

## Design, Synthesis, Docking Studies of Novel Pyrazole-based Scaffolds and Their Evaluation as VEGFR2 Inhibitors in the Treatment of Prostate Cancer.

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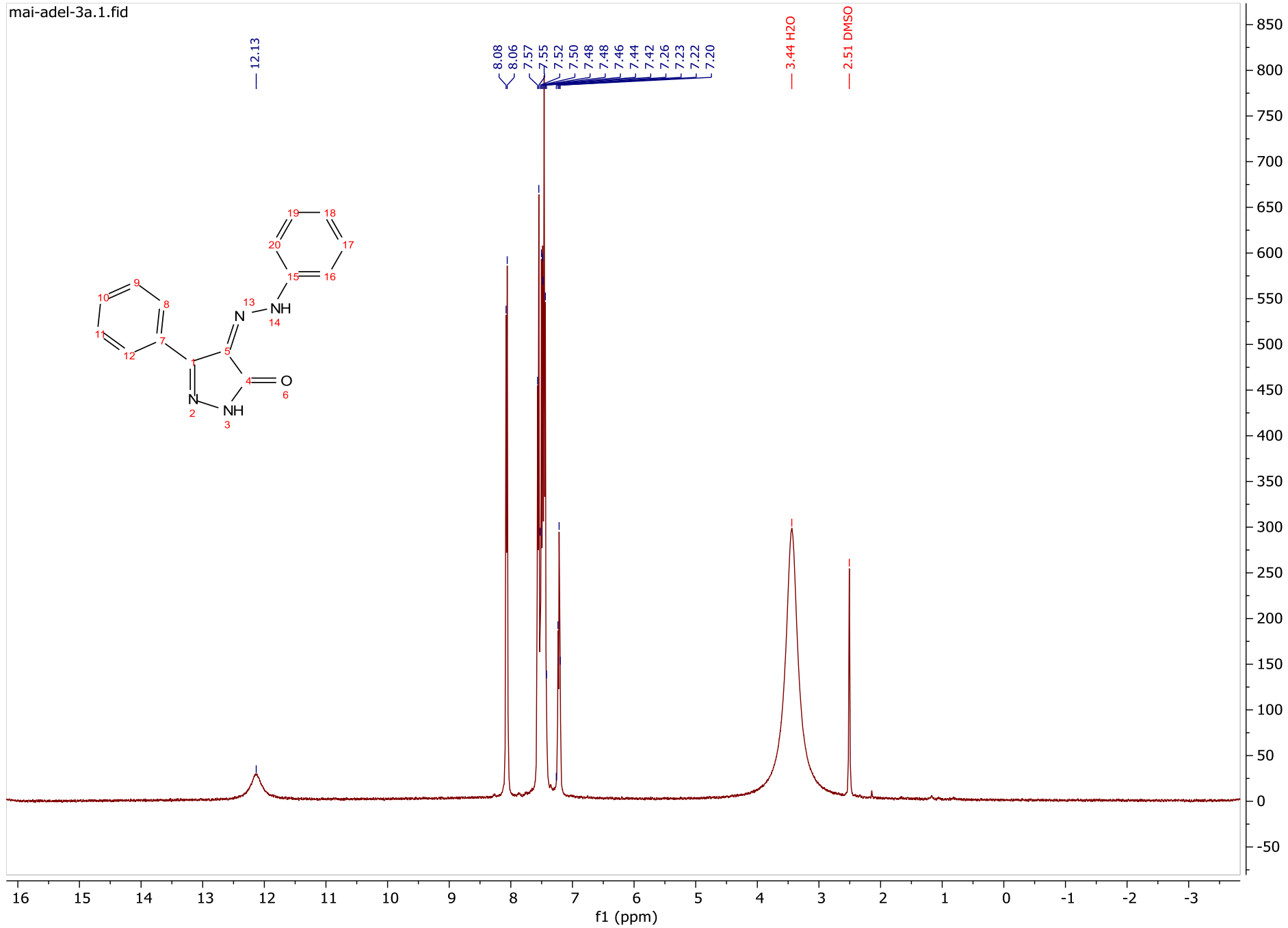
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<sup>3</sup>Department of Chemistry (Biochemistry program), Faculty of Science, Suez Canal University, Ismailia 41522, Egypt; [mohamed\\_nafie@science.suez.edu.eg](mailto:mohamed_nafie@science.suez.edu.eg)

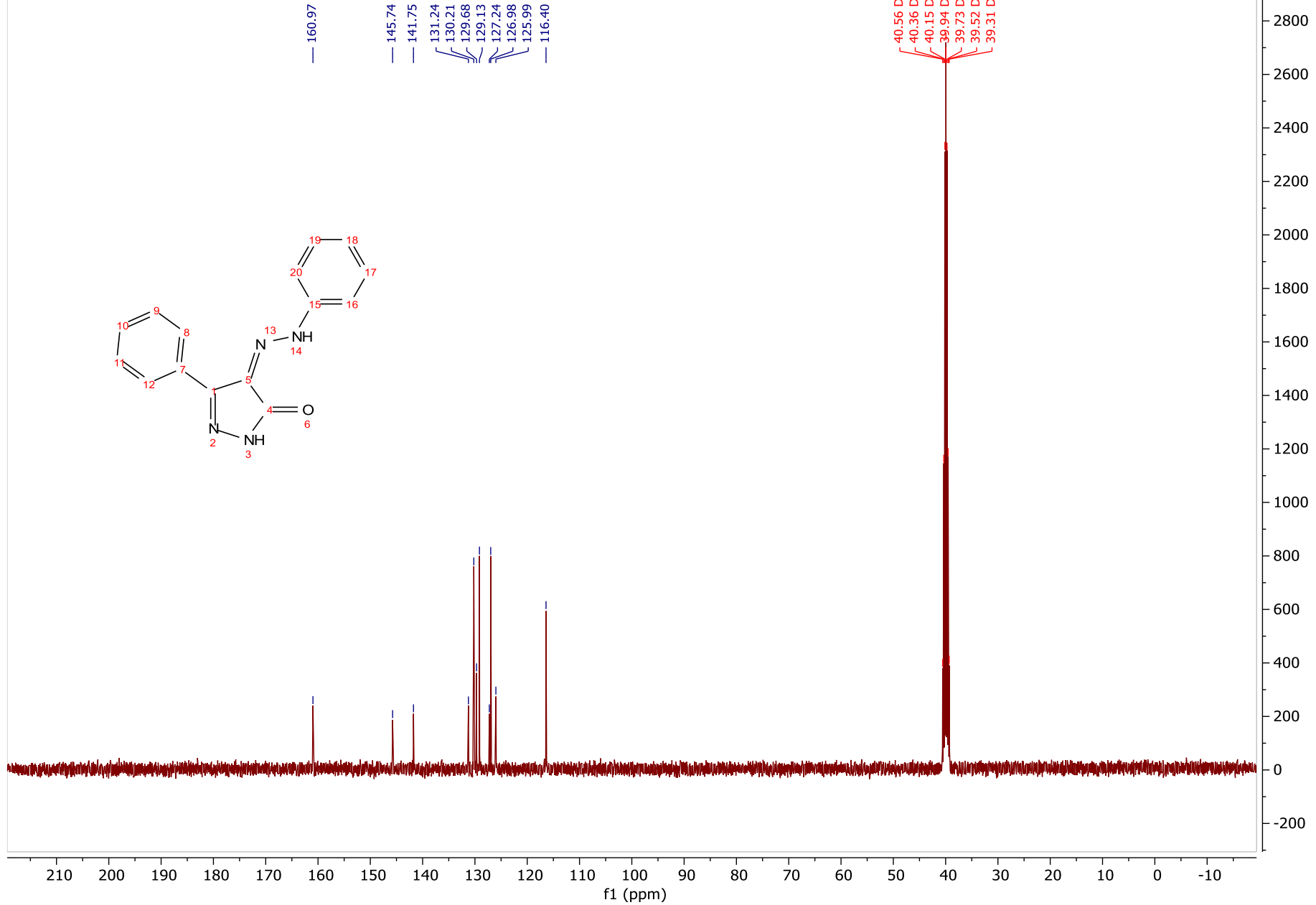
### List of supplementary

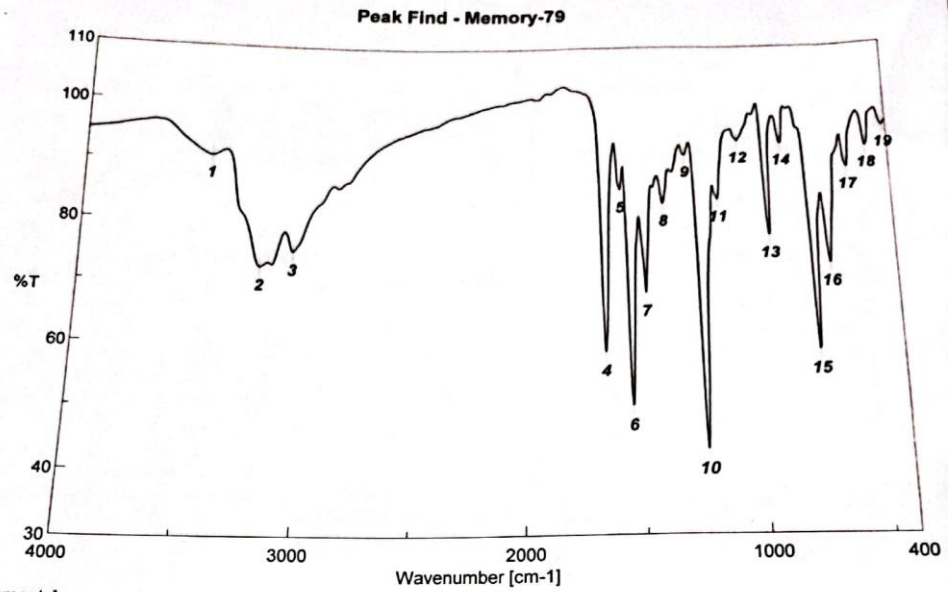
#	Content
1	Supplementary characterization for the tested compounds NMR, IR, Mass
2	Figure S1 & S2 (Docking part)
3	Detailed methodology for the in vivo experiment.

mai-adel-3a.1.fid



mai-adel-3a.2.fid

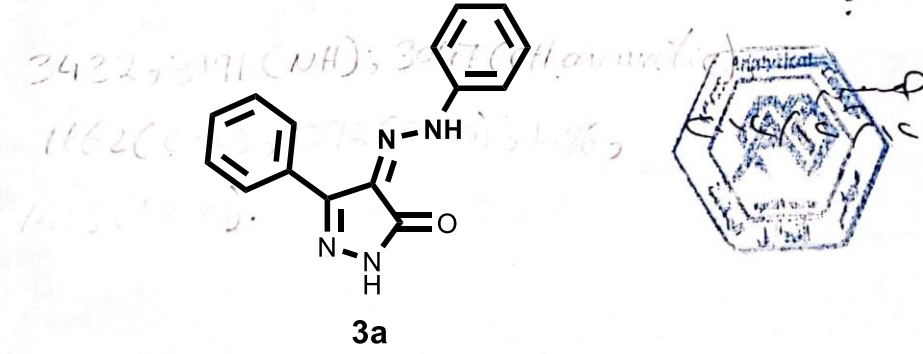




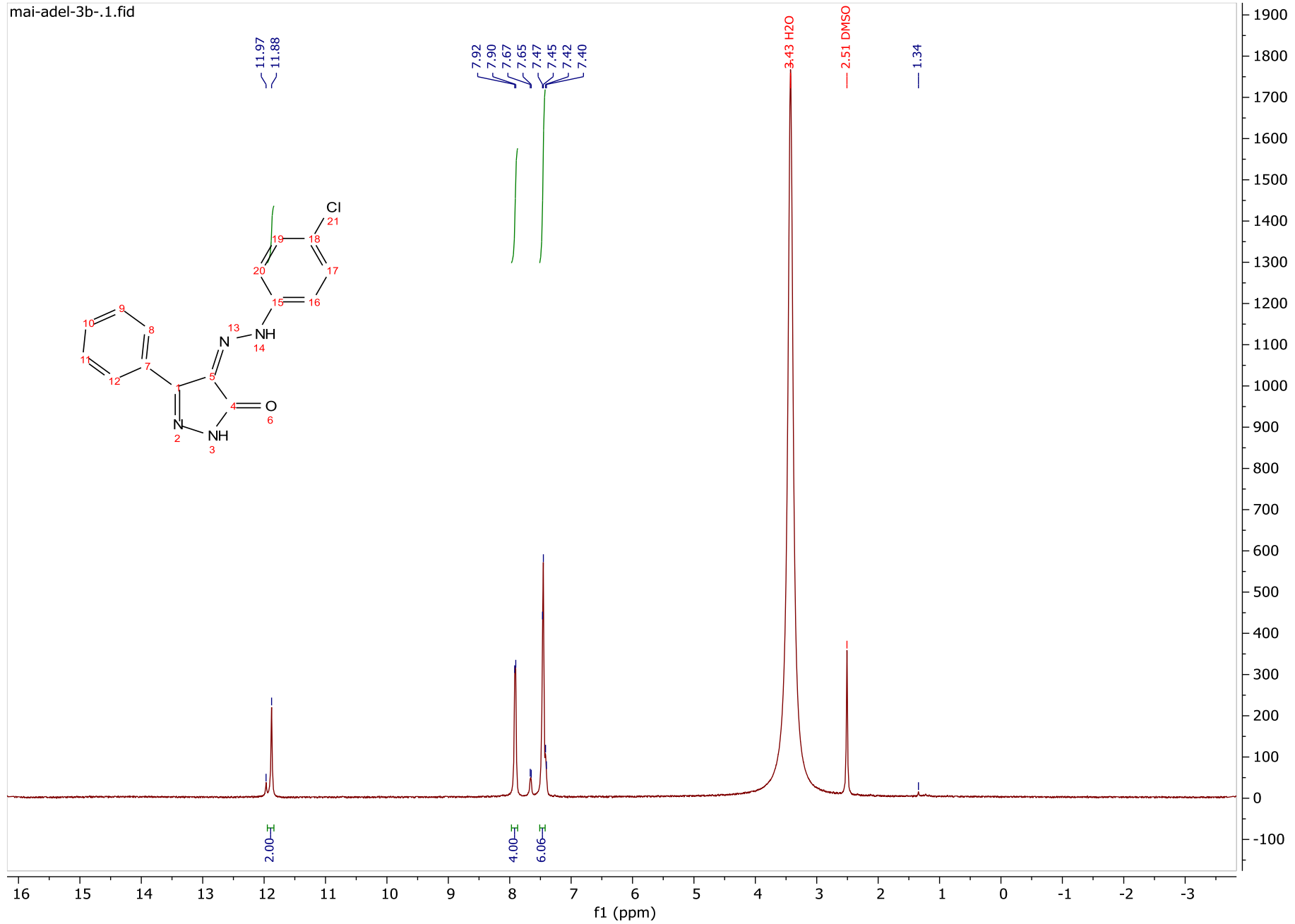
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 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

[ Result of Peak Picking ]

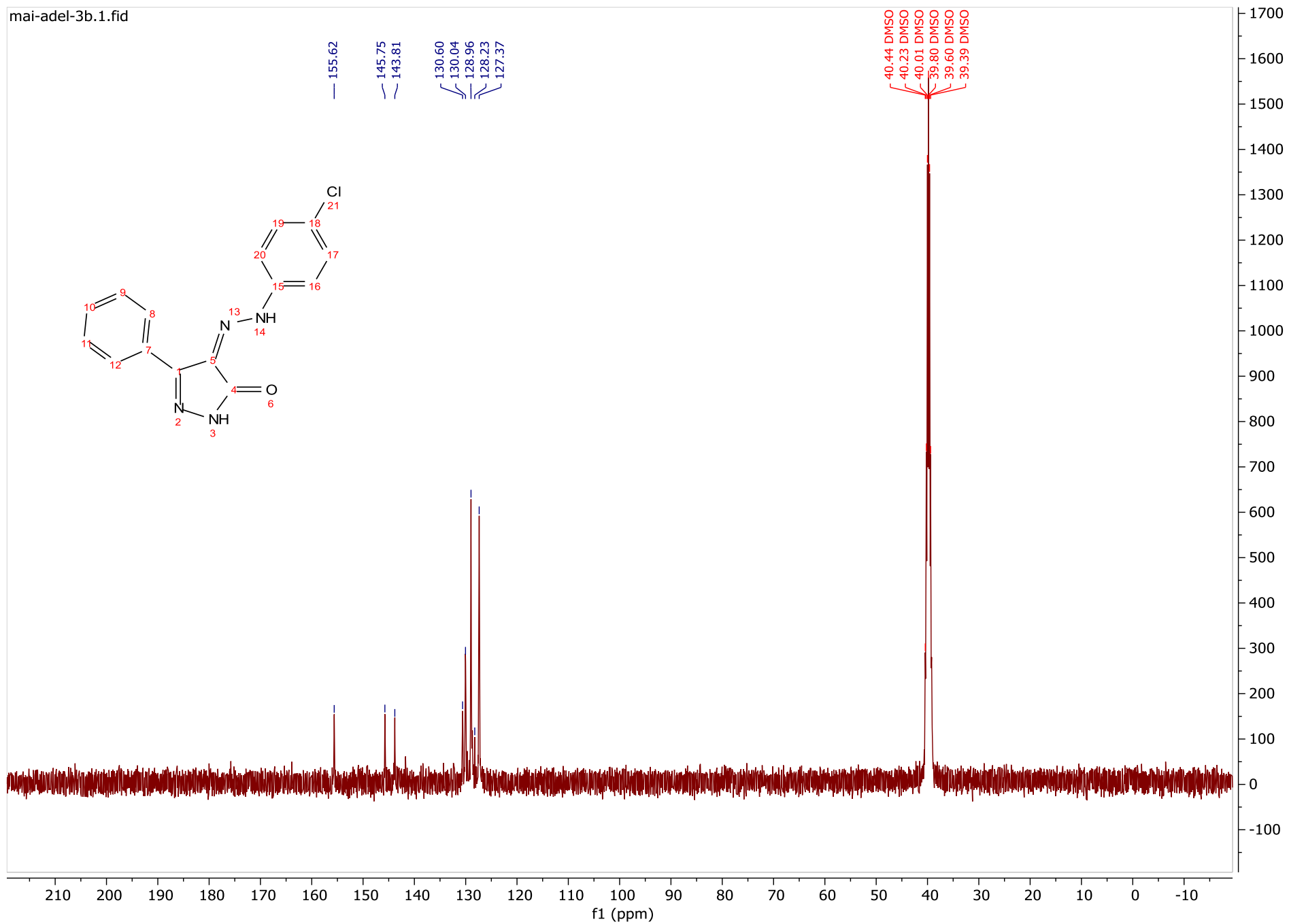
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4	1662.34	57.6248	5	1592.91	84.2517	6	1550.49	49.1939
7	1486.85	66.992	8	1405.85	81.722	9	1304.61	90.1484
10	1243.86	42.4676	11	1164.79	82.2924	12	1066.44	92.5169
13	943.985	76.2029	14	873.596	92.0836	15	755.959	57.5298
16	686.534	71.4918	17	585.29	87.8236	18	496.58	91.4794
19	416.549	95.0176						

