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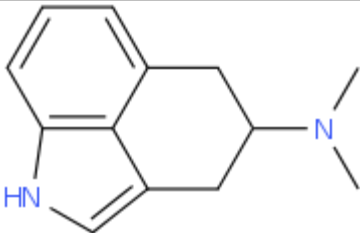
ANALYTICAL REPORT

Ru-28306 (C₁₃H₁₆N₂)

N,N-dimethyl-2-azatricyclo[6.3.1.0^{4,12}]dodeca-1(12),3,8,10-tetraen-6-amine

Remark – other NPS detected: **none**

Sample ID:	1755-17
Sample description:	powder - grey
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	1/13/2017
Date of entry (M/D/Y) into NFL database:	2/23/2017
Report ¹ updates (if any) will be published here:	http://www.policija.si/apps/nfl_response_web/seznam.php

Substance identified - structure ² (base form)	
Systematic name	N,N-dimethyl-2-azatricyclo[6.3.1.0 ^{4,12}]dodeca-1(12),3,8,10-tetraen-6-amine
Other names	N,N-dimethyl-1,3,4,5-tetrahydrobenzo[cd]indol-4-amine, Ru-28306
Formula (per base form)	C ₁₃ H ₁₆ N ₂
M _w (g/mol)	200.29
Salt form/anions detected	HCl
StdInChIKey (for base form)	BQOANWOQEHVATQ-UHFFFAOYSA-N
Other NPS detected	none
Additional info (purity..)	pure by GC-MS, HPLC-TOF, some minor impurities by NMR

¹ 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.

² 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 (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 µm. 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 8.1 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. 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⁻¹; resolution 4cm⁻¹

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 (solid) phase): IR scan range 4000 to 650, resolution 4 cm⁻¹.

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 ₂ Cl ₂	partially
MeOH	soluble
H ₂ O	soluble

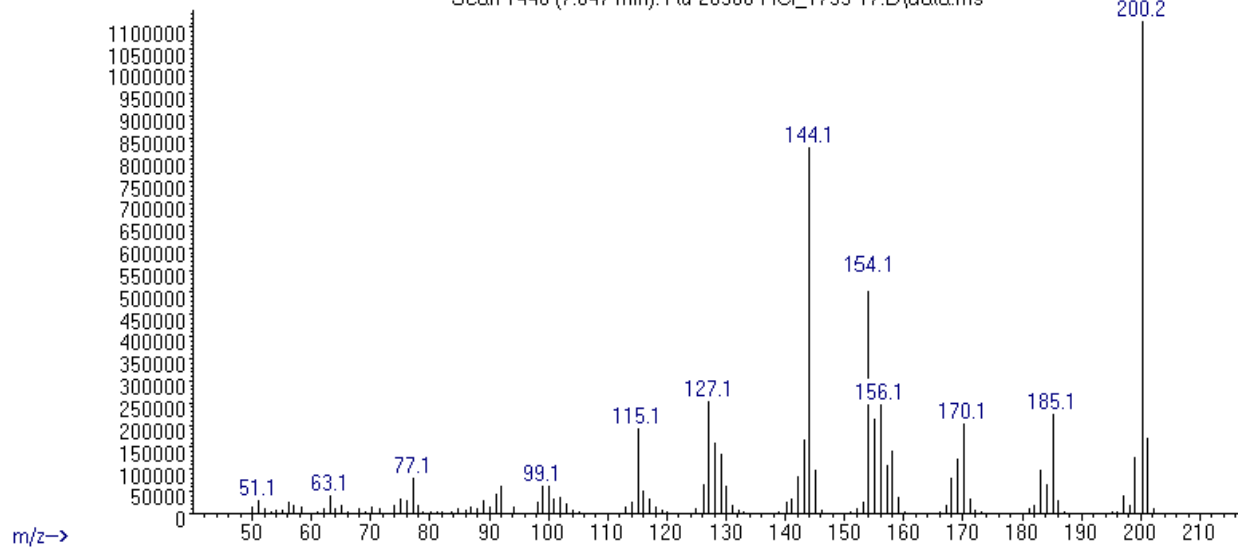
Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 7.05 BP(1): 200; BP(2): 144,BP(3) :154,
HPLC-TOF	+	Exact mass (theoretical): 200.1314; measured value Δppm:-1.02; formula:C13H16N2
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR (in FKKT)	+	
validation		
other		

