

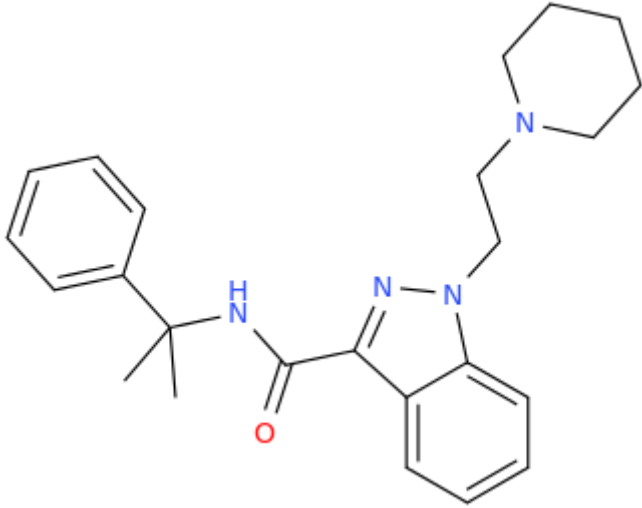
## ANALYTICAL REPORT

CUMYL-PIPETINACA (C<sub>24</sub>H<sub>30</sub>N<sub>4</sub>O)

## N-(2-phenylpropan-2-yl)-1-[2-(piperidin-1-yl)ethyl]-1H-indazole-3-carboxamide

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

Sample ID:	2001-18
Sample description:	liquid - yellow
Sample type:	RM-reference material
Comments:	CHI Lot# 20 907,
Date of entry (DD/MM/YYYY):	22/01/2019

Substance identified-structure <sup>1</sup> (base form)	
Systematic name:	N-(2-phenylpropan-2-yl)-1-[2-(piperidin-1-yl)ethyl]-1H-indazole-3-carboxamide
Other names:	SGT-234; N-(2-Phenylpropan-2-yl)-1-[2-(piperidin-1-yl)ethyl]-1H-indazole-3-carboxamide
Formula (per base form)	C <sub>24</sub> H <sub>30</sub> N <sub>4</sub> O
M <sub>w</sub> (g/mol)	390,53
Salt form:	base
StdInChIKey (per base form)	MTMVUBQQIGZSQB-UHFFFAOYSA-N
Other active cpd. detected	none
Add.info (purity..)	97,8 %

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

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## Report updates

date	comments (explanation)

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## Supporting information

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 14,6 BP(1): 98; BP(2): 99,BP(3) :103,
FTIR-ATR	+	direct measurement
GC-IR (condensed phase)	+	always as base form
HPLC-TOF	+	exact mass theoretical: 390,242 / measured $\Delta$ ppm: -0,82

**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  $\mu$ 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  $\text{cm}^{-1}$ ; resolution 4 $\text{cm}^{-1}$

