

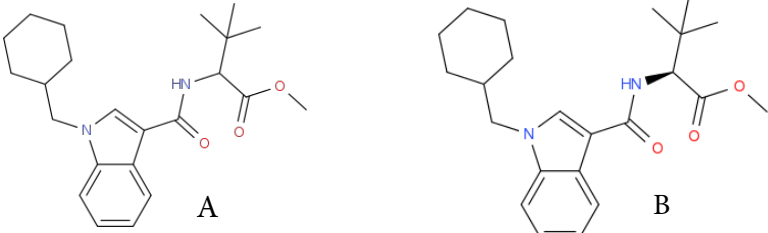
ANALYTICAL REPORT

MDMB-CHMICA (C23H32N2O3)

N-[[1-(cyclohexylmethyl)-1H-indol-3-yl]carbonyl]-3-methyl-valine, methyl ester

Remark – other NPS detected: **none**

Sample ID:	1190-15
Sample description:	powder - off white
Sample type:	test purchase /RESPONSE -purchasing
Comments ¹ :	
Date of entry into NFL database: http://www.policija.si/apps/nfl_response_web/seznam.php	10/14/2015

Substance identified-structure ² (base form)	
Systematic name	methyl 2-[[1-(cyclohexylmethyl)-1H-indol-3-yl]formamido]-3,3-dimethylbutanoate (stereochemistry was not confirmed experimentally in the RESPONSE project (structure A))
Other names	N-[[1-(cyclohexylmethyl)-1H-indol-3-yl]carbonyl]-3-methyl-valine, methyl ester; methyl (2S)-2-[[1-(cyclohexylmethyl)-1H-indol-3-yl]formamido]-3,3-dimethylbutanoate (structure B)
Formula (per base form)	C23H32N2O3
M _w (g/mol)	384,5
Salt form	
StdInChIKey	(A) SRJKCVHWIDFUBO-UHFFFAOYSA-N and SRJKCVHWIDFUBO-HXUWFJFHSA-N (B)
Compound Class	Cannabinoids
Other NPS detected	none
Add.info (purity..)	pure by GC-MS, HPLC-TOF, few % of unidentified organic 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)
18/05/17	Empirical formula corrected.
	Structure (A) where stereochemistry is not defined and corresponding StdInChI key were added.

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.

Supporting information

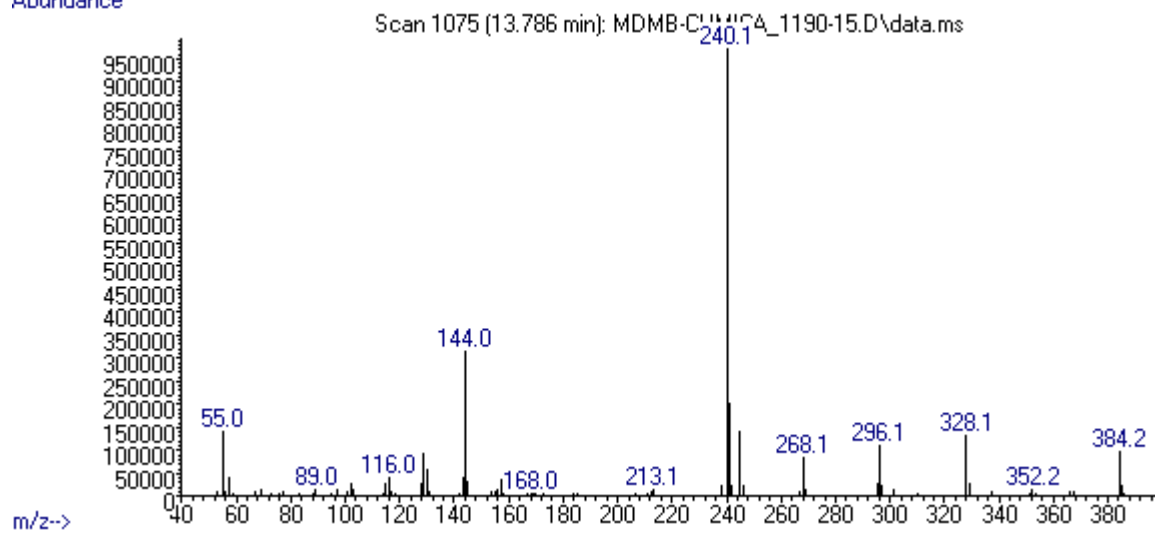
Solubility in	result/remark
CH ₂ Cl ₂	soluble
MeOH	soluble
water	low

