



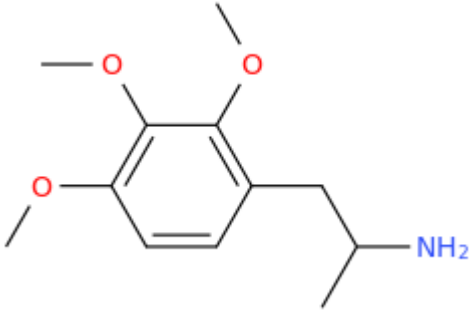
ANALYTICAL REPORT

2,3,4-TMA (C₁₂H₁₉NO₃)

1-(2,3,4-trimethoxyphenyl)propan-2-amine

Remark – other active cpd. detected **none**

Sample ID:	2069-19
Sample description:	powder - white
Sample type:	RM-reference material
Comments:	CAY Lot#0550773-3,
Date of entry (DD/MM/YYYY):	06/06/2019

Substance identified- structure ¹ (base form)	
Systematic name:	1-(2,3,4-trimethoxyphenyl)propan-2-amine
Other names:	TMA-3; 2,3,4-Trimethoxyamphetamine; 2,3,4-trimethoxy- α -methyl-benzeneethanamine
Formula (per base form)	C ₁₂ H ₁₉ NO ₃
M _w (g/mol)	225,29
Salt form:	HCl
StdInChIKey (per base form)	LWDQPPLPHGXYLG-UHFFFAOYSA-N
Other active cpd. detected	none
Add.info (purity..)	≥98%

¹ Created by OPSIN free tool: <http://opsin.ch.cam.ac.uk/> DOI: 10.1021/ci100384d

Report updates

date	comments (explanation)

Supporting information

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 4,61 BP(1): 44; BP(2): 182,BP(3) :167,
FTIR-ATR	+	direct measurement
GC-IR (condensed phase)	+	always as base form
HPLC-TOF	+	exact mass theoretical: 225,1365 / measured Δ ppm: 3,99

1. GC-MS (Agilent): GC-method is RT locked to tetracosane (9.258 min). Injection volume 1 ml and split mode (1:50). Injector temperature: 280 °C. Chromatographic separation: on column HP1-MS (100% dimethylpolysiloxane), length 30 m, internal diameter 0.25 mm, film thickness 0.25 μ m. Carrier gas He: flow-rate 1.2 ml/min. GC oven program: 170 °C for 1 min, followed by heating up to 190 °C at rate 8 °C/min, then heating up to 293 °C at a rate of 18 °C/min, hold for 7.1 min, then heating at 50 °C/min up to 325 °C and finally 6.1 min isothermal. MSD source EI = 70 eV. GC-MS transfer line T= 235°C, source and quadropole temperatures 280°C and 180°C, respectively. Scan range m/z scan range: from 50 (30 until 6 min.) to 550 (300 until 6 min) amu.

2. FTIR-ATR (Perkin Elmer): scan range 4000-400 cm^{-1} ; resolution 4 cm^{-1}

3. GC- (MS)-IR condensed phase (GC-MS (Agilent) & IR (Spectra analyses-Danny)

GC-method: Injection volume 1 ml and split mode (1:5). Injector temperature 280 °C. Chromatographic separation as above **(1)**. Split MS: IR = 1 : 9.

MSD source EI = 70 eV. GC-MS transfer line T= 235°C, source and quadropole temperatures 280°C and 180°C, respectively. Scan range m/z scan range: from 50 (30 until 6 min.) to 550 (300) amu.