**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.

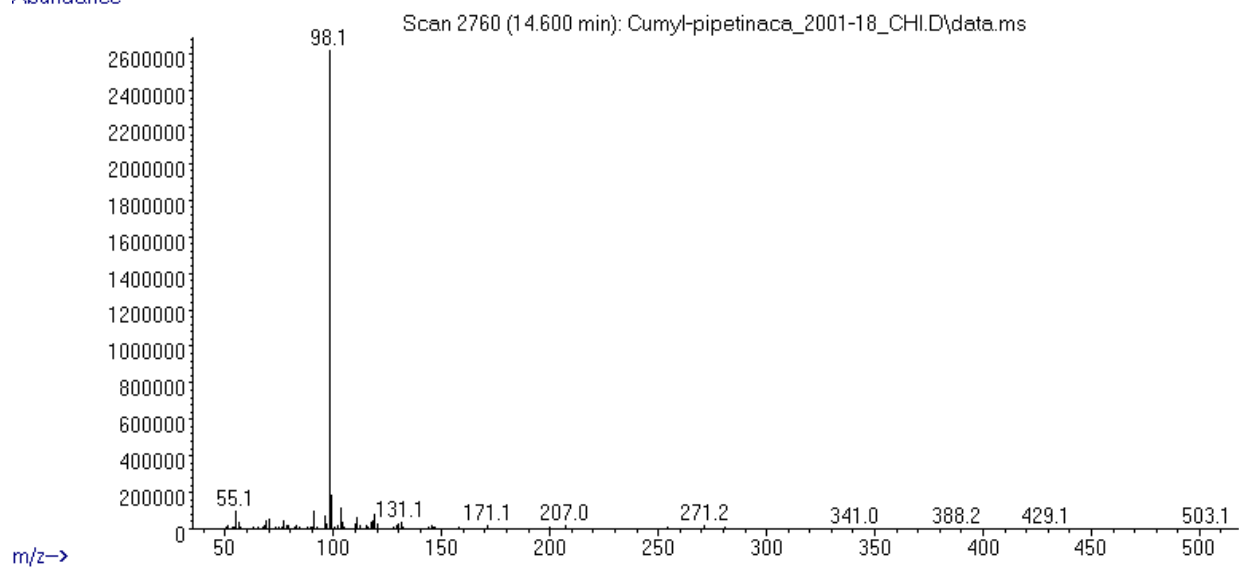
IR (condensed (solid) phase): IR scan range 4000 to 650, resolution 4  $\text{cm}^{-1}$ .

**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  $\mu$ 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 (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.

