

Design, Synthesis, Anticancer Activity, and Solid Lipid Nanoparticle Formulation of Indole-Based Compounds as Novel Pro-Apoptotic Agents Targeting Bcl-2 Protein

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Additional Experimental Detail

- 1) **Table S1:** The Data of Ligand-Docking Studies on Bcl-2 anti-apoptotic protein target.
- 2) **Figure S1:** Standard calibration curve of 8b using HPLC-UV analysis. All data are represented as mean of three independent experiments \pm SEM.
- 3) **Figure S2:** NMR spectra of all new compounds

Table S1: The Data of Ligand-Docking Studies on Bcl-2 anti-apoptotic protein target

Ligand name	MOE docking Scores (S) ^a (Kcal mol ⁻¹)	RMSD ^b (Å)	Ligand-target interaction description [Type; Length (Å); Angle (°); Binding Residues]
3a	-4.2200	1.7838	Hydrogen bonding ; 2.51 Å ; 148.0° ; Asp111-C=O side chain with benzimidazole (NH) ring π - π stacking ; 3.85 Å ; Tyr108 side chain with benzimidazole ring
7b	-4.7696	1.5547	Hydrogen bonding ; 2.98 Å ; 112.5° ; Arg146=NH side chain with 4-MeO
7d	-4.5808	1.7670	Hydrogen bonding ; 2.10 Å ; 149.9° ; Arg146=NH side chain with pyridine (N) ring π -Hydrogen contact ; 4.15 Å ; Leu137 side chain with pyridine ring π -Hydrogen contact ; 4.17 Å ; Leu137 side chain with pyridine ring
8a	-4.7696	1.2117	Hydrogen bonding ; 2.10 Å ; 158.7° ; Arg146=NH side chain with NHC=O linker
8b	-4.8585	1.6829	Hydrogen bonding ; 2.15 Å ; 175.5° ; Arg146=NH side chain with NHC=O linker
8c	-6.2460	1.7374	Hydrogen bonding ; 2.66 Å ; 129.0° ; Arg146=NH side chain with indole (NH) ring π -Hydrogen interaction ; 3.33 Å ; Arg146-NH side chain with indole ring
8d	-6.1846	1.3960	Hydrogen bonding ; 2.08 Å ; 156.0° ; Arg146=NH side chain with NHC=O linker Hydrogen bonding ; 2.53 Å ; 142.5° ; Arg146-NH side chain with NHC=O linker
8f	-5.7070	2.6217	Hydrogen bonding ; 2.30 Å ; 160.1° ; Arg146=NH side chain with NHC=O linker π -Hydrogen interaction ; 3.10 Å ; Arg146-NH side chain with indole ring

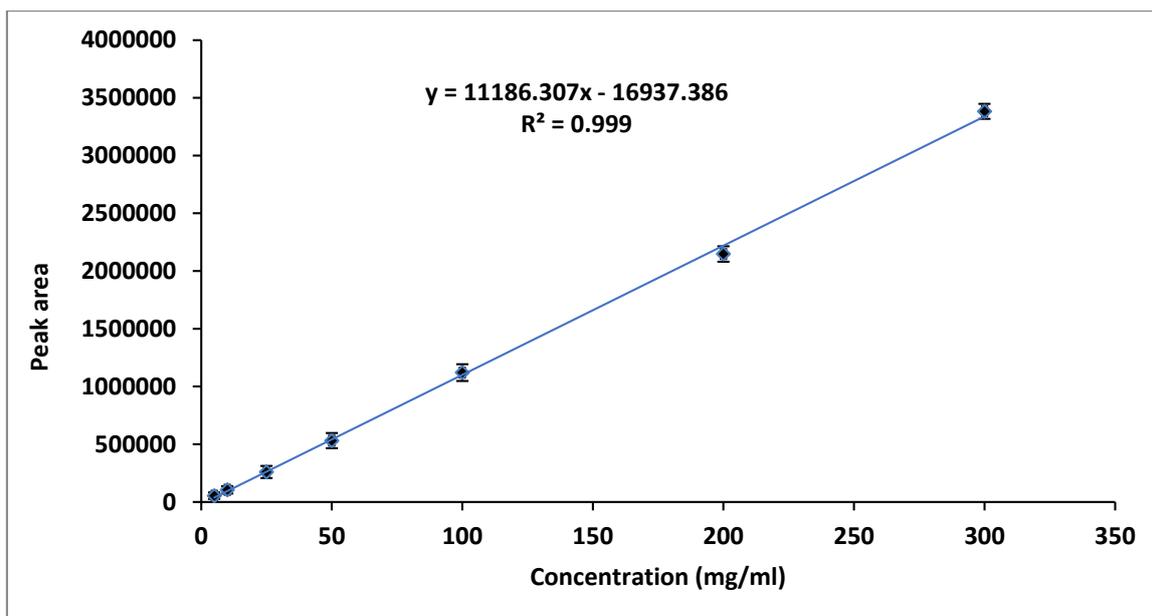
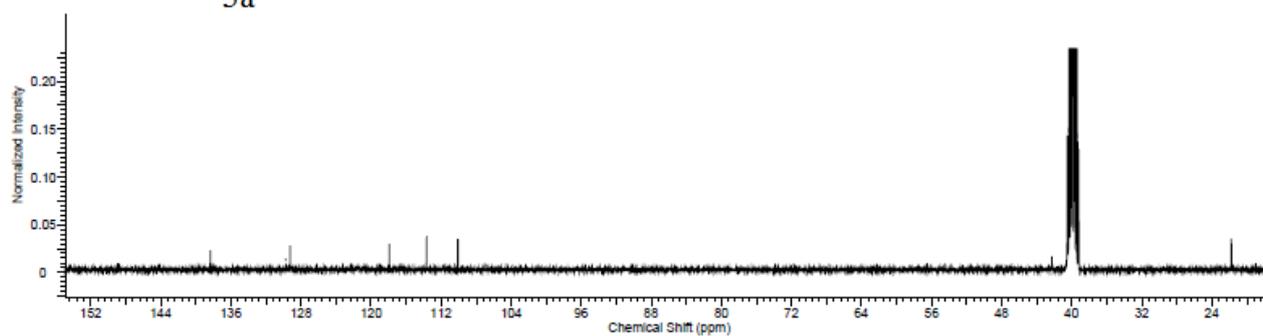
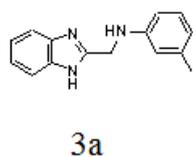
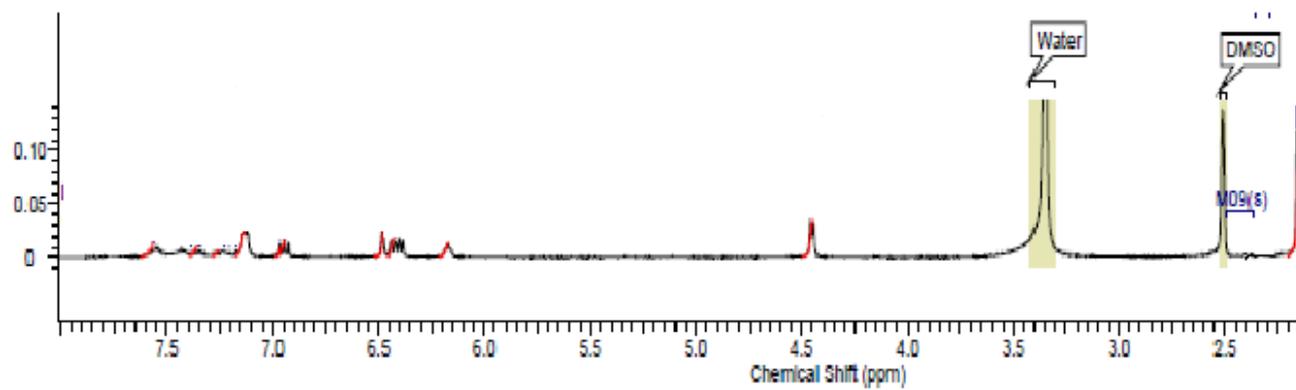
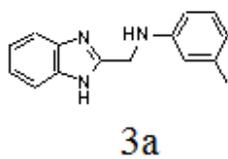
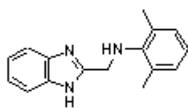


