



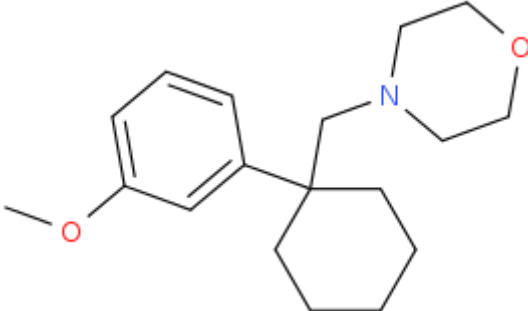
## ANALYTICAL REPORT<sup>1</sup>

### 3-MeO-PCMMo (C<sub>18</sub>H<sub>27</sub>N<sub>2</sub>O<sub>2</sub>)

#### 4-((1-(3-methoxyphenyl)cyclohexyl)methyl)morpholine

Remark – other NPS detected: **none**

Sample ID:	1662-16
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	8/30/2016
Date of entry (M/D/Y) into NFL database:	10/18/2016
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	4-((1-(3-methoxyphenyl)cyclohexyl)methyl)morpholine
Other names	
Formula (per base form)	C <sub>18</sub> H <sub>27</sub> N <sub>2</sub> O <sub>2</sub>
M <sub>w</sub> (g/mol)	289,4
Salt form/anions detected	HCl
StdInChIKey	KVDDTOKOCUZIFC-UHFFFAOYSA-N
Compound Class	Arylcyclohexylamines
Other NPS detected	none
Add.info (purity..)	pure by HPLC-TOF, GC-MS, NMR-some minor impurities

<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)
09/01/2017	Compound class corrected.

### 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

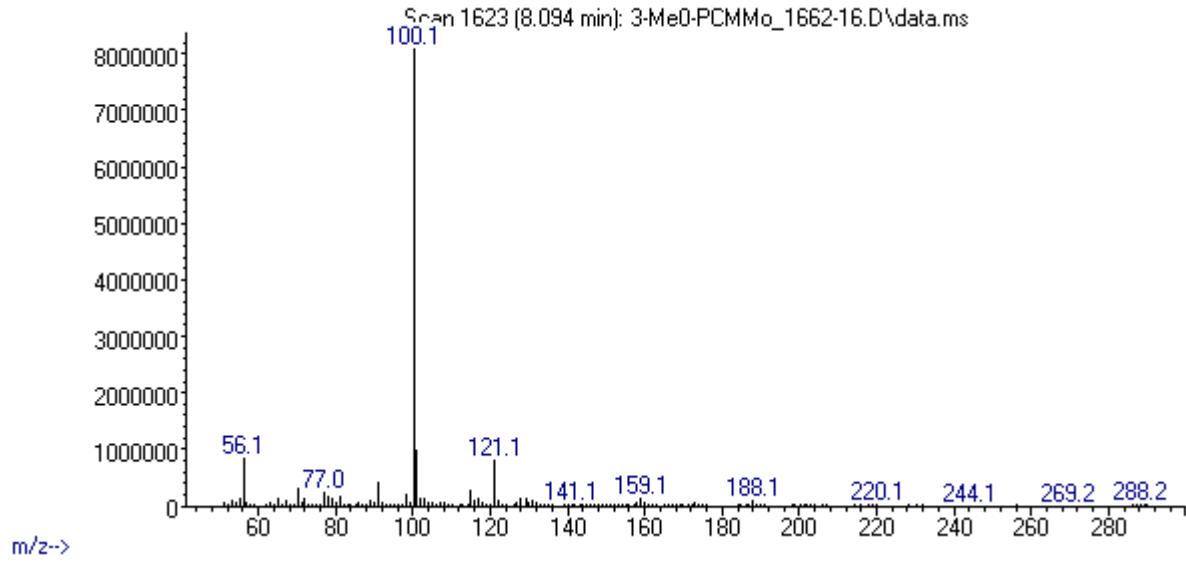
Solubility in	result/remark
CH <sub>2</sub> Cl <sub>2</sub>	soluble
MeOH	soluble
H <sub>2</sub> O	soluble

