	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

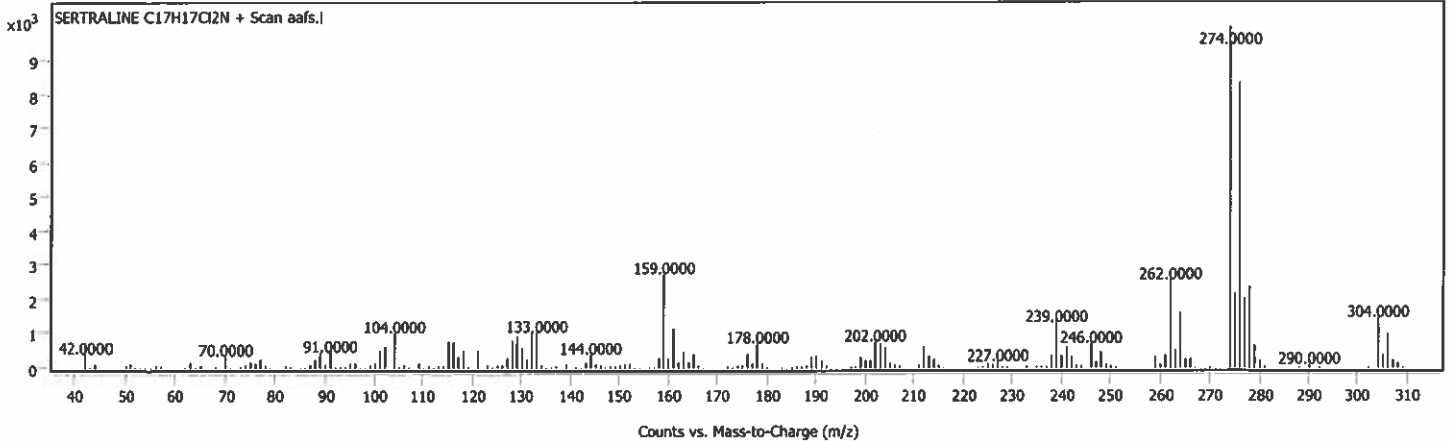
## Observed Spectrum



## Mirror Plot



## Library Spectrum



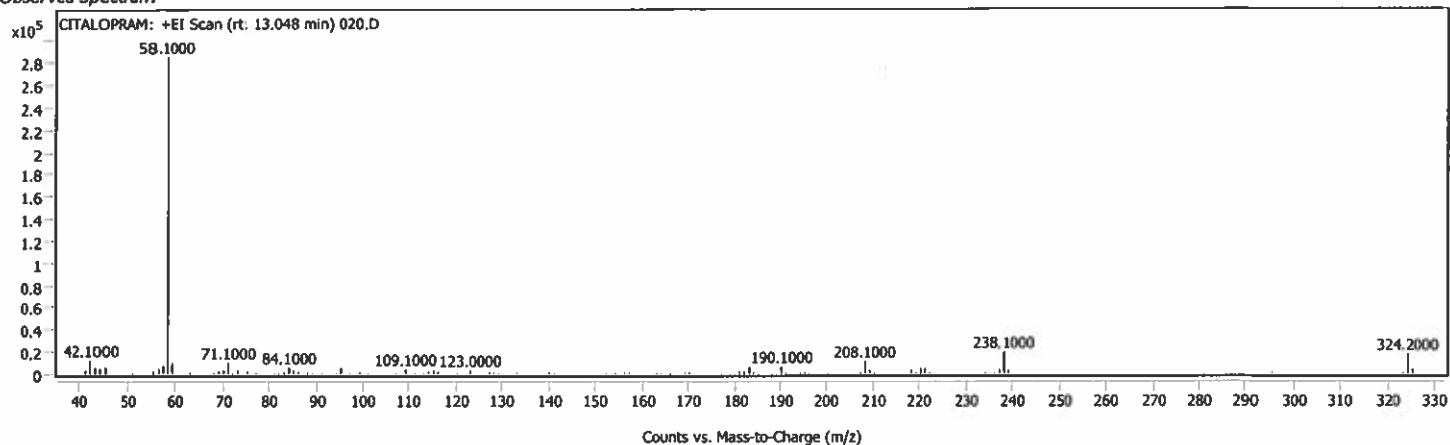
# Analysis Report

+ Scan (rt: 13.048 min)

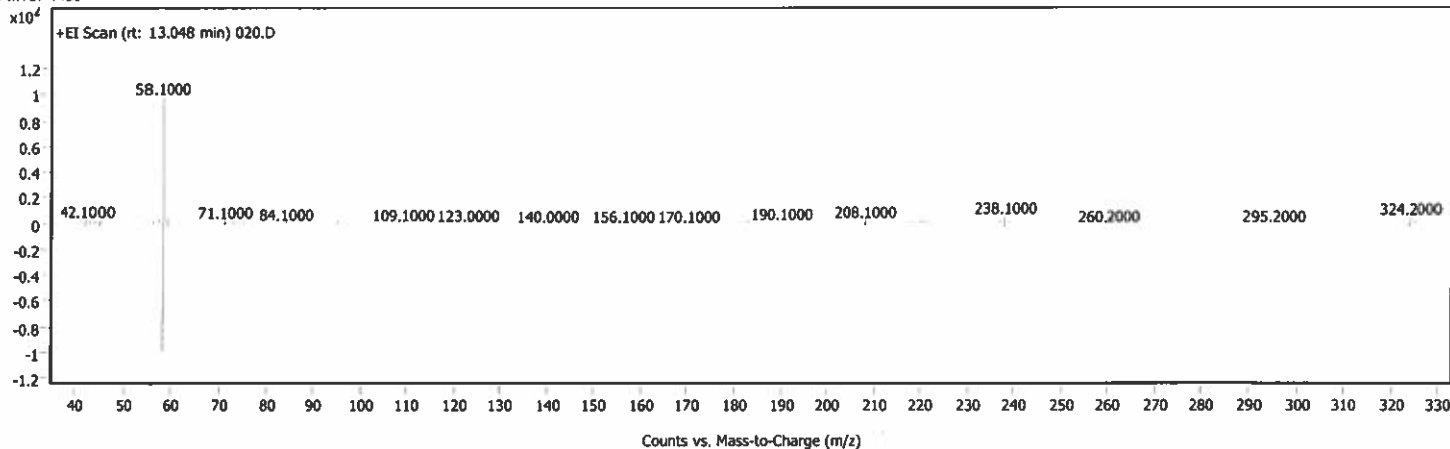
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	93.49	aafs.1	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

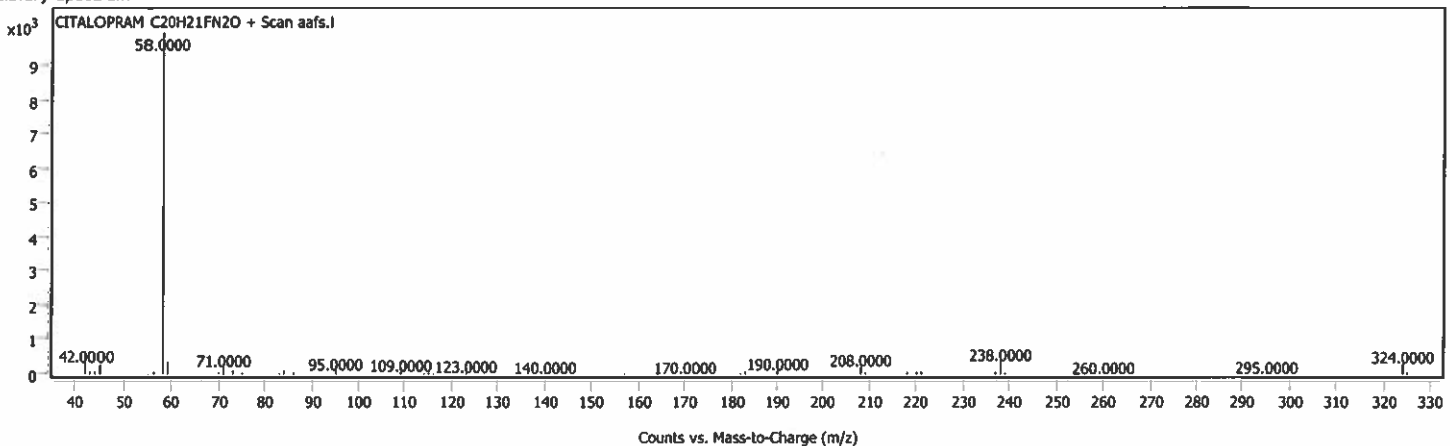
## Observed Spectrum



## Mirror Plot



## Library Spectrum

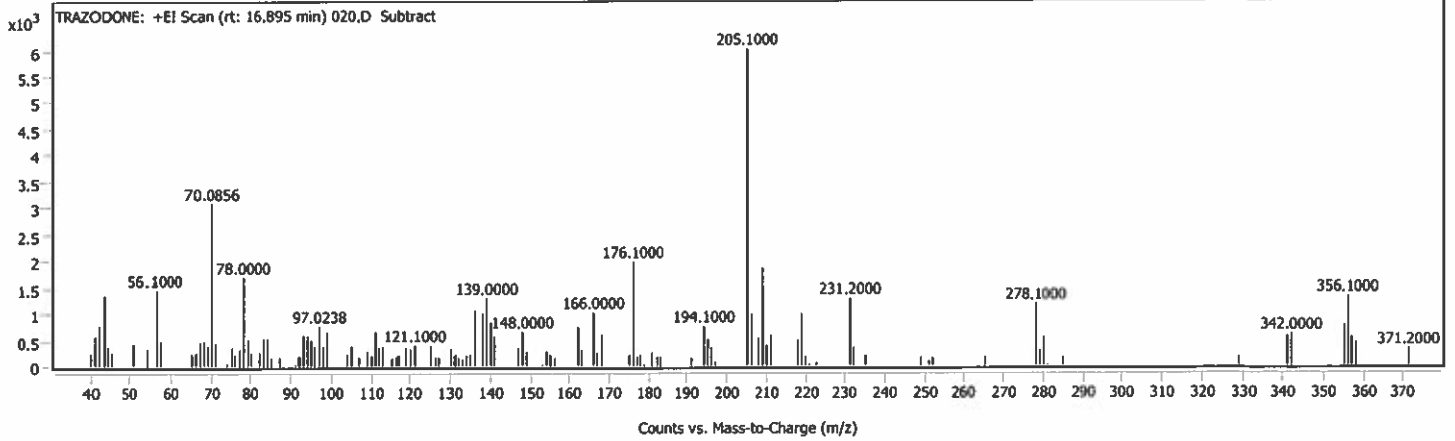


# Analysis Report

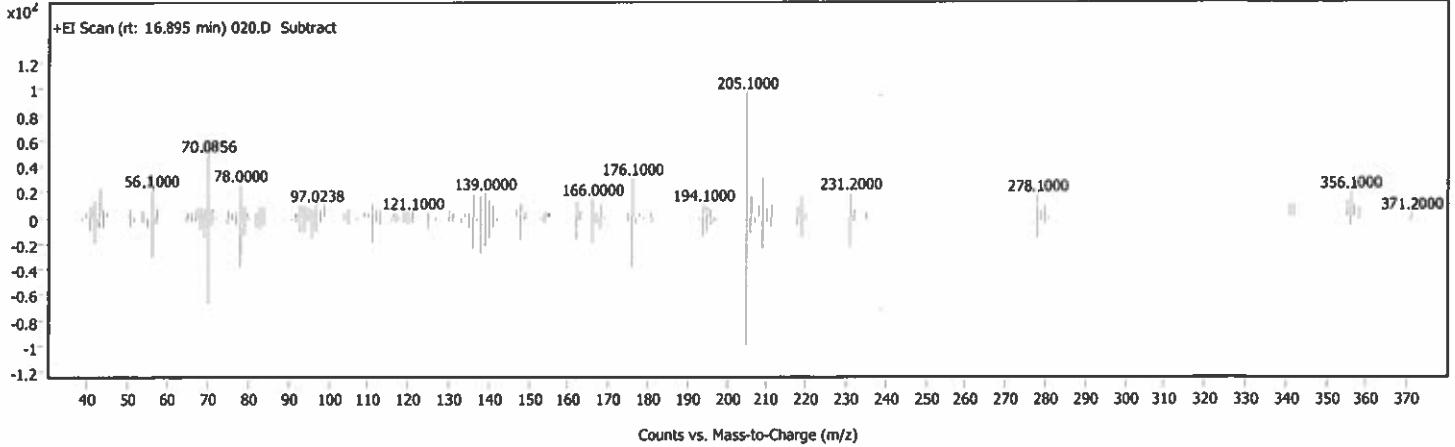
+ Scan (rt: 16.895 min) Sub **TRAZODONE**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		82.12	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

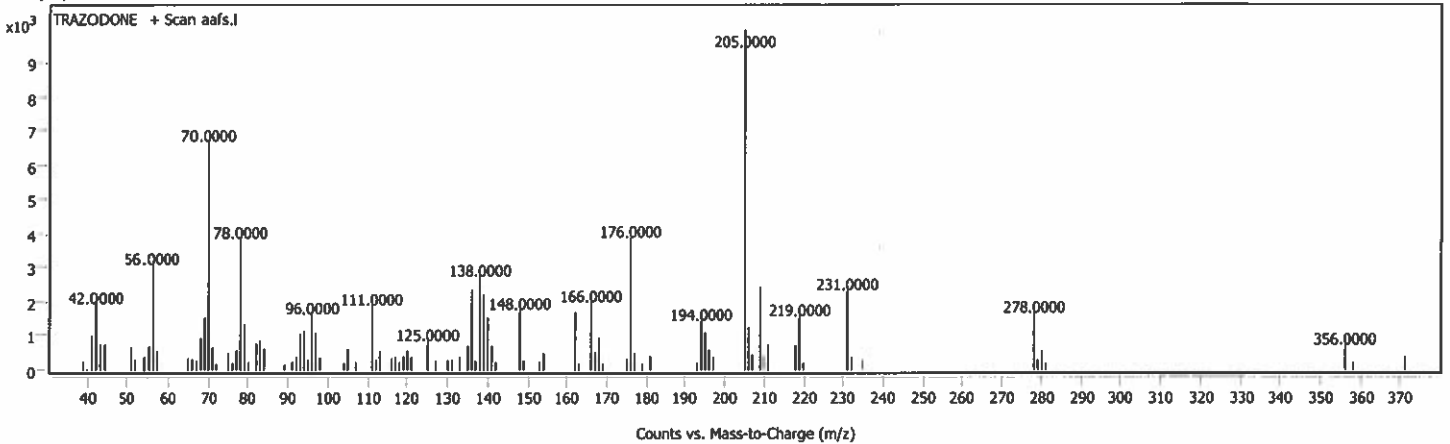
## Observed Spectrum



## Mirror Plot

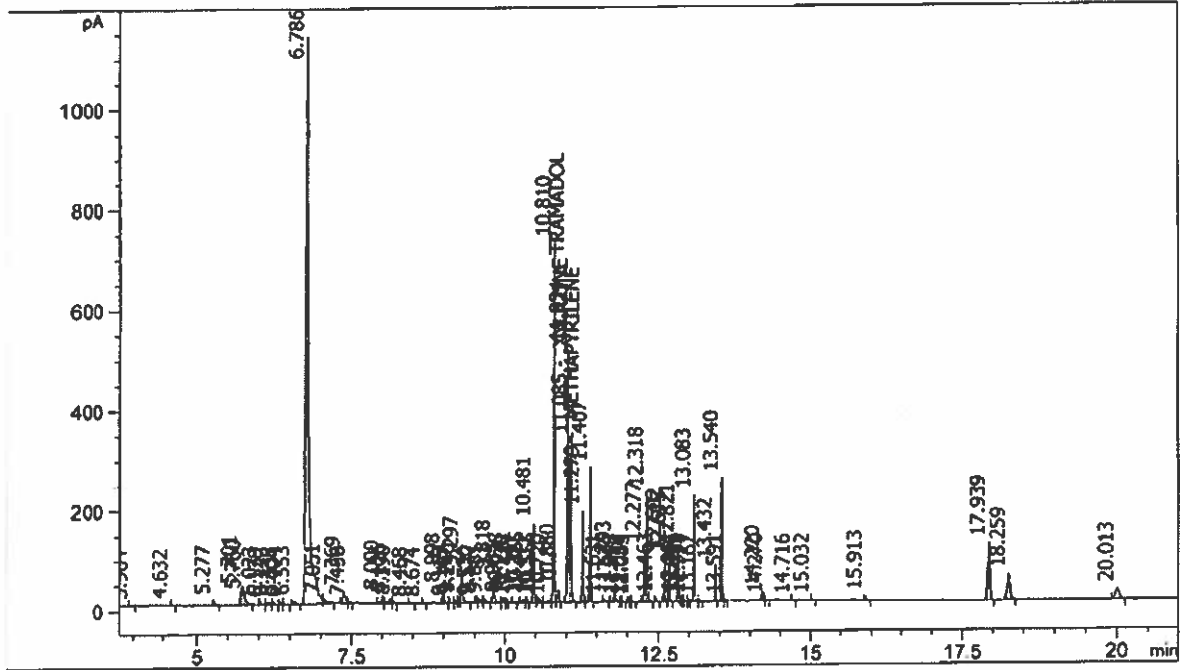


## Library Spectrum



Injection Date: 9/29/2021 1:55:08 AM Seq Line: 31  
 Sample Name: MATRIX 10 W/ 0.1mg/L -> Vial 32  
 Sample Info: *cal + xylazine*  
*KM 11/17/22*

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



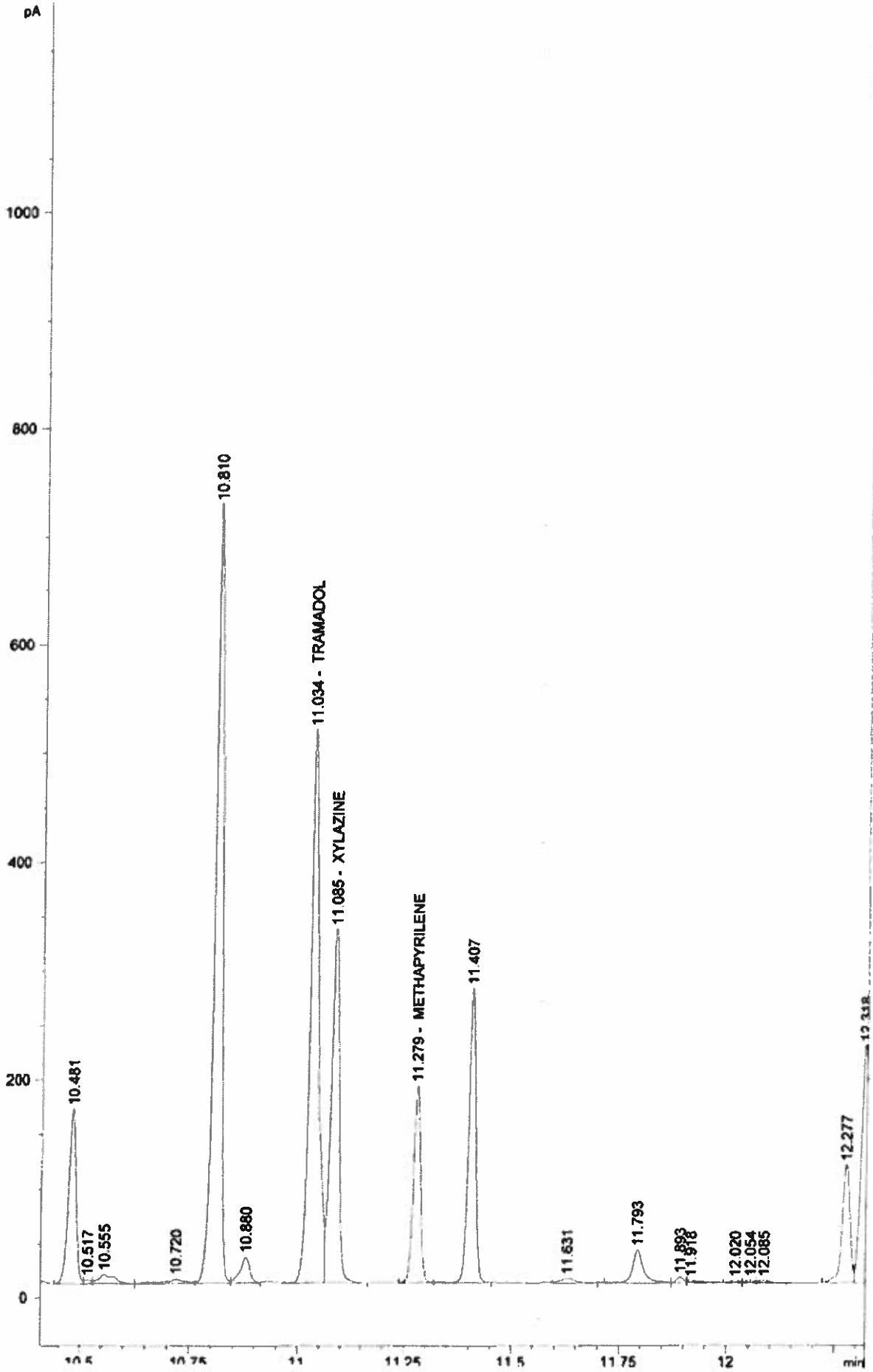
see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.034	11.033	717.094	508.150	0.000000	TRAMADOL RRT - 0.9783
11.085	11.046	429.555	324.796	0.000000	XYLAZINE RRT - 0.9828
11.279	11.278	207.807	180.817	1.000000	METHAPYRILENE

LVR not used in avg of RRTs

Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000031 D)

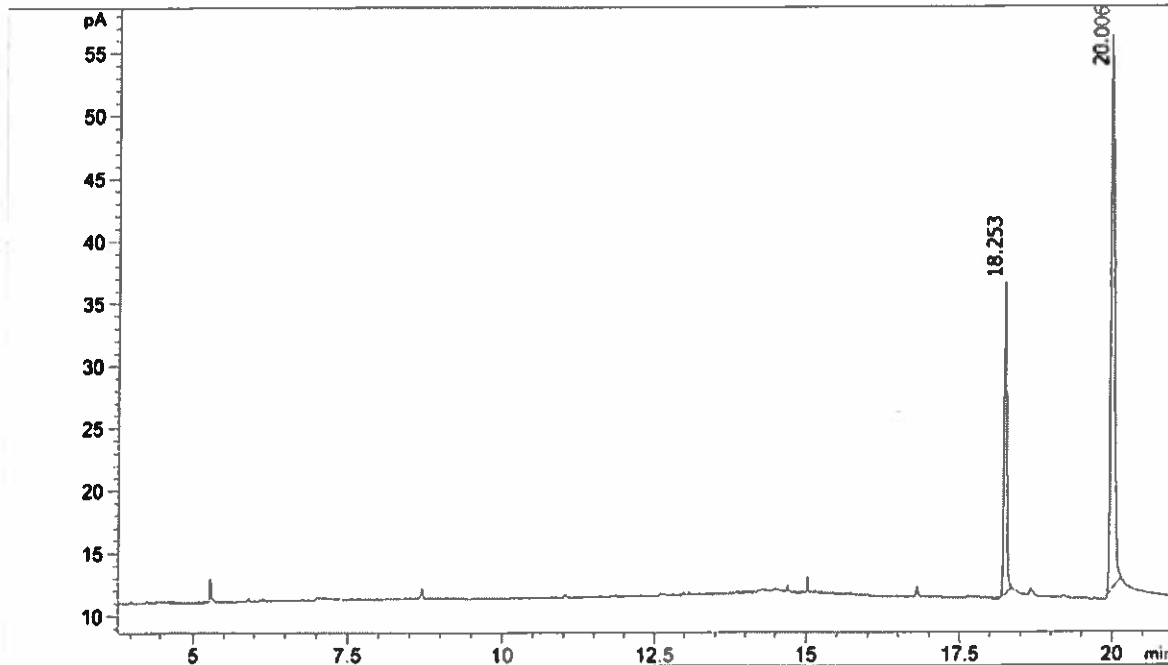




Injection Date: 9/29/2021 2:19:25 AM  
Sample Name: SOLVENT BLANK 2  
Sample Info:

Seq Line: 32  
Vial 33

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



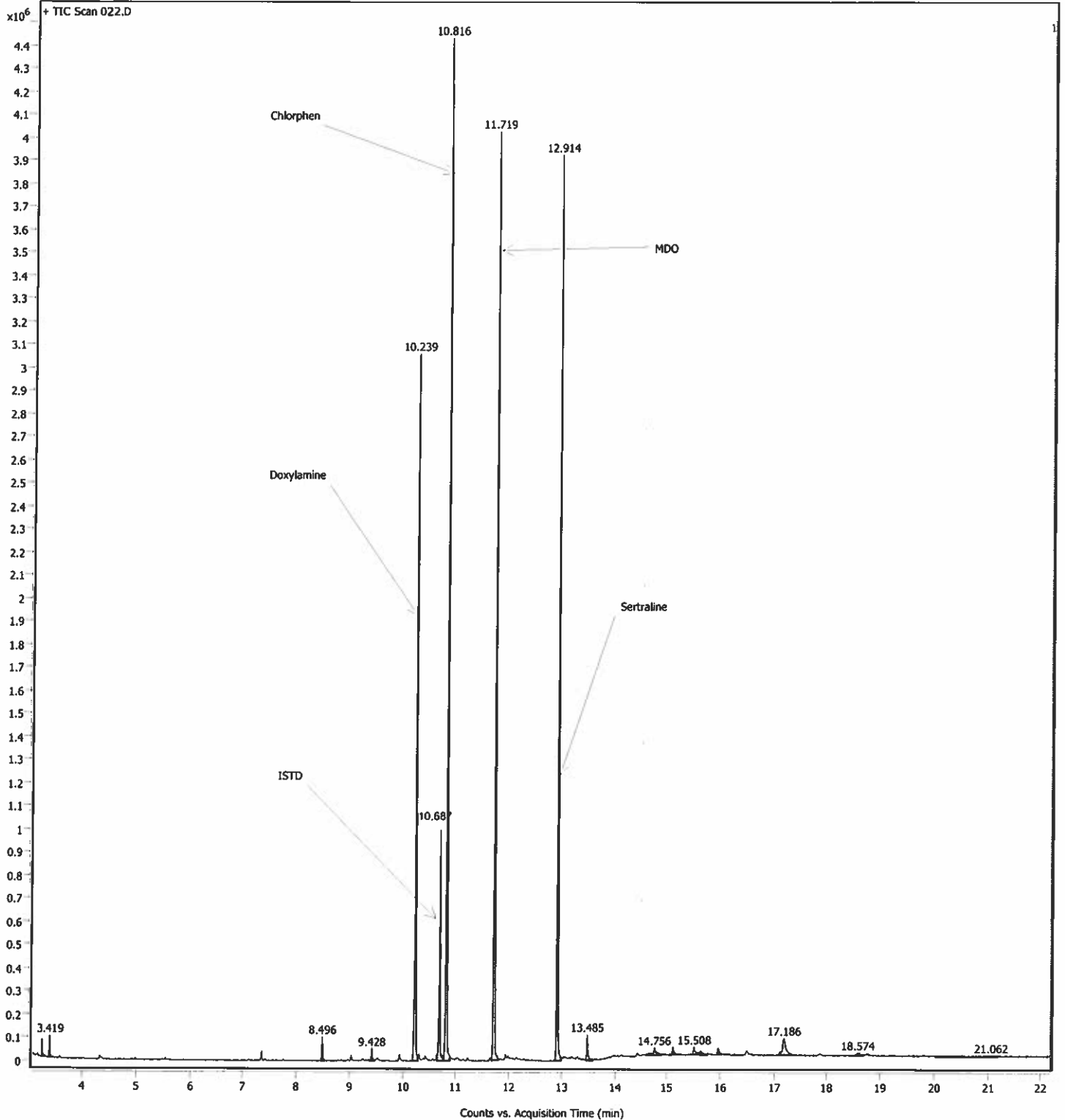
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 2 W/ IH CTL - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\022.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 11:08:49 PM (UTC-04:00)
Position	49	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

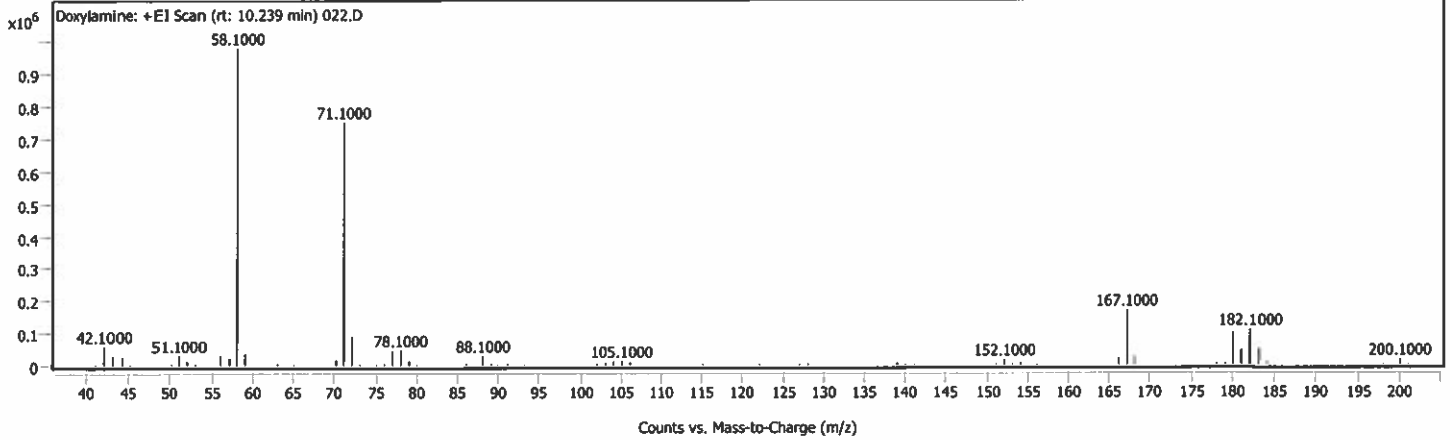
## Sample Spectra

+ Scan (rt: 10.239 min)

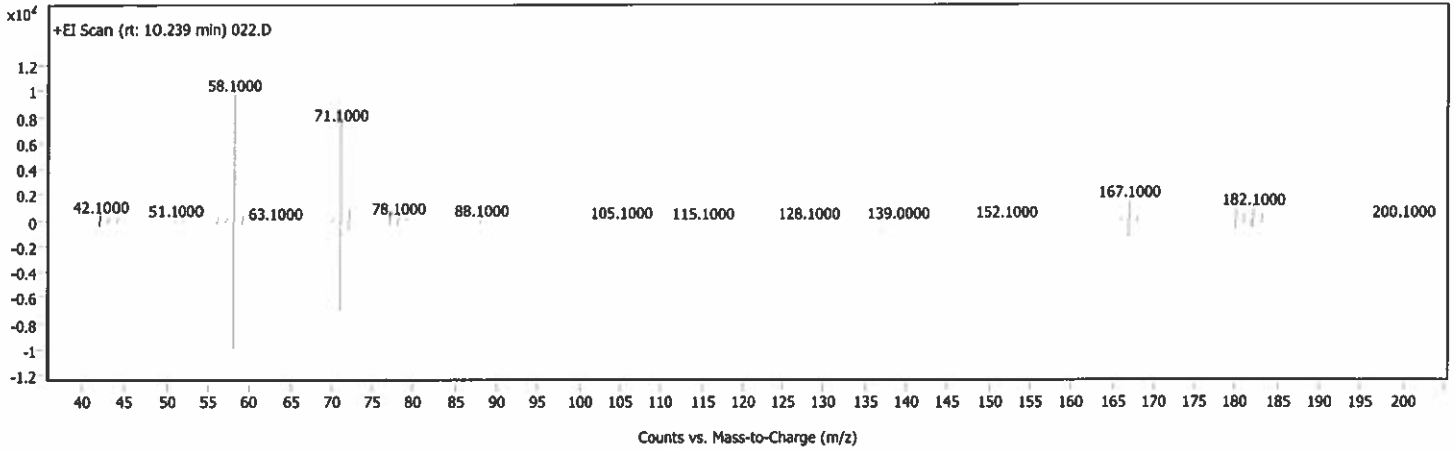
Doxylamine; C<sub>17</sub>H<sub>22</sub>N<sub>2</sub>O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C <sub>17</sub> H <sub>22</sub> N <sub>2</sub> O	98.79	SWGDRUG 3.3.L	MATRIX 2 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

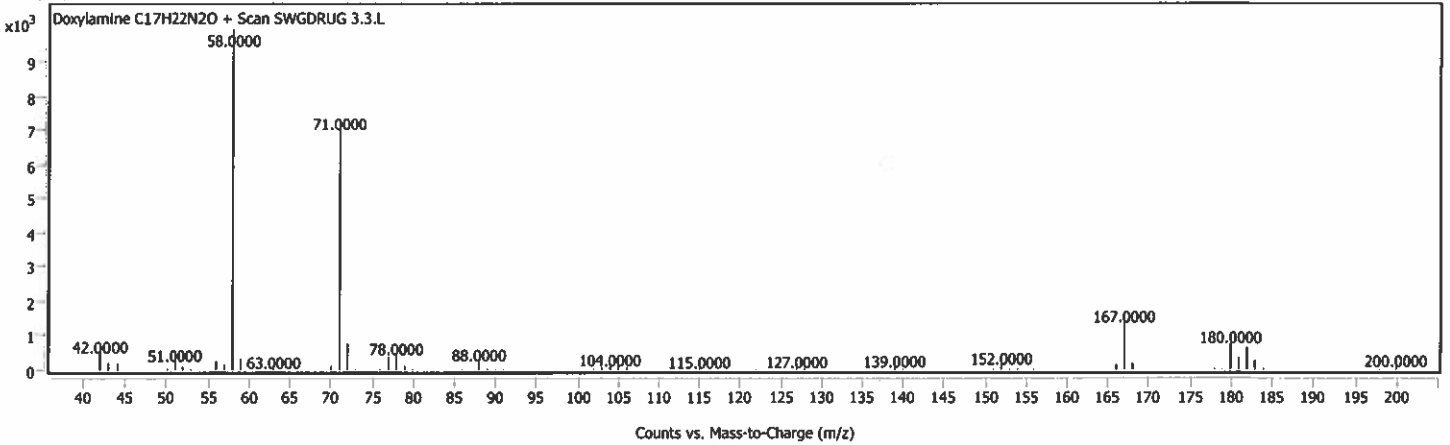
## Observed Spectrum



## Mirror Plot



## Library Spectrum



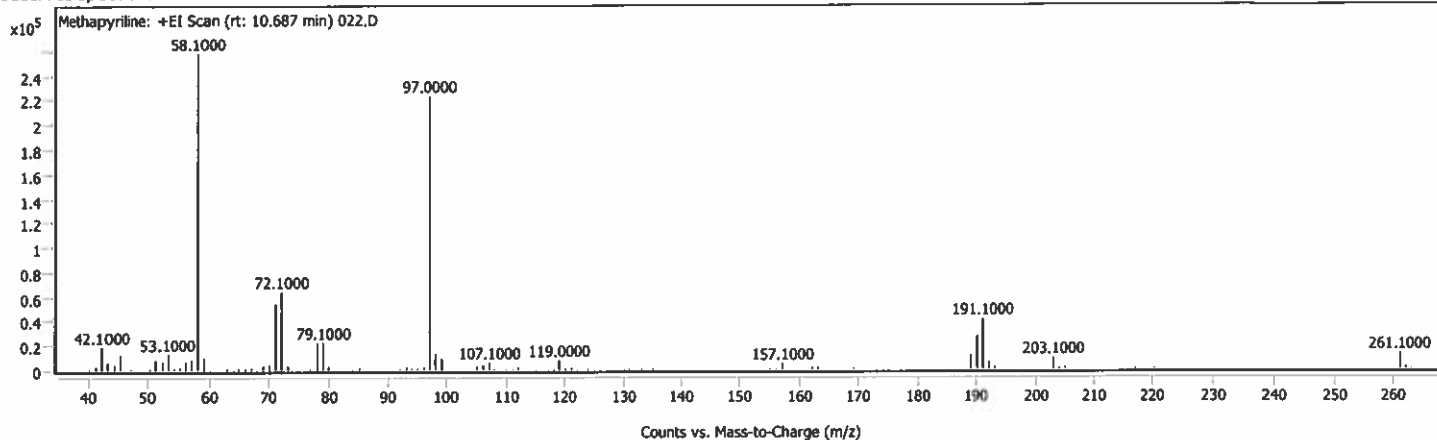
# Analysis Report

+ Scan (rt: 10.687 min)

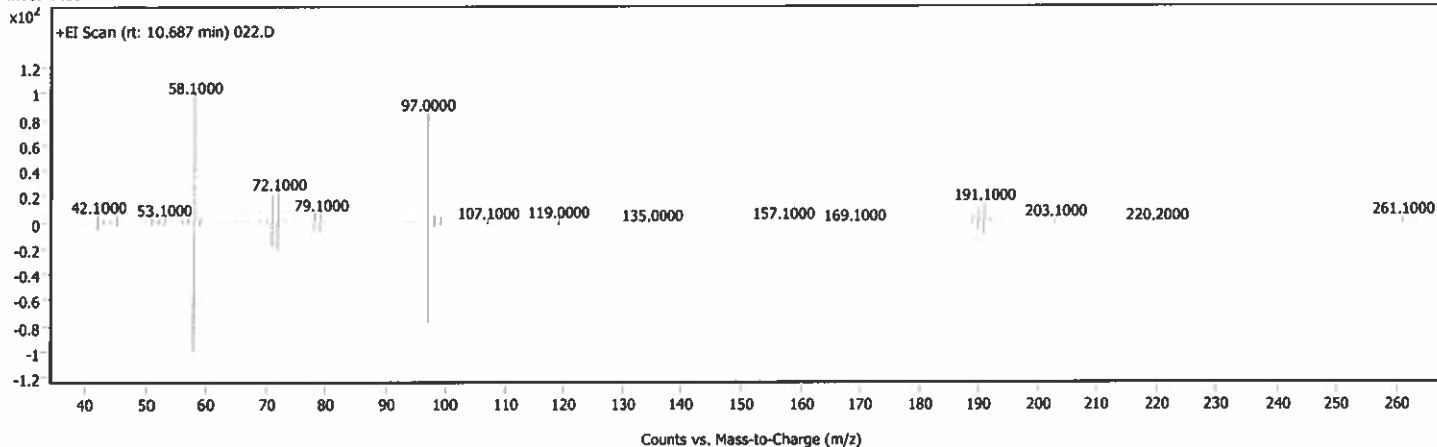
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.26	SWGDRUG 3.3.L	MATRIX 2 W/ TH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

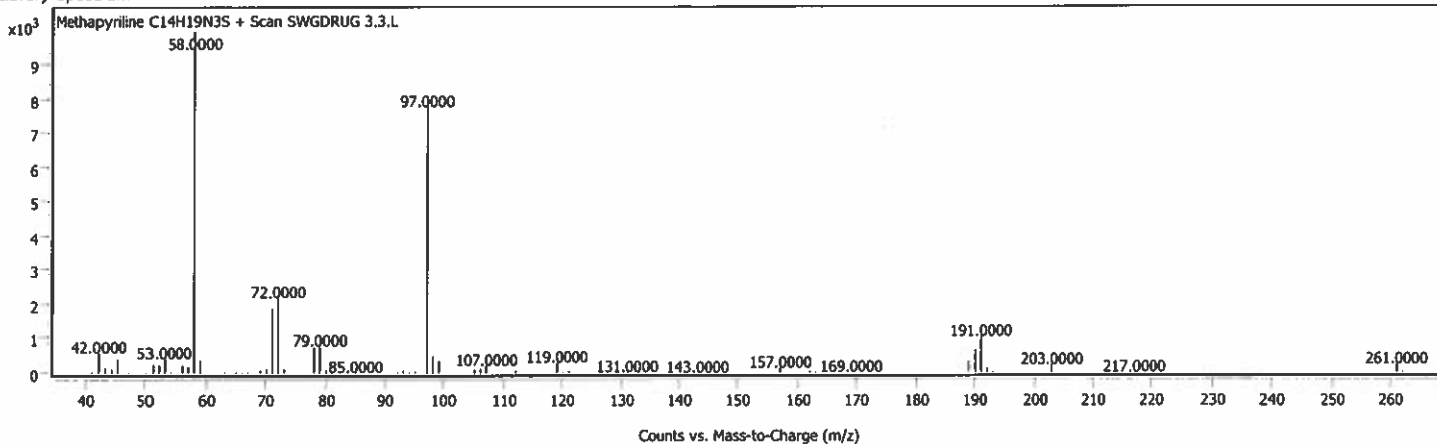
## Observed Spectrum



## Mirror Plot



## Library Spectrum



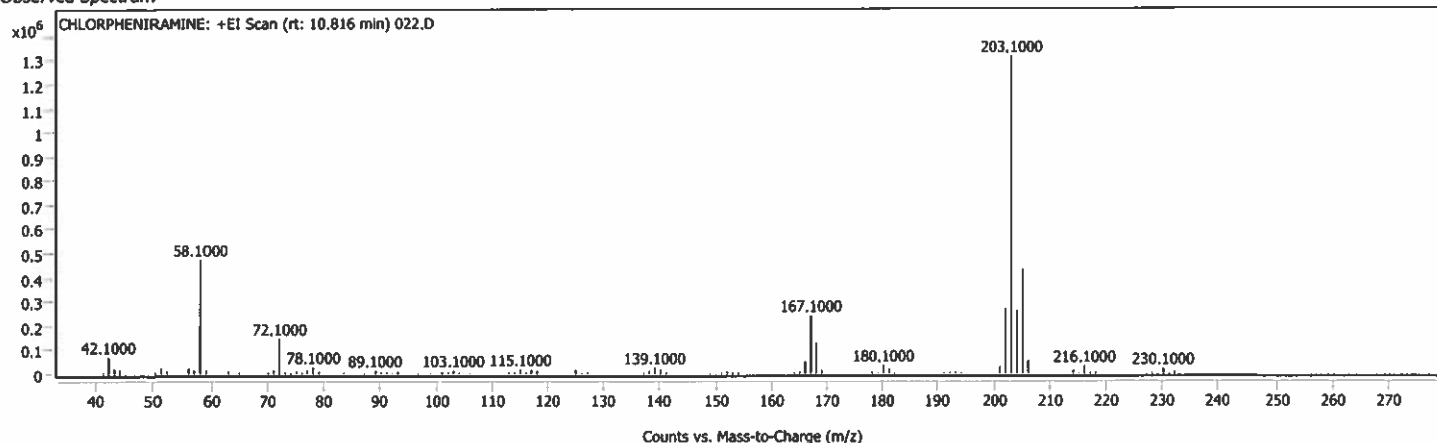
# Analysis Report

+ Scan (rt: 10.816 min)

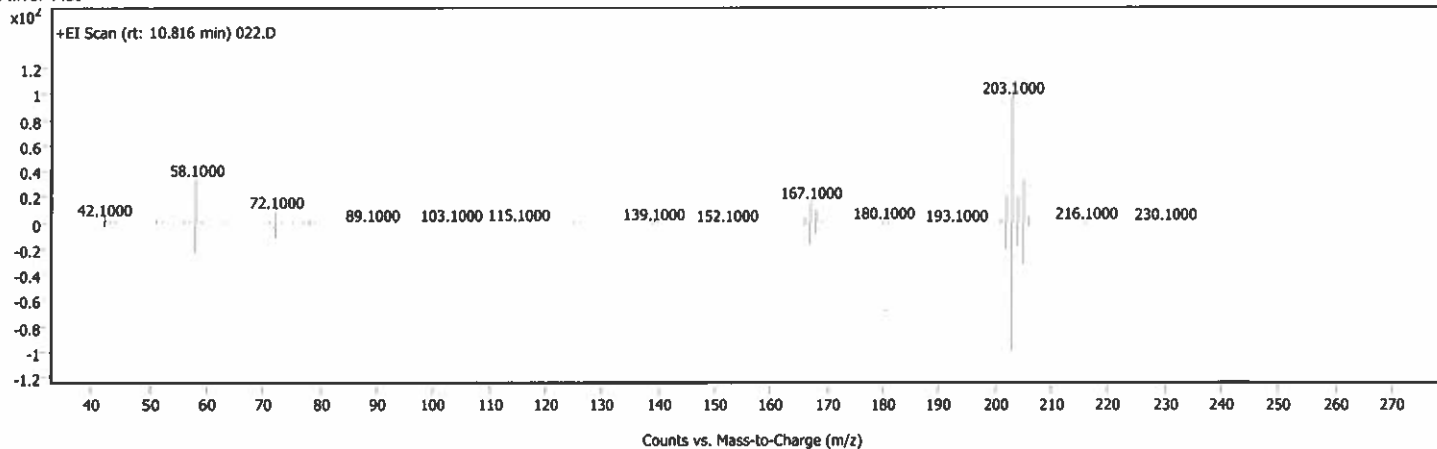
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	99.01	aafs.l	MATRIX 2 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

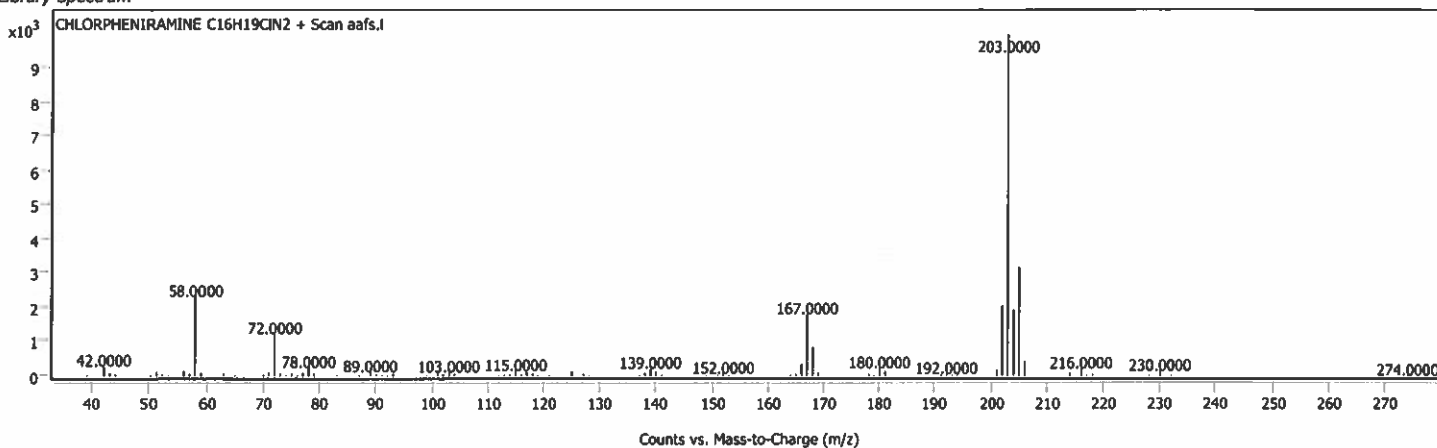
## Observed Spectrum



## Mirror Plot



## Library Spectrum



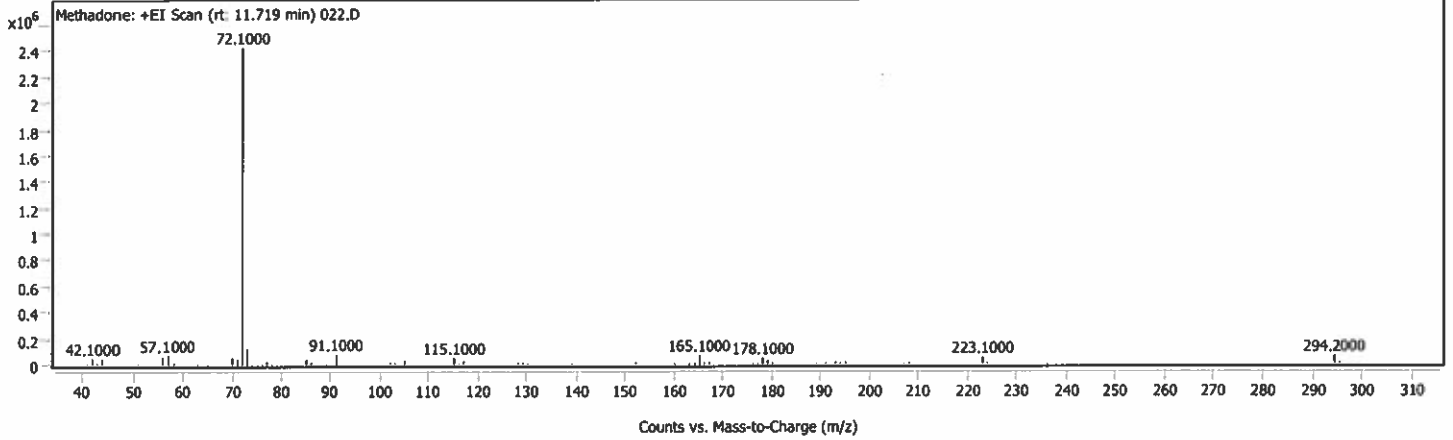
# Analysis Report

+ Scan (rt: 11.719 min)

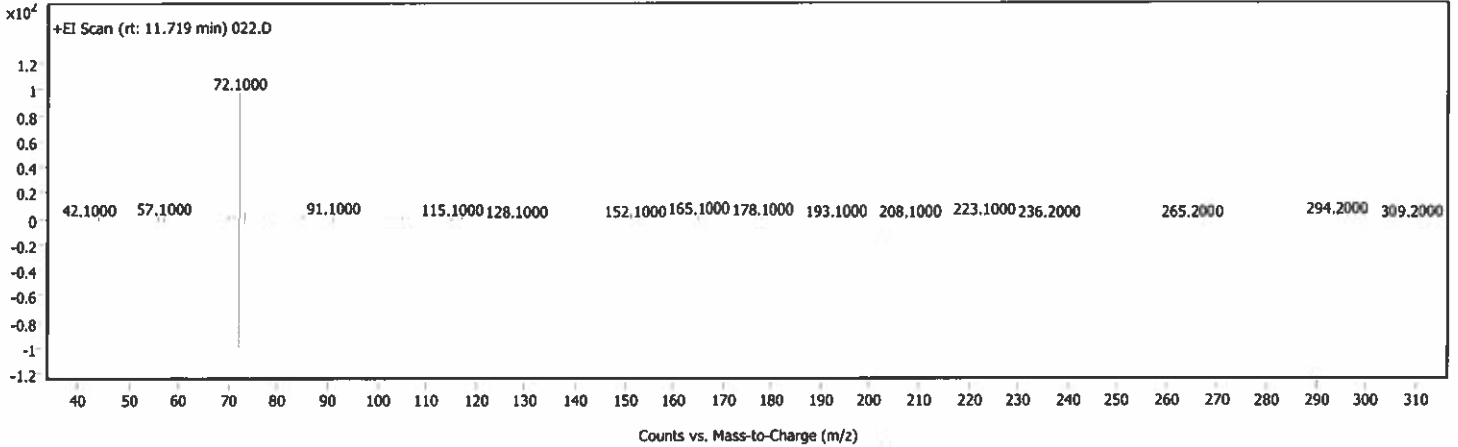
Methadone; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C21H27NO	97.27	SWGDRUG 3.3.L	MATRIX 2 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

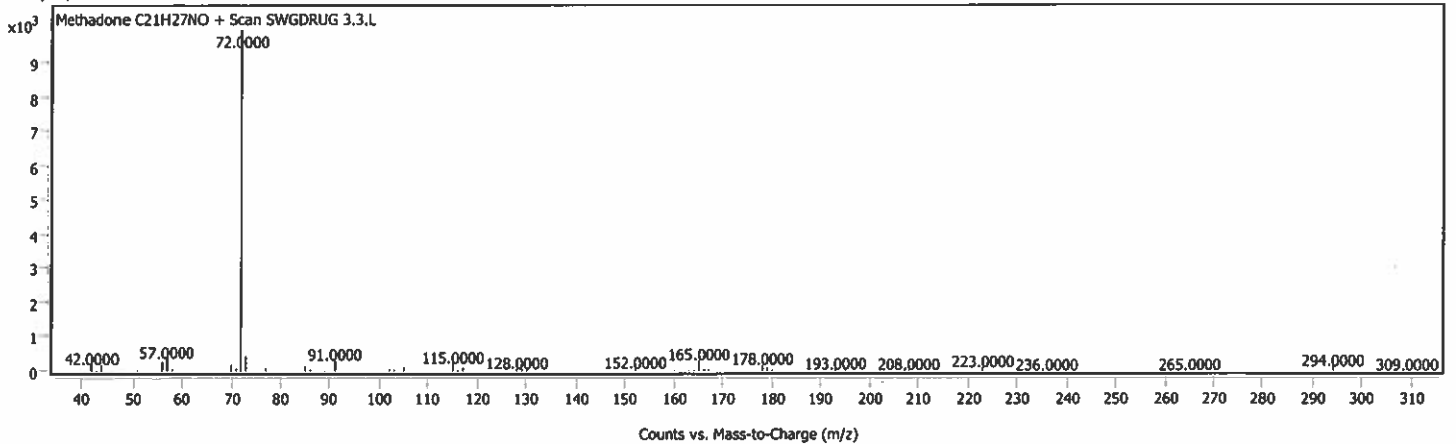
## Observed Spectrum



## Mirror Plot



## Library Spectrum



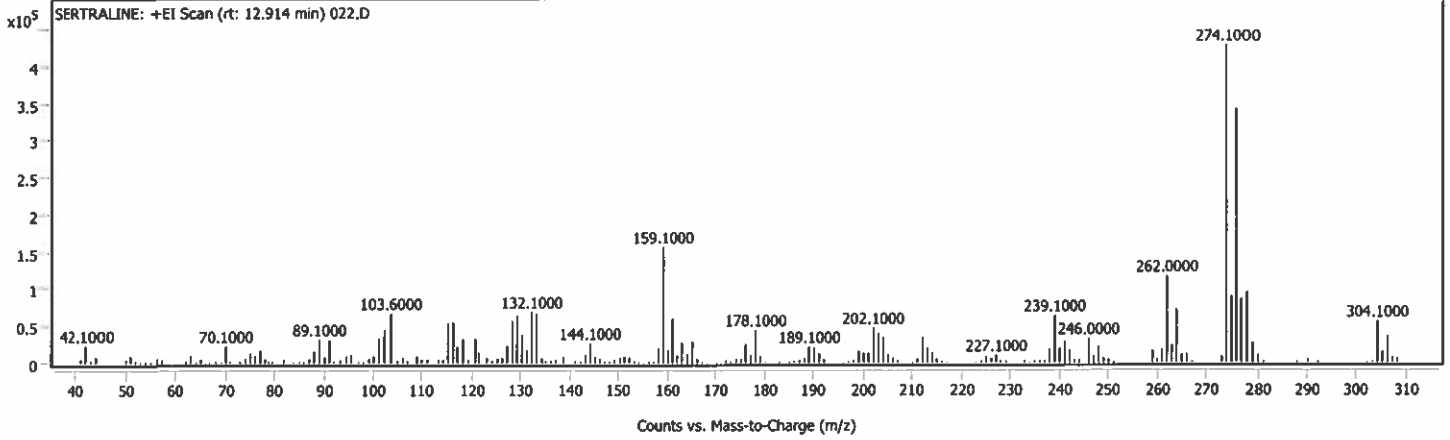
# Analysis Report

+ Scan (rt: 12.914 min)

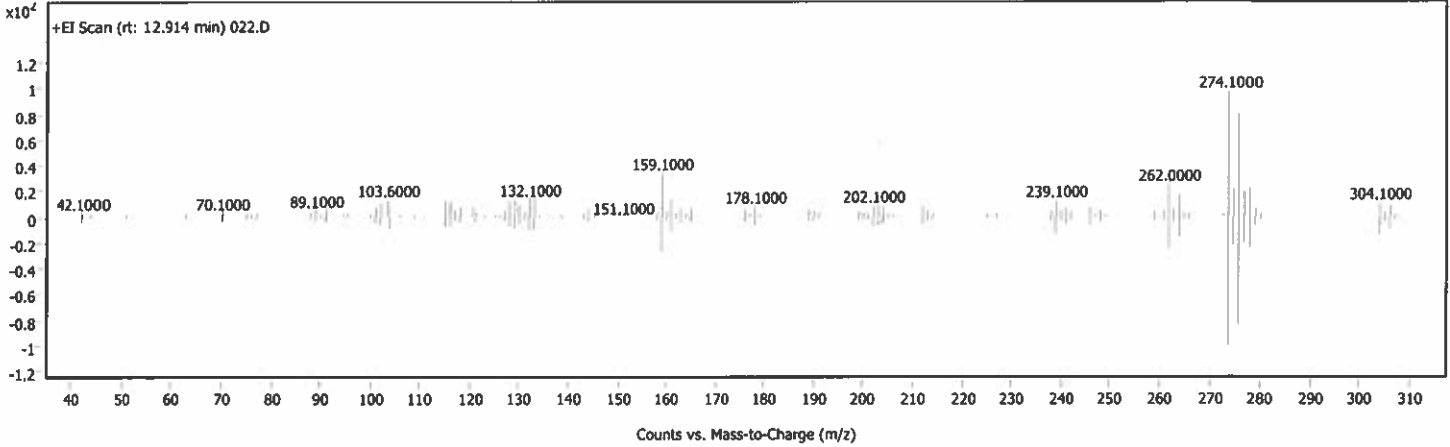
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	91.98	aafs.I	MATRIX 2 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

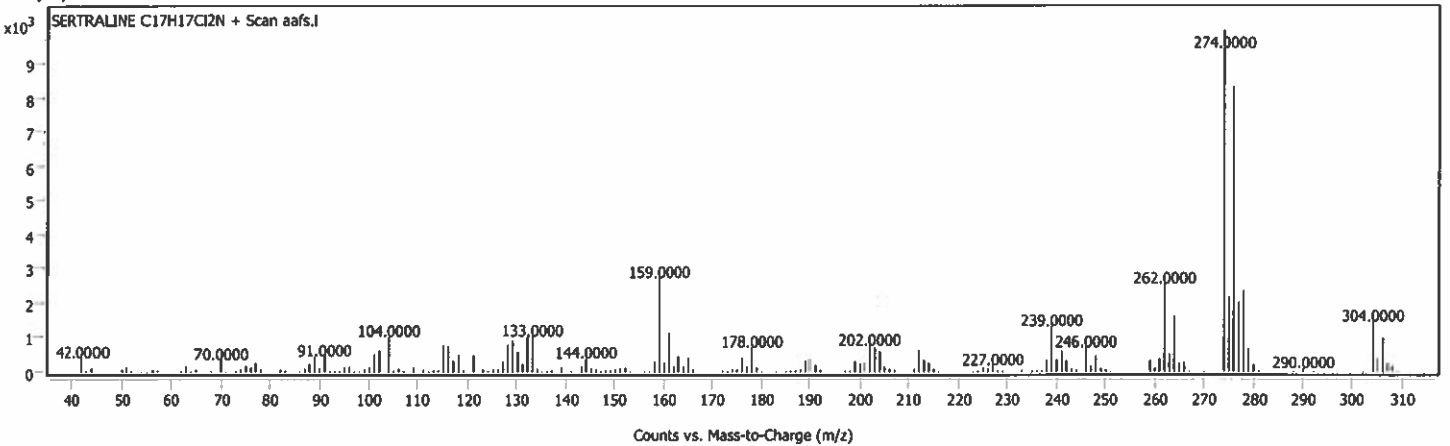
## Observed Spectrum



## Mirror Plot

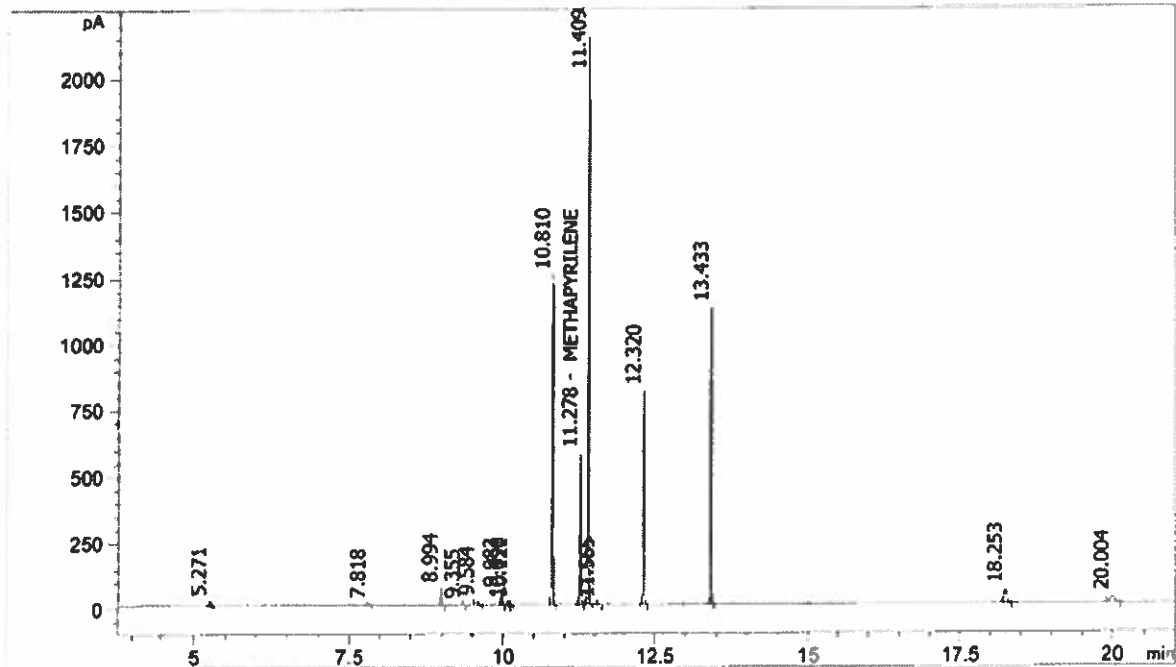


## Library Spectrum



Injection Date: 9/28/2021 11:04:42 PM Seq Line: 24  
 Sample Name: MATRIX 2 W/ IH CTL - -> Vial 25  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	654.601	569.257	1.000000	METHAPYRILENE

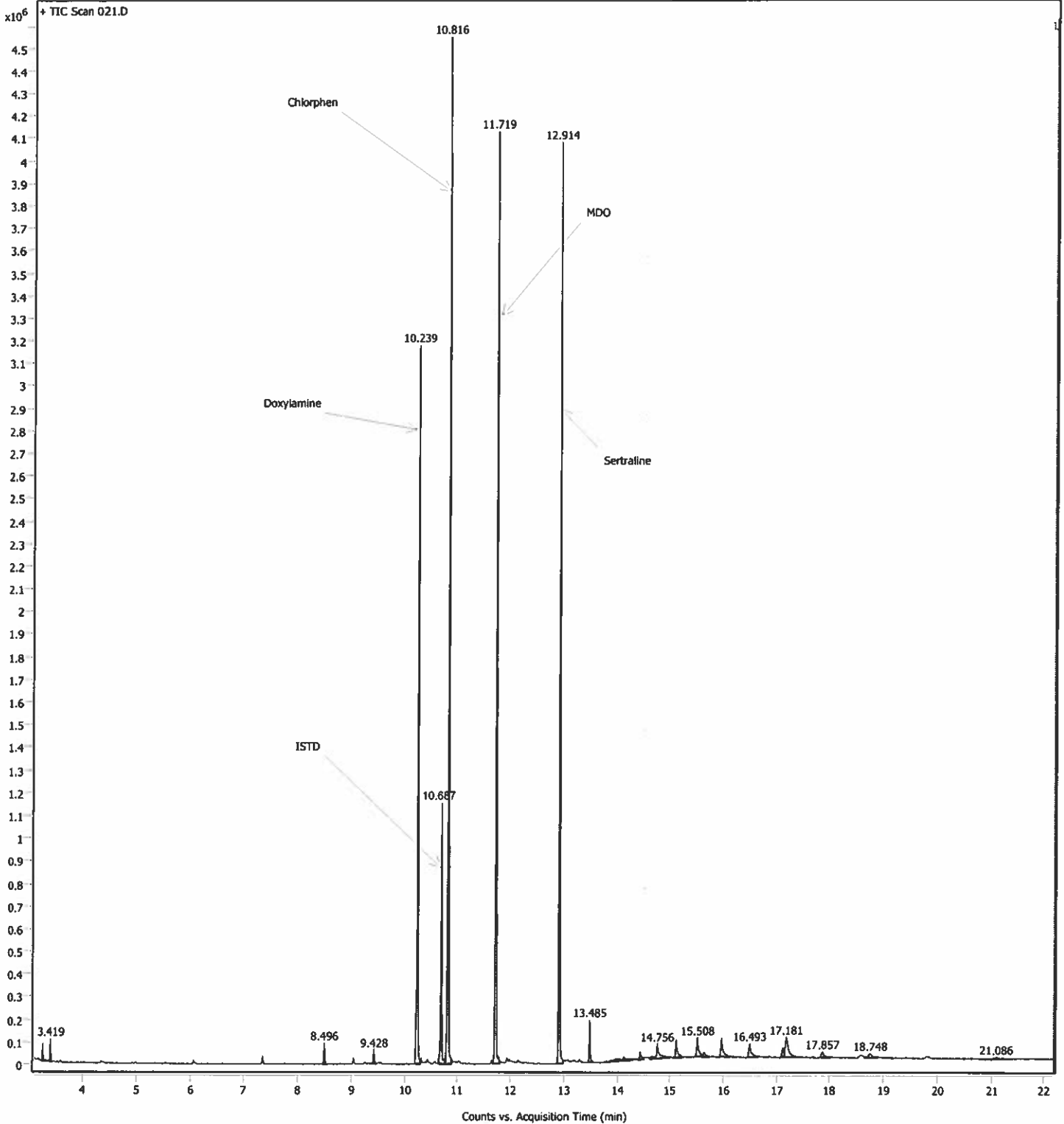


# Analysis Report

## Sample Information

Sample Name	MATRIX 1 W/ IH CTL - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\021.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 10:43:02 PM (UTC-04:00)
Position	48	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

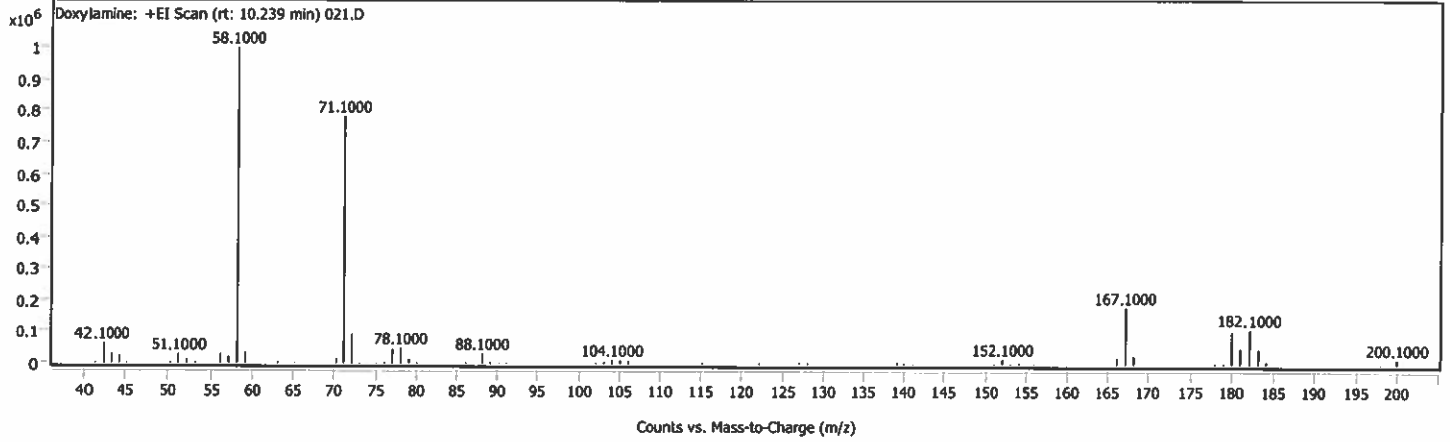
## Sample Spectra

+ Scan (rt: 10.239 min)

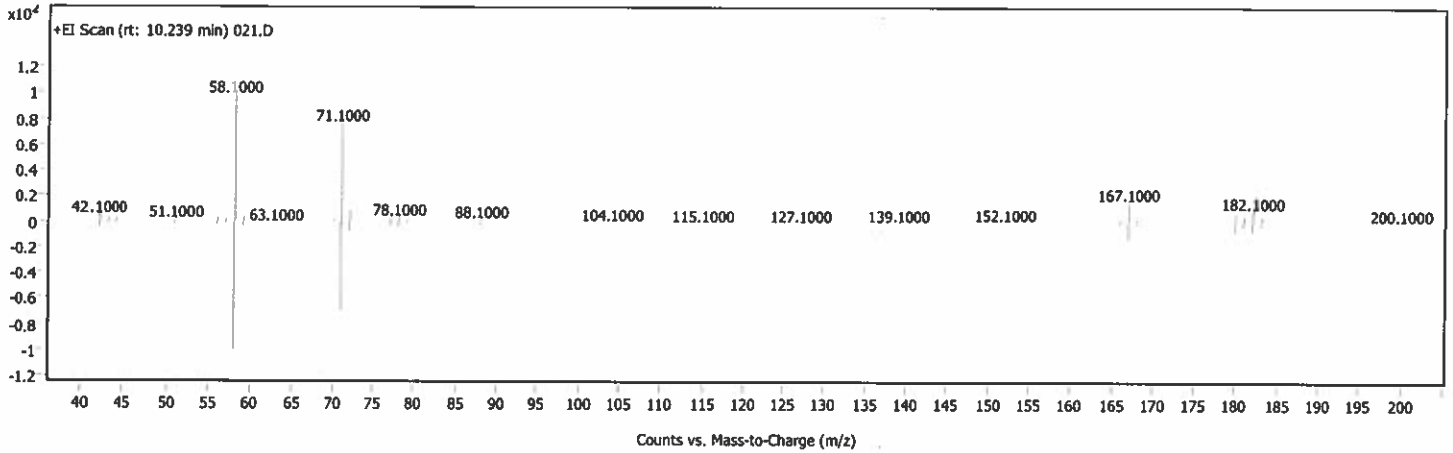
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.91	SWGDRUG 3.3.L	MATRIX 1 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

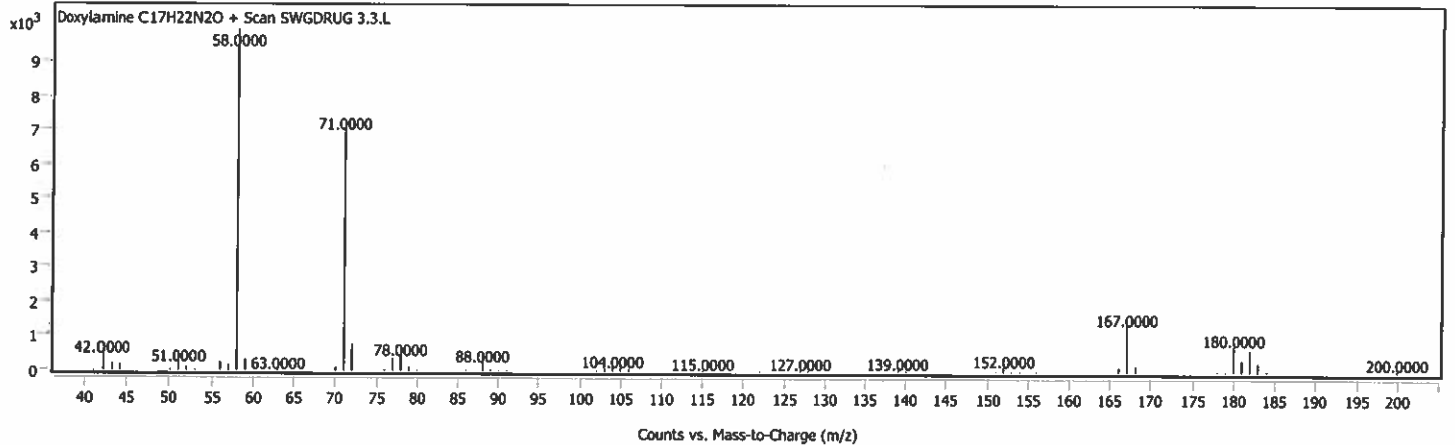
### Observed Spectrum



### Mirror Plot



### Library Spectrum



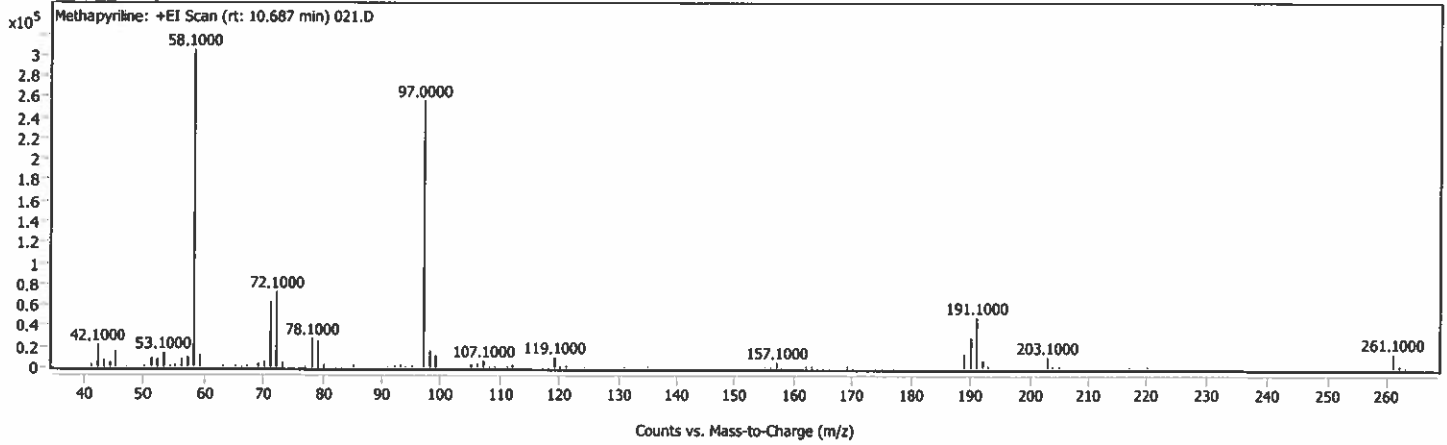
# Analysis Report

+ Scan (rt: 10.687 min)

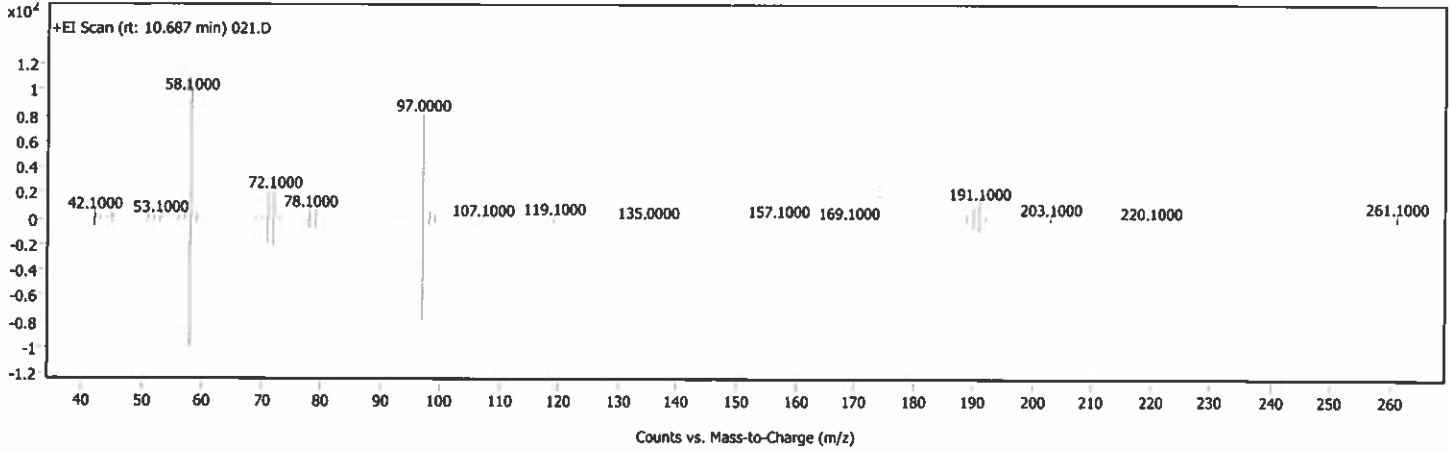
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.45	SWGDRUG 3.3.L	MATRIX 1 W/ IH CTL - BLD	KM/QC DATA IN XLAZINE VALIDATION

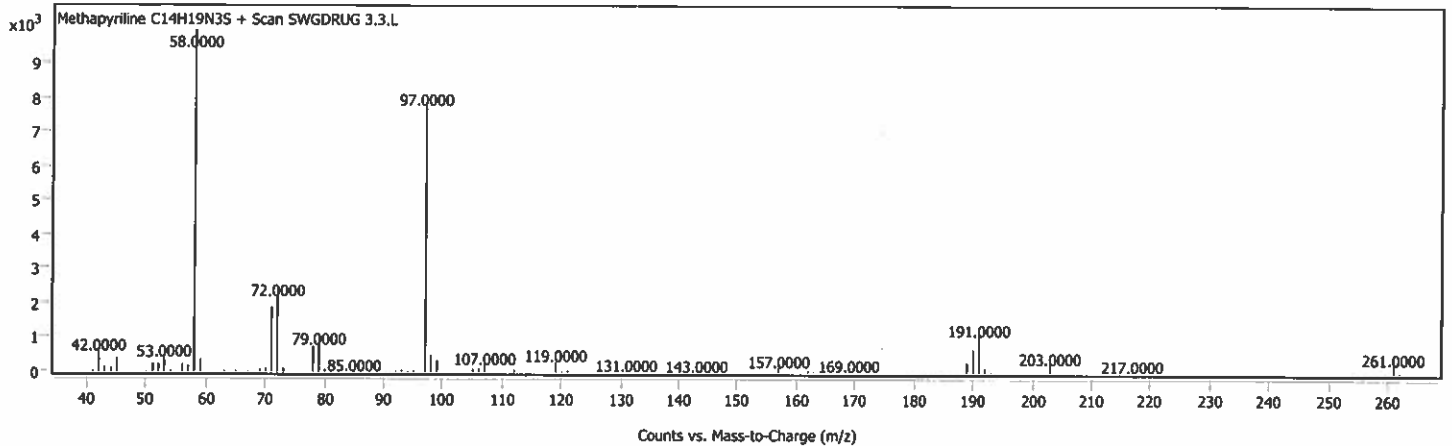
## Observed Spectrum



## Mirror Plot



## Library Spectrum



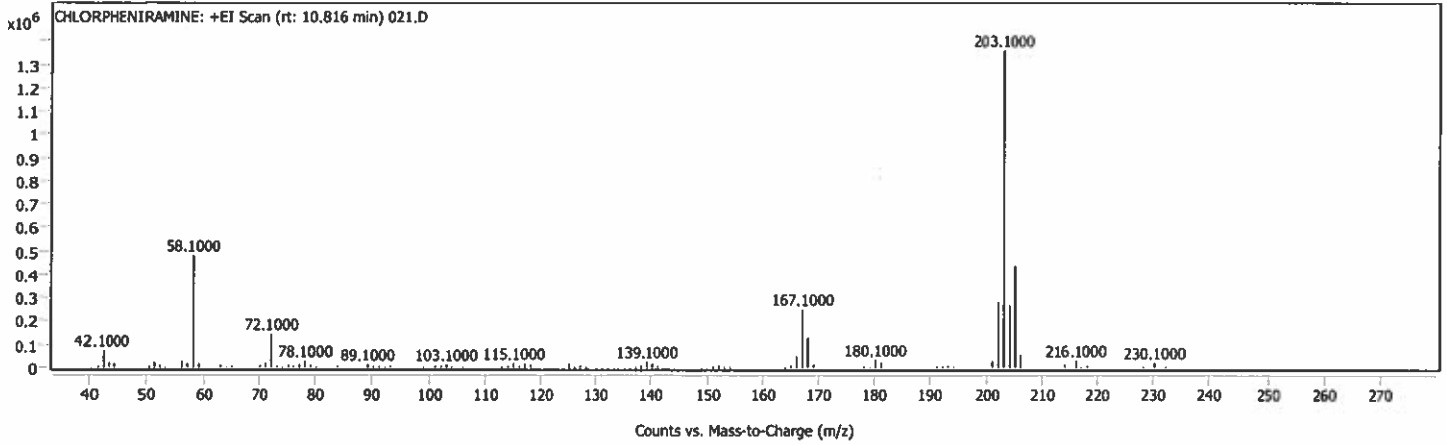
# Analysis Report

+ Scan (rt: 10.816 min)

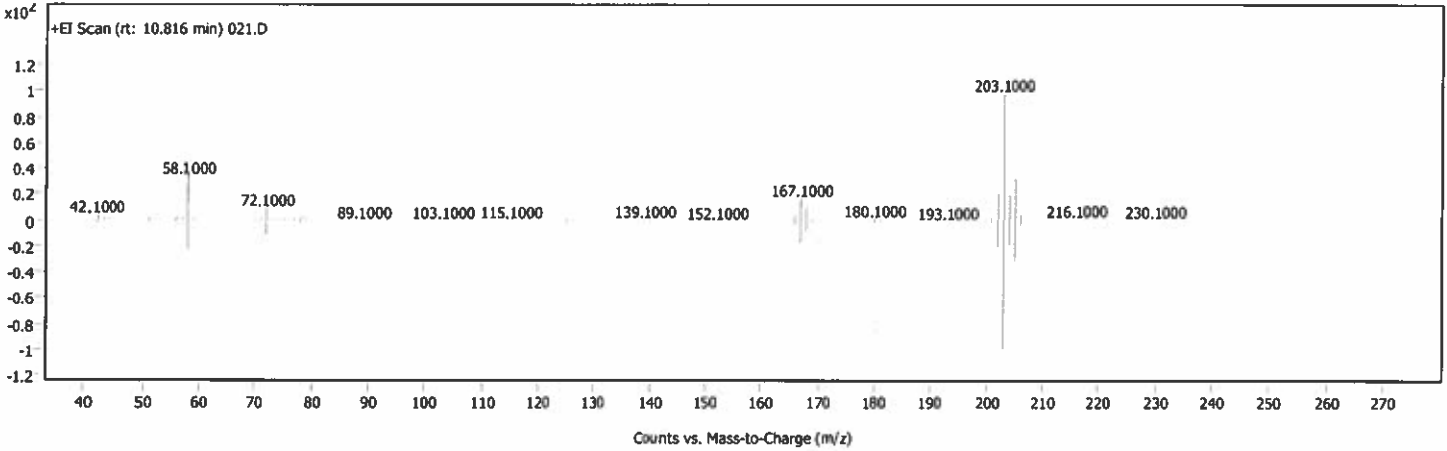
CHLORPHENIRAMINE; C16H19CIN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19CIN2	99.12	aafs.I	MATRIX 1 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

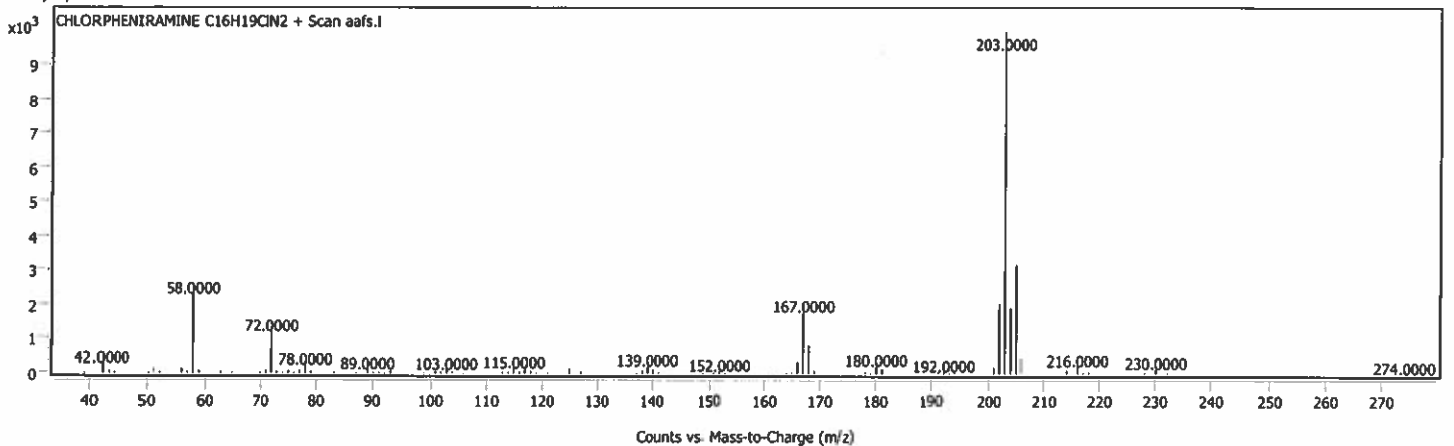
### Observed Spectrum



### Mirror Plot



### Library Spectrum



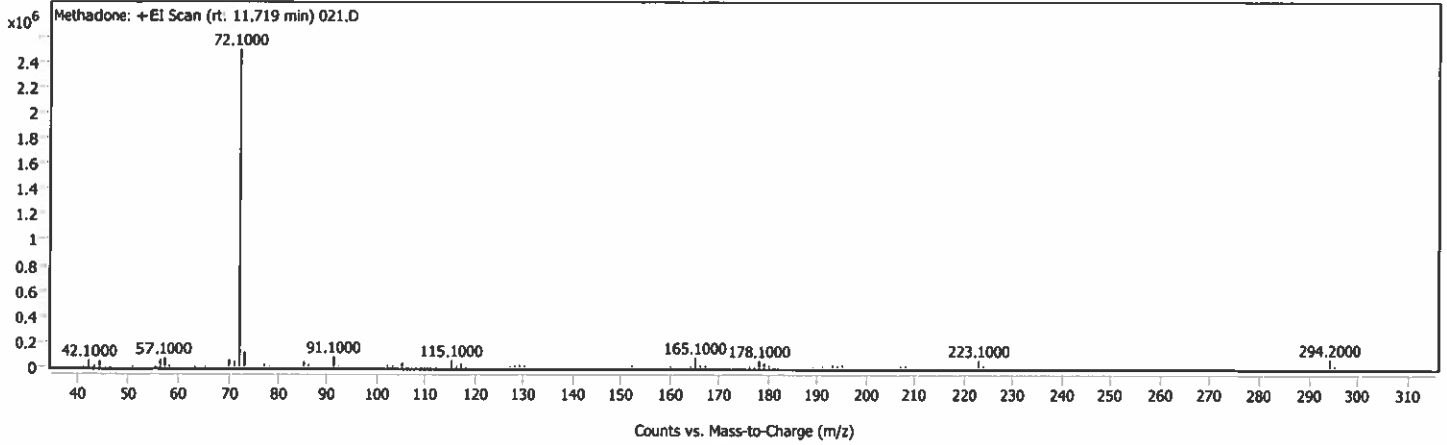
# Analysis Report

+ Scan (rt: 11.719 min)

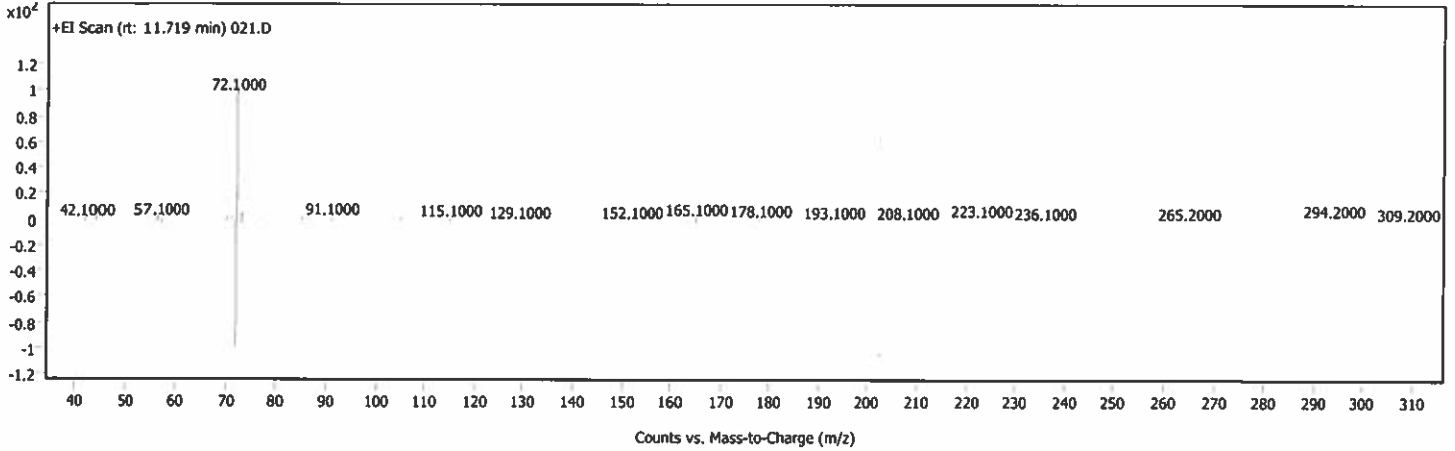
Methadone; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C21H27NO	97.44	SWGDRUG 3.3.L	MATRIX 1 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

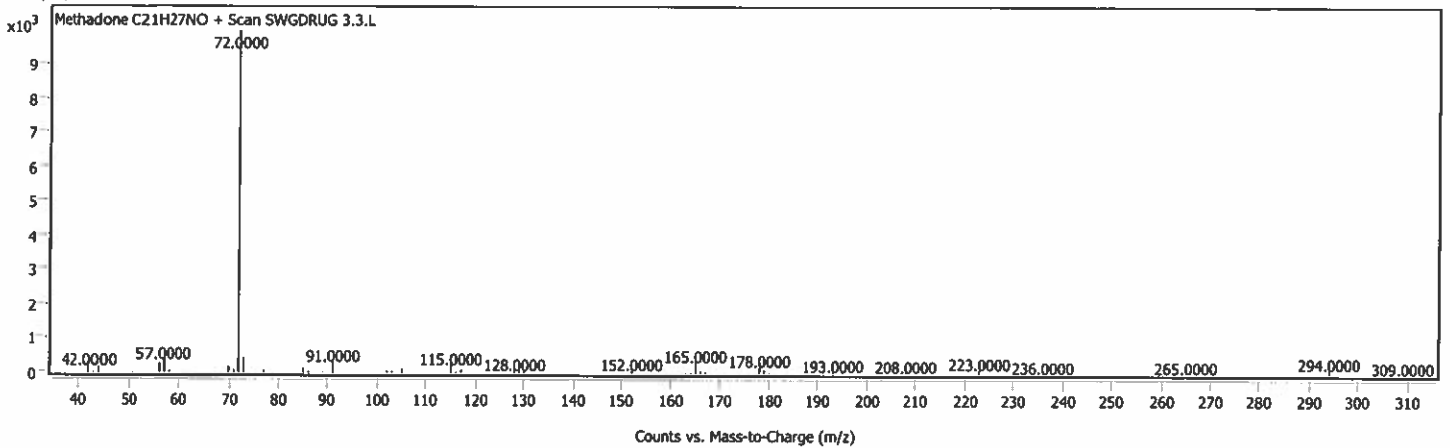
## Observed Spectrum



## Mirror Plot



## Library Spectrum



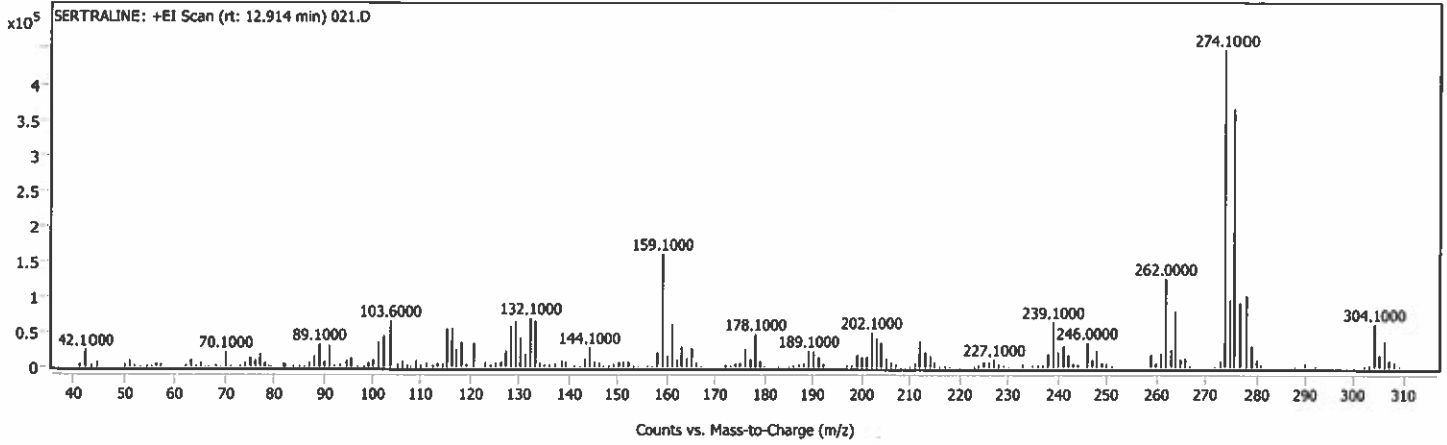
# Analysis Report

+ Scan (rt: 12.914 min)

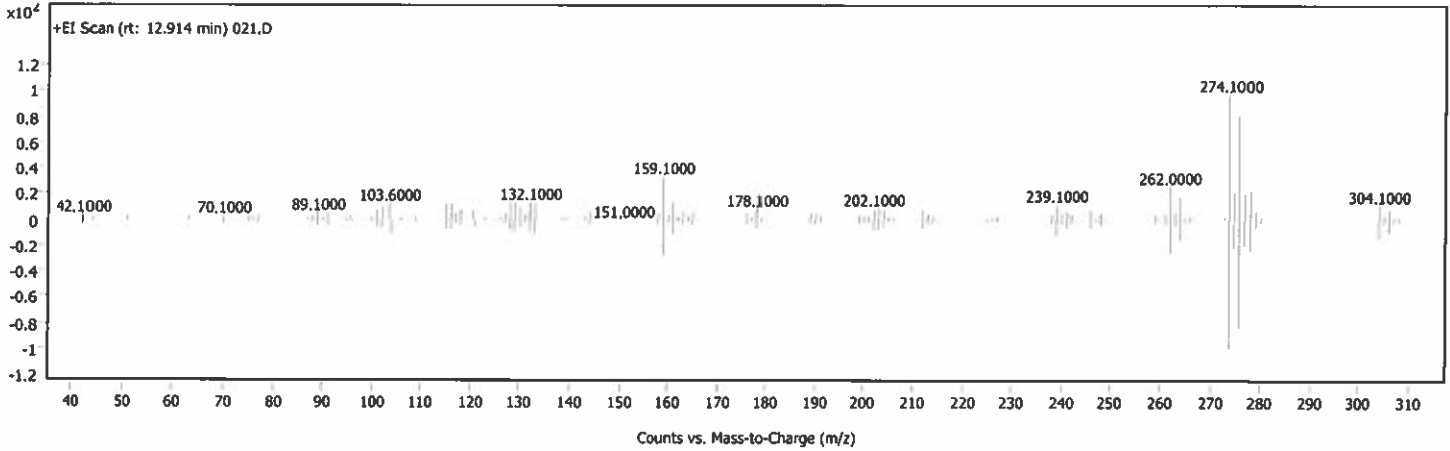
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	92.28	aafs.l	MATRIX 1 W/ IH CTL - BLD	KM/QC DATA IN XYLAZINE VALIDATION

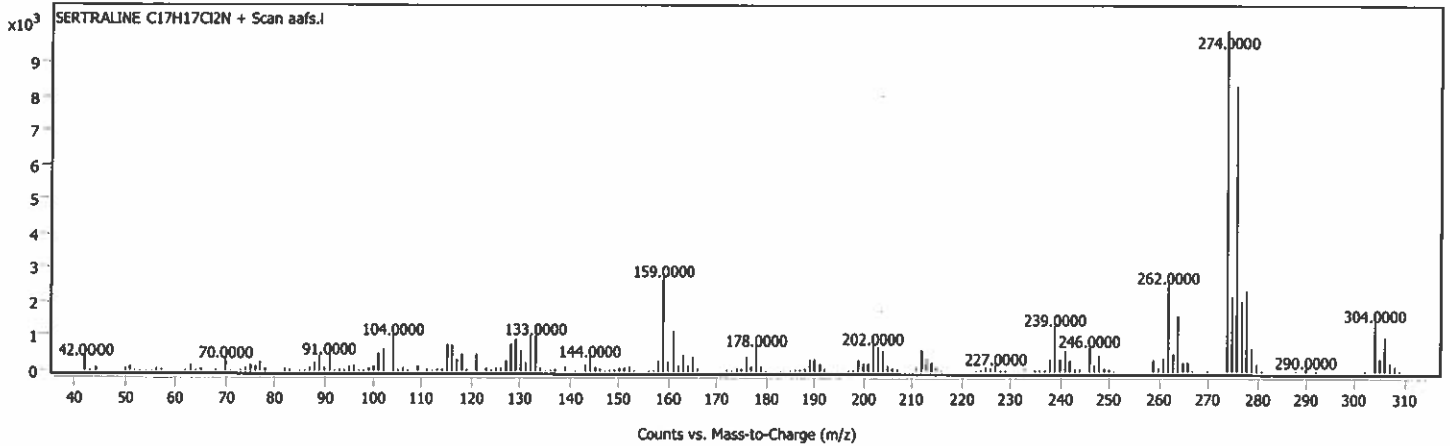
## Observed Spectrum



## Mirror Plot

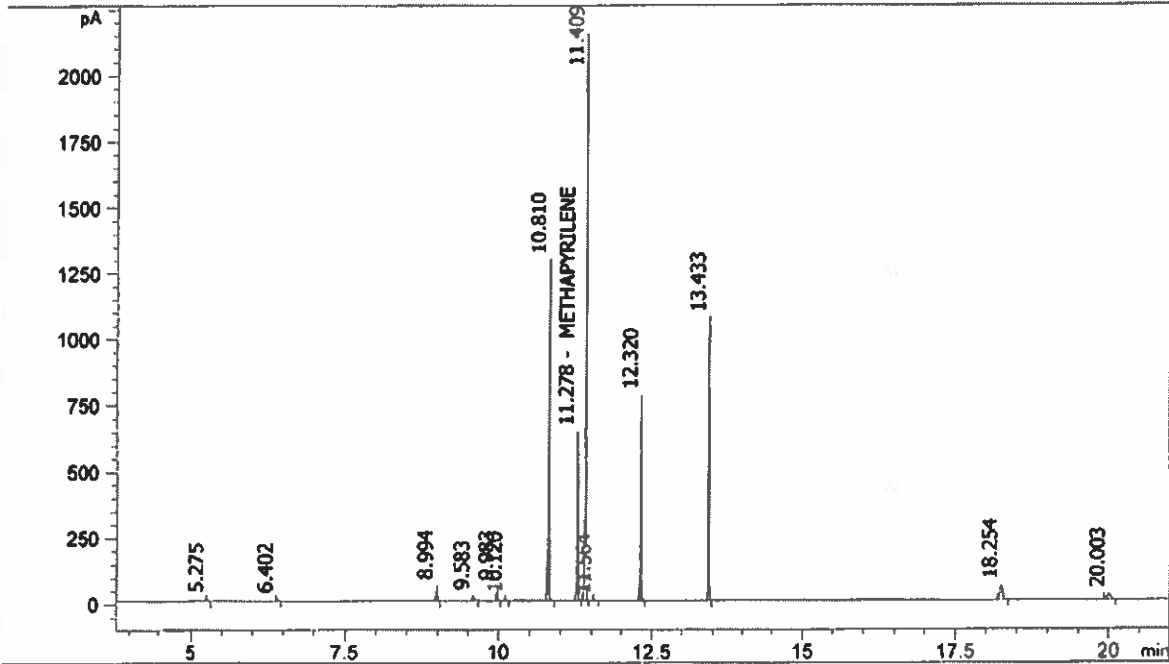


## Library Spectrum



Injection Date: 9/28/2021 10:40:26 PM Seq Line: 23  
 Sample Name: MATRIX 1 W/ IH CTL -> Vial 24  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



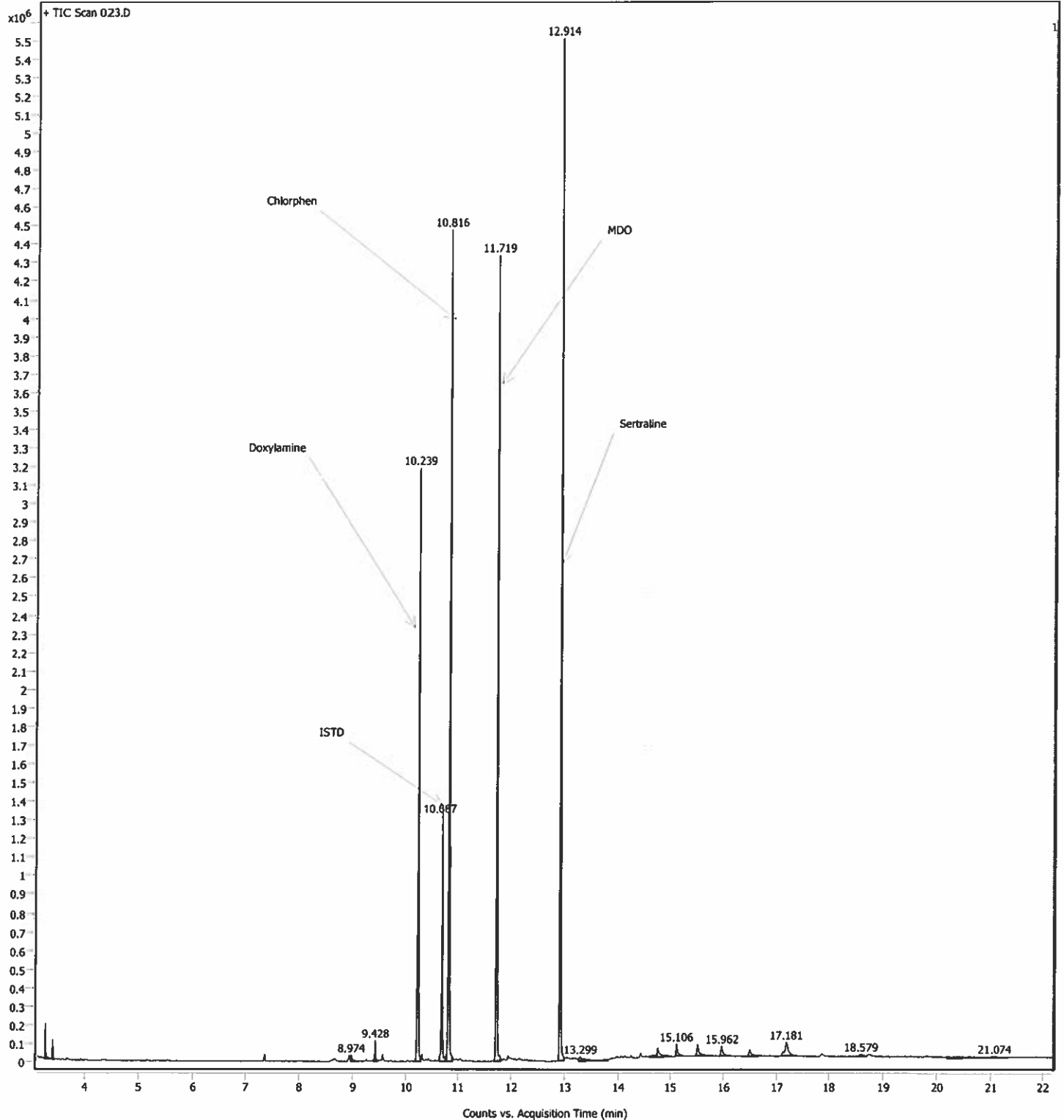
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	747.753	639.091	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 W/ IH CTL - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\023.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 11:34:42 PM (UTC-04:00)
Position	50	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

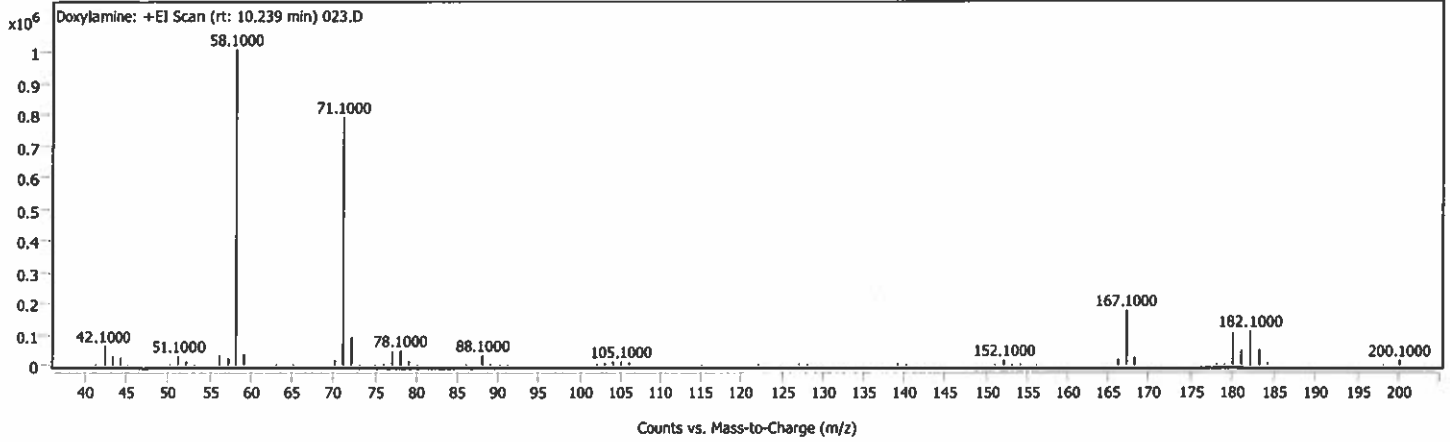
## Sample Spectra

+ Scan (rt: 10.239 min)

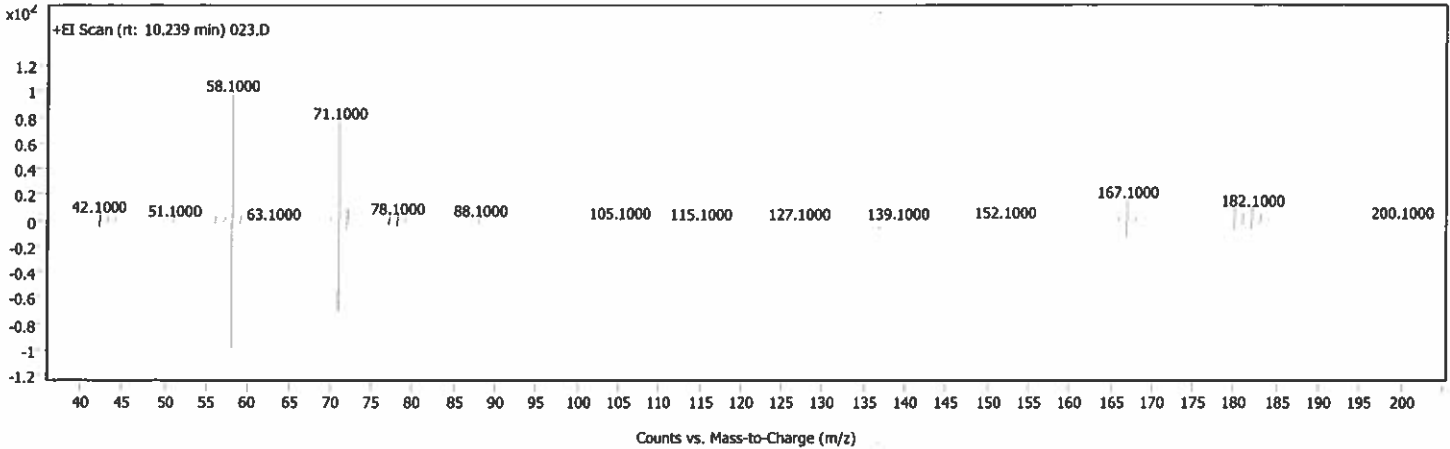
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.80	SWGDRUG 3.3.L	MATRIX 6 W/ IH CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

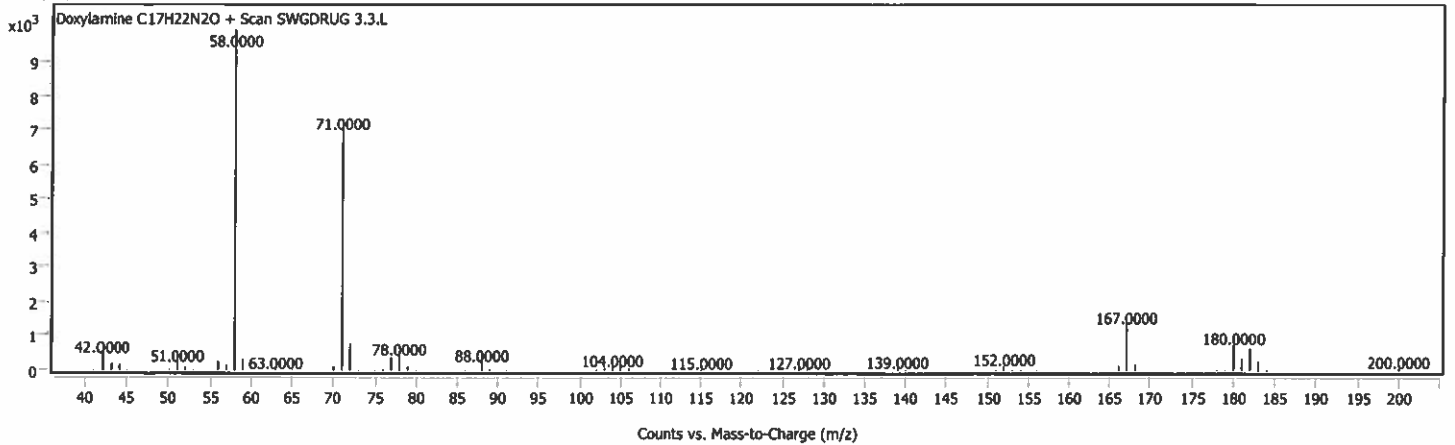
### Observed Spectrum



### Mirror Plot



### Library Spectrum



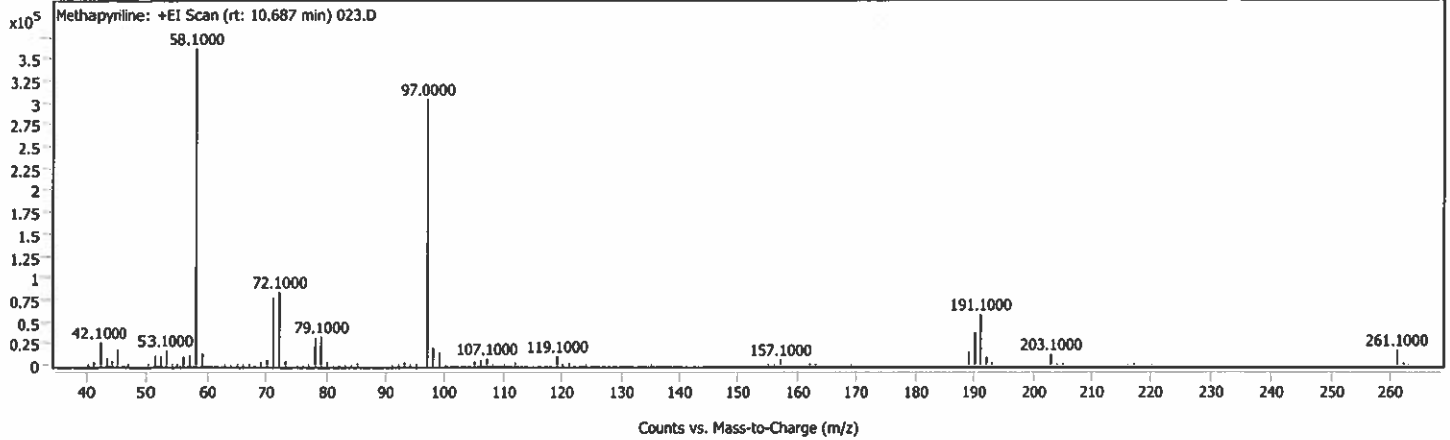
# Analysis Report

+ Scan (rt: 10.687 min)

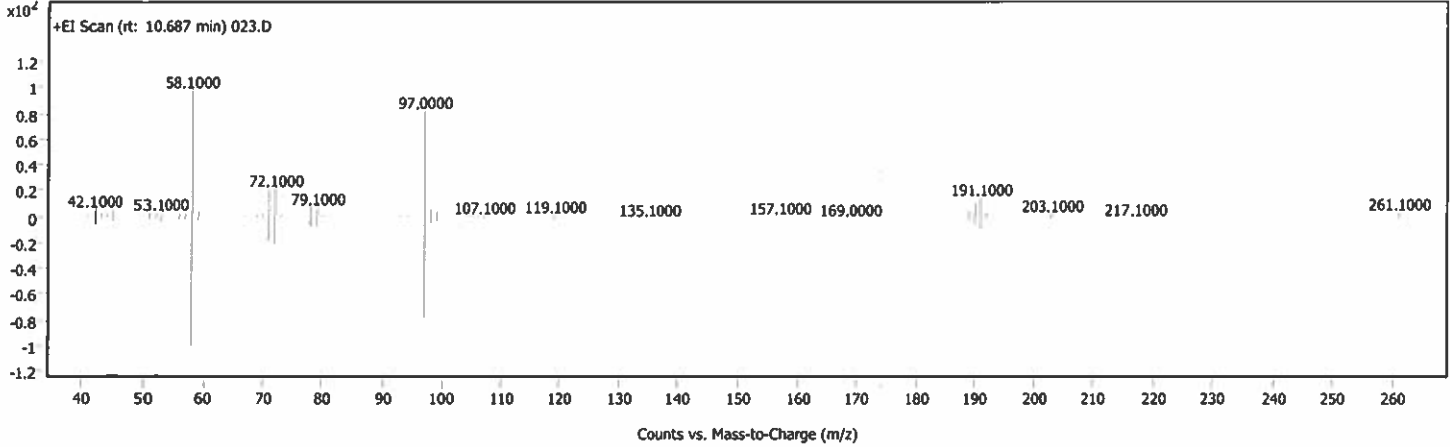
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.59	SWGDRUG 3.3.L	MATRIX 6 W/ IH CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

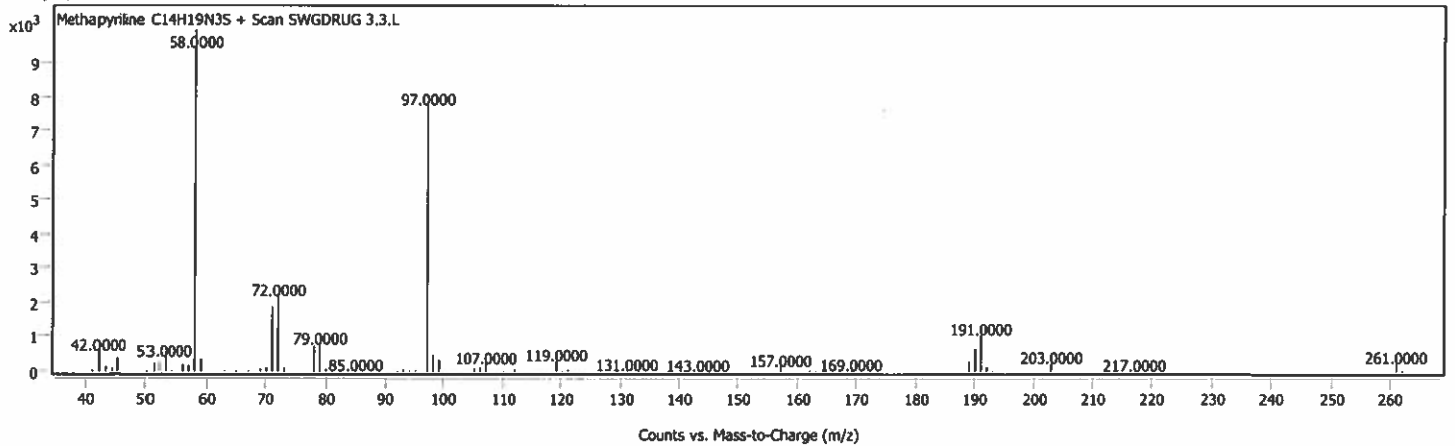
## Observed Spectrum



## Mirror Plot



## Library Spectrum



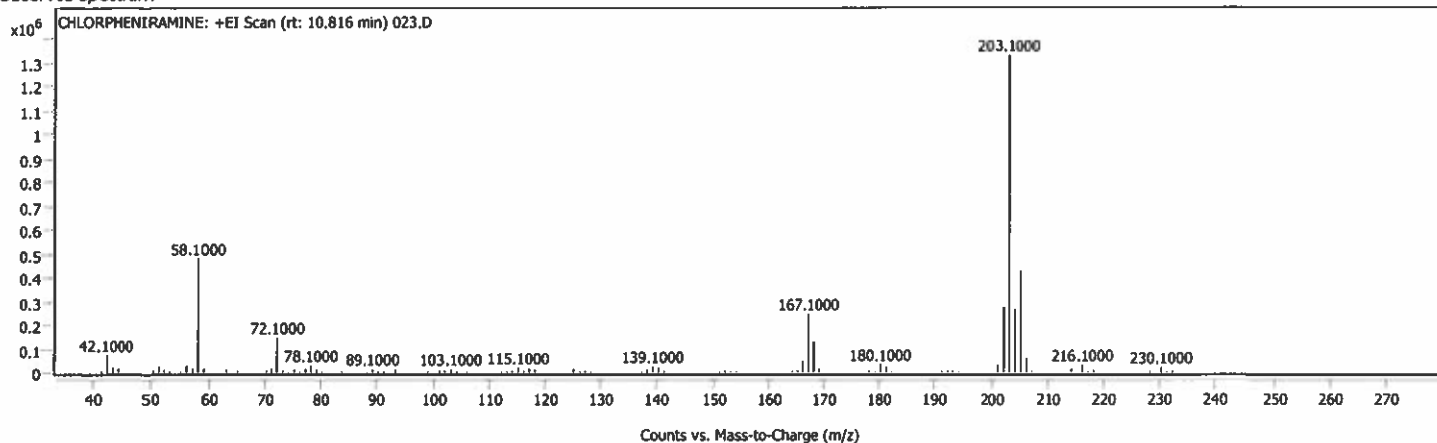
# Analysis Report

+ Scan (rt: 10.816 min)

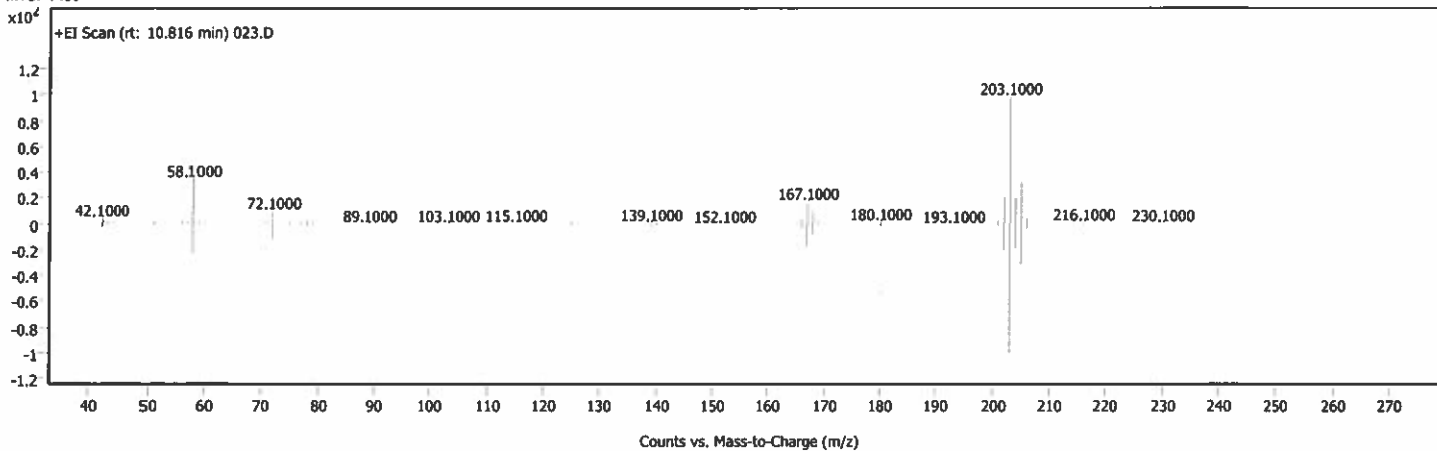
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	99.03	aafs.I	MATRIX 6 W/ IH CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

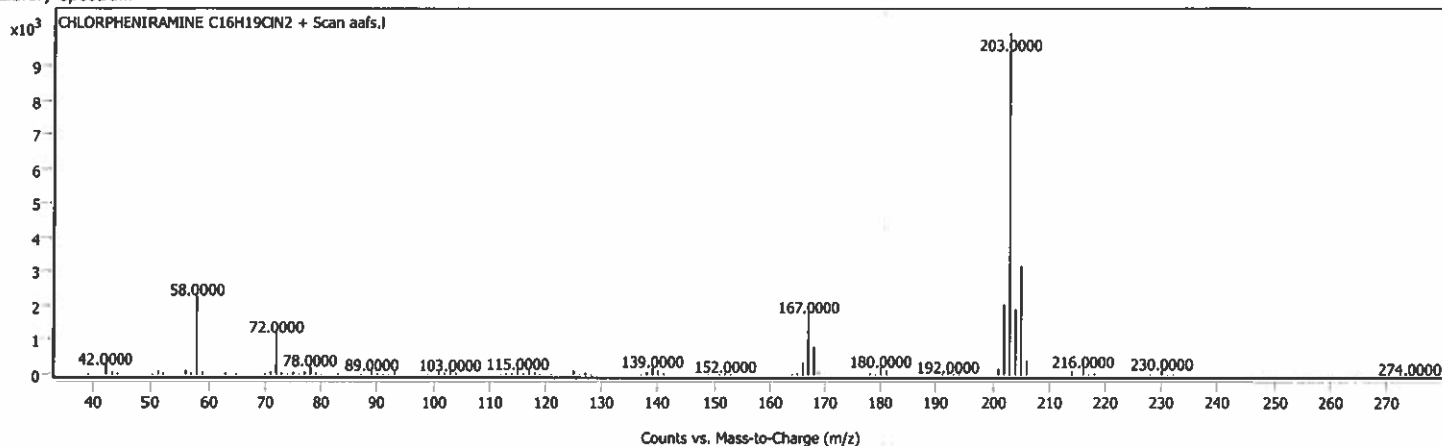
## Observed Spectrum



## Mirror Plot



## Library Spectrum



# Analysis Report

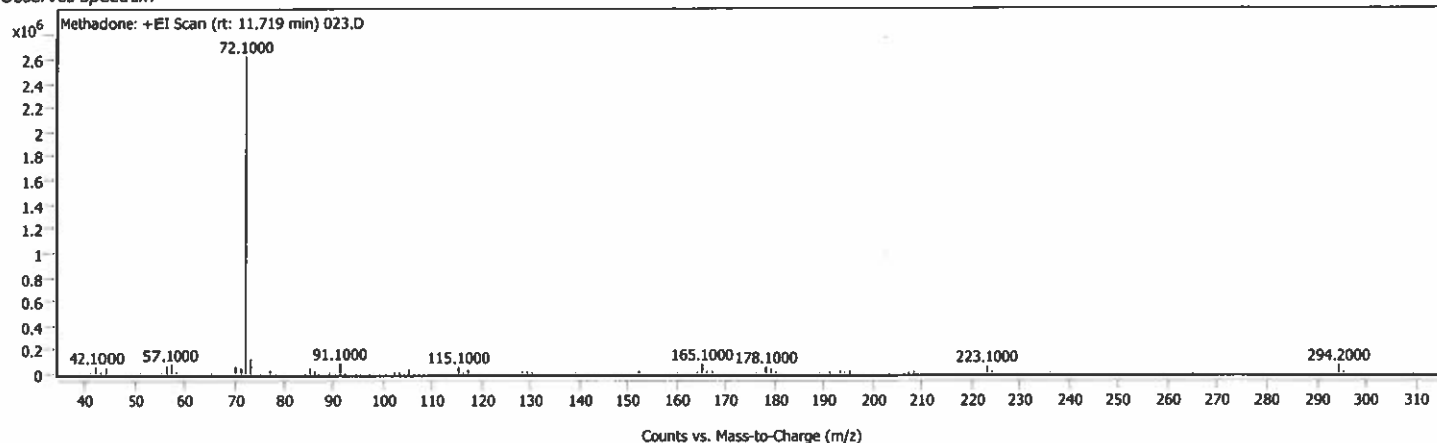


+ Scan (rt: 11.719 min)

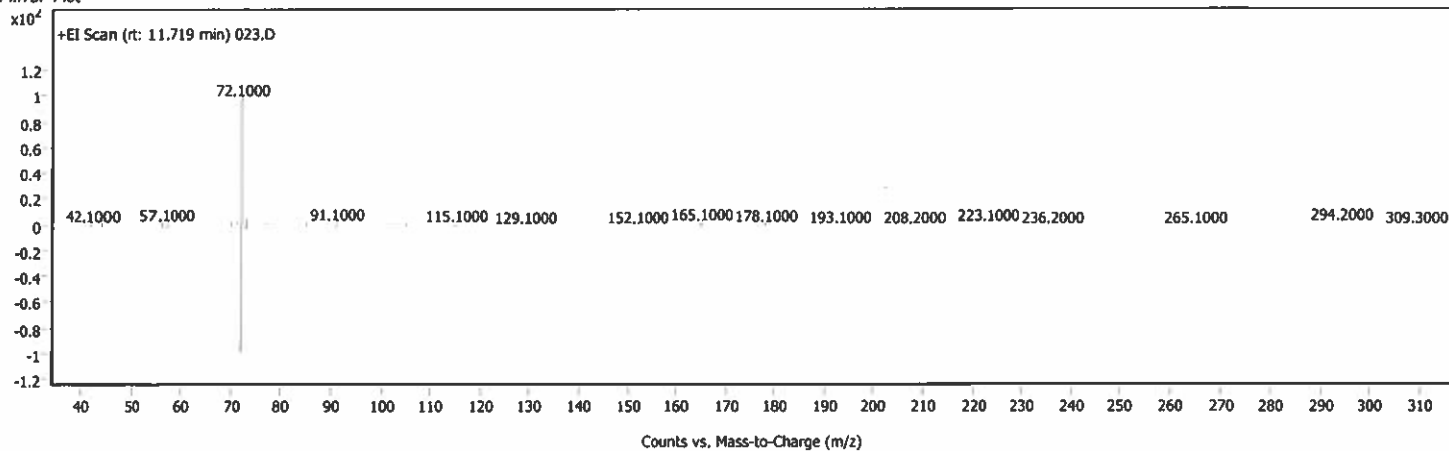
Methadone; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C21H27NO	97.37	SWGDRUG 3.3.L	MATRIX 6 W/ IH CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

