

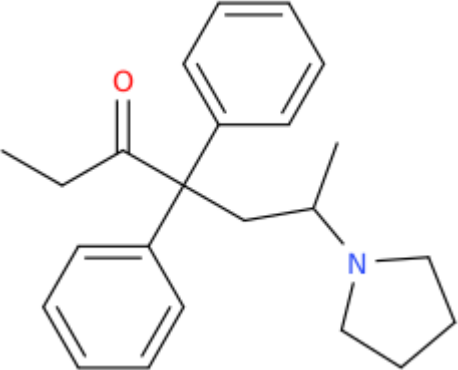
ANALYTICAL REPORT

Dipyanone (C₂₃H₂₉NO)

4,4-diphenyl-6-(pyrrolidin-1-yl)heptan-3-one

Remark – other NPS detected:

Sample ID:	2955-21
Sample description:	powder
Sample type:	test purchase /NFL- purchasing
Date of entry (DD/MM/YYYY) into NFL database:	23/09/2021
Report updates (if any) will be published here:	http://www.policija.si/apps/nfl_response_web/seznam.php

Substance identified - structure ¹ (base form)	
Systematic name	4,4-diphenyl-6-(pyrrolidin-1-yl)heptan-3-one
Other names	Dipipanone pyrrolidinyl
Formula (per base form)	C ₂₃ H ₂₉ NO
M _w (g/mol)	335,49
Salt form/anions detected	HCl
StdInChIKey (per base form)	LJIUPFDRFKFNJE-UHFFFAOYSA-N
Other NPS detected	
Additional info (purity..)	>95% purity of a sample based on 1H NMR spectrum

¹ 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 µl 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 9.1 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 µl 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 KOH 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

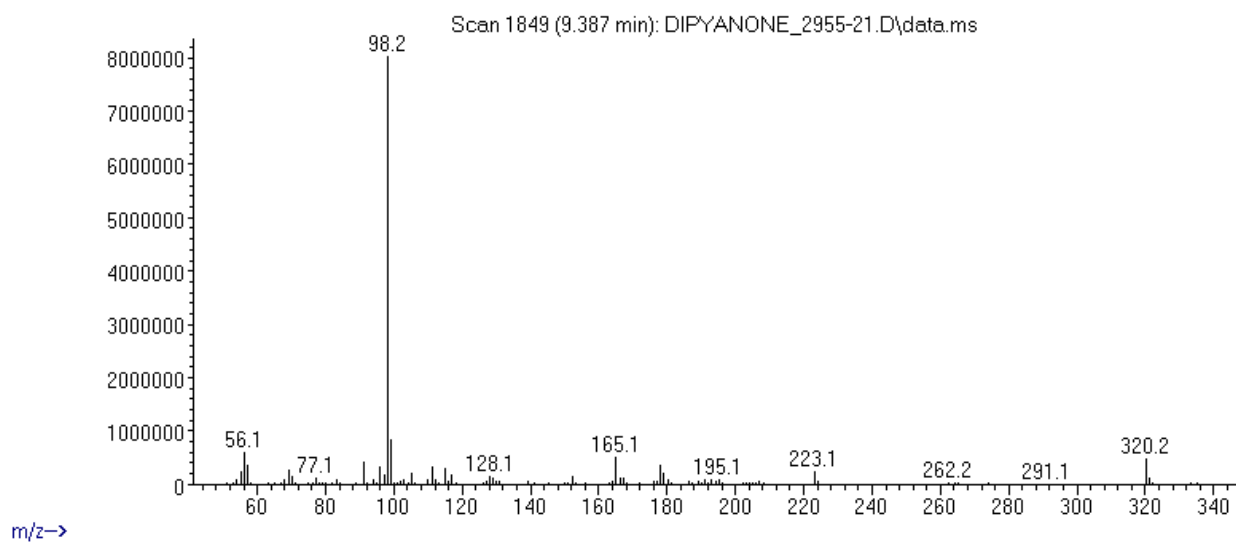
Solubility in	result/remark
CH ₂ Cl ₂	soluble
MeOH	soluble
H ₂ O	partially

Analytical technique:	applied	remarks
GC-MS (EI ionization)	+	NFL GC-RT (min): 9,39 BP(1): 98; BP(2): 99,BP(3) :56,
HPLC-TOF	+	Exact mass (theoretical): 335,2249; measured value Δppm:-0,79; formula:C ₂₃ H ₂₉ NO
FTIR-ATR	+	direct measurement (sample as received)
FTIR (solid phase) always as base form	+	
IC (anions)	+	
NMR (in FKKT)	+	
validation		
other		

