



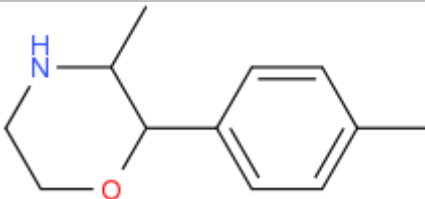
## ANALYTICAL REPORT

### 4-MPH (C<sub>12</sub>H<sub>17</sub>NO)

### 3-methyl-2(p-tolyl)morpholine

Remark – other NPS detected: **none**

Sample ID:	1243-15
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Comments <sup>1</sup> :	
Date of entry into NFL database: <a href="http://www.policija.si/apps/nfl_response_web/seznam.php">http://www.policija.si/apps/nfl_response_web/seznam.php</a>	8/19/2015

Substance identified-structure <sup>2</sup> (base form)	
Systematic name	3-methyl-2(p-tolyl)morpholine
Other names	3-methyl-2-(4-methylphenyl)morpholine, 4-MPM, 4-methylphenmetrazine
Formula (per base form)	C <sub>12</sub> H <sub>17</sub> NO
M <sub>w</sub> (g/mol)	191.27
Salt form	HCl
StdInChIKey	NWNCIXFIIDVRKE-UHFFFAOYSA-N
Compound Class	Others
Other NPS detected	none
Add.info (purity..)	pure by GC, totaly soluble in CH <sub>2</sub> Cl <sub>2</sub> , H <sub>2</sub> O, MeOH /NMR:(a few %) of unidentified organic compounds unidentified organic compound(s).

<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)
29/03/16	Compound class corrected.

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**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<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> .

