

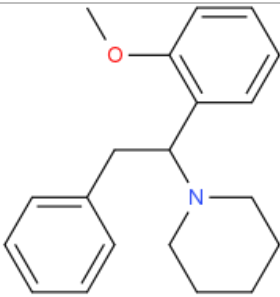
## ANALYTICAL REPORT

### 2-MeO-diphenidine (C<sub>20</sub>H<sub>25</sub>NO)

#### 1-(1-(2-methoxyphenyl)-2-phenylethyl)piperidine

Remark – other NPS detected: **none**

Sample ID:	1163-15A (corresponds to sample 3400-14-3E)
Sample description:	powder - white
Sample type:	seized
Comments:	LJ
Date of entry:	7/17/2014

Substance identified-structure <sup>i</sup> (base form)	
Systematic name	1-(1-(2-methoxyphenyl)-2-phenylethyl)piperidine
Other names	
Formula (per base form)	C <sub>20</sub> H <sub>25</sub> NO
M <sub>w</sub> (g/mol)	295,42
Salt form	HCl
Smiles	<chem>COC1=C(C=CC=C1)C(CC1=CC=CC=C1)N1CCCCC1</chem>
Compound Class	Others
Other NPS detected	none
Add.info (purity..)	pure by GC-MS, HPLC, NMR (not detected impurities are possible)

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.

## Supporting information

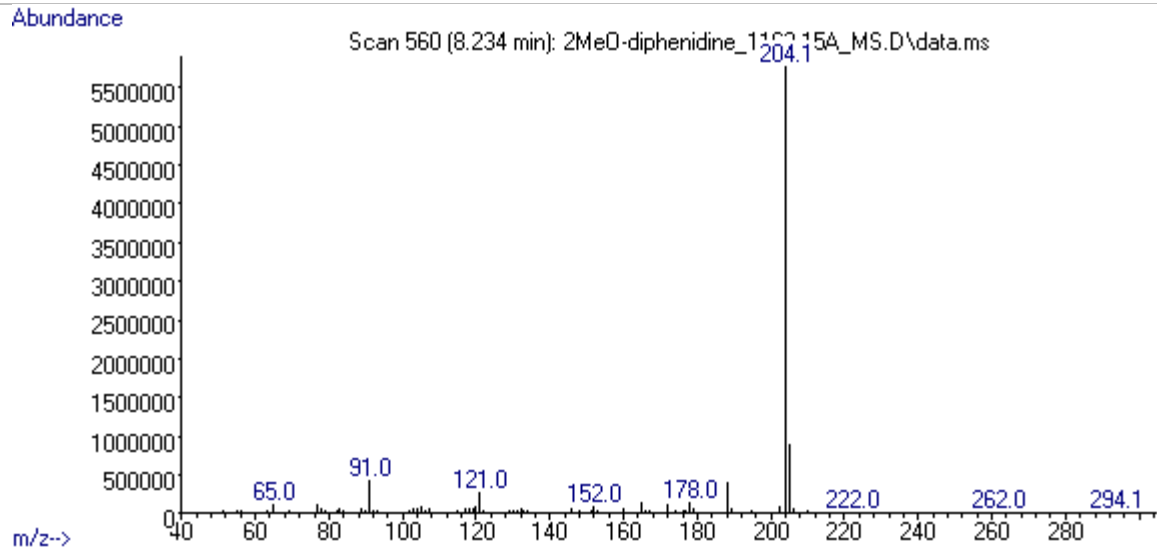
Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): BP(1): 204; BP(2): 205,BP(3) :91,
FTIR-ATR	+	direct measurement
FTIR (condensed phase) always as base form	-	
HPLC-TOF	+	Exact mass (theoretical): 295,1936; measured value $\Delta$ ppm:-0,61; formula:C20H25NO
NMR-confirmed	+	
validation		
other		completely soluble in CH <sub>2</sub> Cl <sub>2</sub> and in MeOH

GC-MS (Agilent): GC-method is RT locked to tetracosane (RT=9.53 min). Injection volume 1 ml and split mode (1:50)for GC-MS instruments and 1:5 for GC-MS-FTIR(condensed phase). Injector temperature: 280 °C. 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, than 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. m/z scan range: from 50 (40) to 550 amu.

