



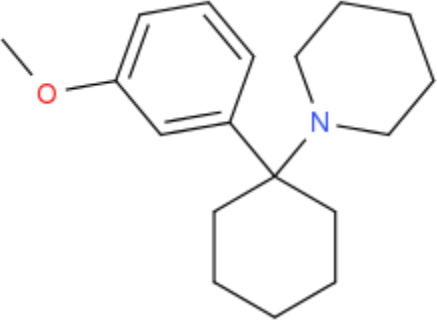
## ANALYTICAL REPORT<sup>1</sup>

### 3-MeO-PCP (C<sub>18</sub>H<sub>27</sub>NO)

#### 1-[1-(3-methoxyphenyl)cyclohexyl]-piperidine

Remark – other NPS detected: **none**

Sample ID:	1262-15
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	9/2/2015
Date of entry (M/D/Y) into NFL database:	9/2/2015
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 (in SFC)- structure <sup>2</sup> (base form)	
Systematic name	1-[1-(3-methoxyphenyl)cyclohexyl]-piperidine
Other names	
Formula (per base form)	C <sub>18</sub> H <sub>27</sub> NO
M <sub>w</sub> (g/mol)	273,42
Salt form	HCl
StdInChIKey	BQQSZHHKGPOXLN-UHFFFAOYSA-N
Compound Class	Arylcyclohexylamines
Other NPS detected	none
Add.info (purity..)	traces of organic impurities by GC. HPLC and 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 (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 (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 (40) to 550 amu.

