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ANALYTICAL REPORT¹

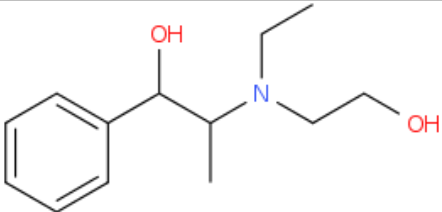
Phenmetetrazine-precursor (

C₁₃H₂₁N₂O₂)

2-[ethyl(2-hydroxyethyl)amino]-1-phenylpropan-1-ol

Remark – other NPS detected:

Sample ID:	1291-15
Sample description:	powder - white
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	9/25/2015
Date of entry (M/D/Y) into NFL database:	11/17/2015
Report updates (if any) will be published here:	http://www.policija.si/apps/nfl_response_web/seznam.php

Substance identified - structure ² (base form)	
Systematic name	2-[ethyl(2-hydroxyethyl)amino]-1-phenylpropan-1-ol
Other names	
Formula (per base form)	C ₁₃ H ₂₁ N ₂ O ₂
M _w (g/mol)	223,32
Salt form/anions detected	HCl
StdInChIKey	PIQWNMHHXJMHFQ-UHFFFAOYSA-N
Compound Class	Others
Other NPS detected	
Add.info (purity..)	HPLC-TOF additional peak with the same monoisotopic mass, trace impurities detected by GC-MS as well

¹ 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.

² Created by OPSIN free tool: <http://opsin.ch.cam.ac.uk/> DOI: 10.1021/ci100384d

Report updates

date	comments (explanation)
09/03/2016	New report issued for this CHEM-ID, due to misinterpretation of analytical data in the previous report (substance was reported as Phenmetetrazine instead of Phenmetetrazine-precursor). The report for Phenmetetrazine one can find under CHEM-ID: 1291-16s
09/08/2016	Typing error corrected (Mw).

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 6.1 min isothermal. MSD source EI = 70 eV. GC-MS transfer line T= 235 °C, source and quadrupole temperatures 280 °C and 180 °C, respectively. Scan range m/z scan range: from 50 (30 until 6 min.) to 550 (300) 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⁻¹; resolution 4cm⁻¹

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 quadrupole 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 phase): IR scan range 4000 to 650, resolution 4 cm⁻¹.

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

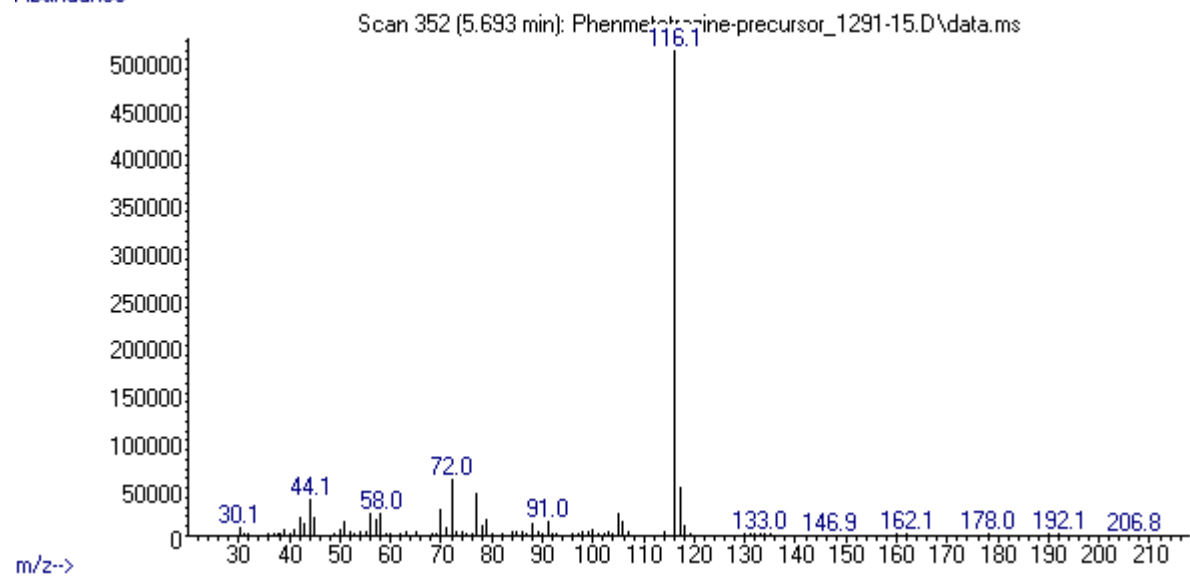
Solubility in	result/remark
CH ₂ Cl ₂	not soluble
MeOH	soluble
H ₂ O	not tested

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 5,69 BP(1): 116; BP(2): 117,BP(3) :72,
HPLC-TOF	+	Exact mass (theoretical): 233,1572; measured value Δppm:-1,41; formula: C ₁₃ H ₂₁ NO ₂
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	+	
IC (anions)	+	
NMR (in FKKT)	+	
validation		
other		

