

Synthesis, Molecular Docking Analysis and *in Vitro* Evaluation of New Heterocyclic Hybrids of 4-Aza-Podophyllotoxin as Potent Cytotoxic Agents

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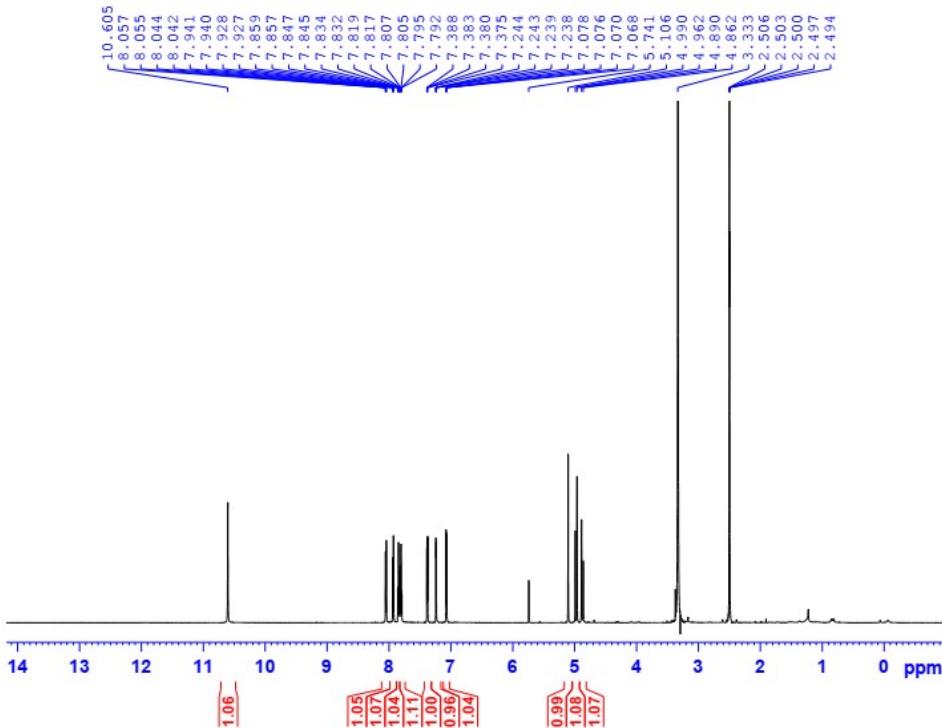
SPECTRUM SECTION

11-(Thiophen-3-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13a)

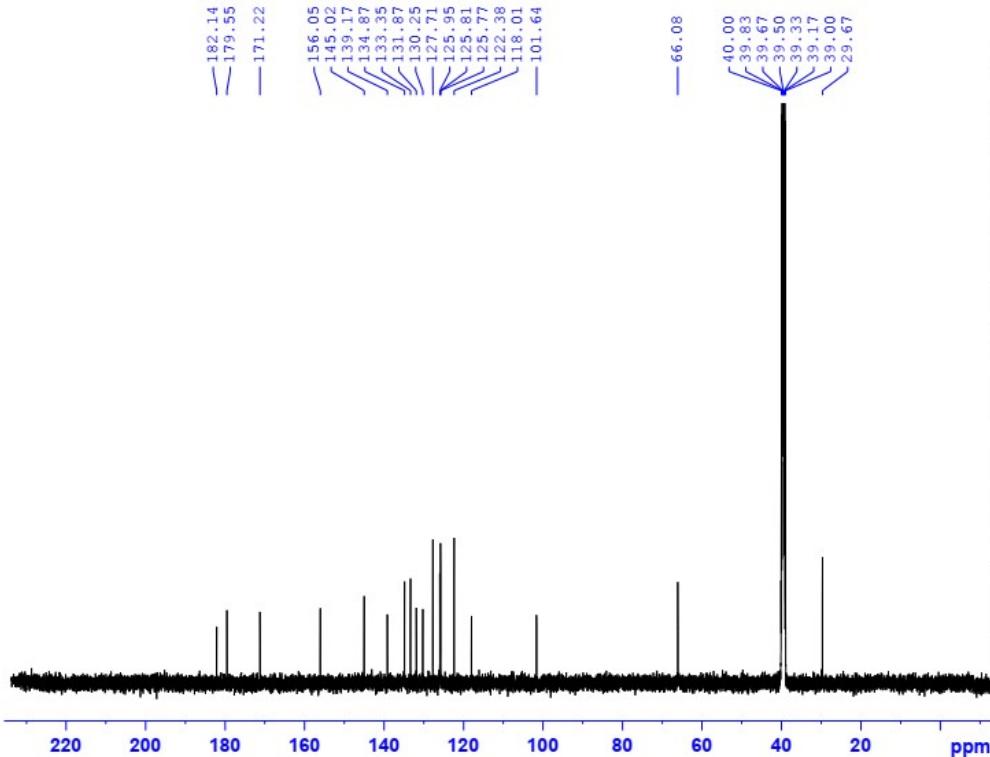
HP245.39-DMSO-1H

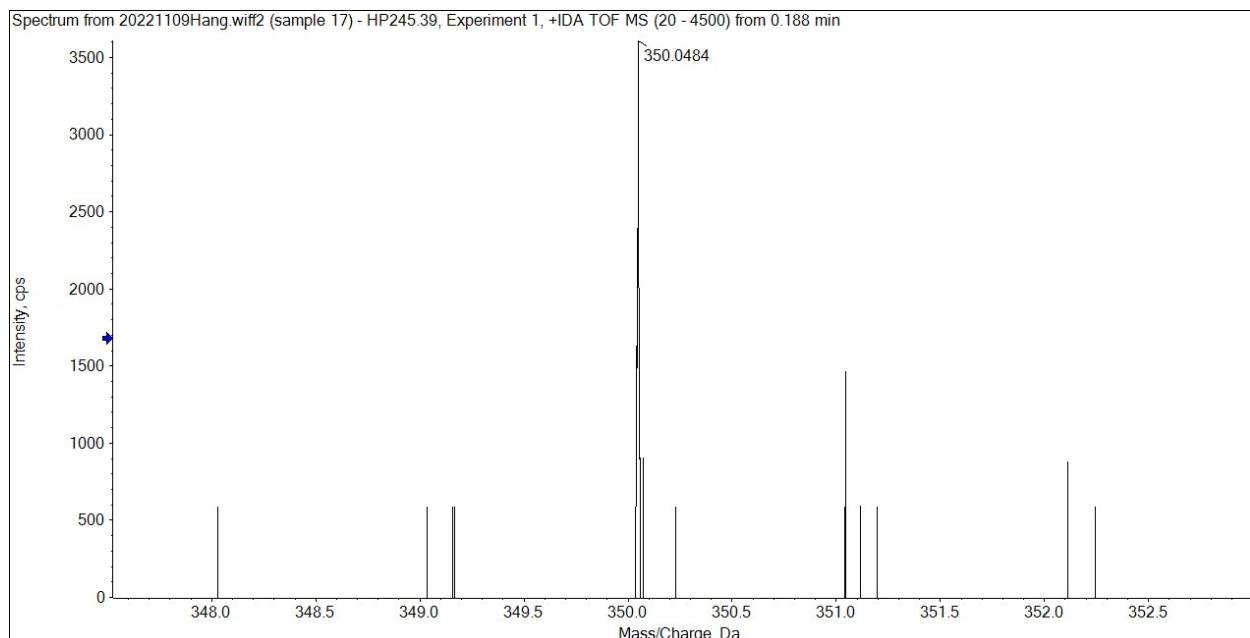


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 PROBHD: Z114607_0862
 PULPROG: zg30
 TD: 65536
 SOLVENT: DMSO
 NS: 16
 DS: 2
 SWH: 11904.762 Hz
 FIDRES: 0.363204 Hz
 AQ: 2.752500 sec
 RG: 97.11844
 DW: 42.000 usec
 DE: 8.71 usec
 TE: 303.2 K
 D1: 1.0000000 sec
 TDO: 1
 SF01: 600.4027075 MHz
 NUC1: 1H
 FO: 3.50 usec
 PI: 10.00 usec
 PLW1: 27.03700066 W
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 PLW1: 1.00



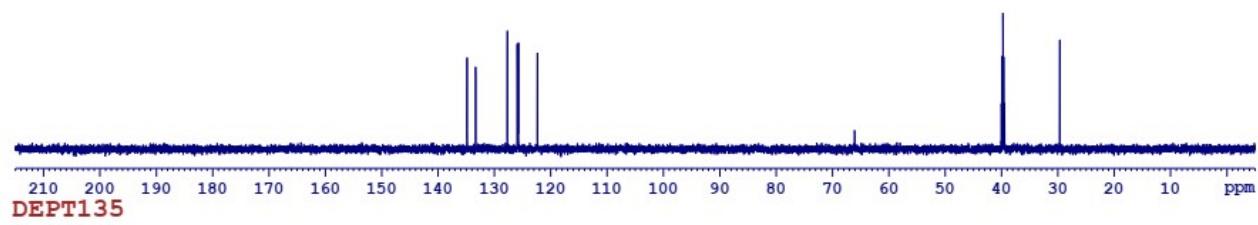
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 PULPROG: zg30
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 SOLVENT: DMSO
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 SWH: 91250.000 Hz
 FIDRES: 0.476327 Hz
 AQ: 1.0485760 sec
 RG: 198.57
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 DE: 6.50 usec
 TE: 303.9 K
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 NUC1: 13C
 PI: 10.00 usec
 PLW1: 88.00000000 W
 ===== CHANNEL f2 ======
 SF02: 500.12410010 MHz
 NUC1: 1H
 CPFG2: 100.000000 Hz
 PCPFG2: 80.00 usec
 PLW2: 22.00000000 W
 PLW12: 0.35764000 W
 PLW13: 0.17989001 W
 F2 - Processing parameters
 SI: 32768
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 WDW: EM
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 PC: 1.40





HP245.39-DMSO-C13CPD&DEPT

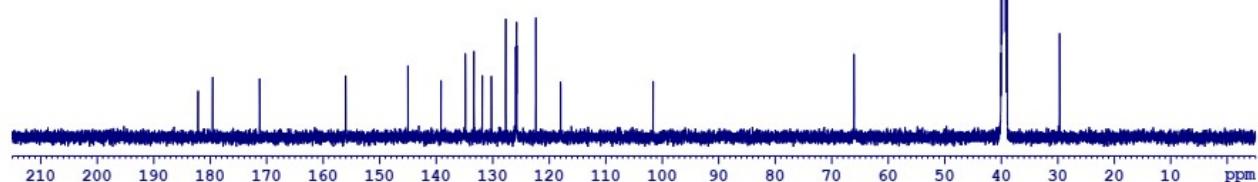
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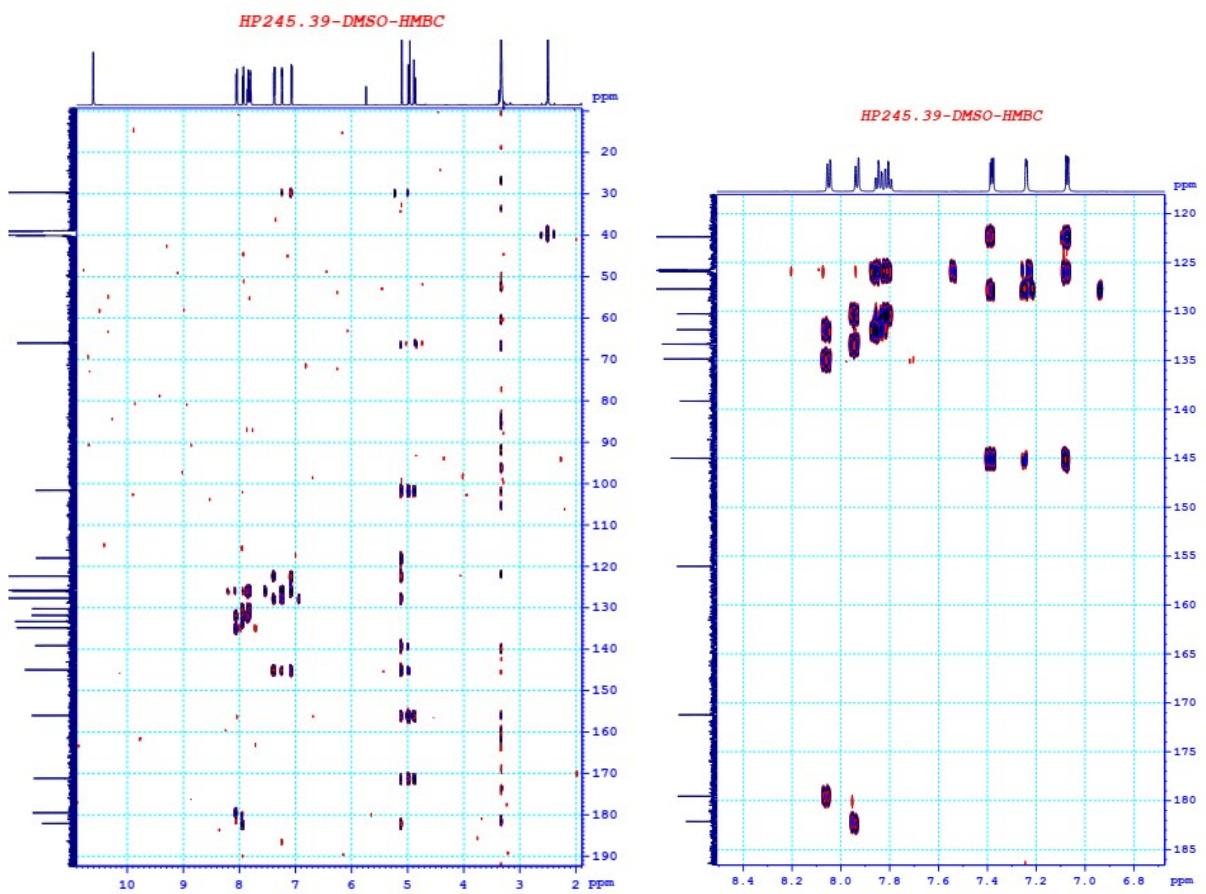
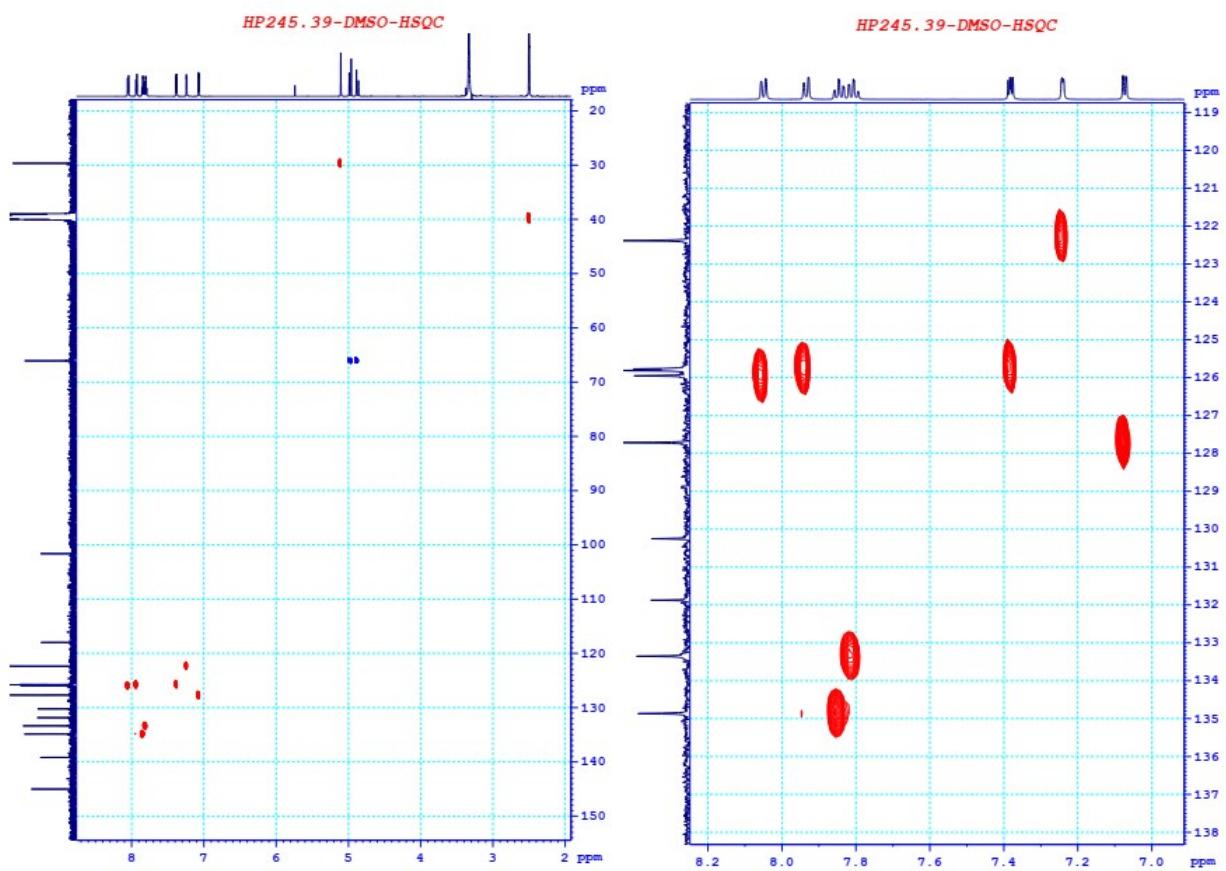