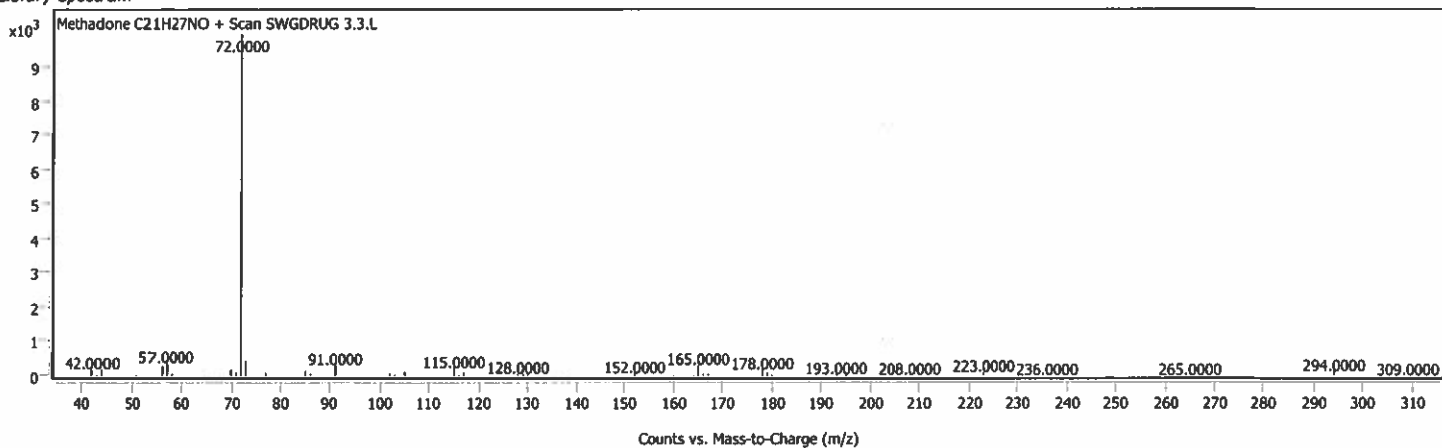
## Observed Spectrum



## Mirror Plot



## Library Spectrum



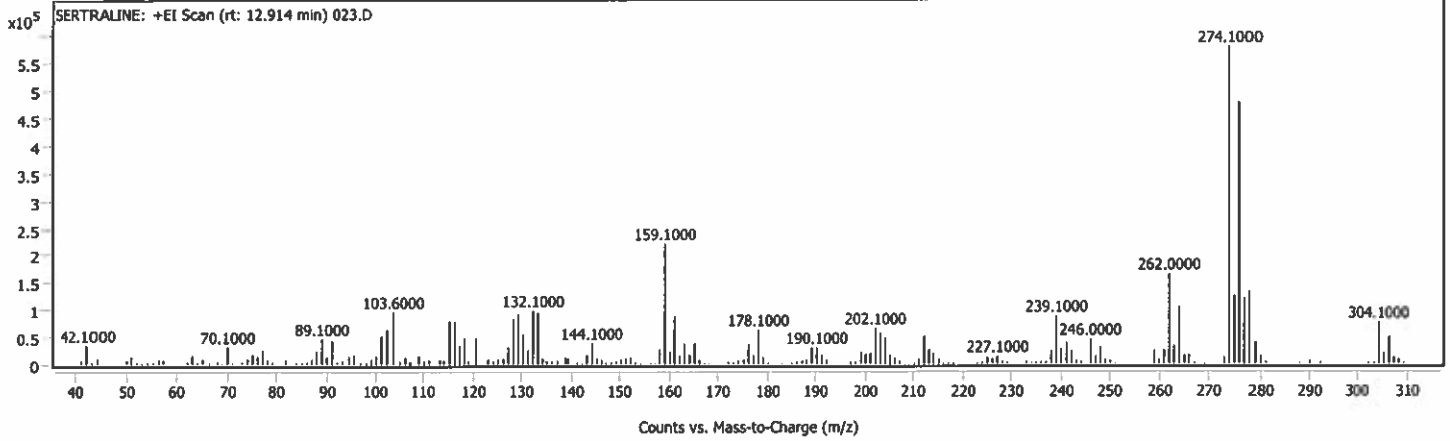
# Analysis Report

+ Scan (rt: 12.914 min)

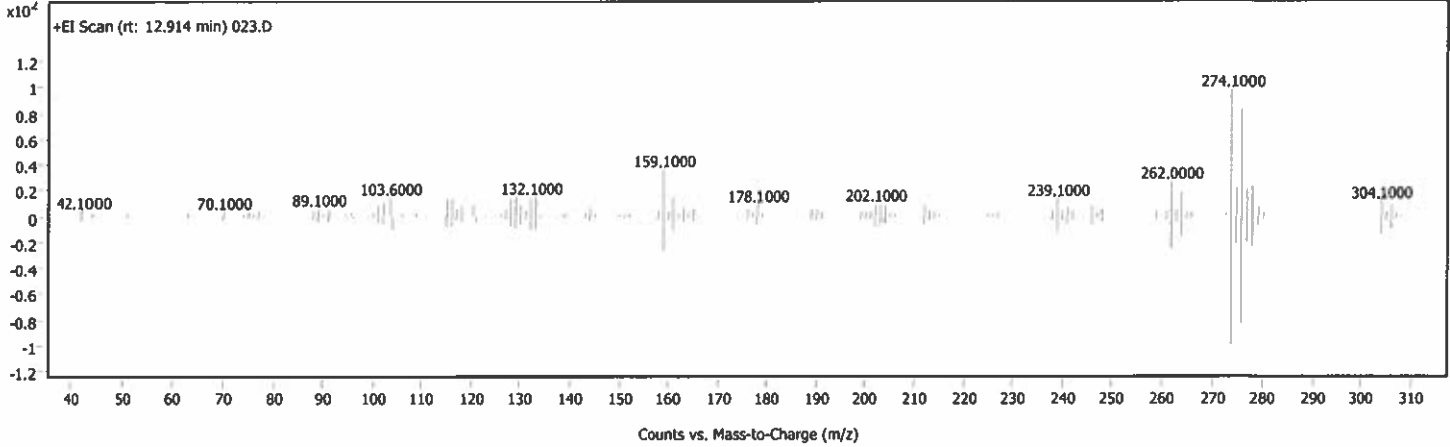
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	91.96	aafs.I	MATRIX 6 W/ IH CTL - URN	KM/QC DATA IN XYLAZINE VALIDATION

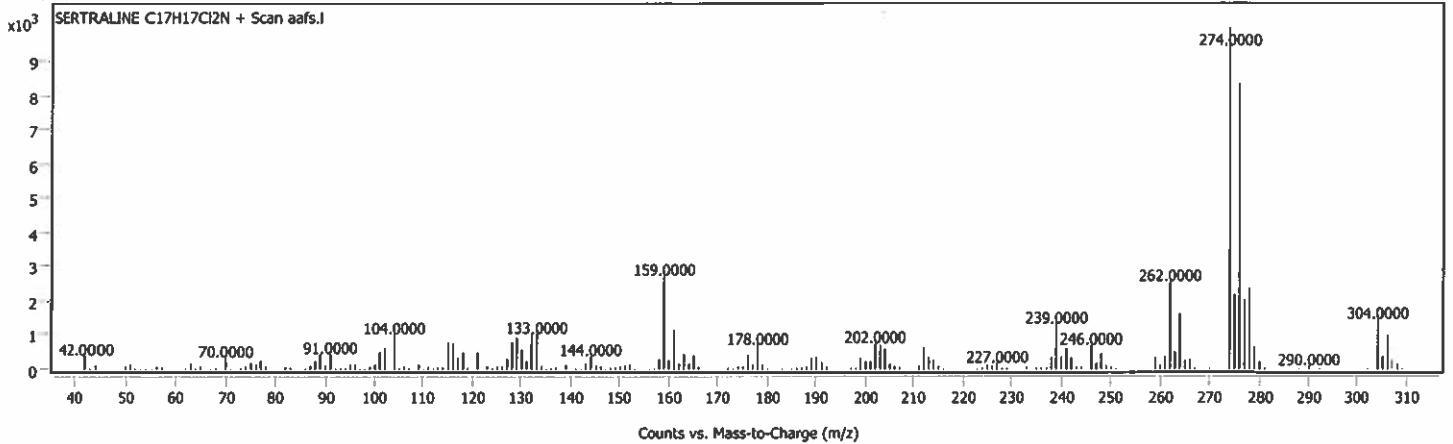
## Observed Spectrum



## Mirror Plot



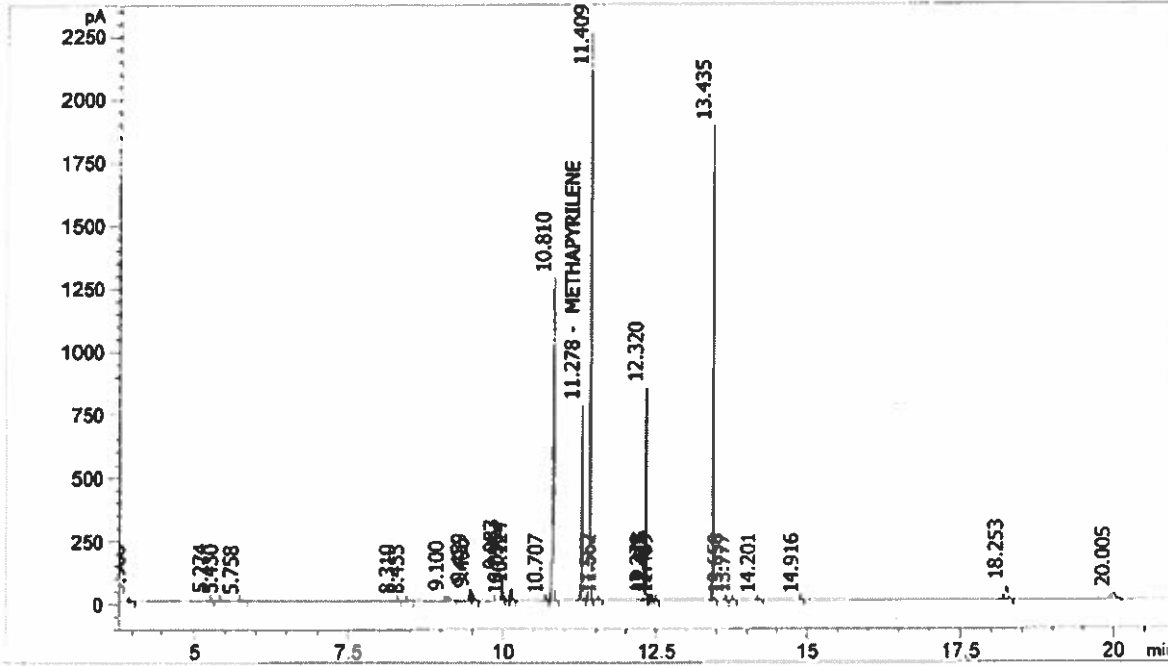
## Library Spectrum



Injection Date: 9/28/2021 11:29:06 PM  
 Sample Name: MATRIX 6 W/ IH CTL - ->  
 Sample Info:

Seq Line: 25  
 Vial 26

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



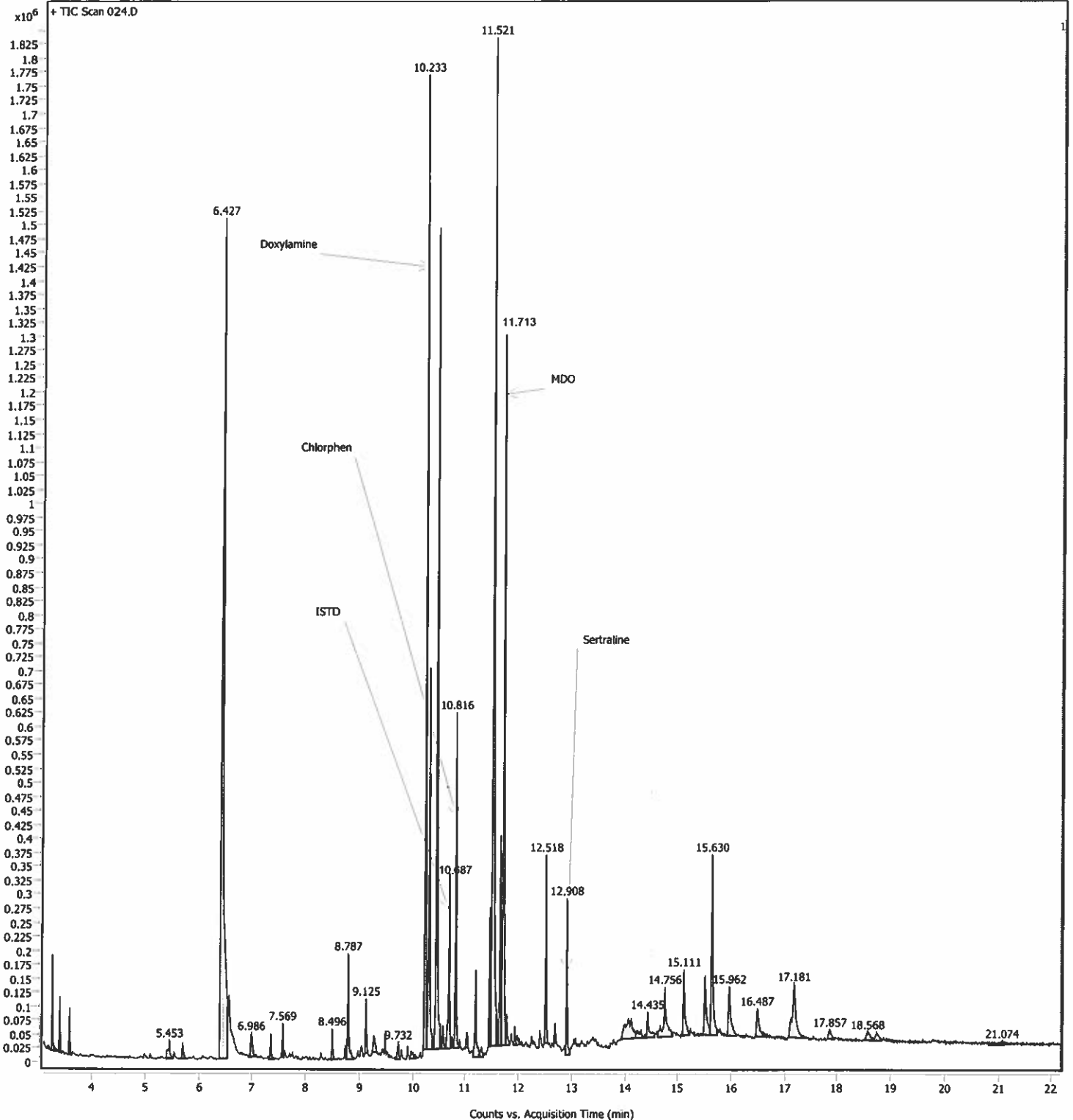
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.278	11.278	913.743	769.209	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 10 W/ IH CTL - LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\024.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 12:00:30 AM (UTC-04:00)
Position	51	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

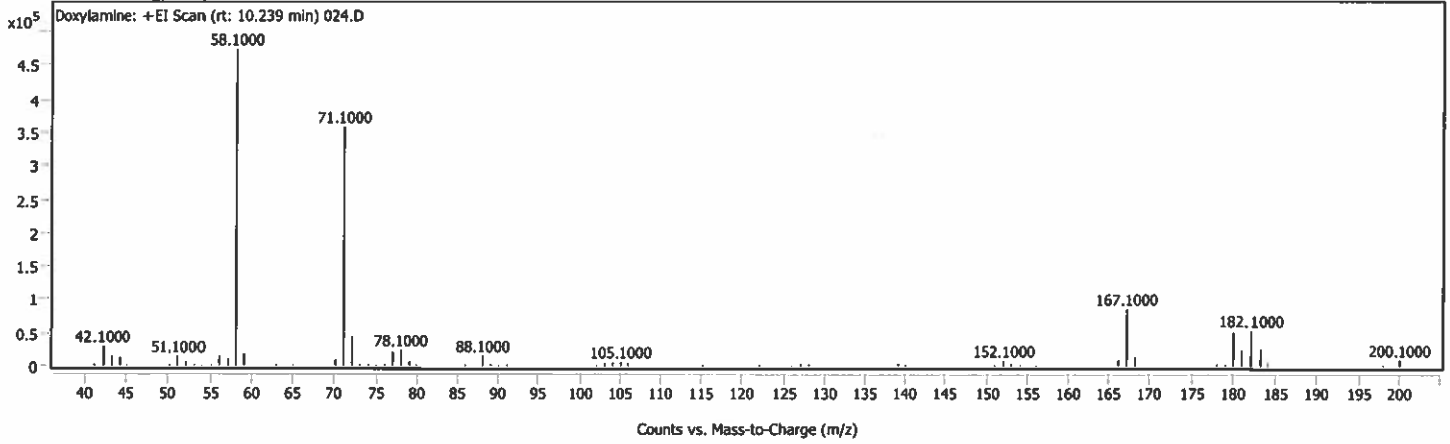
## Sample Spectra

+ Scan (rt: 10.239 min)

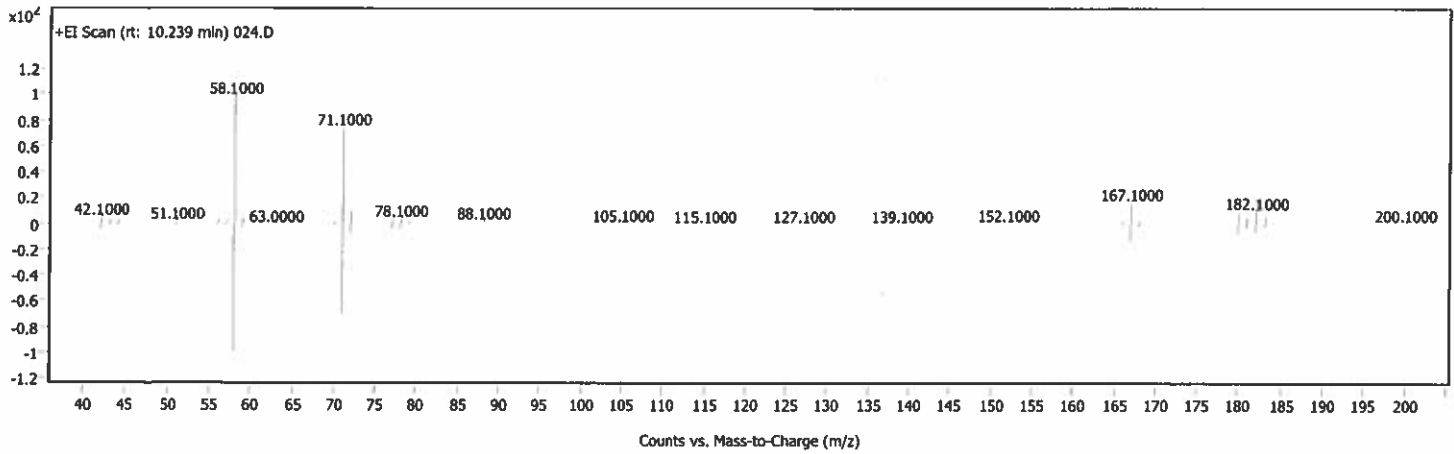
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.23	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

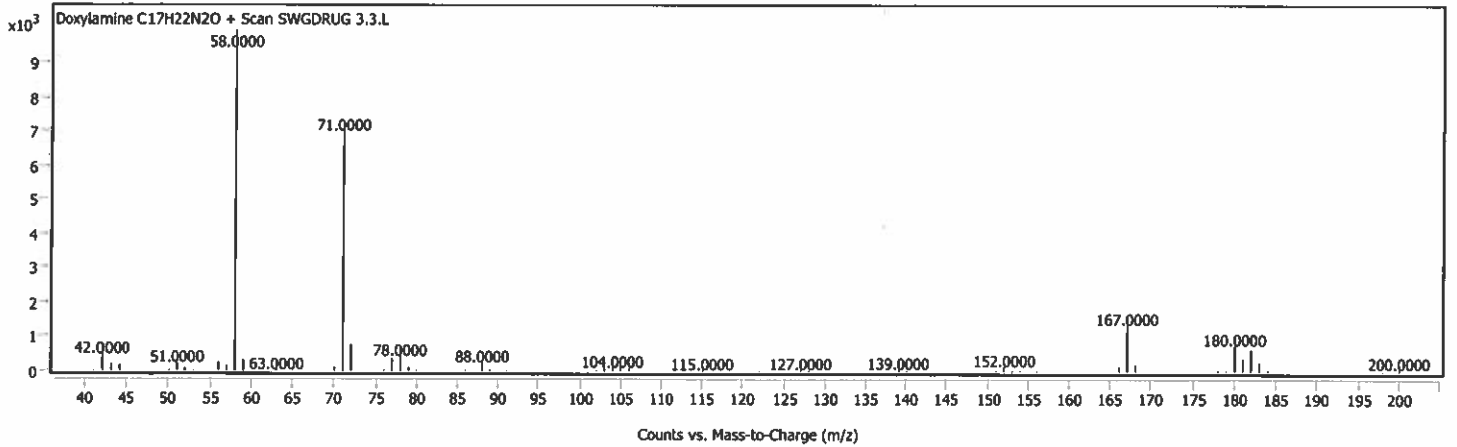
## Observed Spectrum



## Mirror Plot



## Library Spectrum





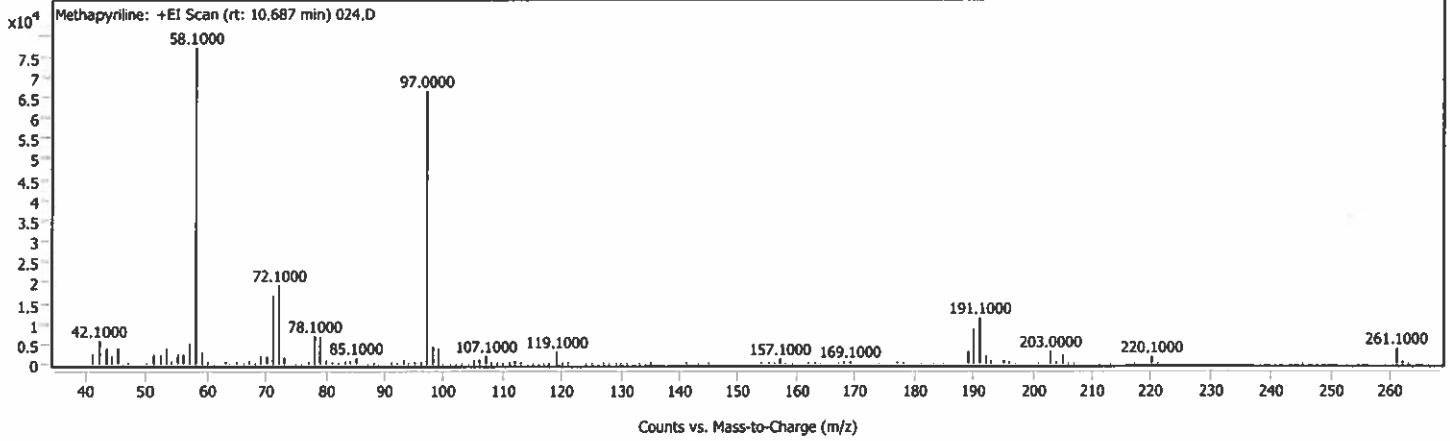
# Analysis Report

+ Scan (rt: 10.687 min)

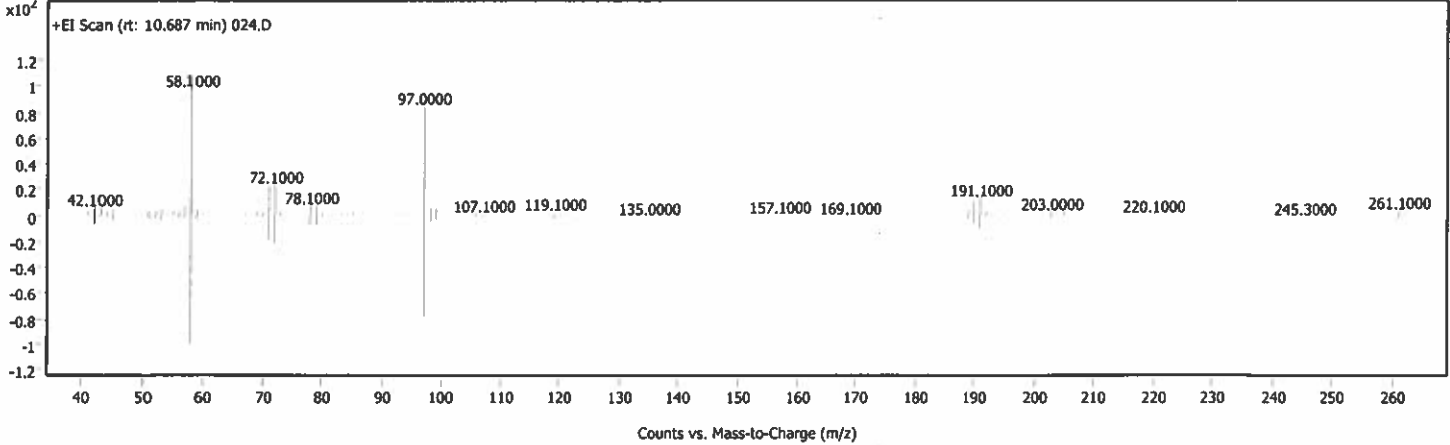
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	92.15	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

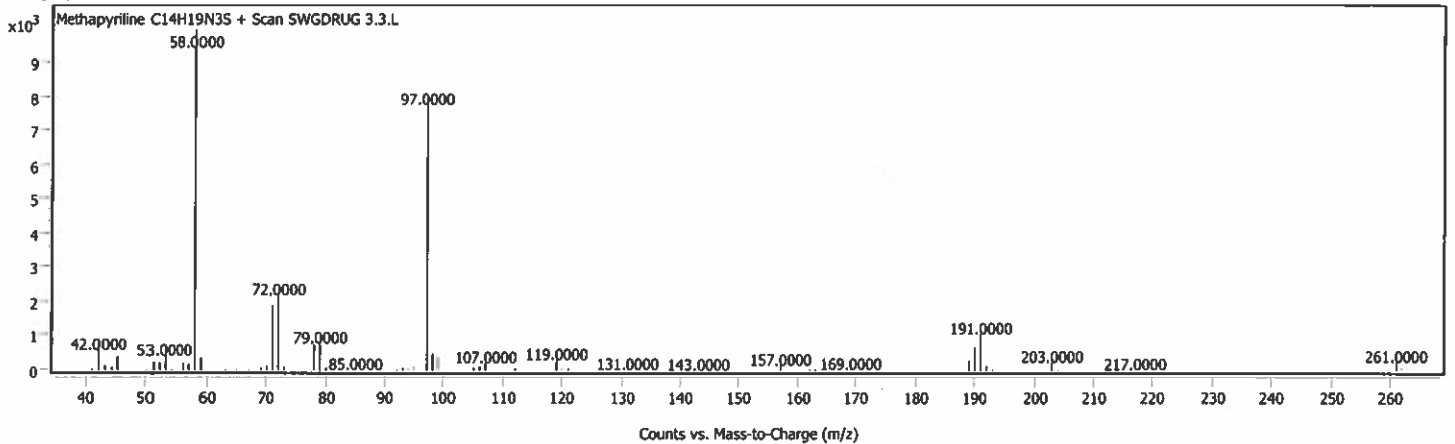
### Observed Spectrum



### Mirror Plot



### Library Spectrum



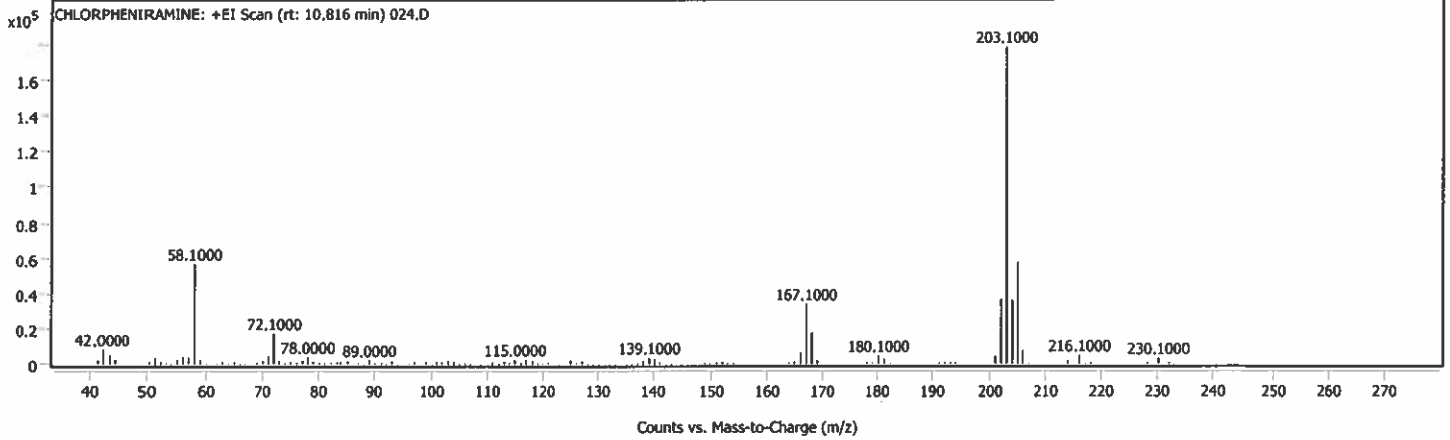
# Analysis Report

+ Scan (rt: 10.816 min)

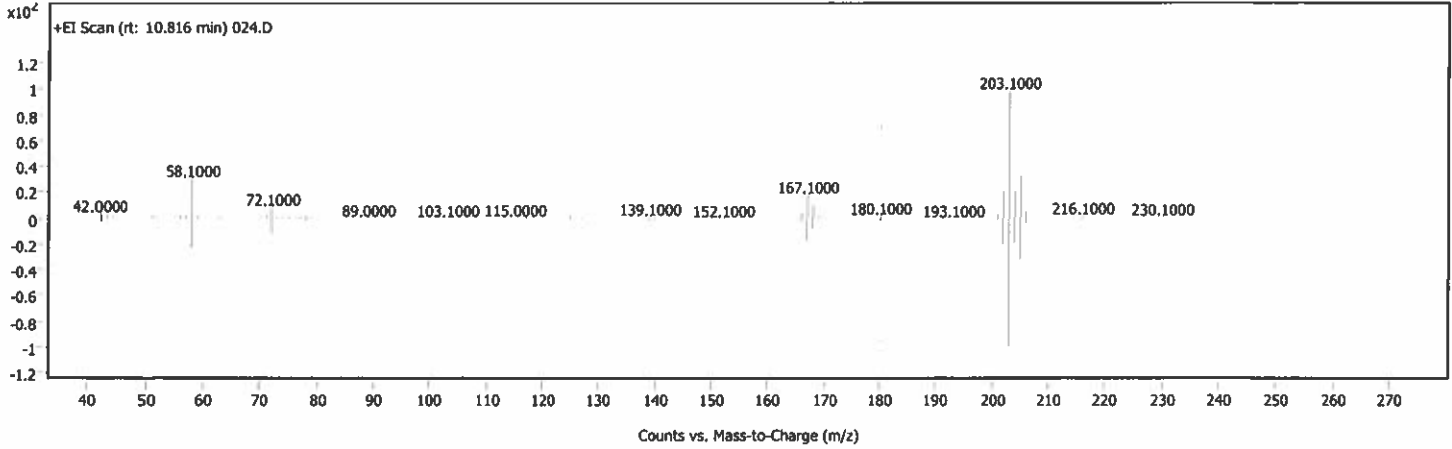
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	98.24	aafs.I	MATRIX 10 W/ IH CTL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

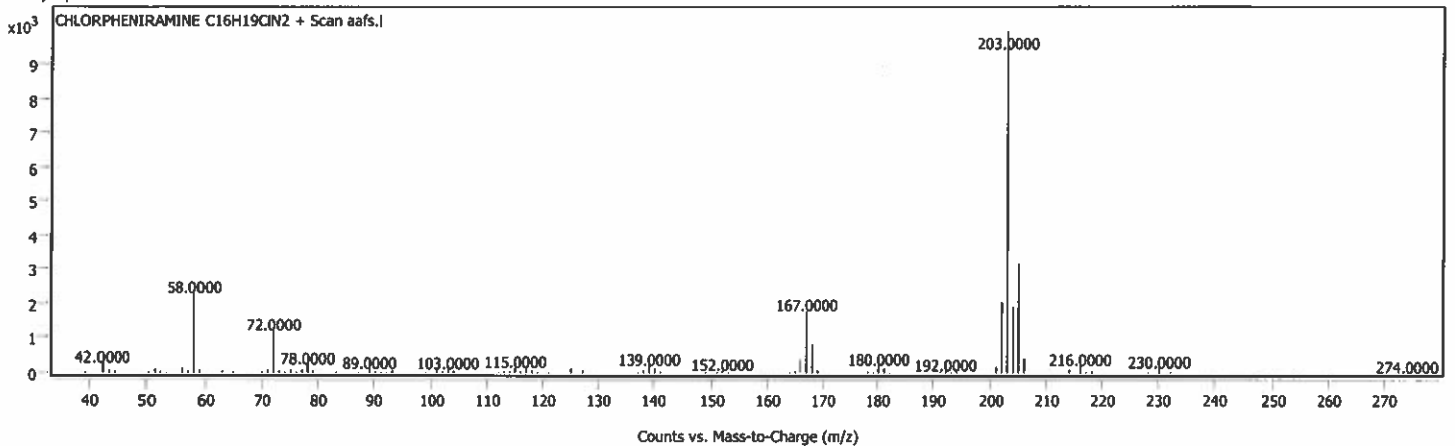
## Observed Spectrum



## Mirror Plot



## Library Spectrum



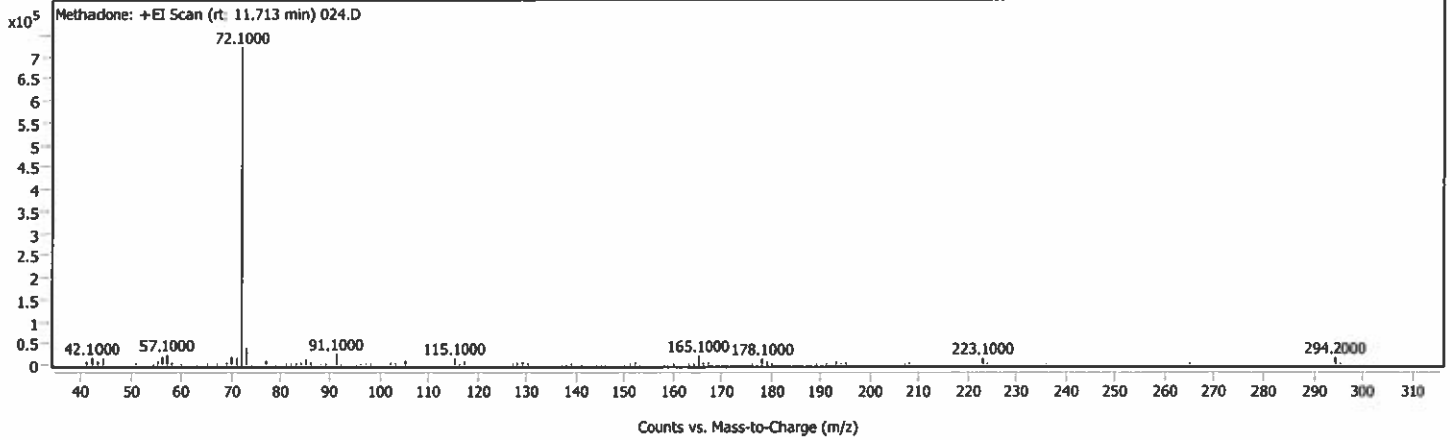
# Analysis Report

+ Scan (rt: 11.713 min)

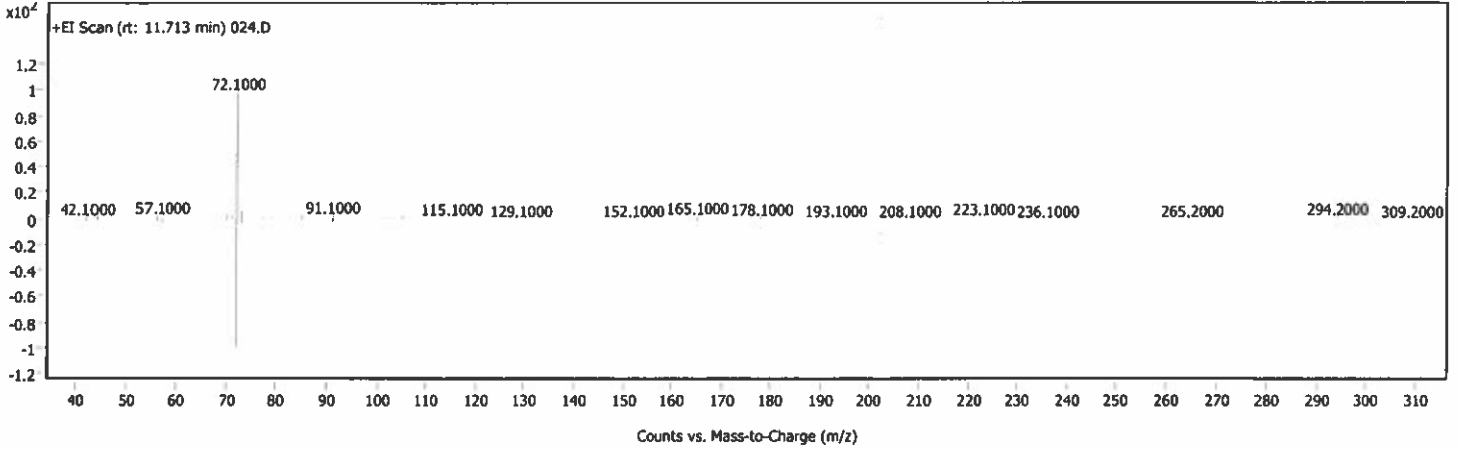
Methadone; C<sub>21</sub>H<sub>27</sub>NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C <sub>21</sub> H <sub>27</sub> NO	93.88	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

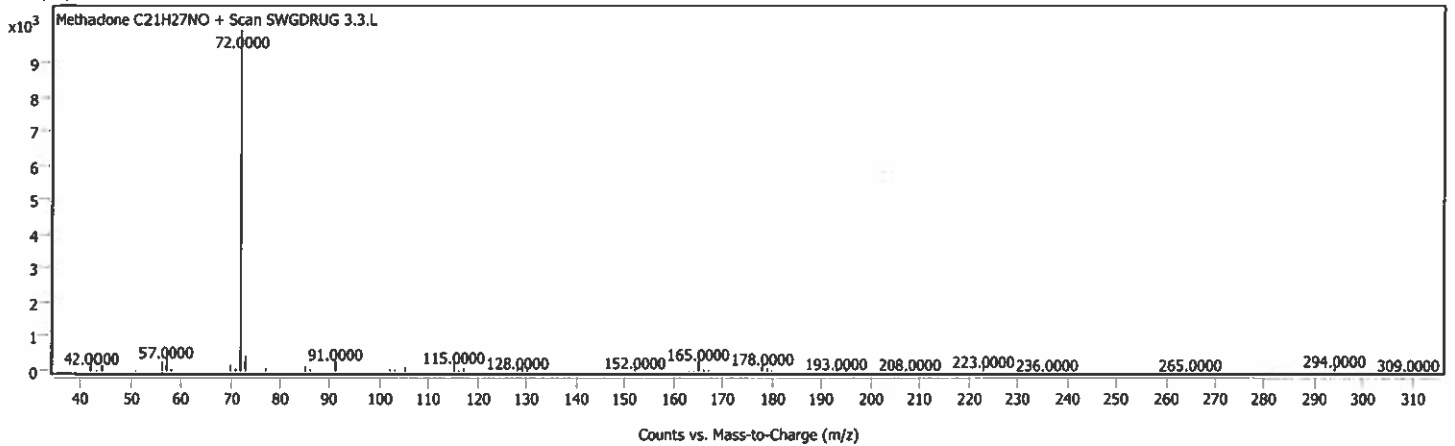
## Observed Spectrum



## Mirror Plot



## Library Spectrum



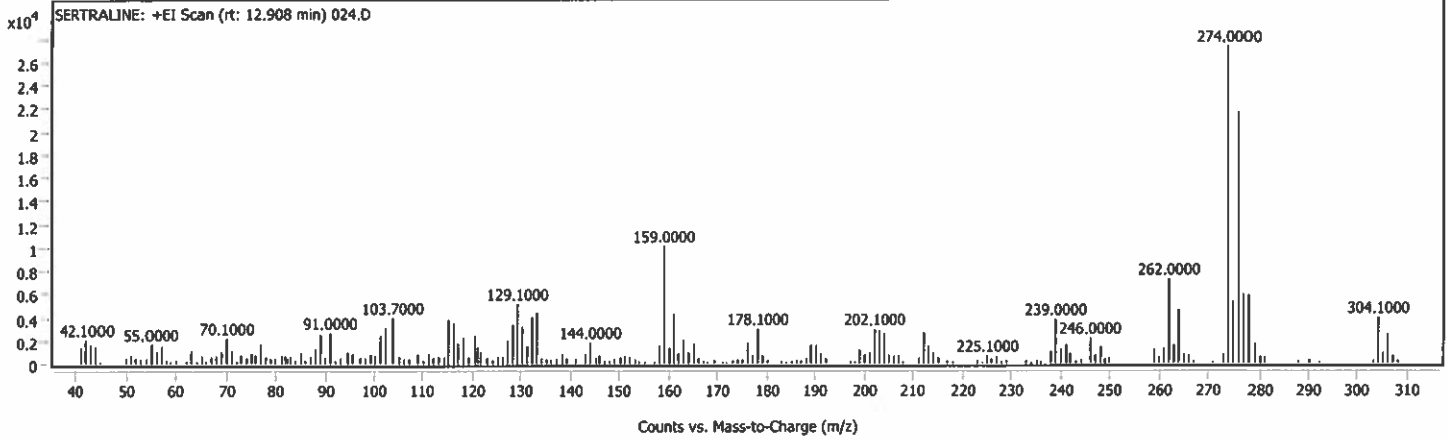
# Analysis Report

+ Scan (rt: 12.908 min)

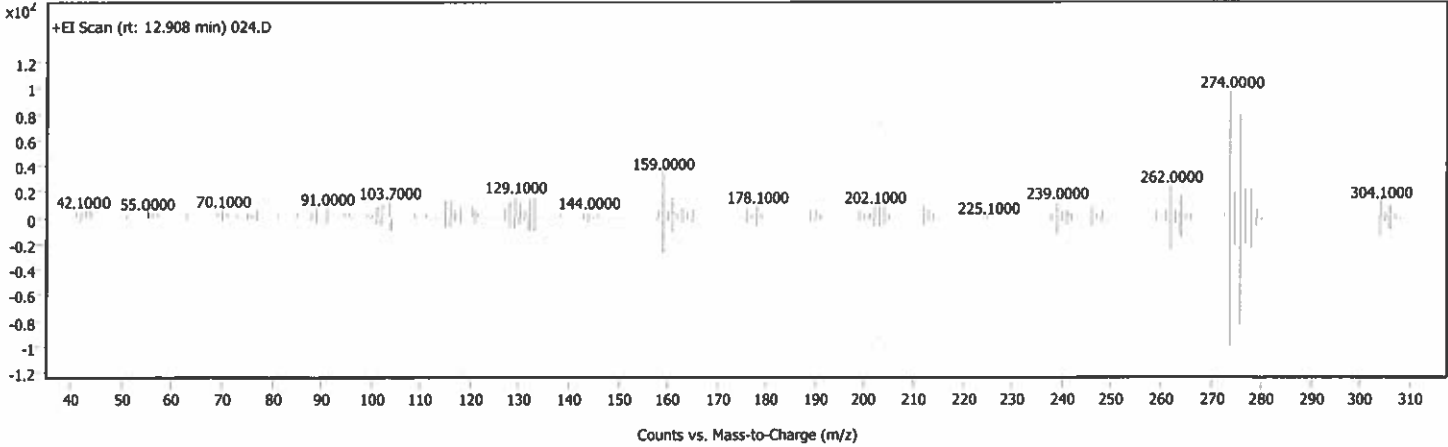
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	88.76	aafs.I	MATRIX 10 W/ IH CTL - LVR	KM/QC DATA IN XYLAZINE VALIDATION

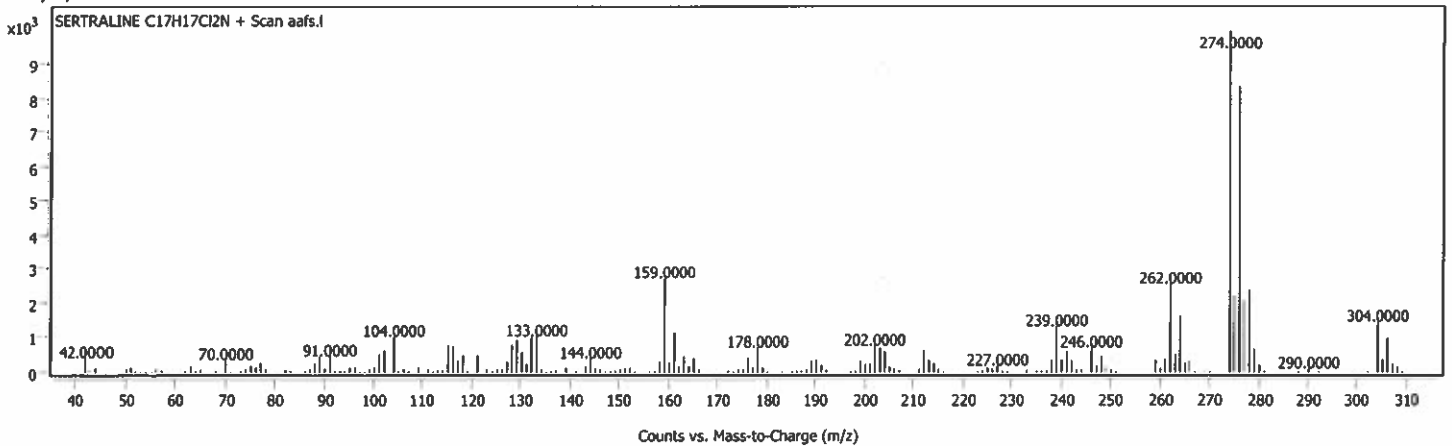
## Observed Spectrum



## Mirror Plot



## Library Spectrum

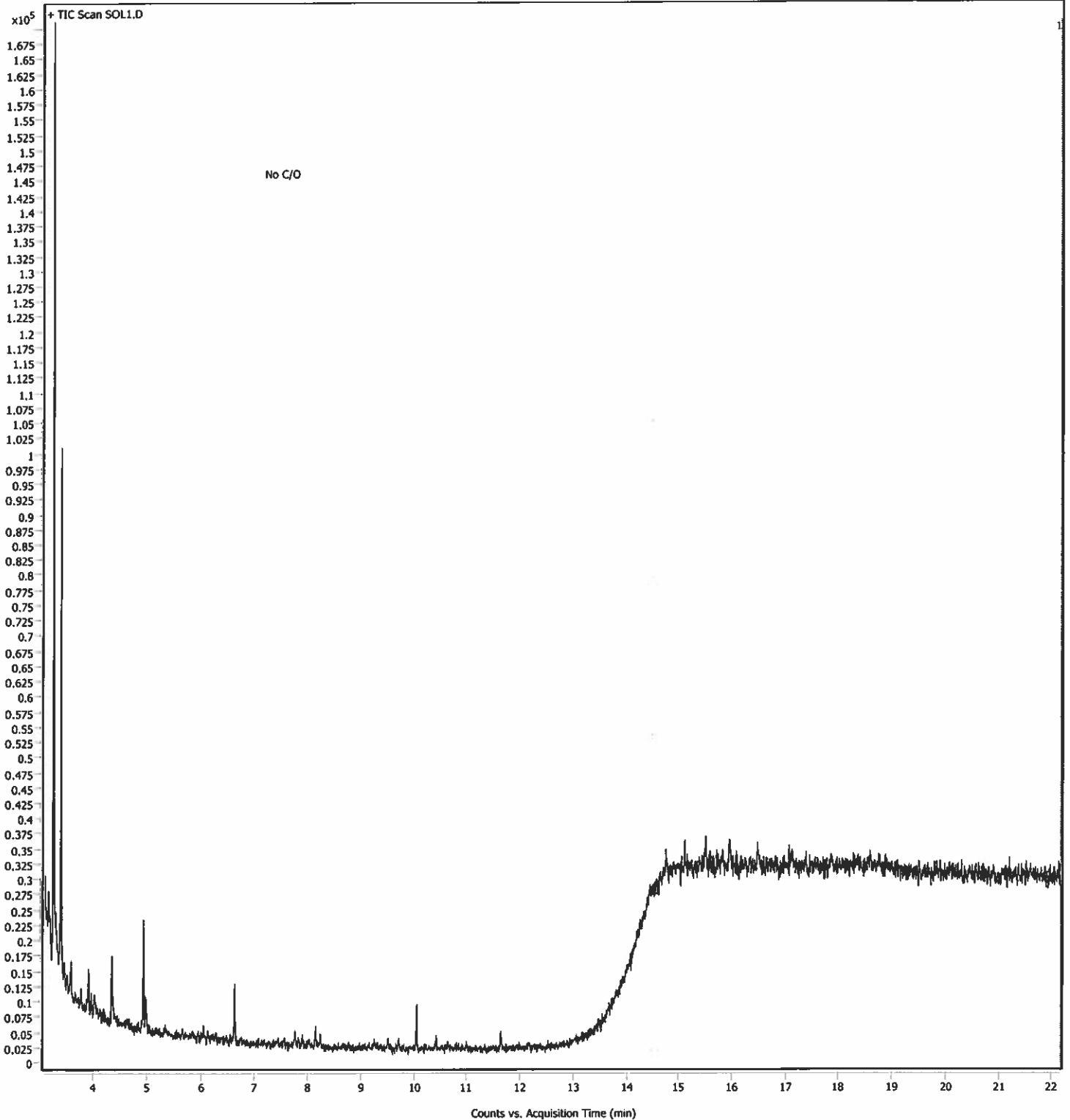


# Analysis Report

## Sample Information

Sample Name	Solvent Blank 1	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\SOL1.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 12:26:18 AM (UTC-04:00)
Position	52	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALIM
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

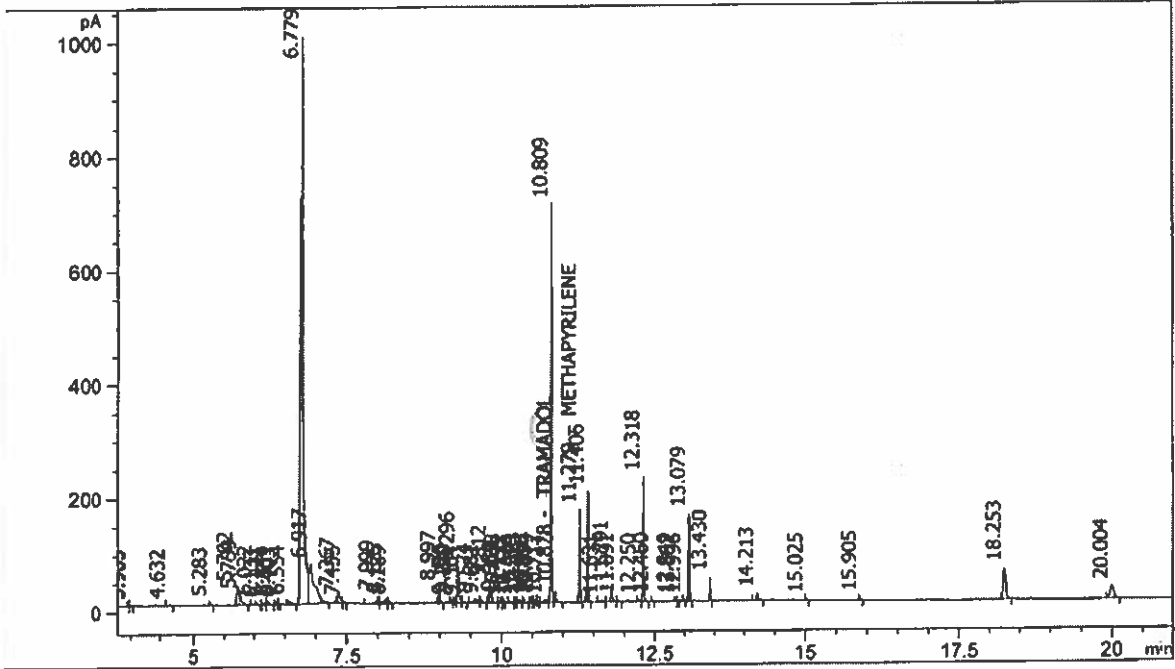
## Sample Chromatograms



Injection Date: 9/28/2021 11:53:23 PM  
 Sample Name: MATRIX 10 W/ IH CTL -->  
 Sample Info:

Seq Line: 26  
 Vial 27

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

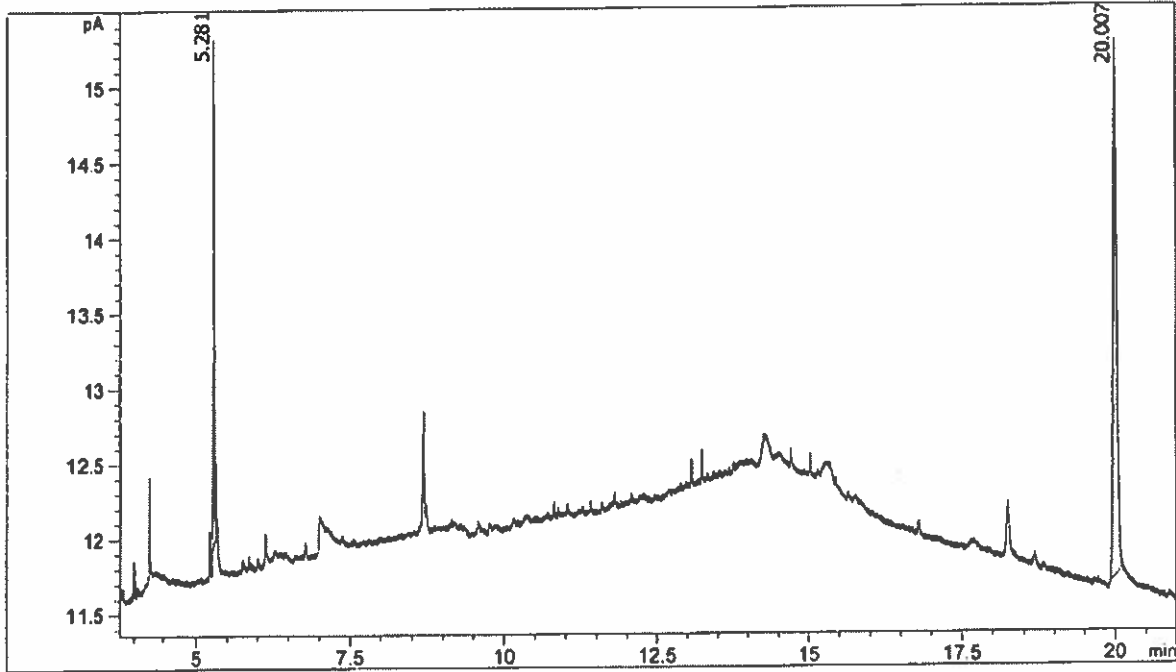


RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
10.878	11.033	36.176	20.639	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
11.279	11.278	191.207	163.954	1.000000	METHAPYRILENE

Injection Date: 9/29/2021 12:17:41 AM  
Sample Name: SOLVENT BLANK 1  
Sample Info:

Seq Line: 27  
Vial 28

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



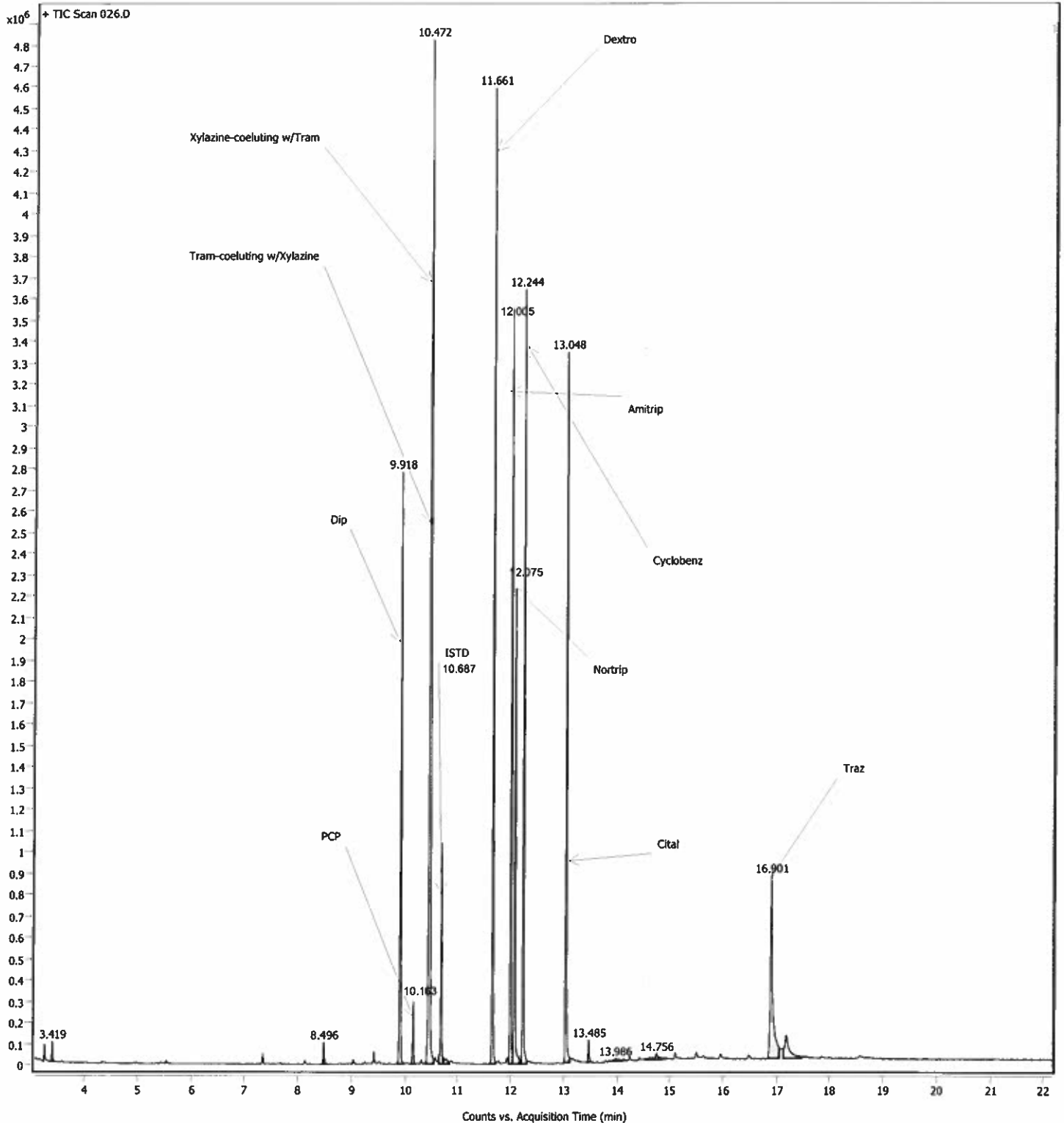
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

<b>Sample Name</b>	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	<b>Data File Path</b>	C:\MassHunter\GCMS\1\data\BASES\092821\026.D
<b>Instrument</b>	#3 - Enhanced	<b>Acq. Time (Local)</b>	9/29/2021 1:18:11 AM (UTC-04:00)
<b>Position</b>	54	<b>Method Path (Acq)</b>	C:\MassHunter\GCMS\1\methods\ALKALI.M
<b>Operator</b>	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

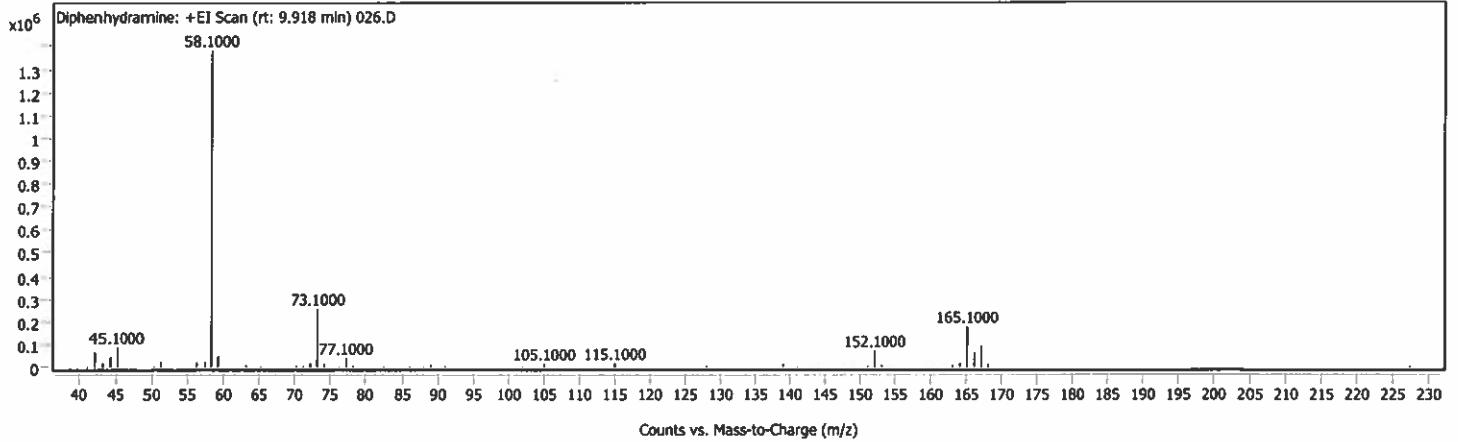
## Sample Spectra

+ Scan (rt: 9.918 min)

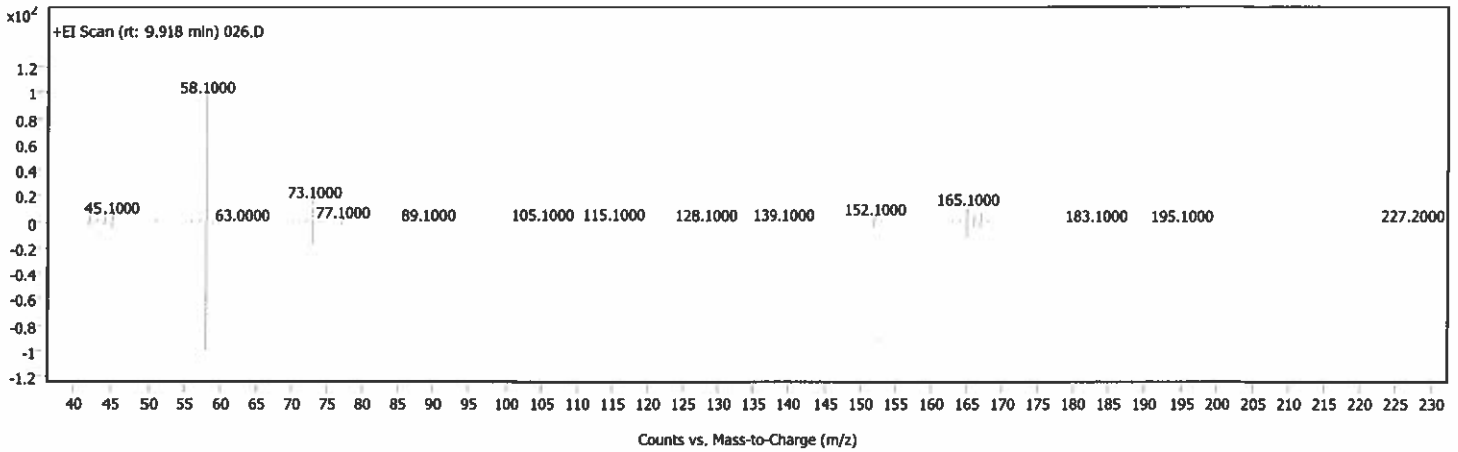
Diphenhydramine; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Diphenhydramine	C17H21NO	98.24	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

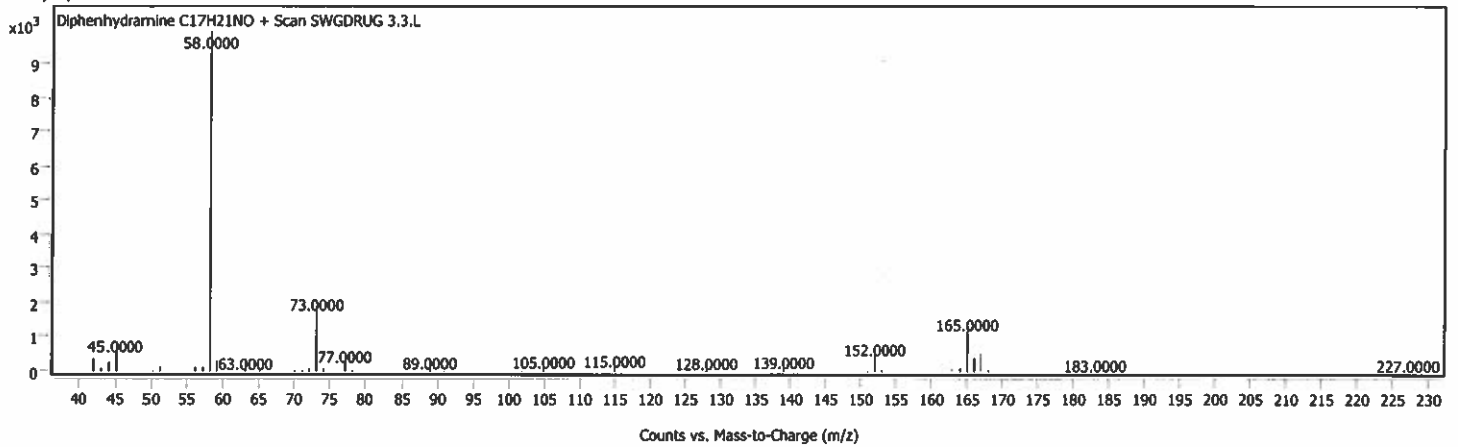
## Observed Spectrum



## Mirror Plot



## Library Spectrum



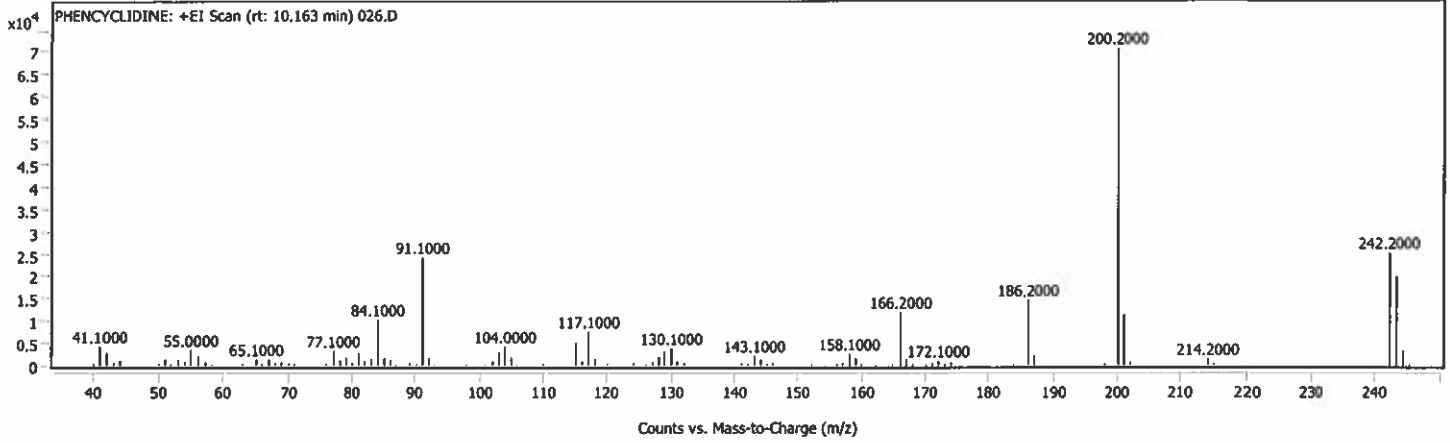
# Analysis Report

+ Scan (rt: 10.163 min)

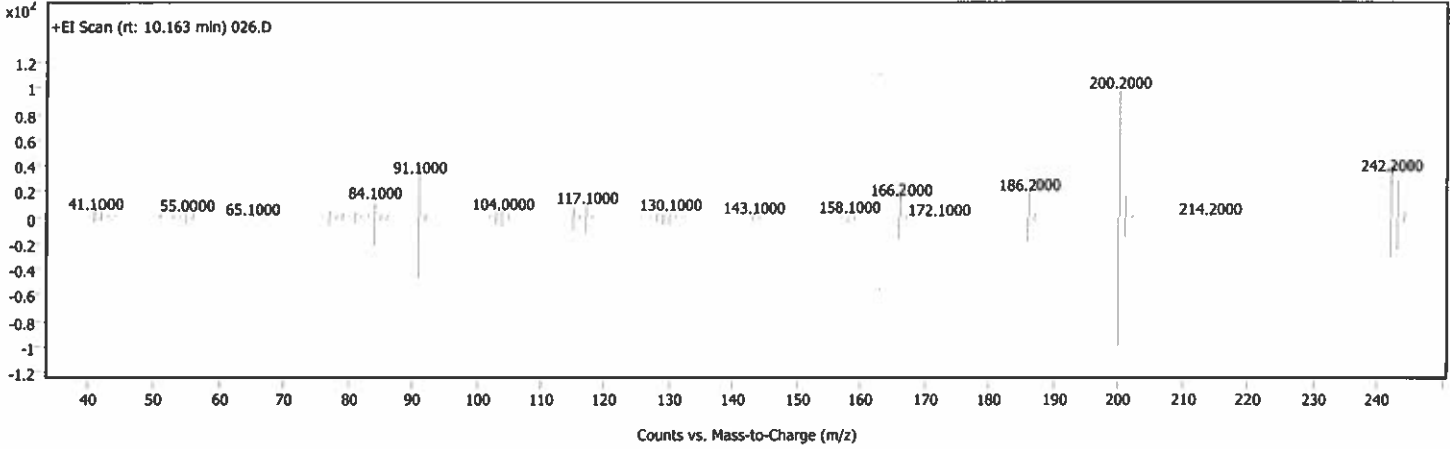
PHENCYCLIDINE; C17H25N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
PHENCYCLIDINE	C17H25N	98.67	aafs.I	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

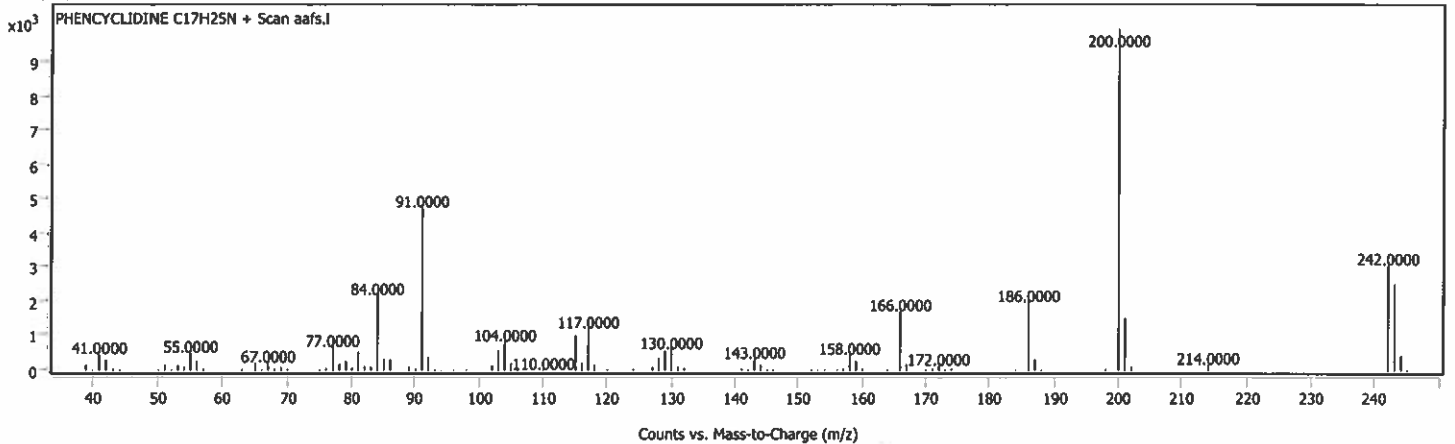
### Observed Spectrum



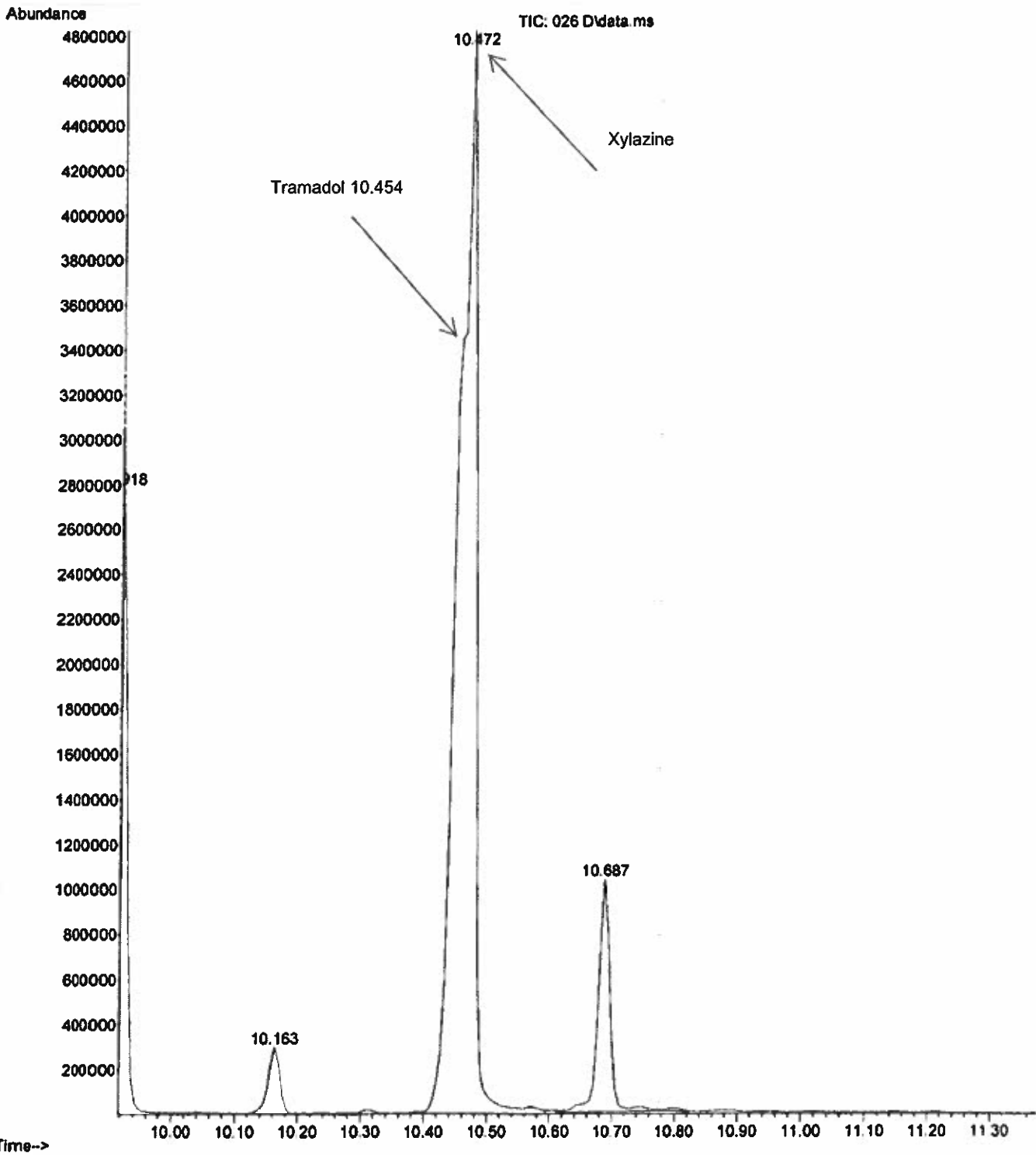
### Mirror Plot



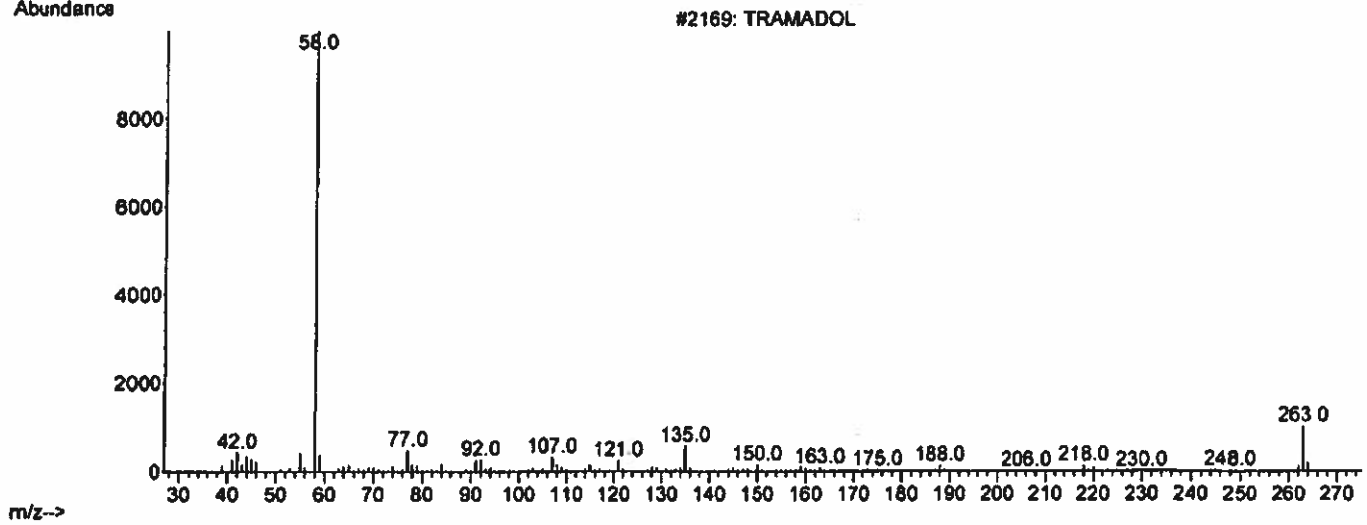
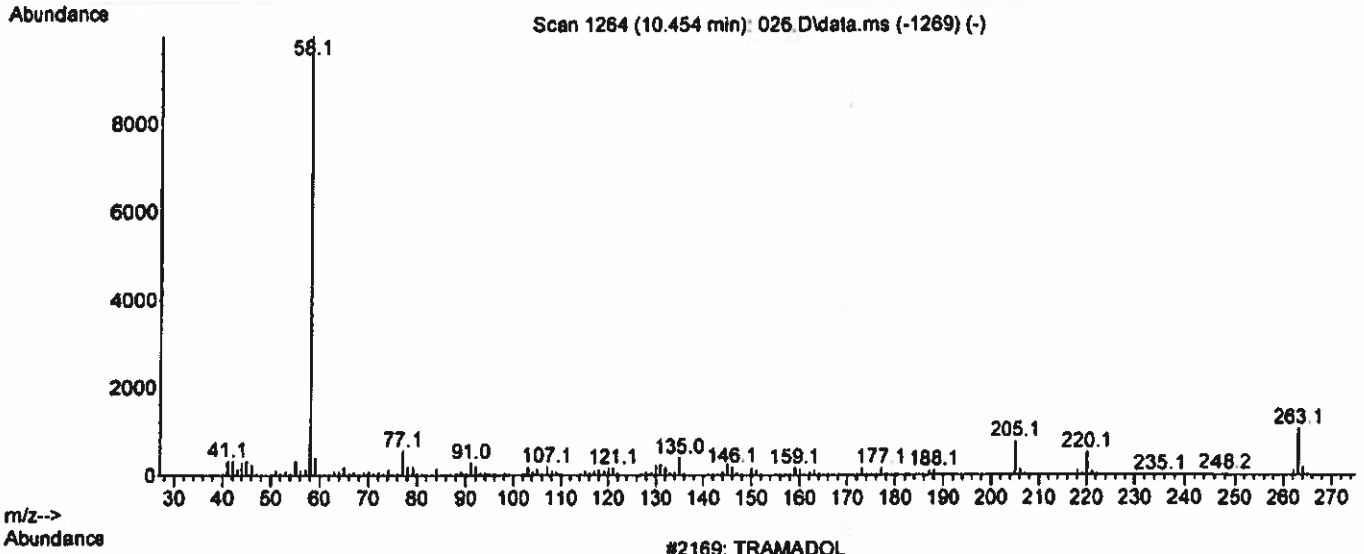
### Library Spectrum



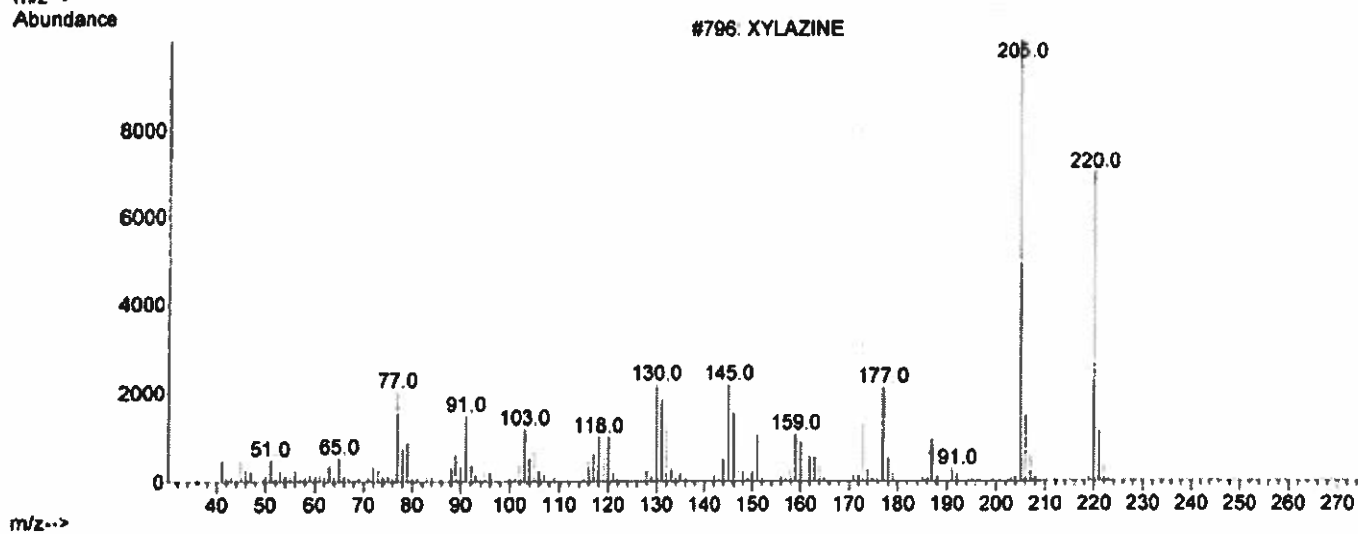
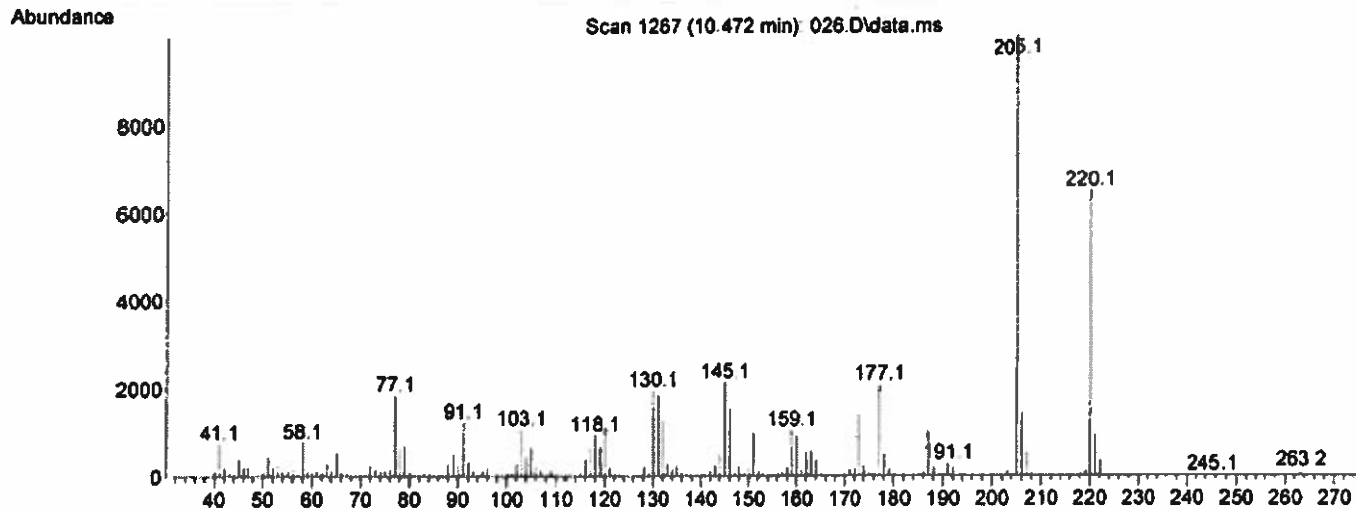
File :C:\Users\TOX\Desktop\092821\026.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 01:18 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 2 W/ SW CTL + XYLAZINE - BLD  
Misc Info : 2.0 mg/L - 400uL of 0.01 mg/mL CTL, plus 400uL 0.1 mg/mL XYLAZINE  
Vial Number: 54



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 95  
ID : TRAMADOL  
Sample Name: MATRIX 2 W/ SW CTL + XYLAZINE - BLD  
File: C:\Users\TOX\Desktop\092821\026.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 01:18  
Vial: 54



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 99  
ID : XYLAZINE  
Sample Name: MATRIX 2 W/ SW CTL + XYLAZINE - BLD  
File: C:\Users\TOX\Desktop\092821\026.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 01:18  
Vial: 54



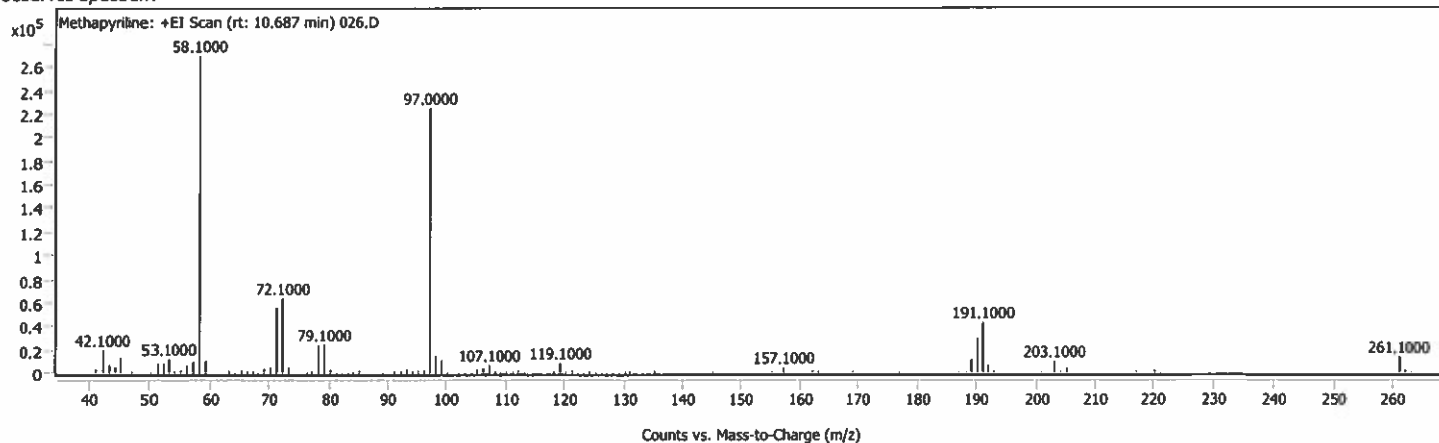
# Analysis Report

+ Scan (rt: 10.687 min)

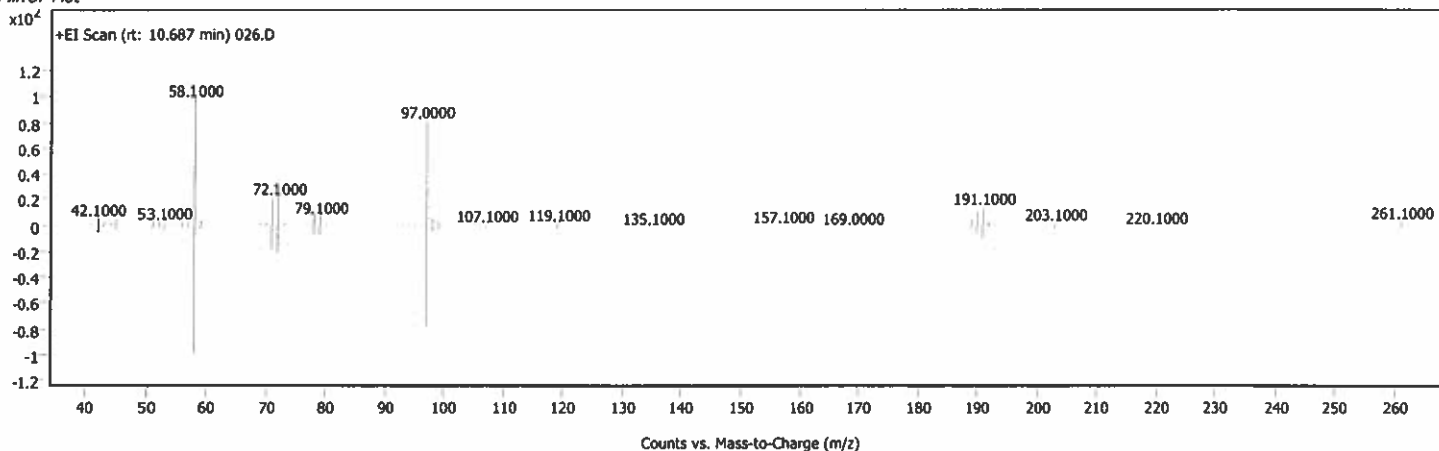
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	97.32	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

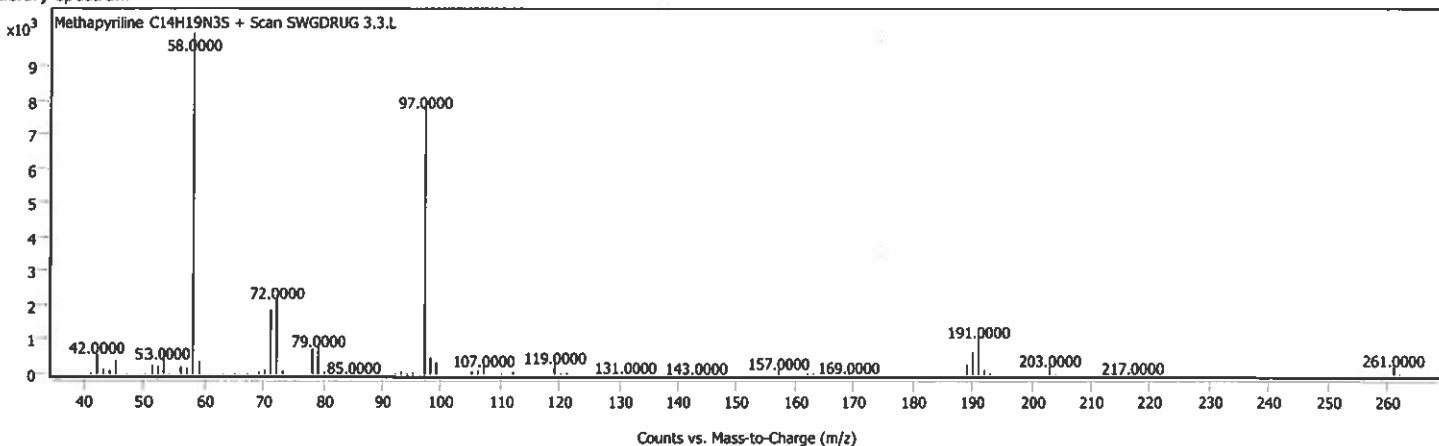
## Observed Spectrum



## Mirror Plot



## Library Spectrum



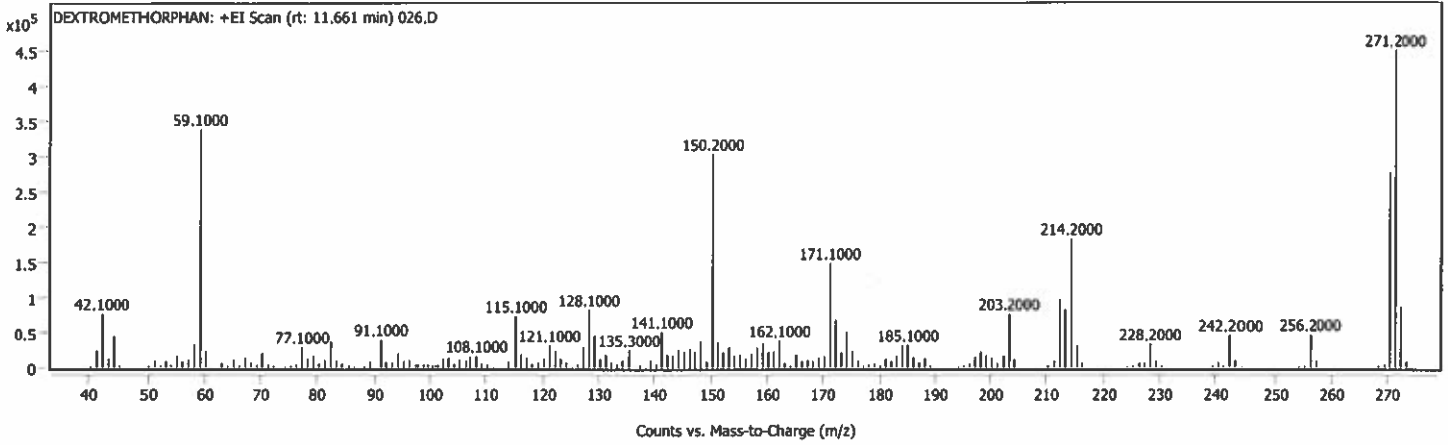
# Analysis Report

+ Scan (rt: 11.661 min)

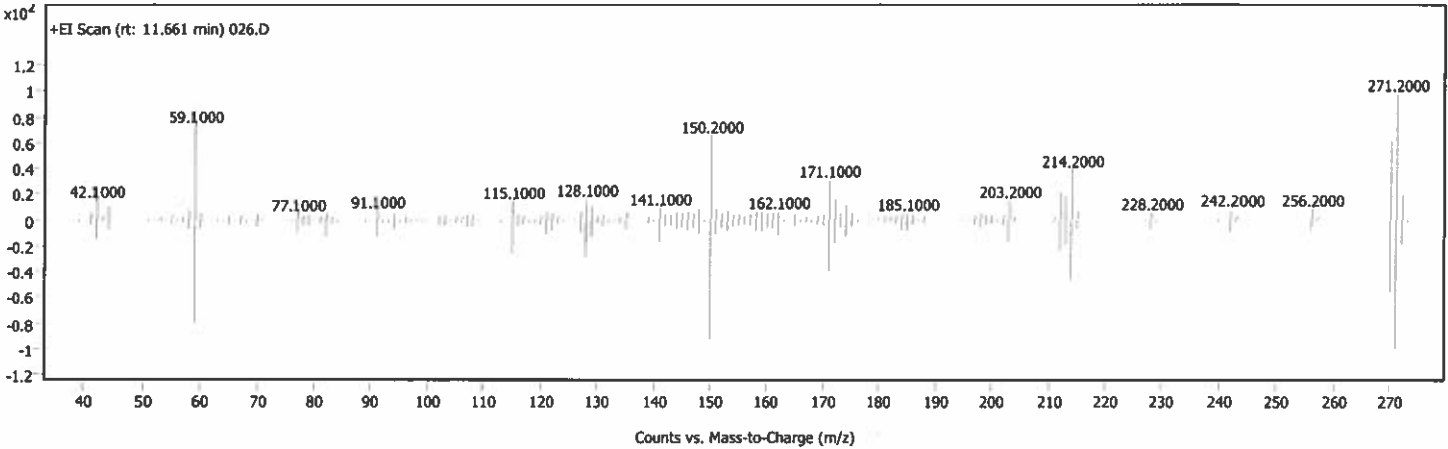
DEXTROMETHORPHAN; C18H25NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	92.25	aafs.1	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

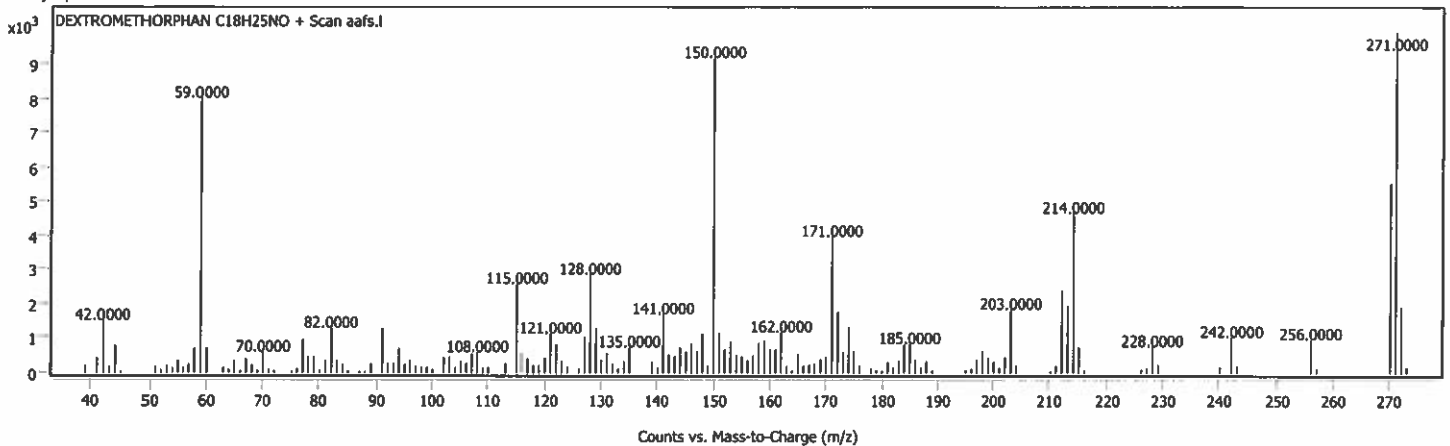
## Observed Spectrum



## Mirror Plot



## Library Spectrum



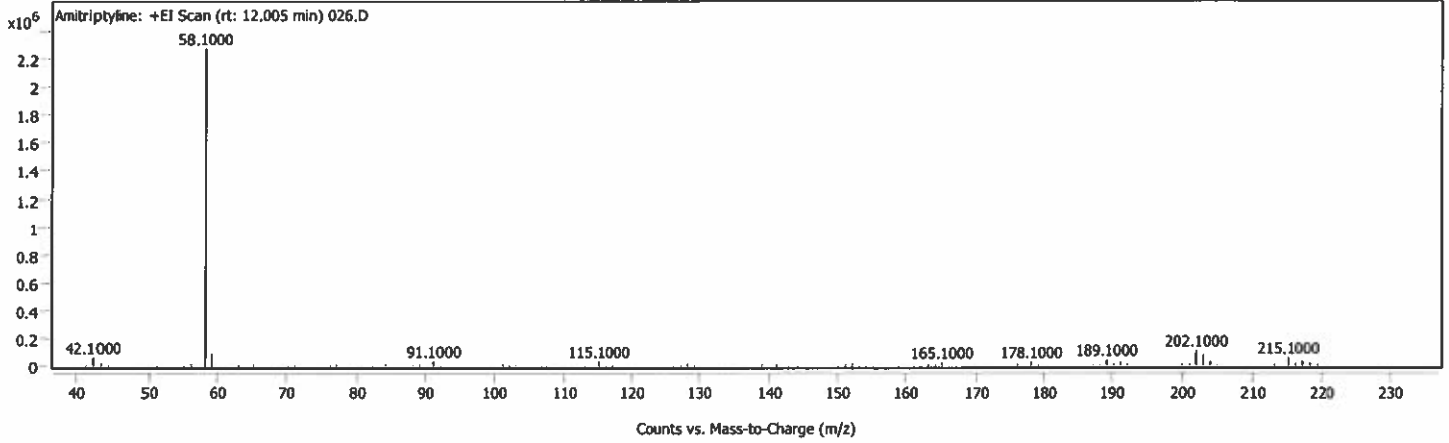
# Analysis Report

+ Scan (rt: 12.005 min)

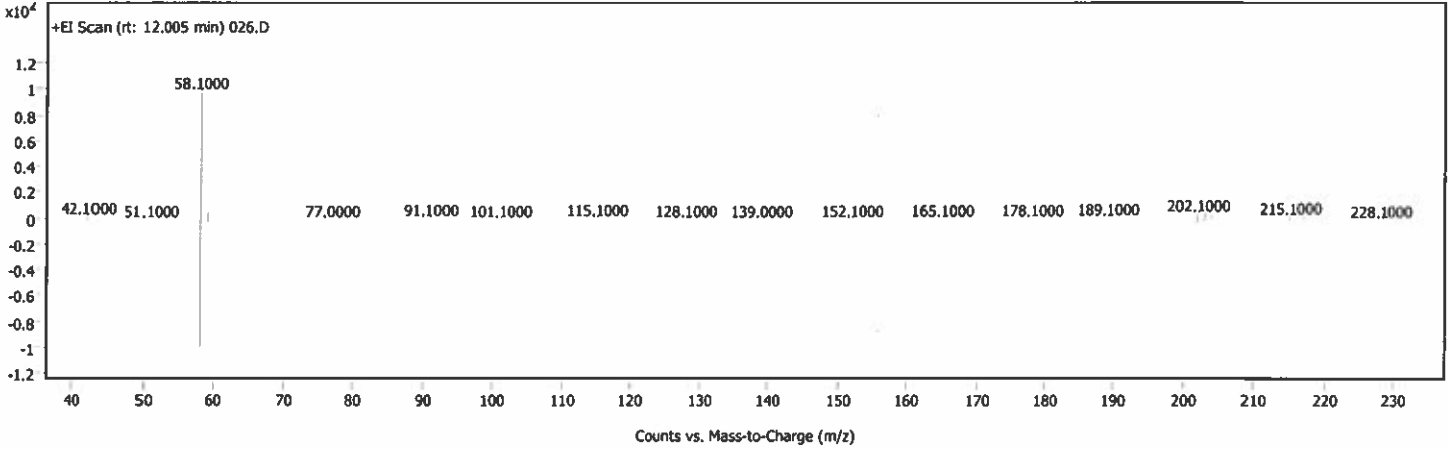
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	98.16	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

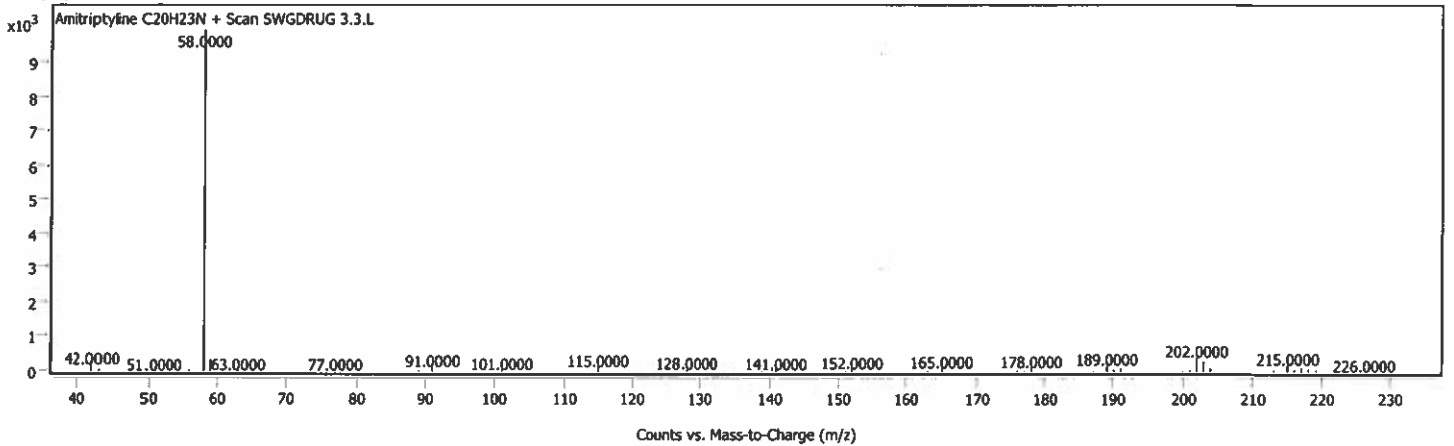
## Observed Spectrum



## Mirror Plot



## Library Spectrum





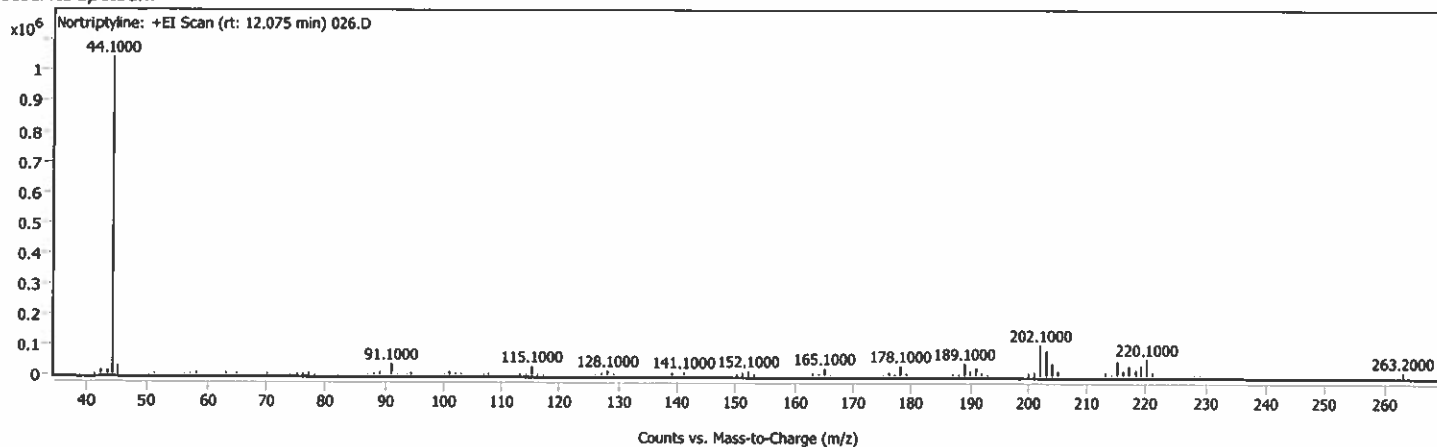
# Analysis Report

+ Scan (rt: 12.075 min)

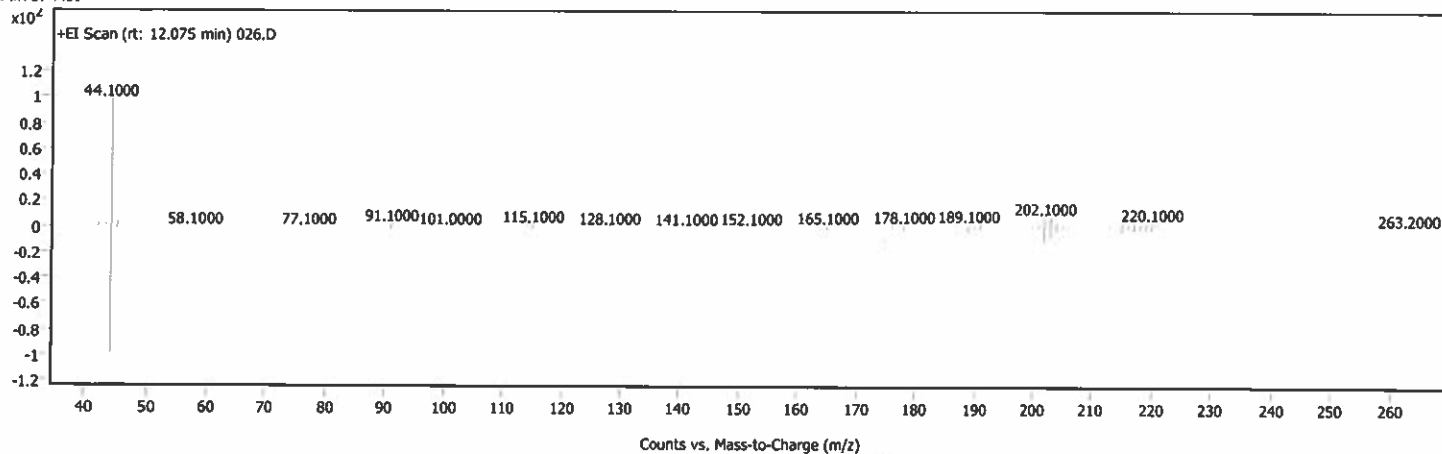
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	97.29	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

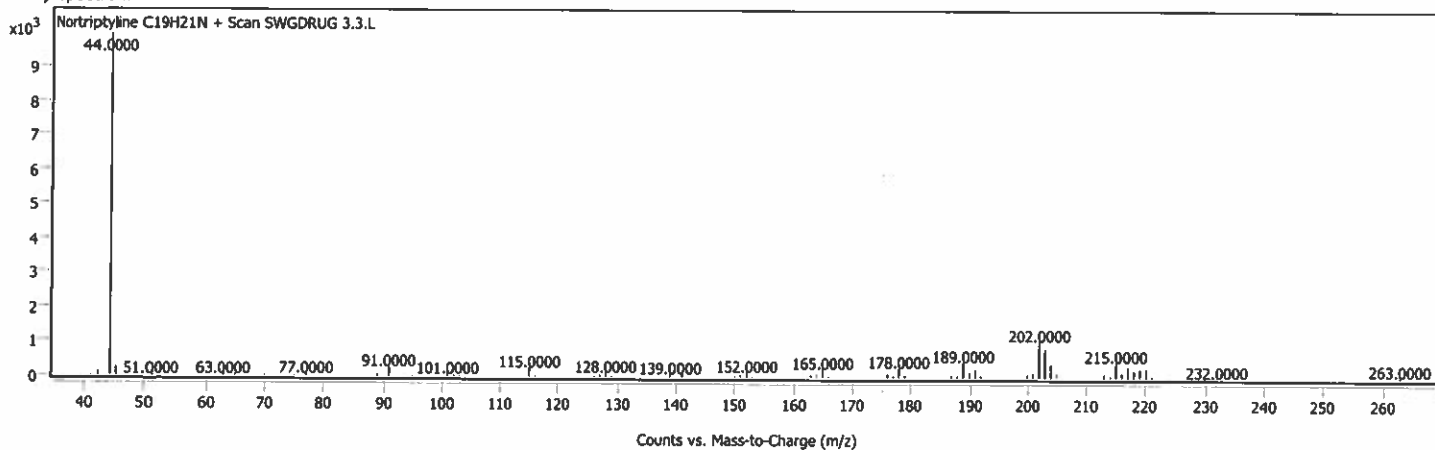
## Observed Spectrum



## Mirror Plot



## Library Spectrum



# Analysis Report

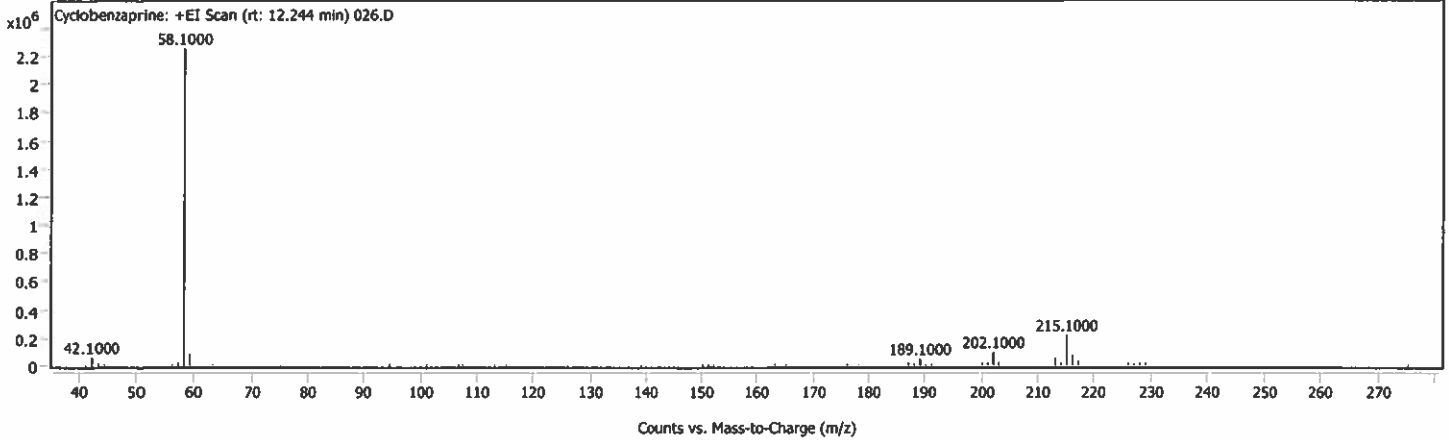


+ Scan (rt: 12.244 min)

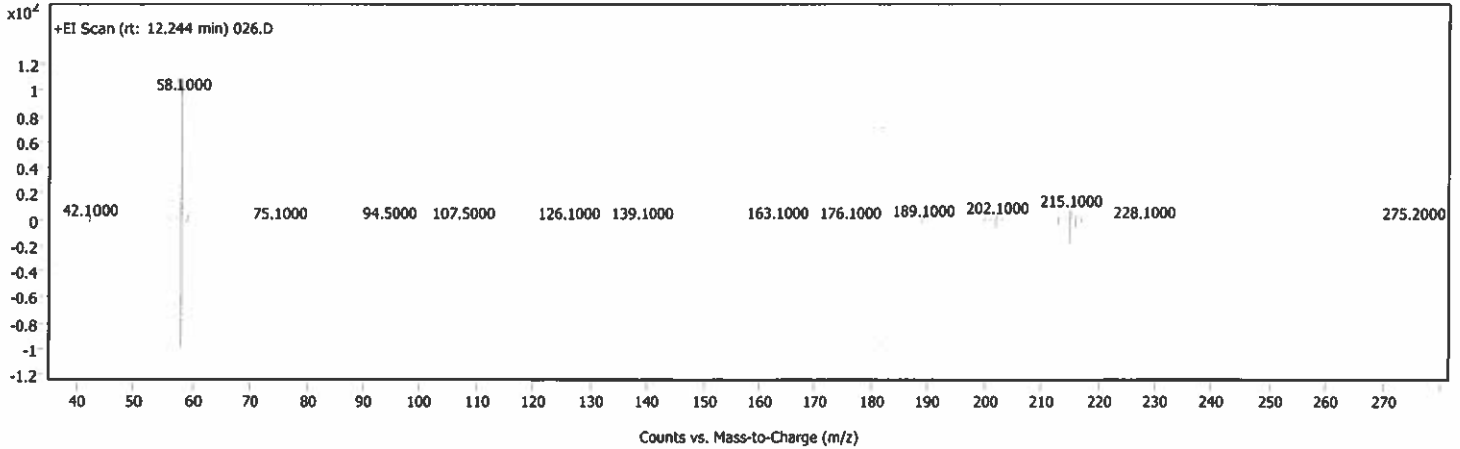
Cyclobenzaprine; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Cyclobenzaprine	C20H21N	96.72	SWGDRUG 3.3.L	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

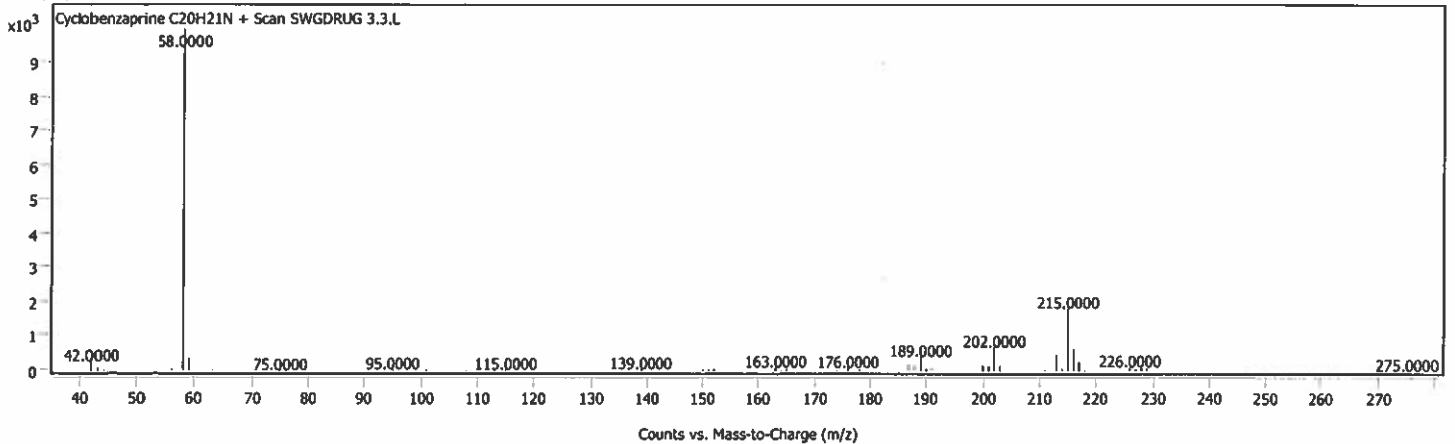
## Observed Spectrum



## Mirror Plot



## Library Spectrum



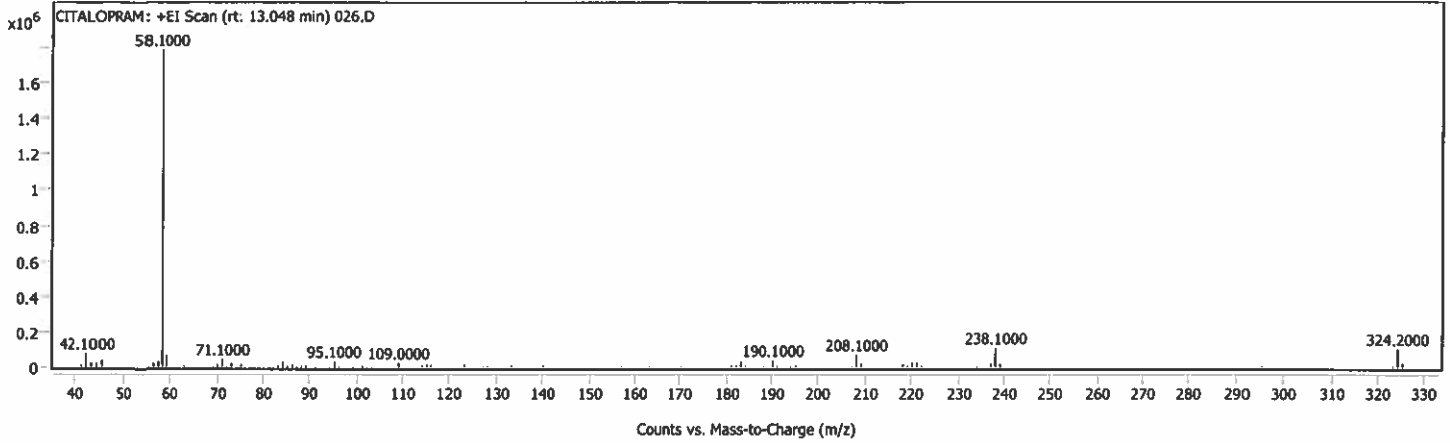
# Analysis Report

+ Scan (rt: 13.048 min)

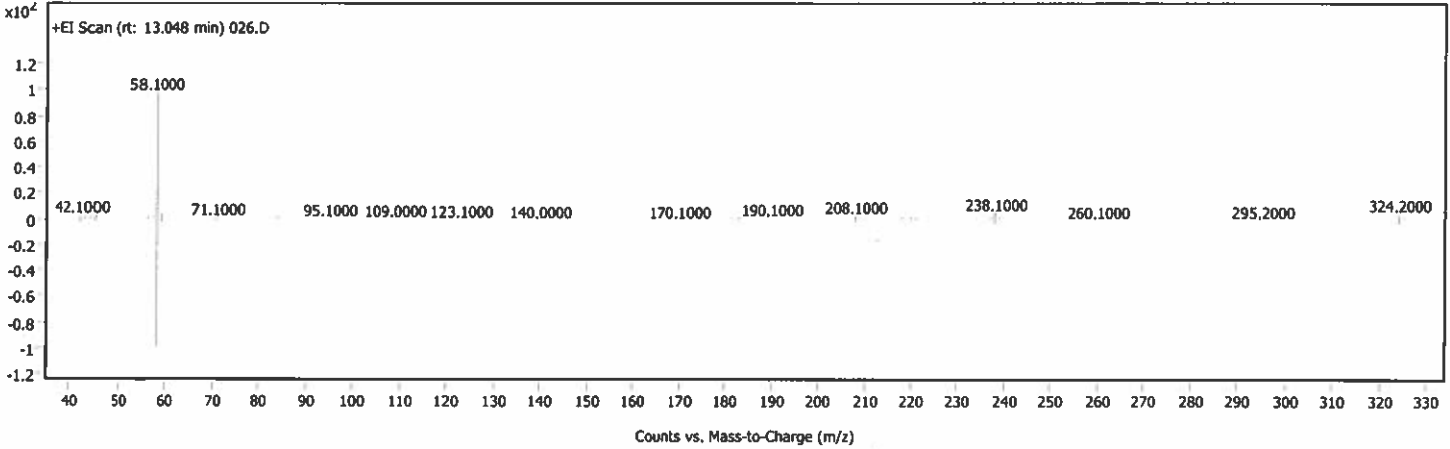
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	96.79	aafs.I	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

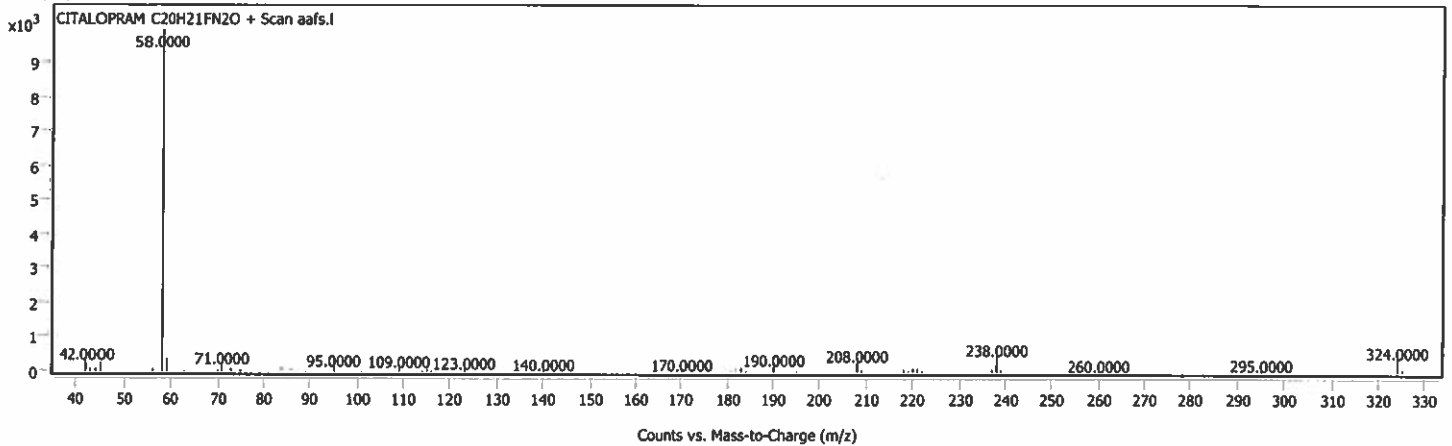
## Observed Spectrum



## Mirror Plot



## Library Spectrum



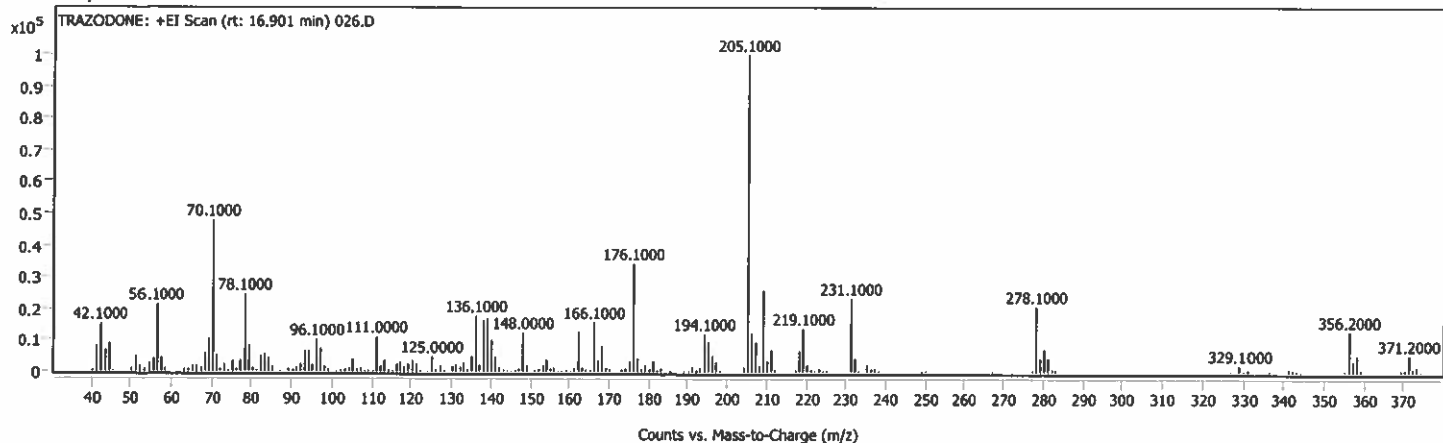
# Analysis Report

+ Scan (rt: 16.901 min)

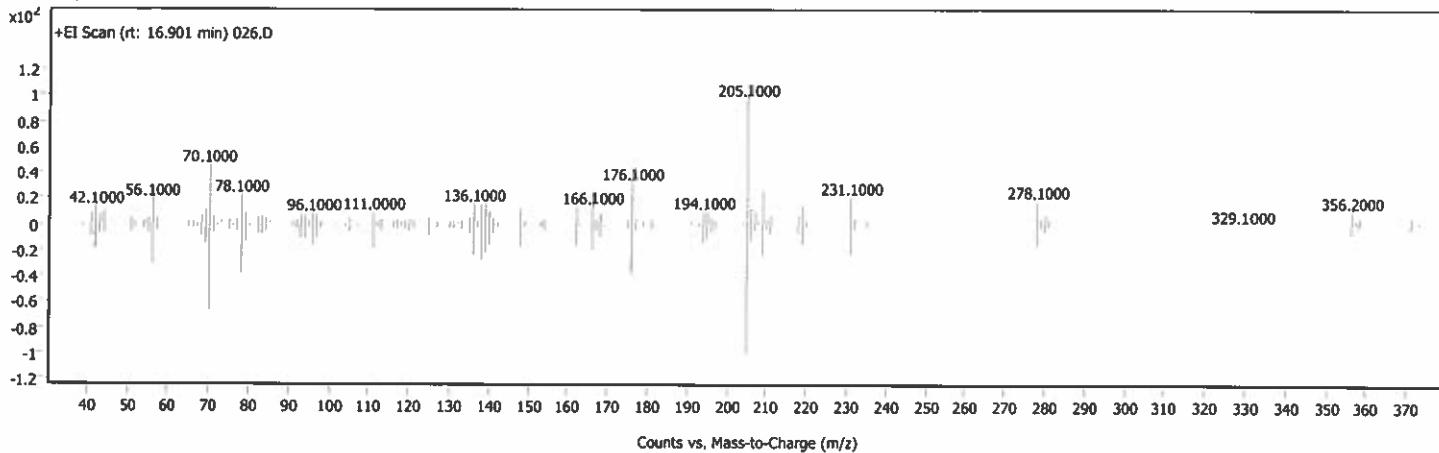
TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		94.29	aafs.i	MATRIX 2 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

