

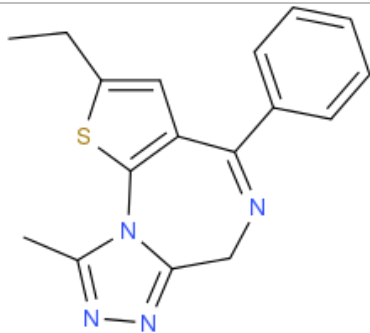
## ANALYTICAL REPORT

### Deschloroetizolam (C<sub>17</sub>H<sub>16</sub>N<sub>4</sub>S)

#### 2-ethyl-9-methyl-4-phenyl-6H-thieno[3,2-f][1,2,4]triazolo[4,3-a][1,4]diazepine

Remark – other NPS detected: **none**

Sample ID:	1174-15
Sample description:	tablet - pink
Sample type:	test purchase
Comments:	RESPONSE -purchasing
Date of entry:	4/17/2015

Substance identified-structure <sup>i</sup> (base form)	
Systematic name	2-ethyl-9-methyl-4-phenyl-6H-thieno[3,2-f][1,2,4]triazolo[4,3-a][1,4]diazepine
Other names	
Formula (per base form)	C <sub>17</sub> H <sub>16</sub> N <sub>4</sub> S
M <sub>w</sub> (g/mol)	308,4
Salt form	base
Smiles	<chem>C(C)C1=CC=2C(=NCC=3N(C2S1)C(=NN3)C)C3=CC=CC=C3</chem>
Compound Class	Benzodiazepines
Other NPS detected	none
Add.info (purity..)	

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): 13,08 BP(1): 279; BP(2): 308,BP(3) :239,
FTIR-ATR	+	dried CHCl <sub>3</sub> extract from tablet
FTIR (condensed phase) always as base form	-	
HPLC-TOF		Exact mass (theoretical): 308,1096; measured value $\Delta$ ppm:-2,43; formula:C <sub>17</sub> H <sub>16</sub> N <sub>4</sub> S
NMR-confirmed	+	partialy
validation		MS spectrum consistent by the one published in EMCDDA EDND database and in the literature: <a href="http://link.springer.com/article/10.1007/s11419-015-0277-6#page-1">http://link.springer.com/article/10.1007/s11419-015-0277-6#page-1</a>
other		

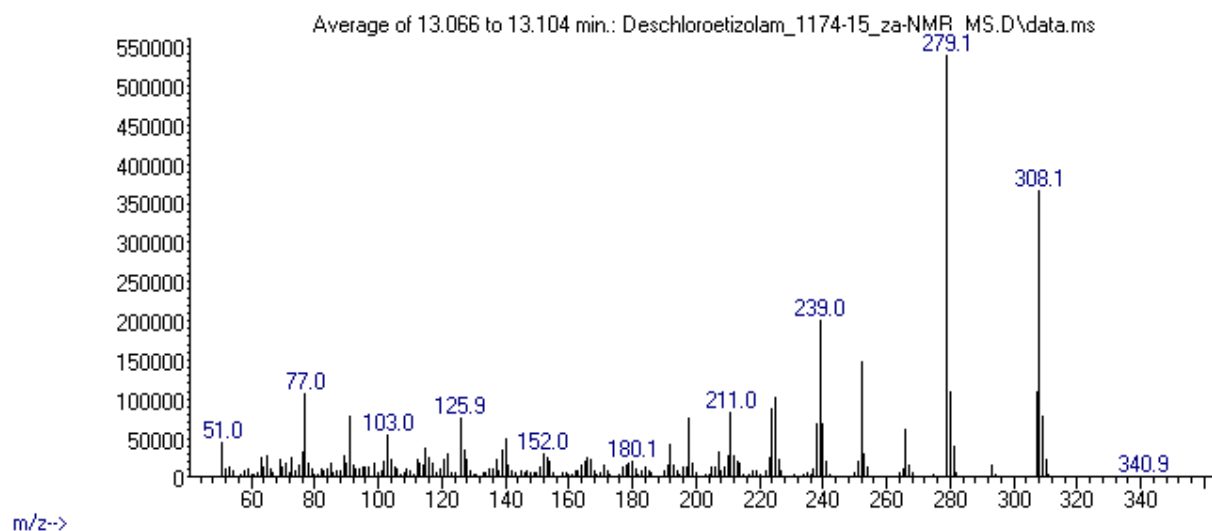
GC-MS (Agilent): GC-method is RT locked to tetracosane (RT=9.53 min). Injection volume 1 ml and split mode (1:50)for GC-MS instruments and 1:5 for GC-MS-FTIR(condensed phase). Injector temperature: 280 °C. Column: HP1-MS (100% dimethylpolysiloxane), length 30 m, internal diameter 0.25 mm, film thickness 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>