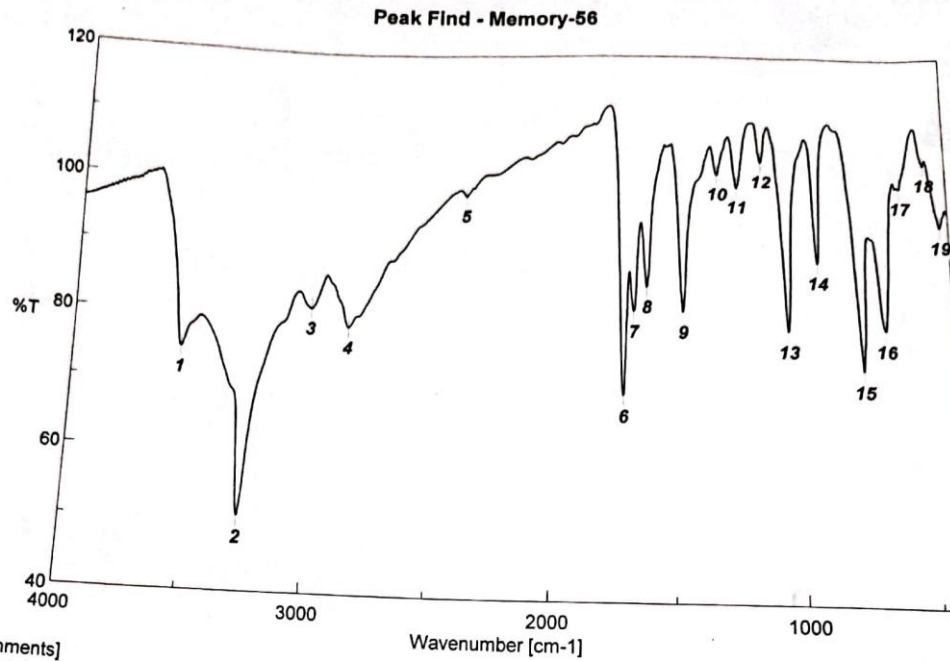
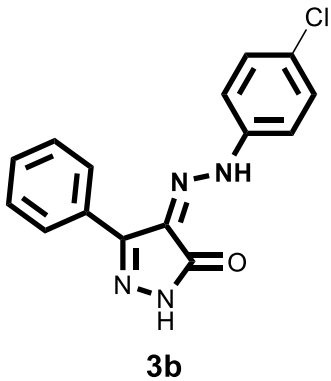


mai-adel-3b-.1.fid



mai-adel-3b.1.fid





[Comments]  
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 Company Micro Analytical Center

Result of Peak Picking ]

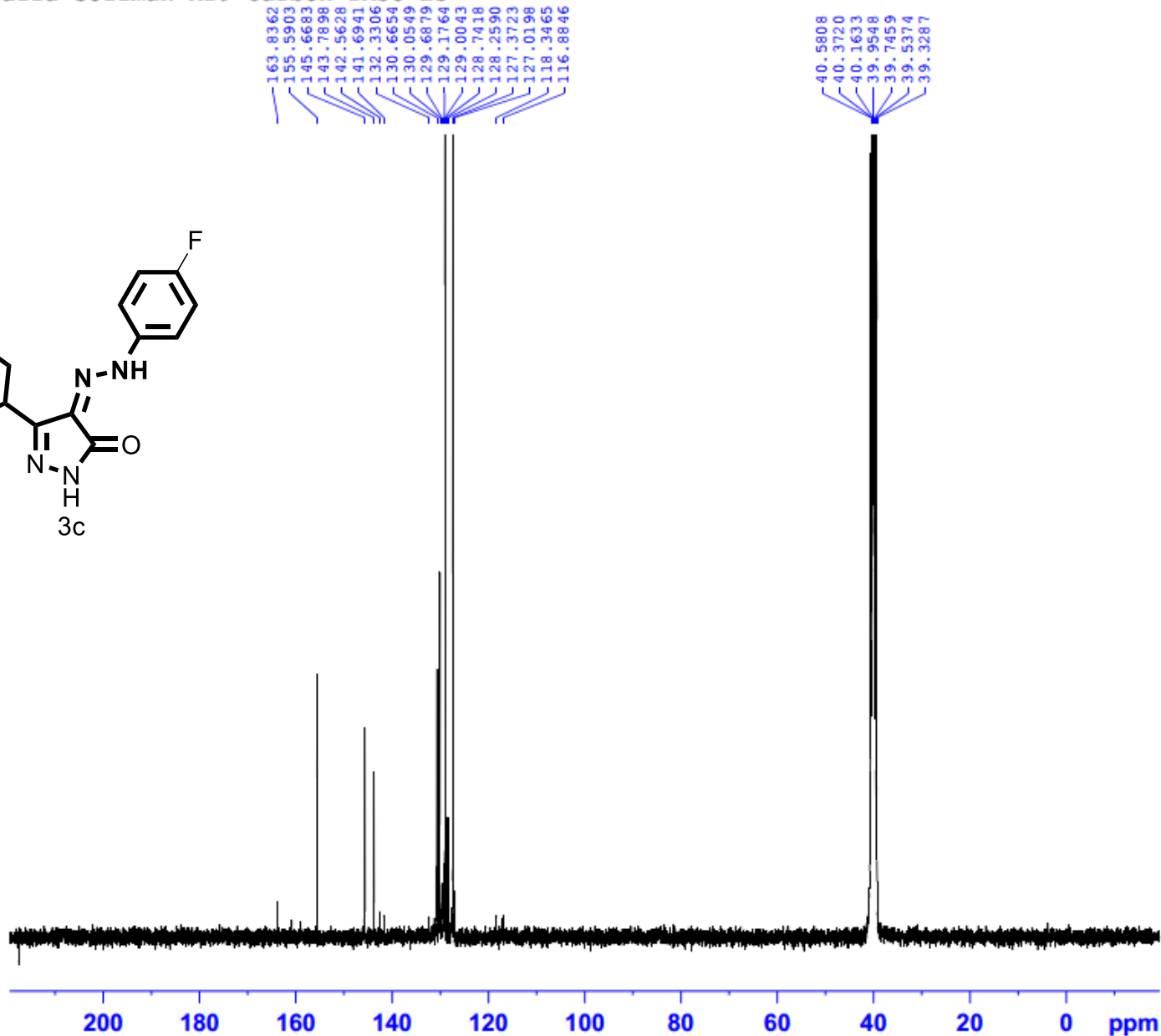
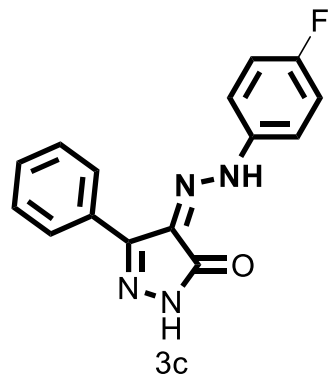
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2848.35	77.8252
1671.02	80.2547
1333.53	100.875
1056.8	77.4236
669.178	77.8957
433.905	93.5379

No.	Position	Intensity
2	3267.79	50.4431
5	2369.12	97.3773
8	1618.95	83.8268
11	1253.5	98.9569
14	932.414	87.636
17	589.147	99.3488

No.	Position	Intensity
3	3007.44	80.5939
6	1711.51	68.1974
9	1474.31	80.1612
12	1153.22	103.023
15	762.709	71.977
18	487.902	103.053



Dalia soliman-MD9-Carbon-DMSO-ES



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 PROCNO 1

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 PULPROG zgpg30  
 TD 65536  
 SOLVENT DMSO  
 NS 2200  
 DS 4  
 SWH 24038.461 Hz  
 FIDRES 0.733596 Hz  
 AQ 1.3631488 sec  
 RG 197.77  
 SW 20.800 usec  
 DE 6.50 usec  
 TE 295.9 K  
 D1 2.0000000 sec  
 D11 0.0300000 sec  
 TD0 1  
 SFO1 100.6404331 MHz  
 NUC1 13C  
 P1 10.00 usec  
 PLM1 47.0000000 W  
 SFO2 400.2016008 MHz  
 NUC2 1H  
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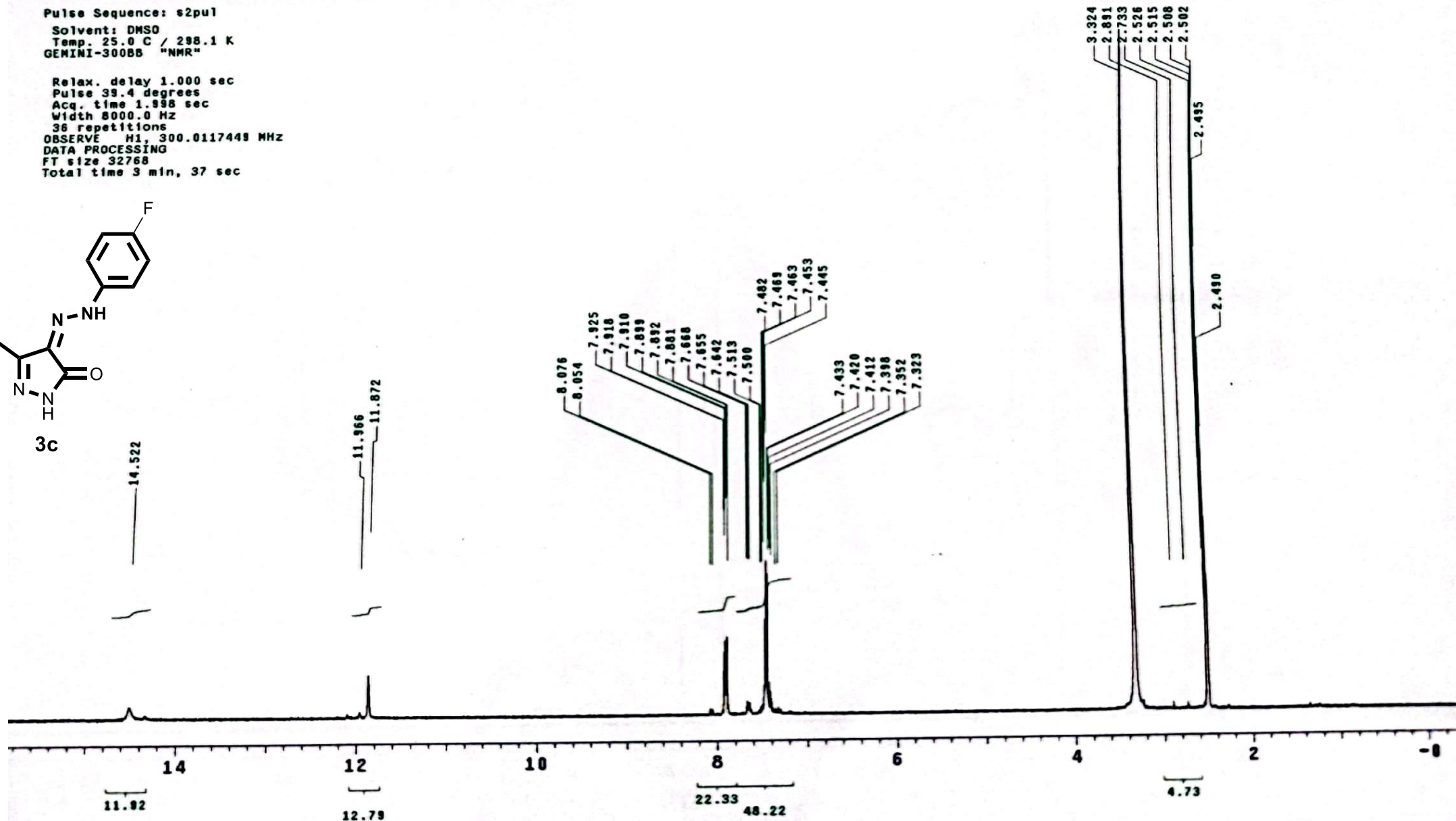
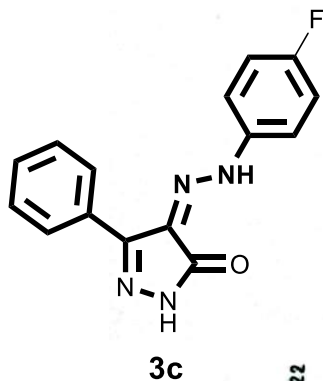
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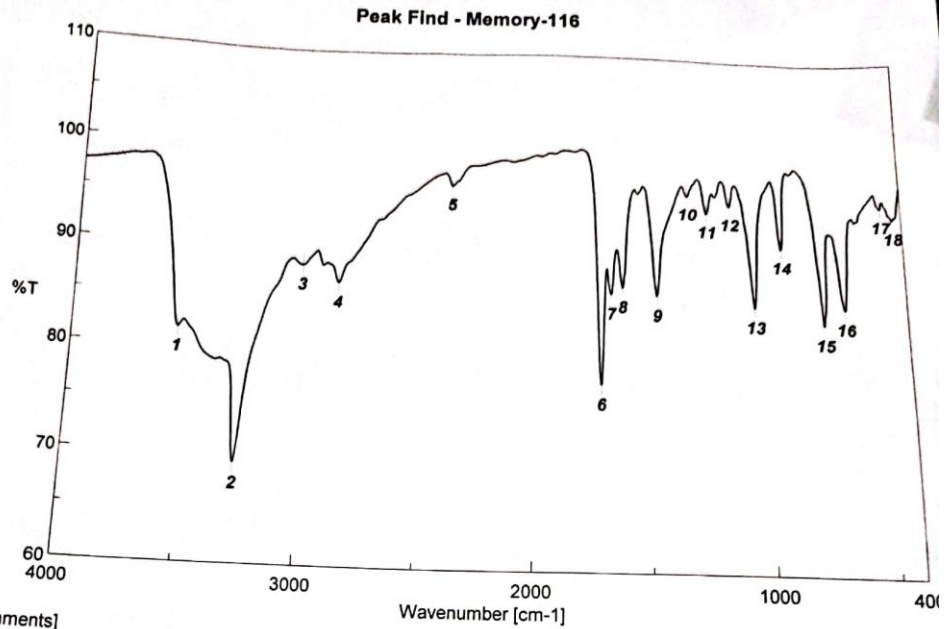
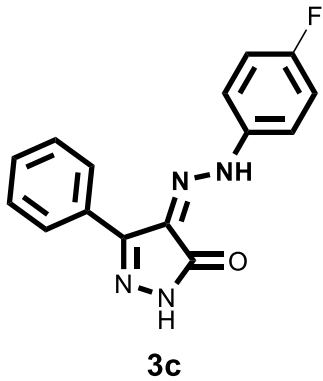


Dr.DallaSoliman-MD9-H1-DMSO-Main.Defence.Chemical.Laboratory

Pulse Sequence: s2pul  
Solvent: DMSO  
Temp. 25.0 C / 298.1 K  
GEMINI-300BB "NMR"

Relax. delay 1.000 sec  
Pulse 39.4 degrees  
Acq. time 1.998 sec  
Width 8000.0 Hz  
36 repetitions  
OBSERVE H1, 300.0117448 MHz  
DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec





[Comments]  
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 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

Result of Peak Picking ]