ANALYTICAL RESULTS

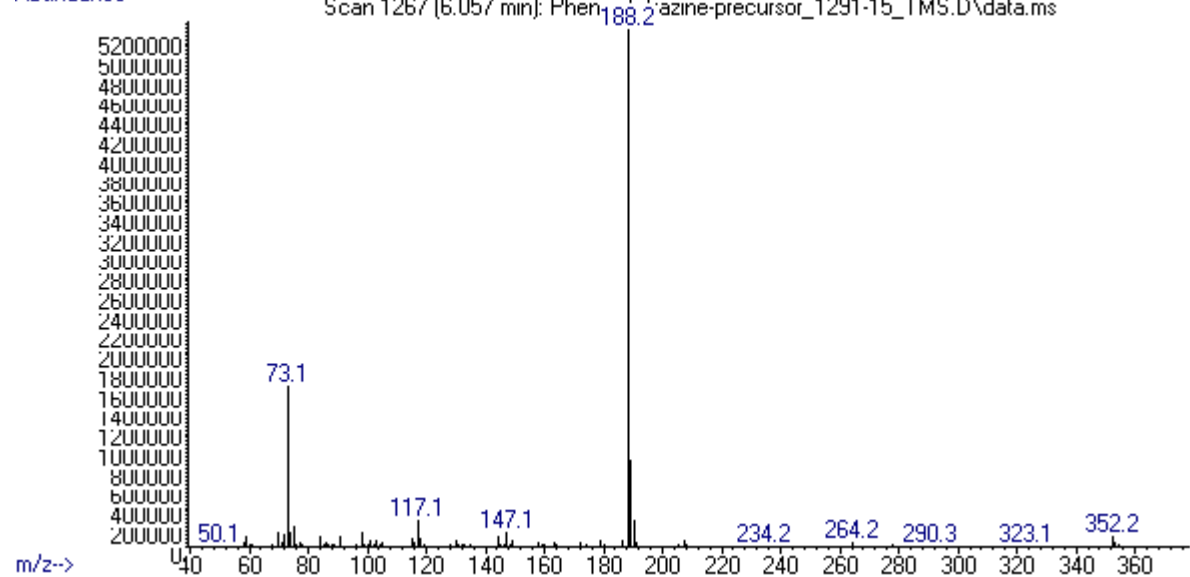
MS (EI+)