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 8,09 BP(1): 100; BP(2): 101,BP(3) :56,
HPLC-TOF	+	Exact mass (theoretical): 289,2042; measured value Δppm:-1,61; formula:C18H27NO2
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR (in FKKT)	+	
validation		
other		

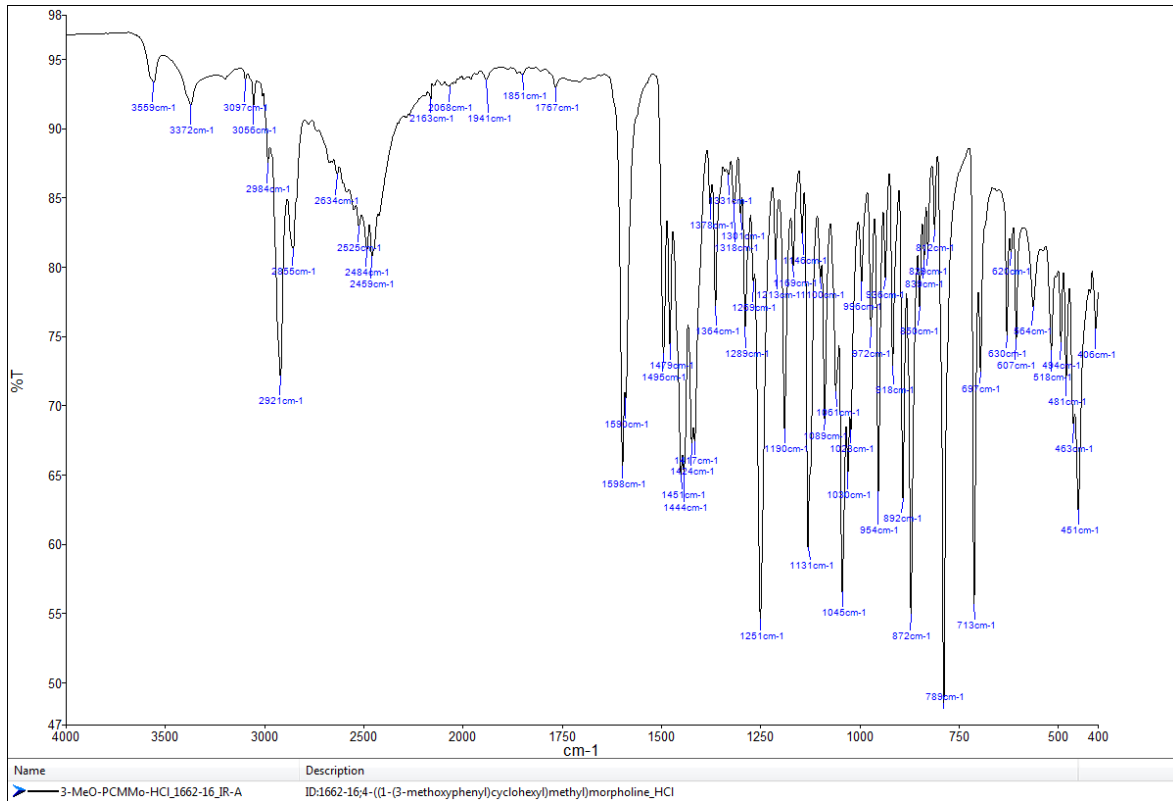
# ANALYTICAL RESULTS

MS (EI)