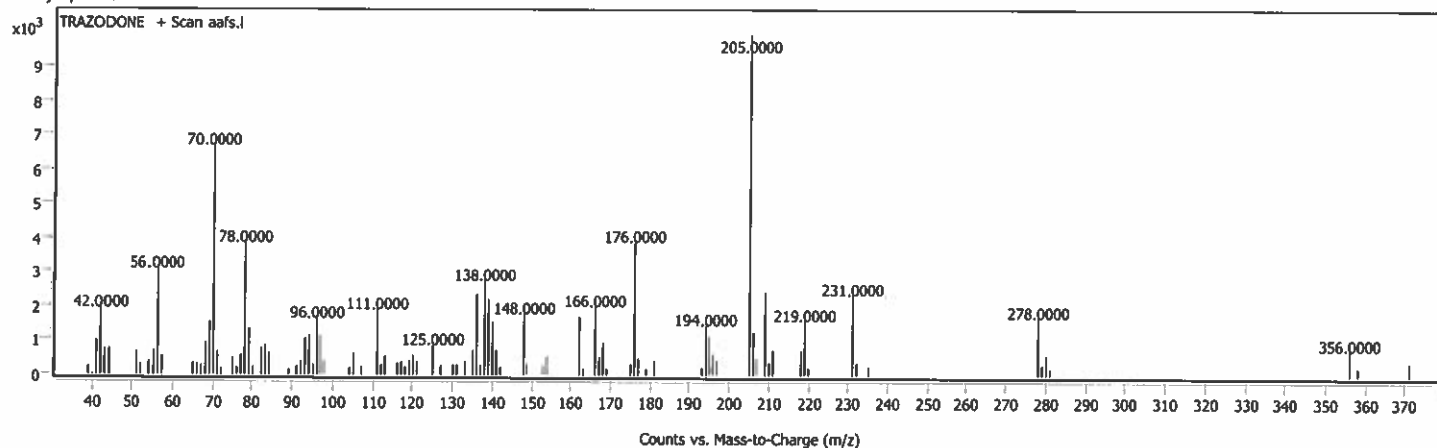
## Observed Spectrum



## Mirror Plot

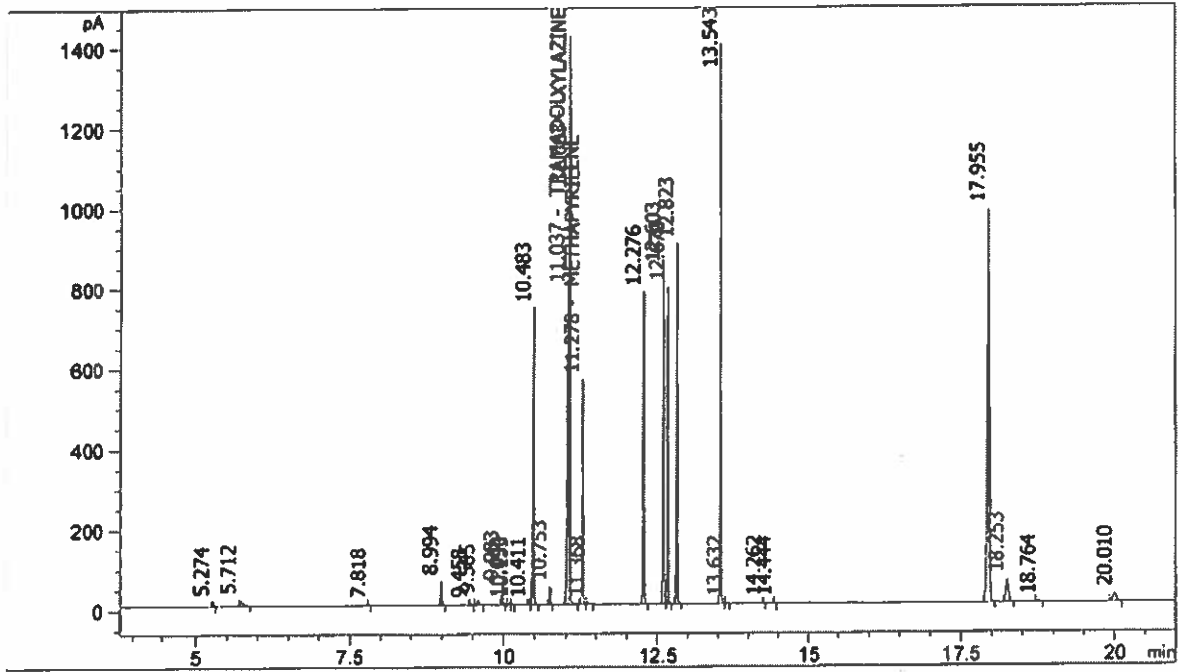


## Library Spectrum



Injection Date: 9/29/2021 1:06:20 AM Seq Line: 29  
 Sample Name: MATRIX 2 W/ SW CTL + -> Vial 30  
 Sample Info: *xy lazine*

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

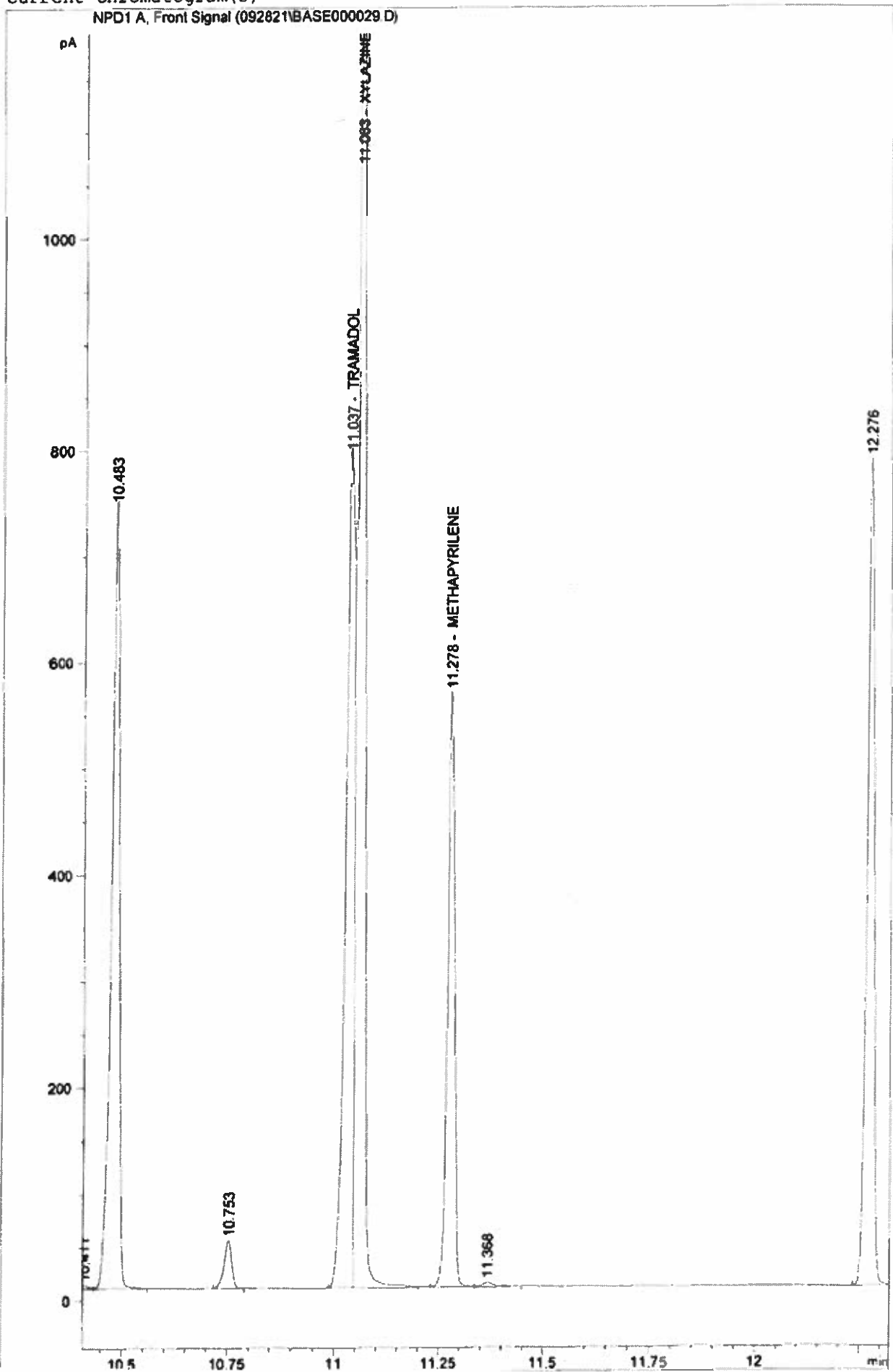


see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.037	11.033	1085.410	785.224	0.000000	TRAMADOL RRT-0.9786
11.063	11.046	1667.894	1410.112	0.000000	XYLAZINE RRT-0.9809
11.278	11.278	654.919	558.788	1.000000	METHAPYRILENE

Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000029.D)

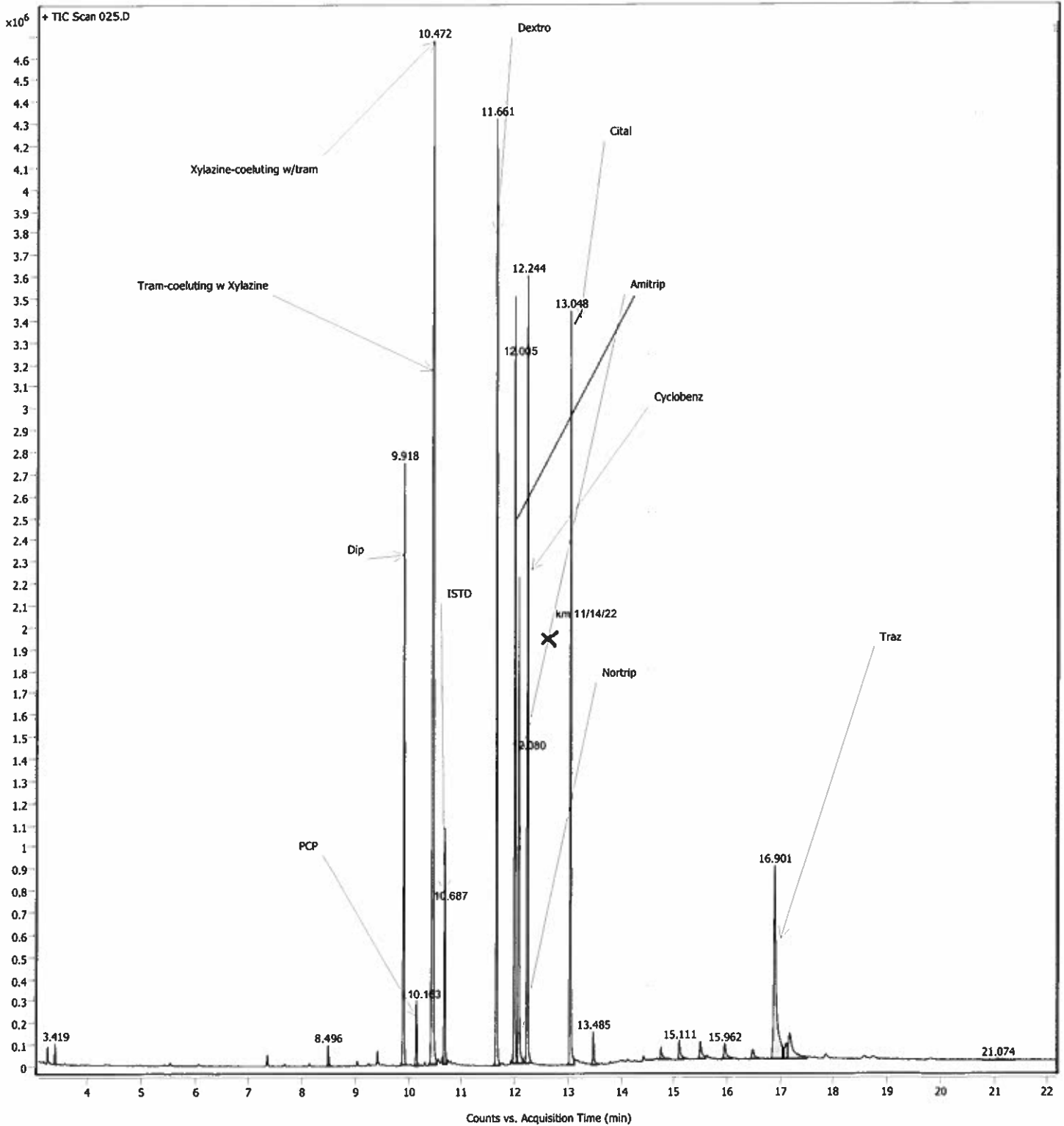


# Analysis Report

## Sample Information

<b>Sample Name</b>	MATRIX 1 W/ SW CTL + XYLAXINE - BLD	<b>Data File Path</b>	C:\MassHunter\GCMS\1\data\BASES\092821\025.D
<b>Instrument</b>	#3 - Enhanced	<b>Acq. Time (Local)</b>	9/29/2021 12:52:10 AM (UTC-04:00)
<b>Position</b>	53	<b>Method Path (Acq)</b>	C:\MassHunter\GCMS\1\methods\ALKALI.M
<b>Operator</b>	KM/QC DATA IN XYLAXINE VALIDATION		

## Sample Chromatograms



# Analysis Report

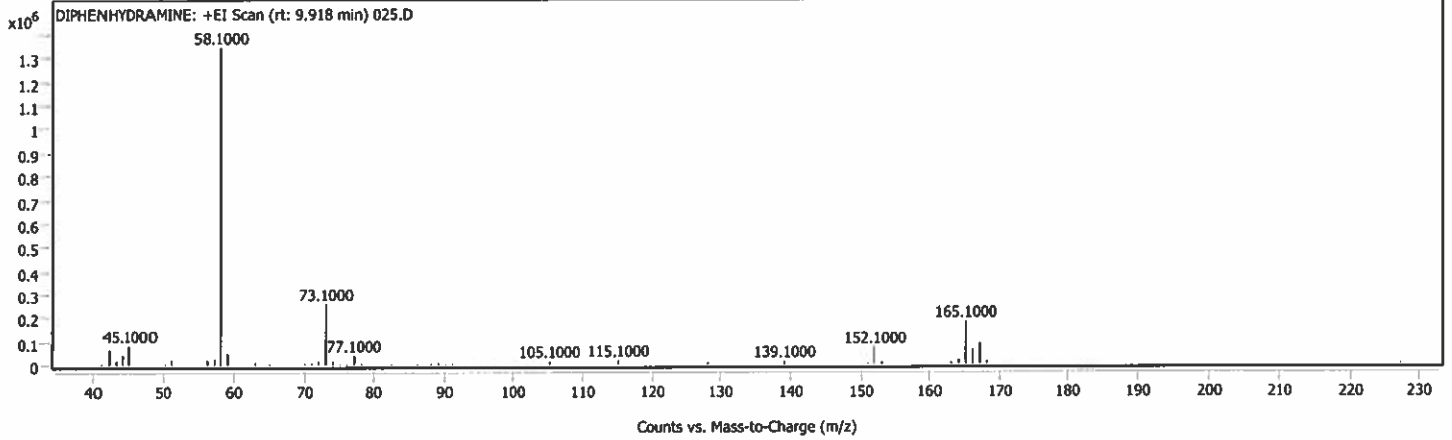
## Sample Spectra

+ Scan (rt: 9.918 min)

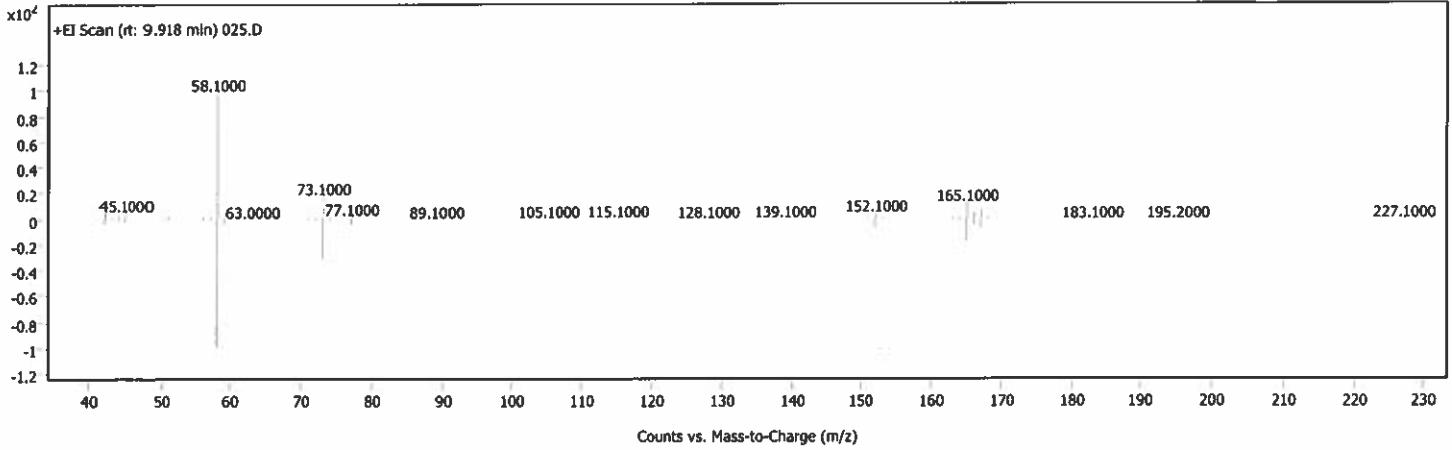
DIPHENHYDRAMINE; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DIPHENHYDRAMINE	C17H21NO	98.31	aafs.l	MATRIX 1 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

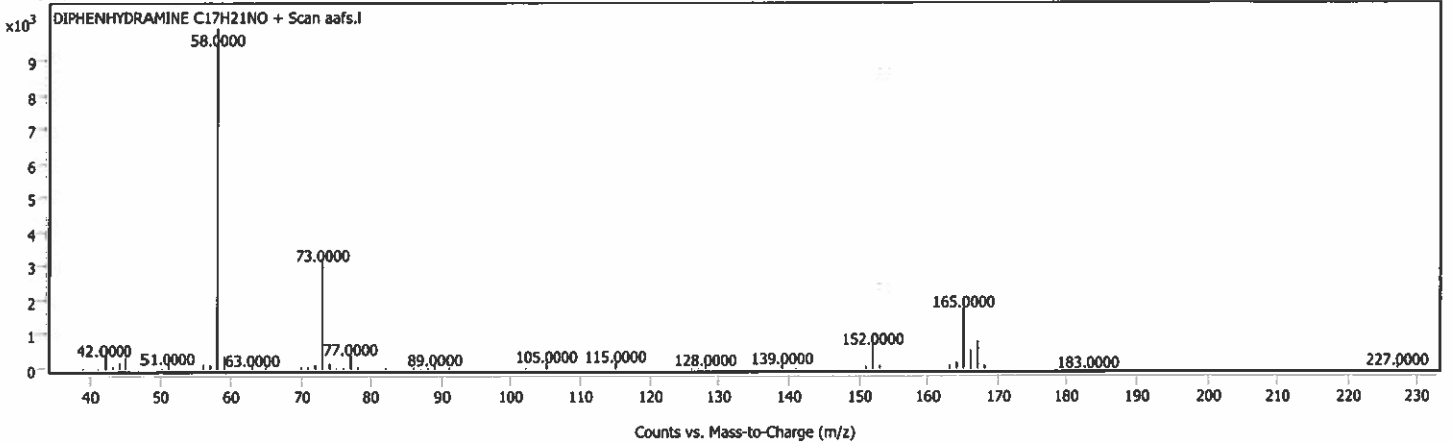
## Observed Spectrum



## Mirror Plot



## Library Spectrum





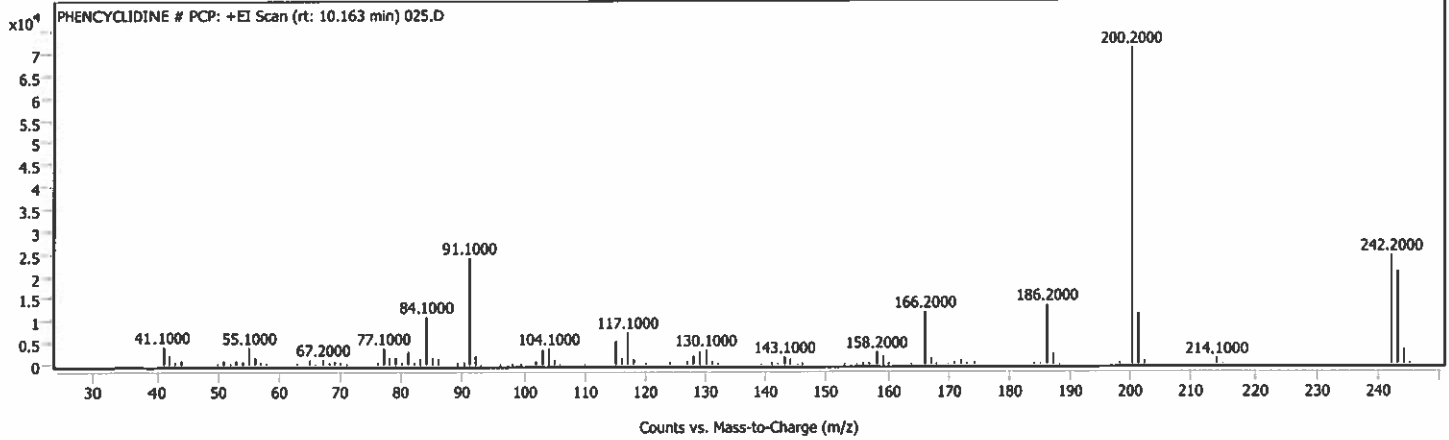
# Analysis Report

+ Scan (rt: 10.163 min)

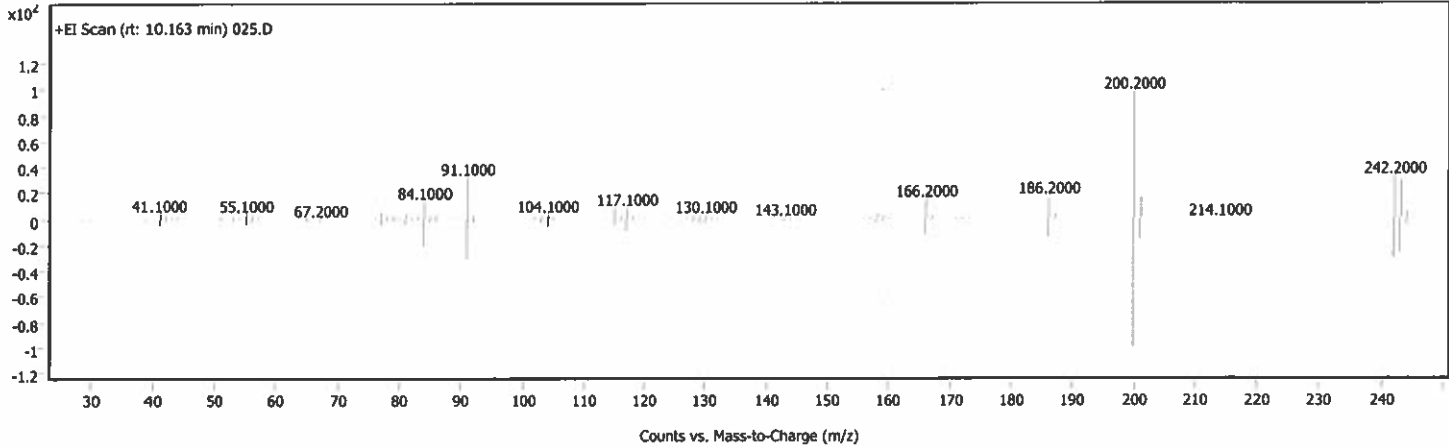
PHENCYCLIDINE # PCP; C17H25N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
PHENCYCLIDINE # PCP	C17H25N	98.37	aafs.I	MATRIX 1 W/ SW CTL + XYLAXINE - 8LD	KM/QC DATA IN XYLAXINE VALIDATION

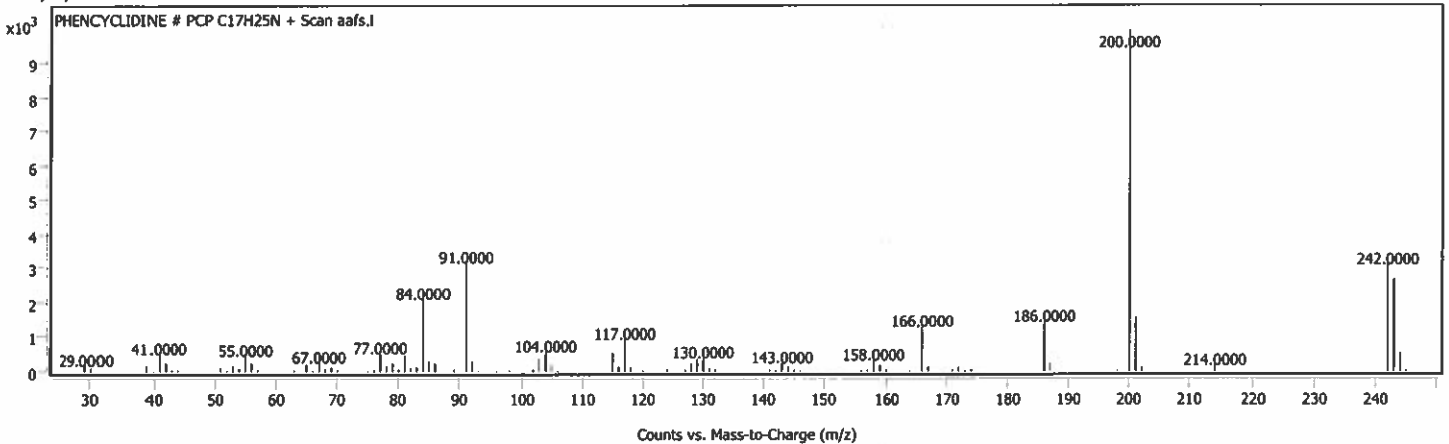
## Observed Spectrum



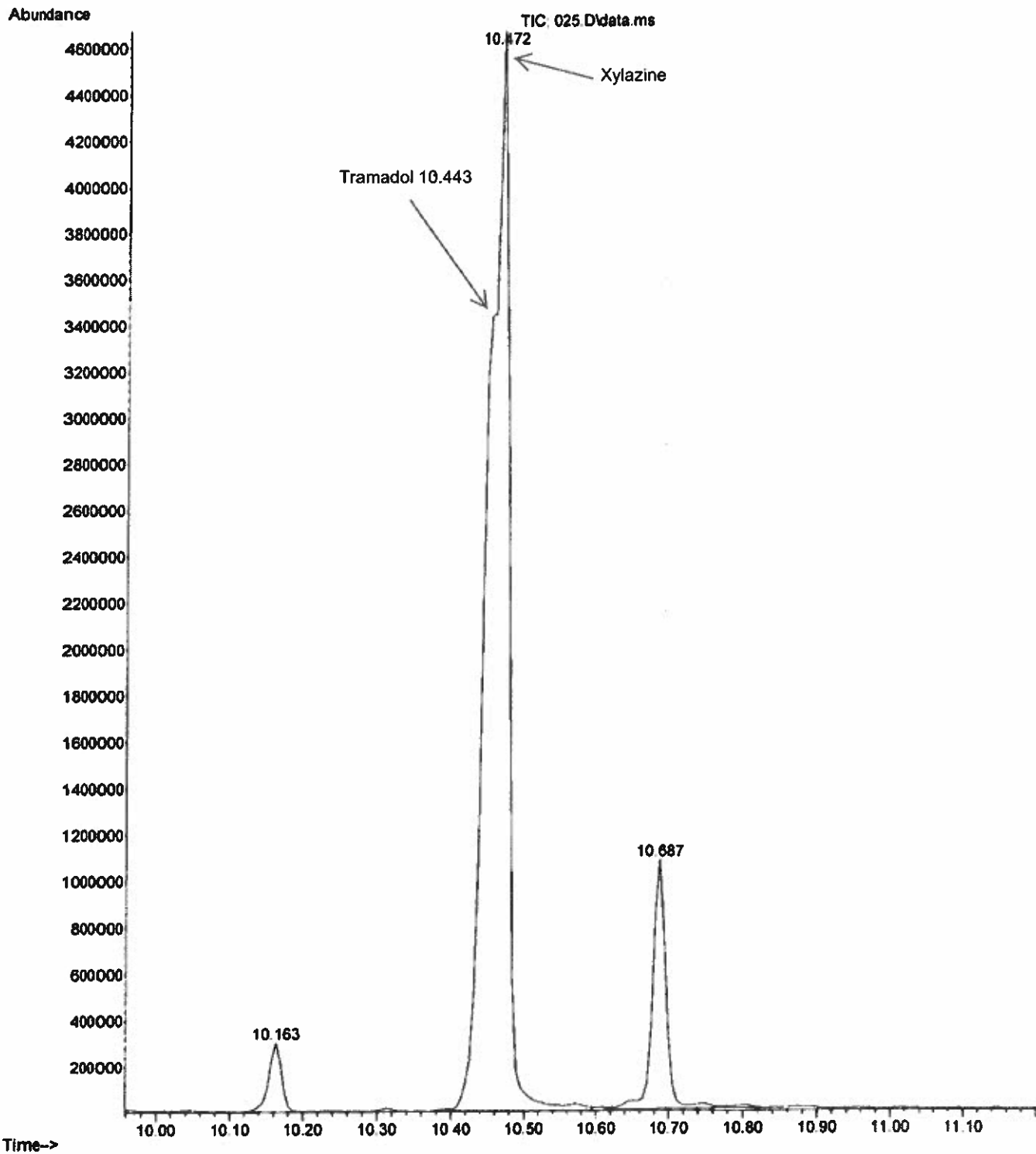
## Mirror Plot



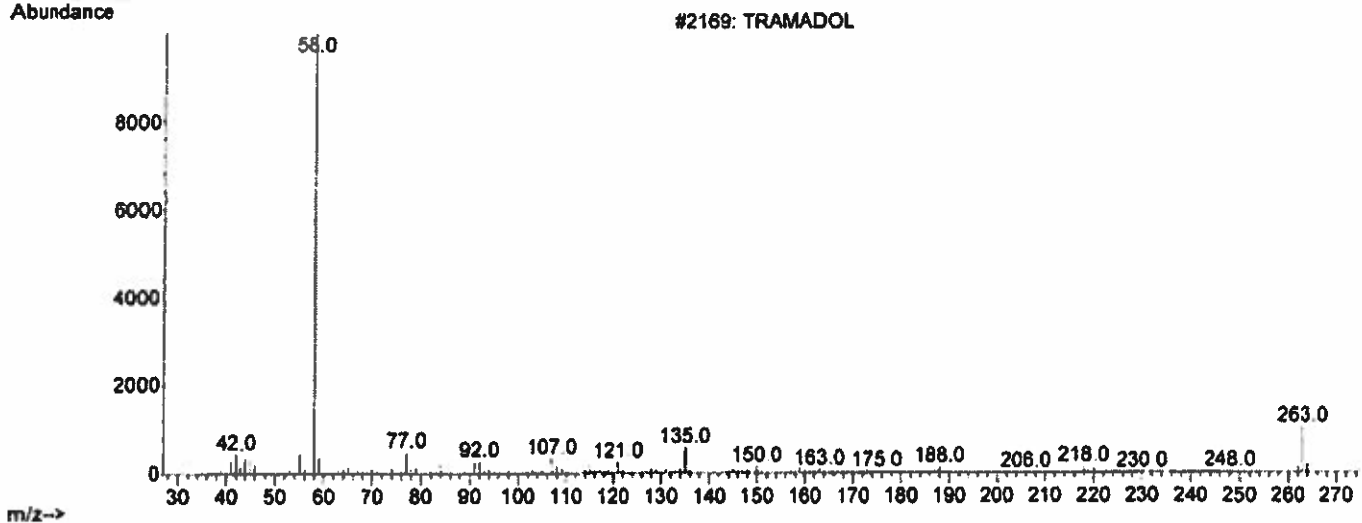
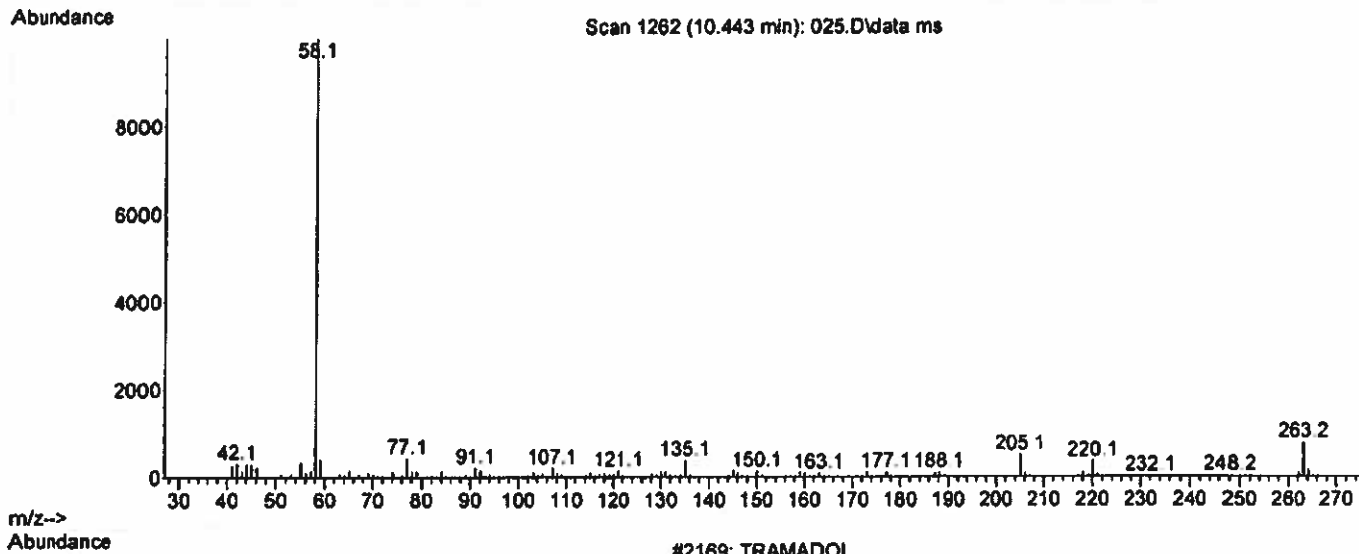
## Library Spectrum



File : C:\Users\TOX\Desktop\092821\025.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 00:52 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 1 W/ SW CTL + XYLAXINE - BLD  
Misc Info : 2.0 mg/L - 400uL of 0.01 mg/mL CTL, plus 400uL 0.1 mg/mL XYLAZINE  
Vial Number: 53



Library Searched : C:\MassHunter\Library\afgs.1  
Quality : 95  
ID : TRAMADOL  
Sample Name: MATRIX 1 W/ SW CTL + XYLAXINE - BLD  
File: C:\Users\TOX\Desktop\092821\025.D  
Operator: KM/QC DATA IN XYLAXINE VALIDATION  
Date Acquired: 29 Sep 2021 00:52  
Vial: 53



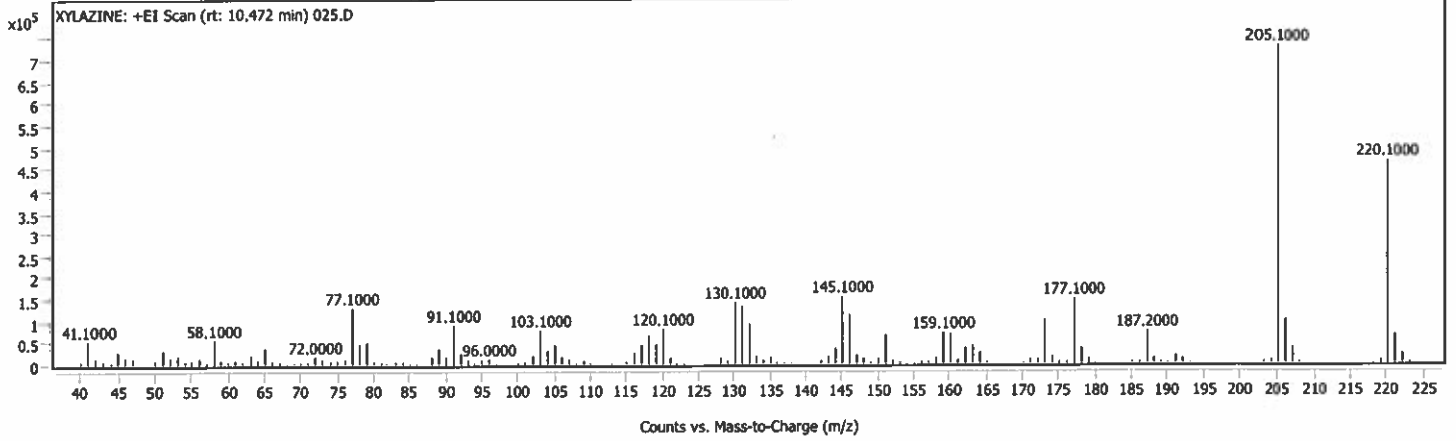
# Analysis Report

+ Scan (rt: 10.472 min)

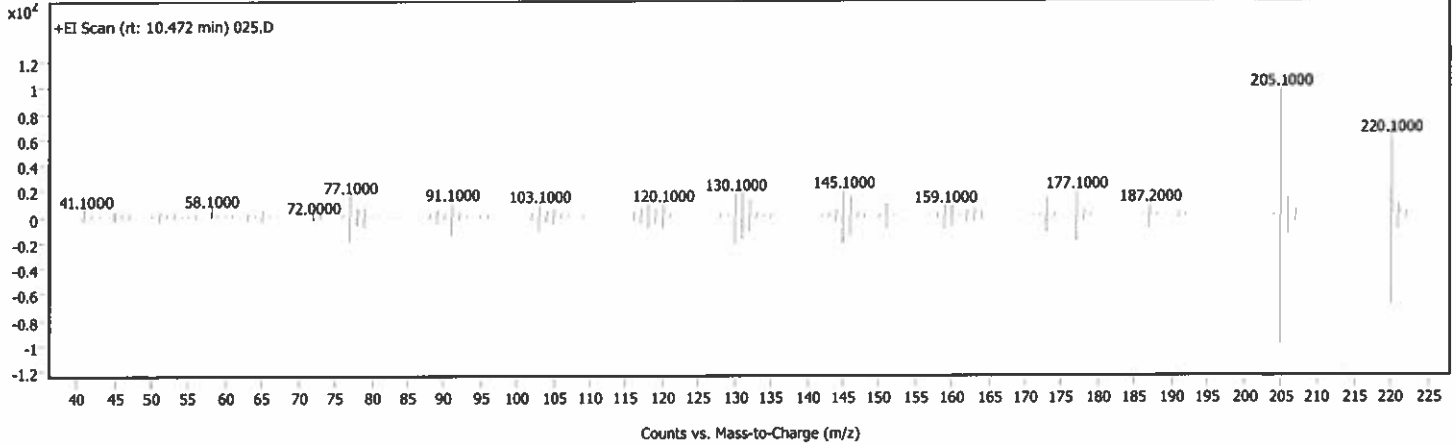
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.28	aafs.l	MATRIX 1 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

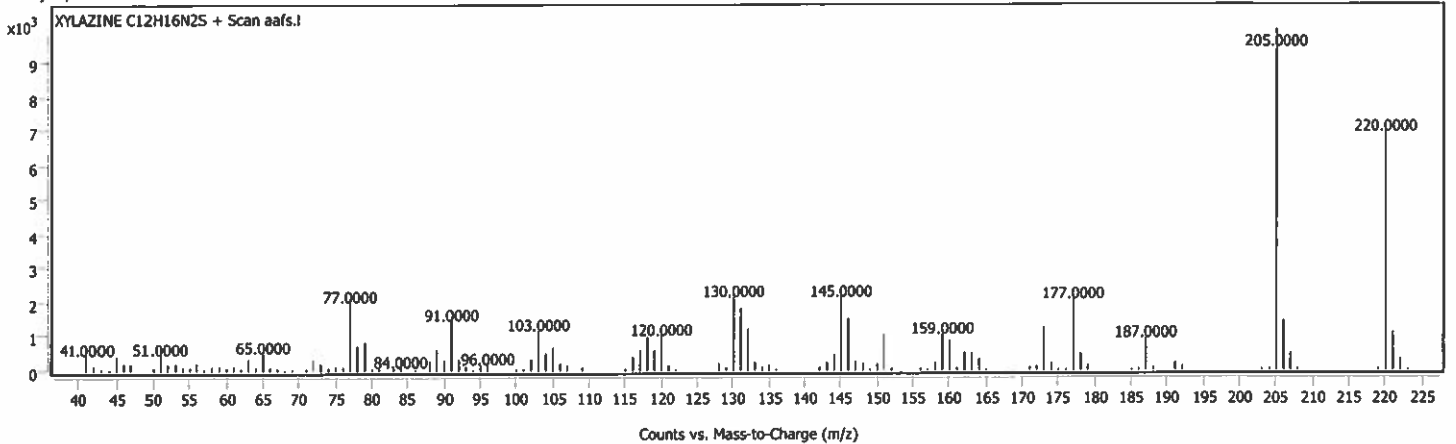
## Observed Spectrum



## Mirror Plot



## Library Spectrum



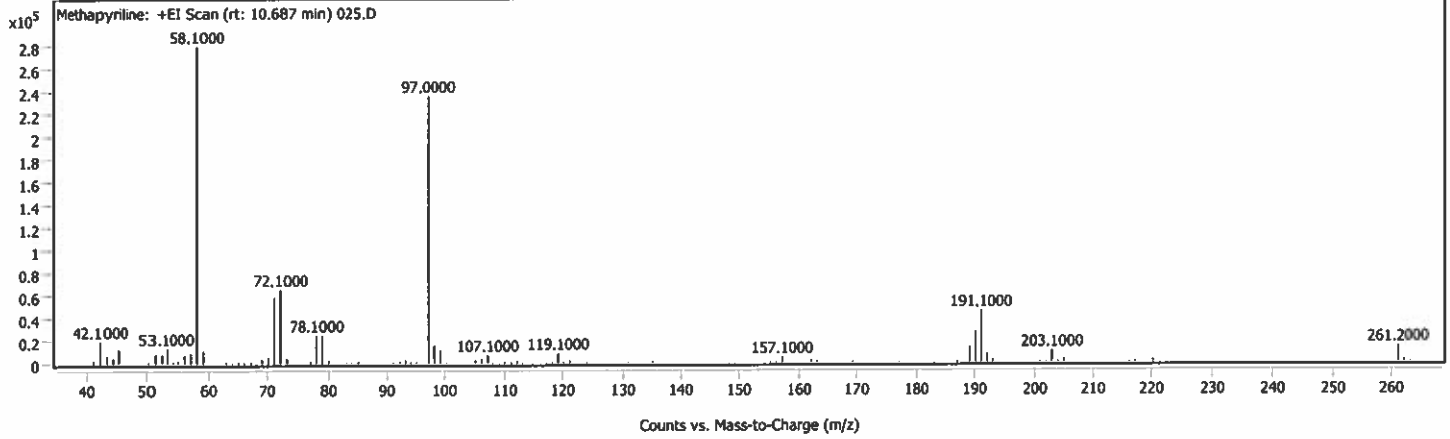
# Analysis Report

+ Scan (rt: 10.687 min)

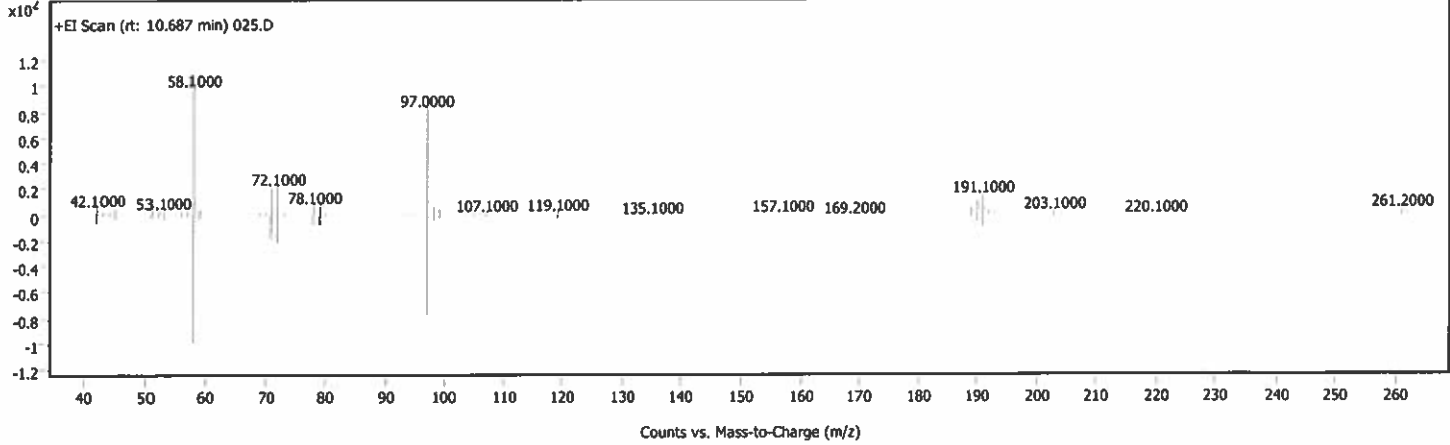
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	97.37	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL + XYLAXINE - BLD	KM/QC DATA IN XYLAXINE VALIDATION

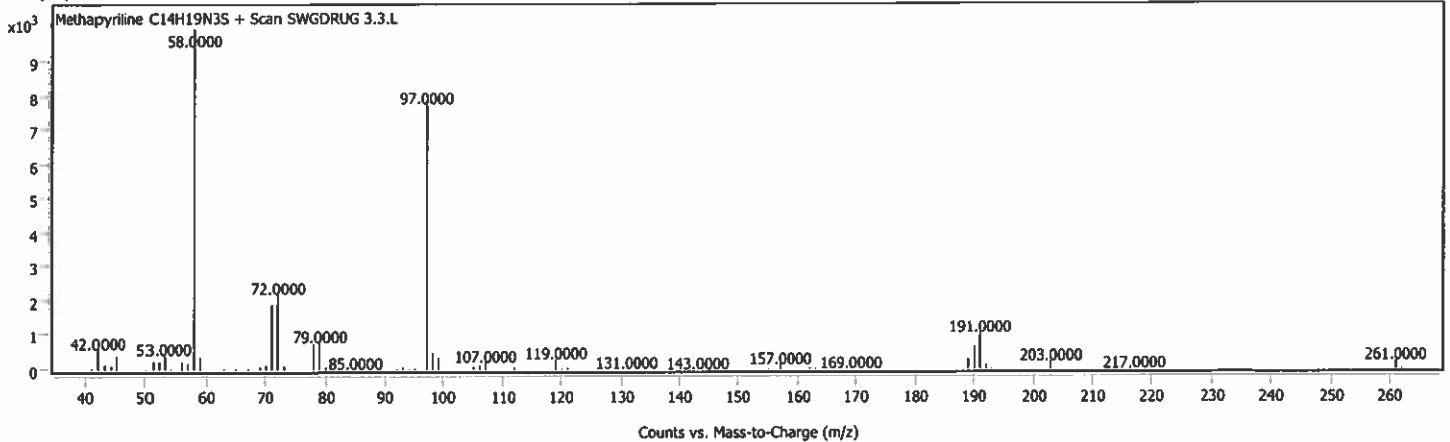
## Observed Spectrum



## Mirror Plot



## Library Spectrum



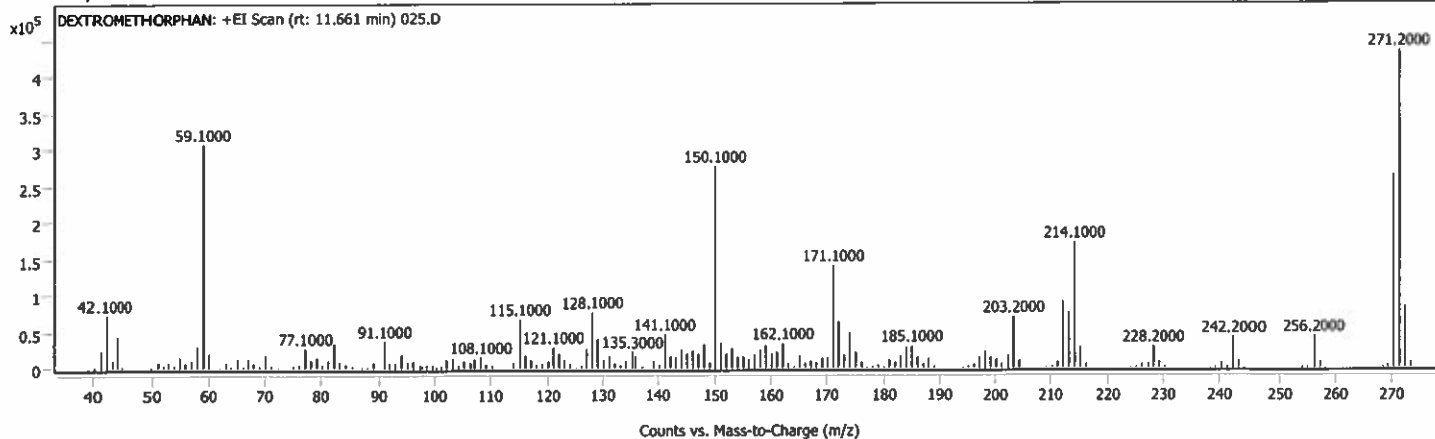
# Analysis Report

+ Scan (rt: 11.661 min)

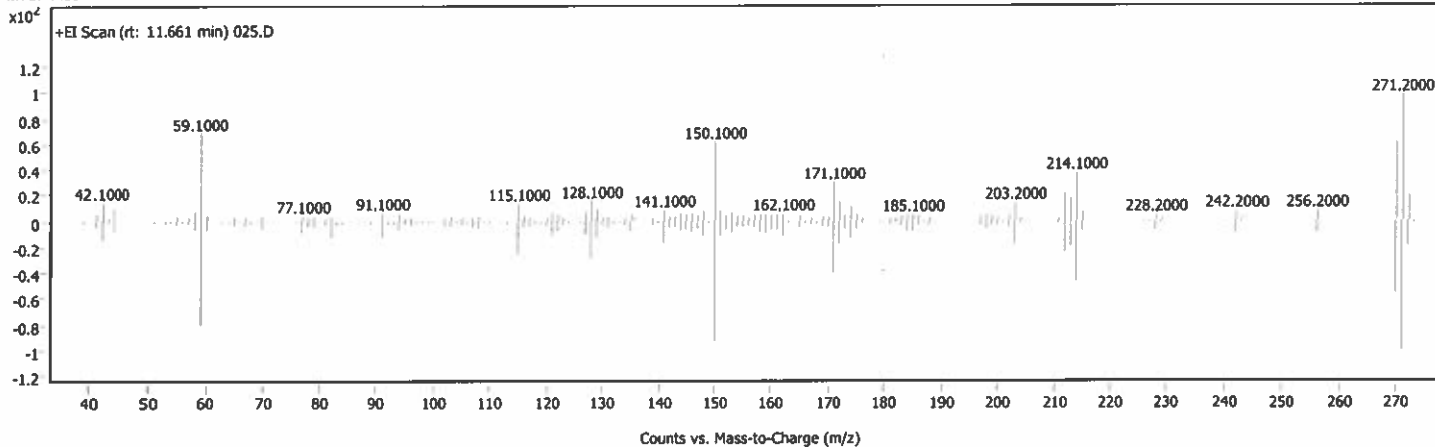
DEXTROMETHORPHAN; C18H25NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	91.99	aafs.l	MATRIX 1 W/ SW CTL + XYLAXINE - BLD	KM/QC DATA IN XYLAXINE VALIDATION

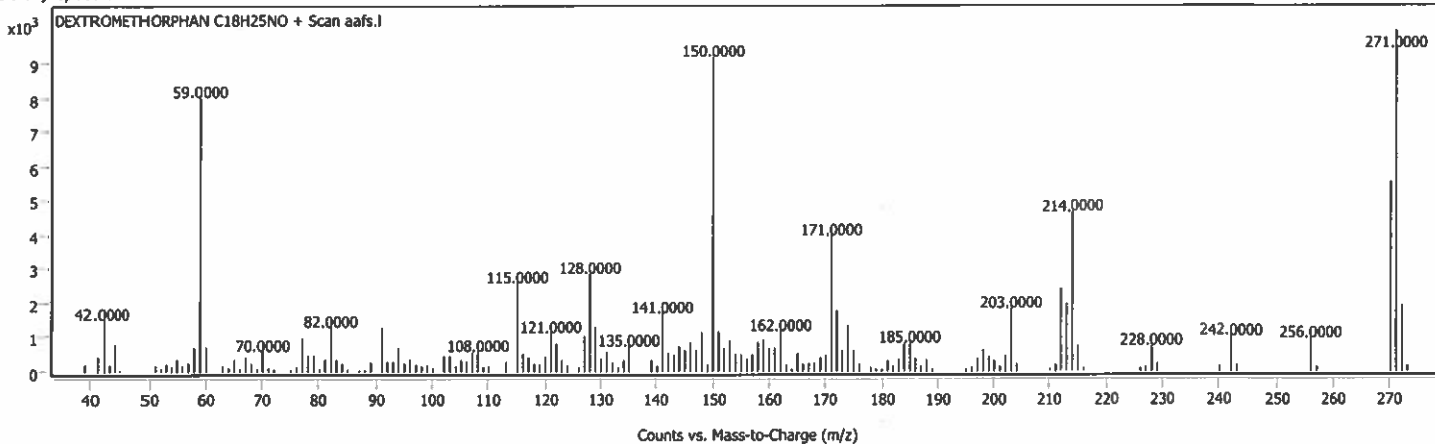
## Observed Spectrum



## Mirror Plot



## Library Spectrum



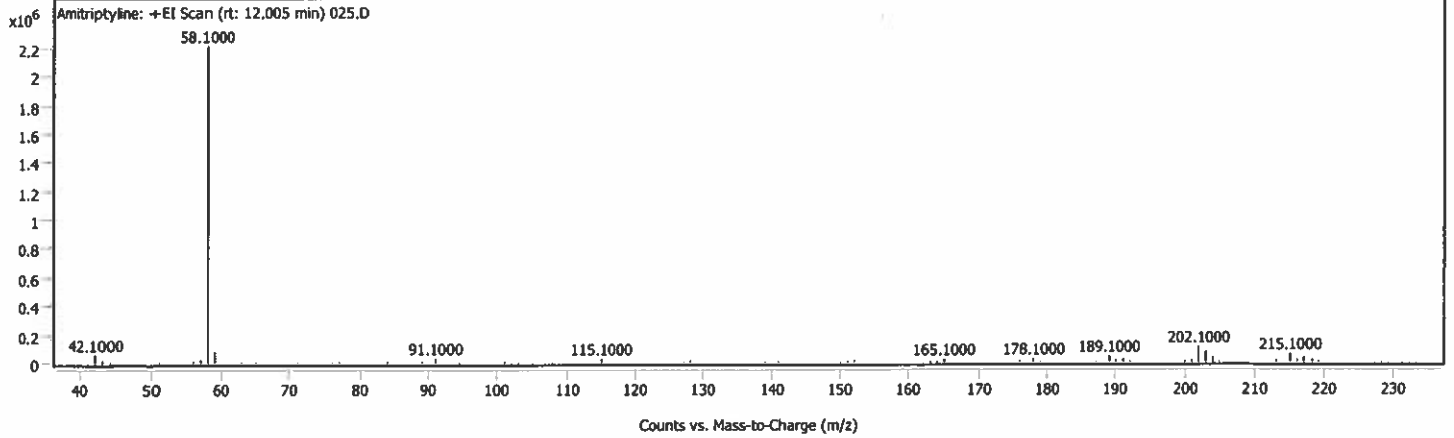
# Analysis Report

+ Scan (rt: 12.005 min)

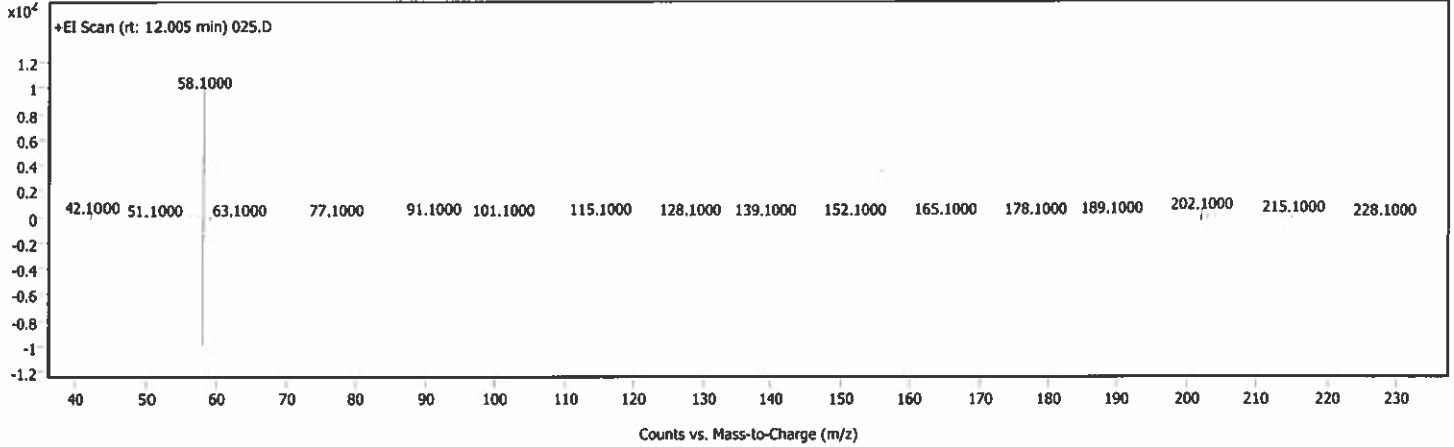
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	97.89	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL + XYLAXINE - BLD	KM/QC DATA IN XYLAXINE VALIDATION

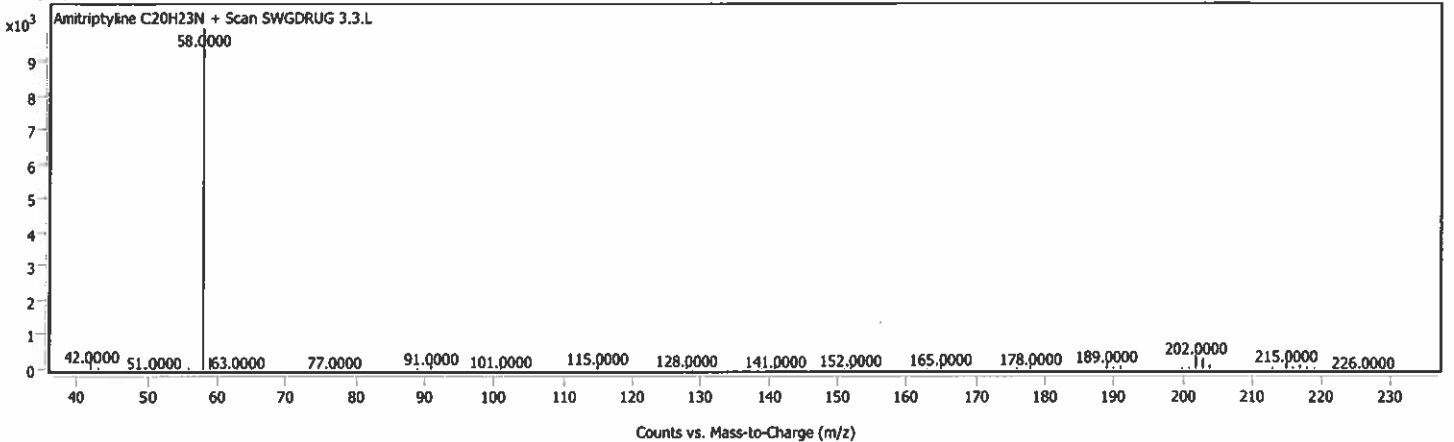
## Observed Spectrum



## Mirror Plot



## Library Spectrum



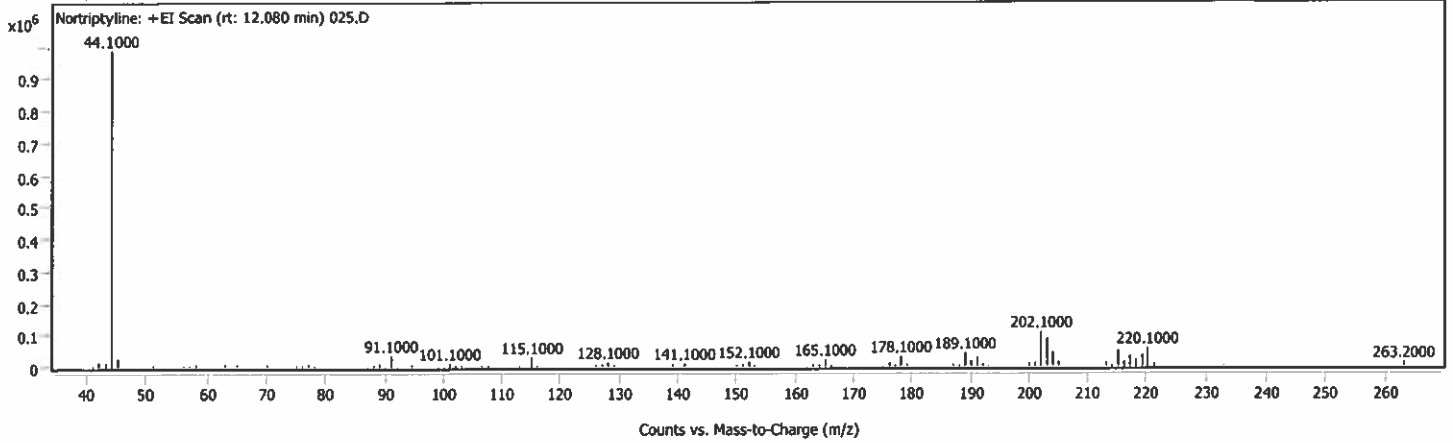
# Analysis Report

+ Scan (rt: 12.080 min)

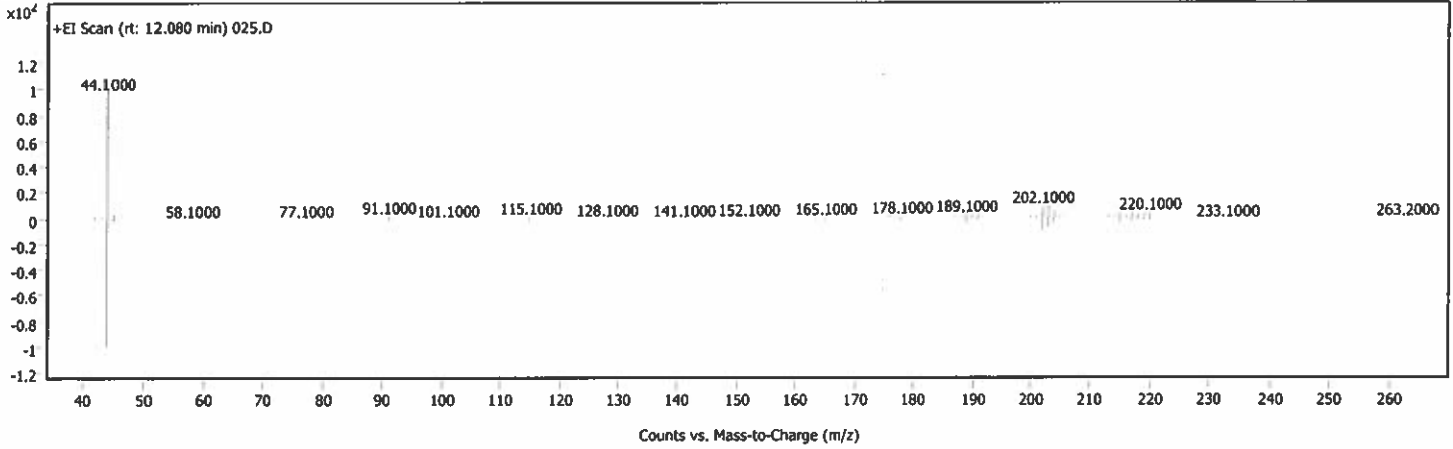
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	97.70	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

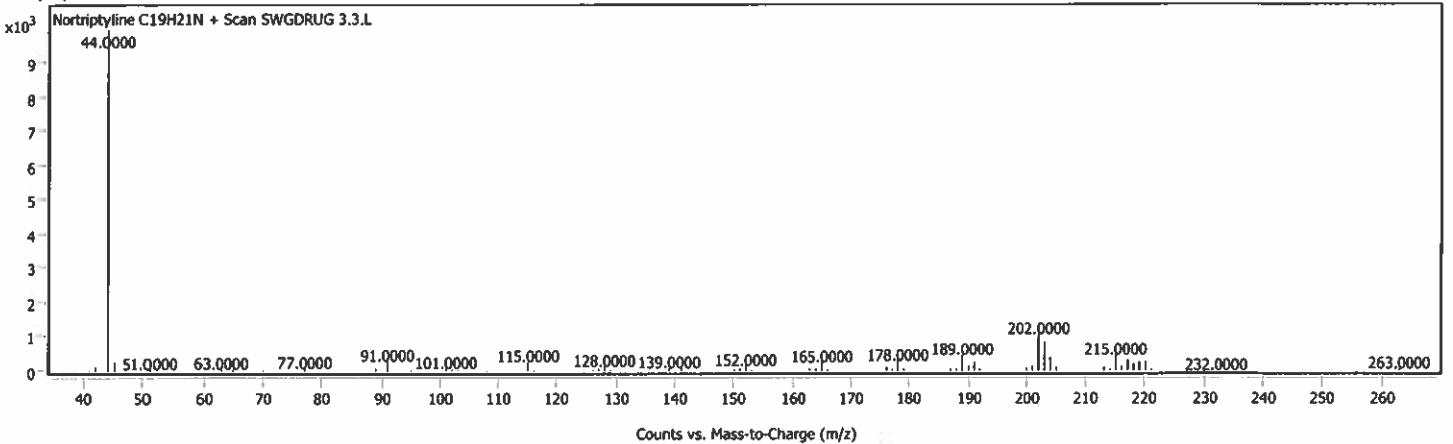
## Observed Spectrum



## Mirror Plot



## Library Spectrum





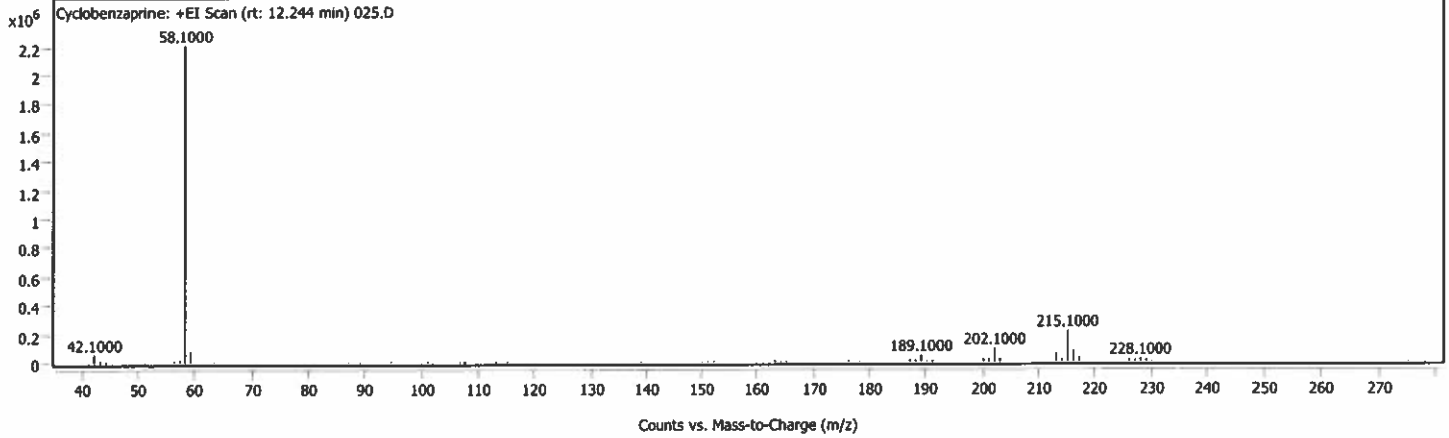
# Analysis Report

+ Scan (rt: 12.244 min)

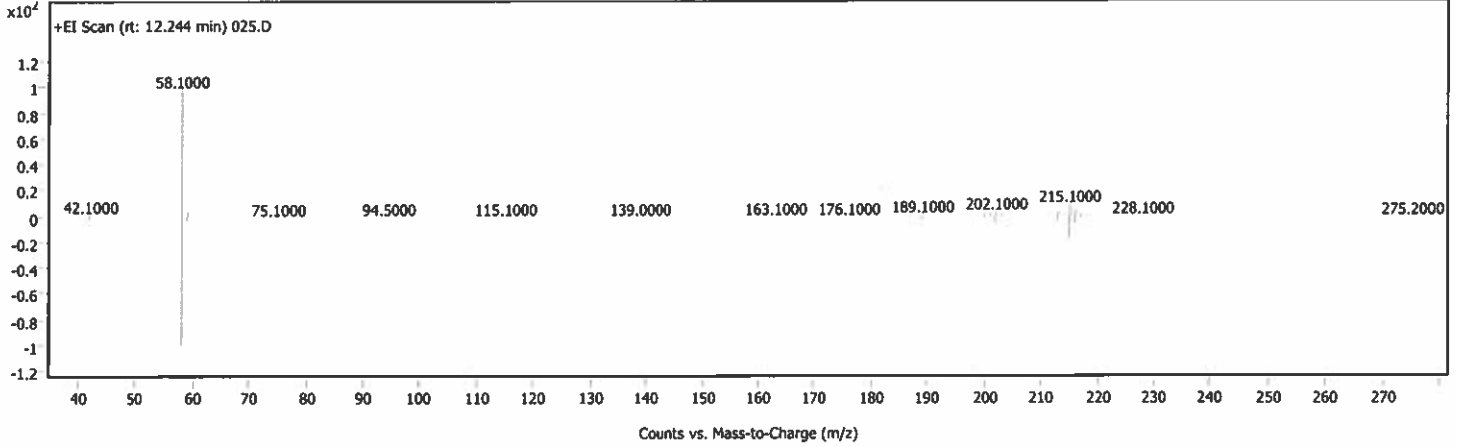
Cyclobenzaprine; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Cyclobenzaprine	C20H21N	96.87	SWGDRUG 3.3.L	MATRIX 1 W/ SW CTL + XYLAXINE - BLD	KM/QC DATA IN XYLAXINE VALIDATION

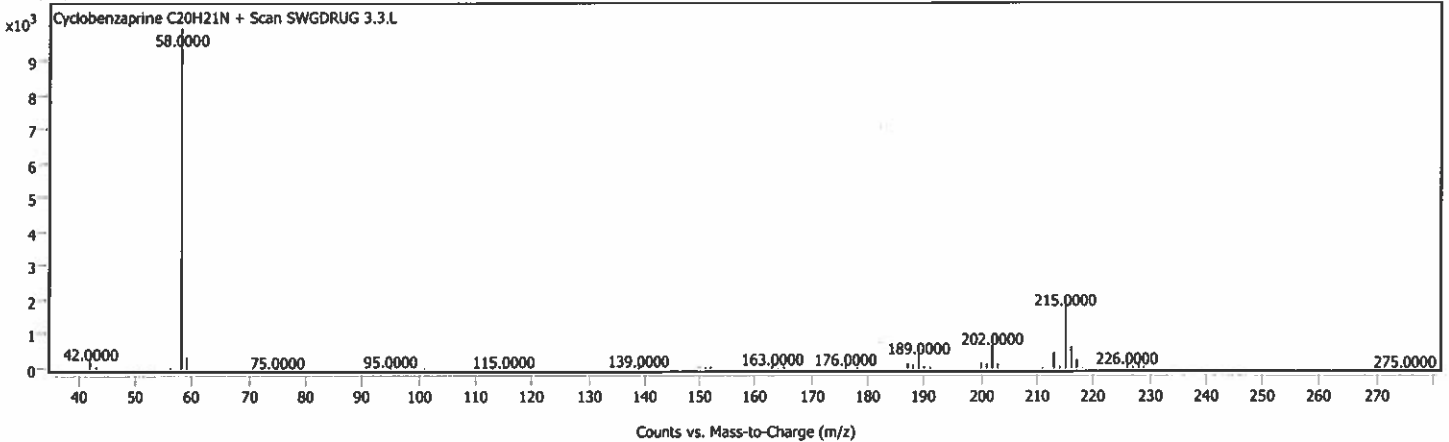
## Observed Spectrum



## Mirror Plot



## Library Spectrum



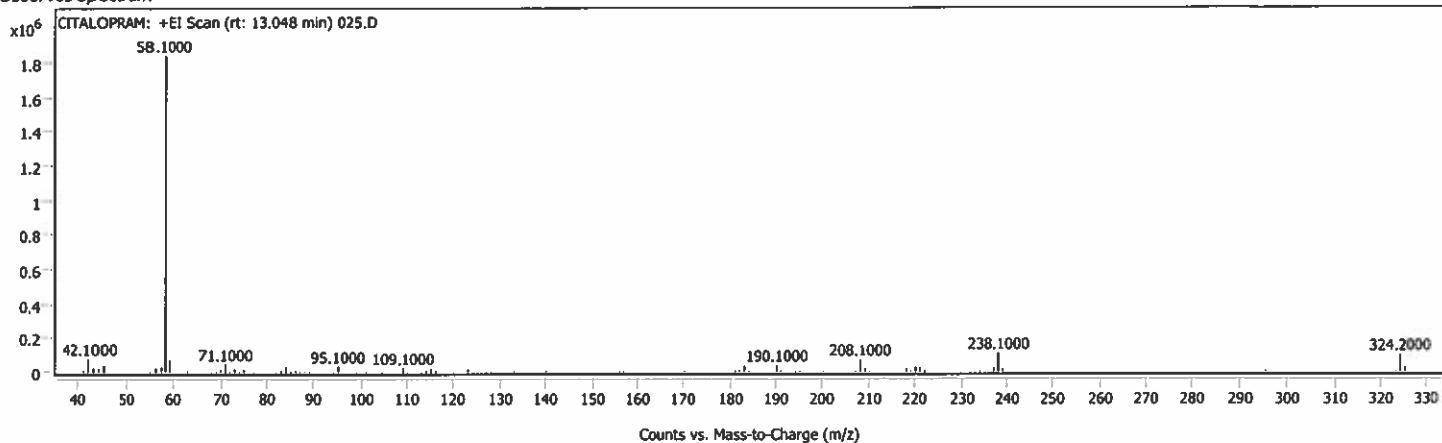
# Analysis Report

+ Scan (rt: 13.048 min)

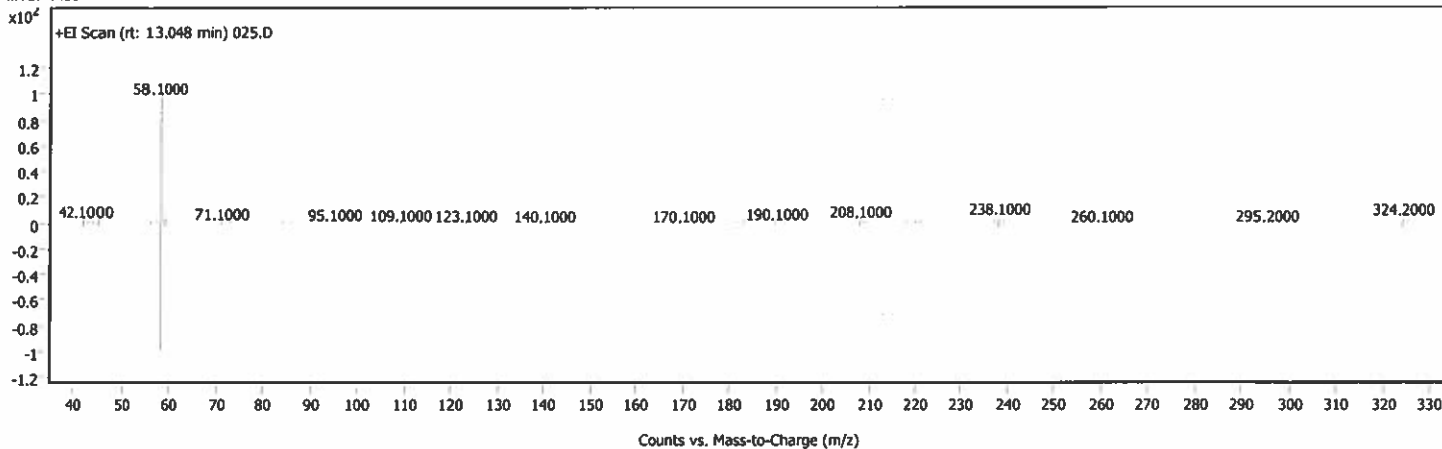
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	96.65	aafs.I	MATRIX 1 W/ SW CTL + XYLAZINE - 8LD	KM/QC DATA IN XYLAZINE VALIDATION

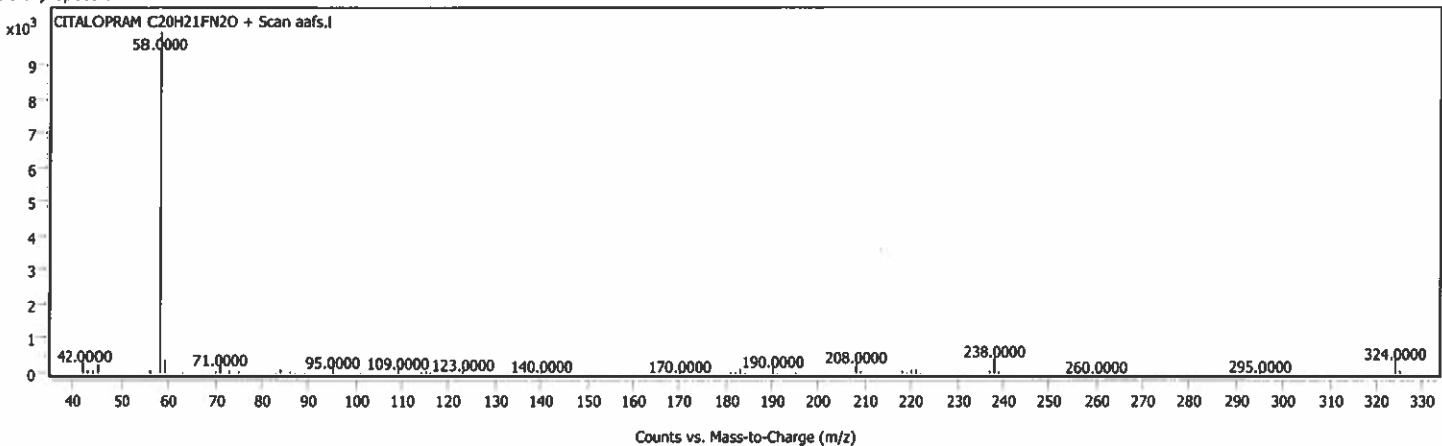
## Observed Spectrum



## Mirror Plot



## Library Spectrum



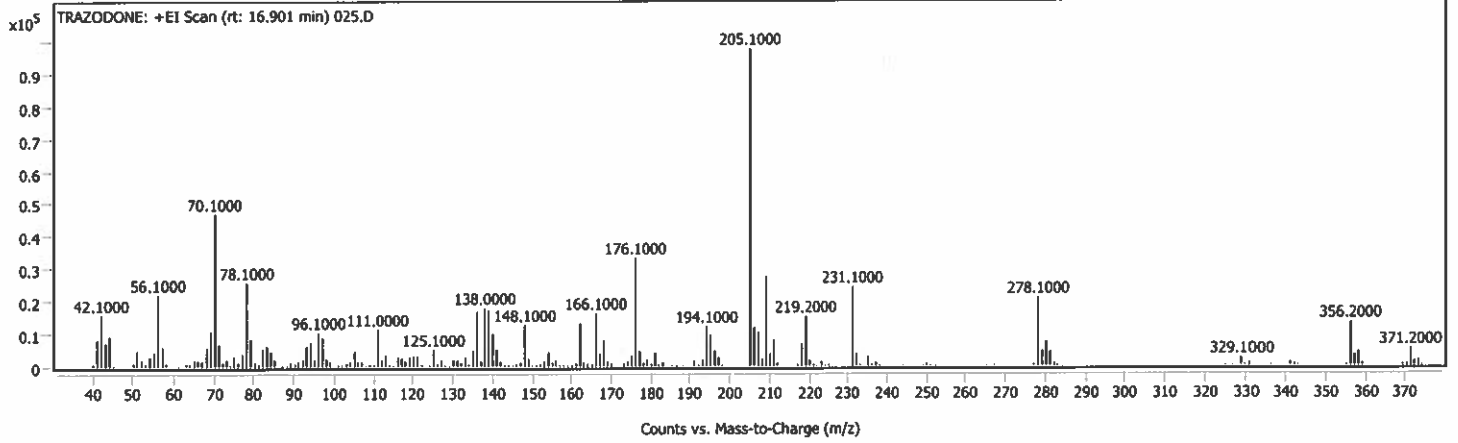
# Analysis Report

+ Scan (rt: 16.901 min)

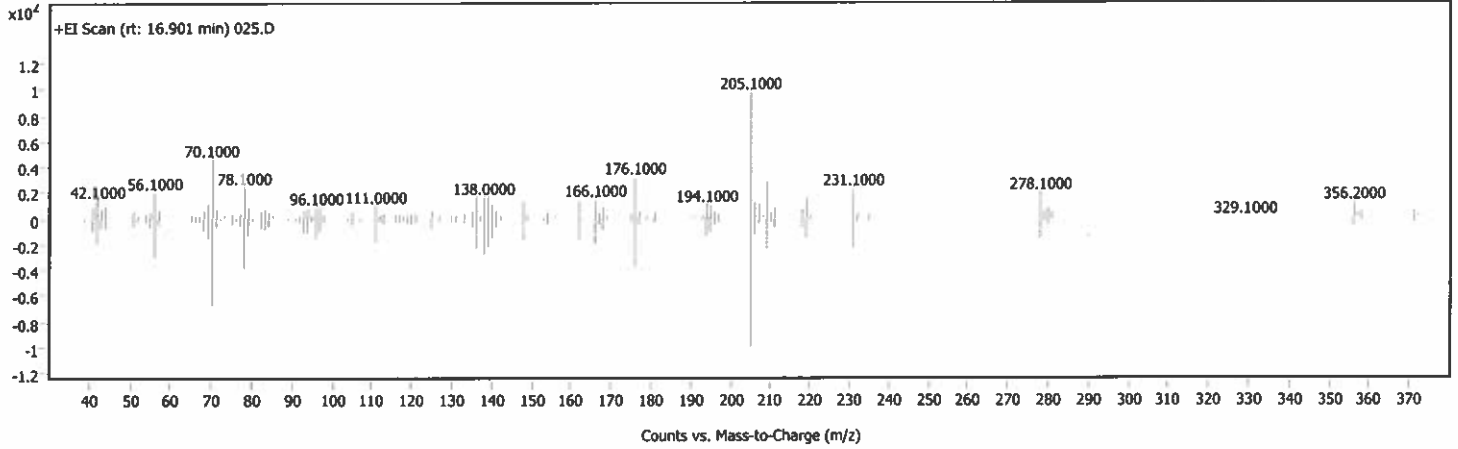
TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		94.10	aafs.l	MATRIX 1 W/ SW CTL + XYLAXINE - BLD	KM/QC DATA IN XYLAXINE VALIDATION

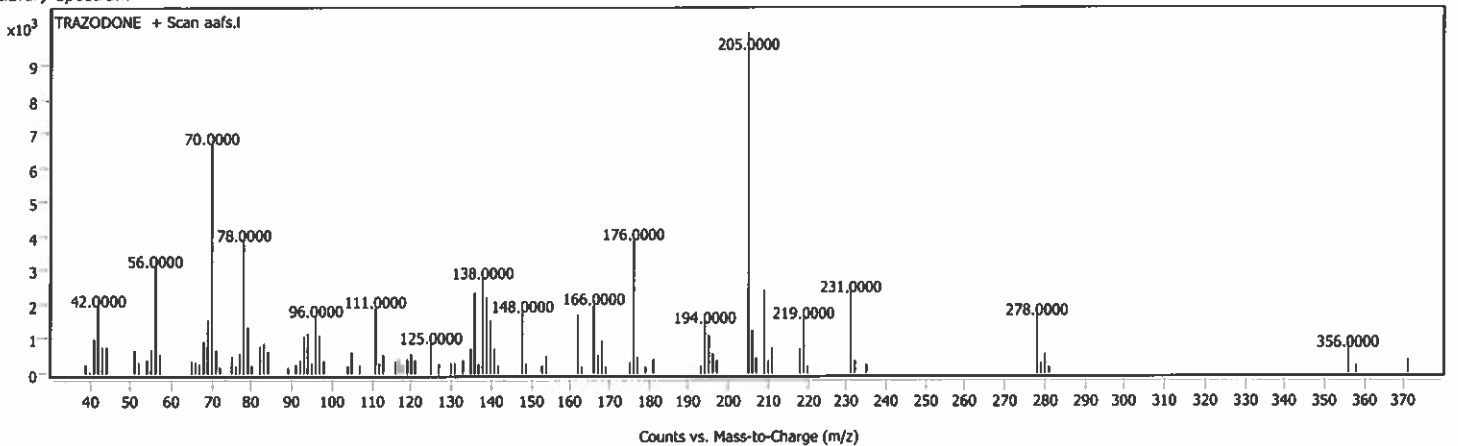
## Observed Spectrum



## Mirror Plot

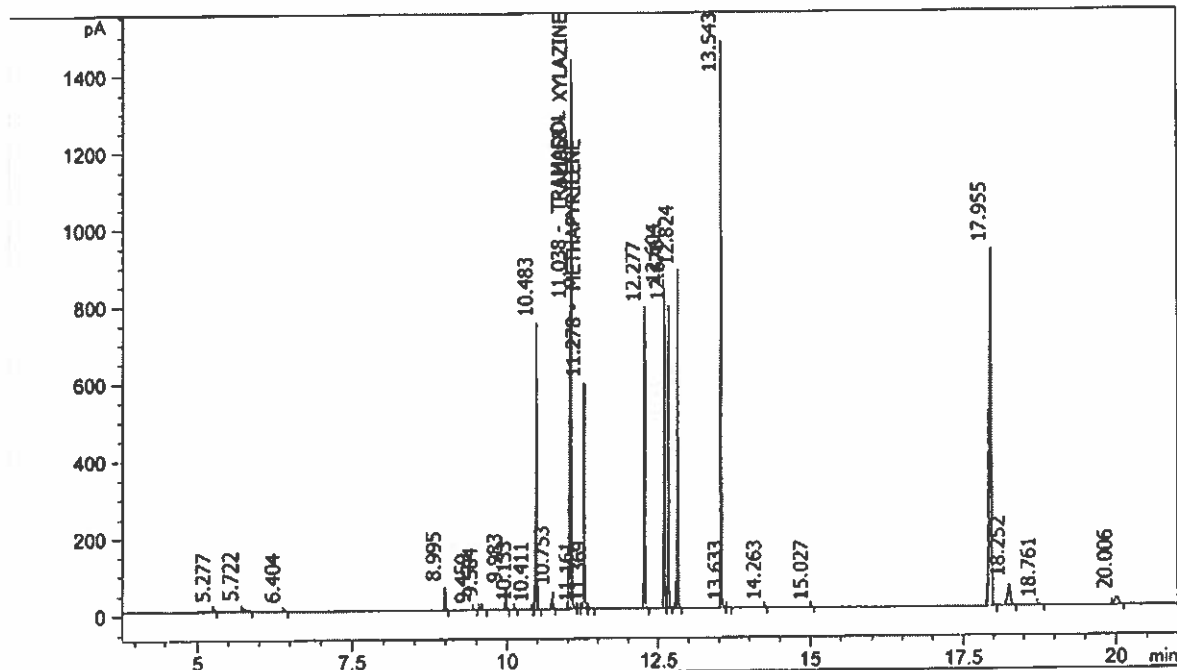


## Library Spectrum



Injection Date: 9/29/2021 12:42:03 AM Seq Line: 28  
 Sample Name: MATRIX 1 W/ SW CTL + -> Vial 29  
 Sample Info: *XYLAZINE KM 11/17/22*

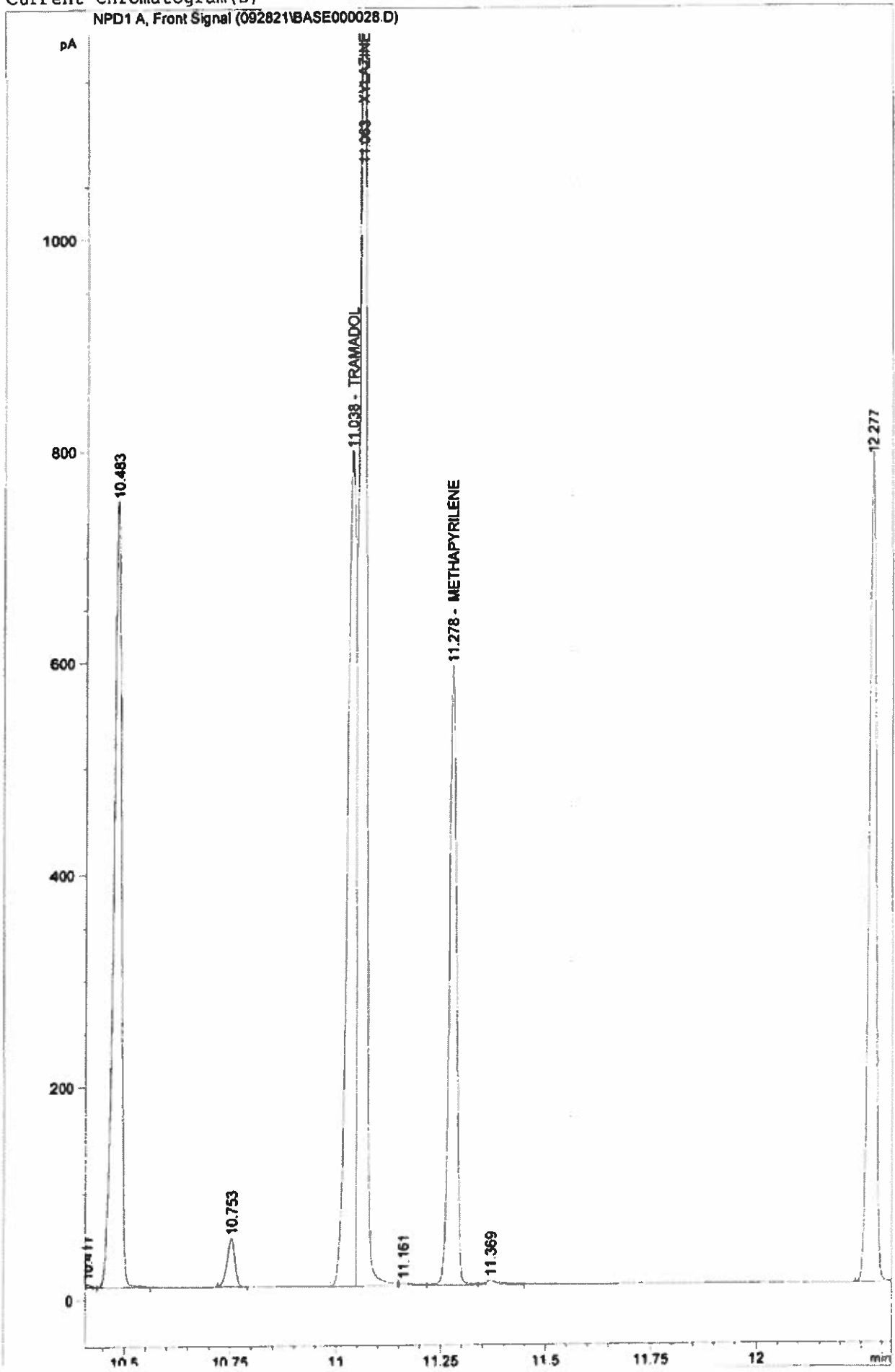
Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



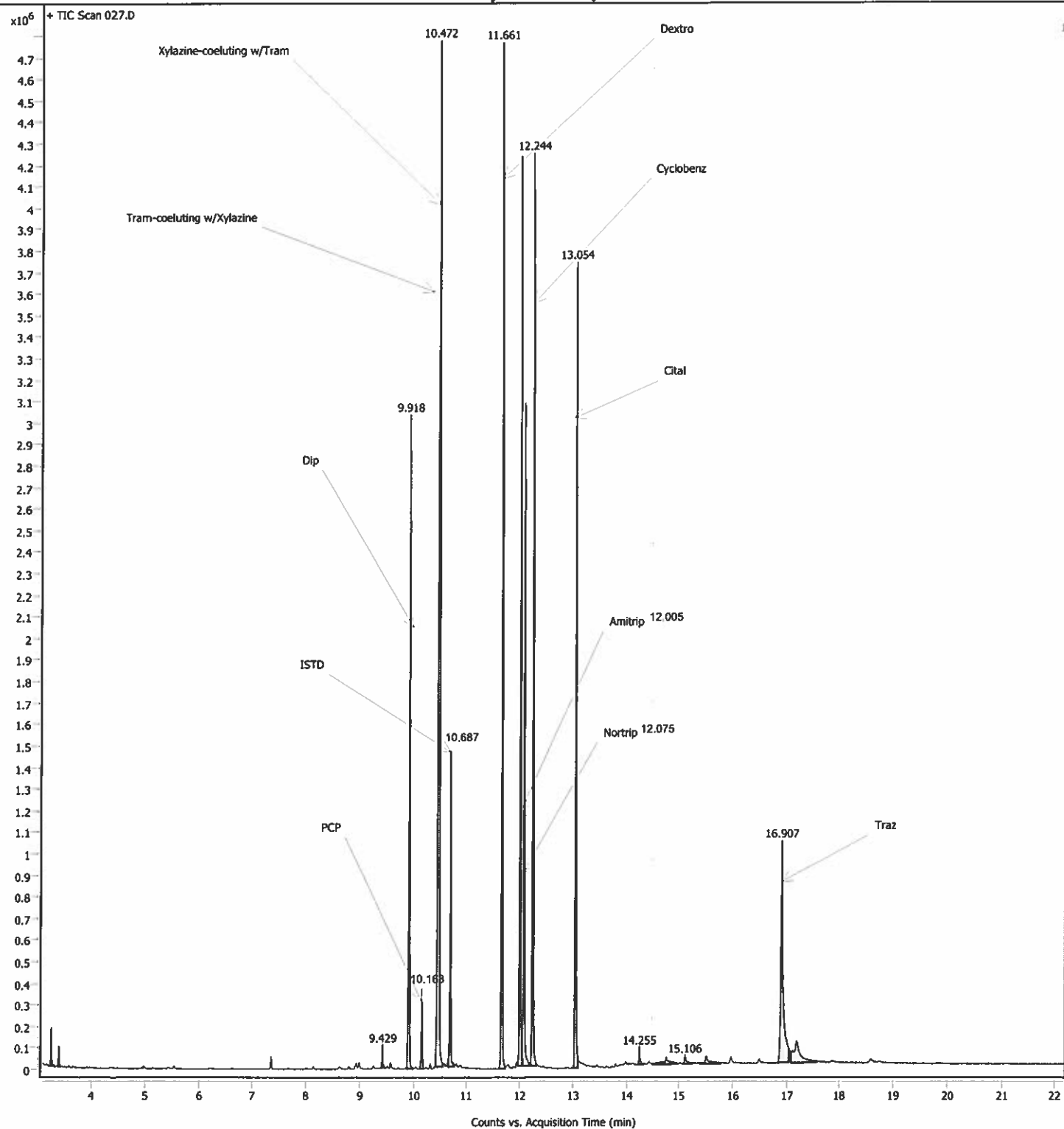
see following page for expanded chromatogram

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.038	11.033	1103.519	786.955	0.000000	TRAMADOL RRT-0.9787
11.063	11.046	1646.089	1424.104	0.000000	XYLAZINE RRT-0.9809
11.278	11.278	687.821	583.640	1.000000	METHAPYRILENE

Current Chromatogram (s)  
NPD1 A, Front Signal (092821\BASE000028.D)



# Analysis Report



# Analysis Report

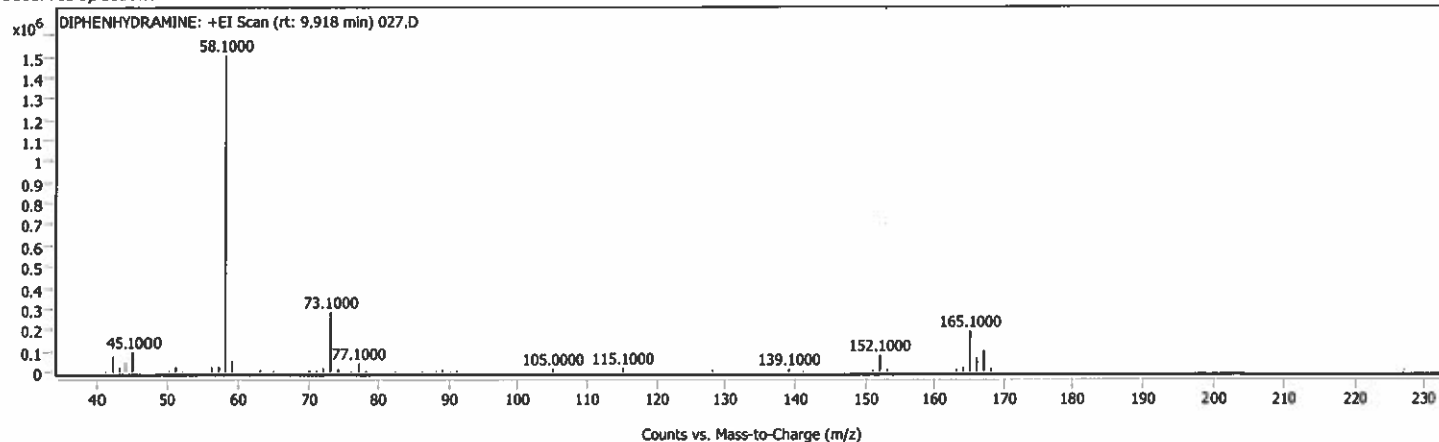
## Sample Spectra

+ Scan (rt: 9.918 min)

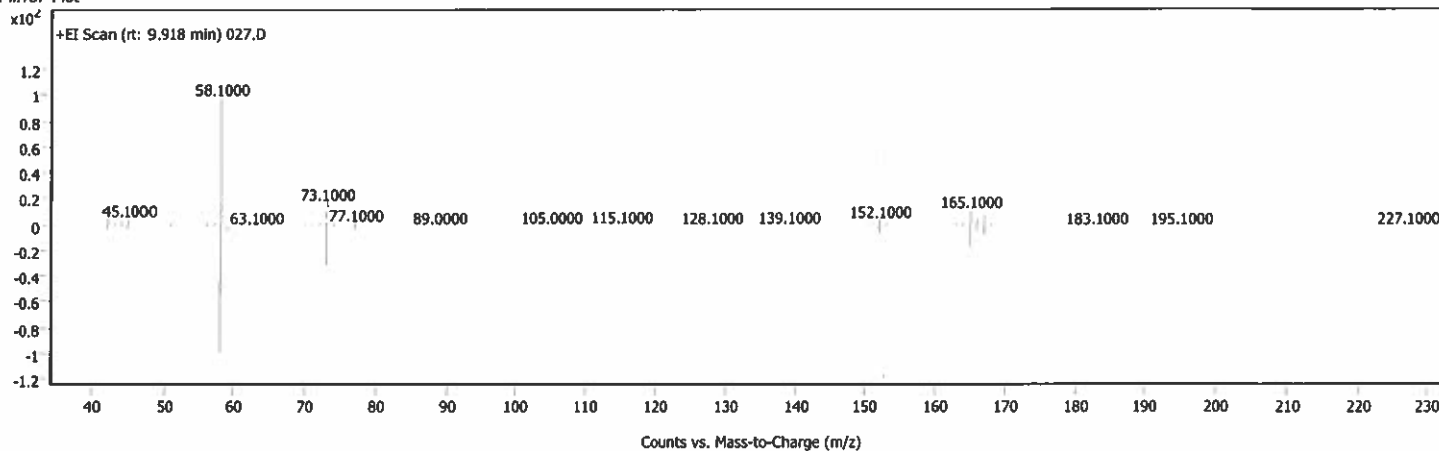
DIPHENHYDRAMINE; C17H21NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DIPHENHYDRAMINE	C17H21NO	98.21	aafs.I	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

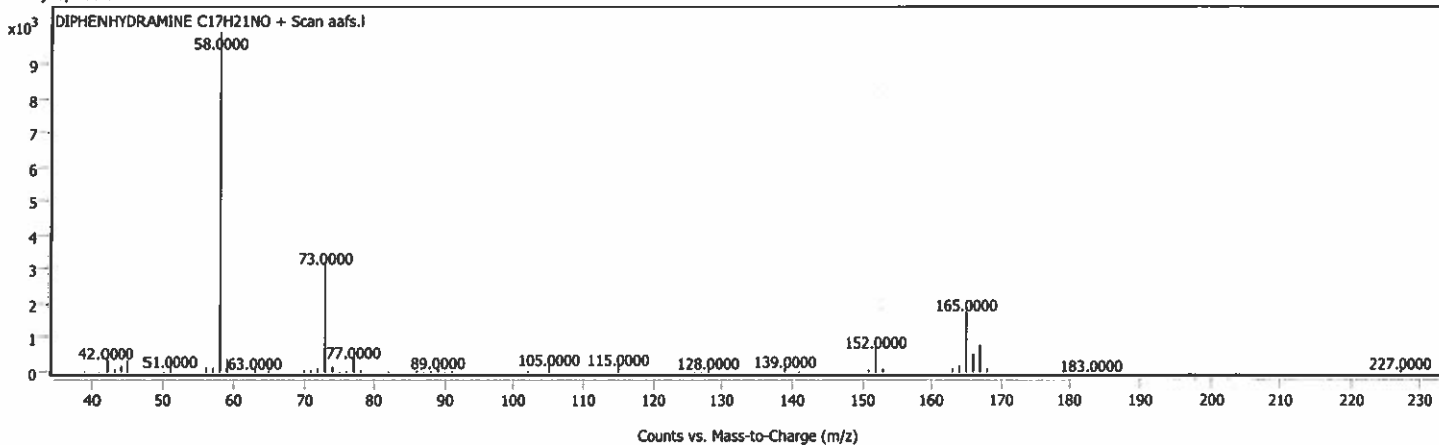
## Observed Spectrum



## Mirror Plot



## Library Spectrum



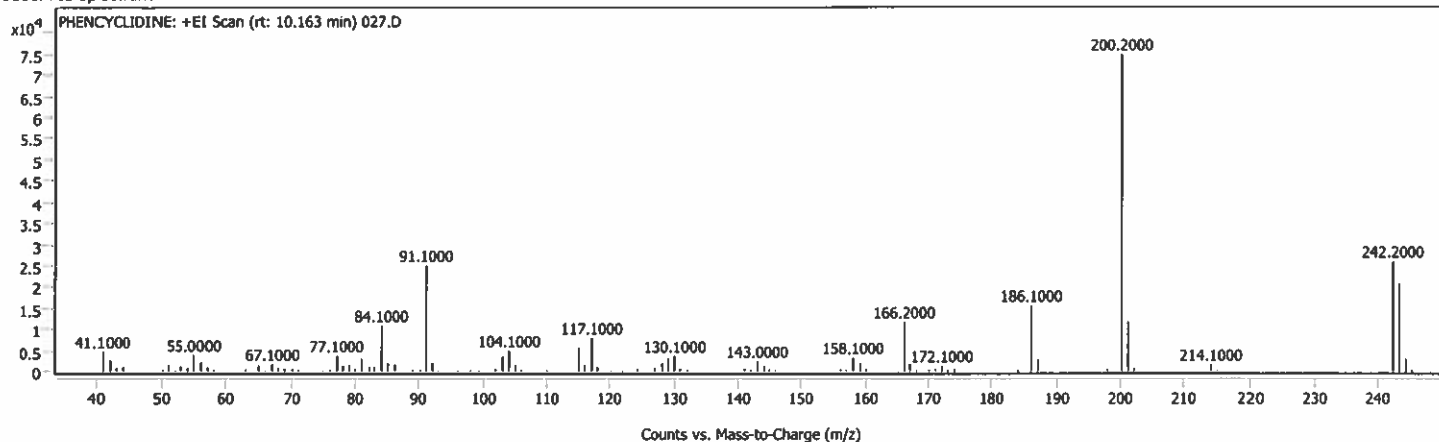
# Analysis Report

+ Scan (rt: 10.163 min)

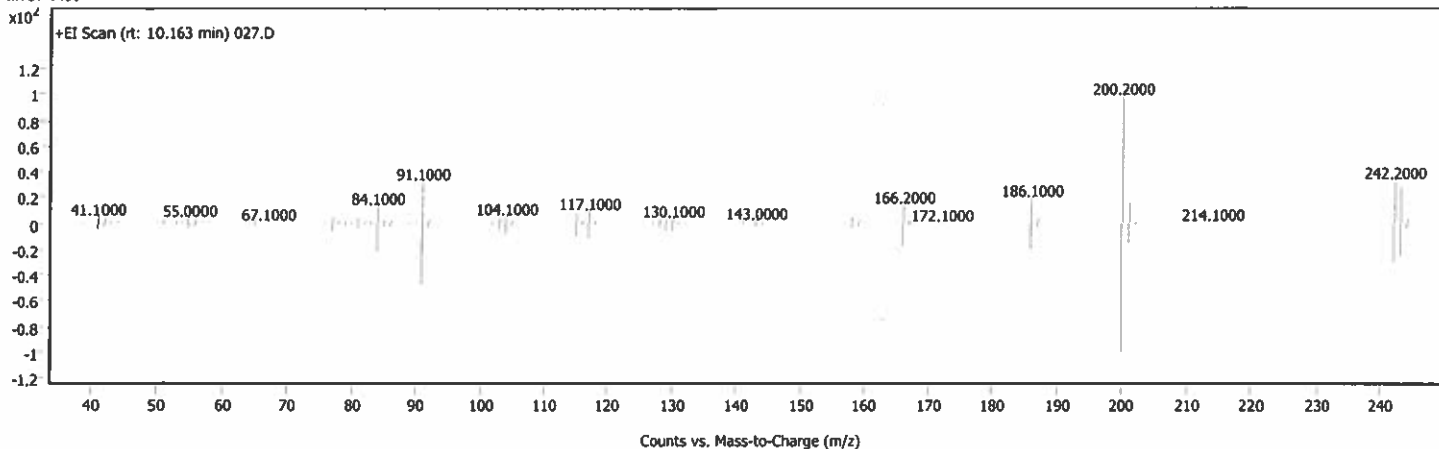
PHENCYCLIDINE; C17H25N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
PHENCYCLIDINE	C17H25N	98.55	aafs.I	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

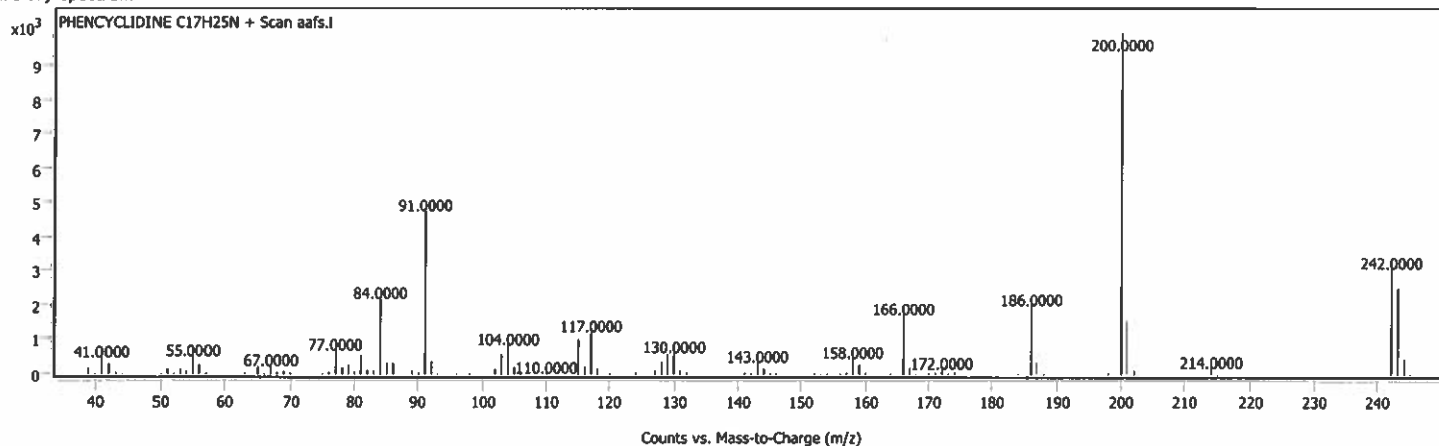
## Observed Spectrum



## Mirror Plot

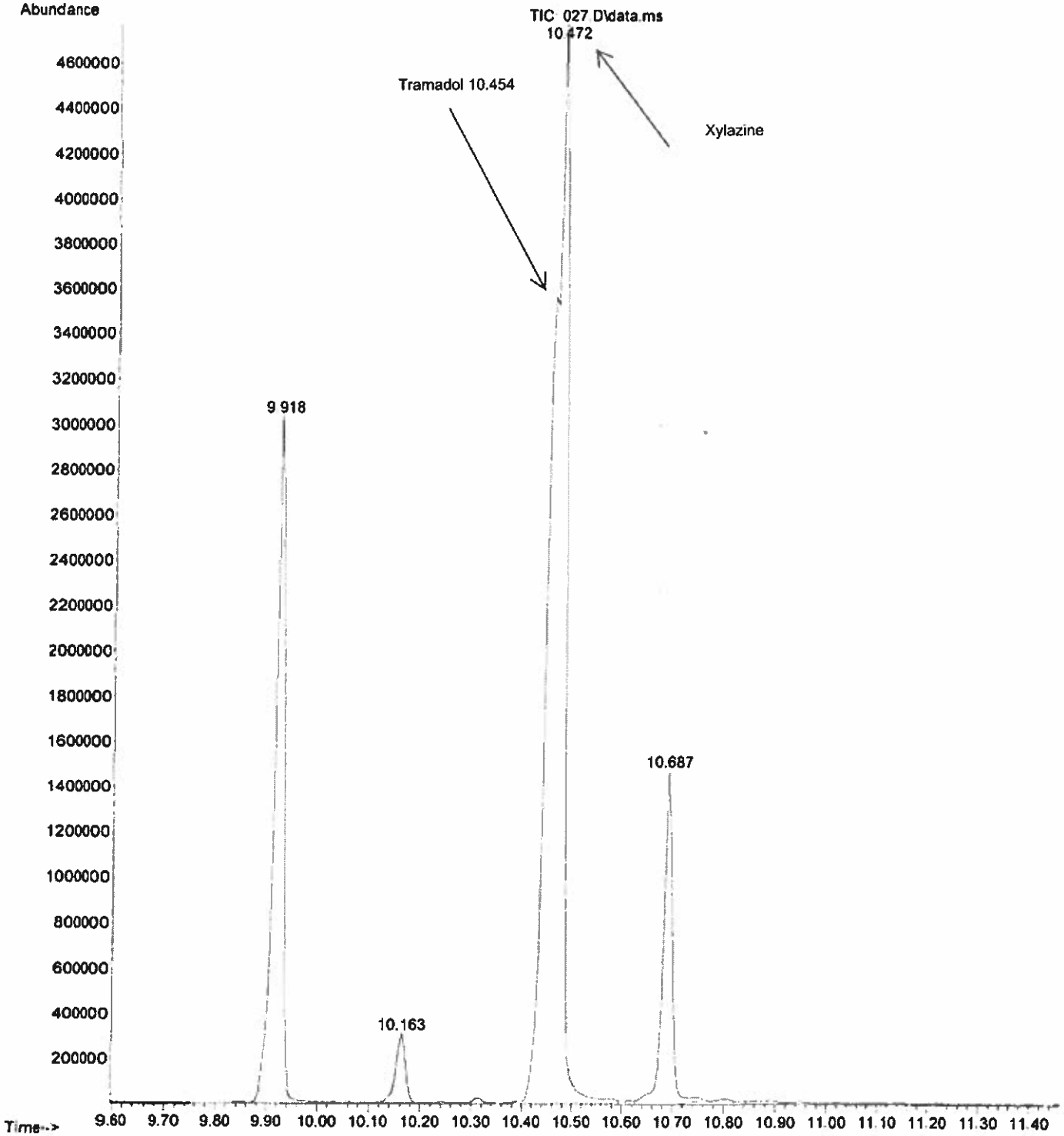


## Library Spectrum

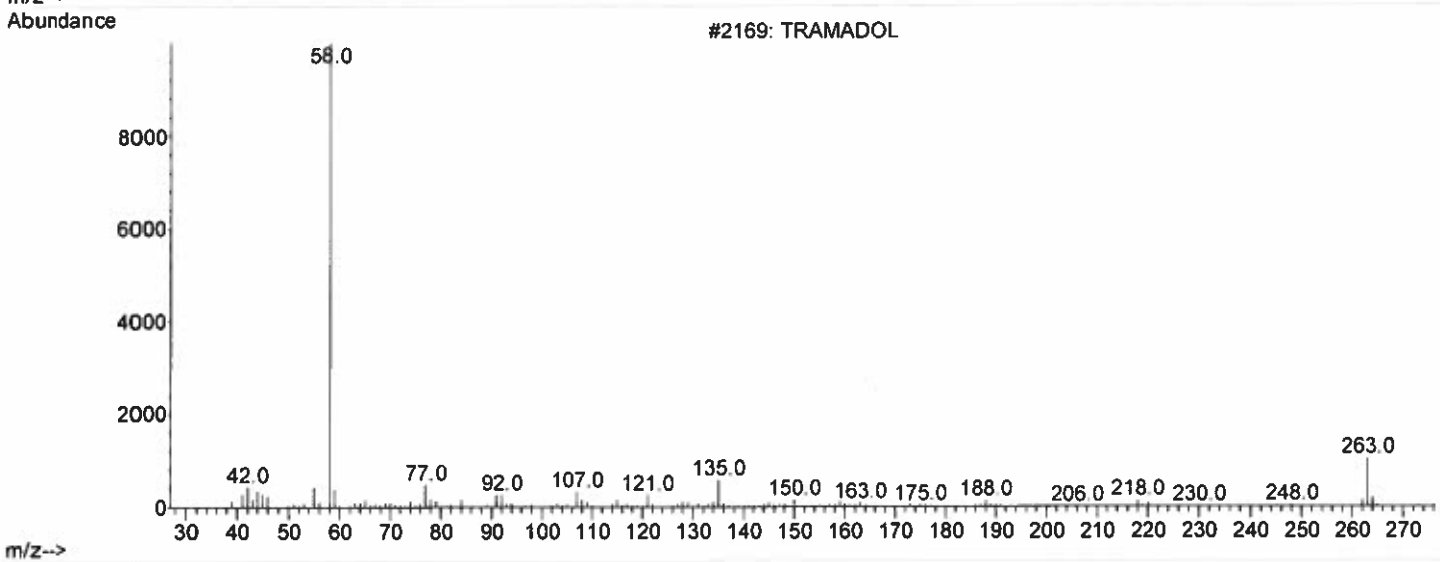
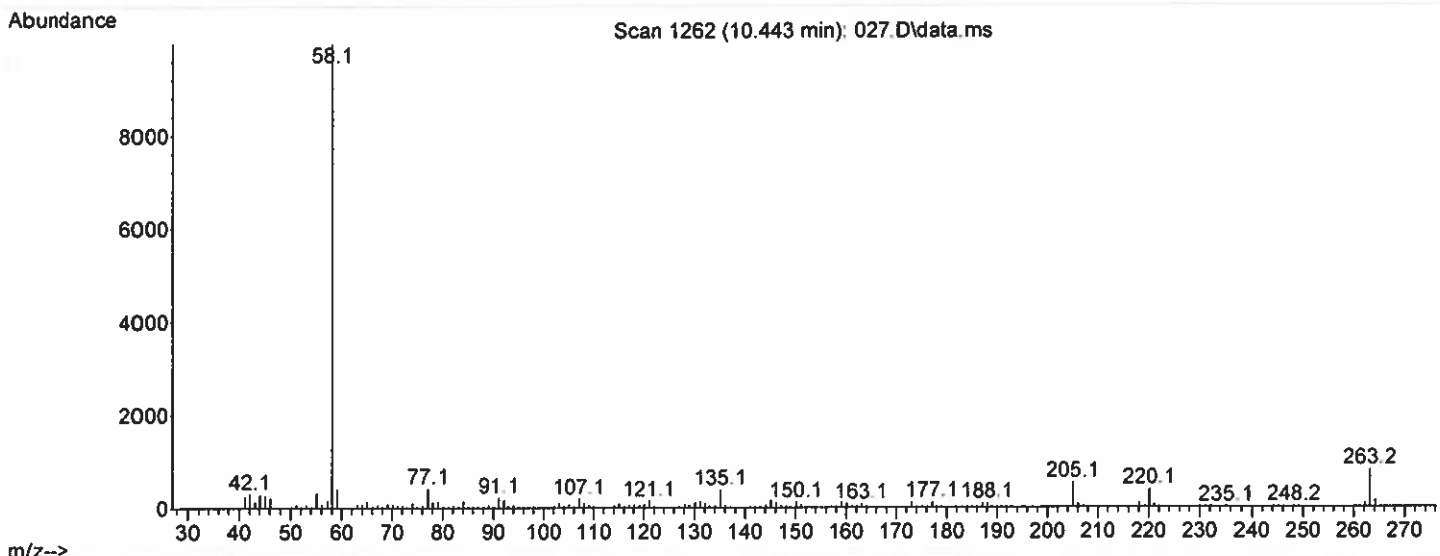




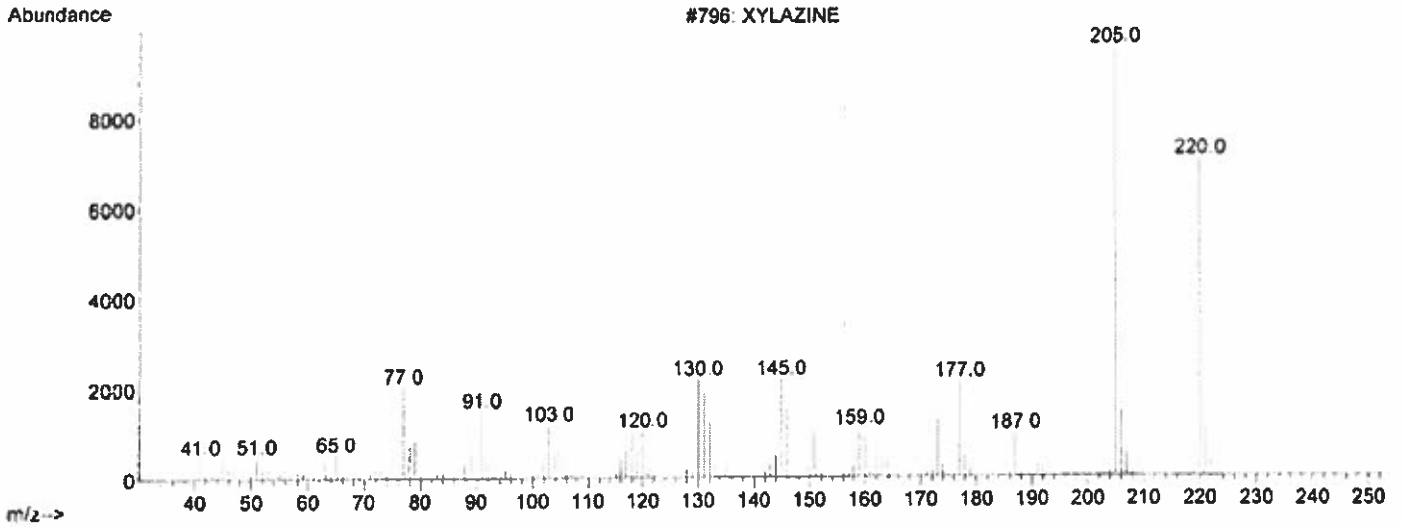
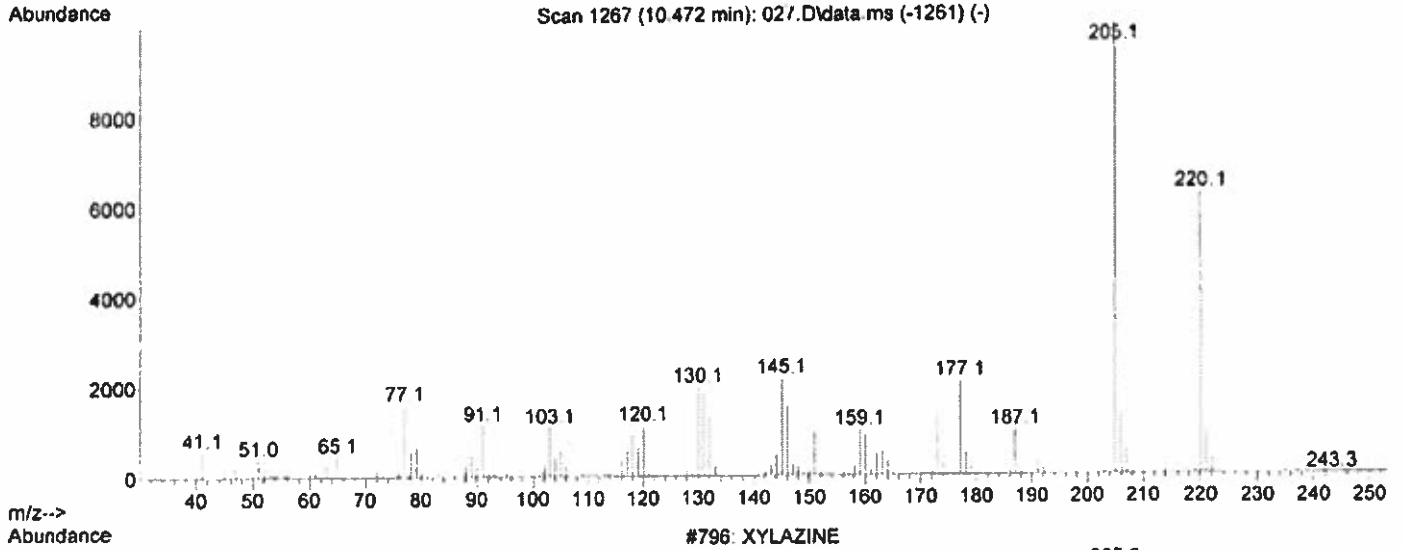
File :C:\Users\TOX\Desktop\092821\027.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 01:43 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 6 W/ SW CTL + XYLAZINE - URN  
Misc Info : 2.0 mg/L - 400uL of 0.01 mg/mL CTL, plus 400uL 0.1 mg/mL XYLAZINE  
Vial Number: 55



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 95  
ID : TRAMADOL  
Sample Name: MATRIX 6 W/ SW CTL + XYLAZINE - URN  
File: C:\Users\TOX\Desktop\092821\027.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 01:43  
Vial: 55



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 99  
ID : XYLAZINE  
Sample Name: MATRIX 6 W/ SW CTL + XYLAZINE - URN  
File: C:\Users\TOX\Desktop\092821\027.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 01:43  
Vial: 55



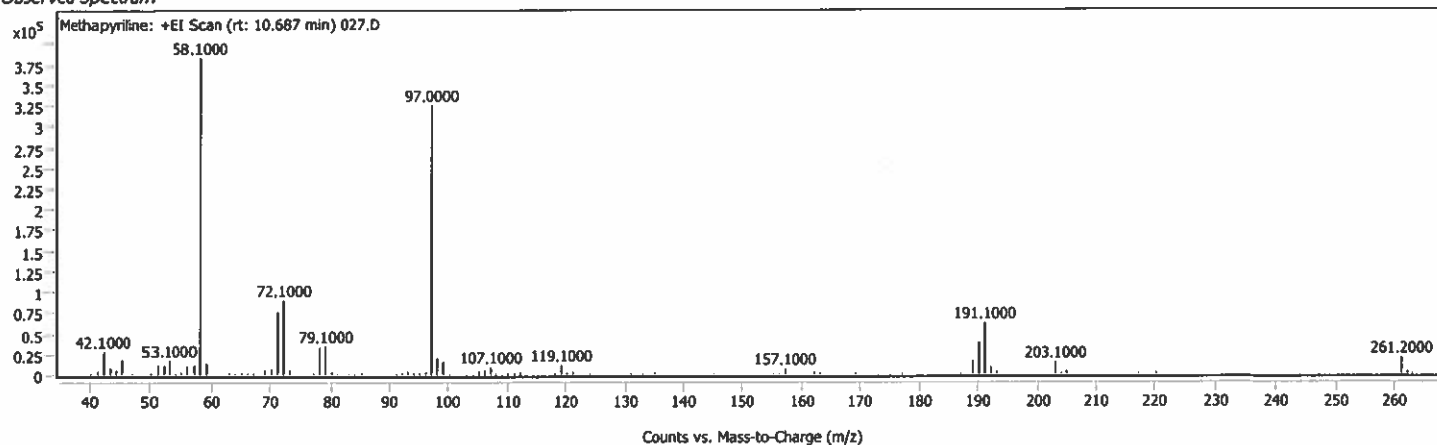
# Analysis Report

+ Scan (rt: 10.687 min)

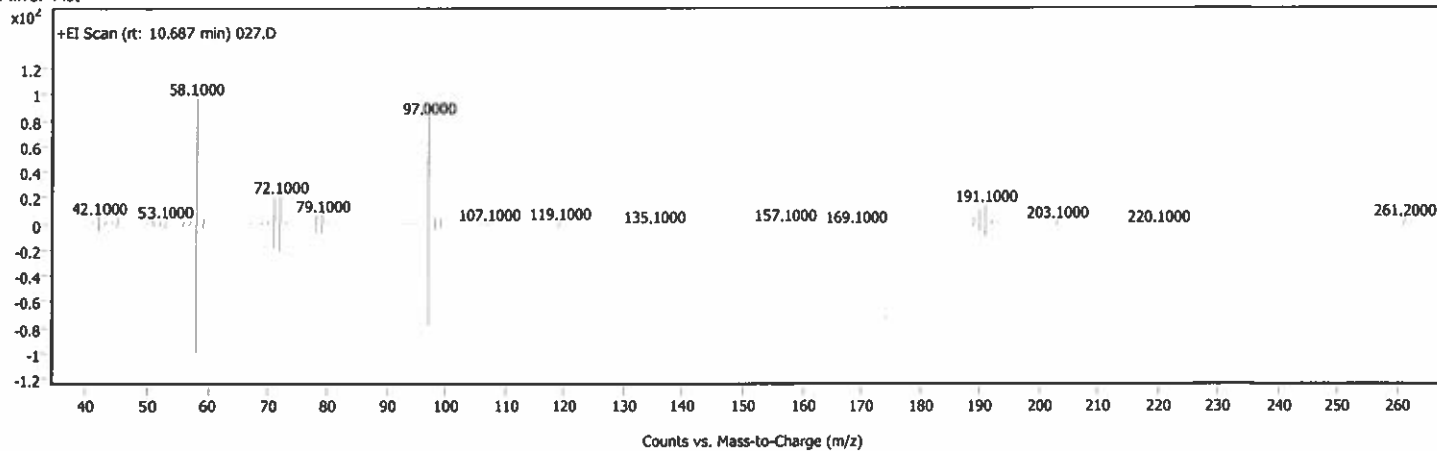
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	97.83	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

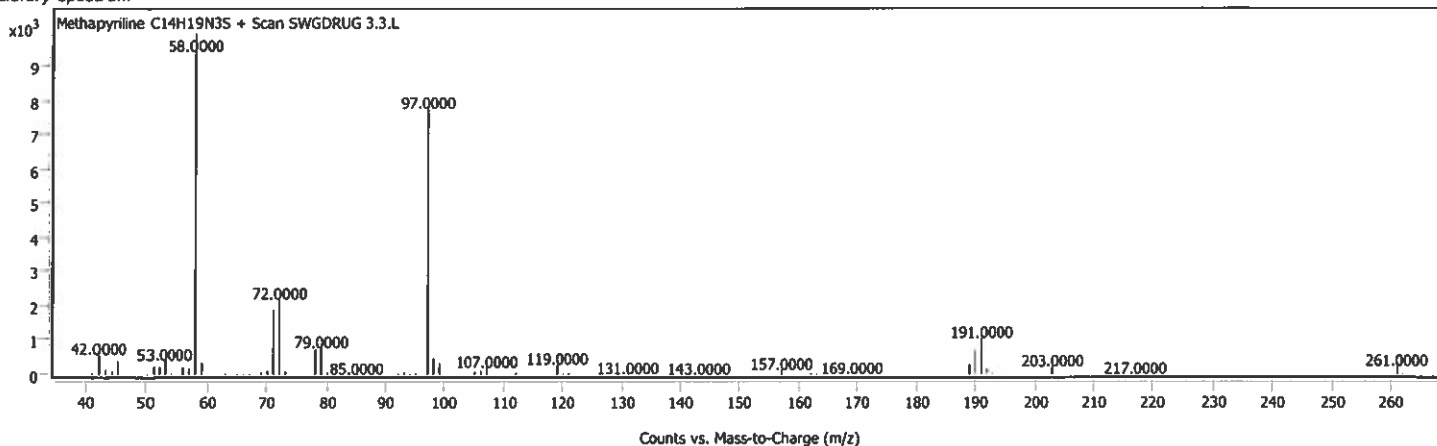
## Observed Spectrum



## Mirror Plot



## Library Spectrum



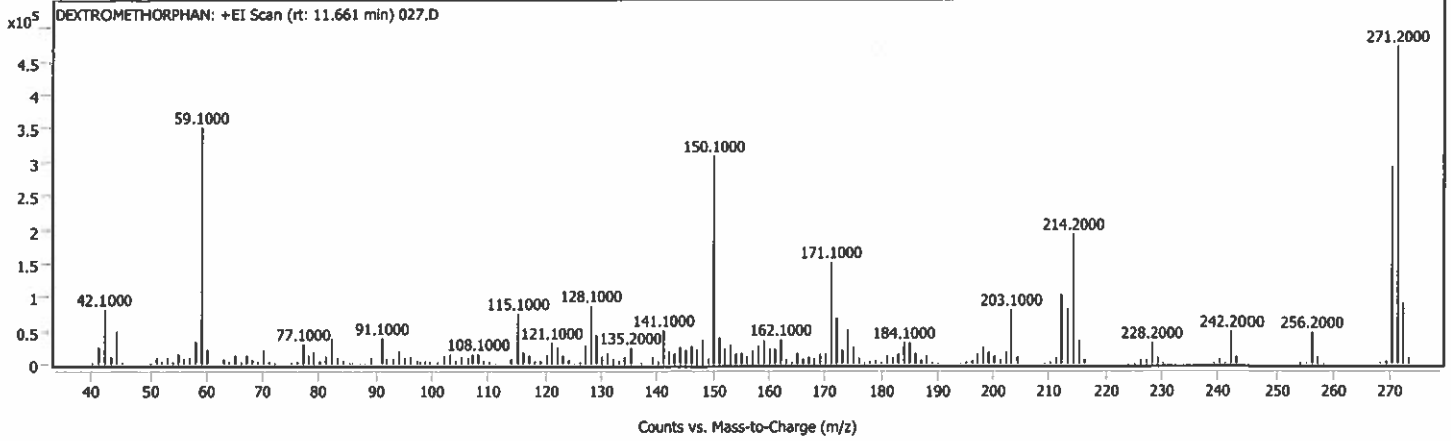
# Analysis Report

+ Scan (rt: 11.661 min)

