



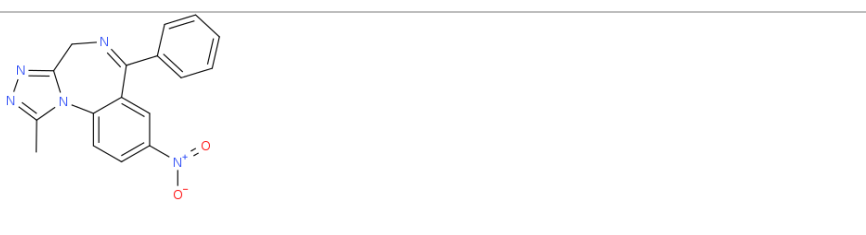
ANALYTICAL REPORT^{1,2}

Nitrazolam (C₁₇H₁₃N₅O₂)

3-methyl-12-nitro-9-phenyl-2,4,5,8-tetraazatricyclo[8.4.0.0.2,6]tetradeca-1(14),3,5,8,10,12-hexaene

Remark – other NPS detected: **none**

Sample ID:	1452-16
Sample description:	powder - yellow-bright
Sample type:	collected /Institute of Forensic medicine, University Freiburg, Germany
Date of sample receipt (M/D/Y):	1/14/2016
Date of entry (M/D/Y) into NFL database:	8/10/2016
Report updates (if any) will be published here:	http://www.policija.si/apps/nfl_response_web/seznam.php

Substance identified - structure ³ (base form)	
Systematic name	3-methyl-12-nitro-9-phenyl-2,4,5,8-tetraazatricyclo[8.4.0.0.2,6]tetradeca-1(14),3,5,8,10,12-hexaene
Other names	1-methyl-8-nitro-6-phenyl-4H-[1,2,4]triazolo[4,3-a][1,4]benzodiazepine
Formula (per base form)	C ₁₇ H ₁₃ N ₅ O ₂
M _w (g/mol)	319,32
Salt form/anions detected	base
StdInChIKey	OYRPNABWTHDOFK-UHFFFAOYSA-N
Compound Class	Benzodiazepines
Other NPS detected	none
Add.info (purity..)	pure by GC-MS, HPLC-tof

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 Acknowledgement: Sample (NMR confirmed) was kindly provided by the Institute of Forensic Medicine, University of Freiburg, Germany (Dr. Verena Angerer). NMR spectra were kindly provided by Dr. Bjoern Moosmann and are enclosed in this report by his permission. Other measurements shown in this report were done in NFL.

³ 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 (9.258 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 µm. Carrier gas He: flow-rate 1.2 ml/min. GC oven program: 170 °C for 1 min, followed by heating up to 190 °C at rate 8 °C/min, then 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 until 6 min) 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 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 (solid) 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

