



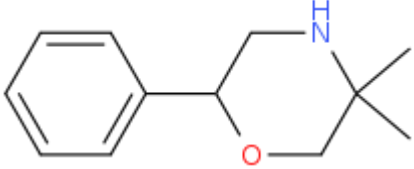
## ANALYTICAL REPORT<sup>1</sup>

**G-130 (C<sub>12</sub>H<sub>17</sub>NO)**

**5,5-dimethyl-2-phenylmorpholine**

Remark – other NPS detected: **none**

Sample ID:	1541-16
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	2/16/2016
Date of entry (M/D/Y) into NFL database:	4/22/2016
Report updates (if any) will be published here:	<a href="http://www.policija.si/apps/nfl_response_web/seznam.php">http://www.policija.si/apps/nfl_response_web/seznam.php</a>

Substance identified - structure <sup>2</sup> (base form)	
Systematic name	5,5-dimethyl-2-phenylmorpholine
Other names	2-Phenyl-5,5-dimethyltetrahydro-1,4-oxazine
Formula (per base form)	C <sub>12</sub> H <sub>17</sub> NO
M <sub>w</sub> (g/mol)	191,27
Salt form/anions detected	hydrochloride
StdInChIKey	KJUOROGOOZJYAI-UHFFFAOYSA-N
Compound Class	Others
Other NPS detected	none
Add.info (purity..)	pure by GC-MS, HPLC-TOF

<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 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) 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 (N<sub>2</sub>) 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 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 (30 until 6 min.) to 550 (300) amu.

IR (condensed phase): IR scan range 4000 to 650, resolution 4 cm<sup>-1</sup>.

**5. IC** (anions) (Thermo Scientific, Dionex ICS 2100), Column: IonPac AS19, 2 x 250mm; Eluent: 10mM from 0 to 10 min, 10-58 mM from 10 to 40min; Flow rate: 0.25 ml/min; Temperature: 30°C; Suppressor: AERS 500 2mm, suppressor current 13mA; Inj. Volume: 25 µl

## Supporting information

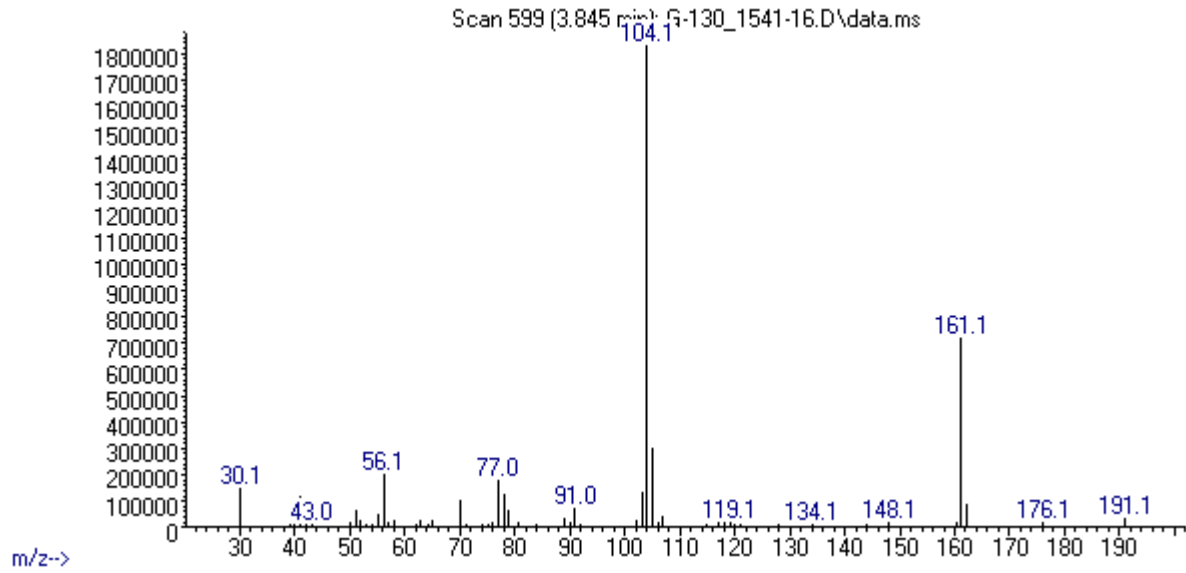
Solubility in	result/remark
CH <sub>2</sub> Cl <sub>2</sub>	soluble
MeOH	soluble
H <sub>2</sub> O	/