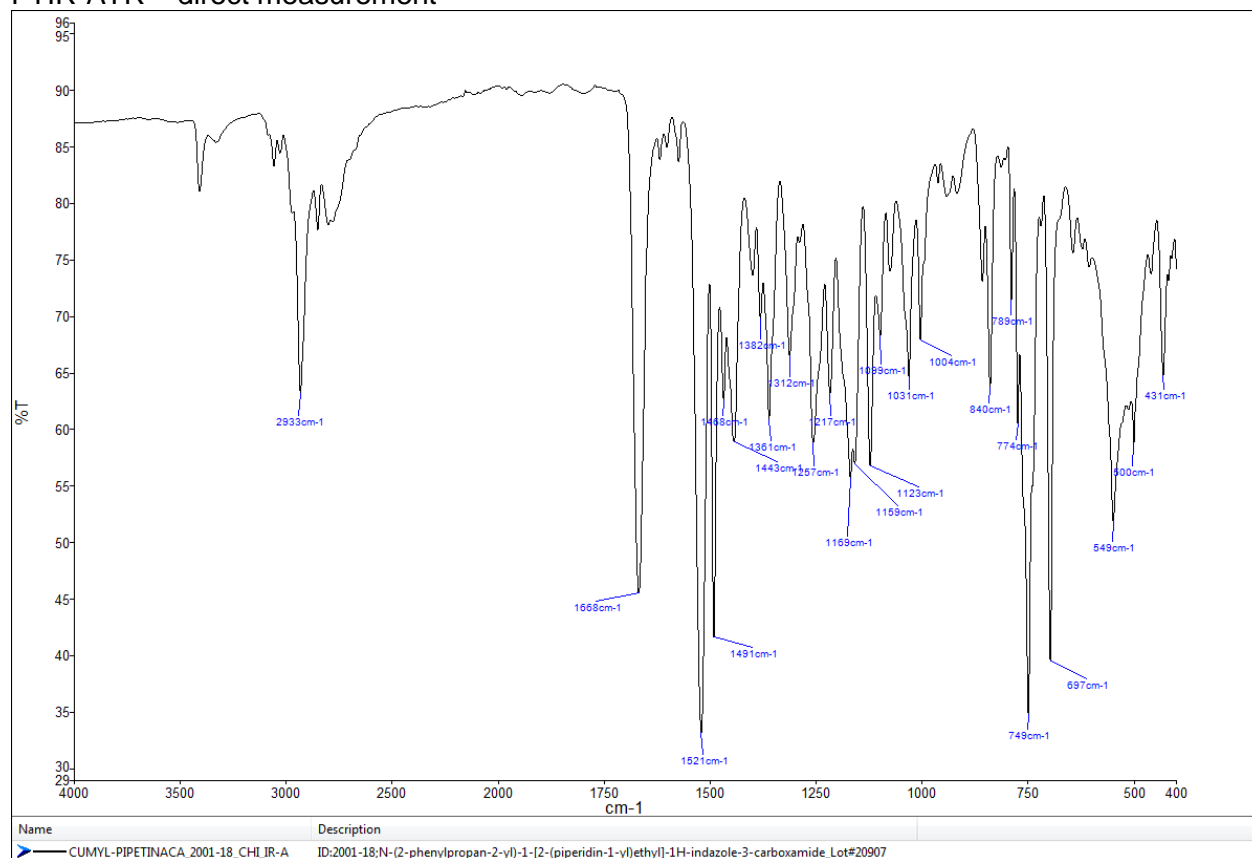
# ANALYTICAL RESULTS

MS (EI)