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 13,79 BP(1): 240; BP(2): 144,BP(3) :241,
HPLC-TOF	+	Exact mass (theoretical): 384,2413; measured value Δppm:0,08; formula:C23H32N2O3
FTIR-ATR	+	direct measurement (note sample contains MeOH)
FTIR (condensed phase) always as base form	+	
NMR	+	
validation		
other		

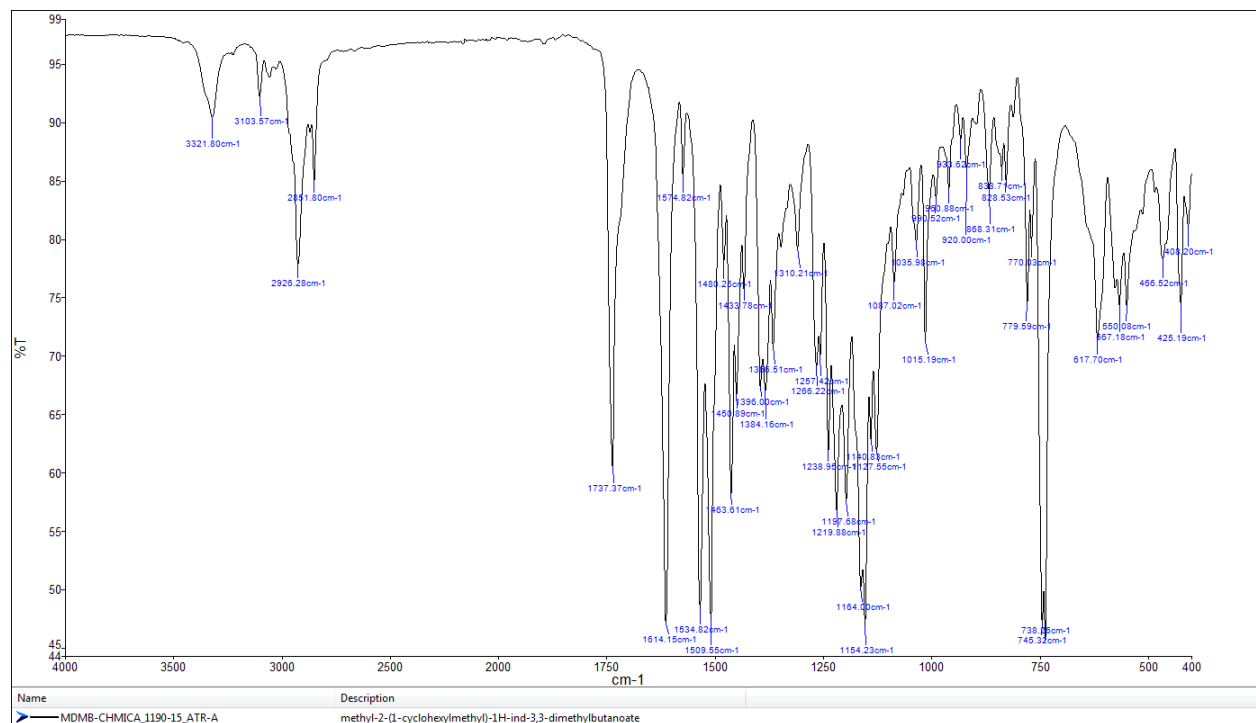
ANALYTICAL RESULTS

MS (EI)

