



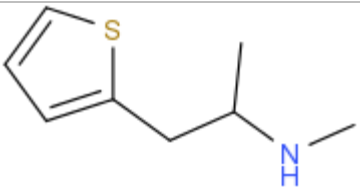
## ANALYTICAL REPORT

**MPA\_Methiopropamine (C<sub>8</sub>H<sub>13</sub>NS)**

**N-Methyl-1-(thiophen-2-yl)propan-2-amine**

Remark – other NPS detected: **none**

Sample ID:	1194-15
Sample description:	powder - white
Sample type:	test purchase (RESPONSE -purchasing)
Comments <sup>1</sup> :	
Date of entry:	9/30/2015

Substance identified- structure <sup>2</sup> (base form)	
Systematic name	N-Methyl-1-(thiophen-2-yl)propan-2-amine
Other names	MPA
Formula (per base form)	C <sub>8</sub> H <sub>13</sub> NS
M <sub>w</sub> (g/mol)	155,26
Salt form	HCl
StdInChIKey	HPHUWHKFQXTZPS-UHFFFAOYSA-N
Compound Class	Arylalkylamines
Other NPS detected	none
Add.info (purity..)	pure by NMR, GC, HPLC

<sup>1</sup> This report has been produced with the financial support of the Prevention of and fight against crime Programme of the European Union (grant agreement number JUST/2013/ISEC/DRUGS/AG/6413). The contents of this report are the sole responsibility of the National Forensic Laboratory and can in no way be taken to reflect the views of the European Commission.

<sup>2</sup> Created by OPSIN free tool: <http://opsin.ch.cam.ac.uk/> DOI: 10.1021/ci100384d

## Report updates

date	comments (explanation)

### Instrumental methods (if applied) in NFL

**1. GC-MS** (Agilent): GC-method is RT locked to tetracosane (RT=9.53 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 mm. Carrier gas He: flow-rate 1.2 ml/min. GC oven program: 170 °C for 1 min, followed by heating up to 293 °C at a rate of 18 °C/min, hold for 6.1 min, then heating at 50 °C/min up to 325 °C and finally 2.8 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 (40) to 550 amu.

**2. 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 µl. 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. 3. FTIR-ATR (Perkin Elmer): scan range 4000-400 cm<sup>-1</sup>; resolution 4 cm<sup>-1</sup>

**3.FTIR-ATR** (Perkin Elmer): scan range 4000-400 cm<sup>-1</sup>; resolution 4cm<sup>-1</sup>

**4. 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 (40) to 550 amu.

IR (condensed phase): IR scan range 4000 to 700, resolution 4 cm<sup>-1</sup>.

## Supporting information

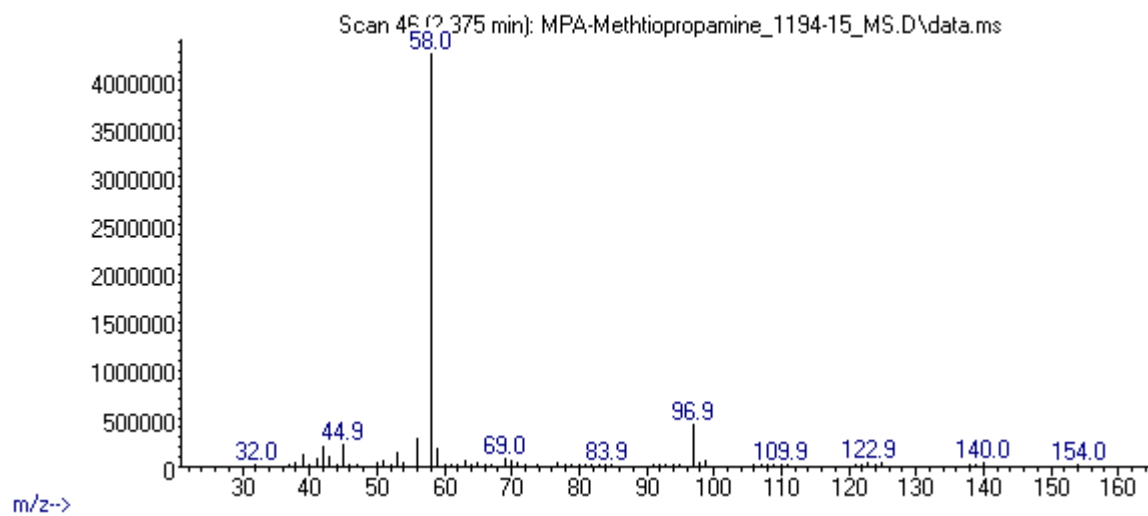
Solubility in	result/remark
CH <sub>2</sub> Cl <sub>2</sub>	soluble
MeOH	soluble
other	not tested