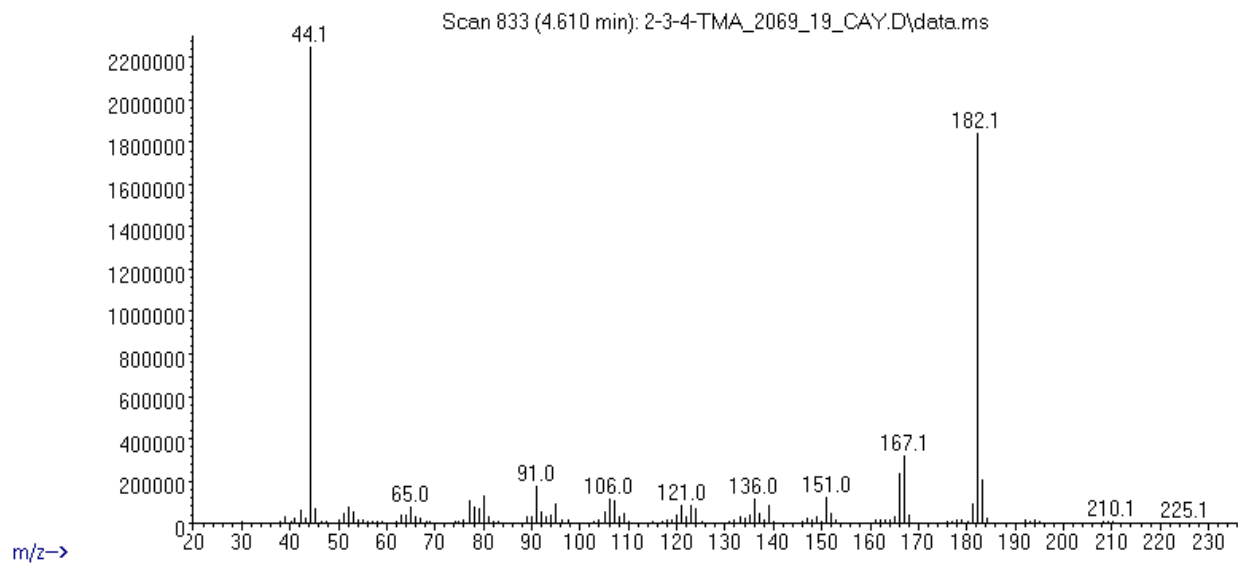
IR (condensed (solid) phase): IR scan range 4000 to 650, resolution 4 cm^{-1} .

4. HPLC-TOF (Agilent): 6230B TOF with Agilent 1260 Infinity HPLC with binary pump, column: Zorbax Eclipse XDB-C18, 50 x 4.6 mm, 1.8 micron. Mobile phases (A) 0.1% formic acid and 1mM ammonium formate in water; (B) 0.1% formic acid in methanol (B). Gradient: starting at 5% B, changing to 40% B over 4 min, then to 70% over 2 min and in 5 min to 100%, hold 1 min and back to 5%, equilibration for 1.7 min. The flow rate: 1.0 ml/min; Injection volume 1 μ l. MS parameters: 2GHz, Extended Dynamic range mode to a maximum of 1700 amu, acquisition rate 1.30 spectra/sec. Sample ionisation: by Agilent Jet Stream technology (Dual AJS ESI). Ion source: positive ion scan mode with mass scanning from 82 to 1000 amu. Other TOF parameters: drying gas (N₂) and sheath temperature 325 °C; drying gas flow rate 6 l/min; sheath gas flow rate 8 l/min; nebulizer 25 psig; Vcap. 4000 V; nozzle 2000 V; skimmer 65 V; fragmentor 175 V and Octopole RF 750 V.