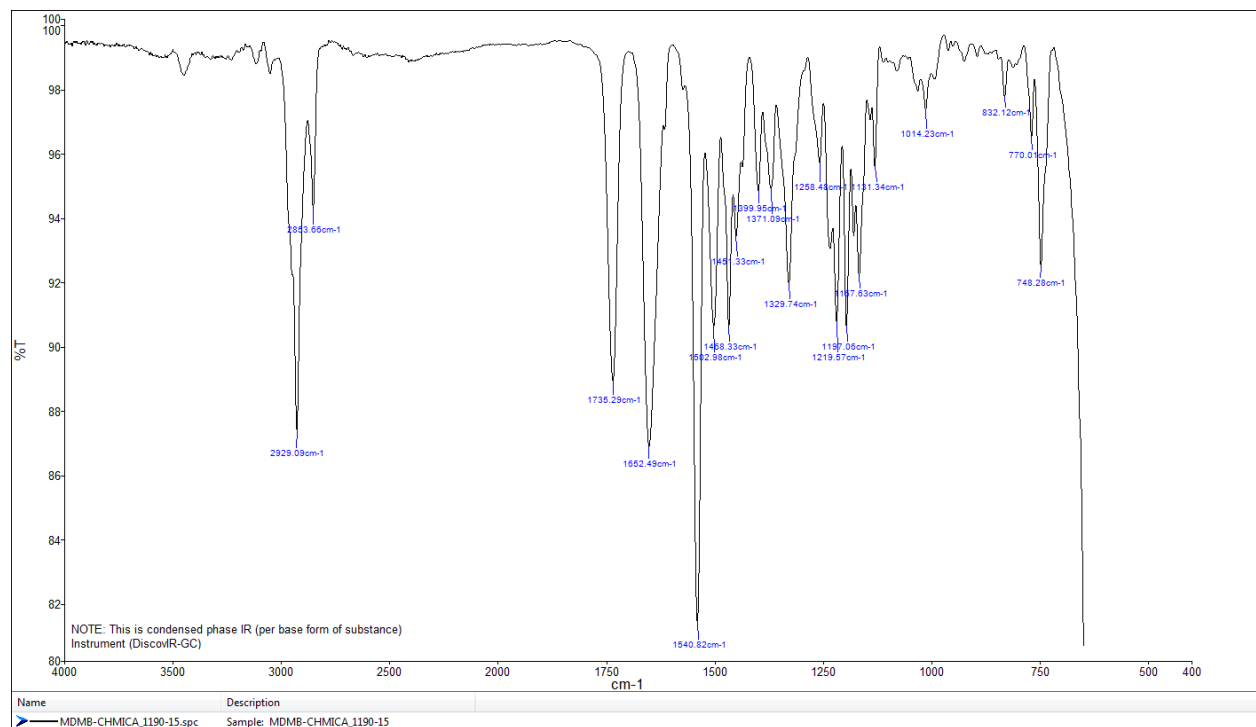
Abundance



FTIR-ATR - direct measurement (note: sample contains few % of organic impurities)



IR (condensed phase)



Target Compound Screening Report

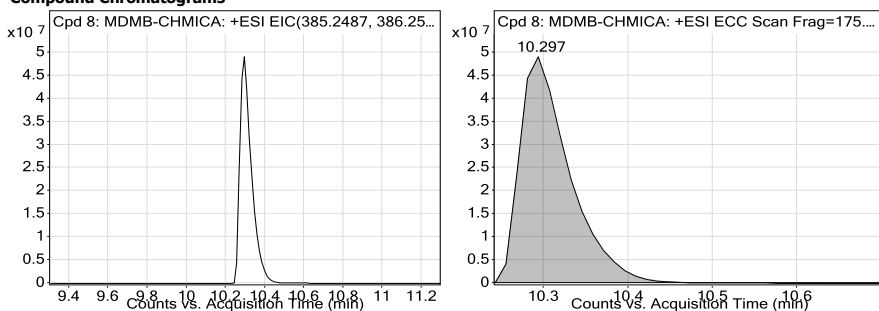
Data File	MDMB-CHMICA_1190-15_TOF.d	Sample Name	MDMB-CHMICA
Sample Type	Sample	Position	P1-C4
Instrument Name	SG13170002	User Name	
Acq Method	droge general-13-5-2015-XDB-C18-ESI-poz.m	Acquired Time	7/8/2015 11:08:06 AM
IRM Calibration Status	Success	DA Method	Droge_Default.m
Comment	extract in MeOH		

Compound Table

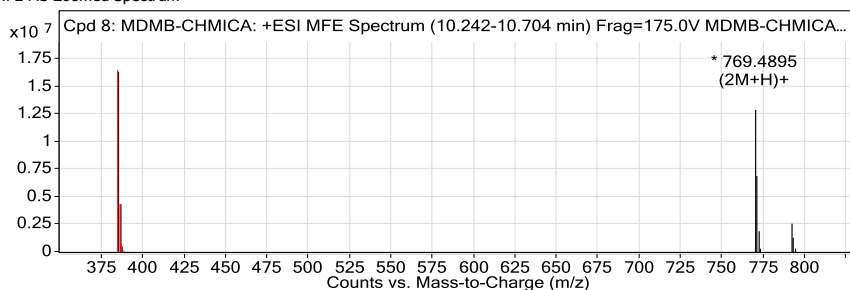
Label	Tgt Name	MFG Formula	Tgt Formula	Obs. RT	Obs. Mass
Cpd 8: MDMB-CHMICA	MDMB-CHMICA	C23 H32 N2 O3	C23 H32 N2 O3	10.297	384.2413

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error	Tgt Formula	Find Cps Algorith
MDMB-CHMICA	385.2485	10.297	384.2413	10.294	C23 H32 N2 O3	384.2413	0.08	C23 H32 N2 O3	Find by Molecular Feature

