

Supporting Information

Marine-Inspired Bis-indoles Possessing Anti-proliferative Activity against Breast Cancer; Design, Synthesis and Biological Evaluation

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Biological Evaluations

1. Anti-proliferative action toward human breast cell lines

The two examined human breast cancer cell lines (MCF-7 and Breast MDA-MB-231), and non-tumorigenic human breast epithelial cell line (MCF-10A) have been obtained from American Type Culture Collection (ATCC). The cells were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% heat inactivated fetal calf serum (GIBCO), penicillin (100 U/ml) and streptomycin (100 µg/ml) at 37 °C in humidified atmosphere containing 5% CO₂. Cells at a concentration of 0.50×10^6 were grown in a 25 cm² flask in 5 ml of culture medium.

The anti-proliferative activity of the tested compounds was measured *in vitro* using the Sulfo-Rhodamine-B stain (SRB) assay. Briefly, Cells were inoculated in 96-well microtiter plate (5×10^4 cells/ well) for 24 h before treatment with the tested compounds to allow attachment of cell to the wall of the plate. Tested compounds were dissolved in DMSO at 1 mg/ml immediately before use and diluted to the appropriate volume just before addition to the cell culture. Different concentrations of tested compounds, doxorubicin and sorafenib were added to the cells (three wells were prepared for each individual dose). Cells were incubated with the compounds for 48 h at 37°C and in atmosphere of 5% CO₂. After 48 h cells were fixed, washed, and stained for 30 min with 0.4% (w/v) SRB dissolved in 1% acetic acid. Unbound dye was removed by four washes with 1% acetic acid, and attached stain was recovered with Tris-EDTA buffer. Color intensity was measured in an ELISA reader. The relation between percent of surviving fraction and drug concentration is plotted to get the survival curve for each cell line. The concentration required for 50% inhibition of cell viability (IC₅₀) was calculated.

2. Cell Cycle Analysis

Breast cancer MCF-7 cells were treated with bis-indoles **7e** and **9a** for 24 h (at their IC₅₀ concentration), and then cells were washed twice with ice-cold phosphate buffered saline (PBS). Subsequently, the treated cells were collected by centrifugation, fixed in ice-cold 70% (*v/v*) ethanol, washed with PBS, re-suspended with 100 µg/mL RNase, stained with 40 µg/mL PI, and analyzed by flow cytometry using FACS Calibur (Becton Dickinson, BD, Franklin Lakes, NJ, USA). The cell cycle distributions were calculated using CellQuest software 5.1 (Becton Dickinson).

3. ELISA Immunoassay

The levels of the pro-apoptotic markers (Bax, caspase-3 and p53) as well as the anti-apoptotic marker Bcl-2 were determined using ELISA colorimetric kits per the manufacturer's instructions. Breast cancer MCF-7 cells were cultured as a monolayer in T-25 flasks and were seeded to attain 30% confluency prior to treatment. Cells were then treated separately with bis-indoles **7e** and **9a** at their IC₅₀ concentrations for 48 h. At the end of treatment, cells were collected via trypsinization and centrifuged at 10,000 rpm. The pellet was then rinsed with PBS and lysed in RIPA lysis buffer at 4 °C for 45 min, then centrifuged at 14,000 rpm for 20 min to remove the cellular debris. Lysates were then collected and stored at -80 °C for later protein determination using Pierce BCA Protein Assay Kit according to manufacturer's recommendations.

The cell lysate was diluted 10 times, and 100 µL (50 mg protein) was added to the wells of four separate microtiter plates for the four ELISA kits that were pre-coated with primary antibodies specific to Bax, Bcl-2, caspase-3 and p53 proteins, respectively. A secondary biotin-linked antibody specific to the protein captured by the primary antibody was further added to bind the captured protein, forming a "sandwich" of specific antibodies around the desired protein in the cell lysate. The streptavidin-HRP complex was then used to bind the biotin-linked secondary antibody through its streptavidin portion. The HRP domain reacted with the added TMB substrate to form a colored product that measured at 450 nm by a plate reader ChroMate-4300 after the reaction was terminated *via* the addition of stop solution.