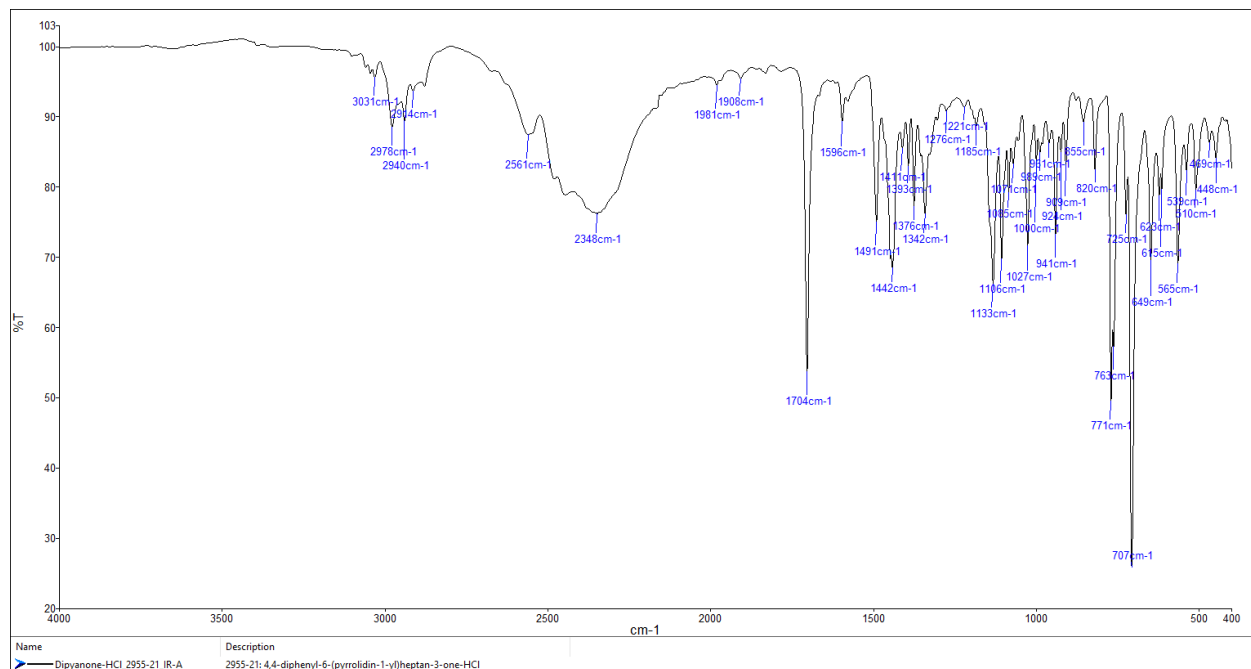
ANALYTICAL RESULTS

MS (EI)