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 2,37 BP(1): 58; BP(2): 97,BP(3) :56,
HPLC-TOF	+	Exact mass (theoretical): 155,0769; measured value Δppm:-2,23; formula:C <sub>8</sub> H <sub>13</sub> N <sub>5</sub>
FTIR-ATR	+	direct measurement
FTIR (condensed phase) always as base form	+	
HPLC-TOF		Exact mass (theoretical): 155,0769; measured value Δppm:-2,23; formula:C <sub>8</sub> H <sub>13</sub> N <sub>5</sub>
NMR	+	
validation		IR-ATR consistent by SWGDRUG lib entry (qM>0.99)
other		

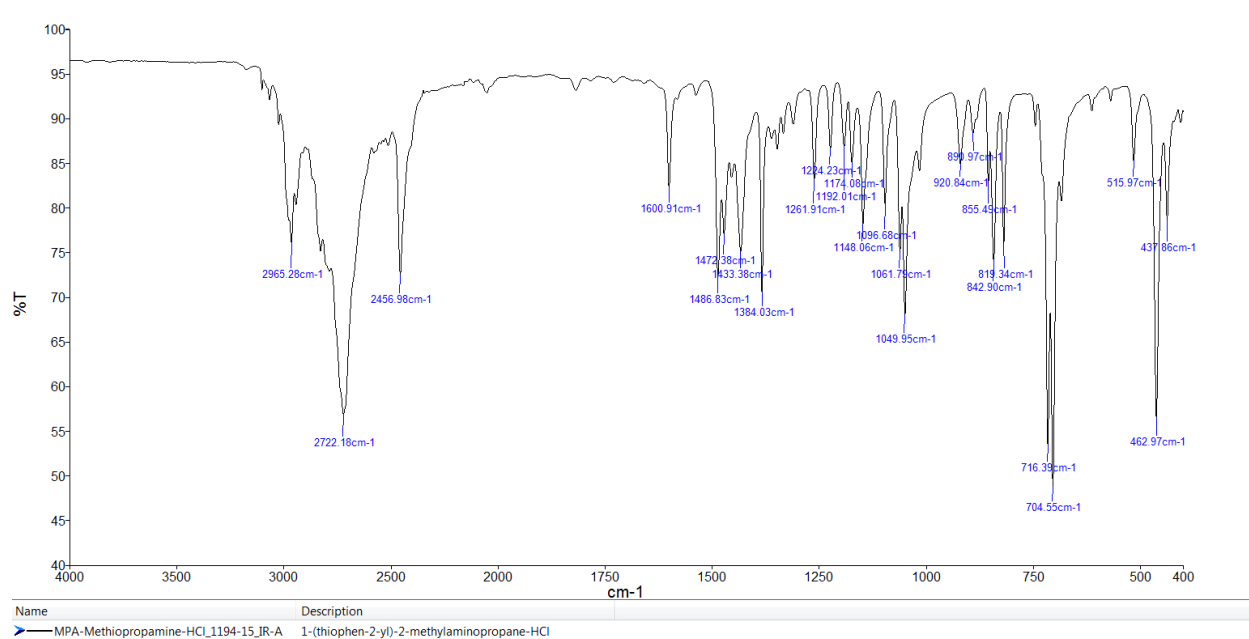
## ANALYTICAL RESULTS

MS (EI)