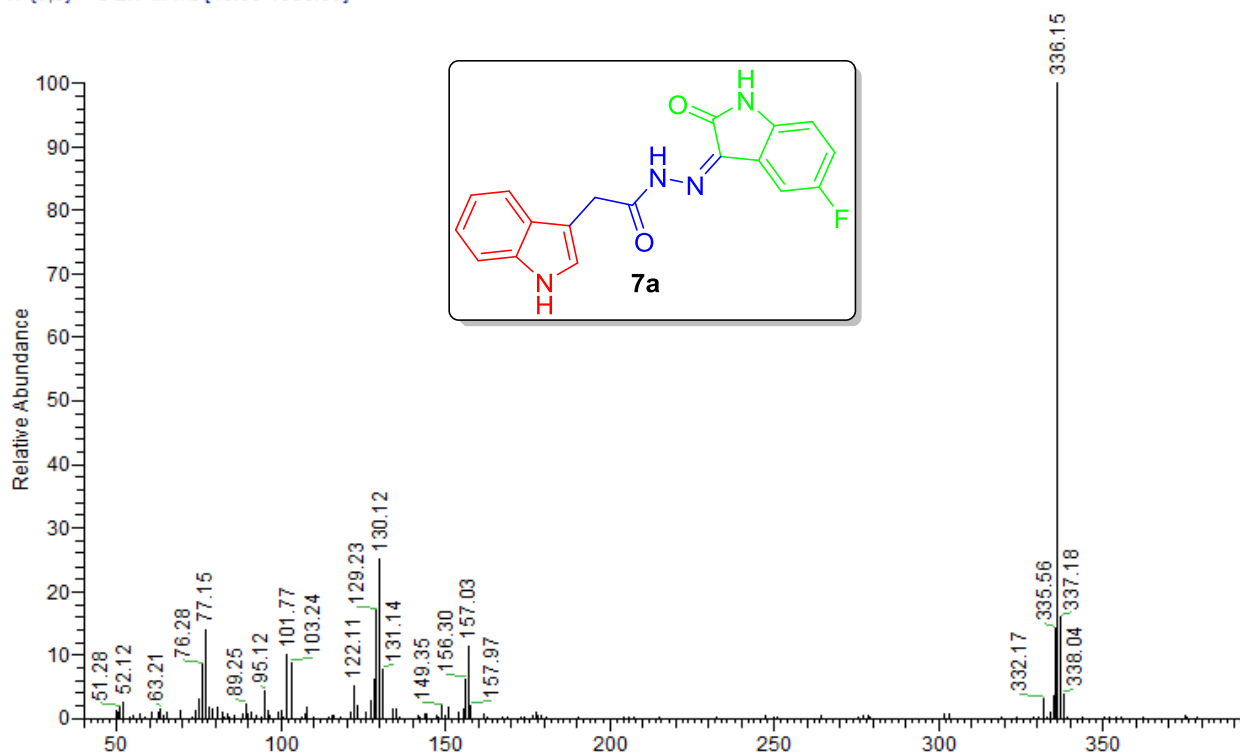
4. Annexin V-FITC Apoptosis Assay

Phosphatidylserine externalization was assayed using Annexin V-FITC/PI apoptosis detection kit (BD Biosciences, USA) according to the manufacturer's instructions. Breast cancer MCF-7 cells were cultured to a monolayer then treated with bis-indoles **7e** and **9a** at their IC₅₀ concentration. Briefly, cells were then harvested *via* trypsinization, and rinsed twice in PBS followed by binding buffer. Moreover, cells were re-suspended in 100 µL of binding buffer with the addition of 1 µL of FITC-Annexin V followed by an incubation period of 30 min at 4 °C. Cells were then rinsed in binding buffer and resuspended in 150 µL of binding buffer with the addition of 1 µL of DAPI (1 µg/µL in PBS). Cells were then analyzed using the flow cytometer BD FACS Canto II and the results were interpreted with FlowJo7.6.4 software (Tree Star, Ashland, OR, USA).