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 3,85 BP(1): 104; BP(2): 161, BP(3) :105,
HPLC-TOF	+	Exact mass (theoretical): 191,131; measured value Δppm: -0,41; formula: C <sub>12</sub> H <sub>17</sub> NO
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR (in FKKT)	+	
validation		
other		

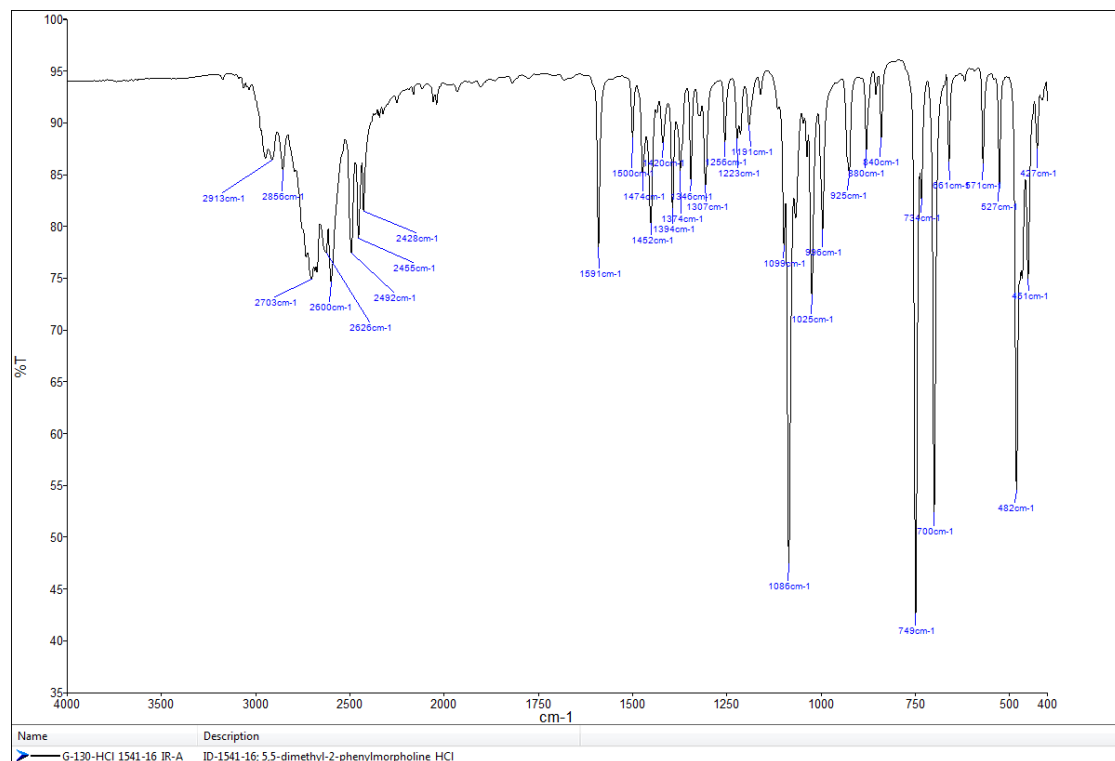
# ANALYTICAL RESULTS

MS (EI)