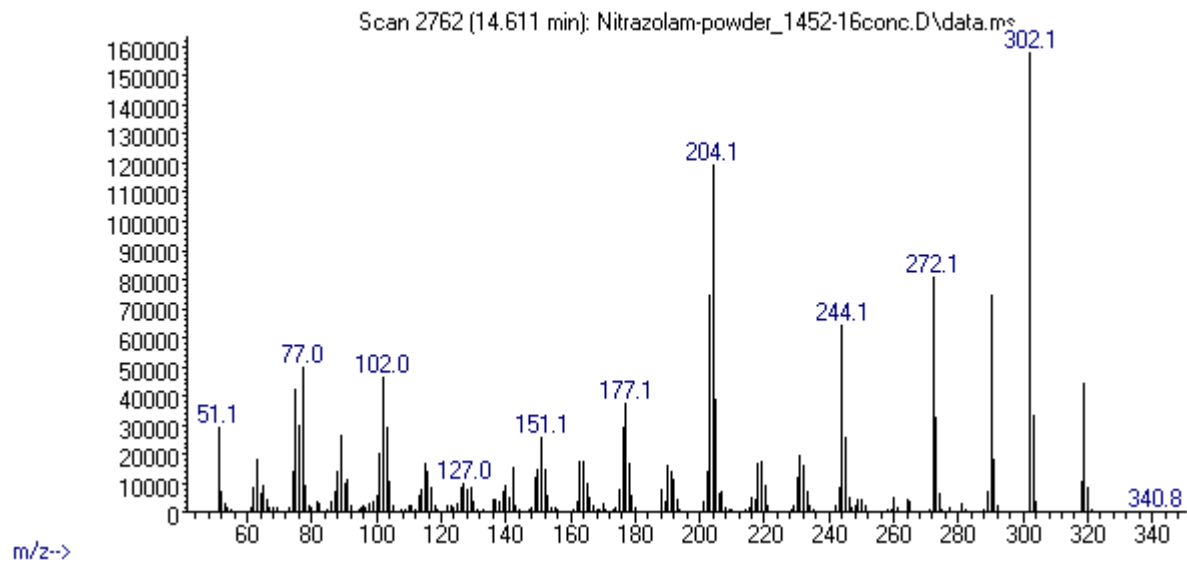
Solubility in	result/remark
CH ₂ Cl ₂	soluble
MeOH	soluble
H ₂ O	soluble

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 14,61 BP(1): 302; BP(2): 204,BP(3) :272,
HPLC-TOF	+	Exact mass (theoretical): 319,1069; measured value Δppm:-0,88; formula:C17H13N5O2
FTIR-ATR	+	direct measurement (sample as received)
FTIR (condensed phase) always as base form	-	
IC (anions)	+	
NMR	+	Kindly provided by Dr. Bjoern Moosmann, Institute of Forensic Medicine, University of Freiburg, Germany
validation		
other		

