



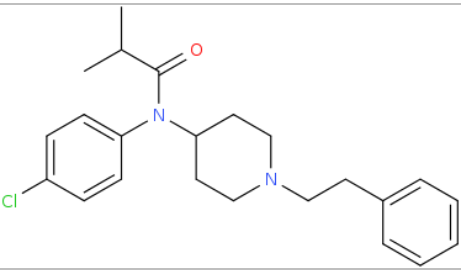
## ANALYTICAL REPORT<sup>1</sup>

### 4-Cl-iBF (C<sub>23</sub>H<sub>29</sub>ClN<sub>2</sub>O)

#### N-(4-chlorophenyl)-N-(1-phenethylpiperidin-4-yl)isobutyramide

Remark – other NPS detected: **none**

Sample ID:	1566-16
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	5/10/2016
Date of entry (M/D/Y) into NFL database:	7/29/2019
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	N-(4-chlorophenyl)-N-(1-phenethylpiperidin-4-yl)isobutyramide
Other names	4-Cl-iBF
Formula (per base form)	C <sub>23</sub> H <sub>29</sub> ClN <sub>2</sub> O
M <sub>w</sub> (g/mol)	384,95
Salt form/anions detected	HCl
StdInChIKey	YWHLYGSHOQKJG-UHFFFAOYSA-N
Compound Class	Opioids
Other NPS detected	none
Add.info (purity..)	not pure by GC-MS, TOF: few % C <sub>19</sub> H <sub>23</sub> ClN <sub>2</sub> Mw=314,16 minor impurities by NMR

<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 (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 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 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 (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 (solid) 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

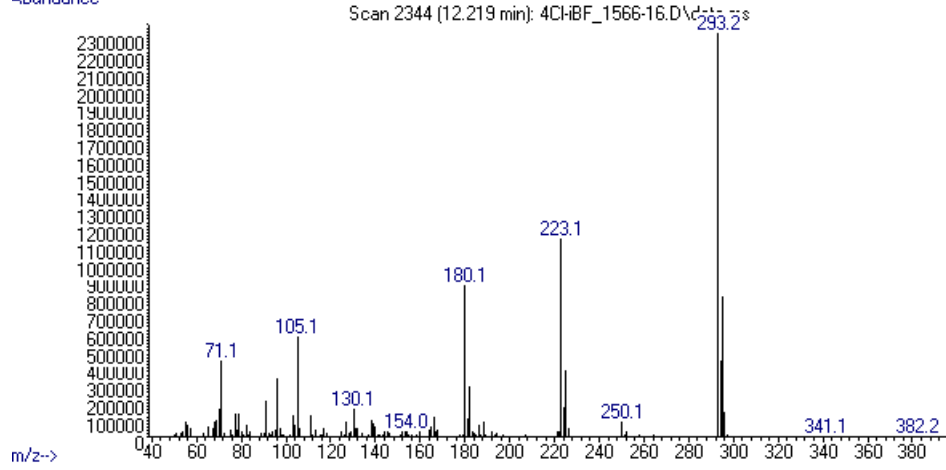
Solubility in	result/remark
CH <sub>2</sub> Cl <sub>2</sub>	partially
MeOH	soluble
H <sub>2</sub> O	partially

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 12,22 BP(1): 293; BP(2): 223,BP(3) :180,
HPLC-TOF	+	Exact mass (theoretical): 384,1968; measured value Δppm:0,03; formula:C <sub>23</sub> H <sub>29</sub> ClN <sub>2</sub> O
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR (in FKKT)	+	
validation		
other		