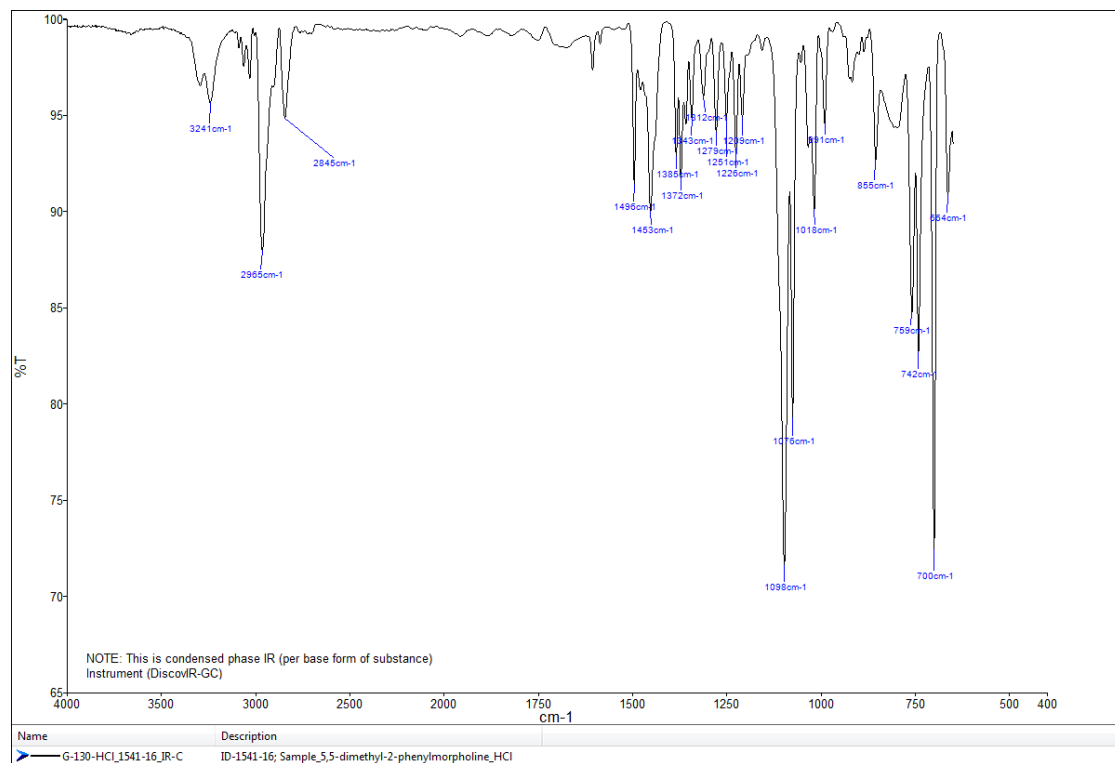
Abundance



## FTIR-ATR - direct measurement (sample as received)



## IR (condensed phase – after chromatographic separation)



# TOF REPORT

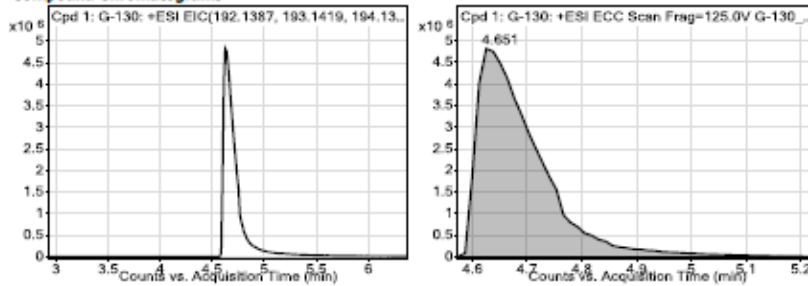
<b>Data File</b>	G-130_1541-16_TOF.d	<b>Sample Name</b>	ID_1541-16
<b>Sample Type</b>	Sample	<b>Position</b>	P1-D9
<b>Instrument Name</b>	62308 TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	general-1512015-XDB-C18-ESI-poz-pod.m	<b>Acquired Time</b>	2/19/2016 12:34:48 PM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Drugs_NFL.m
<b>Comment</b>	extract in MeOH		

**Compound Table**

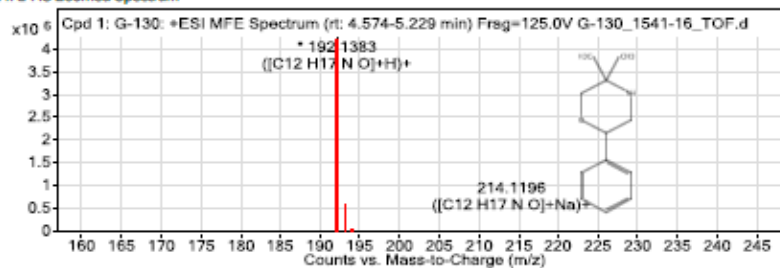
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 1: G-130	G-130	C12 H17 N O	4.651	191.1311

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
G-130	192.1383	4.651	191.1311	4.65	C12 H17 N O	191.131	-0.41