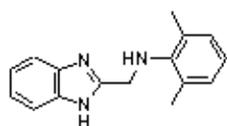
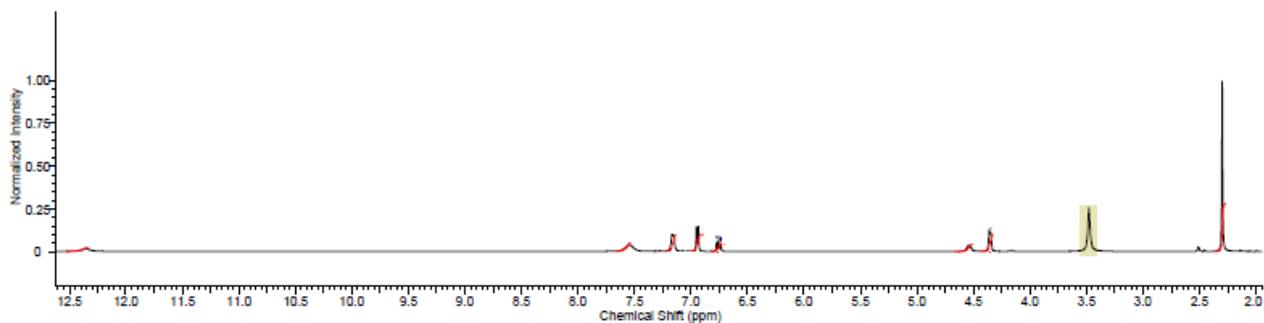
Figure S1: Standard calibration curve of 8b using HPLC-UV analysis. All data are represented as mean of three independent experiments \pm SEM.