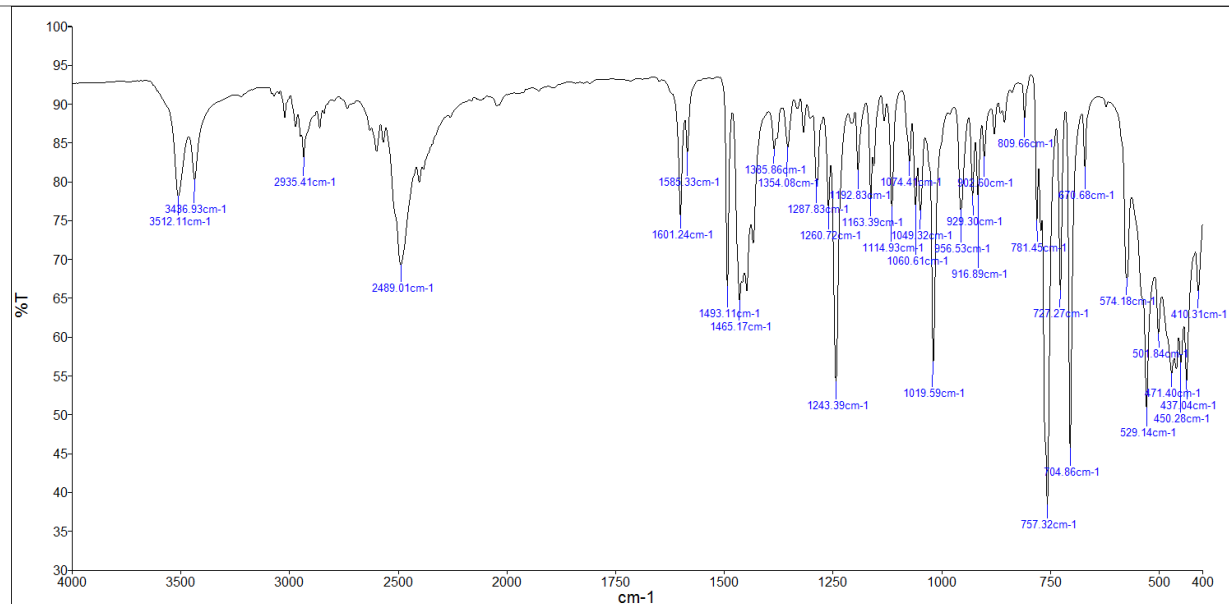
FTIR-ATR (Perkin Elmer): scan range 4000-400 cm<sup>-1</sup>; resolution 4cm<sup>-1</sup>

FTIR (Spectra analyses-Danny): scan range 4000 to 600, resolution 4cm<sup>-1</sup>

### MS spectrum (EI)



# FTIR - ATR



Name	Description
2MeO-diphenidine-HCl_1163-15A_IR-A	1-(1-(2-methoxyphenyl)-2-phenylethyl)piperidine