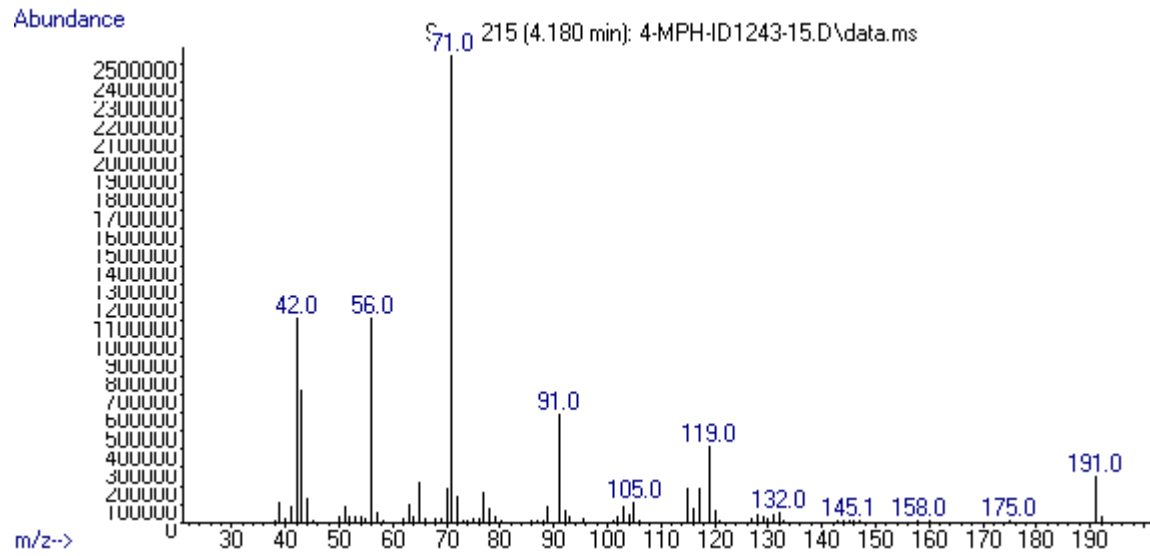
## Supporting information

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

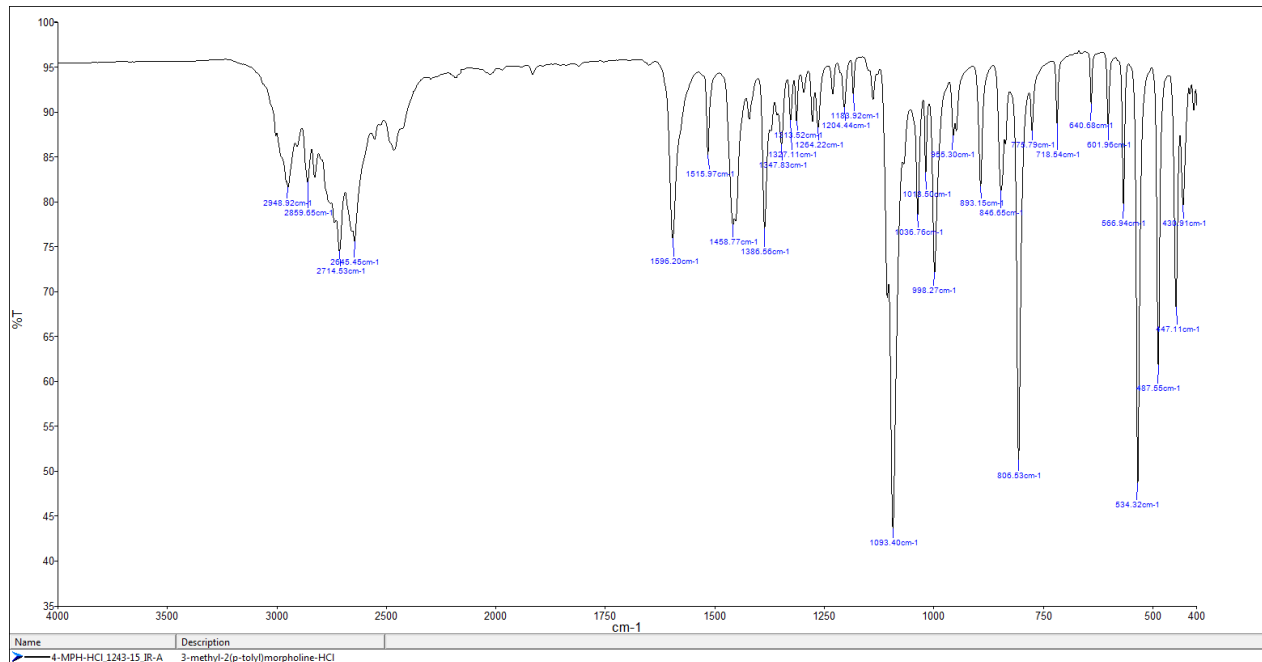
Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 4.17 BP(1): 71; BP(2): 56,BP(3) :42,
HPLC-TOF	+	Exact mass (theoretical): 191.31; measured value Δppm:-0.46; formula:C12H17NO
FTIR-ATR	+	direct measurement
FTIR (condensed phase) always as base form	+	
NMR	+	
validation		
other		

## ANALYTICAL RESULTS

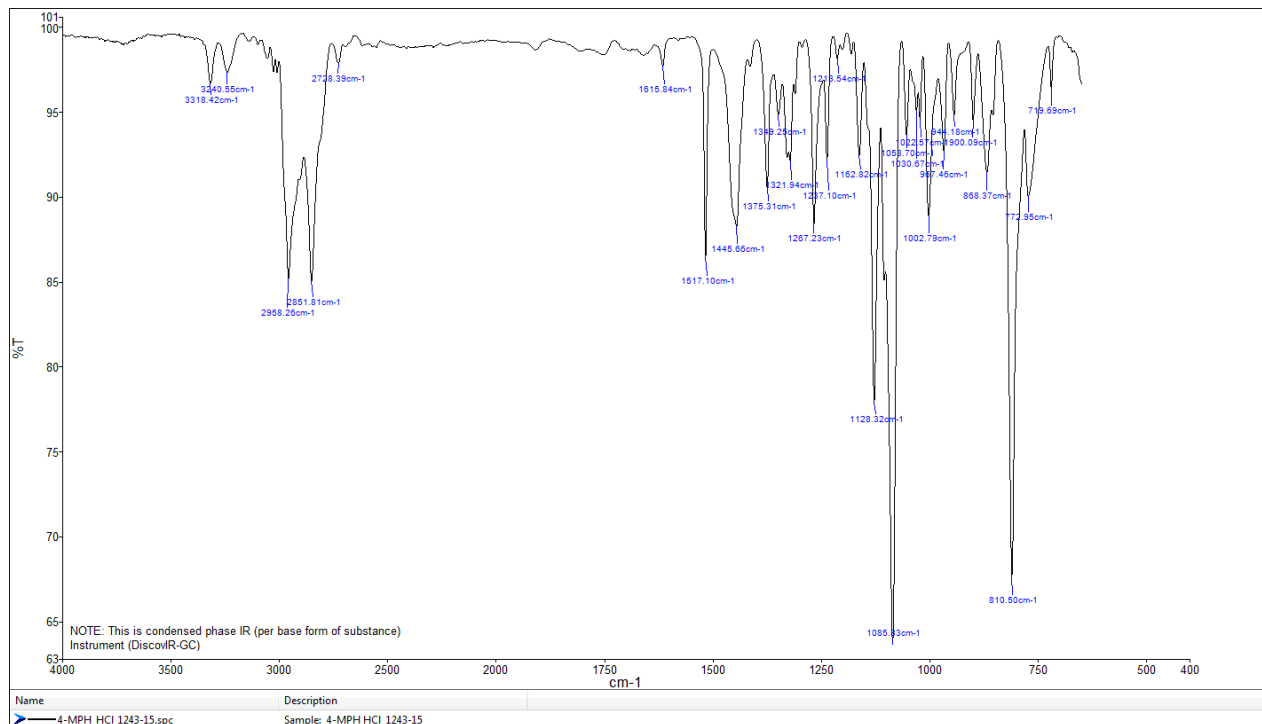
MS (EI)