Abundance

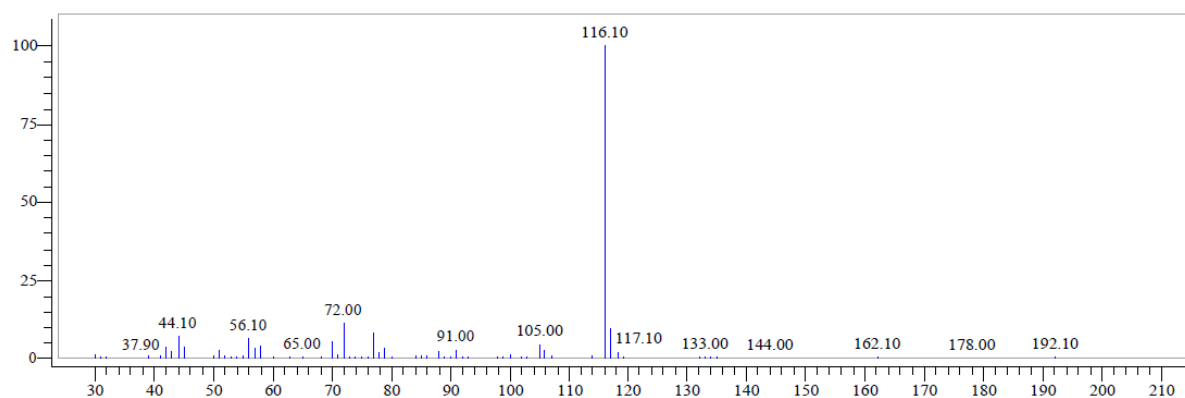


MS (EI+) of TMS derivative

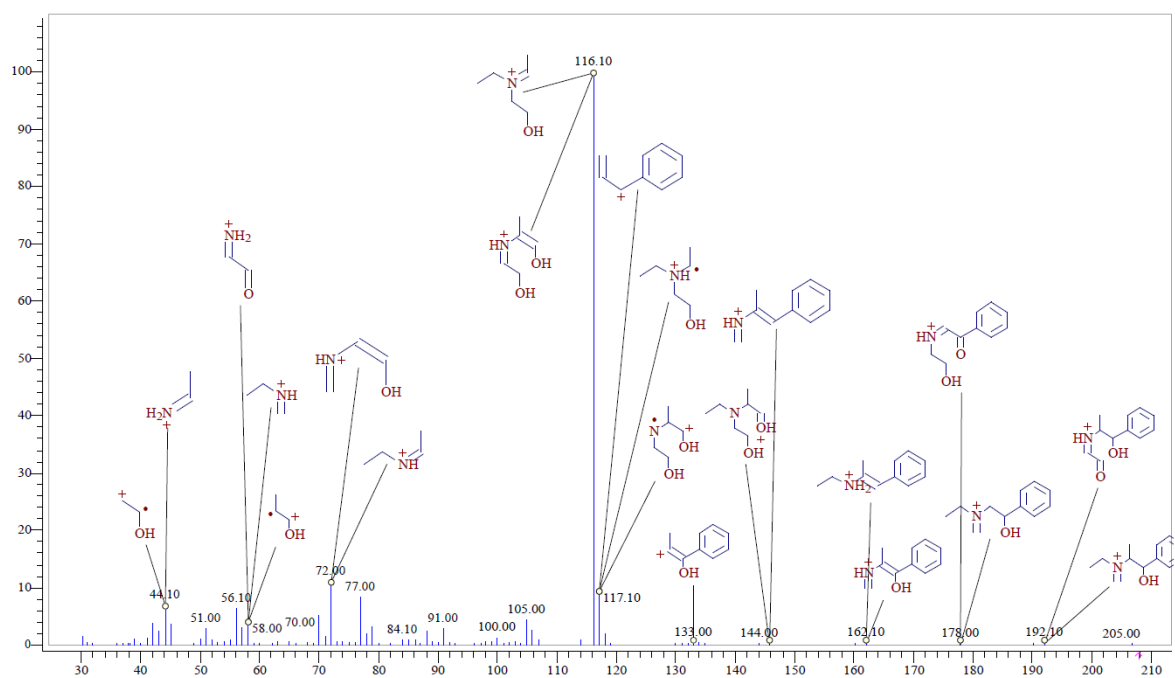
Abundance



MS interpretation - non derivatized sample

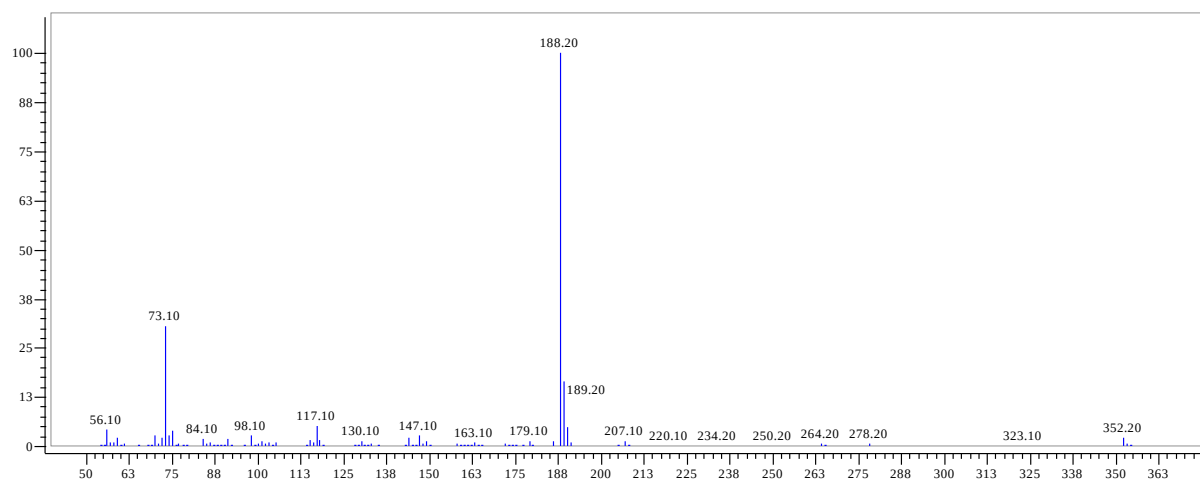


Mass spectrum - observed (measured)