Created by OPSIN free tool: <http://opsin.ch.cam.ac.uk/> DOI: 10.1021/ci100384d

# Target Compound Screening Report

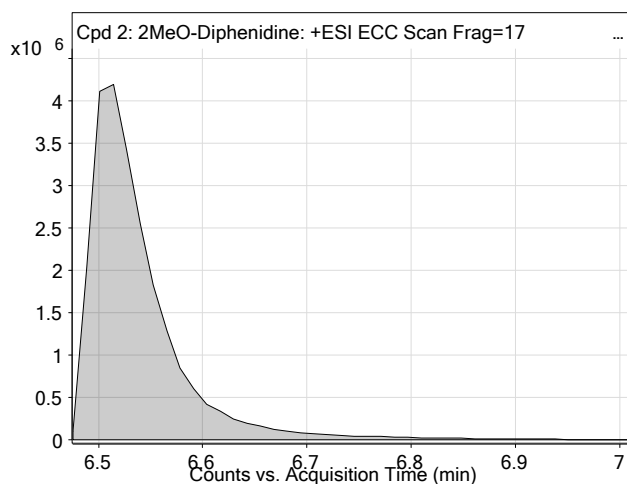
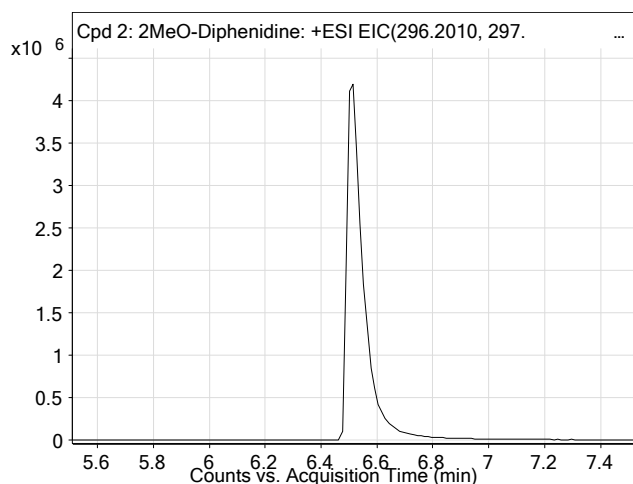
<b>Data File</b>	kben-233-3400_2014_1e.d	<b>Sample Name</b>	vzorec 1e
<b>Sample Type</b>	Sample	<b>Position</b>	P2-E7
<b>Instrument Name</b>	SG13170002	<b>User Name</b>	
<b>Acq Method</b>	droge general-17-9-2014-XDB-C18-ESI-poz.m	<b>Acquired Time</b>	9/18/2014 2:07:55 PM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Droge_Default.m
<b>Comment</b>			

## Compound Table

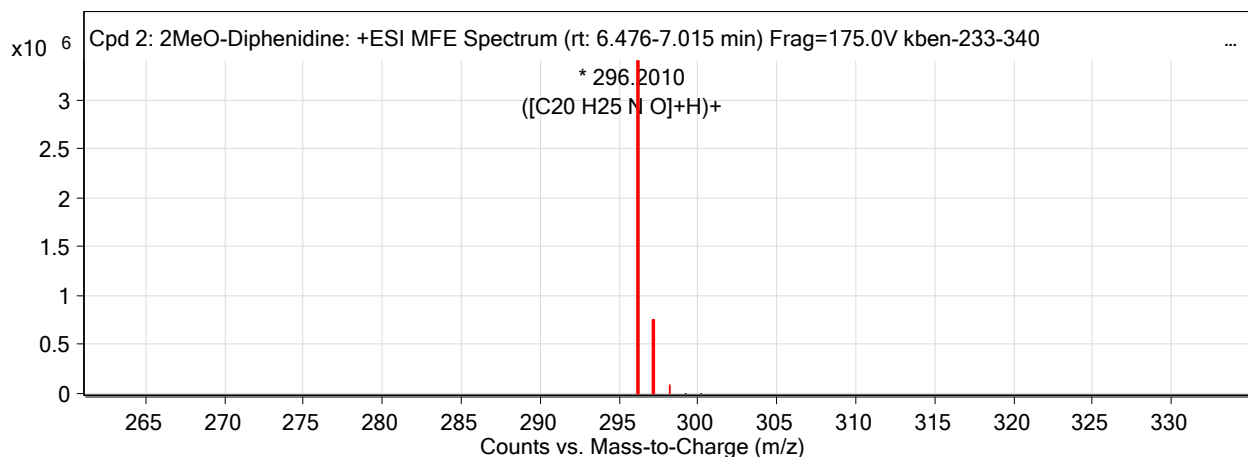
Label	Tgt Name	Obs. RT	Obs. Mass
Cpd 2: 2MeO-Diphenidine	2MeO-Diphenidine	6.515	295.1938