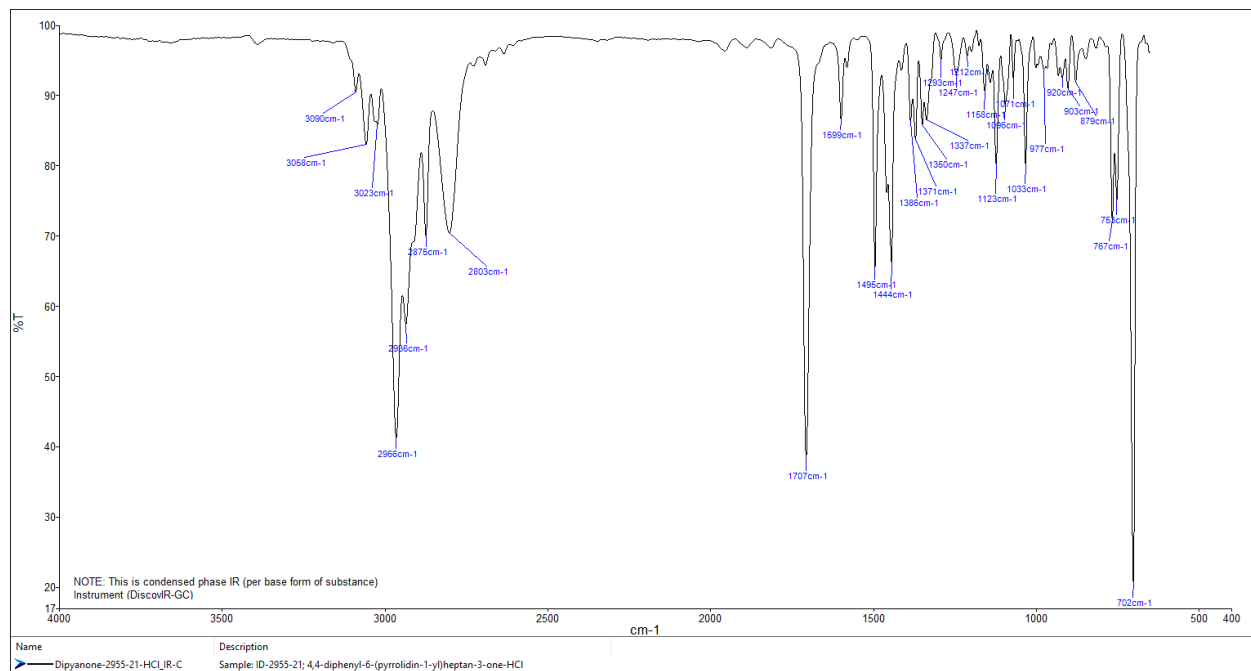
Abundance



FTIR-ATR - direct measurement (sample as received)



IR (solid phase – after chromatographic separation)



TOF REPORT

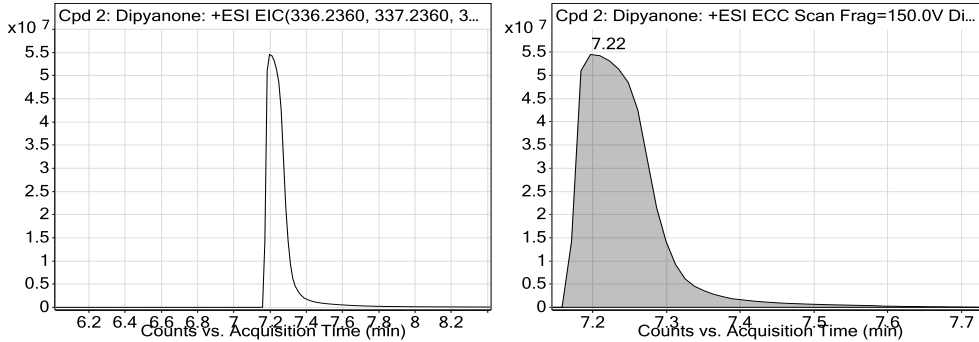
Data File	Dipyanone_2955_21.d	Sample Name	ID-2955-21
Sample Type	Sample	Position	P1-D3
Instrument Name	6230B TOF LC-MS	User Name	TG
Acq Method	general-15_01_2020-XDB-C18-ESI+.m	Acquired Time	5/25/2021 8:38:24 AM
IRM Calibration Status	Success	DA Method	a-Drugs_NFL.m
Comment	MeOH		

Compound Table

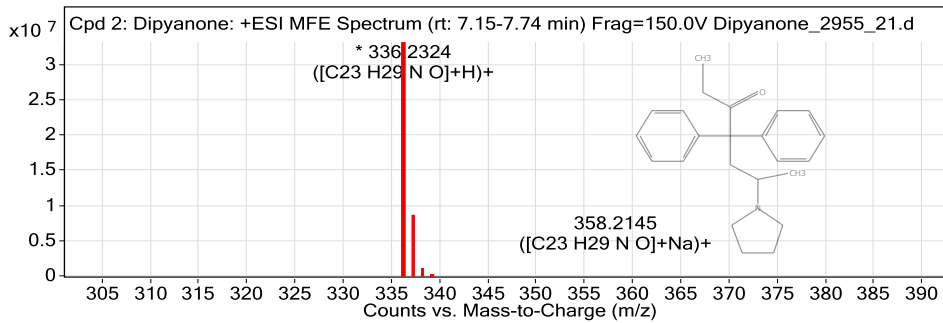
Label	Compound Name	MFG Formula	Obs. RT	Obs. Mass
Cpd 2: Dipyanone	Dipyanone	C23 H29 N O	7.22	335.2252

Name	Obs. m/z	Obs. RT	Obs. Mass	DB RT	DB Formula	DB Mass	DB Mass Error (ppm)
Dipyanone	336.2324	7.22	335.2252	7.22	C23 H29 N O	335.2249	-0.79

