



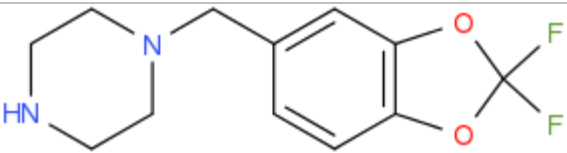
ANALYTICAL REPORT¹

DB-MDBP (C₁₂H₁₄F₂N₂O₂)

1-((2,2-DIFLUOROBENZO[D][1,3]DIOXOL-5-YL)METHYL)PIPERAZINE

Remark – other NPS detected: **none**

Sample ID:	1185-15
Sample description:	powder - white-off
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	6/5/2015
Date of entry (M/D/Y) into NFL database:	7/11/2015
Report 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	1-((2,2-DIFLUOROBENZO[D][1,3]DIOXOL-5-YL)METHYL)PIPERAZINE
Other names	4F-butrylfentanyl
Formula (per base form)	C ₁₂ H ₁₄ F ₂ N ₂ O ₂
M _w (g/mol)	256.25
Salt form	HCl
StdInChIKey	MPZINTHEMFDGTH-UHFFFAOYSA-N
Compound Class	Piperazine derivatives
Other NPS detected	none
Add.info (purity..)	pure by GC, HPLC-TOF, compound is not pure by NMR (see analytical report)

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)
11/04/2018	Compound class corrected

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⁻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 (40) to 550 amu.

IR (condensed 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

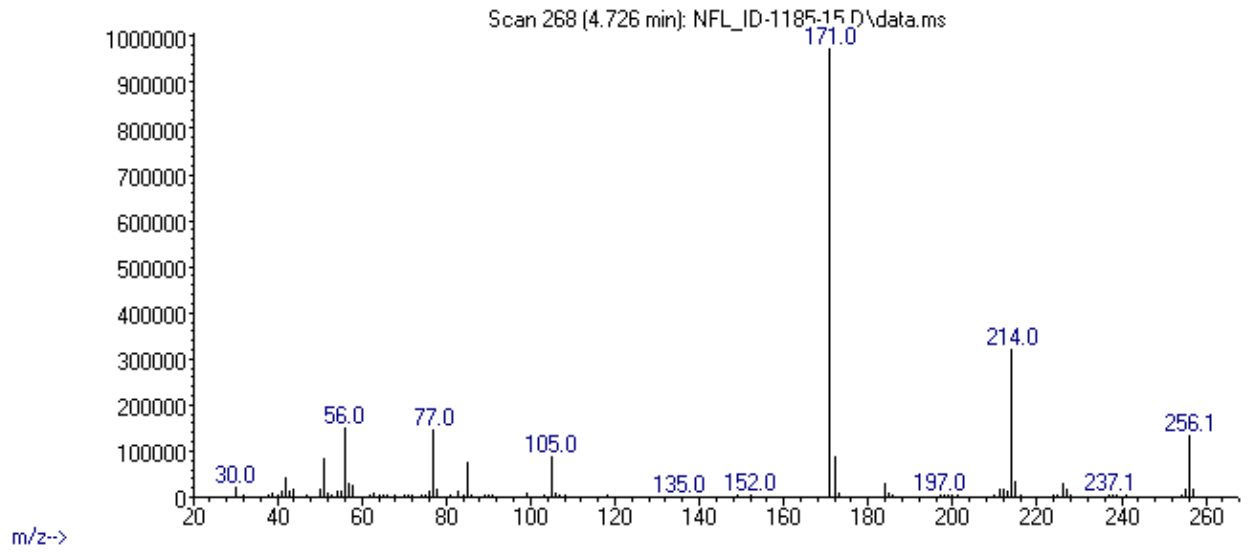
Solubility in	result/remark
CH ₂ Cl ₂	soluble
MeOH	soluble
H ₂ O	not tested soluble

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 4.73 BP(1): 171; BP(2): 214,BP(3) :256,
HPLC-TOF	+	Exact mass (theoretical): 256.1023; measured value Δppm:-1.13; formula:C12H14F2N2O2
FTIR-ATR	+	
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR (FKKT)	+	
validation		
other		