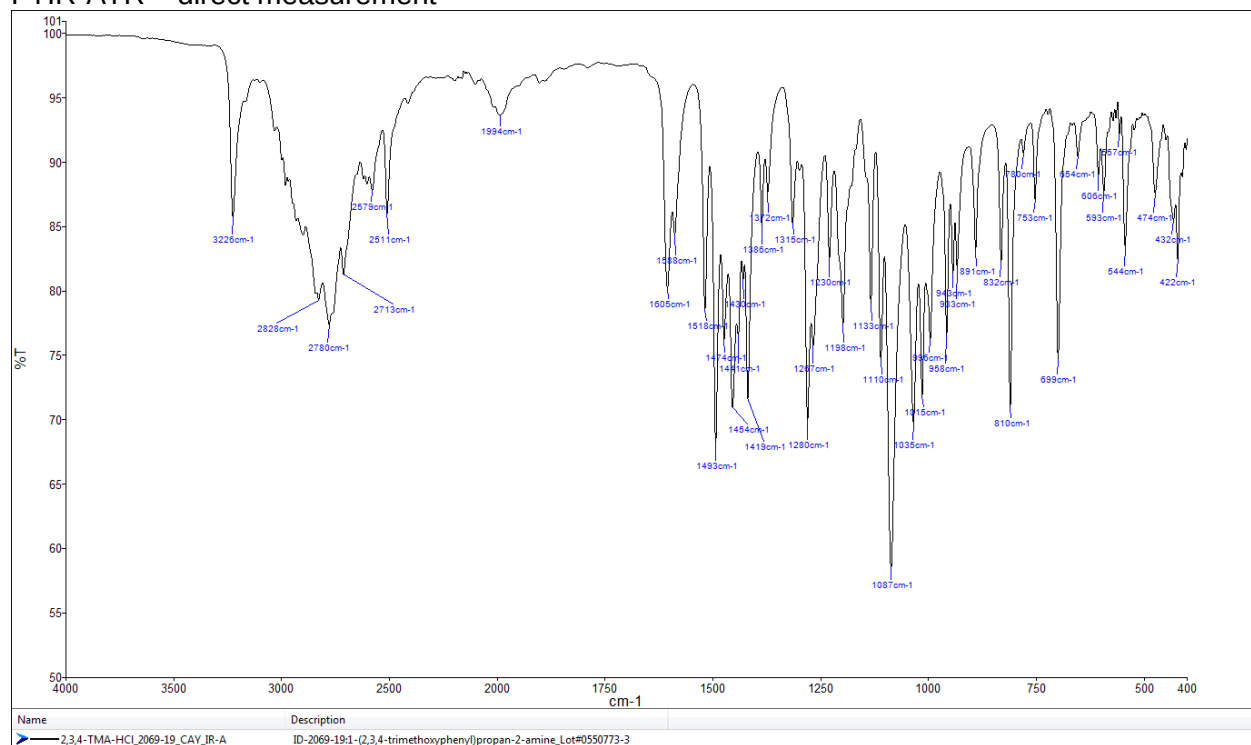
ANALYTICAL RESULTS

MS (EI)