## FTIR-ATR - direct measurement



## IR (condensed phase)



# Target Compound Screening Report

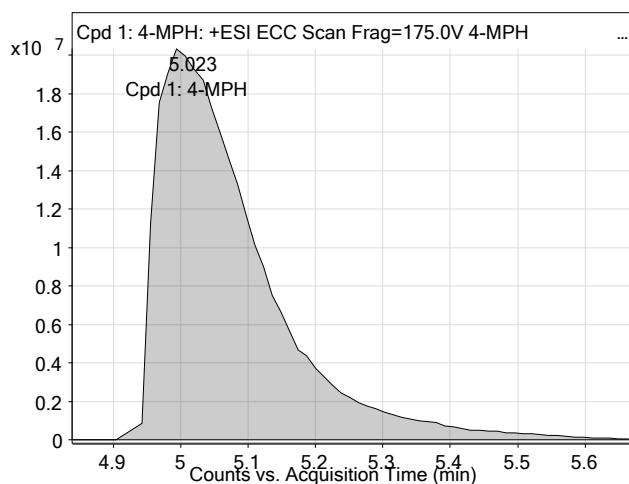
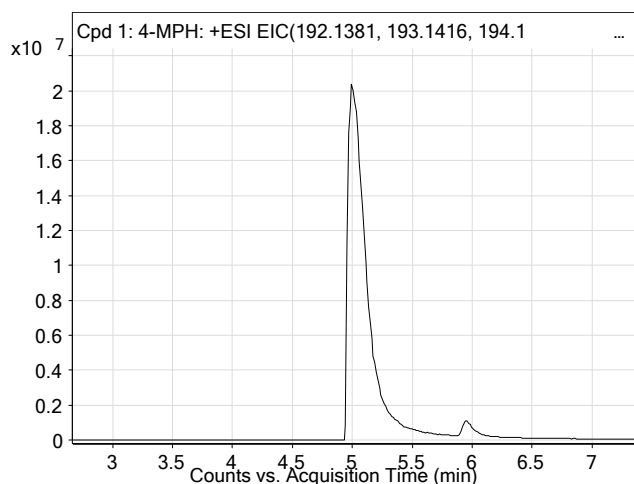
<b>Data File</b>	4-MPH_1243-15_TOF.d	<b>Sample Name</b>	4-MPH
<b>Sample Type</b>	Sample	<b>Position</b>	P2-E1
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	droge general-13-5-2015-XDB-C18-ESI-poz.m	<b>Acquired Time</b>	8/19/2015 12:28:48 PM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Droge_Default.m
<b>Comment</b>	extract in MeOH		