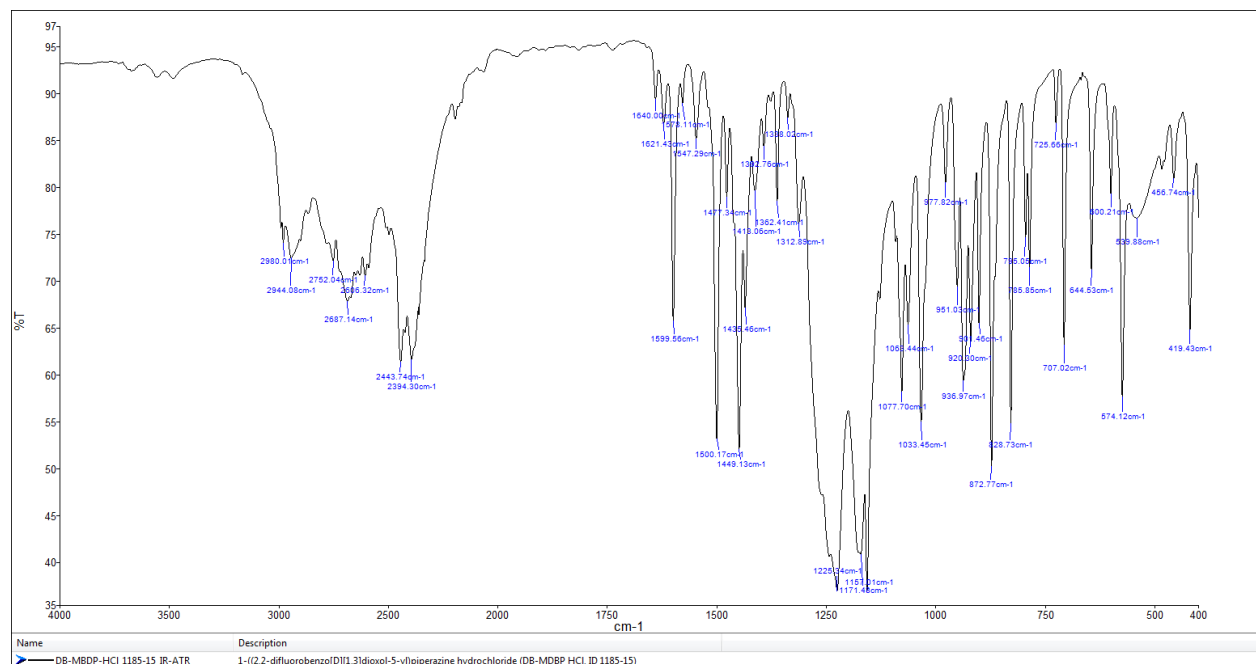
ANALYTICAL RESULTS

MS (EI)