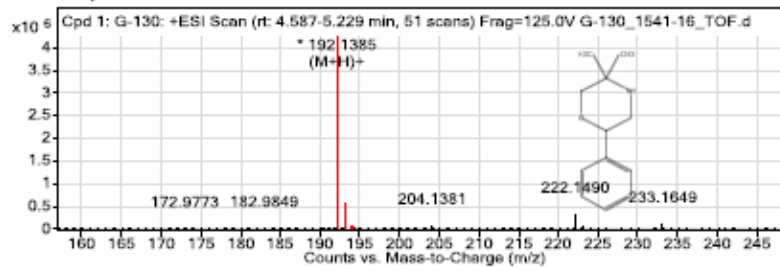
**Compound Chromatograms**



**MFE MS Zoomed Spectrum**



**MS Zoomed Spectrum**



**MS Spectrum Peak List**

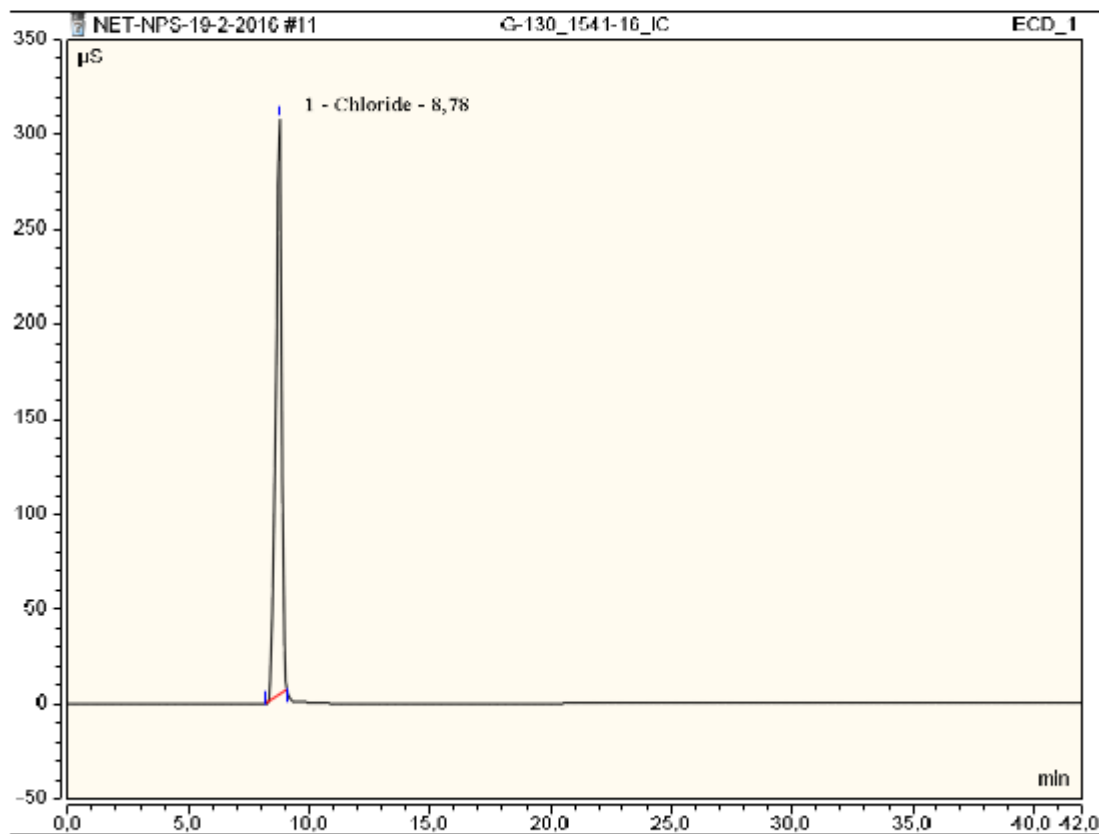
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
192.1383	1	4235083.5	C12 H17 N O	(M+H)+
193.1424	1	532814.67	C12 H17 N O	(M+H)+
194.1444	1	47710.5	C12 H17 N O	(M+H)+
195.146	1	2988.91	C12 H17 N O	(M+H)+
214.1196	1	758.05	C12 H17 N O	(M+Na)+