Figure S2: NMR spectra of all new compounds

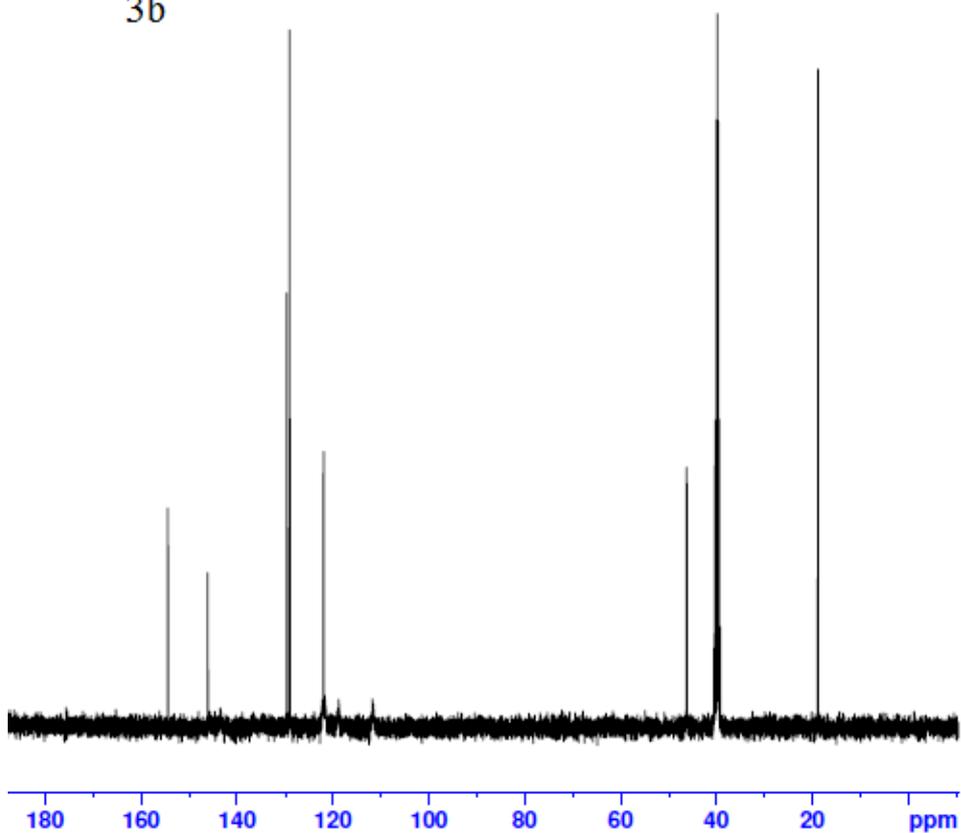




3b



3b



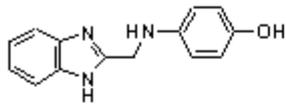
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 SOLVENT DMSO
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 FIDRES 0.366798 Hz
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 RG 205.37
 DW 20.800 usec
 DE 6.50 usec
 TE 299.0 K
 D1 2.0000000 sec
 D11 0.0300000 sec
 TD0 1

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 NUC1 13C
 P1 10.00 usec
 P1W1 47.0000000 W

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 NUC2 1H
 CPDPRG2 waltz16
 PCPD2 90.00 usec
 P1W2 18.0000000 W
 P1W12 0.34722000 W
 P1W13 0.28125000 W

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 GB 0
 PC 1.40



3d

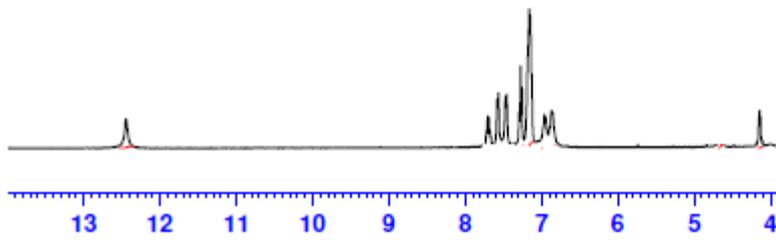


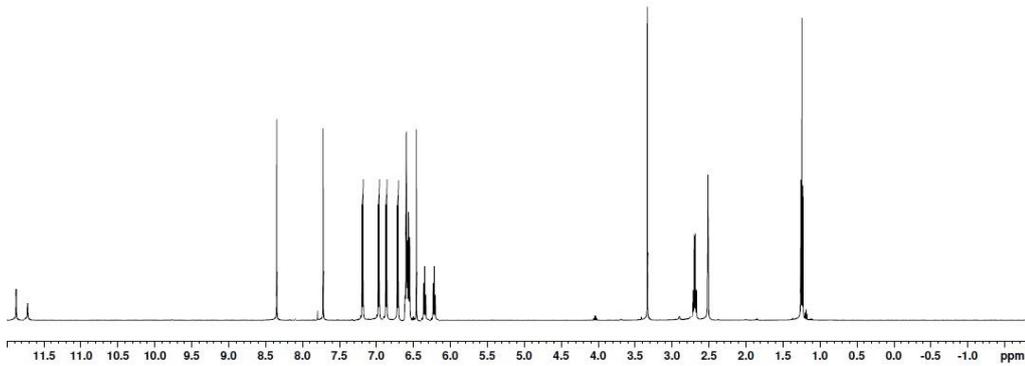
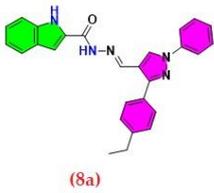
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 P1 12.00 usec
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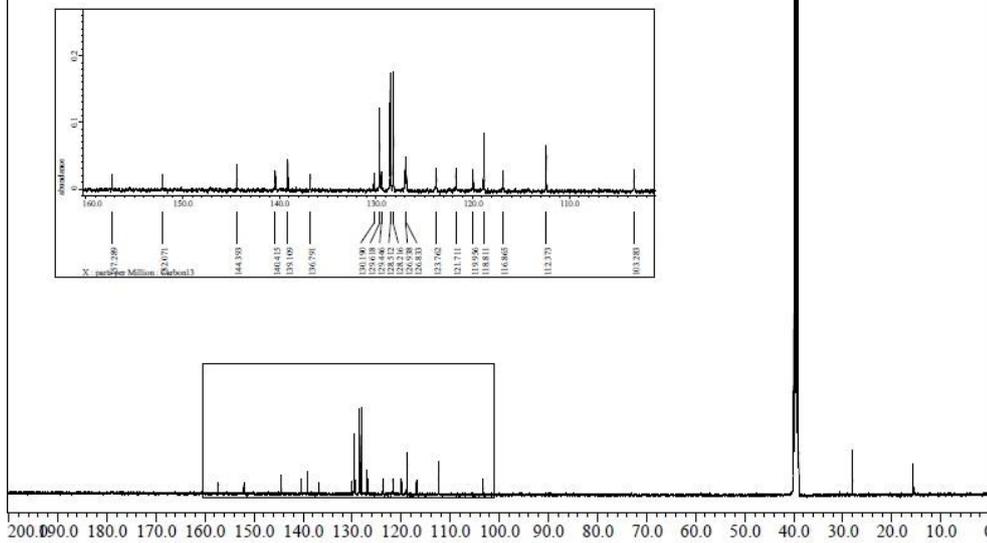
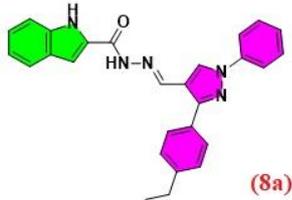
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Current Time = 1-SBP-2019 15:30:47