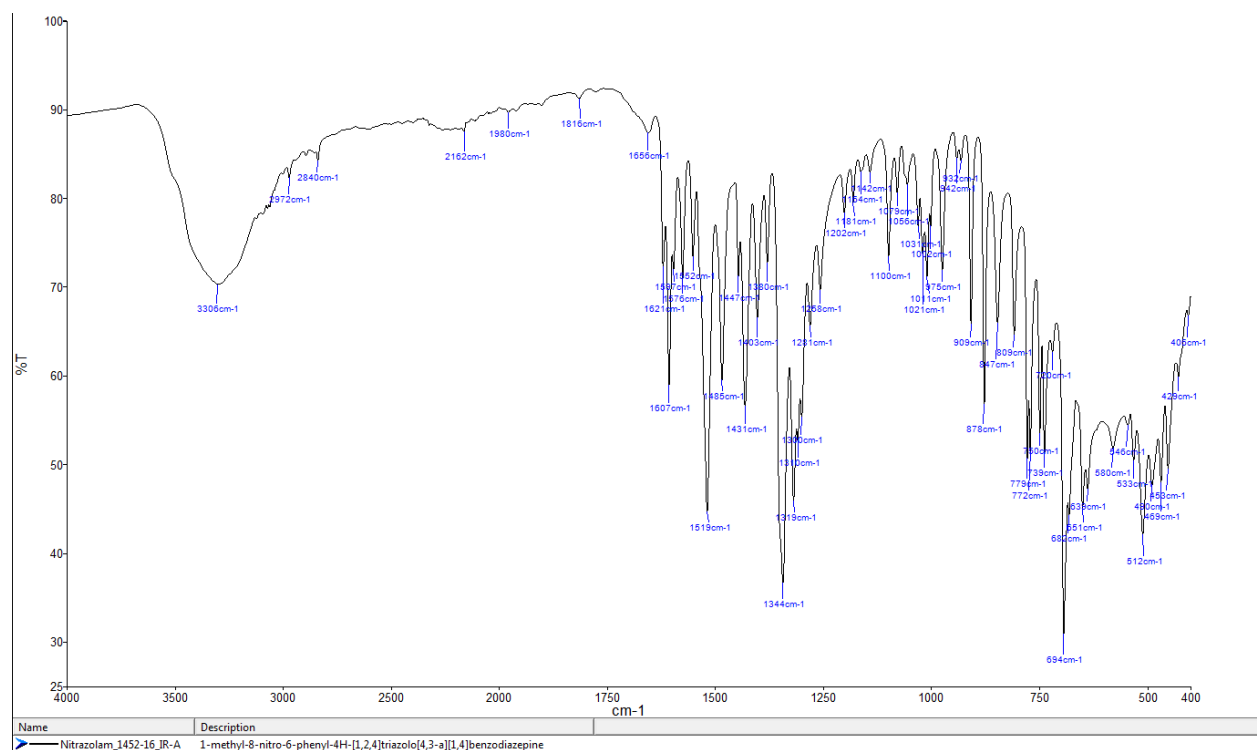
ANALYTICAL RESULTS

MS (EI)

Abundance



FTIR-ATR - direct measurement (sample as received)



TOF REPORT

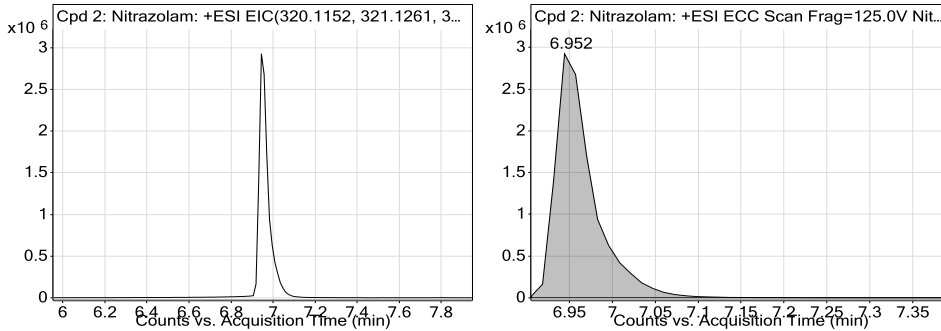
Data File	Nitrazolam_1452-16_TOF.d	Sample Name	ID_1452-16
Sample Type	Sample	Position	P1-F6
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-1512015-XDB-C18-ESI-poz-pod.m	Acquired Time	2/23/2016 12:02:20 PM
IRM Calibration Status	Success	DA Method	Drugs_NFL.m
Comment	extract in MeOH		

Compound Table

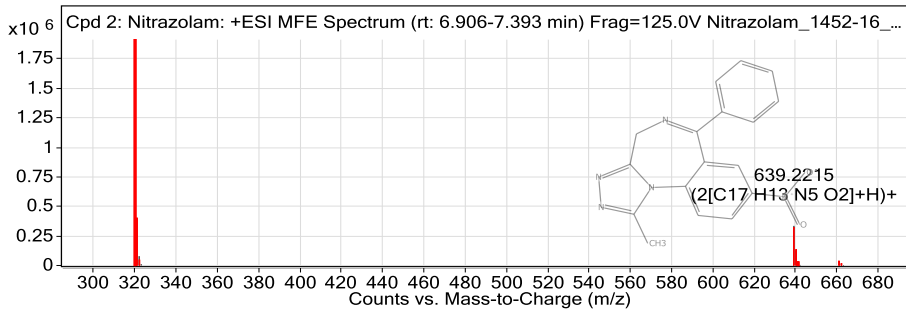
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 2: Nitrazolam	Nitrazolam	C17 H13 N5 O2	6.952	319.1072

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
Nitrazolam	320.1143	6.952	319.1072	6.95	C17 H13 N5 O2	319.1069	-0.88