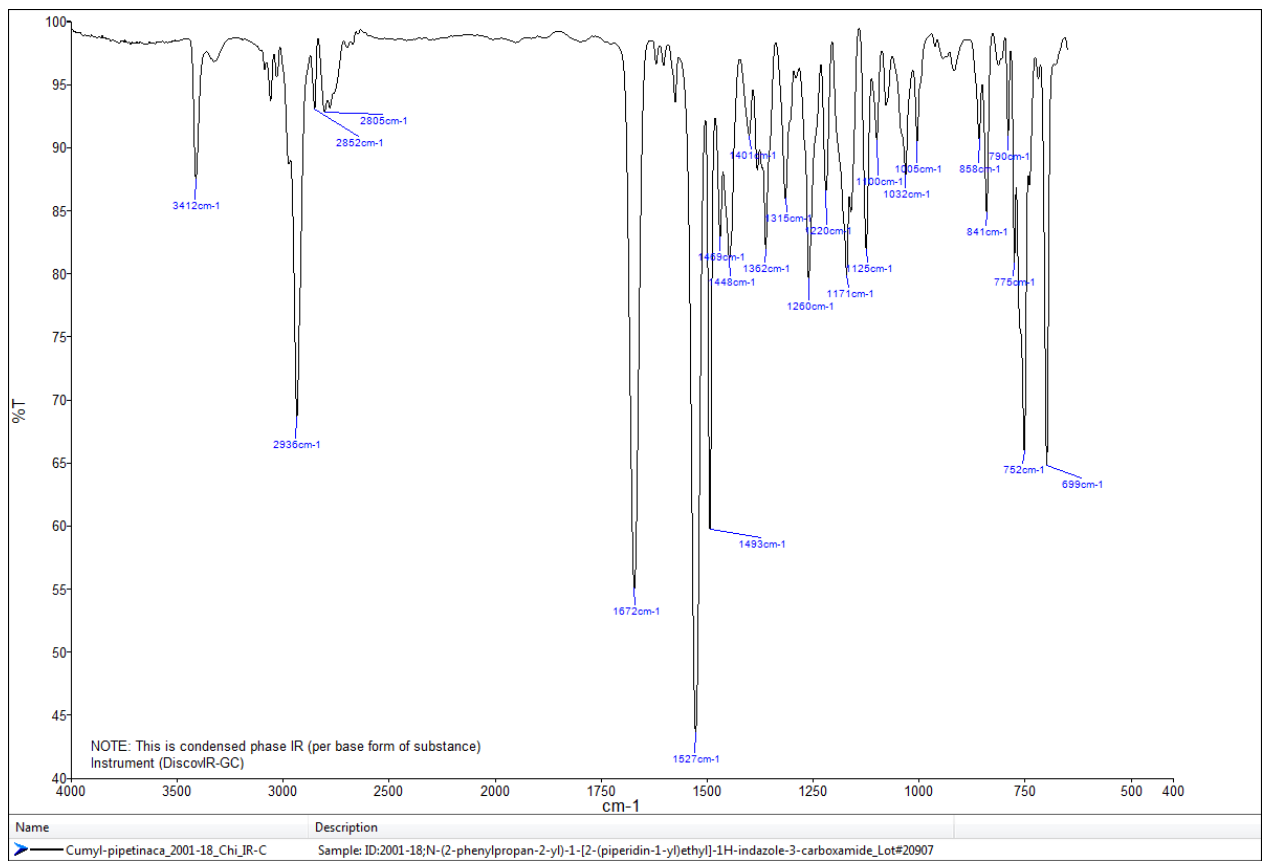
Abundance



# FTIR-ATR – direct measurement



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