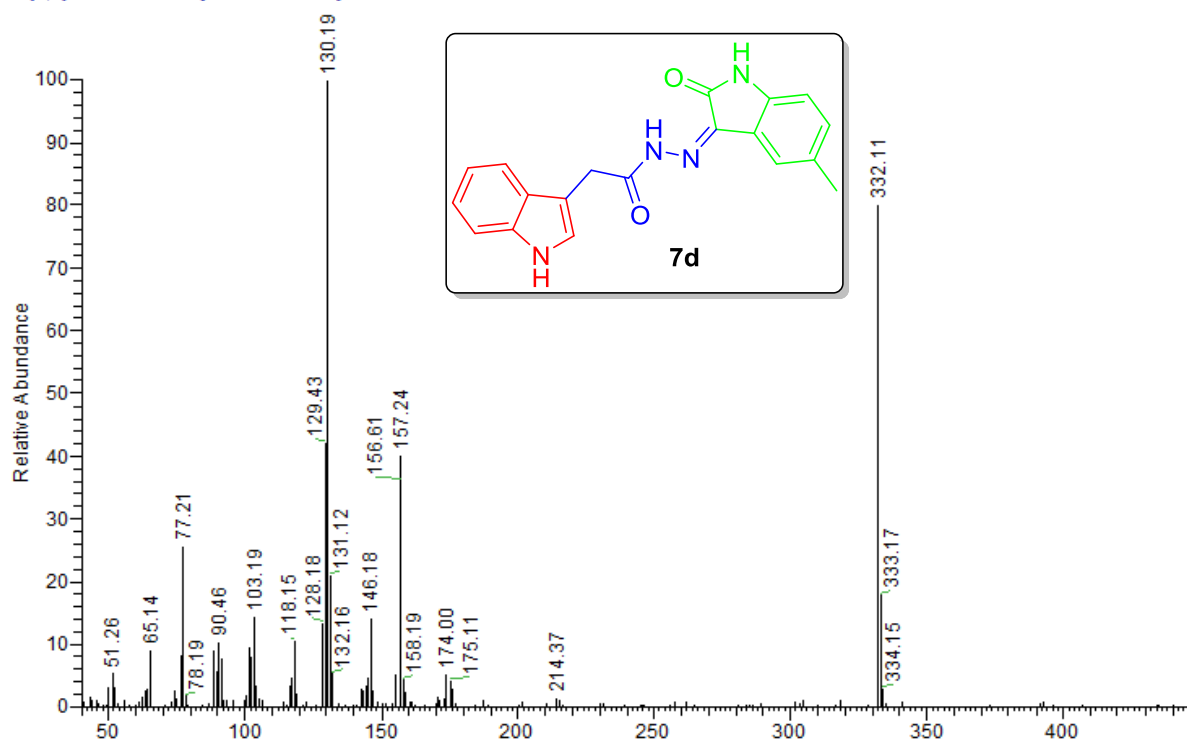
5. CDK2 kinase inhibitory activity

Reaction Biology Corp. Kinase HotSpotSM service (<http://www.reactionbiology.com>) was used for screening of tested compounds. CDK2 Kinase inhibitory activity was assessed by the HotSpot assay platform, which contained specific kinase/substrate pairs along with required cofactors. Base reaction buffer: 20 mM Hepes (pH 7.5), 10 mM MgCl₂, 1 mM EGTA, 0.02% Brij35, 0.02 mg/ml BSA, 0.1 mM Na₃VO₄, 2 mM DTT, 1% DMSO. Testing compounds were dissolved in 100% DMSO to specific concentration. The serial dilution was conducted by Integra Viaflo Assist in DMSO. The reaction mixture containing the examined compound and ³³P-ATP was incubated at room temperature for 2 h and radioactivity was detected by filter-binding method. Kinase activity data were expressed as the percent remaining kinase activity in test samples compared to vehicle (dimethyl sulfoxide) reactions.

