



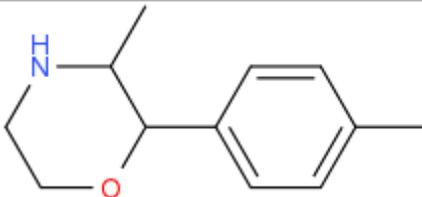
## ANALYTICAL REPORT

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

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

Remark – other NPS detected: **none**

Sample ID:	1294-15
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Comments <sup>1</sup> :	source II
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>	10/14/2015

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, mephenmetrazine, 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-MS, HPLC-TOF,

<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

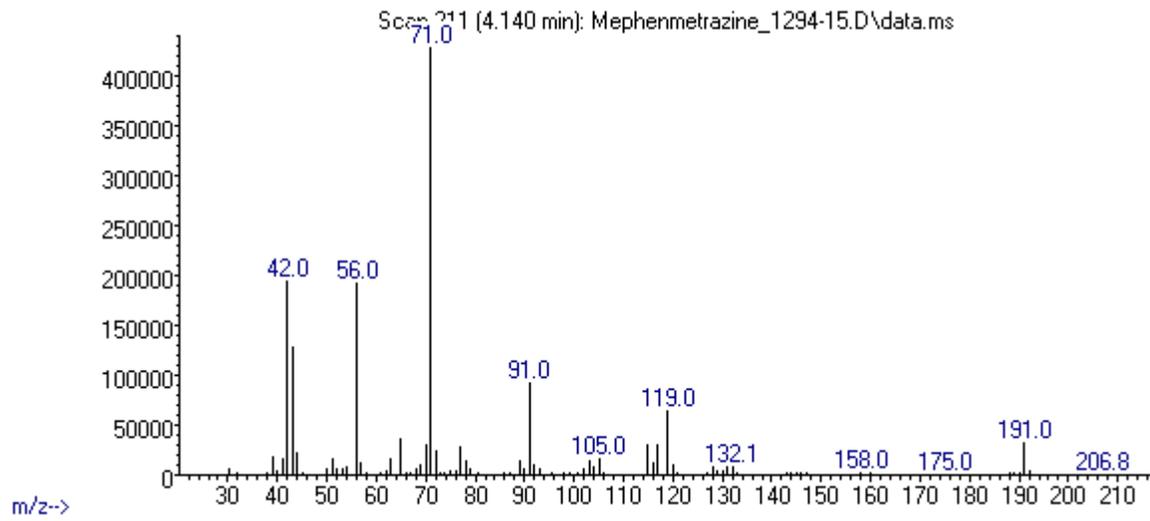
Solubility in	result/remark
CH <sub>2</sub> Cl <sub>2</sub>	soluble
MeOH	soluble
H <sub>2</sub> O	soluble

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 4.14 BP(1): 71; BP(2): 42,BP(3) :56,
HPLC-TOF	+	Exact mass (theoretical): 191.131; measured value Δppm:-1.21; formula:C12H17NO
FTIR-ATR	+	direct measurement
FTIR (condensed phase) always as base form	+	
NMR	-	
validation		FTIR consistent (qm>0.99) with the 4-MPH spectrum of sample confirmed by NMR (ID-1243-15)
other		