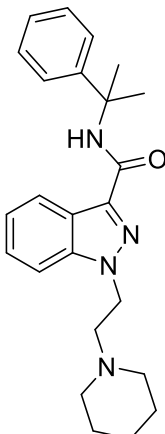


University  
of Ljubljana

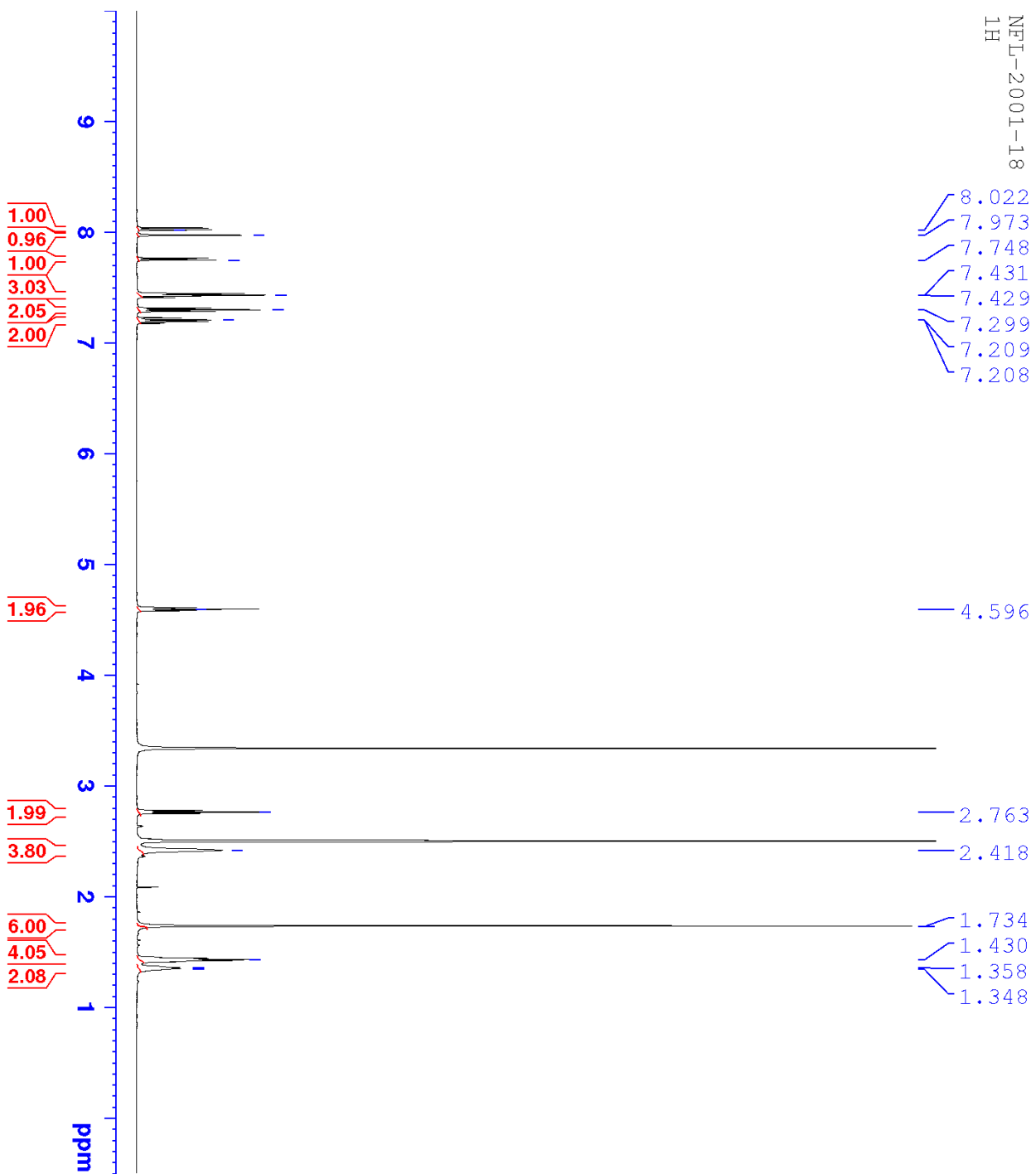
Faculty of Chemistry  
and Chemical Technology