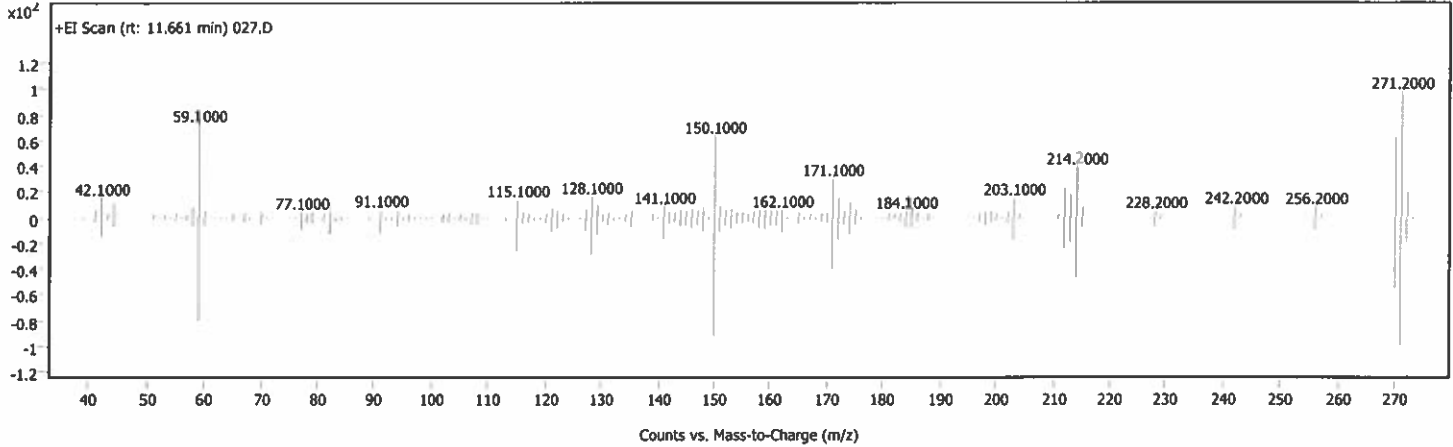
DEXTROMETHORPHAN; C18H25NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	92.20	aafs.I	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

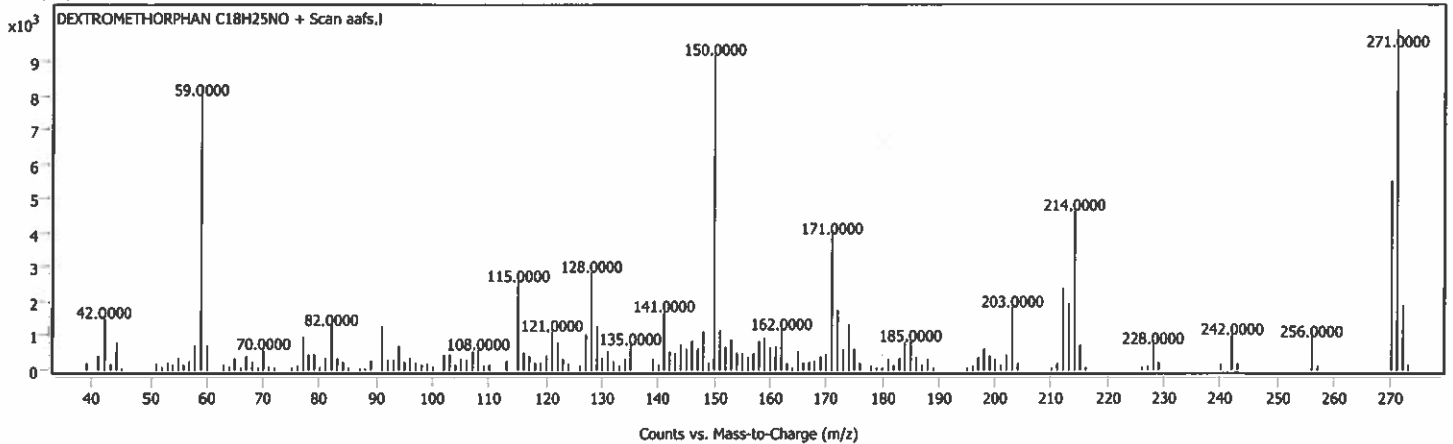
## Observed Spectrum



## Mirror Plot



## Library Spectrum



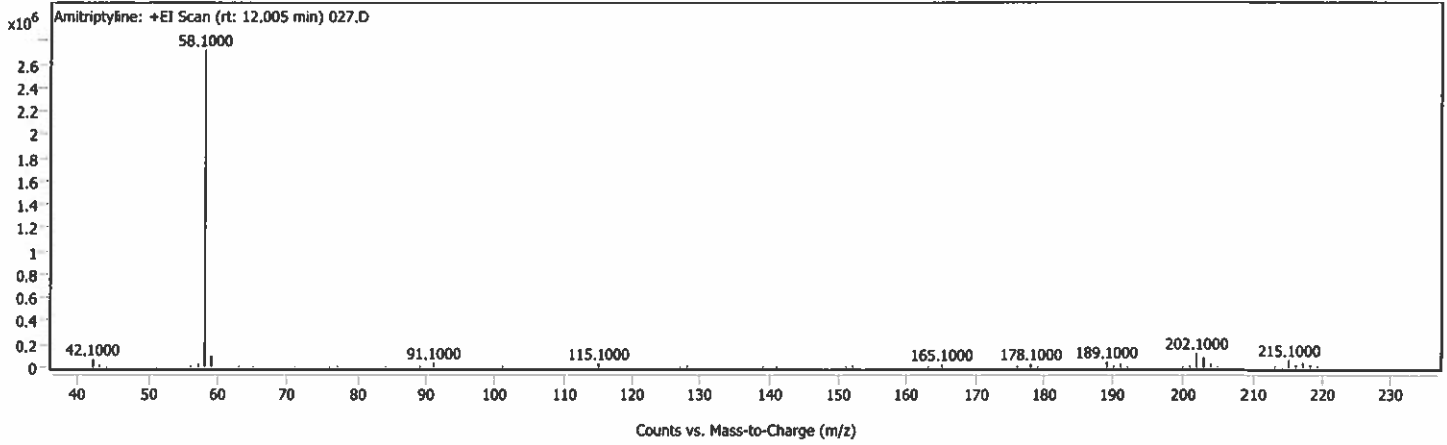
# Analysis Report

+ Scan (rt: 12.005 min)

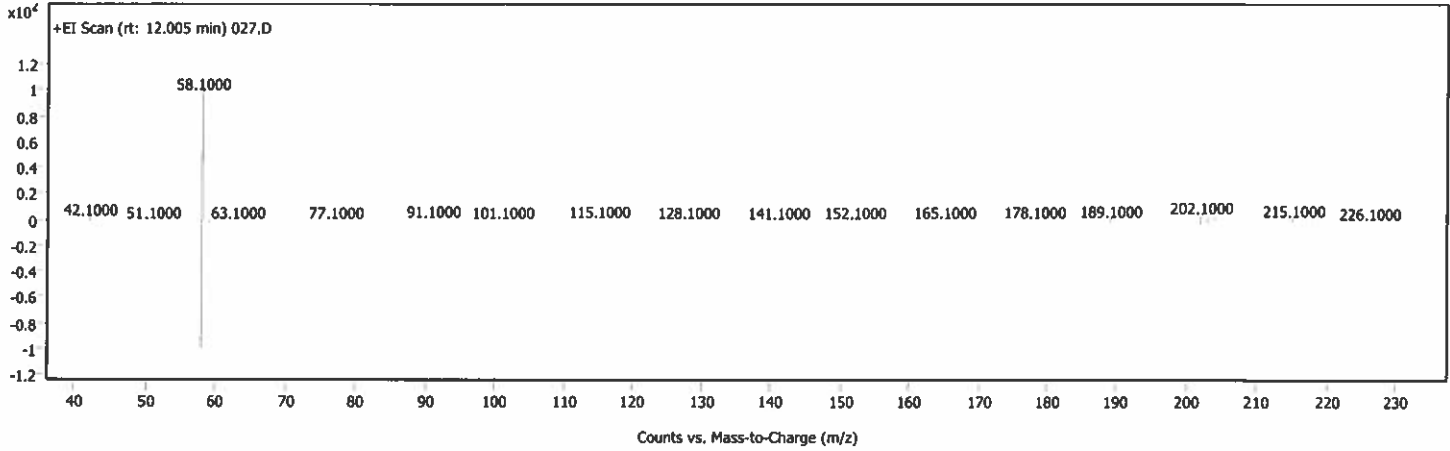
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	97.86	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

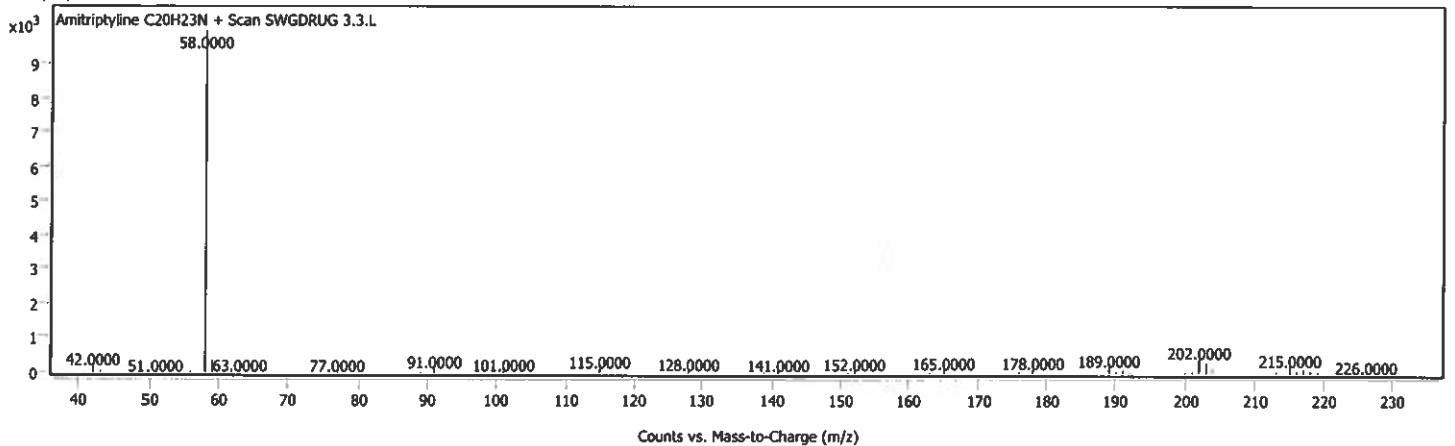
## Observed Spectrum



## Mirror Plot



## Library Spectrum



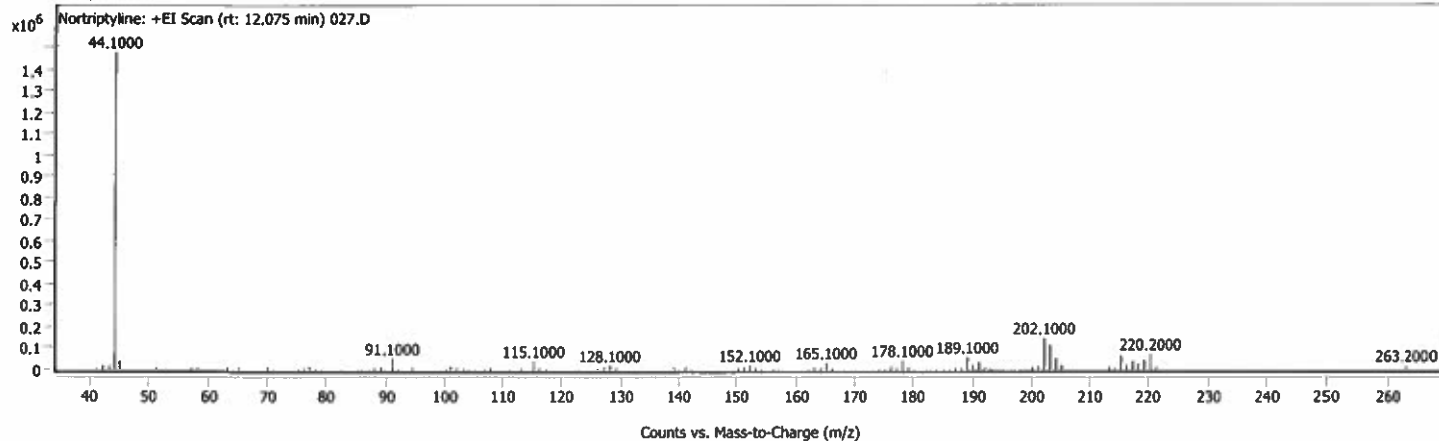
# Analysis Report

+ Scan (rt: 12.075 min)

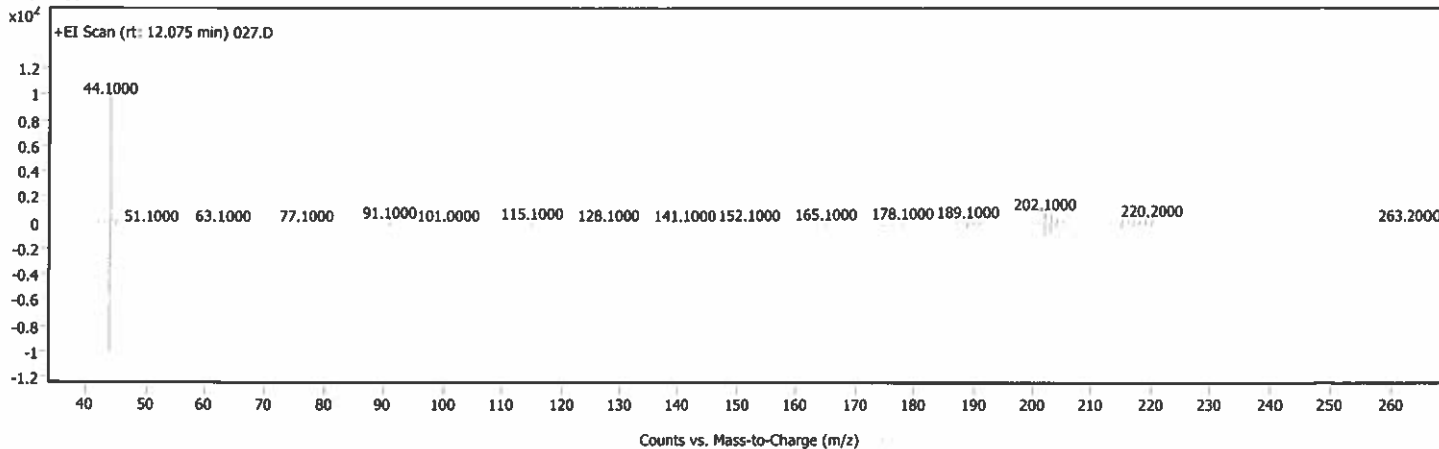
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	97.80	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

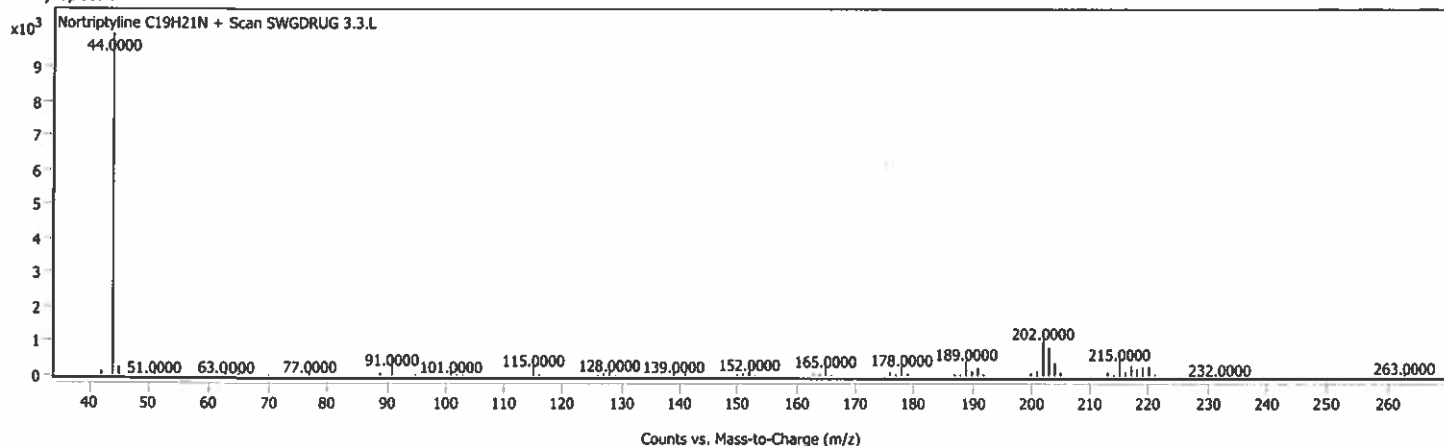
## Observed Spectrum



## Mirror Plot



## Library Spectrum



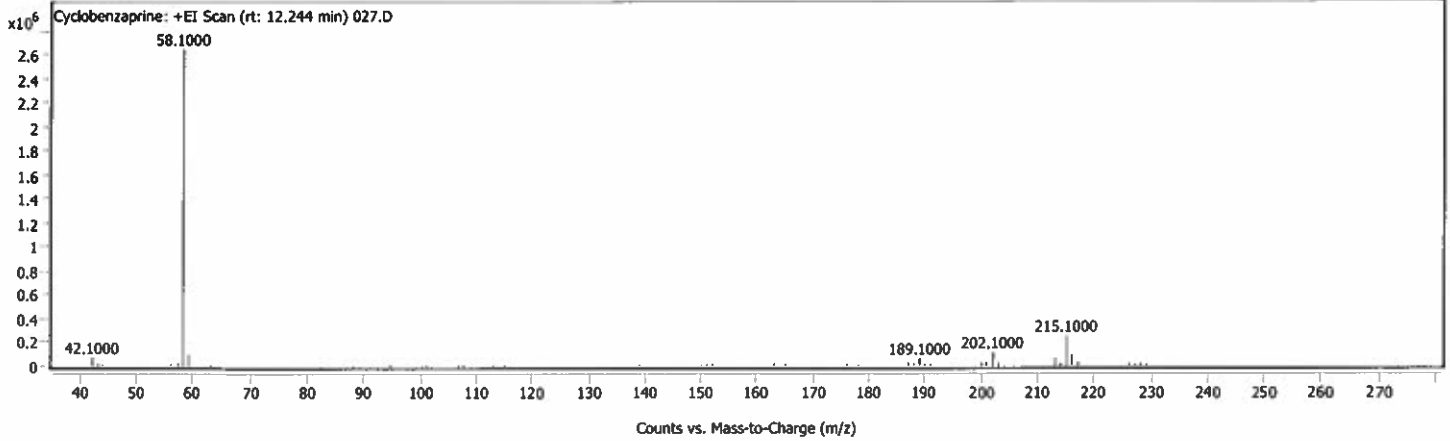
# Analysis Report

+ Scan (rt: 12.244 min)

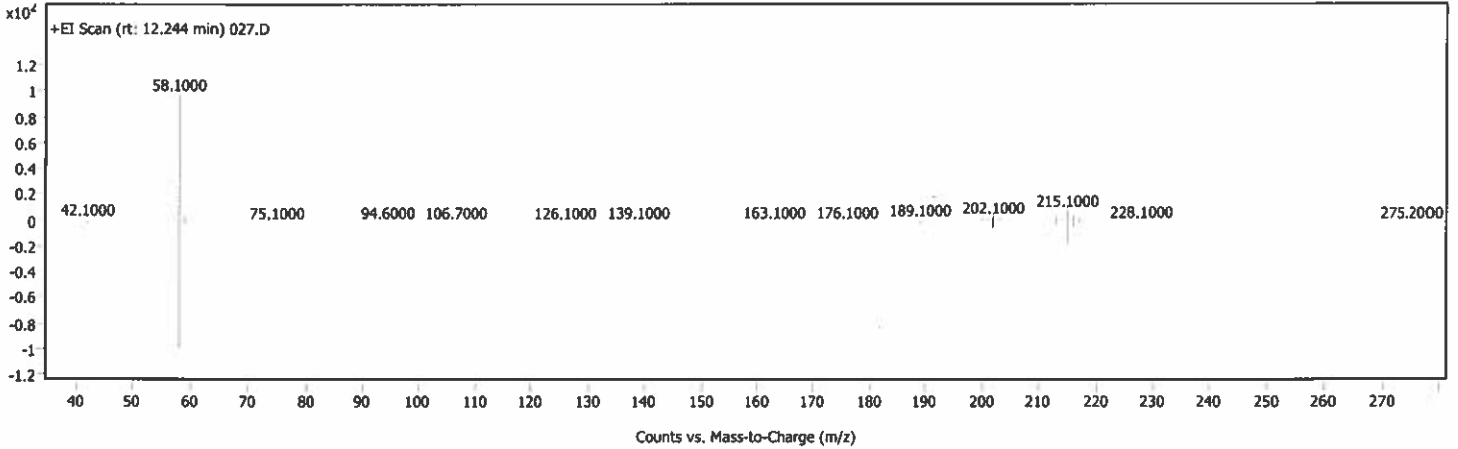
Cyclobenzaprine; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Cyclobenzaprine	C20H21N	96.63	SWGDRUG 3.3.L	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

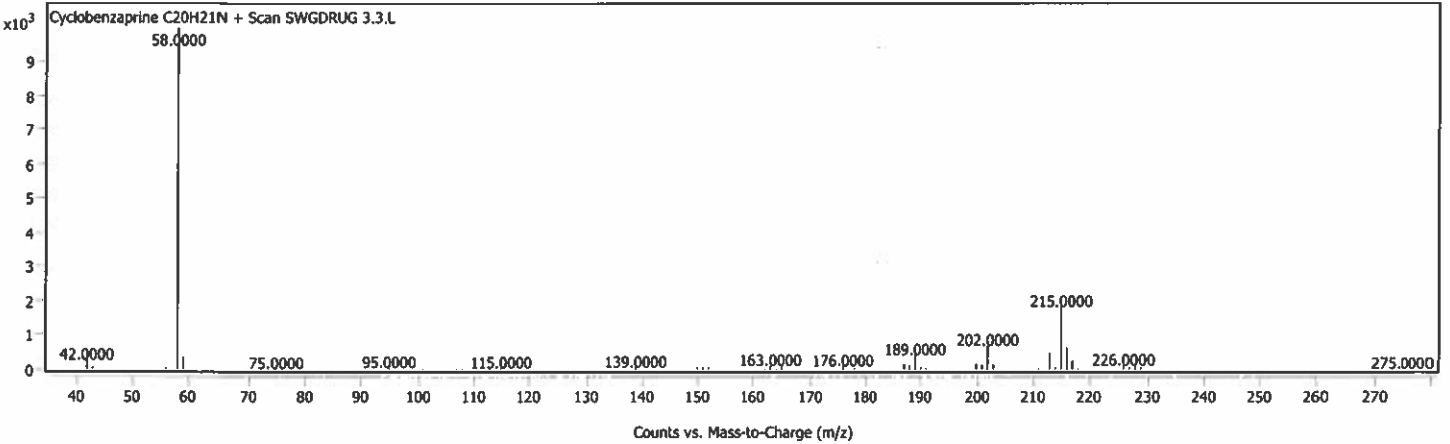
## Observed Spectrum



## Mirror Plot



## Library Spectrum





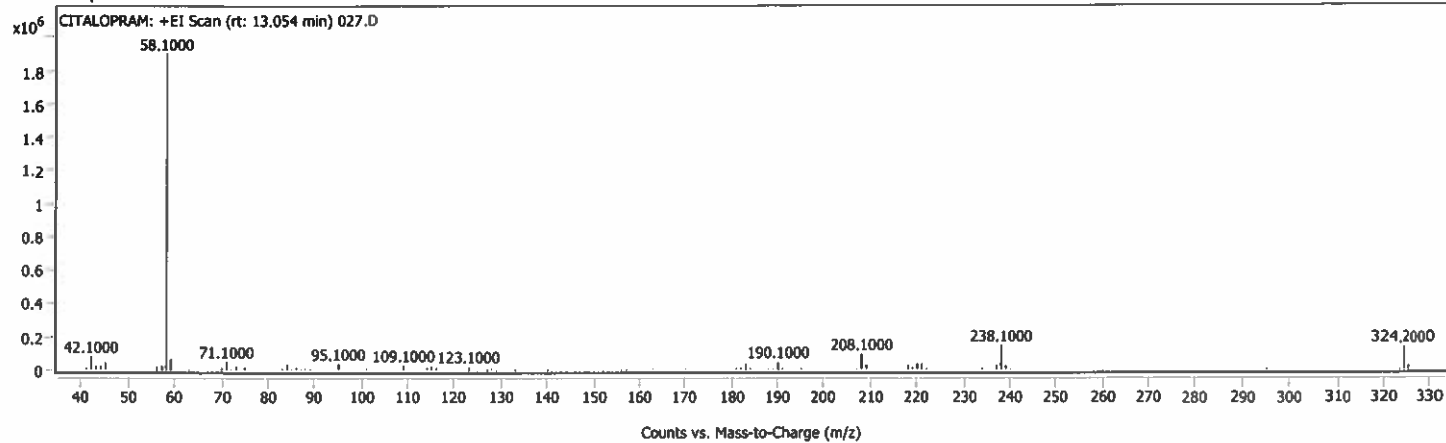
# Analysis Report

+ Scan (rt: 13.054 min)

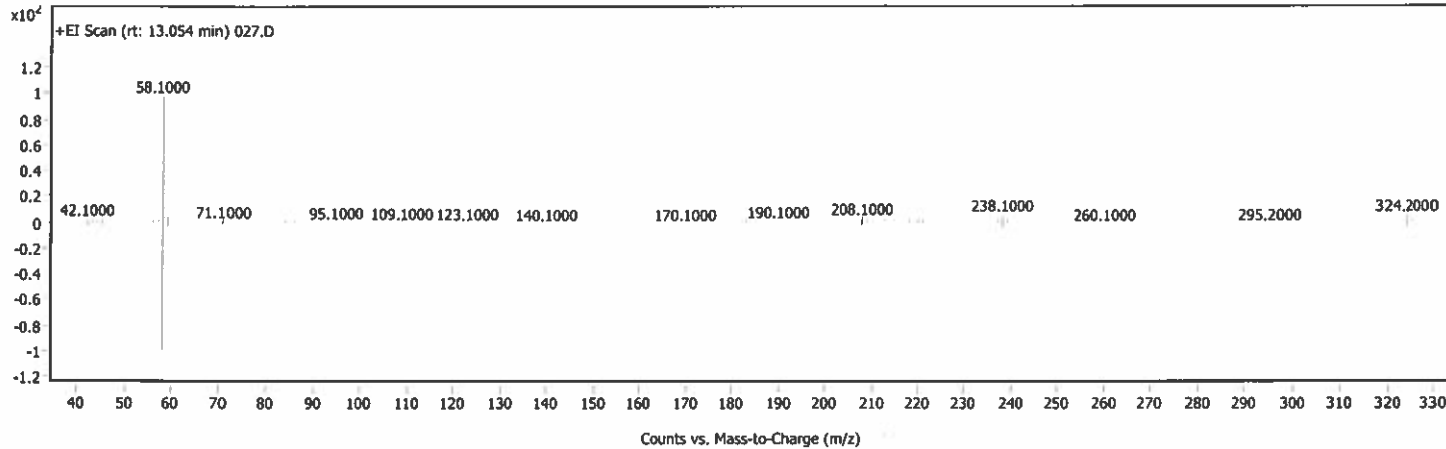
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	96.27	aafs.I	MATRIX 6 W/ SW CTL + XYLAZINE -URN	KM/QC DATA IN XYLAZINE VALIDATION

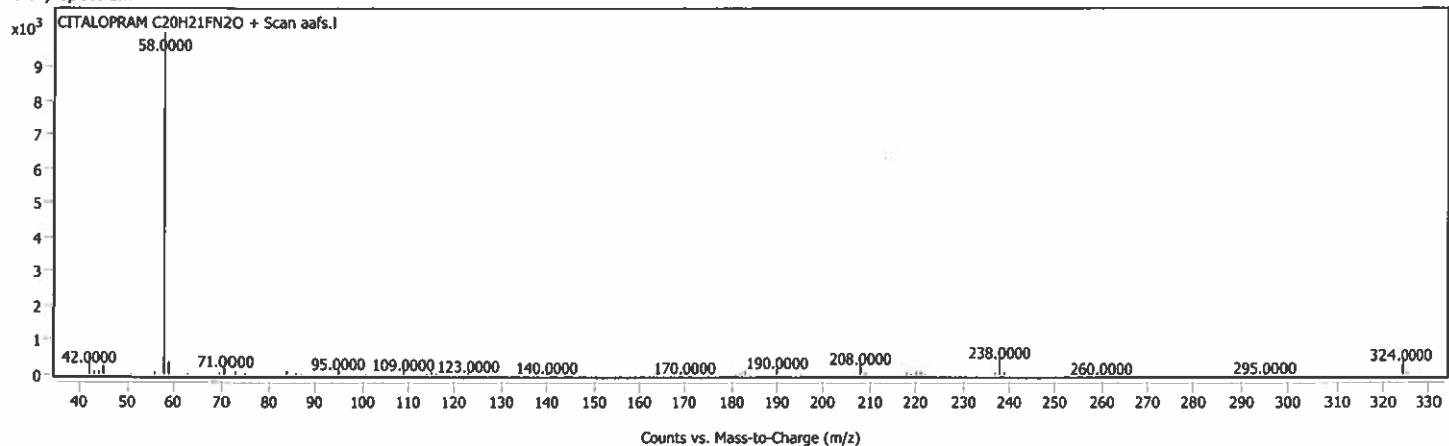
## Observed Spectrum



## Mirror Plot



## Library Spectrum



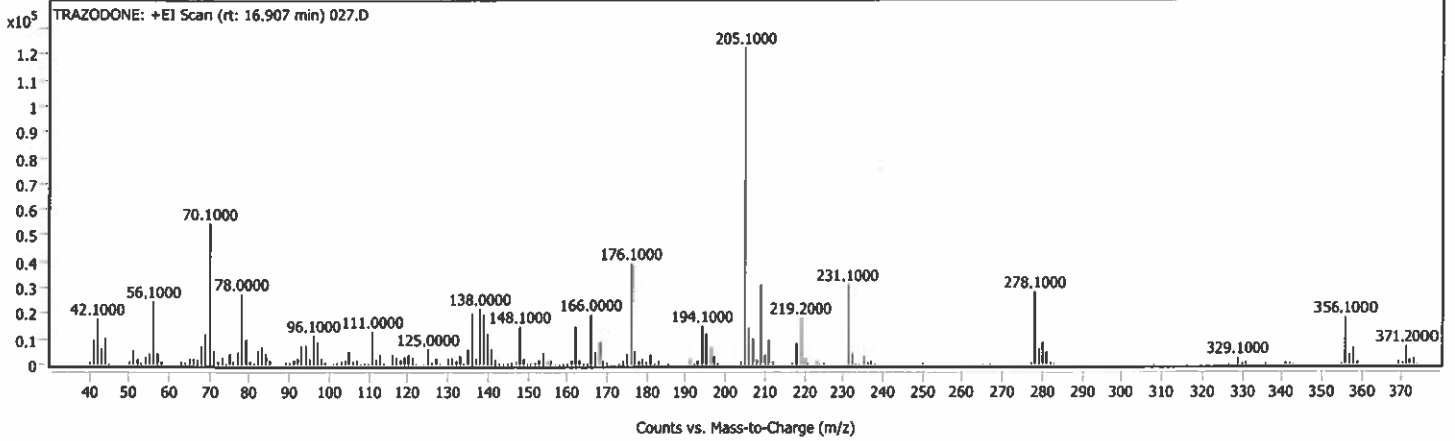
# Analysis Report

+ Scan (rt: 16.907 min)

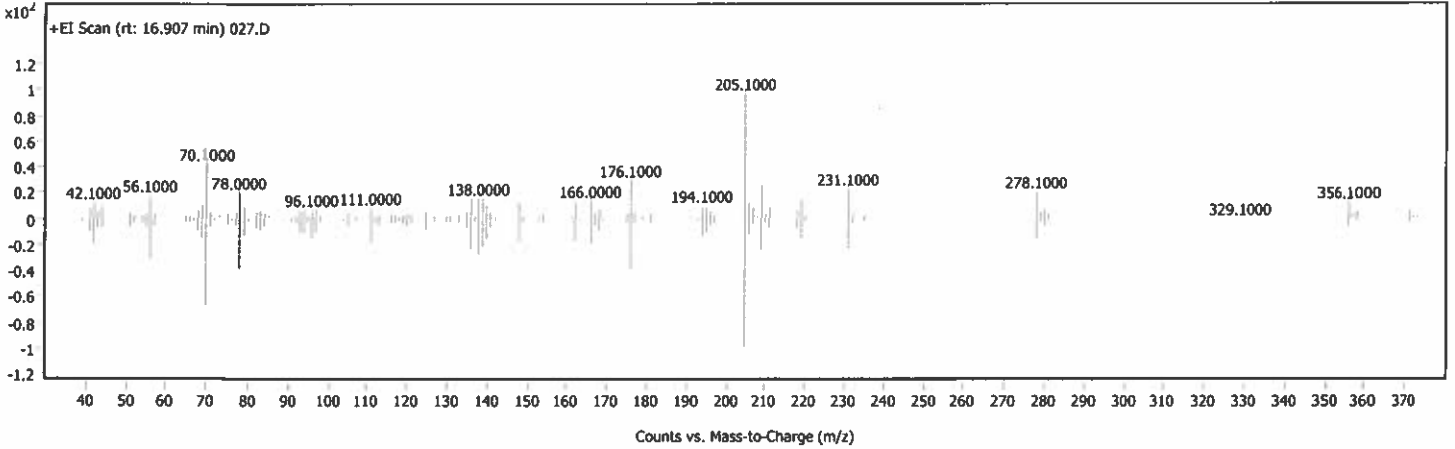
TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		93.56	aafs.1	MATRIX 6 W/ SW CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

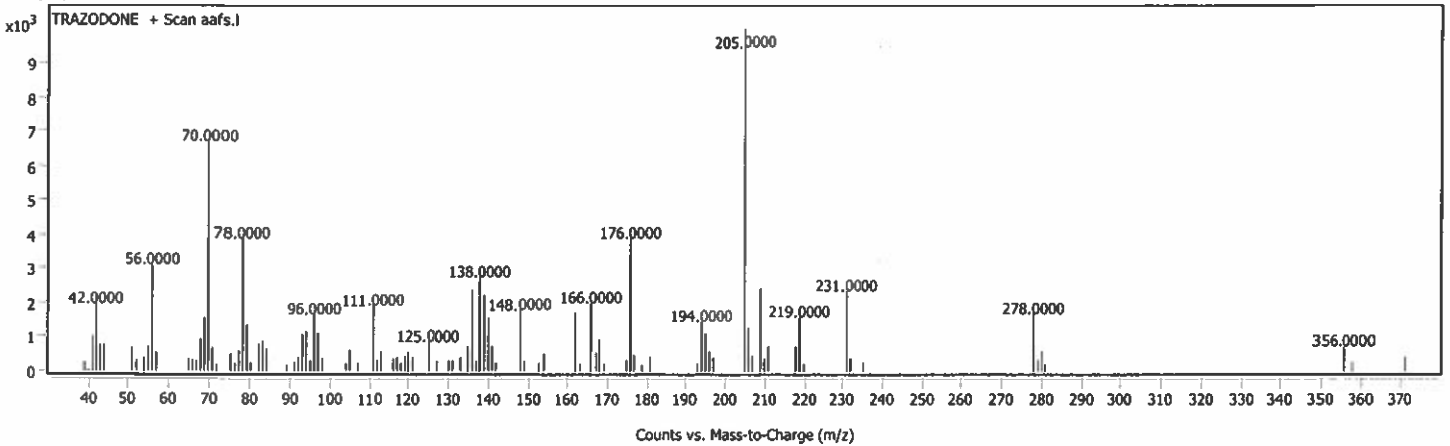
## Observed Spectrum



## Mirror Plot

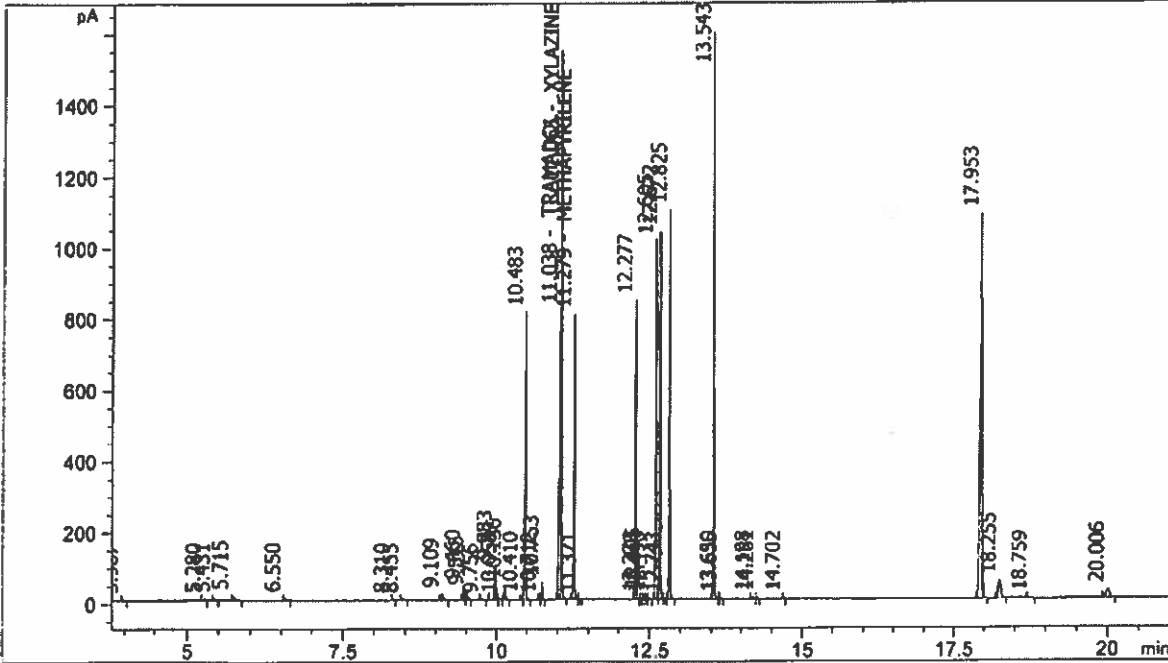


## Library Spectrum



Injection Date: 9/29/2021 1:30:38 AM Seq Line: 30  
 Sample Name: MATRIX 6 W/ SW CTL + -> Vial 31  
 Sample Info: *xyLAZINE* *KM 11/17/22*

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

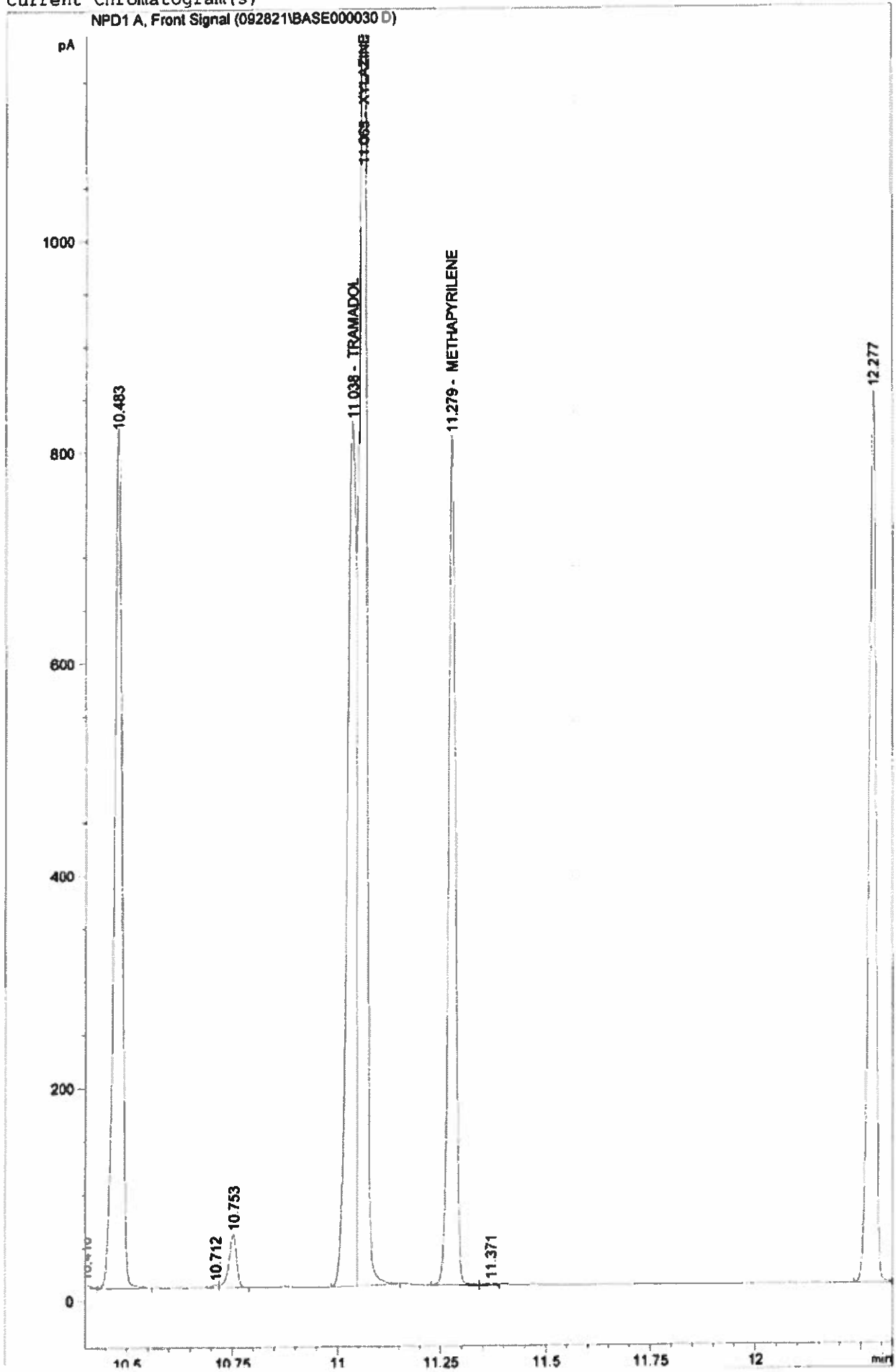


see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.038	11.033	1174.023	814.611	0.00000	TRAMADOL RRT - 0.9786
11.065	11.046	1814.521	1531.839	0.00000	XYLAZINE RRT - 0.9810
11.279	11.278	916.308	800.216	1.00000	METHAPYRILENE

Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000030.D)

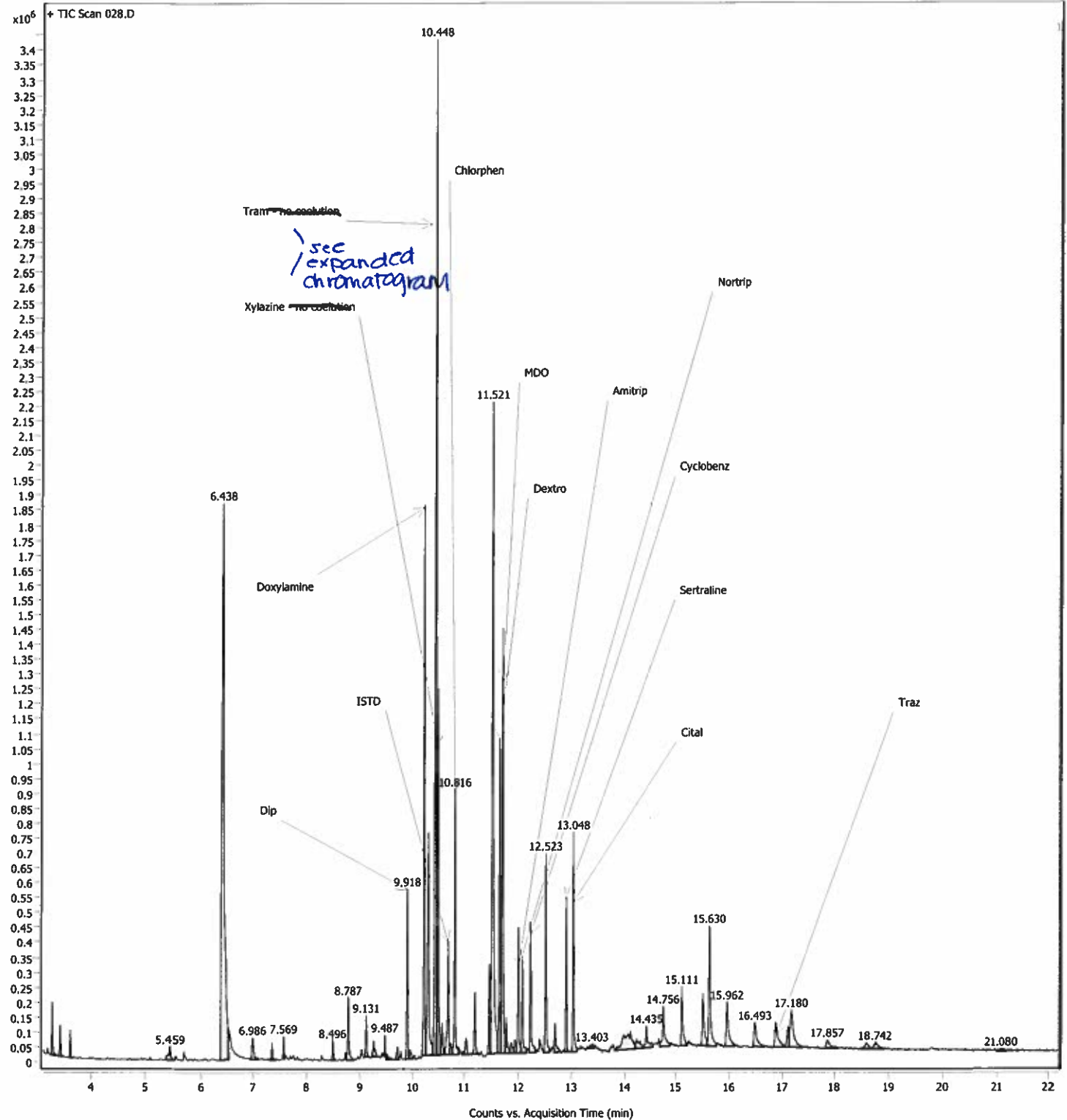


# Analysis Report

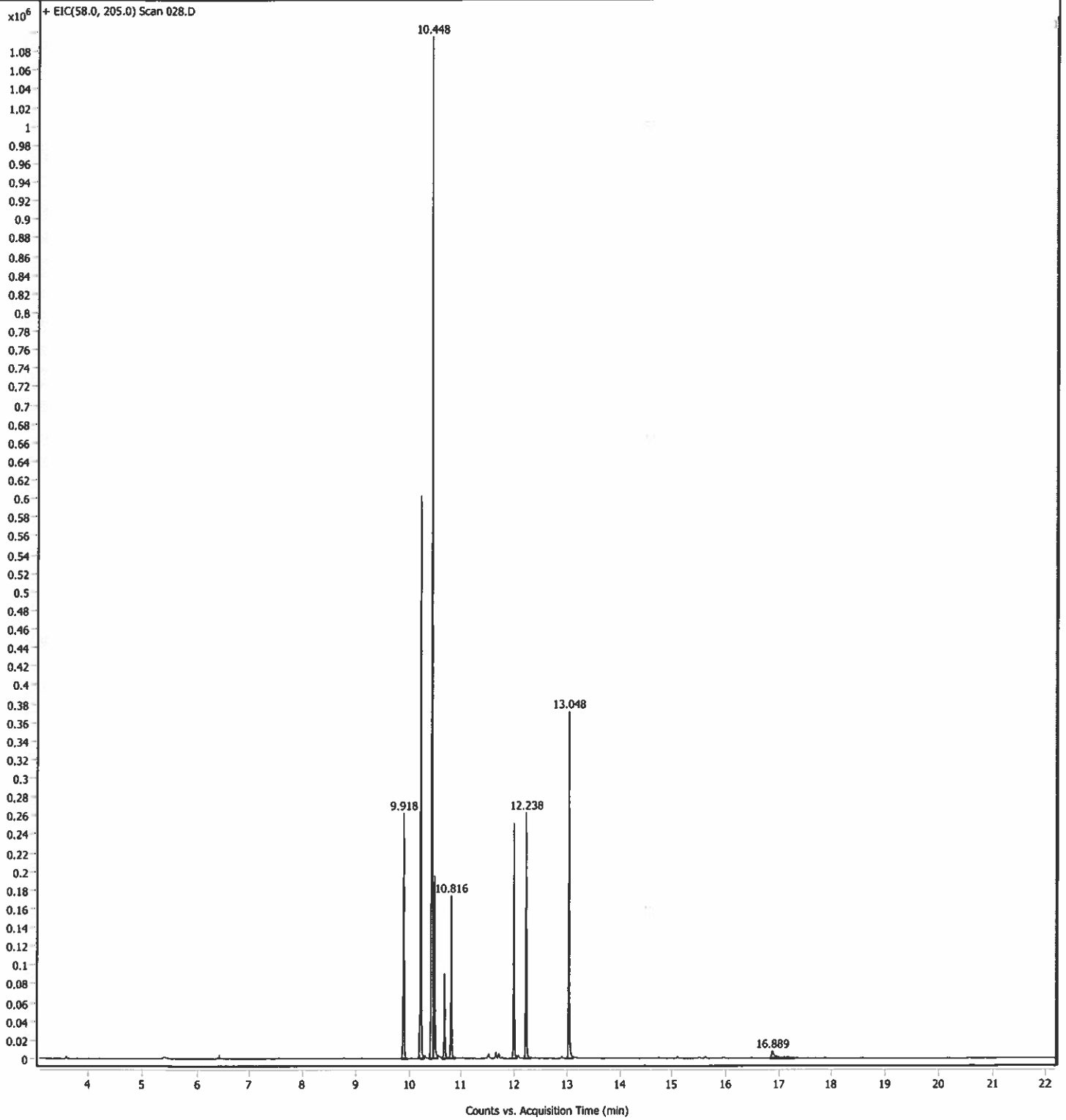
## Sample Information

Sample Name	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\028.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 2:09:52 AM (UTC-04:00)
Position	56	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report



# Analysis Report

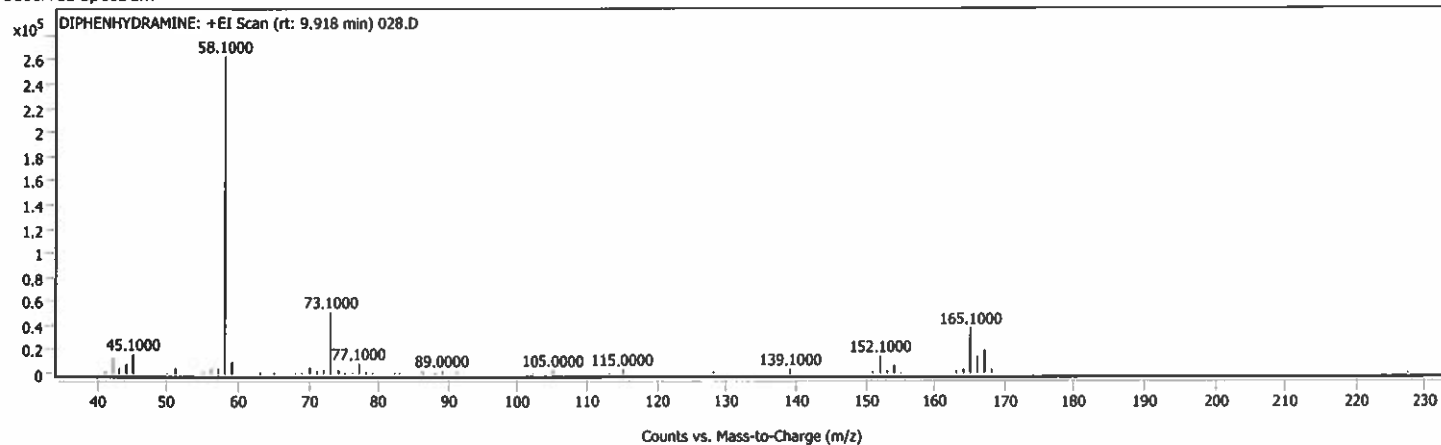
## Sample Spectra

+ Scan (rt: 9.918 min)

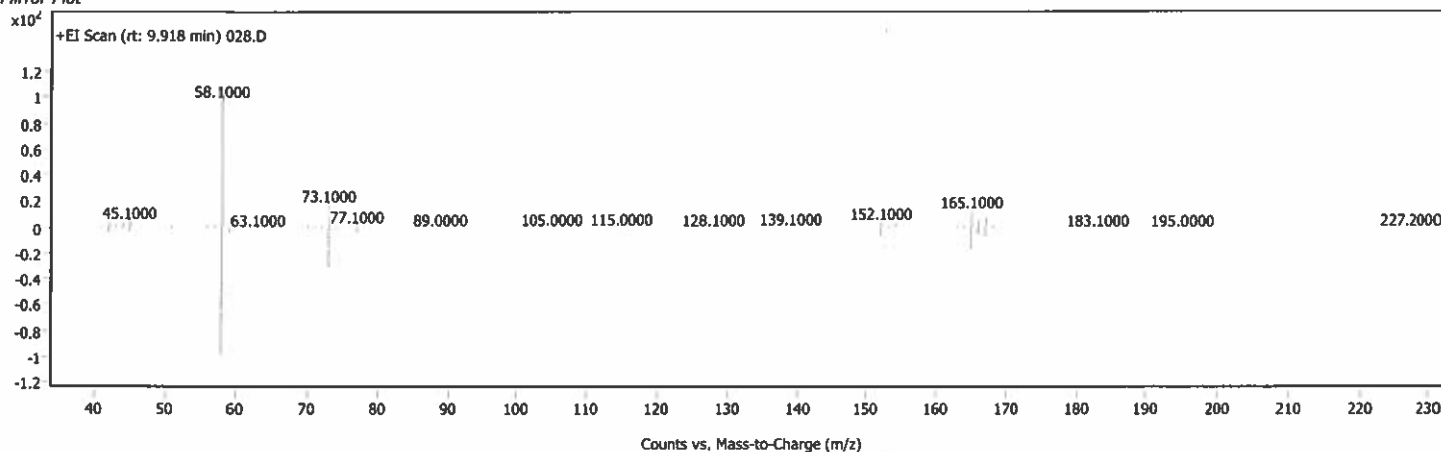
DIPHENHYDRAMINE; C<sub>17</sub>H<sub>21</sub>NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DIPHENHYDRAMINE	C <sub>17</sub> H <sub>21</sub> NO	95.91	aafs.i	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

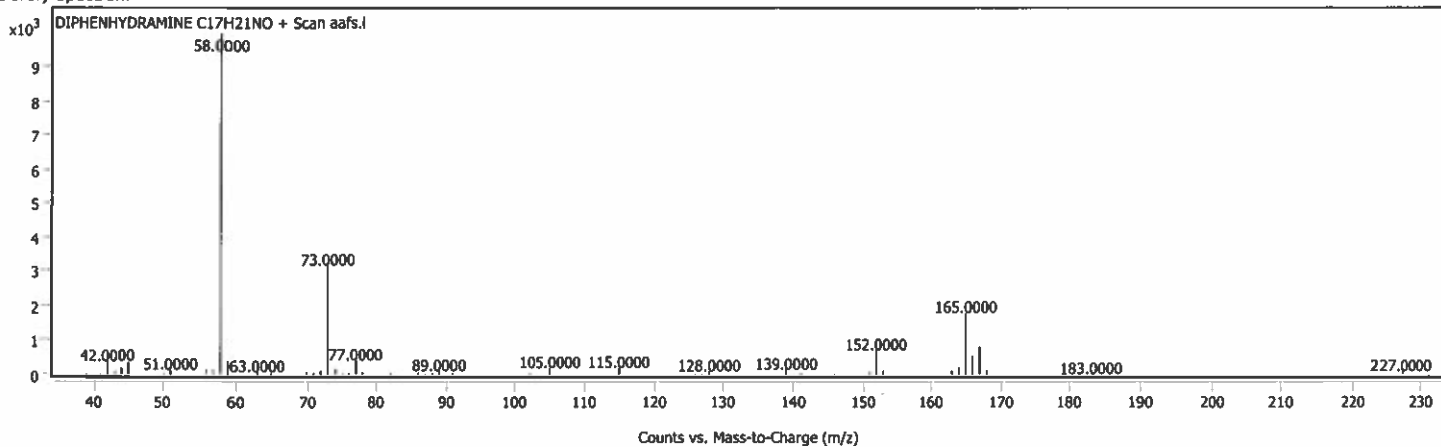
## Observed Spectrum



## Mirror Plot



## Library Spectrum



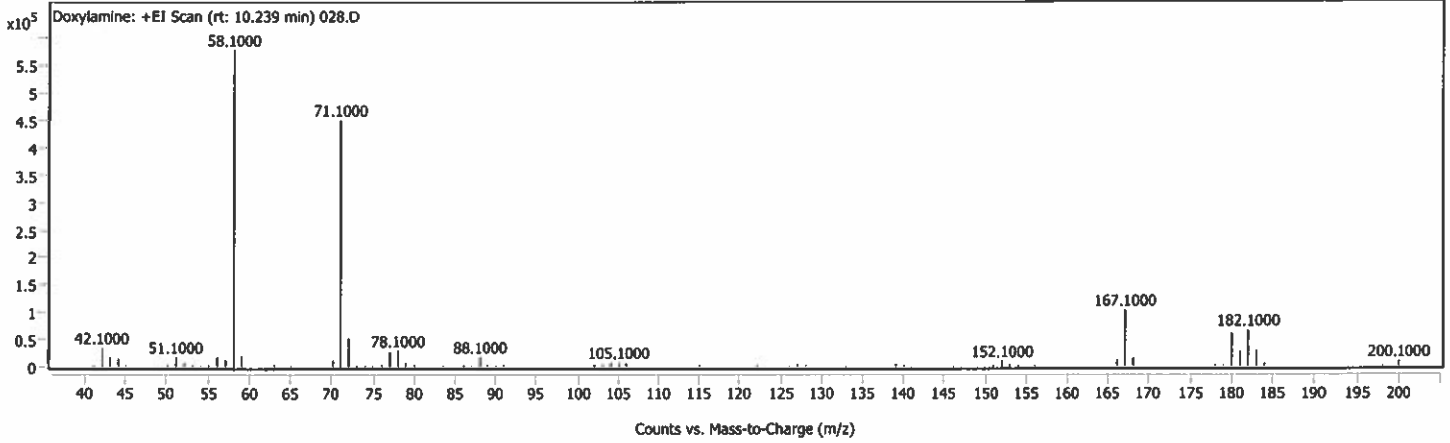
# Analysis Report

+ Scan (rt: 10.239 min)

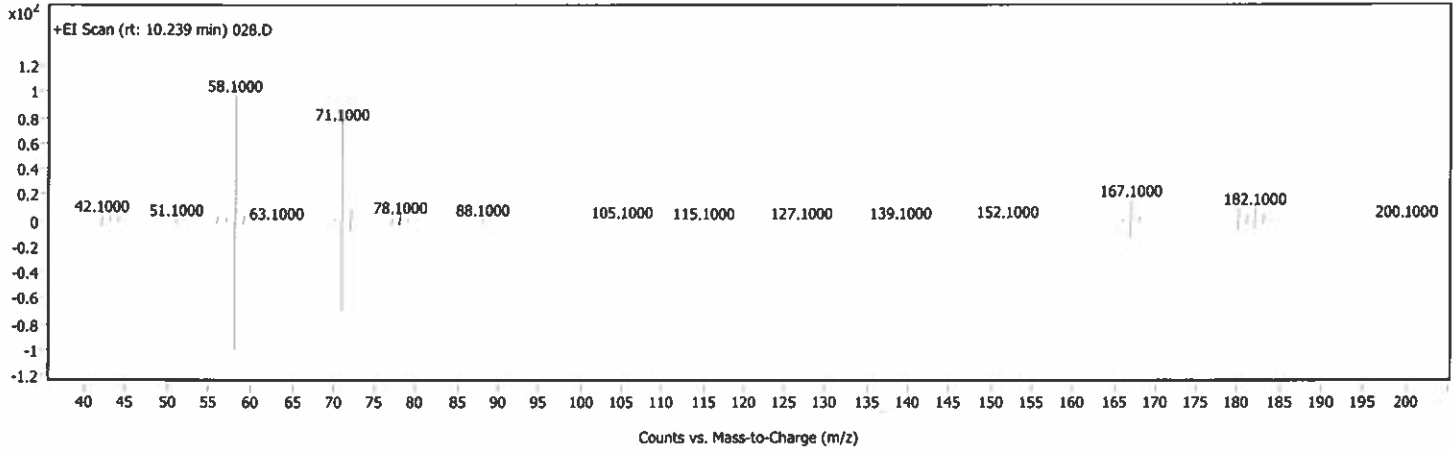
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.61	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

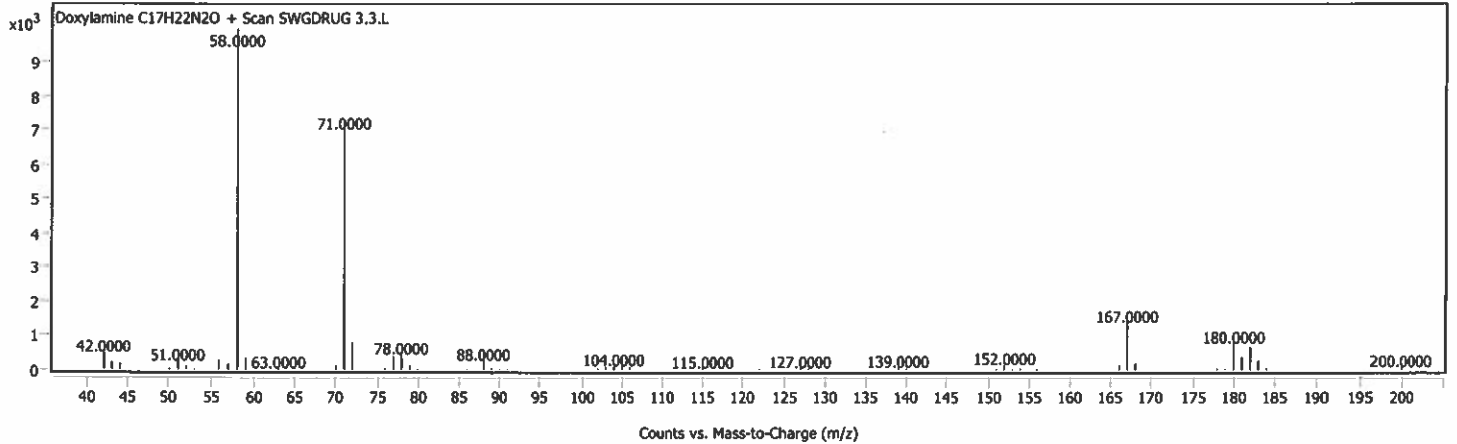
## Observed Spectrum



## Mirror Plot

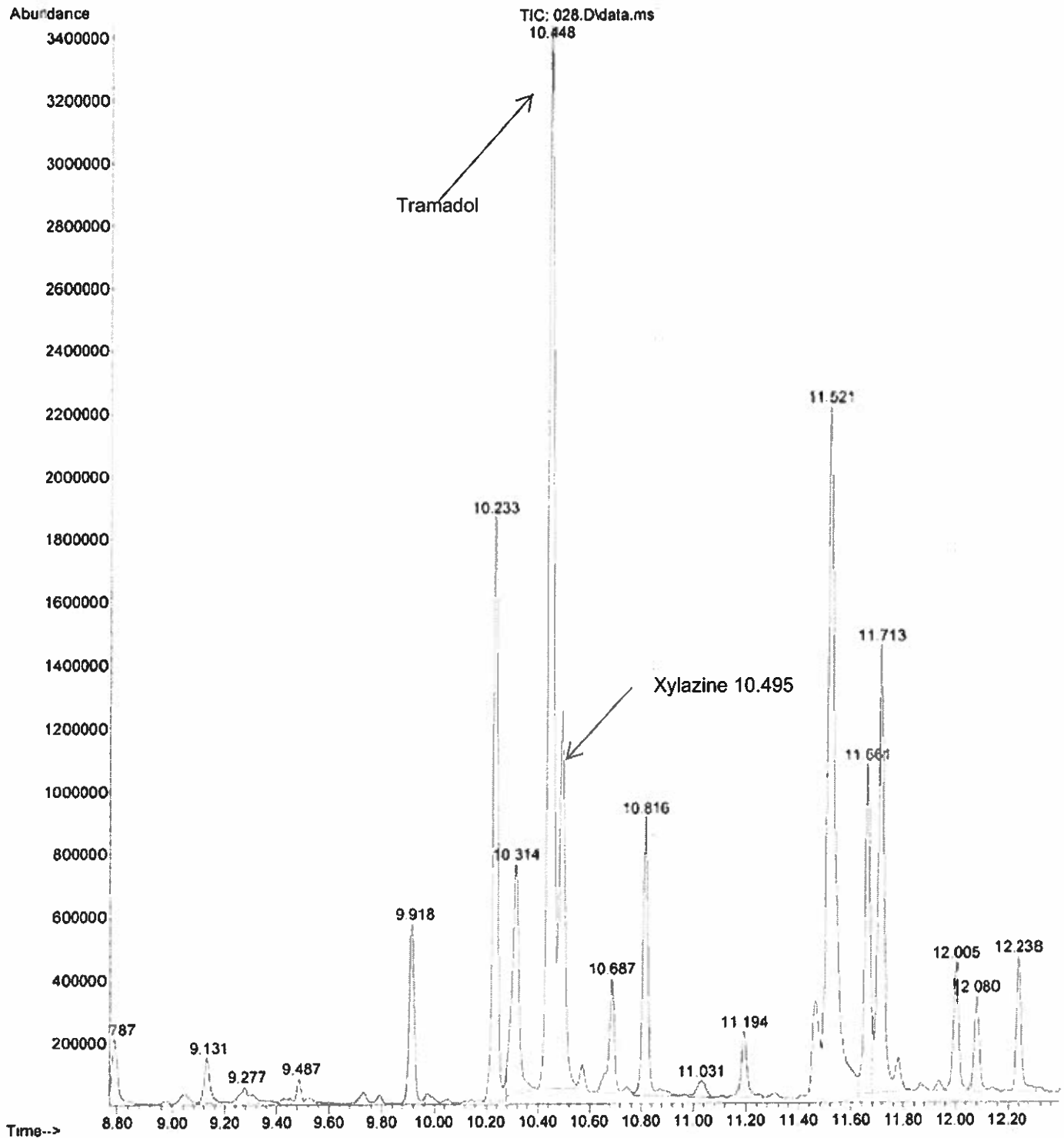


## Library Spectrum

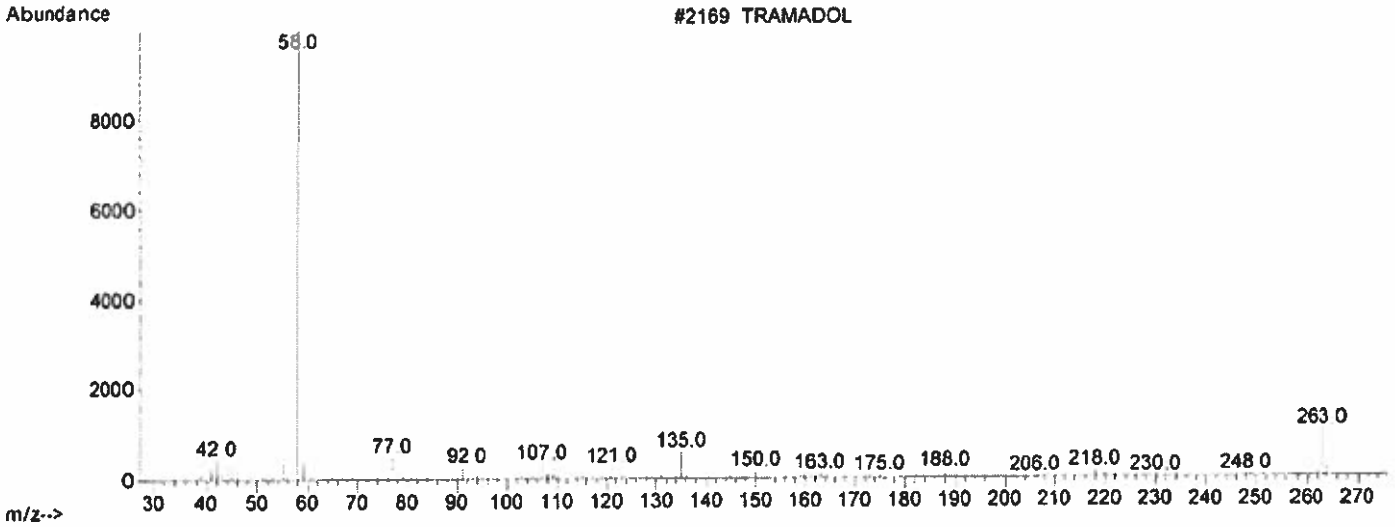
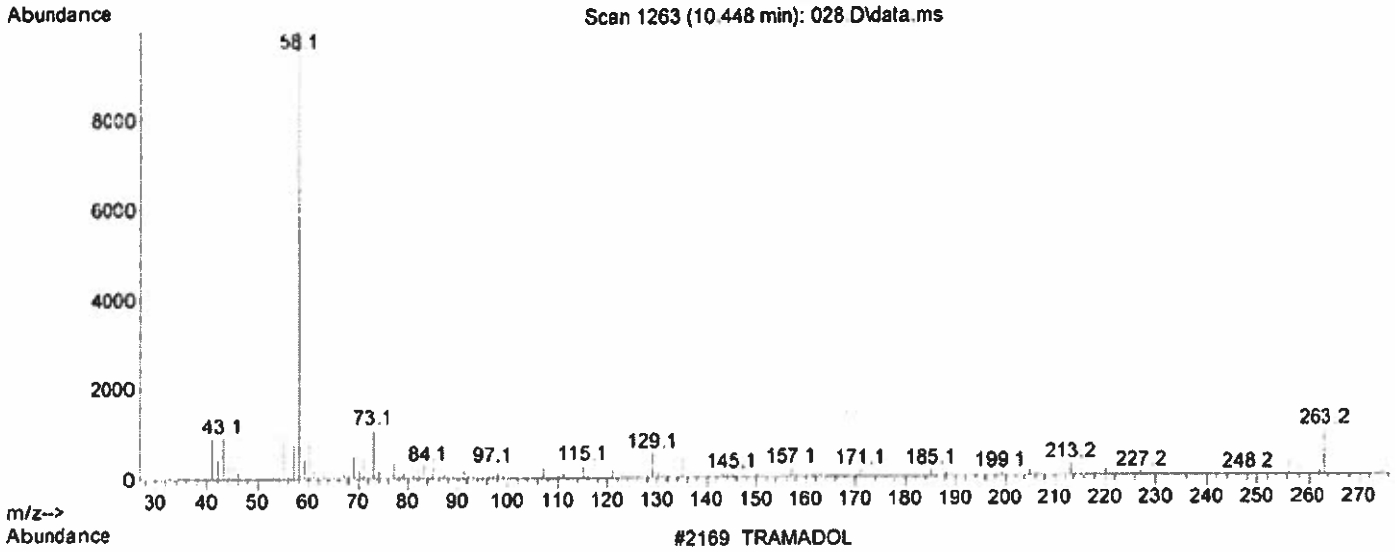




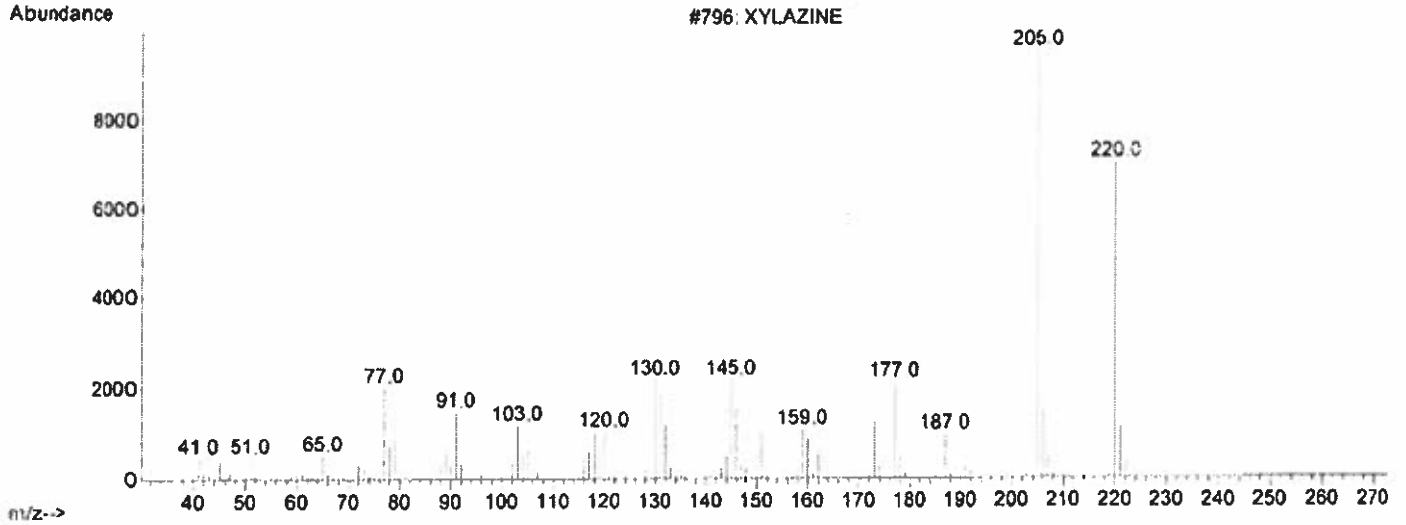
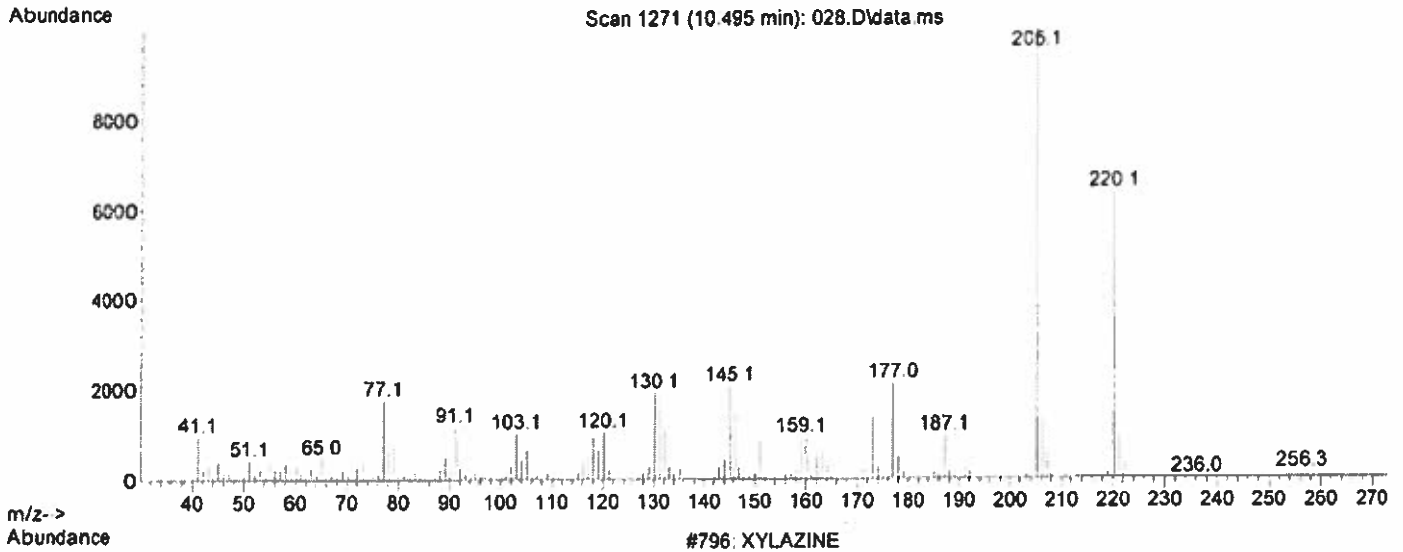
File :C:\Users\TOX\Desktop\092821\028.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 29 Sep 2021 02:09 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR  
Misc Info : 2.0 mg/L - 400uL of 0.01 mg/mL Cal, plus 400uL 0.1 mg/mL XYLAZINE  
Vial Number: 56



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 95  
ID : TRAMADOL  
Sample Name: MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR  
File: C:\Users\TOX\Desktop\092821\028.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 02:09  
Vial: 56



Library Searched : C:\MassHunter\Library\aaafs.1  
Quality : 99  
ID : XYLAZINE  
Sample Name: MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR  
File: C:\Users\TOX\Desktop\092821\028.D  
Operator: KM/QC DATA IN XYLAZINE VALIDATION  
Date Acquired: 29 Sep 2021 02:09  
Vial: 56



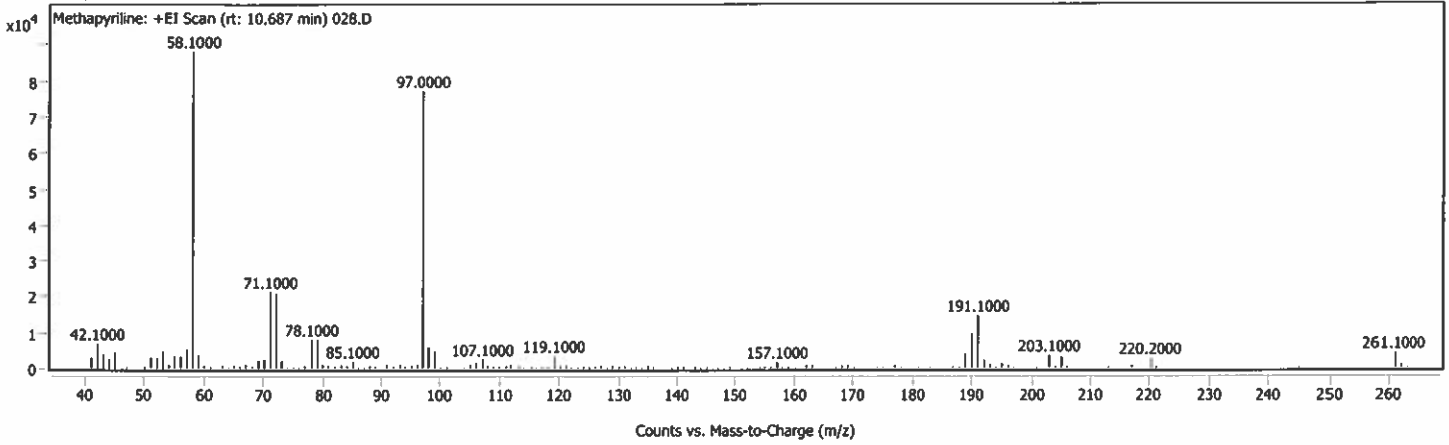
# Analysis Report

+ Scan (rt: 10.687 min)

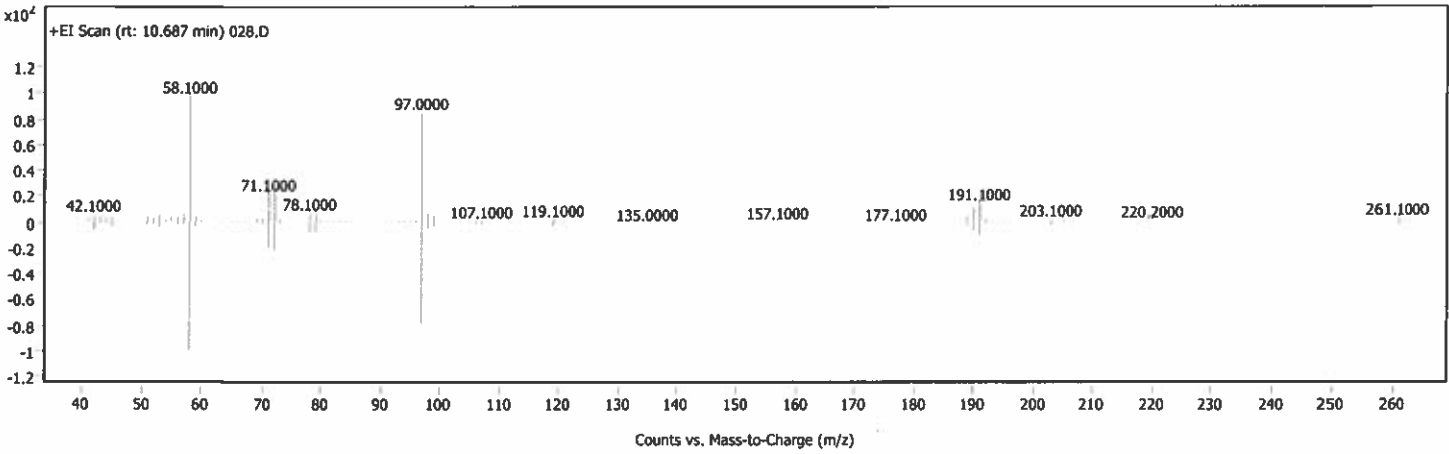
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	91.30	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

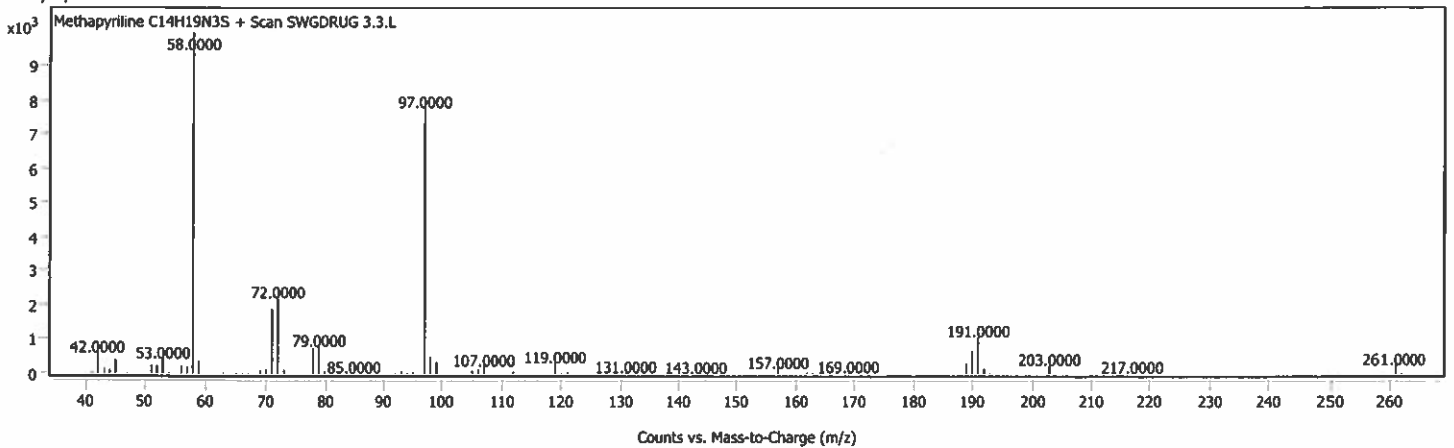
### Observed Spectrum



### Mirror Plot



### Library Spectrum



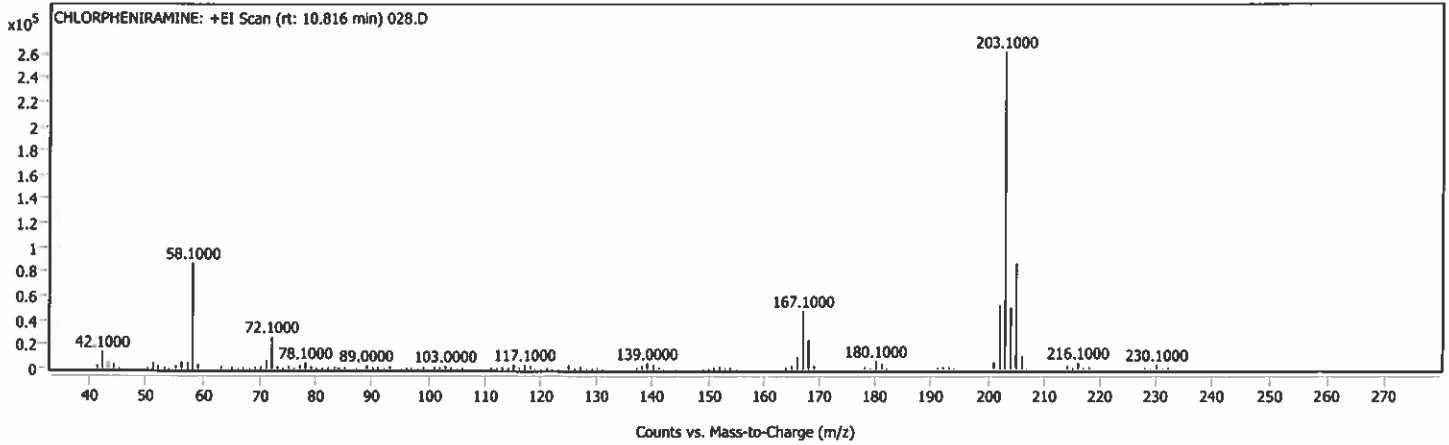
# Analysis Report

+ Scan (rt: 10.816 min)

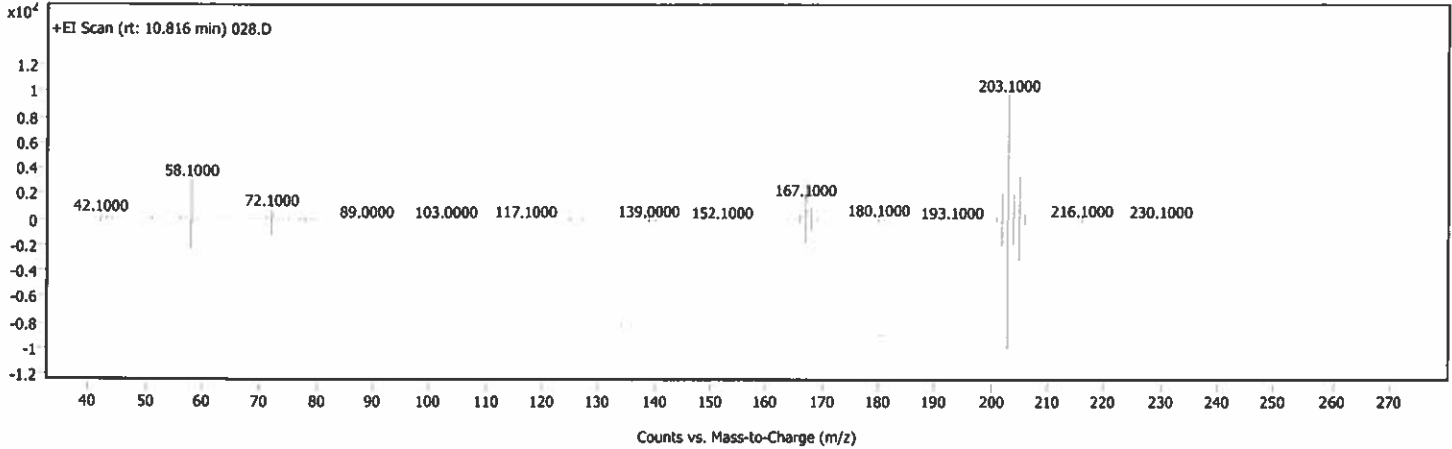
CHLORPHENIRAMINE; C16H19CIN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19CIN2	98.01	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

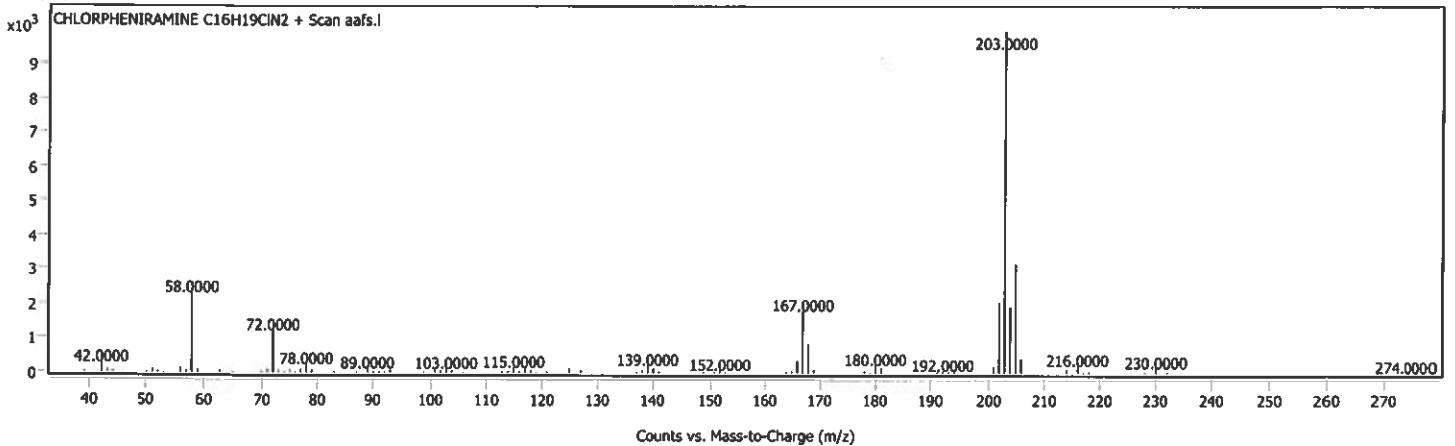
### Observed Spectrum



### Mirror Plot



### Library Spectrum



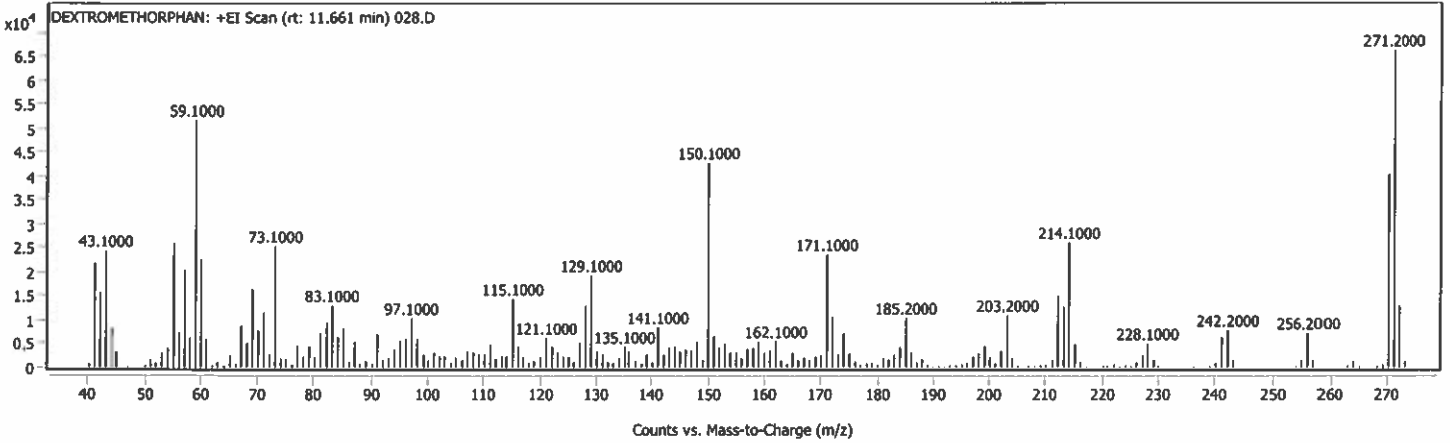
# Analysis Report

+ Scan (rt: 11.661 min)

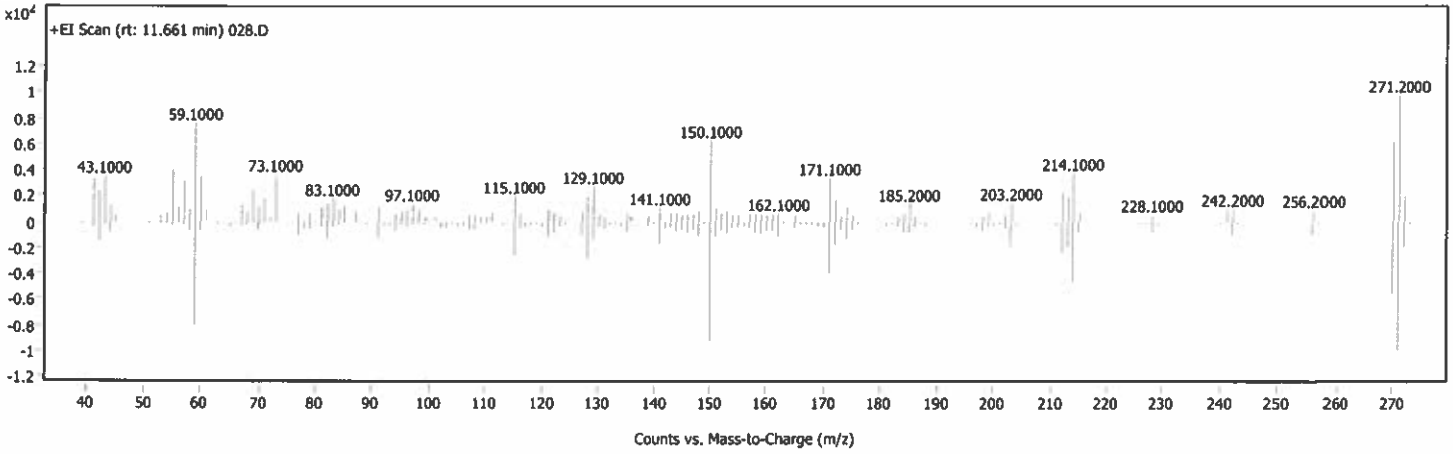
DEXTROMETHORPHAN; C18H25NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
DEXTROMETHORPHAN	C18H25NO	80.14	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

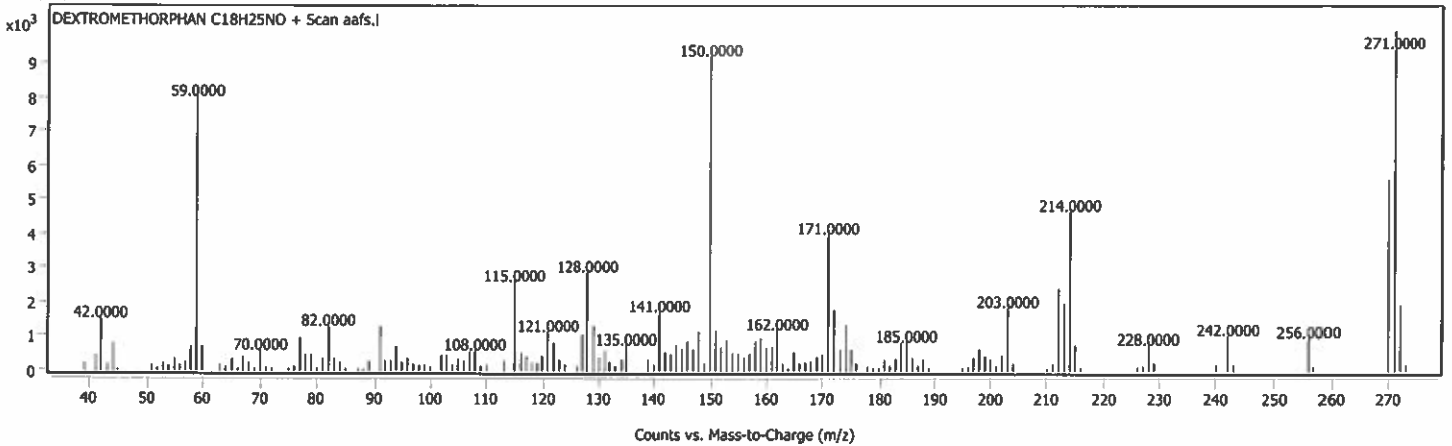
### Observed Spectrum



### Mirror Plot



### Library Spectrum



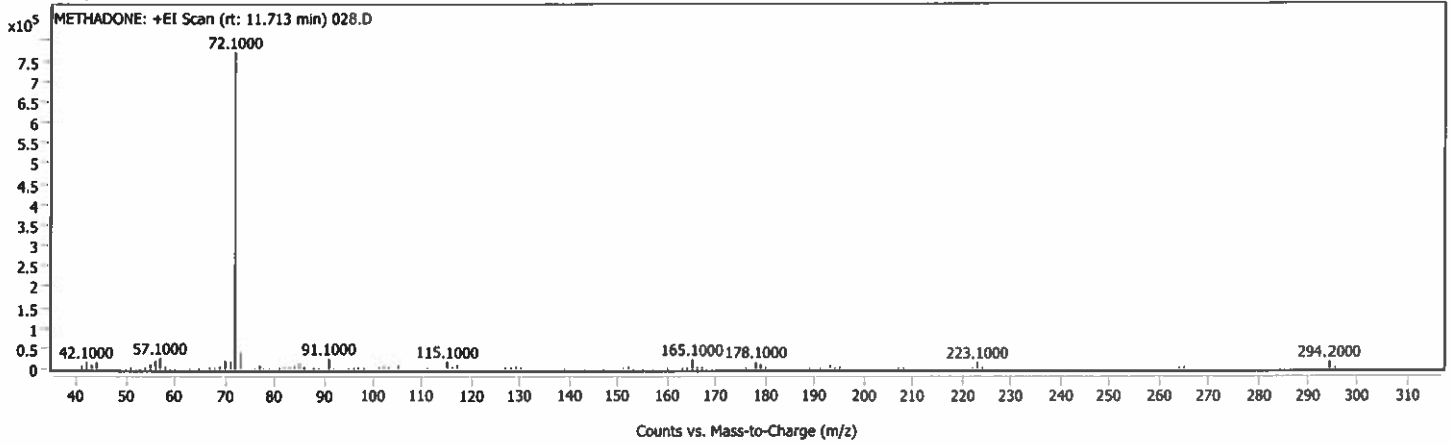
# Analysis Report

+ Scan (rt: 11.713 min)

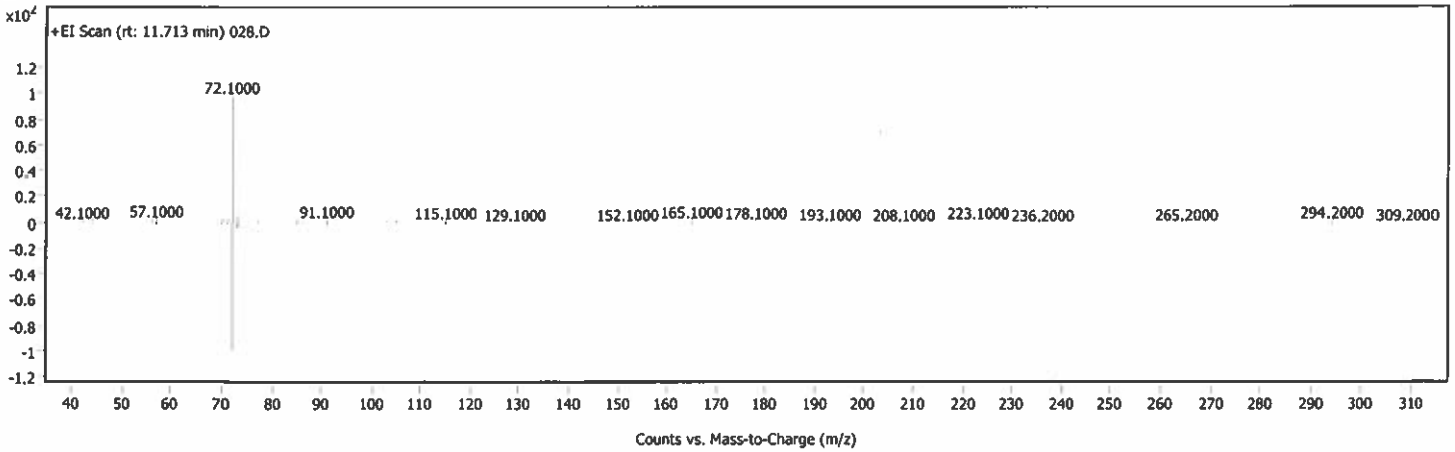
METHADONE; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
METHADONE	C21H27NO	93.19	aafs.l	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

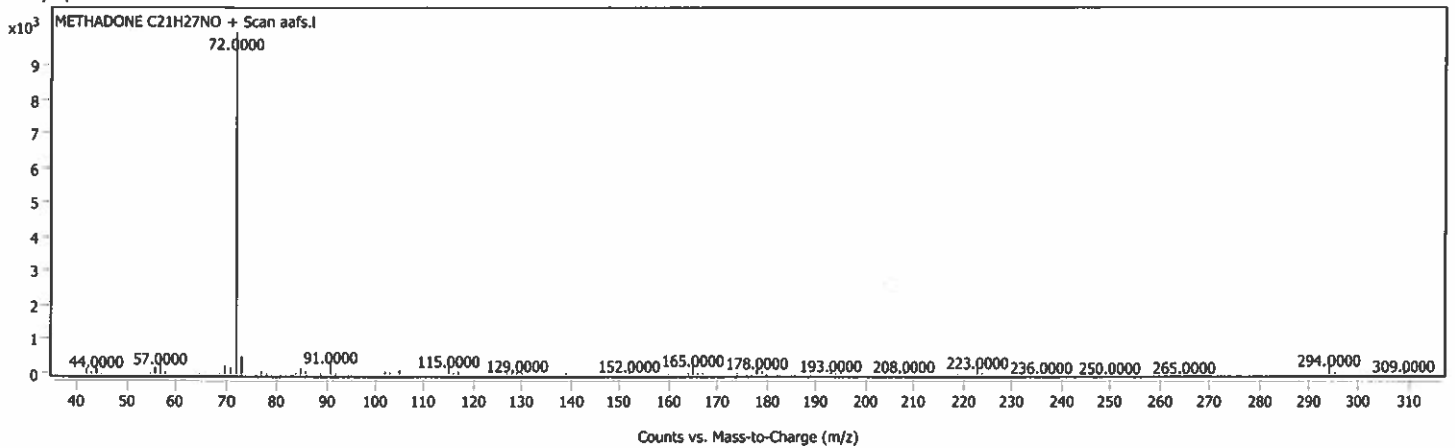
### Observed Spectrum



### Mirror Plot



### Library Spectrum



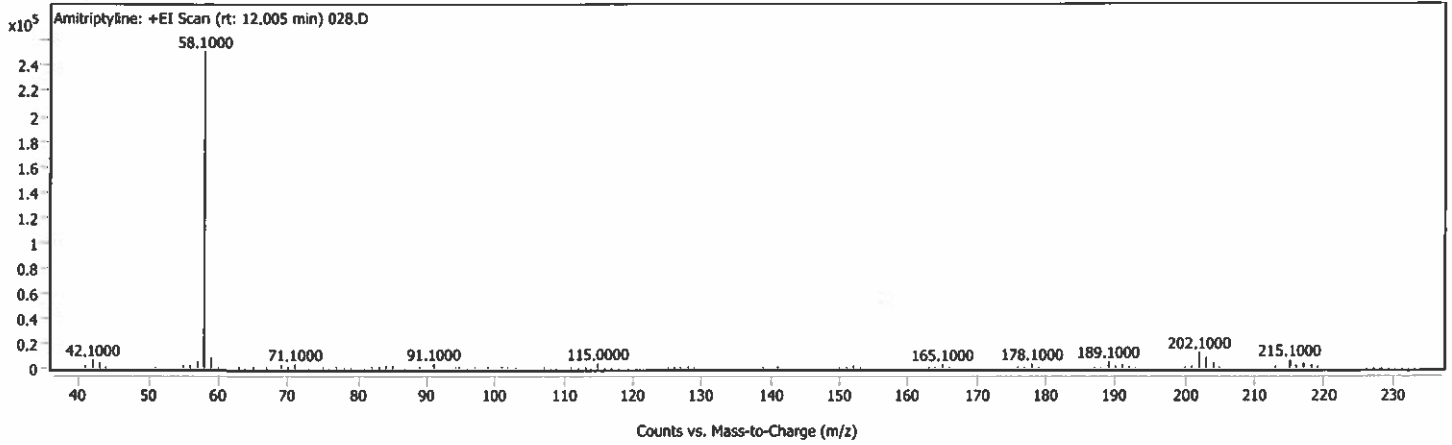
# Analysis Report

+ Scan (rt: 12.005 min)

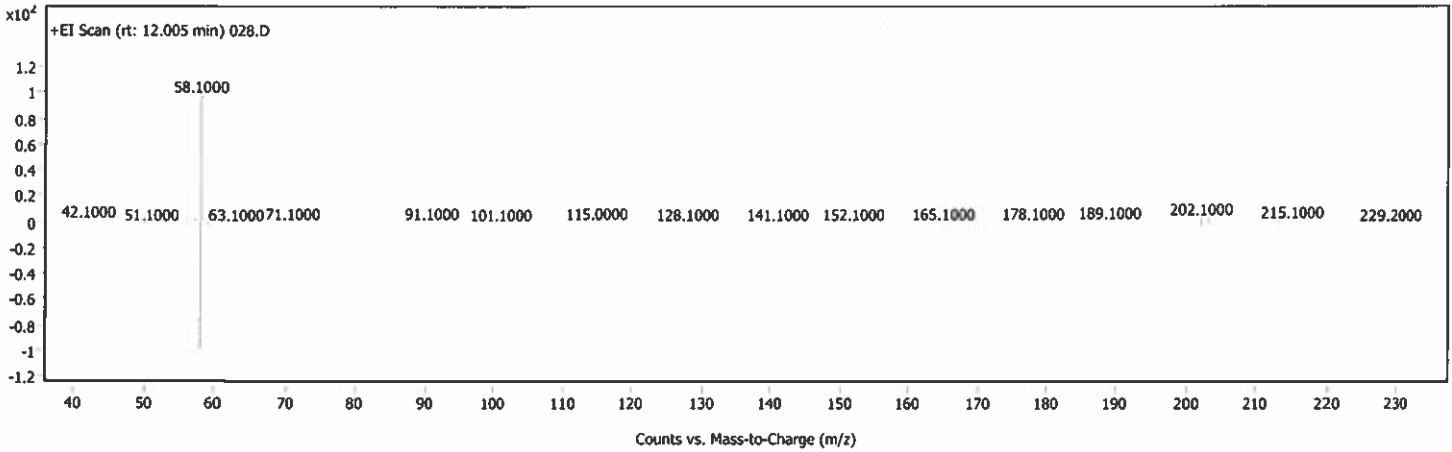
Amitriptyline; C20H23N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Amitriptyline	C20H23N	93.23	SWGDRUG 3.3.1	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

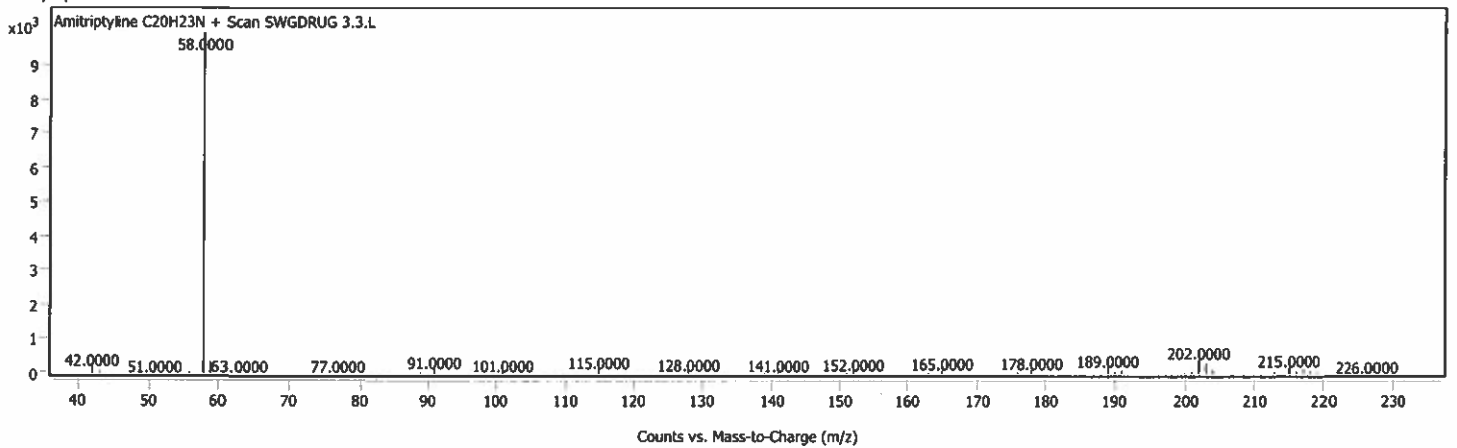
### Observed Spectrum



### Mirror Plot



### Library Spectrum





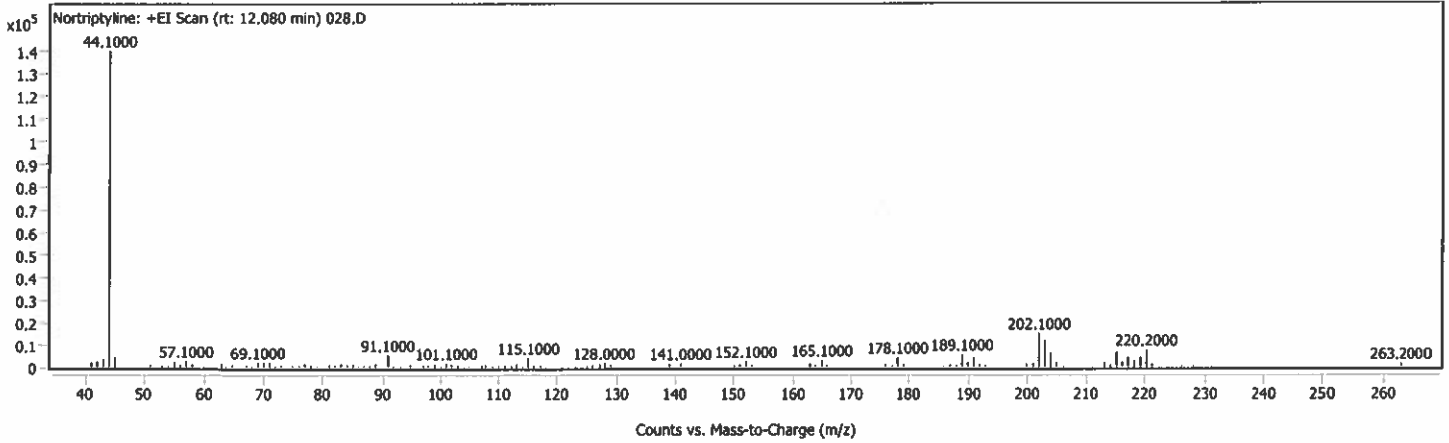
# Analysis Report

+ Scan (rt: 12.080 min)

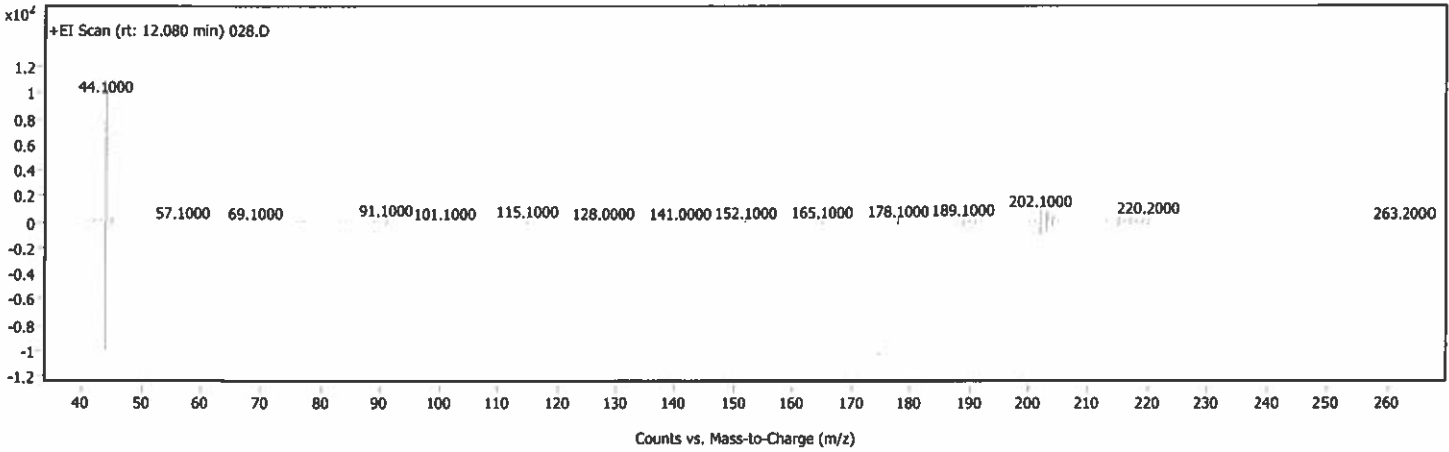
Nortriptyline; C19H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Nortriptyline	C19H21N	92.78	SWGDRUG 3.3.L	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

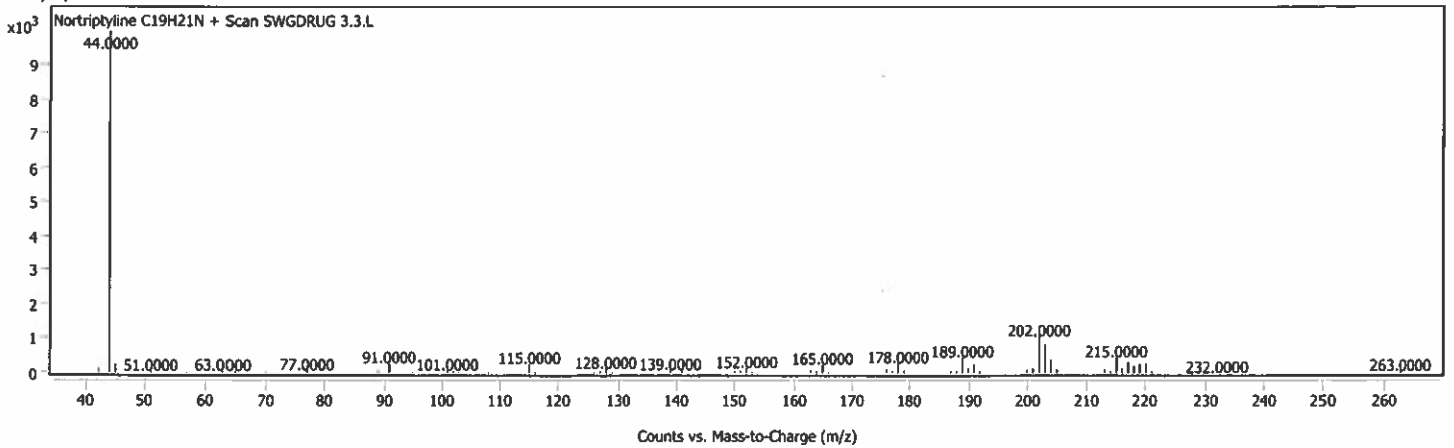
### Observed Spectrum



### Mirror Plot



### Library Spectrum



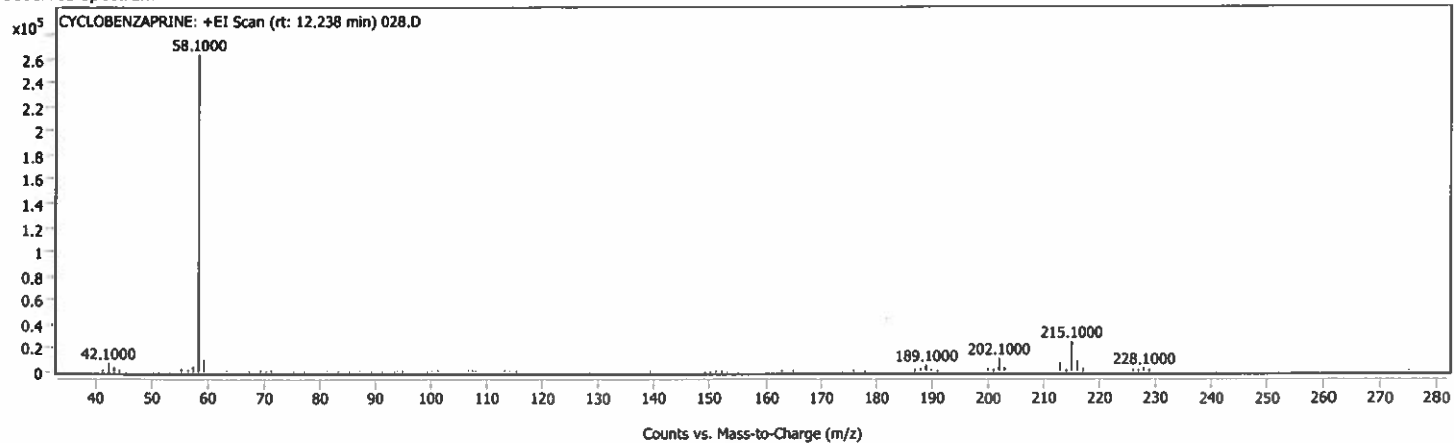
# Analysis Report

+ Scan (rt: 12.238 min)

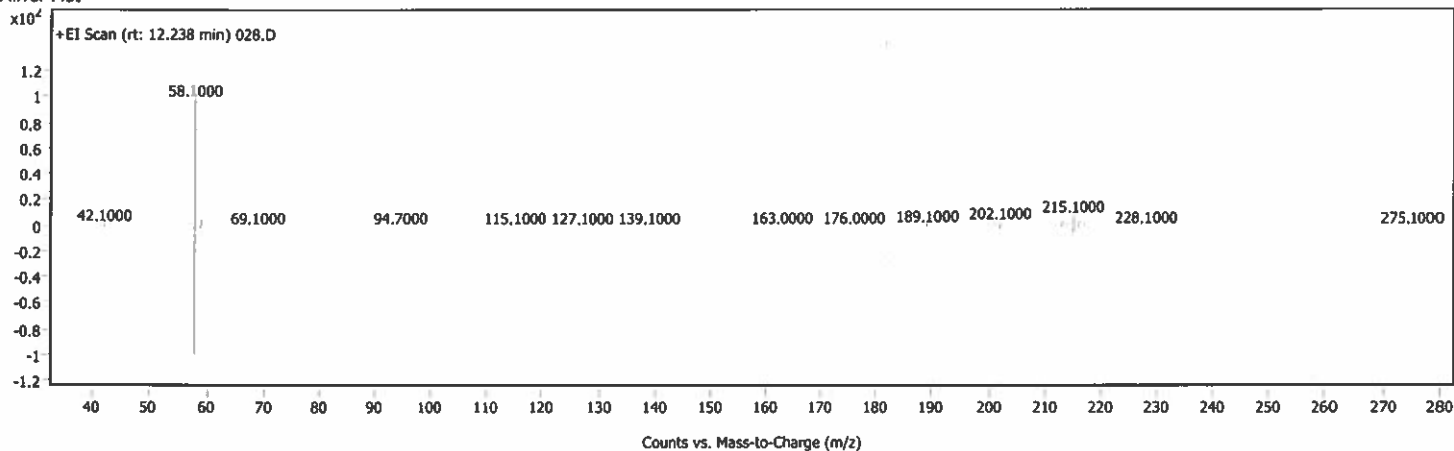
CYCLOBENZAPRINE; C20H21N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CYCLOBENZAPRINE	C20H21N	92.84	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/OC DATA IN XYLAZINE VALIDATION

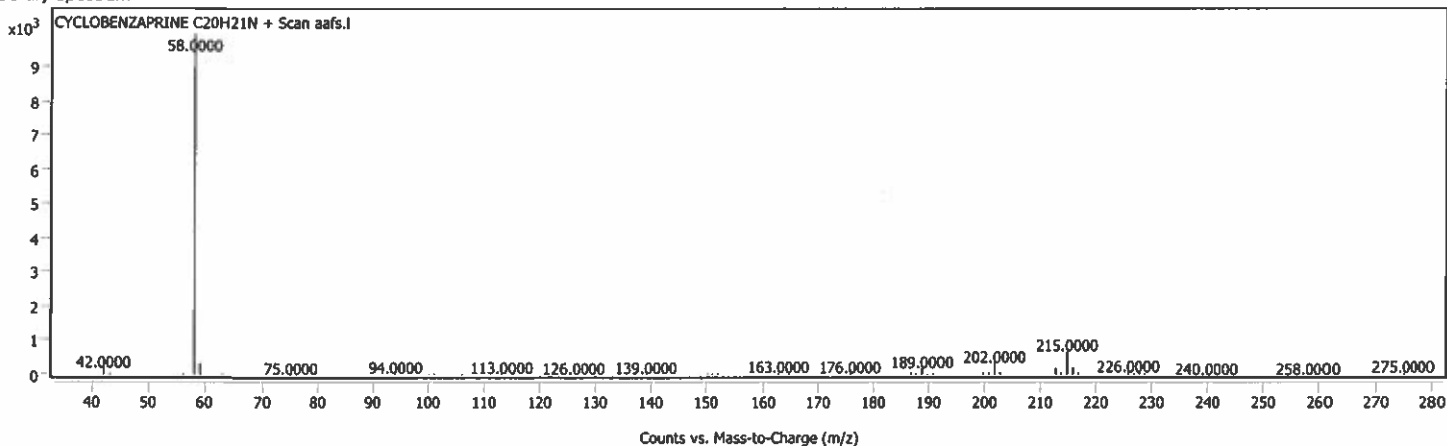
## Observed Spectrum



## Mirror Plot



## Library Spectrum



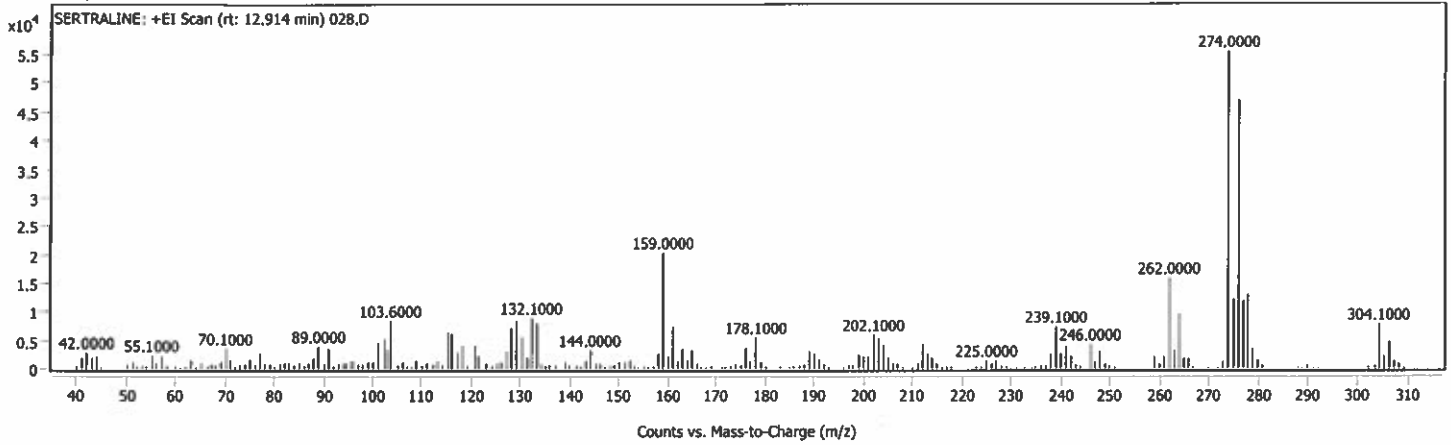
# Analysis Report

+ Scan (rt: 12.914 min)

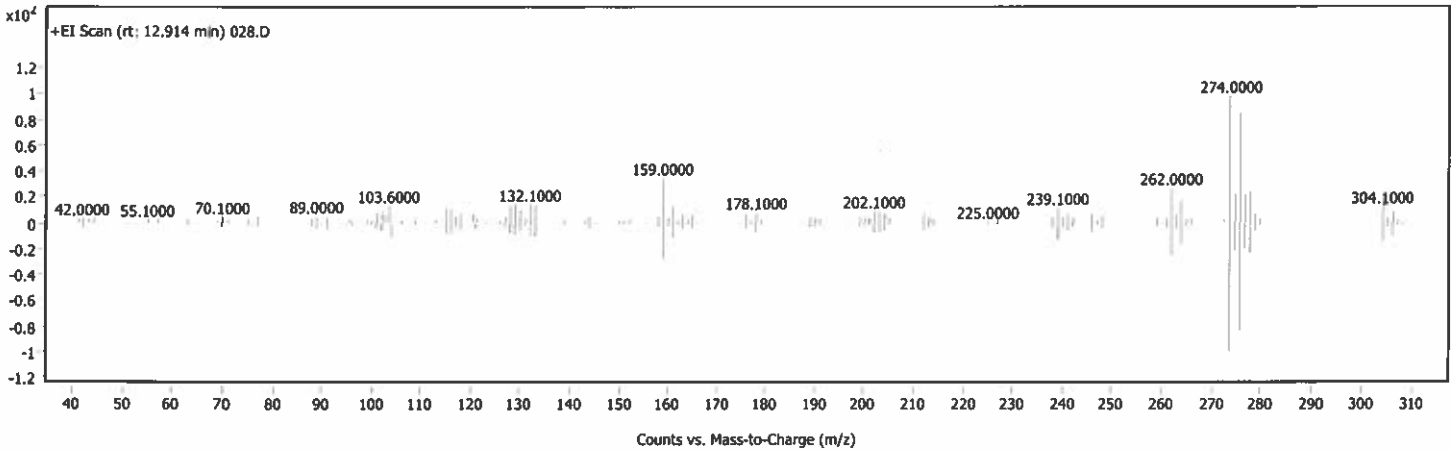
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	90.79	aafs.l	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/OC DATA IN XYLAZINE VALIDATION

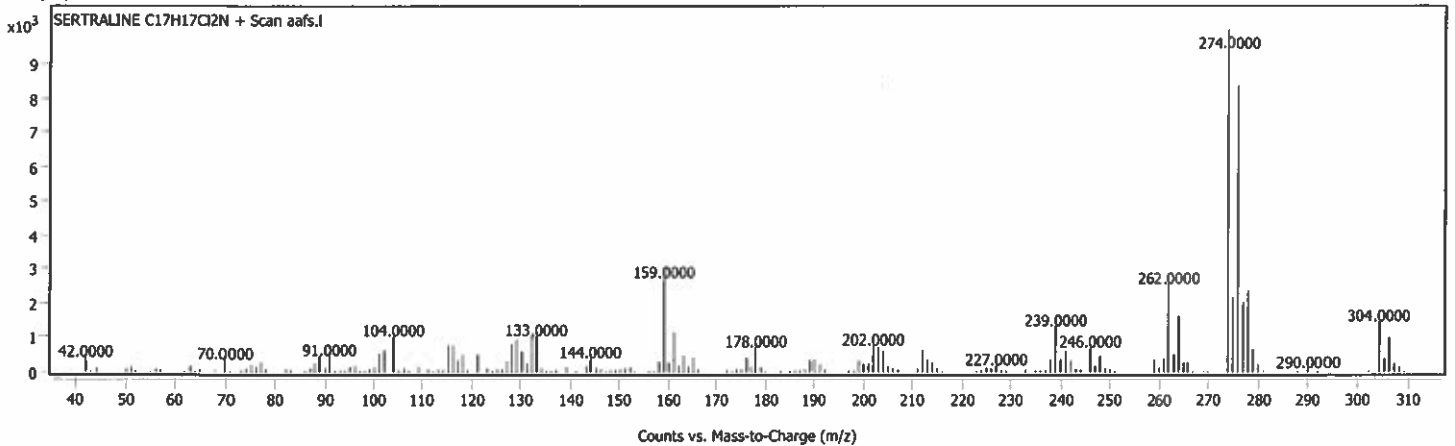
## Observed Spectrum



## Mirror Plot



## Library Spectrum



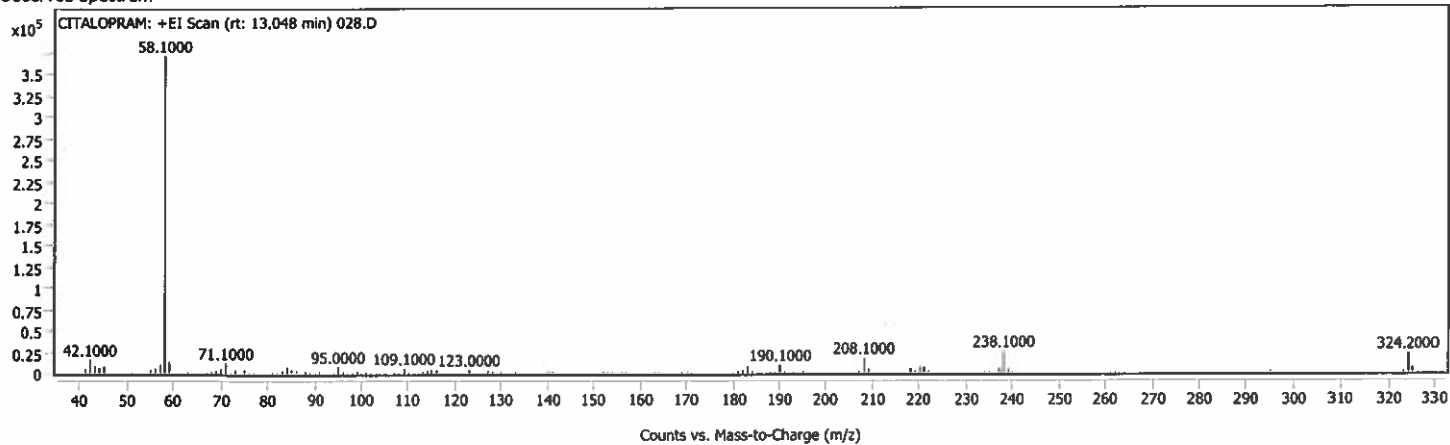
# Analysis Report

+ Scan (rt: 13.048 min)