CH&CH₃



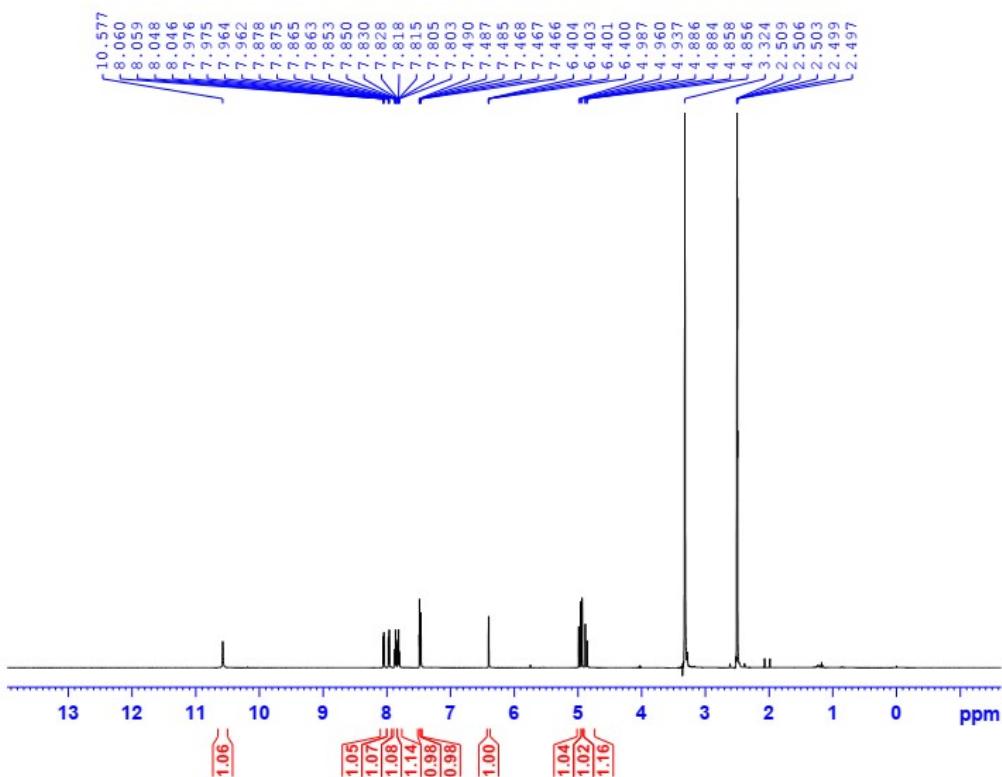
C13CPD





11-(Furan-3-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13b)

TK245.89-DMSO-1H

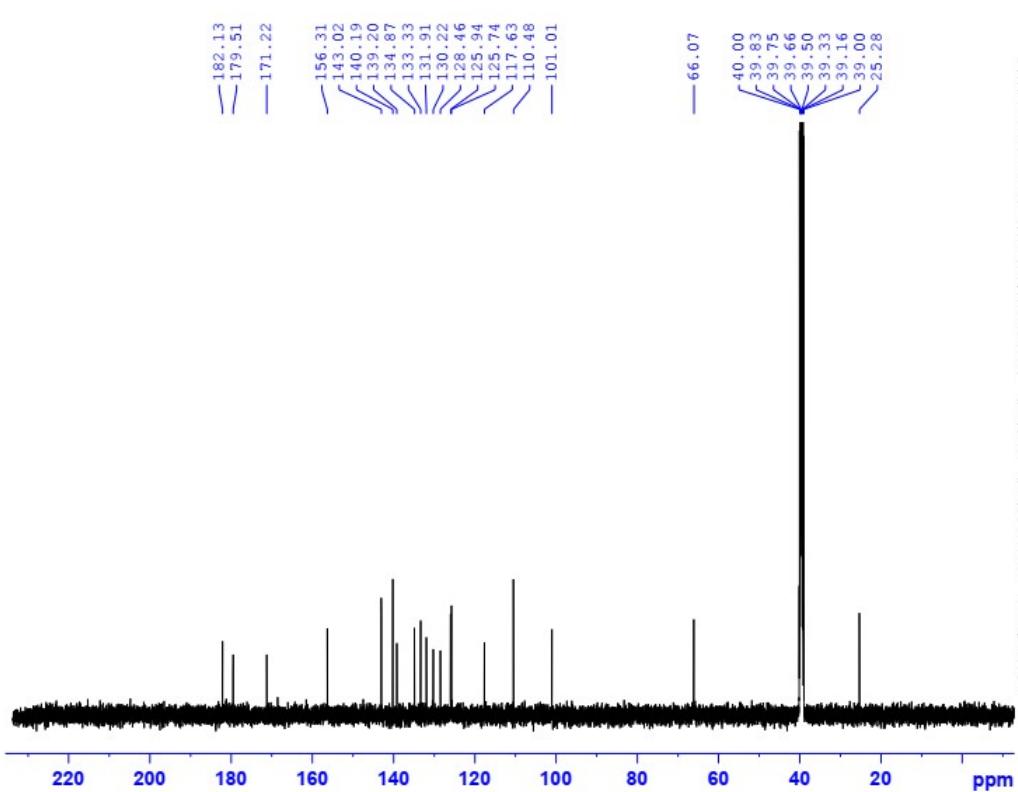


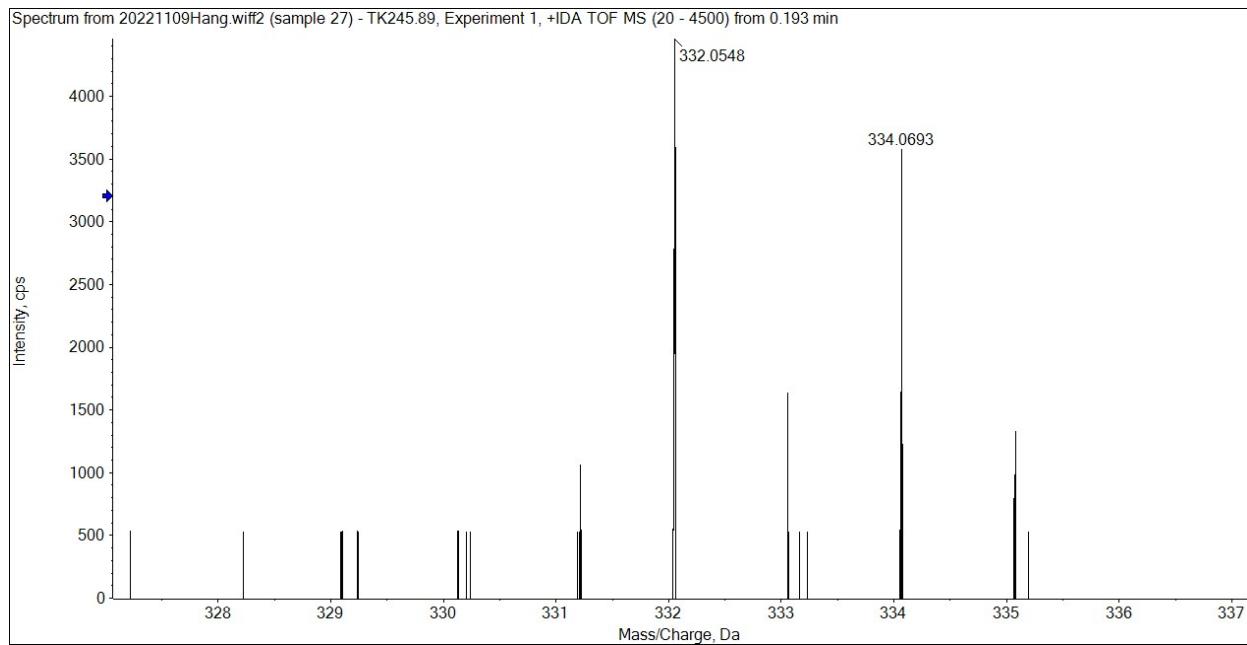
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PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 11904.762 Hz
FIDRES 0.362304 Hz
AQ 2.752512 sec
RG 100.0000
DW 42.000 usec
DE 8.71 usec
TE 303.2 K
D1 1.0000000 sec
TDO 1
SF01 600.4037075 MHz
NUC1 1H
PO 3.50 usec
PL 10.50 usec
PLW1 27.03700066 W

FID - Processing parameters
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PC 1.00

TK245.89-DMSO-C13CPD



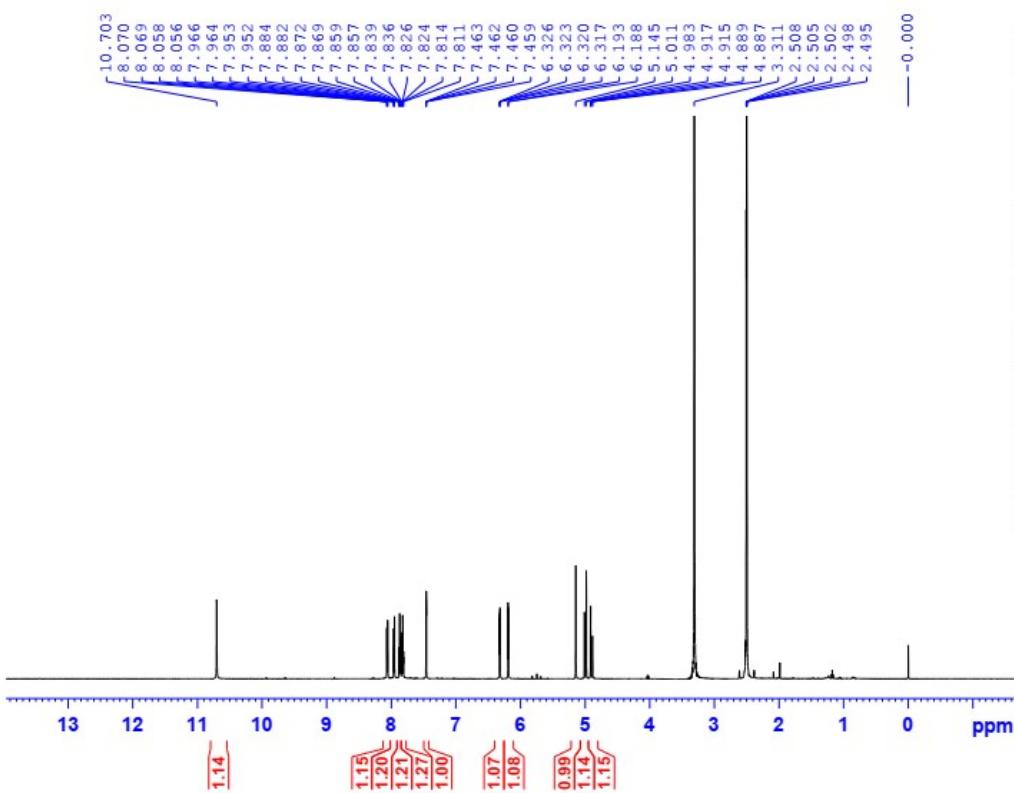


11-(Furan-2-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13c)