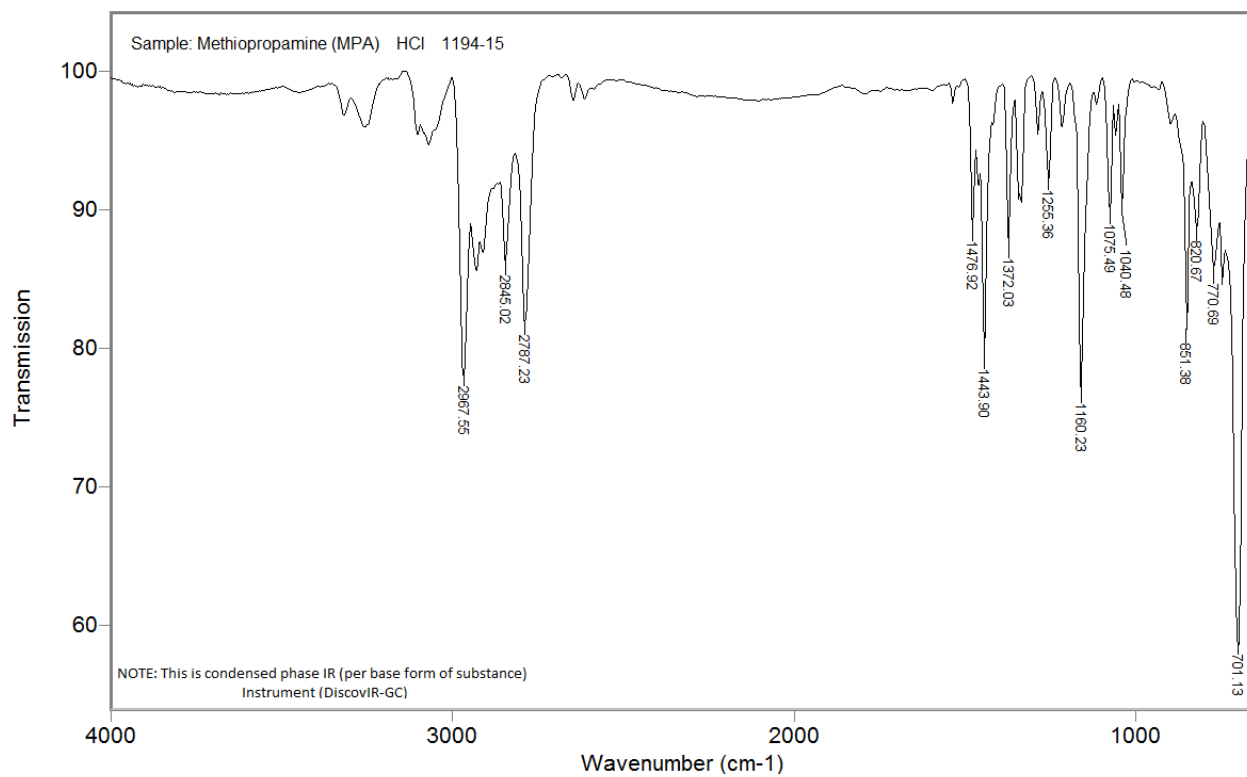
Abundance



## FTIR-ATR



## IR (condensed phase)



# Target Compound Screening Report

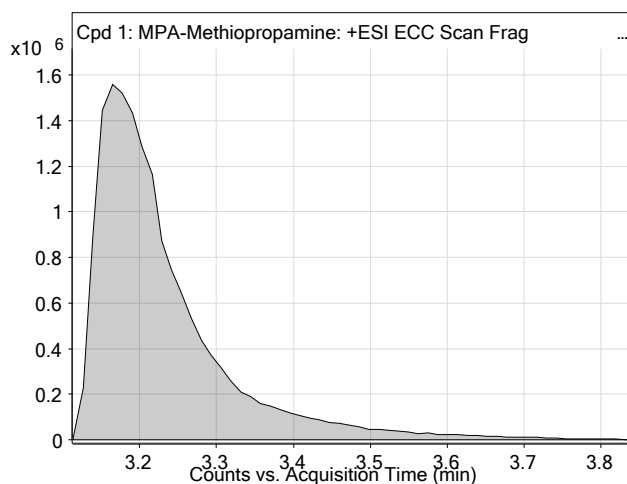
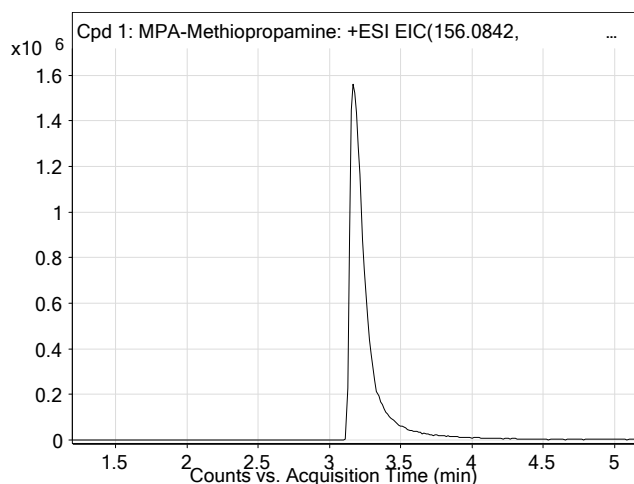
<b>Data File</b>	MPA-Methiopropamine_1194-15_TOF.d	<b>Sample Name</b>	MPA-Methiopropamine
<b>Sample Type</b>	Sample	<b>Position</b>	P1-D4
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	
<b>Acq Method</b>	droge general-13-5-2015-XDB-C18-ESI-poz.m	<b>Acquired Time</b>	7/27/2015 11:07:58 AM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Droge_Default.m
<b>Comment</b>	extract in MeOH		

