



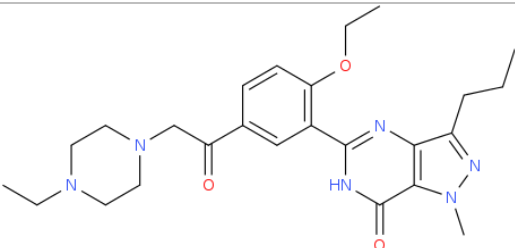
## ANALYTICAL REPORT<sup>1</sup>

### acetildenafilil (C25H34N6O3)

#### 5-{2-ethoxy-5-[2-(4-ethylpiperazin-1-yl)acetyl]phenyl}-1-methyl-3-propyl-1H,6H,7H-pyrazolo[4,3-d]pyrimidin-7-one

Remark – other NPS detected: **none**

Sample ID:	1542-16
Sample description:	powder - brown-yellowish
Sample type:	test purchase /RESPONSE -purchasing
Date of sample receipt (M/D/Y):	3/9/2016
Date of entry (M/D/Y) into NFL database:	3/17/2016
Report updates (if any) will be published here:	<a href="http://www.policija.si/apps/nfl_response_web/seznam.php">http://www.policija.si/apps/nfl_response_web/seznam.php</a>

Substance identified - structure <sup>2</sup> (base form)	
Systematic name	5-{2-ethoxy-5-[2-(4-ethylpiperazin-1-yl)acetyl]phenyl}-1-methyl-3-propyl-1H,6H,7H-pyrazolo[4,3-d]pyrimidin-7-one
Other names	HONGDENAFIL;
Formula (per base form)	C25H34N6O3
M <sub>w</sub> (g/mol)	466,59
Salt form/anions detected	base
StdInChIKey	RRBRQNALHKQCAI-UHFFFAOYSA-N
Compound Class	Piperazine derivates
Other NPS detected	none
Add.info (purity..)	pure by GC-MS not pure, TOF: 10% C27H36N6O5 Mw=524,3

<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)

### 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 quadropole 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 (N<sub>2</sub>) 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 (30 until 6 min.) to 550 (300) amu.

IR (condensed phase): IR scan range 4000 to 650, resolution 4 cm<sup>-1</sup>.

**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

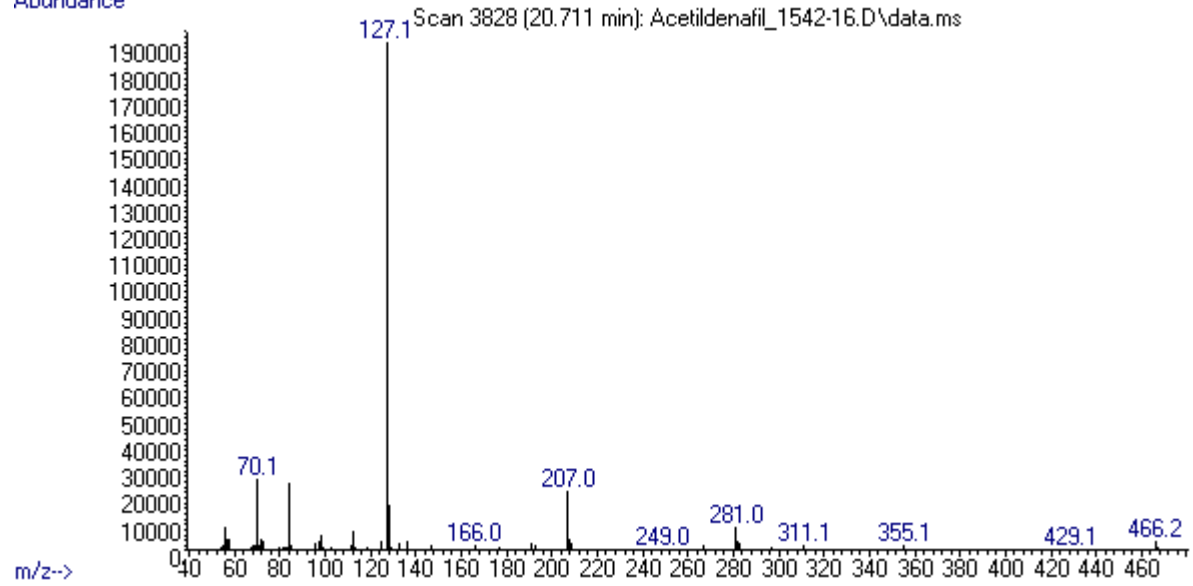
Solubility in	result/remark
CH <sub>2</sub> Cl <sub>2</sub>	soluble
MeOH	soluble
H <sub>2</sub> O	partially

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 20,72 BP(1): 127; BP(2): 70,BP(3) :84,
HPLC-TOF	+	Exact mass (theoretical): 466,2629; measured value Δppm:-0,02; formula:C <sub>25</sub> H <sub>34</sub> N <sub>6</sub> O <sub>3</sub>
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	+	
IC (anions)	-	
NMR (in FKKT)	-	
validation		
other		Confirmed by Cayman and SWGDRUG MS libraries.