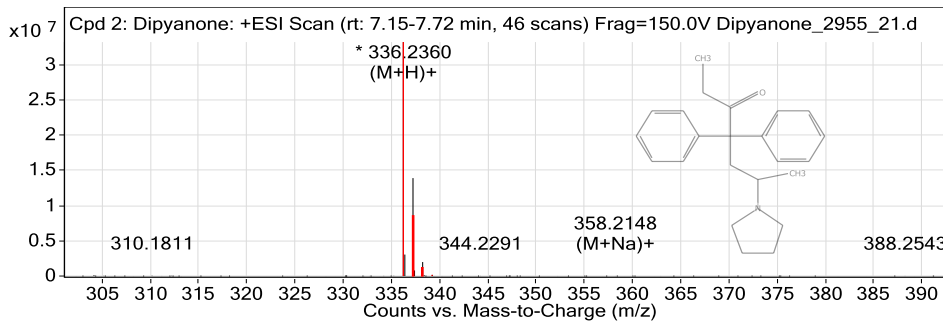
Compound Chromatograms



MFE MS Zoomed Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

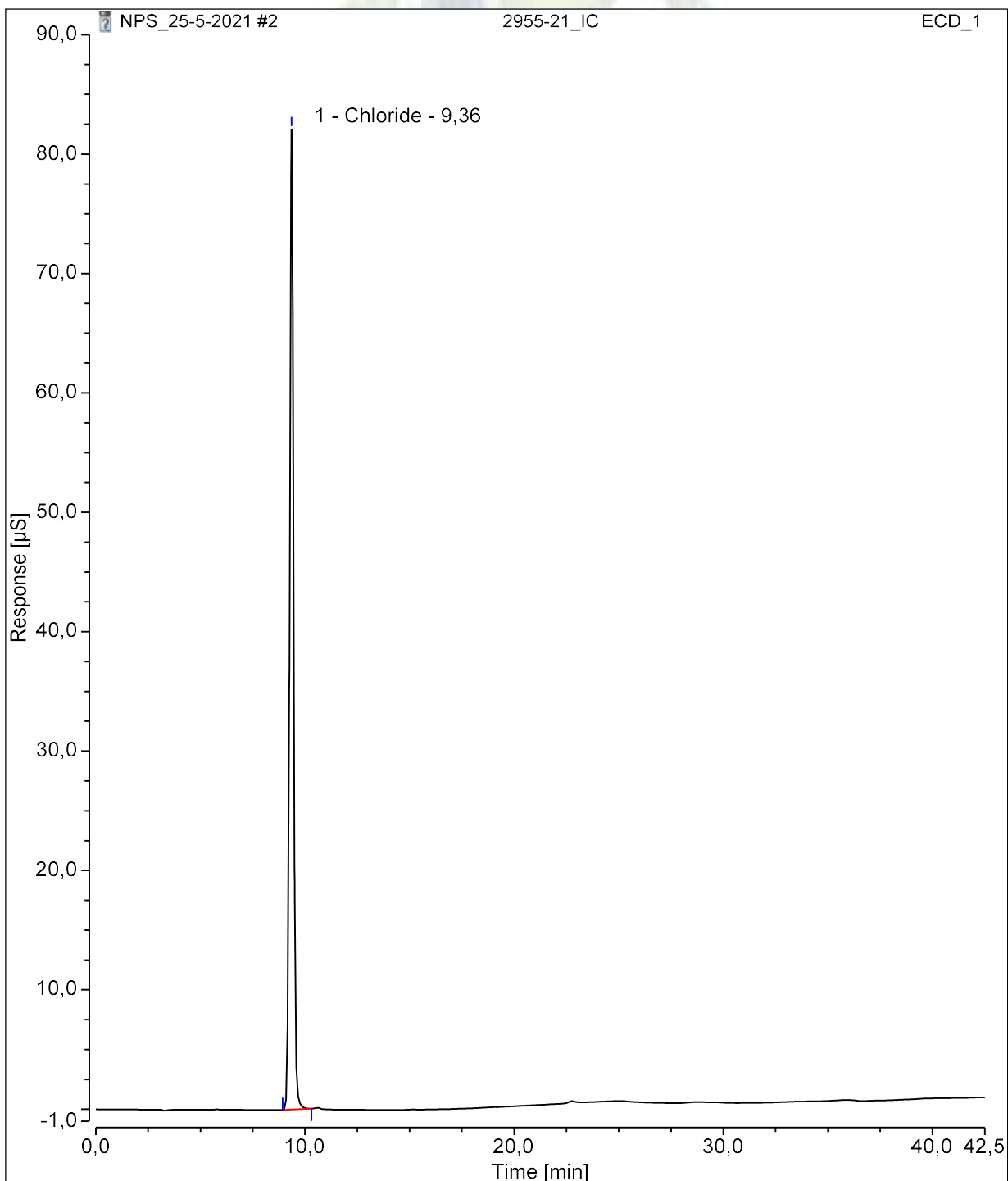
Obs. m/z	Charge	Abund	Formula	Ion/Isotope
336.2324	1	33224840	C23 H29 N O	(M+H)+
337.2358	1	8485043.36	C23 H29 N O	(M+H)+
338.2388	1	1043233.59	C23 H29 N O	(M+H)+
339.2424	1	77956.56	C23 H29 N O	(M+H)+
358.2145	1	9217.8	C23 H29 N O	(M+Na)+