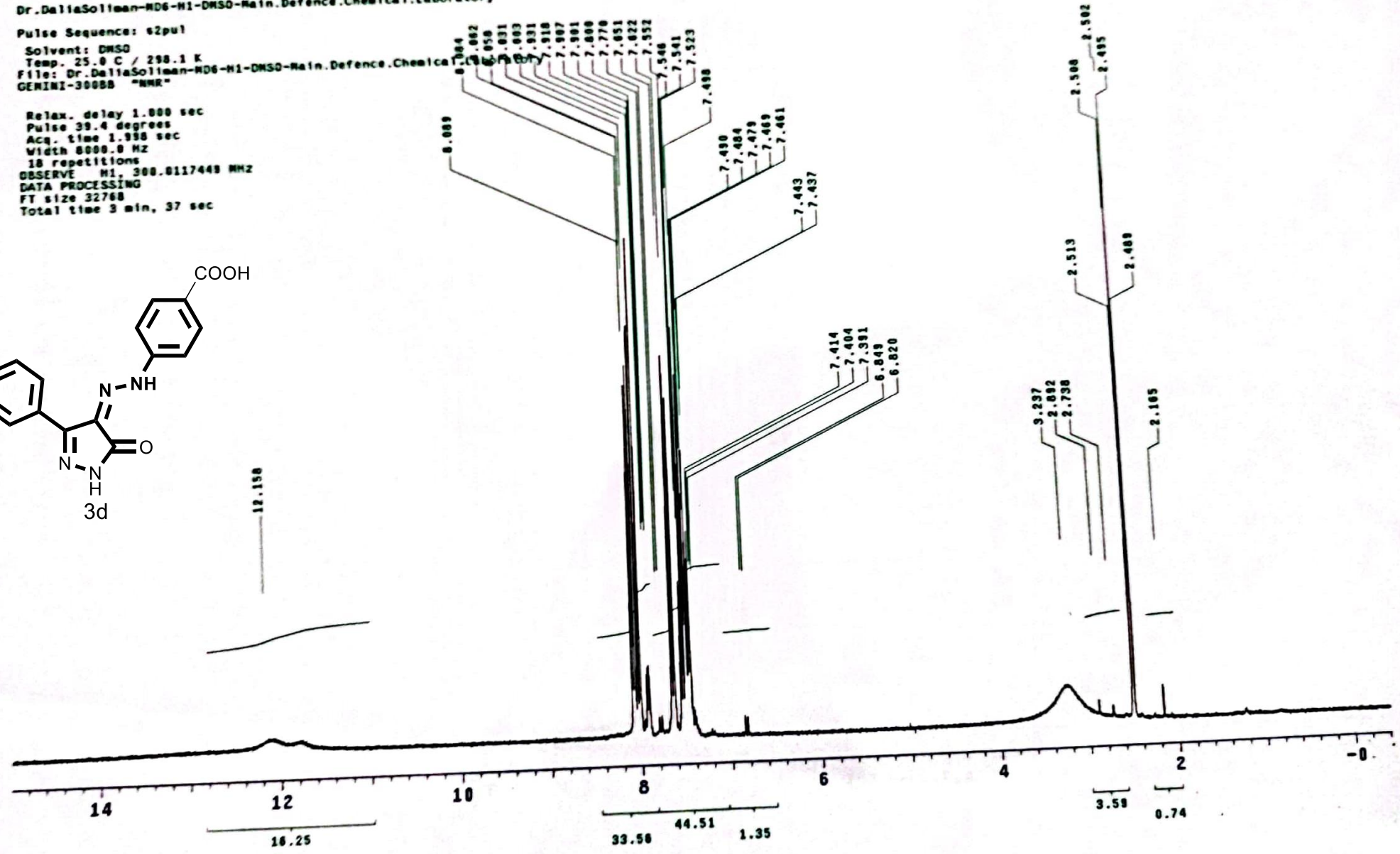
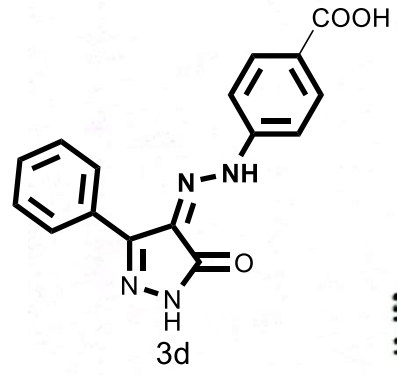
No.	Position	Intensity	No.	Position	Intensity	No.	Position	Intensity
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2	2853.17	86.1914	5	2365.26	95.9361	6	1713.44	76.7895
3	1670.05	85.3349	8	1621.84	85.9855	9	1475.28	85.2198
4	1337.39	95.3479	11	1254.47	93.5837	12	1154.19	94.4152
5	1057.76	84.2929	14	934.342	90.2879	15	761.744	82.8141
6	668.214	84.4367	17	487.902	94.8503	18	438.726	93.6942



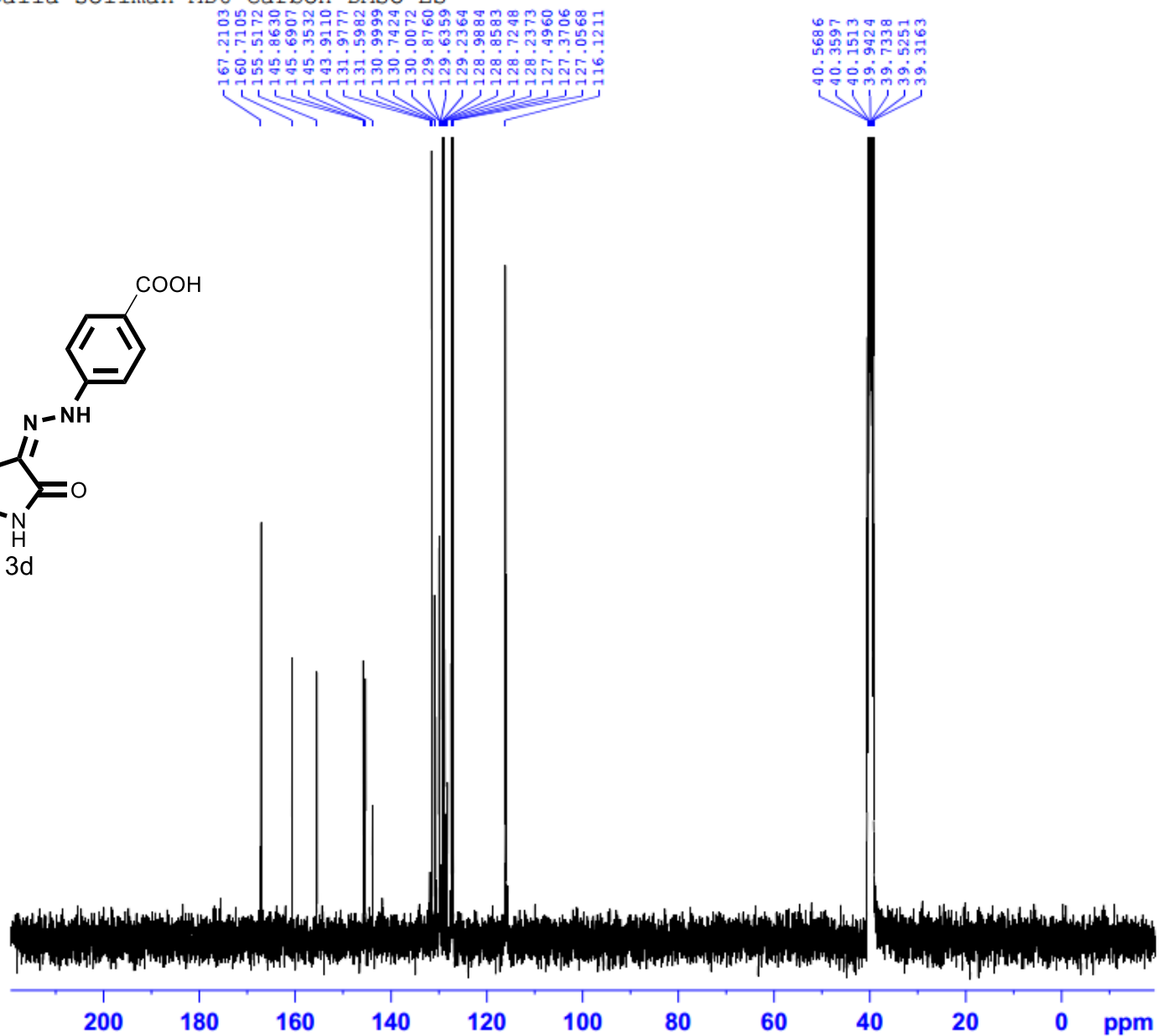
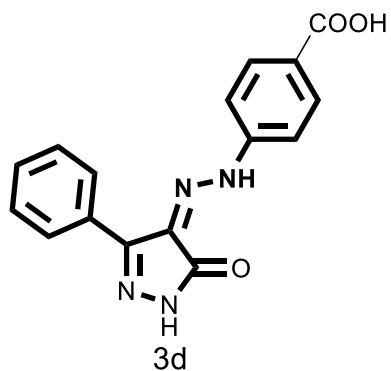
Dr. Dalia Soliman-ND6-H1-DMSO-Rain. Defence. Chemical. Laboratory

Pulse Sequence: s2pul  
Solvent: DMSO  
Temp: 25.0 C / 298.1 K  
File: Dr. Dalia Soliman-ND6-H1-DMSO-Rain. Defence. Chemical. Laboratory  
GENINI-30088 "NMR"

Relax. delay 1.000 sec  
Pulse 39.4 degrees  
Acq. time 1.998 sec  
Width 8000.0 Hz  
18 repetitions  
OBSERVE H1, 300.8117449 MHz  
DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec



Dalia soliman-MD6-Carbon-DMSO-ES

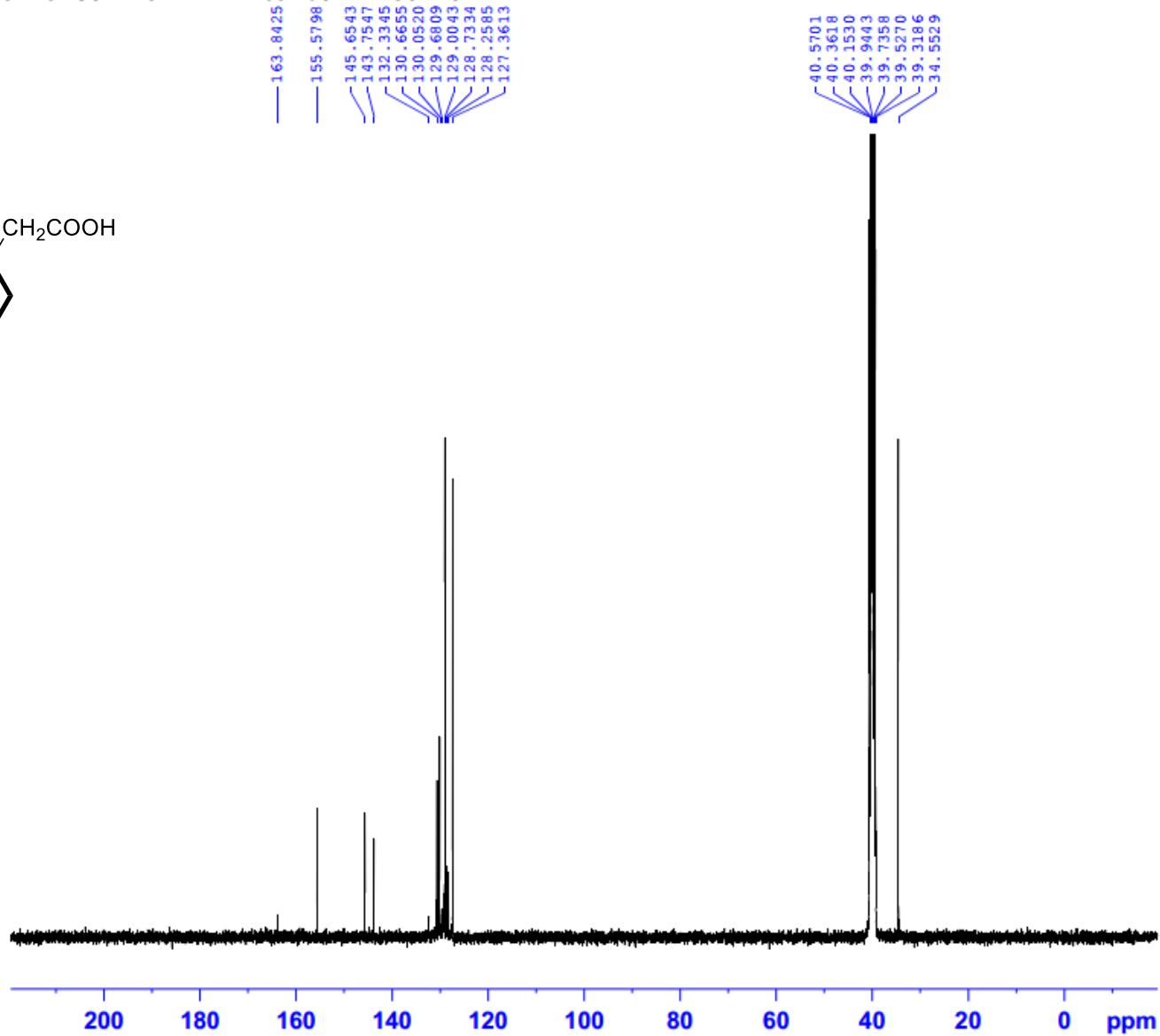
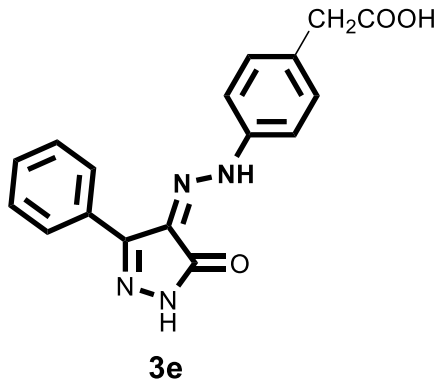


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 PROCNO 1

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 PULPROG zgpg30  
 TD 65536  
 SOLVENT DMSO  
 NS 2200  
 DS 4  
 SWH 24038.461 Hz  
 FIDRES 0.733596 Hz  
 AQ 1.3631488 sec  
 RG 197.77  
 DW 20.800 usec  
 DE 6.50 usec  
 TE 294.1 K  
 D1 2.0000000 sec  
 D11 0.0300000 sec  
 TDD 1  
 SFO1 100.6404331 MHz  
 NUC1 13C  
 P1 10.00 usec  
 PLW1 47.0000000 W  
 SFO2 400.2016008 MHz  
 NUC2 1H  
 CPDPRG2 waltz16  
 PCPD2 90.00 usec  
 PLW2 13.0000000 W  
 PLW12 0.29249999 W  
 PLW13 0.14713000 W

F2 - Processing parameters  
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 SF 100.6303700 MHz  
 MDW EX  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

Dalia soliman-MD11-Carbon-DMSO-ES



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Current Data Parameters
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EXPNO    10
PROCNO   1

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Date_    20220510
Time     22.21 h
INSTRUM  spect
PROBHD   X100618_0945 (
PULPROG  zgpg30
TD       65536
SOLVENT  DMSO
NS       2200
DS       4
SWH      24038.461 Hz
FIDRES   0.733596 Hz
AQ       1.3631488 sec
RG       197.77
DW       20.800 usec
DE       6.50 usec
TE       295.8 K
D1       2.00000000 sec
D11      0.03000000 sec
TDD      1
SFO1     100.6404331 MHz
NUC1     13C
P1       10.00 usec
PLW1     47.00000000 W
SFO2     400.2016008 MHz
NUC2     1H
CPOPRG[2] waltz16
PCPD2    90.00 usec
PLW2     13.00000000 W
PLW12    0.29249999 W
PLW13    0.14713000 W

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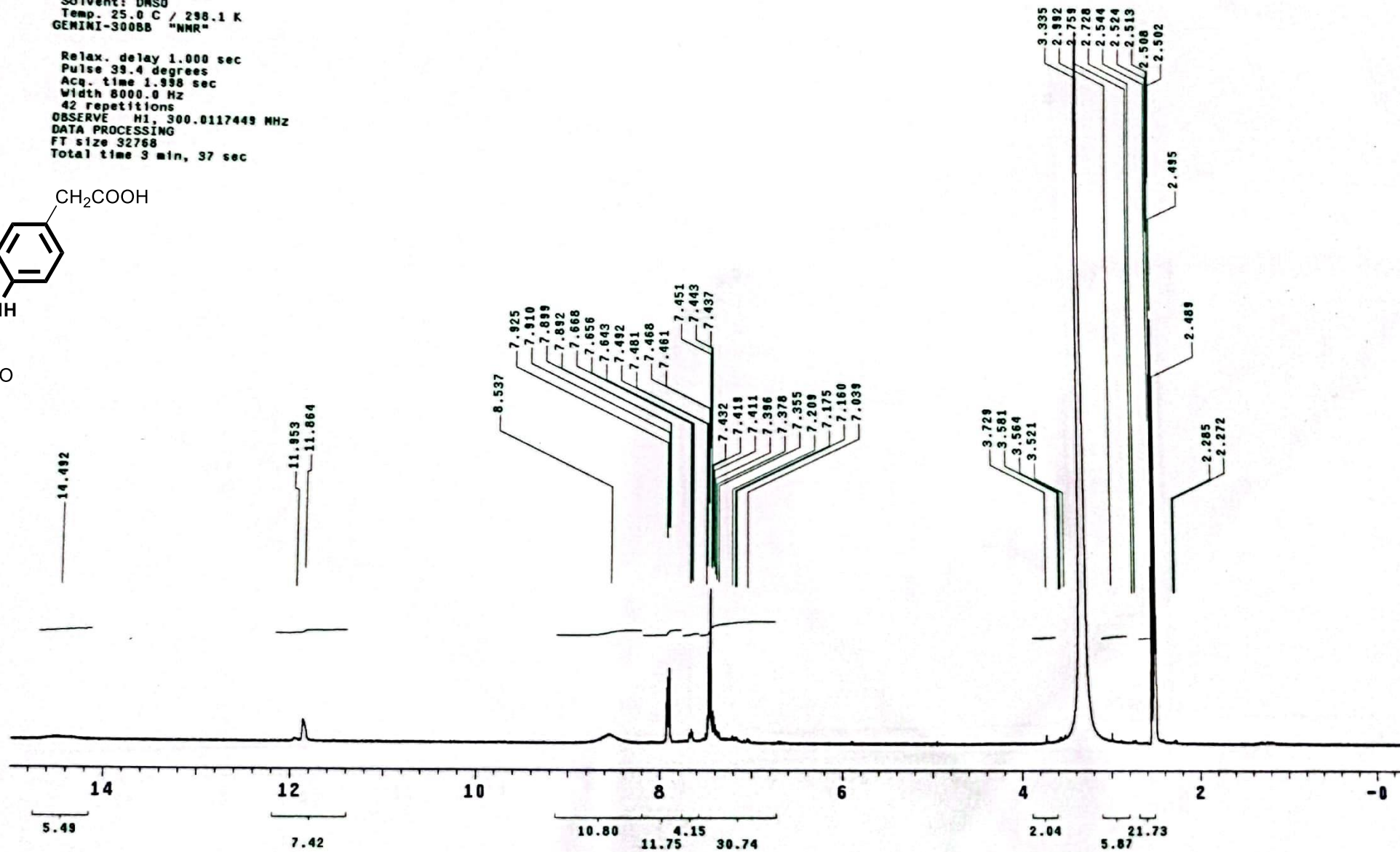
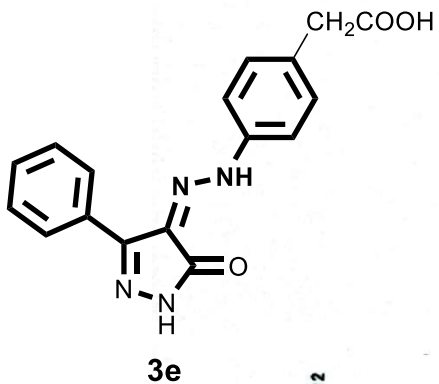