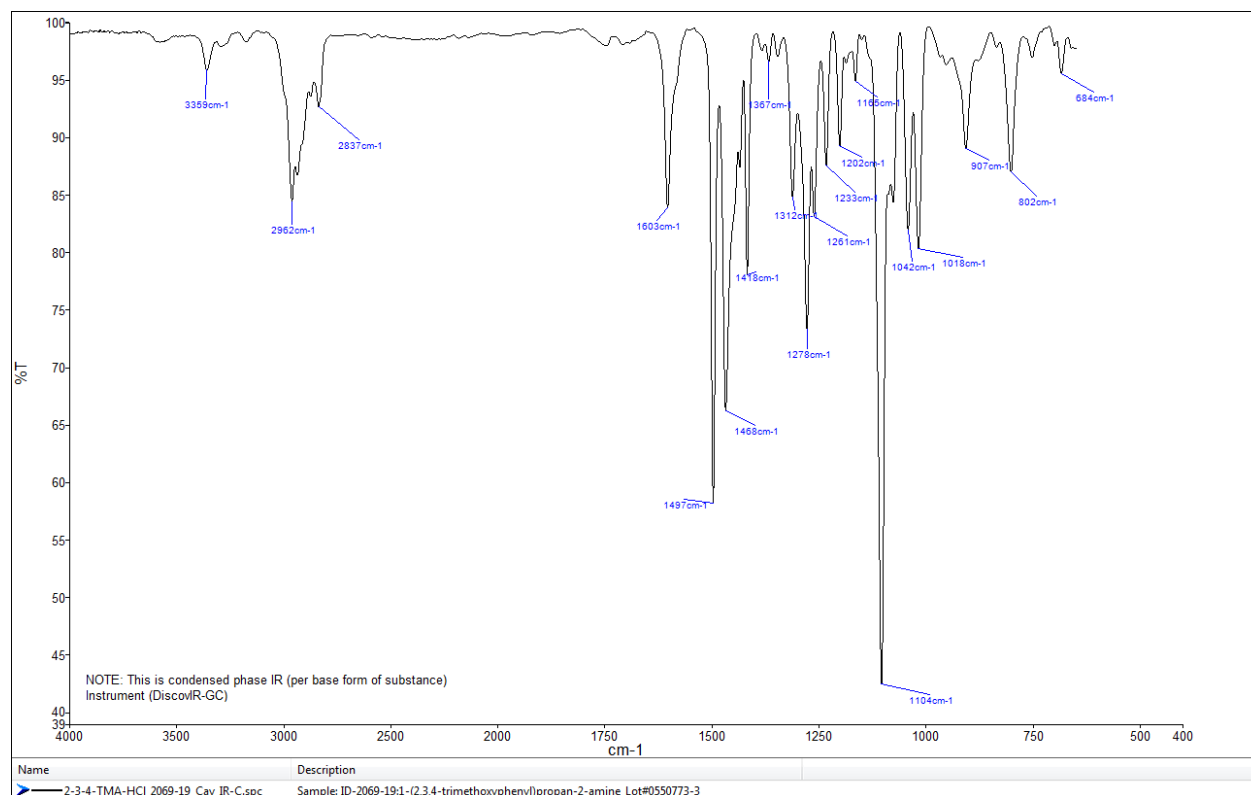
Abundance



FTIR-ATR – direct measurement



IR- (condensed (solid) phase – after chromatographic separation) - spectrum per base form



TOF REPORT

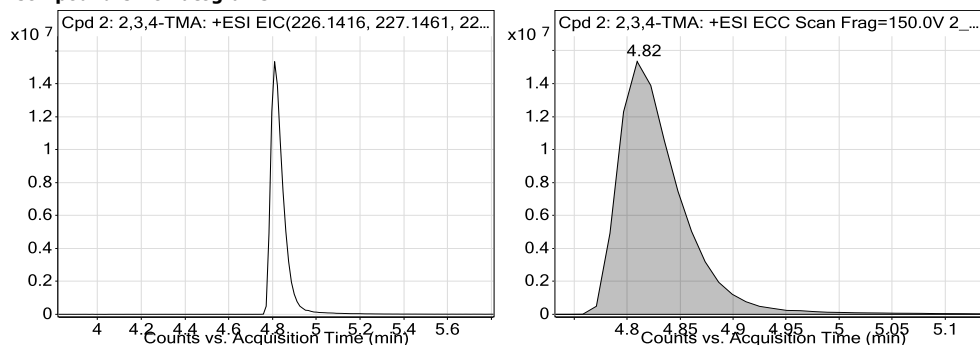
Data File	2_3_4-TMA_2069_19.d	Sample Name	ID-2069-19
Sample Type	Sample	Position	P1-B5
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-8_1_2019-XDB-C18-ESI+.m	Acquired Time	5/14/2019 8:02:21 AM
IRM Calibration Status	Success	DA Method	a-Drugs_NFL.m
Comment	MeOH		

Compound Table

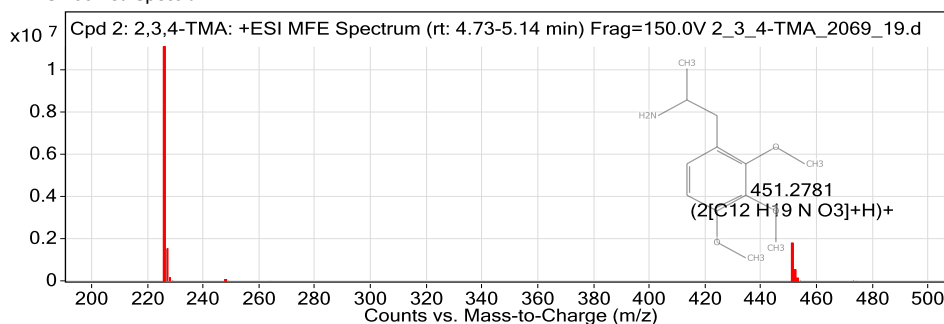
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 2: 2,3,4-TMA	2,3,4-TMA	C12 H19 N O3	4.82	225.1356

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
2,3,4-TMA	226.1431	4.82	225.1356	4.82	C12 H19 N O3	225.1365	3.99