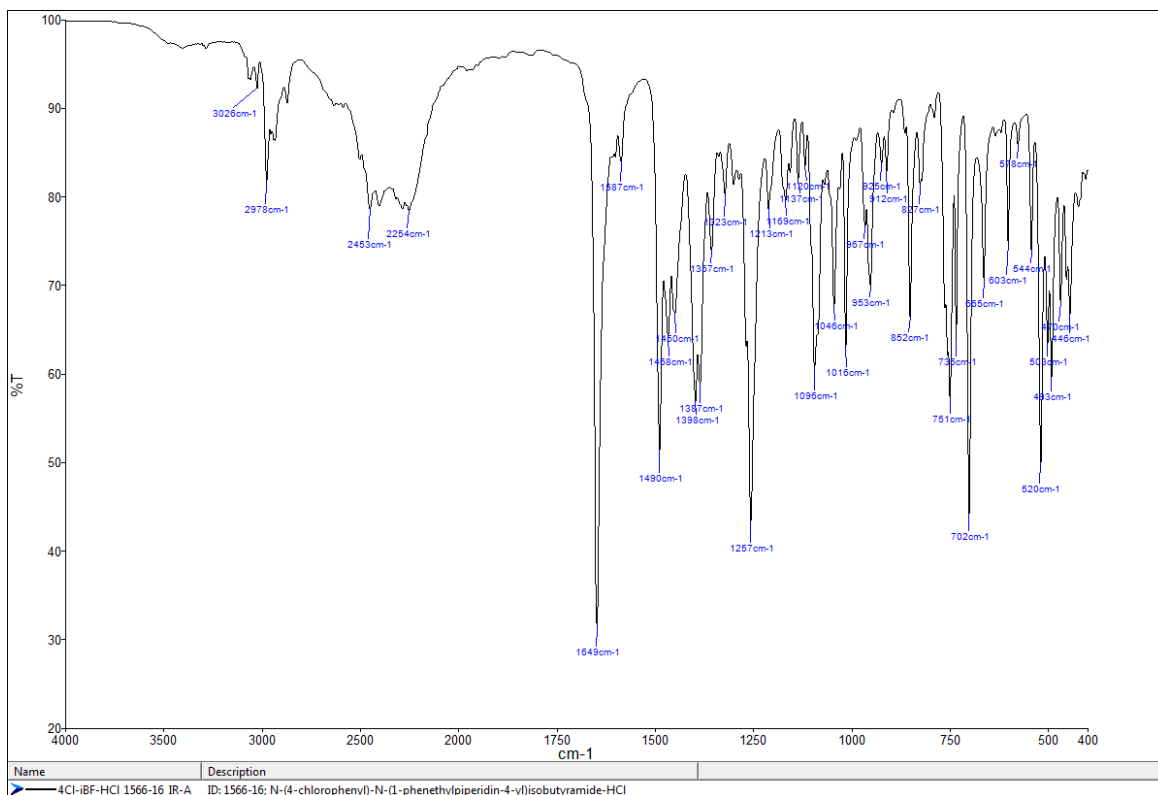
# ANALYTICAL RESULTS

MS (EI)