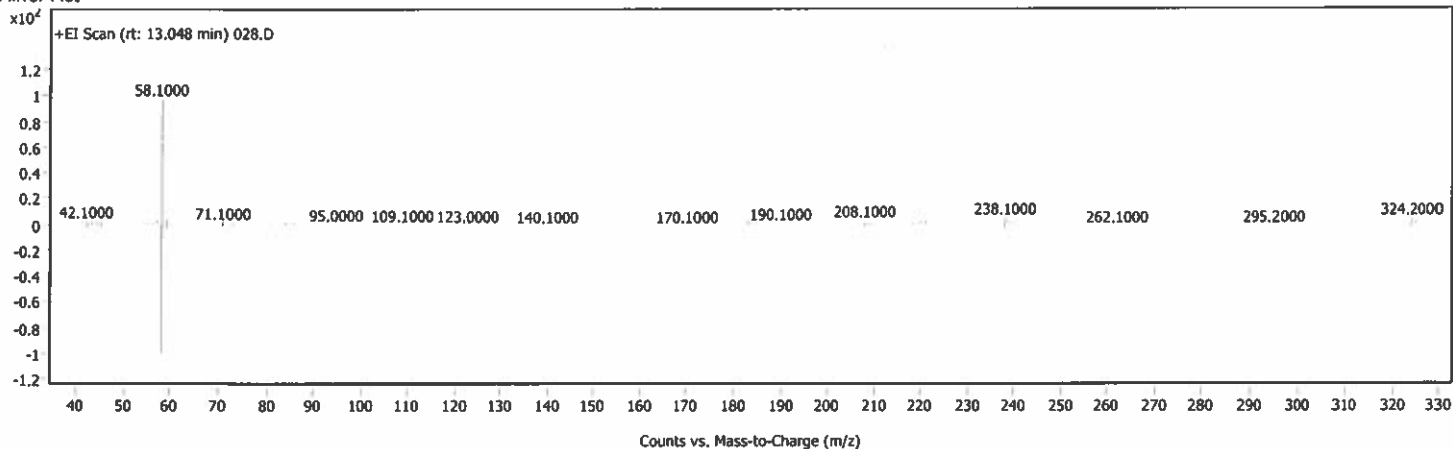
CITALOPRAM; C20H21FN2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CITALOPRAM	C20H21FN2O	93.81	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/OC DATA IN XYLAZINE VALIDATION

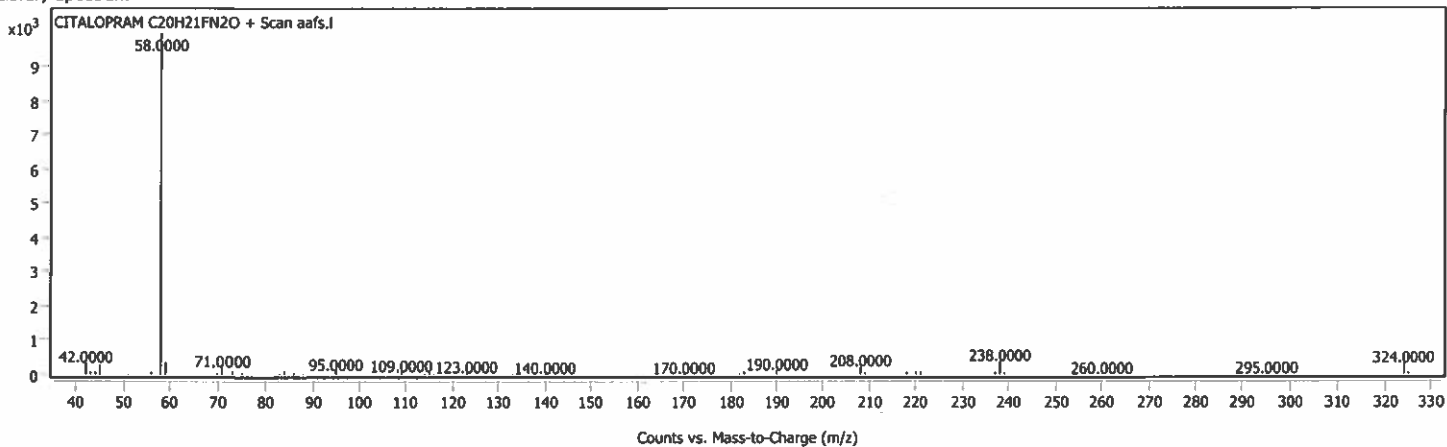
### Observed Spectrum



### Mirror Plot



### Library Spectrum

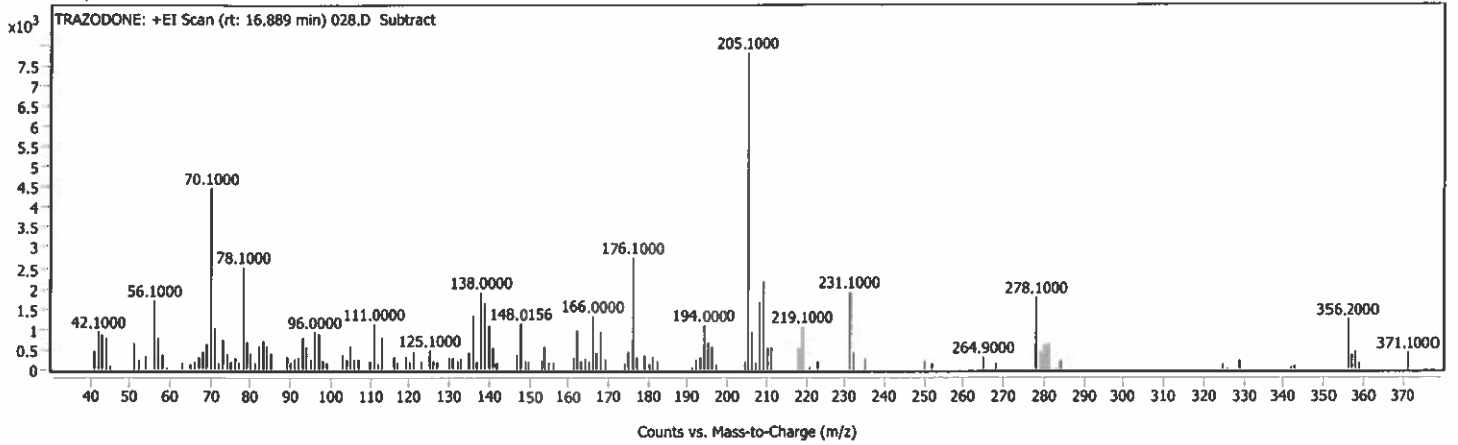


# Analysis Report

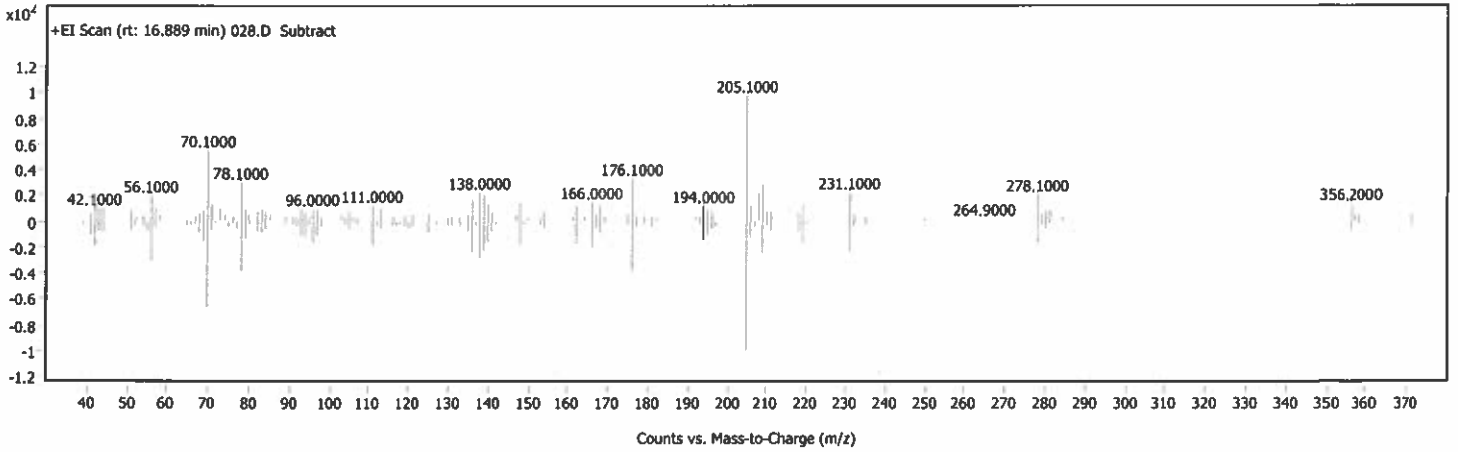
+ Scan (rt: 16.889 min) Sub TRAZODONE

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
TRAZODONE		85.39	aafs.I	MATRIX 10 W/ 0.01 mg/L CAL + XYLAZINE - LVR	KM/OC DATA IN XYLAZINE VALIDATION

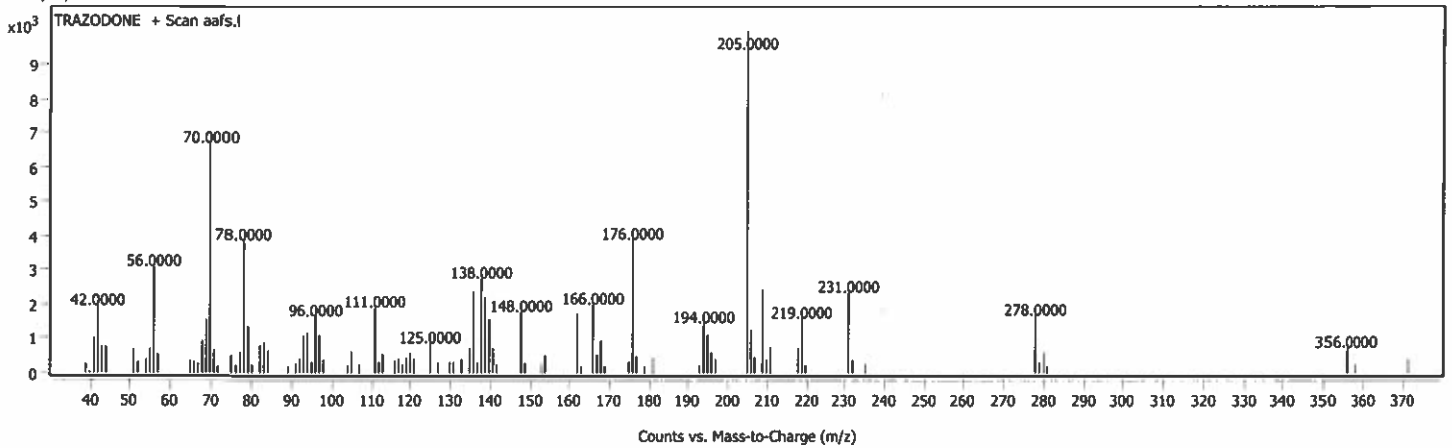
## Observed Spectrum



## Mirror Plot



## Library Spectrum

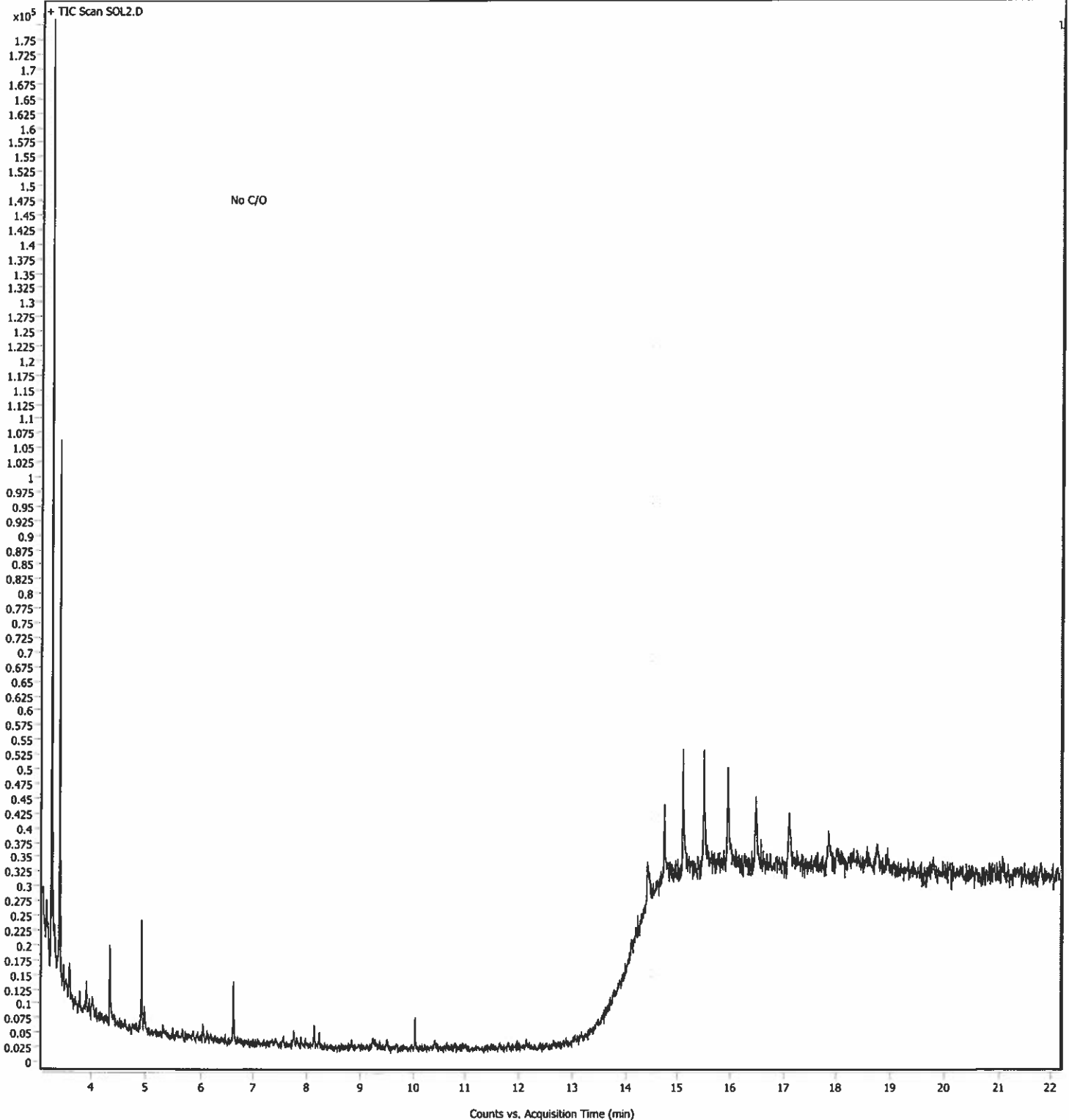


# Analysis Report

## Sample Information

Sample Name	Solvent Blank 2	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\SOL2.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 2:35:40 AM (UTC-04:00)
Position	57	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms

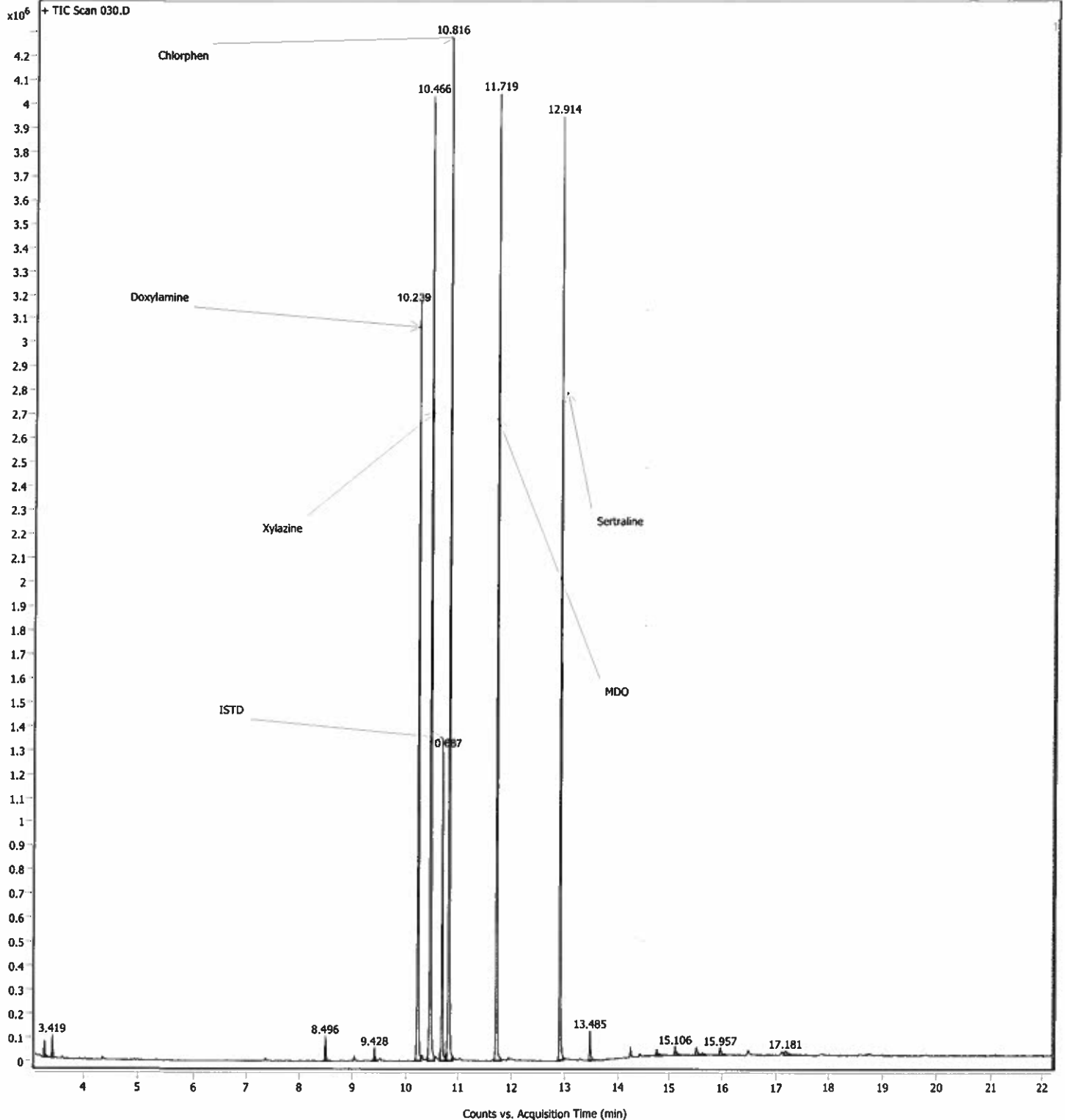


# Analysis Report

## Sample Information

Sample Name	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\030.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 3:27:19 AM (UTC-04:00)
Position	59	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

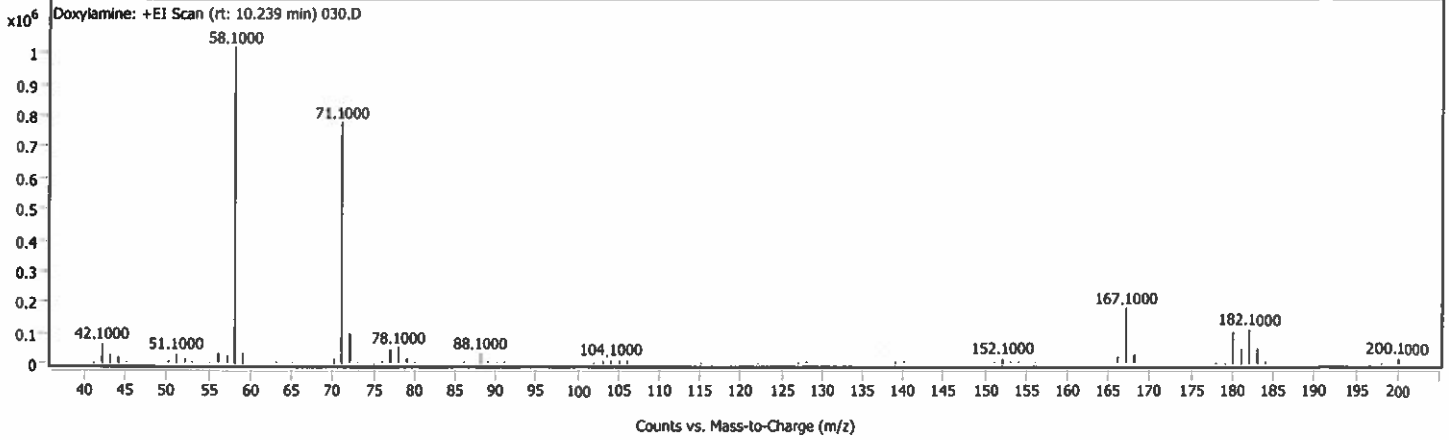
## Sample Spectra

+ Scan (rt: 10.239 min)

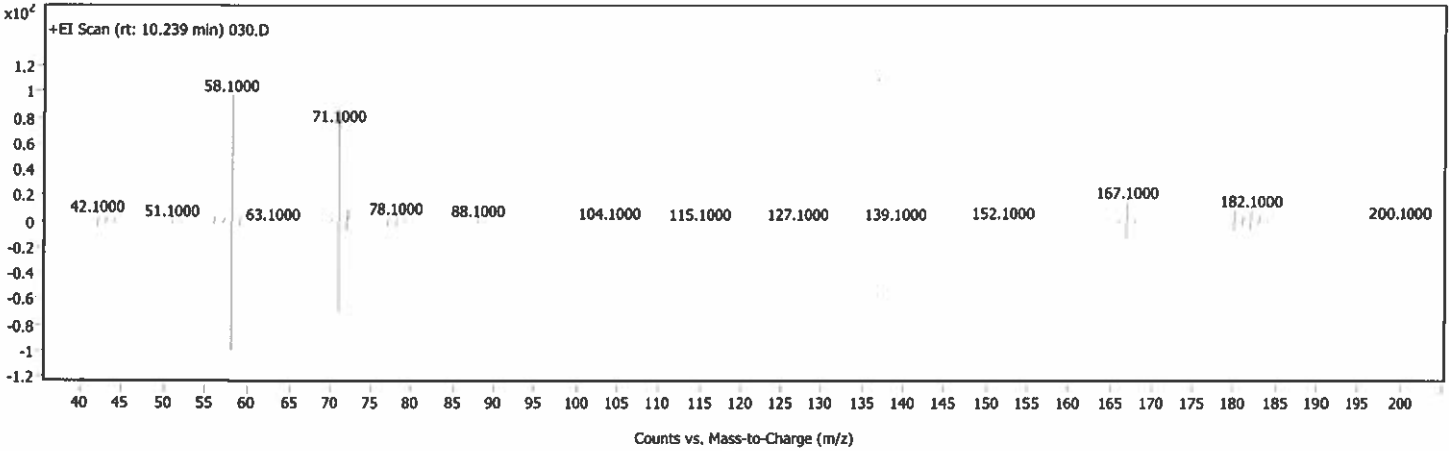
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.97	SWGDRUG 3.3.L	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

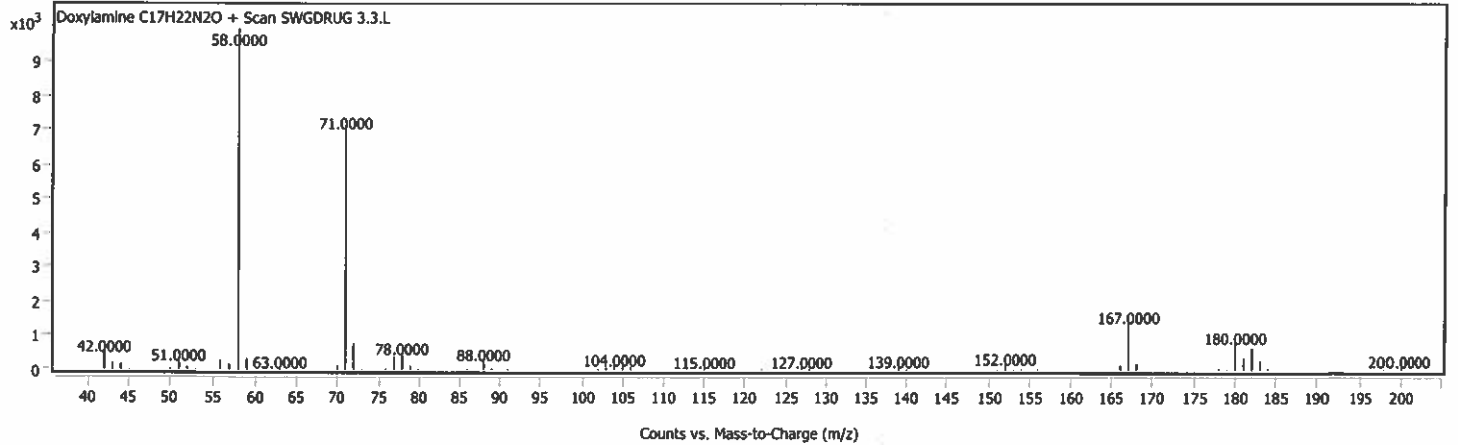
## Observed Spectrum



## Mirror Plot



## Library Spectrum





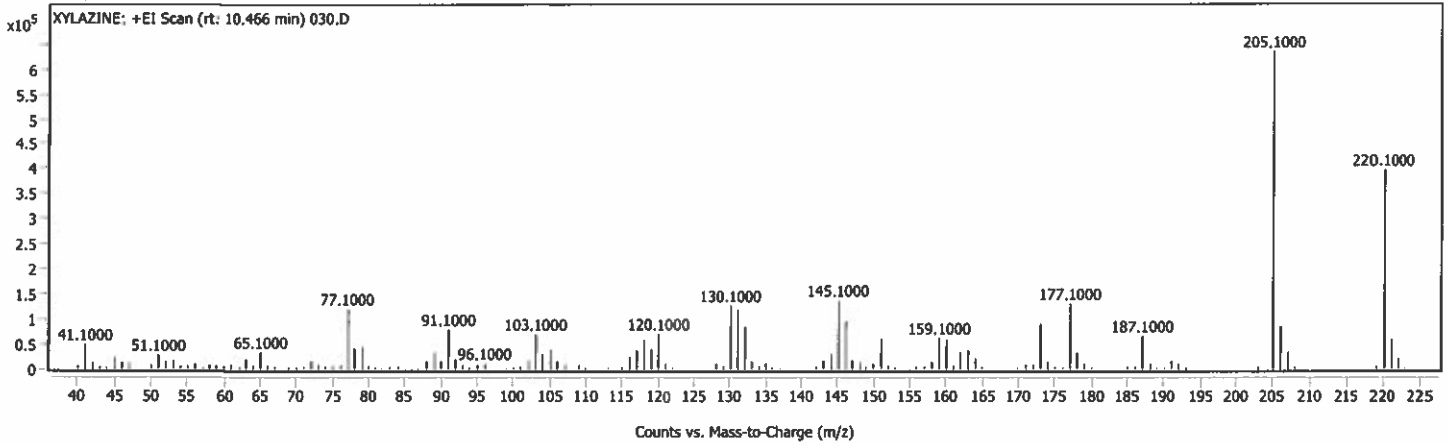
# Analysis Report

+ Scan (rt: 10.466 min)

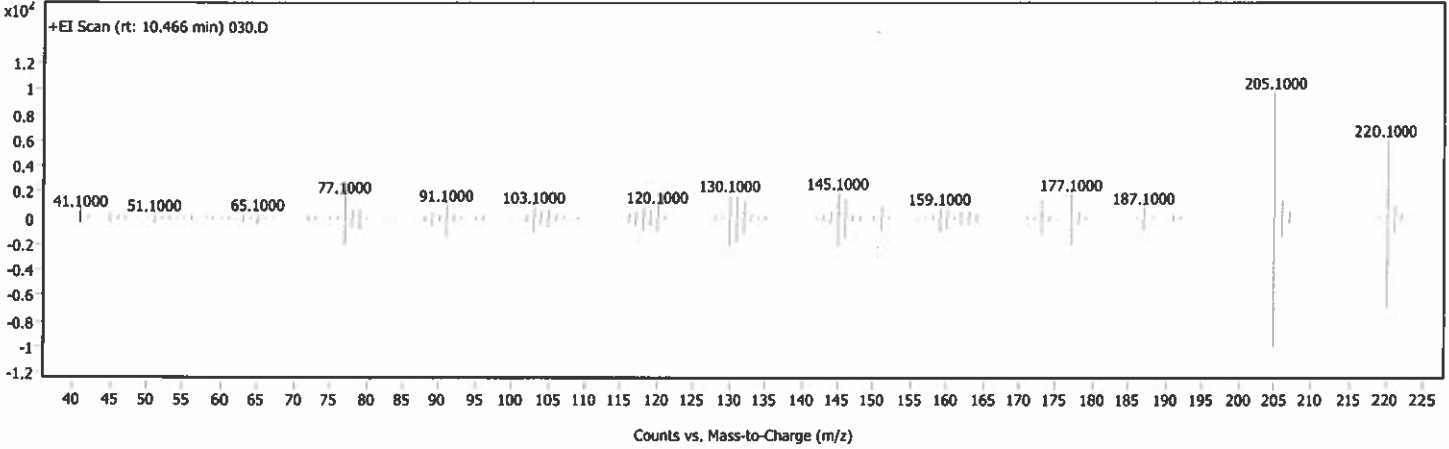
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.66	aafs.l	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

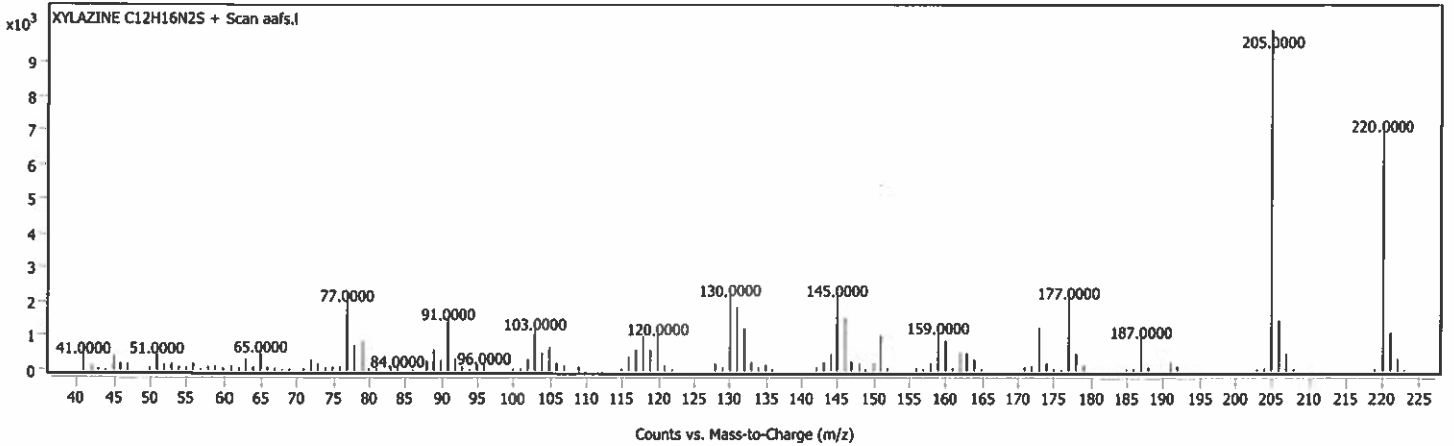
## Observed Spectrum



## Mirror Plot



## Library Spectrum



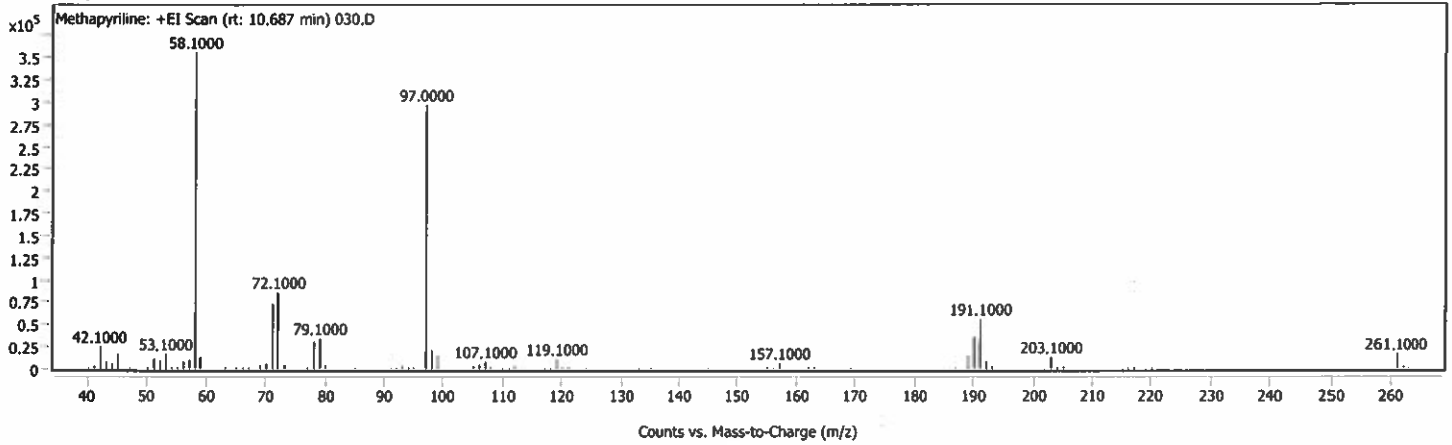
# Analysis Report

+ Scan (rt: 10.687 min)

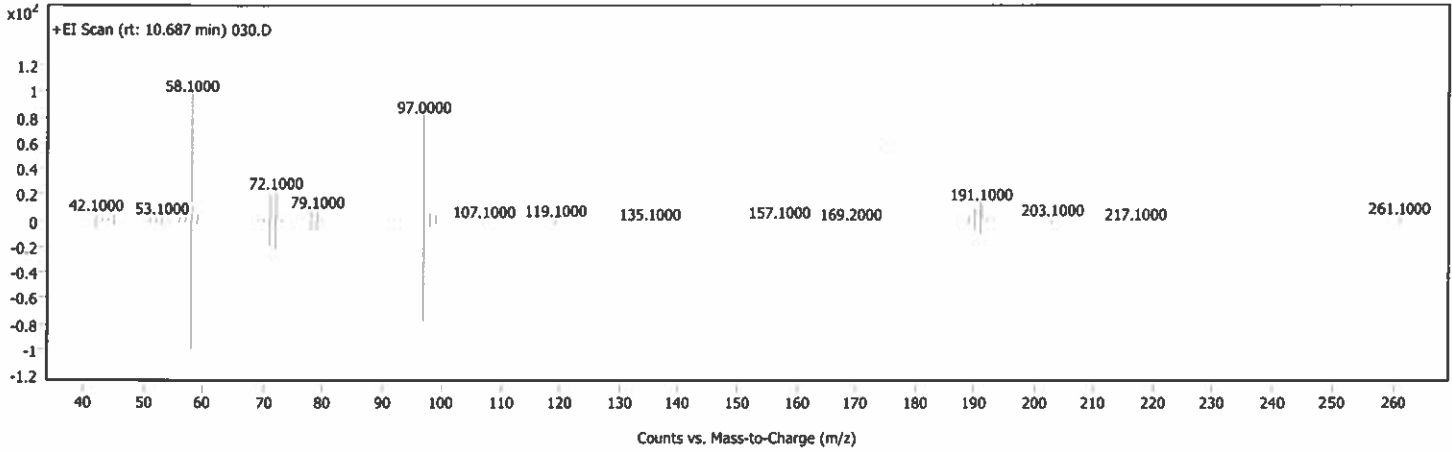
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.39	SWGDRUG 3.3.L	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

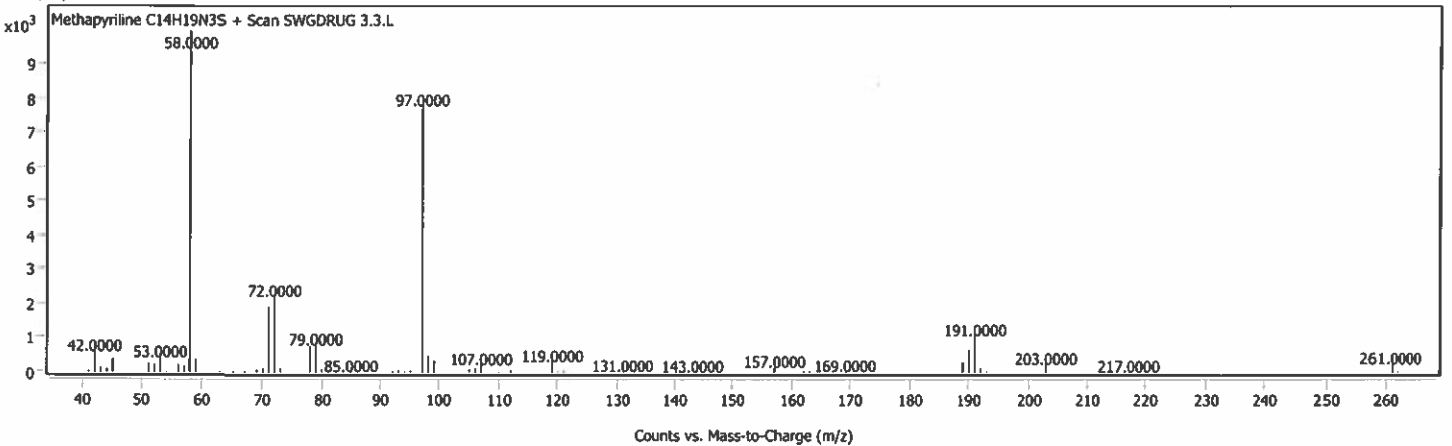
## Observed Spectrum



## Mirror Plot



## Library Spectrum



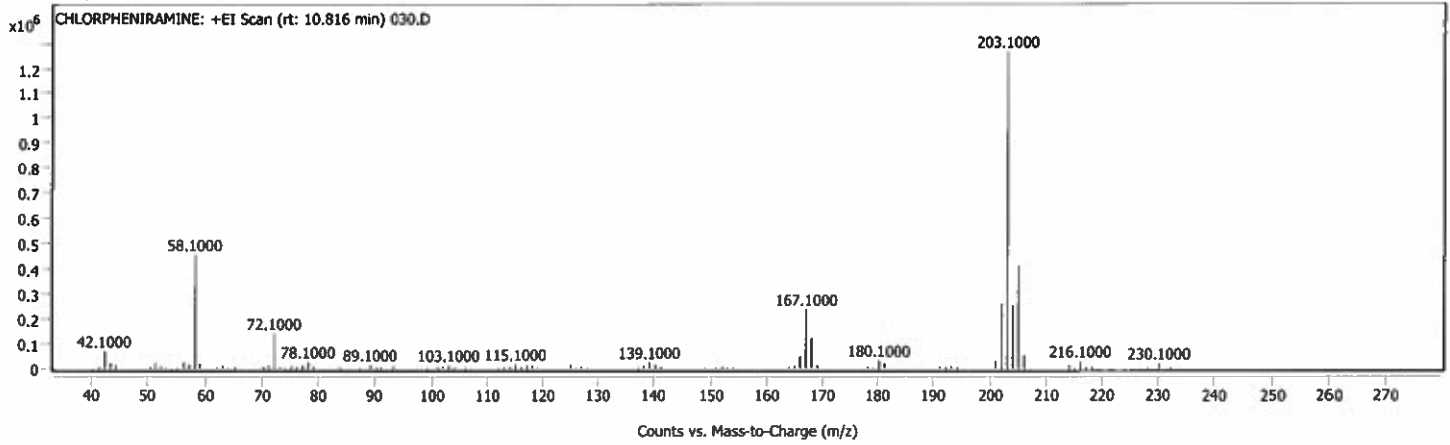
# Analysis Report

+ Scan (rt: 10.816 min)

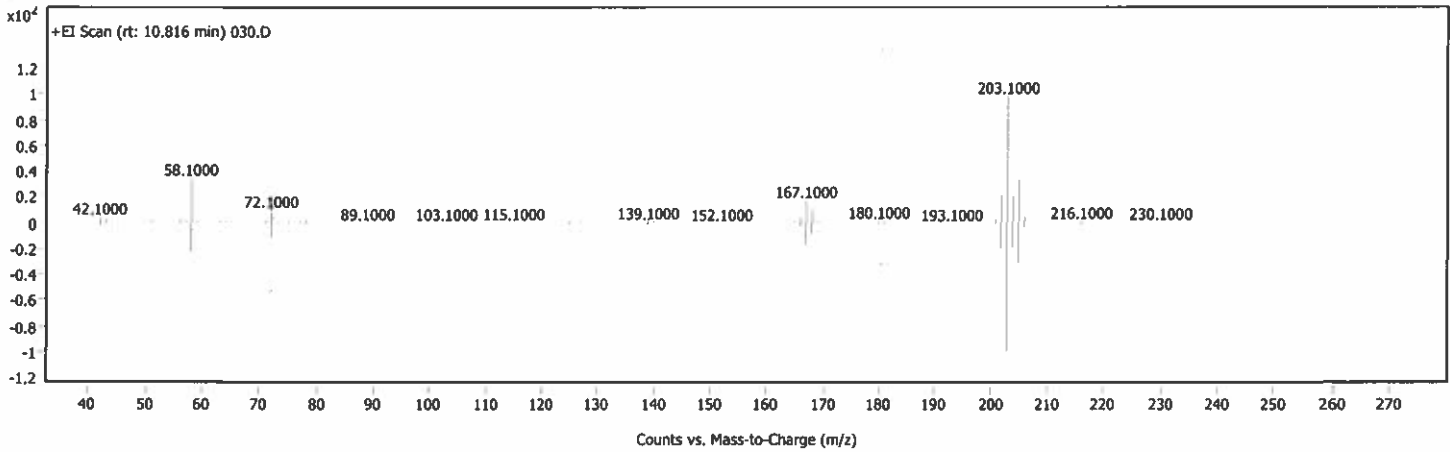
CHLORPHENIRAMINE; C16H19CIN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19CIN2	99.04	aafs.I	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

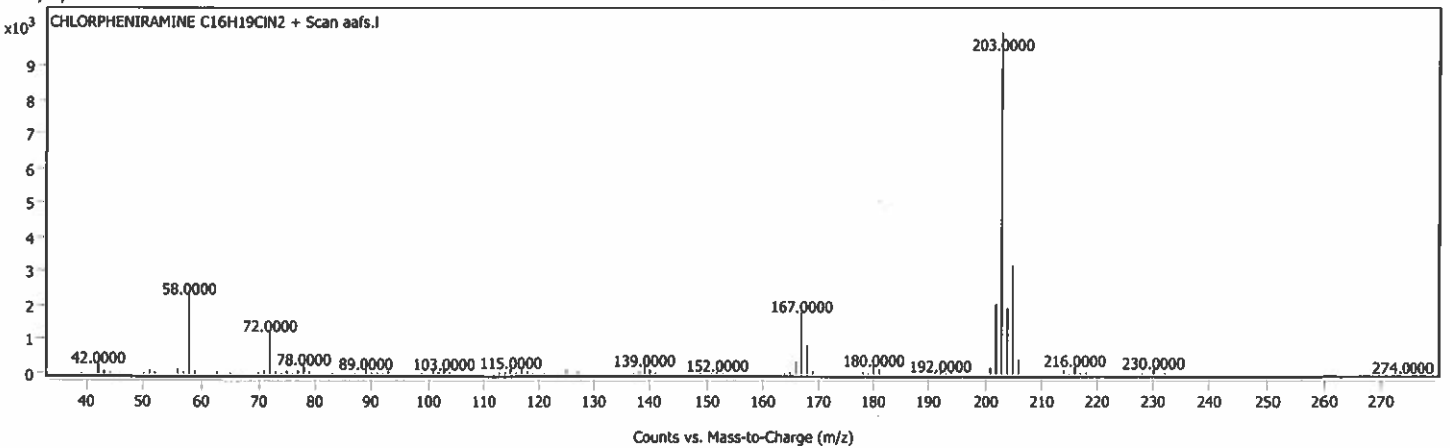
### Observed Spectrum



### Mirror Plot



### Library Spectrum



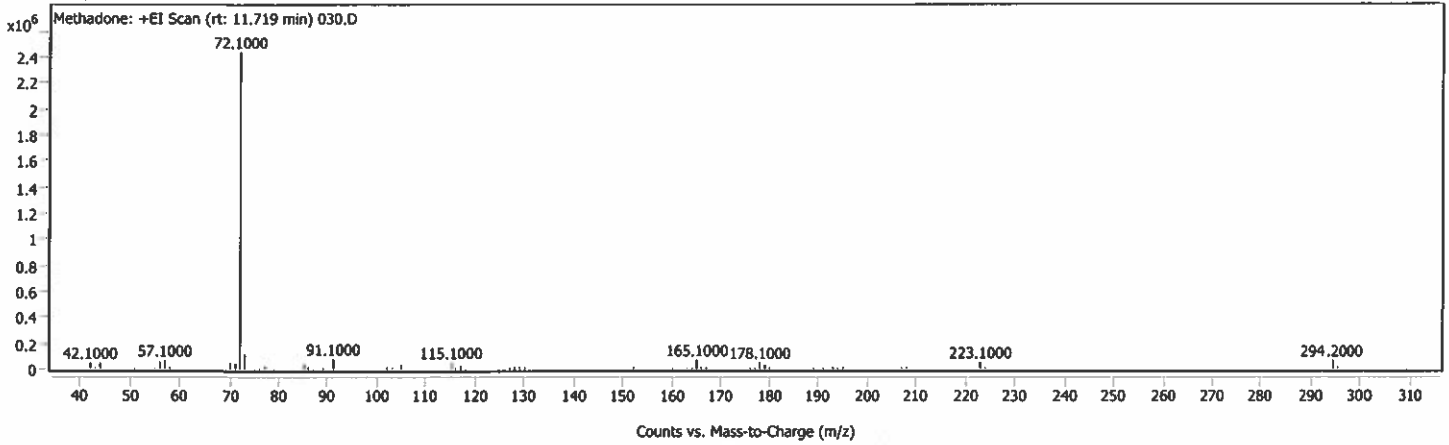
# Analysis Report

+ Scan (rt: 11.719 min)

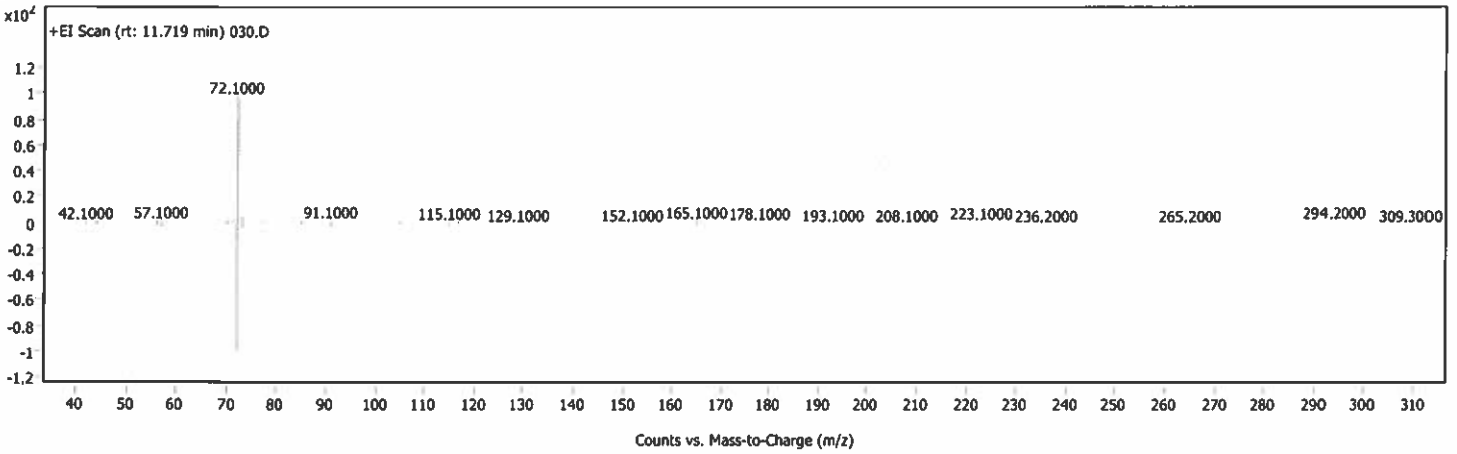
Methadone; C<sub>21</sub>H<sub>27</sub>NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C <sub>21</sub> H <sub>27</sub> NO	97.17	SWGDRUG 3.3.L	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

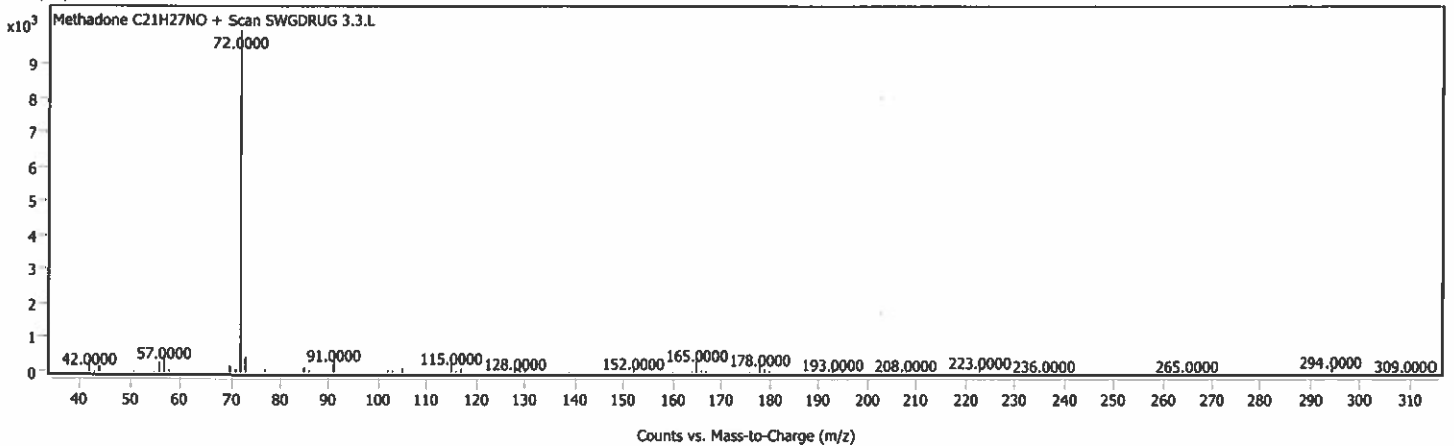
### Observed Spectrum



### Mirror Plot



### Library Spectrum



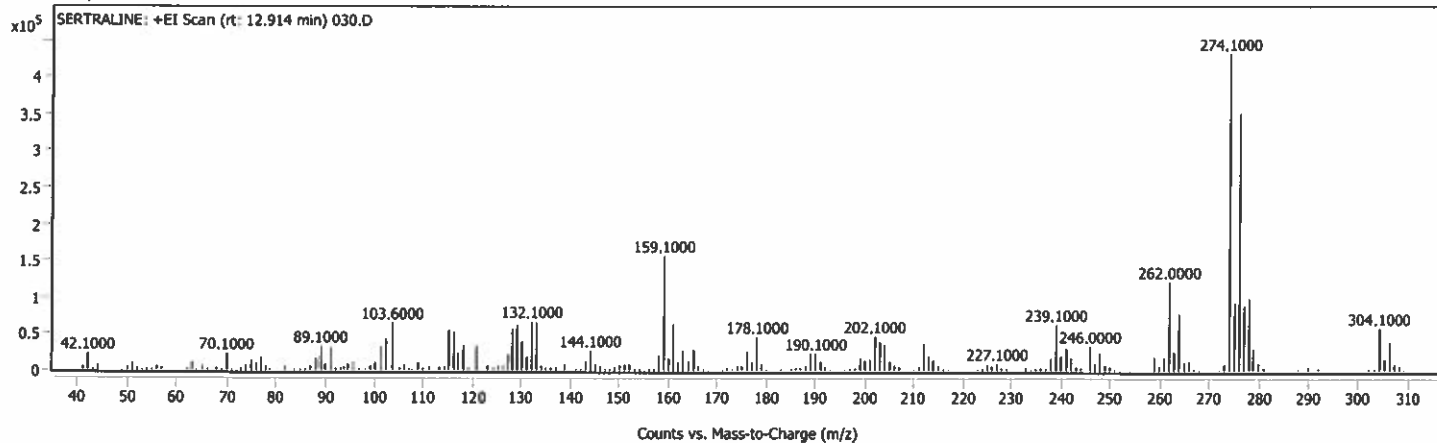
# Analysis Report

+ Scan (rt: 12.914 min)

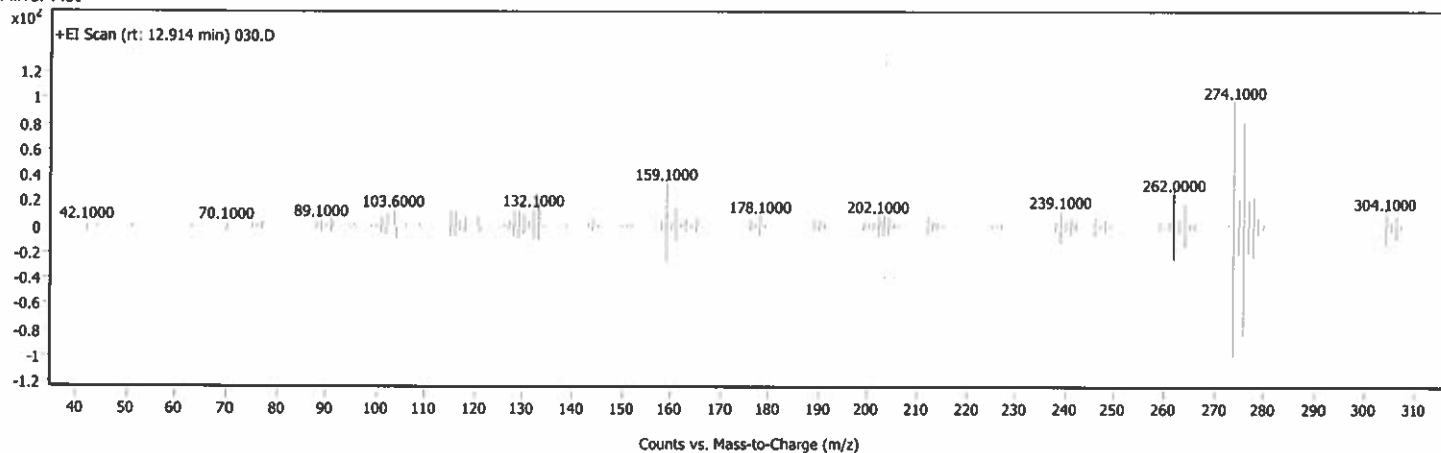
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	92.32	aafs.I	MATRIX 2 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

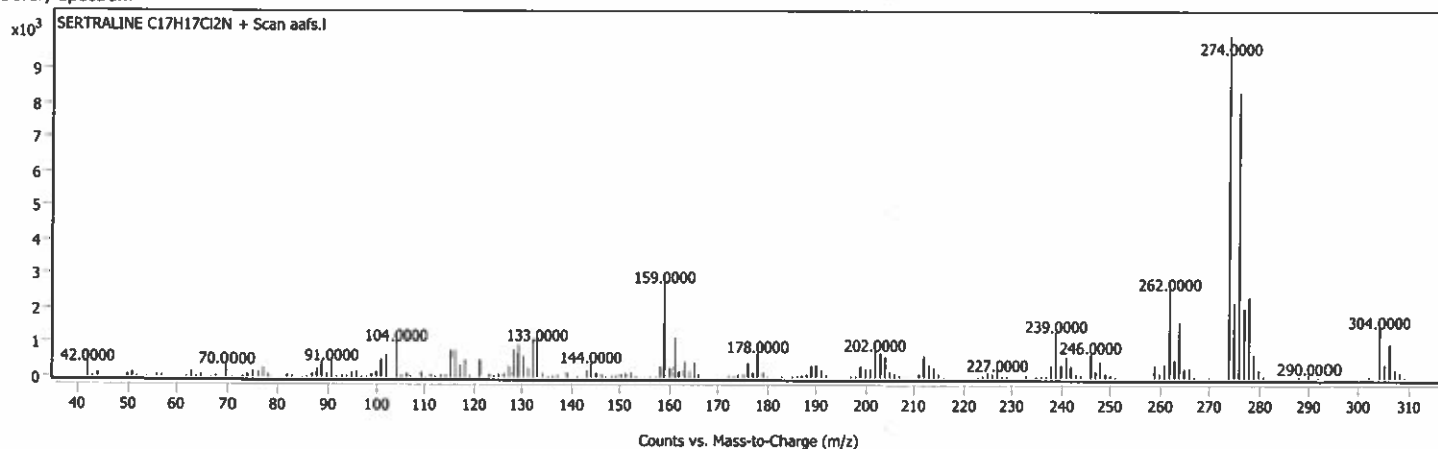
## Observed Spectrum



## Mirror Plot

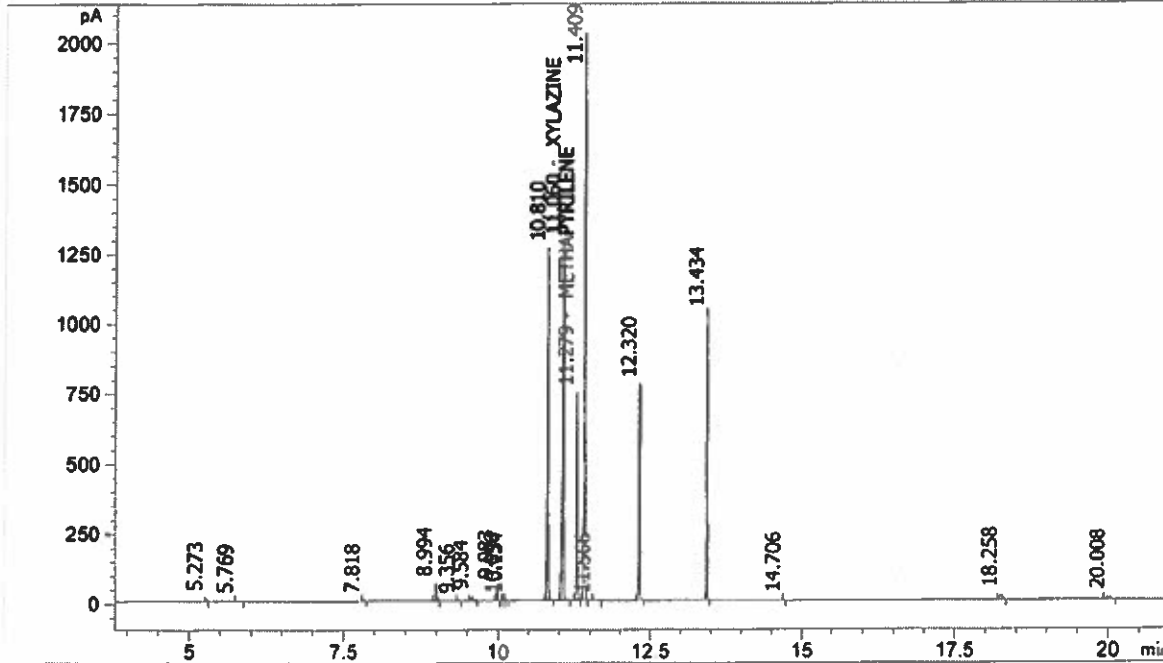


## Library Spectrum



Injection Date: 9/29/2021 3:08:08 AM Seq Line: 34  
 Sample Name: MATRIX 2 W/ IH CTL + -> Vial 35  
 Sample Info: xylazine km 11/17/22

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



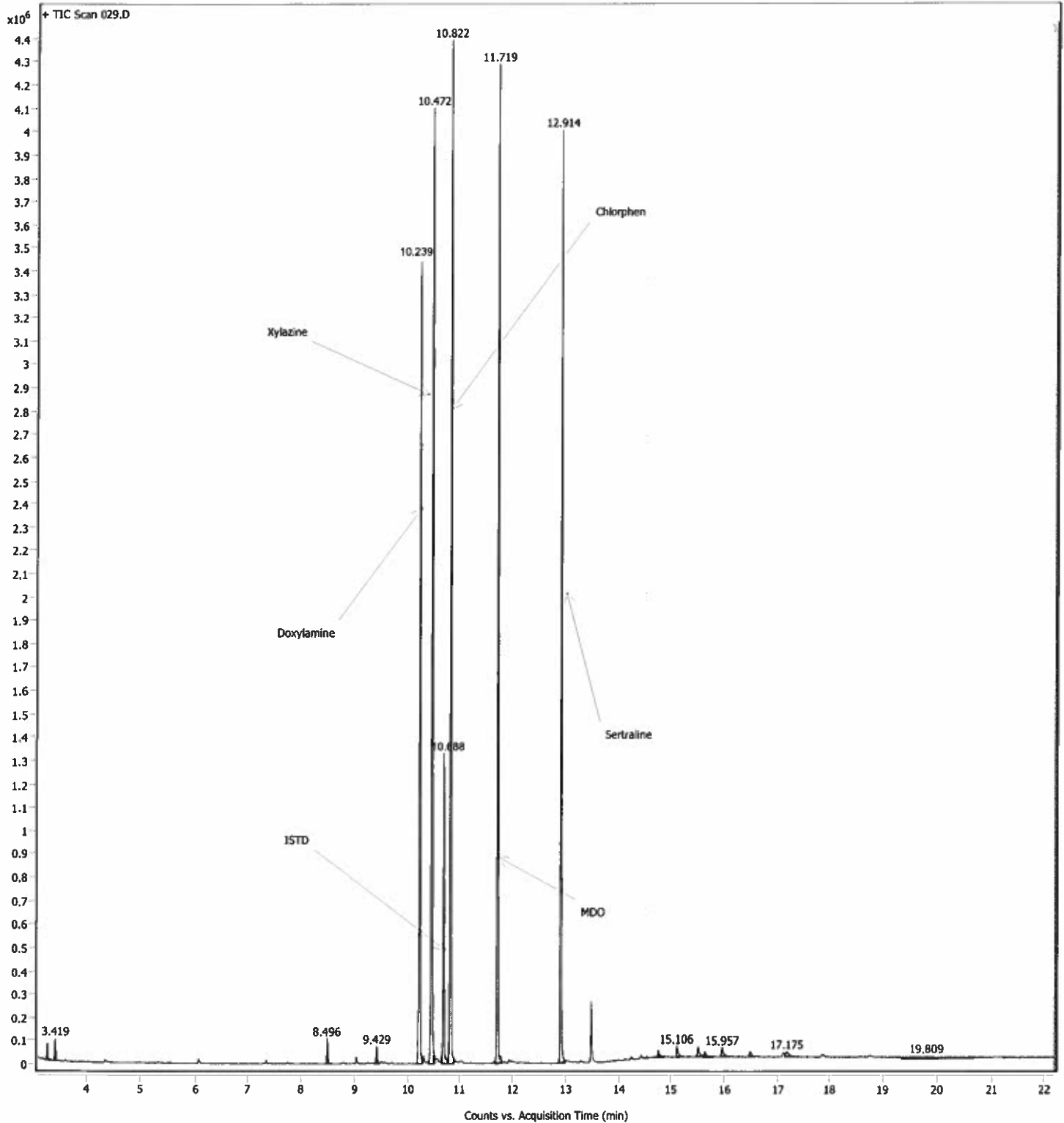
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.060	11.046	1849.875	1285.788	0.000000	XYLAZINE
11.279	11.278	867.967	742.619	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 W/ IH CTL + XYLAZINE - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\029.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 3:01:26 AM (UTC-04:00)
Position	58	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

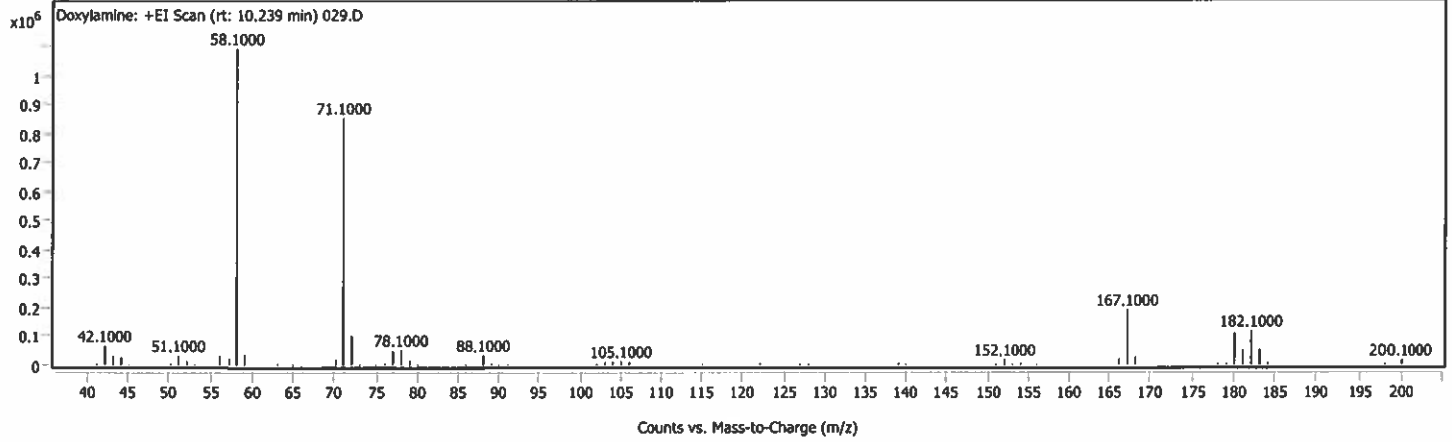
## Sample Spectra

+ Scan (rt: 10.239 min)

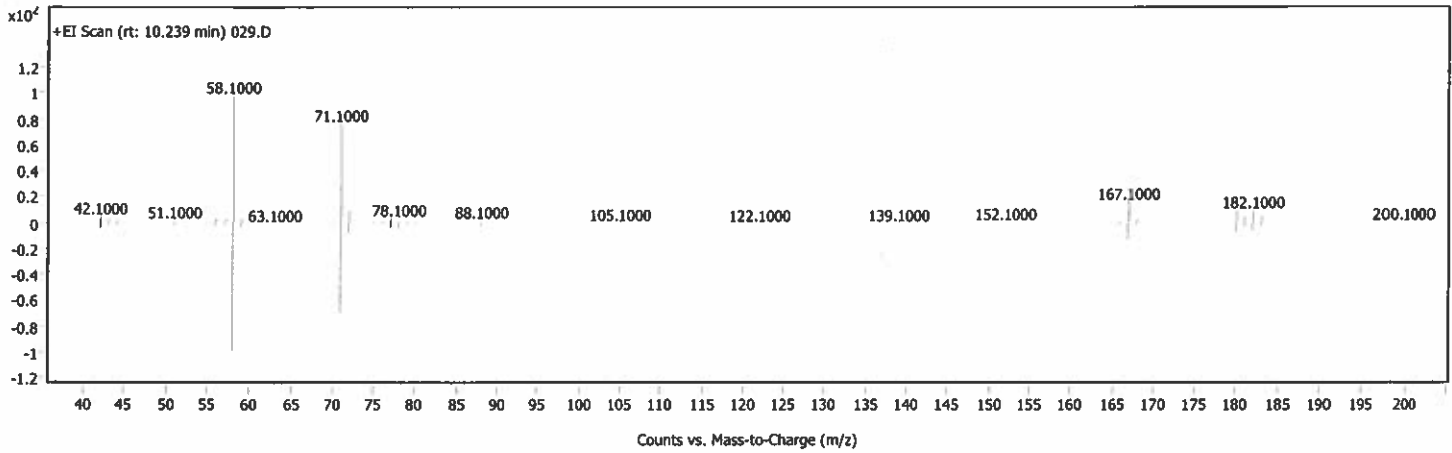
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.94	SWGDRUG 3.3.L	MATRIX 1 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

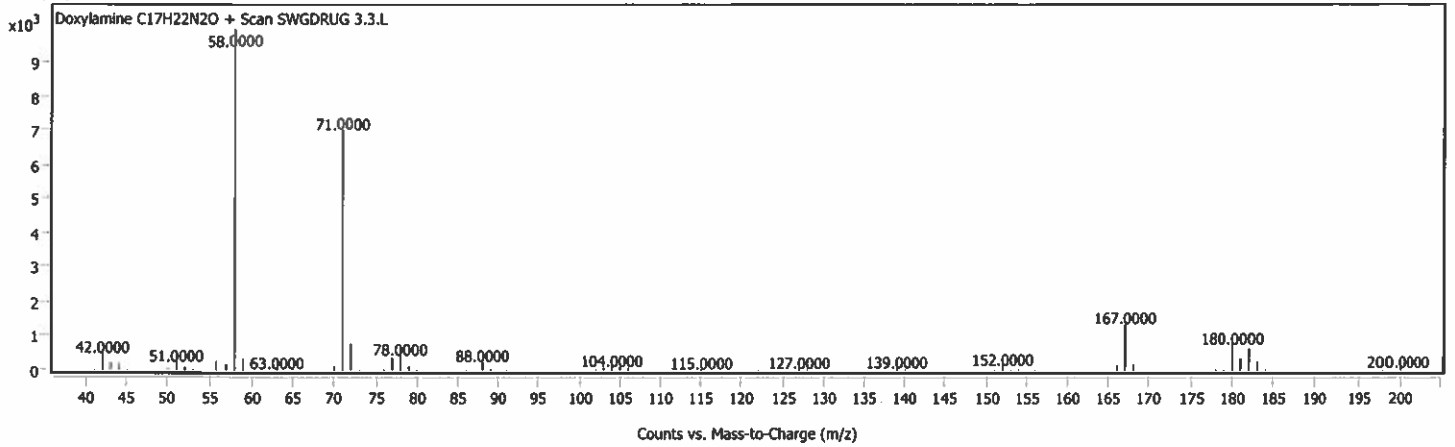
## Observed Spectrum



## Mirror Plot



## Library Spectrum





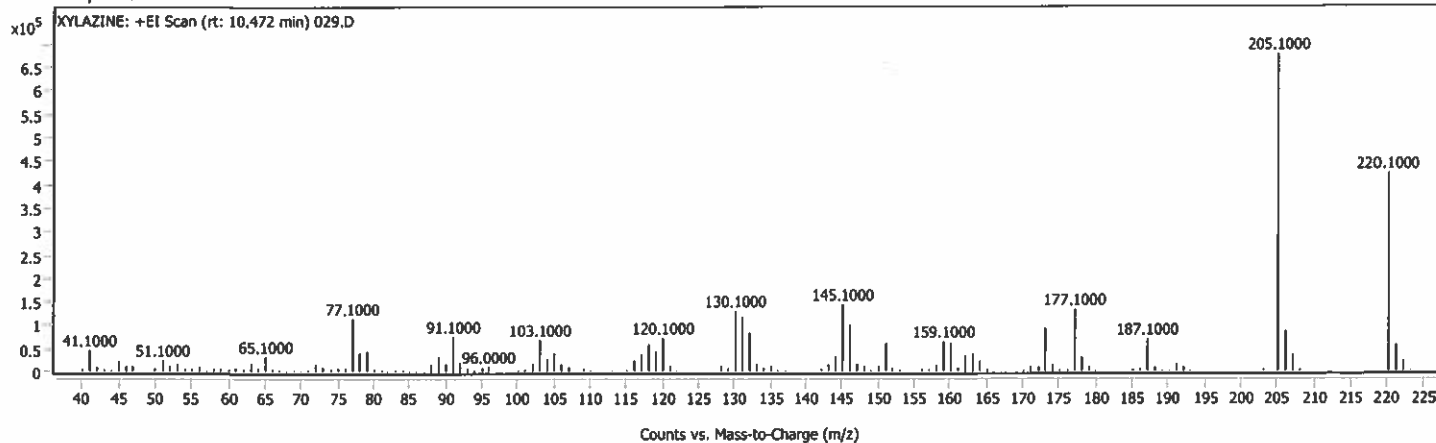
# Analysis Report

+ Scan (rt: 10.472 min)

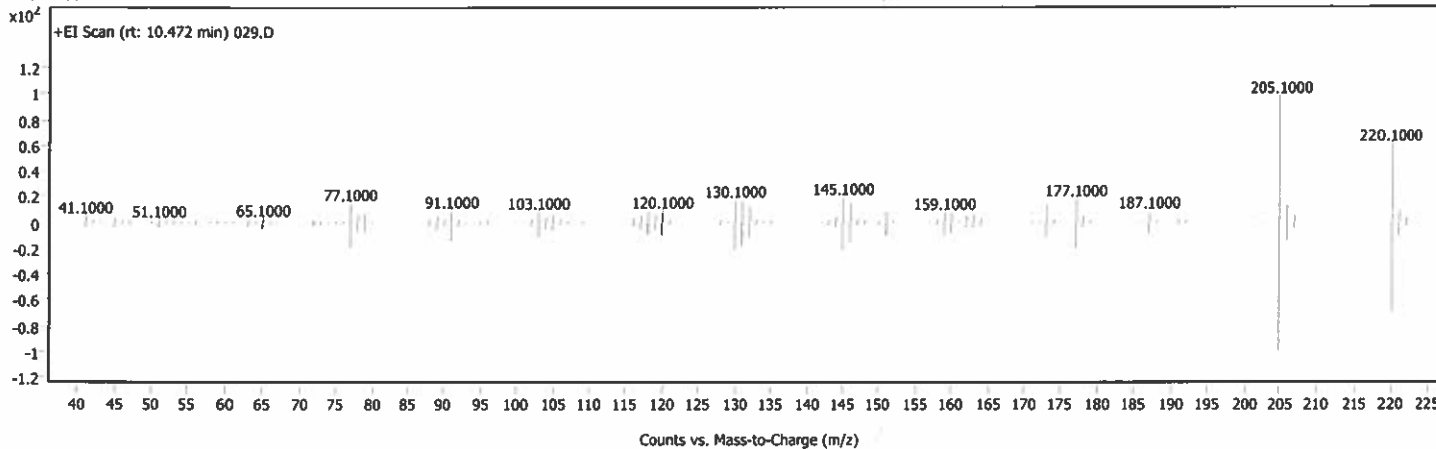
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.58	aafs.l	MATRIX 1 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

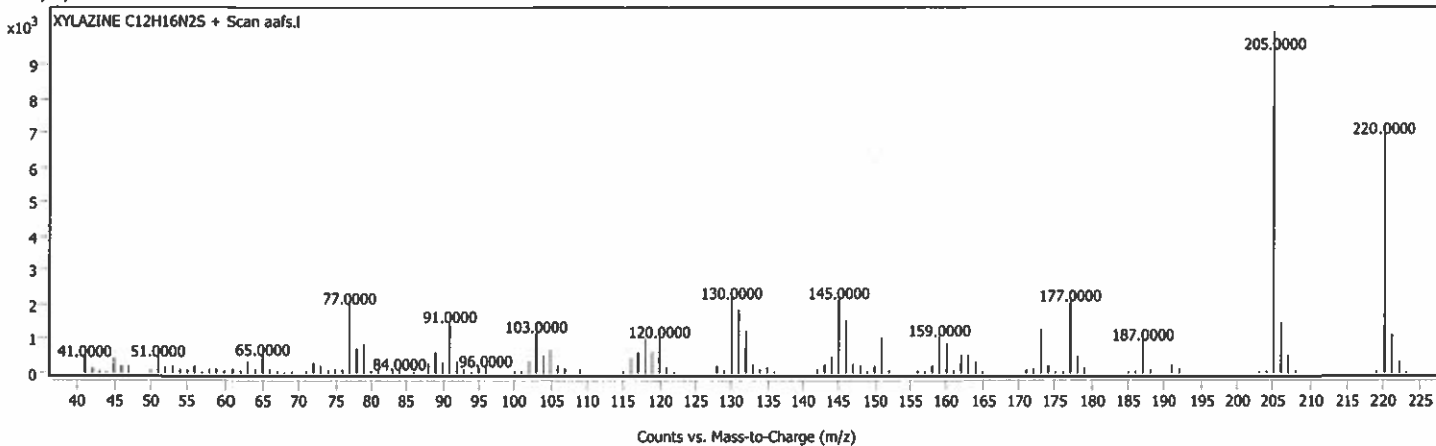
## Observed Spectrum



## Mirror Plot



## Library Spectrum



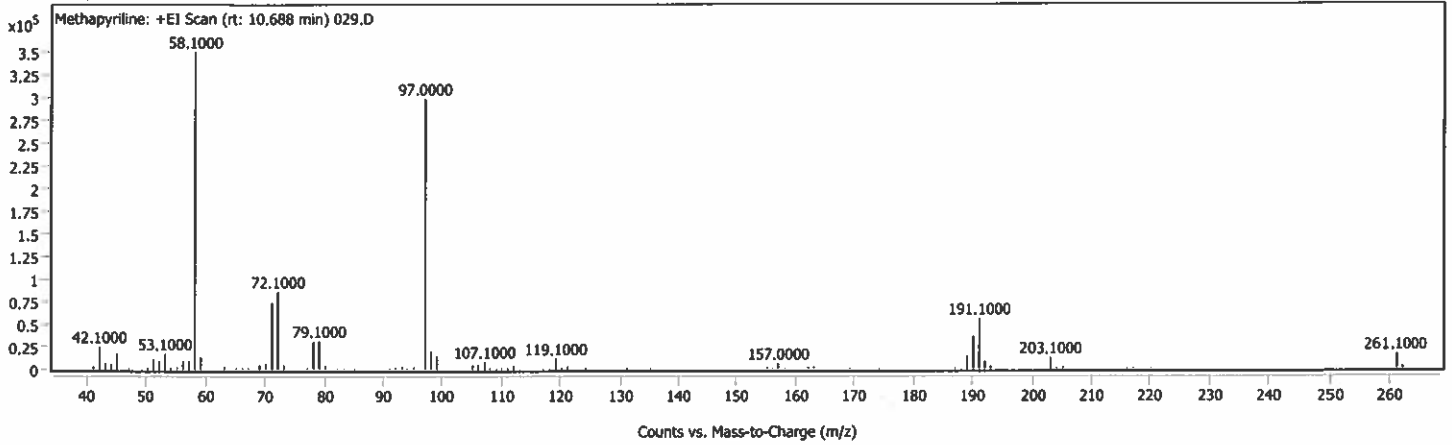
# Analysis Report

+ Scan (rt: 10.688 min)

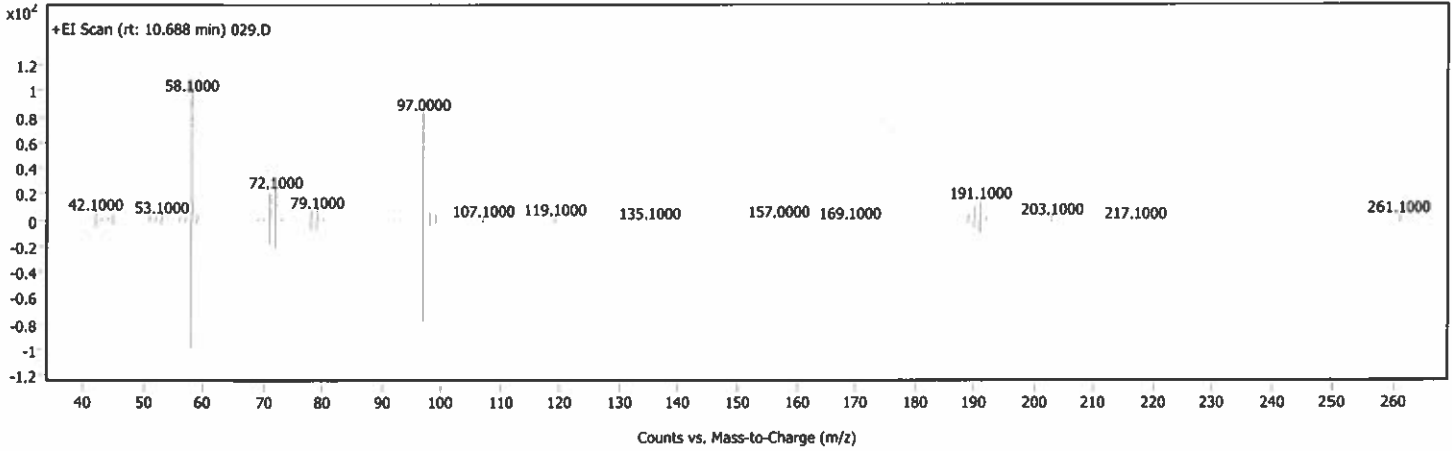
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.49	SWGDRUG 3.3.L	MATRIX 1 W/ 1H CTL + XYLAZINE - 8LD	KM/QC DATA IN XYLAZINE VALIDATION

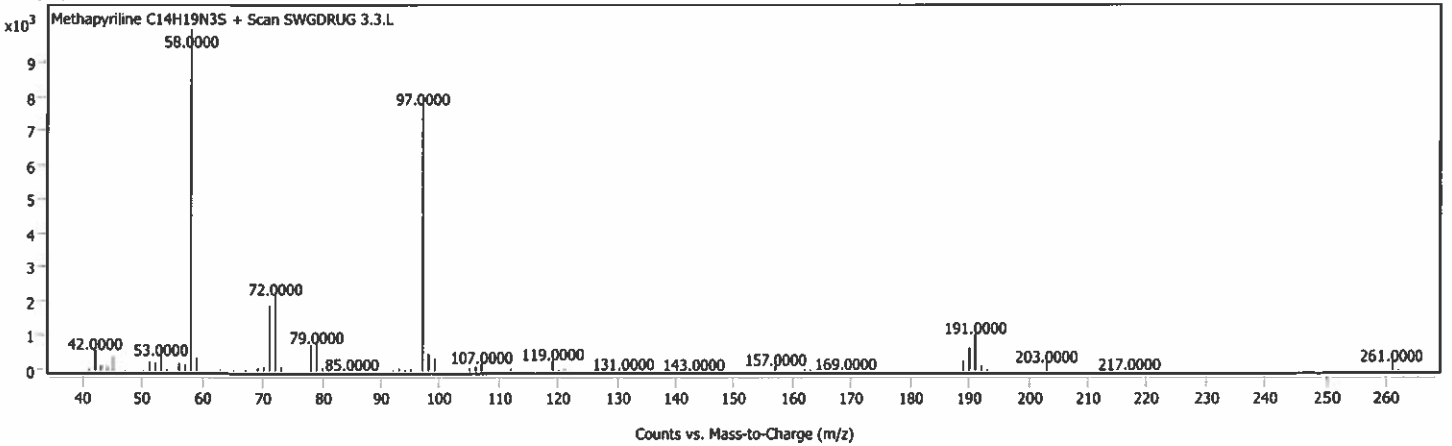
## Observed Spectrum



## Mirror Plot



## Library Spectrum



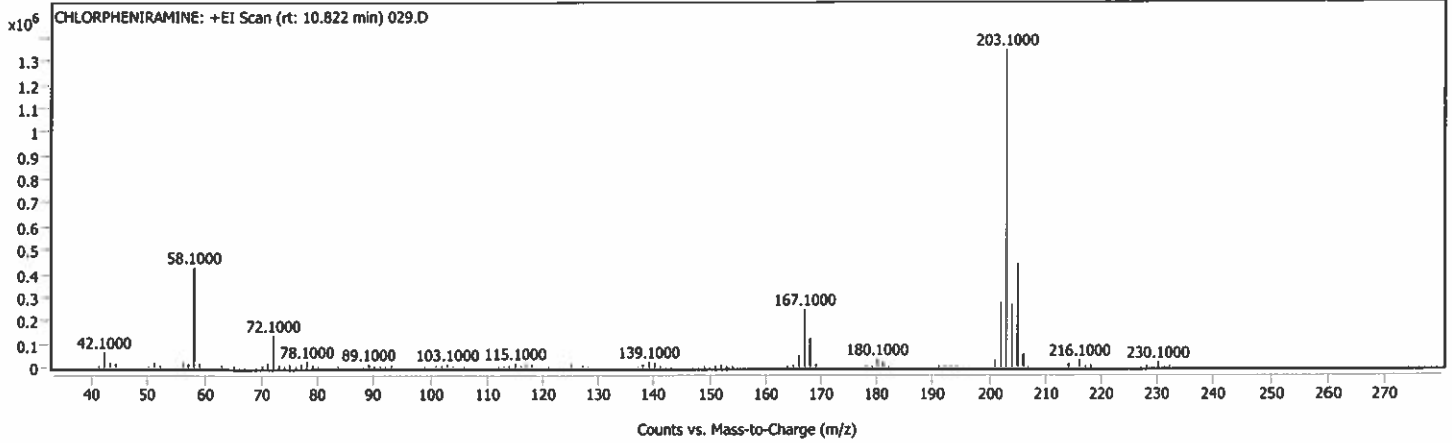
# Analysis Report

+ Scan (rt: 10.822 min)

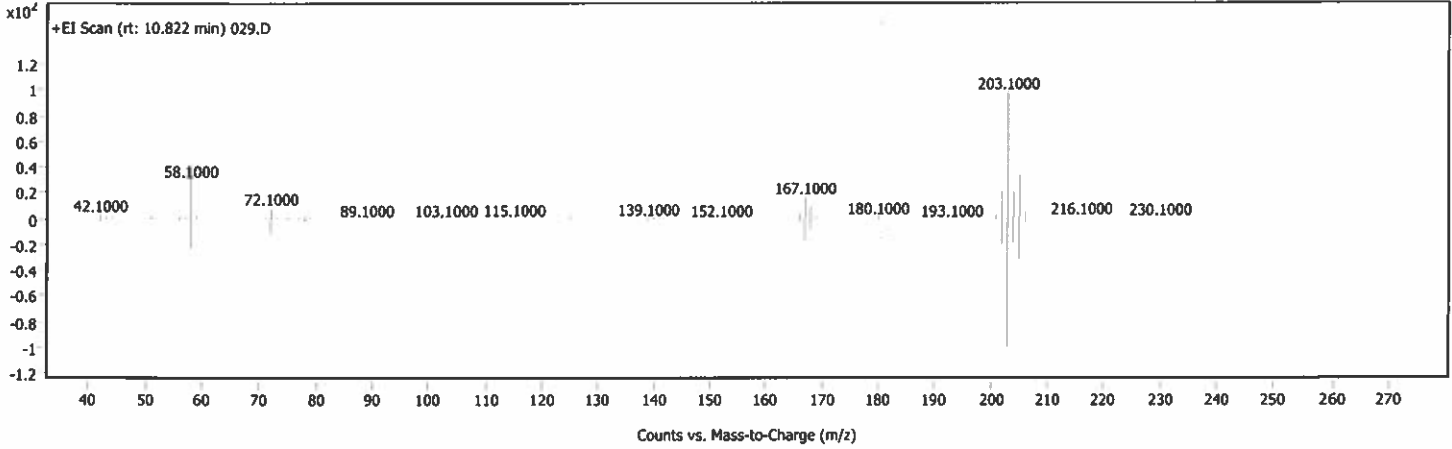
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	99.11	aafs.I	MATRIX 1 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

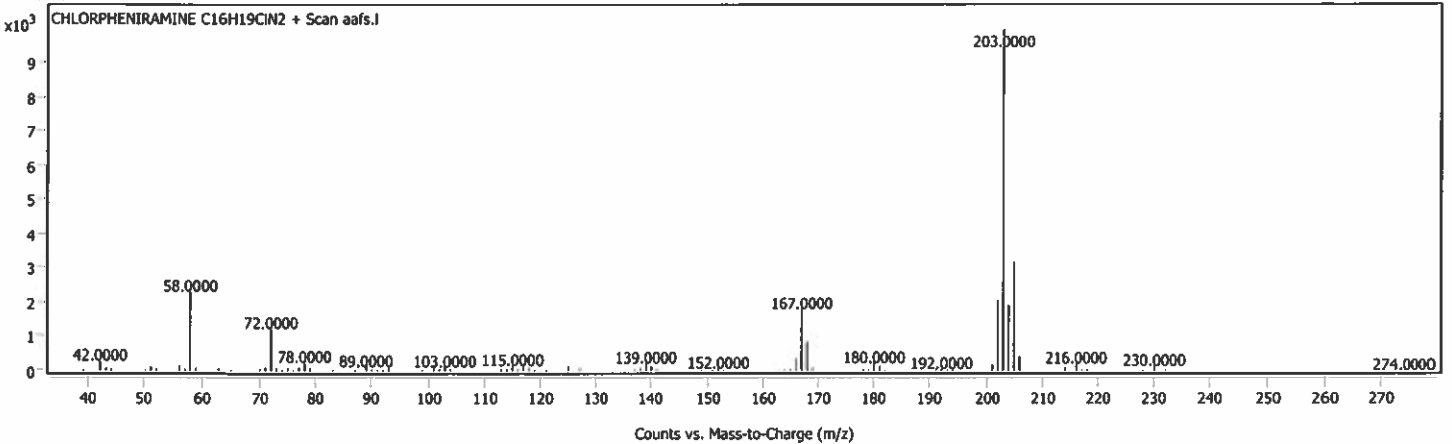
## Observed Spectrum



## Mirror Plot



## Library Spectrum



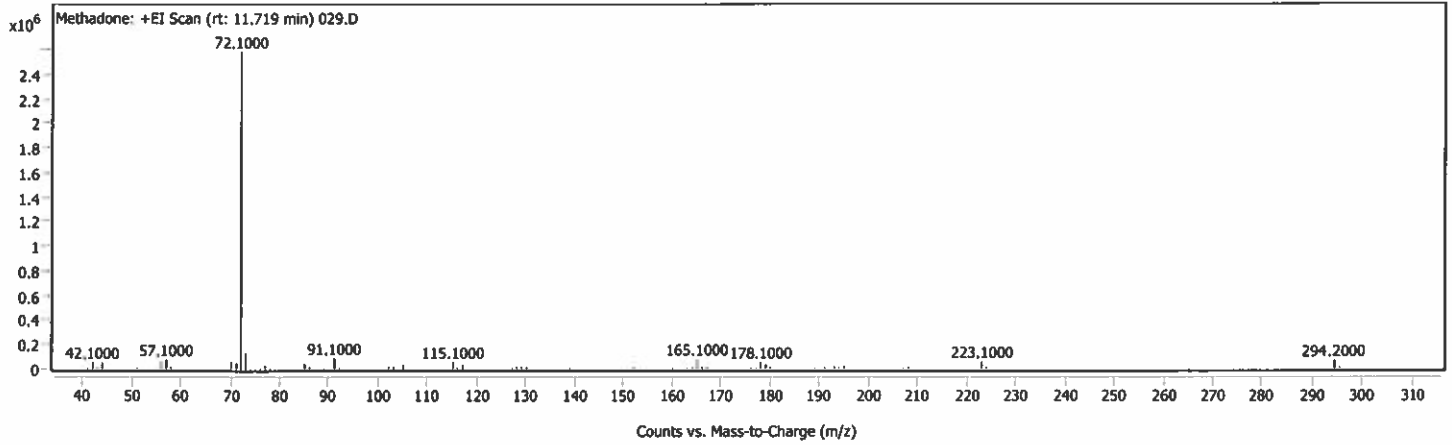
# Analysis Report

+ Scan (rt: 11.719 min)

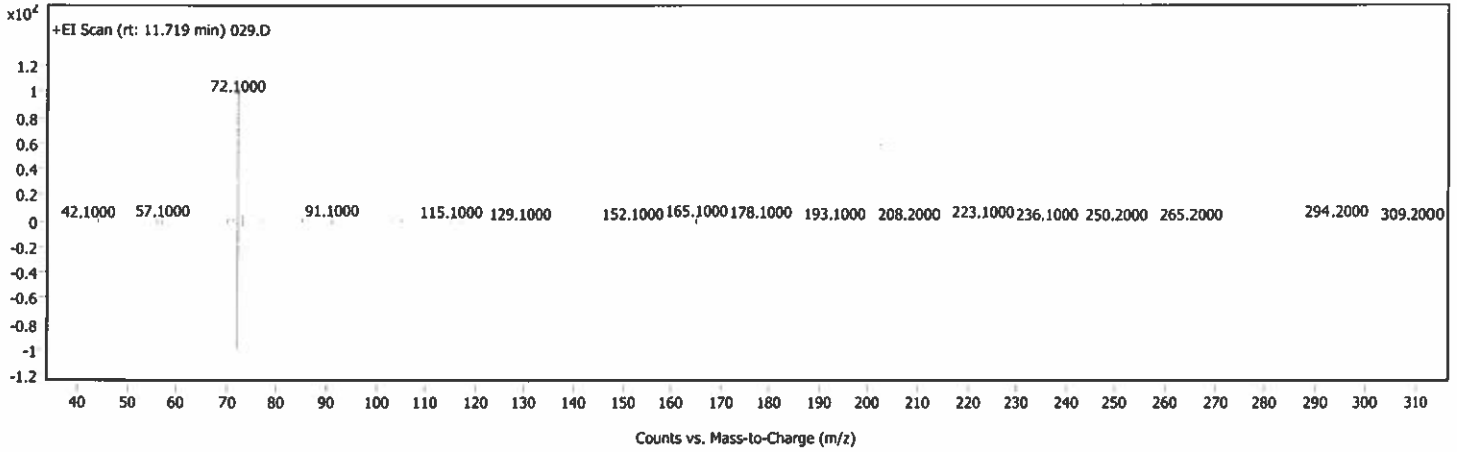
Methadone; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C21H27NO	97.38	SWGDRUG 3.3.L	MATRIX 1 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

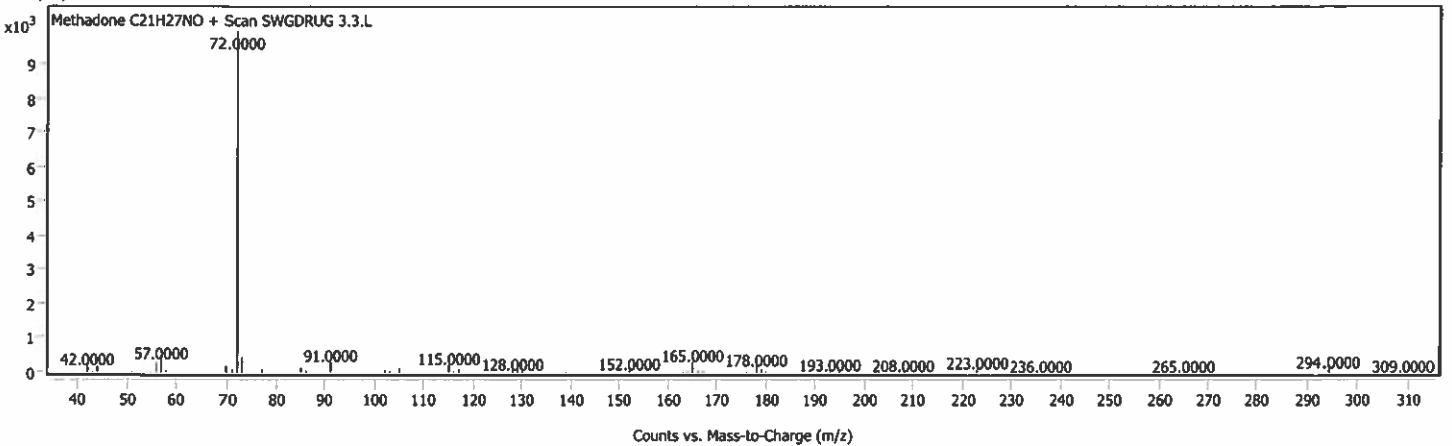
### Observed Spectrum



### Mirror Plot



### Library Spectrum



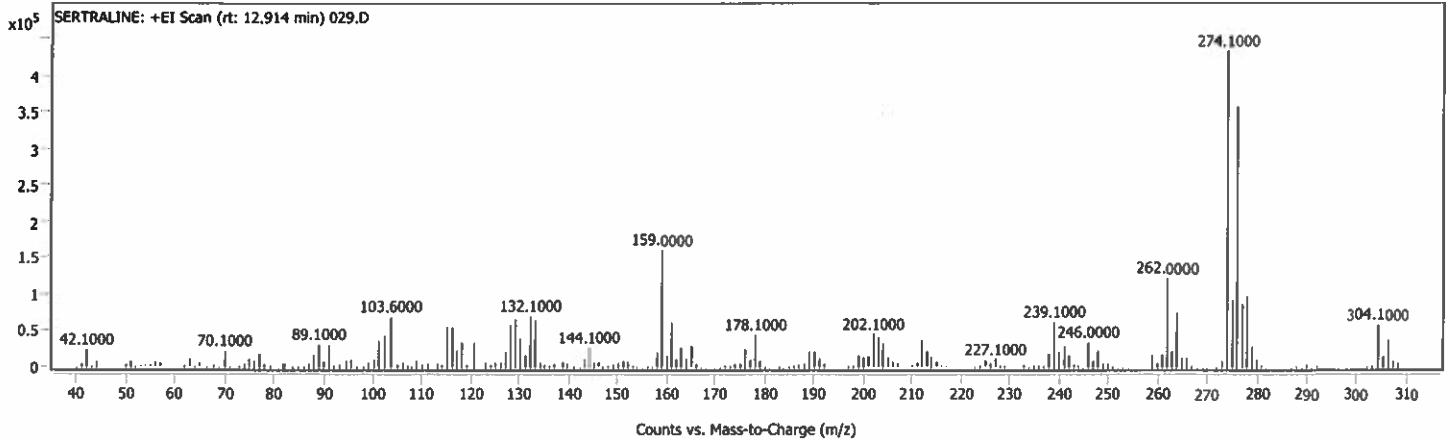
# Analysis Report

+ Scan (rt: 12.914 min)

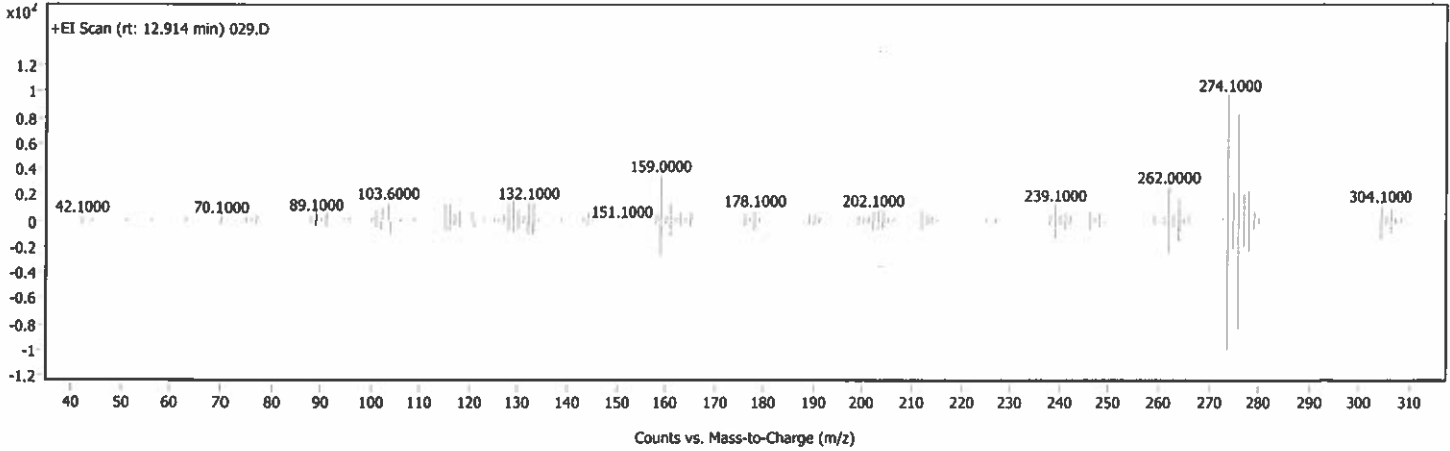
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	92.27	aafs.l	MATRIX 1 W/ IH CTL + XYLAZINE - BLD	KM/QC DATA IN XYLAZINE VALIDATION

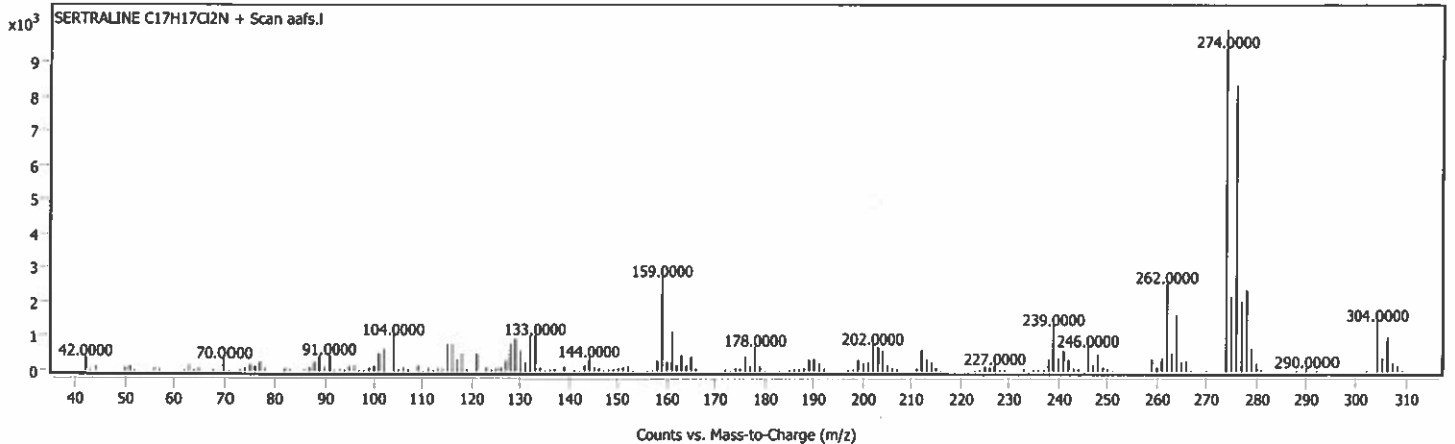
## Observed Spectrum



## Mirror Plot

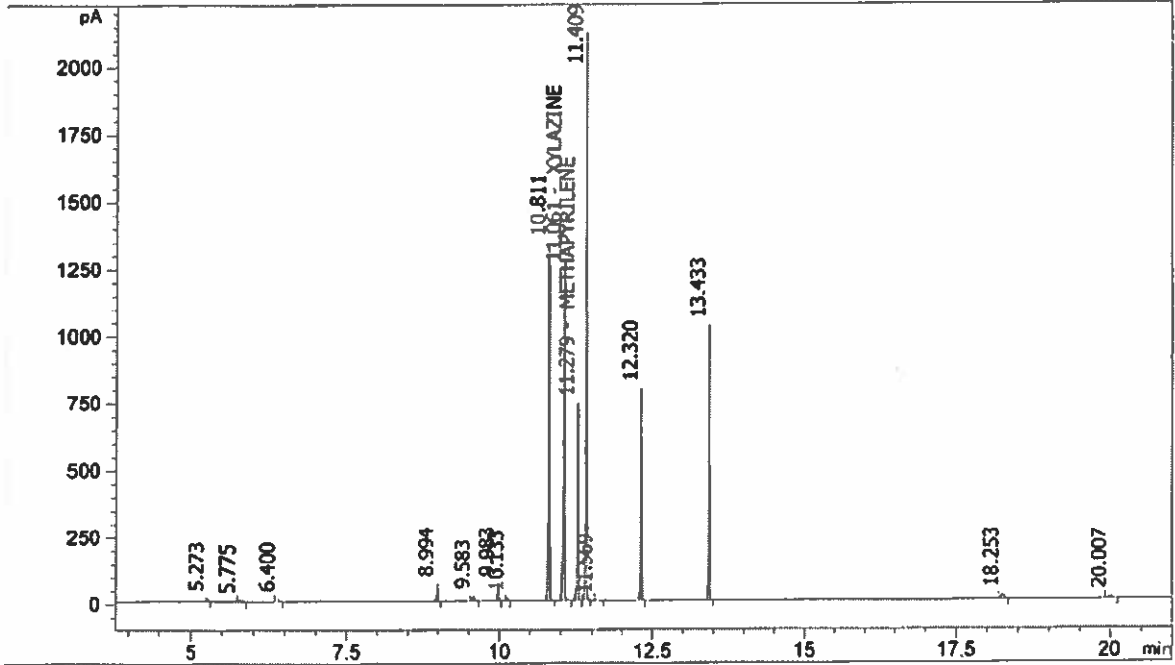


## Library Spectrum



Injection Date: 9/29/2021 2:43:44 AM Seq Line: 33  
 Sample Name: MATRIX 1 W/ IH CTL + -> Vial 34  
 Sample Info: Xylazine KM 11/17/22

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



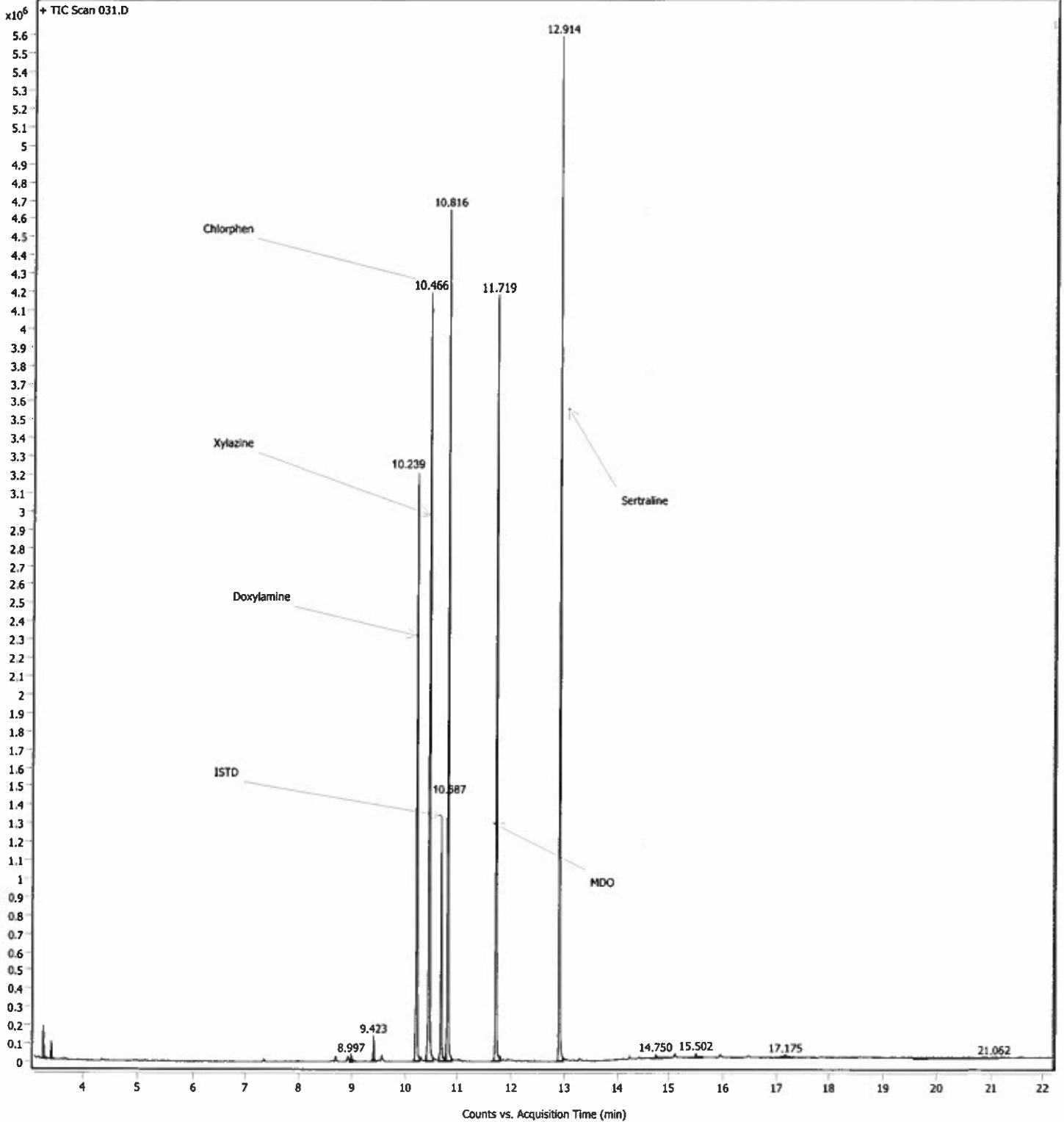
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.061	11.046	1873.649	1242.294	0.000000	XYLAZINE RRT - 0.9807
11.279	11.278	851.543	738.420	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 W/ 1H CTL + XYLAZINE - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\031.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 3:53:06 AM (UTC-04:00)
Position	60	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

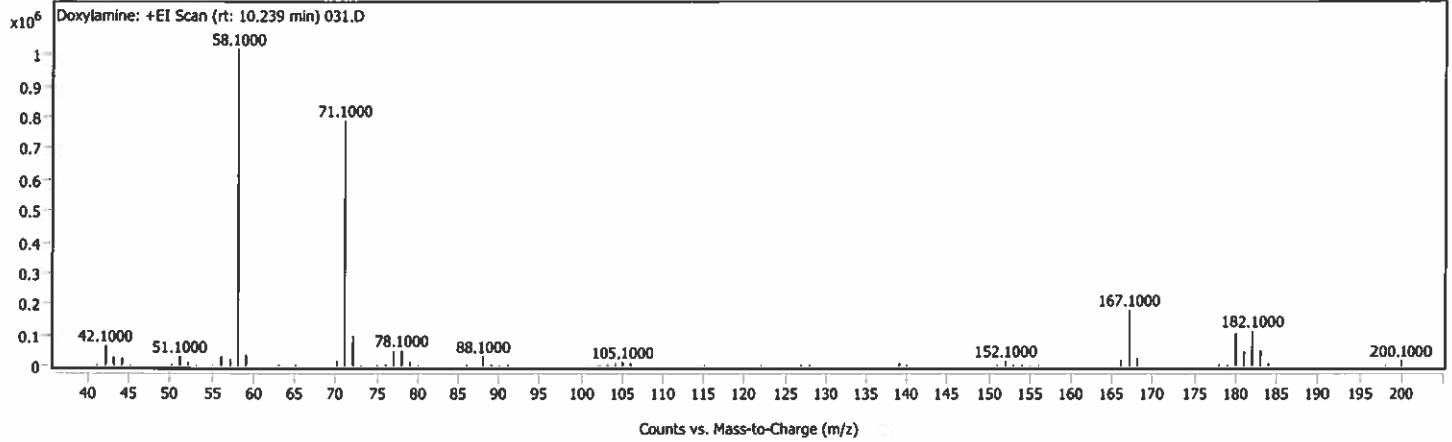
## Sample Spectra

+ Scan (rt: 10.239 min)

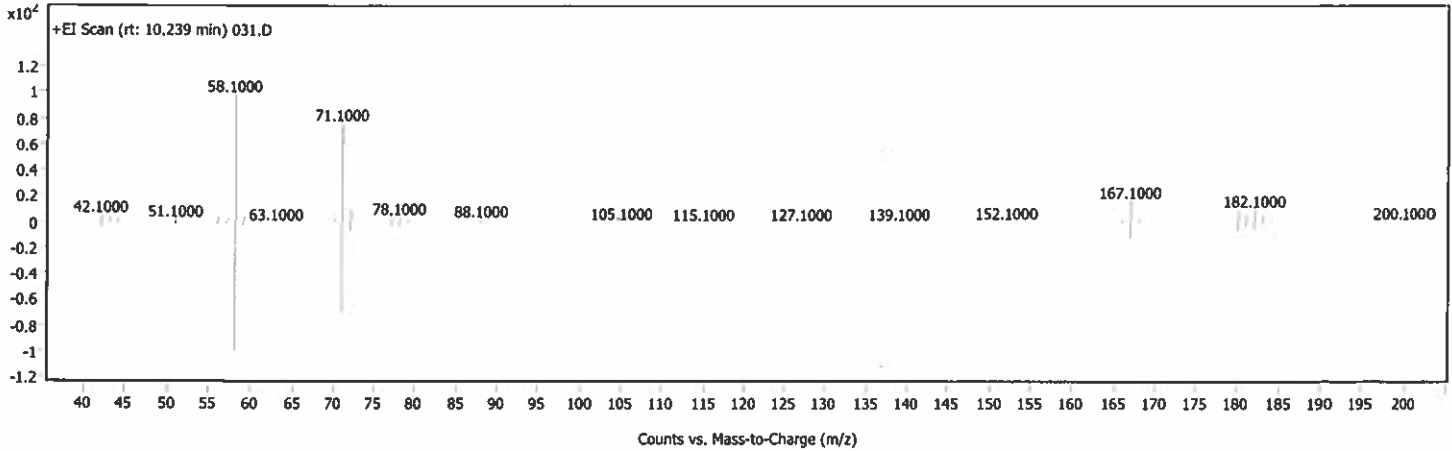
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.75	SWGDRUG 3.3.L	MATRIX 6 W/ IH CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

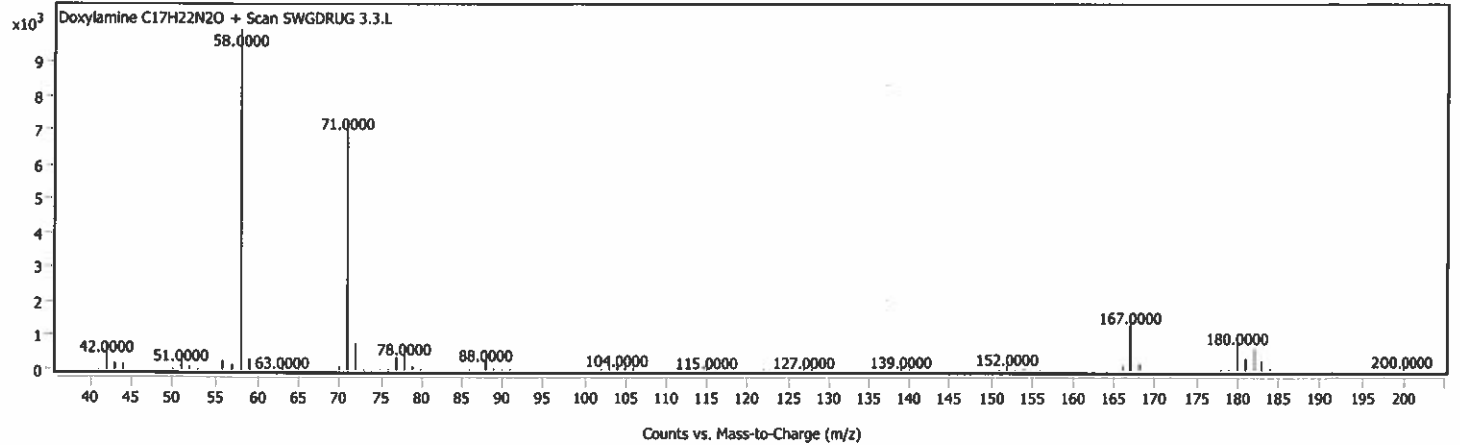
## Observed Spectrum



## Mirror Plot



## Library Spectrum





# Analysis Report

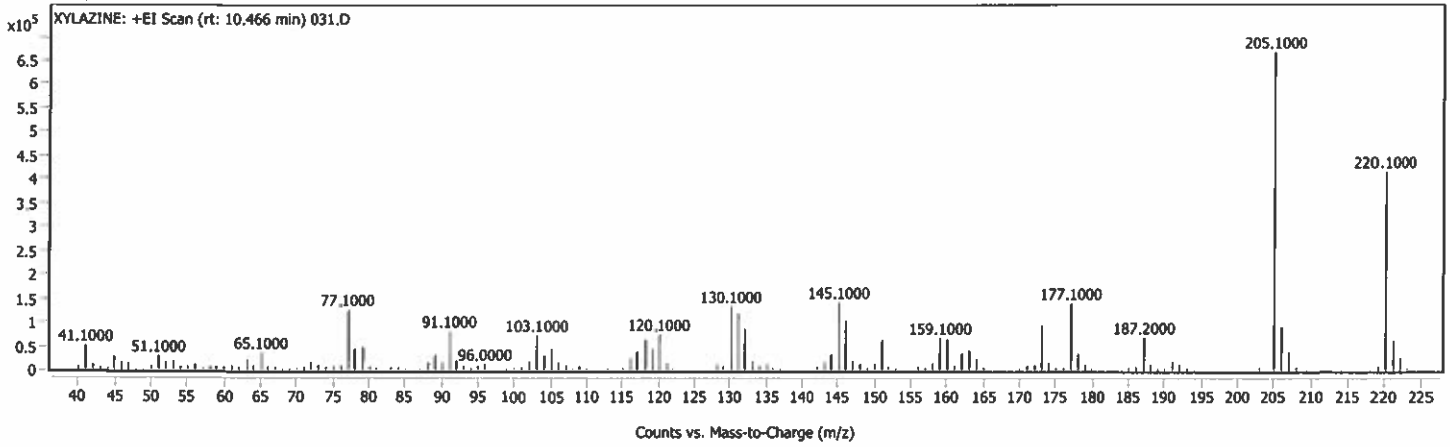


+ Scan (rt: 10.466 min)

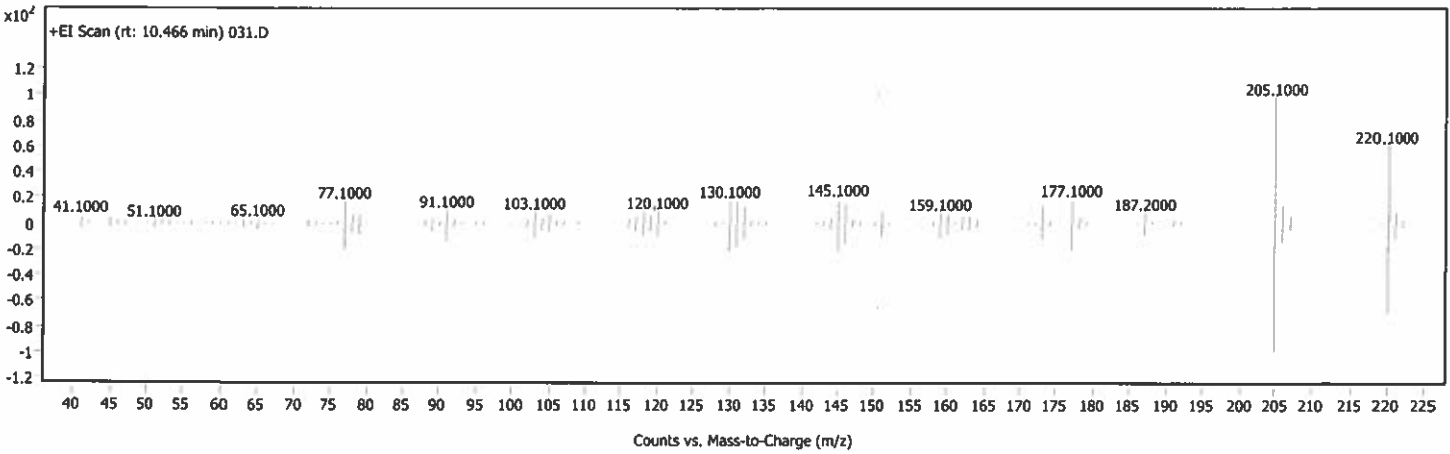
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.58	aafs.l	MATRIX 6 W/ IH CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

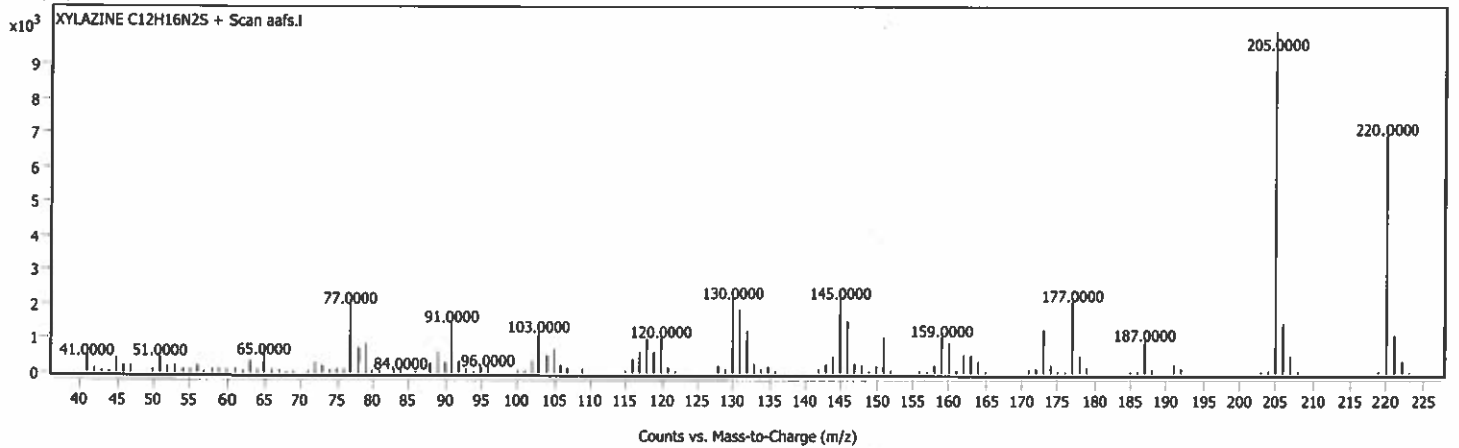
## Observed Spectrum



## Mirror Plot



## Library Spectrum



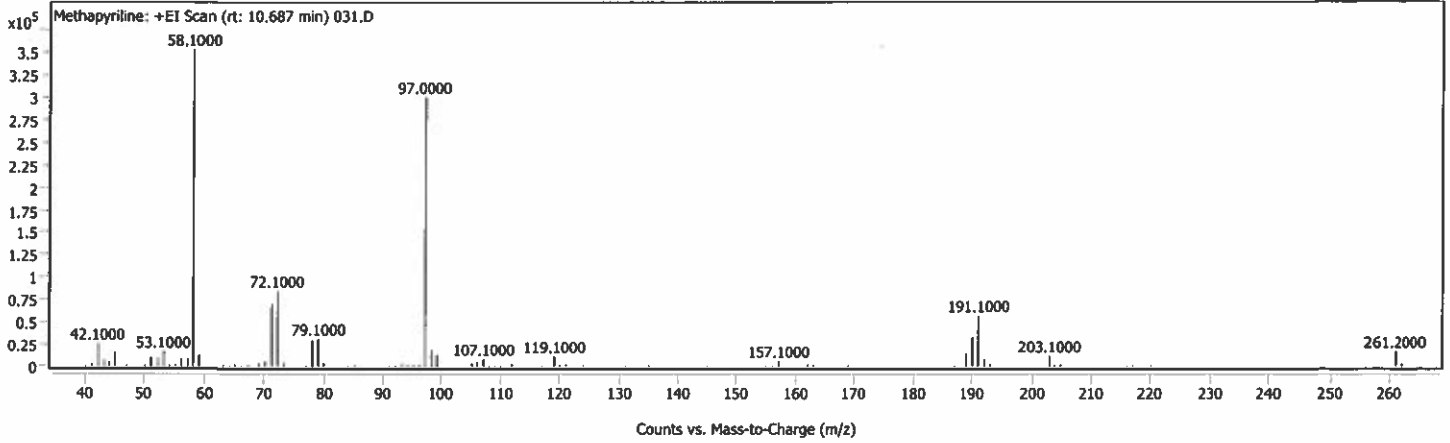
# Analysis Report

+ Scan (rt: 10.687 min)

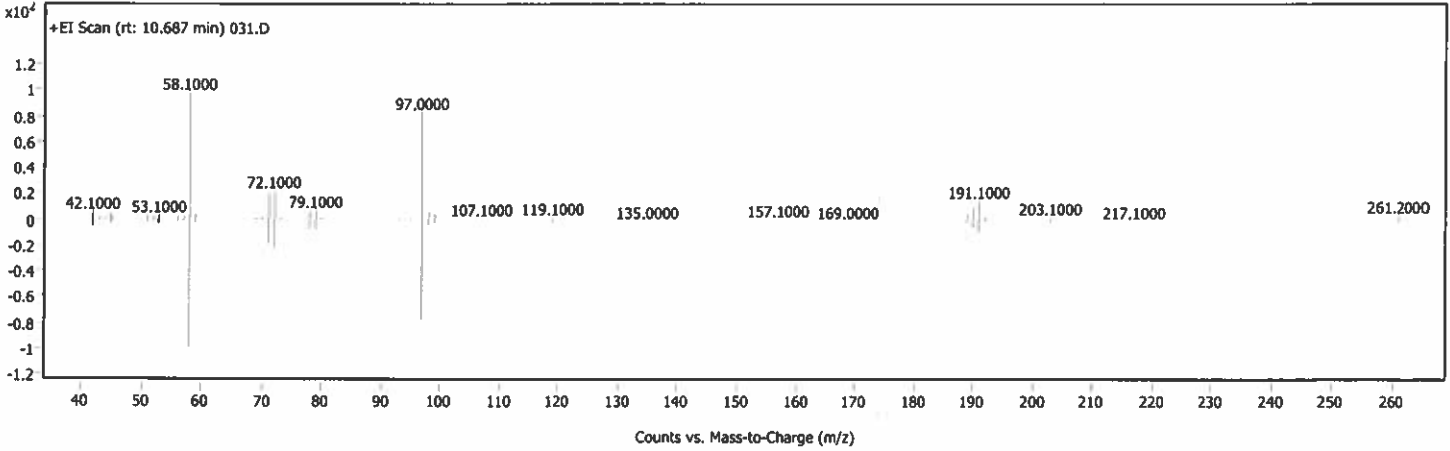
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.64	SWGDRUG 3.3.L	MATRIX 6 W/ IH CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