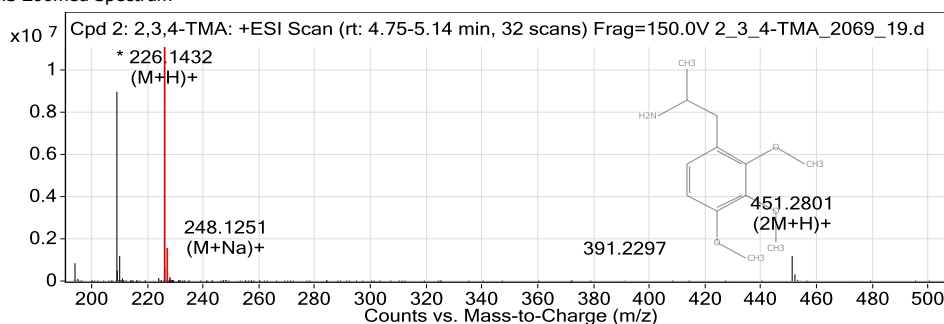
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
226.1431	1	11066632	C12 H19 N O3	(M+H)+
227.1447	1	1508017.89	C12 H19 N O3	(M+H)+
228.1469	1	161165.41	C12 H19 N O3	(M+H)+
229.15	1	14459.12	C12 H19 N O3	(M+H)+
248.123	1	50174.8	C12 H19 N O3	(M+Na)+
451.2781	1	1805327.5	C12 H19 N O3	(2M+H)+
452.2814	1	484655.43	C12 H19 N O3	(2M+H)+
453.2837	1	84737.05	C12 H19 N O3	(2M+H)+
454.2857	1	11782.06	C12 H19 N O3	(2M+H)+
473.2603	1	15234.68	C12 H19 N O3	(2M+Na)+

--- End Of Report ---

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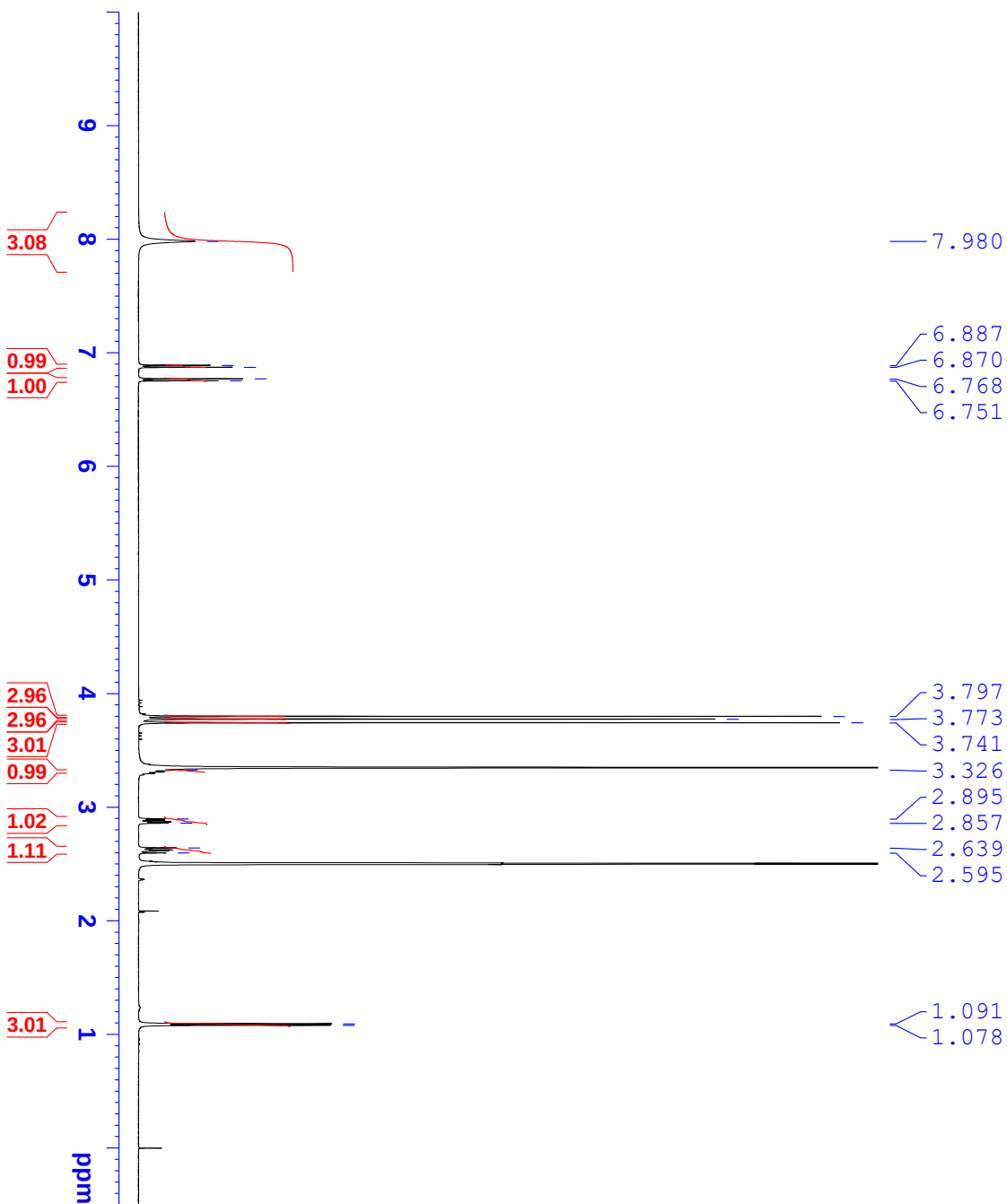
University
of Ljubljana
Faculty of Chemistry
and Chemical Technology