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Dim Size     = 13107
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Dim Units    = [ppm]
Dimensions   = X
Site         = JNM-ECAS500II
Spectrometer = DELTA2_NMR

Field Strength = 11.7473579[T] (500 MHz)
X_Acq_Duration = 1.4548992[s]
X_Domain      = 15
X_Freq        = 500.15991521[MHz]
X_Offset      = 6[ppm]
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X_Resolution  = 0.68733284[Hz]
X_Sweep       = 11.26126126[kHz]
X_Sweep_Clippped = 9.00900001[kHz]
Irr_Domain    = Proton
Irr_Freq      = 500.15991521[MHz]
Irr_Offset    = 5.0[ppm]
Tri_Domain    = Proton
Tri_Freq      = 500.15991521[MHz]
Tri_Offset    = 5.0[ppm]
Clipped       = FALSE
Scans         = 40
Total Scans   = 40

Relaxation_Delay = 5[s]
Recov_Gain       = 44
Temp_Get         = 20.7[dC]
X_Acq_Width     = 14.5[us]
X_Acq_Time      = 1.4548992[s]
X_Angle         = 45[deg]
X_Attn          = 4.9[dB]
X_Pulse         = 7[us]
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Tri_Mode        = Off
Densit_Preset   = FLSB
Initial_Wait    = 1[s]
Repetition_Time = 6.4548992[s]
  
```



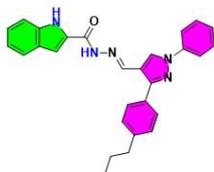
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Author       = delta
Experiment   = carbon_jmp
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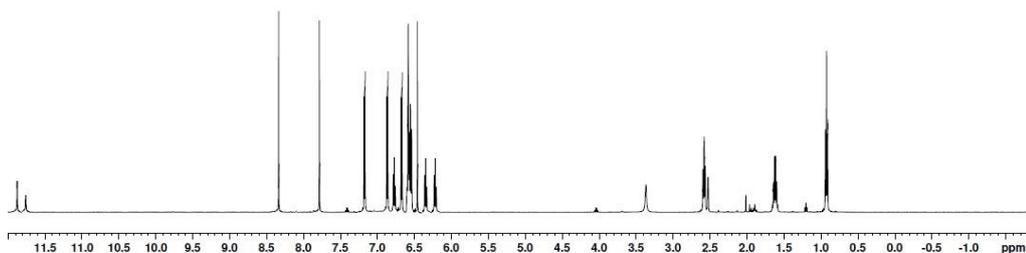
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Spectrometer = DELTA2_NMR

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X_Domain      = 13C
X_Freq        = 125.76529768[MHz]
X_Offset      = 100[ppm]
X_Points      = 32768
X_Prescans    = 4
X_Resolution  = 1.19959034[Hz]
X_Sweep       = 39.3081761[kHz]
X_Sweep_Clippped = 31.44654088[kHz]
Irr_Domain    = Proton
Irr_Freq      = 500.15991521[MHz]
Irr_Offset    = 5.0[ppm]
Clipped       = TRUE
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Total Scans   = 3700

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Recov_Gain       = 58
Temp_Get         = 20.2[dC]
X_90_Width      = 52.78[us]
X_Acq_Time      = 0.83361792[s]
X_Angle         = 30[deg]
X_Attn          = 9.7[dB]
X_Pulse         = 17.59333333[us]
Irr_Attn_Dec    = 20.948[dB]
Irr_Attn_Noe    = 20.948[dB]
Irr_Mode        = WALTZ
Irr_Pwidth     = 92[us]
Decoupling      = TRUE
Initial_Wait    = 1[s]
Noe              = TRUE
Noe Time        = 2[s]
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```



(8b)



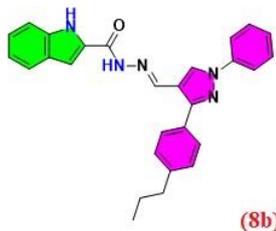
Mansoura University