## Compound Table

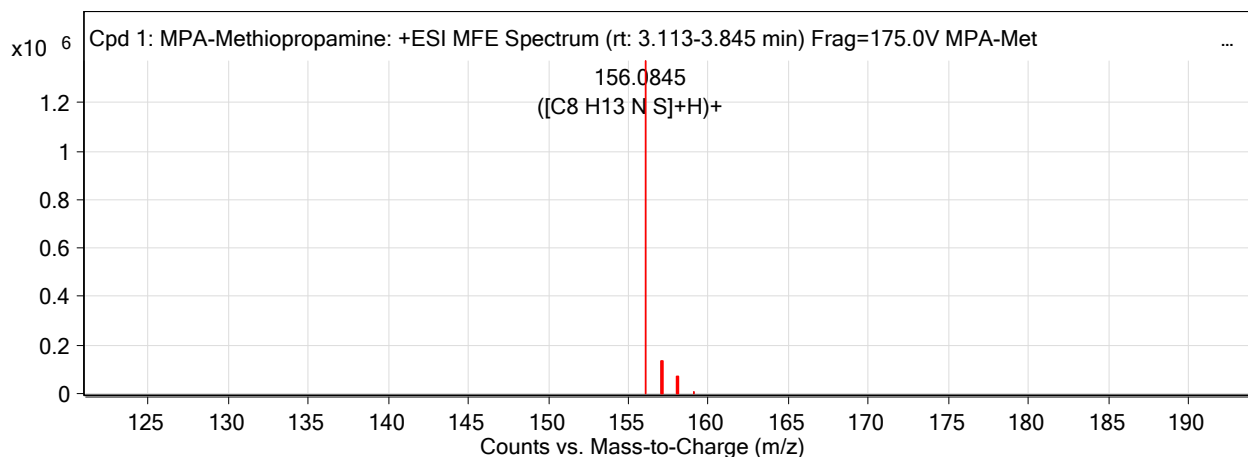
Label	Tgt Name	Obs. RT	Obs. Mass
Cpd 1: MPA-Methiopropamine	MPA-Methiopropamine	3.182	155.0772

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)	Find Cpds Algorithm
MPA-Methiopropamine	156.0845	3.182	155.0772	3.182	C8 H13 N S	155.0769	-2.23	Find by Molecular Feature