R E P O R T

Contract No.	C1714-17-460078 (Republic of Slovenia, Ministry of the Interior, POLICE)
Sample ID:	2069-19
Received date:	May 20, 2019
Our notebook code:	NFL-2069-19
NMR sample preparation:	2.9 mg dissolved in 0.7 mL DMSO- d_6
NMR experiments:	^1H , ^{13}C , ^1H - ^1H <i>gs</i> -COSY, ^1H - ^{13}C <i>gs</i> -HSQC, ^1H - ^{13}C <i>gs</i> -HMBC, ^1H - ^{15}N <i>gs</i> -HMBC
Proposed structure with formula, exact mass, molecular weight:	Chemical Formula: $\text{C}_{12}\text{H}_{20}\text{NO}_3^+$ Exact Mass: 226,1438 Molecular Weight: 226,2955
Chemical name:	<i>N</i> -protonated 1-(2,3,4-trimethoxyphenyl)propan-2-amine
Comments:	- Structure elucidation based on 1D and 2D NMR spectra and HRMS. - >96% purity of a sample based on ^1H NMR spectrum.
Supporting information:	Copies of ^1H and ^{13}C NMR spectra, ^1H and ^{13}C FIDs.
Principal investigator:	Prof. Dr. Janez Košmrlj
Date of report:	May 28, 2019