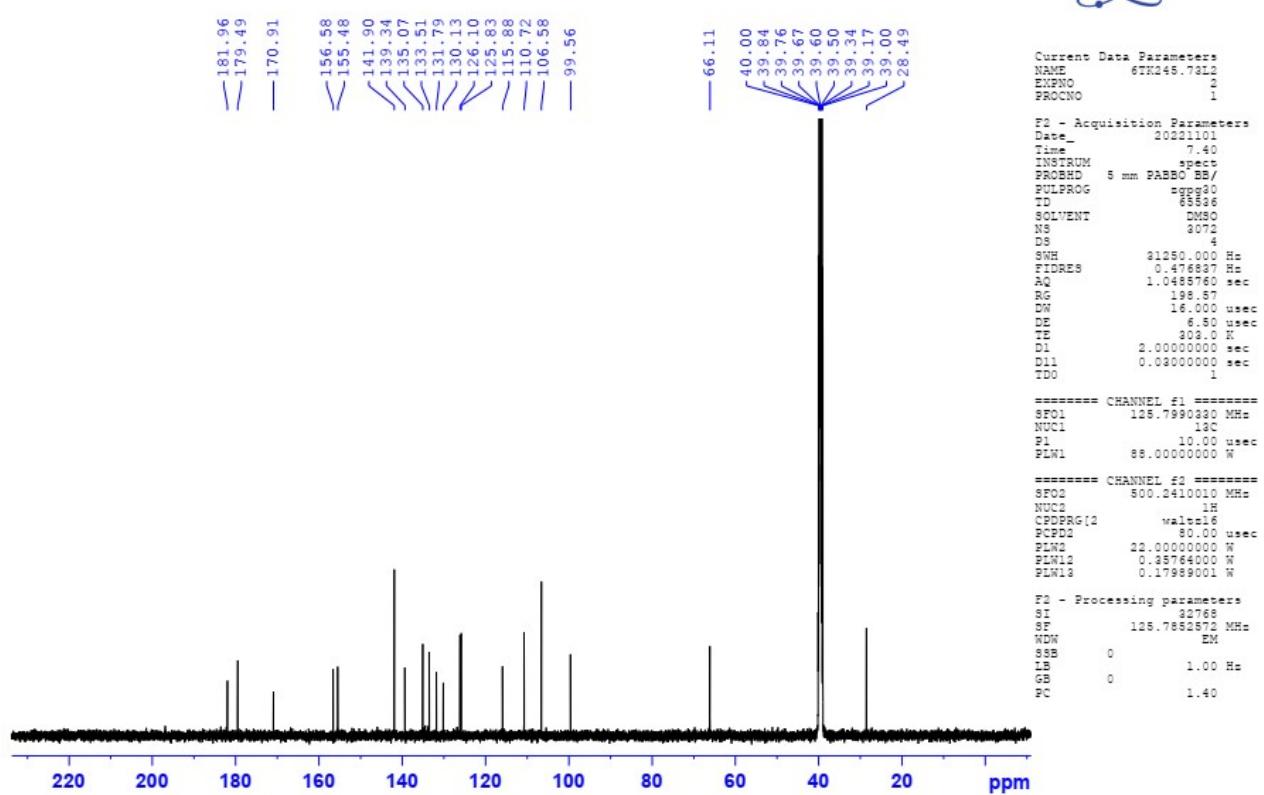
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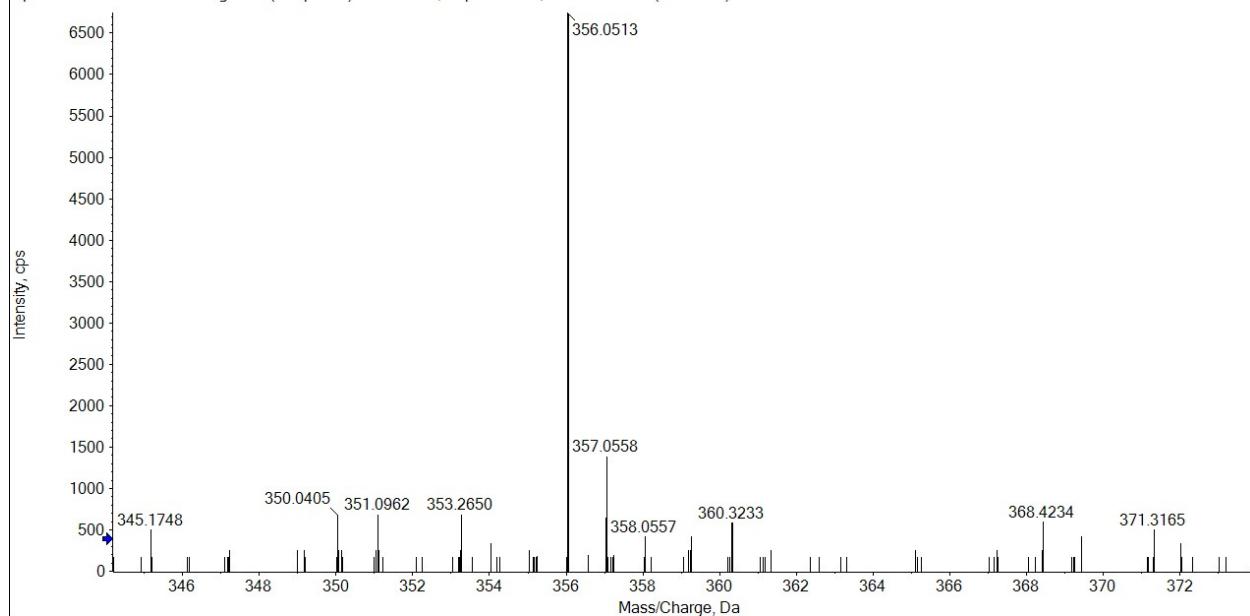
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 PULPROG zg30
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 SOLVENT DMSO
 NS 16
 DS 2
 SWH 11904.762 Hz
 FIDRES 0.2623204 Hz
 AQ 2.7525120 sec
 RG 101
 DW 42.000 usec
 DE 8.71 usec
 TE 303.2 K
 D1 1.0000000 sec
 TDO 1
 SF01 600.4037075 MHz
 NUC1 1H
 P0 3.50 usec
 PI 10.50 usec
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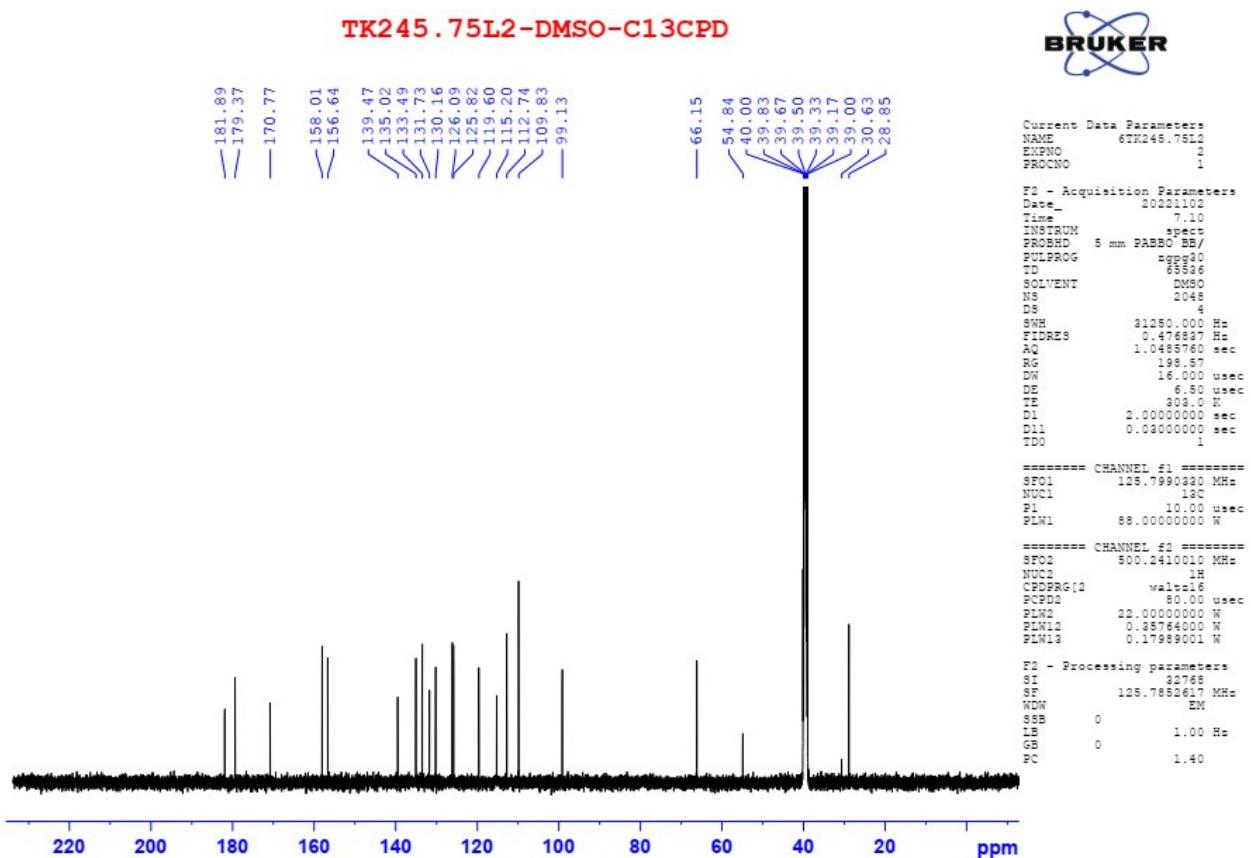
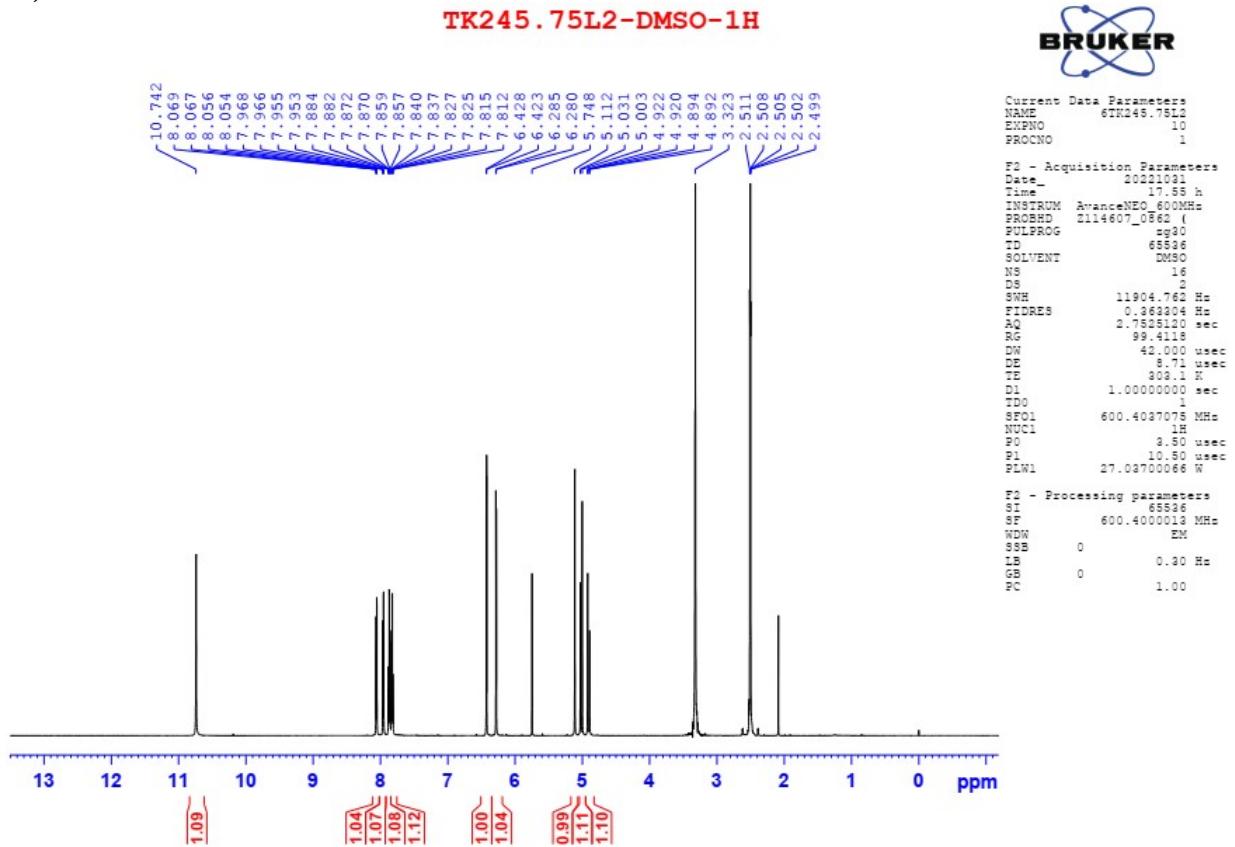
TK245.73L2-DMSO-C13CPD

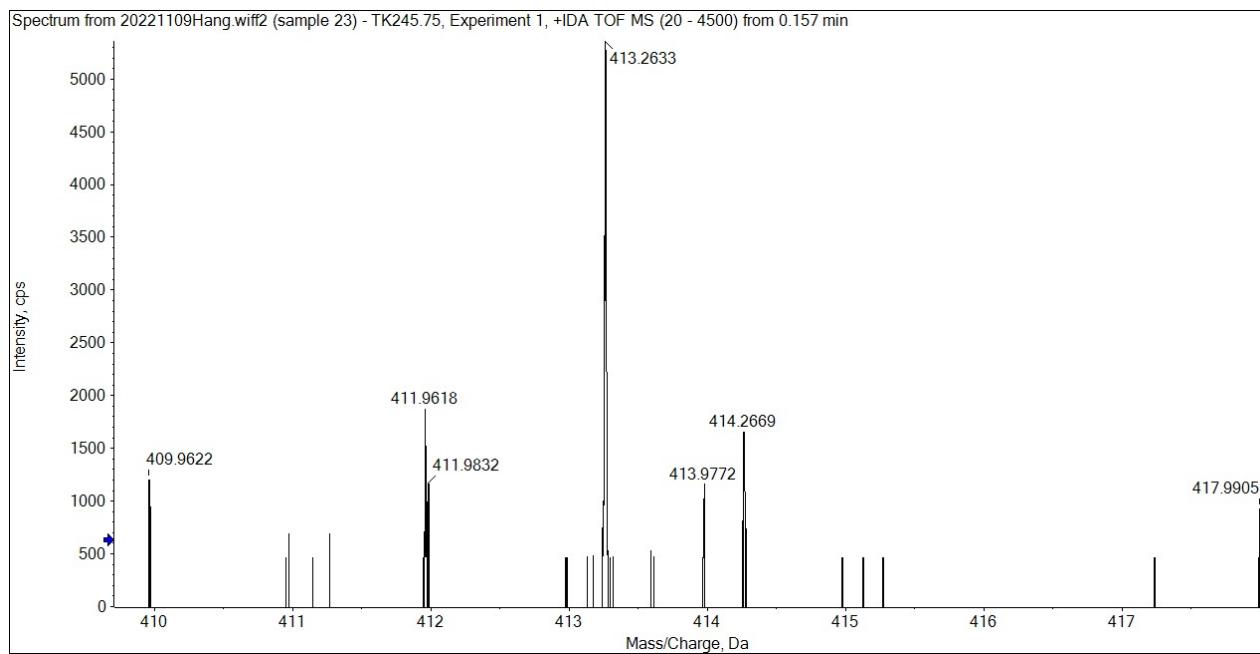


Spectrum from 20221109Hang.wiff2 (sample 21) - HP245.73, Experiment 1, +IDA TOF MS (20 - 4500) from 0.356 min



11-(5-Bromofuran-2-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13d)





11-(Benzo[b]thiophen-3-yl)-4,11-dihydrobenzo[g]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13e)