--- End Of Report ---

### Peak Integration Report

Sample Name:	G-130_1541-16_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	22-feb-2016 / 18:24	Run Time:	42,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}^*\text{min}$	Height $\mu\text{S}$	Amount mg/L
1,00	8,78	Chloride	BMB	84,76	302,99	n.a.
TOTAL:				84,76	302,99	0,00



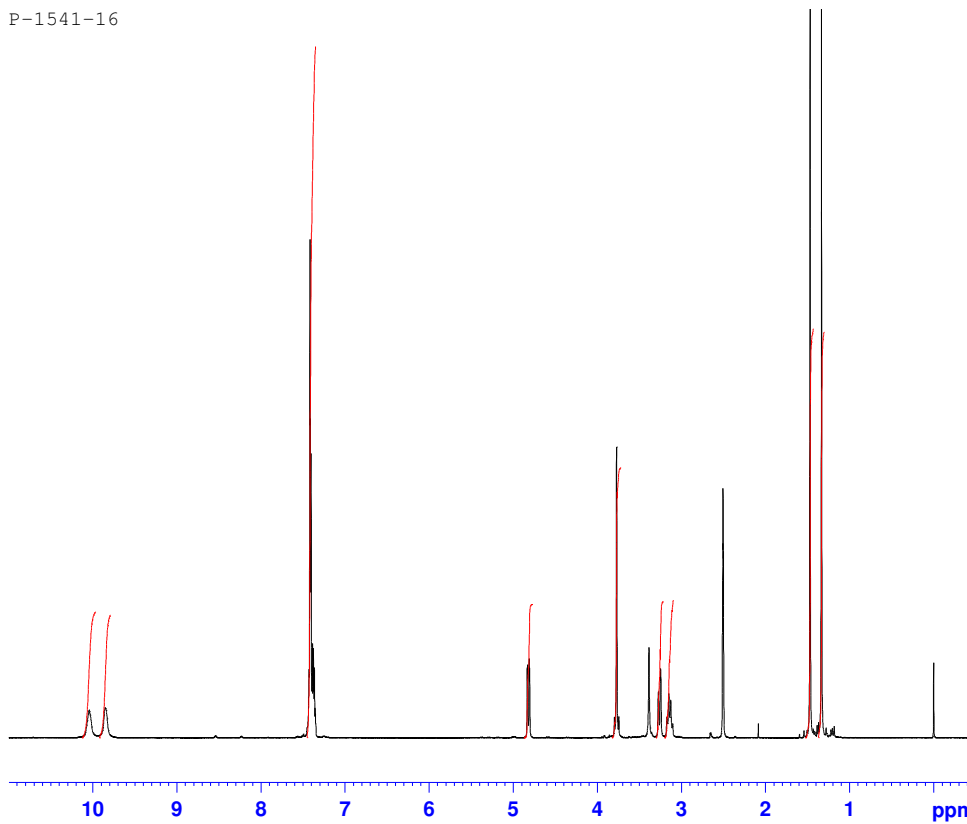


## REPORT

Sample ID:	<b>1541-16</b>
Our notebook code:	P-1541-16
NMR sample preparation:	15 mg dissolved in 0.7 mL DMSO- <i>d</i> <sub>6</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:	
Chemical name:	5,5-dimethyl-2-phenylmorpholin-4-ium cation
Comments:	- Structure elucidation based on 1D and 2D NMR spectra - Sample contains some minor impurities as evident in <sup>1</sup> H 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:	April 21, 2016



P-1541-16



Current Data Parameters  
 NAME P-1541-16  
 EXPNO 1  
 PROCNO 1

F2 - Acquisition Parameters  
 Date\_ 20160413  
 Time 0.58  
 INSTRUM spect  
 PROBHD 5 mm PABBO BB-  
 PULPROG zg30  
 TD 65536  
 SOLVENT DMSO  
 NS 16  
 DS 2  
 SWH 10000.000 Hz  
 FIDRES 0.152588 Hz  
 AQ 3.2768500 sec  
 RG 71.8  
 DW 50.000 usec  
 DE 6.50 usec  
 TE 300.0 K  
 D1 1.00000000 sec  
 TD0 1