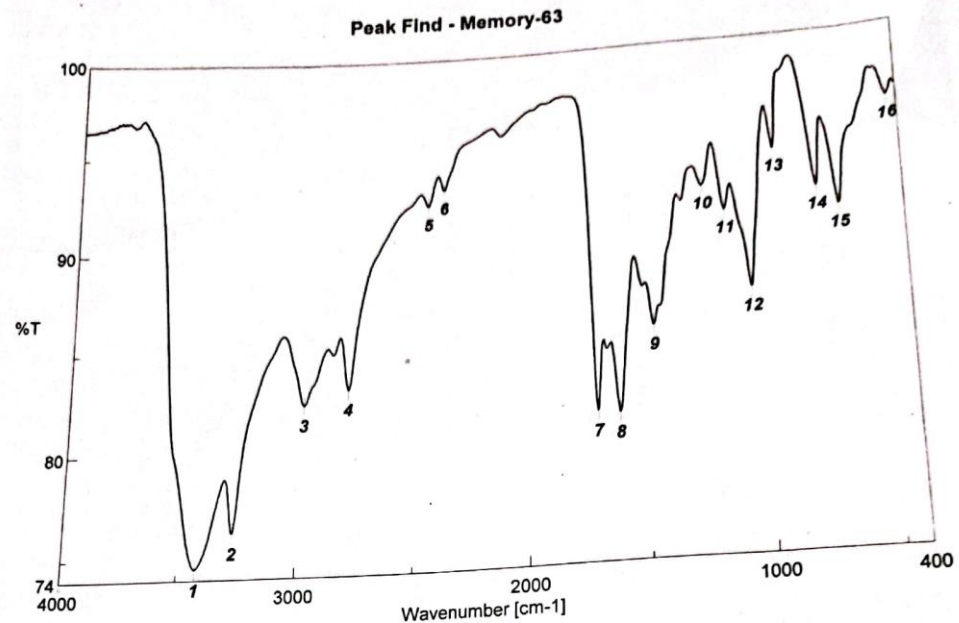
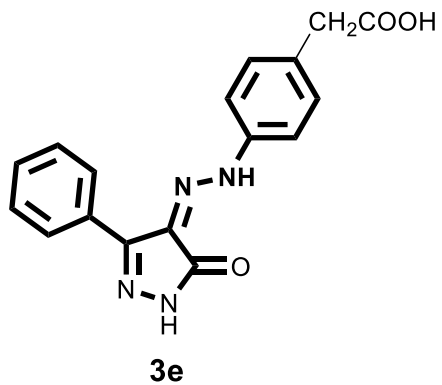
Dr. Dalia Soliman-ND11-H1-DMSO-Main. Defence. Chemical. Laboratory

Pulse Sequence: s2pu1

Solvent: DMSO  
Temp. 25.0 C / 298.1 K  
GEMINI-300BB "NMR"

Relax. delay 1.000 sec  
Pulse 39.4 degrees  
Acq. time 1.998 sec  
Width 8000.0 Hz  
42 repetitions  
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DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec





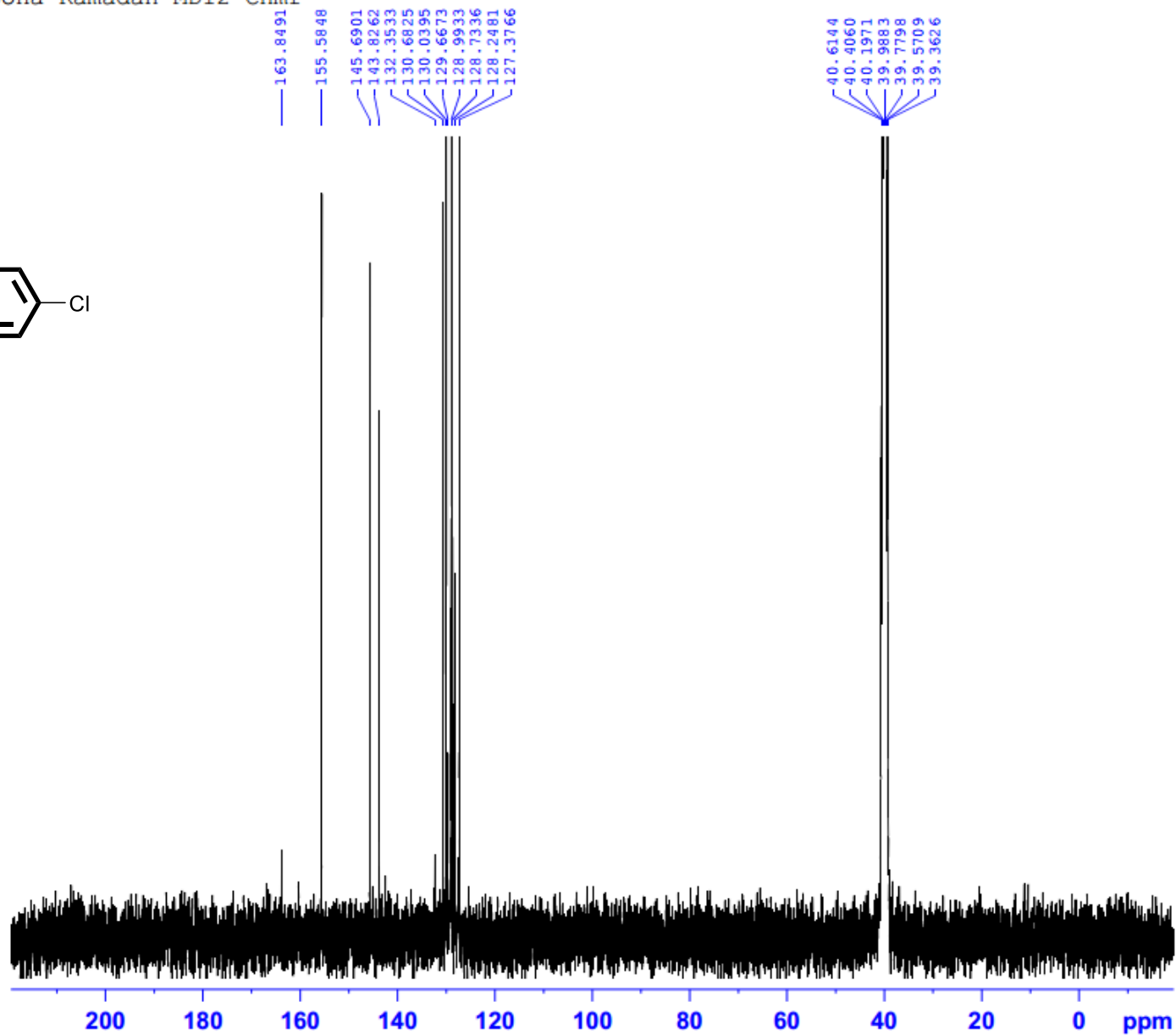
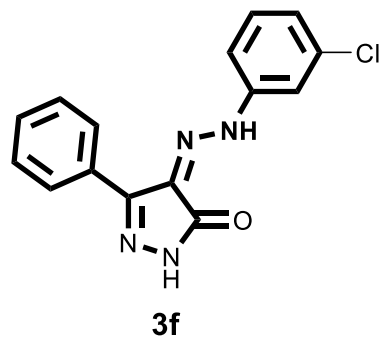
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 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

[ Result of Peak Picking ]

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4	2782.78	82.9325	5	2439.51	92.0784	6	2369.12	92.8806
7	1709.59	81.1071	8	1617.98	80.9596	9	1470.46	85.0764
10	1247.72	92.0117	11	1156.12	90.6488	12	1055.84	86.6001
13	935.306	93.7165	14	759.816	91.5939	15	669.178	90.5794
16	439.69	96.0167						



Soha Ramadan-MD12-Cnmr



Current Data Parameters  
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EXPNO 10  
PROCNO 1

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PULPROG zgpg30  
TD 65536  
SOLVENT DMSO  
NS 2200  
DS 4  
SWH 24038.461 Hz  
FIDRES 0.733596 Hz  
AQ 1.3631488 sec  
RG 197.77  
DW 20.800 usec  
DE 6.50 usec  
TE 300.1 K  
D1 2.00000000 sec  
D11 0.03000000 sec  
TDO 1  
SFO1 100.6404331 MHz  
NUC1 13C  
P1 10.00 usec  
PLW1 47.00000000 W  
SFO2 400.2016008 MHz  
NUC2 1H  
CPDPRG[2] waltz16  
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PLW2 13.00000000 W  
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PLW13 0.14713000 W

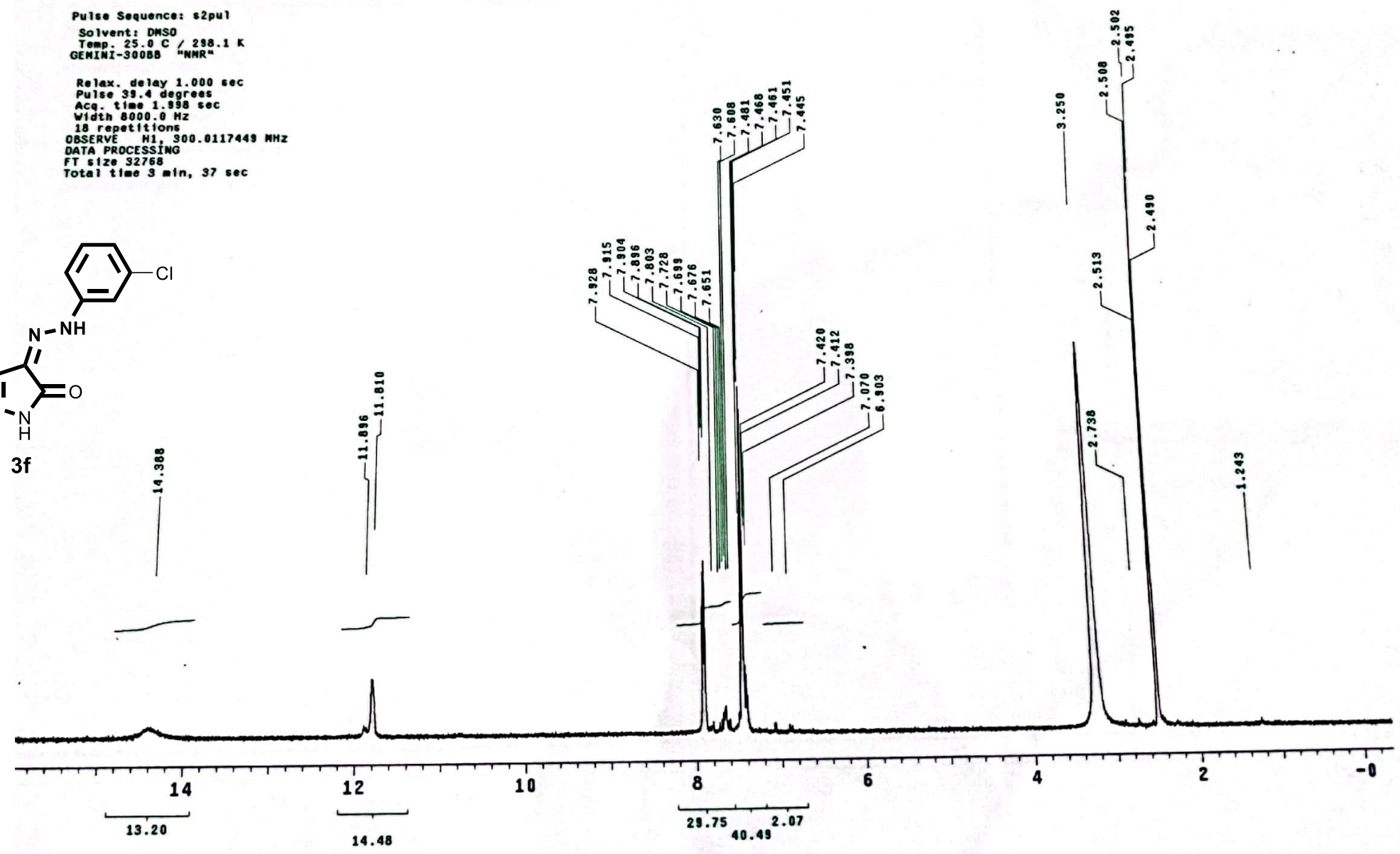
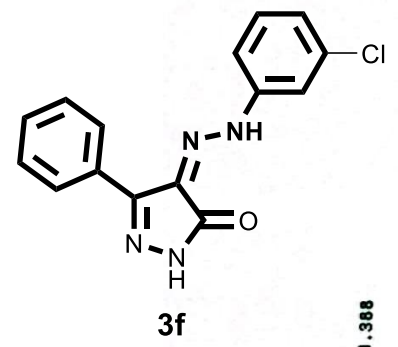
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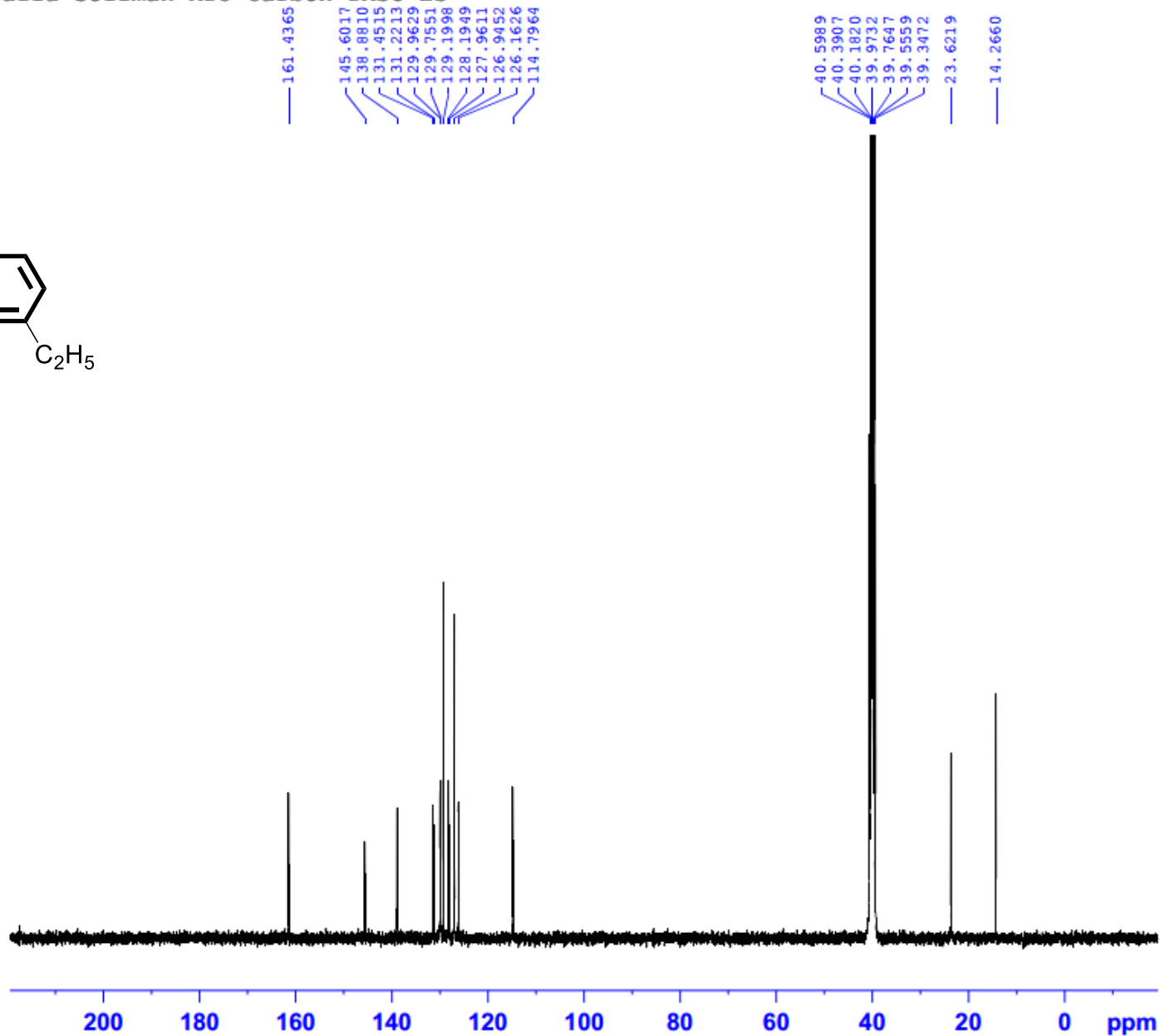
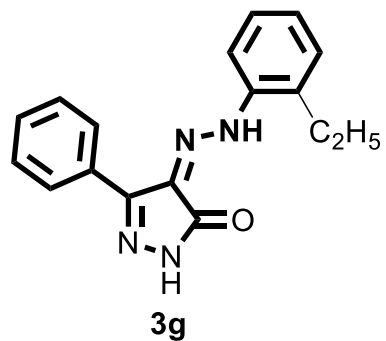
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Solvent: DMSO  
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GENI-30088 "NMR"

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Pulse 39.4 degrees  
Acq. time 1.998 sec  
Width 8000.0 Hz  
18 repetitions  
OBSERVE H1, 300.0117449 MHz  
DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec