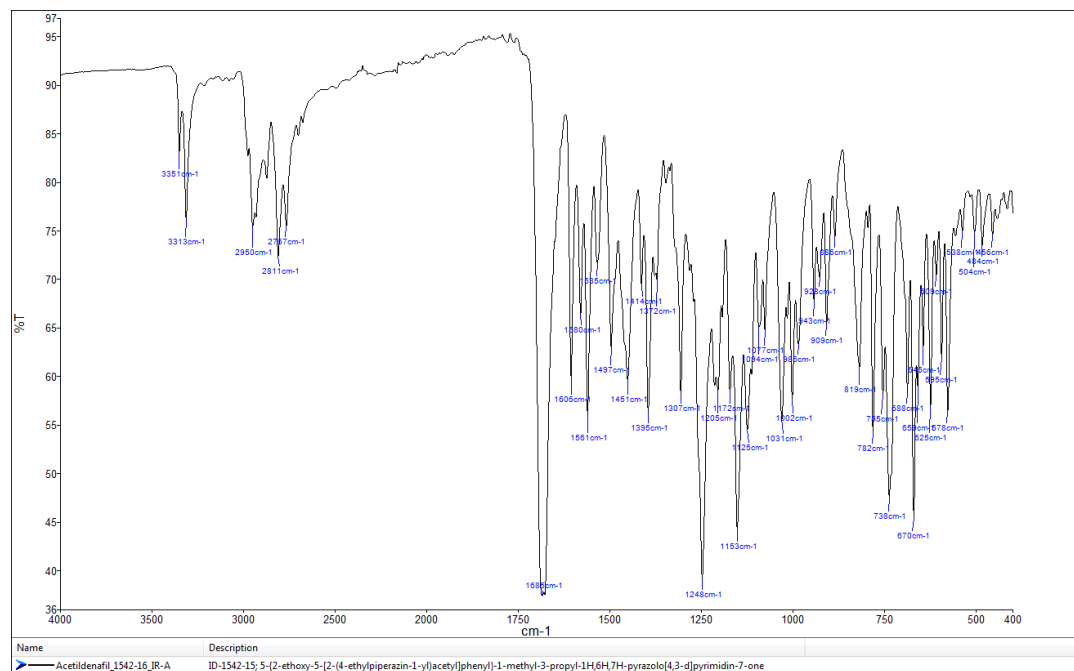
# ANALYTICAL RESULTS

MS (EI)