IR (condensed 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

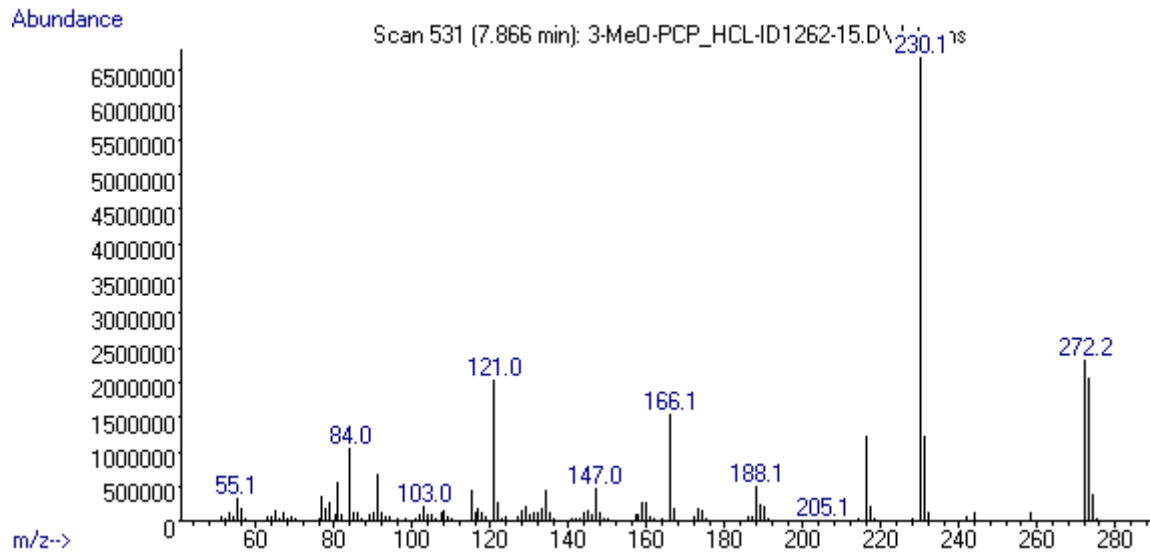
## Supporting information

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

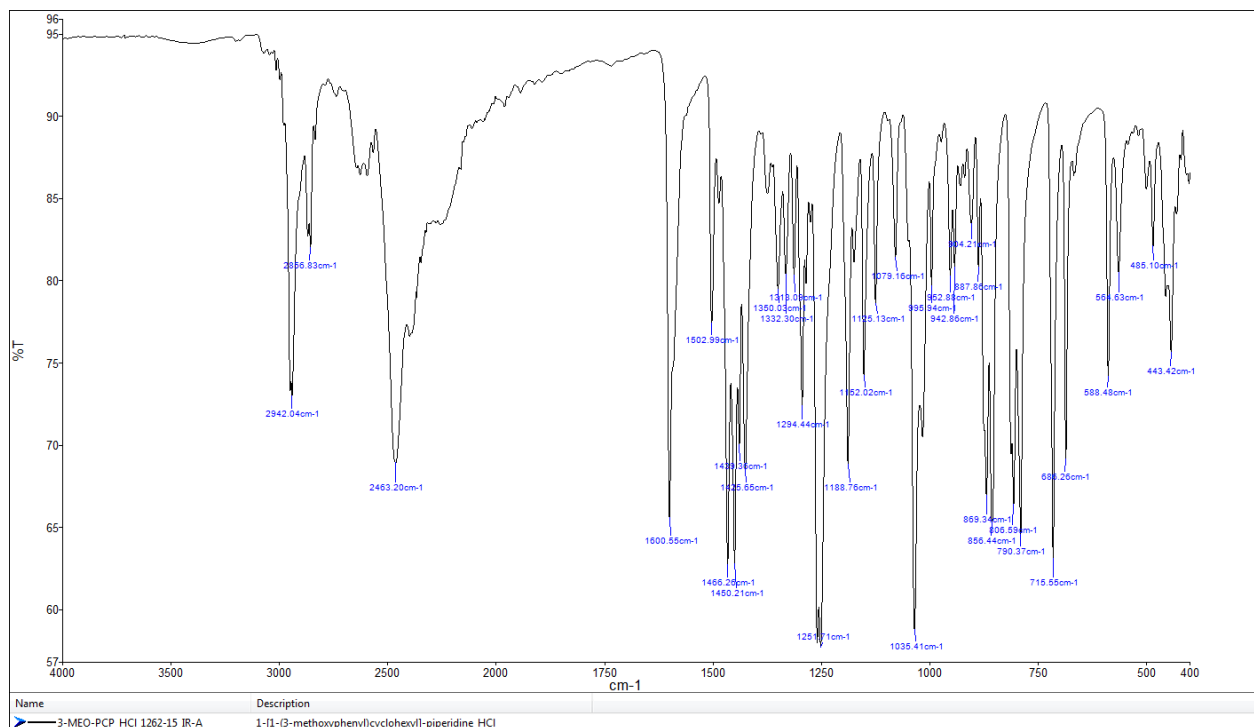
Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 7,87 BP(1): 230; BP(2): 272,BP(3) :273,
HPLC-TOF	+	Exact mass (theoretical): 273,2093; measured value Δppm:-1,15; formula:C18H27NO
FTIR-ATR	+	direct measurement
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR	+	
validation		MS consistent with the one in 2015 ENFSI library(qM =99)
other		IR condensed phase in good agreement with the one reported in EMCDDAs EDND database by German NFP (assessment was based on a visual comparison of spectra)

# ANALYTICAL RESULTS

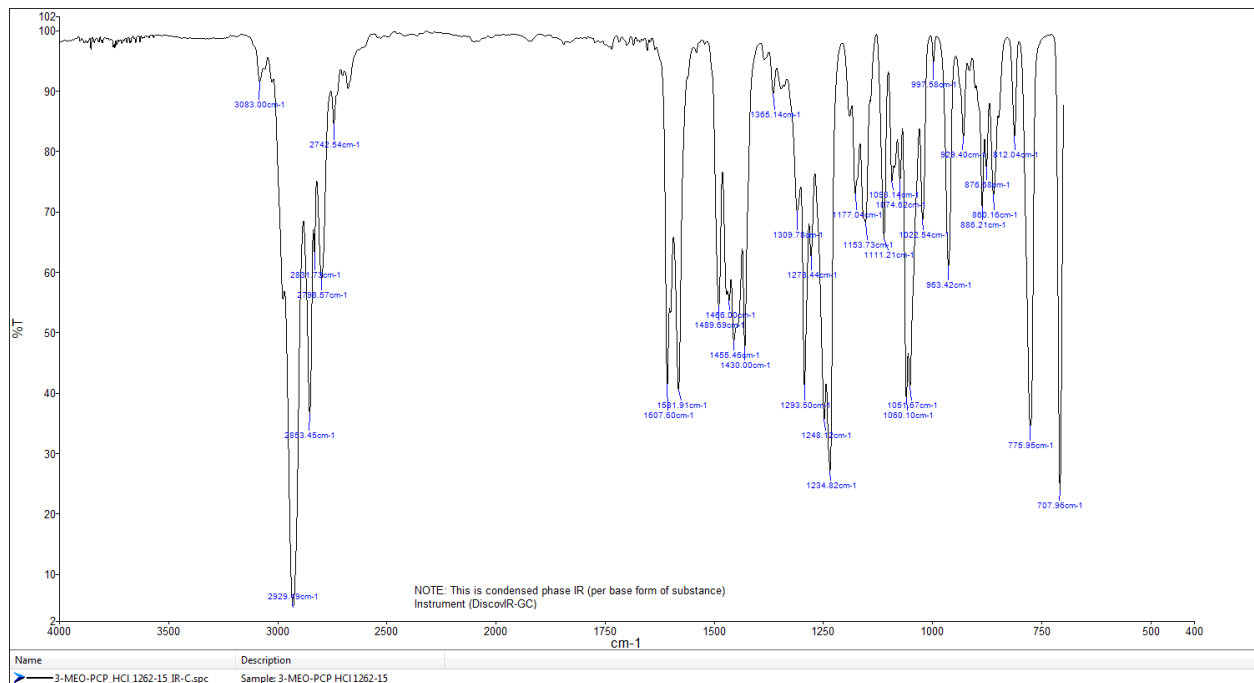
MS (EI)