Dalia soliman-MD5-Carbon-DMSO-ES



```

Current Data Parameters
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EXPNO    10
PROCNO    1

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PROBHD    Z106618_0945 (
PULPROG   zgpg30
TD         65536
SOLVENT   DMSO
NS         2200
DS         4
SWH        24038.461 Hz
FIDRES     0.733596 Hz
AQ         1.3631488 sec
RG         197.77
DW         20.800 usec
DE         6.50 usec
TE         295.8 K
D1         2.0000000 sec
d11        0.0300000 sec
TDO        1
SFO1       100.6404331 MHz
NUC1       13C
P1         10.00 usec
PLW1       47.0000000 W
SFO2       400.2016008 MHz
NUC2       1H
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PLW13     0.14713000 W

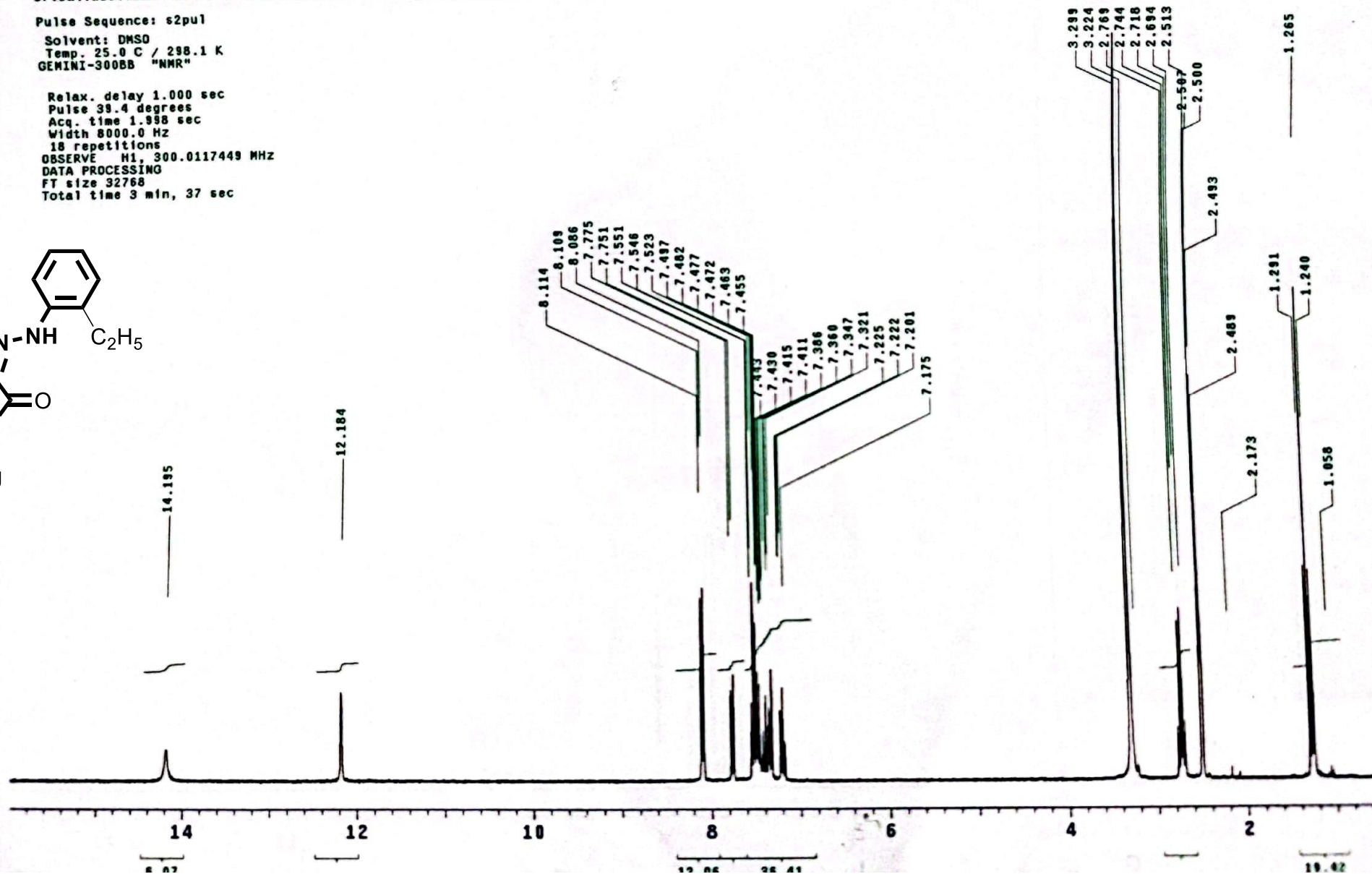
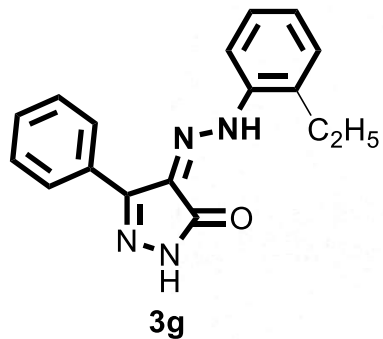
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Dr. Dalila Soliman-MD5-H1-DMSO-Main.Defence.Chemical.Laboratory

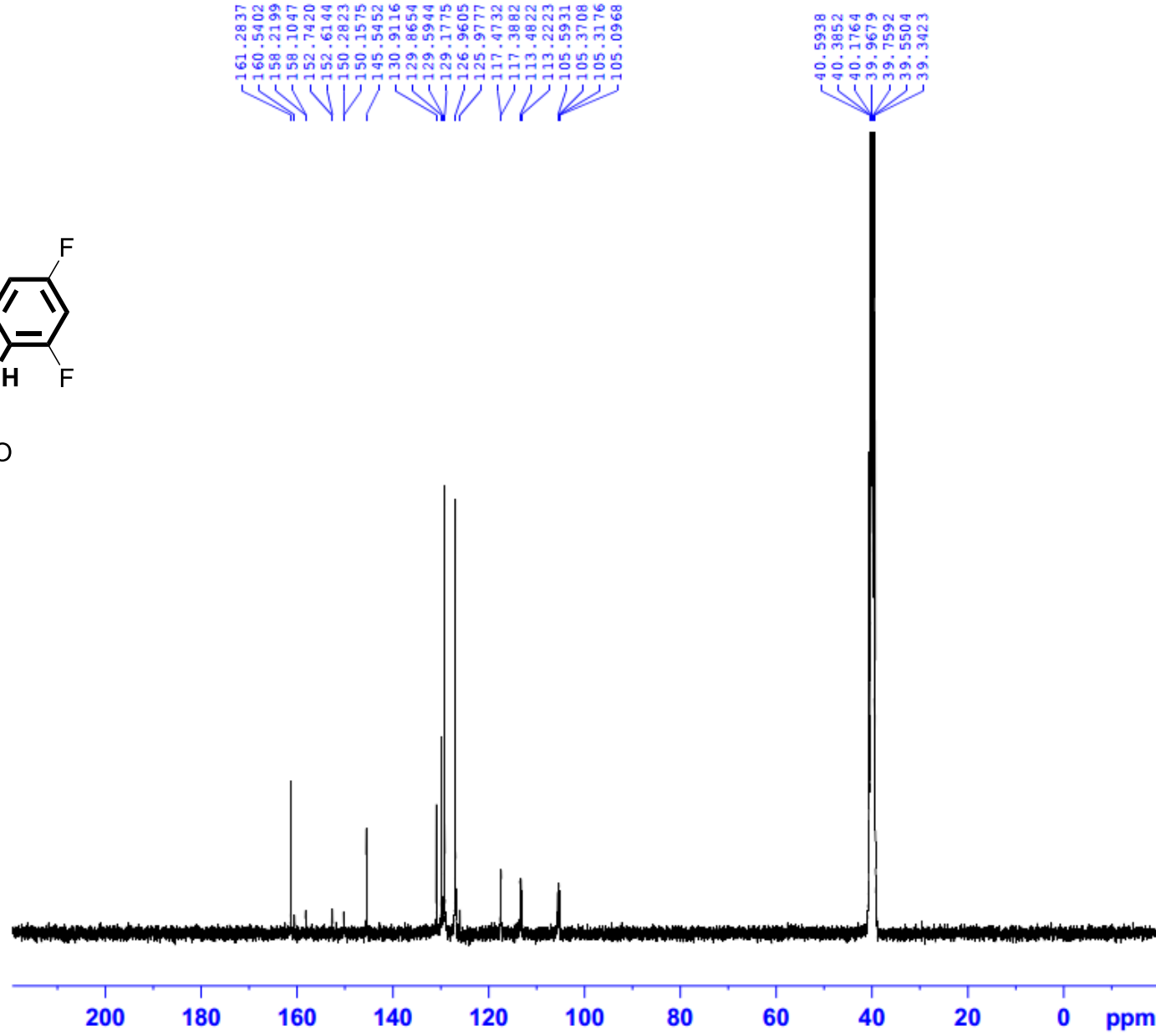
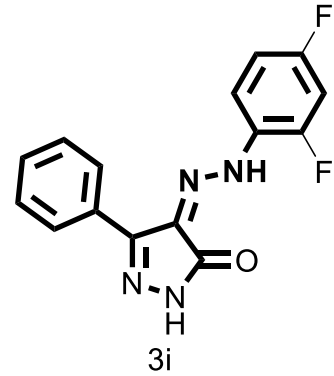
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Solvent: DMSO  
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GEMINI-300BB "NMR"

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Pulse 39.4 degrees  
Acq. time 1.998 sec  
Width 8000.0 Hz  
18 repetitions  
OBSERVE H1, 300.0117449 MHz  
DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec



Dalia soliman-MD1-Carbon-DMSO-ES



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158.2199  
158.1047  
152.7420  
152.6144  
150.2823  
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129.8654  
129.5944  
129.1775  
126.9605  
125.9777  
117.4732  
117.3882  
113.8822  
113.2223  
105.5931  
105.3708  
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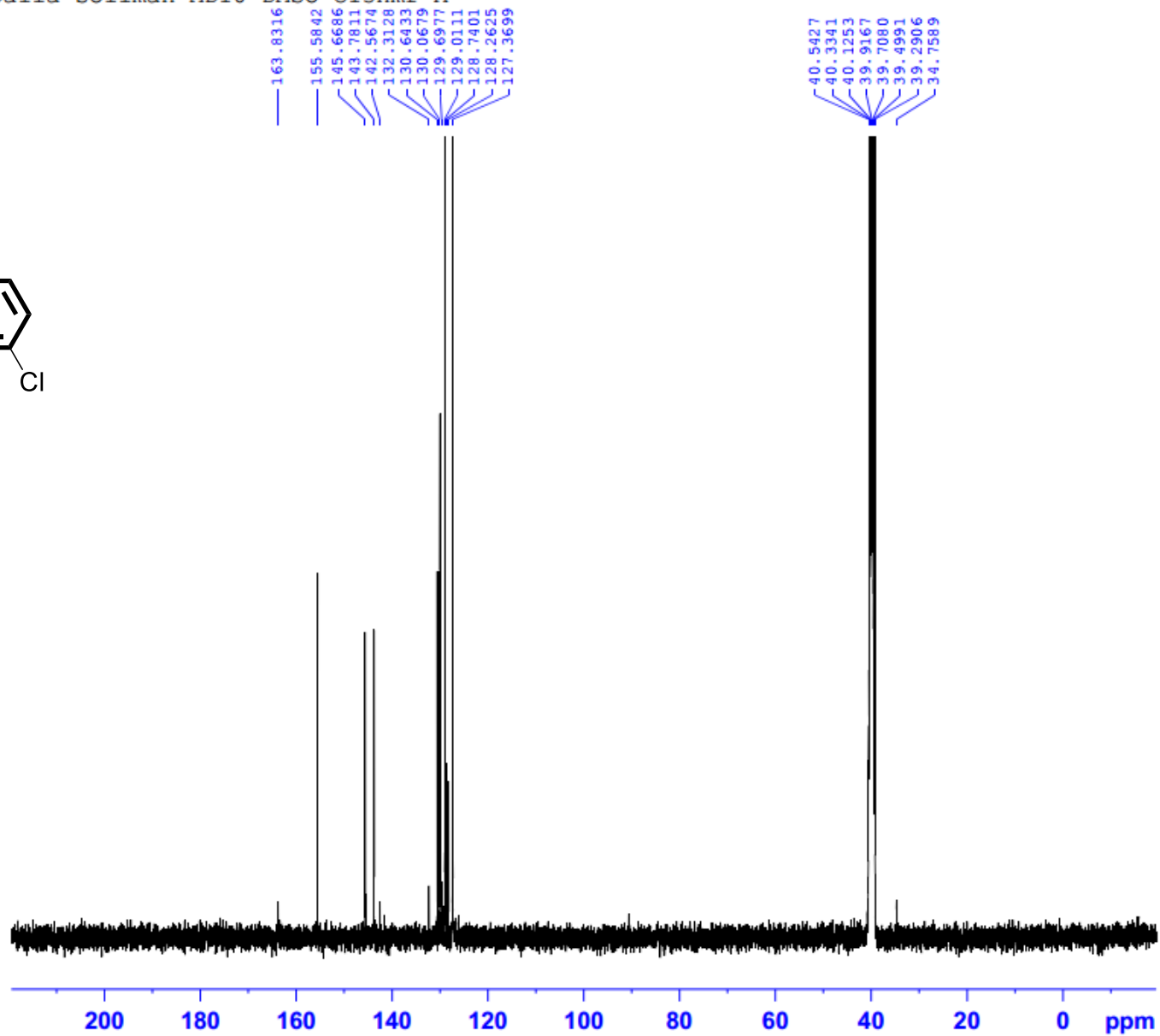
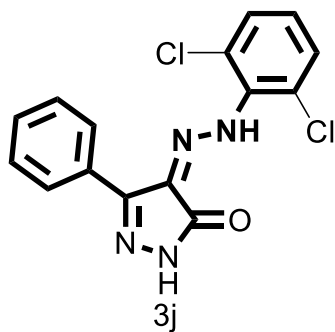


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PROCNO 1

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PULPROG zgpg30  
TD 65536  
SOLVENT DMSO  
NS 2200  
DS 4  
SWH 24038.461 Hz  
FIDRES 0.733596 Hz  
AQ 1.3631488 sec  
RG 197.77  
DW 20.800 usec  
DE 6.50 usec  
TE 296.0 K  
D1 2.0000000 sec  
D11 0.0300000 sec  
TD0 1  
SFO1 100.6404331 MHz  
NUC1 13C  
P1 10.00 usec  
PLM1 47.0000000 W  
SFO2 400.2016008 MHz  
NUC2 1H  
CPOPRG[2] waltz16  
PCPD2 90.00 usec  
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PLM12 0.29249999 W  
PLM13 0.14713000 W

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GB 0  
PC 1.40

Dalia soliman-MD10-DMSO-C13nmr-A



Current Data Parameters  
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 PROCNO 1

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 TD 65536  
 SOLVENT DMSO  
 NS 2200  
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 SWH 24038.461 Hz  
 FIDRES 0.733596 Hz  
 AQ 1.3631488 sec  
 RG 197.77  
 DW 20.800 usec  
 DE 6.50 usec  
 TE 294.0 K  
 D1 2.0000000 sec  
 D11 0.0300000 sec  
 TDO 1  
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 NUC1 13C  
 P1 10.00 usec  
 PLW1 47.0000000 W  
 SFO2 400.2016008 MHz  
 NUC2 1H  
 CPDPRG2 waltz16  
 PCPD2 90.00 usec  
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 PLW13 0.14713000 W

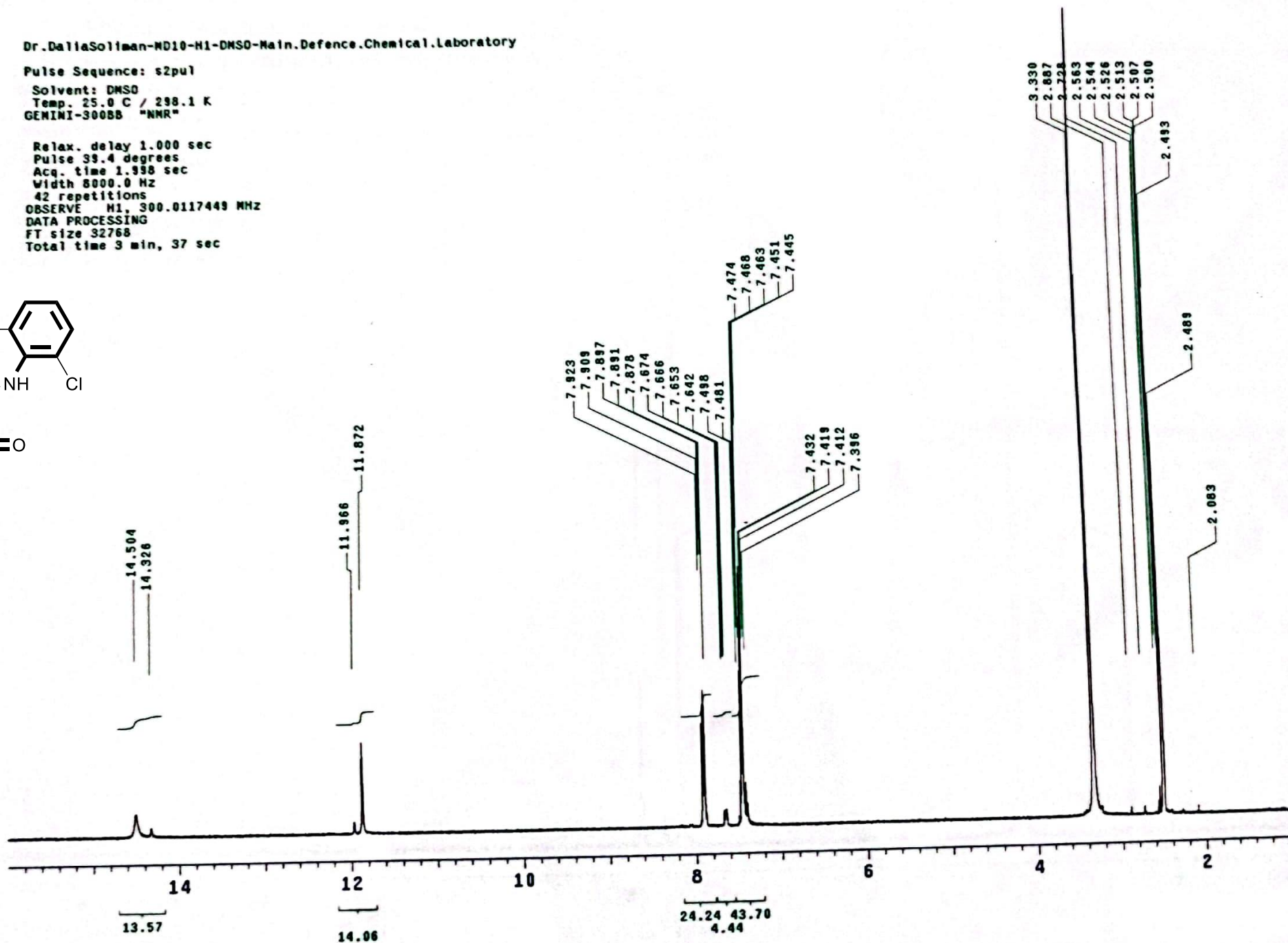
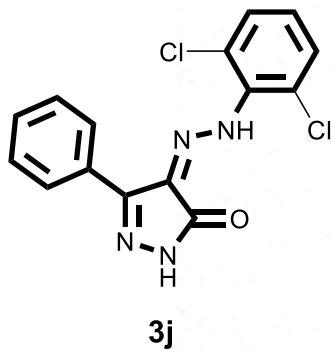
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 WDW EM  
 SSB 0  
 LB 1.00 Hz  
 GB 0  