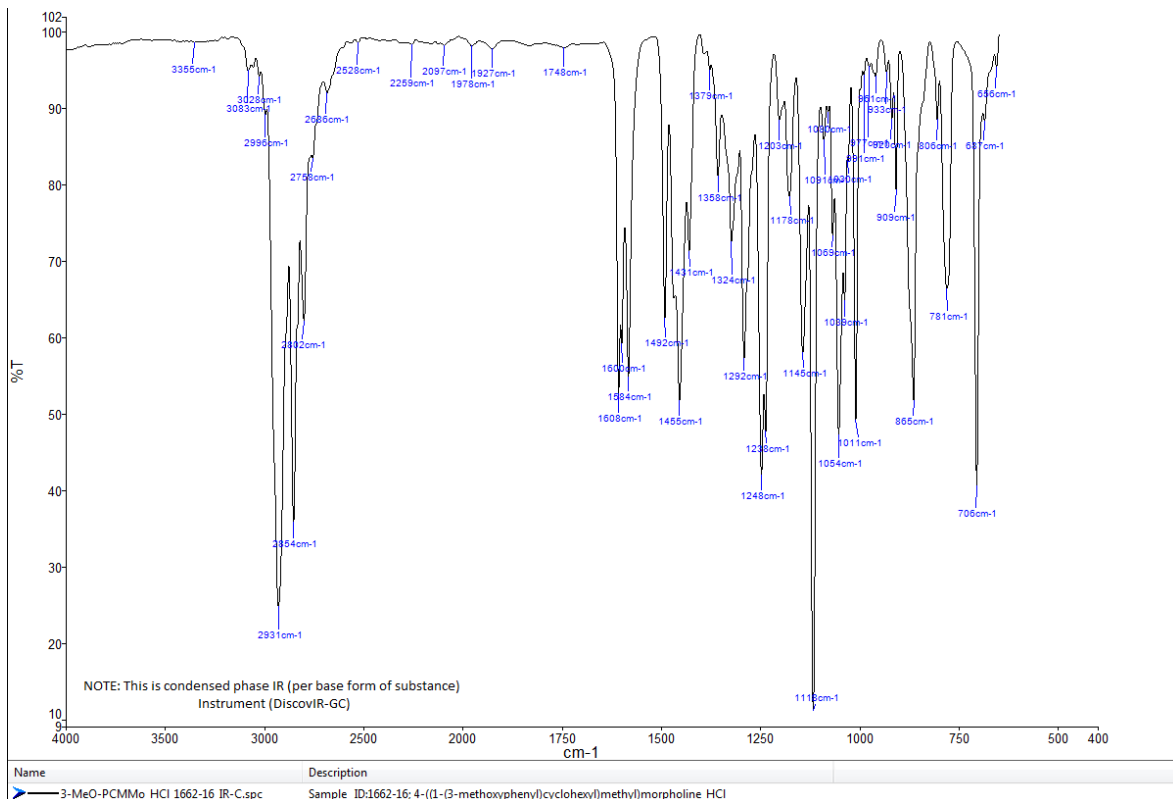
Abundance



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



## IR (condensed phase – after chromatographic separation)



# TOF REPORT

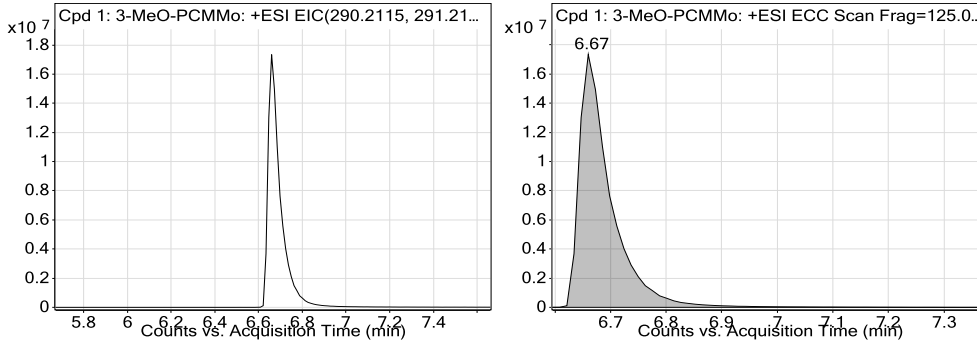
<b>Data File</b>	3-MeO-PCMMo_1662-16_TOF.d	<b>Sample Name</b>	ID_1662-16
<b>Sample Type</b>	Sample	<b>Position</b>	P1-A5
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	general-24_08_2016-XDB-C18-ESI-poz-soft.m	<b>Acquired Time</b>	9/1/2016 8:31:10 AM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Drugs_NFL.m
<b>Comment</b>	extract in MeOH		