## Compound Table

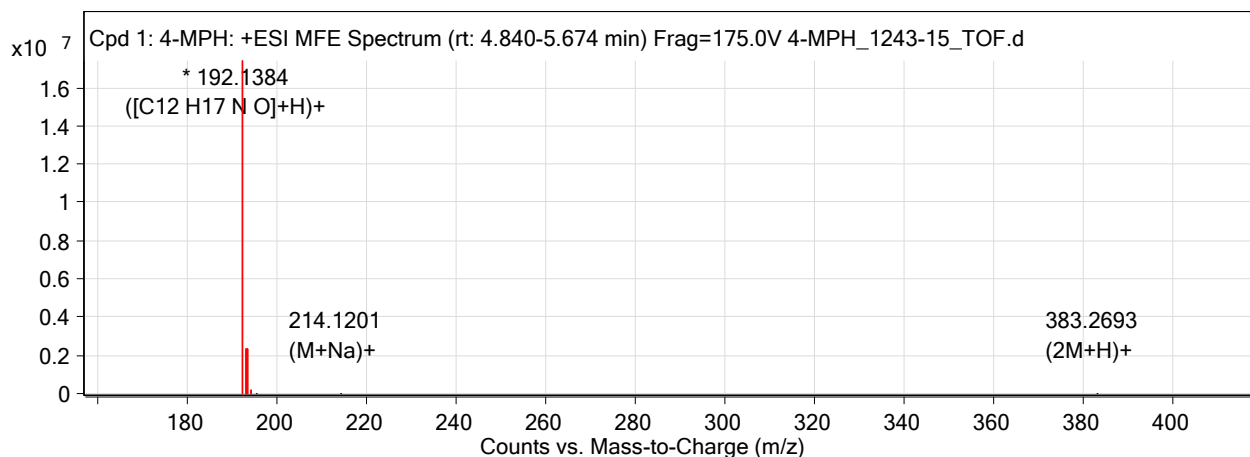
Label	Tgt Name	Obs. RT	Obs. Mass
Cpd 1: 4-MPH	4-MPH	5.023	191.1311

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)	Find Cpds Algorithm
4-MPH	192.1384	5.023	191.1311	5.02	C12 H17 N O	191.131	-0.46	Find by Molecular Feature

## Compound Chromatograms



## MFE MS Zoomed Spectrum

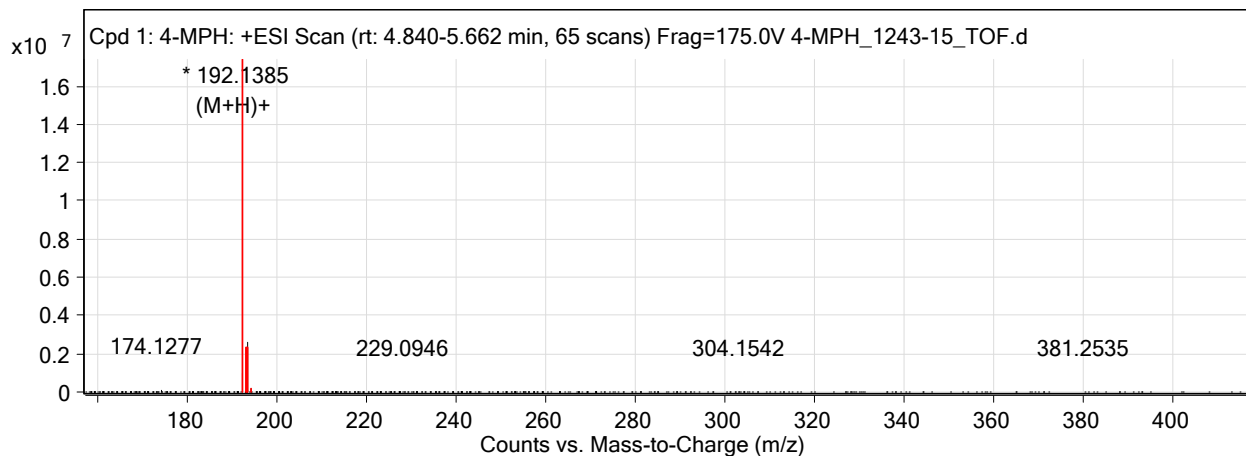


## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
192.1384	1	17392268	C12 H17 N O	(M+H)+
193.1417	1	2319382.39	C12 H17 N O	(M+H)+
194.1447	1	179934.52	C12 H17 N O	(M+H)+
195.1466	1	4395.84	C12 H17 N O	(M+H)+
214.1201	1	5997.31		(M+Na)+
383.2693	1	2583.29		(2M+H)+