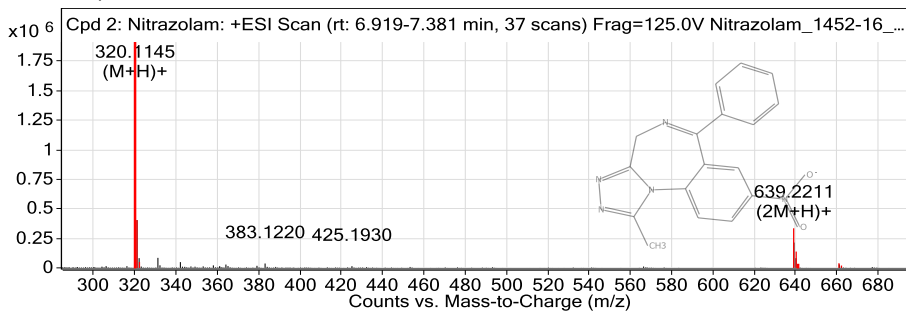
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

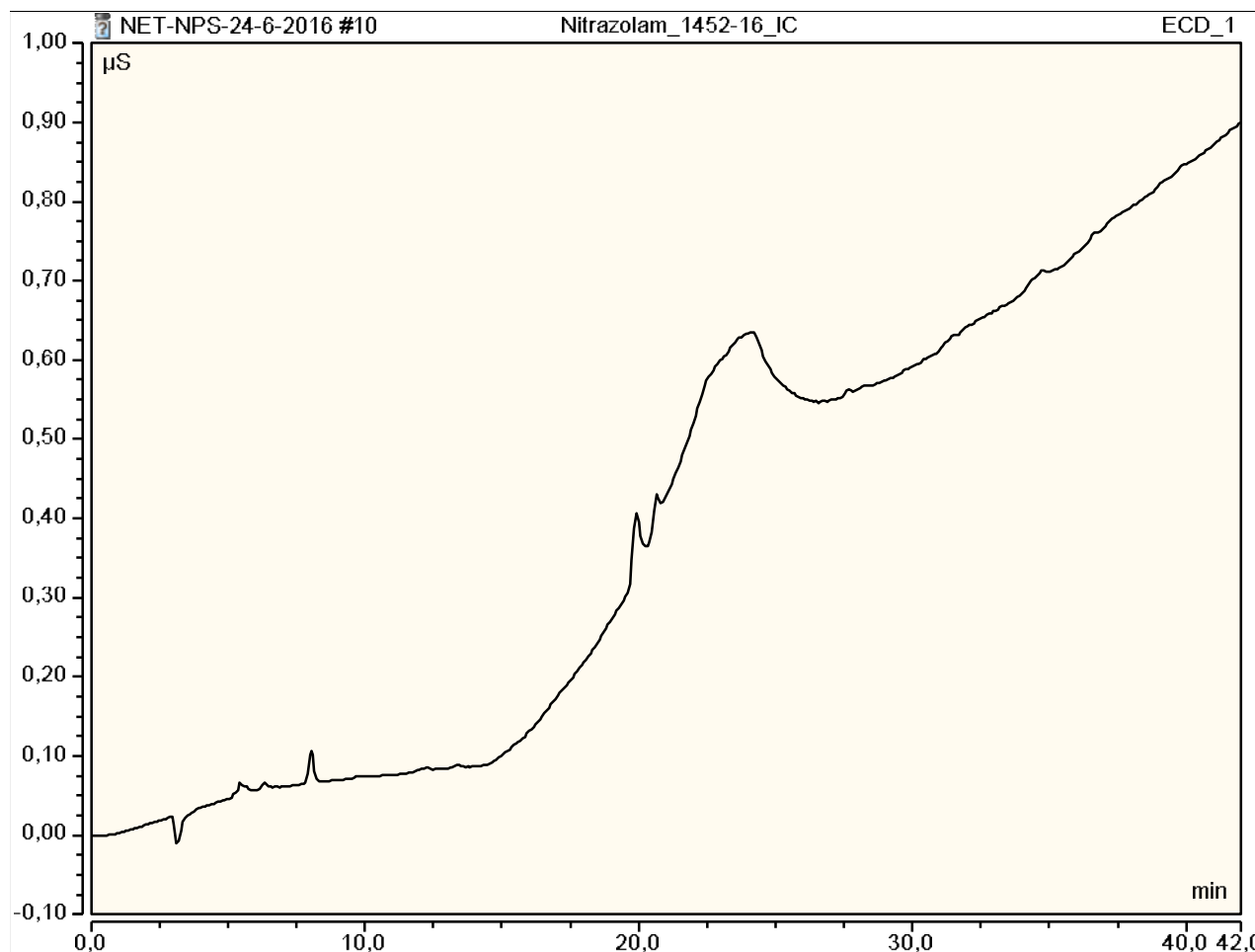
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
320.1143	1	1912323.25	C17 H13 N5 O2	(M+H)+
321.1179	1	400376.43	C17 H13 N5 O2	(M+H)+
322.124	1	79021.76	C17 H13 N5 O2	(M+H)+
323.1288	1	10668.9	C17 H13 N5 O2	(M+H)+
639.2215	1	335495.69	C17 H13 N5 O2	(2M+H)+
640.224	1	127864.77	C17 H13 N5 O2	(2M+H)+
641.2267	1	27965.17	C17 H13 N5 O2	(2M+H)+
642.2303	1	4548.14	C17 H13 N5 O2	(2M+H)+
661.2027	1	37709.03	C17 H13 N5 O2	(2M+Na)+
662.2052	1	15094.34	C17 H13 N5 O2	(2M+Na)+