ANALYTICAL RESULTS

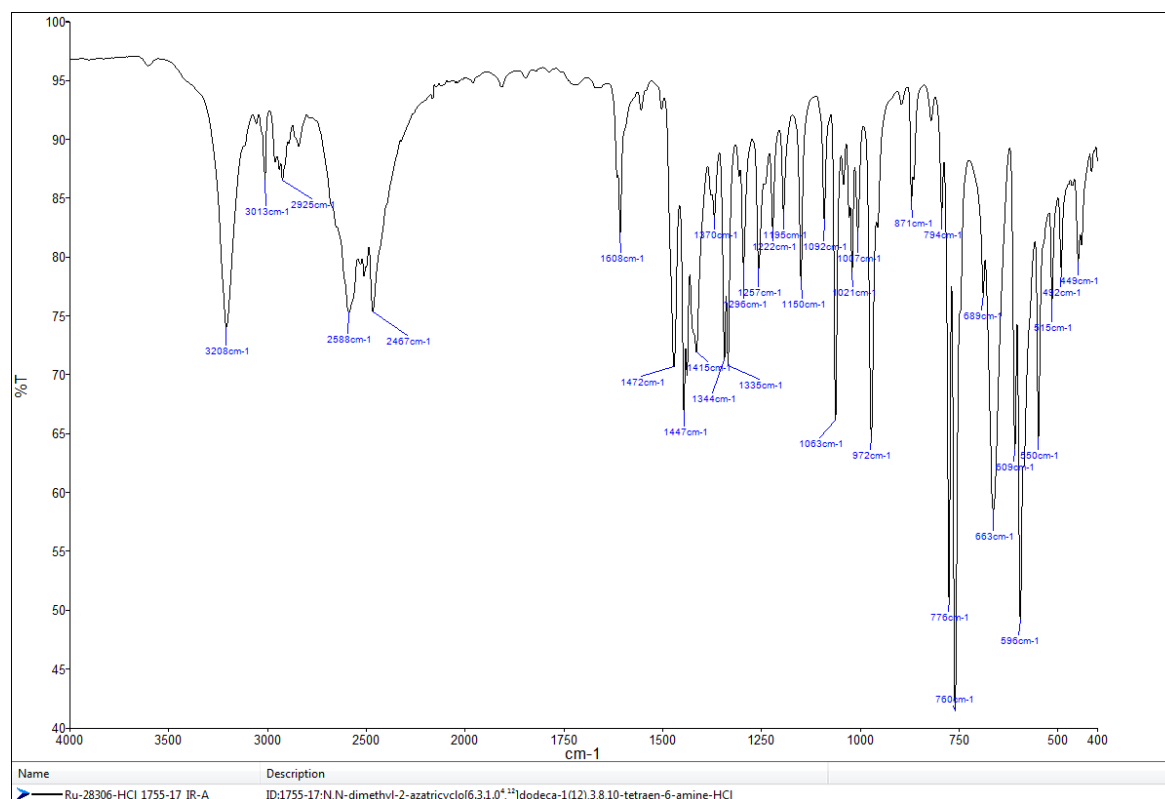
MS (EI)