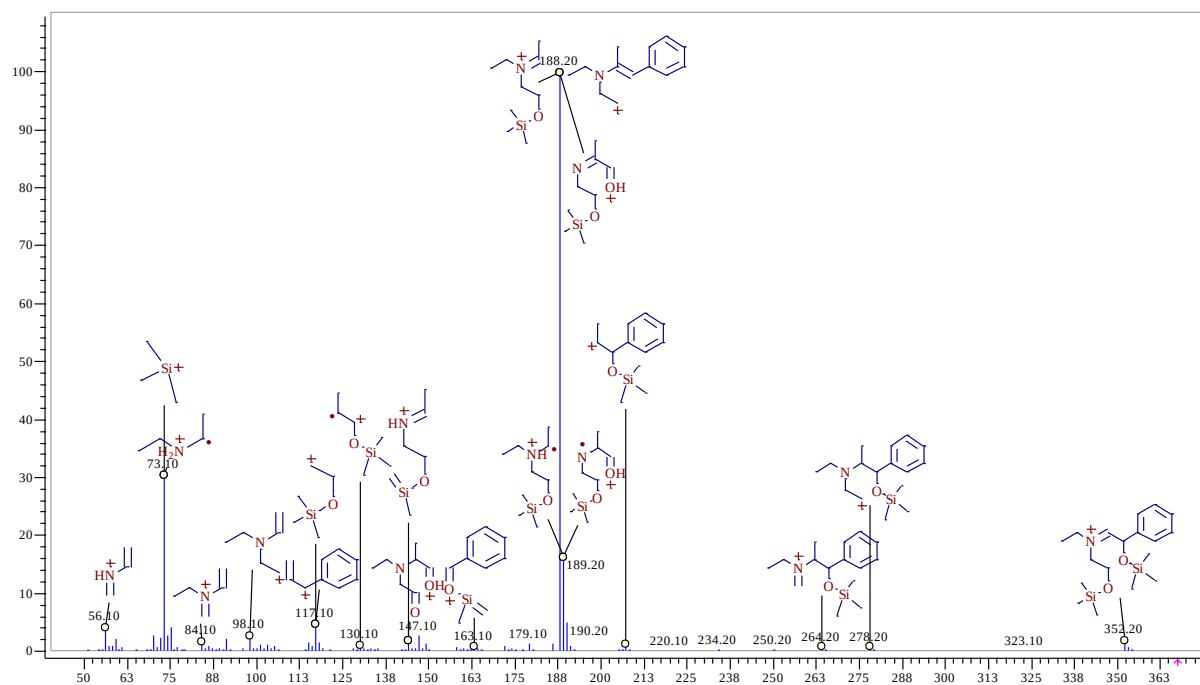


Mass spectrum interpretation by Thermo Scientific Mass Frontier (7.0) software

MS interpretation - TMS derivative

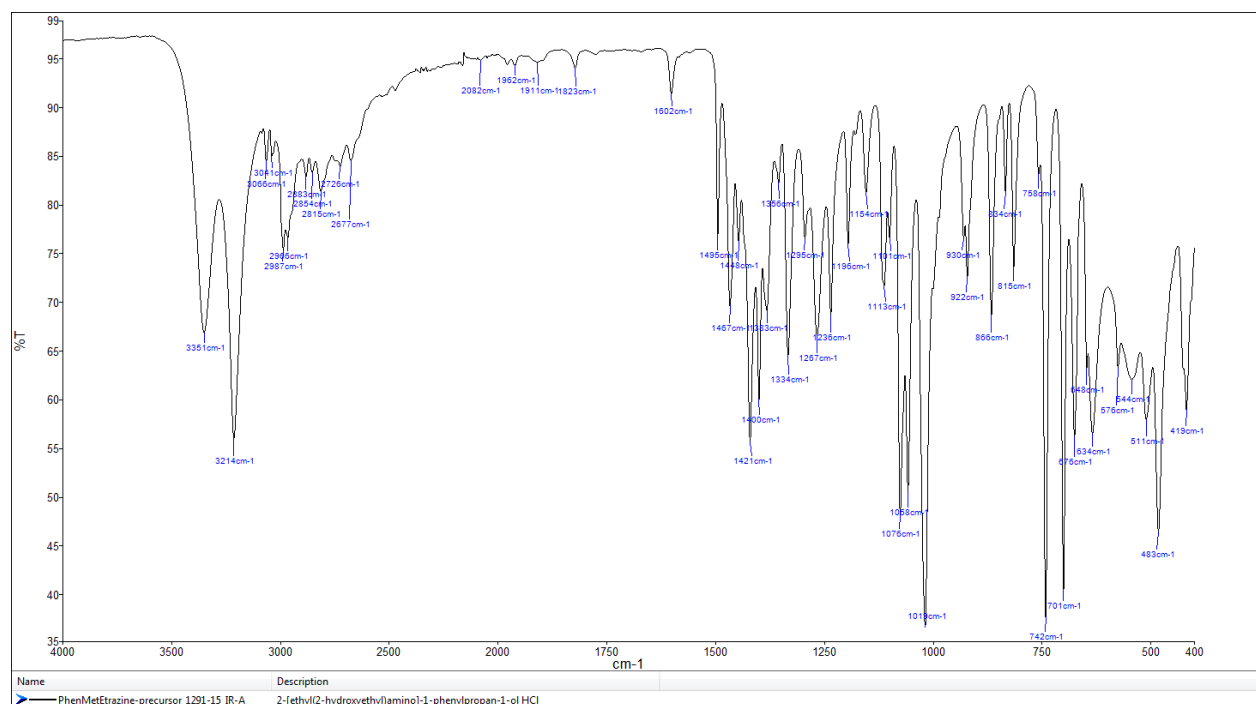


Mass spectrum - observed (measured)