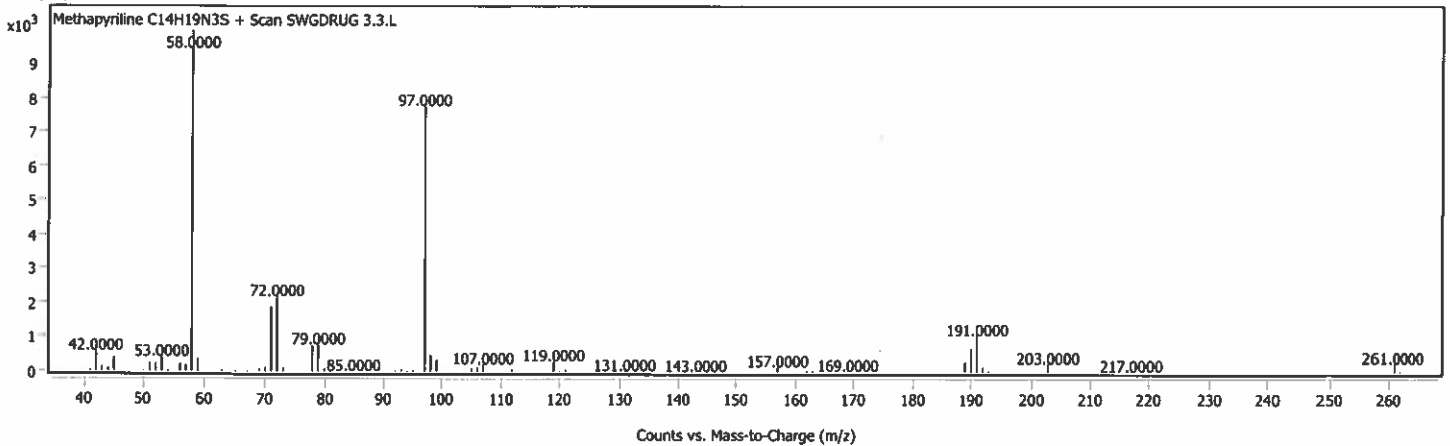
### Observed Spectrum



### Mirror Plot



### Library Spectrum



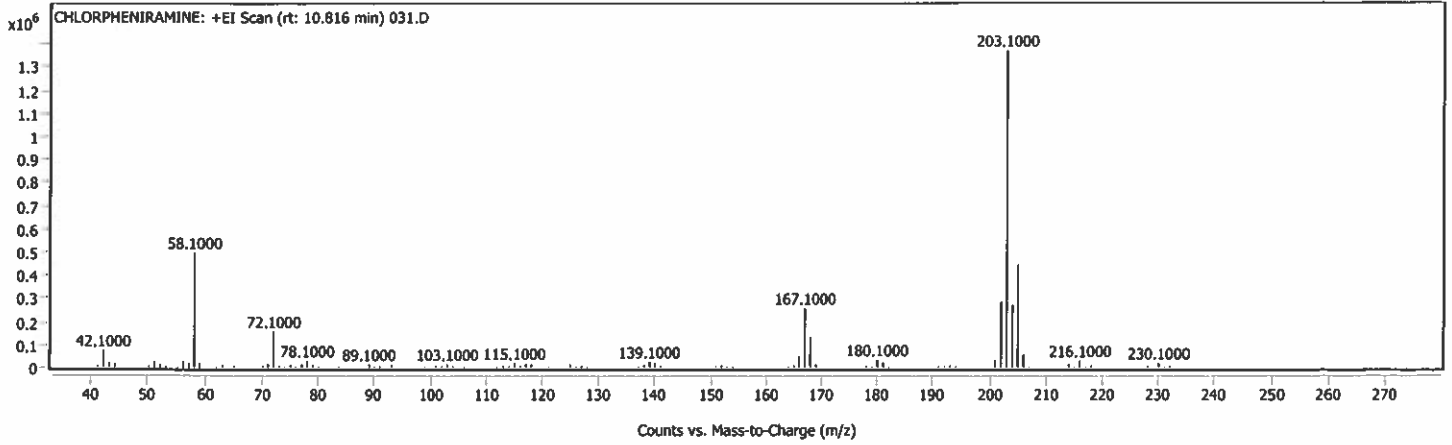
# Analysis Report

+ Scan (rt: 10.816 min)

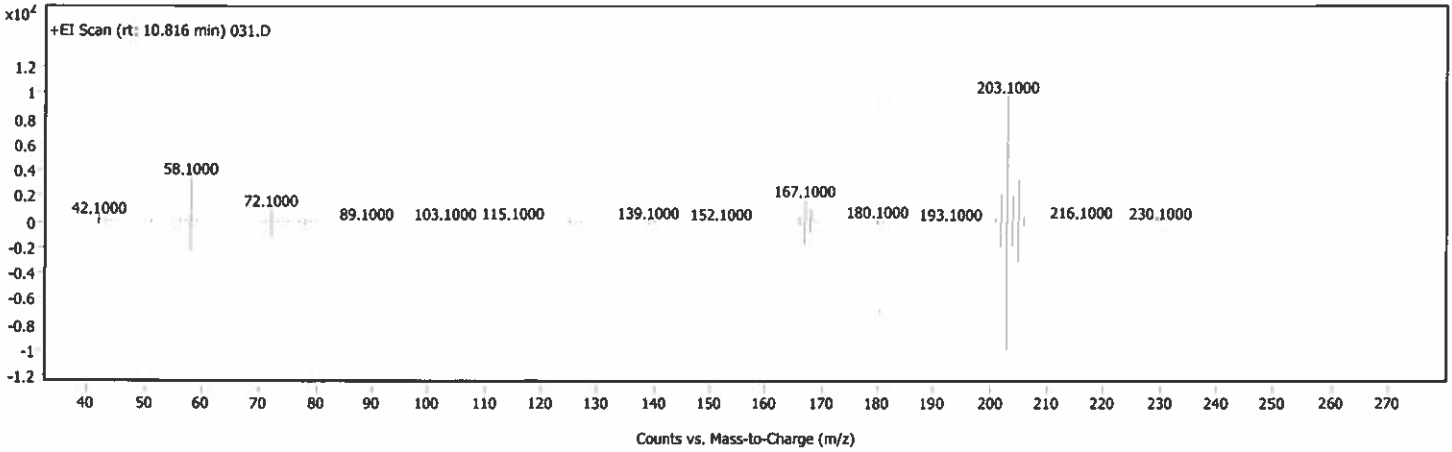
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	99.02	aafs.I	MATRIX 6 W/ IH CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

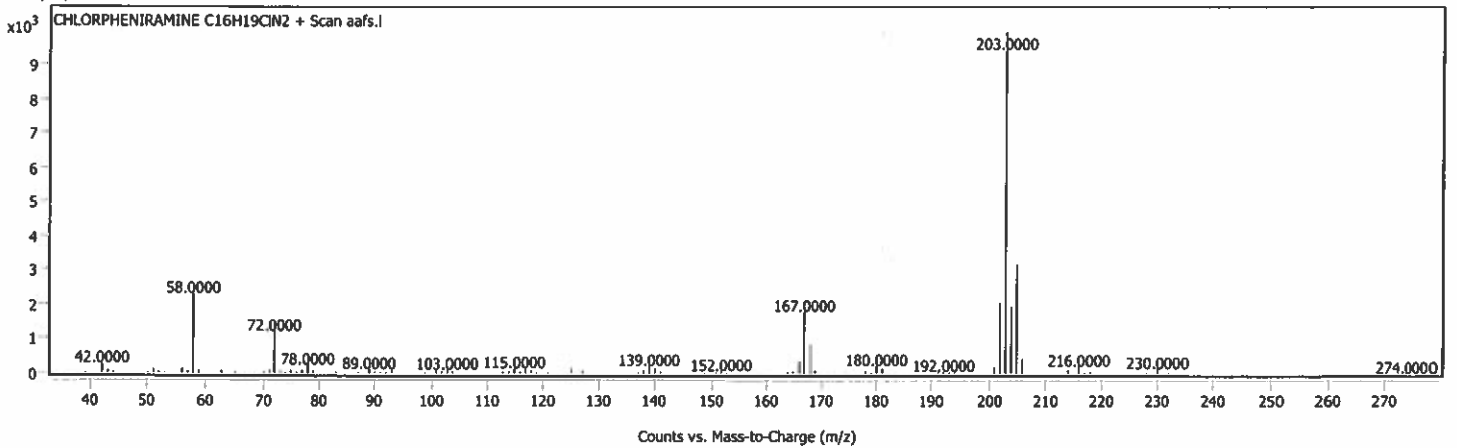
## Observed Spectrum



## Mirror Plot



## Library Spectrum



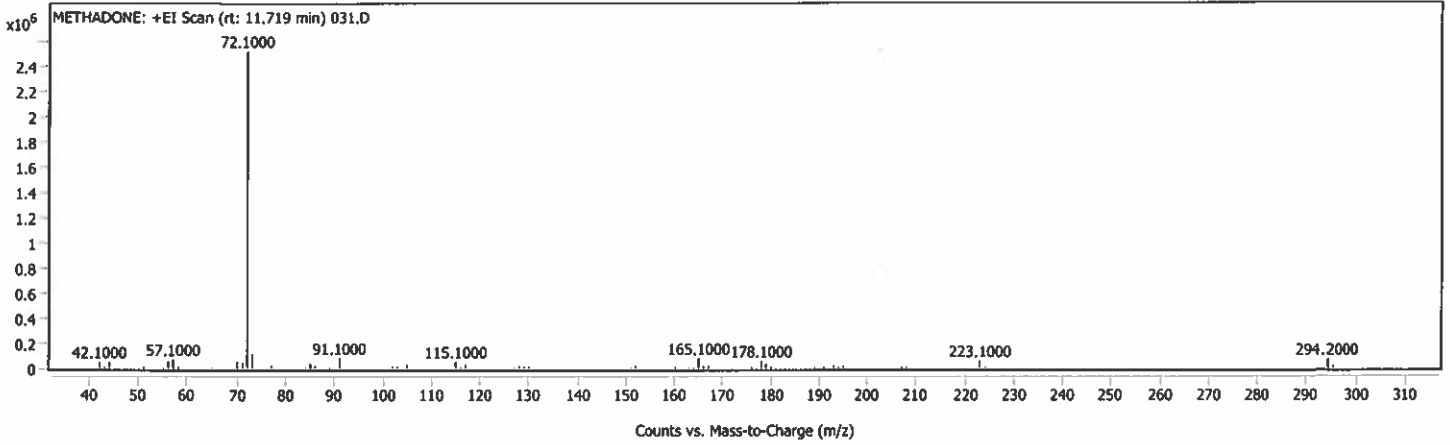
# Analysis Report

+ Scan (rt: 11.719 min)

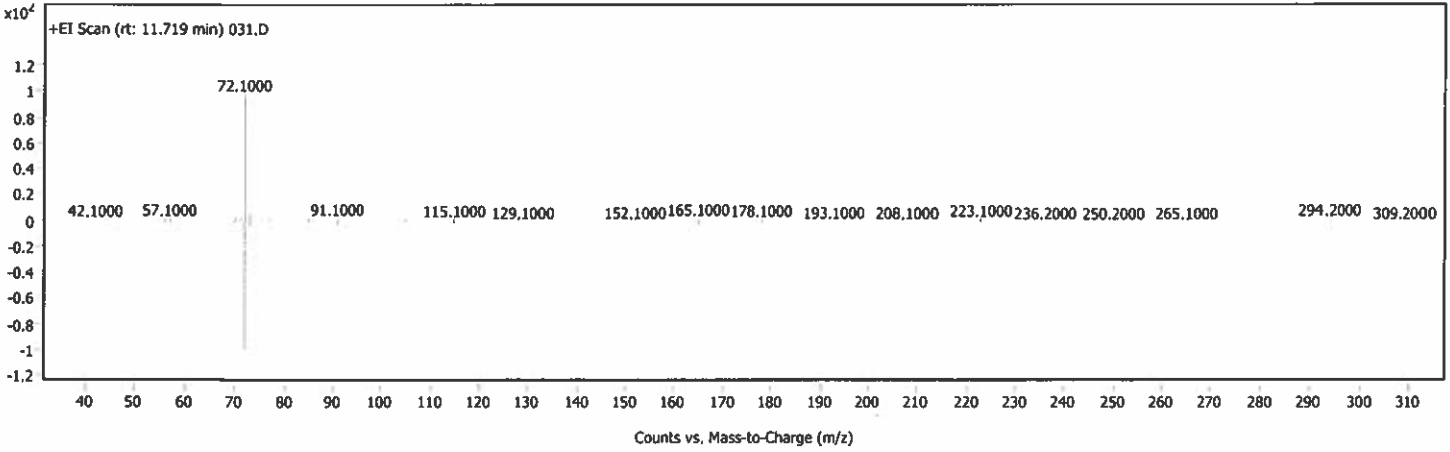
METHADONE; C<sub>21</sub>H<sub>27</sub>NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
METHADONE	C <sub>21</sub> H <sub>27</sub> NO	97.07	aafs.l	MATRIX 6 W/ IH CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

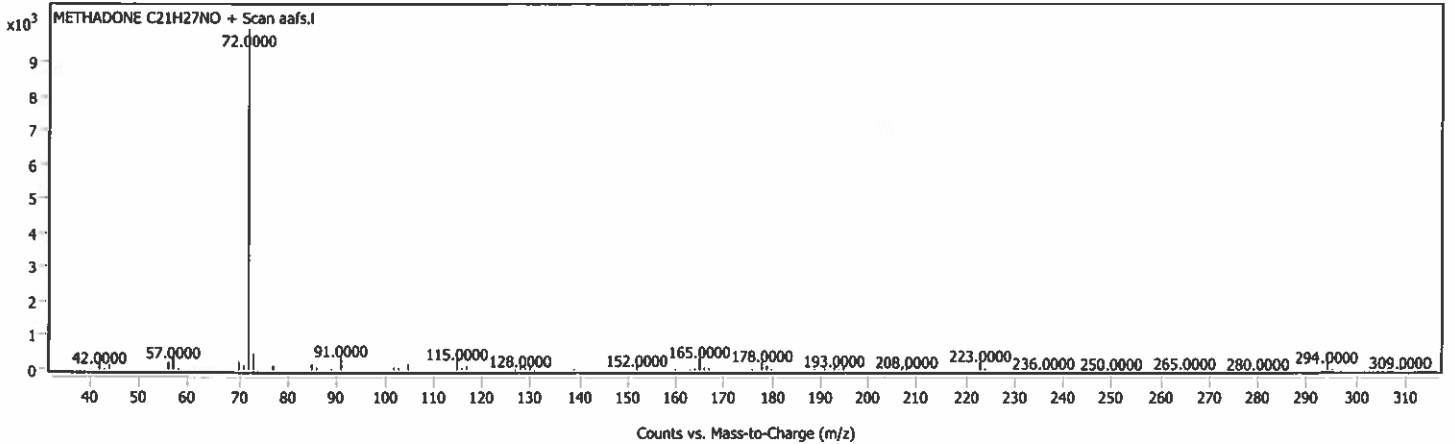
### Observed Spectrum



### Mirror Plot



### Library Spectrum



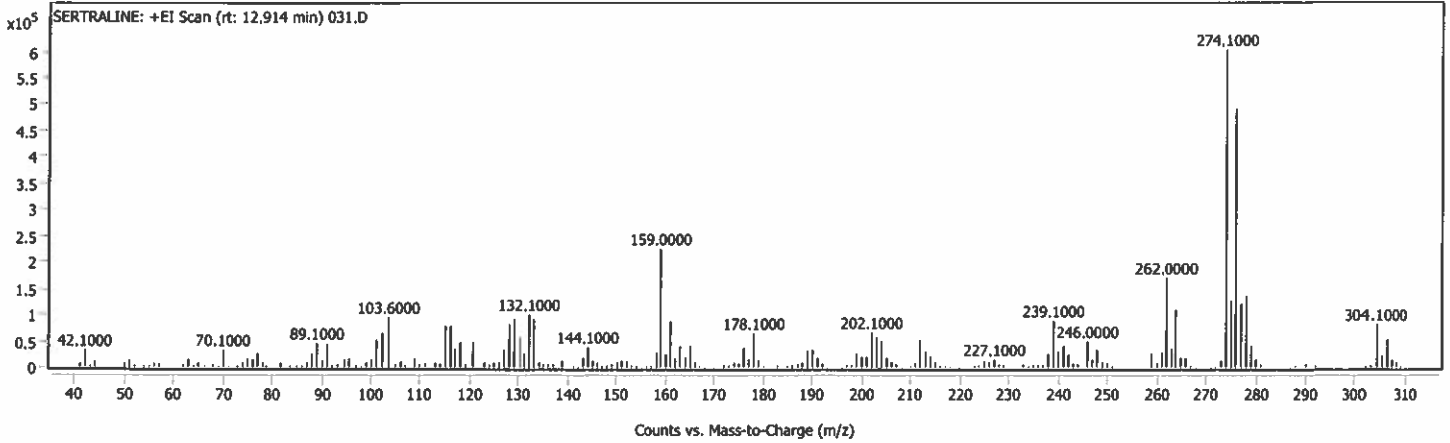
# Analysis Report

+ Scan (rt: 12.914 min)

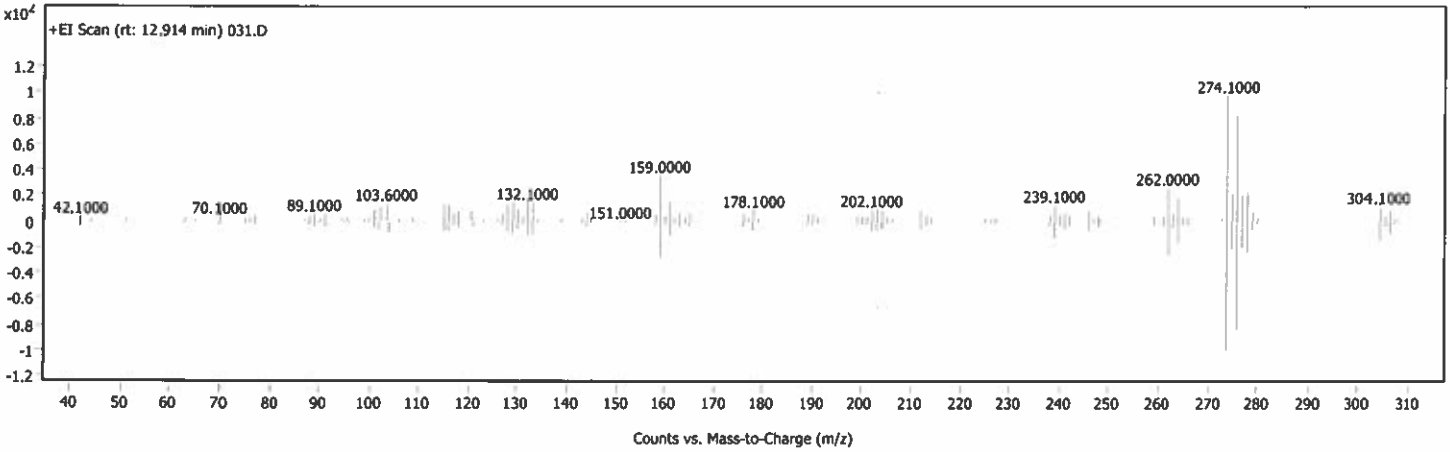
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	91.95	aafs.l	MATRIX 6 W/ IH CTL + XYLAZINE - URN	KM/QC DATA IN XYLAZINE VALIDATION

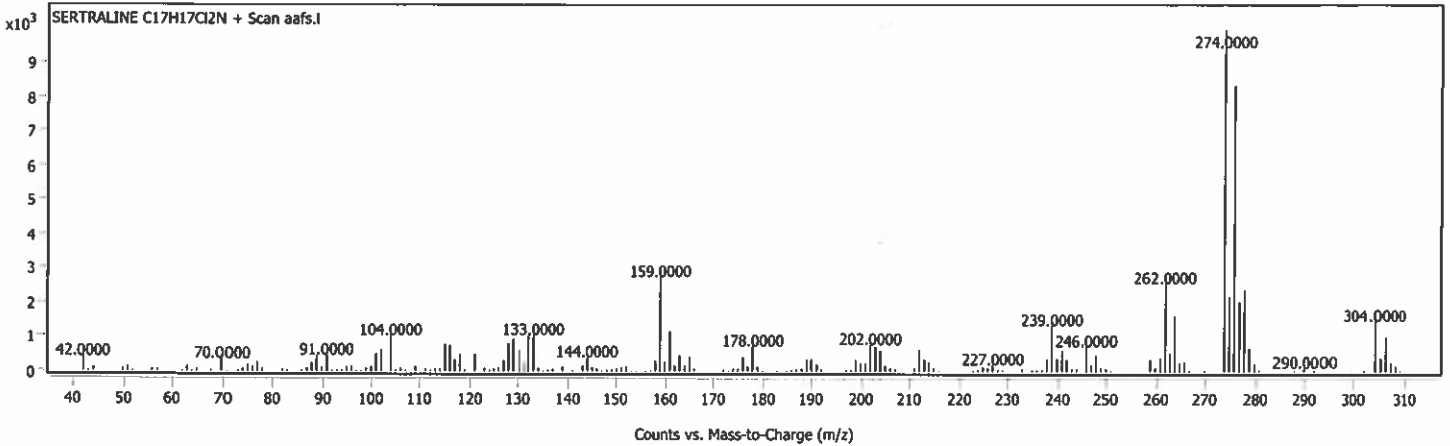
## Observed Spectrum



## Mirror Plot

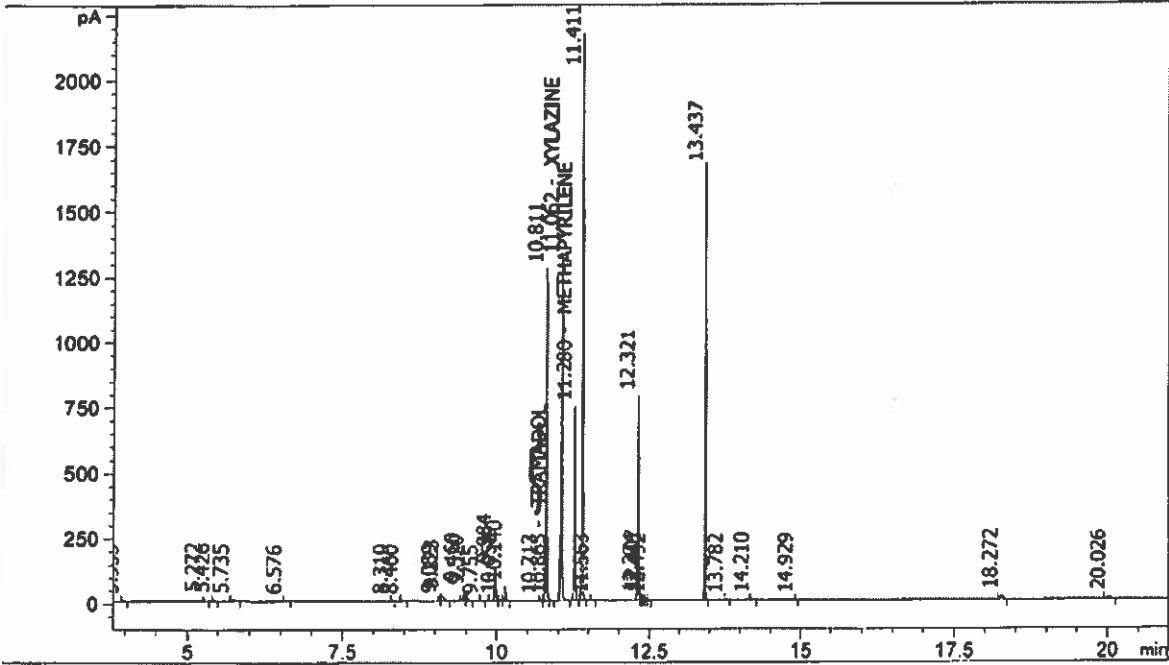


## Library Spectrum



Injection Date: 9/29/2021 3:32:25 AM Seq Line: 35  
 Sample Name: MATRIX 6 W/ IH CTL + -> Vial 36  
 Sample Info: xylazine 11/7/22

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



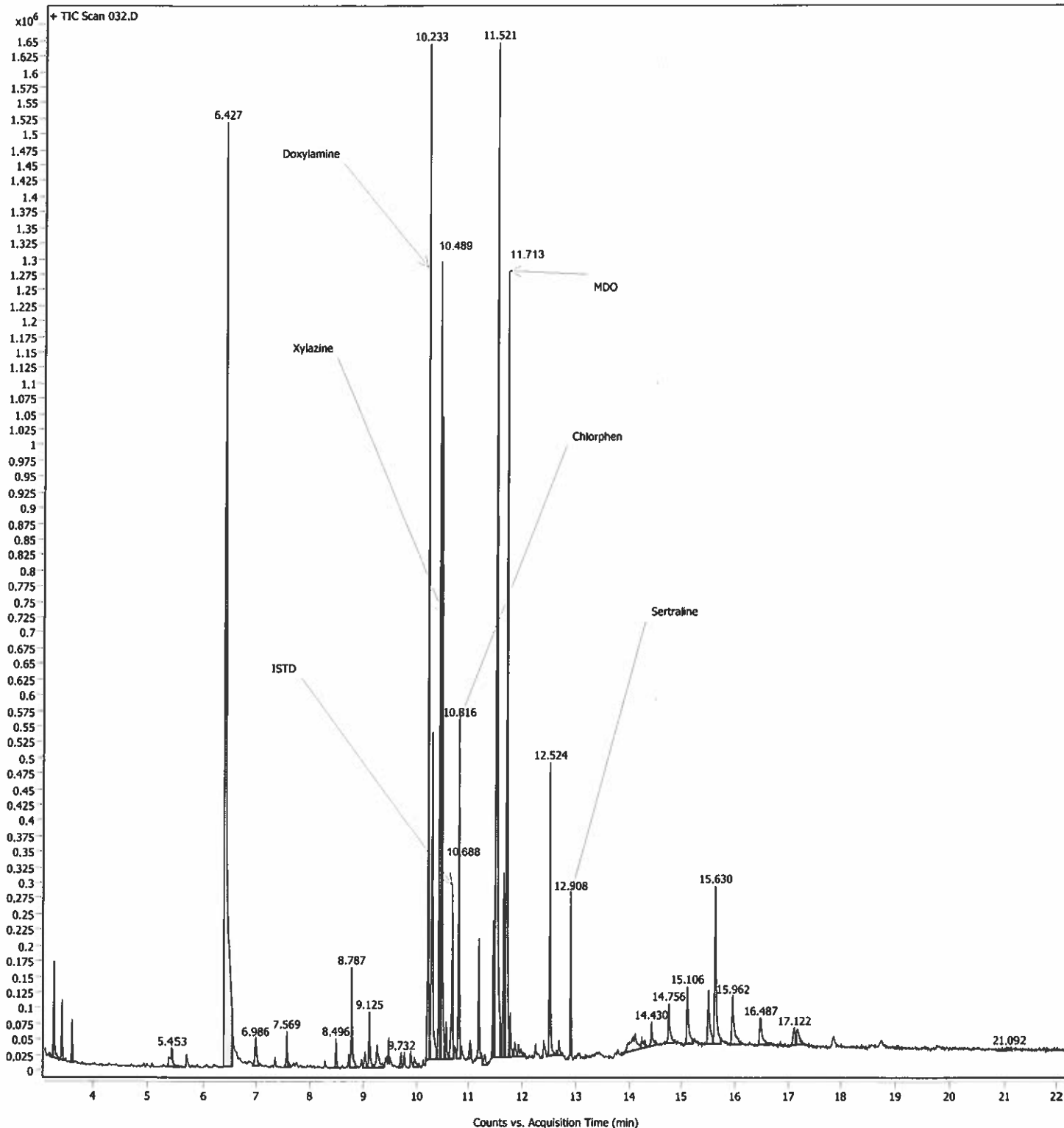
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
10.865	11.033	5.261	2.911	0.00000	TRAMADOL ND
11.062	11.046	1934.194	1307.255	0.00000	XYLAZINE RRT - 0.9807
11.280	11.278	869.681	740.347	1.00000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 10 W/ IH CTL + XYLAZINE Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\032.D
Instrument	#3 - Enhanced	Acq. Time (Local) 9/29/2021 4:18:53 AM (UTC-04:00)
Position	61	Method Path (Acq) C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION	

## Sample Chromatograms



# Analysis Report

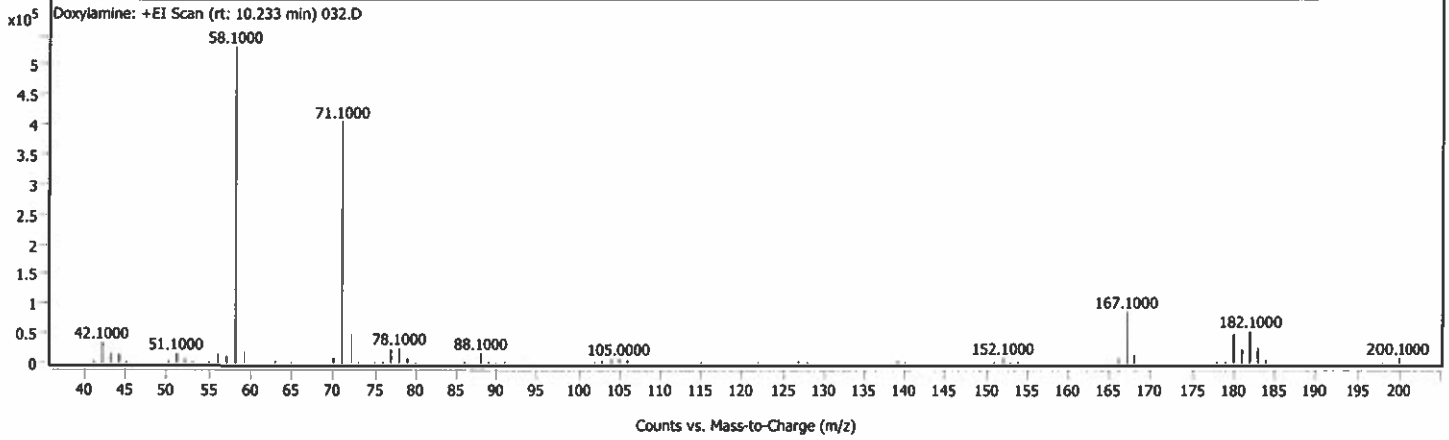
## Sample Spectra

+ Scan (rt: 10.233 min)

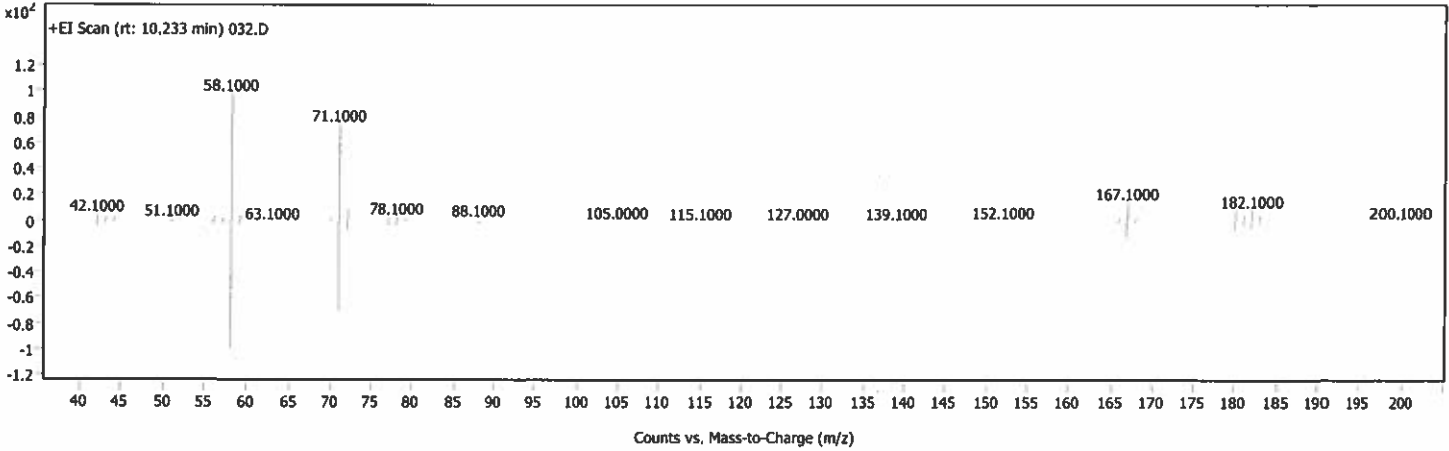
Doxylamine; C17H22N2O

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Doxylamine	C17H22N2O	98.89	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

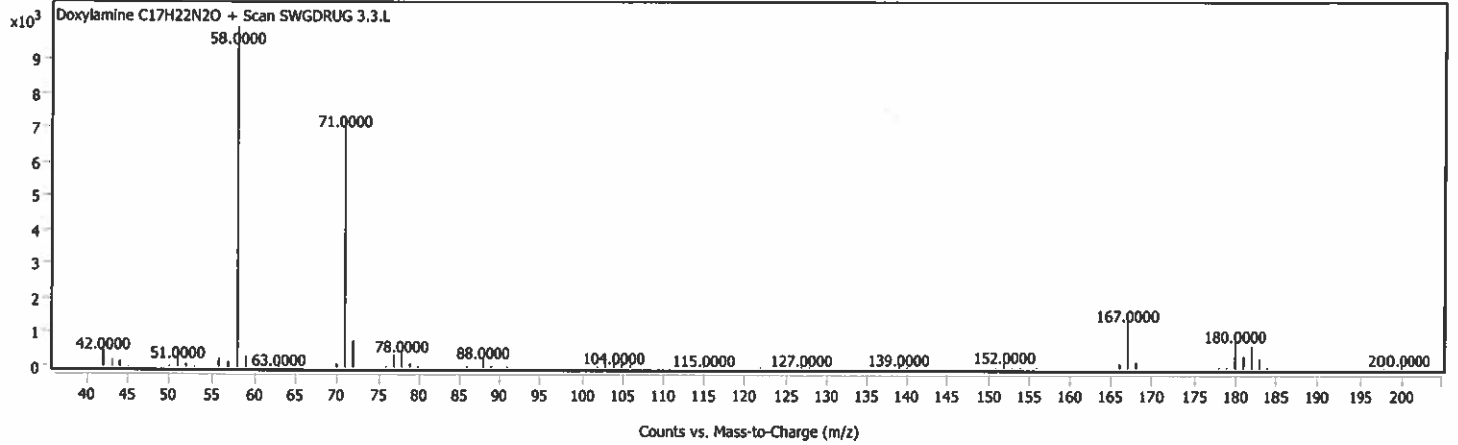
## Observed Spectrum



## Mirror Plot



## Library Spectrum





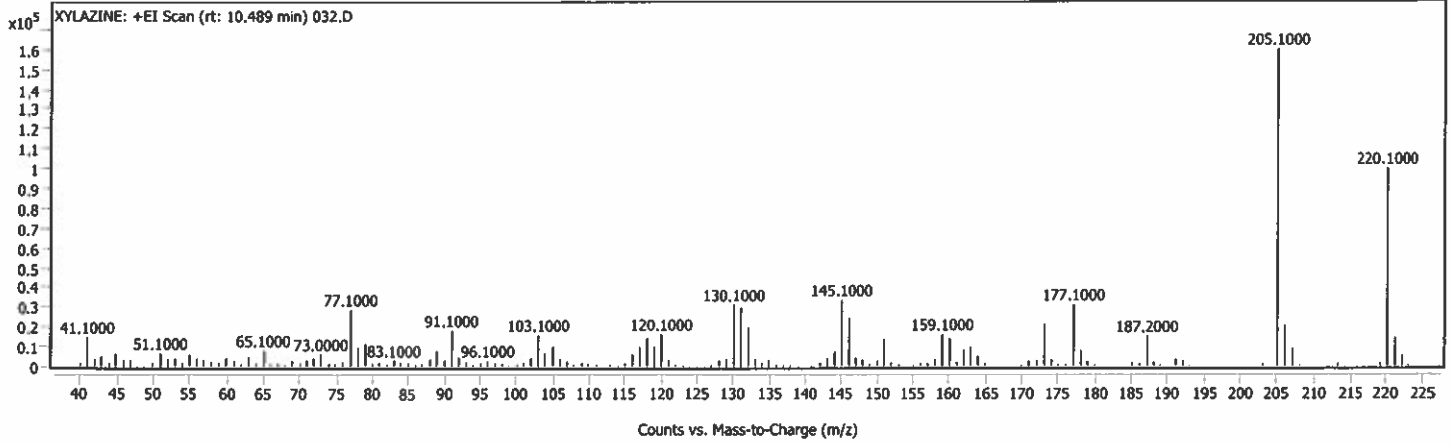
# Analysis Report

+ Scan (rt: 10.489 min)

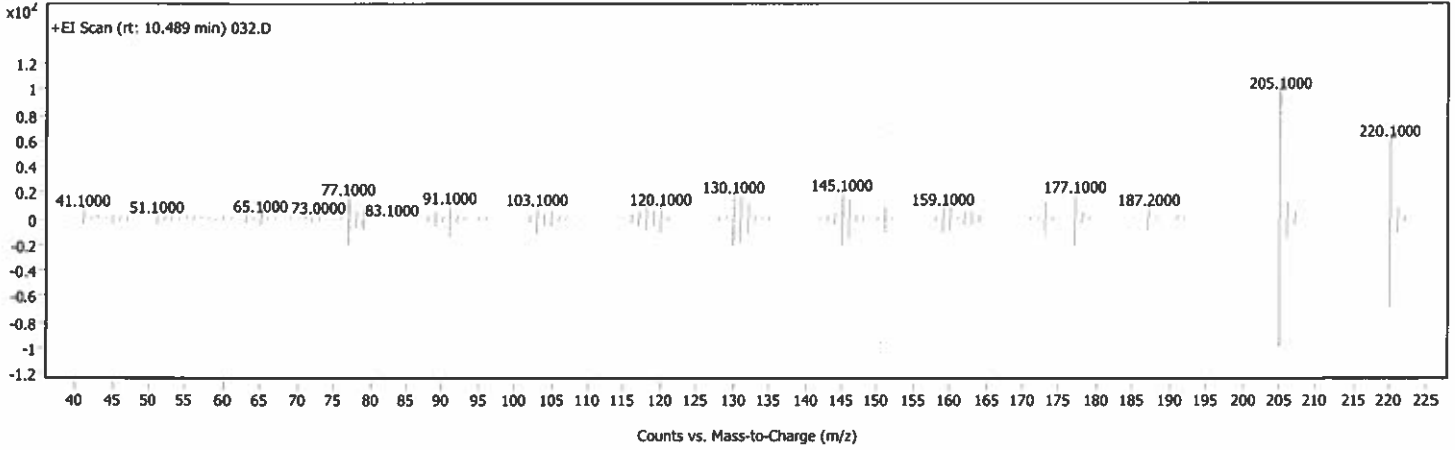
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	96.57	aafs.l	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

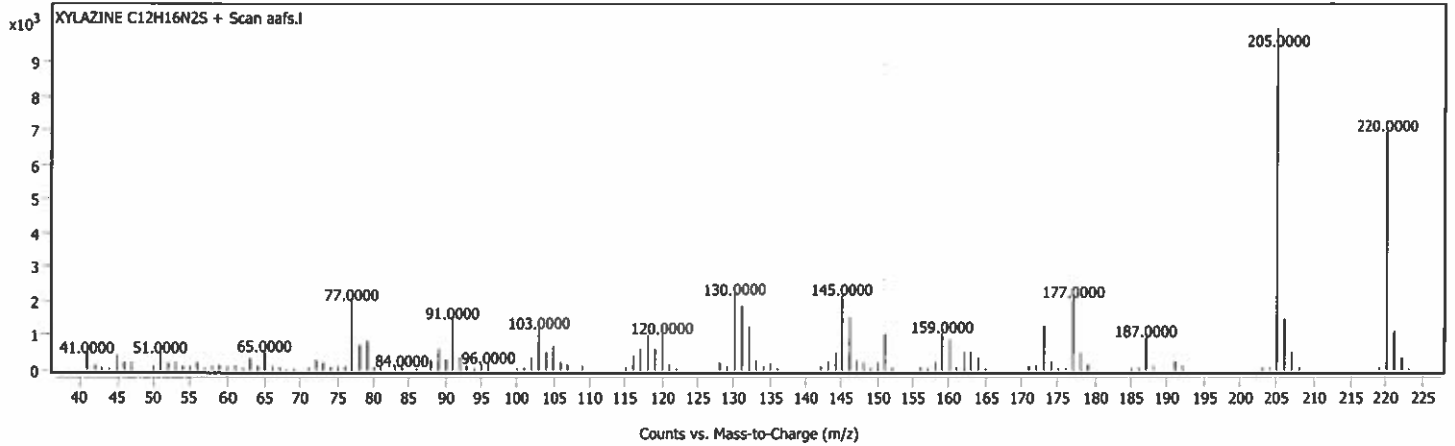
## Observed Spectrum



## Mirror Plot



## Library Spectrum



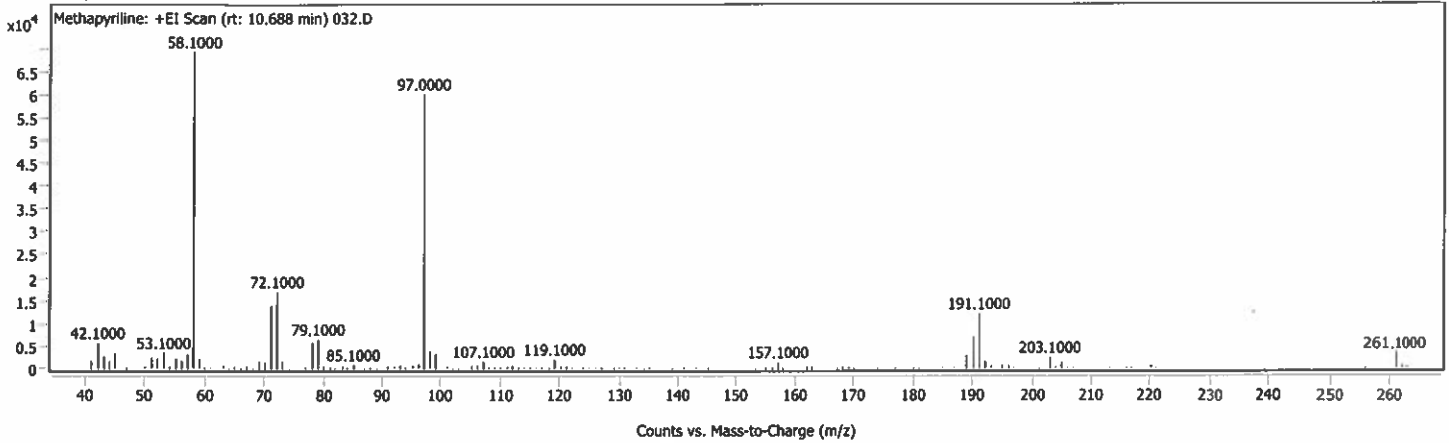
# Analysis Report

+ Scan (rt: 10.688 min)

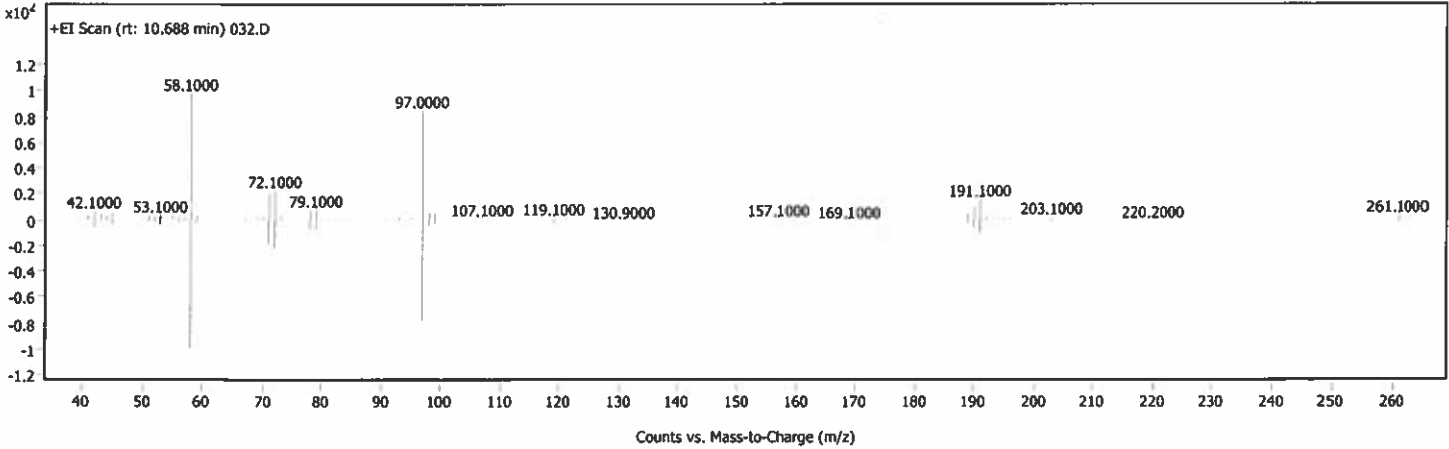
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	93.41	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

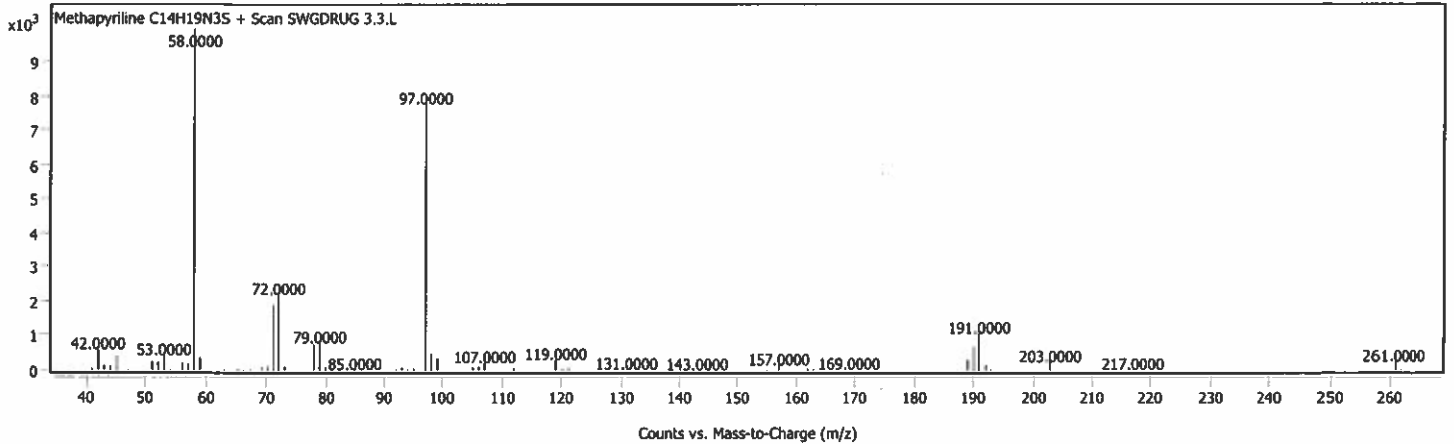
### Observed Spectrum



### Mirror Plot



### Library Spectrum



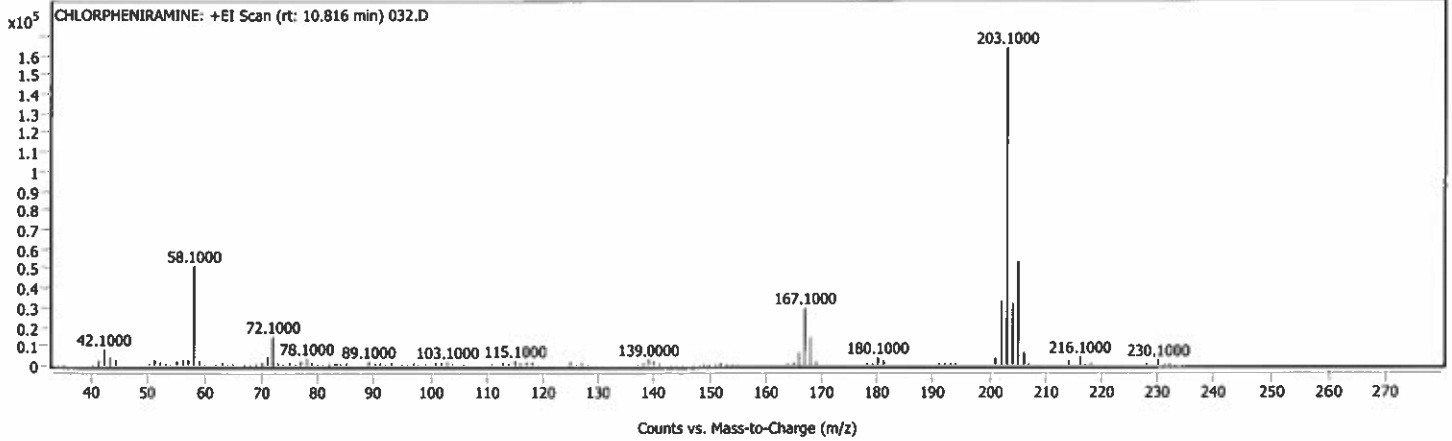
# Analysis Report

+ Scan (rt: 10.816 min)

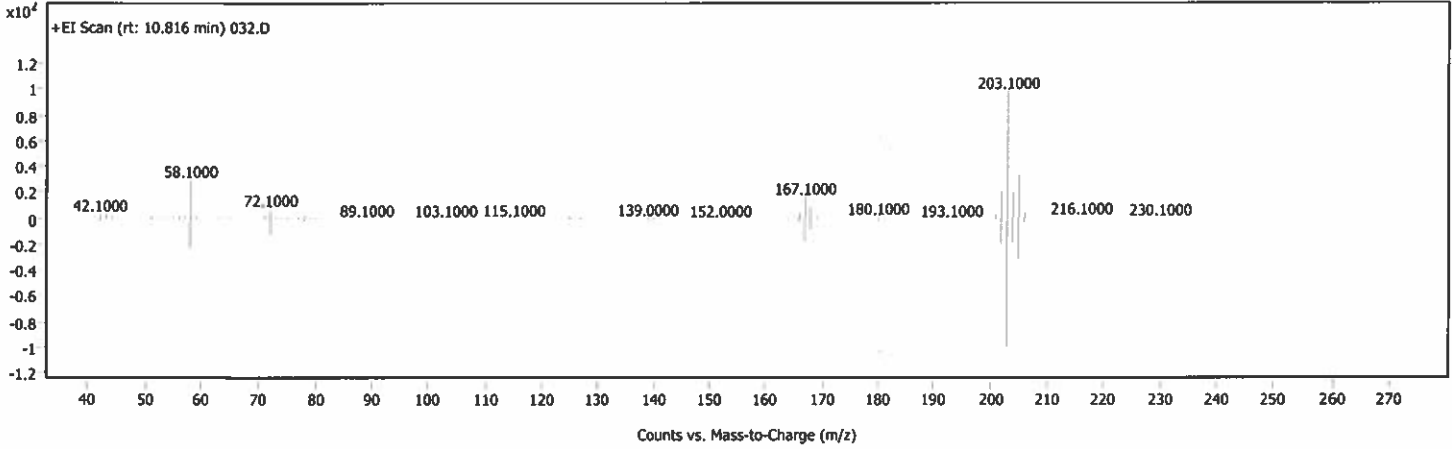
CHLORPHENIRAMINE; C16H19ClN2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CHLORPHENIRAMINE	C16H19ClN2	97.83	aafs.l	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

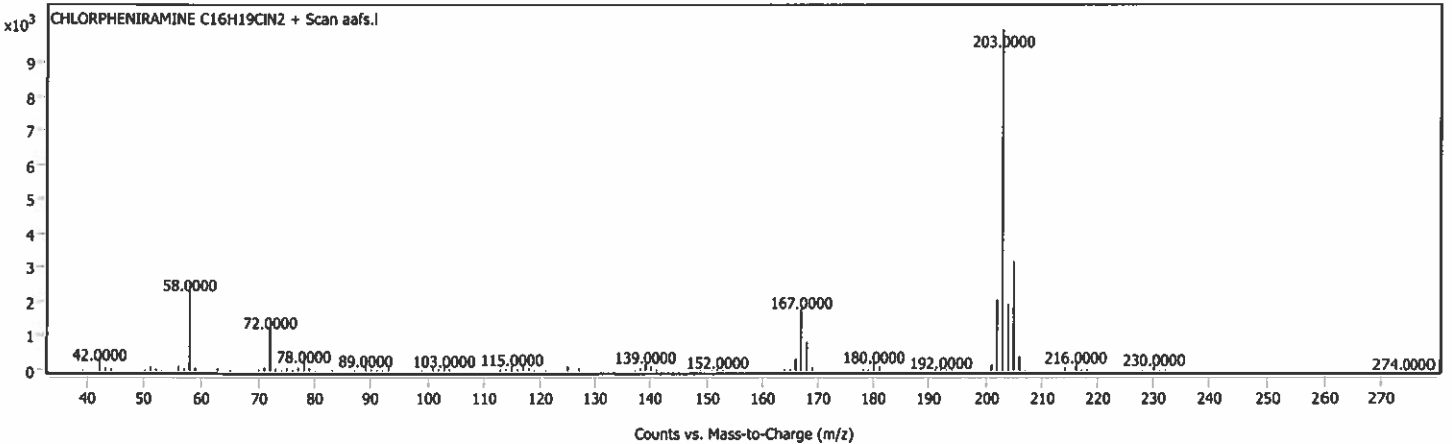
### Observed Spectrum



### Mirror Plot



### Library Spectrum



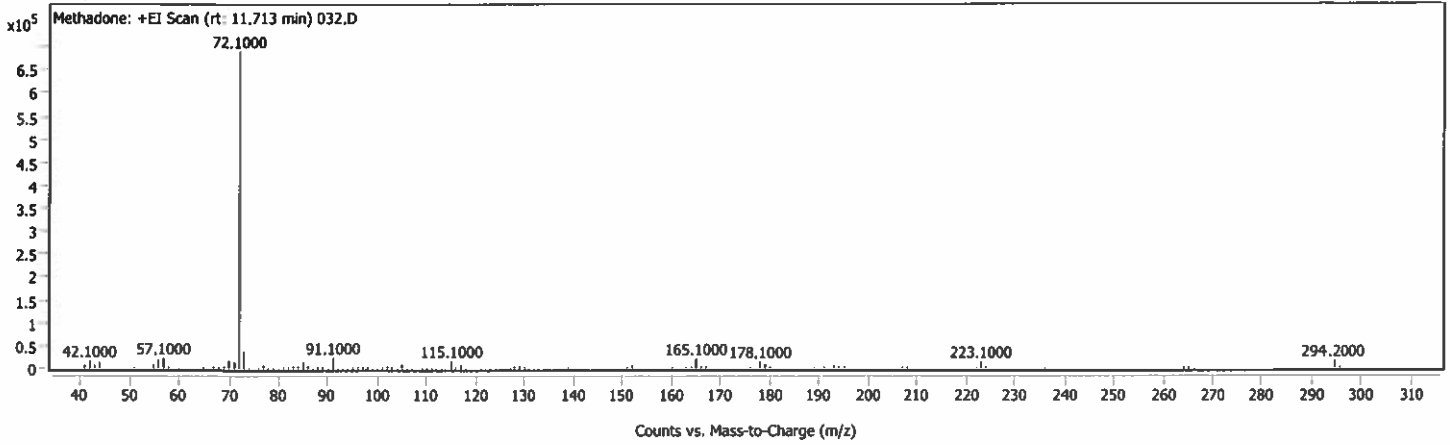
# Analysis Report

+ Scan (rt: 11.713 min)

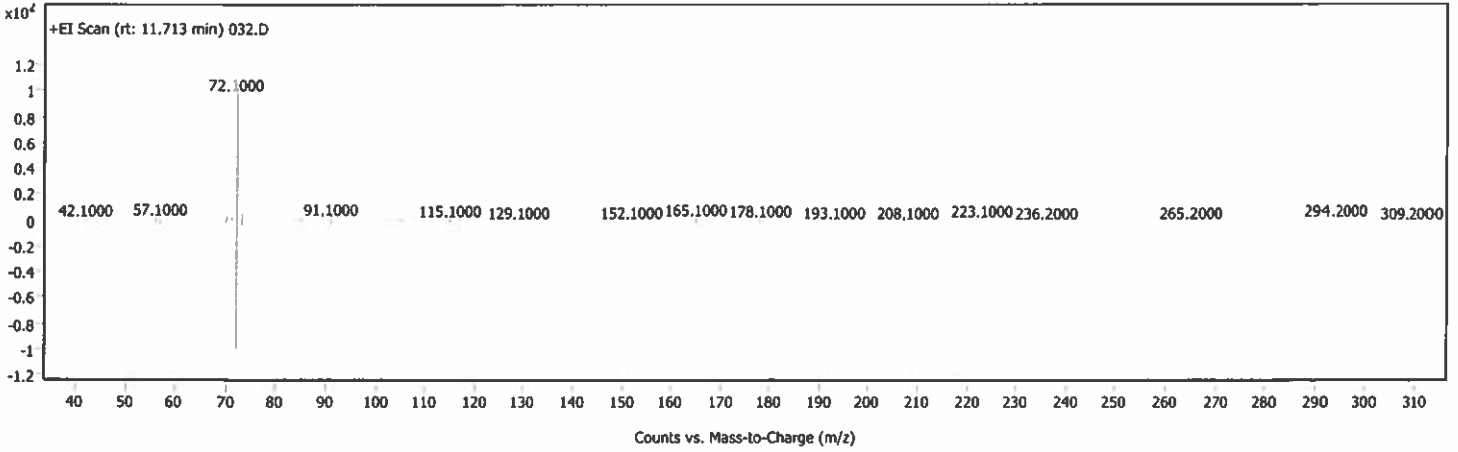
Methadone; C21H27NO

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methadone	C21H27NO	92.80	SWGDRUG 3.3.L	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/OC DATA IN XYLAZINE VALIDATION

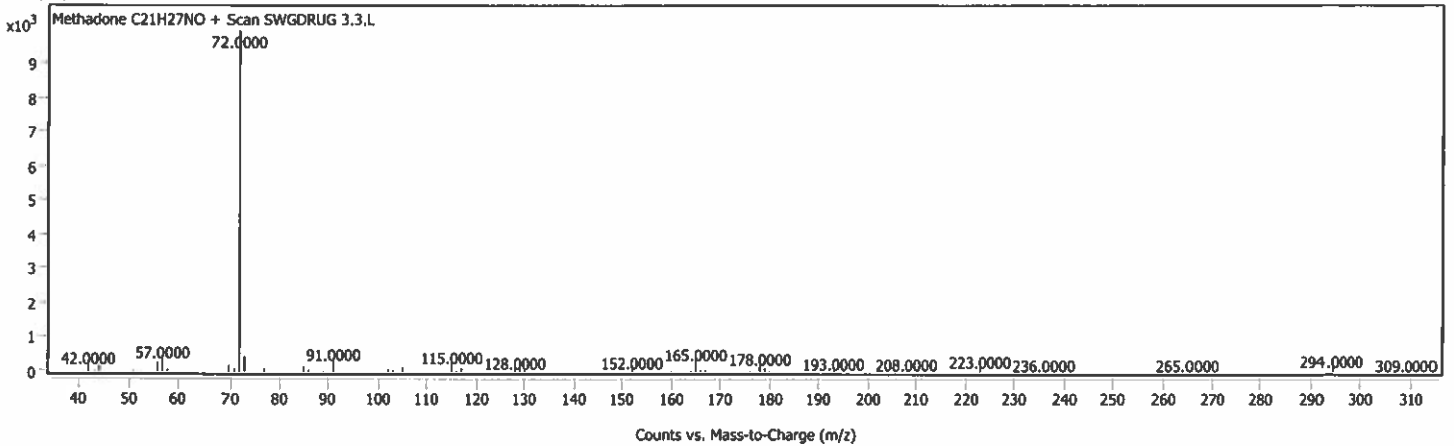
### Observed Spectrum



### Mirror Plot



### Library Spectrum



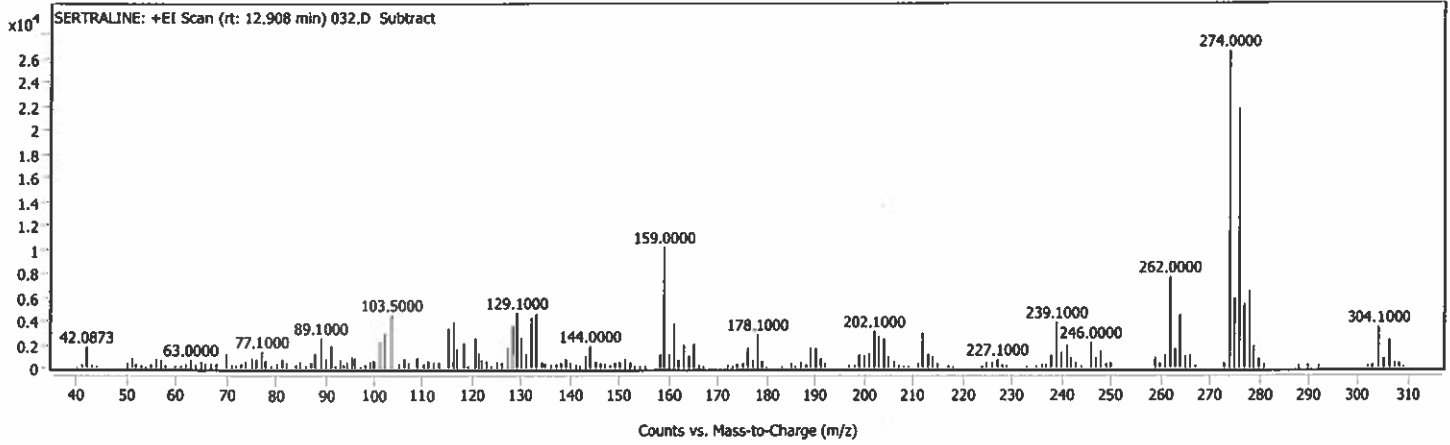
# Analysis Report

+ Scan (rt: 12.908 min) Sub

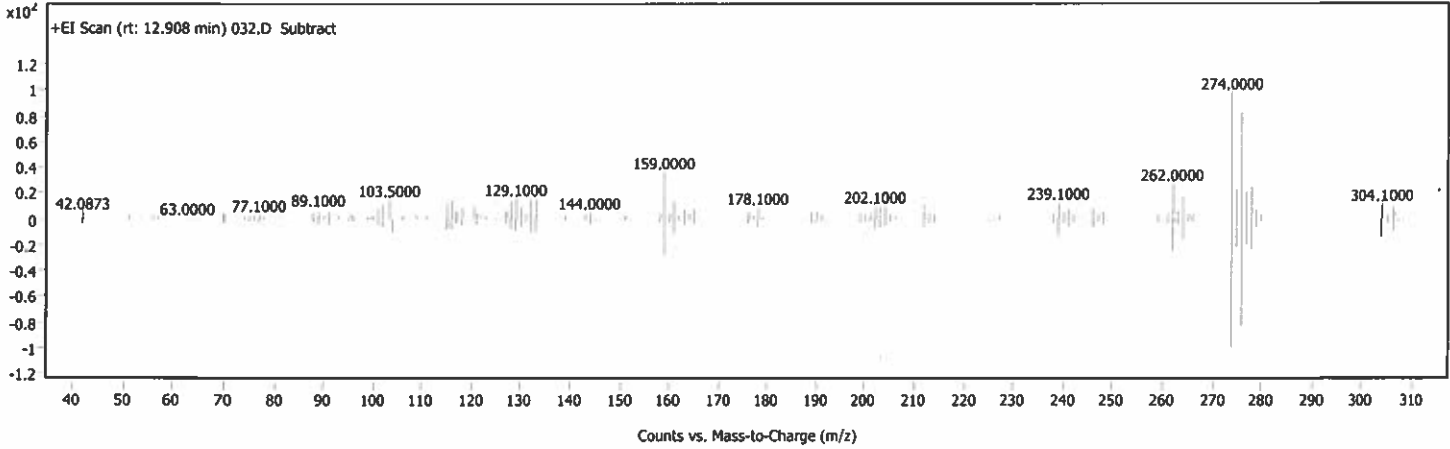
SERTRALINE; C17H17Cl2N

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
SERTRALINE	C17H17Cl2N	90.42	aafs.I	MATRIX 10 W/ IH CTL + XYLAZINE - LVR	KM/QC DATA IN XYLAZINE VALIDATION

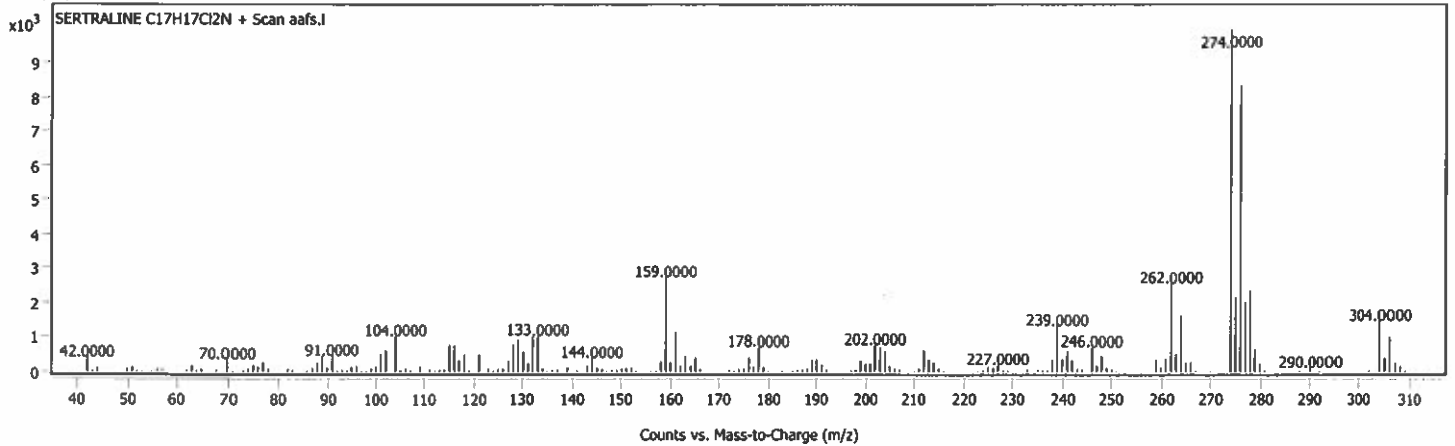
## Observed Spectrum



## Mirror Plot

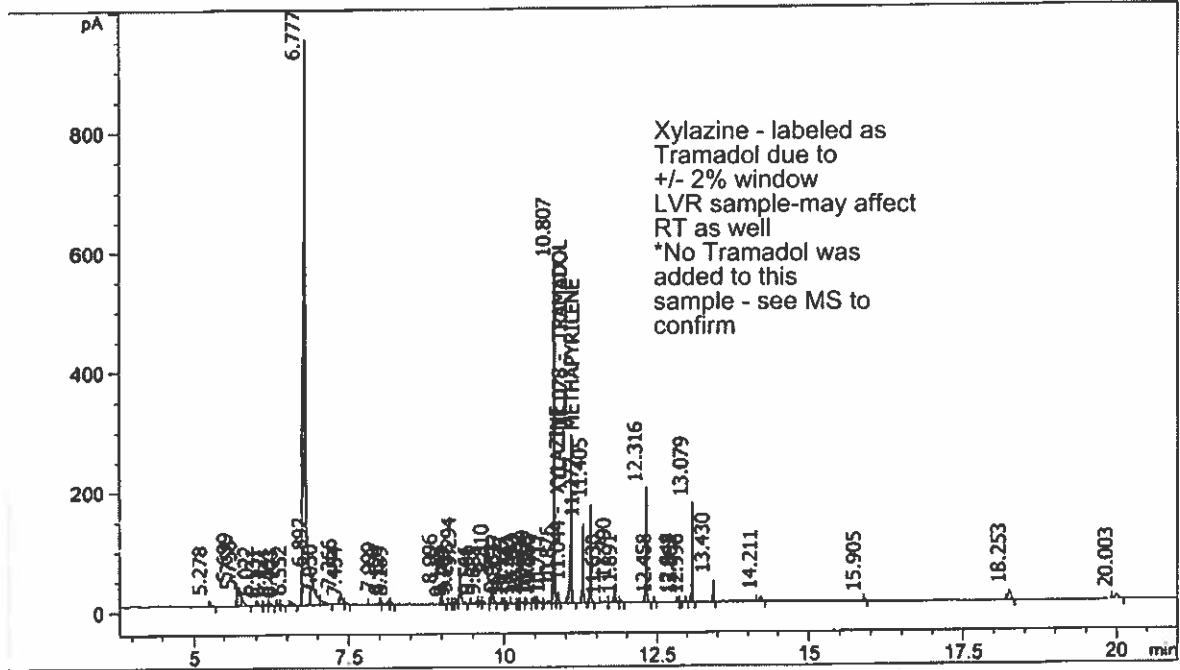


## Library Spectrum



Injection Date: 9/29/2021 3:56:40 AM Seq Line: 36  
 Sample Name: MATRIX 10 W/ IH CTL +-> Vial 37  
 Sample Info: *XYLAZINE KM N/17/22*

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



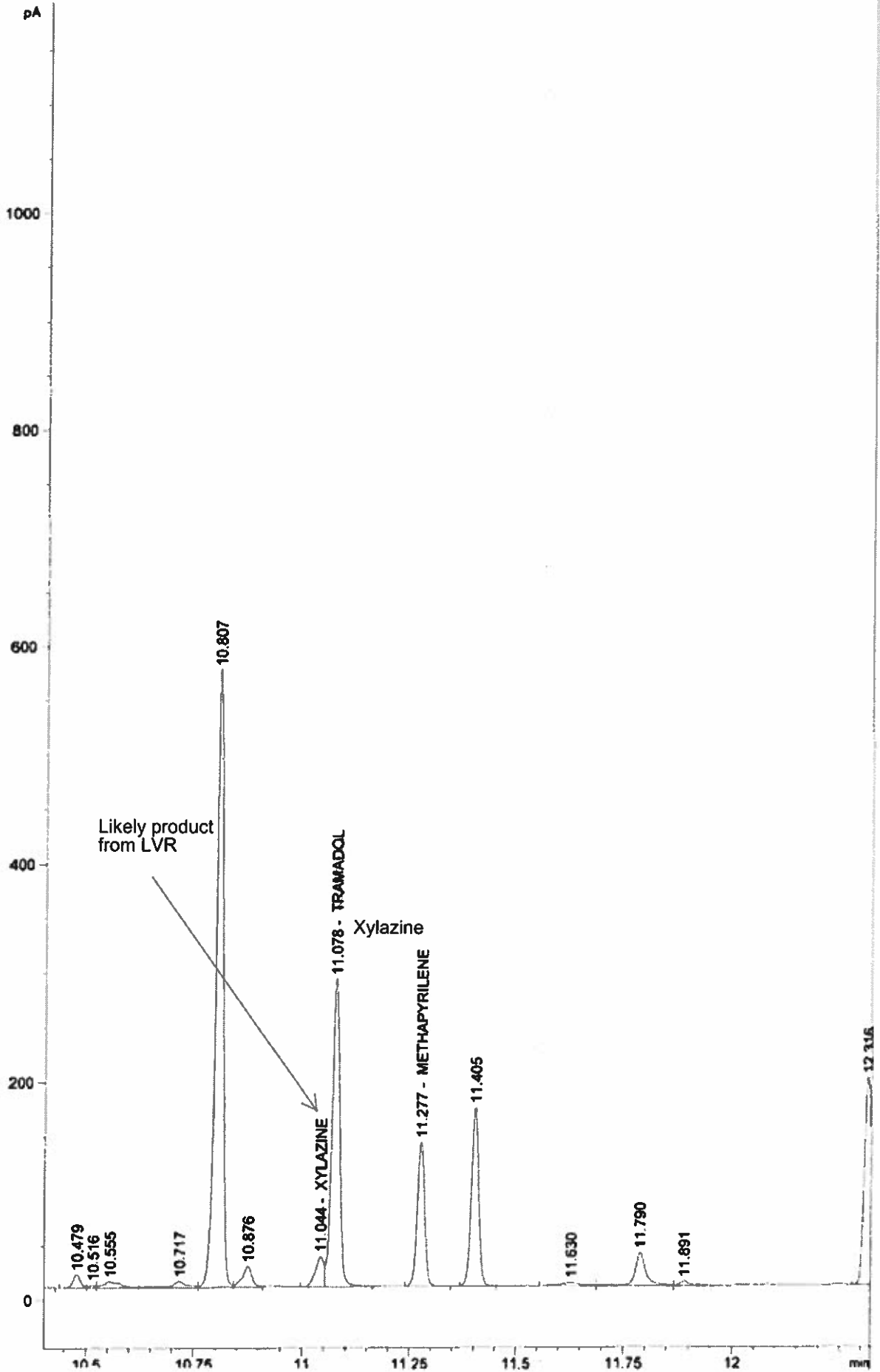
see expanded chromatogram on following page

RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
11.044	11.046	37.204	27.497	0.000000	XYLAZINE
11.078	11.033	350.607	281.564	0.000000	TRAMADOL RRT -0.9824
11.277	11.278	161.030	132.303	1.000000	METHAPYRILENE

LVR sample not included in average RRTs

Current Chromatogram(s)

NPD1 A, Front Signal (092821\BASE000036 D)

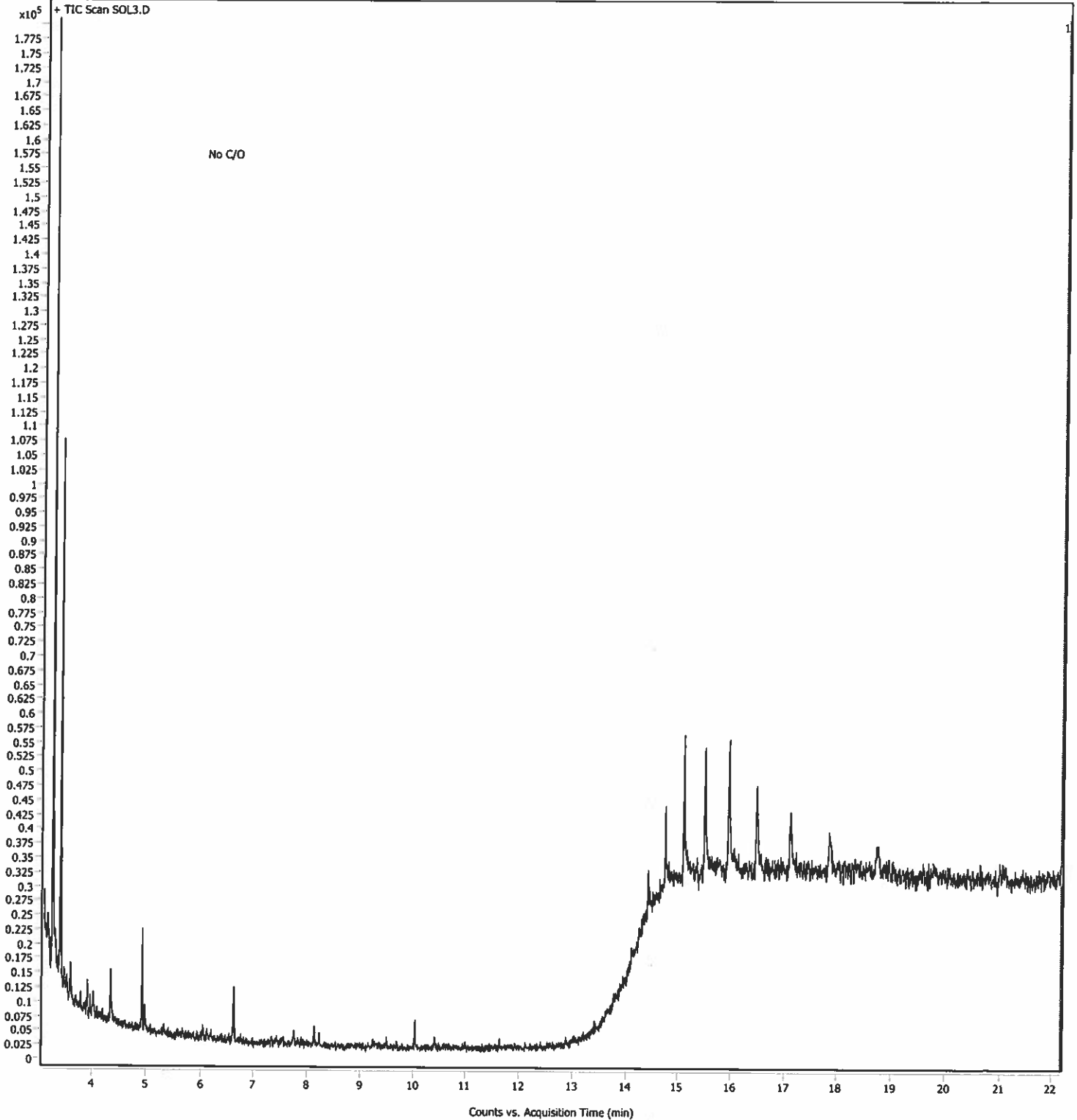


# Analysis Report

## Sample Information

Sample Name	Solvent Blank 3	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\SOL3.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 4:44:48 AM (UTC-04:00)
Position	62	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





Injection Date: 9/29/2021 4:21:01 AM

Seq Line: 37

Sample Name: SOLVENT BLANK 3

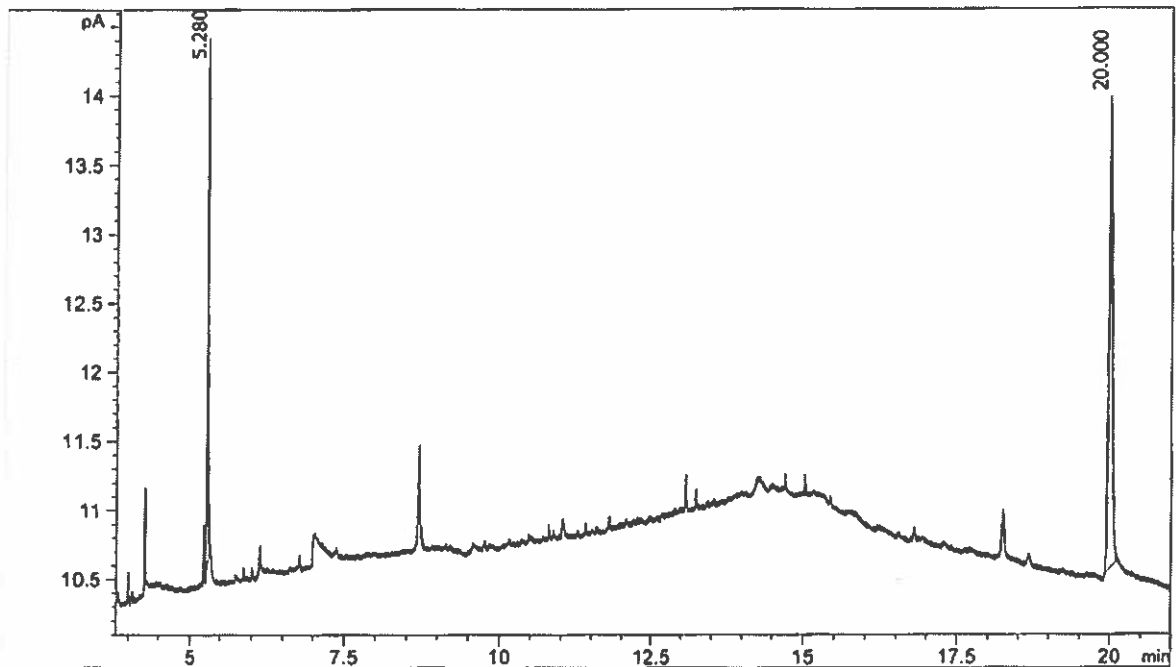
Vial 38

Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M

Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M

KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

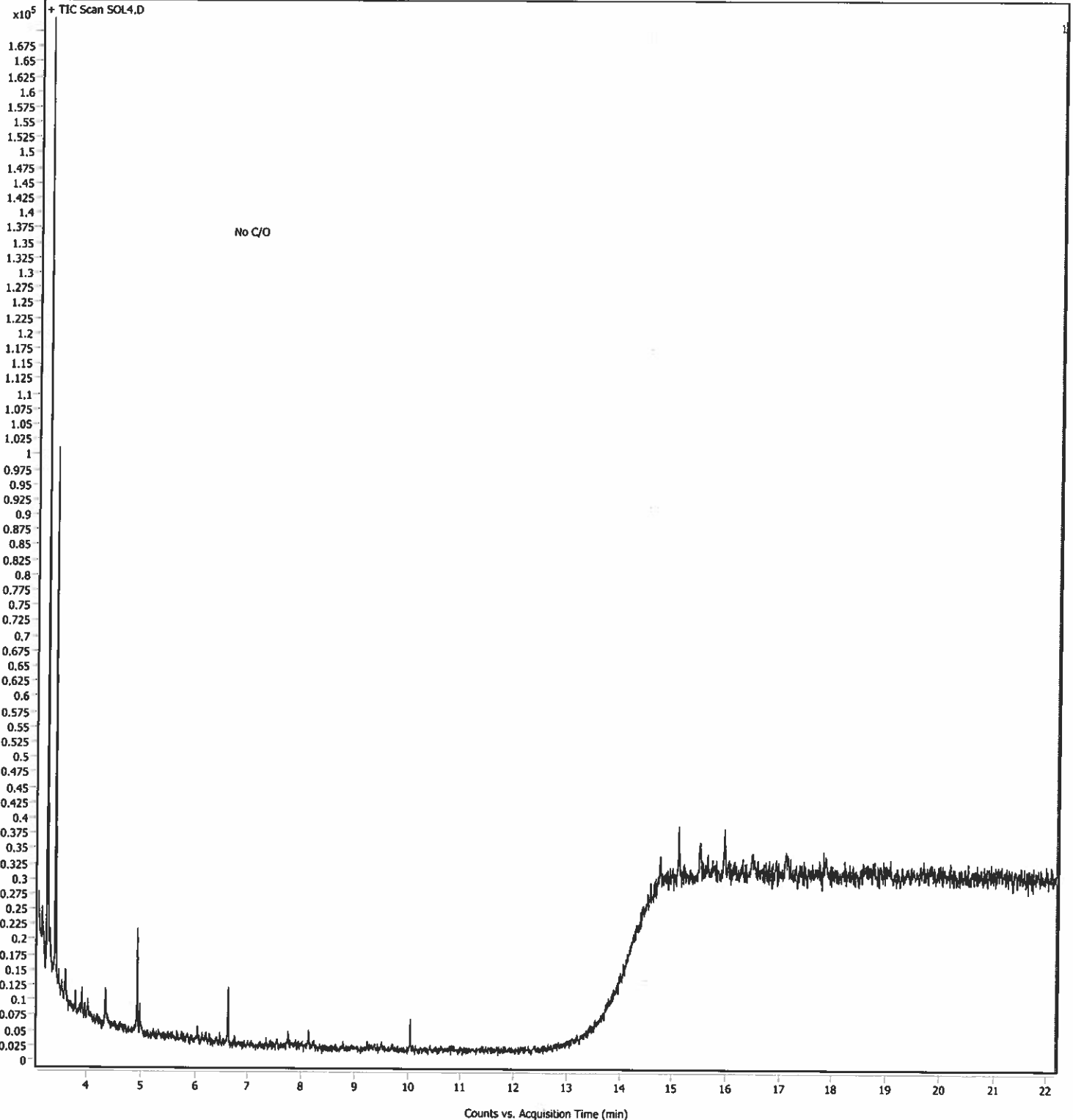
# Analysis Report



## Sample Information

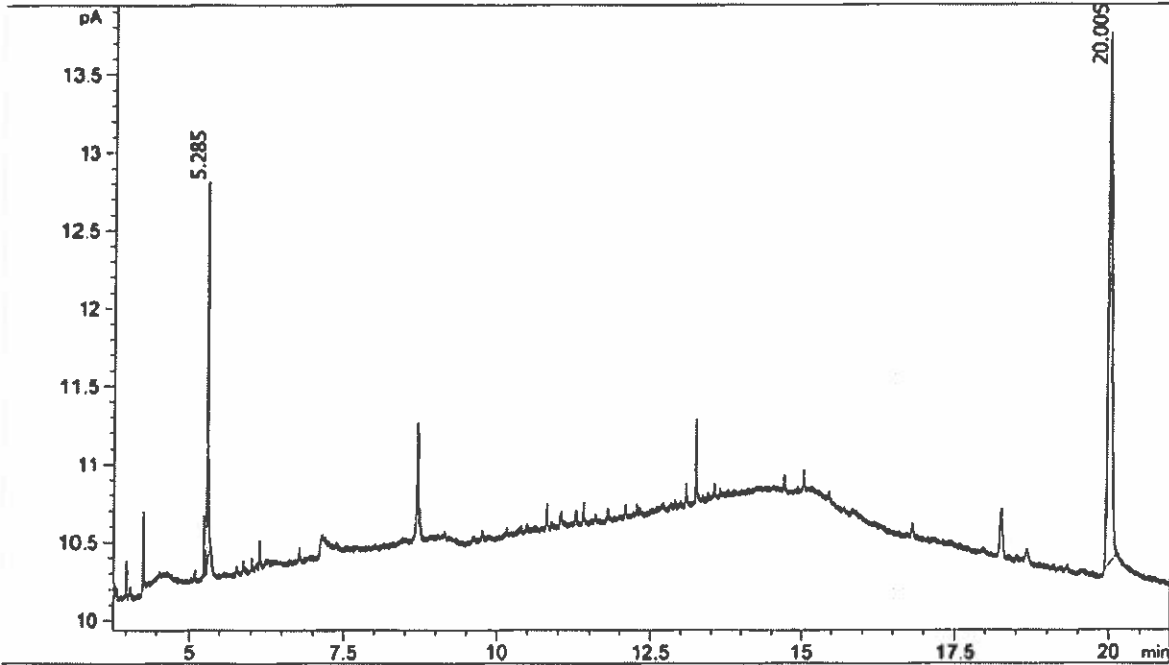
Sample Name	Solvent Blank 4	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\SOL4.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/29/2021 5:10:36 AM (UTC-04:00)
Position	63	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALIM
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



=====  
Injection Date: 9/29/2021 4:45:16 AM Seq Line: 38  
Sample Name: SOLVENT BLANK 4 Vial 39  
Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



RT (min)	Exp RT (min)	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

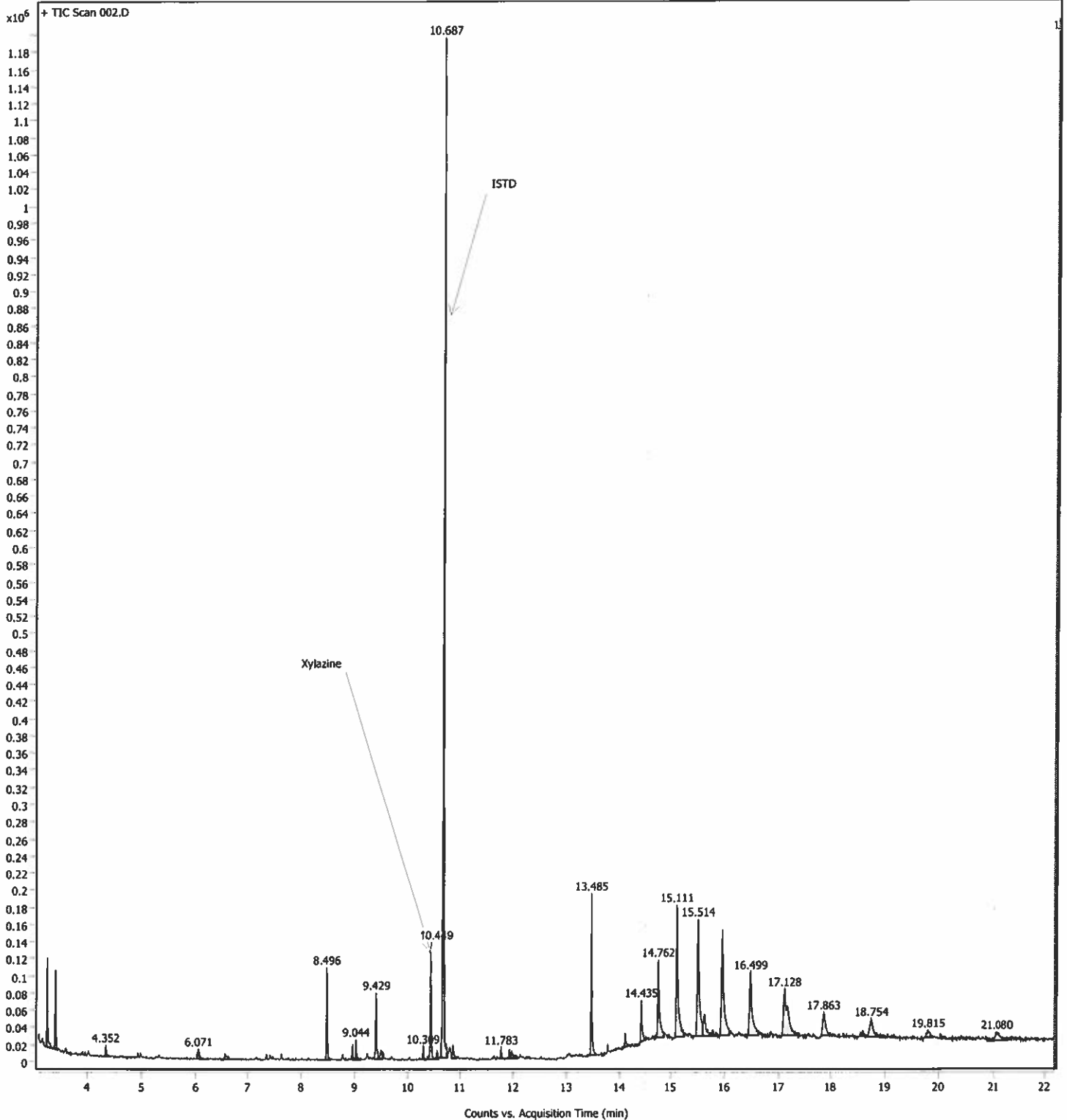
## Sample Information

**Sample Name** MATRIX 2 0.10 mg/L - BLD  
**Instrument** #3 - Enhanced  
**Position** 28  
**Operator** KM/QC DATA IN XYLAZINE VALIDATION

**Data File Path**  
**Acq. Time (Local)**  
**Method Path (Acq)**

C:\MassHunter\GCMS\1\data\BASES\092821\002.D  
9/28/2021 2:06:40 PM (UTC-04:00)  
C:\MassHunter\GCMS\1\methods\ALKALI.M

## Sample Chromatograms



# Analysis Report

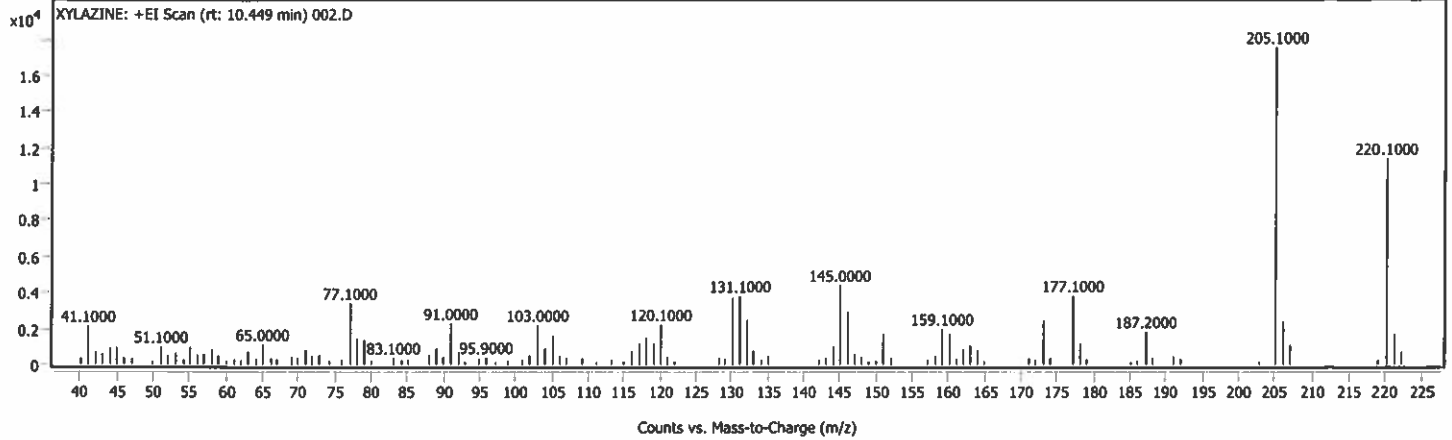
## Sample Spectra

+ Scan (rt: 10.449 min)

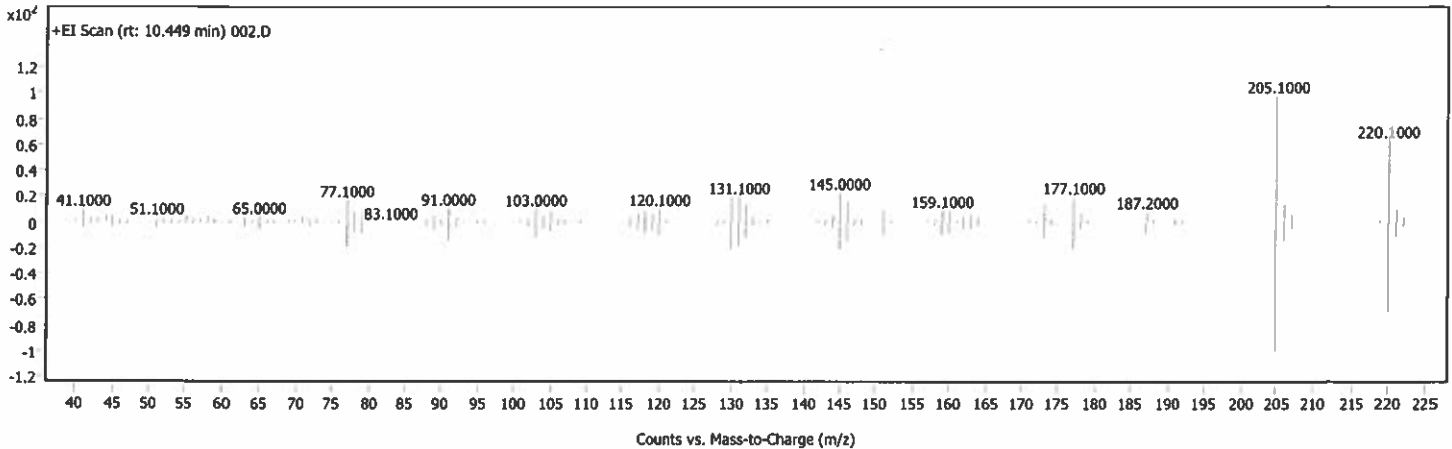
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	95.79	aafs.l	MATRIX 2 0.10 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

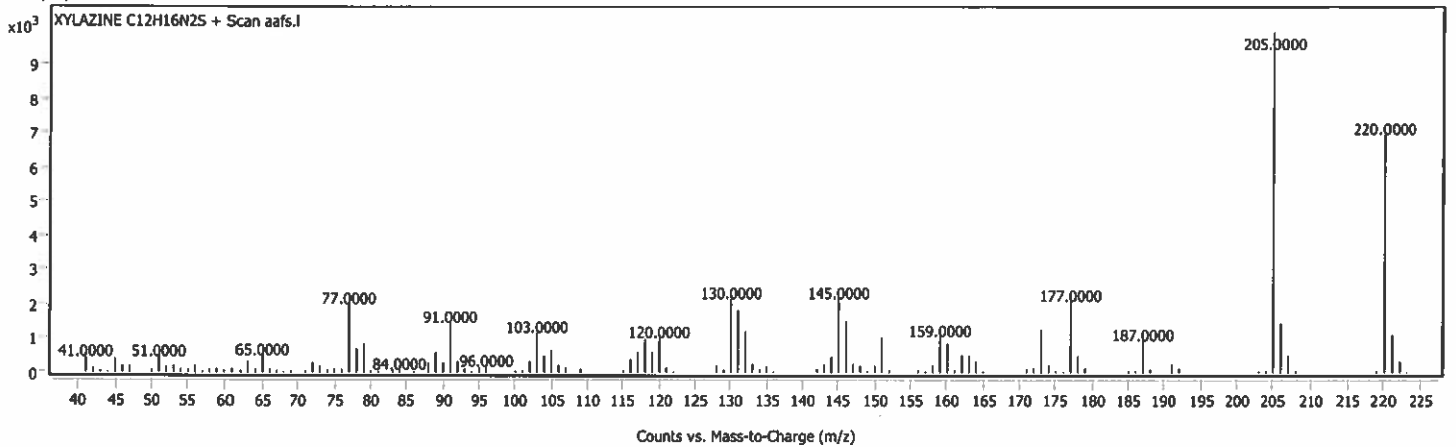
## Observed Spectrum



## Mirror Plot



## Library Spectrum



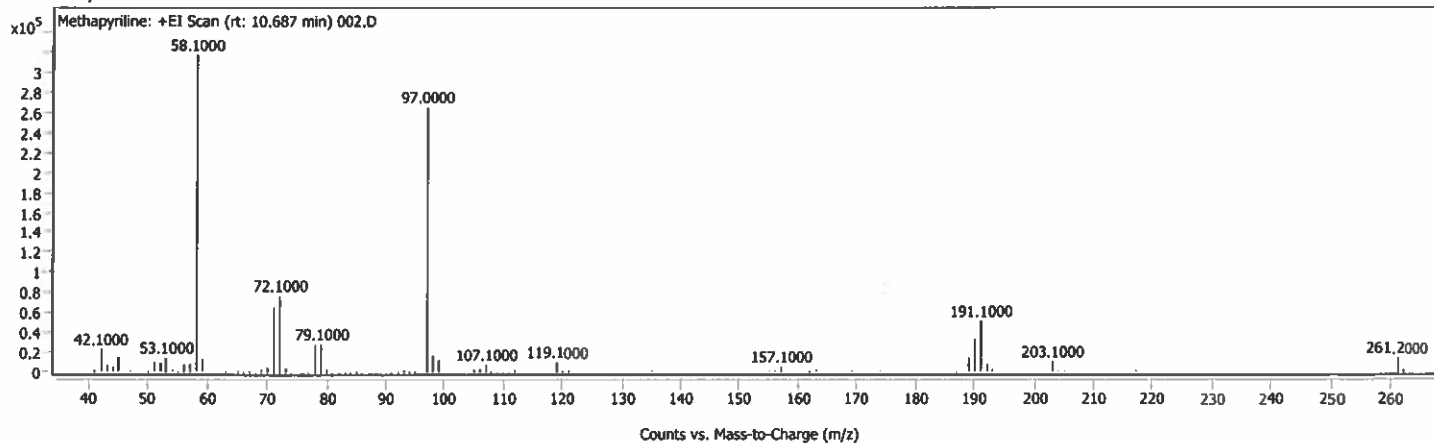
# Analysis Report

+ Scan (rt: 10.687 min)

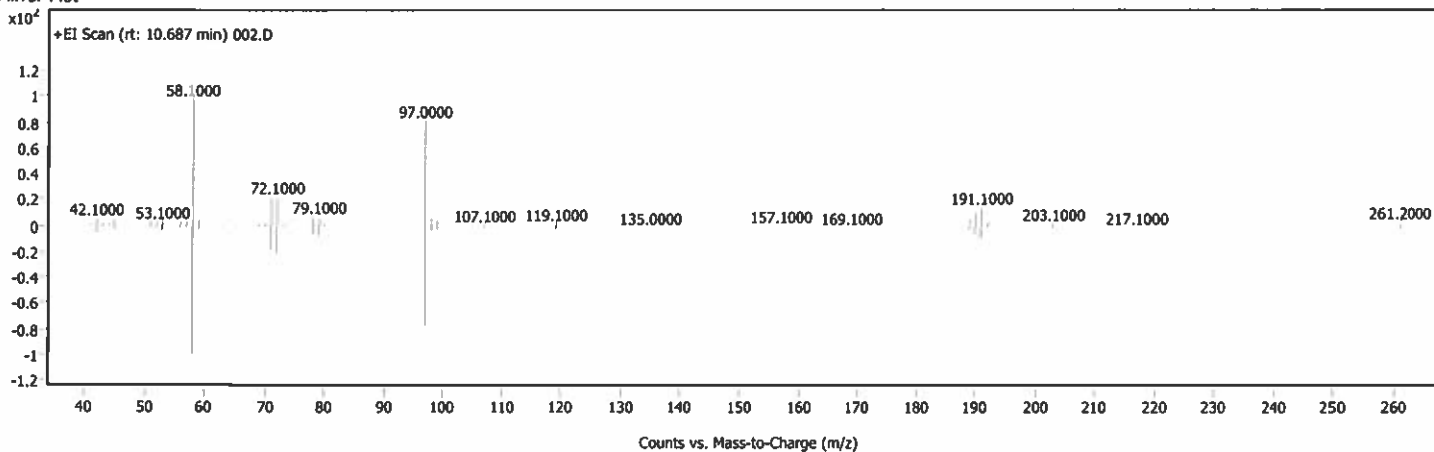
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.83	SWGDRUG 3.3.L	MATRIX 2 0.10 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

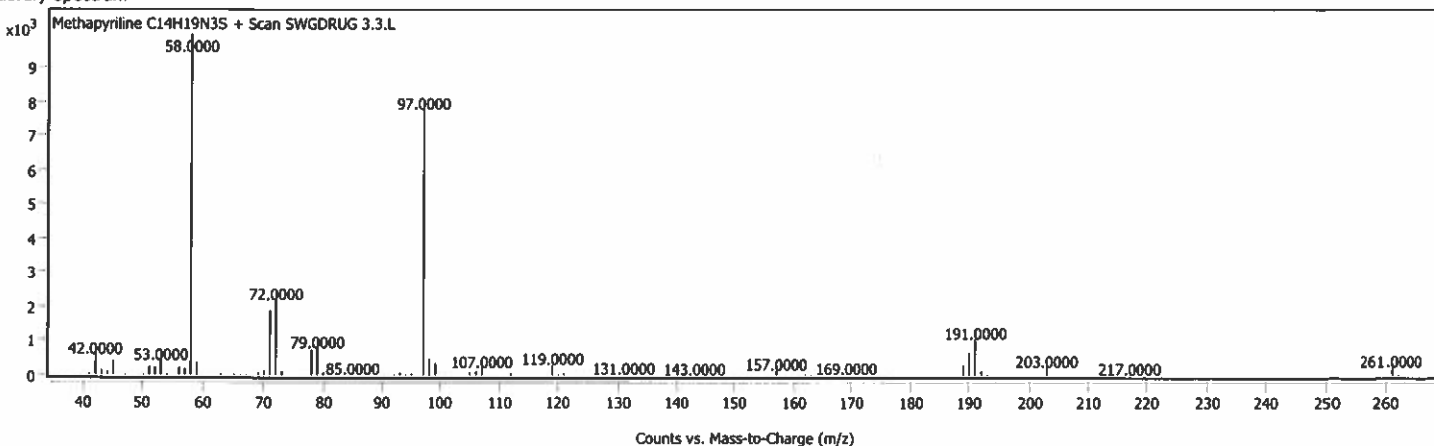
## Observed Spectrum



## Mirror Plot

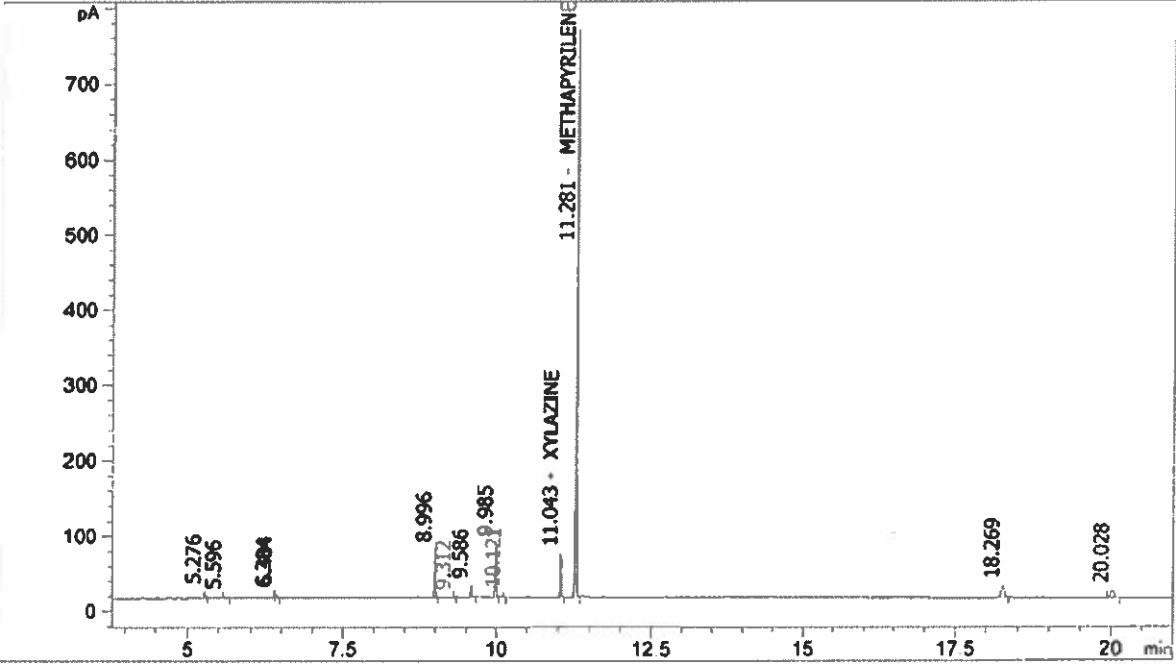


## Library Spectrum



Injection Date: 9/28/2021 2:34:07 PM Seq Line: 3  
 Sample Name: MATRIX 2 0.10 mg/L - -> Vial 3  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



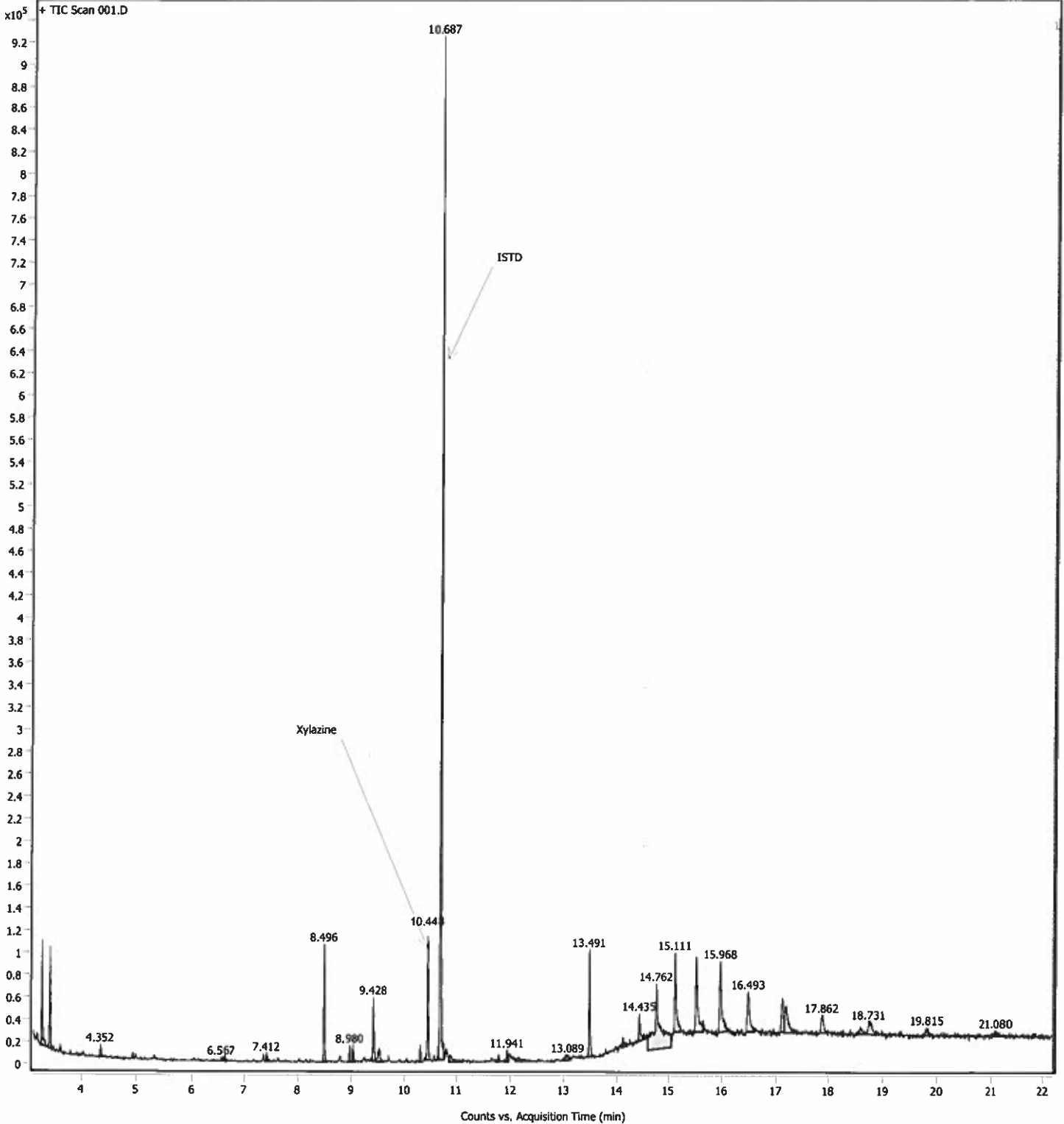
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.043	11.046	70.356	58.838	0.000000	XYLAZINE RRT -0.9789
11.281	11.278	890.595	749.848	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 0.10 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\001.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 1:40:54 PM (UTC-04:00)
Position	27	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

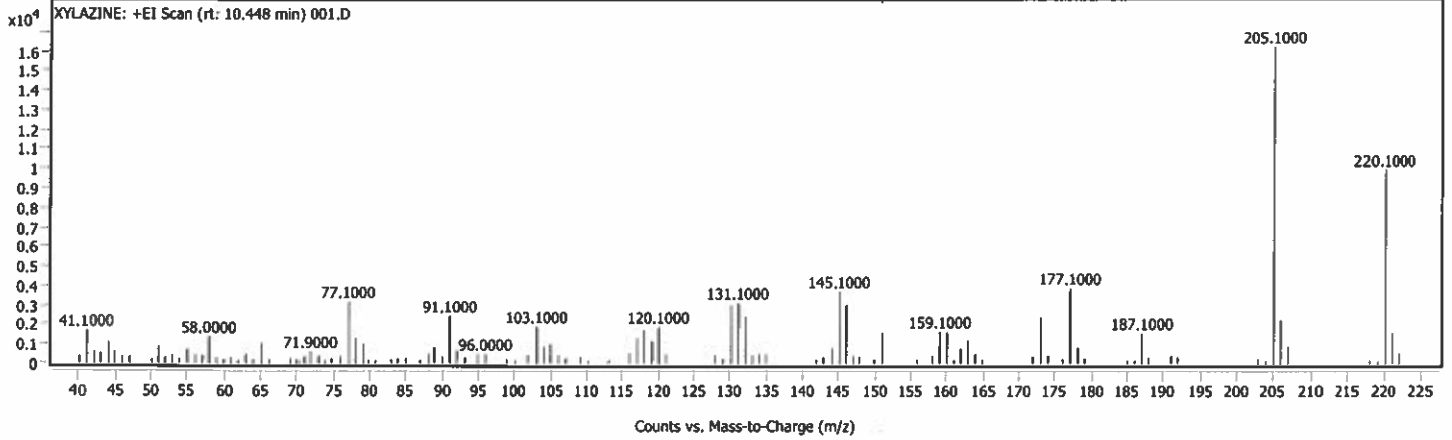
## Sample Spectra

+ Scan (rt: 10.448 min)

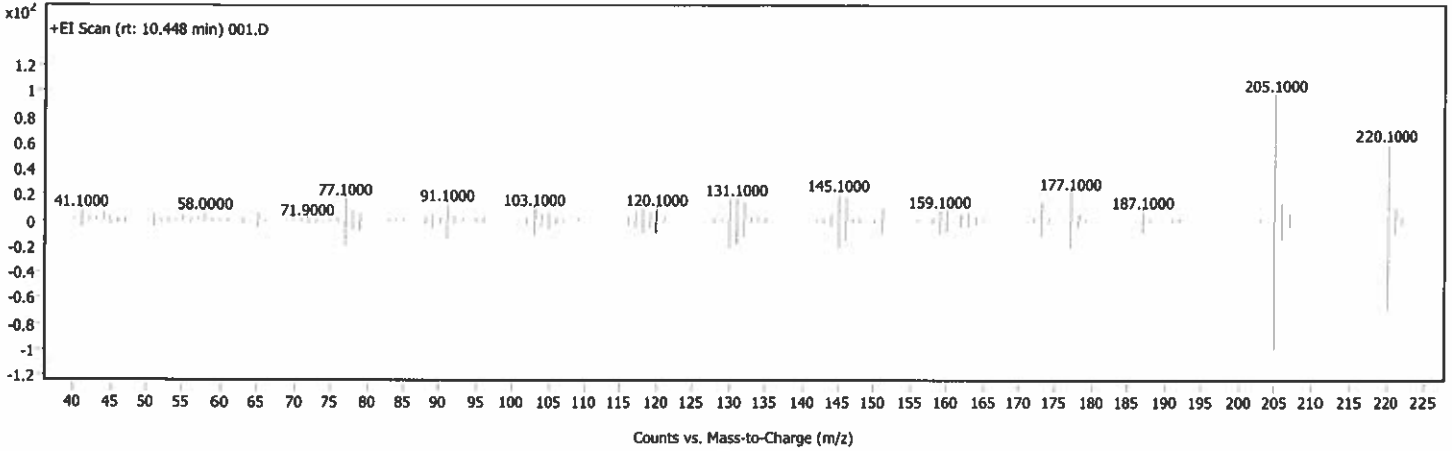
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	95.94	aafs.l	MATRIX 1 0.10 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

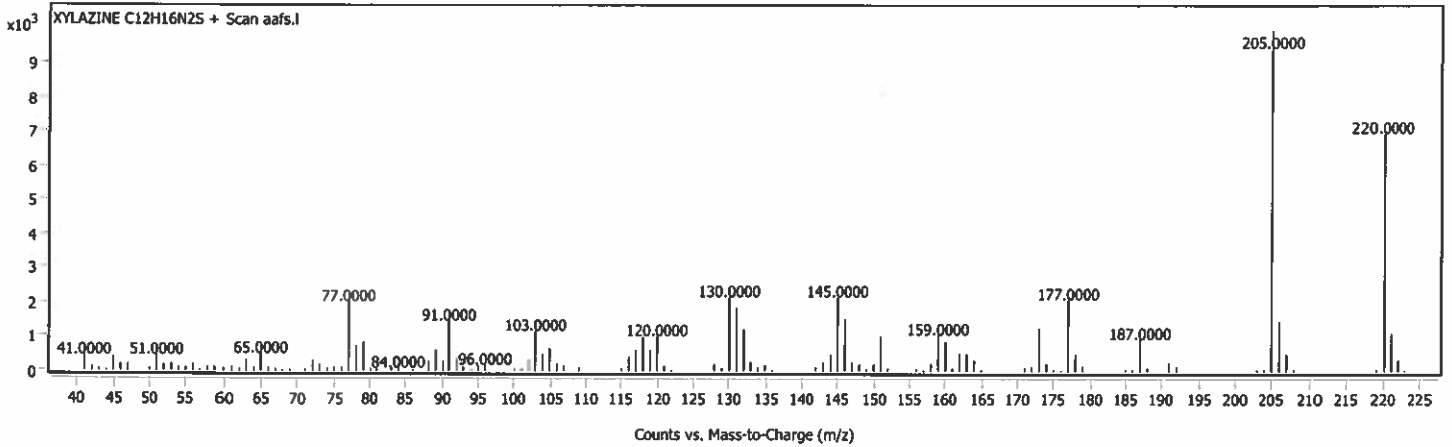
### Observed Spectrum



### Mirror Plot



### Library Spectrum



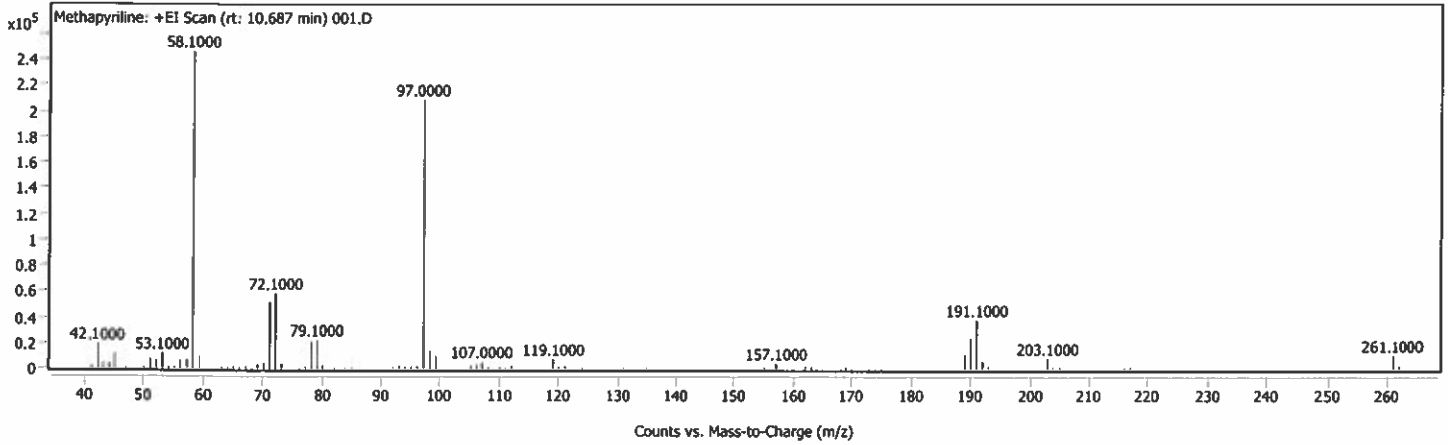
# Analysis Report

+ Scan (rt: 10.687 min)

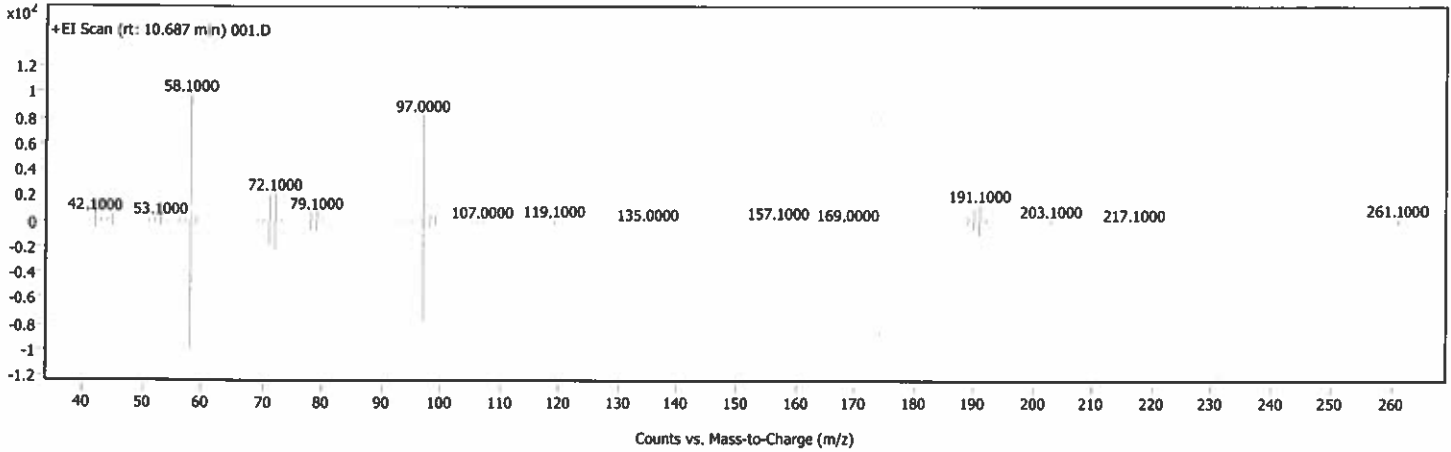
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.73	SWGDRUG 3.3.L	MATRIX 1 0.10 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

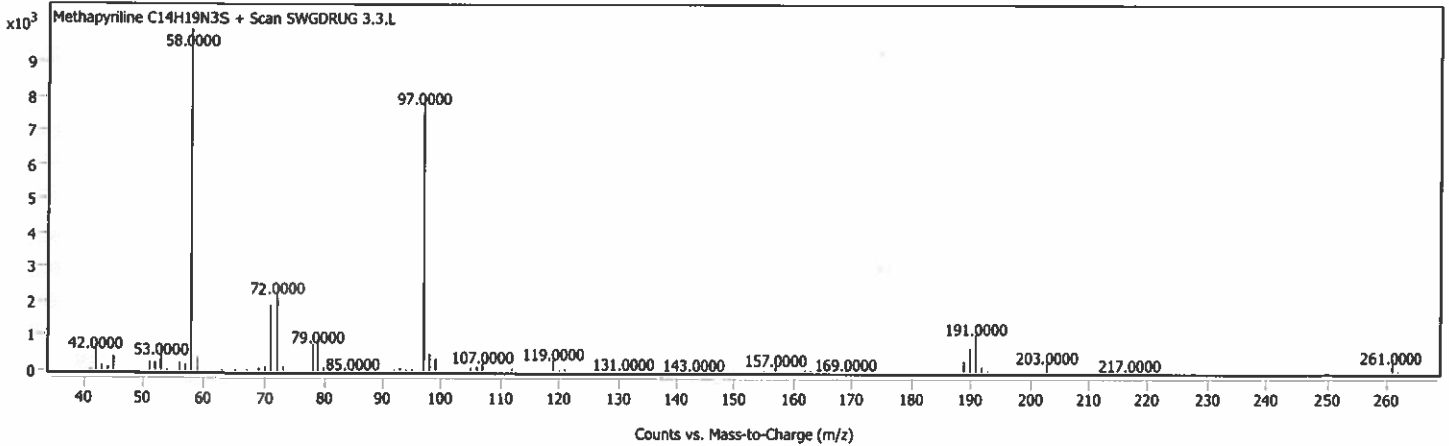
## Observed Spectrum



## Mirror Plot

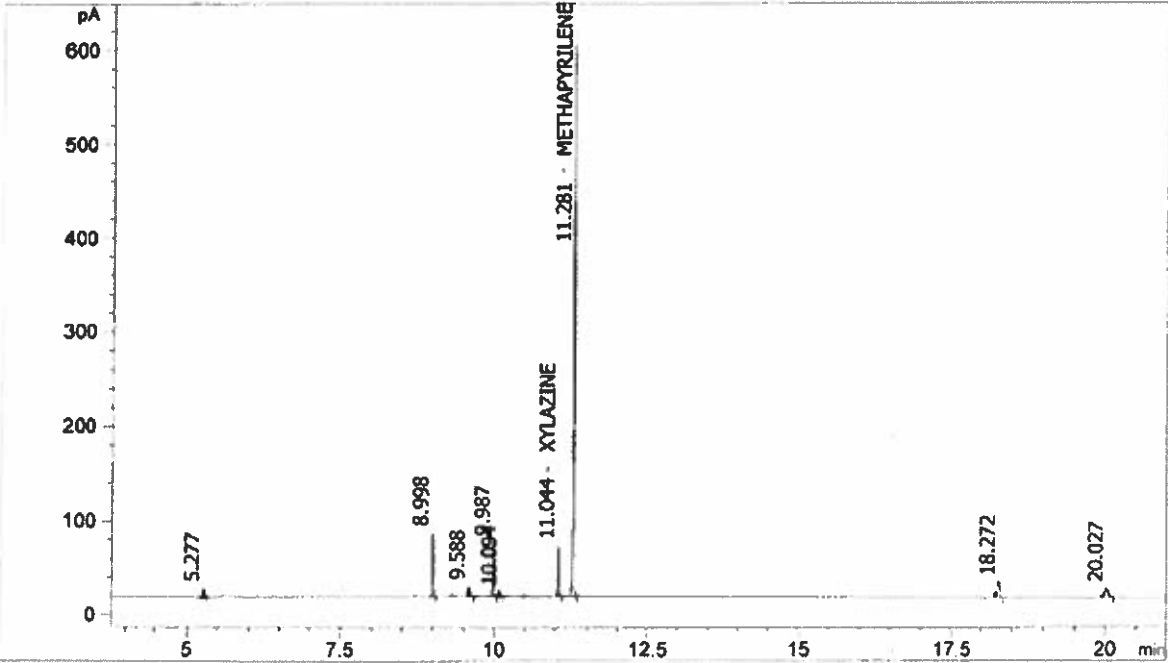


## Library Spectrum



Injection Date: 9/28/2021 2:09:50 PM Seq Line: 2  
 Sample Name: MATRIX 1 0.10 mg/L - -> Vial 2  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



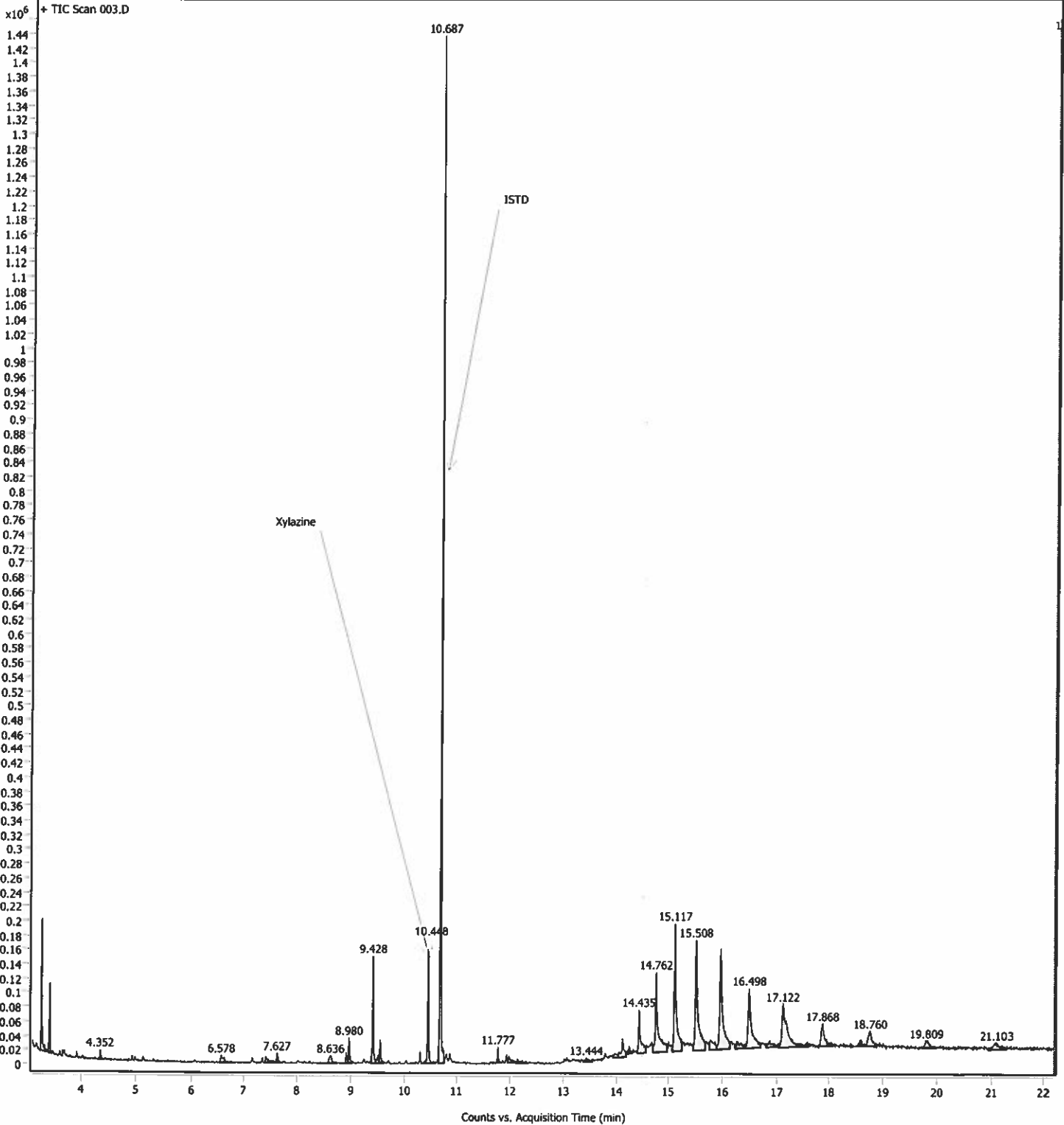
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.044	11.046	66.004	54.125	0.000000	XYLAZINE RRT - 0.9790
11.281	11.278	695.777	596.023	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 0.10 mg/L - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\003.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 2:32:32 PM (UTC-04:00)
Position	29	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

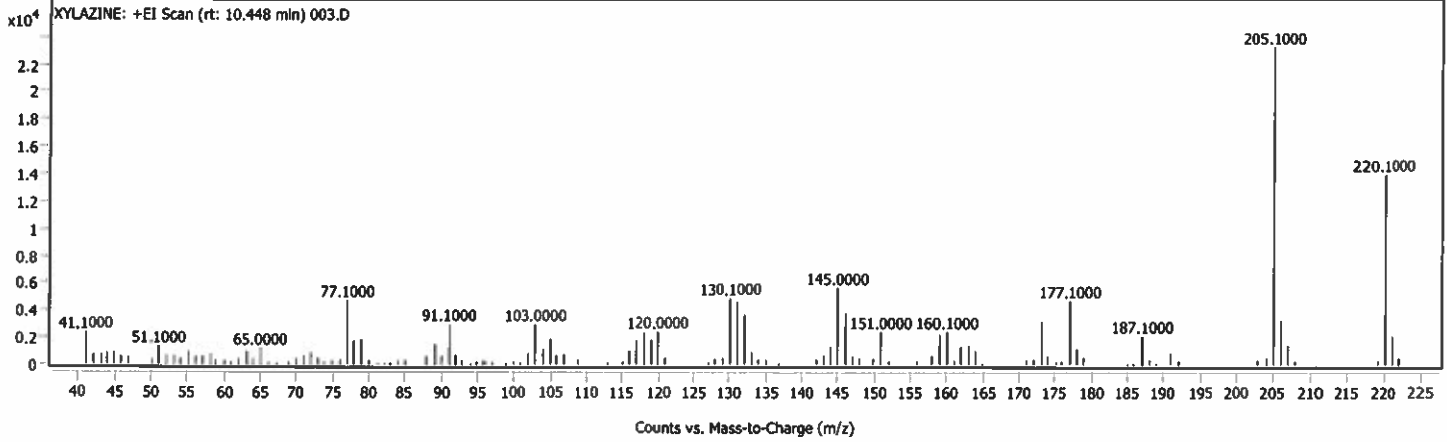
## Sample Spectra

+ Scan (rt: 10.448 min)