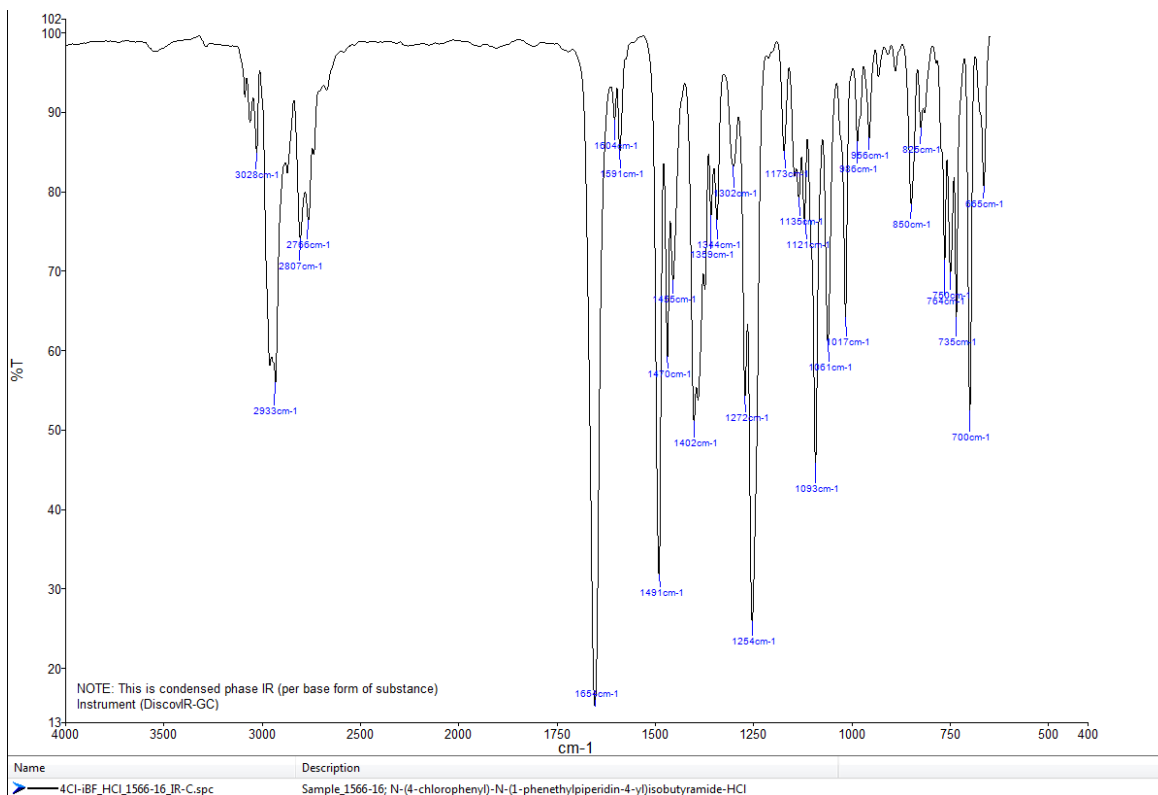
Abundance



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



### IR (condensed phase – after chromatographic separation)



# TOF REPORT

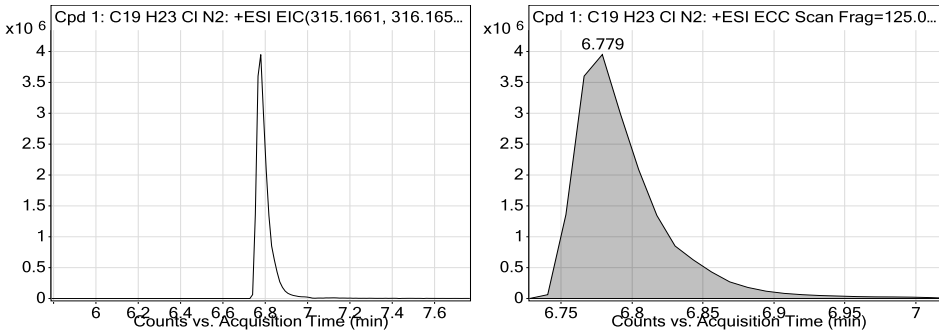
<b>Data File</b>	4Cl-iBF_1566-16_TOF.d	<b>Sample Name</b>	ID_1566-16
<b>Sample Type</b>	Sample	<b>Position</b>	P1-C3
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	general-1512015-XDB-C18-ESI-poz-pod.m	<b>Acquired Time</b>	5/18/2016 10:46:59 AM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Drugs_NFL.m
<b>Comment</b>	extract in MeOH		