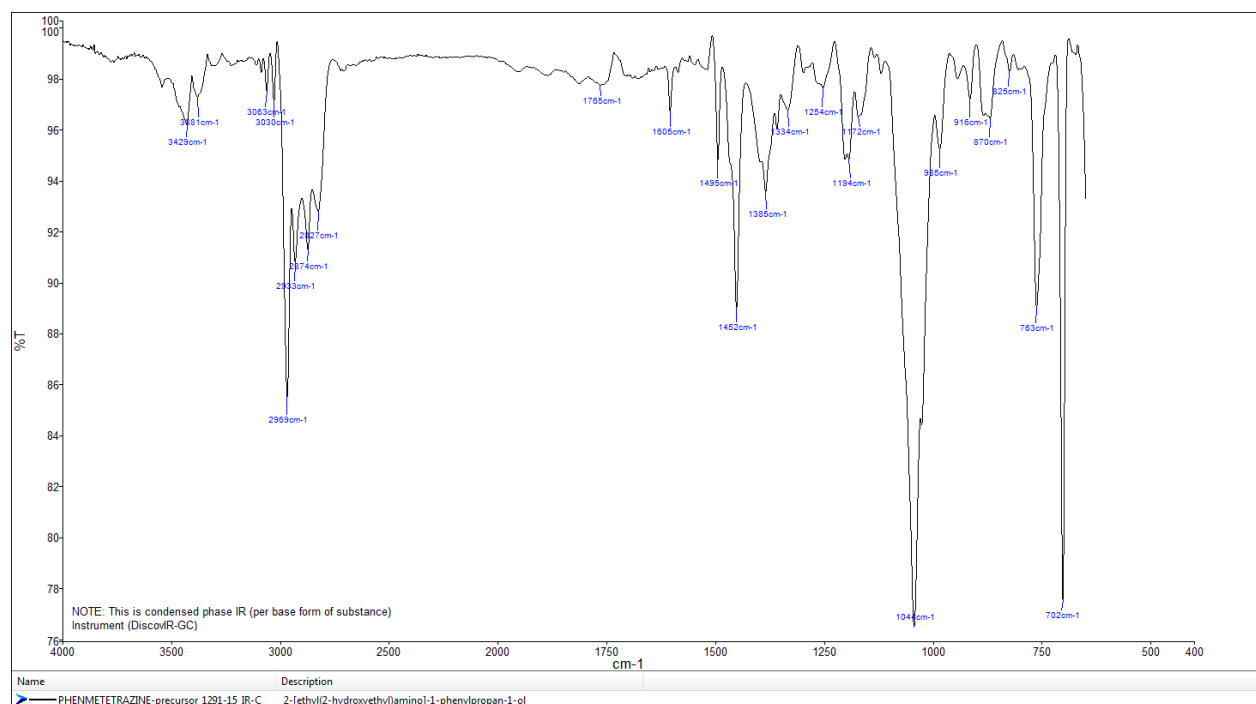


Mass spectrum interpretation by Thermo Scientific Mass Frontier (7.0) software

FTIR-ATR - direct measurement (sample as received)



IR (condensed phase – after chromatographic separation)



TOF REPORT

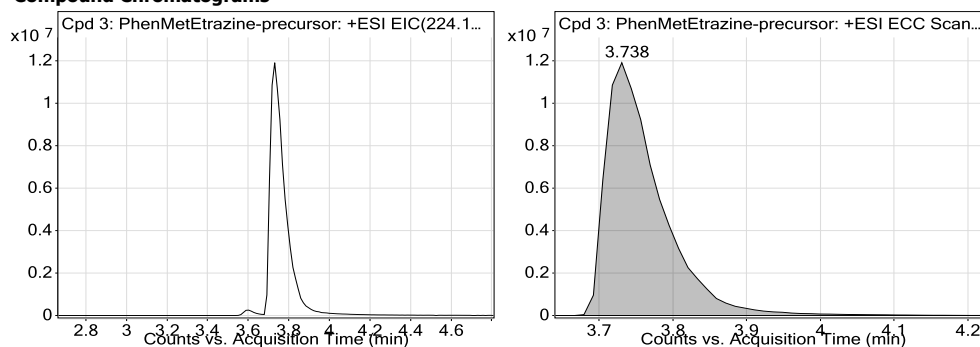
Data File	PhenMetEtrazine-precursor_1291-15_TOF.d	Sample Name	PhenMetEtrazine
Sample Type	Sample	Position	P2-B6
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-28052015-XDB-C18-ESI-poz.m	Acquired Time	9/28/2015 1:04:34 PM
IRM Calibration Status	Success	DA Method	Drugs_NFL.m
Comment	extract in MeOH		

Compound Table

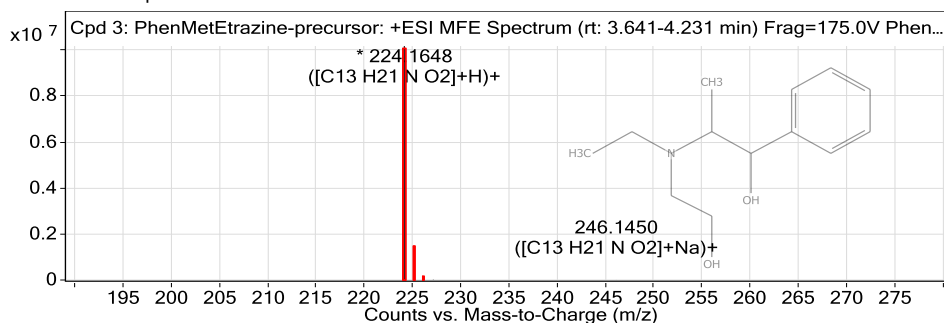
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 3: PhenMetEtrazine-precursor	PhenMetEtrazine-precursor	C13 H21 N O2	3.738	223.1575