Abundance

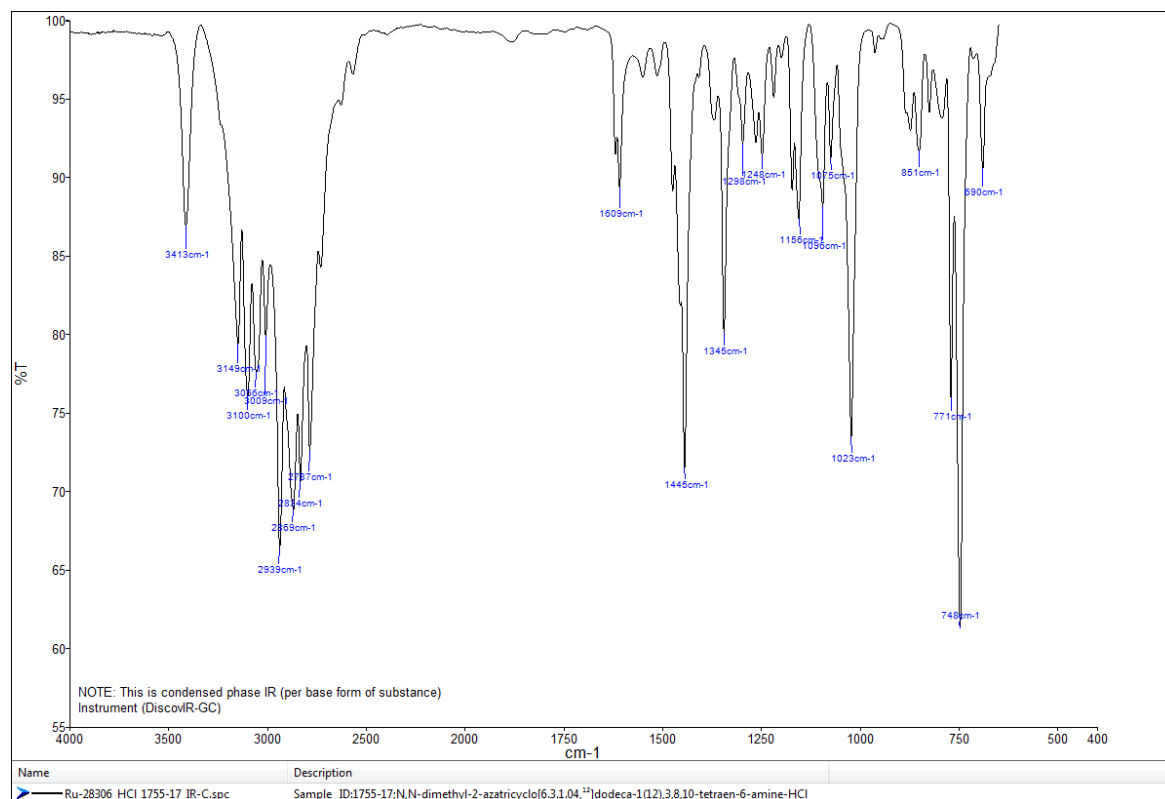
Scan 1440 (7.047 min): Ru-28306-HCl_1755-17.D\data.ms



FTIR-ATR - direct measurement (sample as received)



IR (condensed phase – after chromatographic separation)



TOF REPORT

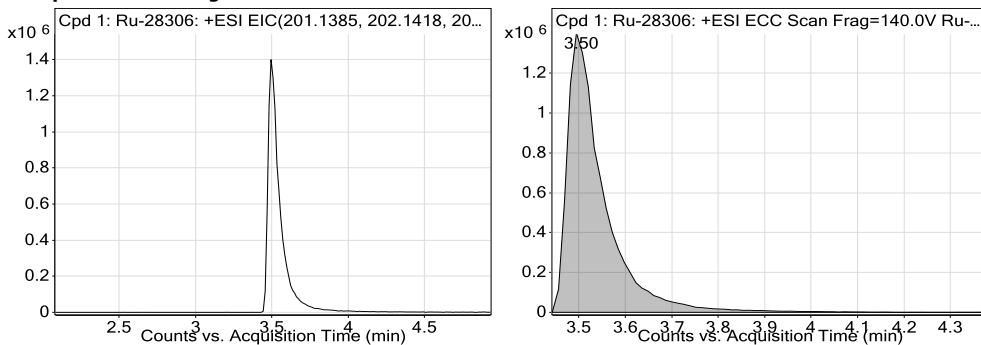
Data File	Ru-28306_1755-17.d	Sample Name	1755-17
Sample Type	Sample	Position	P1-C2
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-20_12_2016-XDB-C18-ESI-poz-soft.m	Acquired Time	1/31/2017 9:34:53 AM
IRM Calibration Status	Success	DA Method	Drugs_NFL.m
Comment	MeOH		

Compound Table

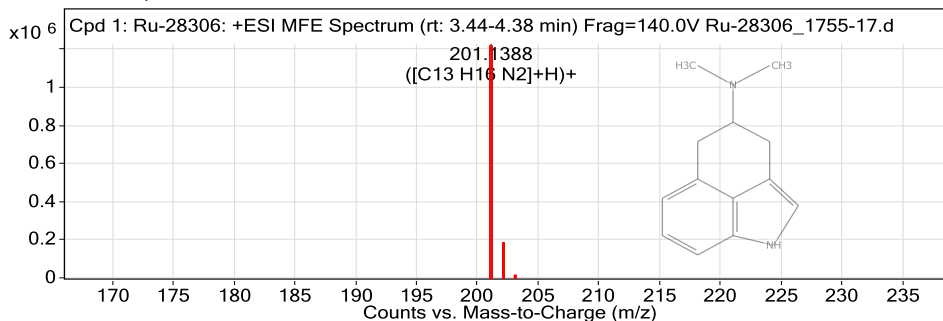
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 1: Ru-28306	Ru-28306	C13 H16 N2	3.5	200.1316