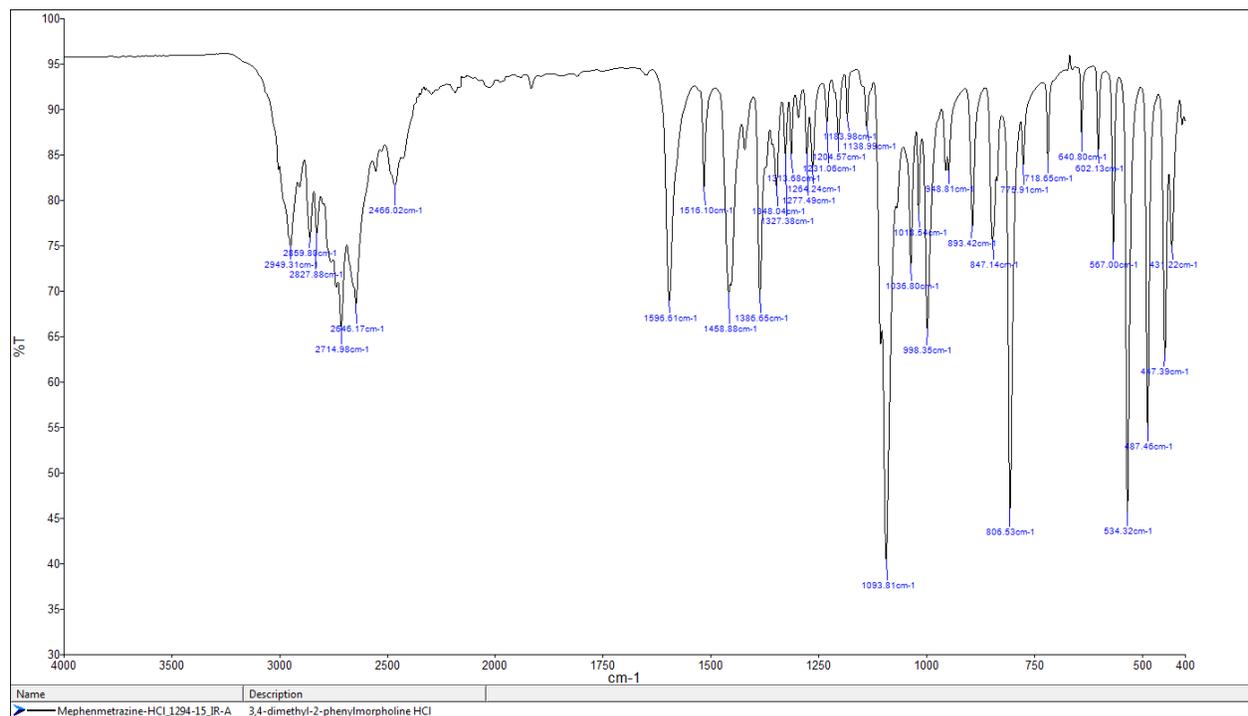
## ANALYTICAL RESULTS

MS (EI)