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
PhenMetEtrazine-precursor	224.1648	3.738	223.1575	3.74	C13 H21 N O2	223.1572	-1.41

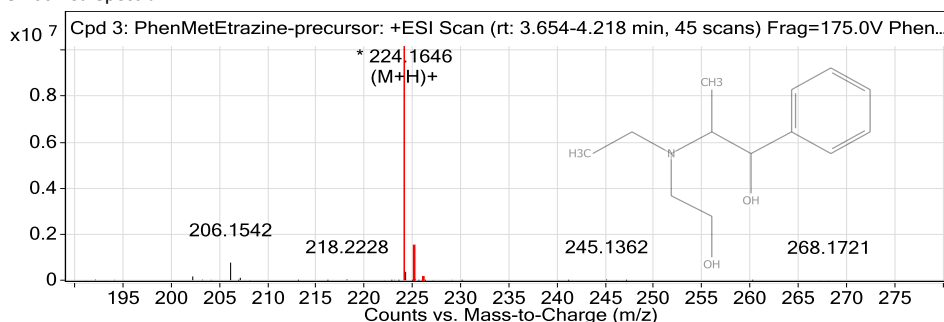
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
224.1648	1	10169786	C13 H21 N O2	(M+H)+
225.1684	1	1389628.44	C13 H21 N O2	(M+H)+
226.1709	1	124739.23	C13 H21 N O2	(M+H)+
227.1734	1	10269.39	C13 H21 N O2	(M+H)+
246.145	1	1593.94	C13 H21 N O2	(M+Na)+

--- End Of Report ---

TOF REPORT

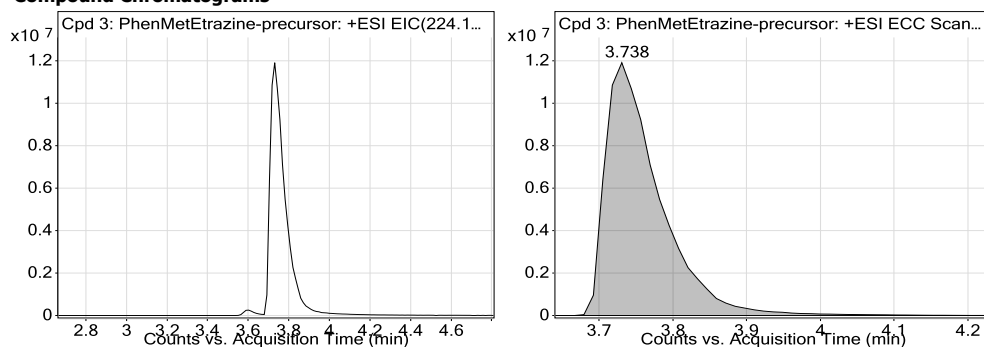
Data File	PhenMetEtrazine-precursor_1291-15_TOF.d	Sample Name	PhenMetEtrazine
Sample Type	Sample	Position	P2-B6
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-28052015-XDB-C18-ESI-poz.m	Acquired Time	9/28/2015 1:04:34 PM
IRM Calibration Status	Success	DA Method	Drugs_NFL.m
Comment	extract in MeOH		

Compound Table

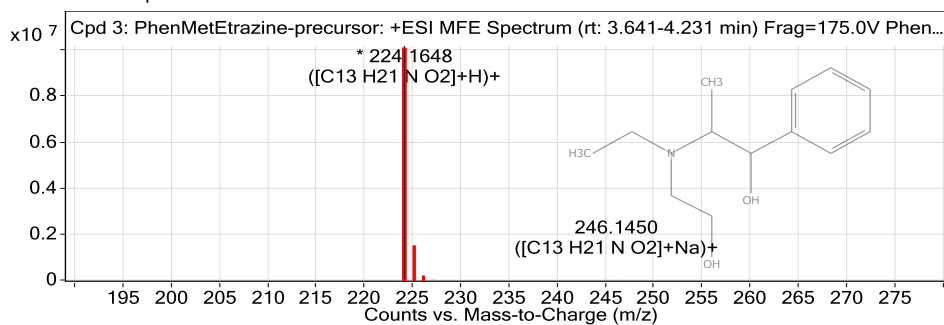
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 3: PhenMetEtrazine-precursor	PhenMetEtrazine-precursor	C13 H21 N O2	3.738	223.1575

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
PhenMetEtrazine-precursor	224.1648	3.738	223.1575	3.74	C13 H21 N O2	223.1572	-1.41