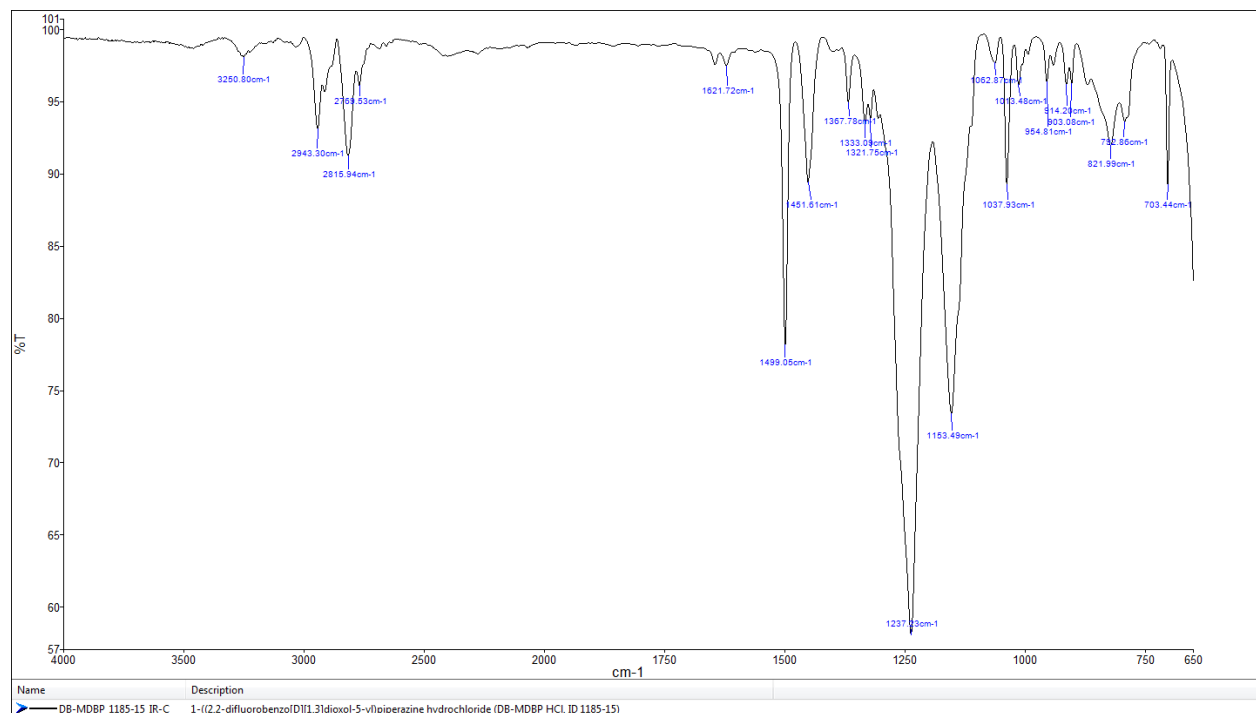
Abundance



FTIR-ATR - direct measurement



IR (condensed phase)



Target Compound Screening Report

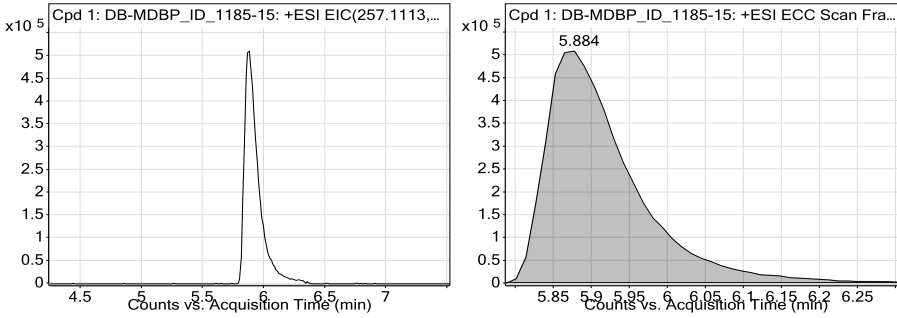
Data File	DB-MDBP_1185-15_TOF.d	Sample Name	DB-MDBP
Sample Type	Sample	Position	P1-E6
Instrument Name	SG13170002	User Name	
Acq Method	droge general-13-5-2015-XDB-C18-ESI-poz.m	Acquired Time	6/8/2015 10:55:49 AM
IRM Calibration Status	Success	DA Method	Droge_Default.m
Comment	extract in MeOH		

Compound Table

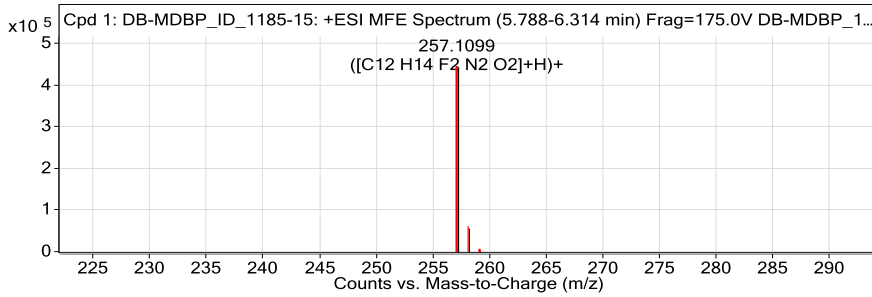
Label	Tgt Name	MFG Formula	Tgt Formula	Obs. RT	Obs. Mass
Cpd 1: DB-MDBP_ID_1185-15	DB-MDBP_ID_1185-15	C12 H14 F2 N2 O2	C12 H14 F2 N2 O2	5.884	256.1026