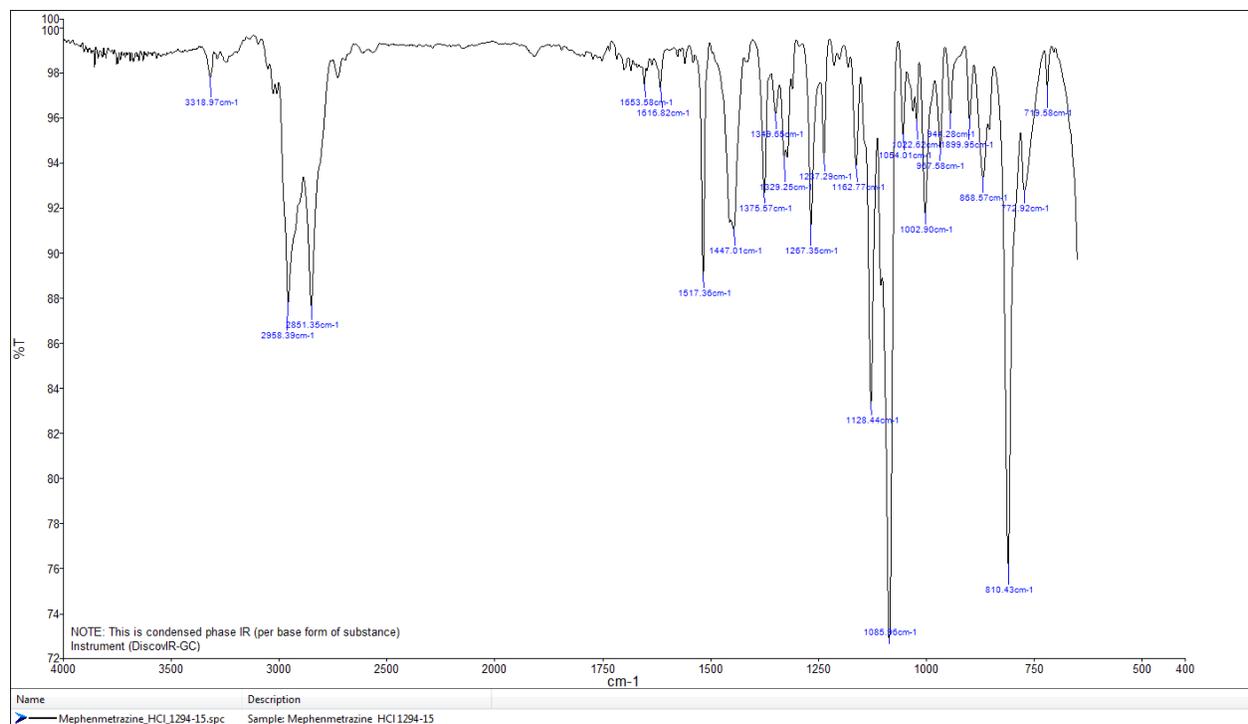
Abundance



## FTIR-ATR - direct measurement



## IR (condensed phase)



# TOF REPORT

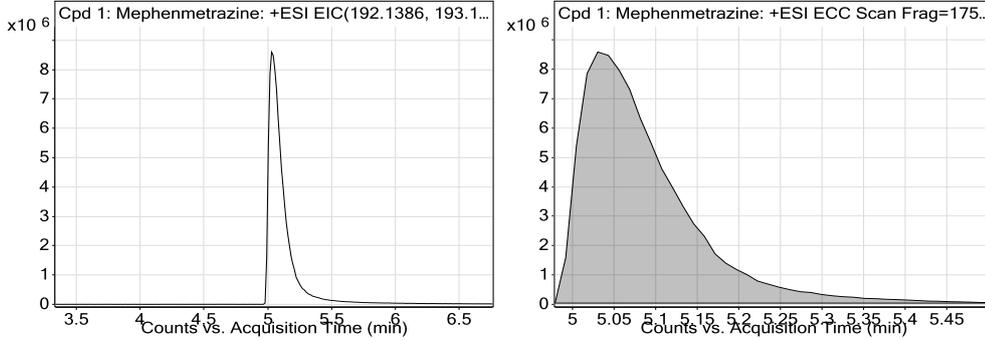
<b>Data File</b>	Mephenmetrazine-1294-15_TOF.d	<b>Sample Name</b>	Mephenmetrazine
<b>Sample Type</b>	Sample	<b>Position</b>	P2-B9
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	general-28052015-XDB-C18-ESI-poz.m	<b>Acquired Time</b>	9/28/2015 1:49:57 PM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Droge_Default.m
<b>Comment</b>	extract in MeOH		

## Compound Table

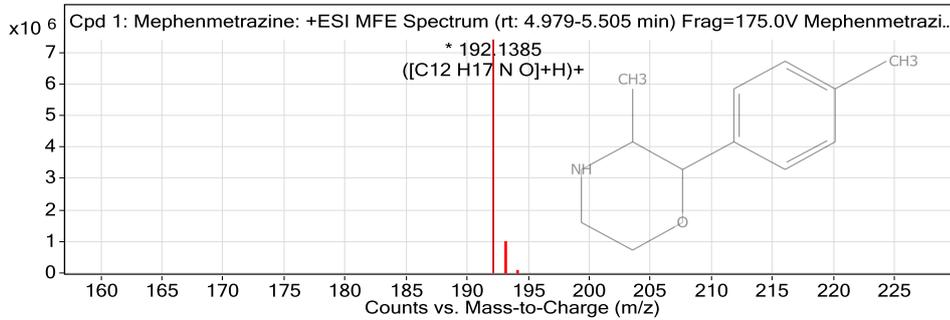
Label	Compound Name	Obs. RT	Obs. Mass
Cpd 1: Mephenmetrazine	Mephenmetrazine	5.05	191.1312

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
Mephenmetrazine	192.1385	5.05	191.1312	5.05	C12 H17 N O	191.131	-1.21

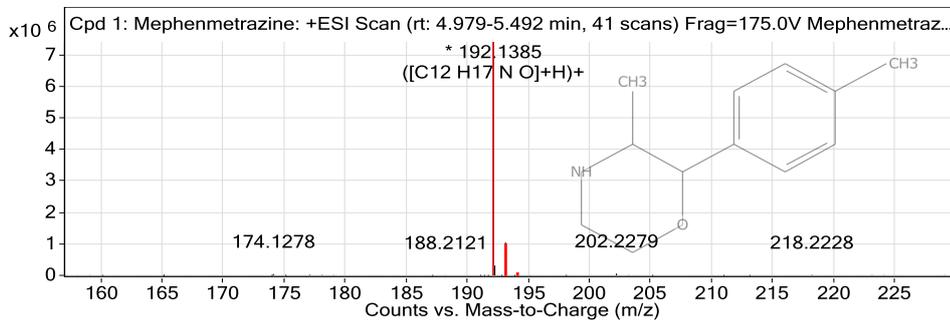
## Compound Chromatograms



## MFE MS Zoomed Spectrum



## MS Zoomed Spectrum



## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
192.1385	1	7402563.5	C12 H17 N O	(M+H)+
193.1421	1	1007947.69	C12 H17 N O	(M+H)+
194.1446	1	72009.59	C12 H17 N O	(M+H)+
195.1462	1	3421.98	C12 H17 N O	(M+H)+

--- End Of Report ---

### Peak Integration Report

Sample Name:	Mephenemtrazine_1294-15_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	30-sep-2015 / 13:54	Run Time:	41,99

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}^*\text{min}$	Height $\mu\text{S}$	Amount mg/L
1,00	9,74	Chloride	BMB	17,58	72,36	n.a.
TOTAL:				17,58	72,36	0,00