## FTIR-ATR - direct measurement



## IR (condensed phase)



# Target Compound Screening Report

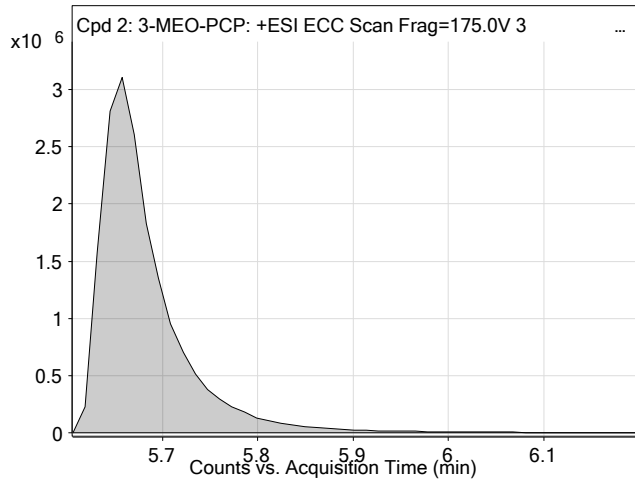
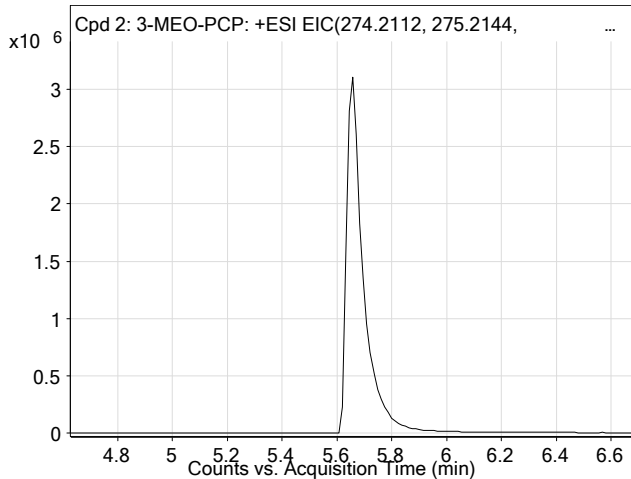
<b>Data File</b>	3-MEO-PCP_1262-15_TOF.d	<b>Sample Name</b>	1262-15
<b>Sample Type</b>	Sample	<b>Position</b>	P1-B4
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	droge general-13-5-2015-XDB-C18-ESI-poz.m	<b>Acquired Time</b>	9/2/2015 2:54:35 PM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Droge_Default.m
<b>Comment</b>	extract in MeOH		