JEOL
ECA-500 II

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 Experiment = proton.jpg
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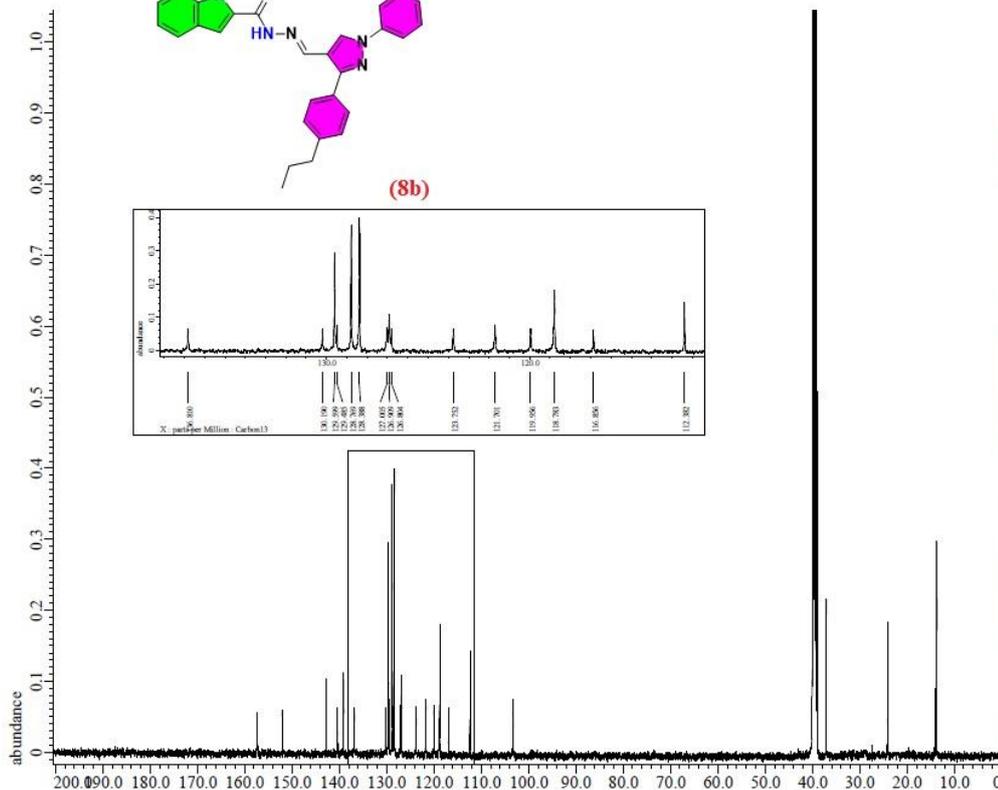
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 Spectrometer = DBLTA2_NMR

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 X_Resolution = 0.68733264 [Hz]
 X_Sweep = 11.26126126 [kHz]
 X_Sweep_Clipped = 9.00900901 [kHz]
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 Ix1_Freq = 500.15991521 [MHz]
 Ix1_Offset = 5.0 [ppm]
 Ix1_Domain = Proton
 Ix1_Freq = 500.15991521 [MHz]
 Ix1_Offset = 5.0 [ppm]
 Clipped = FALSE
 Scans = 40
 Total Scans = 40

Relaxation_Delay = 5 [s]
 Recvx_Gain = 36
 Temp_Get = 21.1 [dC]
 X_90_Width = 14.1 [us]
 X_Acq_Time = 1.4548992 [s]
 X_Angle = 45 [deg]
 X_Atn = 4.9 [dB]
 X_Pulse = 7.25 [us]
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 Repetition_Time = 6.4548992 [s]



(8b)



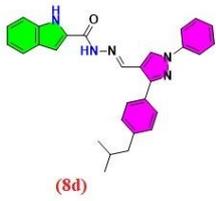
Mansoura University
JEOL
ECA-500 II

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 Spectrometer = DBLTA2_NMR

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 X_Domain = 13C
 X_Freq = 125.76529768 [MHz]
 X_Offset = 100 [ppm]
 X_Points = 32768
 X_Prescans = 4
 X_Resolution = 1.19959034 [Hz]
 X_Sweep = 39.3081761 [kHz]
 X_Sweep_Clipped = 31.44654068 [kHz]
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 Ix1_Freq = 500.15991521 [MHz]
 Ix1_Offset = 5.0 [ppm]
 Clipped = FALSE
 Scans = 1099
 Total Scans = 1099

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 Recvx_Gain = 58
 Temp_Get = 18.7 [dC]
 X_90_Width = 52.78 [us]
 X_Acq_Time = 0.83361792 [s]
 X_Angle = 30 [deg]
 X_Atn = 9.7 [dB]
 X_Pulse = 17.59333333 [us]
 Ix1_Atn_Dec = 20.948 [dB]
 Ix1_Atn_Noise = 20.948 [dB]
 Ix1_Noise = WALTZ
 Ix1_Fwidth = 92 [us]
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 Initial_Wait = 1 [s]
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 Repetition Time = 2.83361792 [s]



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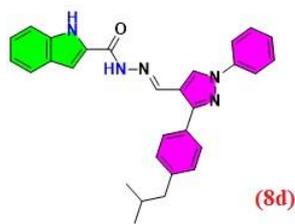
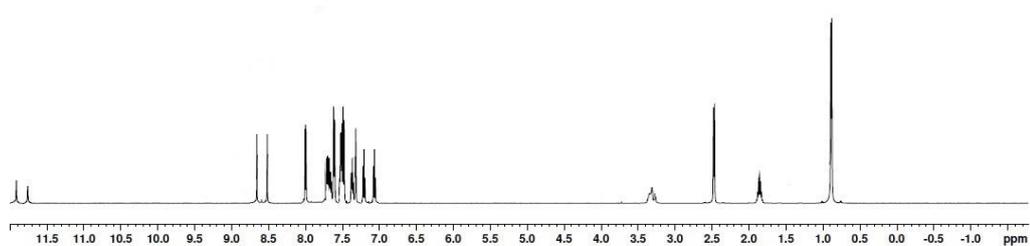
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X_Sweep_Clipped = 9.00900901[kHz]
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X_Acq_Time = 1.4548992 [s]
X_Angle = 45 [deg]
X_Atn = 4.9 [dB]
X_Pulse = 7.25 [us]
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Densit_Preset = FLG2B
Initial_Wait = 1 [s]
Repetition_Time = 6.4548992 [s]