HP245.68-DMSO-1H



Current Data Parameters
NAME 6HP245.68
EXPNO 10
PROCNO 1

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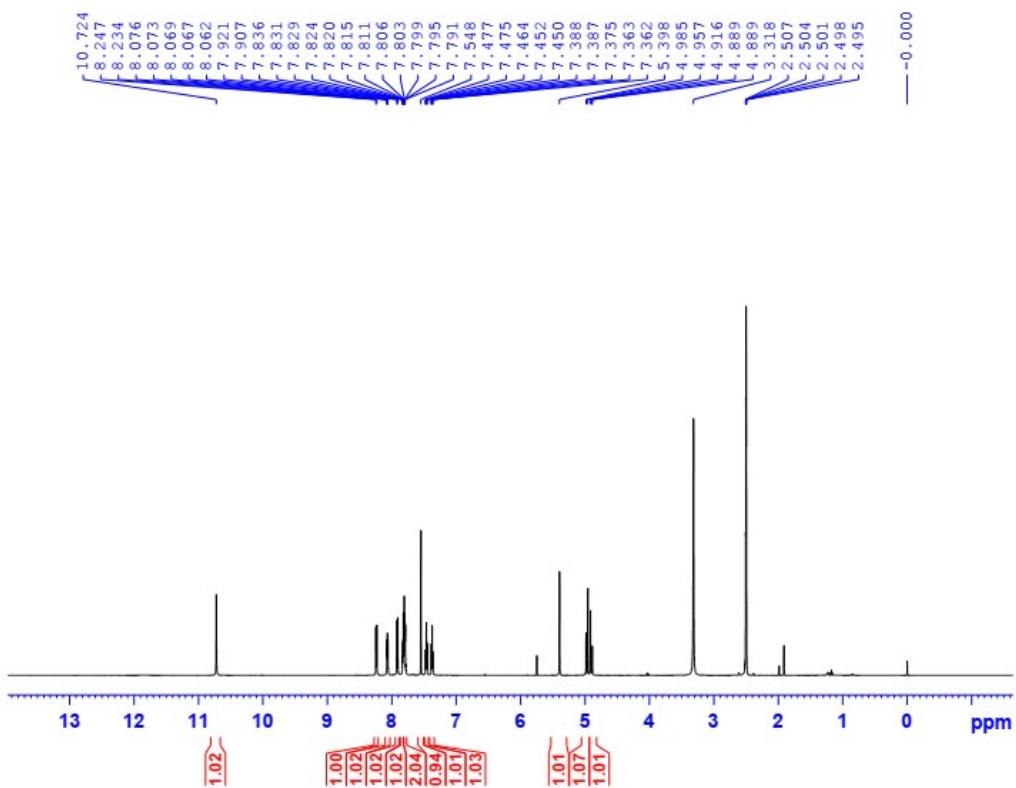
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PROBHD 2114607_0562 (
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TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 11904.760 Hz
FIDRES 0.383004 Hz
AQ 2.755210 sec
RG 101
DW 42.000 used
DE 8.71 used
TE 302.1 used
D1 1.0000000 sec
TDO
SF01 600.4037075 MHz
NUC1 1H
PS 3.50 used
P1 10.50 used
P2 10.50 used

```

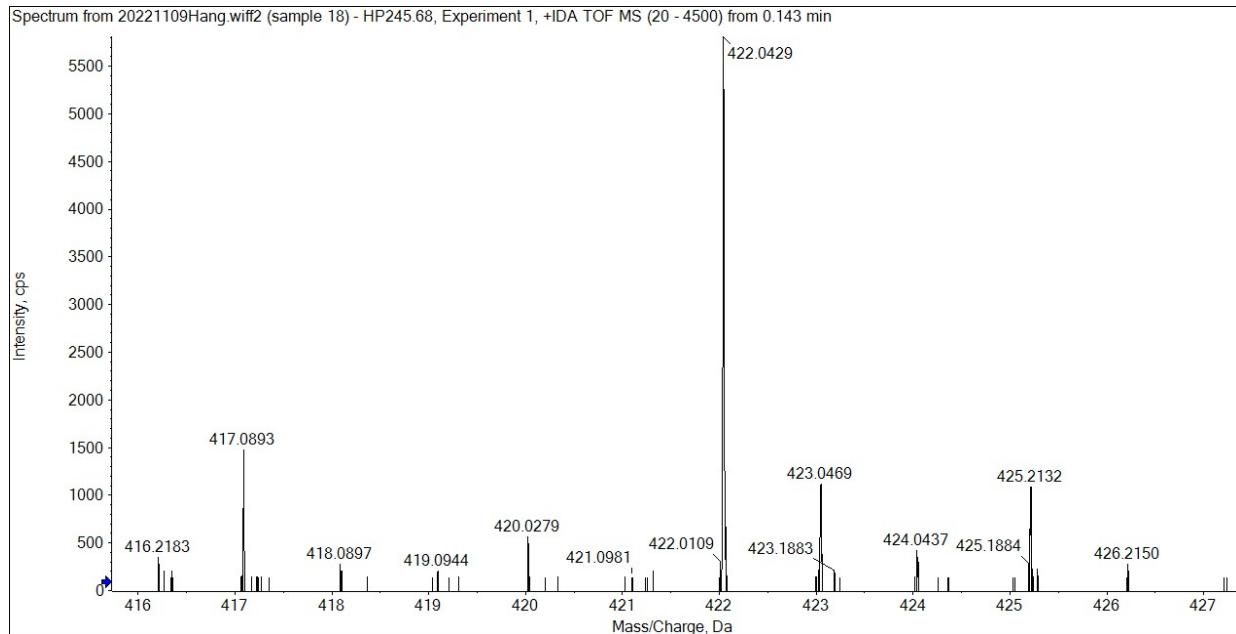
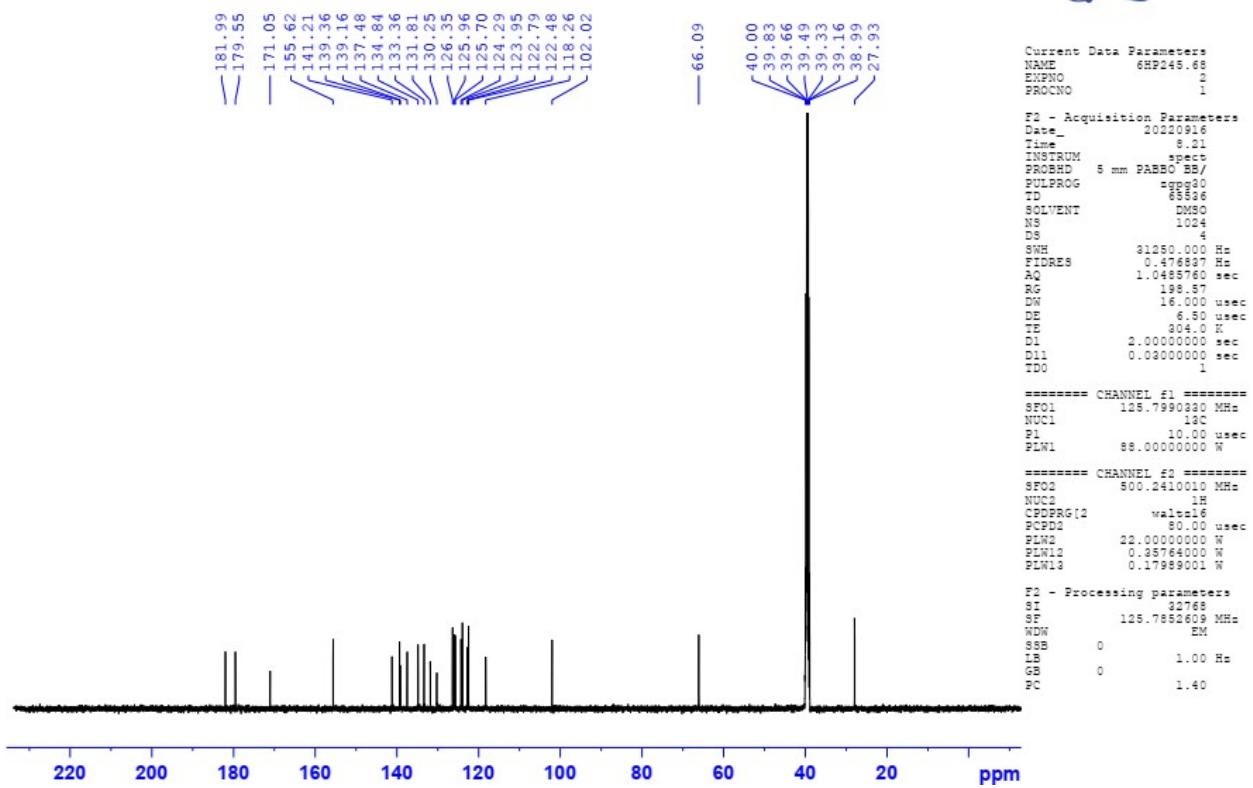
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F2 - Processing parameters
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GB          0
PC          1.00

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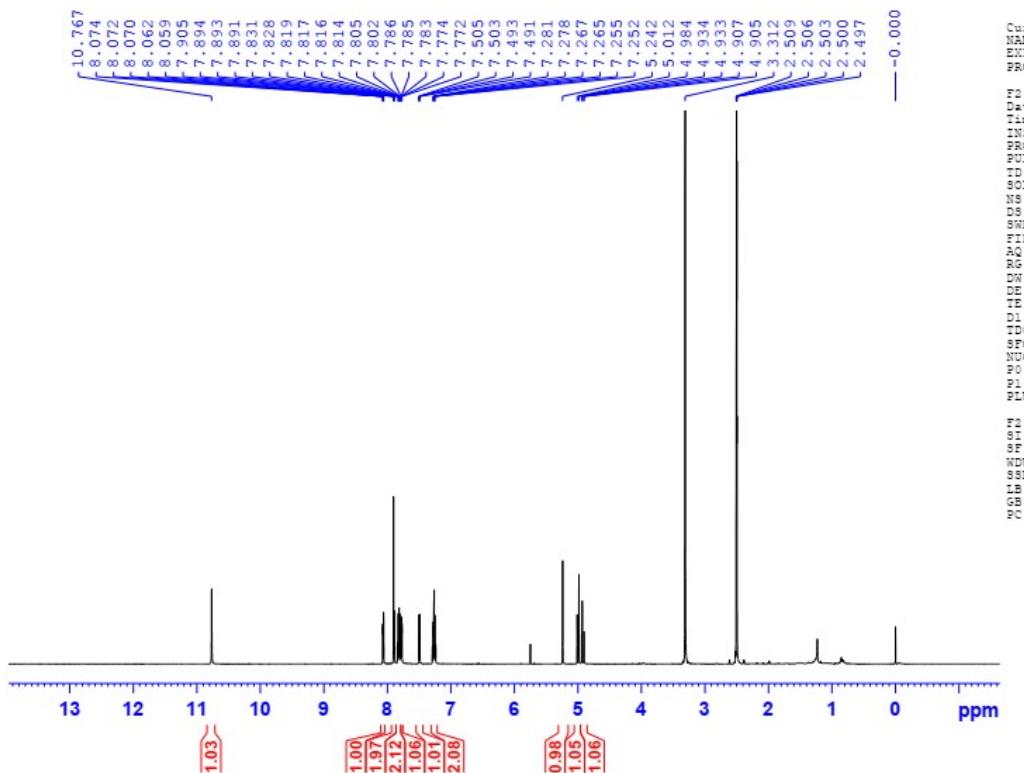


HP245 . 68-DMSO-C13CPD



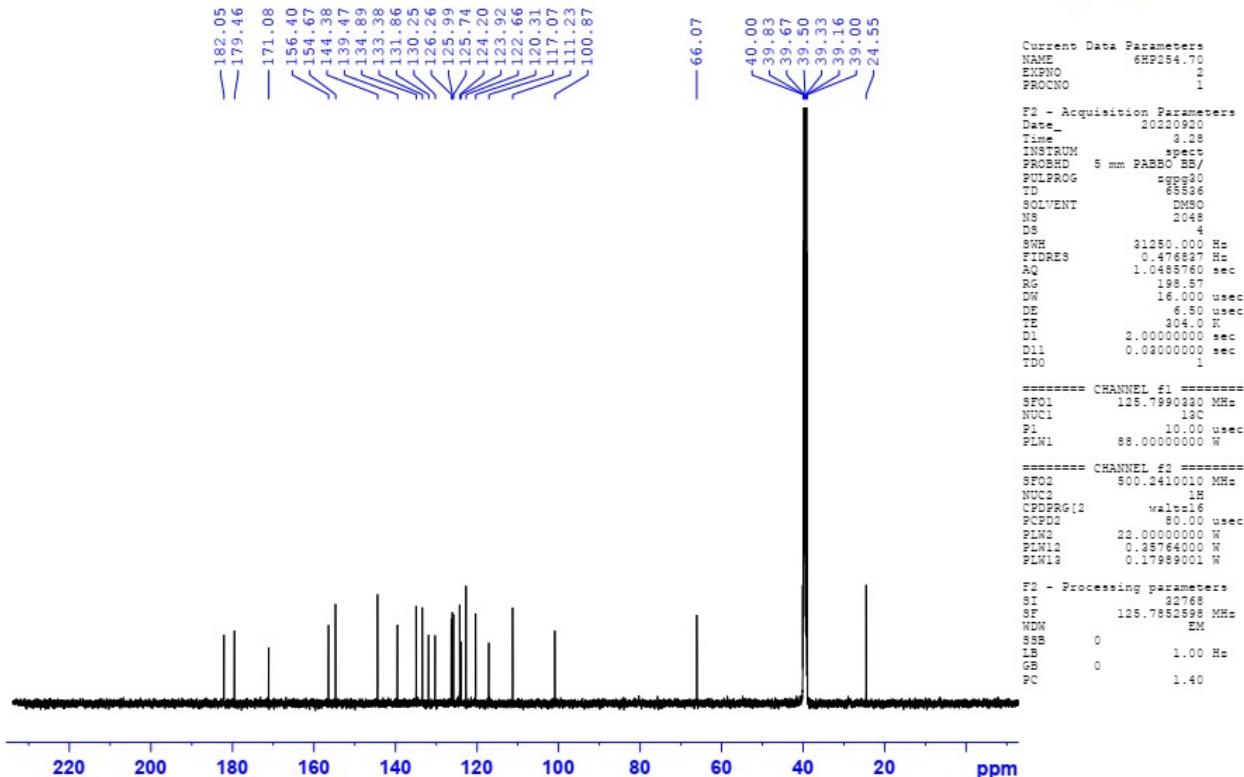
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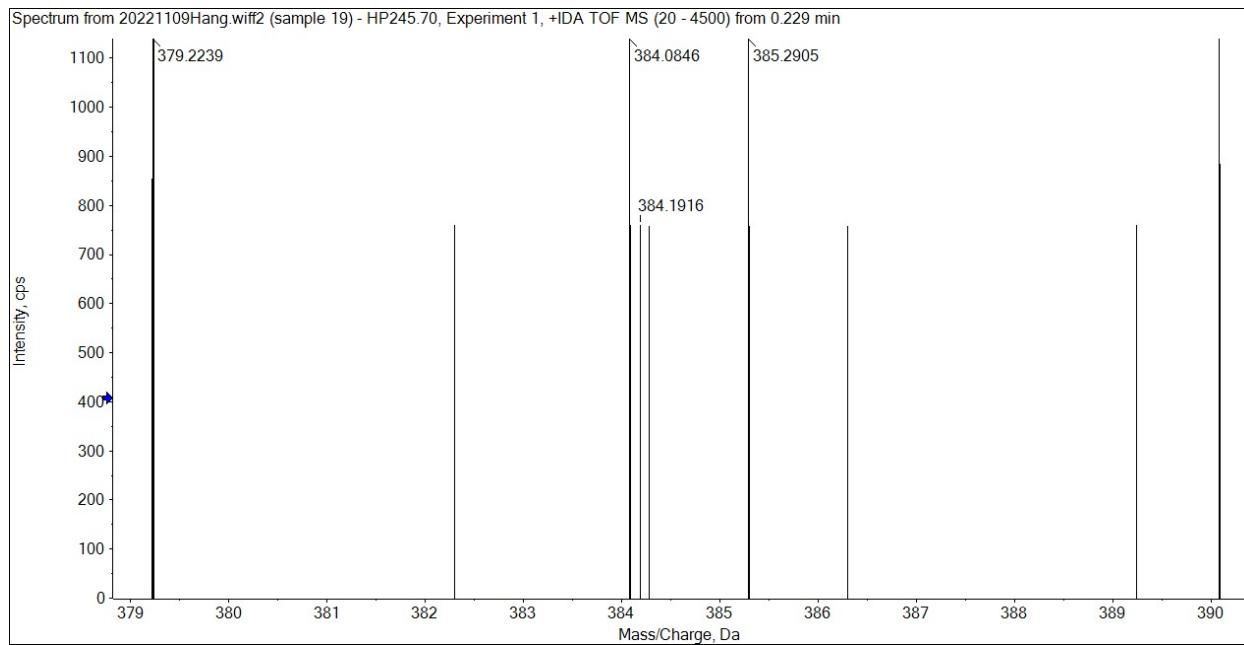
HP254.70-DMSO-1H



HP254 .70=DMSO=C13CPD

BRUKER





11-(Benzo[b]thiophen-2-yl)-4,11-dihydrobenzo[g]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13g)

HP245.72-DMSO-1H



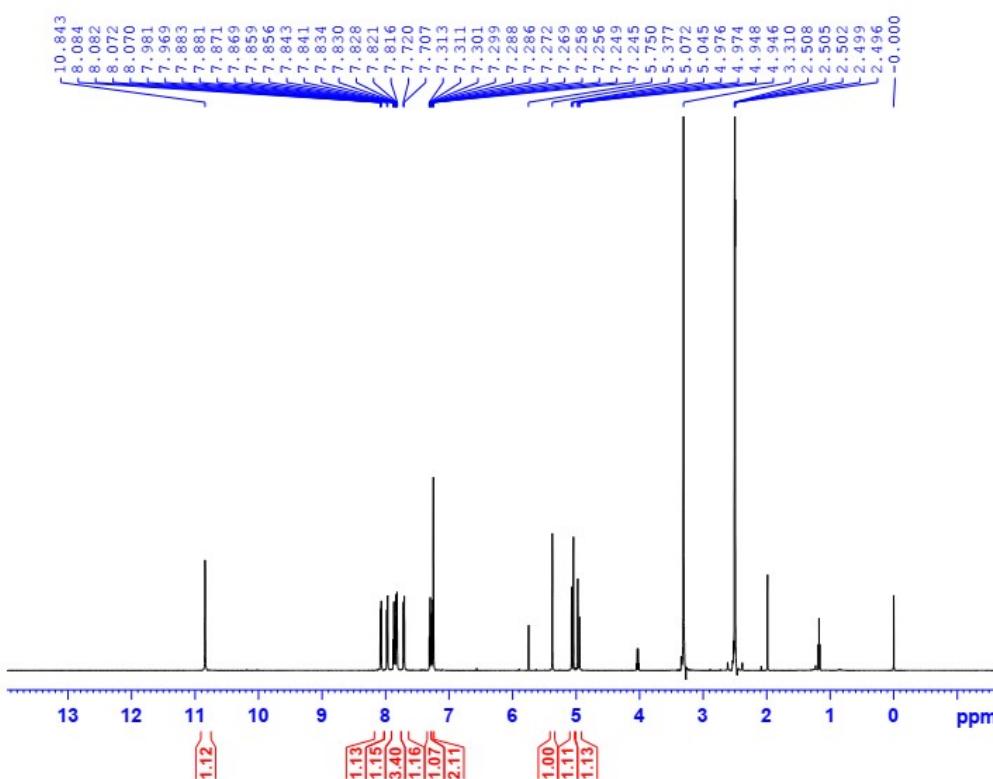
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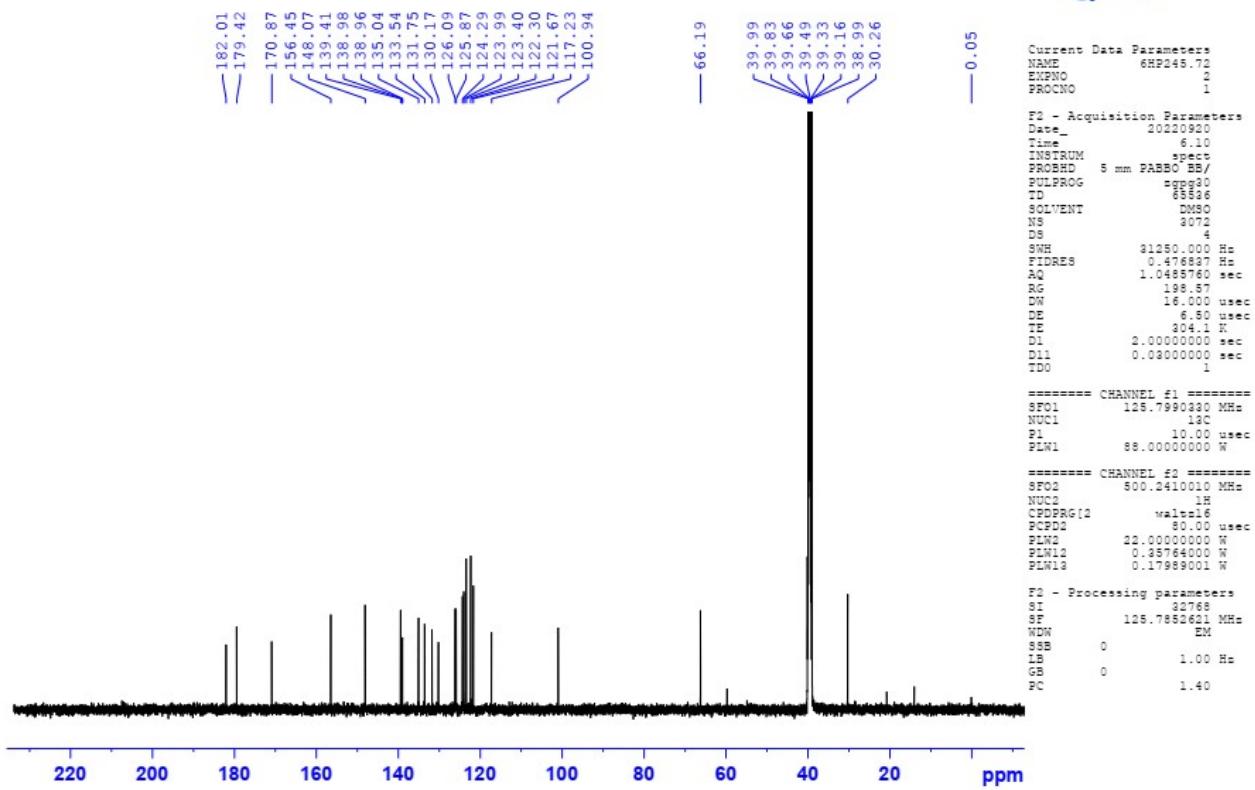
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TD        65536
SOLVENT    DMSO
NS           16
DS            2
SWH       11804.762 Hz
ETR        0.3832840 sec
AQ        2.7852100 sec
RG           150
TE        42.00 usec
TM        6.71 usec
TEC       303.1 usec
D1        1.0000000 sec