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)	Find Cmps Algorithm
2MeO-Diphenidine	296.201	6.515	295.1938	6.545	C20 H25 N O	295.1936	-0.61	Find by Molecular Feature

## Compound Chromatograms



## MFE MS Zoomed Spectrum

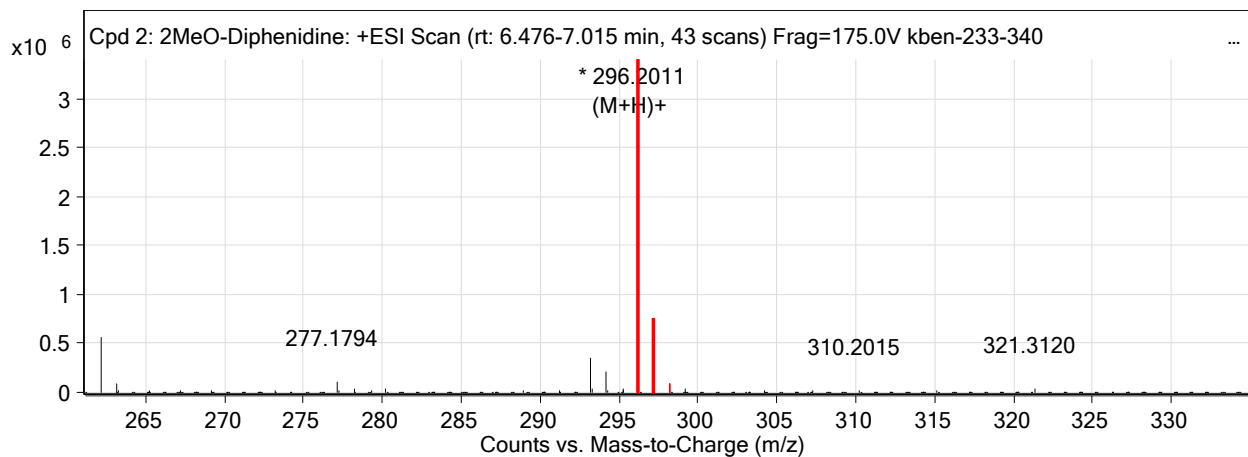


## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
296.201	1	3394337	C20 H25 N O	(M+H)+
297.2048	1	706808.43	C20 H25 N O	(M+H)+
298.2072	1	67544.2	C20 H25 N O	(M+H)+
299.2089	1	4820.21	C20 H25 N O	(M+H)+
300.2072	1	341.62	C20 H25 N O	(M+H)+