 PC 1.40

Dr. Dalia Soliman-MD10-H1-DMSO-Main. Defence. Chemical. Laboratory

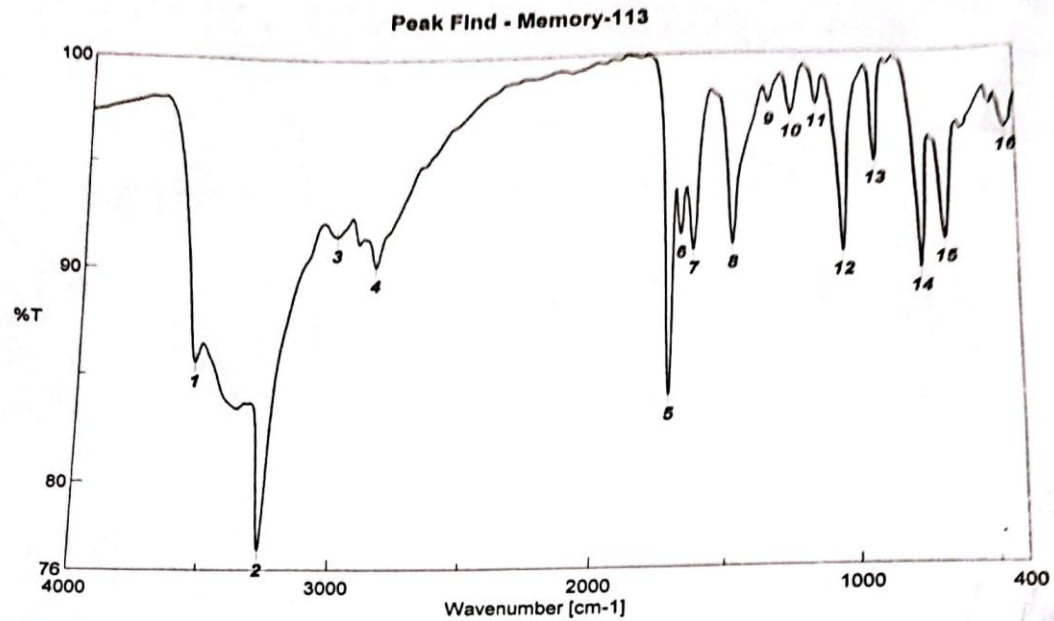
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Solvent: DMSO  
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DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec



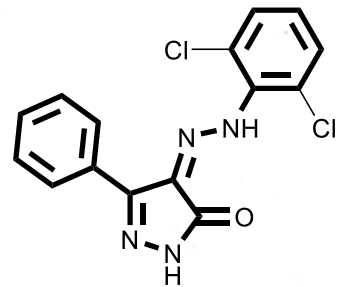




[Comments]  
 Sample name MD 10  
 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

[ Result of Peak Picking ]

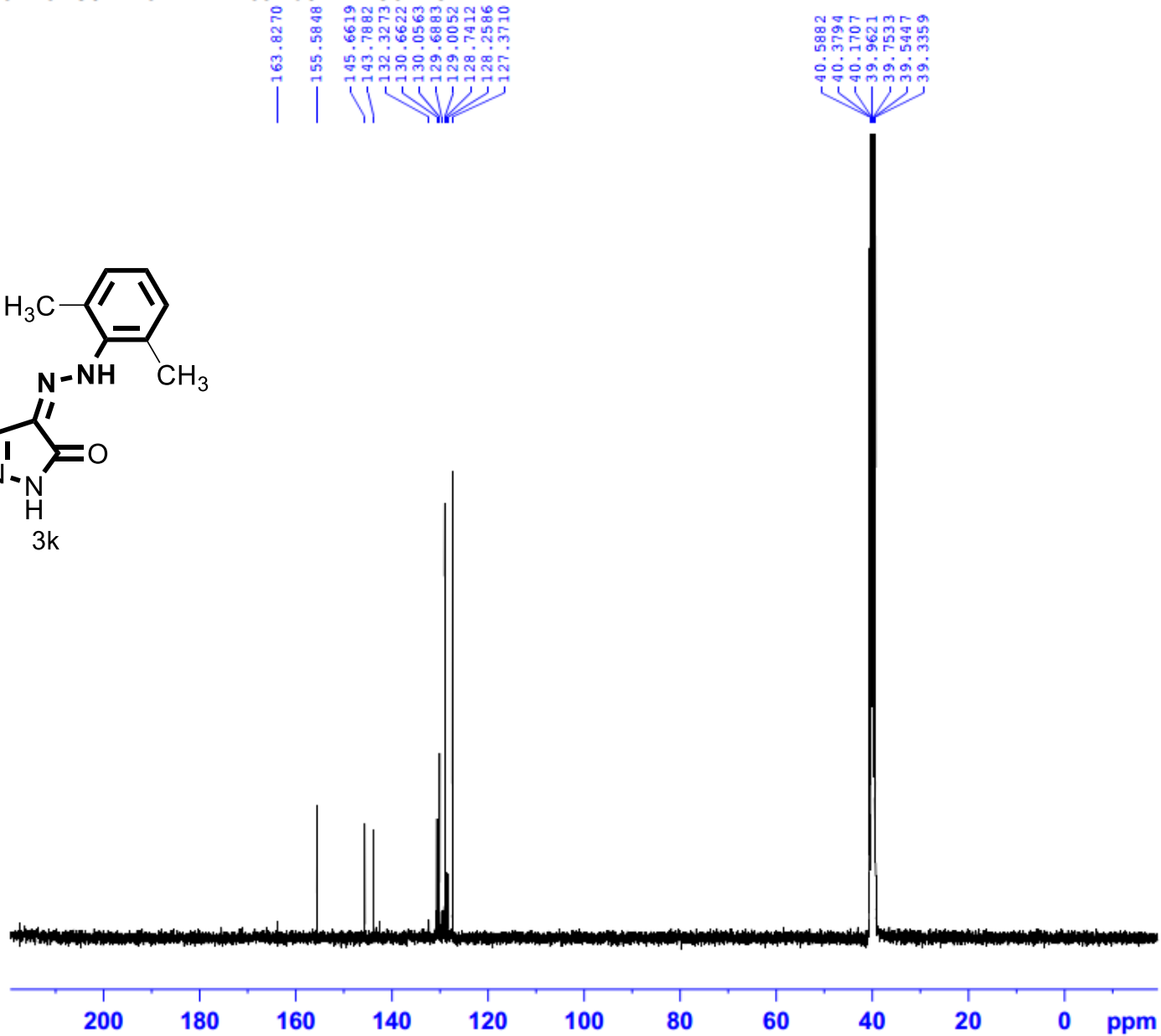
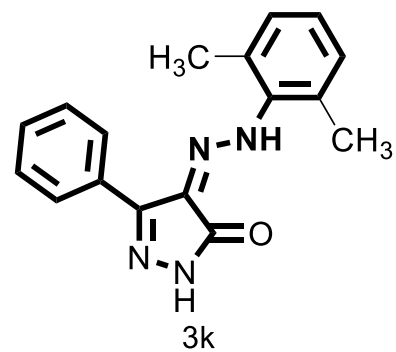
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4	2853.17	89.9321	5	1714.41	83.6458	6	1670.05	91.0971
7	1622.8	90.3407	8	1475.28	90.5472	9	1338.36	97.5105
10	1254.47	96.8972	11	1155.15	97.401	12	1057.76	90.1593
13	933.378	94.5632	14	762.709	89.3672	15	669.178	90.7441
16	434.869	96.0791						



3j



Dalia soliman-MD7-Carbon-DMSO-ES



Current Data Parameters  
NAME Dalia soliman-MD7-Carbon-DMSO-ES  
EXPNO 10  
PROCNO 1

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SOLVENT DMSO  
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DS 4  
SWH 24038.461 Hz  
FIDRES 0.733596 Hz  
AQ 1.3631488 sec  
RG 197.77  
EW 20.800 usec  
DE 6.50 usec  
TE 295.8 K  
D1 2.00000000 sec  
D11 0.03000000 sec  
TD0 1  
SFO1 100.6404331 MHz  
NUC1 13C  
P1 10.00 usec  
PLM1 47.00000000 W  
SFO2 400.2016008 MHz  
NUC2 1H  
CPDPRG2 waltz16  
PCPD2 90.00 usec  
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PLM12 0.29249999 W  
PLM13 0.14713000 W

F2 - Processing parameters  
SI 32768  
SF 100.6303700 MHz  
WDW EM  
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GB 0  
PC 1.40



Dr. DaliaSoliman-ND7-H1-DMSO-Main.Defence  
 .Chemical.Laboratory

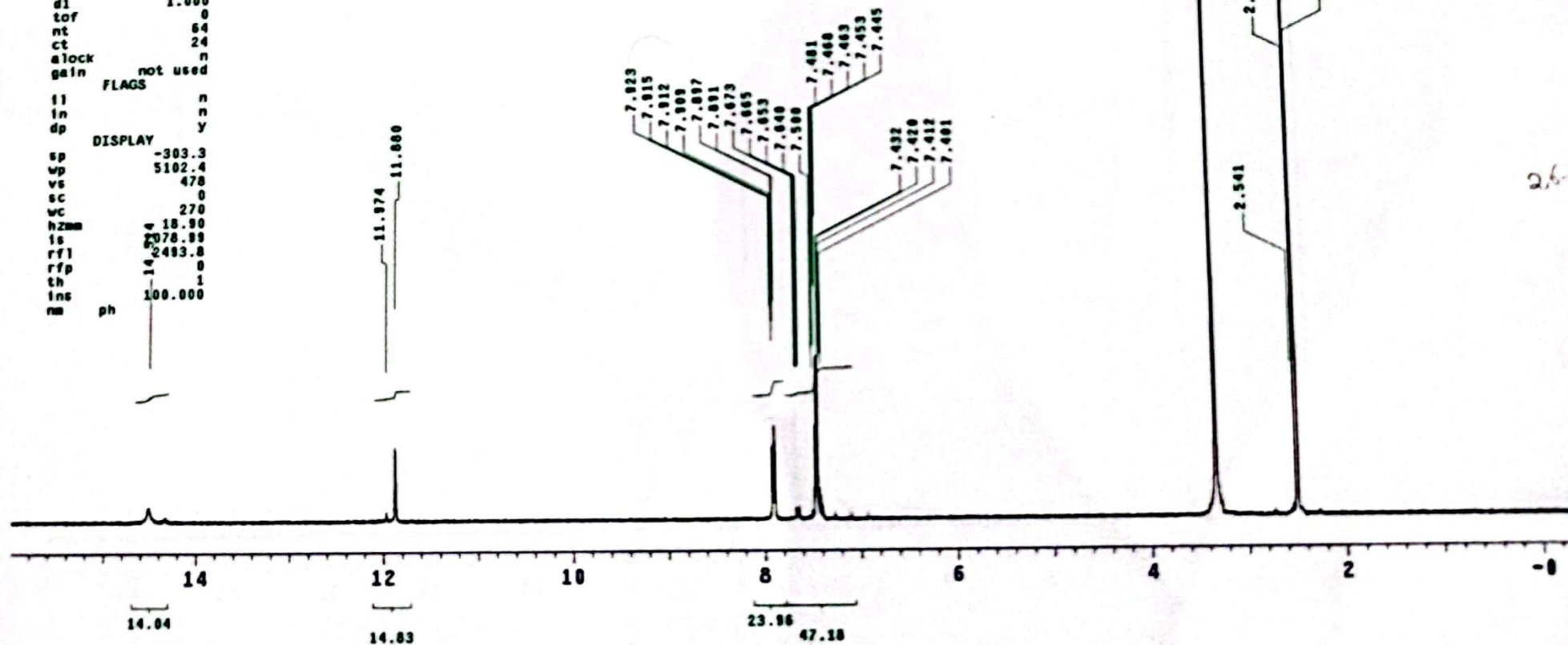
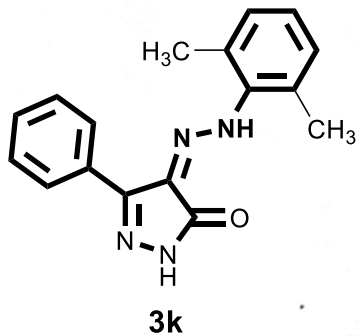
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 vnmr1/M.Afif13/Dr.~ dof 0  
 DaliaSoliman-ND7-H~ dm nnn  
 l-DMSO-Main.Defenc~ dnm c  
 e.Chemical.Laborat~ dmf 200  
 ory.fid temp 25.0

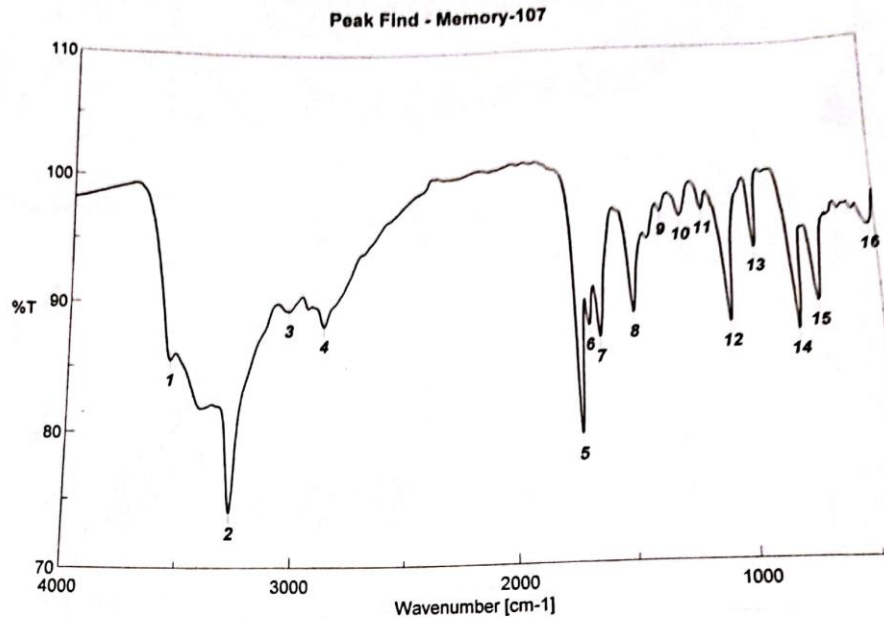
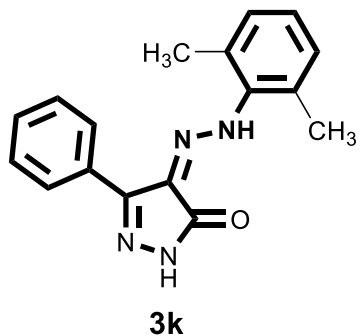
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 at 1.898 fn not used  
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 fb 4400 wexp  
 bs 6 wbs wft aph  
 tpwr 52 wnt  
 pw 7.0  
 dl 1.000  
 tof 0  
 nt 64  
 ct 24  
 alock n  
 gain not used

FLAGS  
 ll n  
 ln n  
 dp y

DISPLAY  
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 vs 478  
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 hzmm 18.90  
 ls 2078.99  
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 rfp 0  
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 nm ph



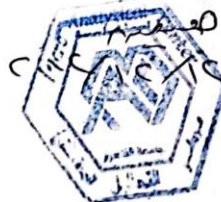
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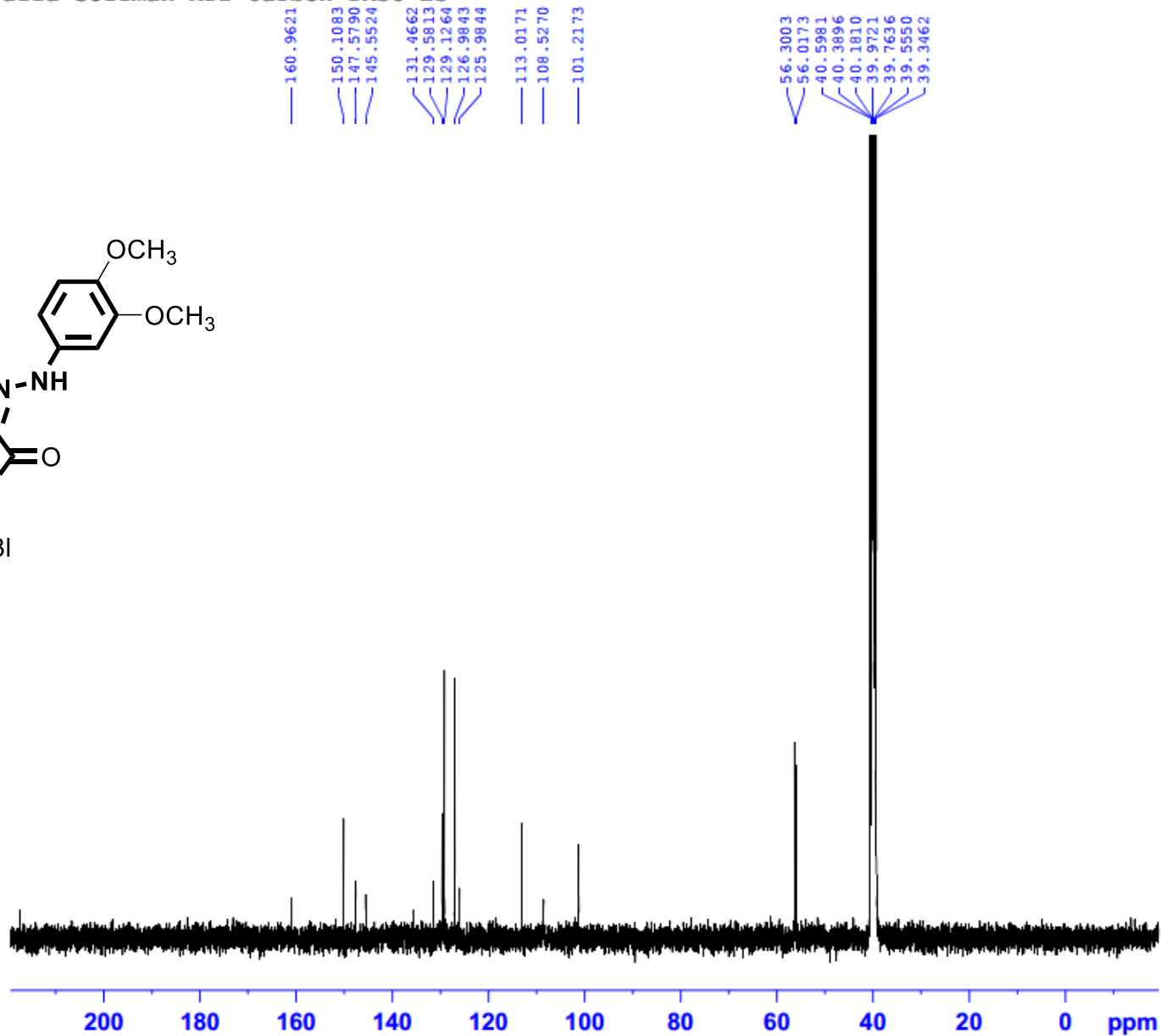
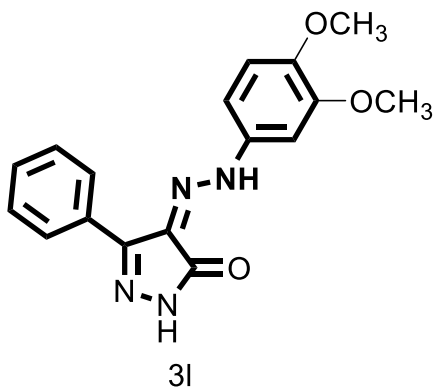
[Comments]  
 Sample name MD7  
 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

[ Result of Peak Picking ]

No.	Position	Intensity	No.	Position	Intensity	No.	Position	Intensity
1	3540.67	85.3871	2	3267.79	73.9667	3	3010.34	89.1254
4	2852.2	87.8833	5	1712.48	79.1504	6	1669.09	87.2955
7	1622.8	86.3832	8	1475.28	88.1398	9	1339.32	96.0209
10	1253.5	95.6034	11	1155.15	96.0779	12	1056.8	87.1685
13	933.378	92.9489	14	762.709	86.4947	15	670.142	88.6502
16	427.155	94.5296						



Dalia soliman-MD2-Carbon-DMSO-ES