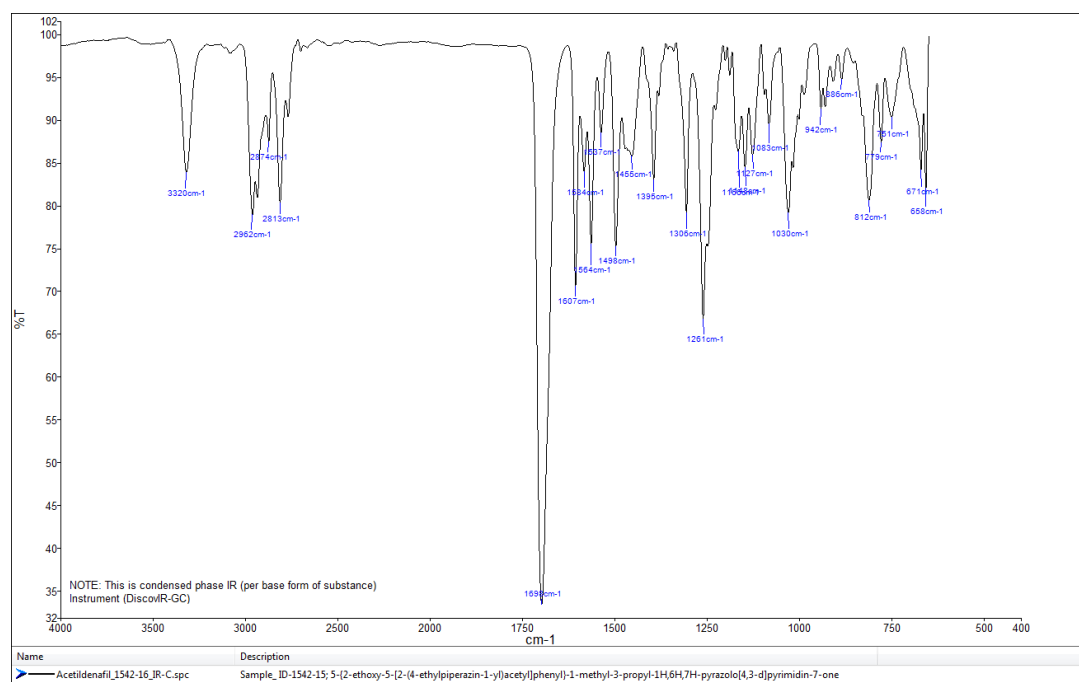
Abundance



## FTIR-ATR - direct measurement (sample as received)



## IR (condensed phase – after chromatographic separation)



# TOF REPORT

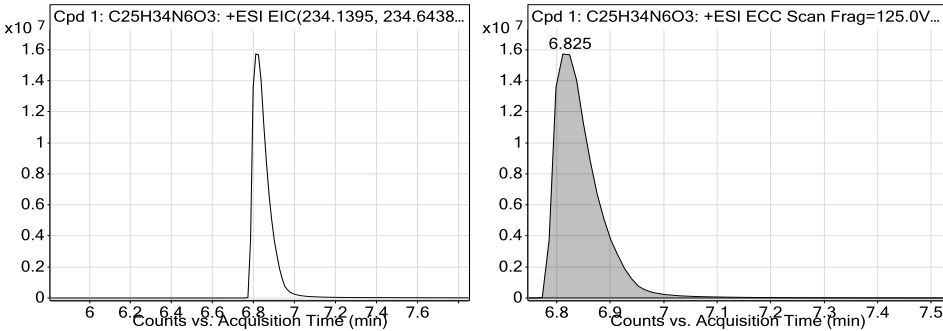
<b>Data File</b>	Acetildenafil-1542-16_TOF.d	<b>Sample Name</b>	ID_1542-16
<b>Sample Type</b>	Sample	<b>Position</b>	P1-E5
<b>Instrument Name</b>	6230B TOF LC-MS	<b>User Name</b>	TG
<b>Acq Method</b>	general-1512015-XDB-C18-ESI-poz-pod.m	<b>Acquired Time</b>	3/13/2016 11:08:59 AM
<b>IRM Calibration Status</b>	Success	<b>DA Method</b>	Drugs_NFL.m
<b>Comment</b>	extract in MeOH		

**Compound Table**

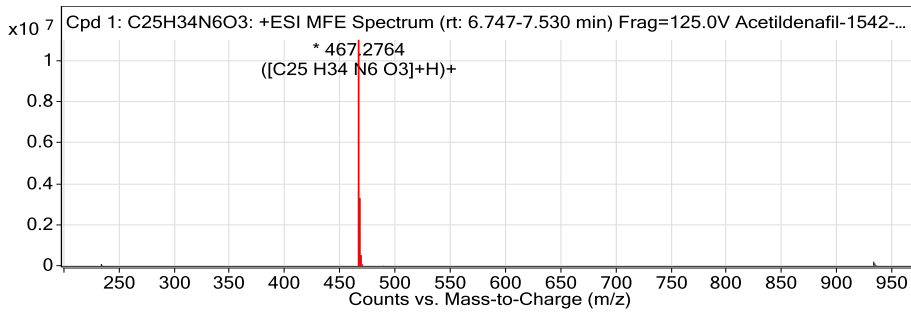
Label	Compound Name	Obs. RT	Obs. Mass
Cpd 1: C25H34N6O3	C25H34N6O3	6.825	466.2692
Cpd 2: C27H36N6O5	C27H36N6O5	6.965	524.2748

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
<b>C25H34N6O3</b>	467.2764	6.825	466.2692	6.83	C25 H34 N6 O3	466.2692	-0.02