### Compound Table

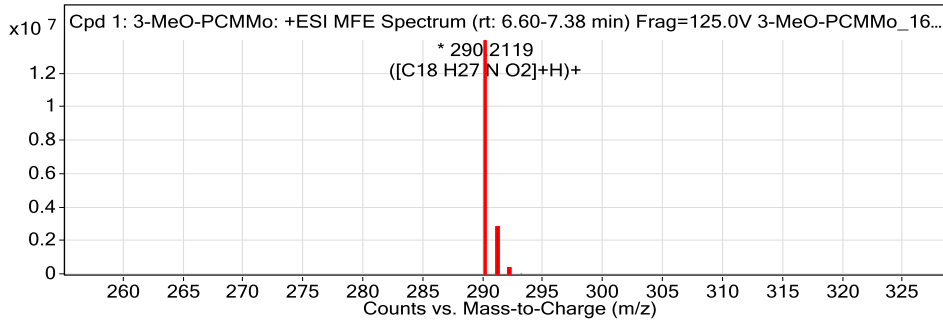
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 1: 3-MeO-PCMMo	3-MeO-PCMMo	C18 H27 N O2	6.67	289.2046

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
3-MeO-PCMMo	290.2119	6.67	289.2046	6.67	C18 H27 N O2	289.2042	-1.61