```

Current Data Parameters
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EXPNO    10
PROCNO   1

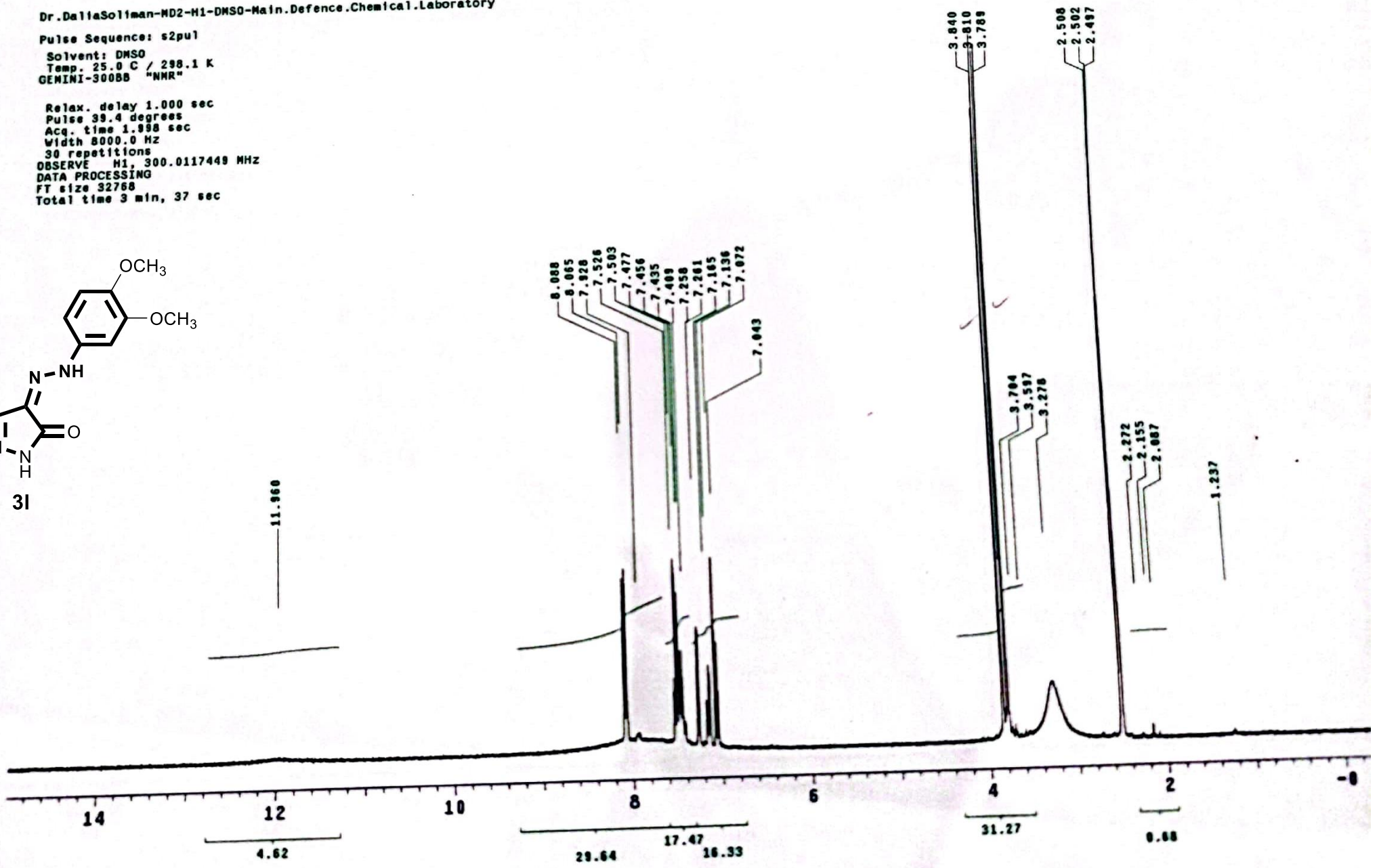
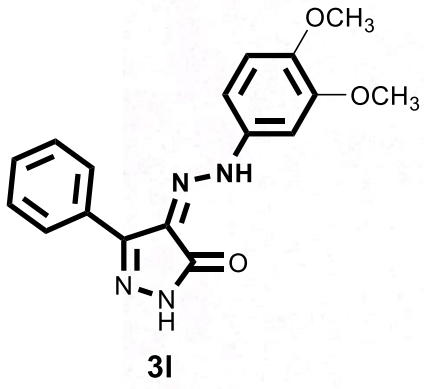
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SOLVENT  DMSO
NS       2200
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FIDRES   0.733596 Hz
AQ       1.3631488 sec
RG       197.77
DW       20.800 usec
DE       6.50 usec
TE       295.7 K
D1       2.0000000 sec
D11      0.03000000 sec
TQ0      1
SFO1     100.6404331 MHz
NUC1     13C
P1       10.00 usec
PLM1     47.0000000 W
SFO2     400.2016008 MHz
NUC2     1H
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PLM13    0.14713000 W

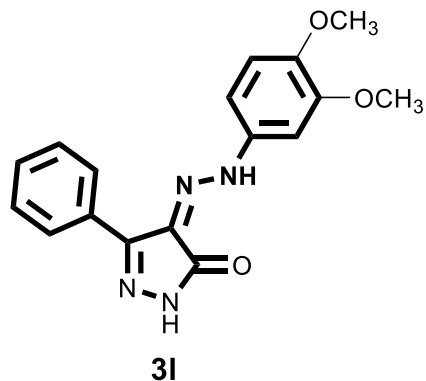
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PC       1.40
    
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Dr. Dalia Soliman-MD2-H1-DMSO-Main Defence Chemical Laboratory

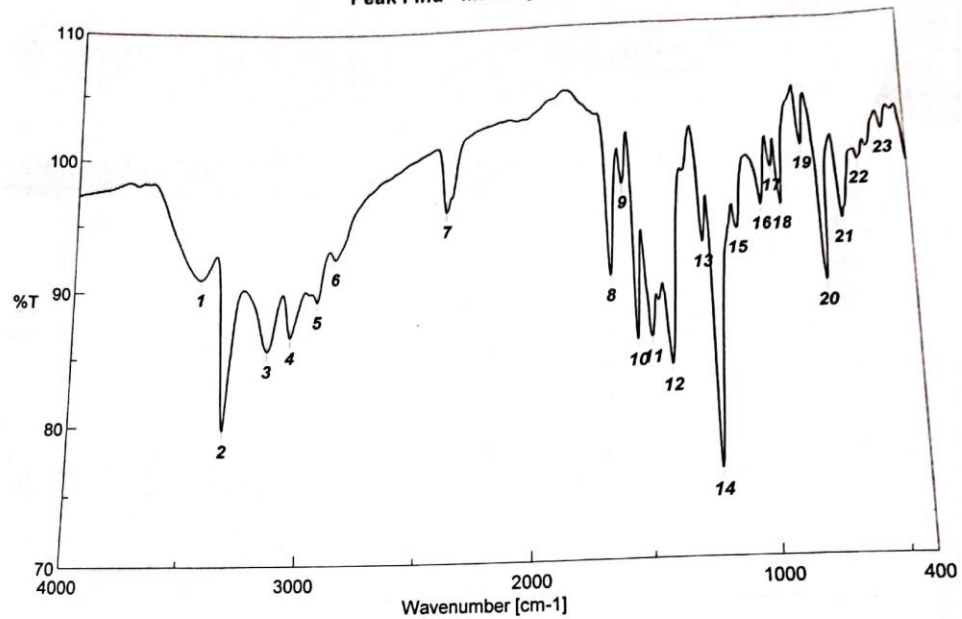
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Solvent: DMSO  
Temp. 25.0 C / 298.1 K  
GEMINI-300BB "NMR"

Relax. delay 1.000 sec  
Pulse 39.4 degrees  
Acq. time 1.998 sec  
Width 8000.0 Hz  
30 repetitions  
OBSERVE H1, 300.0117449 MHz  
DATA PROCESSING  
FT size 32768  
Total time 3 min, 37 sec





Peak Find - Memory-37

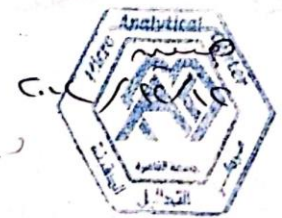


[Comments]  
 Sample name M.SD 2  
 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

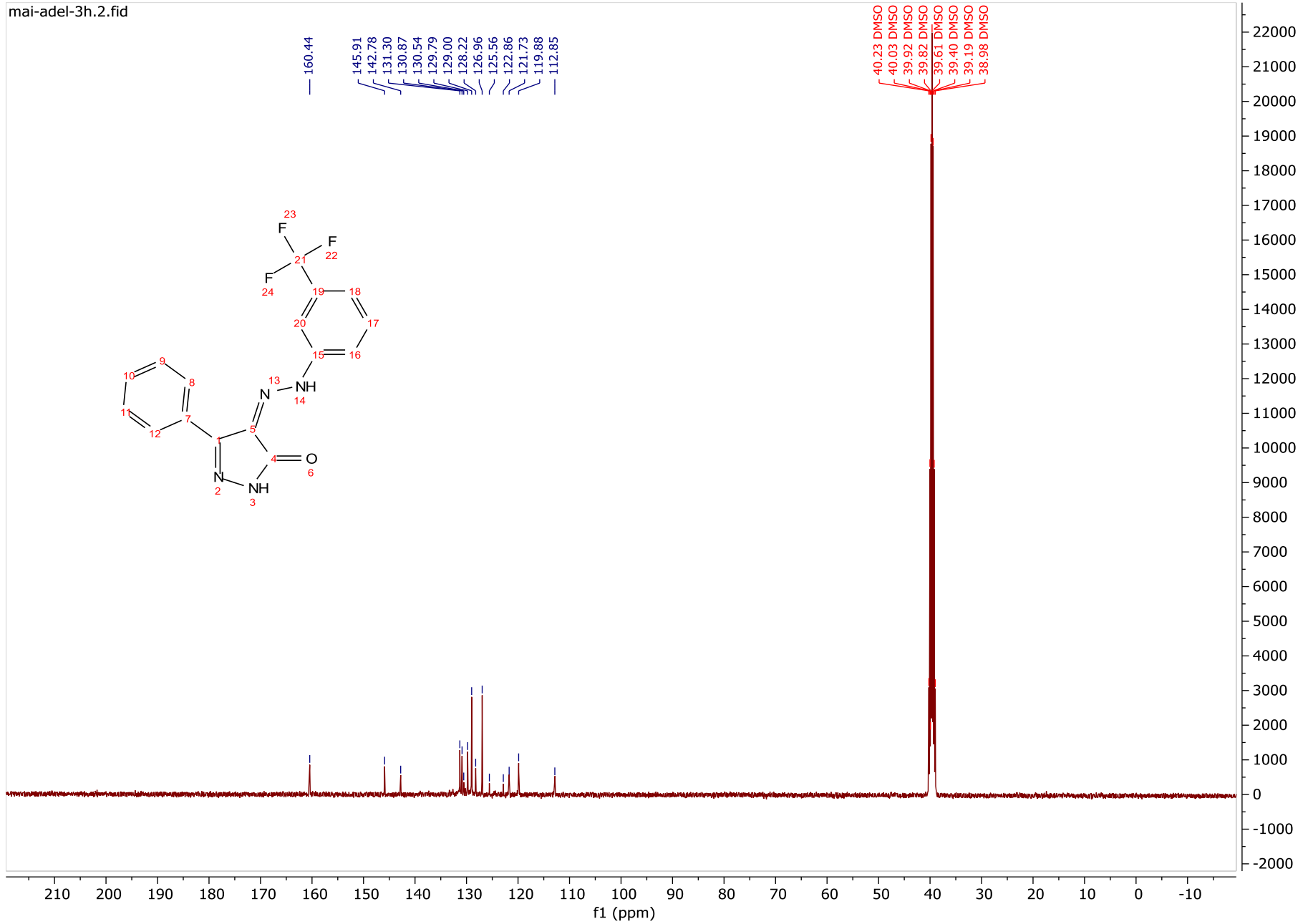
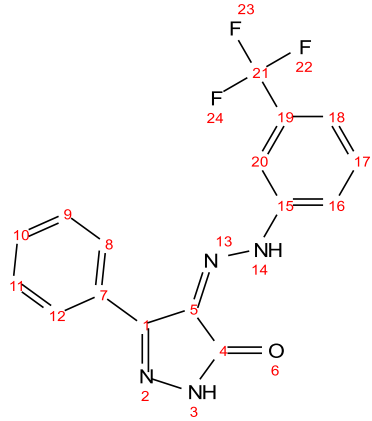
Result of Peak Picking ]

No.	Position	Intensity	No.	Position	Intensity	No.	Position	Intensity
1	3439.42	90.9026	2	3325.64	79.7092	3	3141.47	85.4807
4	3042.16	86.4484	5	2927.41	89.0458	6	2849.31	92.228
7	2360.44	95.4371	8	1658.48	90.0128	9	1604.48	96.9844
10	1550.49	85.2869	11	1490.7	85.4495	12	1409.71	83.3804
13	1273.75	92.2371	14	1221.68	75.9868	15	1127.19	93.2243
16	1021.12	94.9852	17	970.983	97.8989	18	936.271	94.9693
19	836.955	99.5774	20	761.744	89.184	21	680.749	93.833
22	602.646	98.3362	23	493.688	100.607			

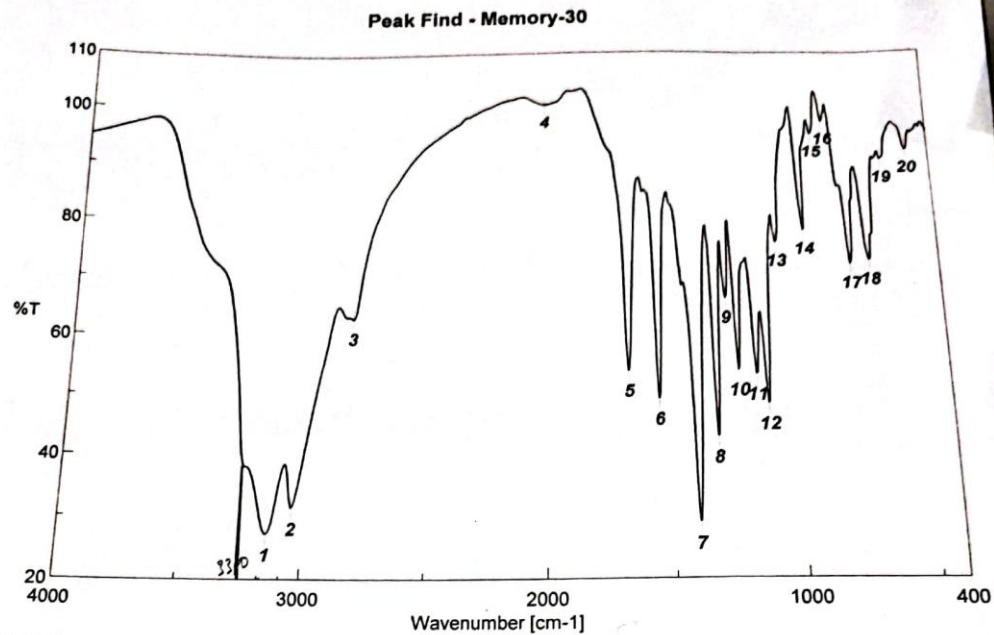
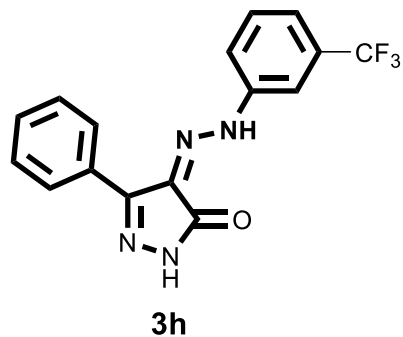
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mai-adel-3h.2.fid