NFL-2069-19
¹H



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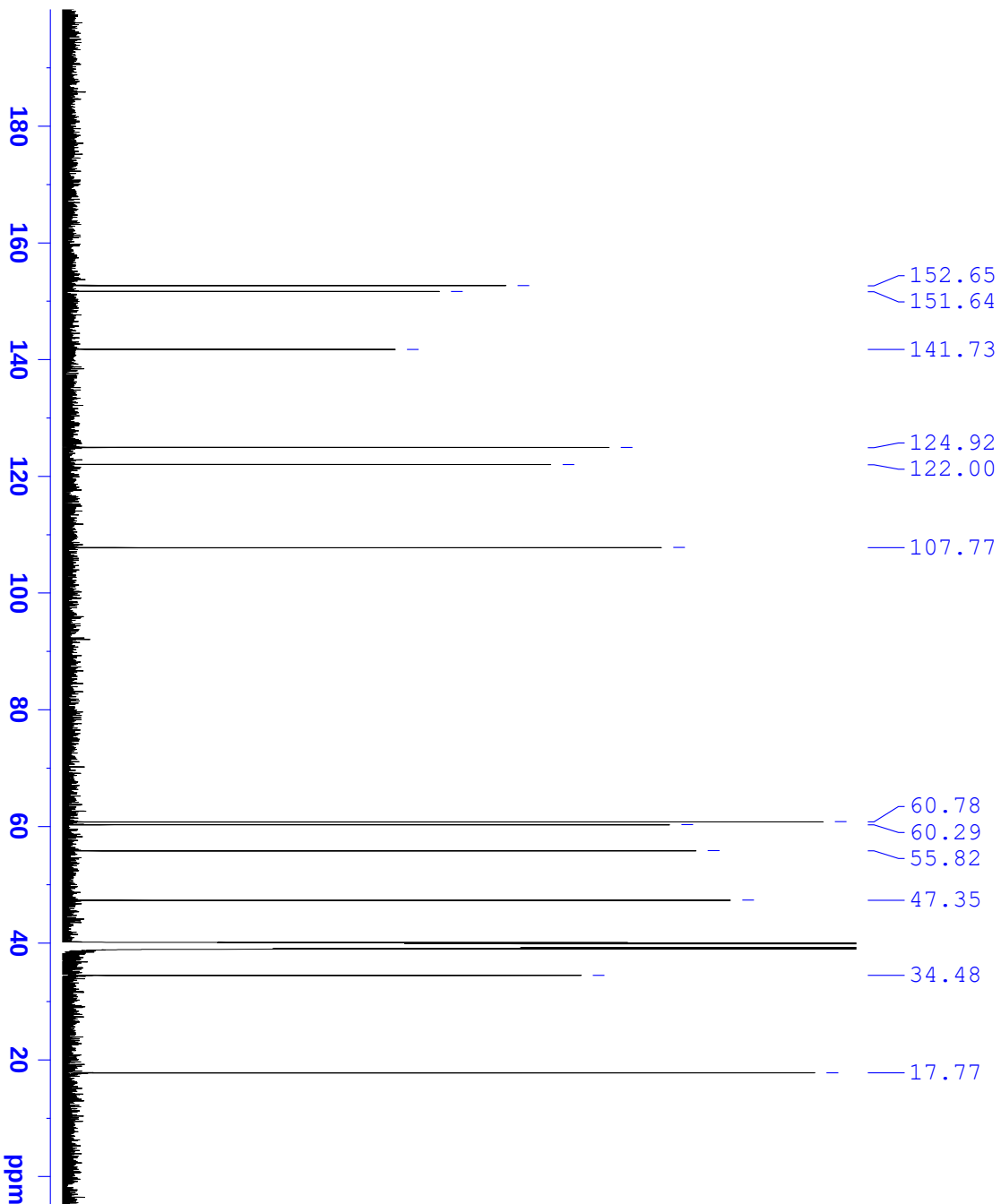
Current Data Parameters
NAME      NFL-2069-19
EXPNO     1
PROCNO    1

F2 - Acquisition Parameters
Date_     20190520
Time      15.31
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zg30
TD         65536
SOLVENT   DMSO
NS         64
DS         2
SWH        10000.000 Hz
FIDRES     0.152588 Hz
AQ         3.276799 sec
RG         128
DW         50.000 usec
DE         6.50 usec
TE         296.0 K
D1         2.00000000 sec
TD0        1

===== CHANNEL f1 =====
SFO1      500.130885 MHz
NUC1      1H
P1        8.70 usec
PLW1      26.00000000 W

F2 - Processing parameters
SI        65536
SF        500.1300047 MHz
WDW       EM
SSB       0
LB        0.30 Hz
GB        0
PC        1.00
  
```

NFL-2069-19
13C



Current Data Parameters
NAME NFL-2069-19
EXPNO 2
PROCNO 1

F2 - Acquisition Parameters
Date_ 20190520
Time 18.33
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
FIDRES 0.454131 Hz
AQ 1.1010048 sec
RG 2050
DW 16.800 usec
DE 6.50 usec
TE 296.0 K
D1 1.0000000 sec
D11 0.0300000 sec
TD0 1

===== CHANNEL f1 =====
SFO1 125.7703637 MHz
NUC1 13C
P1 8.70 usec
PLW1 122.0000000 W

===== CHANNEL f2 =====
SFO2 500.132005 MHz
NUC2 1H
CPDPRG12 waltz16
PCPD2 80.00 usec
PLW2 26.00000000 W
PLW12 0.30046001 W
PLW13 0.15113001 W

F2 - Processing parameters
SI 32768
SF 125.7578452 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40