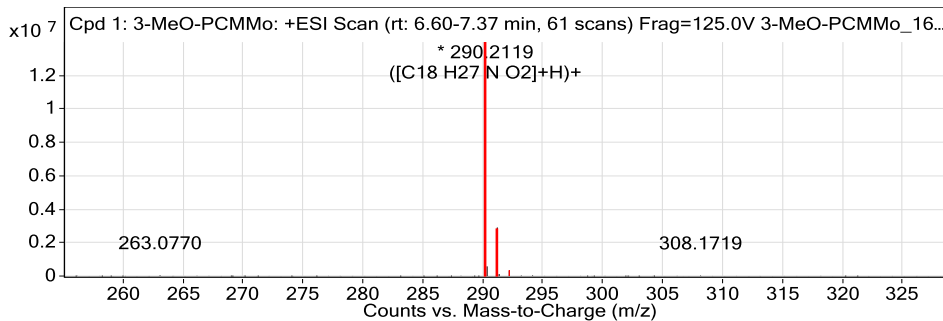
### Compound Chromatograms



### MFE MS Zoomed Spectrum



### MS Zoomed Spectrum



### MS Spectrum Peak List

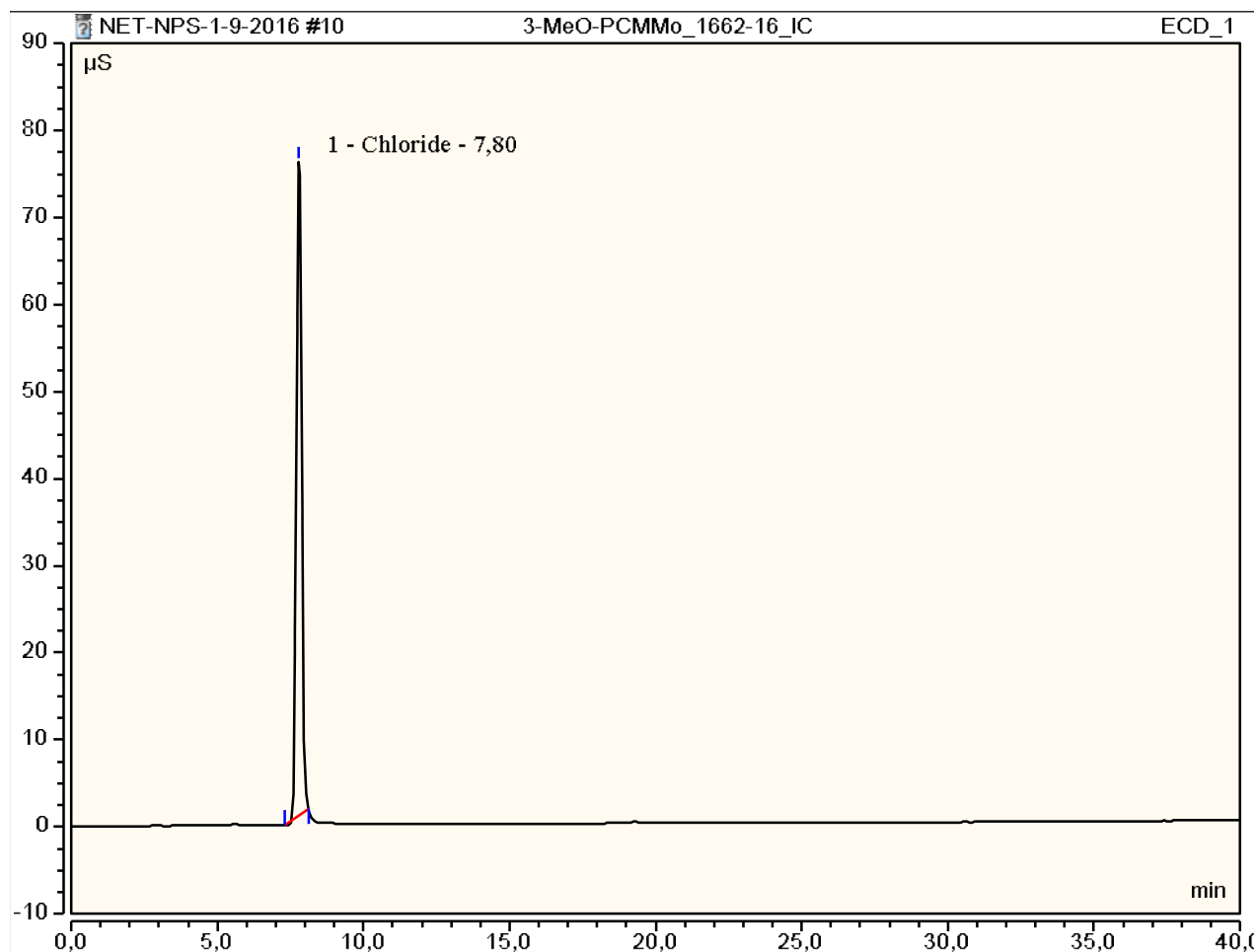
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
290.2119	1	13982980	C18 H27 N O2	(M+H)+
291.2152	1	2762841.27	C18 H27 N O2	(M+H)+
292.2183	1	280577.41	C18 H27 N O2	(M+H)+
293.2207	1	23456.76	C18 H27 N O2	(M+H)+
294.2225	1	1492.78	C18 H27 N O2	(M+H)+

--- End Of Report ---

### Peak Integration Report

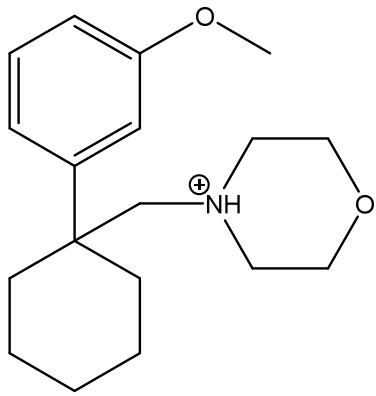
Sample Name:	3-MeO-PCMMo_1662-16_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	01-sep-2016 / 17:08	Run Time:	42,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}\cdot\text{min}$	Height $\mu\text{S}$	Amount mg/L
1,00	7,80	Chloride	BMB	15,12	75,08	n.a.
TOTAL:				15,12	75,08	0,00





## REPORT

Sample ID:	<b>1662-16</b>
Our notebook code:	P-1662-16
NMR sample preparation:	15 mg dissolved in 0.7 mL CDCl <sub>3</sub>
NMR experiments:	<sup>1</sup> H, <sup>13</sup> C, <sup>1</sup> H- <sup>1</sup> H <i>gs</i> -COSY, <sup>1</sup> H- <sup>13</sup> C <i>gs</i> -HSQC, <sup>1</sup> H- <sup>13</sup> C <i>gs</i> -HMBC, <sup>1</sup> H- <sup>15</sup> N <i>gs</i> -HMBC.
Proposed structure:	
Chemical name:	4-((1-(3-methoxyphenyl)cyclohexyl)methyl)morpholin-4-ium cation
Comments:	- Structure elucidation based on 1D and 2D NMR spectra - Sample is not pure as evident by NMR; it contains some minor impurities
Supporting information:	Copies of <sup>1</sup> H and <sup>13</sup> C NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	October 12, 2016