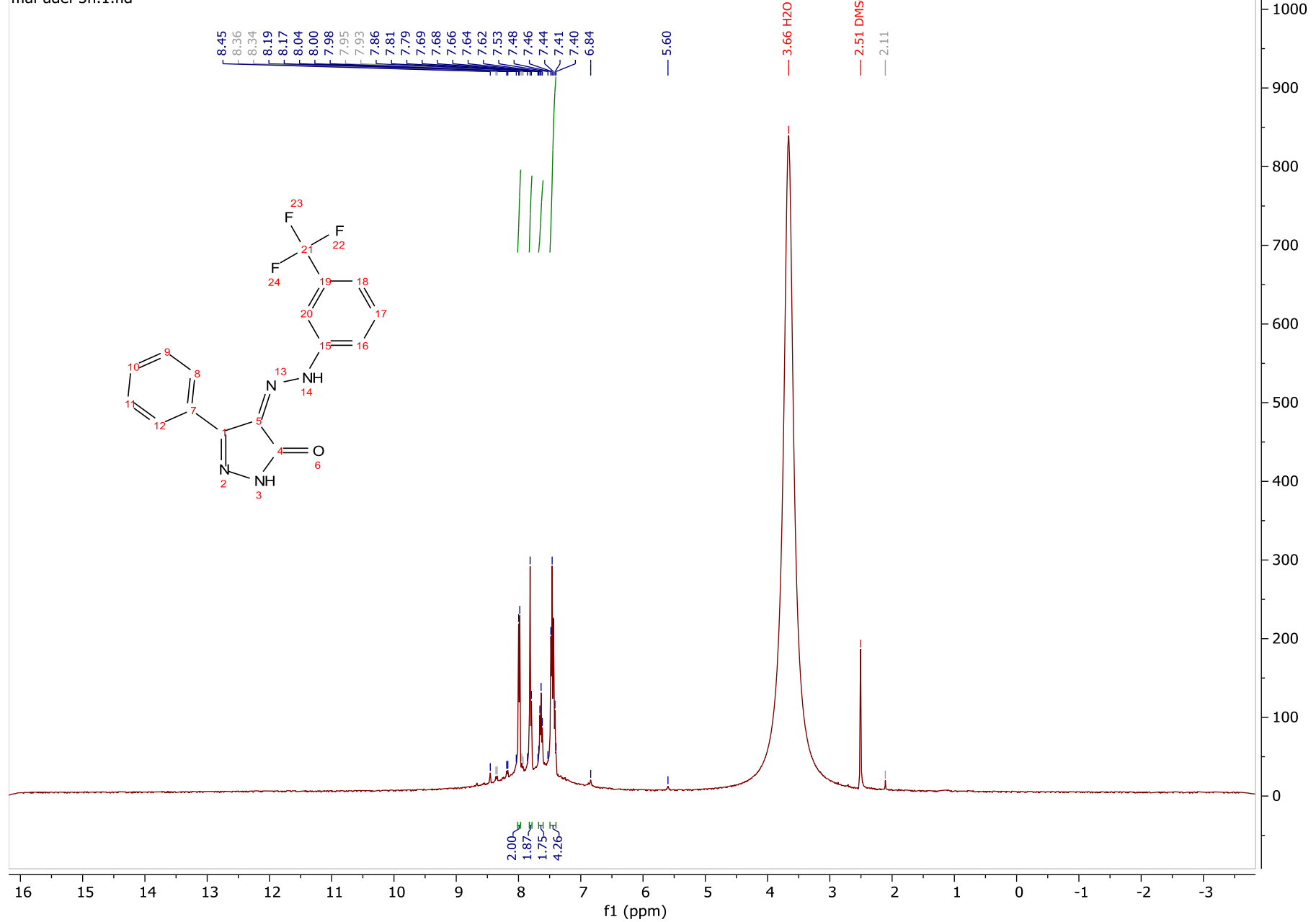
[Comments]  
 Sample name MSD3  
 Comment 12/12/2012  
 User IR  
 Division IR  
 Company Micro Analytical Center

Result of Peak Picking ]

No.	Position	Intensity	No.	Position	Intensity	No.	Position	Intensity
1	3142.44	27.1806	2	3043.12	31.2856	3	2814.6	62.0183
2	2017.18	99.9552	5	1675.84	52.5383	6	1556.27	47.8968
3	1405.85	28.5557	8	1327.75	41.7668	9	1283.39	64.3712
4	1240	52.4221	11	1168.65	51.6675	12	1126.22	46.9187
5	1066.44	74.0867	14	947.842	76.2686	15	889.023	94.1178
6	836.955	96.529	17	763.673	70.3229	18	683.642	71.0352
7	603.61	89.4393	20	492.723	91.2829			



mai-adel-3h.1.fid





### **Docking study**

Automated docking studies were carried out using Glide (SP) scoring function Maestro10.1 Schrodinger, (2015-Release-4).

### **Ligand Preparation**

LigPrep module, was used for geometrical refining of chemical structures. LigPrep is intended to set up premium 3D structures with accurate chiralities. Original states of ionization were retained; tautomers and conformations were generated by the Monte Carlo method as implemented in MacroModel version 9.8, 2010, using OPLS-2005 force field. The generated conformers were subsequently minimized using truncated Newton conjugate gradient (TNCG) minimization up to 500 iterations. The conformers with an energy difference of 30 kcal/mol as compared to the global energy minimum conformer were retained. The conformational searches were carried out for aqueous solution using the generalized born/solvent accessible surface (GB/SA) continuum solvation model.

### **Protein Preparation.**

Protein preparation wizard of Maestro software was used for protein preparation. The selected chains were edited for missing hydrogens and for assigning proper bond orders. The H-bonds were optimized using sample orientations. All the polar hydrogens were displayed. All the crystallographic waters without hydrogen bond interactions with protein residues were removed. Finally, the non-hydrogen atoms of the protein structure was minimized to the default Root Mean Square Deviation (RMSD) value of 0.3.

### **Receptor Grid Generation.**

From the defined receptor, the co-crystallized ligand was separated from the active site of receptor chain. The atoms were of size equal to Van der Waals radii of 1.0 $\text{\AA}$  while the partial atomic charge was less than 0.25 defaults. The active site represents an enclosing box at the centroid of the workspace ligand. Following this protocol, a grid centered on the ligand was generated using the default Glide settings. All ligands were docked into this grid structure.

### **Molecular Docking Analysis.**

On a defined receptor grid, flexible docking was performed using the precision (SP) feature of Glide module, version 5.6, 2010. The constraints to defined ligand-receptor interactions were not set. The structure output format was set to pose viewer file so as to view the output of the resulting docking studies from pose viewer.

### Acknowledgements

In this work, docking studies were carried out using the Glide, Maestro10.1 Schrodinger, (2015-Release-4).

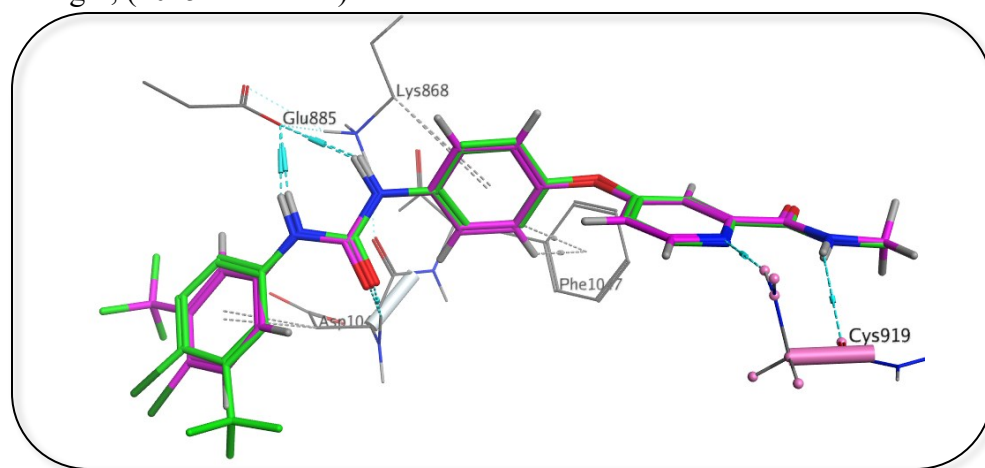


Figure S1 3D Representation of the superimposition of the docking pose (green) and the co-crystallized (purple) of sorafenib in the VEGFR-2 active site with RMSD of 0.21Å.

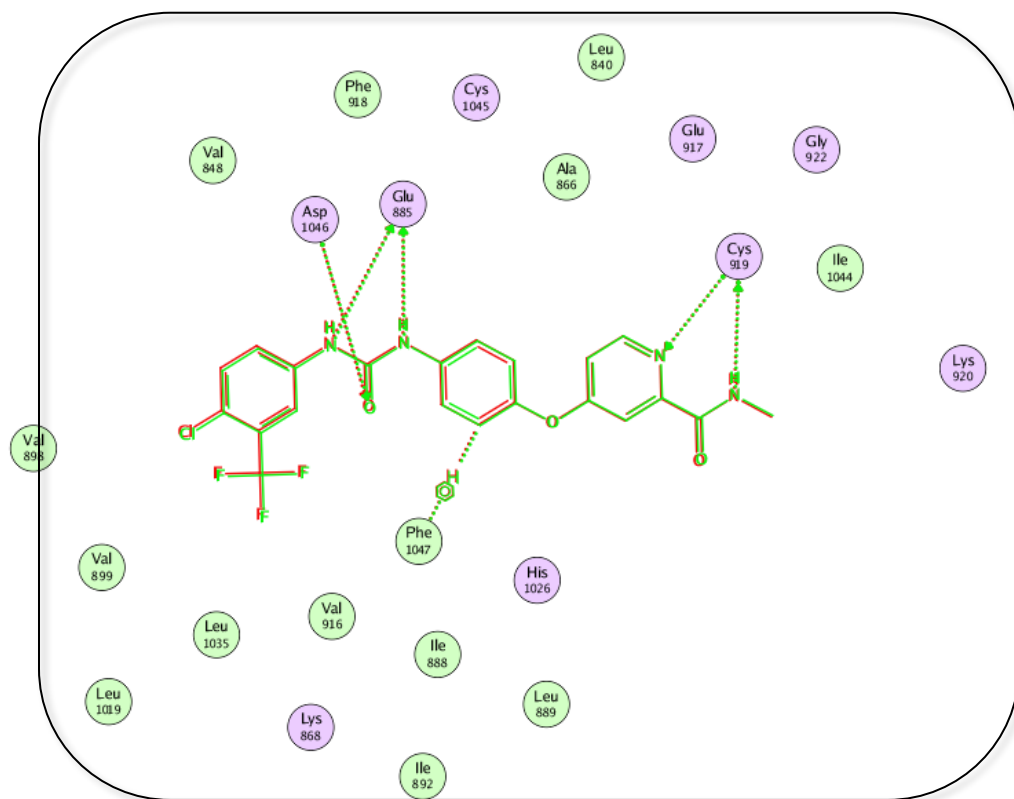


Figure S2 2D Representation of the superimposition of the docking pose (green) and the co-crystallized (purple) of sorafenib in the VEGFR-2 active site with RMSD of 0.21Å.

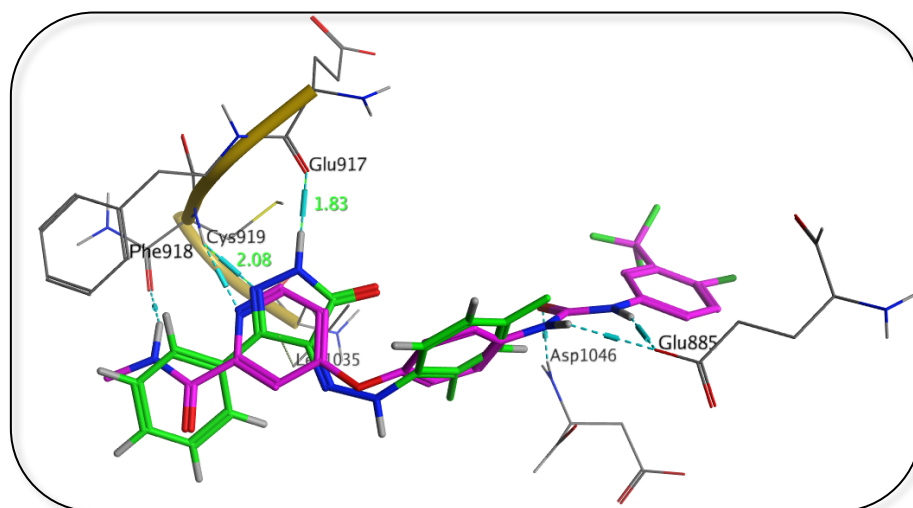


Figure S3 Superimposition of the docking pose of **3i** (green) and the co-crystallized (purple) of sorafenib in the VEGFR-2 active site.

## ***In Vivo* Assay**

### **Animals and tumor cell line**

Adult female Swiss albino mice purchased from Faculty of Pharmacy, Suez Canal University, Ismailia, Egypt, with an average body weight of (18-23) g was used. Mice were housed under constant conditions of 12 h light/dark cycle in a temperature under conditions of controlled humidity ( $22 \pm 2$  °C), with free access to standard laboratory mice food and water. All procedures related to care and maintenance of the animals were performed according to the international guiding principles for animal research and approved by Faculty of Science, Suez Canal University bioethics and animal ethics committee (Approval number REC107/2022).

Solid Ehrlich carcinoma (SEC) were got from the National Cancer Institute (Cairo University, Egypt). The tumor cell line was proliferated in mice through serial intraperitoneal (I.P.) transplantation of a volume of 0.2 mL physiological saline containing  $1 \times 10^6$  viable cells for 24 h. SEC cells were collected 7 days after I.P. implantation. The harvested cells were diluted with saline to obtain a concentration of  $5 \times 10^6$  viable SEC cells/mL. A volume of 0.2 mL saline contains  $1 \times 10^6$  SEC cells that were I.P. implanted into each normal mouse. SEC cells ( $1 \times 10^6$  tumor cells/mouse) were implanted subcutaneously into the right thigh of the hind limb.

The experimental animals were randomly divided into four groups. Group 1 served as the normal saline control. Group 2 served as the SEC control ( $1 \times 10^6$  cells/mouse). Group 3 served as the compound-treated group (6 mg/kg B.Wt., I.P.). Group 4 received the standard anticancer drug of Sorafenib (6 mg/kg BW, I.P.) and is considered as a reference control. Body weight and survival were recorded daily until the 24<sup>th</sup> day in both treated and control groups. At the end of experiment, anesthetized animals were then sacrificed for evaluation of the antitumor activity and histopathological examination.

### **Antitumor potentiality**

It includes tumor volume, weight, and tumor inhibition ration (TIR%). Time interval measurements of tumor volume using digital Vernier caliper (Tricle Brand, Shanghai, China). Measure tumor length and width using clipper and then calculate tumor volume using formulations  $V = (L \times W \times W)/2$ , where V is tumor volume, W is tumor width, L is tumor length. While TIR% was calculated according to the following equation  $\frac{Tumor\ volume\ (Control) - Tumor\ volume\ (treated)}{Tumor\ volume\ (control)} \times 100$ .

### **Histopathological study**

Specimens of liver-sacrificed mice were fixed in 10% saline formalin. The fixed liver specimens were dehydrated in ascending series of ethyl alcohol and embedded in paraffin. Sections at 5 mm thicknesses were stained with hematoxylin and eosin and examined under light microscopy

## References

- [1] M.S. Nafie, K. Arafa, N.K. Sedky, A.A. Alakhdar, R.K. Arafa, Triaryl dicationic DNA minor-groove binders with antioxidant activity display cytotoxicity and induce apoptosis in breast cancer, *Chemico-Biological Interactions*. 324 (2020) 109087. <https://doi.org/10.1016/j.cbi.2020.109087>.
- [2] M. Dicato, L. Plawny, M. Diederich, Anemia in cancer, *Annals of Oncology*. 21 (2010) vii167–vii172. <https://doi.org/10.1093/annonc/mdq284>.
- [3] H.S.A. ElZahabi, M.S. Nafie, D. Osman, N.H. Elghazawy, D.H. Soliman, A.A.H. EL-Helby, R.K. Arafa, Design, synthesis and evaluation of new quinazolin-4-one derivatives as apoptotic enhancers and autophagy inhibitors with potent antitumor activity, *European Journal of Medicinal Chemistry*. 222 (2021) 113609. <https://doi.org/10.1016/j.ejmech.2021.113609>.
- [4] A.T.A. Boraie, E.H. Eltamany, I.A.I. Ali, S.M. Gebriel, M.S. Nafie, Synthesis of new substituted pyridine derivatives as potent anti-liver cancer agents through apoptosis induction: In vitro, in vivo, and in silico integrated approaches, *Bioorganic Chemistry*. 111 (2021) 104877. <https://doi.org/10.1016/j.bioorg.2021.104877>.