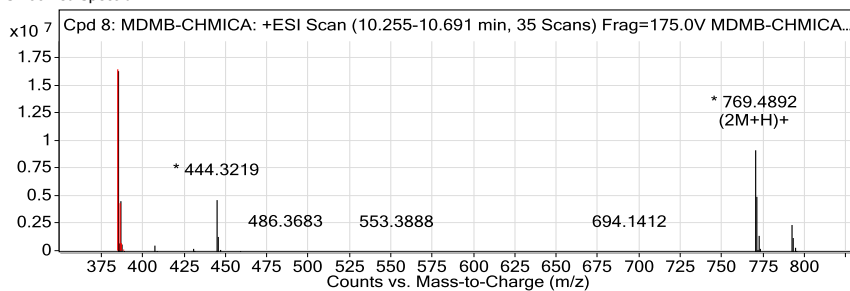
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



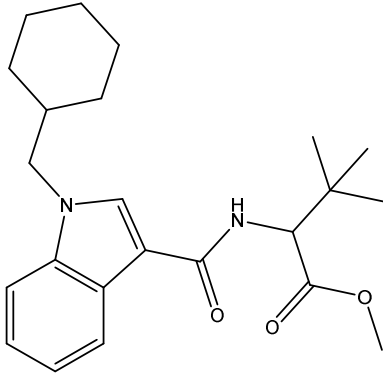
MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
385.2485	1	16352723	C23 H32 N2 O3	(M+H)+
386.2518	1	4397796.08	C23 H32 N2 O3	(M+H)+
387.2552	1	590485.43	C23 H32 N2 O3	(M+H)+
769.4895	1	12944640		(2M+H)+
770.4928	1	6915544.8		(2M+H)+
771.496	1	1947154.22		(2M+H)+
772.4996	1	358547.83		(2M+H)+
791.4715	1	2652121.5		(2M+Na)+
792.4754	1	1389992.18		(2M+Na)+
793.4784	1	372498.4		(2M+Na)+

--- End Of Report ---



REPORT

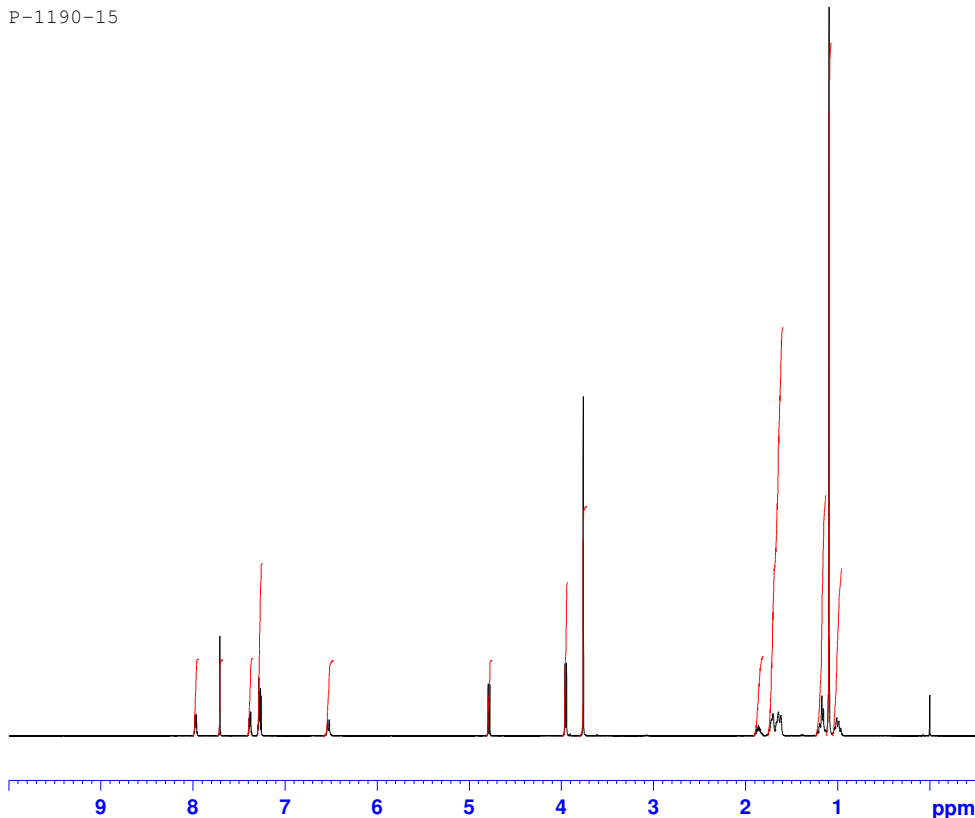
Sample ID:	1190-15
Our notebook code:	P-1190-15
NMR sample preparation:	15 mg dissolved in 0.7 mL CDCl ₃
NMR experiments:	¹ H, ¹³ C, ¹ H- ¹ H <i>gs</i> -COSY, ¹ H- ¹³ C <i>gs</i> -HSQC, ¹ H- ¹³ C <i>gs</i> -HMBC, ¹ H- ¹⁵ N <i>gs</i> -HMBC.
Proposed structure with chemical name:	 <p>methyl 2-(1-(cyclohexylmethyl)-1<i>H</i>-indole-3-carboxamido)-3,3-dimethylbutanoate</p>
Comments:	<ul style="list-style-type: none"> - Structure elucidation based on 1D and 2D NMR spectra - Compound is pure by NMR, containing a small amount (a few %) of unidentified organic compound(s).
Supporting information:	Copies of ¹ H and ¹³ C NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	October 6, 2015