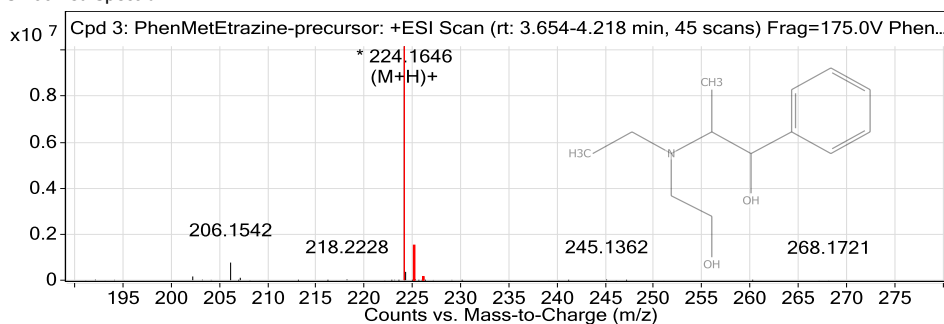
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

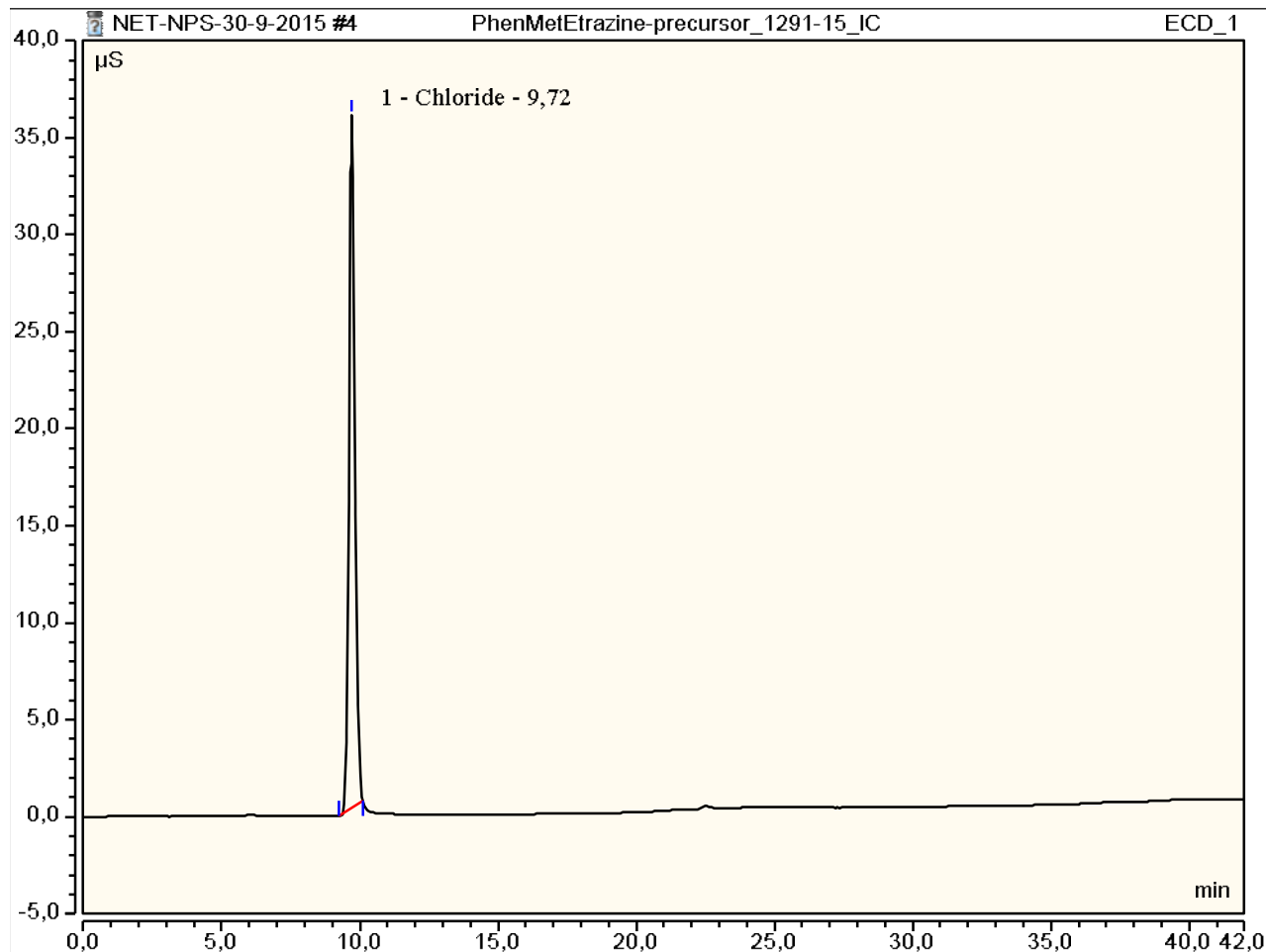
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
224.1648	1	10169786	C13 H21 N O2	(M+H)+
225.1684	1	1389628.44	C13 H21 N O2	(M+H)+
226.1709	1	124739.23	C13 H21 N O2	(M+H)+
227.1734	1	10269.39	C13 H21 N O2	(M+H)+
246.145	1	1593.94	C13 H21 N O2	(M+Na)+