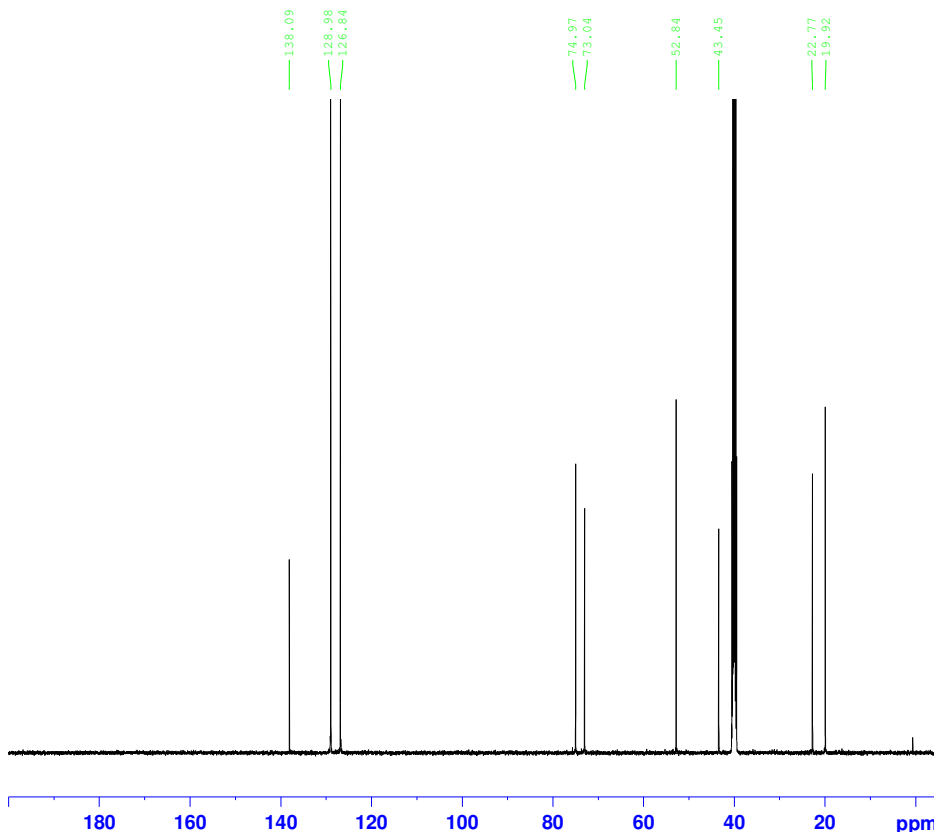
===== CHANNEL f1 =====  
 SFO1 500.1330885 MHz  
 NUC1 1H  
 P1 8.90 usec  
 PLW1 26.00000000 W

F2 - Processing parameters  
 SI 65536  
 SF 500.1300011 MHz  
 WDW EM  
 SSB 0  
 LB 0.30 Hz  
 GB 0  
 PC 1.00

10 9 8 7 6 5 4 3 2 1 ppm

0.93 0.91 5.13 0.99 2.00 1.01 1.02 3.04 3.01

P-1541-16



Current Data Parameters  
 NAME P-1541-16  
 EXPNO 3  
 PROCNO 1

F2 - Acquisition Parameters  
 Date\_ 20160413  
 Time 2.57  
 INSTRUM spect  
 PROBHD 5 mm PABBO BB-  
 PULPROG zgpg30  
 TD 65536  
 SOLVENT DMSO  
 NS 3072  
 DS 4  
 SWH 29761.904 Hz  
 FIDRES 0.454131 Hz  
 AQ 1.1010048 sec  
 RG 2050  
 DW 16.800 usec  
 DE 6.50 usec  
 TE 300.0 K  
 D1 1.00000000 sec  
 D11 0.03000000 sec  
 TD0 1

===== CHANNEL f1 =====  
 SFO1 125.7703637 MHz  
 NUC1 13C  
 P1 9.00 usec  
 PLW1 122.00000000 W

===== CHANNEL f2 =====  
 SFO2 500.1320005 MHz  
 NUC2 1H  
 CPDPRG[2] waltz16  
 PCPD2 80.00 usec  
 PLW2 26.00000000 W  
 PLW12 0.32179001 W  
 PLW13 0.16186000 W

F2 - Processing parameters  
 SI 32768  
 SF 125.7579245 MHz  
 WDW EM  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

180 160 140 120 100 80 60 40 20 ppm