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error	Tgt Formula	Find Cps Alaorith Find by Molecula r Feature
DB-MDBP_ID_1185-15	257.1099	5.884	256.1026	5.884	C12 H14 F2 N2 O2	256.1023	-1.13	C12 H14 F2 N2 O2	

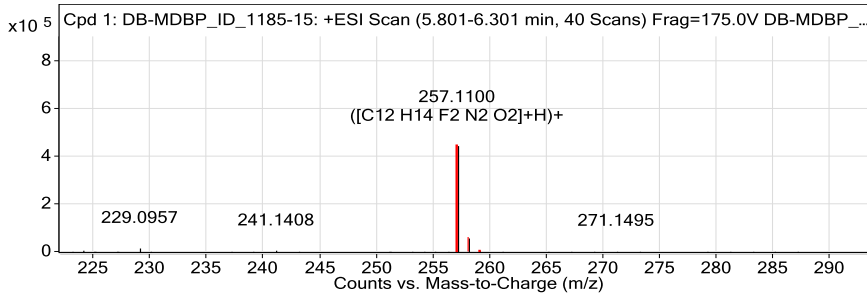
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



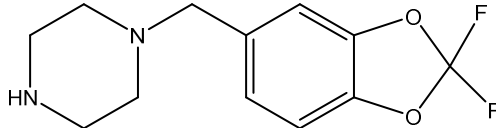
MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
257.1099	1	446082.75	C12 H14 F2 N2 O2	(M+H)+
258.1128	1	58936.41	C12 H14 F2 N2 O2	(M+H)+
259.1158	1	5765.24	C12 H14 F2 N2 O2	(M+H)+