```



```

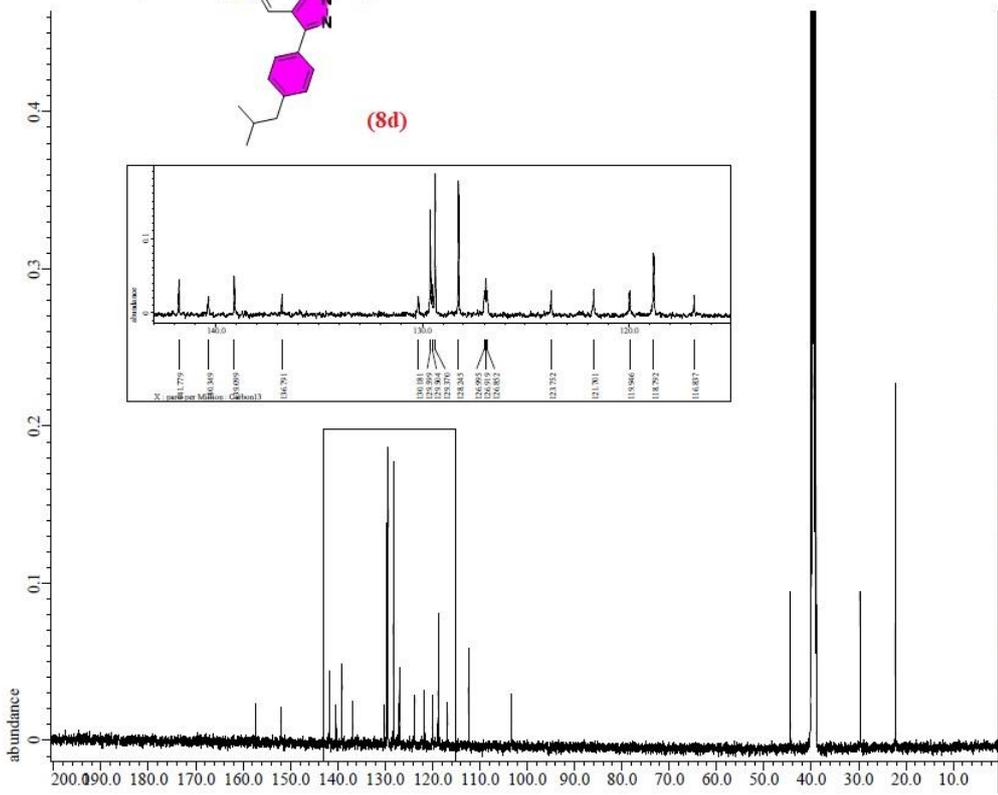
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Experiment = carbon_jmp
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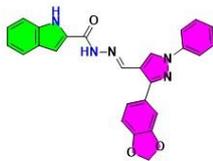
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X_Offset = 100 [ppm]
X_Points = 32768
X_Prescans = 4
X_Resolution = 1.19959034 [Hz]
X_Sweep = 39.3081761 [kHz]
X_Sweep_Clipped = 31.44654088 [kHz]
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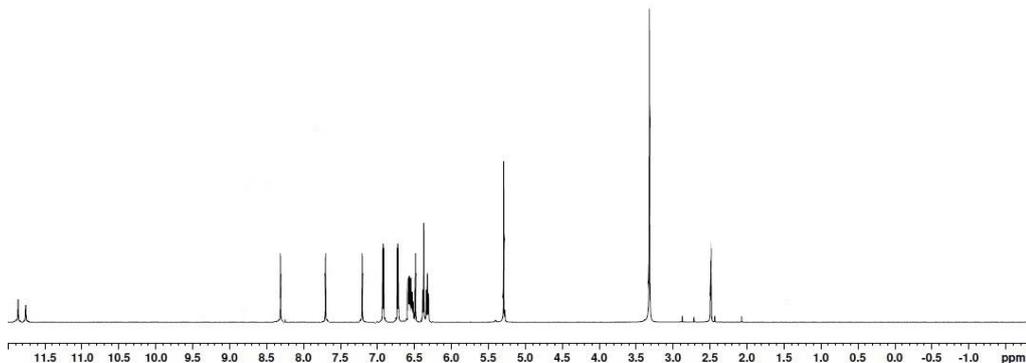
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X_Angle = 30 [deg]
X_Atn = 9.7 [dB]
X_Pulse = 17.59333333 [us]
Irx_Atn_Dec = 20.948 [dB]
Irx_Atn_NoE = 20.948 [dB]
Irx_Noise = WALTZ
Irx_Width = 92 [us]
Decoupling = TRUB
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Noe_Time = 2 [s]
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```





(8e)



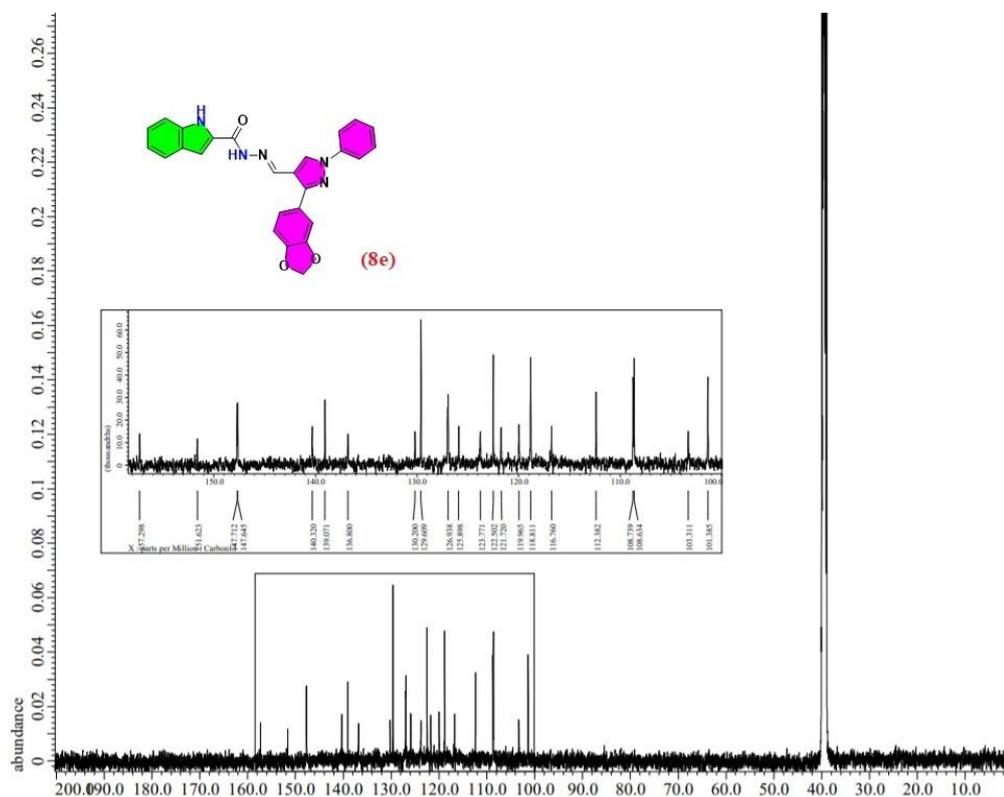
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Experiment   = proton.jpg
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Revision Time = 30-JUL-2019 14:10:03
Current Time  = 30-JUL-2019 14:11:10

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Dim Title    = Proton
Dim Units    = [ppm]
Dimensions   = X
Site         = JNM-ECA500II
Spectrometer = DELTA2_NMR

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X_Acq Duration = 1.4548992[s]
X_Domain      = 1H
X_Freq        = 500.15991521[MHz]
X_Offset      = 6[ppm]
X_Points      = 14264
X_Prescans    = 1
X_Resolution  = 0.68733264[Hz]
X_Sweep       = 11.26126126[kHz]
X_Sweep Clipped = 9.00909091[kHz]
Irr_Domain    = Proton
Irr_Freq      = 500.15991521[MHz]
Irr_Offset    = 5.0[ppm]
Irr_Domain    = Proton
Tri_Freq      = 500.15991521[MHz]
Tri_Offset    = 5.0[ppm]
Clipped       = FALSE
Scans         = 40
Total Scans   = 40

Relaxation Delay = 5[s]
Recvr Gain      = 36
Temp_Get       = 21.1[dc]
X_90_Width     = 14.3[us]
X_Acq Time     = 1.4548992[s]
X_Angle        = 45[deg]
X_Atn         = 4.9[db]
X_Pulse        = 7.25[us]
Irr_Atn_Dec   = Off
Irr_Atn_Noise = Off
Dens1_Preset  = FALSB
Initial Wait   = 1[s]
Repetition Time = 6.4548992[s]
  