## Compound Chromatograms



## MFE MS Zoomed Spectrum

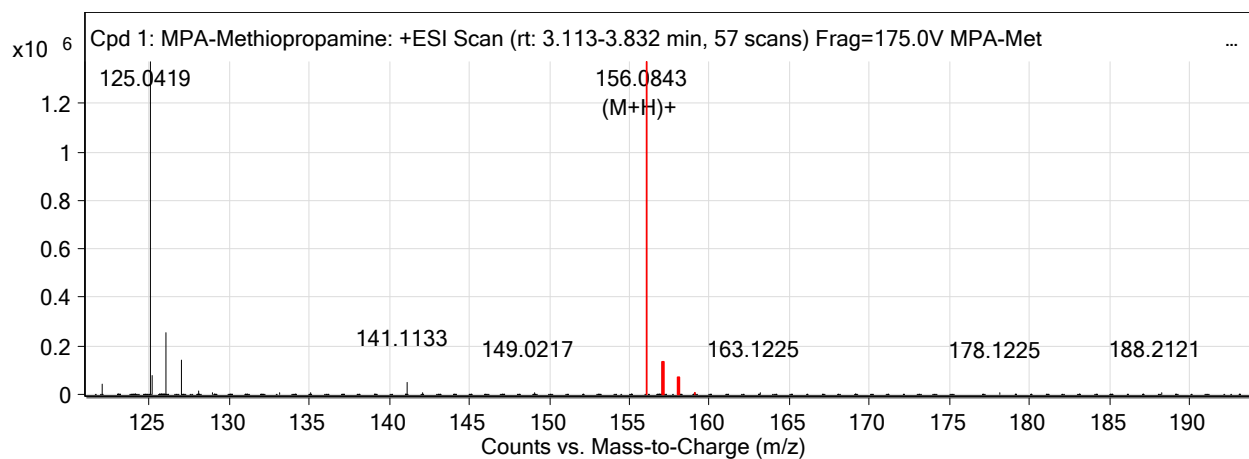


## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
156.0845	1	1372734.88	C8 H13 N S	(M+H)+
157.0873	1	131786.03	C8 H13 N S	(M+H)+
158.0808	1	61929.11	C8 H13 N S	(M+H)+
159.0834	1	5460.93	C8 H13 N S	(M+H)+

## MS Zoomed Spectrum

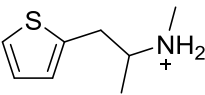
# Target Compound Screening Report



--- End Of Report ---

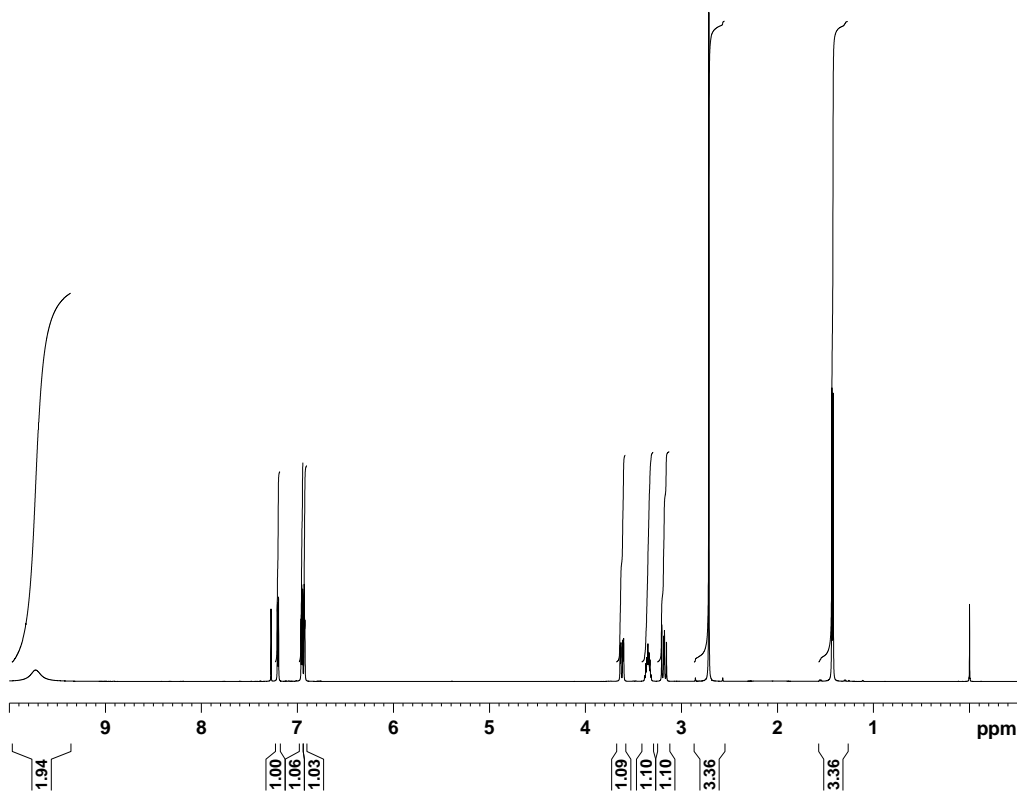


## REPORT

Sample ID:	<b>1194-15</b>
Our notebook code:	P-1194-15
NMR sample preparation:	15 mg dissolved in 0.7 mL CDCl <sub>3</sub>
NMR experiments:	<sup>1</sup> H, <sup>13</sup> C, <sup>1</sup> H- <sup>1</sup> H <i>gs</i> -COSY, <sup>1</sup> H- <sup>13</sup> C <i>gs</i> -HSQC, <sup>1</sup> H- <sup>13</sup> C <i>gs</i> -HMBC, <sup>1</sup> H- <sup>15</sup> N <i>gs</i> -HMBC.
Proposed structure with chemical name:	 N-methyl-1-(thiophen-2-yl)propan-2-aminium ion
Comments:	- Structure elucidation based on 1D and 2D NMR spectra - Compound is pure by NMR
Supporting information:	Copies of <sup>1</sup> H and <sup>13</sup> C NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	September 28, 2015