--- End Of Report ---

Peak Integration Report

Sample Name:	2955-21_IC	Inj. Vol.:	25,00
Injection Type:	Unknown	Dilution Factor:	1,0000
Program:	ANIONI	Operator:	Admin
Inj. Date / Time:	25-May-2021 / 09:09	Run Time:	43,00

No.	Time min	Peak Name	Peak Type	Area $\mu\text{S}\cdot\text{min}$	Height μS	Amount mg/L
1	9,36	Chloride	BMB	17,716	82,101	n.a.
TOTAL:				17,72	82,10	0,0

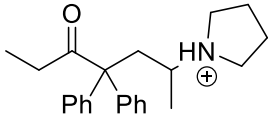


University
of Ljubljana

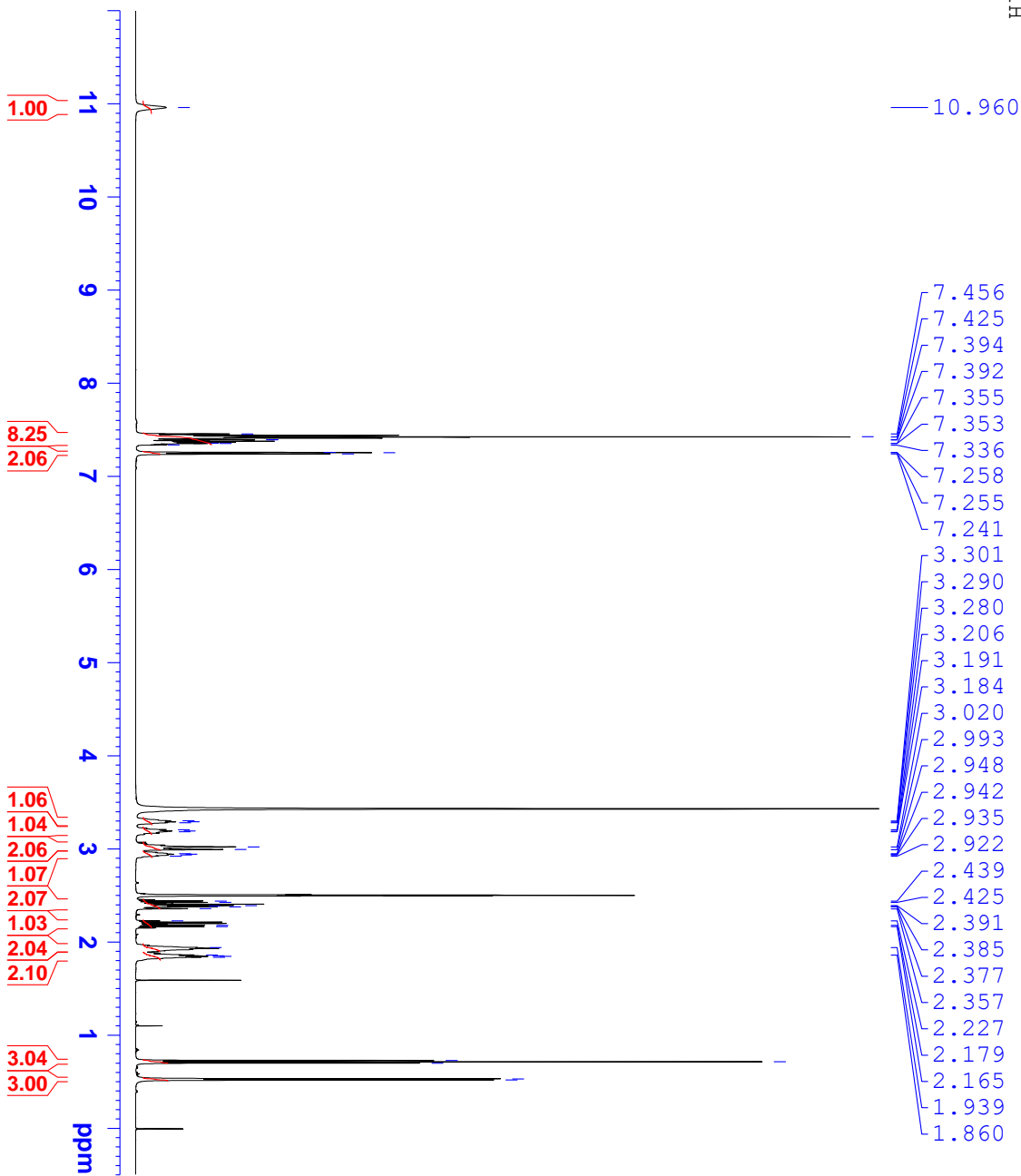
Faculty of Chemistry
and Chemical Technology