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
Ru-28306	201.1388	3.5	200.1316	3.5	C13 H16 N2	200.1314	-1.02

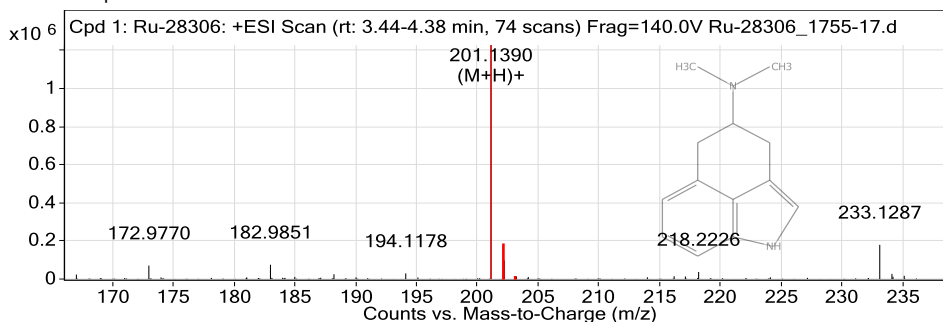
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

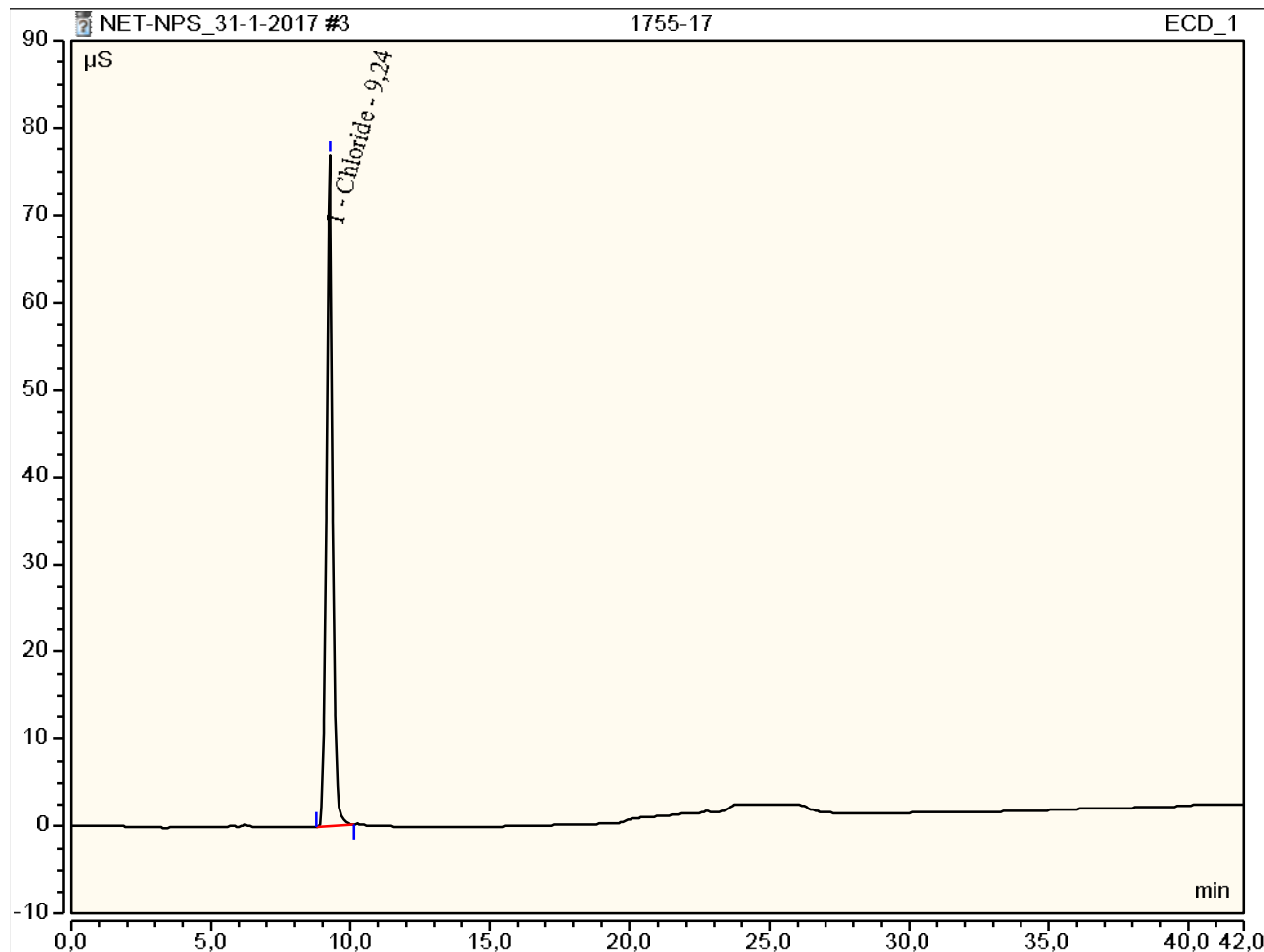
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
201.1388	1	1227783.88	C13 H16 N2	(M+H)+
202.142	1	168620.07	C13 H16 N2	(M+H)+
203.145	1	11354.55	C13 H16 N2	(M+H)+
204.1447	1	523.71	C13 H16 N2	(M+H)+