```



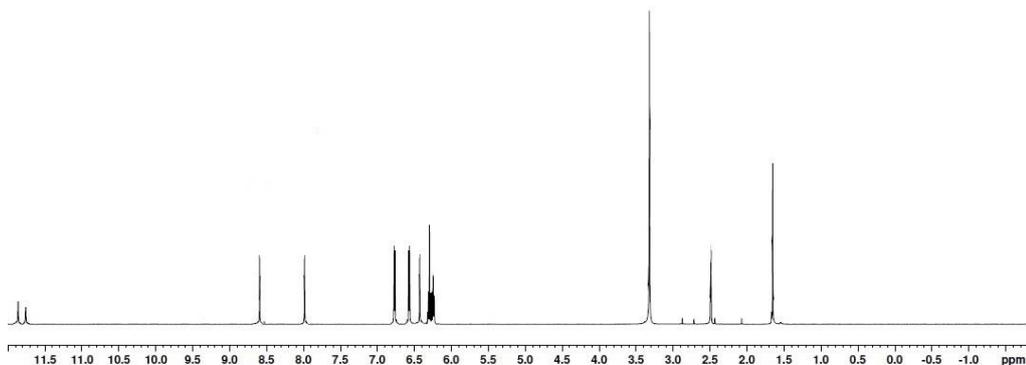
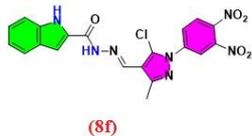
```

Filename      = Manar Nagy_8_carbon-1-1
Author       = delta
Experiment   = carbon.jpg
Sample Id    = Manar Nagy_8
Solvent      = DMSO-D6
Creation Time = 18-AUG-2019 20:20:43
Revision Time = 19-AUG-2019 09:09:26
Current Time  = 19-AUG-2019 09:10:27

Comment      = single pulse decoupled
Data Format   = 1D REAL
Dim Size     = 26214
Dim Title    = Carbon13
Dim Units    = [ppm]
Dimensions   = X
Site         = JNM-ECA500II
Spectrometer = DELTA2_NMR

Field Strength = 11.7473579[T] (500[MHz])
X_Acq Duration = 0.83361792[s]
X_Domain      = 13C
X_Freq        = 125.76529768[MHz]
X_Offset      = 100[ppm]
X_Points      = 32768
X_Prescans    = 4
X_Resolution  = 1.19959034[Hz]
X_Sweep       = 39.3081761[kHz]
X_Sweep Clipped = 31.44654088[kHz]
Irr_Domain    = Proton
Irr_Freq      = 500.15991521[MHz]
Irr_Offset    = 5.0[ppm]
Clipped       = TRUE
Scans         = 2500
Total Scans   = 2500

Relaxation Delay = 2[s]
Recvr Gain      = 58
Temp_Get       = 18.3[dc]
X_90_Width     = 52.78[us]
X_Acq Time     = 0.83361792[s]
X_Angle        = 30[deg]
X_Atn         = 9.7[db]
X_Pulse        = 17.59333333[us]
Irr_Atn_Dec   = 20.948[db]
Irr_Atn_Noise = 20.948[db]
Irr_Noise     = WALTZ
Irr_Pwidth    = 92[us]
Decoupling    = TRUE
Initial Wait   = 1[s]
Noe           = TRUE
Noe Time      = 2[s]
Repetition Time = 2.83361792[s]
  