## Compound Table

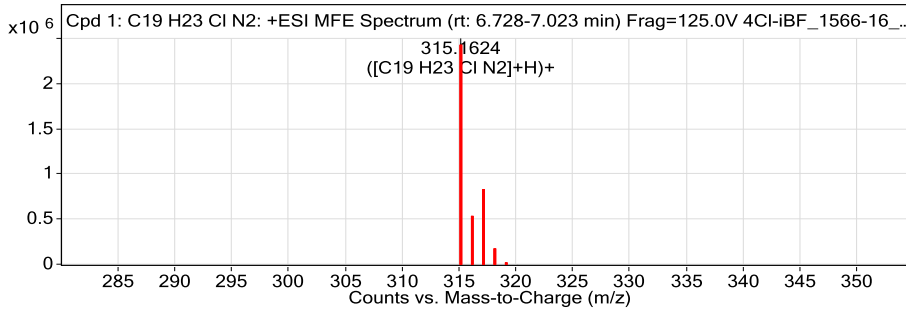
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 1: C19 H23 Cl N2		C19 H23 Cl N2	6.779	314.1553
Cpd 2: C23H29ClN2O (received as 4Cl-iBF)	C23H29ClN2O (received as 4Cl-iBF)	C23 H29 Cl N2 O	7.052	384.1968

Obs. m/z	Obs. RT	Obs. Mass
315.1624	6.779	314.1553

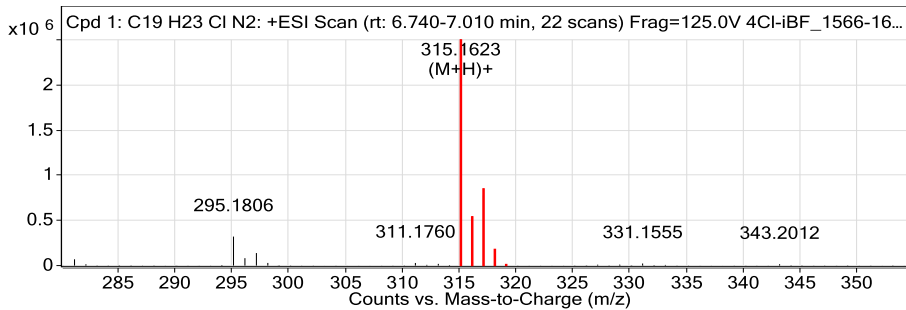
## Compound Chromatograms



## MFE MS Zoomed Spectrum



## MS Zoomed Spectrum



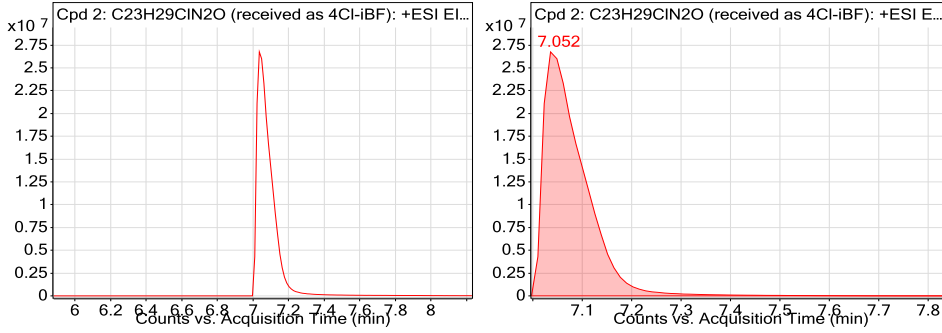
## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
315.1624	1	2500553.25	C19 H23 Cl N2	(M+H)+
316.1662	1	506972.66	C19 H23 Cl N2	(M+H)+
317.1607	1	775647.01	C19 H23 Cl N2	(M+H)+
318.1631	1	161665.08	C19 H23 Cl N2	(M+H)+
319.1655	1	16244.34	C19 H23 Cl N2	(M+H)+
320.1703	1	965.79	C19 H23 Cl N2	(M+H)+