--- End Of Report ---

Peak Integration Report

Sample Name:	Nitrazolam_1452-16_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	kemija
Inj. Date / Time:	24-jun-2016 / 18:30	Run Time:	42,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}\cdot\text{min}$	Height μS	Amount n.a.
		TOTAL:		0,00	0,00	0,00



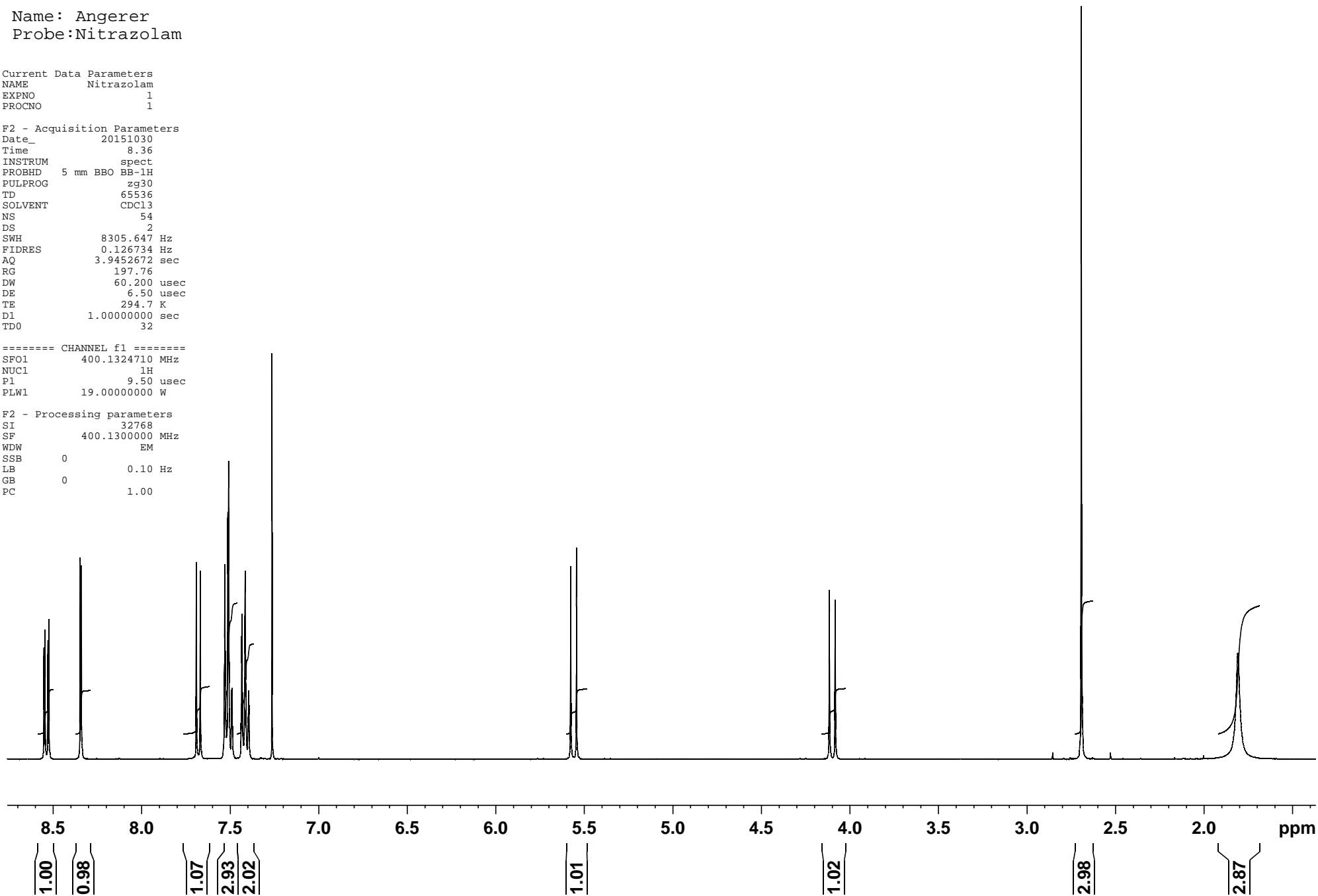
Name: Angerer
Probe: Nitrazolam

Current Data Parameters
NAME Nitrazolam
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20151030
Time 8.36
INSTRUM spect
PROBHD 5 mm BBO BB-1H
PULPROG zg30
TD 65536
SOLVENT CDCl3
NS 54
DS 2
SWH 8305.647 Hz
FIDRES 0.126734 Hz
AQ 3.9452672 sec
RG 197.76
DW 60.200 usec
DE 6.50 usec
TE 294.7 K
D1 1.0000000 sec
TD0 32

==== CHANNEL f1 =====
SF01 400.1324710 MHz
NUC1 1H
P1 9.50 usec
PLW1 19.00000000 W