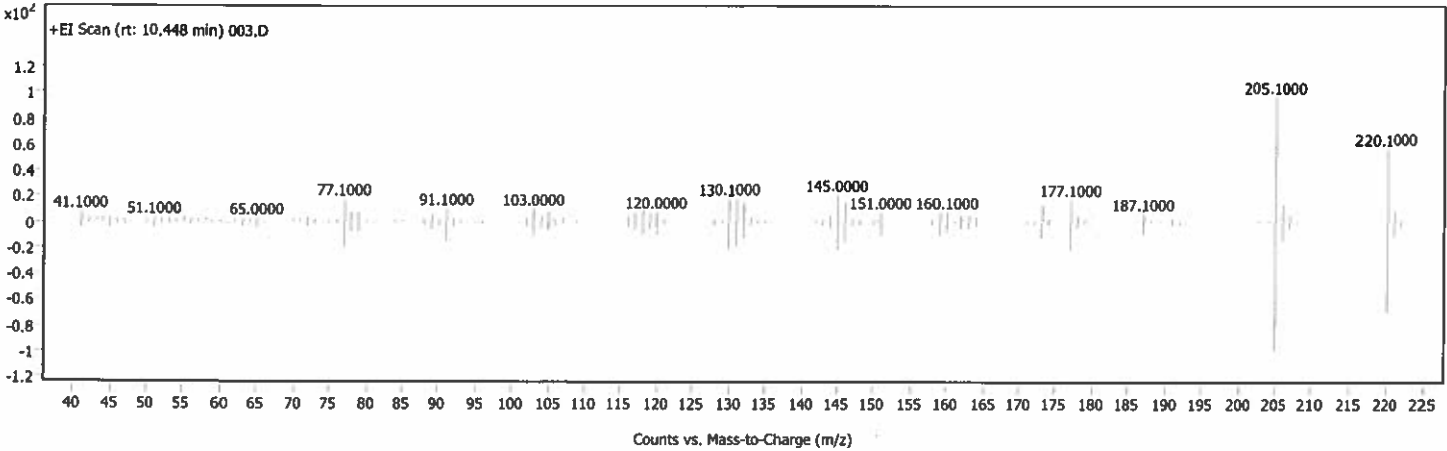
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	96.57	aafs.l	MATRIX 6 0.10 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

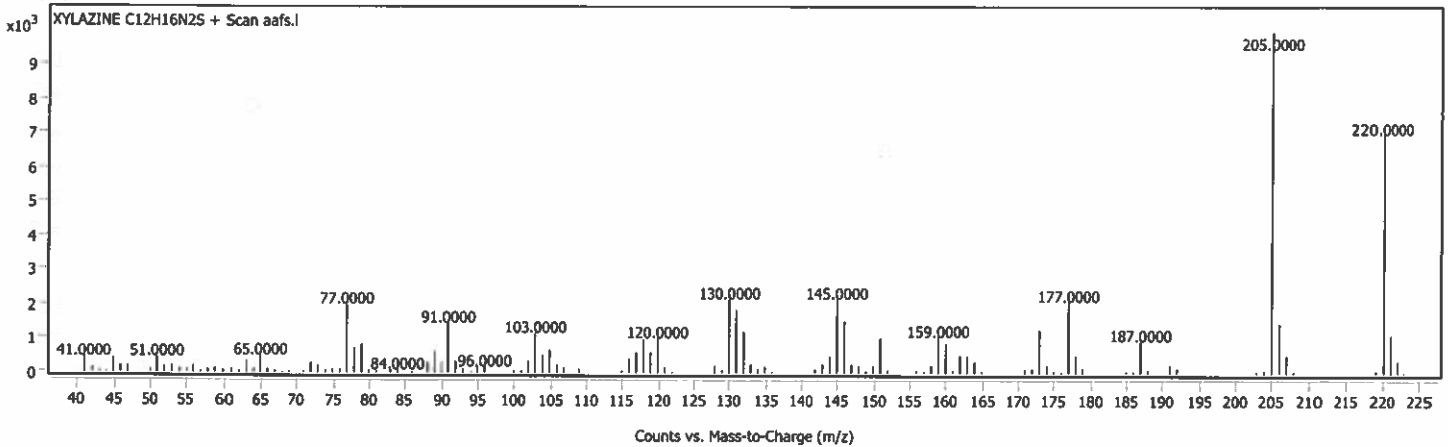
## Observed Spectrum



## Mirror Plot



## Library Spectrum



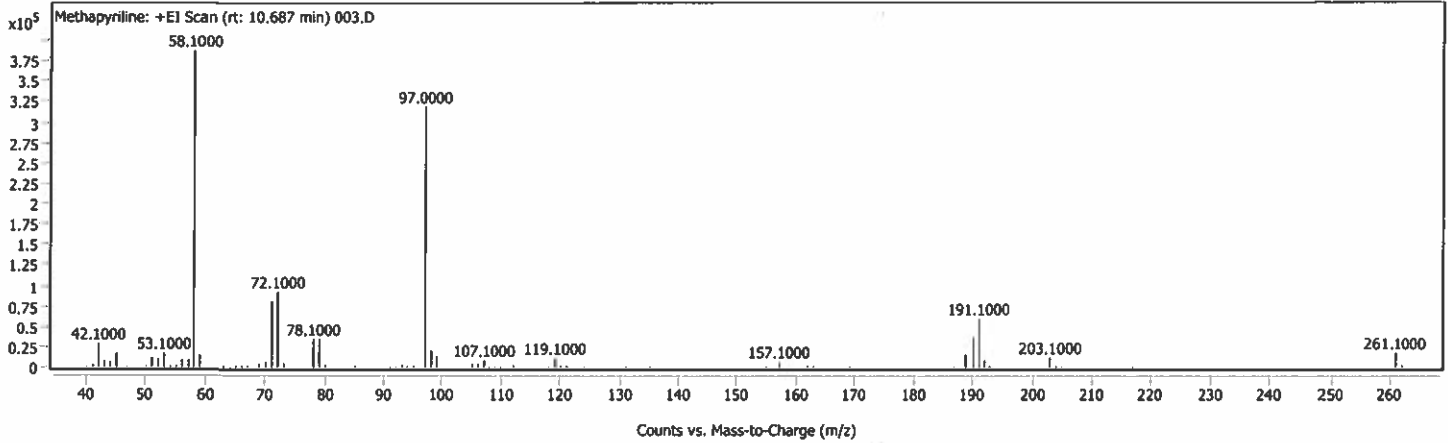
# Analysis Report

+ Scan (rt: 10.687 min)

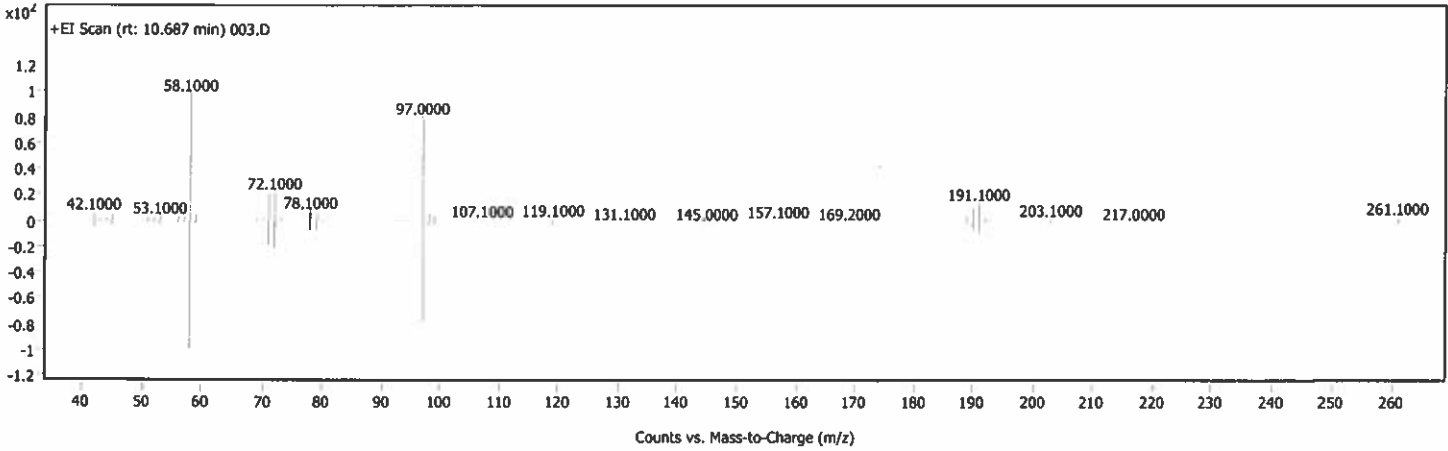
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.93	SWGDRUG 3.3.L	MATRIX 6 0.10 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

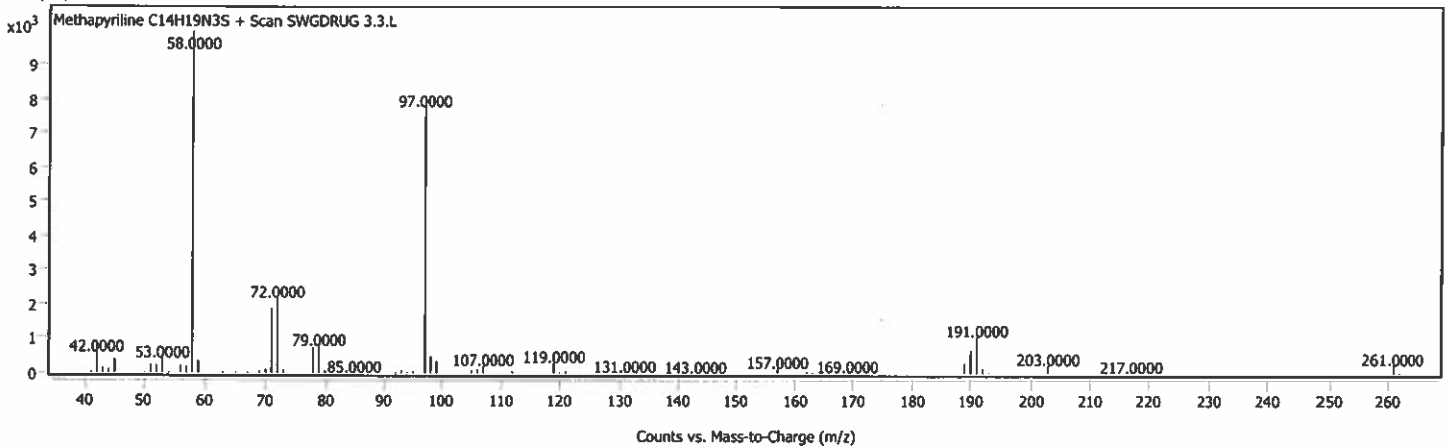
### Observed Spectrum



### Mirror Plot



### Library Spectrum

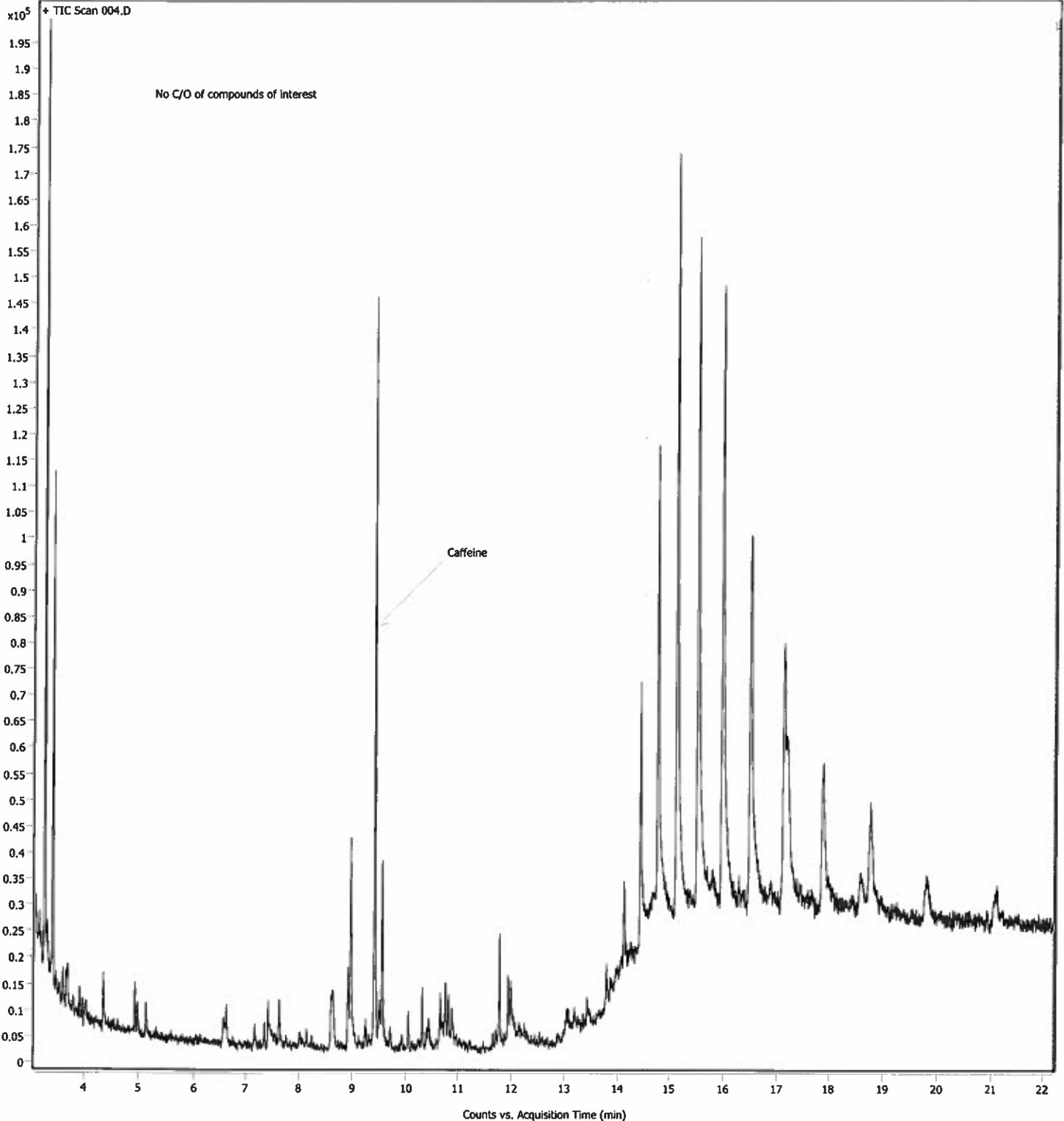


# Analysis Report

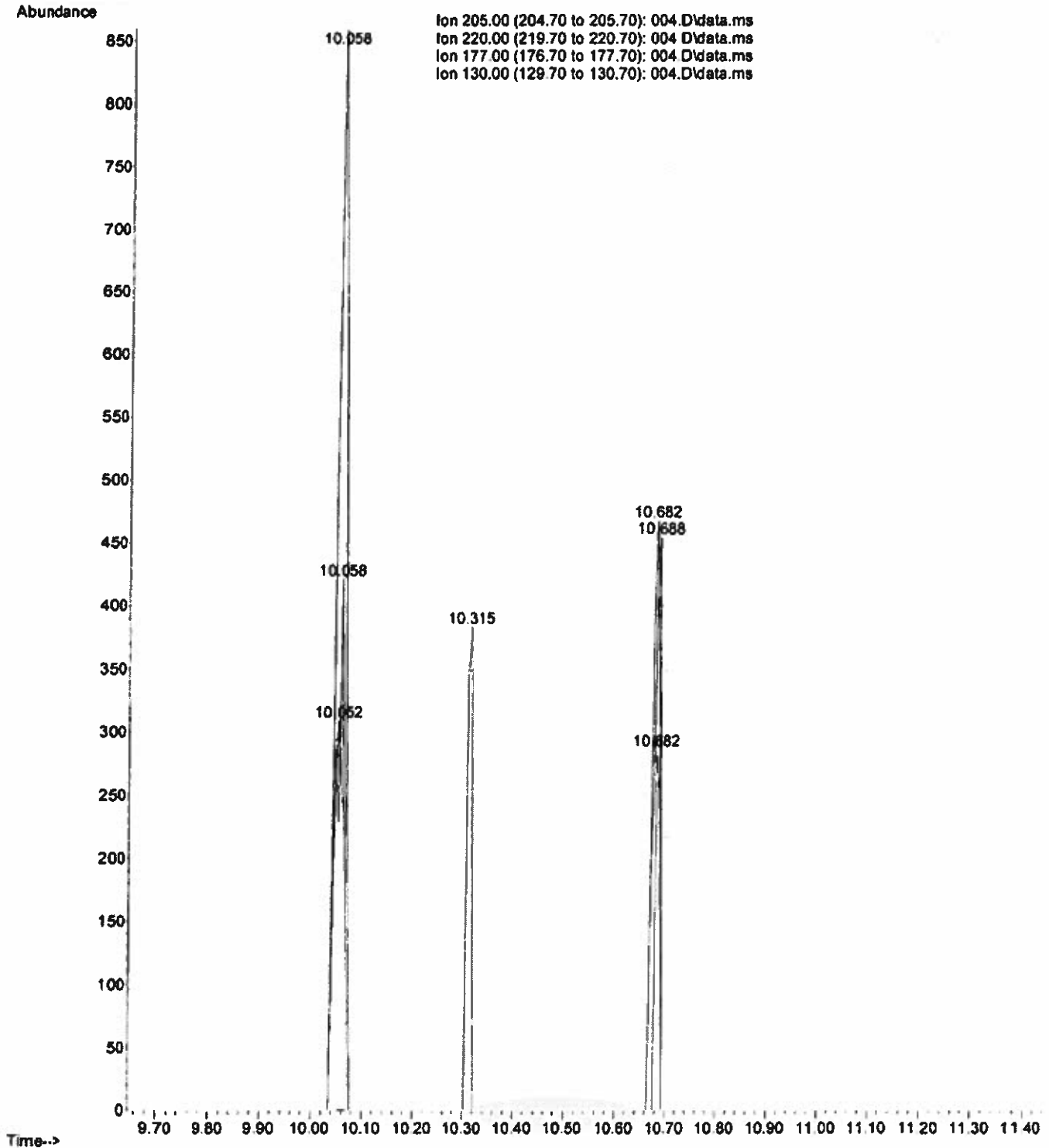
## Sample Information

Sample Name	MATRIX BLANK after 0.1 mg/L	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\004.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 2:58:20 PM (UTC-04:00)
Position	30	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



File :C:\Users\TOX\Desktop\092821\004.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 28 Sep 2021 14:58 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK after 0.1 mg/L  
Misc Info :  
Vial Number: 30





# Analysis Report

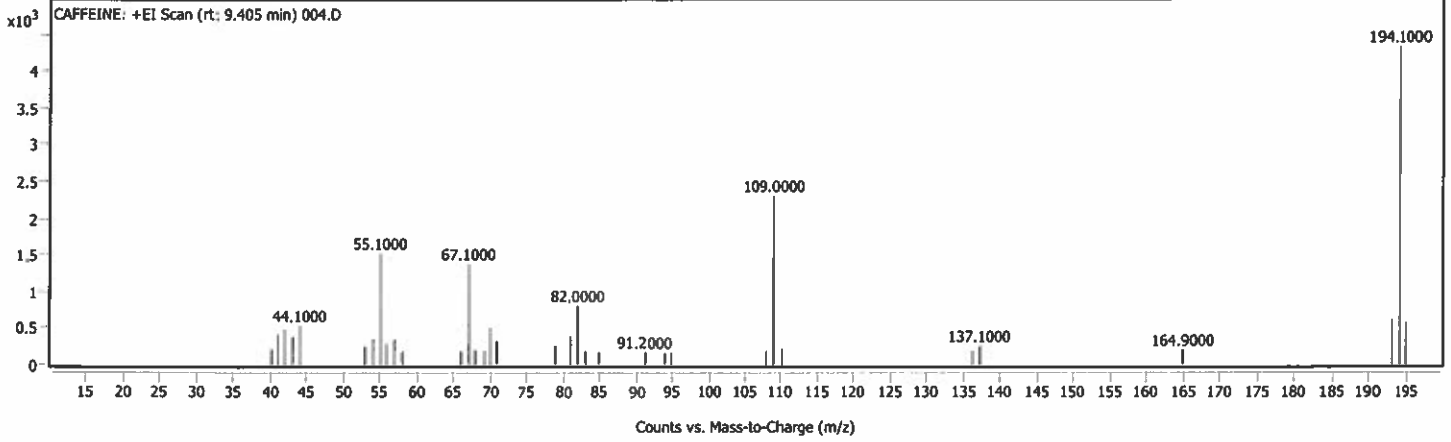
## Sample Spectra

+ Scan (rt: 9.405 min)

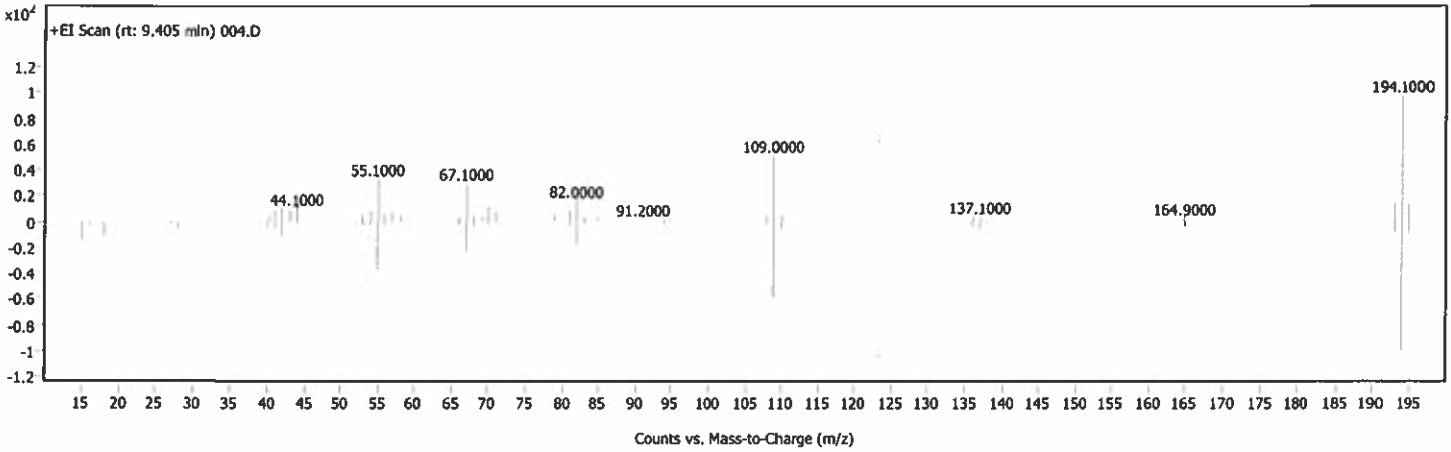
CAFFEINE; C8H10N4O2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CAFFEINE	C8H10N4O2	91.71	aafs.1	MATRIX BLANK after 0.1 mg/L	KM/QC DATA IN XYLAZINE VALIDATION

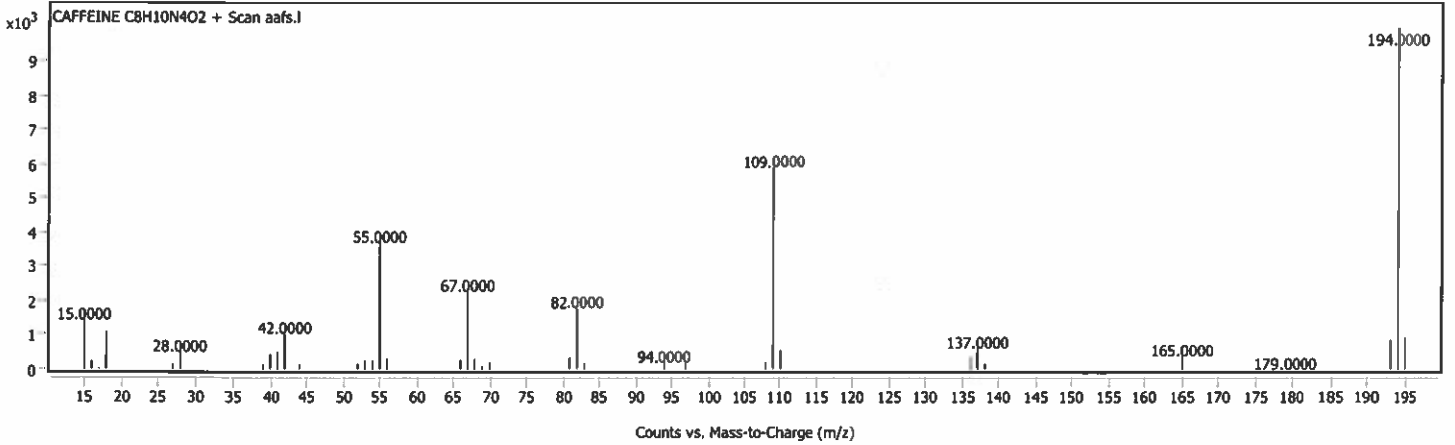
### Observed Spectrum



### Mirror Plot



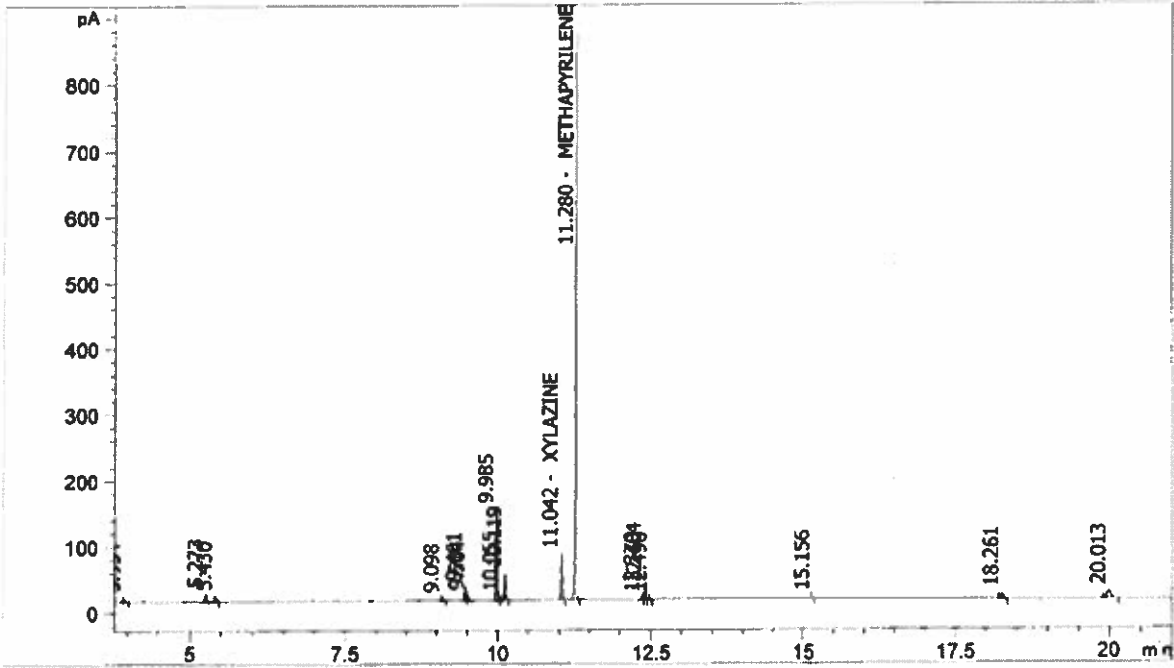
### Library Spectrum



Injection Date: 9/28/2021 2:58:28 PM  
 Sample Name: MATRIX 6 0.10 mg/L - ->  
 Sample Info:

Seq Line: 4  
 Vial 4

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

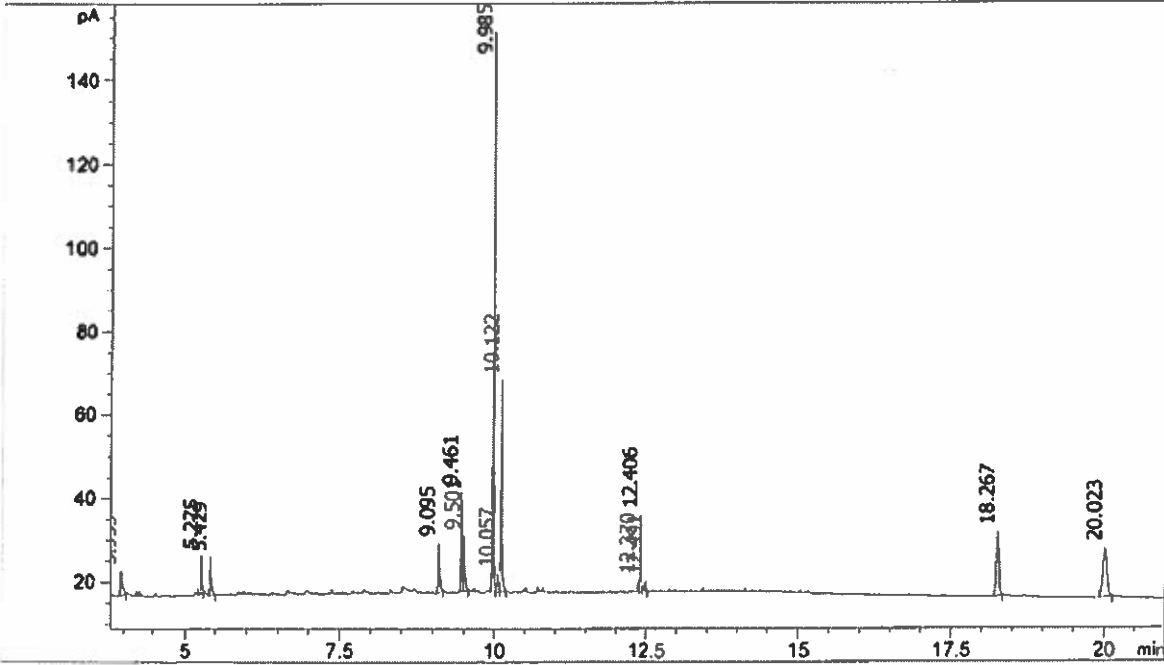


RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.042	11.046	84.260	70.081	0.000000	XYLAZINE RRT-0.9789
11.280	11.278	1015.278	853.502	1.000000	METHAPYRILENE

Injection Date: 9/28/2021 3:22:45 PM  
 Sample Name: MATRIX BLANK after 0.->  
 Sample Info:

Seq Line: 5  
 Vial 5

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



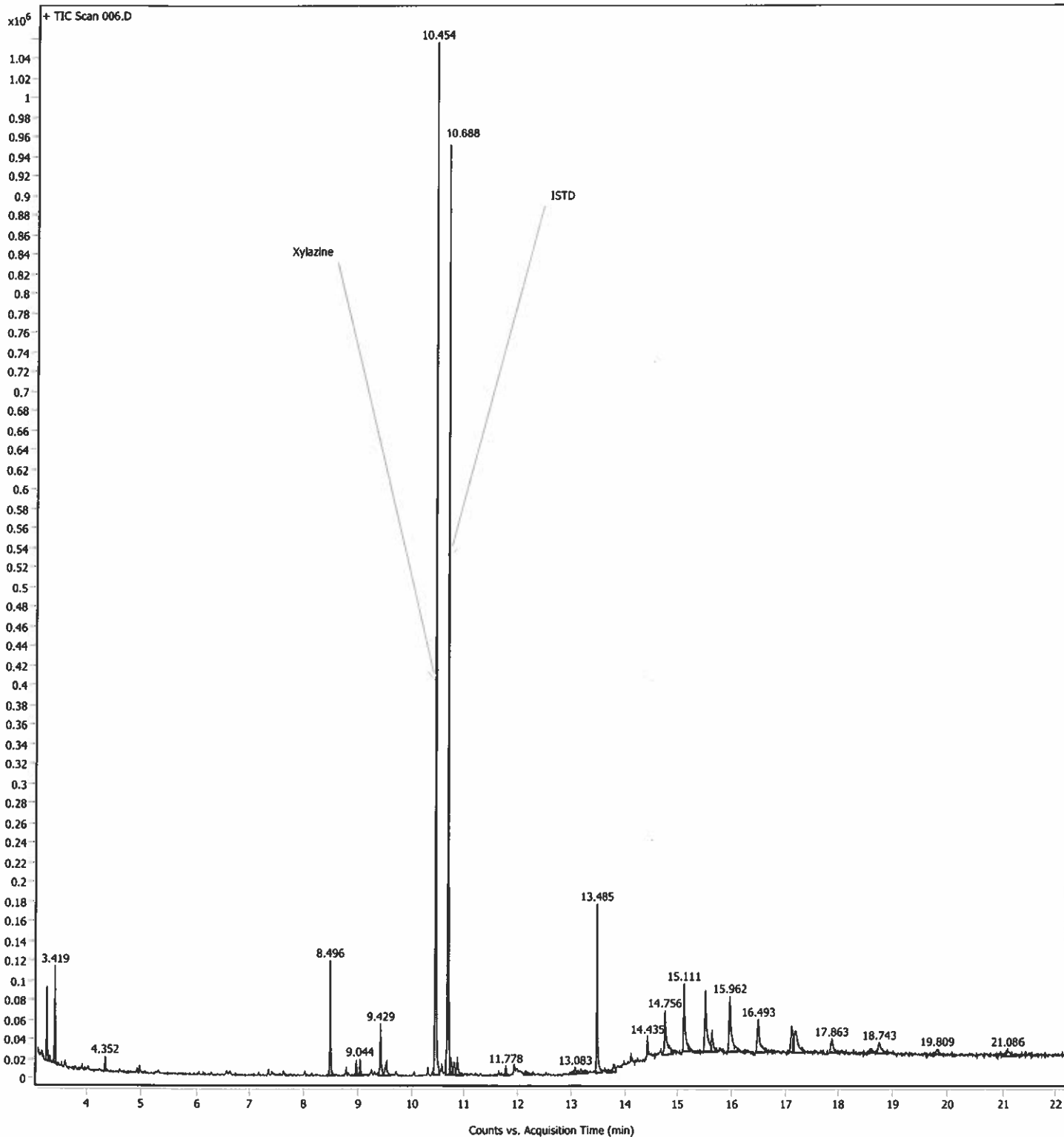
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 2 0.50 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\006.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 3:49:58 PM (UTC-04:00)
Position	32	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

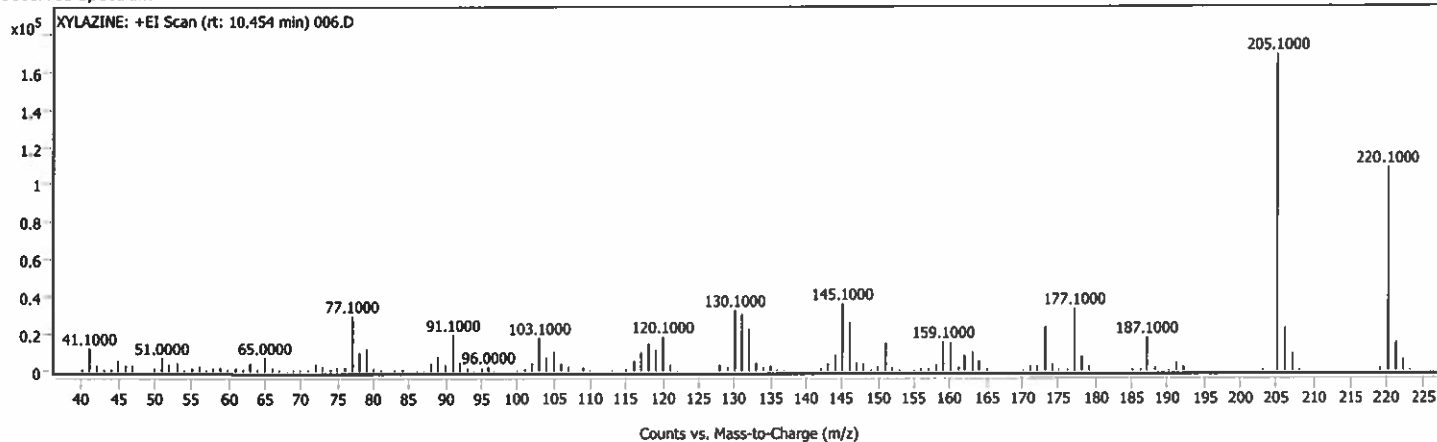
## Sample Spectra

+ Scan (rt: 10.454 min)

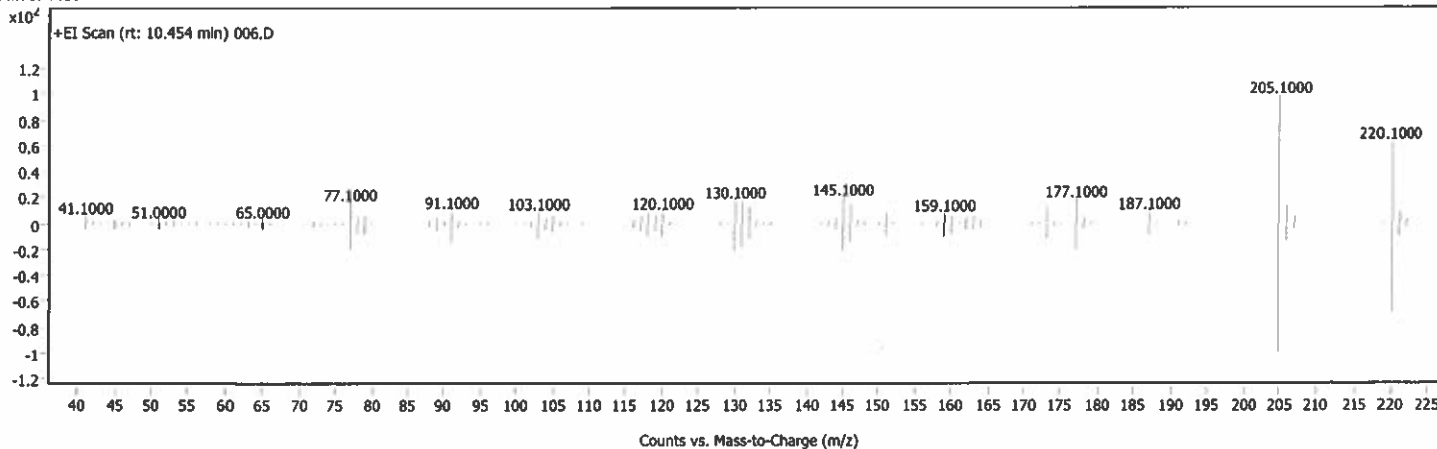
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	97.51	aafs.l	MATRIX 2 0.50 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

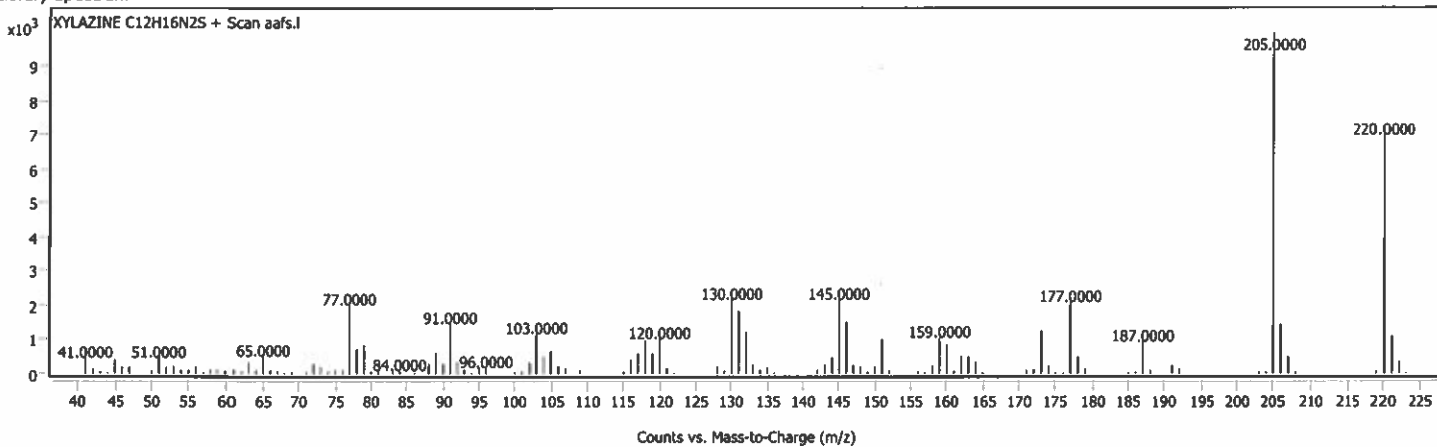
## Observed Spectrum



## Mirror Plot



## Library Spectrum



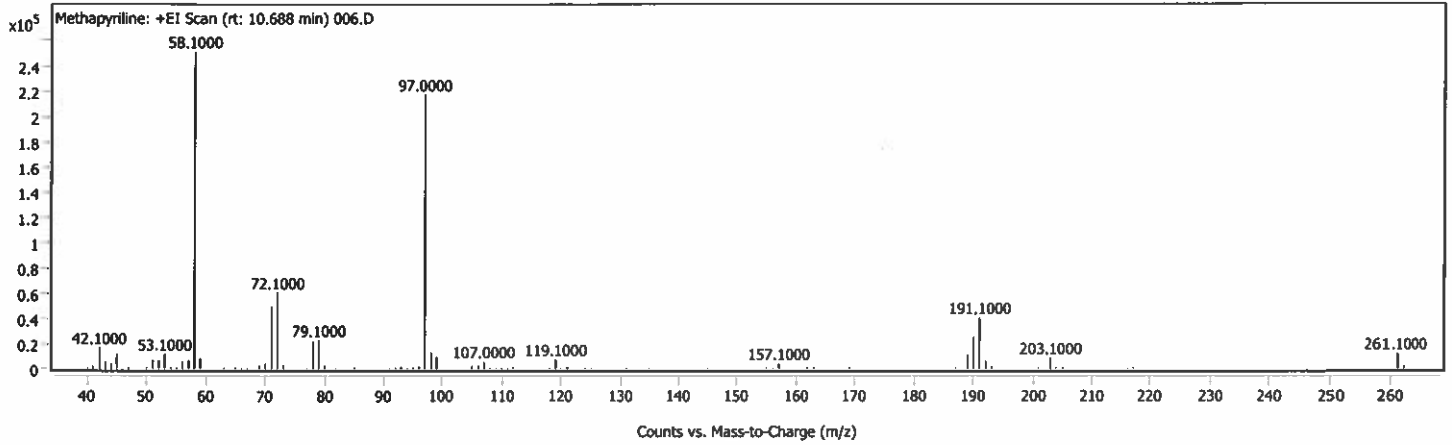
# Analysis Report

+ Scan (rt: 10.688 min)

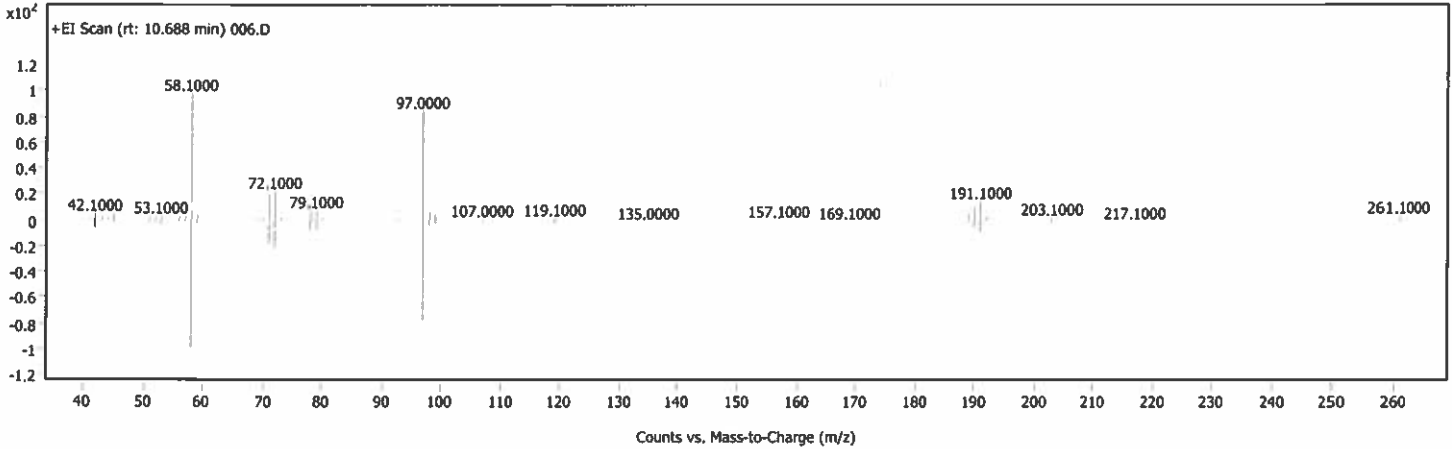
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.43	SWGDRUG 3.3.L	MATRIX 2 0.50 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

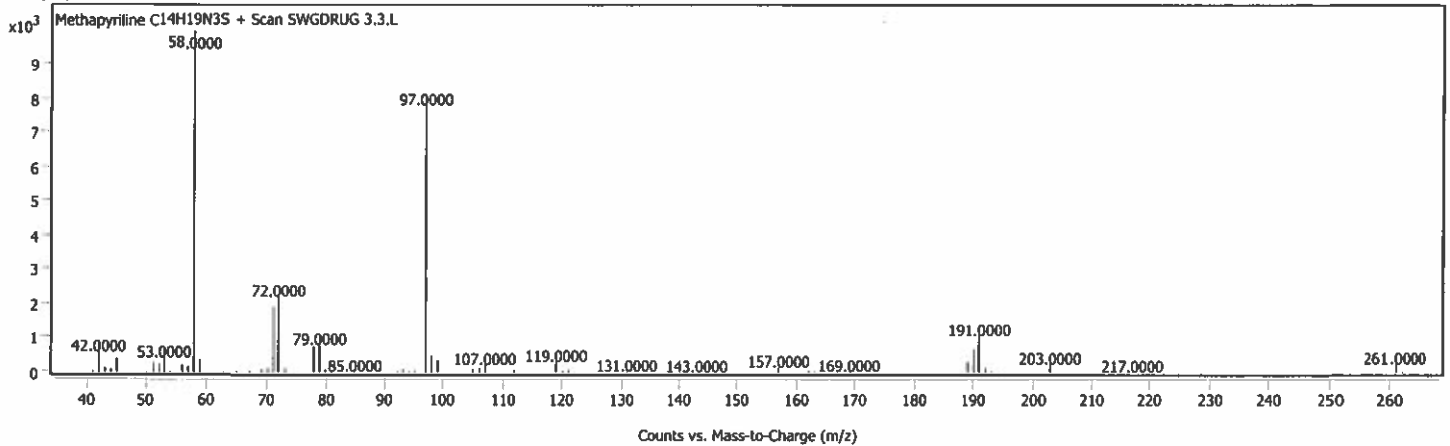
## Observed Spectrum



## Mirror Plot



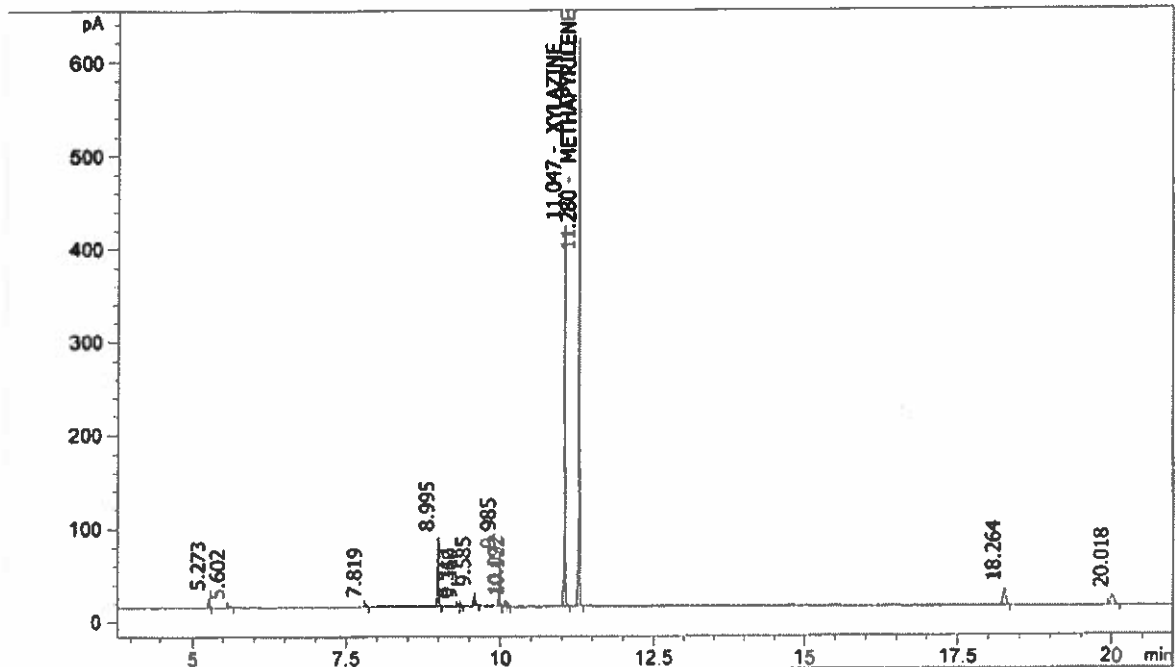
## Library Spectrum



Injection Date: 9/28/2021 4:11:26 PM  
 Sample Name: MATRIX 2 0.50 mg/L - ->  
 Sample Info:

Seq Line: 7  
 Vial 7

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



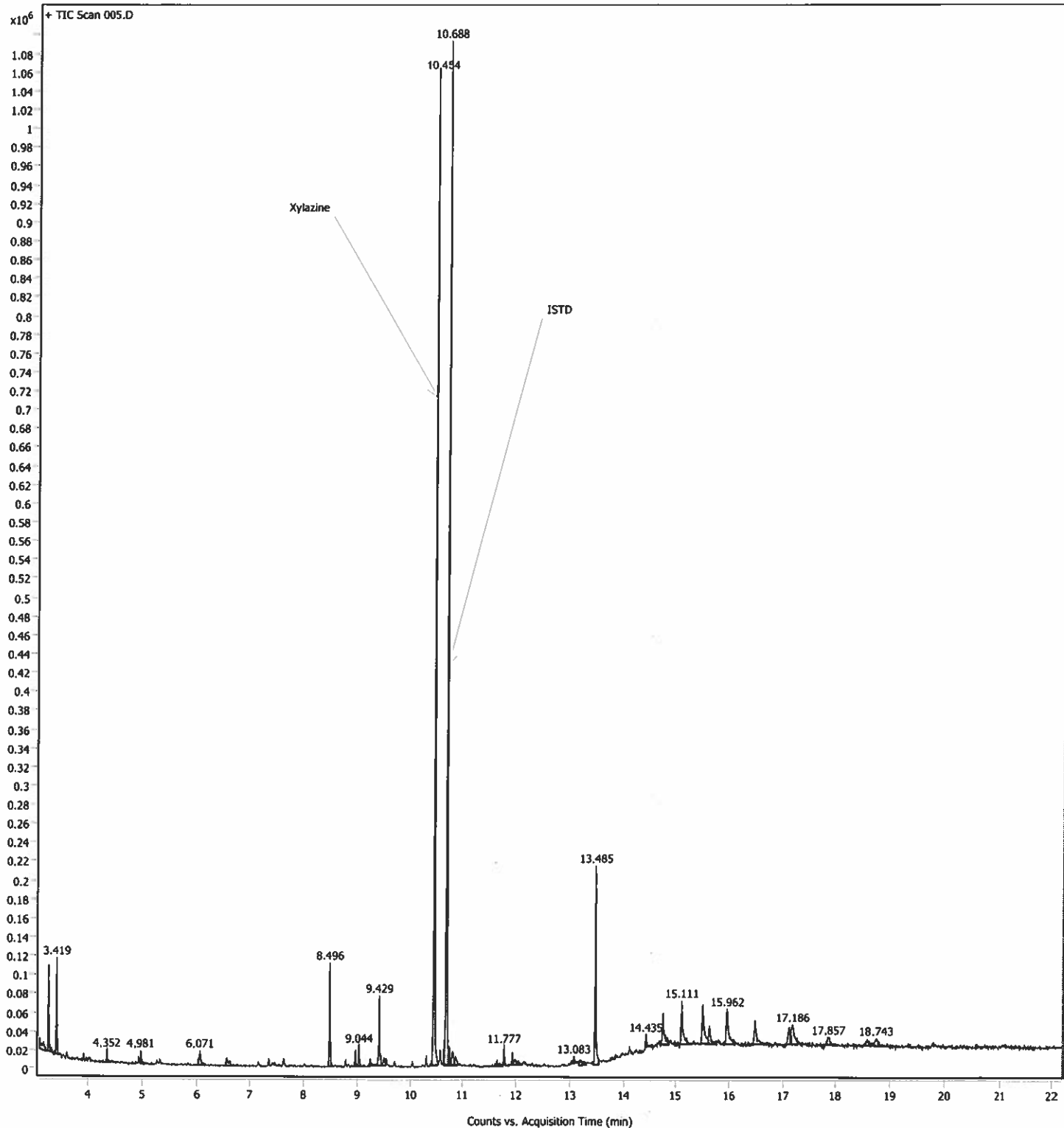
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.047	11.046	474.613	407.068	0.000000	XYLAZINE RRT - 0.9793
11.280	11.278	697.688	603.915	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 0.50 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\005.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 3:24:07 PM (UTC-04:00)
Position	31	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

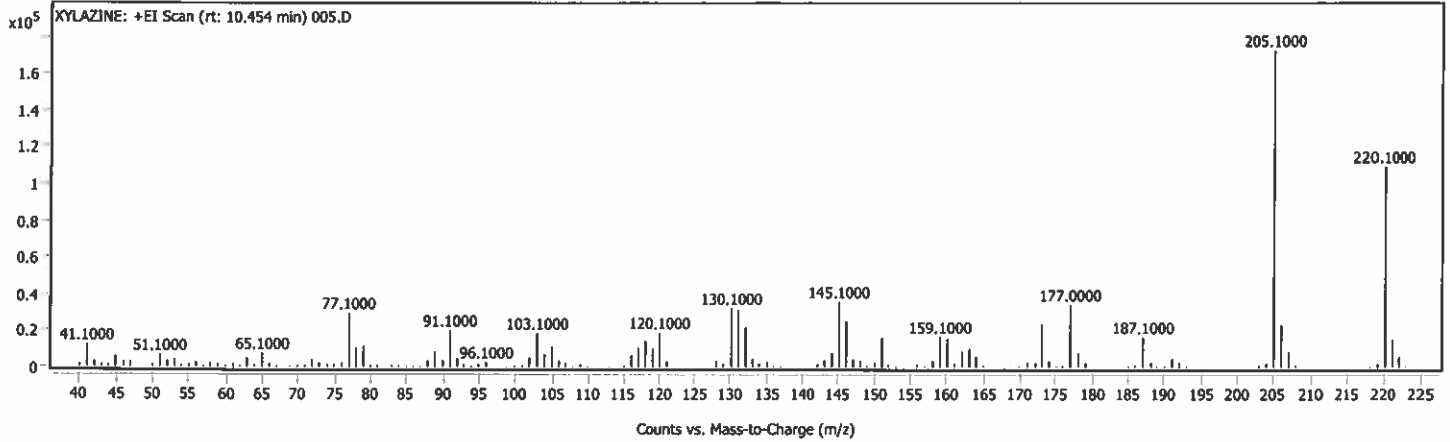
## Sample Spectra

+ Scan (rt: 10.454 min)

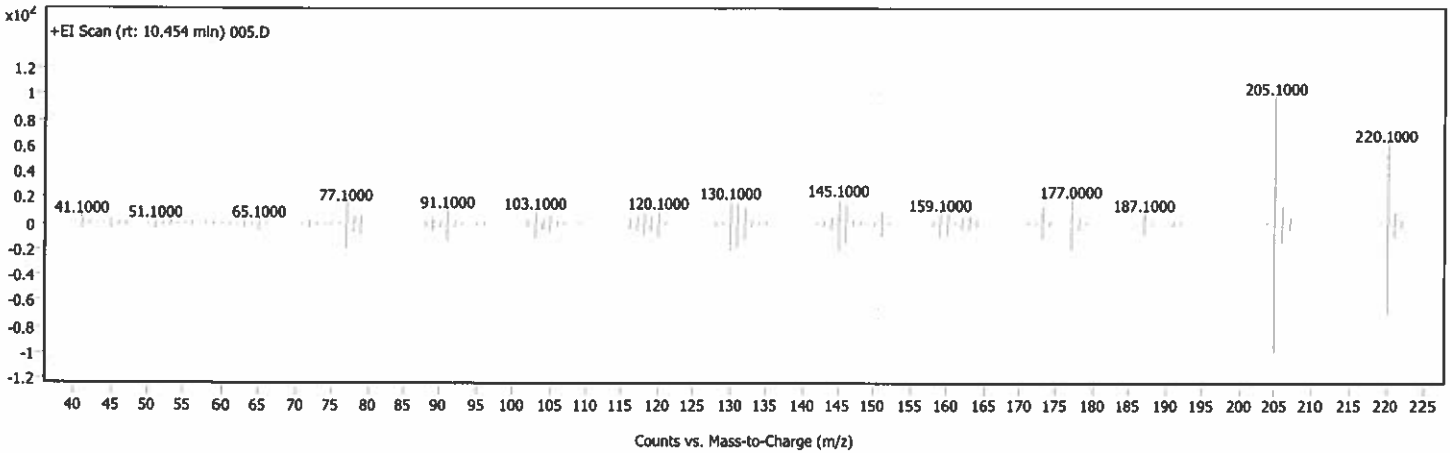
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	97.56	aafs.l	MATRIX 1 0.50 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

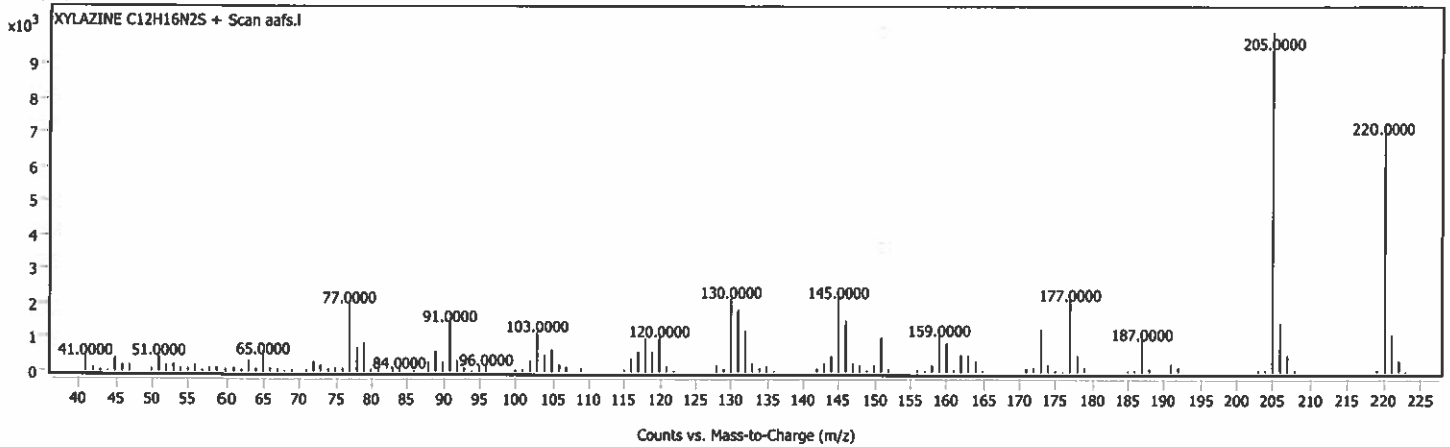
### Observed Spectrum



### Mirror Plot



### Library Spectrum



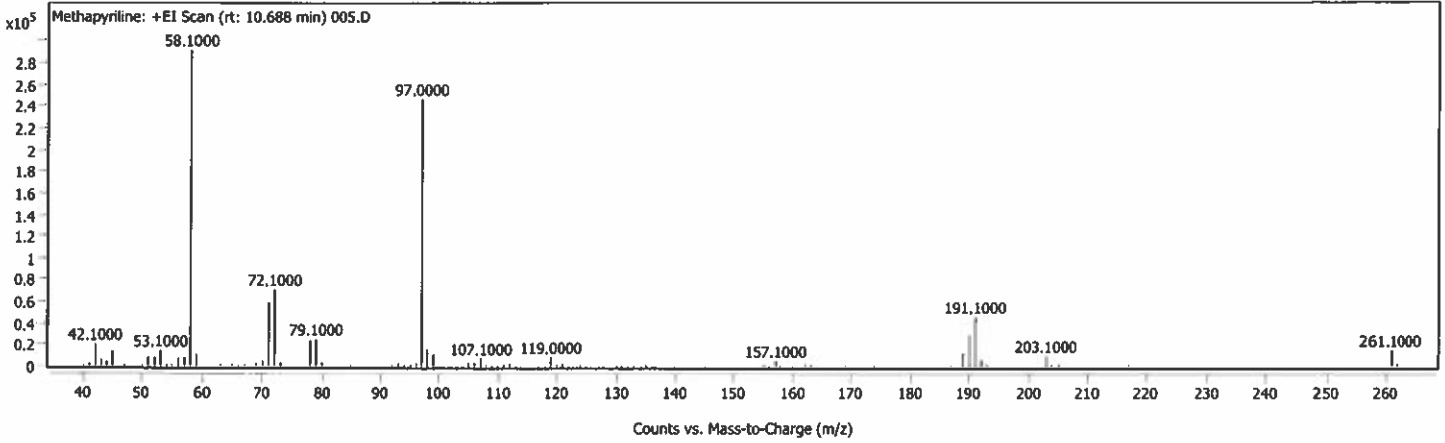
# Analysis Report

+ Scan (rt: 10.688 min)

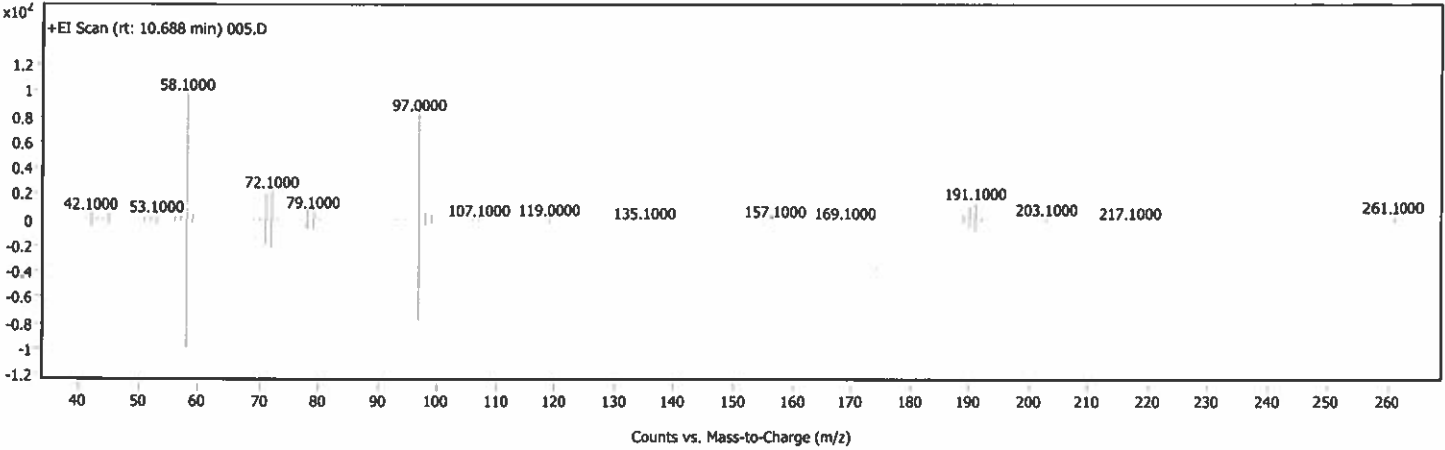
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.36	SWGDRUG 3.3.L	MATRIX 1 0.50 mg/L - BLD	KM/QC DATA IN XLAZINE VALIDATION

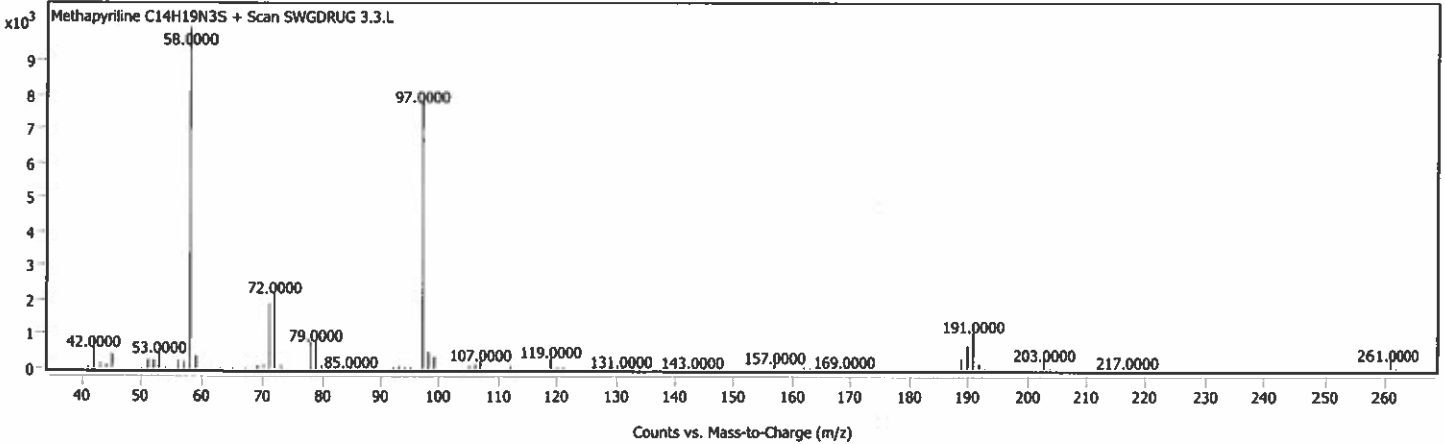
## Observed Spectrum



## Mirror Plot



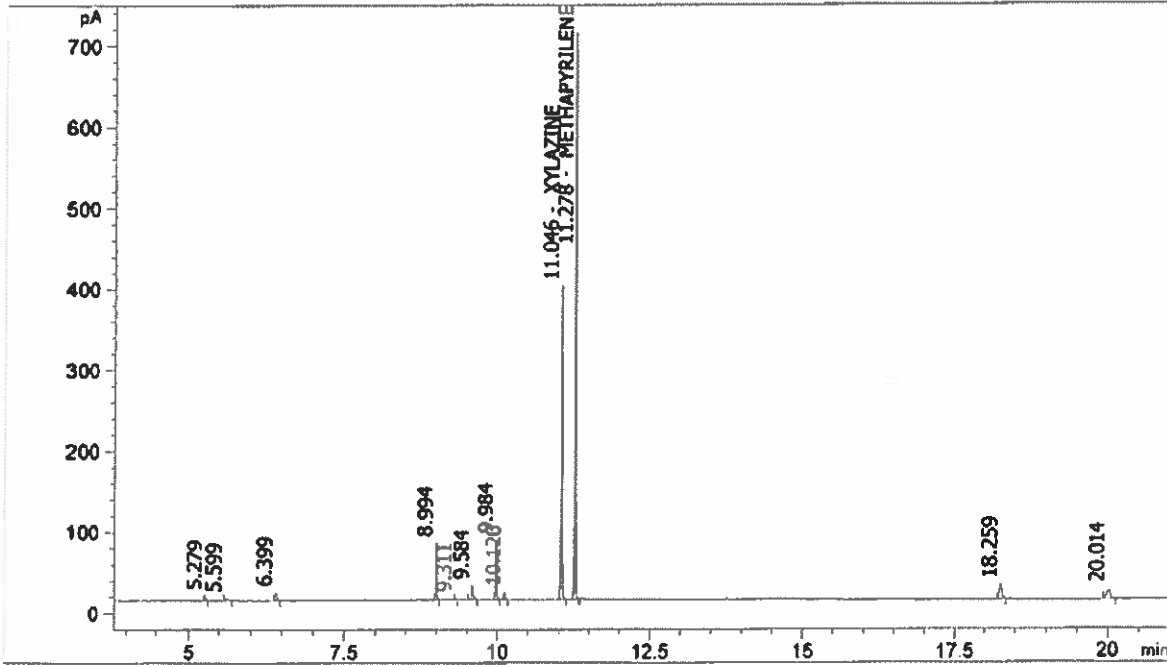
## Library Spectrum



Injection Date: 9/28/2021 3:47:05 PM  
 Sample Name: MATRIX 1 0.50 mg/L - ->  
 Sample Info:

Seq Line: 6  
 Vial 6

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.046	11.046	472.345	386.736	0.000000	XYLAZINE RRT - 0.9794
11.278	11.278	820.587	696.468	1.000000	METHAPYRILENE

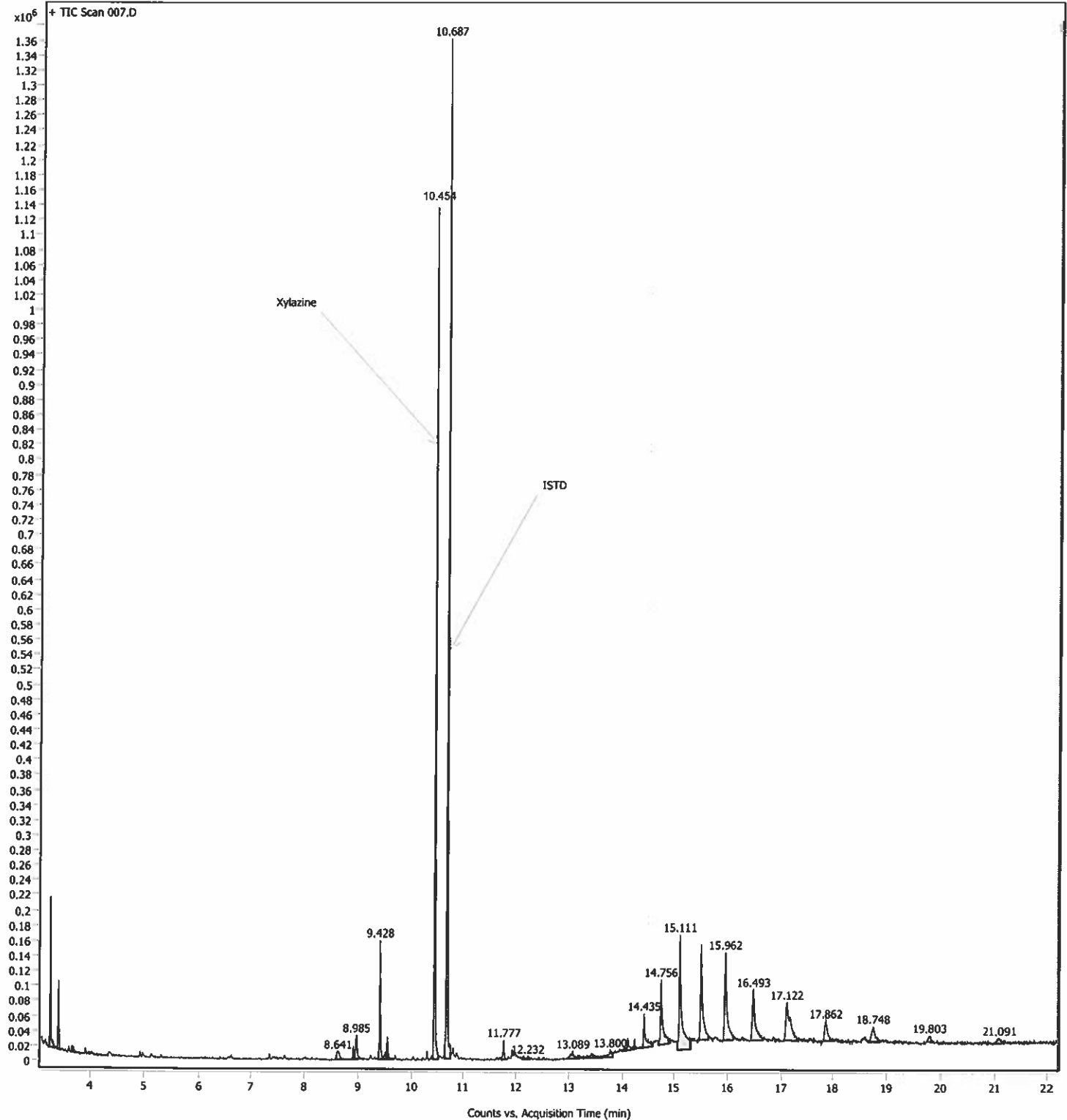
\*RTs for Xylazine + Methapyrilene set from sample

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 0.50 mg/L - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\007.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 4:15:44 PM (UTC-04:00)
Position	33	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

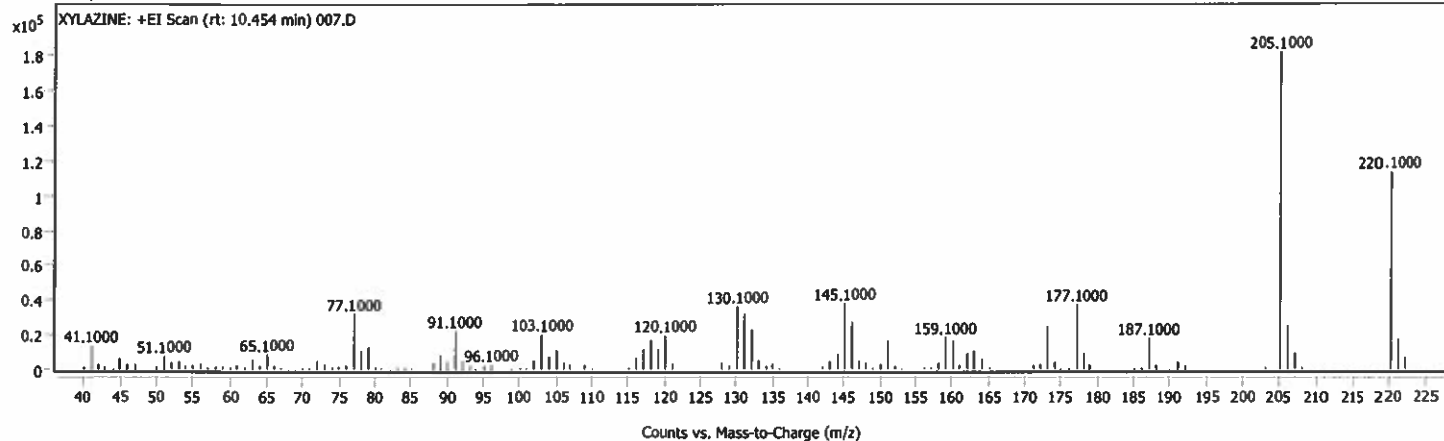
## Sample Spectra

+ Scan (rt: 10.454 min)

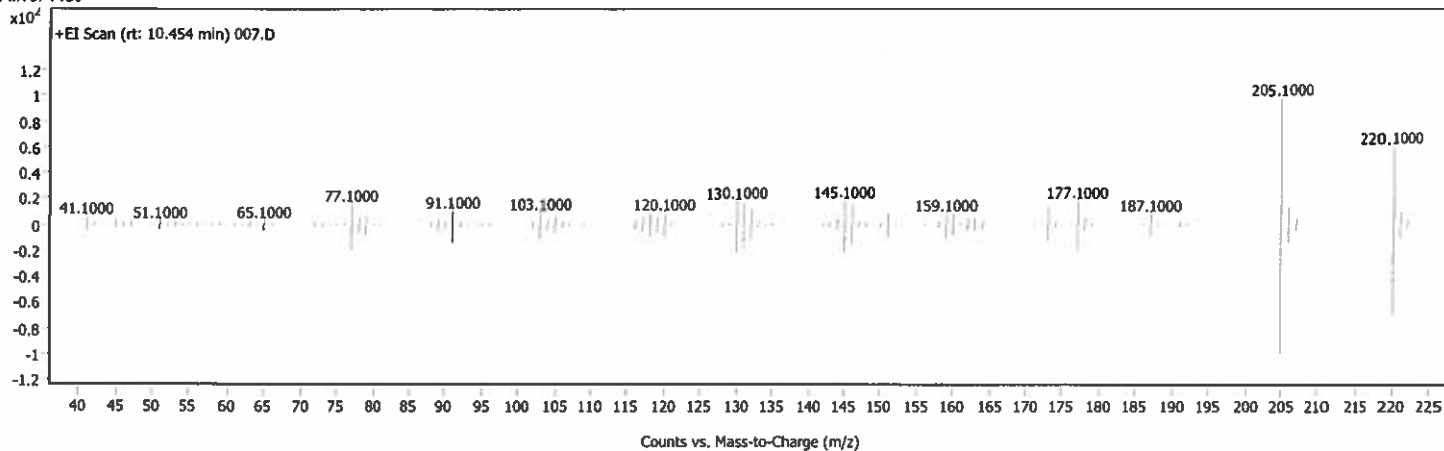
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	97.65	aafs.l	MATRIX 6 0.50 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

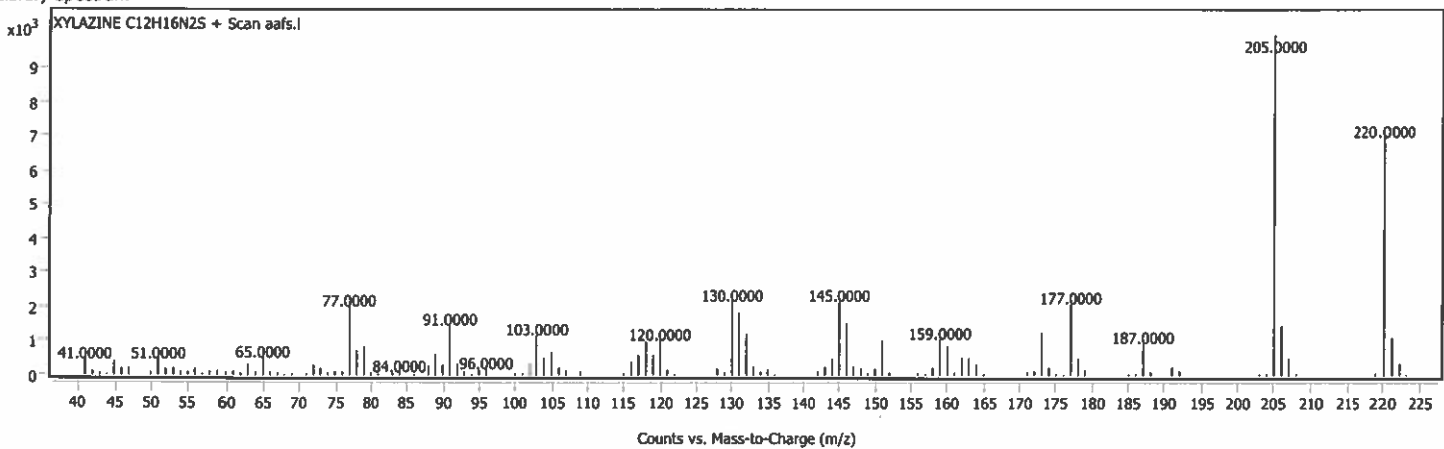
## Observed Spectrum



## Mirror Plot



## Library Spectrum



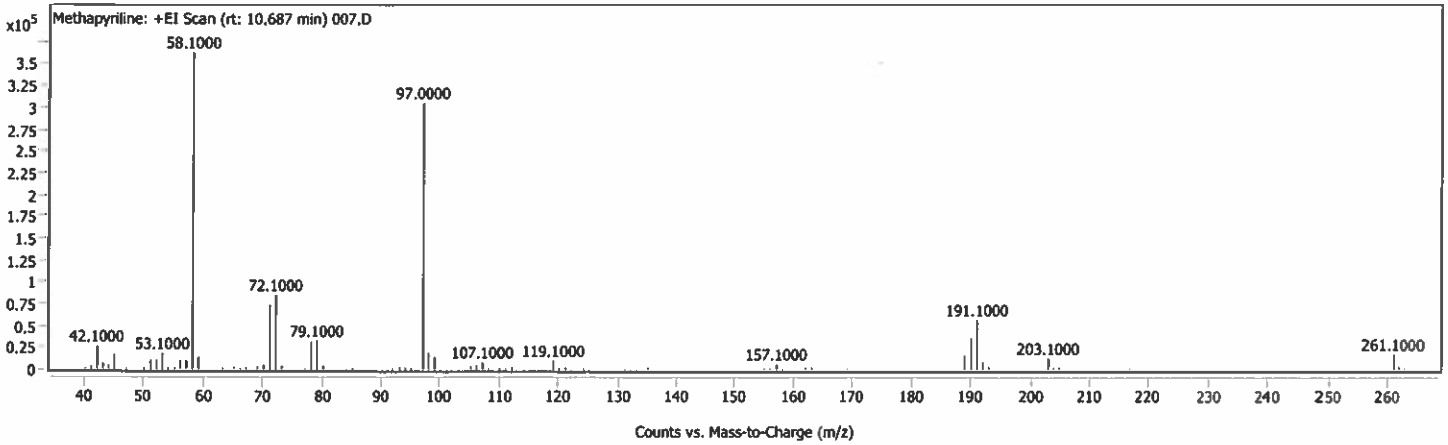
# Analysis Report

+ Scan (rt: 10.687 min)

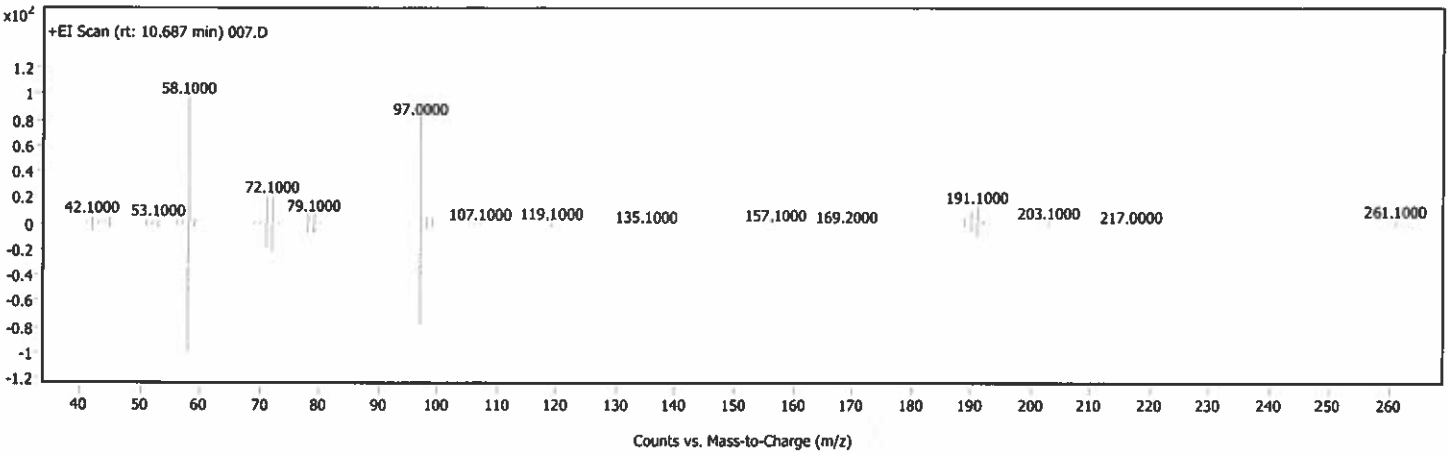
**Methapyriline; C14H19N3S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.75	SWGDRUG 3.3.L	MATRIX 6 0.50 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

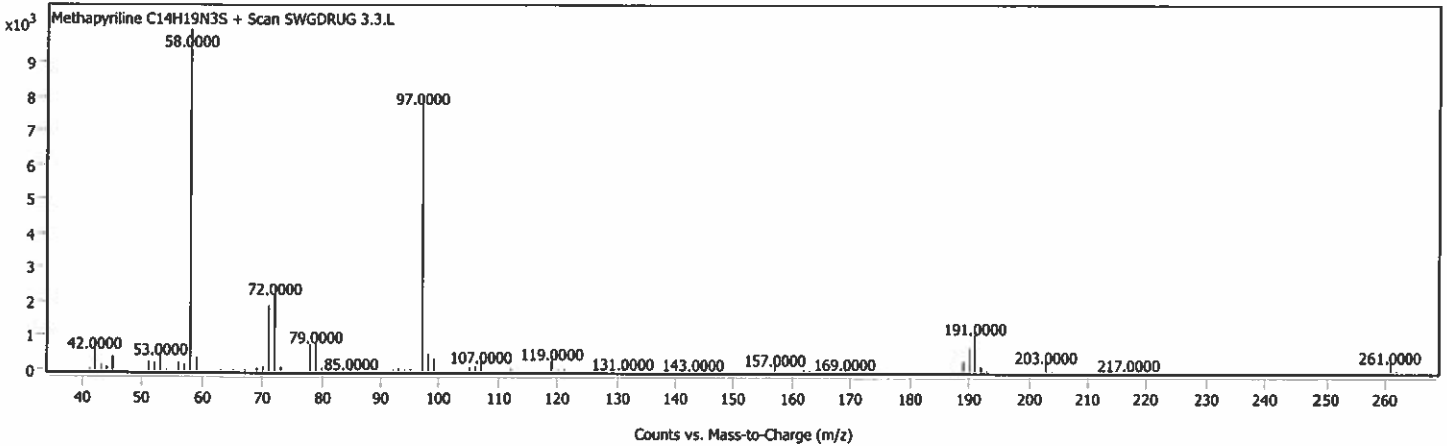
### Observed Spectrum



### Mirror Plot



### Library Spectrum

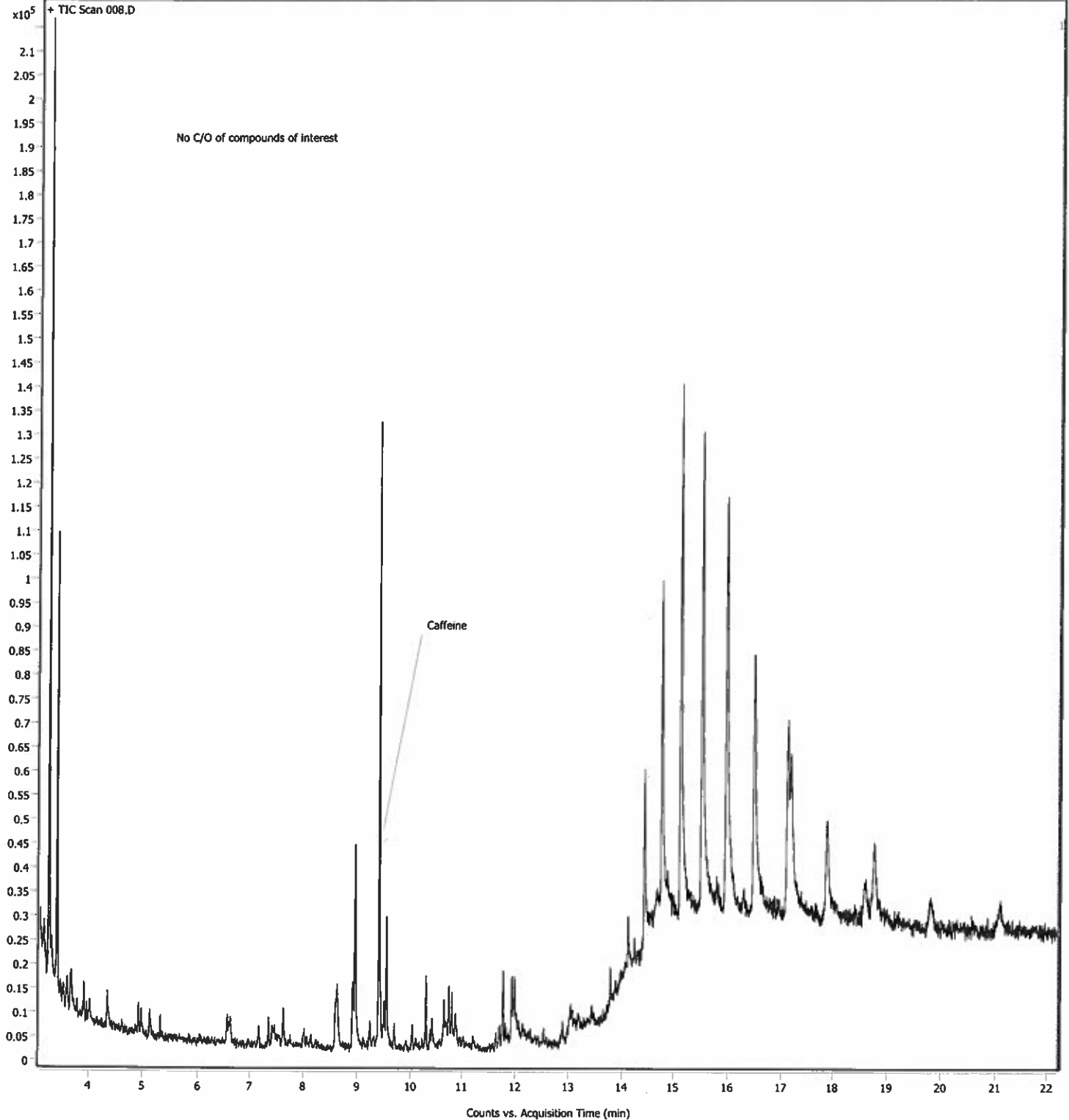


# Analysis Report

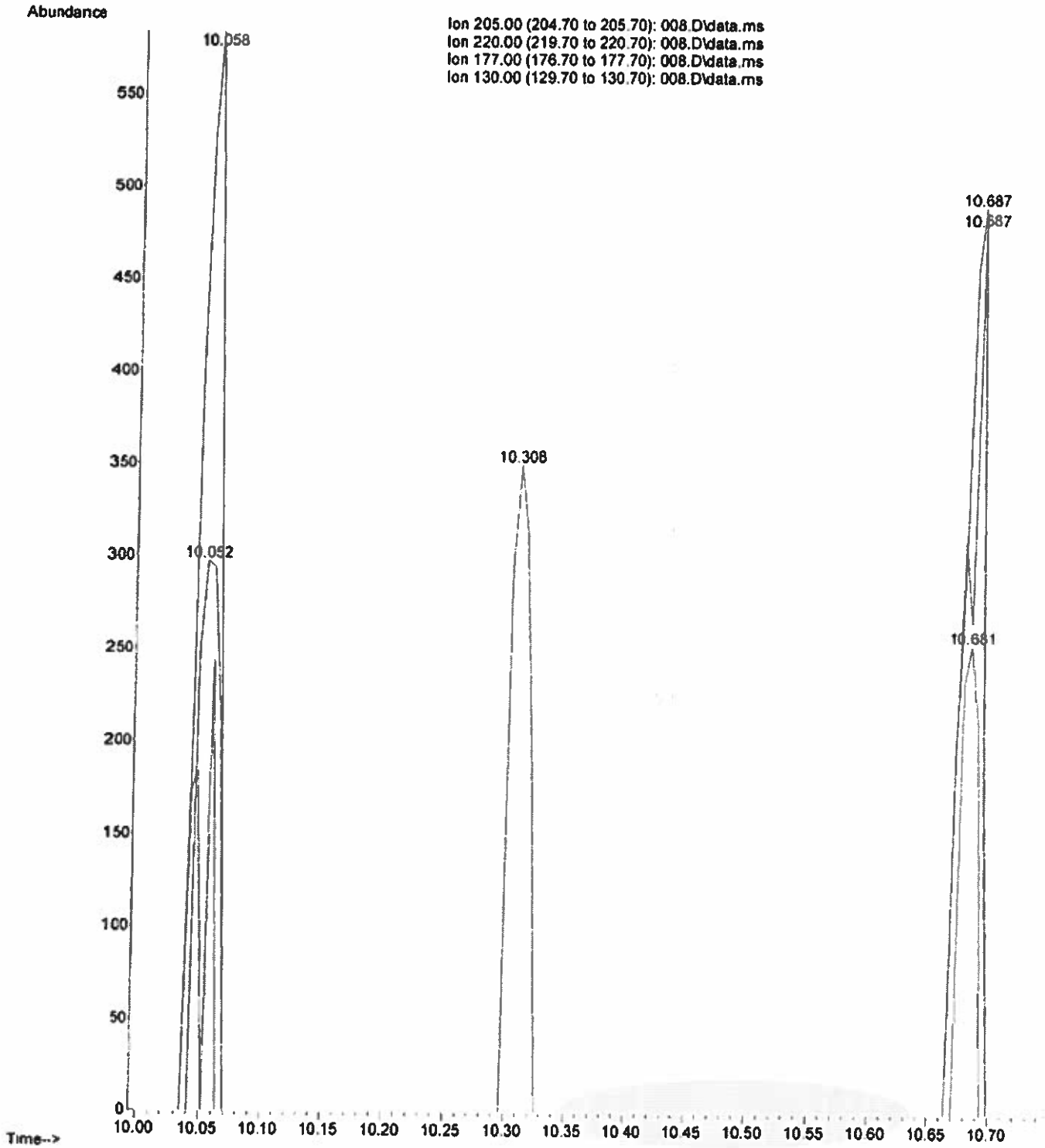
## Sample Information

Sample Name	MATRIX BLANK after 0.5 mg/L	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\008.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 4:41:32 PM (UTC-04:00)
Position	34	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



File :C:\Users\TOX\Desktop\092821\008.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 28 Sep 2021 16:41 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK after 0.5 mg/L  
Misc Info :  
Vial Number: 34





# Analysis Report

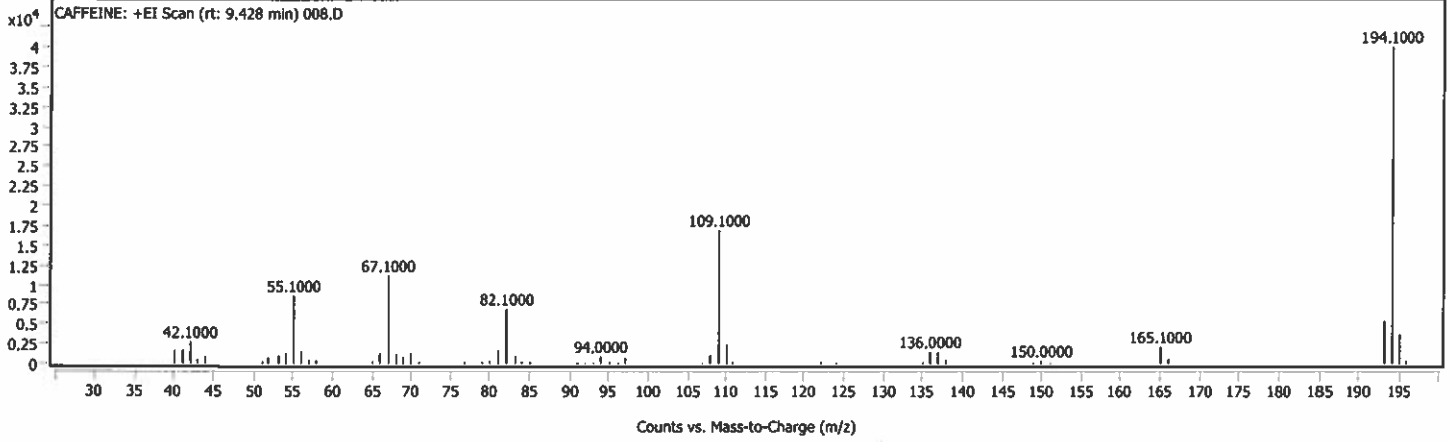
## Sample Spectra

+ Scan (rt: 9.428 min)

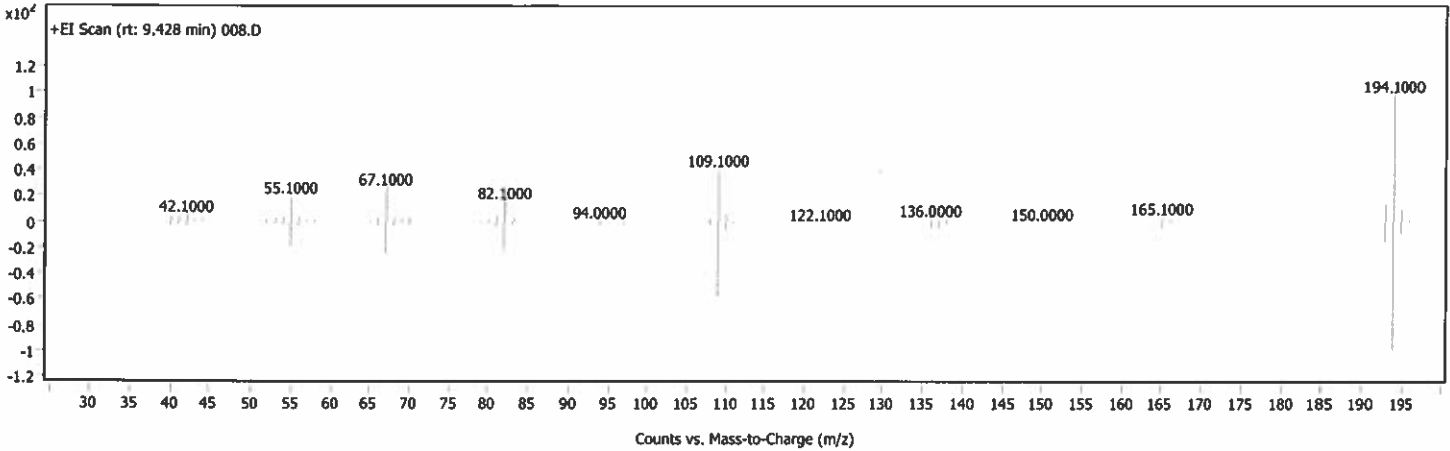
CAFFEINE; C8H10N4O2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CAFFEINE	C8H10N4O2	98.12	aafs.l	MATRIX BLANK after 0.5 mg/L	KM/QC DATA IN XYLAZINE VALIDATION

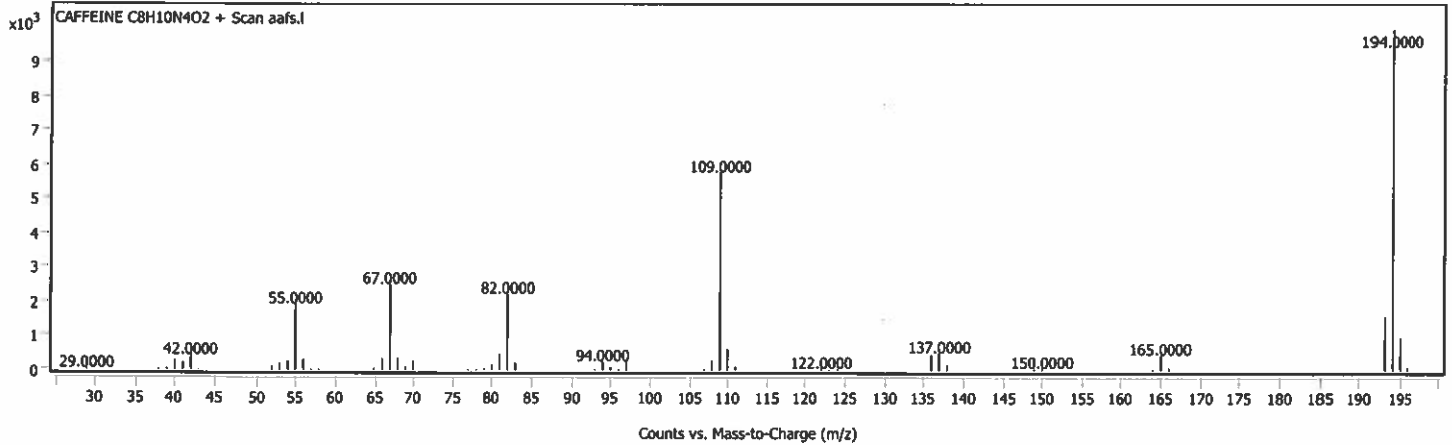
## Observed Spectrum



## Mirror Plot

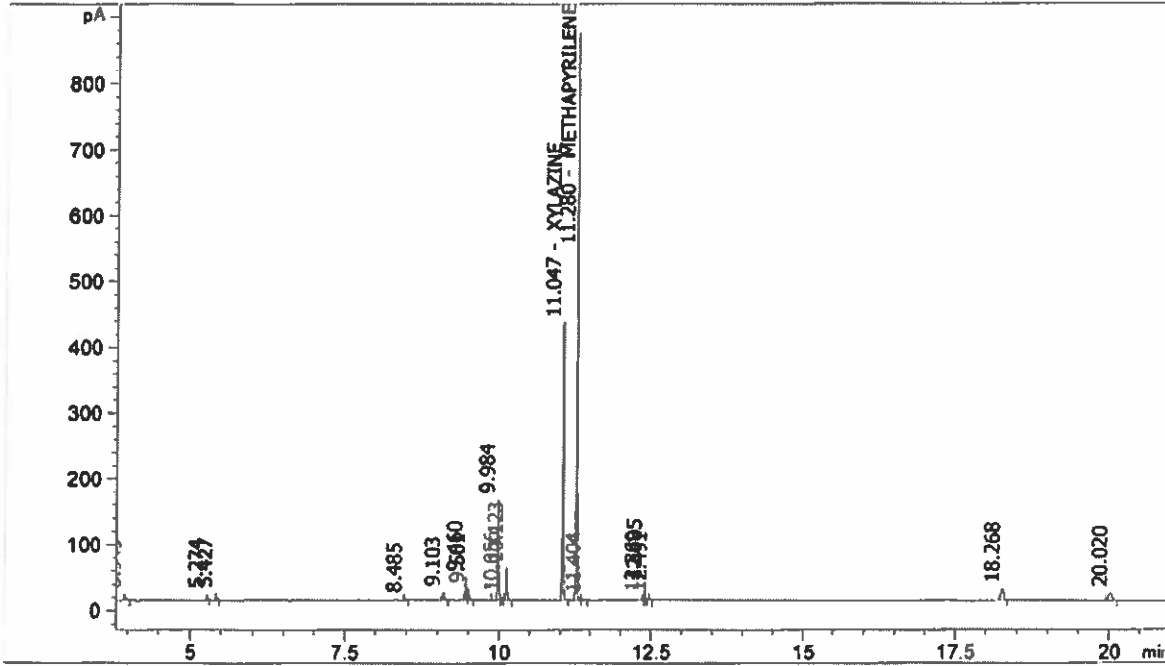


## Library Spectrum



Injection Date: 9/28/2021 4:35:44 PM Seq Line: 8  
 Sample Name: MATRIX 6 0.50 mg/L - -> Vial 8  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

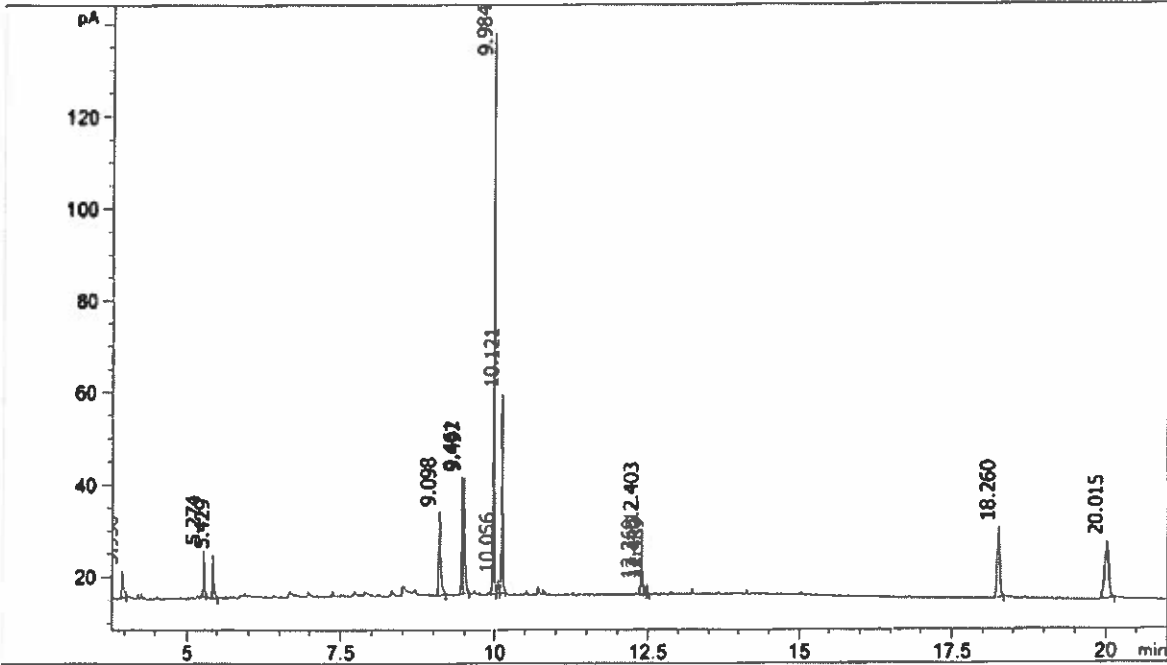


RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.047	11.046	509.258	420.059	0.000000	XYLAZINE RRT-0.9793
11.280	11.278	1025.839	859.222	1.000000	METHAPYRILENE

Injection Date: 9/28/2021 5:00:01 PM  
Sample Name: MATRIX BLANK after 0.->  
Sample Info:

Seq Line: 9  
Vial 9

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
KM/QC DATA IN XYLAZINE VALIDATION



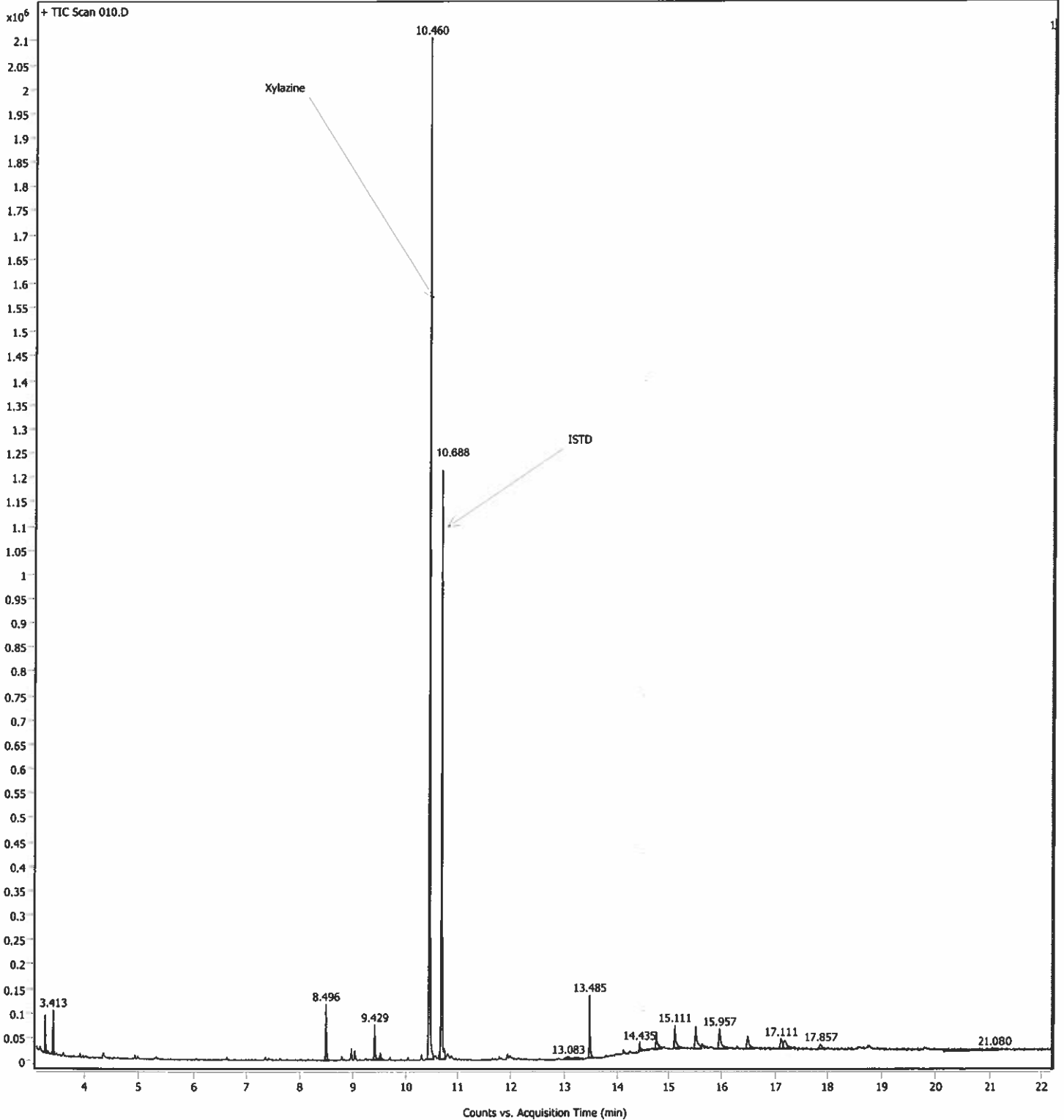
RT (min)	Exp RT (min)	Area	Height	Amount (mg/L)	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 2 1.0 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\010.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 5:33:13 PM (UTC-04:00)
Position	36	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

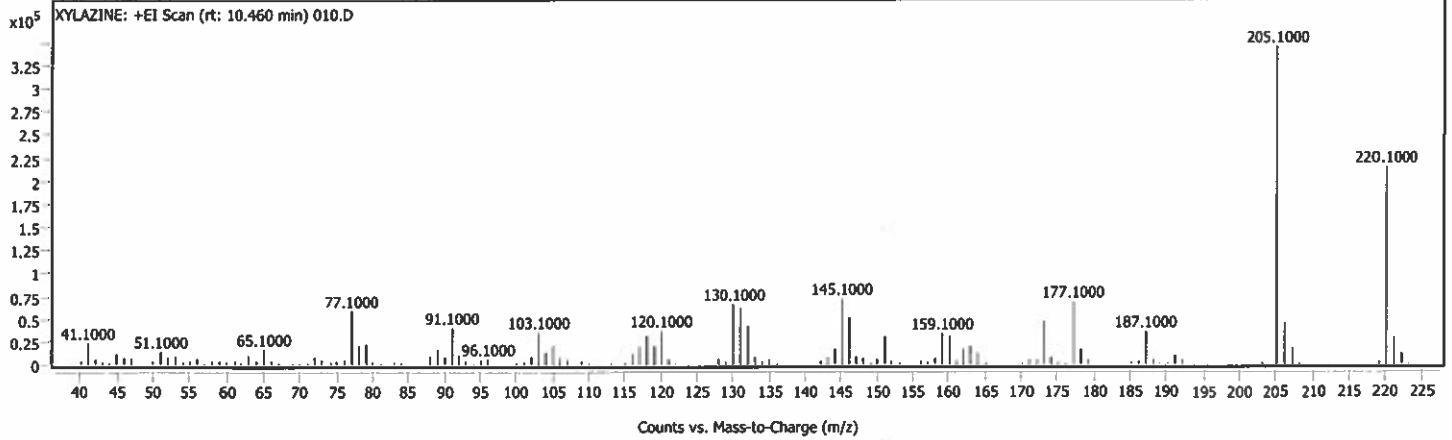
## Sample Spectra

+ Scan (rt: 10.460 min)

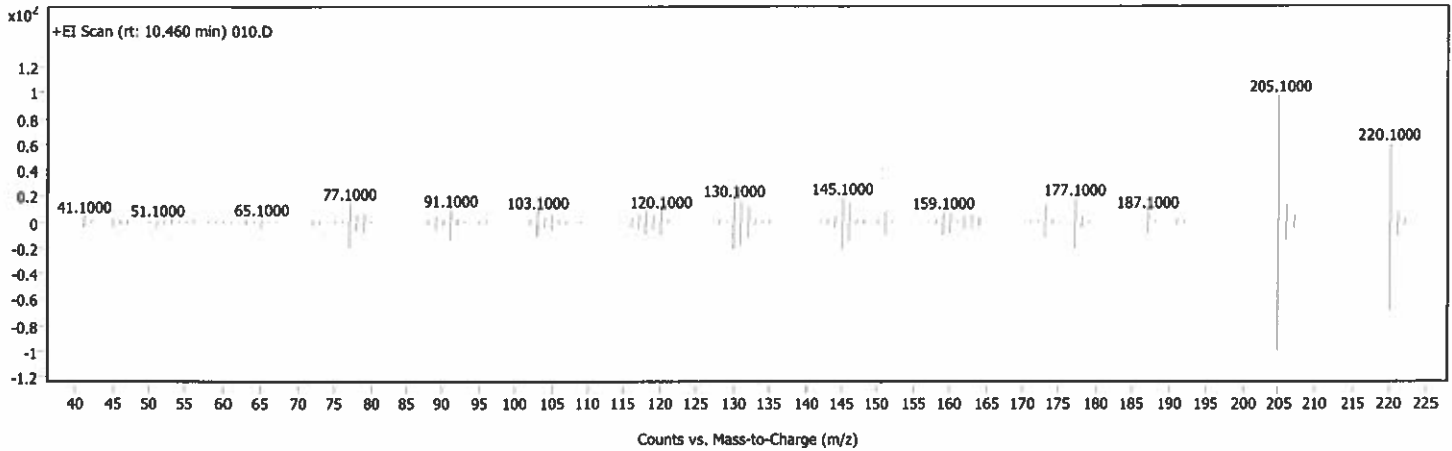
XYLAZINE; C<sub>12</sub>H<sub>16</sub>N<sub>2</sub>S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C <sub>12</sub> H <sub>16</sub> N <sub>2</sub> S	97.67	aafs.l	MATRIX 2 1.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

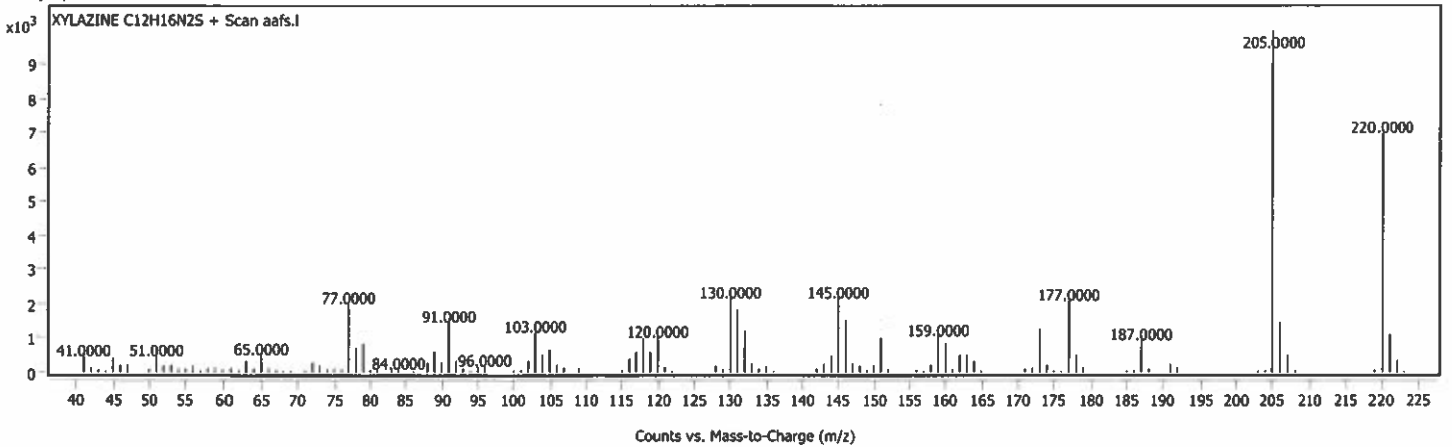
## Observed Spectrum



## Mirror Plot



## Library Spectrum



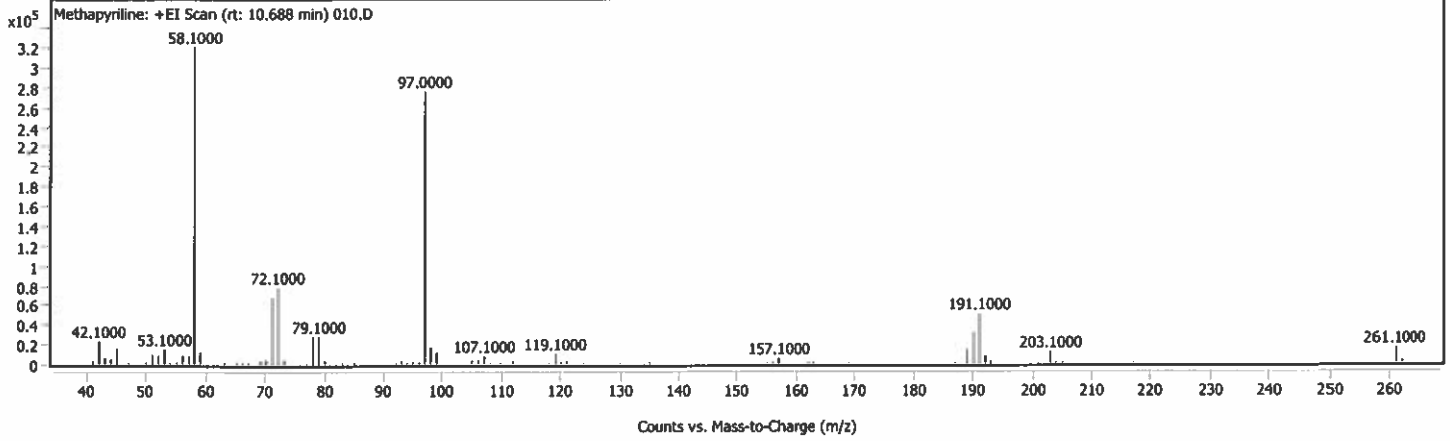
# Analysis Report

+ Scan (rt: 10.688 min)

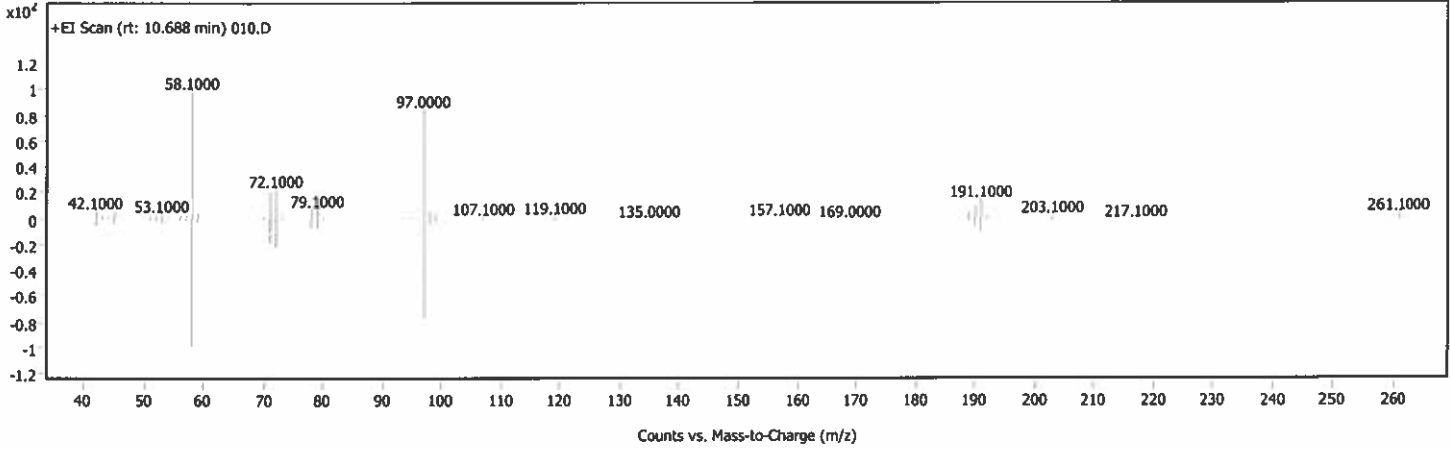
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.45	SWGDRUG 3.3.L	MATRIX 2 1.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

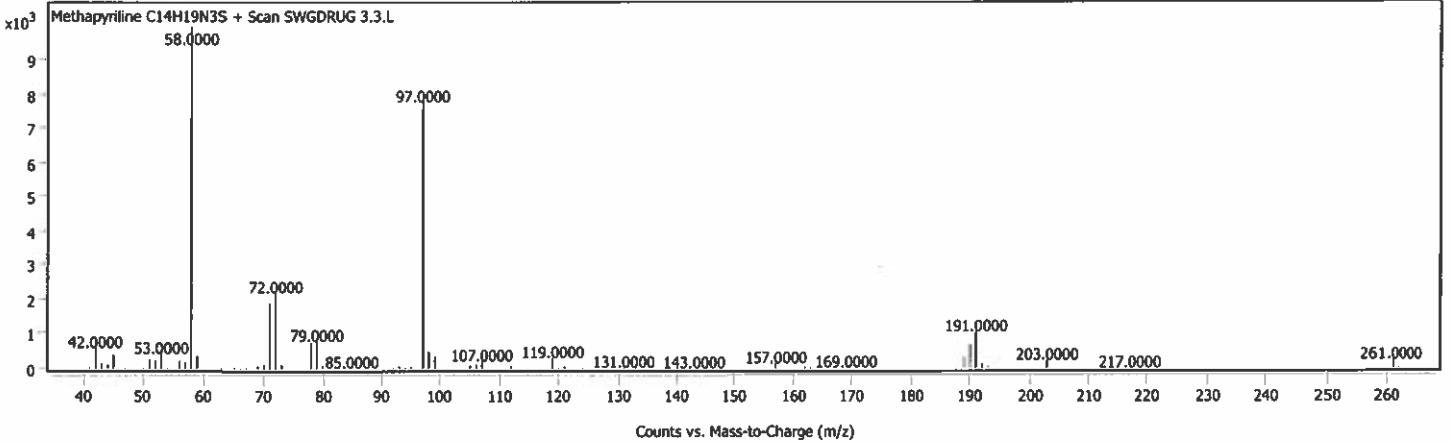
### Observed Spectrum



### Mirror Plot



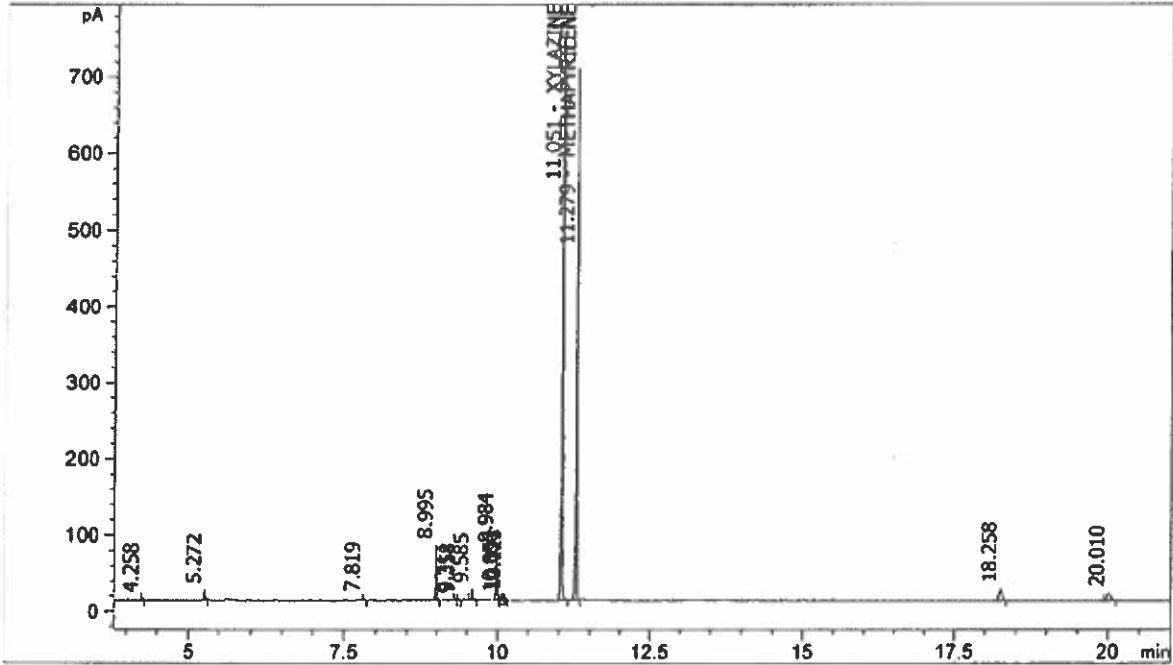
### Library Spectrum



Injection Date: 9/28/2021 5:48:39 PM  
 Sample Name: MATRIX 2 1.0 mg/L - BLD  
 Sample Info:

Seq Line: 11  
 Vial 11

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



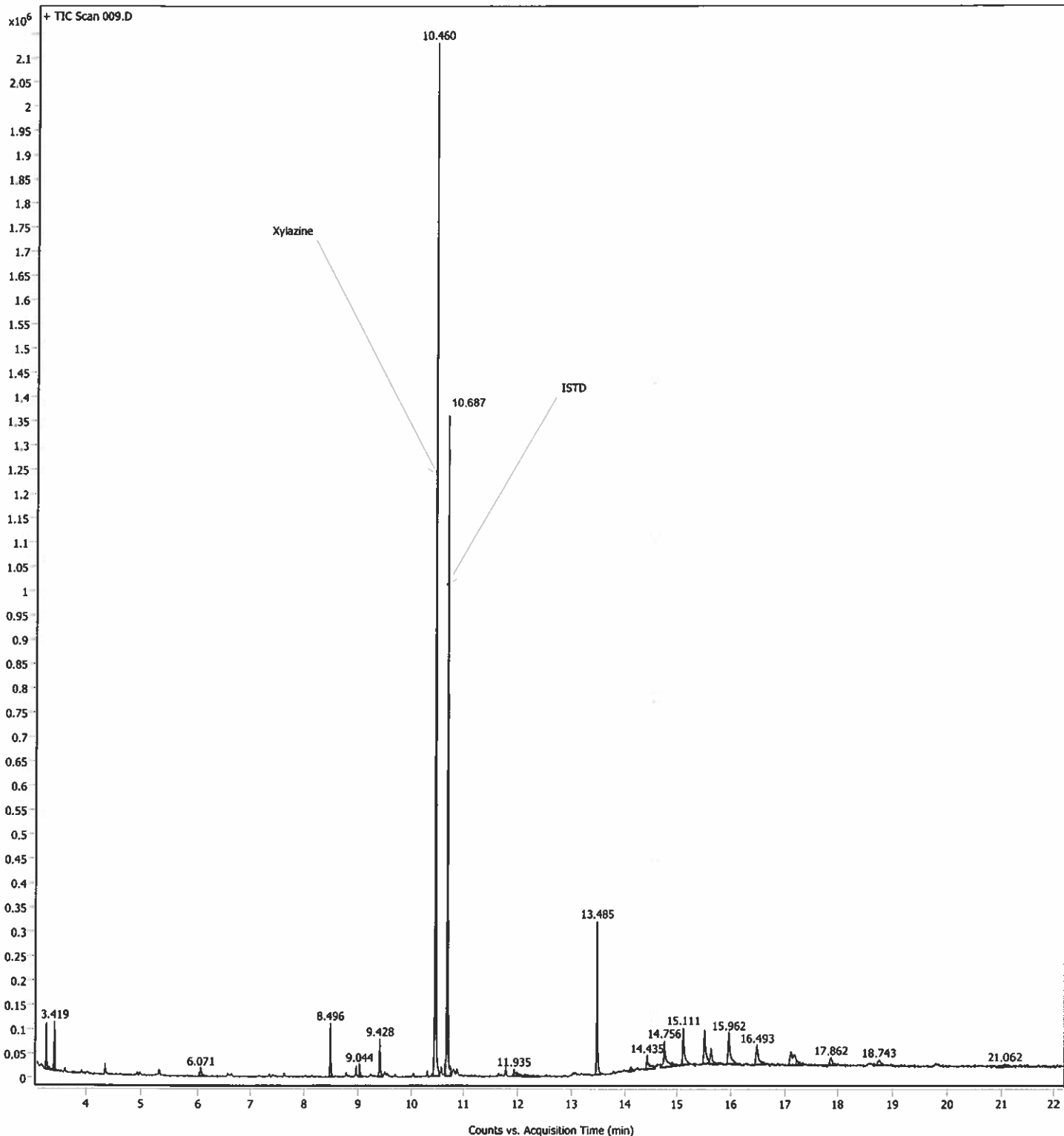
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.051	11.046	934.438	740.777	0.000000	XYLAZINE RRT -0.9798
11.279	11.278	810.992	695.724	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 1.0 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASE5\092821\009.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 5:07:26 PM (UTC-04:00)
Position	35	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