--- End Of Report ---

Peak Integration Report

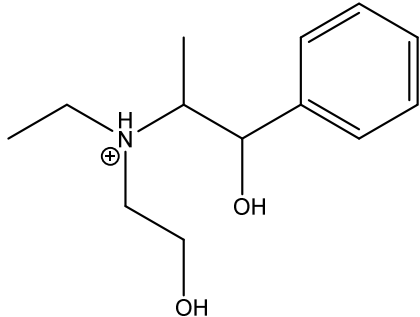
Sample Name:	PhenMetEtrazine-precursor_1291-15_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	30-sep-2015 / 11:26	Run Time:	41,98

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}\cdot\text{min}$	Height μS	Amount mg/L
1,00	9,72	Chloride	BMB	8,67	35,71	n.a.
		TOTAL:		8,67	35,71	0,00





REPORT

Sample ID:	1291-15
Our notebook code:	P-1291-15
NMR sample preparation:	15 mg dissolved in 0.7 mL DMSO- d_6
NMR experiments:	^1H , ^{13}C .
Proposed structure:	
Chemical name:	N-ethyl-1-hydroxy-N-(2-hydroxyethyl)-1-phenylpropan-2-aminium cation
Comments:	- Structure elucidation based on 1D NMR spectra - Sample is a mixture as evident by NMR, it contains another isomer (possibly diastereoisomer or other similar compound) in similar amounts - as obvious from the ^{13}C NMR (where most of the signals are doubled) - and additionally probably also a third compound (evident in ^1H NMR spectrum with redundant signals at 3.8, 3.5 and 3.1).
Supporting information:	Copies of ^1H and ^{13}C NMR spectra
Author:	Prof. Dr. Janez Košmrlj, Doc. Dr. Krištof Kranjc
Date of report:	March 9, 2016

P-1291-15

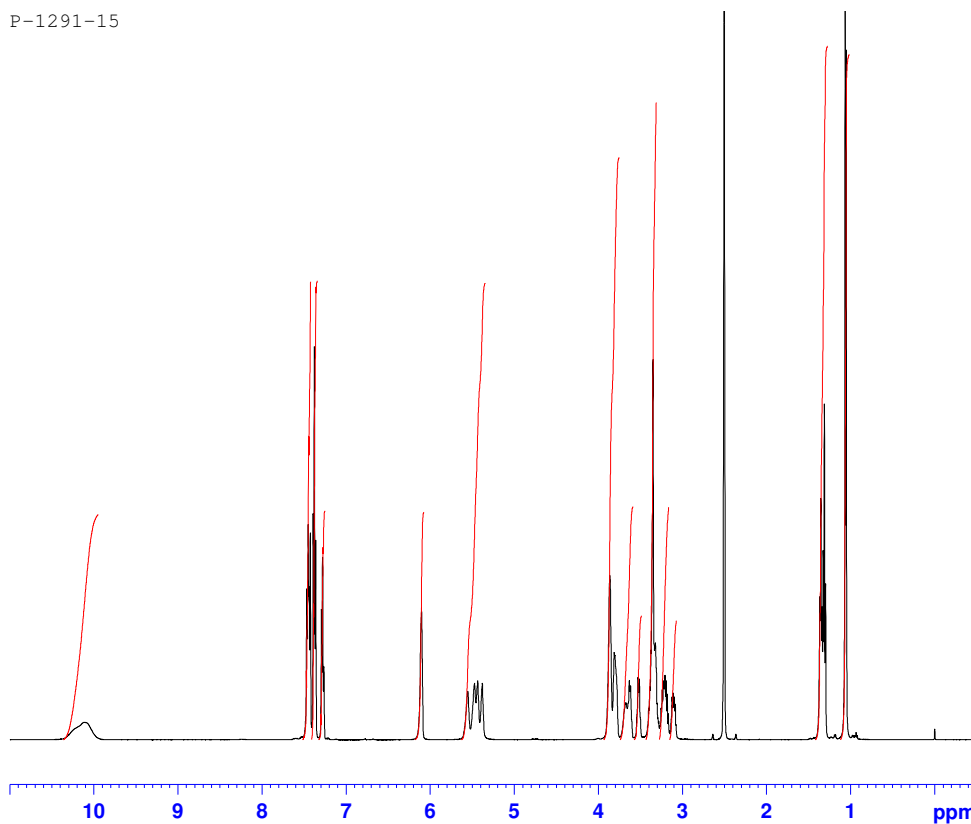


Current Data Parameters
NAME P-1291-15
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20151129
Time 7.26
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 10000.000 Hz
FIDRES 0.152588 Hz
AQ 3.2768500 sec
RG 71.8
DW 50.000 usec
DE 6.50 usec
TE 296.0 K
D1 1.00000000 sec
TD0 1

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

F2 - Processing parameters
SI 65536
SF 500.1300027 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



P-1291-15



Current Data Parameters
NAME P-1291-15
EXPNO 2
PROCNO 1

F2 - Acquisition Parameters
Date_ 20151129
Time 9.55
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
SOLVENT DMSO
NS 4096
DS 4
SWH 29761.904 Hz
FIDRES 0.454131 Hz
AQ 1.1010048 sec
RG 1820
DW 16.800 usec
DE 6.50 usec
TE 296.0 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