F2 - Processing parameters
SI 32768
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.10 Hz
GB 0
PC 1.00



Name: Angerer
Probe: Nitrazolam

Current Data Parameters
NAME Nitrazolam
EXPNO 5
PROCNO 1

F2 - Acquisition Parameters
Date_ 20151030
Time 11.05
INSTRUM spect
PROBHD 5 mm BBO BB-1H
PULPROG zgpg30
TD 50248
SOLVENT CDCl3
NS 1452
DS 4
SWH 25000.000 Hz
FIDRES 0.497532 Hz
AQ 1.0049599 sec
RG 197.76
DW 20.000 usec
DE 6.50 usec
TE 295.0 K
D1 1.00000000 sec
D11 0.03000000 sec
TD0 2000

==== CHANNEL f1 =====
SFO1 100.6228298 MHz
NUC1 13C
P1 9.00 usec
PLW1 45.60400009 W

==== CHANNEL f2 =====
SFO2 400.1316005 MHz
NUC2 1H
CPDPRG[2] waltz16
PCPD2 90.00 usec
PLW2 19.00000000 W
PLW12 0.28382999 W
PLW13 0.22990000 W

F2 - Processing parameters
SI 32768
SF 100.6127743 MHz
WDW EM
SSB 0
LB 2.00 Hz
GB 0
PC 1.40