P-1190-15



Current Data Parameters
 NAME P-1190-15
 EXPNO 1
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20150811
 Time 0.40
 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 64
 DW 48.400 usec
 DE 6.50 usec
 TE 298.5 K
 D1 1.00000000 sec



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

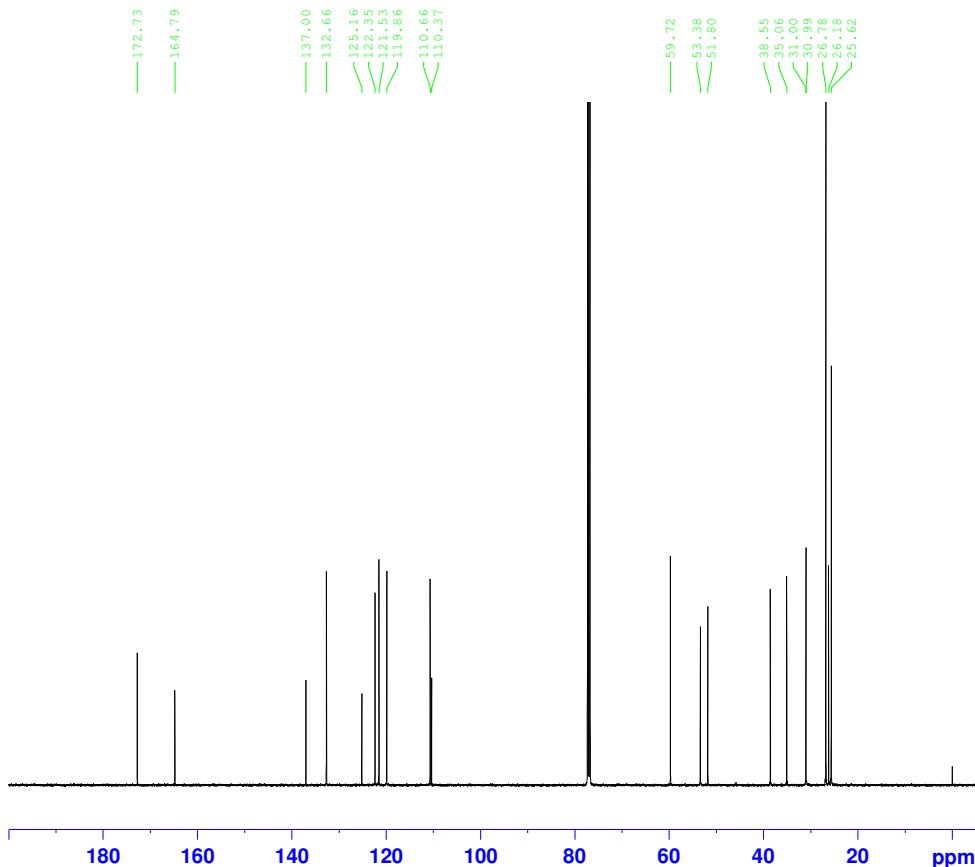
F2 - Processing parameters
 SI 65536
 SF 500.1300135 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00

P-1190-15



Current Data Parameters
 NAME P-1190-15
 EXPNO 4
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20150811
 Time 4.07
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zgpg30
 TD 65536
 SOLVENT CDCl3
 NS 5120
 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 299.1 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 125.7577897 MHz
 WDW EM
 SSB 0
 LB 1.00 Hz
 GB 0
 PC 1.40