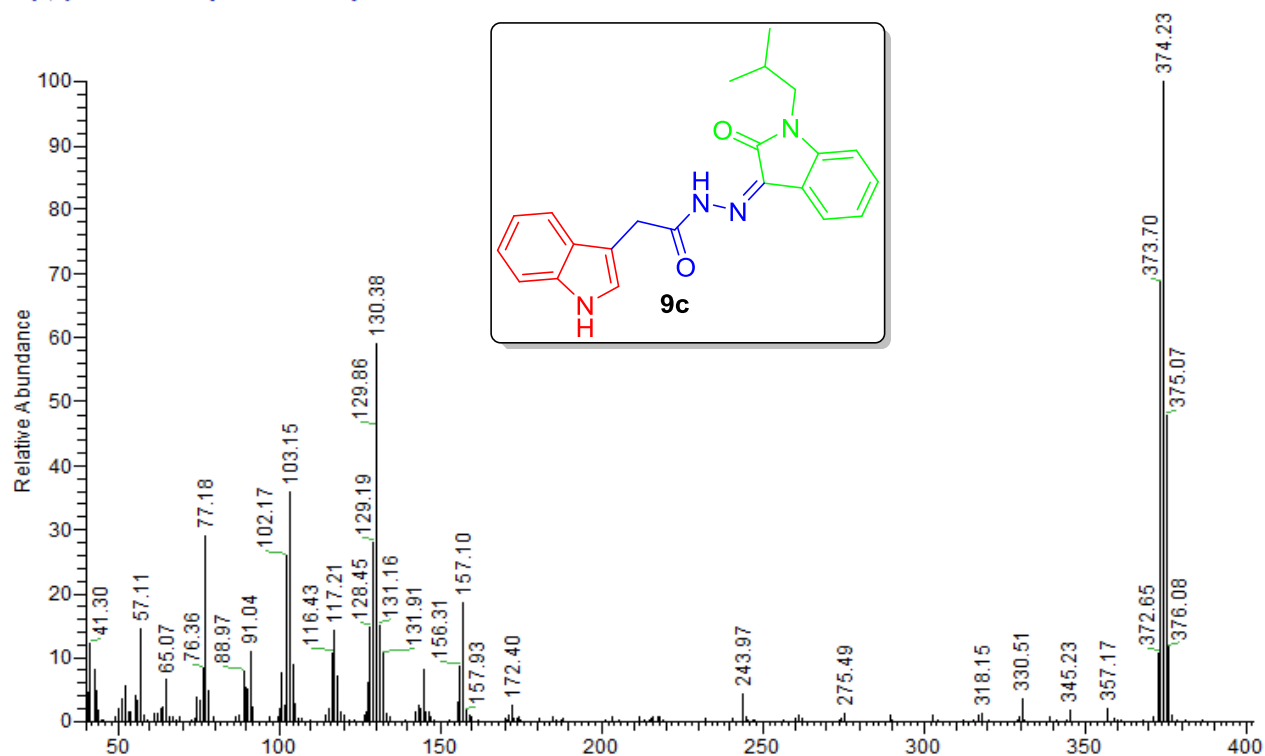
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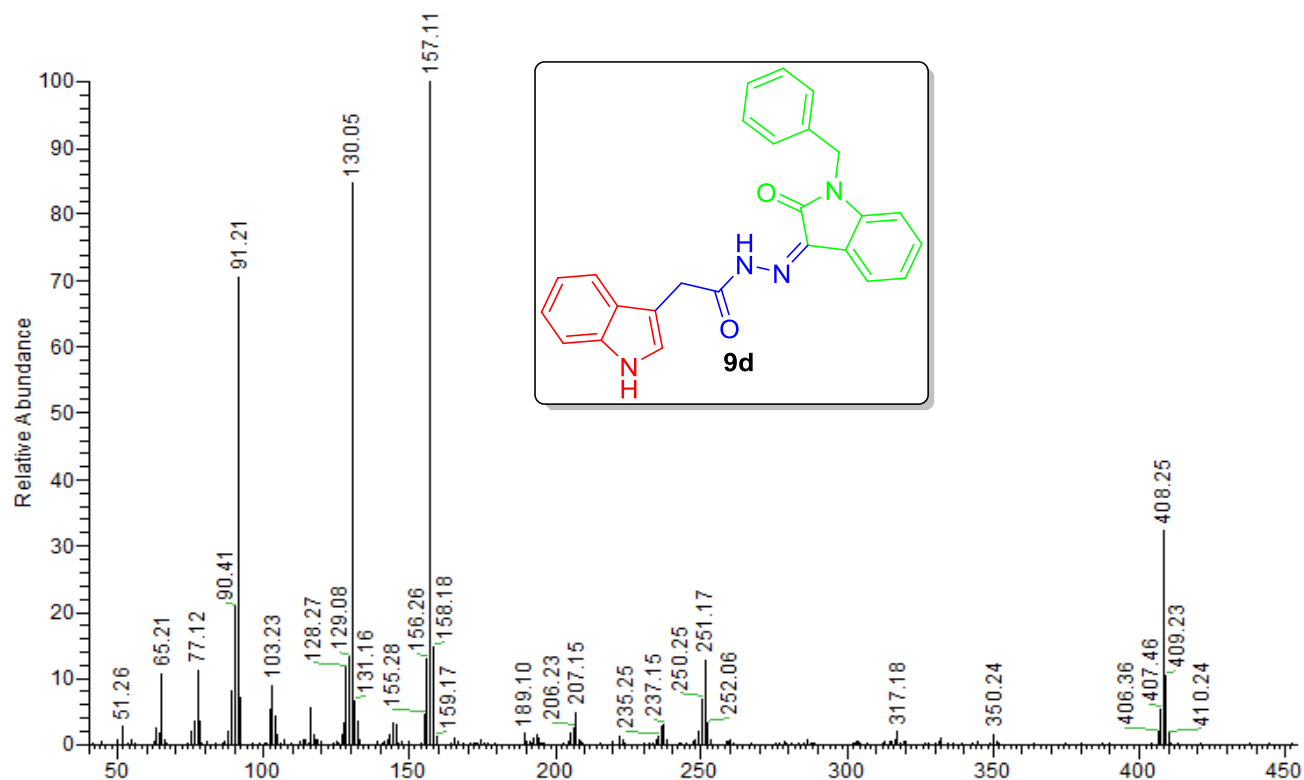
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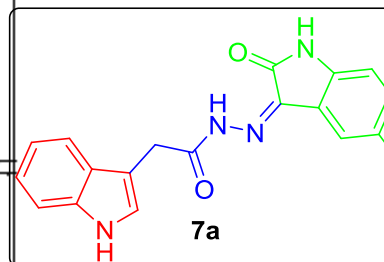