## MS Zoomed Spectrum

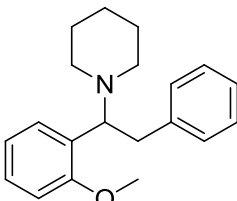
# Target Compound Screening Report



--- End Of Report ---



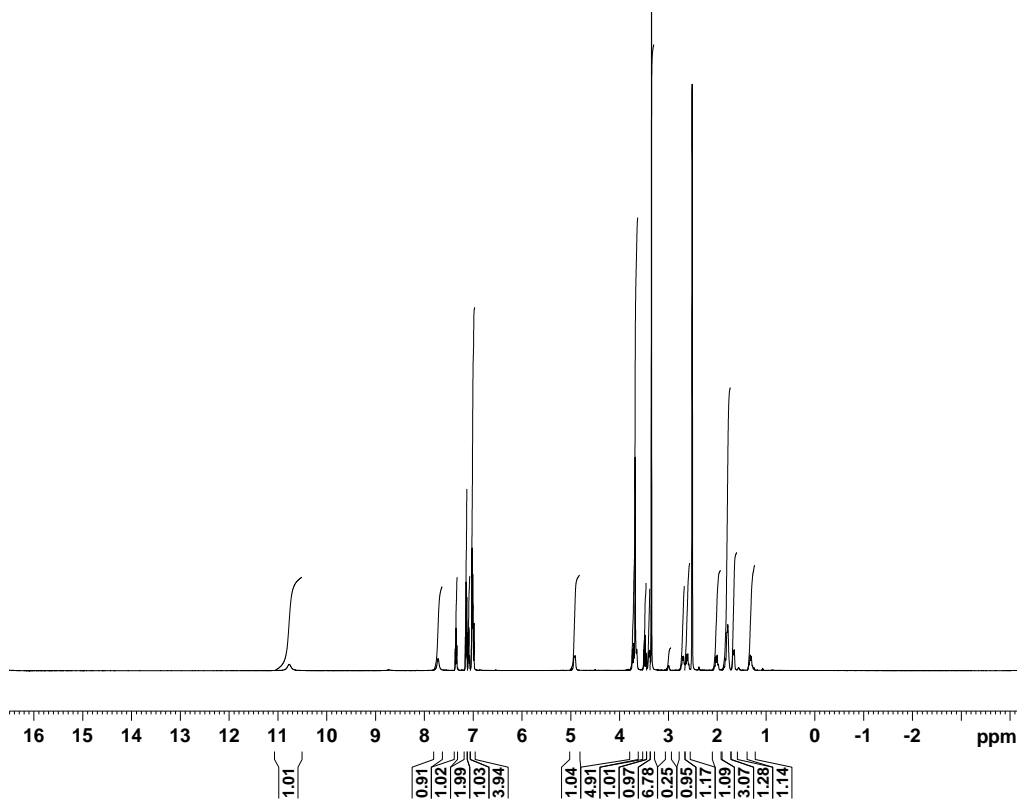
## REPORT

Sample ID:	<b>233-3400-14-1E</b>
Our notebook code:	P-233-3400-14-1E
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 with chemical name:	 <p style="text-align: right;">HCl</p> <p>1-(1-(2-methoxyphenyl)-2-phenylethyl)piperidine hydrochloride</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</li> </ul>
Supporting information:	Copies of $^1\text{H}$ and $^{13}\text{C}$ NMR spectra
Author:	Prof. Dr. Janez Košmrlj
Date of report:	April 30, 2015

P-233-3400-14-1E  
1H



Current Data Parameters  
NAME P-233-3400-14-1E  
EXFNO 3  
PROCNO 1  
F2 - Acquisition Parameters  
Date 20150131  
Time 15.37  
INSTRUM spect  
PROBHD 5 mm FAPBO BB-  
FULPROG zg30  
TD 65536  
SOLVENT DMSO  
NS 16  
DS 2  
SWH 10330.578 Hz  
FIDRES 0.157632 Hz  
AQ 3.1719923 sec  
RG 114  
DW 48.400 usec  
DE 6.50 usec  
TE 297.5 K  
D1 1.00000000 sec  
===== CHANNEL f1 =====  
NUC1 1H  
P1 8.90 usec  
PLW1 26.00000000 W  
SF01 500.1330885 MHz  
F2 - Processing parameters  
SI 65536  
SF 500.1300000 MHz  
WDW EM  
SSB 0  
LB 0.30 Hz  
GB 0  
PC 1.00



P-233-3400-14-1E  
13C



Current Data Parameters  
NAME P-233-3400-14-1E  
EXFNO 4  
PROCNO 1  
F2 - Acquisition Parameters  
Date 20150131  
Time 18.44  
INSTRUM spect  
PROBHD 5 mm FAPBO BB-  
FULPROG 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  
SF01 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  
SF02 500.1320005 MHz  
F2 - Processing parameters  
SI 32768  
SF 125.7578493 MHz  
WDW EM  
SSB 0  
LB 1.00 Hz  
GB 0  
PC 1.40