## Compound Table

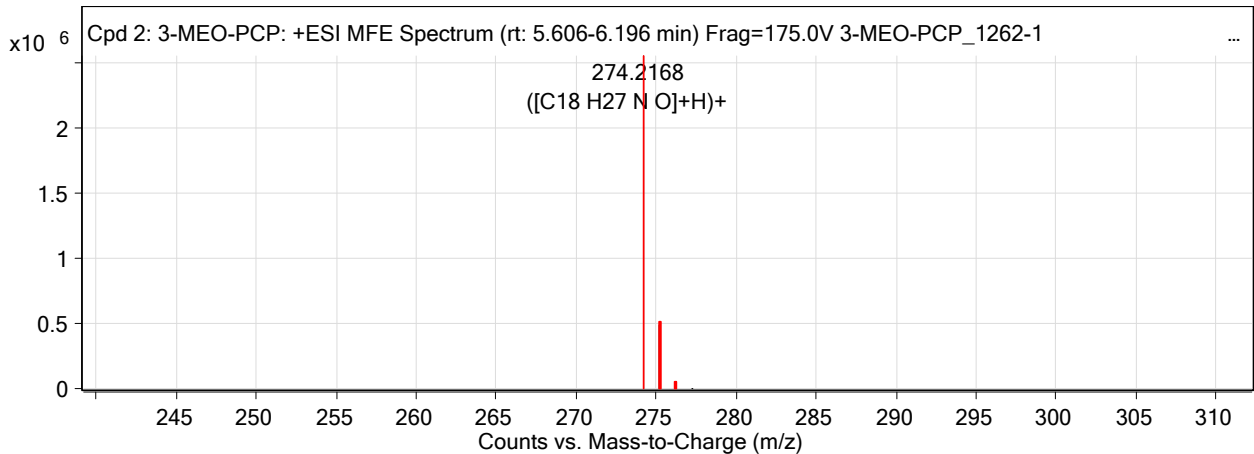
Label	Tgt Name	Obs. RT	Obs. Mass
Cpd 2: 3-MEO-PCP	3-MEO-PCP	5.659	273.2096

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)	Find Cpds Algorithm
3-MEO-PCP	274.2168	5.659	273.2096	5.659	C18 H27 N O	273.2093	-1.15	Find by Molecular Feature

## Compound Chromatograms



## MFE MS Zoomed Spectrum

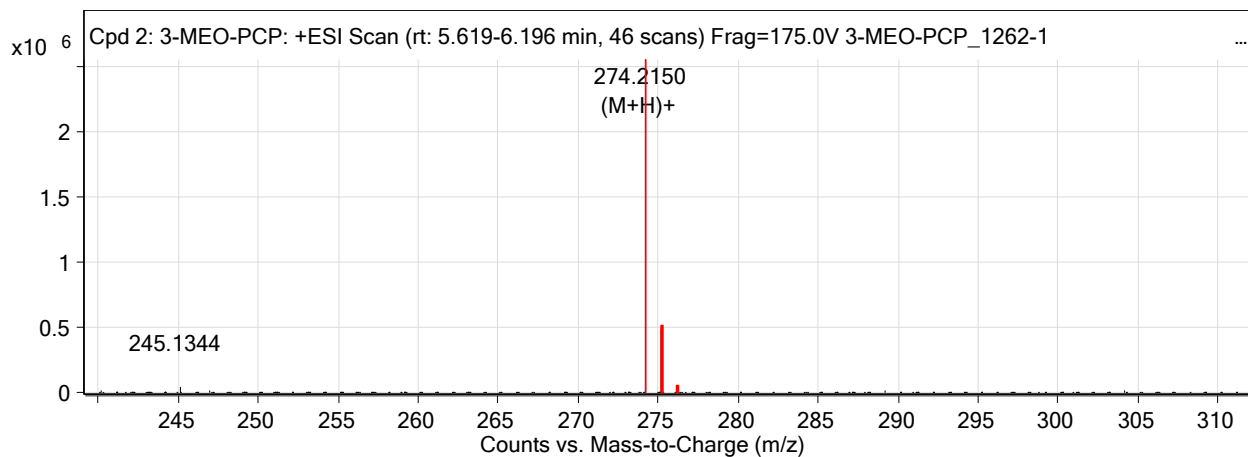


## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
274.2168	1	2552933.25	C18 H27 N O	(M+H)+
275.2204	1	490776.64	C18 H27 N O	(M+H)+
276.2232	1	47931.34	C18 H27 N O	(M+H)+
277.2257	1	3798.46	C18 H27 N O	(M+H)+