TDO          1
SF01      600.4027075 MHz
NUC1         1H
PO           3.50 usec
PL          10.50 usec
PLW1     27.0870066 W

F2 - Processing parameters
SI           65536
SF        600.4000031 MHz
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LB           0.30 Hz
GSB           0
GSC           1.00

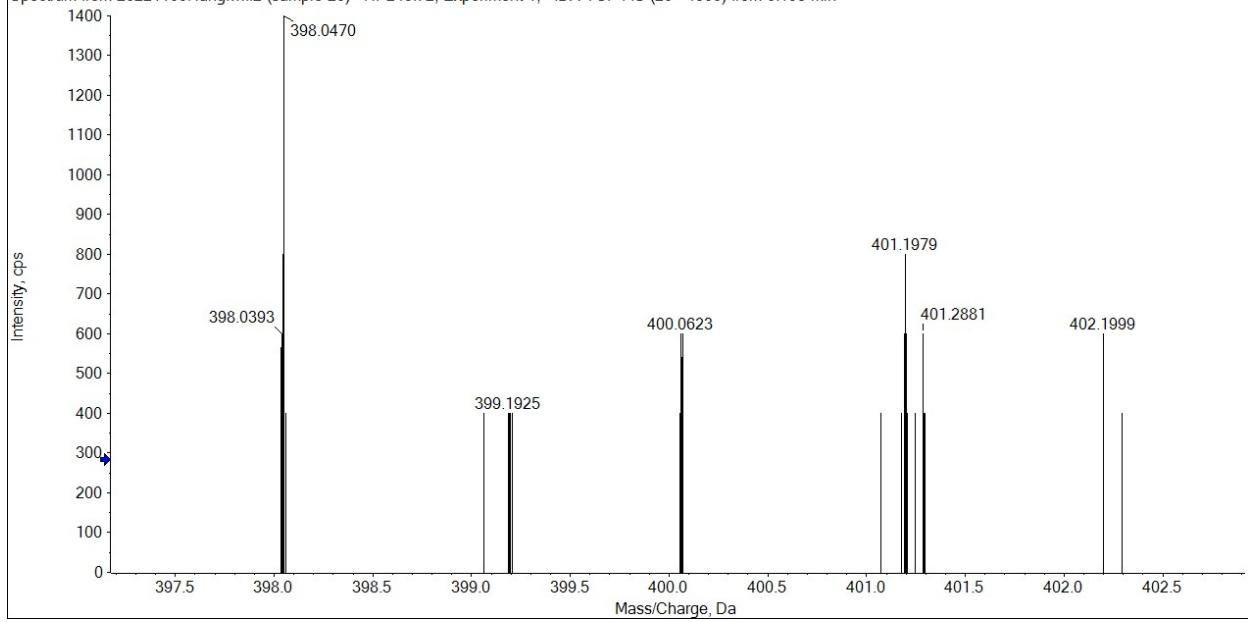
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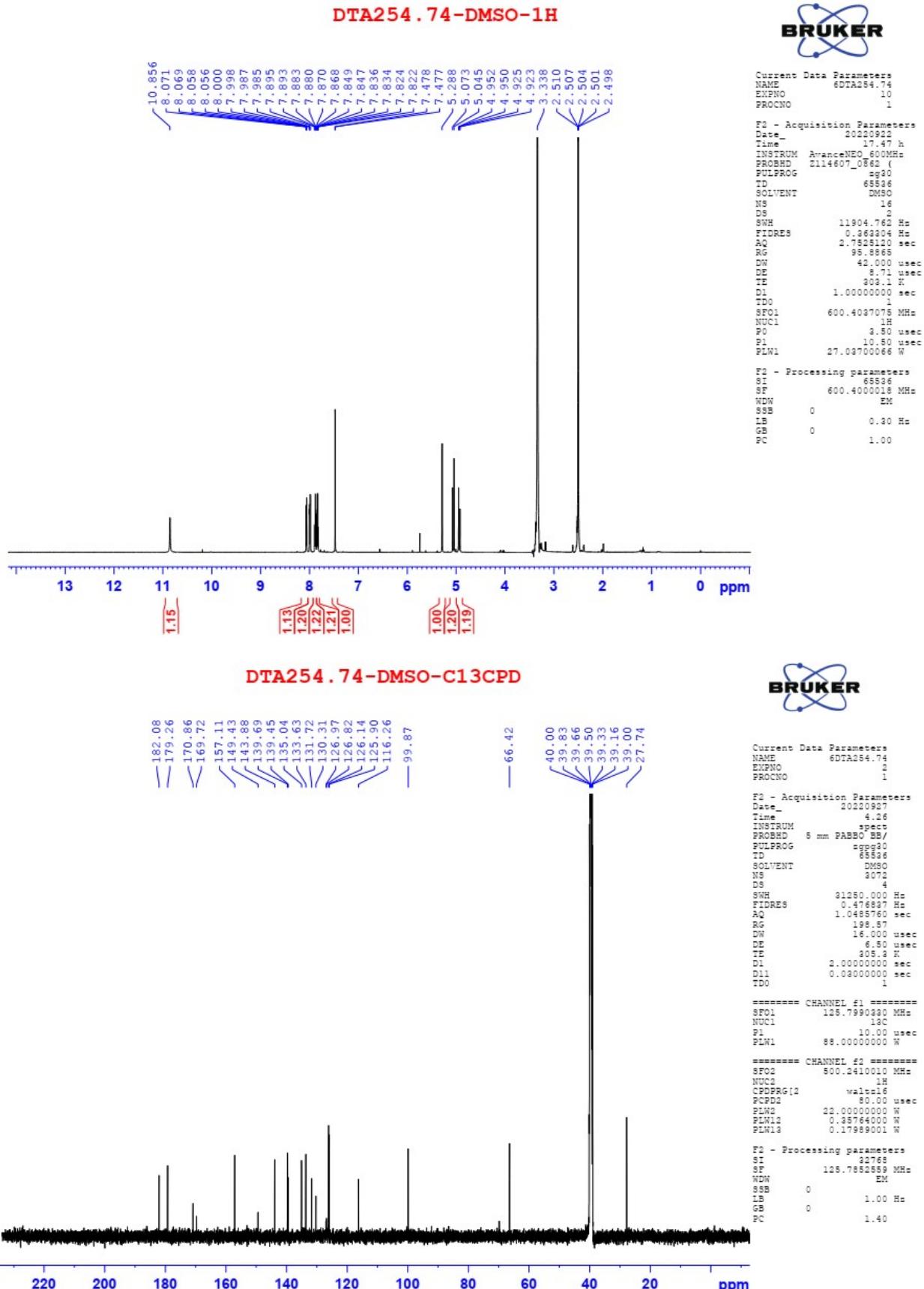
HP245.72-DMSO-C13CPD

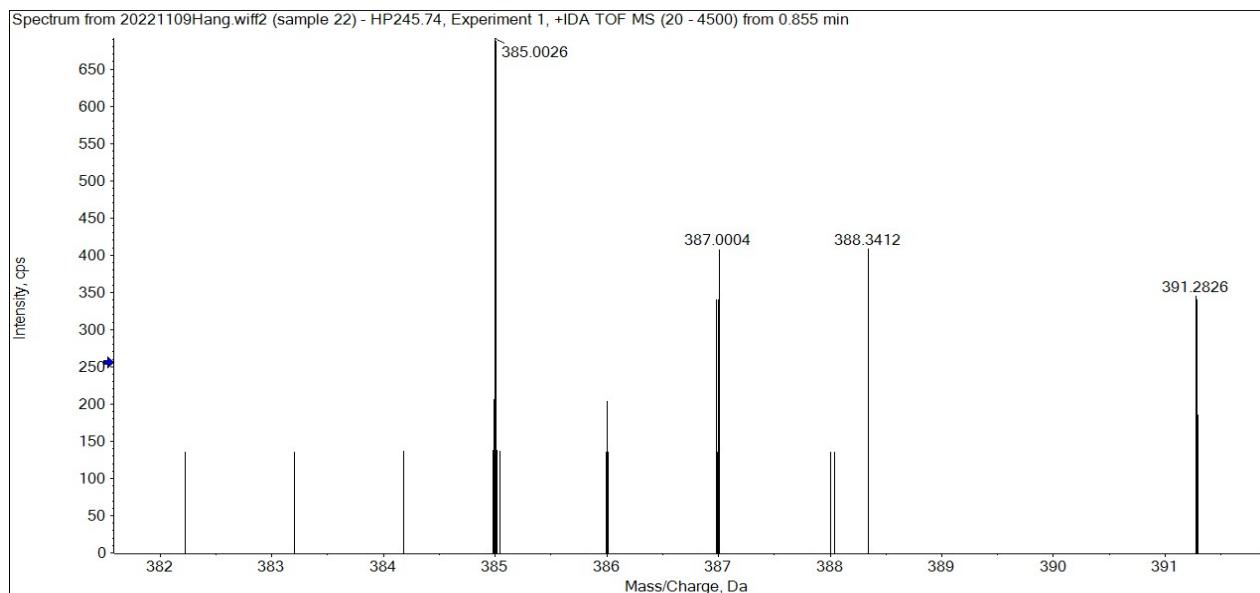


Spectrum from 20221109Hang.wiff2 (sample 20) - HP245.72, Experiment 1, +IDA TOF MS (20 - 4500) from 0.193 min



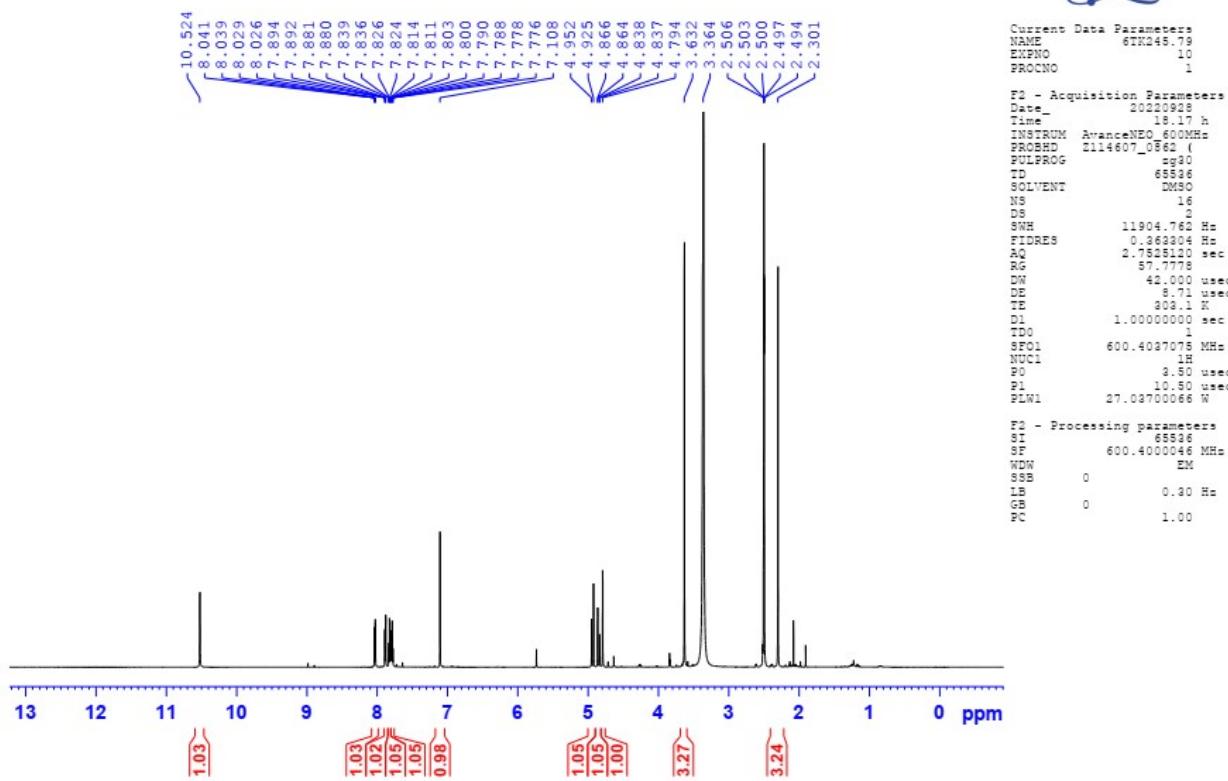
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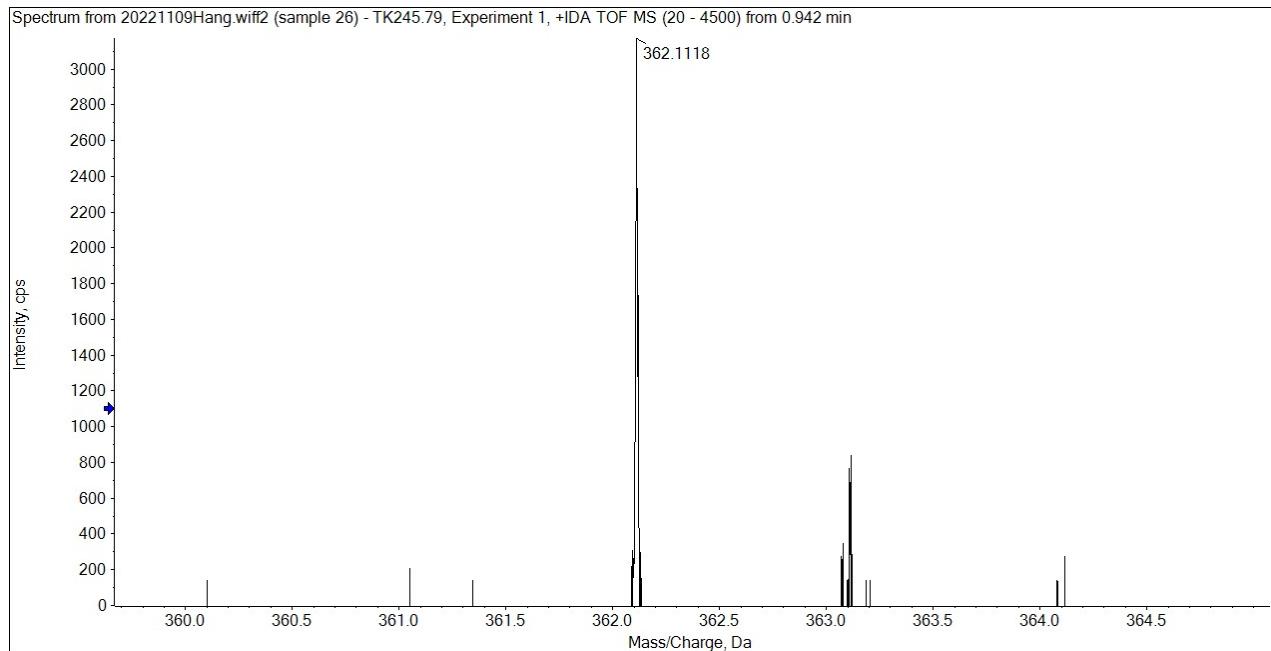
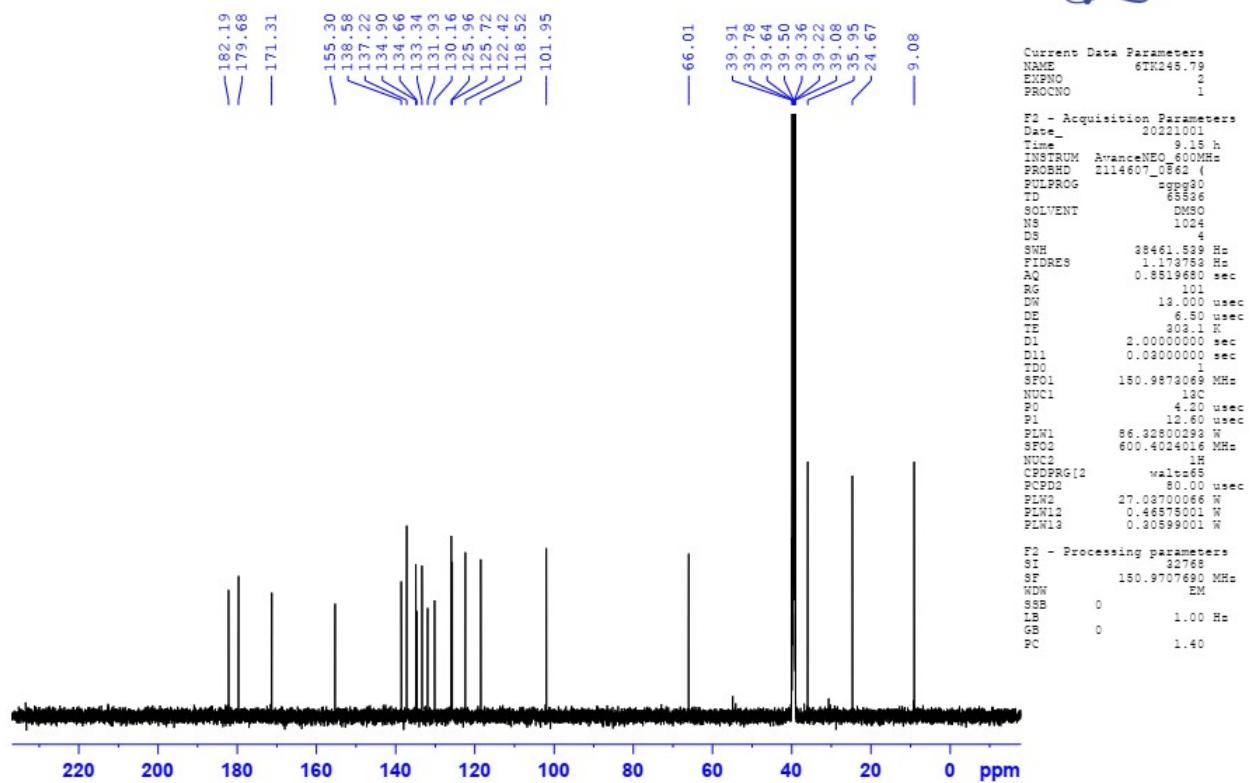


11-(1,5-Dimethyl-1H-pyrazol-4-yl)-4,11-dihydrobenzo[g]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13i)

TK245 . 79-DMSO-1H

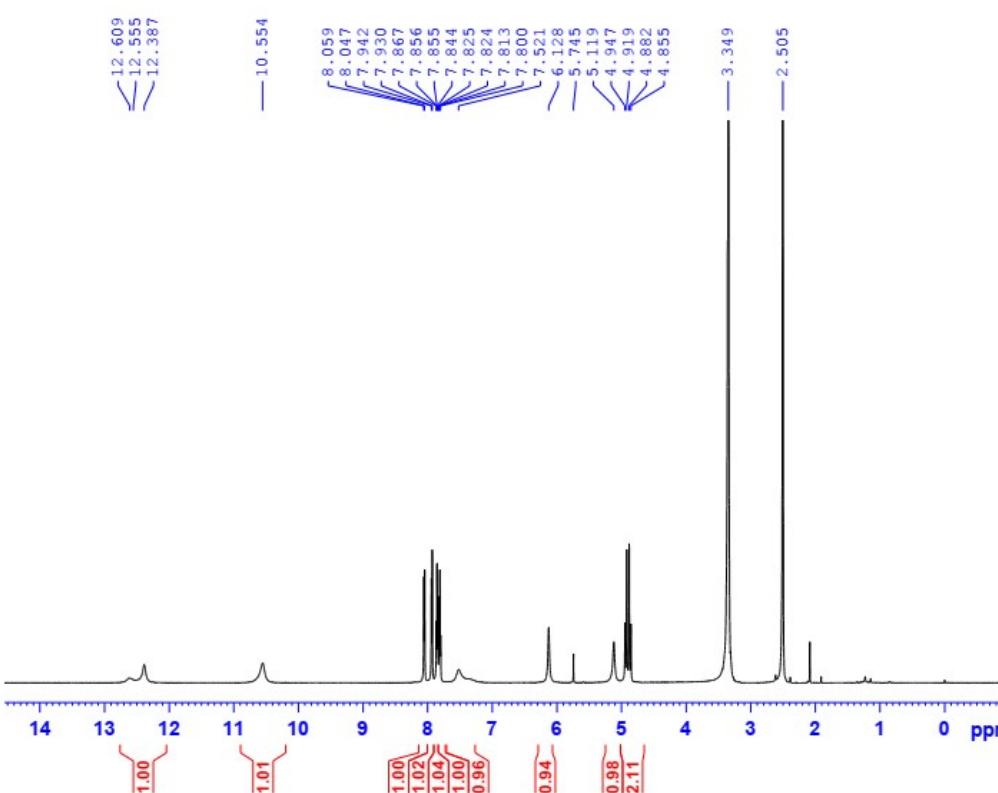


TK245.79-DMSO-C13CPD



11-(1H-pyrazol-5-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13j)

TK254 . 62-DMSO-1H

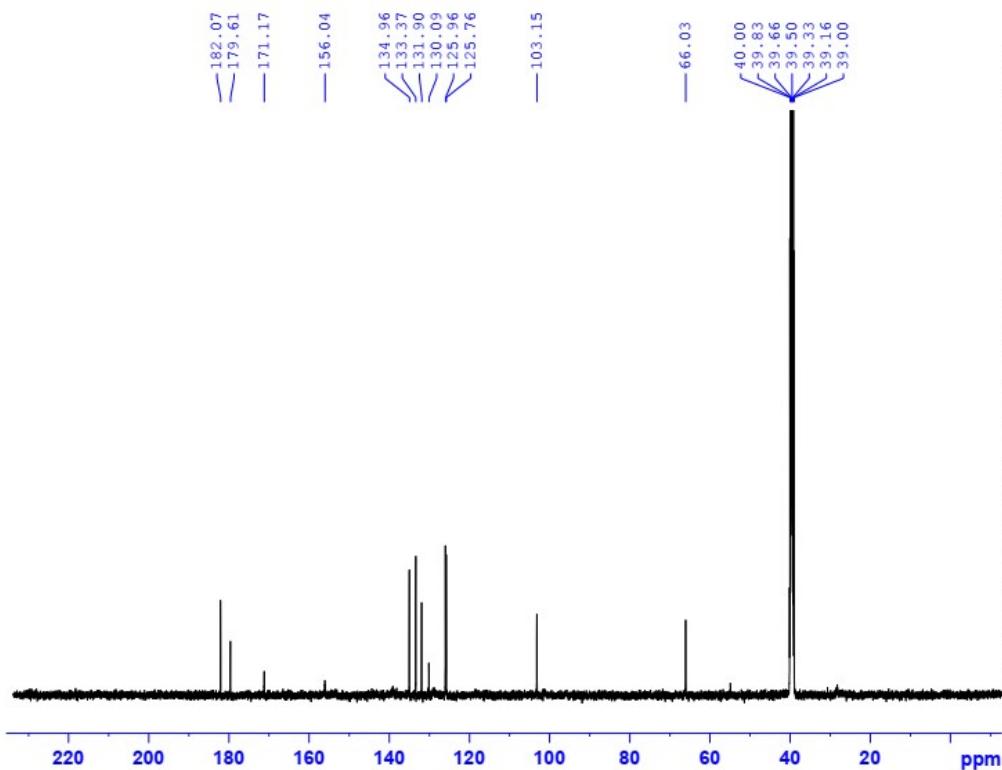


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PROCNO 1

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PROBHD Z114607_0862
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 11904.762 Hz
FIDRES 0.263304 Hz
AQ 2.7525120 sec
RG 94.8107
DW 42.000 usec
DE 8.71 usec
TE 303.1 K
D1 1.0000000 sec
TDO 1
SF01 600.4027075 MHz
NUC1 1H
PO 3.50 usec
PI 10.50 usec
PLW1 27.03700066 W

F2 - Processing parameters
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SSB 0
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GB 0