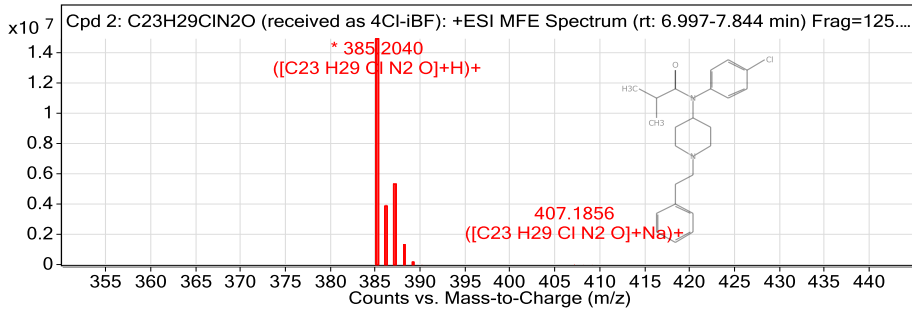
Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
C23H29ClN2O (received as 4Cl-iBF)	385.204	7.052	384.1968	7.05	C23 H29 Cl N2 O	384.1968	0.03

## Compound Chromatograms

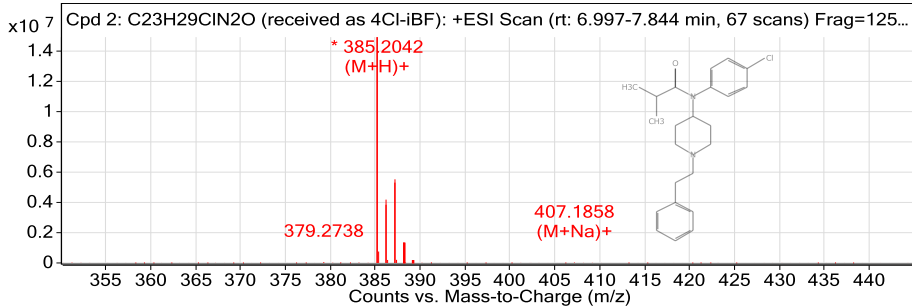
# TOF REPORT



## MFE MS Zoomed Spectrum



## MS Zoomed Spectrum



## MS Spectrum Peak List

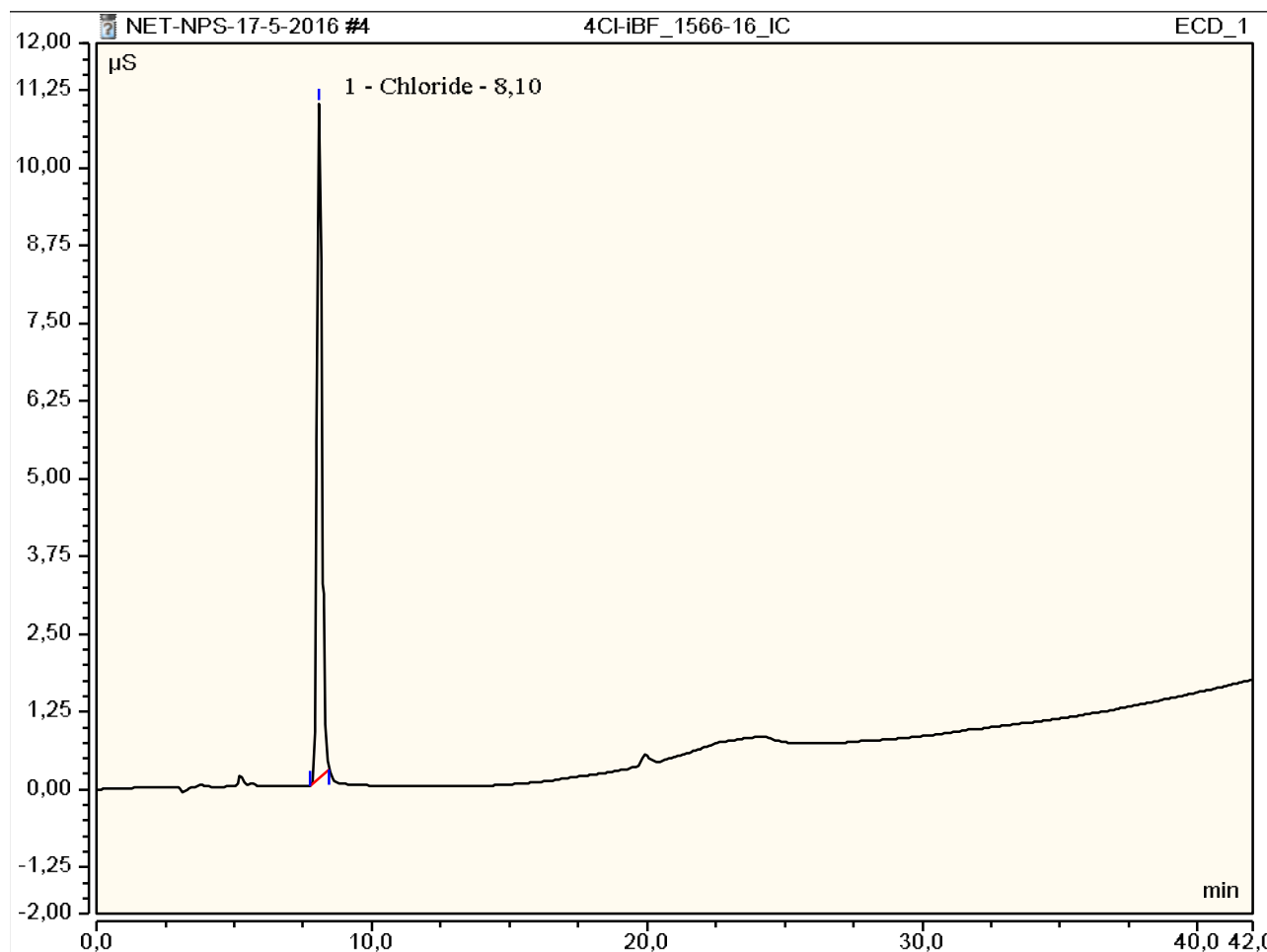
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
385.204	1	14964119	C23 H29 Cl N2 O	(M+H)+
386.2074	1	3961208.19	C23 H29 Cl N2 O	(M+H)+
387.2021	1	5285509.74	C23 H29 Cl N2 O	(M+H)+
388.2052	1	1222368.55	C23 H29 Cl N2 O	(M+H)+
389.208	1	154050.47	C23 H29 Cl N2 O	(M+H)+
390.21	1	12914.58	C23 H29 Cl N2 O	(M+H)+
407.1856	1	20608.94	C23 H29 Cl N2 O	(M+Na)+
408.1889	1	5252.37	C23 H29 Cl N2 O	(M+Na)+
409.1835	1	7035.13	C23 H29 Cl N2 O	(M+Na)+
410.1866	1	1749.34	C23 H29 Cl N2 O	(M+Na)+

--- End Of Report ---