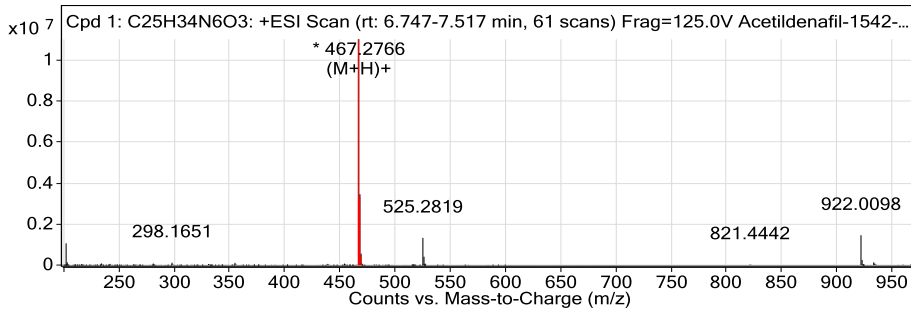
**Compound Chromatograms**



**MFE MS Zoomed Spectrum**



**MS Zoomed Spectrum**



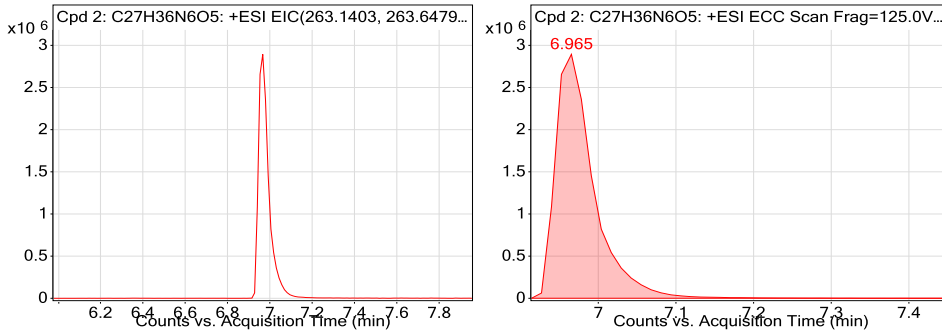
**MS Spectrum Peak List**

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
234.143	2	82664.06		(M+2H)+2
234.6442	2	27909.28		(M+2H)+2
467.2764	1	11007167	C25 H34 N6 O3	(M+H)+
468.2797	1	3297568.22	C25 H34 N6 O3	(M+H)+
469.2831	1	514451.74	C25 H34 N6 O3	(M+H)+
470.2849	1	57804.95	C25 H34 N6 O3	(M+H)+
489.2583	1	10039.62		(M+Na)+
933.546	1	200111.52		(2M+H)+
934.5487	1	110892.42		(2M+H)+
935.5506	1	31782.99		(2M+H)+

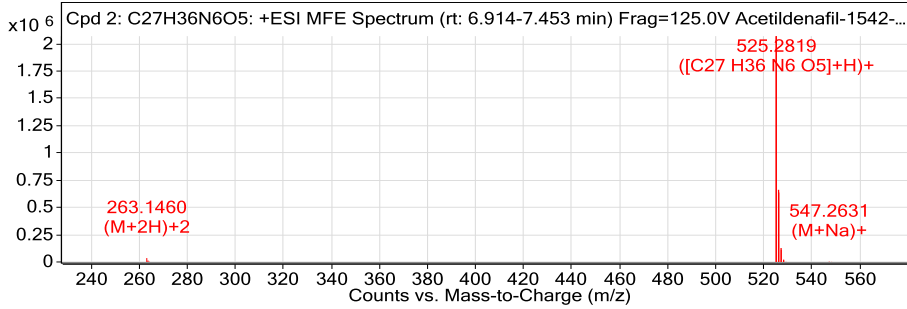
Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
<b>C27H36N6O5</b>	525.2819	6.965	524.2748	6.97	C27 H36 N6 O5	524.2747	-0.22

# TOF REPORT

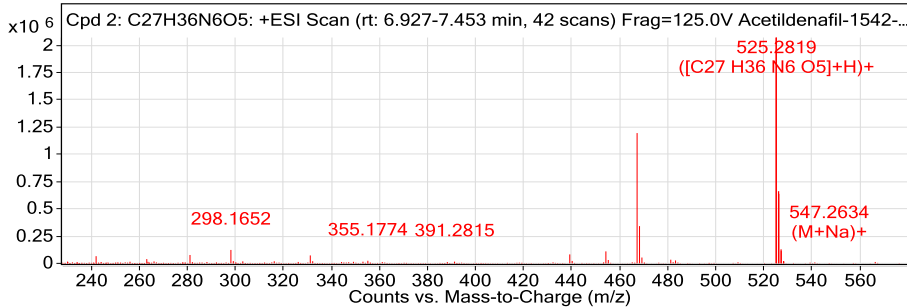
## Compound Chromatograms



## MFE MS Zoomed Spectrum



## MS Zoomed Spectrum



## MS Spectrum Peak List

Obs. m/z	Charge	Abund	Formula	Ion/Isotope
263.146	2	35610.11		(M+2H)+2
263.6468	2	13324.66		(M+2H)+2
264.1489	2	3130.62		(M+2H)+2
525.2819	1	2070495.5	C27 H36 N6 O5	(M+H)+
526.2858	1	637237.15	C27 H36 N6 O5	(M+H)+
527.2878	1	111810.69	C27 H36 N6 O5	(M+H)+
528.2894	1	14260.5	C27 H36 N6 O5	(M+H)+
529.2911	1	1450.93	C27 H36 N6 O5	(M+H)+
547.2631	1	3142.35		(M+Na)+
548.2681	1	1019.06		(M+Na)+

--- End Of Report ---