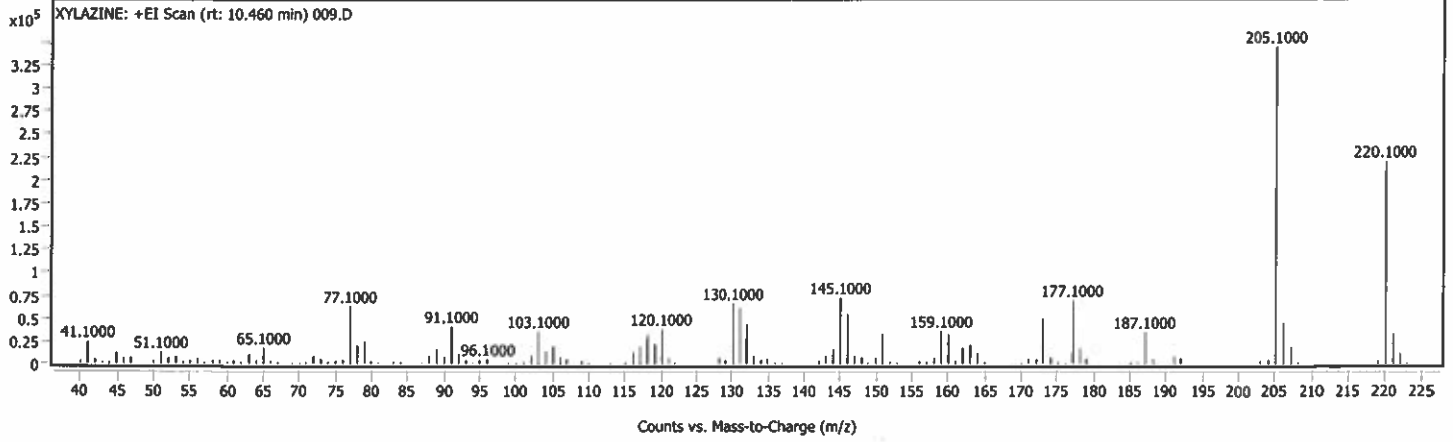
## Sample Spectra

+ Scan (rt: 10.460 min)

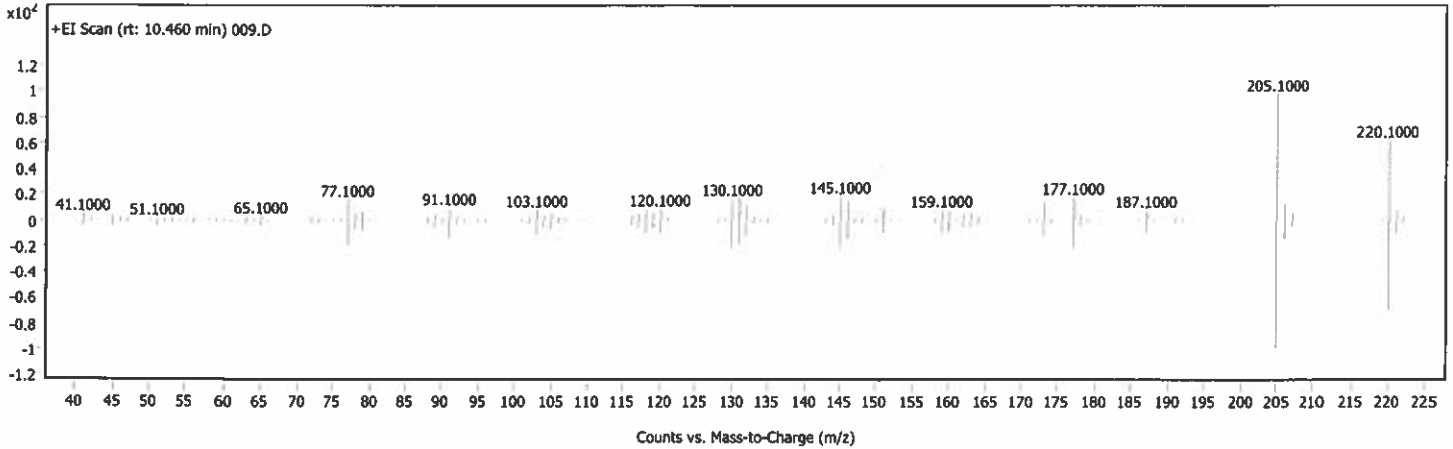
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.75	aafs.l	MATRIX 1 1.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

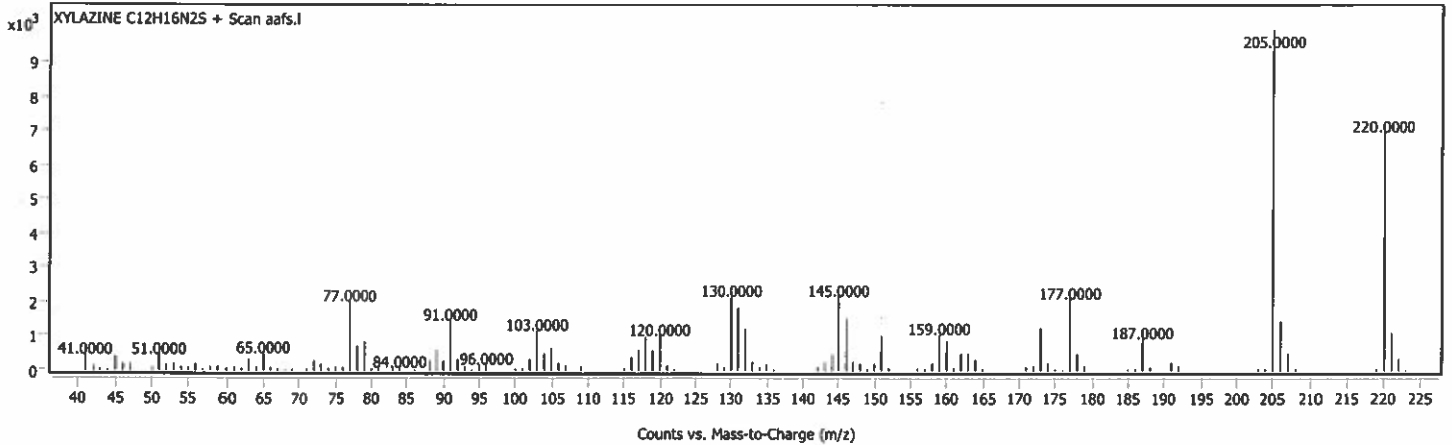
### Observed Spectrum



### Mirror Plot



### Library Spectrum



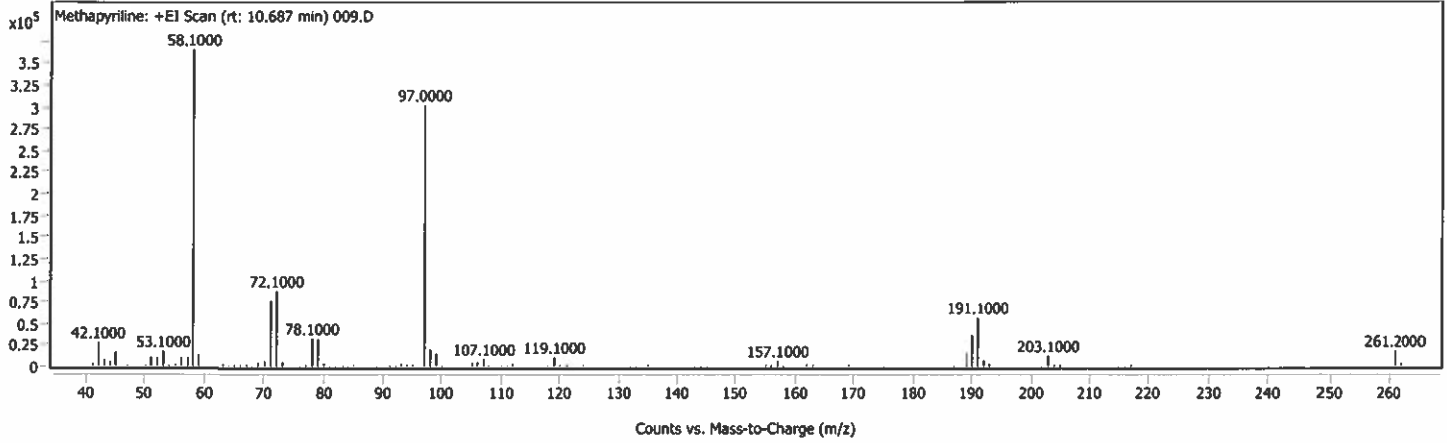
# Analysis Report

+ Scan (rt: 10.687 min)

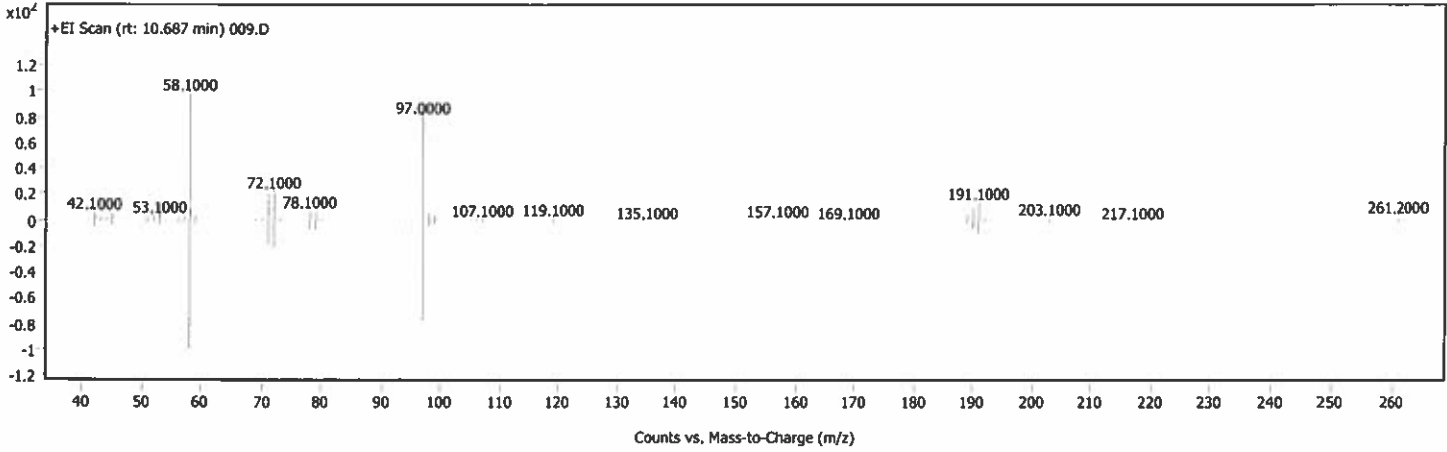
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.76	SWGDRUG 3.3.L	MATRIX 1 1.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

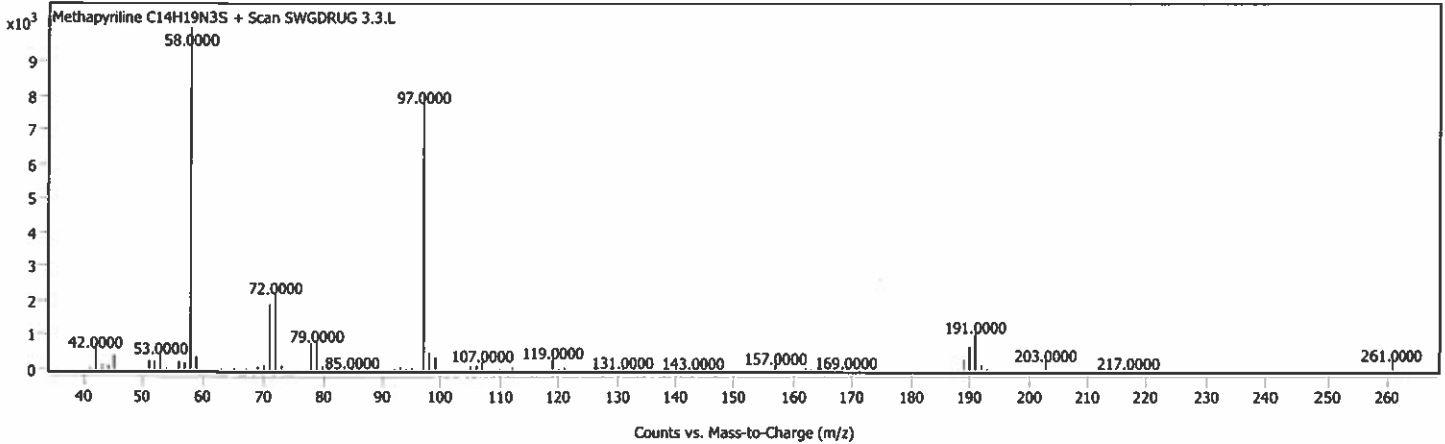
## Observed Spectrum



## Mirror Plot



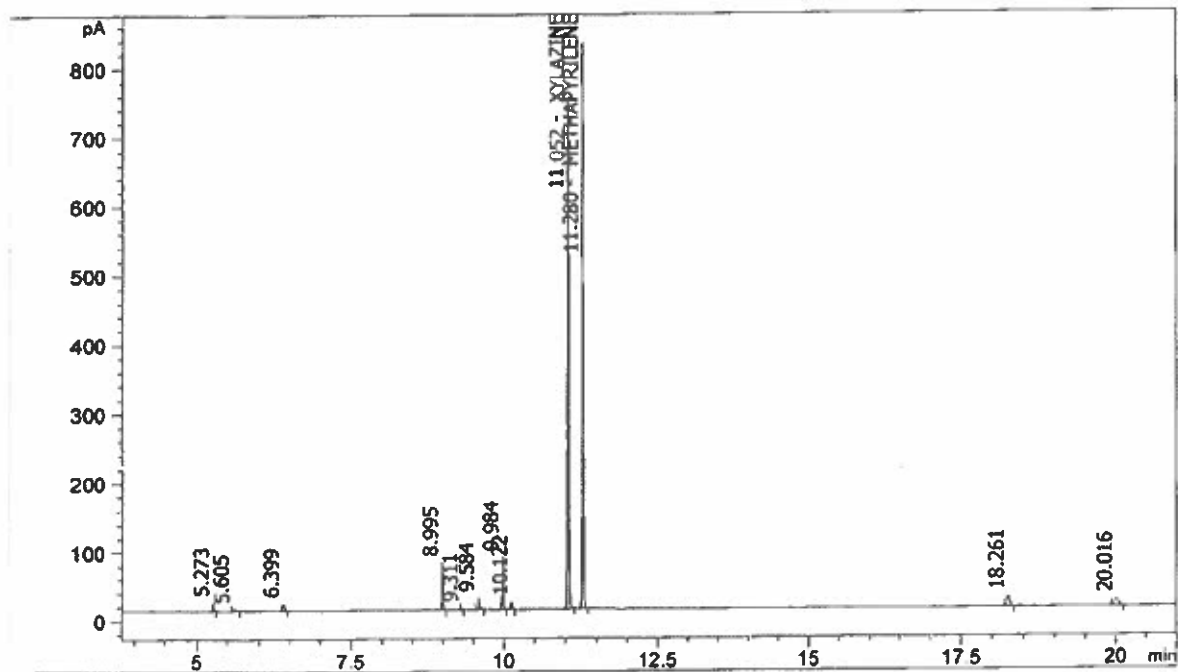
## Library Spectrum



Injection Date: 9/28/2021 5:24:23 PM  
 Sample Name: MATRIX 1 1.0 mg/L - BLD  
 Sample Info:

Seq Line: 10  
 Vial 10

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



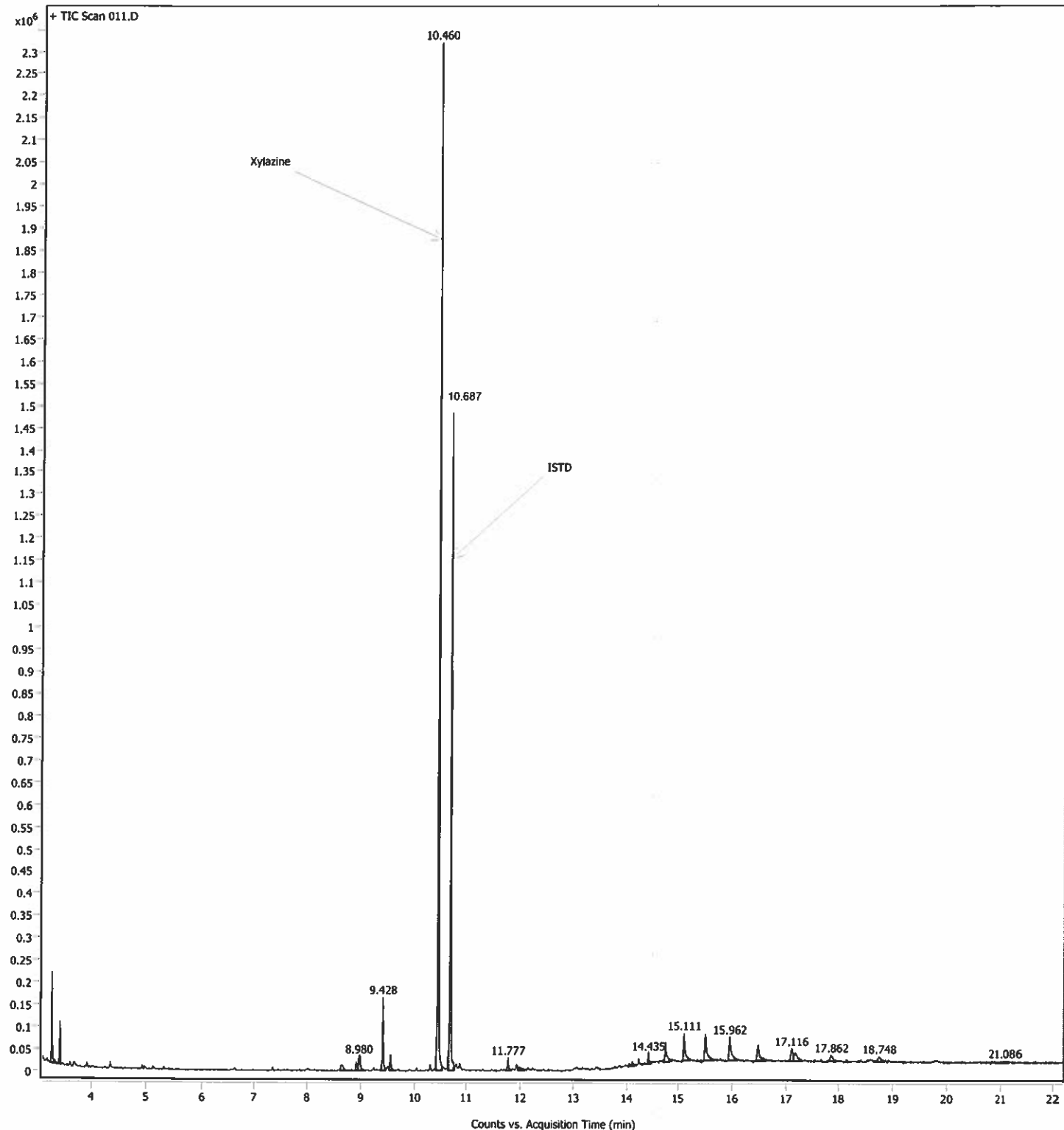
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.052	11.046	955.597	742.159	0.000000	XYLAZINE RRT - 0.9798
11.280	11.278	954.907	818.491	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 1.0 mg/L - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\011.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 5:59:01 PM (UTC-04:00)
Position	37	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

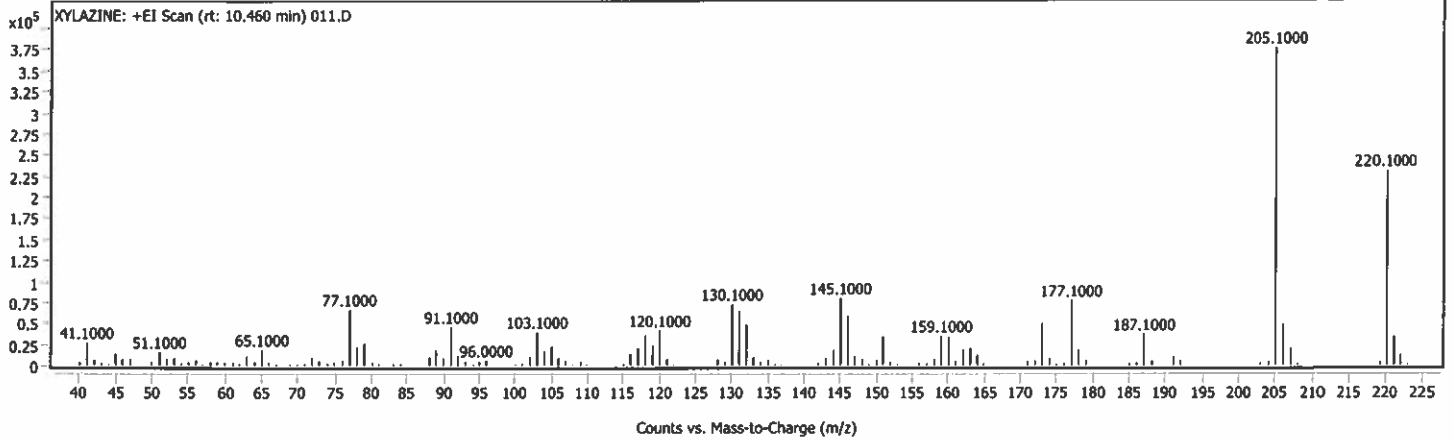
## Sample Spectra

+ Scan (rt: 10.460 min)

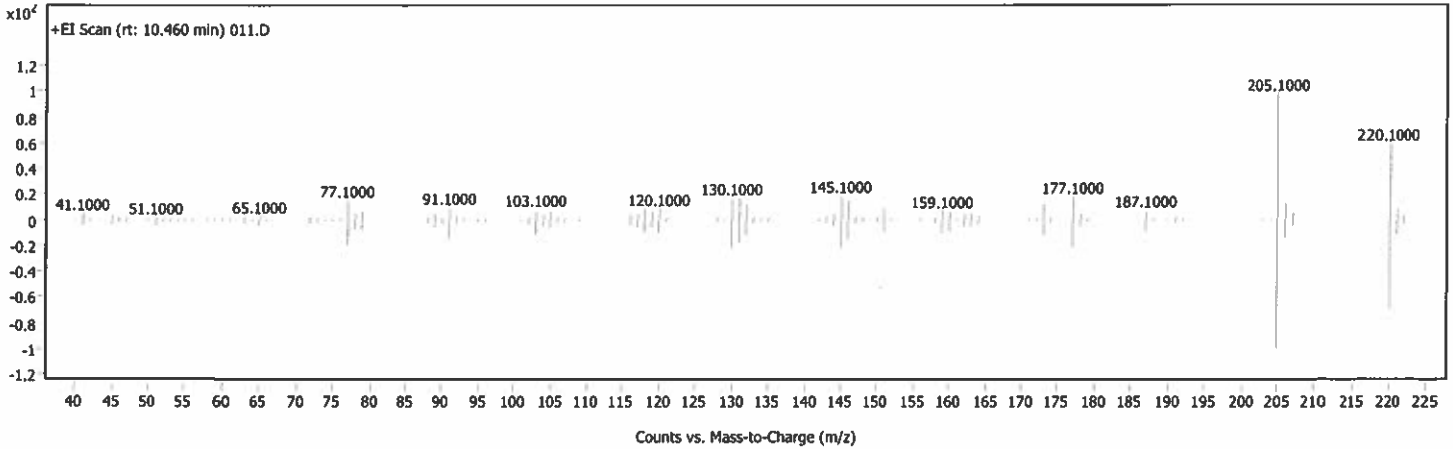
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.83	aafs.i	MATRIX 6 1.0 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

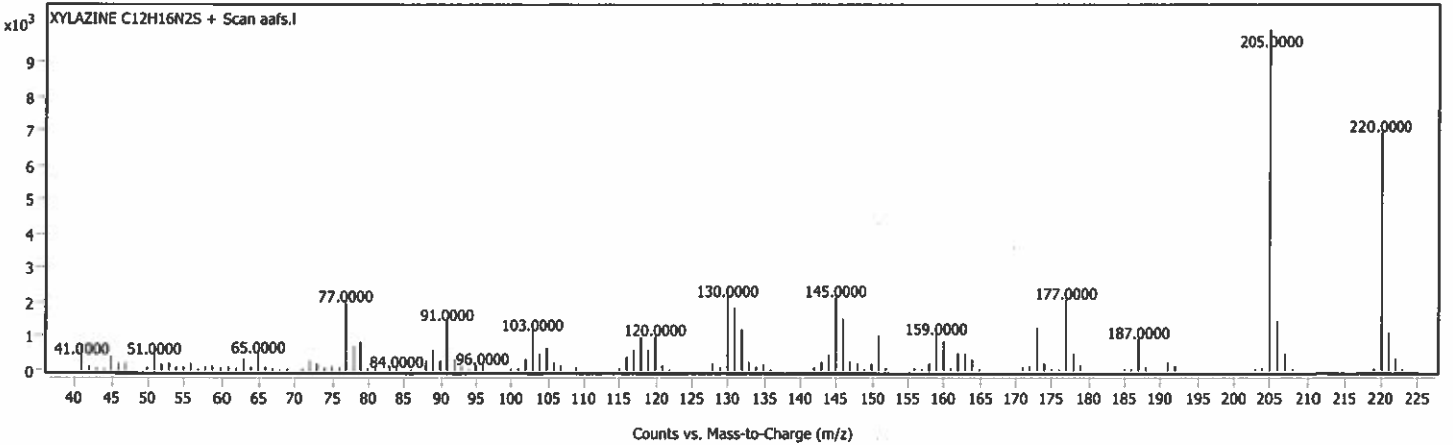
### Observed Spectrum



### Mirror Plot



### Library Spectrum



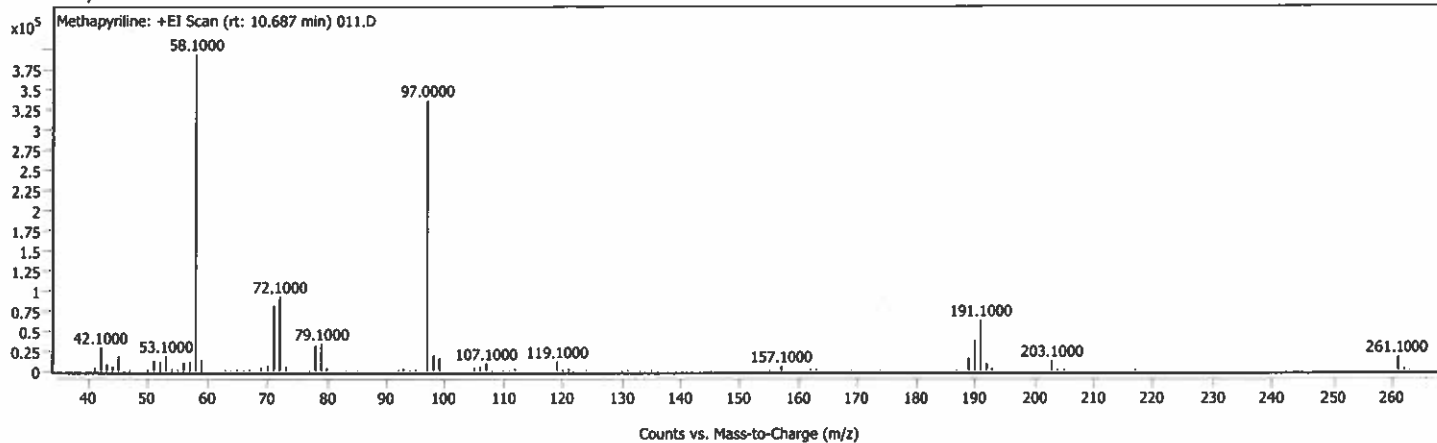
# Analysis Report

+ Scan (rt: 10.687 min)

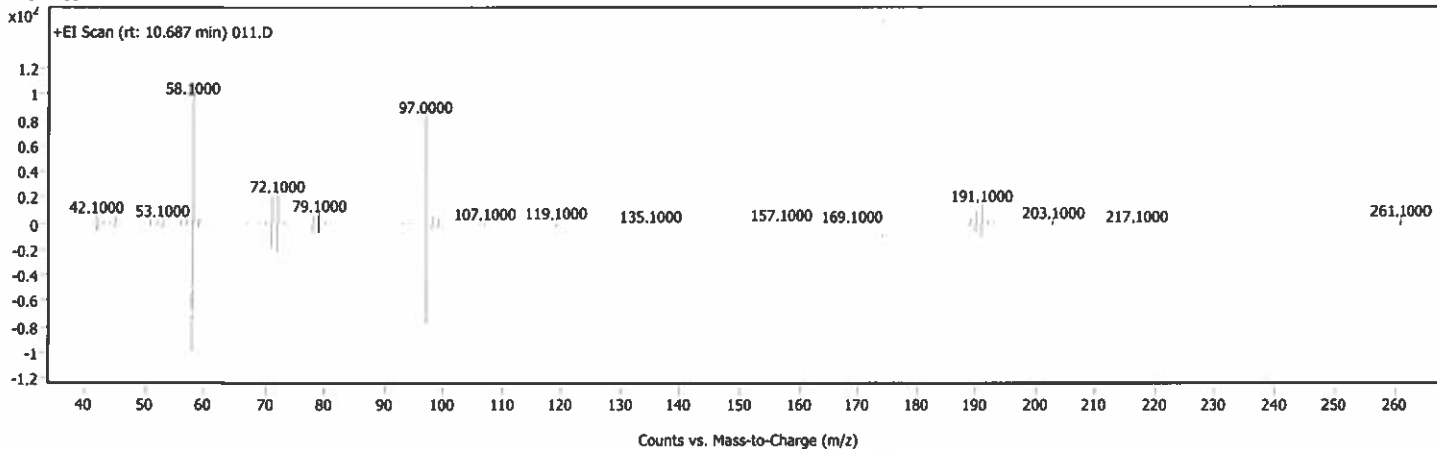
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.63	SWGDRUG 3.3.L	MATRIX 6 1.0 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

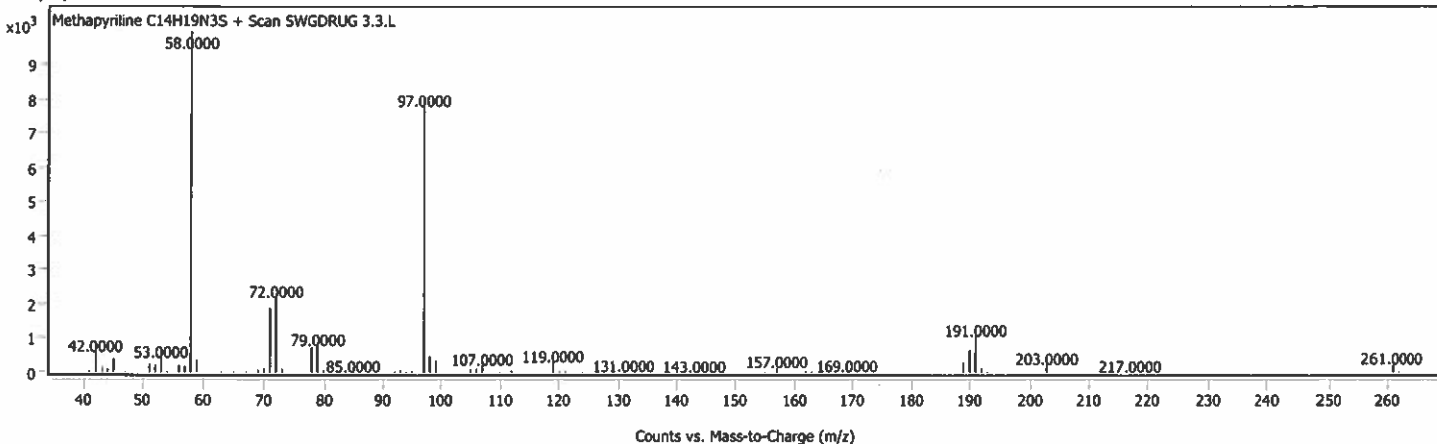
## Observed Spectrum



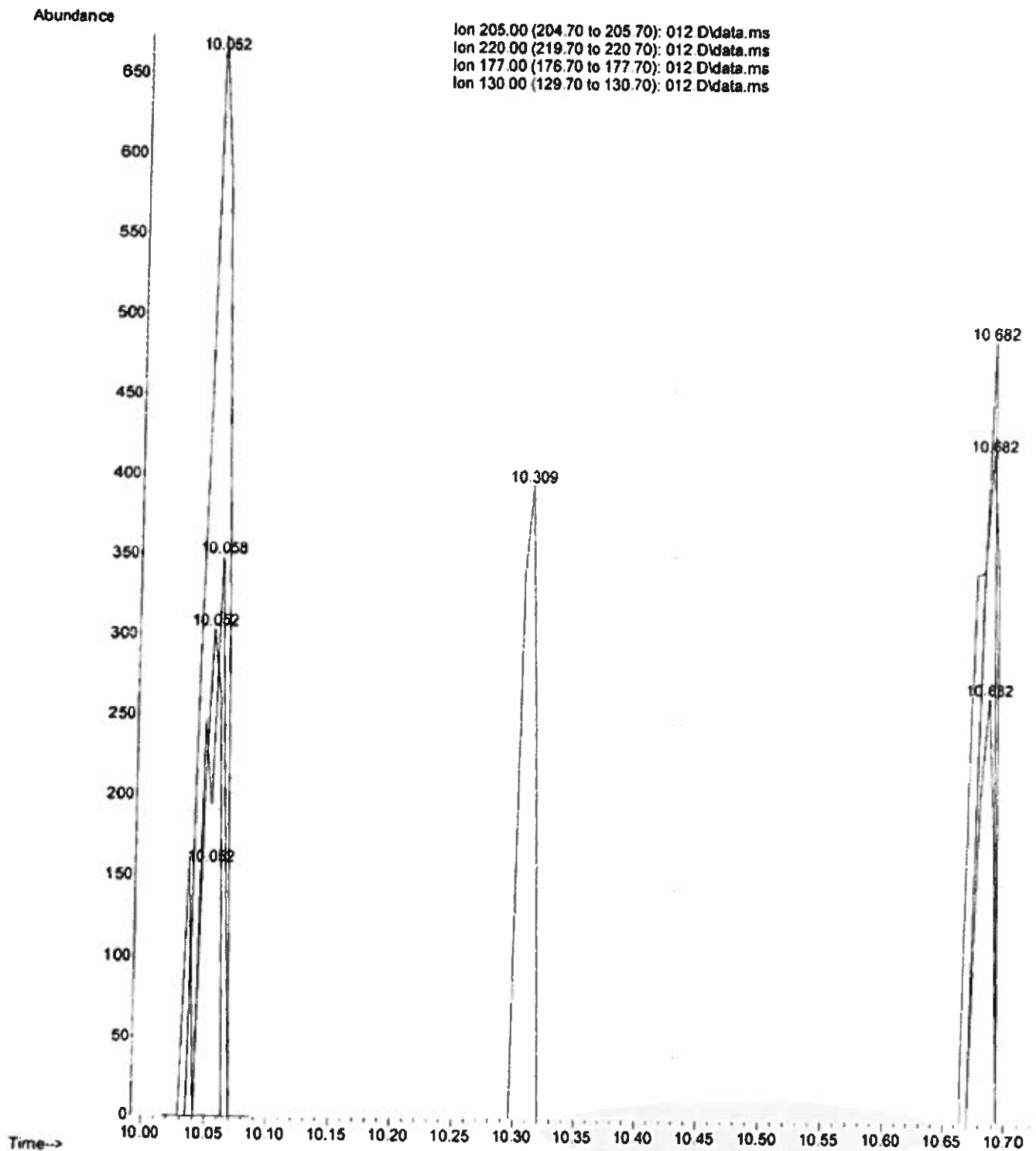
## Mirror Plot



## Library Spectrum



File :C:\Users\TOX\Desktop\092821\012.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 28 Sep 2021 18:24 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK after 1.0 mg/L  
Misc Info :  
Vial Number: 38

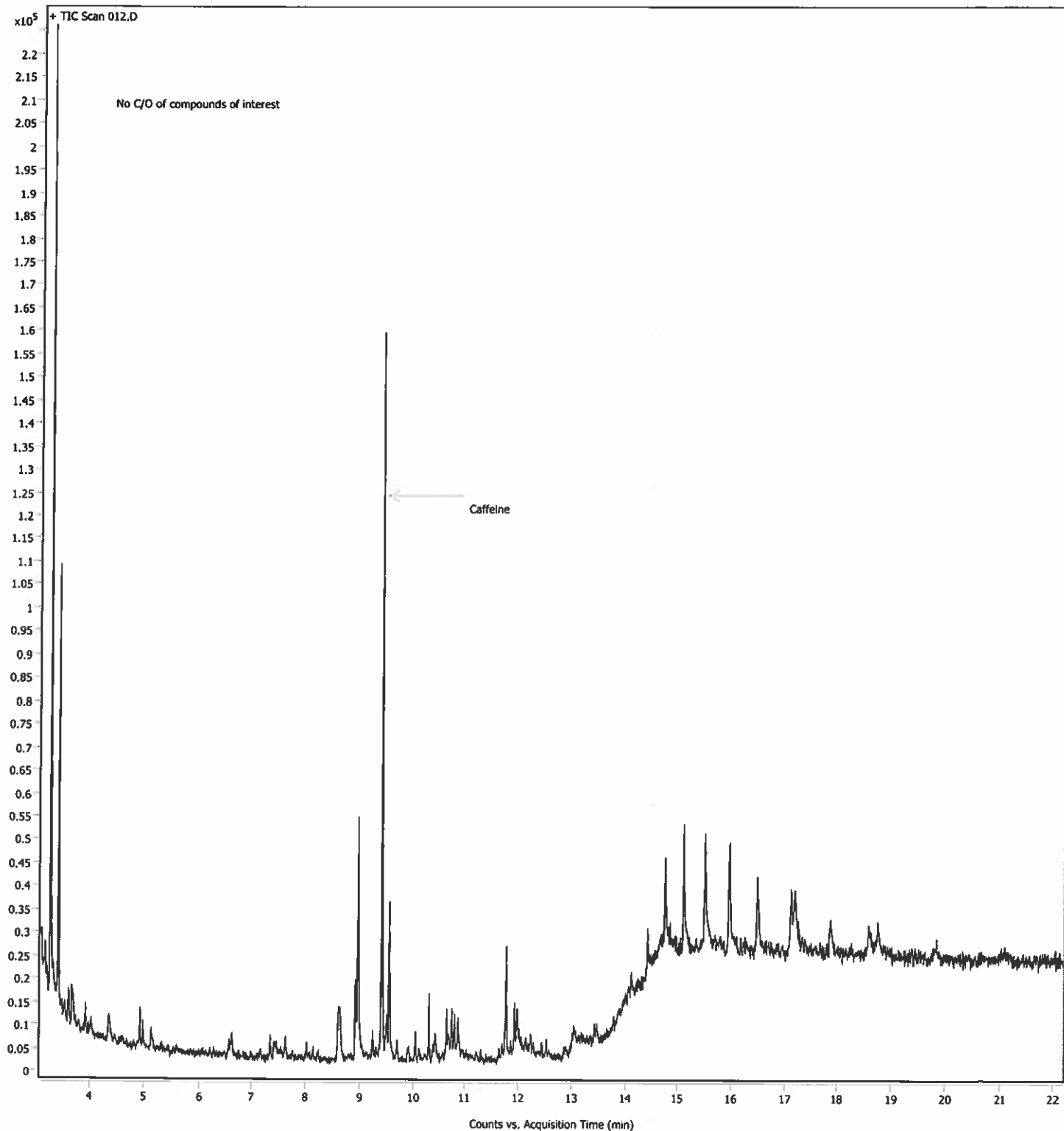


# Analysis Report

## Sample Information

Sample Name	MATRIX BLANK after 1.0 mg/L	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\012.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 6:24:54 PM (UTC-04:00)
Position	38	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

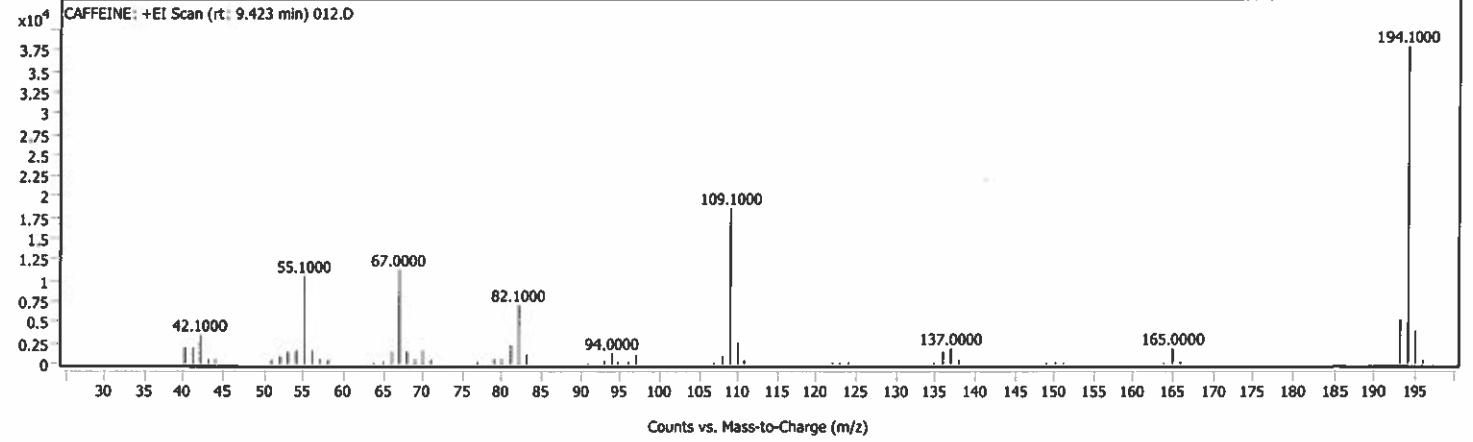
## Sample Spectra

+ Scan (rt: 9.423 min)

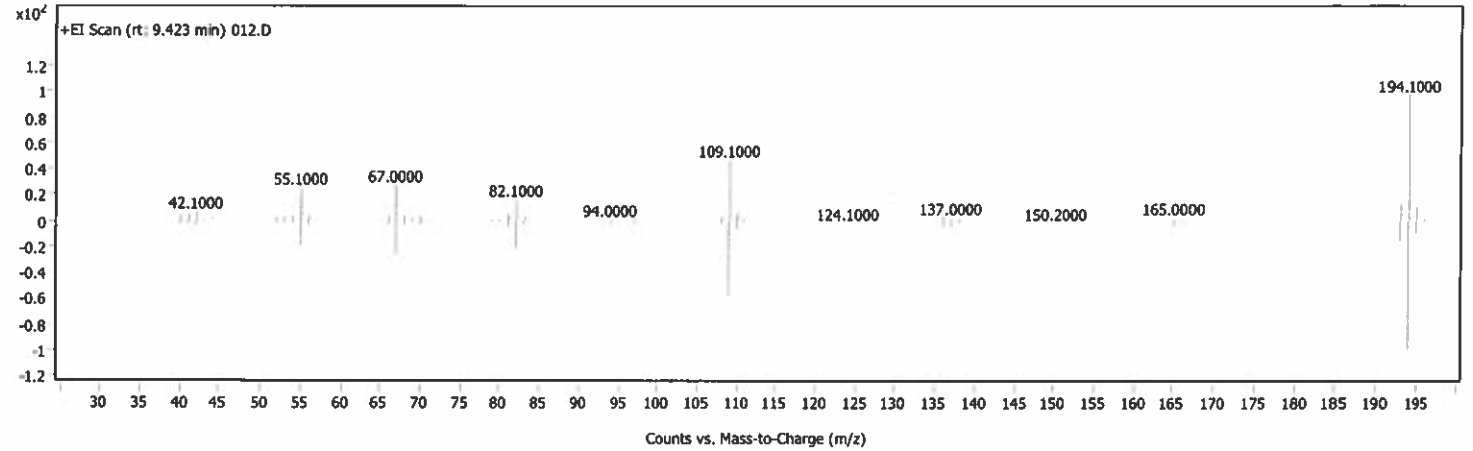
CAFFEINE; C8H10N4O2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CAFFEINE	C8H10N4O2	98.67	aafs.l	MATRIX BLANK after 1.0 mg/L	KM/QC DATA IN XYLAZINE VALIDATION

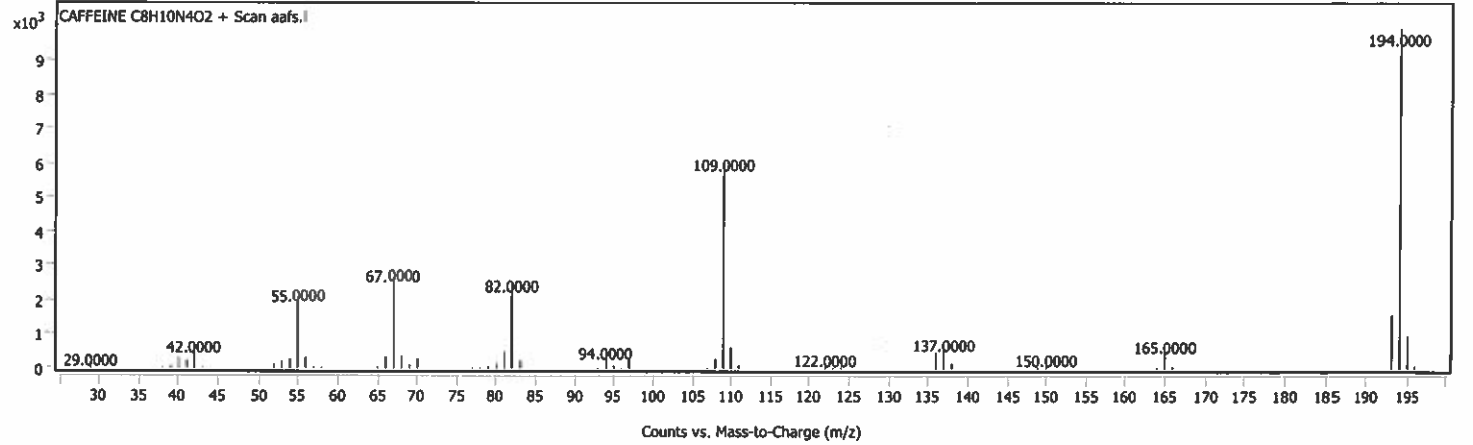
## Observed Spectrum



## Mirror Plot

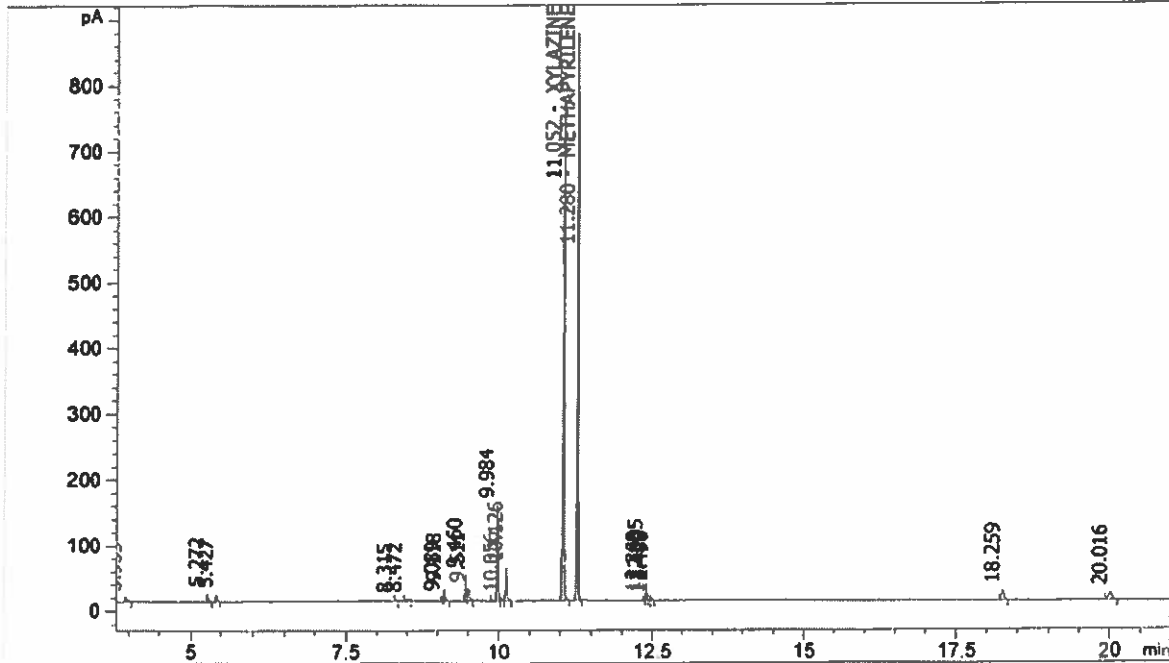


## Library Spectrum



Injection Date: 9/28/2021 6:12:56 PM Seq Line: 12  
 Sample Name: MATRIX 6 1.0 mg/L - URN Vial 12  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION

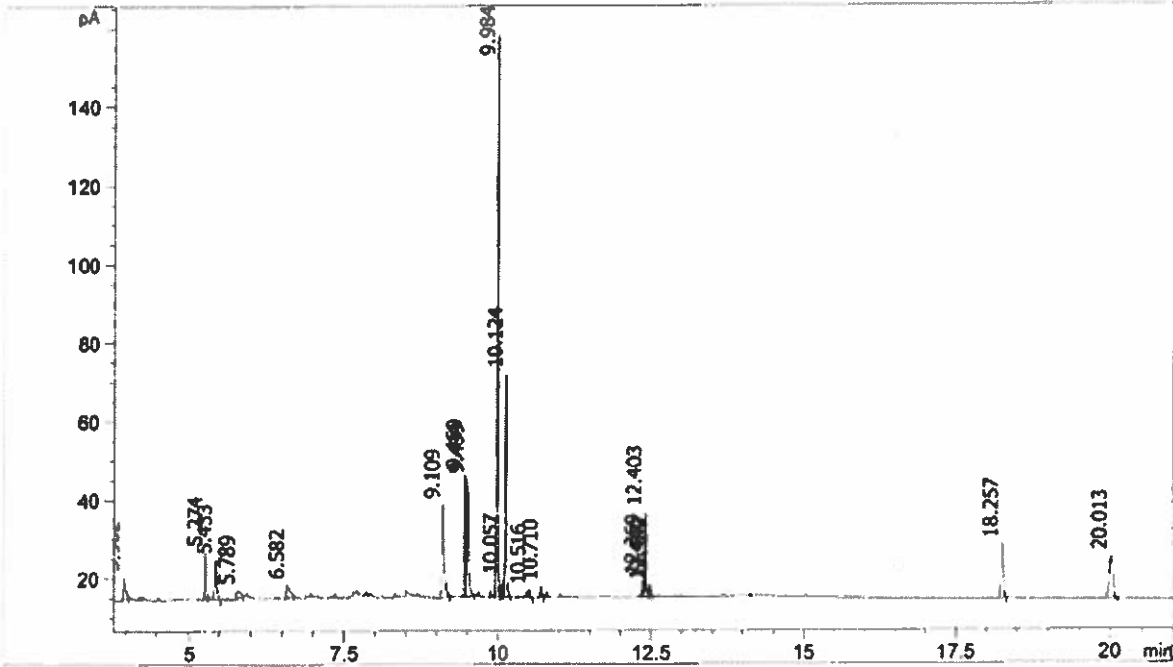


RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.052	11.046	994.796	755.586	0.000000	XYLAZINE RRT-0.9798
11.280	11.278	1031.261	862.071	1.000000	METHAPYRILENE

Injection Date: 9/28/2021 6:37:18 PM  
 Sample Name: MATRIX BLANK after 1.->  
 Sample Info:

Seq Line: 13  
 Vial 13

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



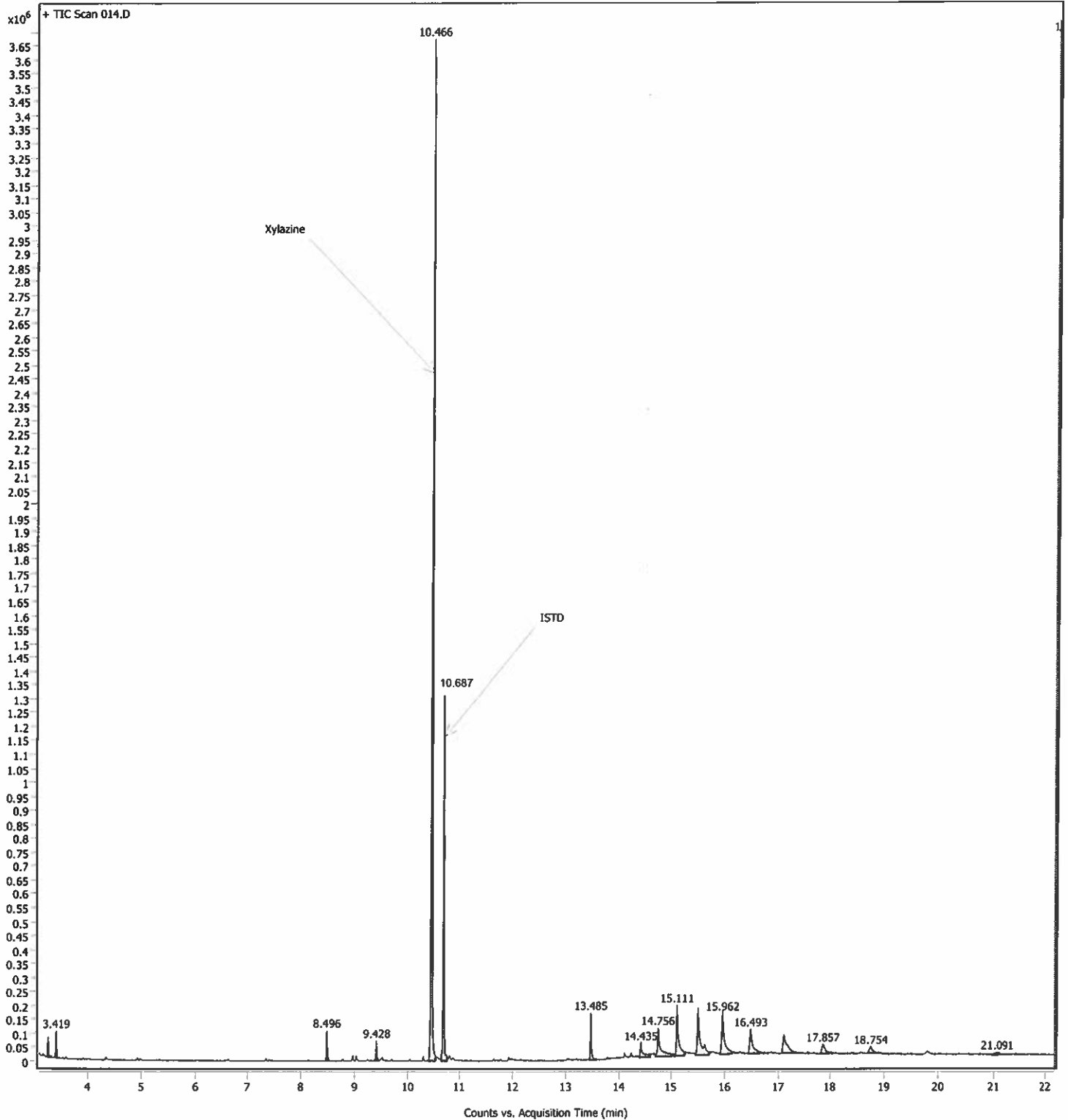
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 2 2.0 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\014.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 7:16:26 PM (UTC-04:00)
Position	40	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

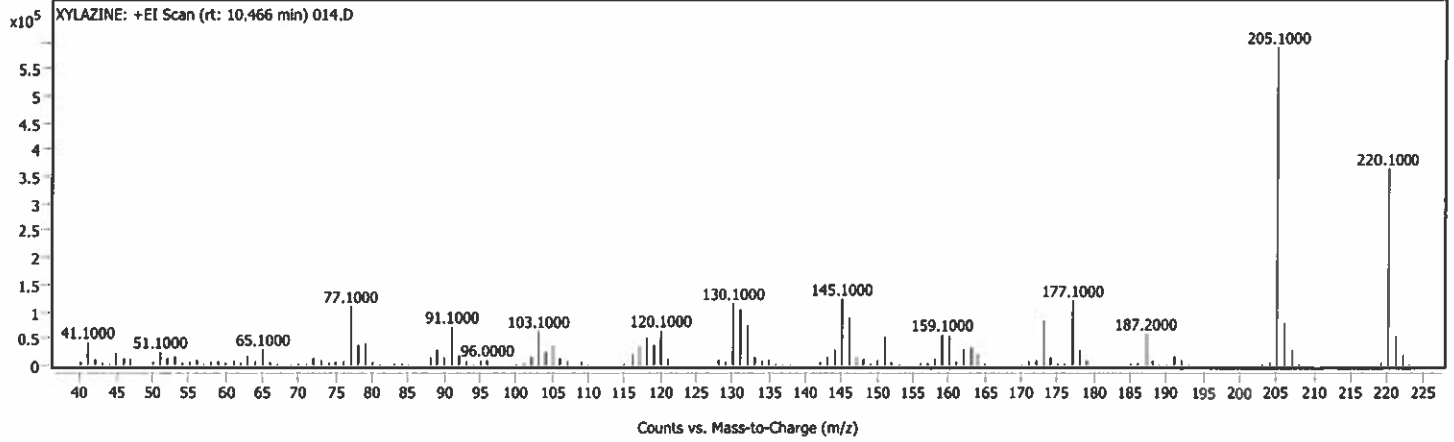
## Sample Spectra

+ Scan (rt: 10.466 min)

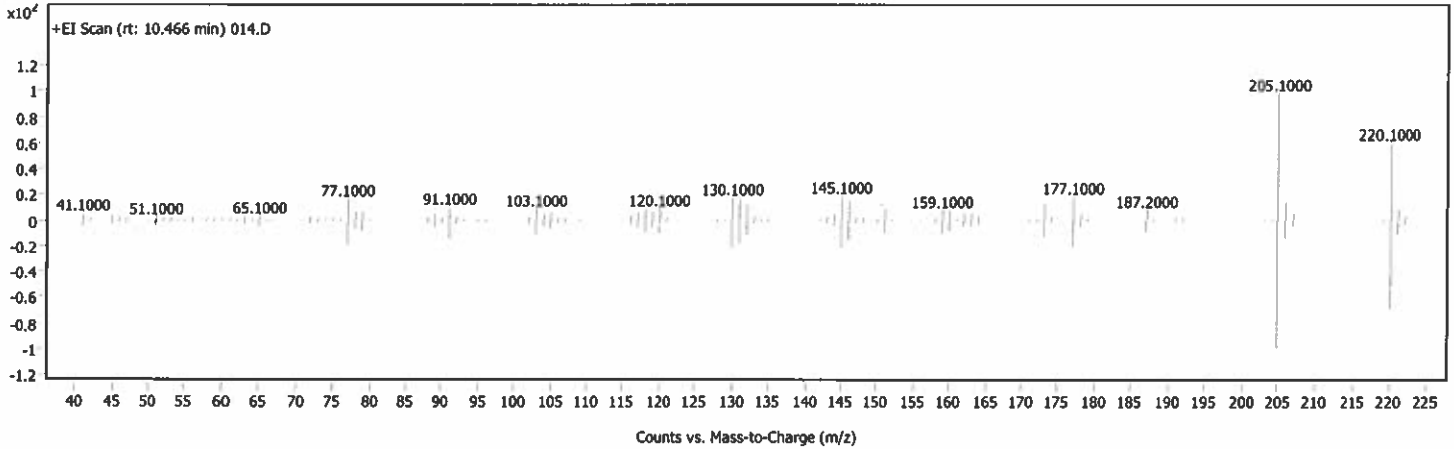
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.74	aafs.I	MATRIX 2 2.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

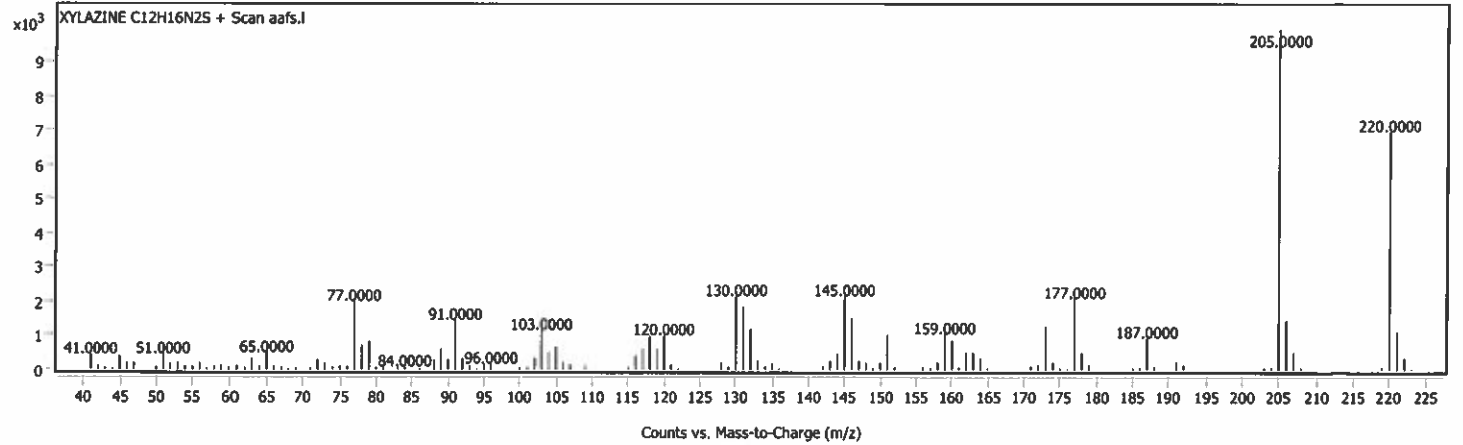
## Observed Spectrum



## Mirror Plot



## Library Spectrum



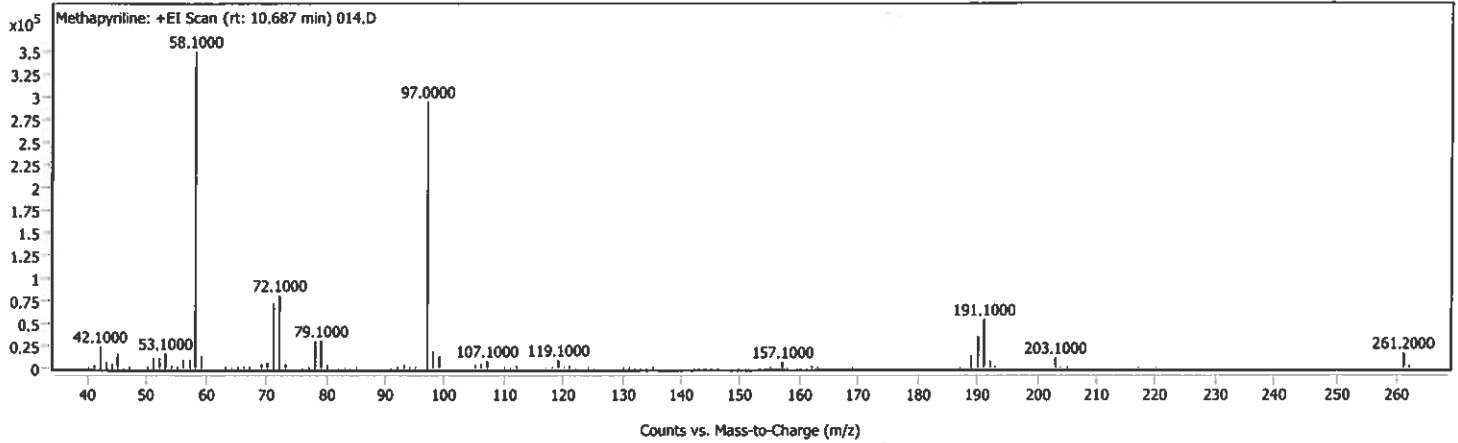
# Analysis Report

+ Scan (rt: 10.687 min)

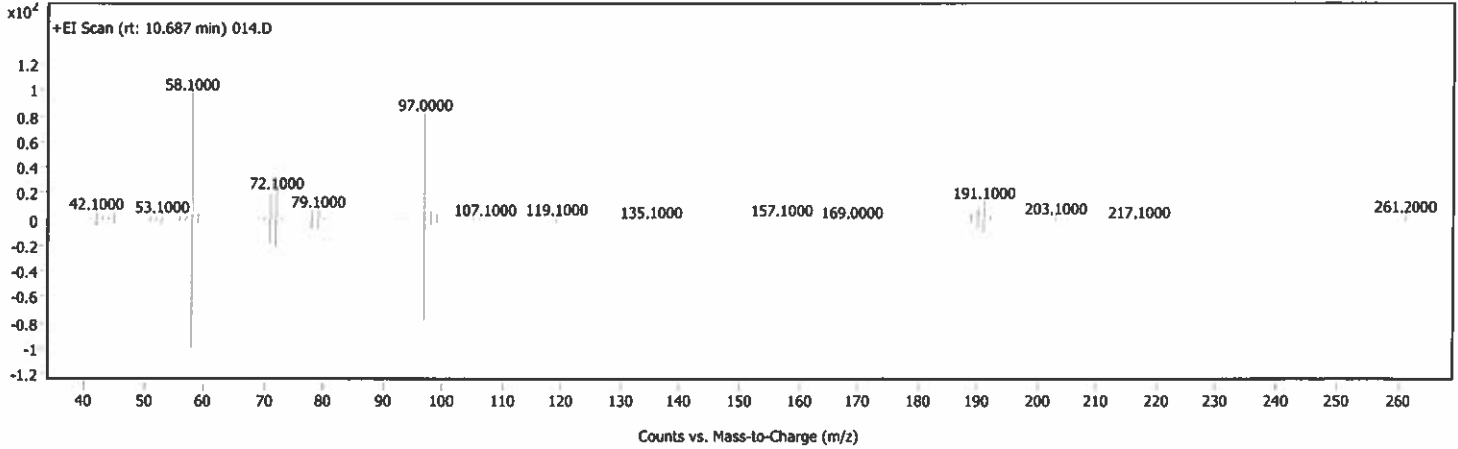
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.56	SWGDRUG 3.3.L	MATRIX 2 2.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

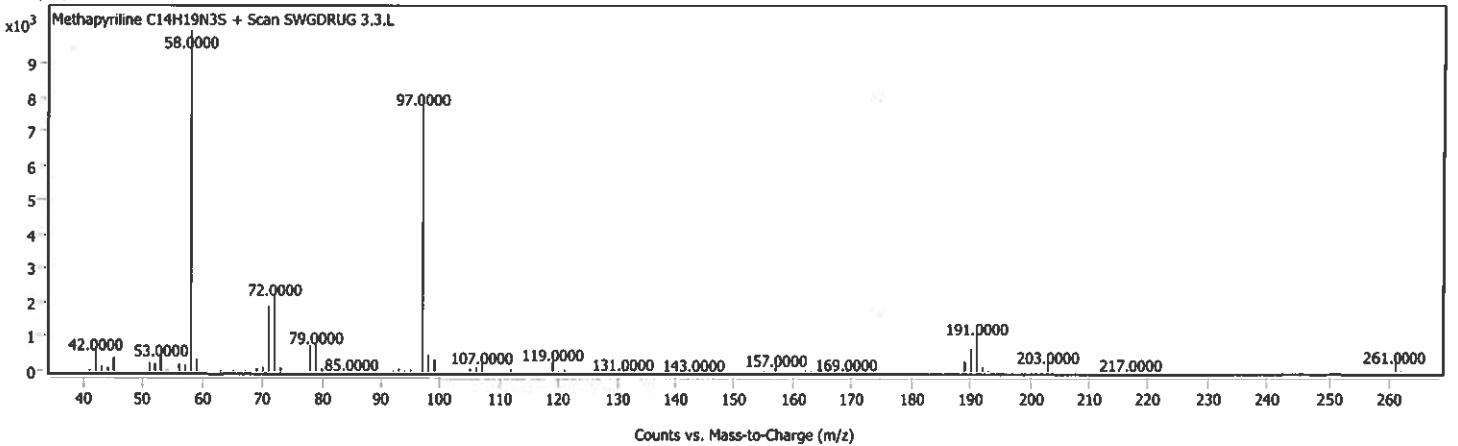
## Observed Spectrum



## Mirror Plot

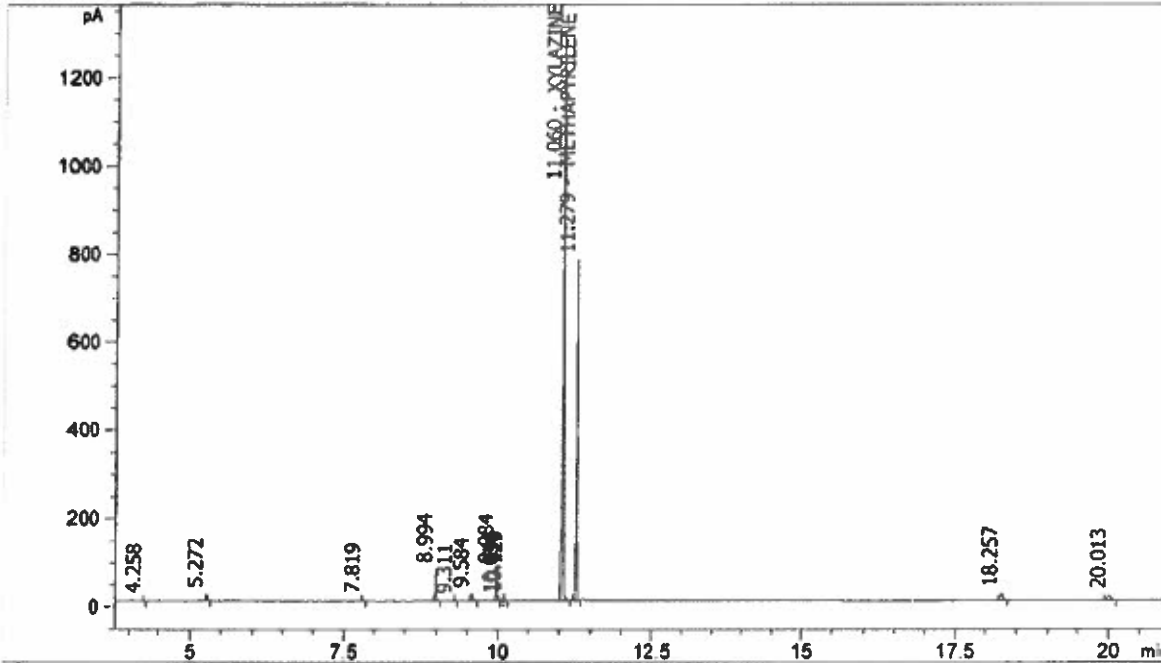


## Library Spectrum



Injection Date: 9/28/2021 7:25:52 PM Seq Line: 15  
 Sample Name: MATRIX 2 2.0 mg/L - BLD Vial 15  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



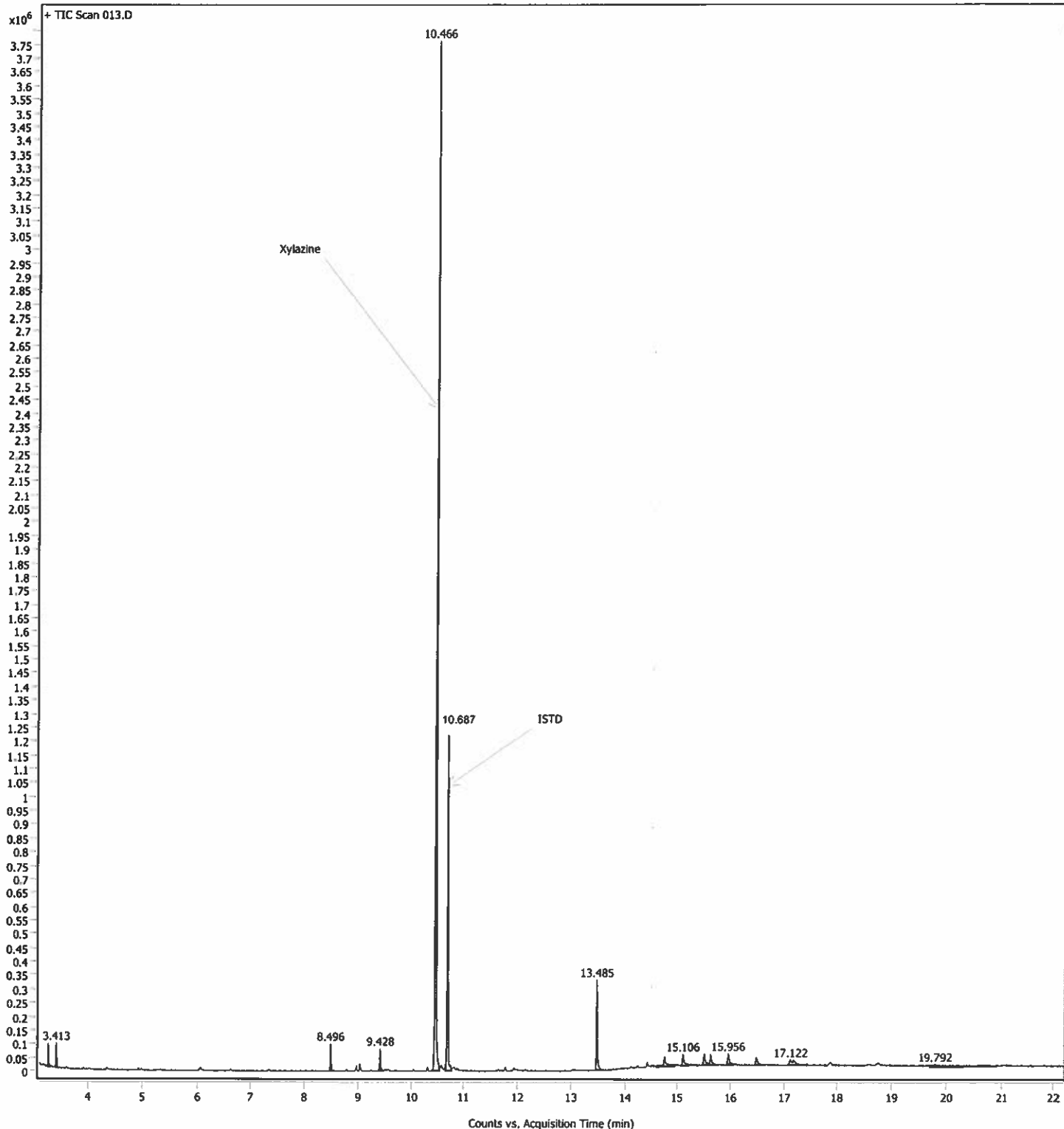
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.060	11.046	1891.334	1272.978	0.000000	XYLAZINE RRT - 0.9806
11.279	11.278	920.167	773.041	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 1 2.0 mg/L - BLD	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\013.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 6:50:41 PM (UTC-04:00)
Position	39	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms





# Analysis Report

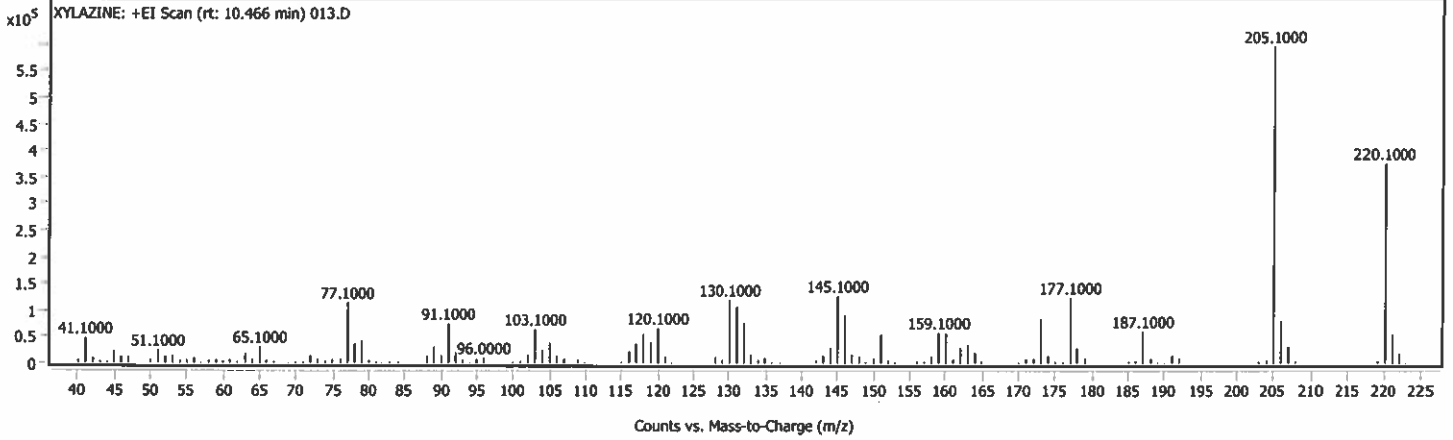
## Sample Spectra

+ Scan (rt: 10.466 min)

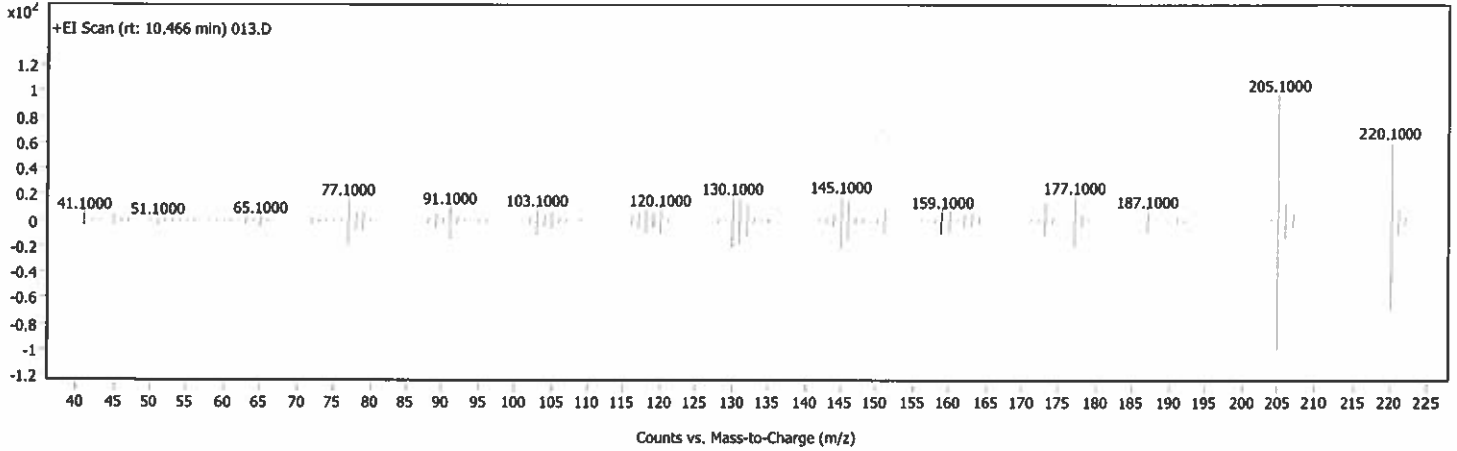
XYLAZINE; C12H16N2S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.76	aafs.I	MATRIX 1 2.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

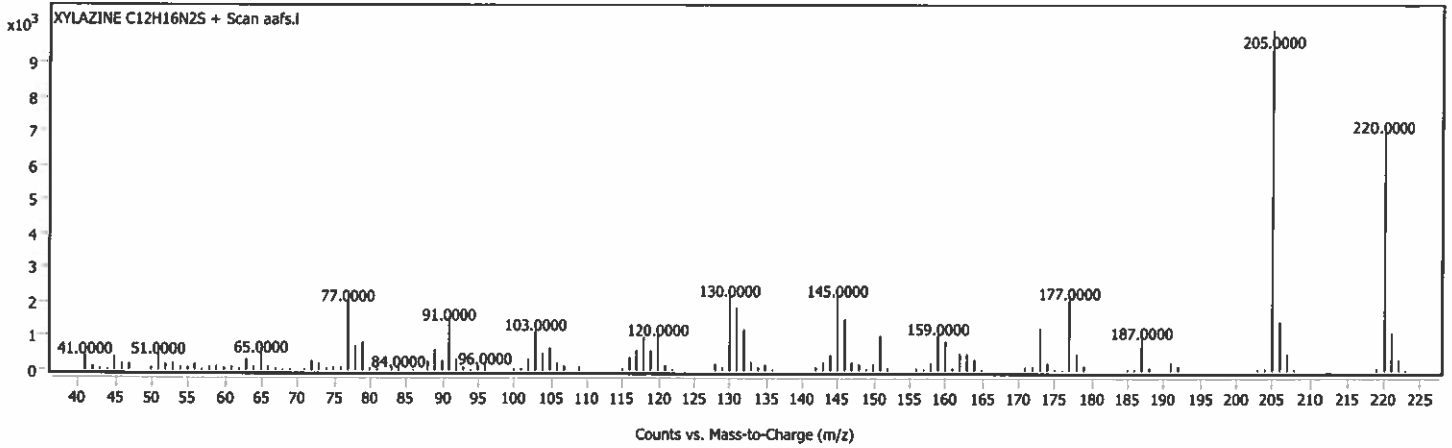
## Observed Spectrum



## Mirror Plot



## Library Spectrum



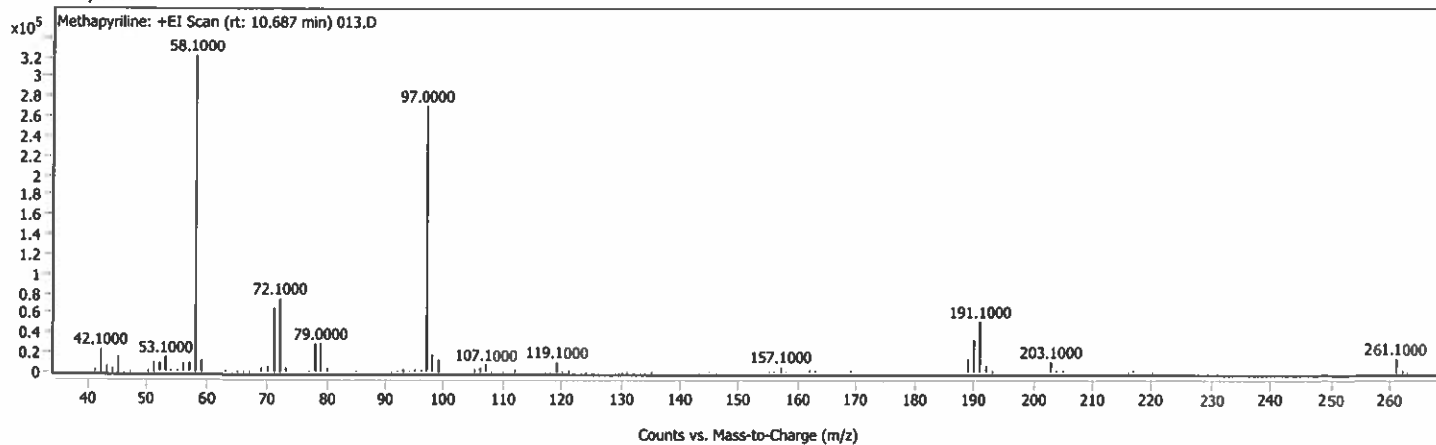
# Analysis Report

+ Scan (rt: 10.687 min)

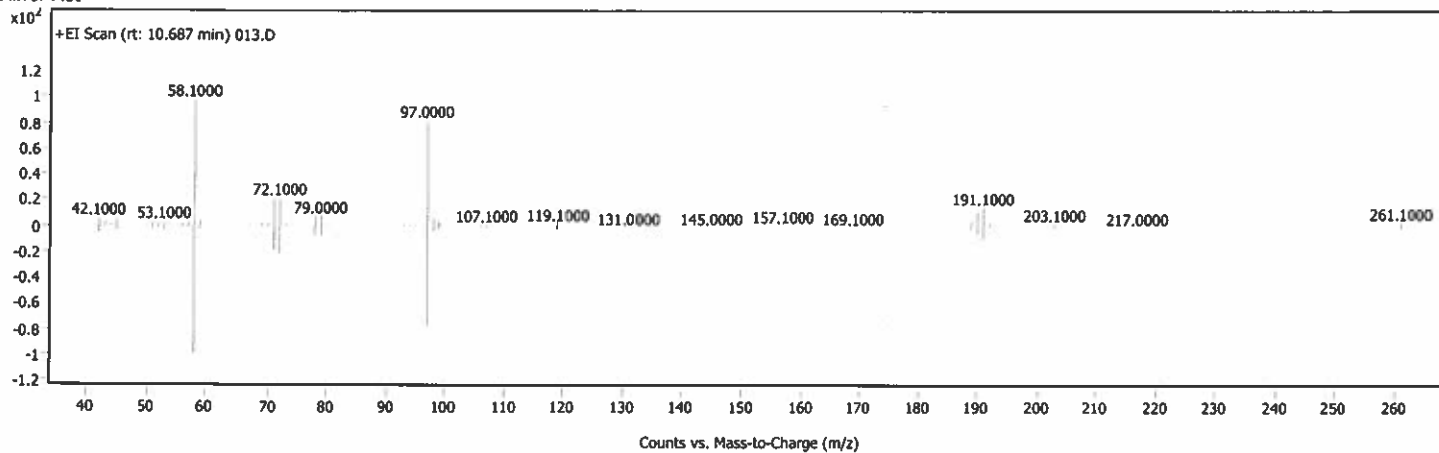
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.22	SWGDRUG 3.3.L	MATRIX 1 2.0 mg/L - BLD	KM/QC DATA IN XYLAZINE VALIDATION

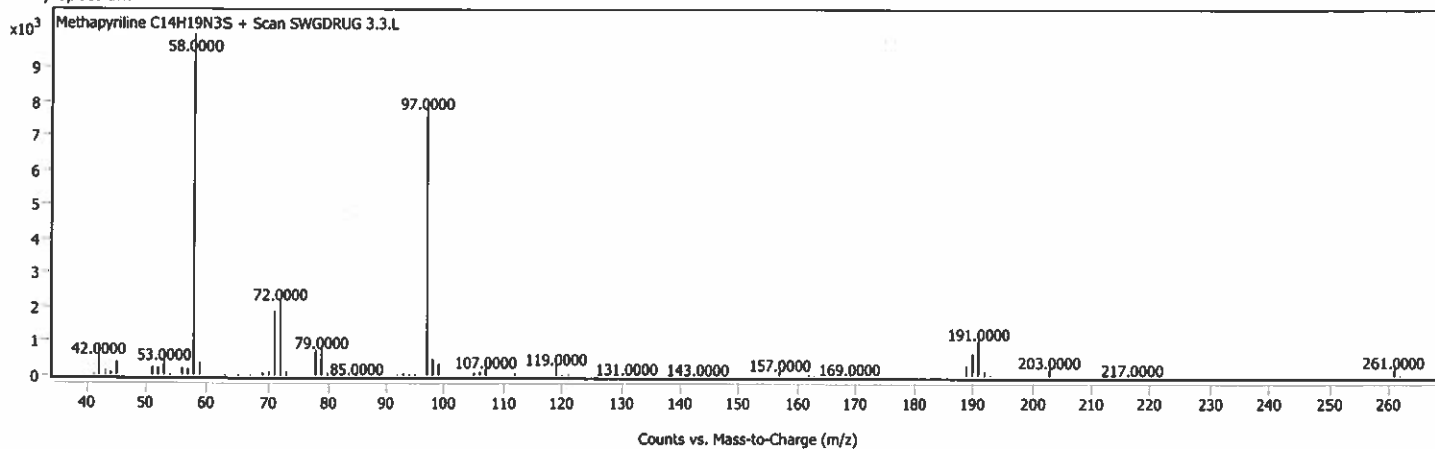
## Observed Spectrum



## Mirror Plot



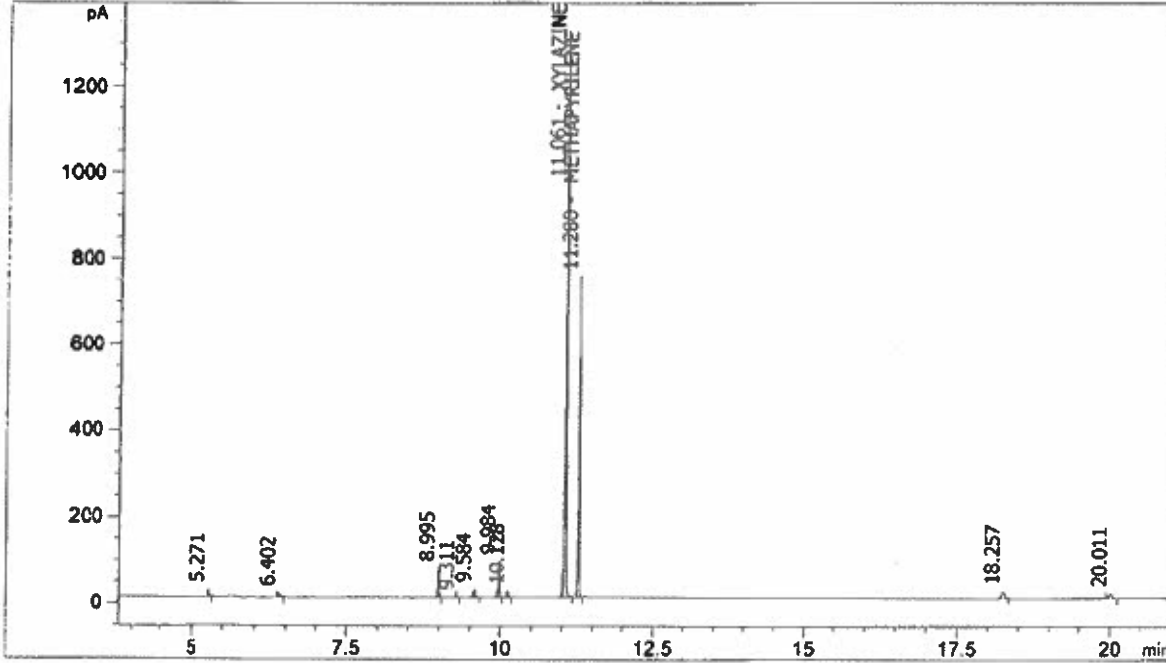
## Library Spectrum



=====

Injection Date: 9/28/2021 7:01:35 PM Seq Line: 14  
 Sample Name: MATRIX 1 2.0 mg/L - BLD Vial 14  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



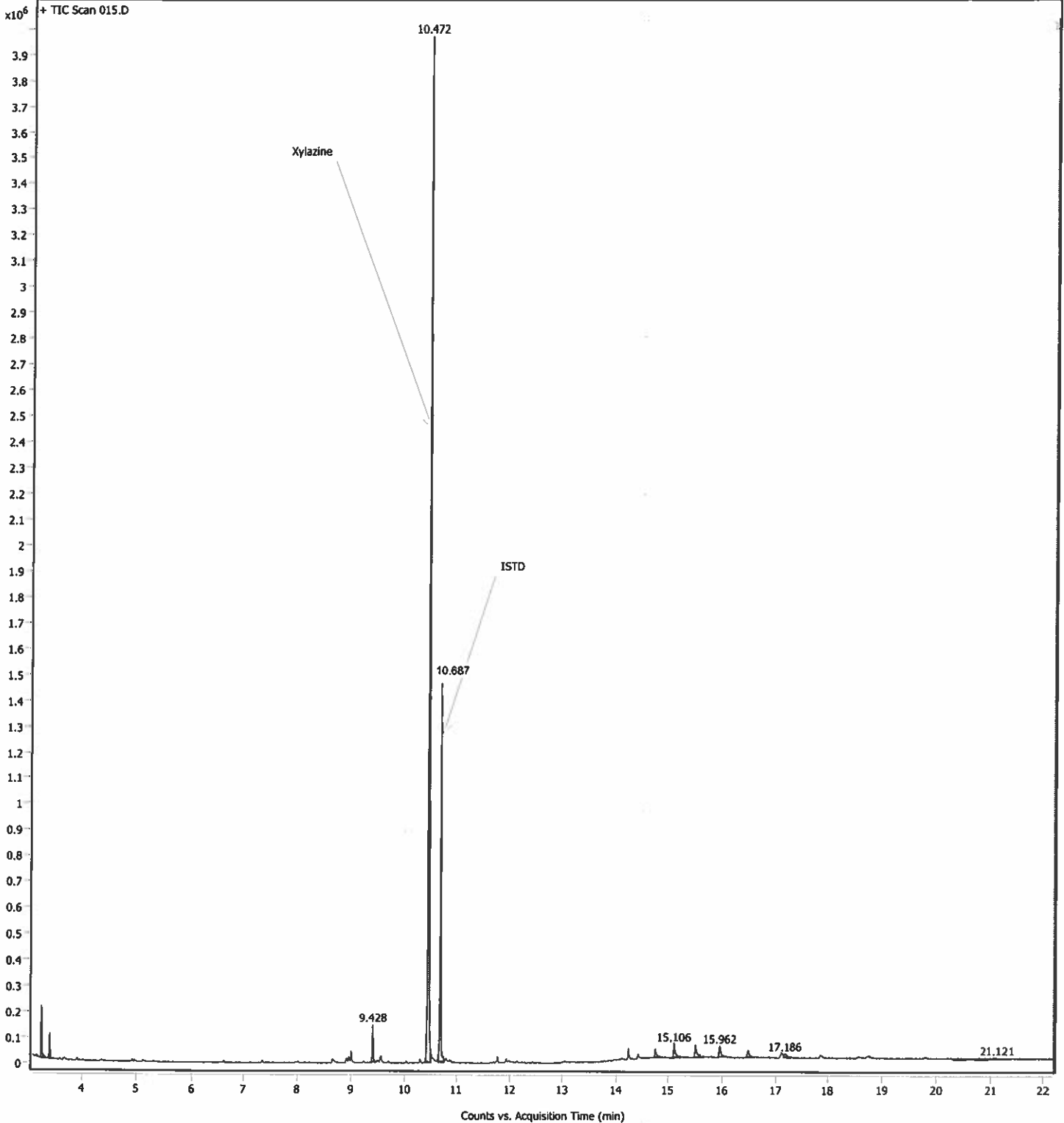
RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.061	11.046	1961.487	1308.153	0.000000	XYLAZINE RRT - 0.9806
11.280	11.278	879.732	743.979	1.000000	METHAPYRILENE

# Analysis Report

## Sample Information

Sample Name	MATRIX 6 2.0 mg/L - URN	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\015.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 7:42:20 PM (UTC-04:00)
Position	41	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/OC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



# Analysis Report

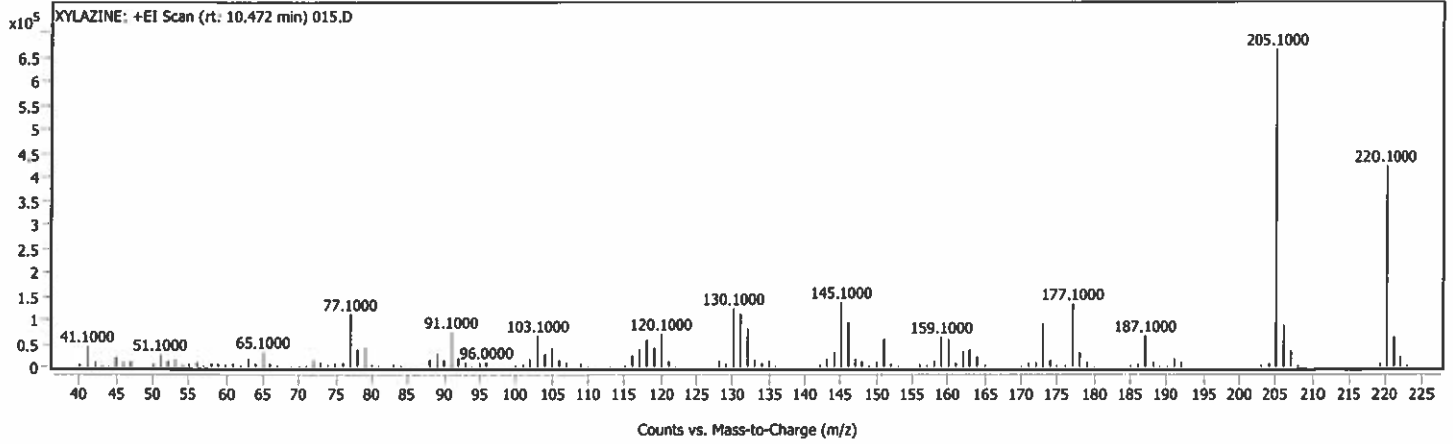
## Sample Spectra

+ Scan (rt: 10.472 min)

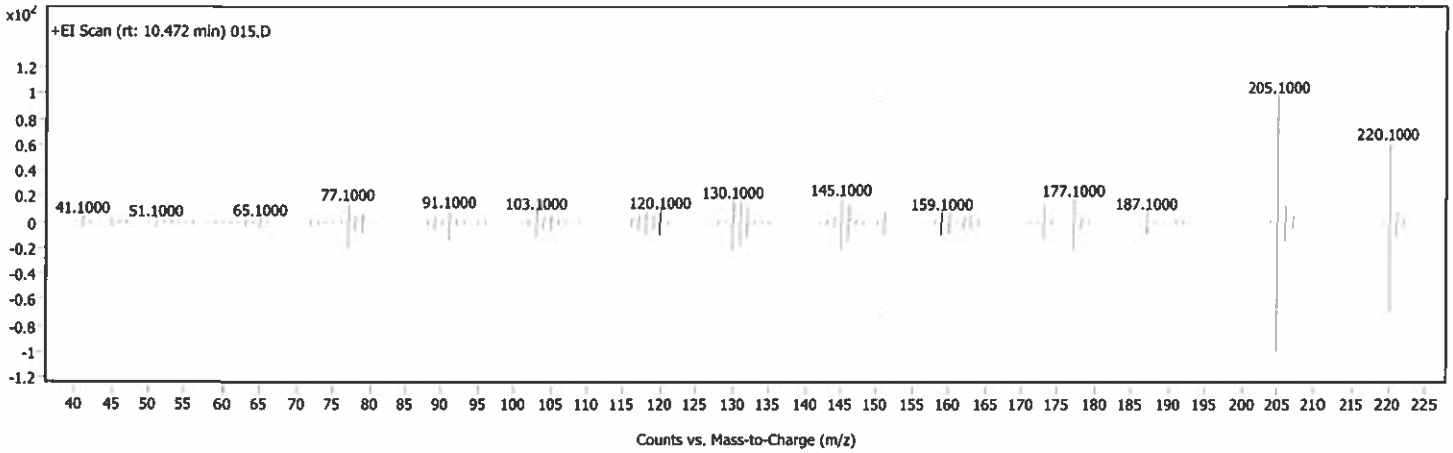
**XYLAZINE; C12H16N2S**

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
XYLAZINE	C12H16N2S	97.79	aafs.l	MATRIX 6 2.0 mg/L - URN	KM/QC DATA IN XYLAZINE VALIDATION

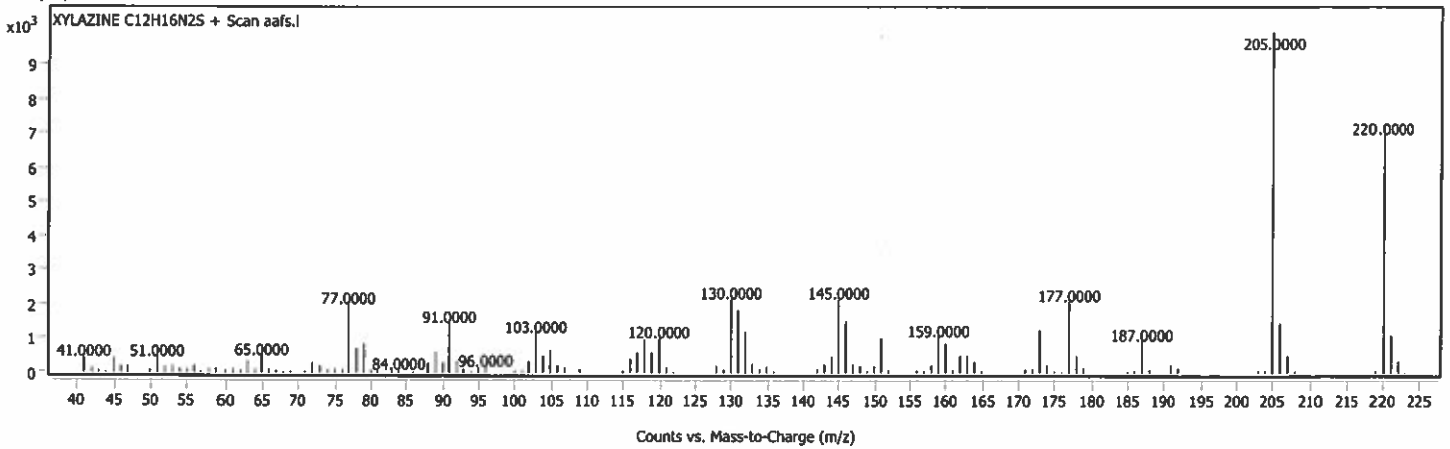
### Observed Spectrum



### Mirror Plot



### Library Spectrum



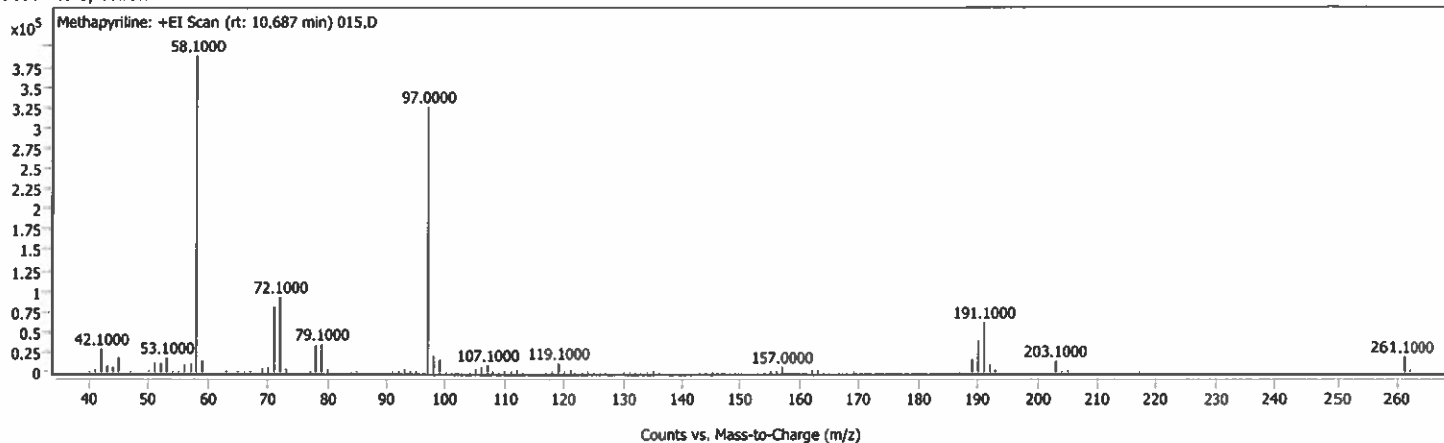
# Analysis Report

+ Scan (rt: 10.687 min)

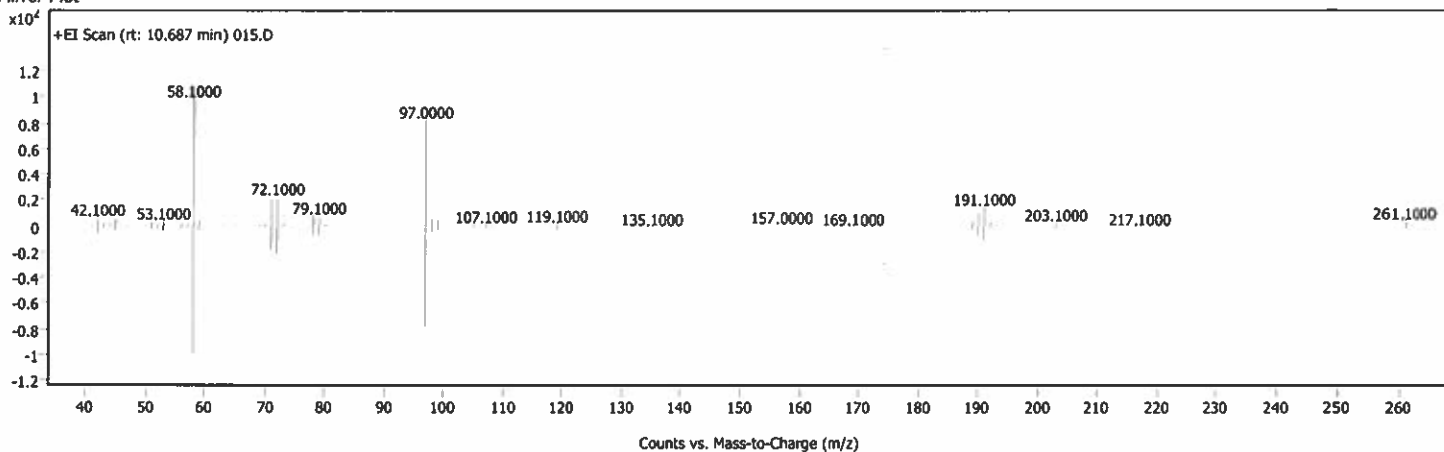
Methapyriline; C14H19N3S

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
Methapyriline	C14H19N3S	98.53	SWGDRUG 3.3.L	MATRIX 6 2.0 mg/L - URN	KM/OC DATA IN XYLAZINE VALIDATION

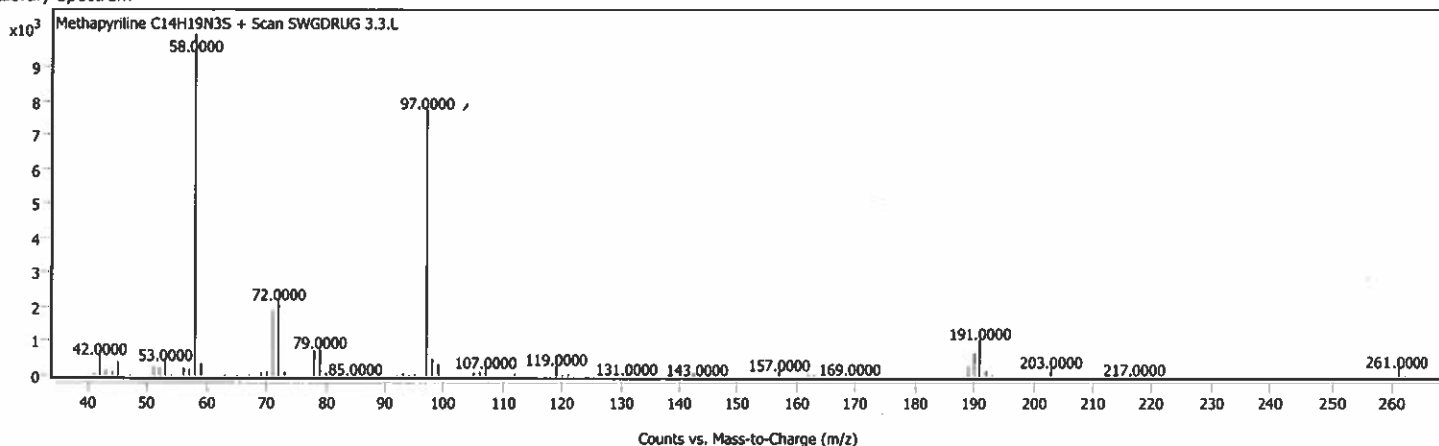
## Observed Spectrum



## Mirror Plot



## Library Spectrum

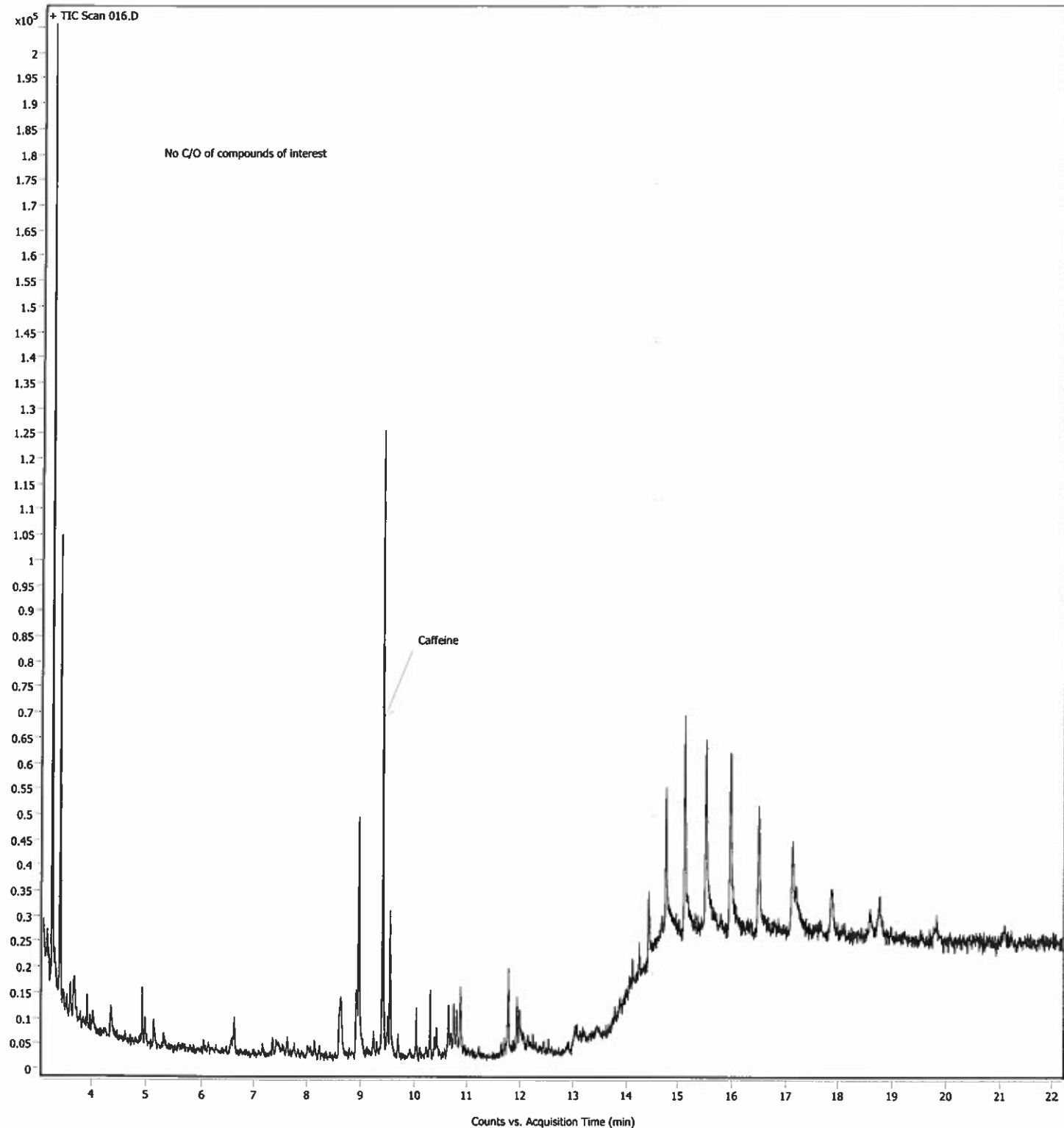


# Analysis Report

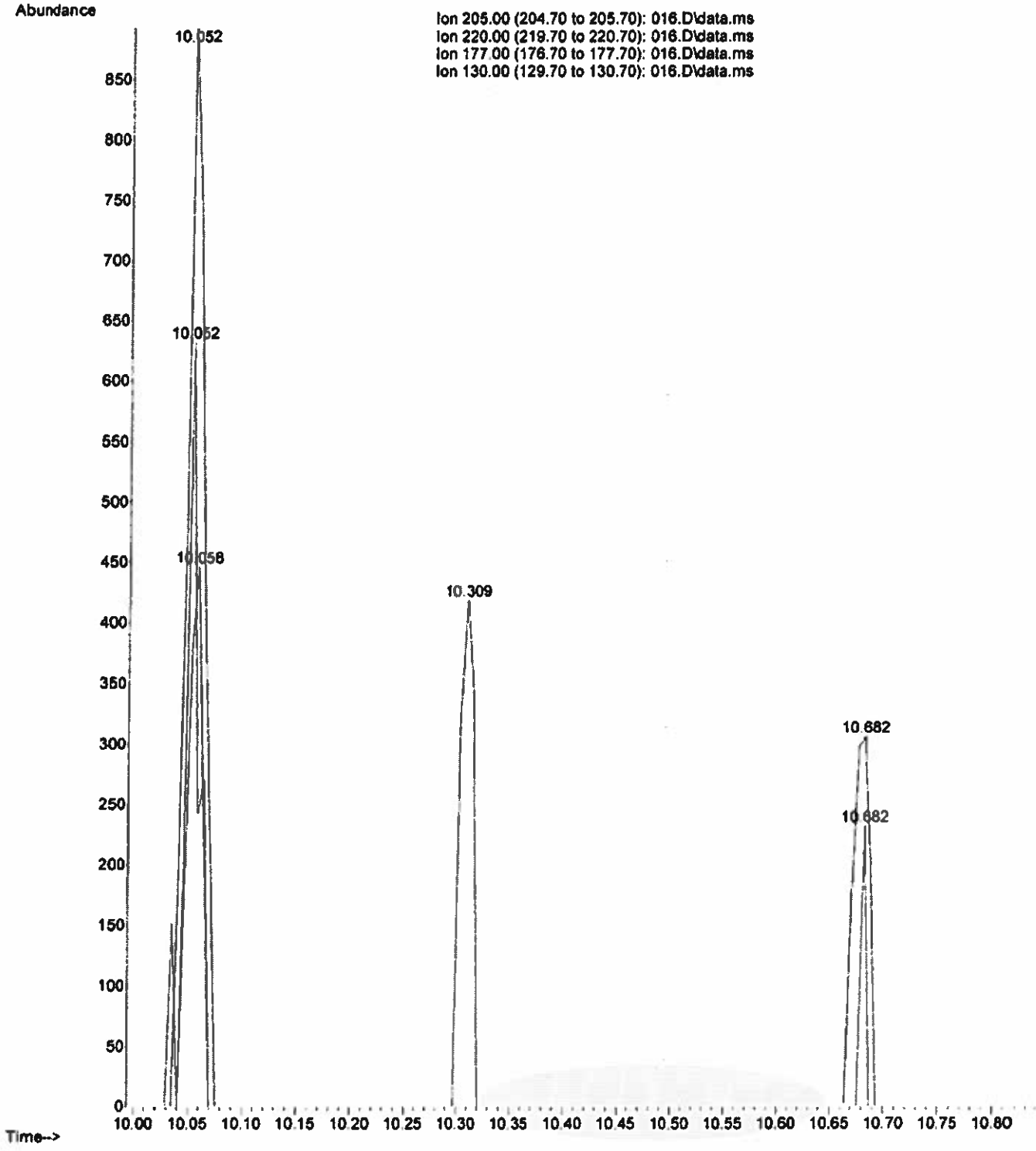
## Sample Information

Sample Name	MATRIX BLANK after 2.0 mg/L	Data File Path	C:\MassHunter\GCMS\1\data\BASES\092821\016.D
Instrument	#3 - Enhanced	Acq. Time (Local)	9/28/2021 8:08:07 PM (UTC-04:00)
Position	42	Method Path (Acq)	C:\MassHunter\GCMS\1\methods\ALKALI.M
Operator	KM/QC DATA IN XYLAZINE VALIDATION		

## Sample Chromatograms



File :C:\Users\TOX\Desktop\092821\016.D  
Operator : KM/QC DATA IN XYLAZINE VALIDATION  
Acquired : 28 Sep 2021 20:08 using AcqMethod ALKALI.M  
Instrument : #3 - Enhanced  
Sample Name: MATRIX BLANK after 2.0 mg/L  
Misc Info :  
Vial Number: 42





# Analysis Report

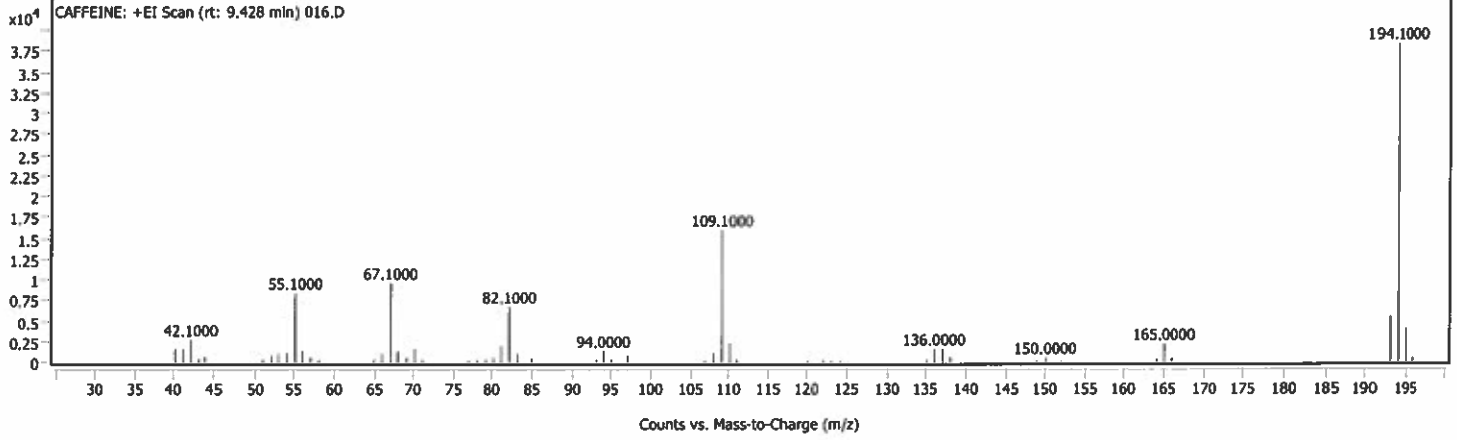
## Sample Spectra

+ Scan (rt: 9.428 min)

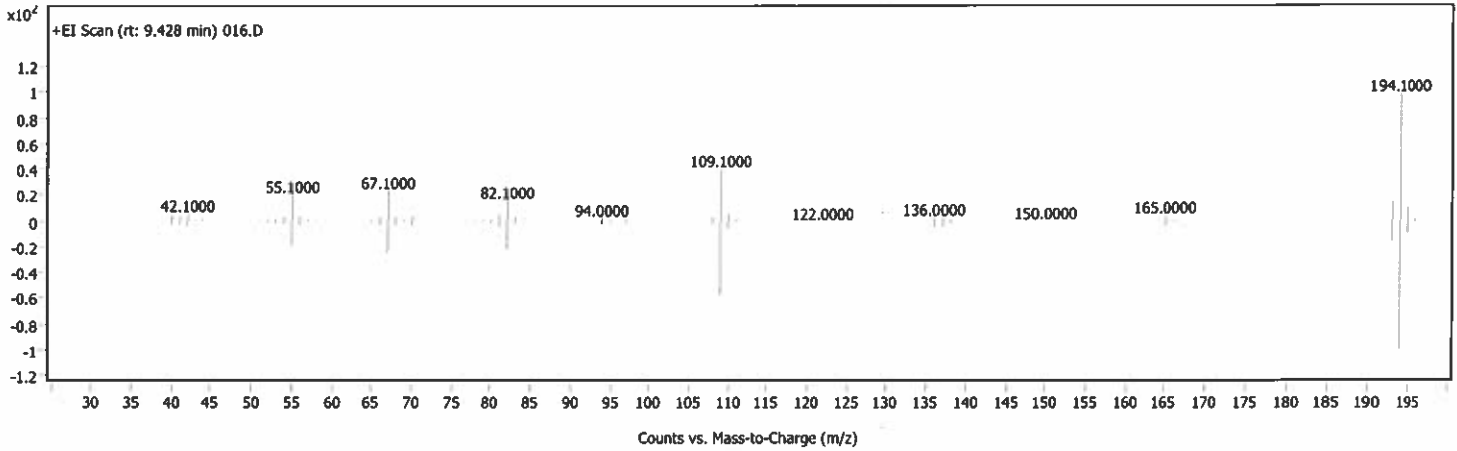
CAFFEINE; C8H10N4O2

Name	Formula	Score (Lib)	Lib/DB	Sample Name	Operator
CAFFEINE	C8H10N4O2	98.67	aafs.l	MATRIX BLANK after 2.0 mg/L	KM/QC DATA IN XYLAZINE VALIDATION

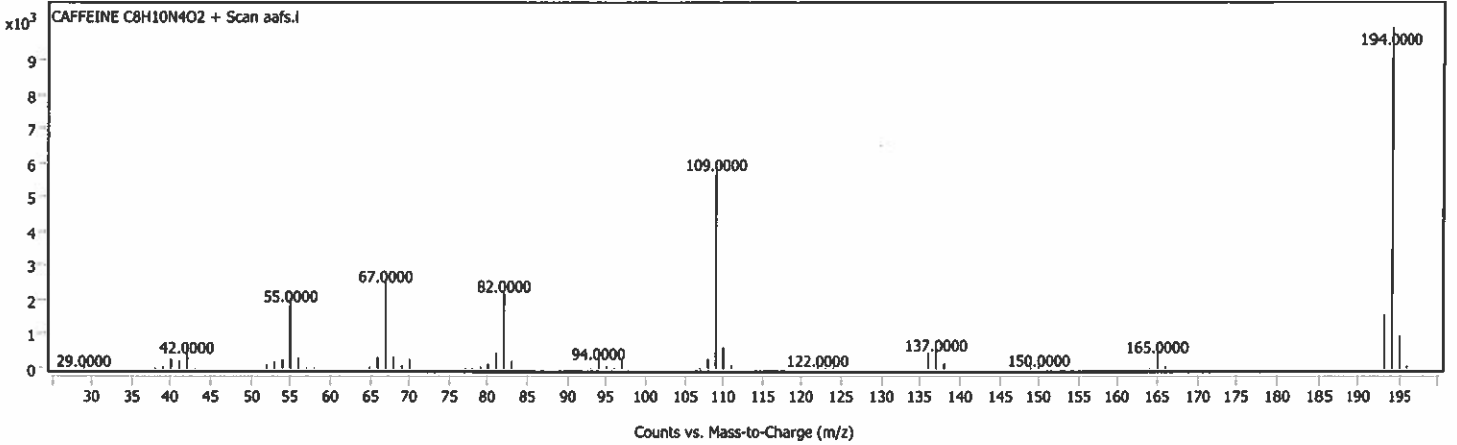
### Observed Spectrum



### Mirror Plot

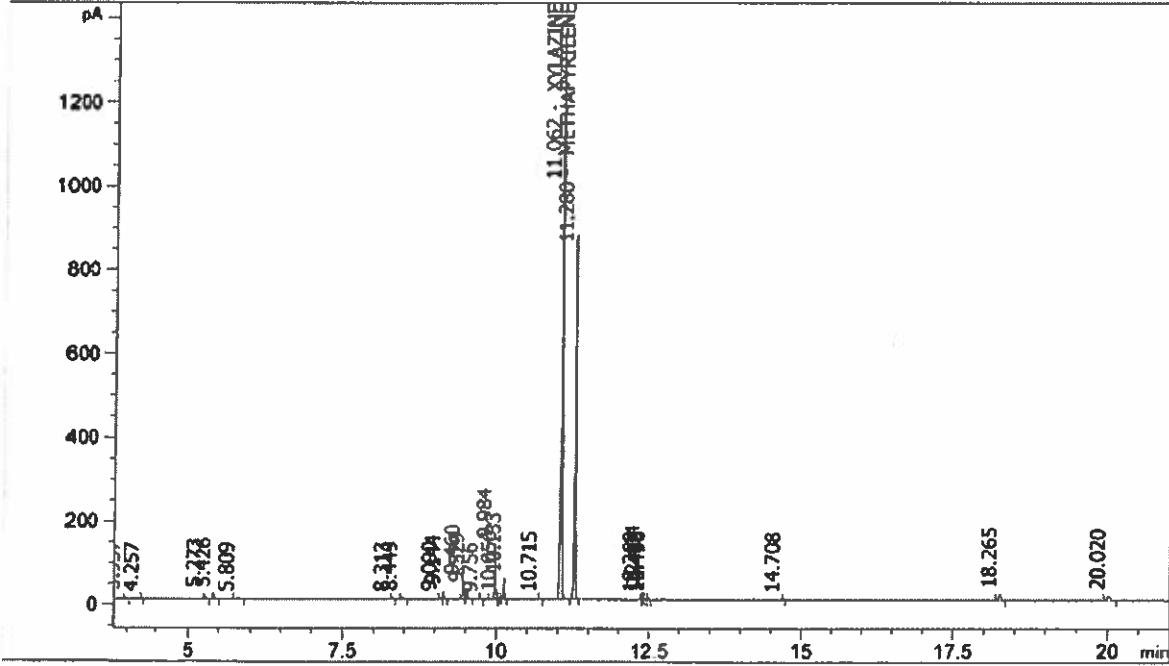


### Library Spectrum



Injection Date: 9/28/2021 7:50:15 PM Seq Line: 16  
 Sample Name: MATRIX 6 2.0 mg/L - URN Vial 16  
 Sample Info:

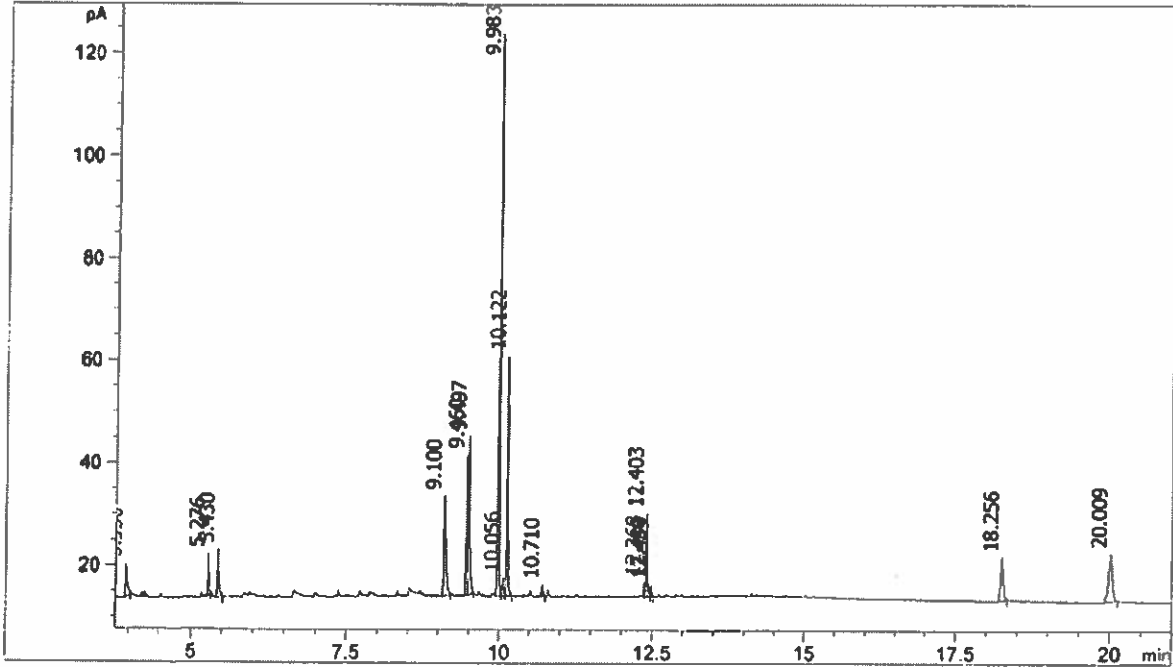
Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
11.062	11.046	2057.273	1355.428	0.000000	XYLAZINE RRT - 0.9807
11.280	11.278	1019.985	866.136	1.000000	METHAPYRILENE

Injection Date: 9/28/2021 8:14:33 PM Seq Line: 17  
 Sample Name: MATRIX BLANK after 2.-> Vial 17  
 Sample Info:

Acq. Method: C:\CHEM32\1\METHODS\ALKALI.M  
 Analysis Method: C:\CHEM32\1\METHODS\092821KMVALI2.M  
 KM/QC DATA IN XYLAZINE VALIDATION



RT [min]	Exp RT [min]	Area	Height	Amount mg/L	Compound
0.000	11.033	0.000	0.000	0.000000	TRAMADOL
0.000	11.046	0.000	0.000	0.000000	XYLAZINE
0.000	11.278	0.000	0.000	0.000000	METHAPYRILENE