### Peak Integration Report

Sample Name:	4Cl-iBF_1566-16_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	17-maj-2016 / 14:21	Run Time:	42,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}\cdot\text{min}$	Height $\mu\text{S}$	Amount mg/L
1,00	8,10	Chloride	BMB	2,29	10,84	n.a.
TOTAL:				2,29	10,84	0,00







## REPORT

Sample ID:	<b>1566-16</b>
Our notebook code:	P-1566-16
NMR sample preparation:	15 mg dissolved in 0.7 mL DMSO- $d_6$
NMR experiments:	$^1\text{H}$ , $^{13}\text{C}$ , $^1\text{H}$ - $^1\text{H}$ <i>gs</i> -COSY, $^1\text{H}$ - $^{13}\text{C}$ <i>gs</i> -HSQC, $^1\text{H}$ - $^{13}\text{C}$ <i>gs</i> -HMBC, $^1\text{H}$ - $^{15}\text{N}$ <i>gs</i> -HMBC.
Proposed structure:	
Chemical name:	4-( <i>N</i> -(4-chlorophenyl)isobutyramido)-1-phenethylpiperidin-1-ium cation
Comments:	- Structure elucidation based on 1D and 2D NMR spectra - Sample contains some minor impurities as evident by NMR.
Supporting information:	Copies of $^1\text{H}$ and $^{13}\text{C}$ NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	July 27, 2016