```



```

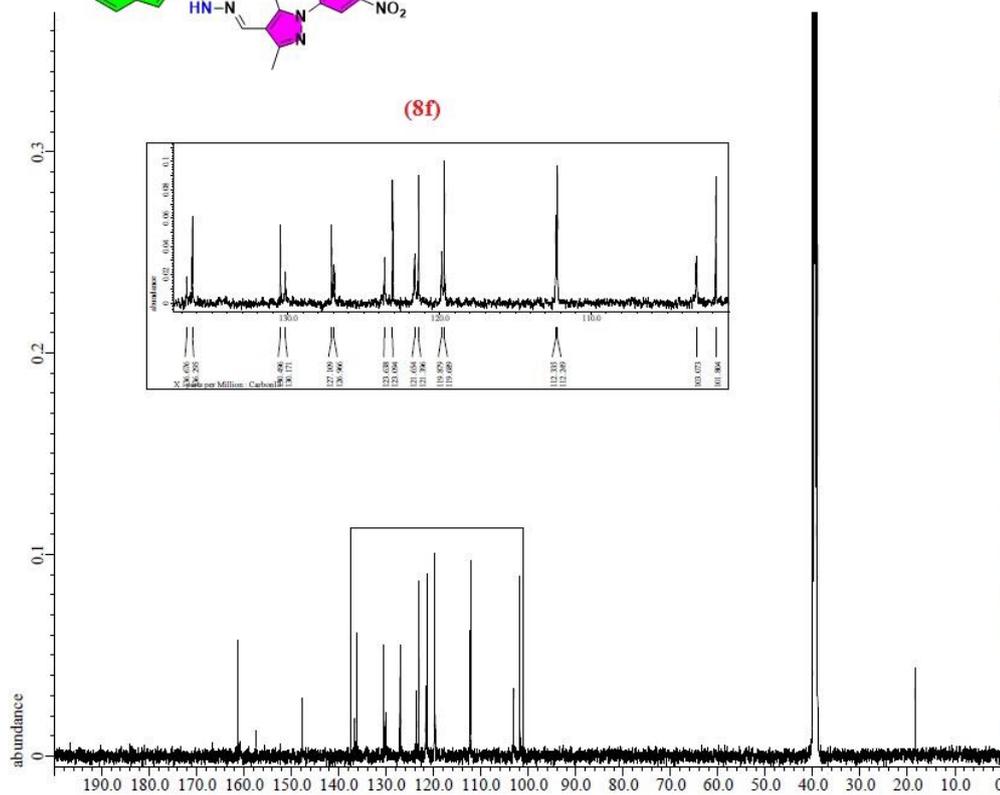
Filename = Manar Nagy10_proton-1-11
Author = delta
Experiment = proton JMP
Sample Id = Manar Nagy_11
Solvent = DMSO-D6
Creation Time = 30-JUL-2019 15:03:21
Revision Time = 30-JUL-2019 15:10:47
Current Time = 30-JUL-2019 15:11:10

Comment = single pulse
Data Format = 1D CDEXX
Dim Size = 13107
Dim Title = Proton
Dim Units = [ppm]
Dimensions = X
Site = JNM-BCA500II
Spectrometer = DBLFA2_NMR

Field Strength = 11.7473579[T] (500 MHz)
X_Acq Duration = 1.4548992[s]
X_Domain = 1H
X_Freq = 500.15991521[MHz]
X_Offset = 6 [ppm]
X_Points = 14384
X_Prescans = 1
X_Resolution = 0.6873284[Hz]
X_Sweep = 11.26126126[kHz]
X_Sweep Clipped = 9.0090901[kHz]
Irr_Domain = Proton
Irr_Freq = 500.15991521[MHz]
Irr_Offset = 5.0 [ppm]
Irr_Domain = Proton
Tri_Domain = Proton
Tri_Freq = 500.15991521[MHz]
Tri_Offset = 5.0 [ppm]
Clipped = FALSE
Scans = 40
Total Scans = 40

Relaxation Delay = 5 [s]
Recvr Gain = 36
Temp Set = 21.1[dC]
X_90 Width = 13 [us]
X_Acq Time = 1.4548992[s]
X_Angle = 45 [deg]
X_Atn = 4.9 [dB]
X_Pulse = 7.25 [us]
Irr_Mode = Off
Tri_Mode = Off
Dens1 Preset = FALSB
Initial Wait = 1 [s]
Repetition Time = 6.4548992[s]

```



```

Filename = Manar Nagy_11_carbon-1-1
Author = delta
Experiment = carbon JMP
Sample Id = Manar Nagy_11
Solvent = DMSO-D6
Creation Time = 10-SEP-2019 13:58:07
Revision Time = 15-SEP-2019 15:04:31
Current Time = 15-SEP-2019 15:05:11

Comment = single pulse decoupled g
Data Format = 1D REAL
Dim Size = 26214
Dim Title = Carbon13
Dim Units = [ppm]
Dimensions = X
Site = JNM-BCA500II
Spectrometer = DBLFA2_NMR

Field Strength = 11.7473579[T] (500 MHz)
X_Acq Duration = 0.83361792[s]
X_Domain = 13C
X_Freq = 125.76529768[MHz]
X_Offset = 100 [ppm]
X_Points = 32768
X_Prescans = 4
X_Resolution = 1.19959034[Hz]
X_Sweep = 39.3081761[kHz]
X_Sweep Clipped = 31.44654088[kHz]
Irr_Domain = Proton
Irr_Freq = 500.15991521[MHz]
Irr_Offset = 5.0 [ppm]
Clipped = TRUE
Scans = 2402
Total Scans = 2402

Relaxation Delay = 2 [s]
Recvr Gain = 58
Temp Set = 19.9[dC]
X_90 Width = 52.78 [us]
X_Acq Time = 0.83361792[s]
X_Angle = 30 [deg]
X_Atn = 9.7 [dB]
X_Pulse = 17.59333333 [us]
Irr_Atn Dec = 20.948 [dB]
Irr_Atn Noe = 20.948 [dB]
Irr_Noise = WJLSE
Irr_Pwidth = 92 [us]
Decoupling = TRUB
Initial Wait = 1 [s]
Noe = TRUB
Noe Time = 2 [s]
Repetition Time = 2.83361792[s]

```