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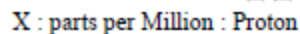


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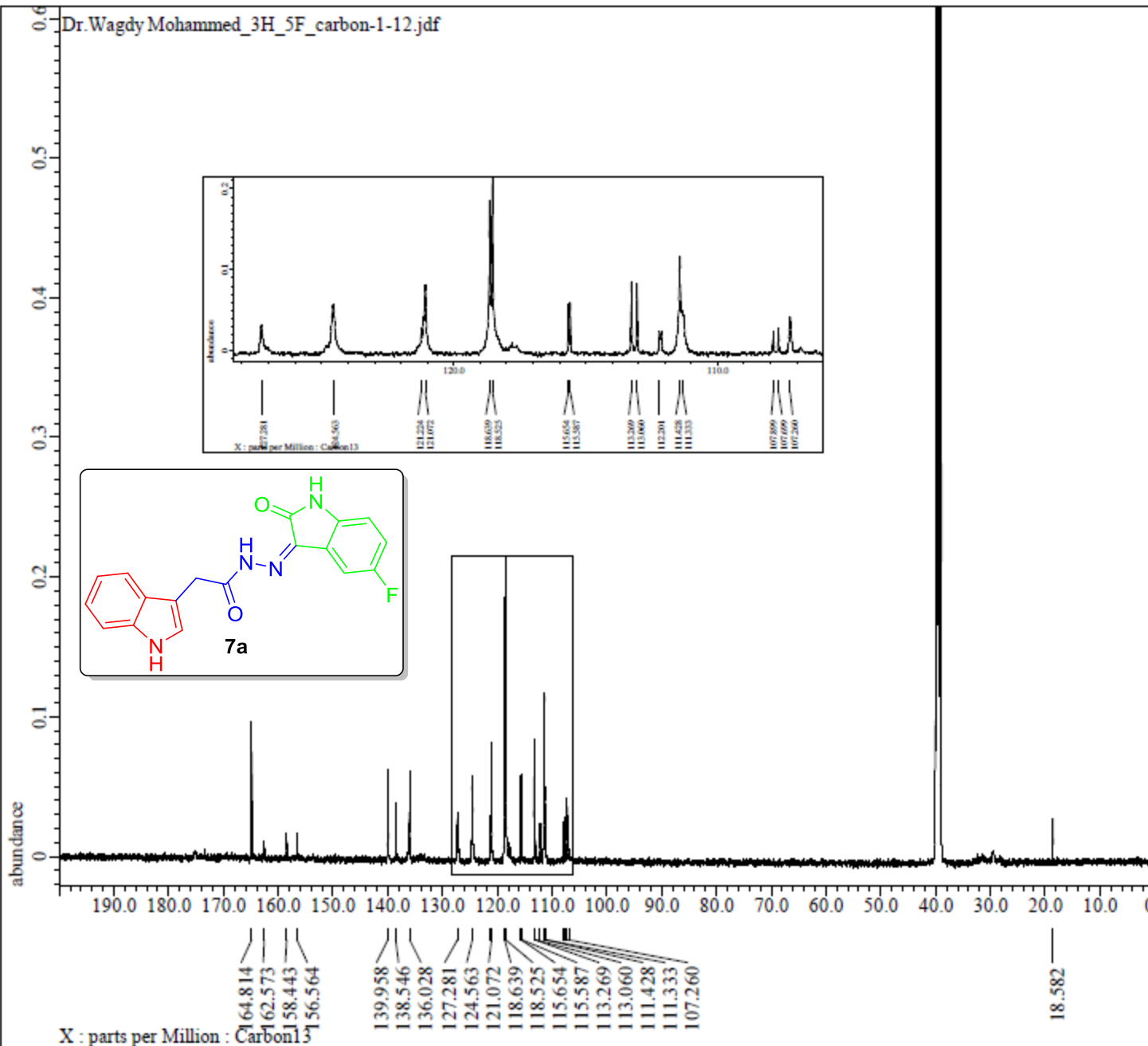
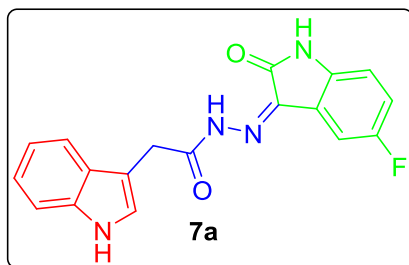
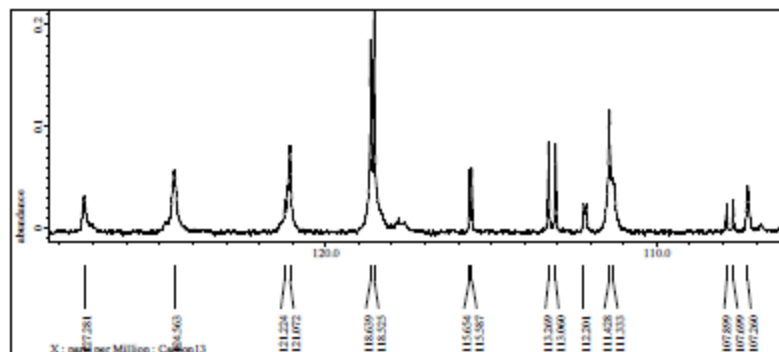