R E P O R T

Contract No.	C1714-21-460153 (Republic of Slovenia, Ministry of the Interior, POLICE)
Sample ID:	2955-21
Received date:	September 6, 2021
Our notebook code:	NFL-2955-21
NMR sample preparation:	19.7 mg dissolved in 0.7 mL DMSO- <i>d</i> ₆
NMR experiments:	¹ H, ¹³ C, ¹ H- ¹ H <i>gs</i> -COSY, ¹ H- ¹³ C <i>gs</i> -HSQC, ¹ H- ¹³ C <i>gs</i> -HMBC, ¹ H- ¹⁵ N <i>gs</i> -HMBC
Proposed structure with formula, exact mass, molecular weight:	 <p>Chemical Formula: C₂₃H₃₀NO⁺ Exact Mass: 336,2322 Molecular Weight: 336,4985</p>
Chemical name:	<i>N</i> -protonated 4,4-diphenyl-6-(pyrrolidin-1-yl)heptan-3-one
Comments:	- Structure elucidation based on 1D and 2D NMR spectra and HRMS. ->95% purity of a sample based on ¹ H NMR spectrum.
Supporting information:	Copies of ¹ H and ¹³ C NMR spectra, ¹ H and ¹³ C FIDs.
Principal investigator:	Prof. Dr. Janez Košmrlj
Date of report:	September 18, 2021