P-1566-16

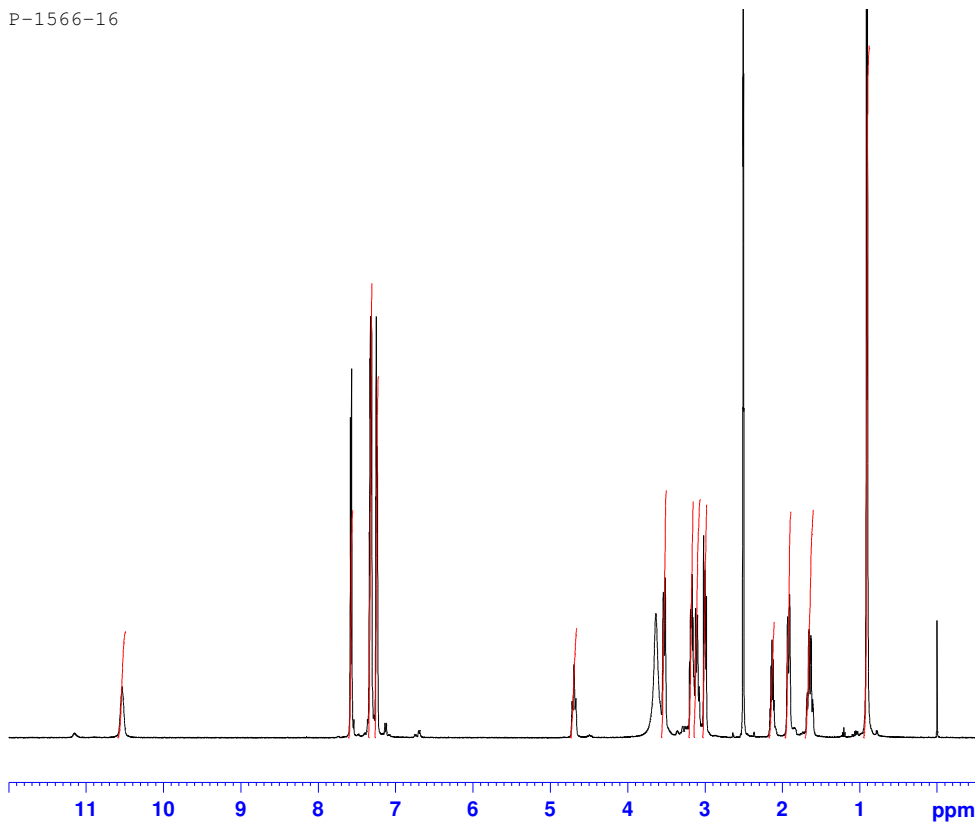


Current Data Parameters  
NAME P-1566-16  
EXPNO 1  
PROCNO 1

F2 - Acquisition Parameters  
Date\_ 20160531  
Time 0.22  
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 203  
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 500.1300012 MHz  
WDW EM  
SSB 0  
LB 0.30 Hz  
GB 0  
PC 1.00



P-1566-16



Current Data Parameters  
NAME P-1566-16  
EXPNO 3  
PROCNO 1

F2 - Acquisition Parameters  
Date\_ 20160705  
Time 21.14  
INSTRUM spect  
PROBHD 5 mm PABBO BB-  
PULPROG zgpg30  
TD 65536  
SOLVENT DMSO  
NS 3072  
DS 4  
SWH 29761.904 Hz  
FIDRES 0.484131 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 125.7577885 MHz  
WDW EM  
SSB 0  
LB 1.00 Hz  
GB 0  
PC 1.40