## MS Zoomed Spectrum

# Target Compound Screening Report

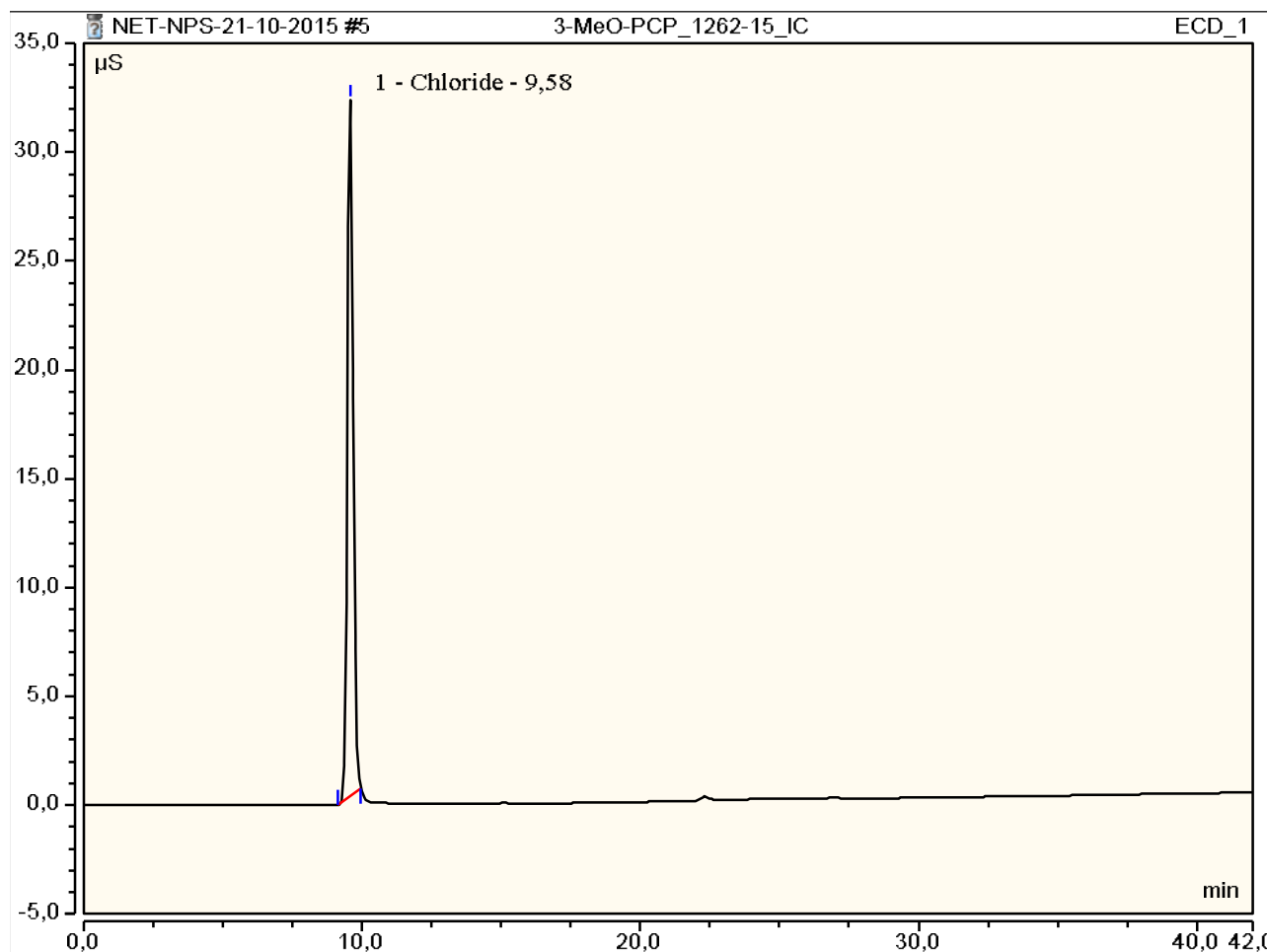


--- End Of Report ---

### Peak Integration Report

Sample Name:	3-MeO-PCP_1262-15_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	21-okt-2015 / 17:27	Run Time:	42,00