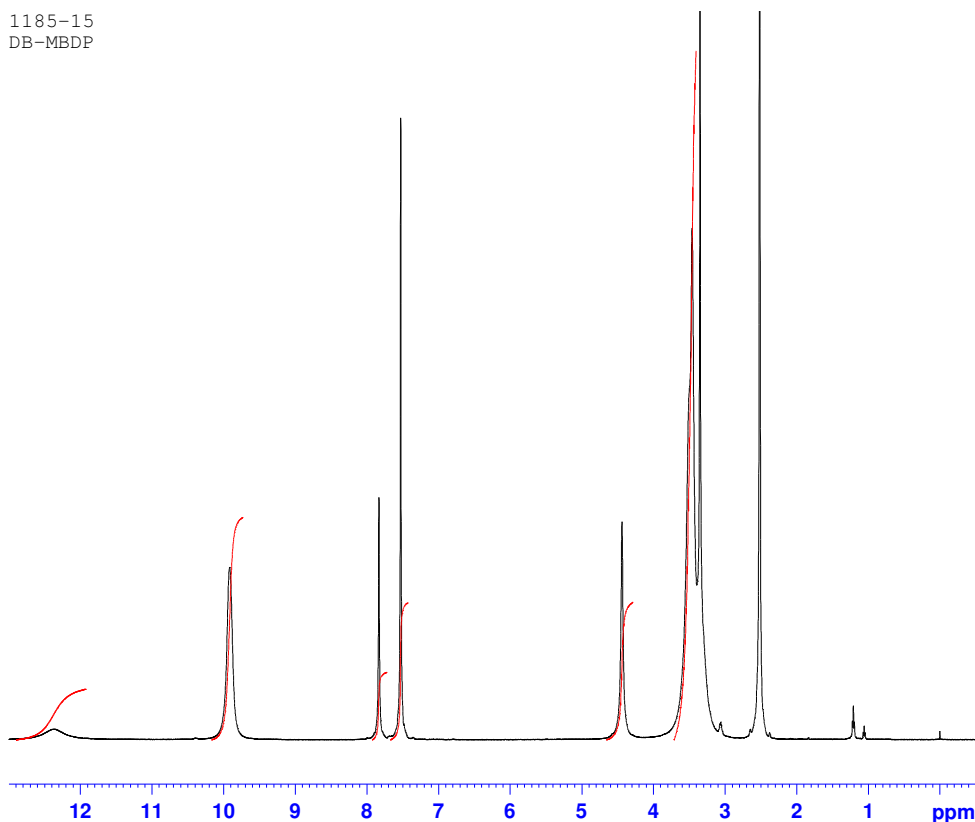
--- End Of Report ---



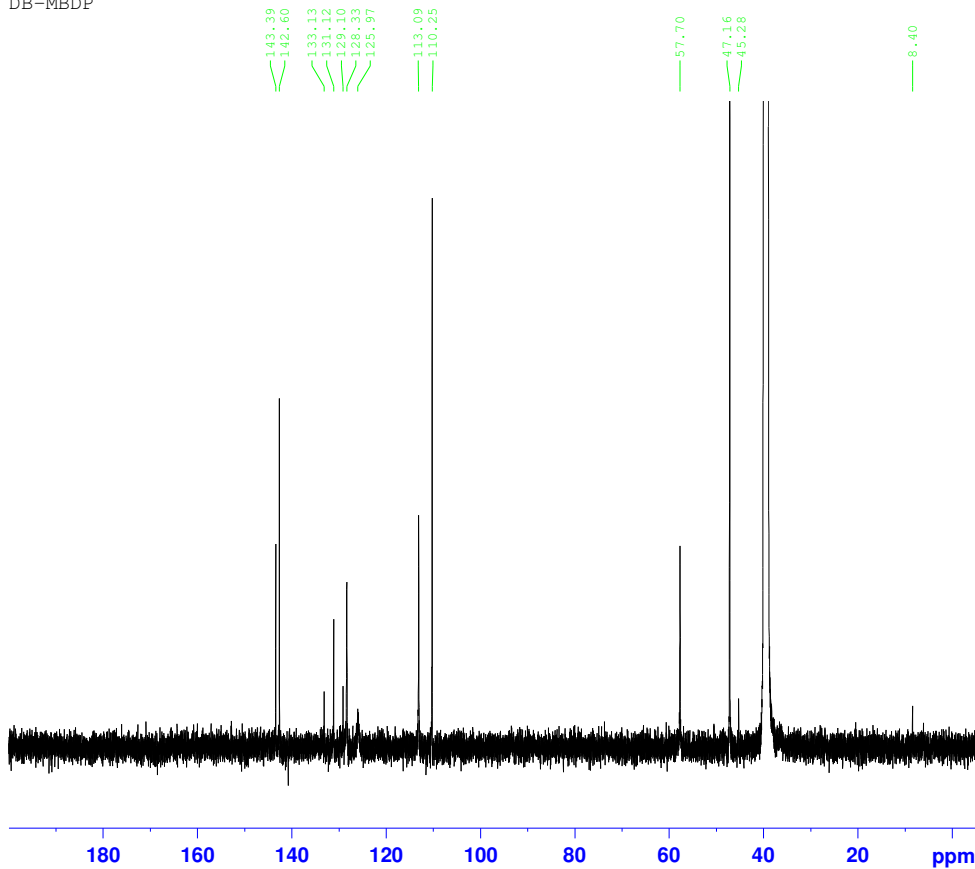
REPORT

Sample ID:	1185-15
Our notebook code:	P-1185-15
NMR sample preparation:	15 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.
Proposed structure:	
Chemical name:	1-((2,2-difluorobenzo[d][1,3]dioxol-5-yl)methyl)piperazine
Comments:	<ul style="list-style-type: none"> - Structure elucidation based on 1D and 2D NMR spectra - Compound is not pure by NMR, it contains organic impurities (evident in ^1H NMR signals around 1.0 ppm) and also possibly some other impurities (inorganic cations?) that cause line broadening in ^1H NMR spectrum, consequently significantly diminishing its use for structural determination. - According to NMR spectra this sample is identical to the sample P-1252-15.
Supporting information:	Copies of ^1H and ^{13}C NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	November 19, 2015

1185-15
DB-MBDP



1185-15
DB-MBDP



Current Data Parameters
NAME P-1185-15
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20150708
Time 14.38
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 10330.578 Hz
FIDRES 0.157632 Hz
AQ 3.1719923 sec
RG 80.6
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
SFO1 500.1330885 MHz

F2 - Processing parameters
SI 65536
SF 500.1299969 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



Current Data Parameters
NAME P-1185-15
EXPNO 5
PROCNO 1

F2 - Acquisition Parameters
Date_ 20150708
Time 15.59
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
SOLVENT DMSO
NS 1536
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
SFO1 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
SFO2 500.1320005 MHz

F2 - Processing parameters
SI 32768
SF 125.7578519 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40