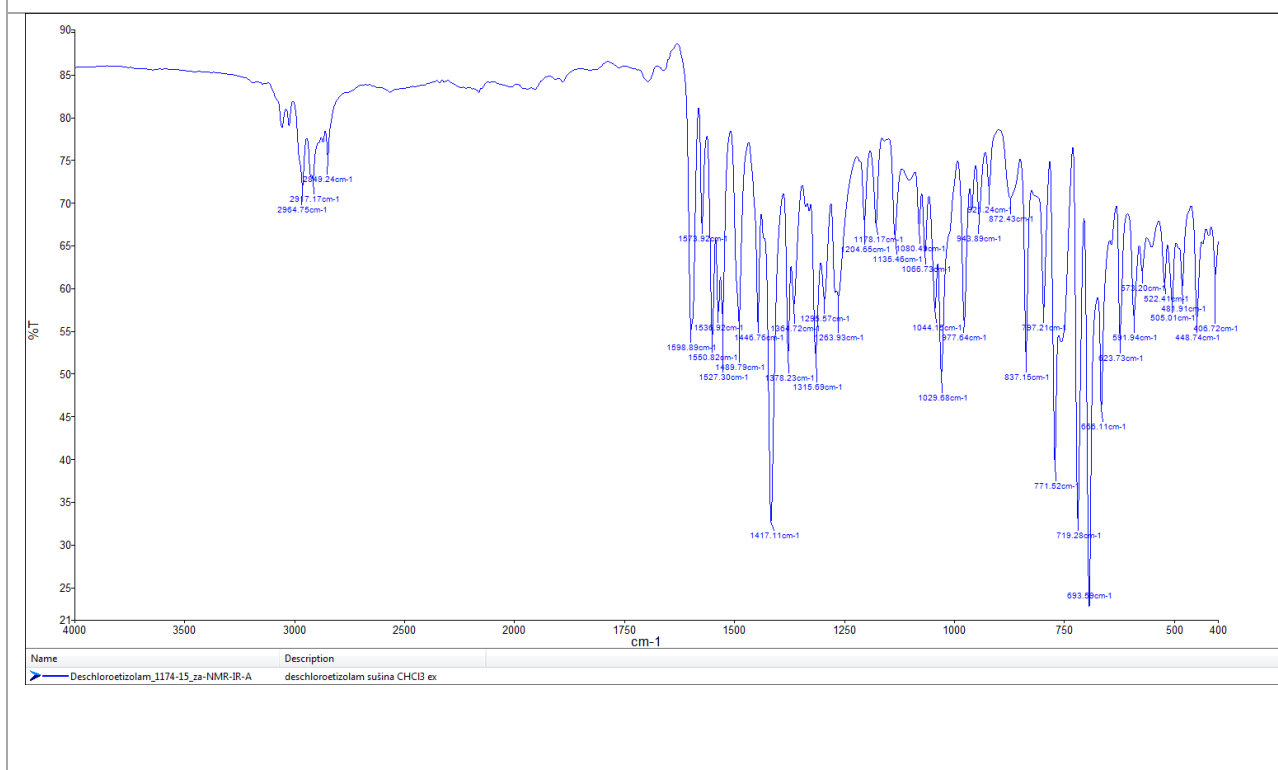
FTIR (Spectra analyses-Danny): scan range 4000 to 600, resolution 4cm<sup>-1</sup>

## MS spectrum (EI)

Abundance



## FTIR - ATR



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

# Target Compound Screening Report

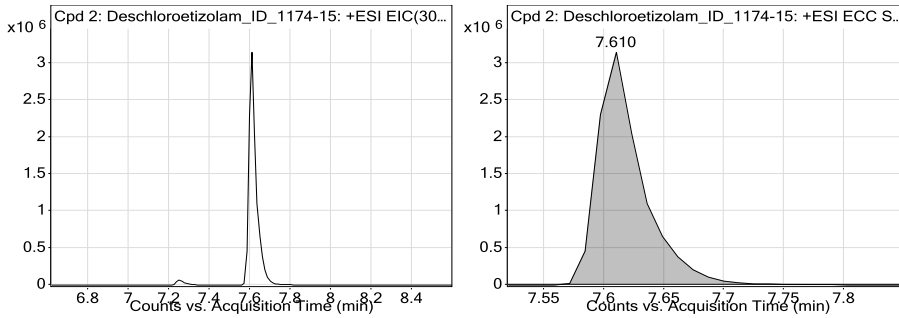
<b>Data File</b>	Deschloroetizolam_ID_1174-15_TOF.d	<b>Sample Name</b>	Deschloroetizolam
<b>Sample Type</b>	Sample	<b>Position</b>	P1-F7
<b>Instrument Name</b>	SG13170002	<b>User Name</b>	
<b>Acq Method</b>	droge general-2-4-2015-XDB-C18-ESI-poz.m	<b>Acquired Time</b>	4/10/2015 10:03:30 AM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Droge_Default.m
<b>Comment</b>	red tablet, extract in MeOH		