PC 1.00

TK254 . 62-DMSO-C13CPD



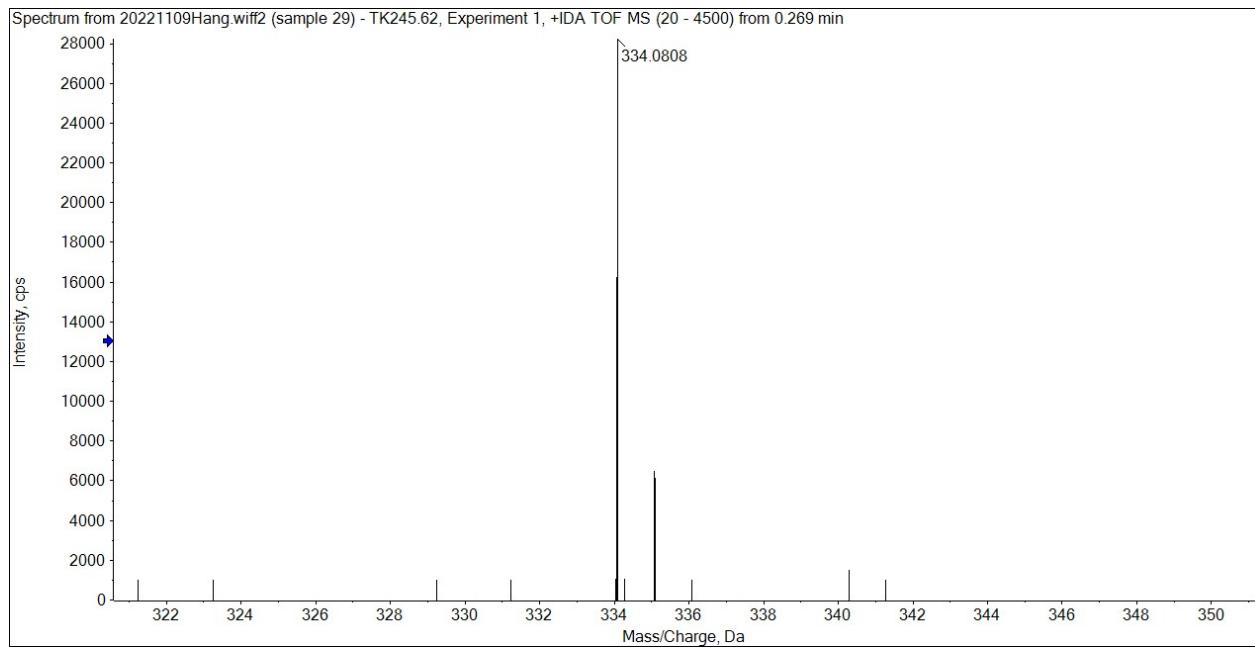
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EXPNO 10
PROCNO 1

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PULPROG zg30
TD 65536
SOLVENT DMSO
NS 2048
DS 4
SWH 81250.000 Hz
FIDRES 0.476587 Hz
AQ 1.0485760 sec
RG 198.57
DW 16.000 usec
DE 6.50 usec
TE 303.0 K
D1 2.0000000 sec
D11 0.03000000 sec
TDO 1

===== CHANNEL f1 =====
SF01 125.7990320 MHz
NUC1 13C
PI 10.00 usec
PLW1 88.00000000 W

===== CHANNEL f2 =====
SF02 600.2410010 MHz
NUC2 1H
CPDPRG[2] Waltz16
PCPD2 80.00 usec
PLW2 22.00000000 W
PLW12 0.38764000 W
PLW13 0.17989001 W

F2 - Processing parameters
SI 32768
SF 125.7852593 MHz
NDW 0 EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40



**11-(2-Methoxypyridin-4-yl)-4,11-dihydrobenzo[g]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione
(13k)**

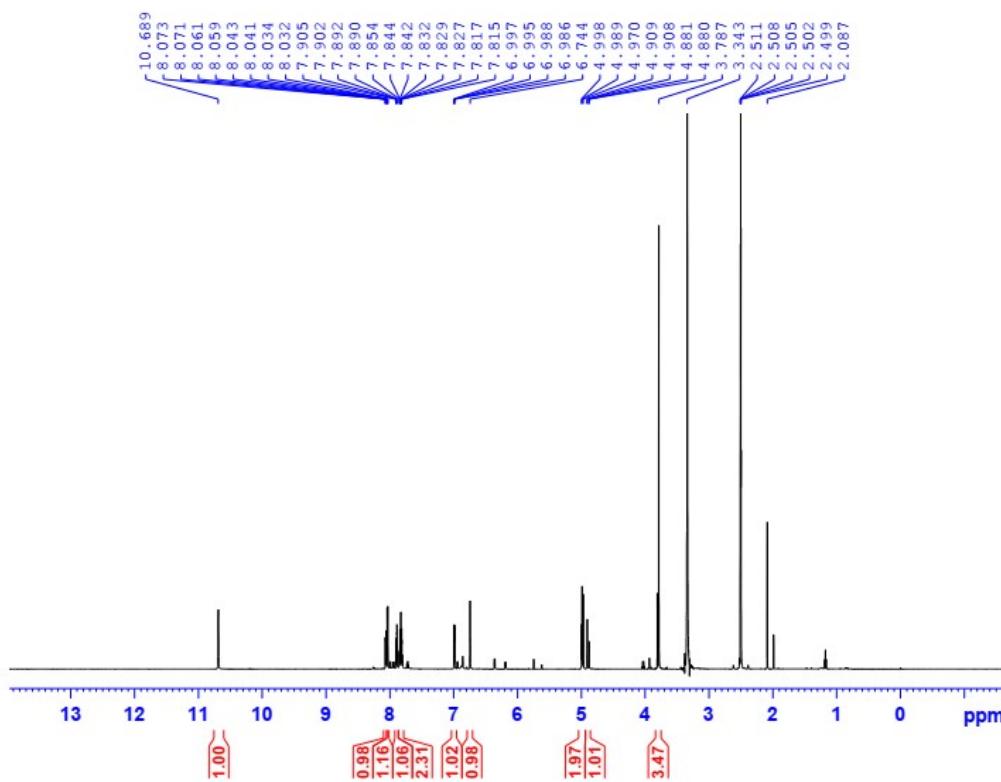
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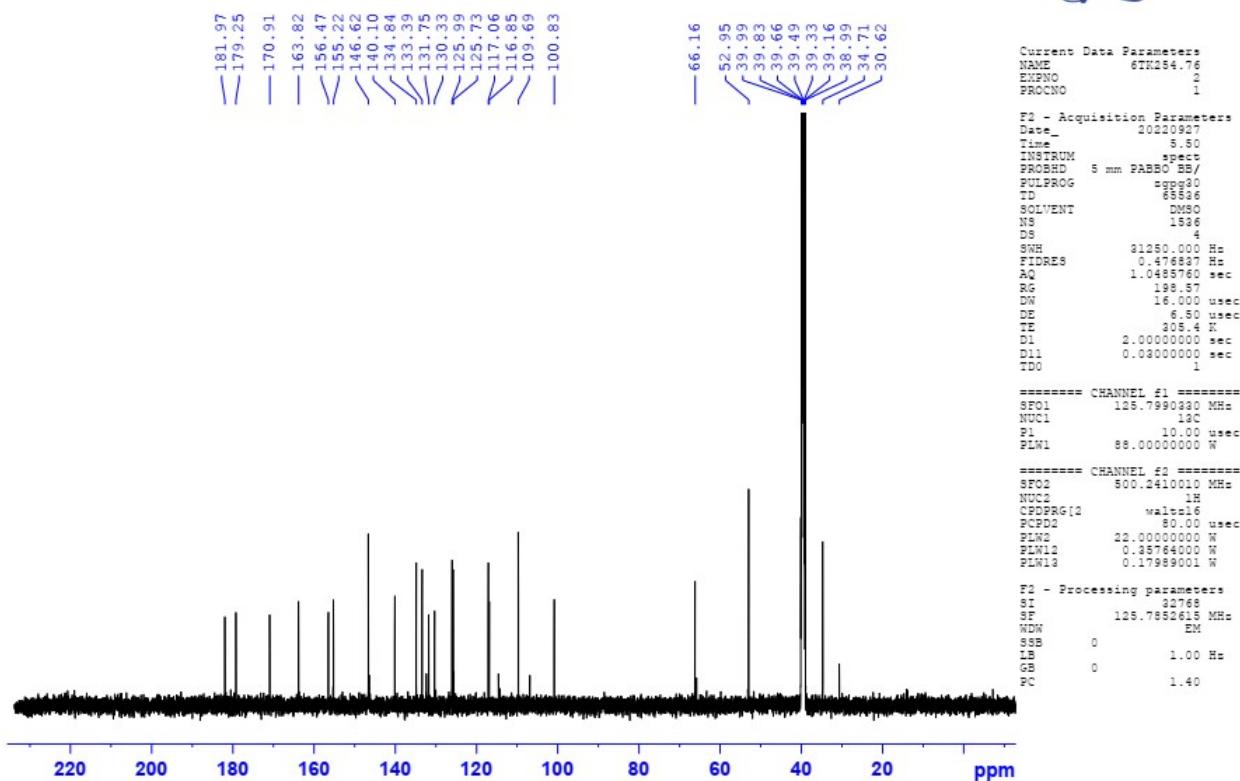
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EXPNO 10
PROCNO 1

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PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 0
SWH 11904.762 Hz
FIDRES 0.363204 Hz
AQ 2.7525120 sec
RG 101
DW 42.000 usec
DE 8.71 usec
TM 303.2 usec
D1 1.0000000 sec
TD0 600.4037078 MHz
SFO1 600.4037078 MHz
NUC1 1H
SS 3.80 usec
SI 10.00 usec
PLW1 27.037000968 Hz

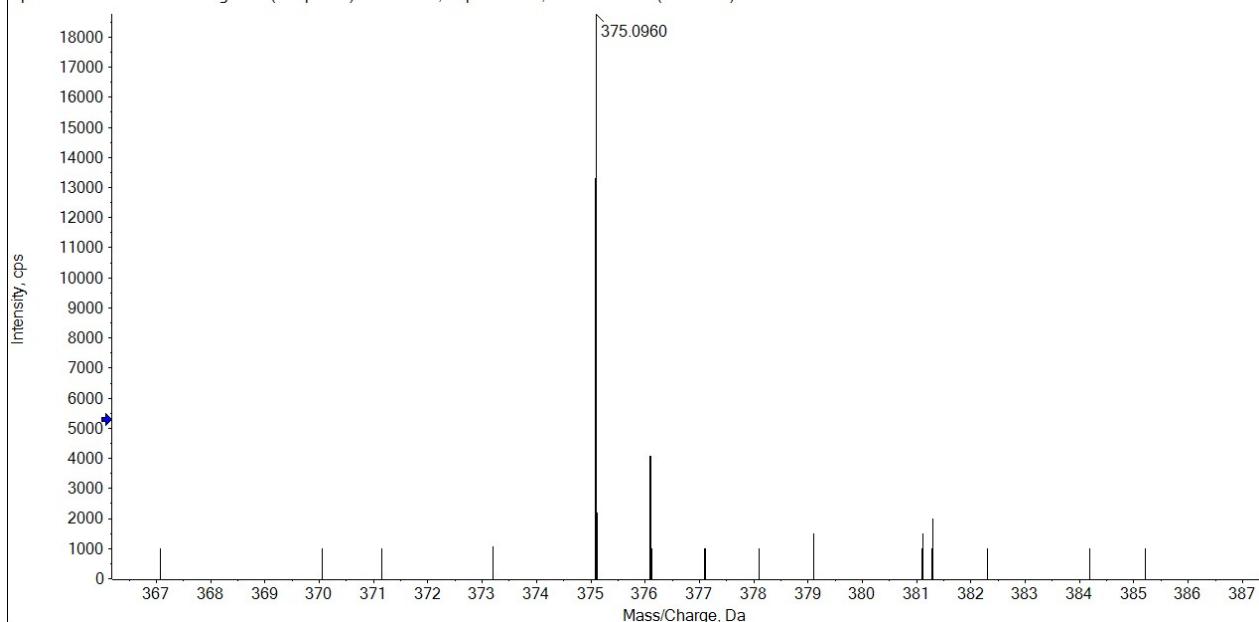
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WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



TK254 . 76-DMSO-C13CPD

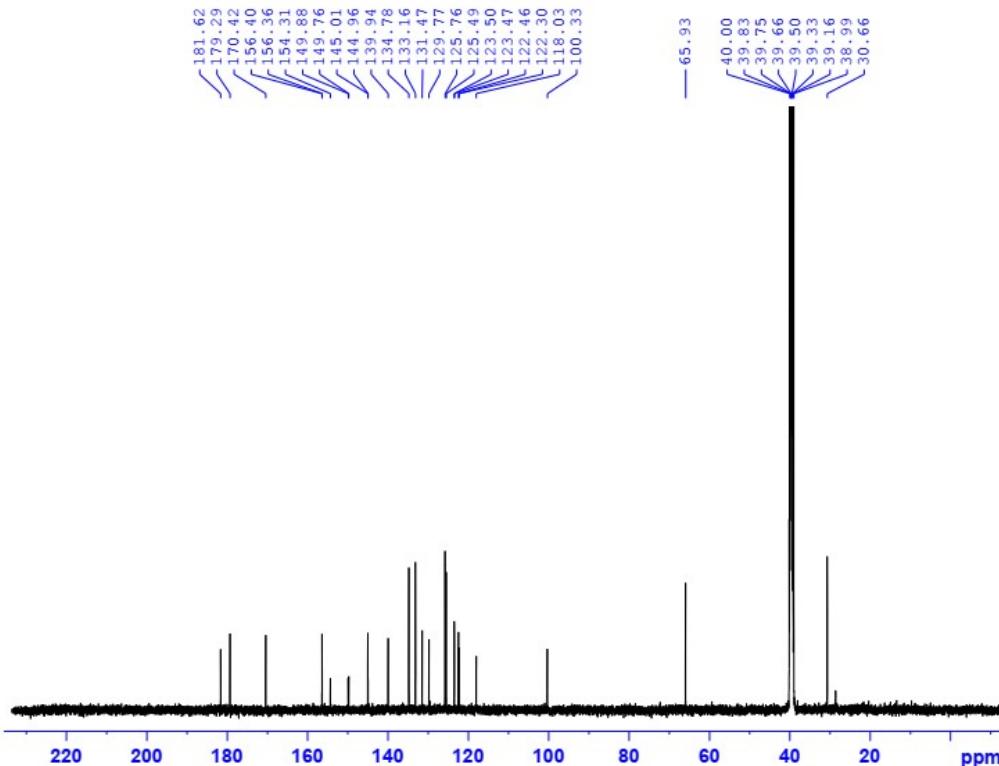
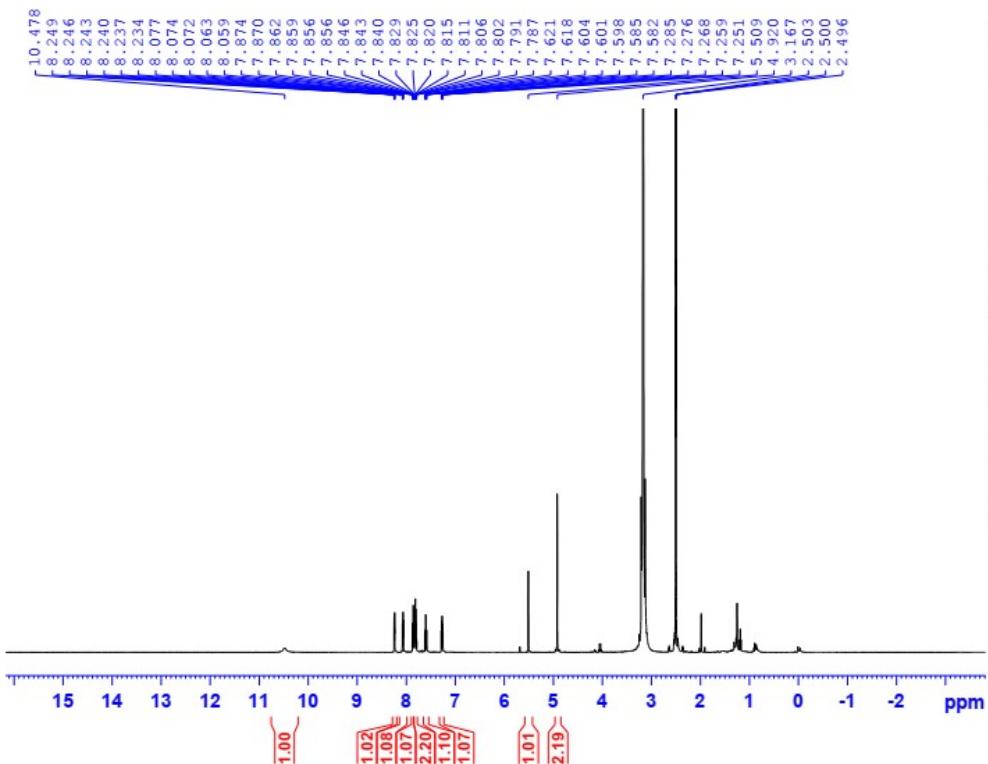