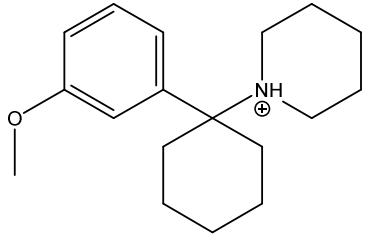
No.	Time min	Peak Name	Peak Type	Area $\mu\text{S} \cdot \text{min}$	Height $\mu\text{S}$	Amount mg/L
1,00	9,58	Chloride	BMB	7,73	32,02	n.a.
TOTAL:				7,73	32,02	0,00







## REPORT

Sample ID:	<b>1262-15</b>
Our notebook code:	P-1262-15
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.
Proposed structure:	
Chemical name:	1-(1-(3-methoxyphenyl)cyclohexyl)piperidin-1-ium
Comments:	- Structure elucidation based on 1D and 2D NMR spectra - Compound is not pure by NMR; possibly containing a few % of unknown organic impurities (signals in $^{13}\text{C}$ NMR at 43.5, 41.3, 26.4 and 22.1 ppm).
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:	November 6, 2015

P-1262-15

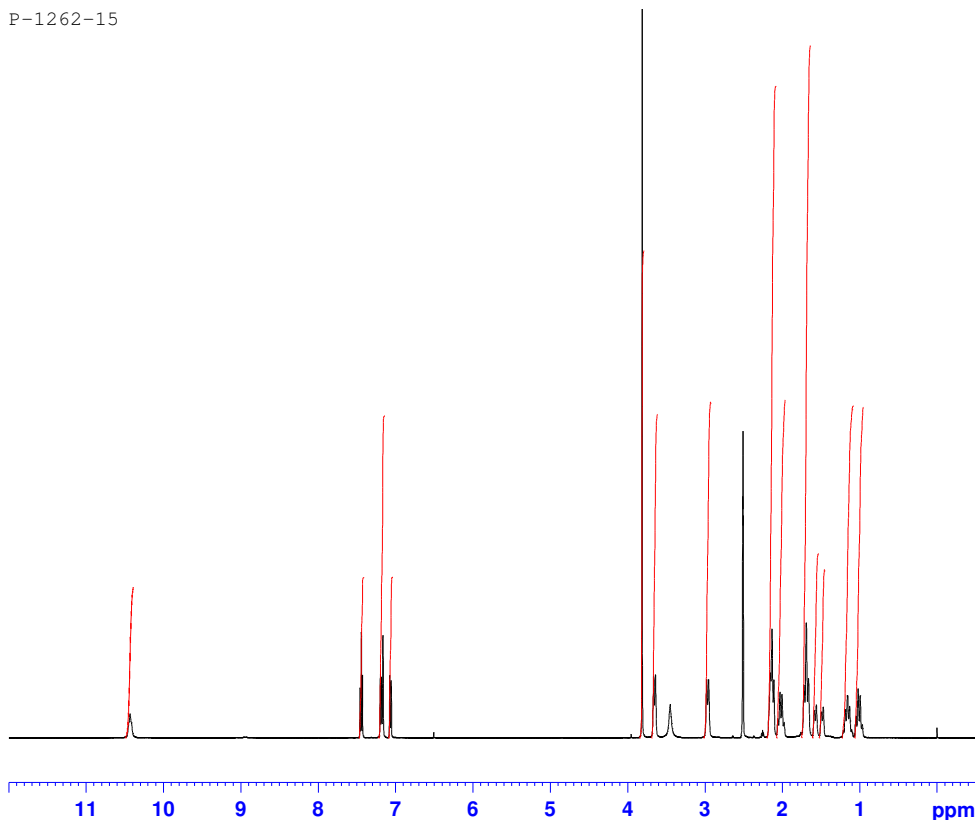


Current Data Parameters  
 NAME P-1262-15  
 EXPNO 1  
 PROCNO 1

F2 - Acquisition Parameters  
 Date\_ 20151101  
 Time 16.52  
 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 64  
 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 500.1300012 MHz  
 WDW EM  
 SSB 0  
 LB 0.30 Hz  
 GB 0  
 PC 1.00



P-1262-15



Current Data Parameters  
 NAME P-1262-15  
 EXPNO 4  
 PROCNO 1

F2 - Acquisition Parameters  
 Date\_ 20151101  
 Time 20.19  
 INSTRUM spect  
 PROBHD 5 mm PABBO BB-  
 PULPROG zgpg30  
 TD 65536  
 SOLVENT DMSO  
 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 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 125.7577890 MHz  
 WDW EM  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