# Target Compound Screening Report

## MS Zoomed Spectrum

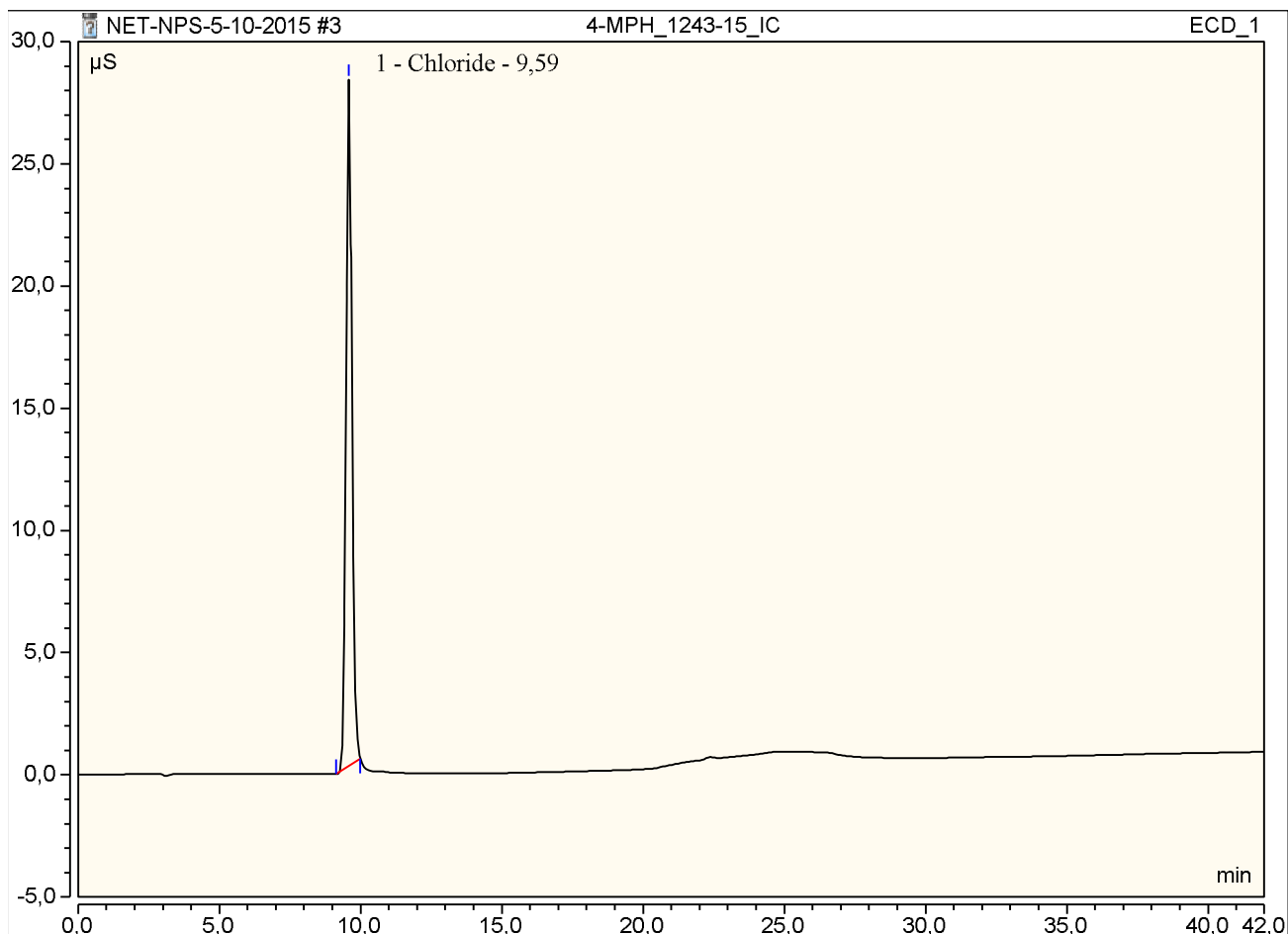


--- End Of Report ---

### Peak Integration Report

Sample Name:	4-MPH_1243-15_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	05-okt-2015 / 15:27	Run Time:	41,99

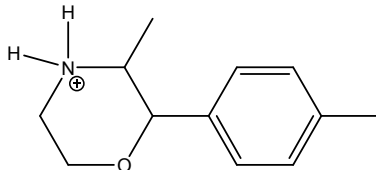
No.	Time min	Peak Name	Peak Type	Area $\mu\text{S} \cdot \text{min}$	Height $\mu\text{S}$	Amount mg/L
1,00	9,59	Chloride	BMB	6,75	28,09	n.a.
		TOTAL:		6,75	28,09	0,00