P-1662-16



```

Current Data Parameters
NAME          p-1662-16
EXPNO         1
PROCNO        1

F2 - Acquisition Parameters
Date_         20160909
Time          6.05
INSTRUM       spect
PROBHD        5 mm PABBO BB-
PULPROG       zg30
TD            65536
SOLVENT       CDCl3
NS            16
DS            2
SWH           10000.000 Hz
FIDRES        0.152588 Hz
AQ            3.2768500 sec
RG            80.6
DW            50.000 usec
DE            6.50 usec
TE            300.0 K
D1            1.00000000 sec
TD0           1

```

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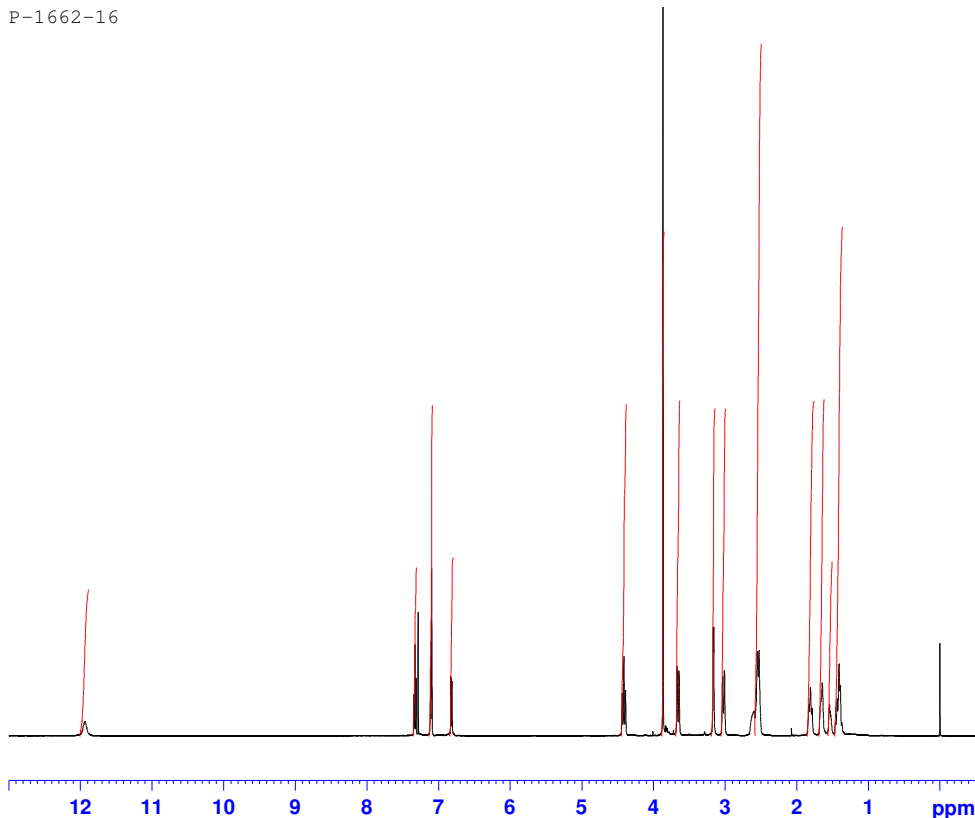
===== CHANNEL f1 =====
SFO1          500.1330885 MHz
NUC1           1H
P1             8.90 usec
PLW1          26.00000000 W

```

```

F2 - Processing parameters
SI            65536
SF            500.1300002 MHz
WDW           EM
SSB           0
LB            0.30 Hz
GB            0
PC            1.00

```



P-1662-16



```

Current Data Parameters
NAME          P-1662-16
EXPNO         3
PROCNO        1

F2 - Acquisition Parameters
Date_         20160909
Time          8.03
INSTRUM       spect
PROBHD        5 mm PABBO BB-
PULPROG       zgpg30
TD            65536
SOLVENT       CDCl3
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            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

```