--- End Of Report ---

Peak Integration Report

Sample Name:	1755-17	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	31-jan-2017 / 11:46	Run Time:	42,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S} \cdot \text{min}$	Height μS	Amount mg/L
1,00	9,24	Chloride	BMB	19,89	76,74	n.a.
		TOTAL:		19,89	76,74	0,00





REPORT

Sample ID:	1755-17
Our notebook code:	P-1755-17
NMR sample preparation:	15 mg dissolved in 0.7 mL DMSO- <i>d</i> ₆
NMR experiments:	¹ H, ¹³ C, ¹ H- ¹ H <i>gs</i> -COSY, ¹ H- ¹³ C <i>gs</i> -HSQC, ¹ H- ¹³ C <i>gs</i> -HMBC, ¹ H- ¹⁵ N <i>gs</i> -HMBC.
Proposed structure:	
Chemical name:	<i>N,N</i> -dimethyl-1,3,4,5-tetrahydrobenzo[<i>cd</i>]indol-4-aminium cation
Comments:	- Structure elucidation based on 1D and 2D NMR spectra - Sample is mostly pure according to the NMR, there are just some minor signals of impurities observed in the upfield part of ¹H NMR spectrum.
Supporting information:	Copies of ¹ H and ¹³ C NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	February 22, 2017

P-1755-17

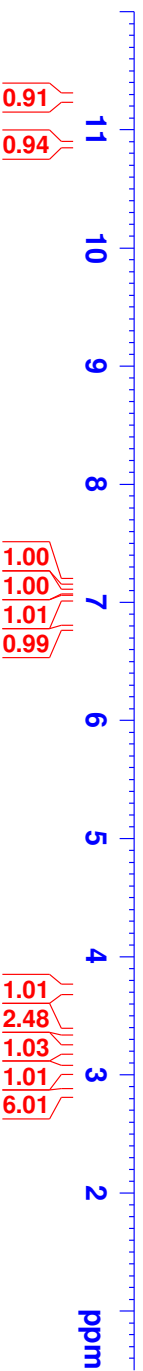


Current Data Parameters
NAME P-1755-17
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20170209
Time 21.27
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 101
DW 50.000 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

===== CHANNEL f1 =====
SFO1 500.1330885 MHz
NUC1 1H
P1 8.60 usec
PIW1 26.00000000 W

F2 - Processing parameters
SI 65536
SF 500.130039 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00





Current Data Parameters
NAME P-1755-17
EXPNO 3
PROCNO 1

F2 - Acquisition Parameters
Date_ 20170209
Time 23.25
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
ID 63536
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 298.0 K
D1 1.00000000 sec
D11 0.03000000 sec
TD0 1

===== CHANNEL f1 =====
SF01 125.7703637 MHz
NUC1 13C
P1 8.70 usec
PLW1 122.00000000 W

===== CHANNEL f2 =====
SF02 500.1320005 MHz
NUC2 1H
CPDPRGf2 waltz16
PCPD2 80.00 usec
PLW2 26.00000000 W
PLW12 0.30046001 W
PLW13 0.15113001 W

F2 - Processing parameters
SI 32768
SF 125.7579262 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40