## REPORT

Sample ID:	<b>1243-15</b>
Our notebook code:	P-1243-15
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.
Proposed structure with chemical name:	 <p>3-methyl-2-(<i>p</i>-tolyl)morpholin-4-ium</p>
Comments:	<ul style="list-style-type: none"> <li>- Structure elucidation based on 1D and 2D NMR spectra</li> <li>- Compound is pure by NMR, containing a small amount (a few %) of unidentified organic compound(s).</li> </ul>
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 6, 2015

P-1243-15



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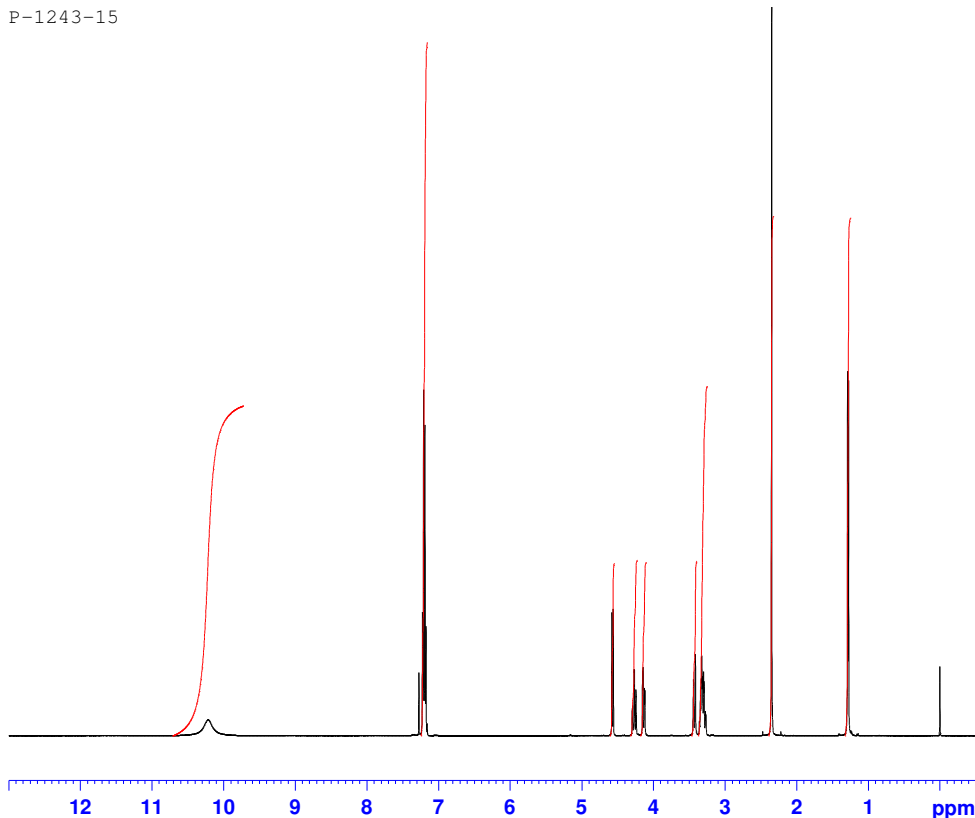
Current Data Parameters
NAME          p-1243-15
EXPNO         1
PROCNO        1

F2 - Acquisition Parameters
Date_         20151003
Time          15.10
INSTRUM       spect
PROBHD        5 mm PABBO BB-
PULPROG       zg30
TD            65536
SOLVENT       CDC13
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            295.0 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            500.1300063 MHz
WDW           EM
SSB           0
LB            0.30 Hz
GB            0
PC            1.00

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P-1243-15



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Current Data Parameters
NAME          p-1243-15
EXPNO         3
PROCNO        1

F2 - Acquisition Parameters
Date_         20151003
Time          20.18
INSTRUM       spect
PROBHD        5 mm PABBO BB-
PULPROG       zgpg30
TD            65536
SOLVENT       CDC13
NS            8192
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            296.2 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            125.7577896 MHz
WDW           EM
SSB           0
LB            1.00 Hz
GB            0
PC            1.40

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