P-1194-15



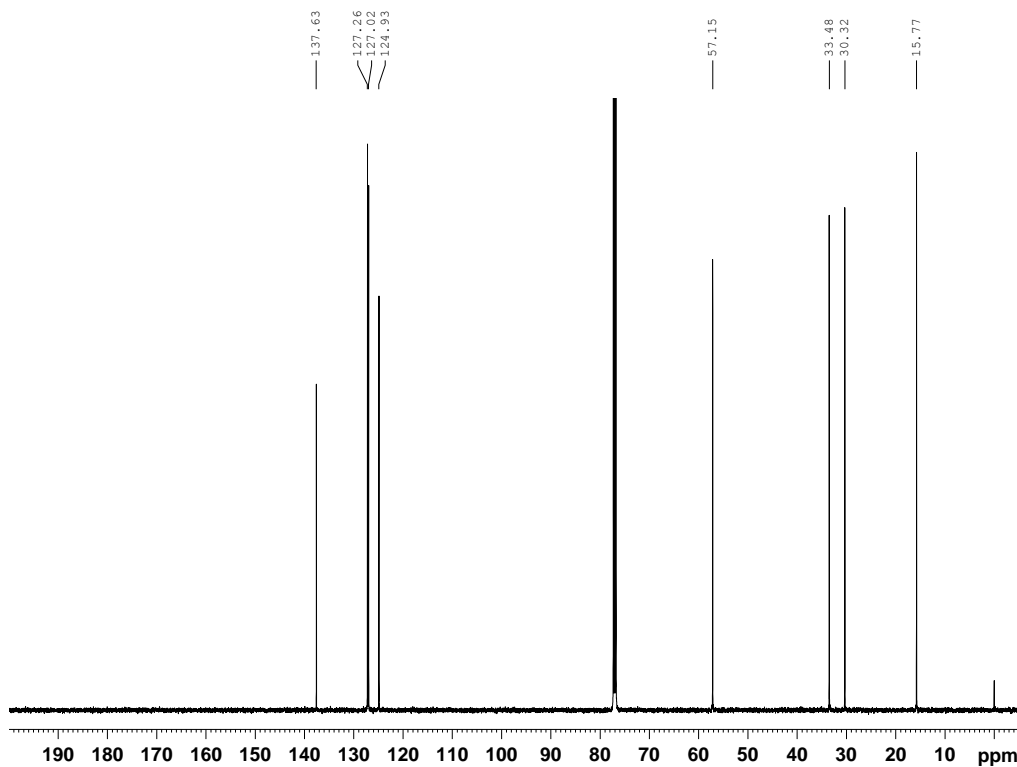
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Current Data Parameters
NAME P-1194-15
EXFNO 1
PROCNO 1

F2 - Acquisition Parameters
Date 20150822
Time 1.24
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CDCl3
NS 16
DS 2
SWH 10330.578 Hz
FIDRES 0.157632 Hz
AQ 3.1719923 sec
RG 90.5
DW 48.400 usec
DE 6.50 usec
TE 296.0 K
D1 1.00000000 sec

===== CHANNEL f1 =====
NUC1 1H
P1 8.90 usec
PLW1 26.00000000 W
SF01 500.1330885 MHz

F2 - Processing parameters
SI 65536
SF 500.1300059 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00
```

P-1194-15



```
Current Data Parameters
NAME P-1194-15
EXFNO 3
PROCNO 1

F2 - Acquisition Parameters
Date 20150822
Time 3.59
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
SOLVENT CDCl3
NS 4096
DS 4
SWH 29761.904 Hz
FIDRES 0.454131 Hz
AQ 1.1010548 sec
RG 2050
DW 16.800 usec
DE 6.50 usec
TE 296.0 K
D1 1.00000000 sec
D11 0.03000000 sec

===== CHANNEL f1 =====
NUC1 13C
P1 9.00 usec
PLW1 122.00000000 W
SF01 125.7703637 MHz

===== CHANNEL f2 =====
CPDPRG2 waltz16
NUC2 1H
PCPD2 80.00 usec
PLW2 26.00000000 W
PLW12 0.32179001 W
PLW13 0.20595001 W
SF02 500.1320005 MHz

F2 - Processing parameters
SI 32768
SF 125.7577906 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40
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