## Compound Table

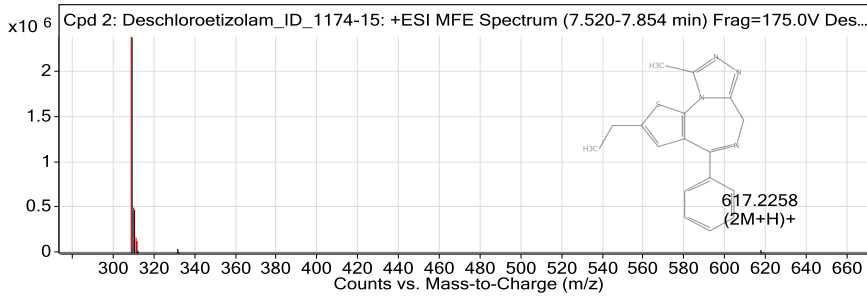
Label	Tgt Name	MFG Formula	Tgt Formula	Obs. RT	Obs. Mass
Cpd 2: Deschloroetizolam_ID_1174-15	Deschloroetizolam_ID_1174-15	C17 H16 N4 S	C17 H16 N4 S	7.61	308.1103

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error	Tgt Formula	Find Cpds Algorith
<b>Deschloroetizolam_ID_1174-15</b>	309.1177	7.61	308.1103	7.61	C17 H16 N4 S	308.1096	-2.43	C17 H16 N4 S	Find by Molecular Feature

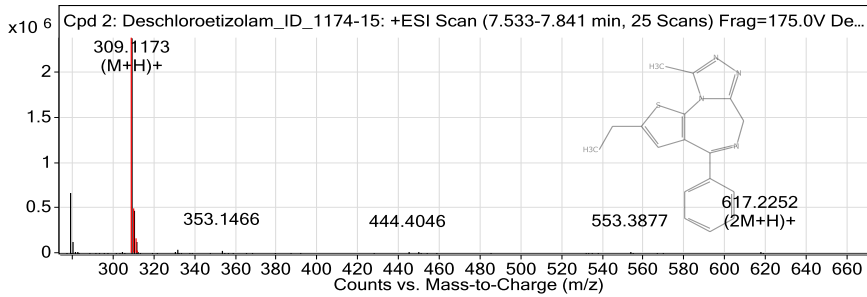
## Compound Chromatograms



## MFE MS Zoomed Spectrum



## MS Zoomed Spectrum



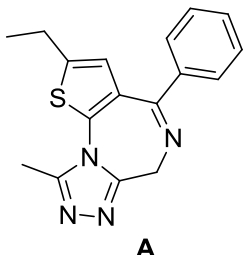
## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
309.1177	1	2386072	C17 H16 N4 S	(M+H)+
310.1199	1	481698.52	C17 H16 N4 S	(M+H)+
311.116	1	141143	C17 H16 N4 S	(M+H)+
312.1171	1	22869.43	C17 H16 N4 S	(M+H)+
331.0991	1	45489.21		(M+Na)+
332.1025	1	10623.43		(M+Na)+
333.0947	1	3885.45		(M+Na)+
617.2258	1	34732.21		(2M+H)+
618.2278	1	13298.98		(2M+H)+
619.2266	1	5903.71		(2M+H)+

--- End Of Report ---



## REPORT

Sample ID:	<b>1174-15</b>
Our notebook code:	P-1174-15
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, <sup>1</sup> H- <sup>15</sup> N <i>gs</i> -HMBC.
Proposed structure with chemical name:	
Comments:	<p>- Two resonances of impurities are detected in <sup>1</sup>H NMR (0.08 ppm and 1.18 ppm) and <sup>13</sup>C NMR (-1.02 ppm and 28.67 ppm) (see the supporting information).</p> <p>- 1D and 2D NMR spectra confirm some parts of the suggested structure (<b>A</b>):</p> <div style="text-align: center;">  <p><b>A</b></p> </div> <p>i.e.: a) the appearance of <sup>1</sup>H NMR with the corresponding integration, and b) the number of <sup>13</sup>C NMR resonances with the corresponding chemical shifts are both in agreement with structure <b>A</b>.</p> <p>The lack of protons in the fused ring system and thus the lack of connectivity pattern in 2D NMR spectra prevent unambiguous structure elucidation.</p>
Supporting information:	Copies of <sup>1</sup> H and <sup>13</sup> C NMR spectra in CDCl <sub>3</sub>
Author:	Prof. Dr. Janez Košmrlj
Date of report:	June 20, 2015