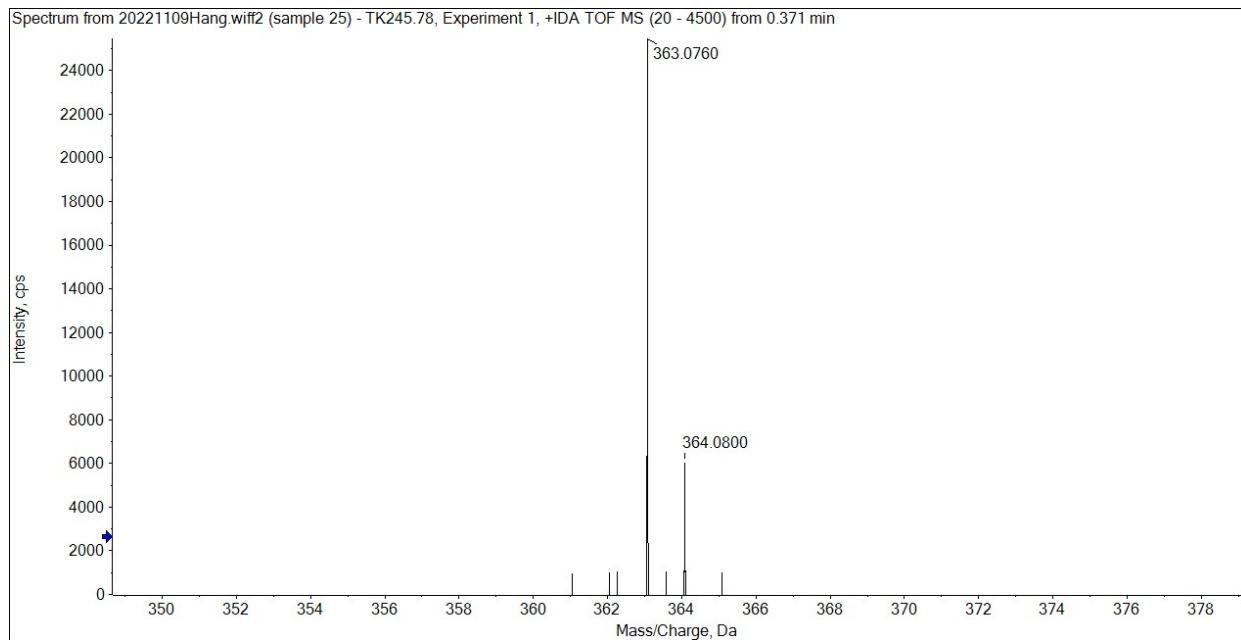
Spectrum from 20221109Hang.wiff2 (sample 24) - TK245.76, Experiment 1, +IDA TOF MS (20 - 4500) from 0.167 min



**11-(3-Fluoropyridin-2-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione
(13l)**

TK245.78.1-DMSO-1H





11-(2-Methoxy-5-(trifluoromethyl)pyridin-3-yl)-4,11-dihydrobenzo[*g*]furo[3,4-*b*]quinoline-1,5,10(3*H*)-trione (13m)

TK245.96-DMSO-1H



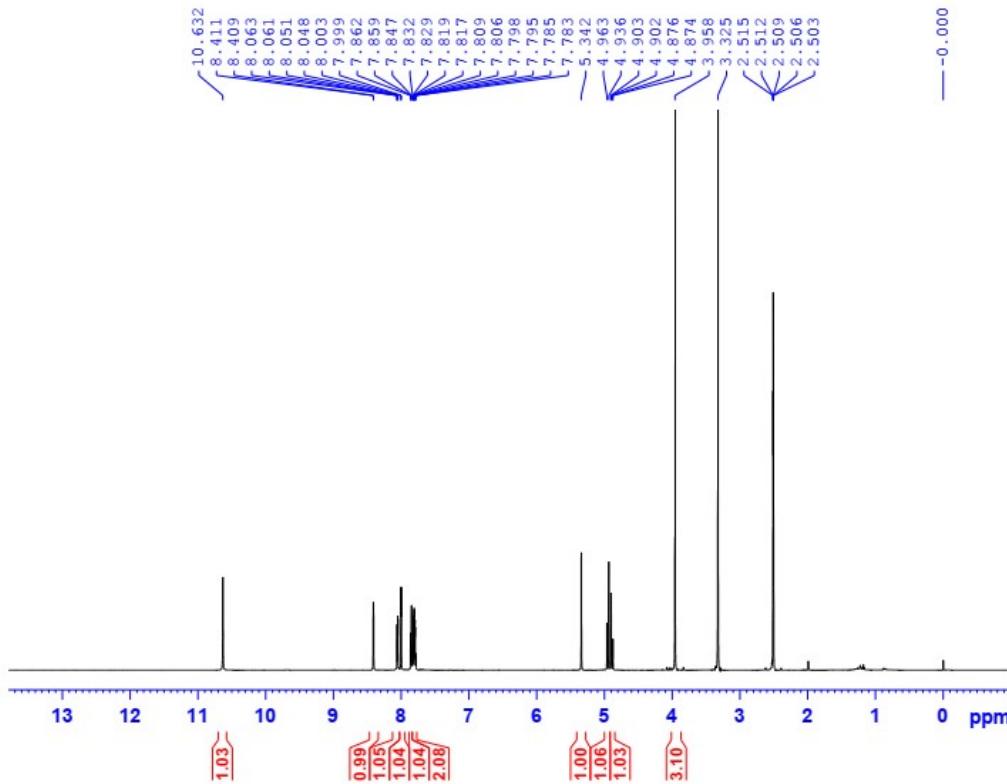
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Current Data Parameters
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PROCNO        1

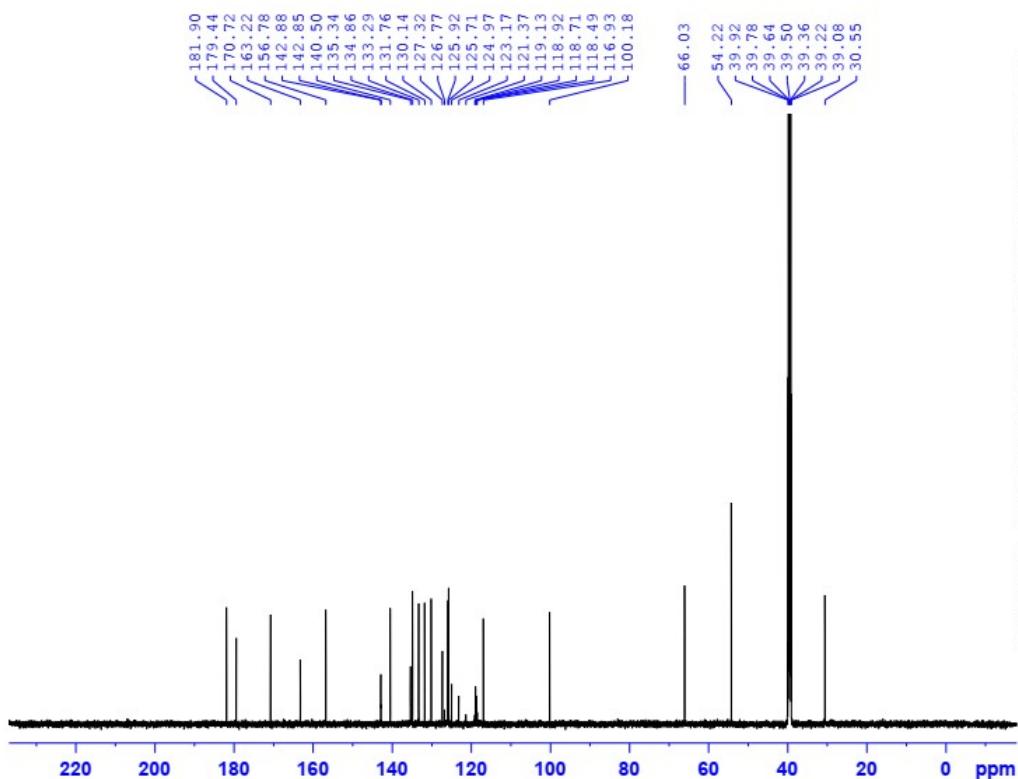
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Time       18.41 h
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PROBHD    C114607_0862
PULPROG   zg30
TD        65536
SOLVENT    DMSO
NS           16
DS            2
SWH        11904.762 Hz
FIDRES   0.083304 Hz
AQ        7.625000 sec
RG        2.752500 sec
TE        40.000 usec
DE        8.71 usec
TE        303.1 K
DL        1.00000000 sec
TDO        1
SF01      600.4037075 MHz
NUC1        1H
PO        3.50 usec
PL        10.50 usec
PLW1      27.03700066 W

FD - Processing parameters
SI           65536
SF        600.39999915 MHz
WDW        EM
SSB         0
LB        0.30 Hz
GB         0
PC        1.00

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TK245 . 96-DMSO-C13CPD



Spectrum from 20221109Hang.wiff2 (sample 28) - TK245.96, Experiment 1, +IDA TOF MS (20 - 4500) from 0.237 min

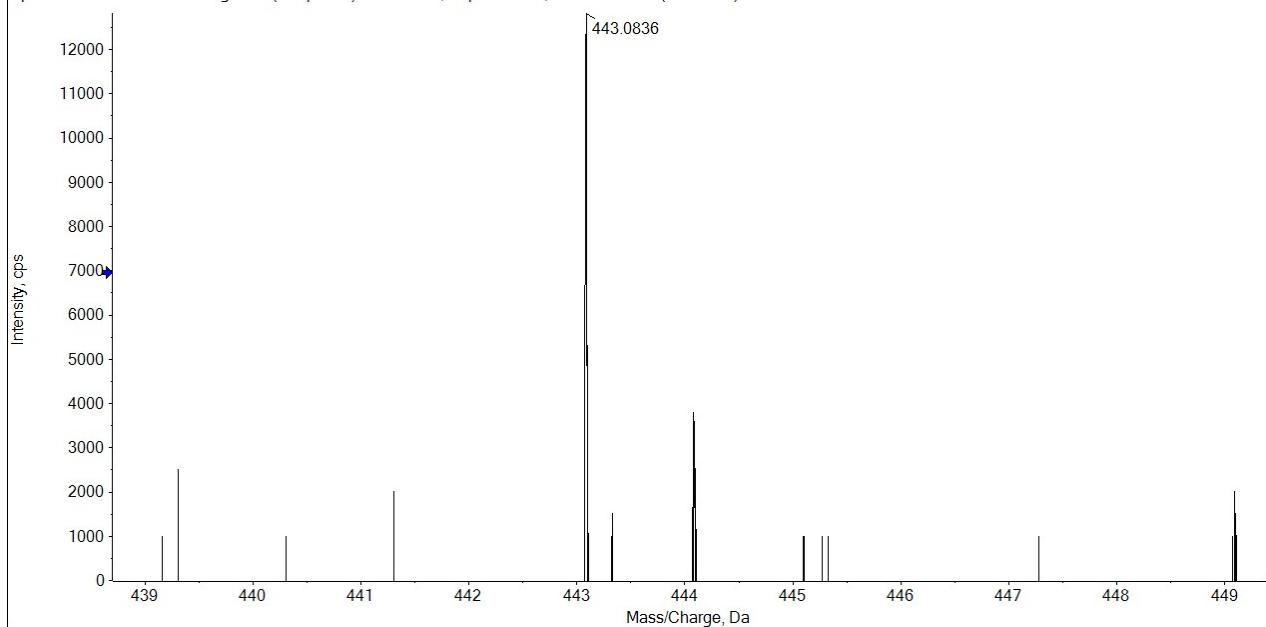


Table S1. Estimated of binding energy of docking complexes for **13a-m**, podophyllotoxin and colchicine.*