NFL-2955-21
1H



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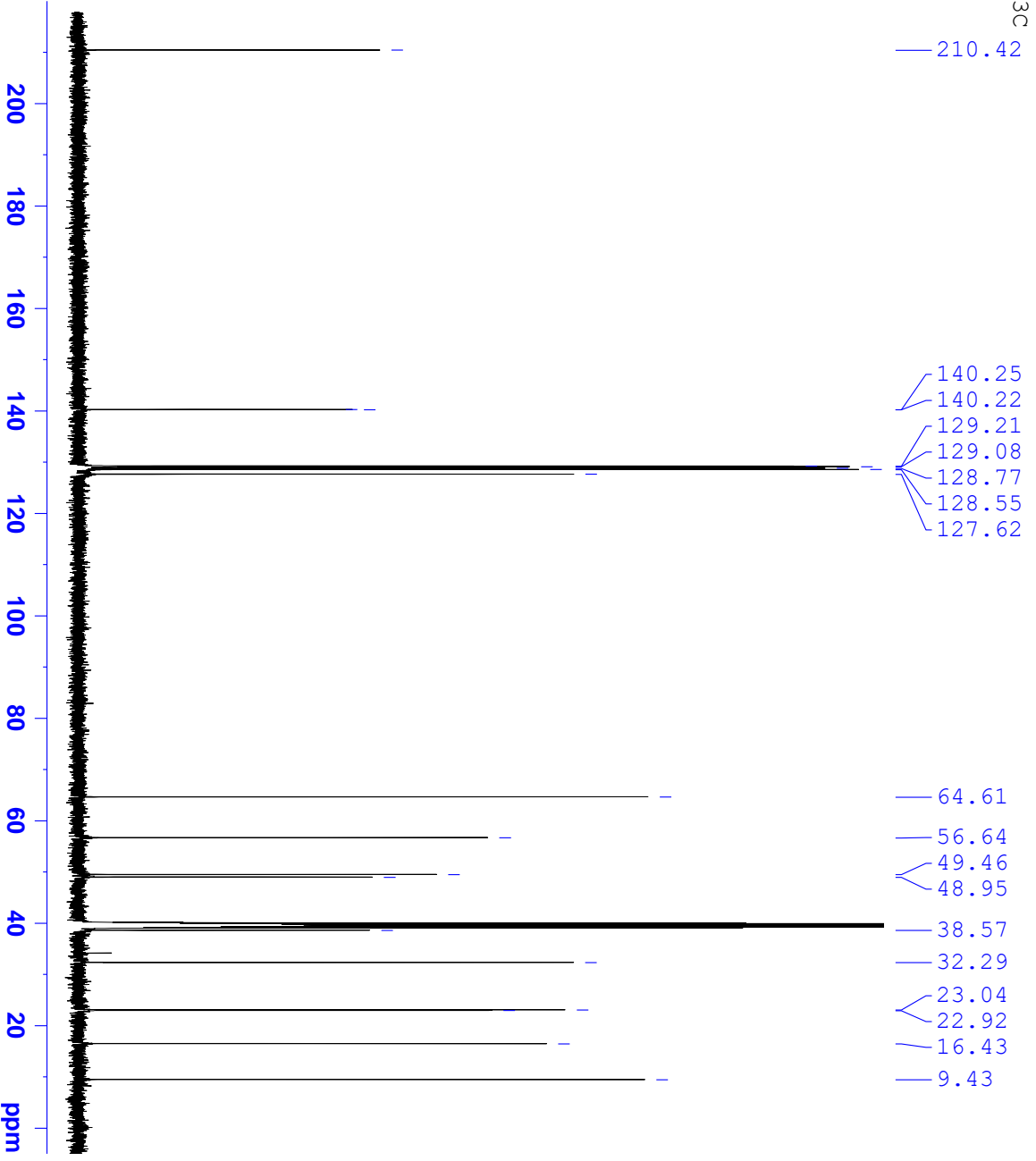
Current Data Parameters
NAME          NFL-2955-21
EXPNO         1
PROCNO       1

F2 - Acquisition Parameters
Date_         20210908
Time          0.11
INSTRUM      spect
PROBHD       5 mm PABBI 1H/
PULPROG      zg30
TD           65536
SOLVENT      DMSO
NS           16
DS           2
SWH          10000.000 Hz
FIDRES       0.152588 Hz
AQ           3.2767999 sec
RG           57
DW           50.000 usec
DE           6.50 usec
TE           299.0 K
D1           1.00000000 sec
TD0          1

===== CHANNEL f1 =====
SFO1         500.1330885 MHz
NUC1         1H
P1           7.10 usec
PLM1         15.00000000 W

F2 - Processing parameters
SI           65536
SF           500.1300043 MHz
WDW          EM
SSB          0
LB           0.30 Hz
GB           0
PC           1.00
  
```

NFL-2955-21
13C



Current Data Parameters
 NAME NFL-2955-21
 EXPNO 2
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20210908
 Time 2.55

INSTRUM spect
 PROBHD 5 mm PABBI 1H/
 PULPROG zgpg30
 TD 65536
 SOLVENT DMSO
 NS 3072
 DS 4

SWH 29761.904 Hz
 FIDRES 0.454131 Hz
 AQ 1.1010048 sec
 RG 2050
 DW 16.800 usec
 DE 6.50 usec
 TE 299.4 K

D1 2.00000000 sec
 D11 0.03000000 sec
 TD0 1

==== CHANNEL f1 =====
 SFO1 125.7703637 MHz
 NUC1 13C
 P1 14.00 usec
 PLW1 114.0000000 W

==== CHANNEL f2 =====
 SFO2 500.1320005 MHz
 NUC2 1H
 CPDPRG12 waltz16
 PCPD2 80.00 usec
 PLW2 16.00000000 W
 PLW12 0.12250000 W
 PLW13 0.06161700 W

F2 - Processing parameters
 SI 32768
 SF 125.7578516 MHz
 WDW EM
 SSB 0
 LB 1.00 Hz
 GB 0
 PC 1.40