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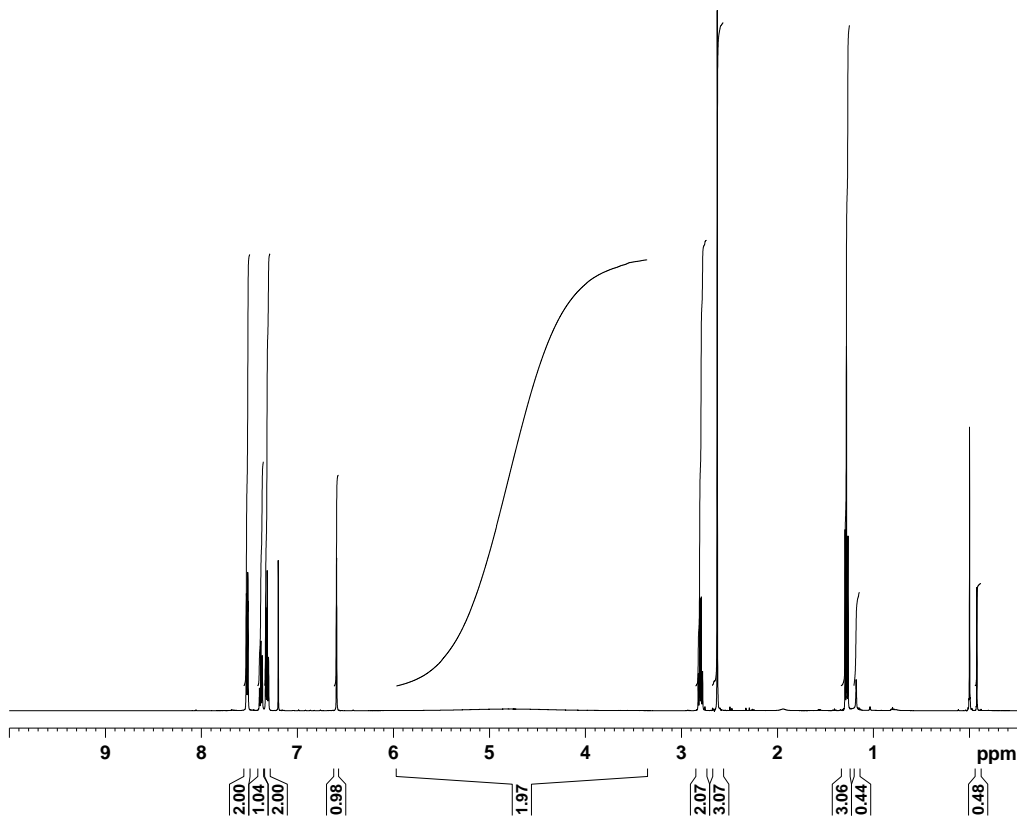


Current Data Parameters  
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 EXFNO 1  
 PROCNO 1

F2 - Acquisition Parameters  
 Date 20150424  
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 PROBHD 5 mm PABBO BB-  
 PULPROG zg30  
 TD 65536  
 SOLVENT CDCl3  
 NS 16  
 DS 2  
 SWH 10330.578 Hz  
 FIDRES 0.157632 Hz  
 AQ 3.1719923 sec  
 RG 71.8  
 DW 48.400 usec  
 DE 6.50 usec  
 TE 296.0 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 500.1300433 MHz  
 WDW EM  
 SSB 0  
 LB 0.30 Hz  
 GB 0  
 PC 1.00



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Current Data Parameters  
 NAME P-DESCHLOROETIZOLAM  
 EXFNO 2  
 PROCNO 1

F2 - Acquisition Parameters  
 Date 20150424  
 Time 8.48  
 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 1.1010548 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  
 SFO1 125.7703637 MHz

===== CHANNEL f2 =====  
 CPDPRG2 waltz16  
 NUC2 1H  
 FCPD2 80.00 usec  
 PLW2 26.00000000 W  
 PLW12 0.32179001 W  
 PLW13 0.20595001 W  
 SFO2 500.1320005 MHz

F2 - Processing parameters  
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 SF 125.7579188 MHz  
 WDW EM  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