Compounds	Binding energy	Hbond	Hphob	VwInt	Eintl	Dsolv	SolEl	mfScore	dTSsc
13a	-16.38	-3.064	-6.433	-25.42	17.12	10.230	11.46	-75.37	0.9014
13b	-13.79	-2.741	-5.399	-25.26	16.97	9.797	10.81	-76.93	0.8058
13c	-14.55	-2.851	-5.708	-24.54	15.64	9.732	11.74	-75.70	0.8152
13d	-15.29	-2.934	-6.015	-24.47	18.18	9.686	10.38	-80.66	0.8435
13e	-14.09	-1.143	-5.254	-26.30	20.77	10.580	12.15	-88.50	0.9600
13f	-16.45	-3.189	-6.794	-25.84	21.89	12.820	10.85	-93.06	0.9395
13g	-15.08	-2.684	-5.401	-25.16	14.61	10.260	15.23	-87.20	1.0070
13h	-17.86	-3.206	-7.434	-27.08	9.12	9.591	11.77	-102.91	0.9055
13i	-13.83	-1.905	-6.337	-24.90	11.67	14.110	18.42	-79.77	1.2020
13j	-15.64	-2.877	-5.179	-24.42	15.74	9.646	10.97	-80.76	0.7939
13k	-18.45	-3.159	-7.445	-27.87	8.42	9.007	11.80	-103.10	0.8863
13l	-15.78	-3.070	-6.207	-23.93	19.26	10.880	9.36	-71.60	0.8769
13m	-16.80	-1.996	-6.526	-26.86	9.32	10.360	10.88	-74.05	0.9259
Podophyllotoxin	-18.68	-1.611	-7.660	-28.16	9.21	10.143	10.68	-103.40	0.9565
Colchicine	-19.18	-1.418	-8.605	-28.34	9.71	10.690	9.05	-104.00	0.9576

*Binding energy of tubulin-ligands (kcal/mol) computed via GBSA/MM-type scoring function; Hbond: Hbond is Hydrogen Bond energy; Hphob: the hydrophobic energy in exposing a surface to water; VwInt: the van der Waals interaction energy (sum of gc and gh van der waals); Eintl: internal conformation energy of the ligand; Dsolv: the desolvation of exposed H-bond donors and acceptors; SolEl: the solvation electrostatics energy change upon binding; mfScore: the potential of mean force score; mfScore: the potential of mean force score; dTSsc: Loss of entropy by the rotatable protein side-chains.