## R E P O R T

Contract No.	C1714-17-460078 (Republic of Slovenia, Ministry of the Interior, POLICE)
Sample ID:	<b>2001-18</b>
Received date:	November 29, 2018
Our notebook code:	NFL-2001-18
NMR sample preparation:	5.0 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, <sup>19</sup> F
Proposed structure with formula, exact mass, molecular weight:	 <p>Chemical Formula: C<sub>24</sub>H<sub>30</sub>N<sub>4</sub>O Exact Mass: 390,2420 Molecular Weight: 390,5310</p>
Chemical name:	<i>N</i> -(2-phenylpropan-2-yl)-1-(2-(piperidin-1-yl)ethyl)-1H-indazole-3-carboxamide
Comments:	- Structure elucidation based on 1D and 2D NMR spectra and HRMS. - >98% purity of a sample based on <sup>1</sup> H NMR spectrum
Supporting information:	Copies of <sup>1</sup> H and <sup>13</sup> C NMR spectra, <sup>1</sup> H and <sup>13</sup> C FIDs.
Principal investigator:	Prof. Dr. Janez Košmrlj
Date of report:	December 13, 2018

NFL-2001-18  
1H



Current Data Parameters  
NAME NFL-2001-18  
EXPNO 1  
PROCNO 1

F2 - Acquisition Parameters  
Date\_ 20181130  
Time 0.03

INSTRUM spect  
PROBHD 5 mm PABBO BB-  
PULPROG zg30

TD 65536  
SOLVENT DMSO  
NS 32

DS 2  
SWH 10000.000 Hz  
FIDRES 0.152588 Hz

AQ 3.2767999 sec  
RG 114  
DE 50.000 usec  
TE 296.0 K

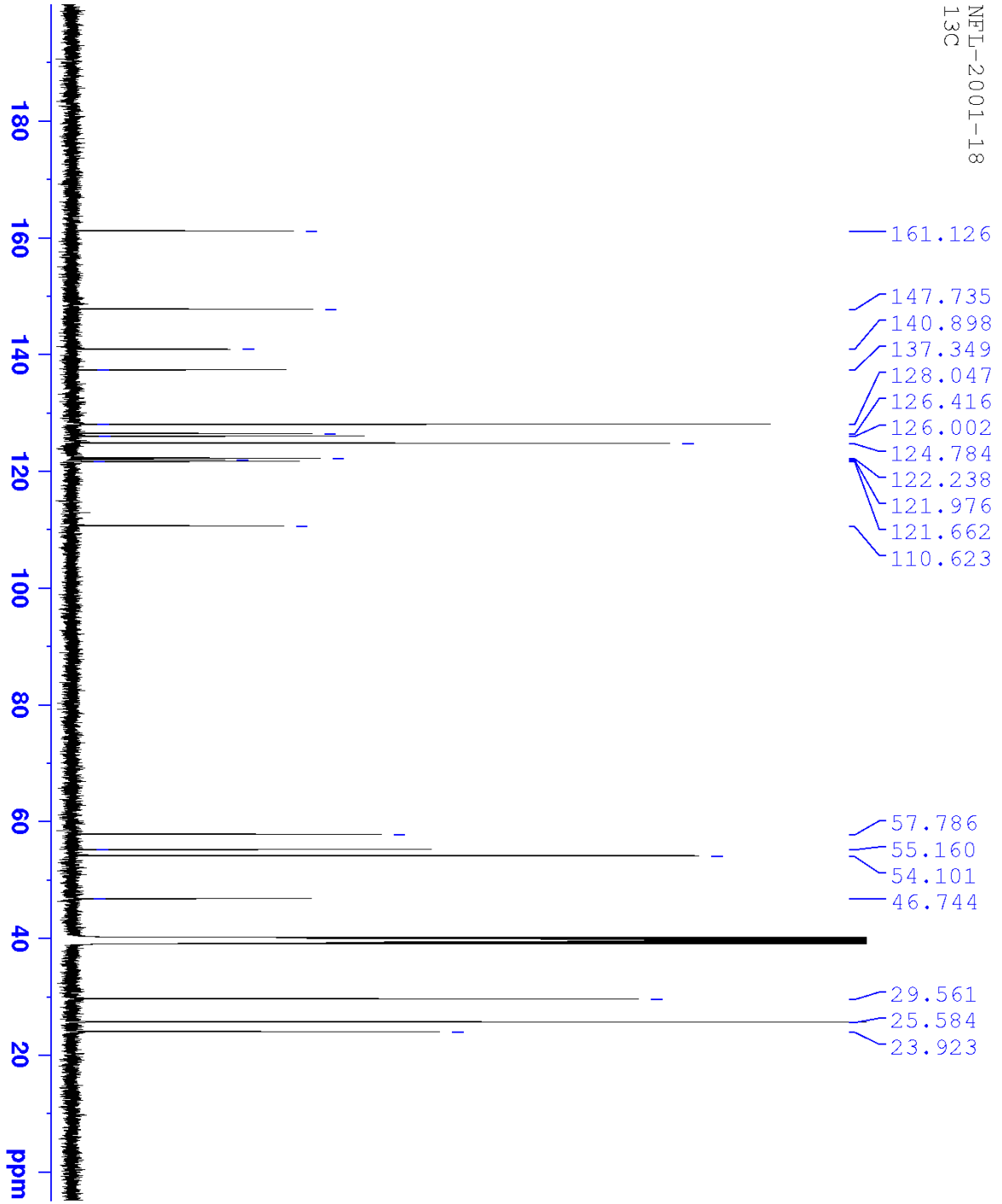
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TD0 1

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NUC1 1H  
P1 8.70 usec  
PLM1 26.00000000 W

F2 - Processing parameters  
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SF 500.1300041 MHz  
WDW EM  
SSB 0

LB 0.30 Hz  
GB 0  
PC 1.00

NFL-2001-18  
13C



Current Data Parameters  
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EXPNO 3  
PROCNO 1

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Date\_ 20181130  
Time 4.01

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PULPROG zgpg30  
TD 65536  
SOLVENT DMSO  
NS 4096  
DS 4

SWH 29761.904 Hz  
FIDRES 0.454131 Hz  
AQ 1.101048 sec  
RG 2050  
DW 16.800 usec  
DE 6.50 usec  
TE 296.0 K  
D1 2.00000000 sec  
D11 0.03000000 sec  
TD0 1

CHANNEL F1 -----  
SFO1 125.7703637 MHz  
NUC1 13C  
P1 8.70 usec  
PLM1 122.00000000 W

CHANNEL F2 -----  
SFO2 506.1320005 MHz  
NUC2 1H  
CPDPRG12 waltz16  
PCPDZ 80.00 usec  
PLM2 26.00000000 W  
PLM12 0.30046001 W  
PLM13 0.15113001 W

F2 - Processing parameters  
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WDW EM  
SSB 0  
LB 1.00 Hz  
GB 0  
PC 1.40