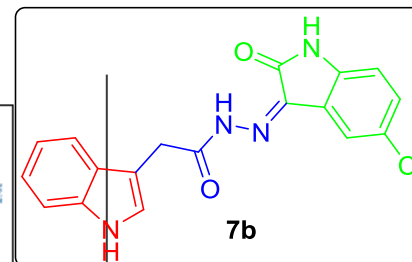
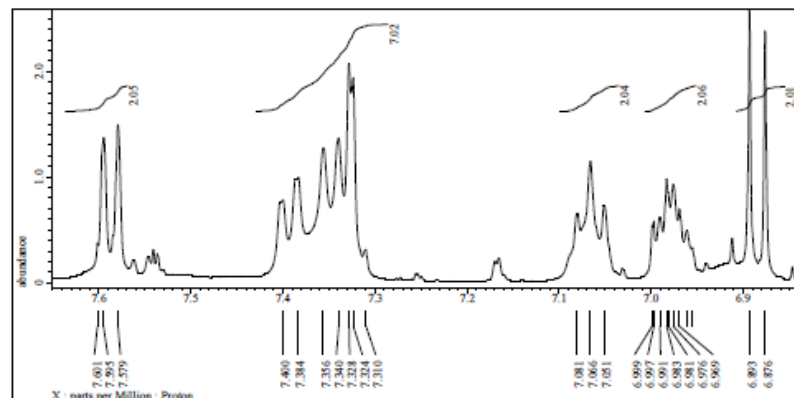
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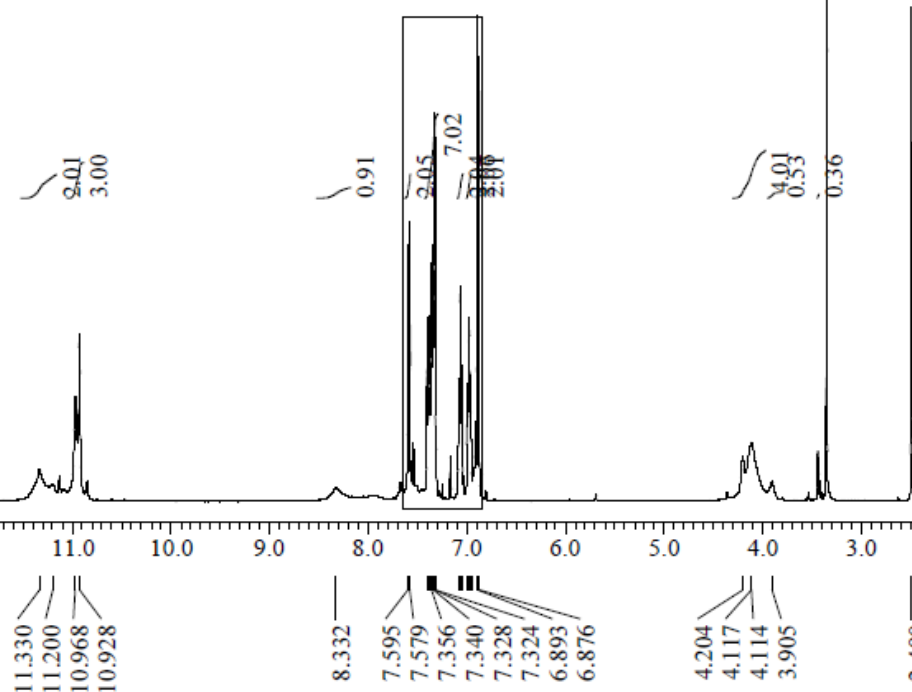
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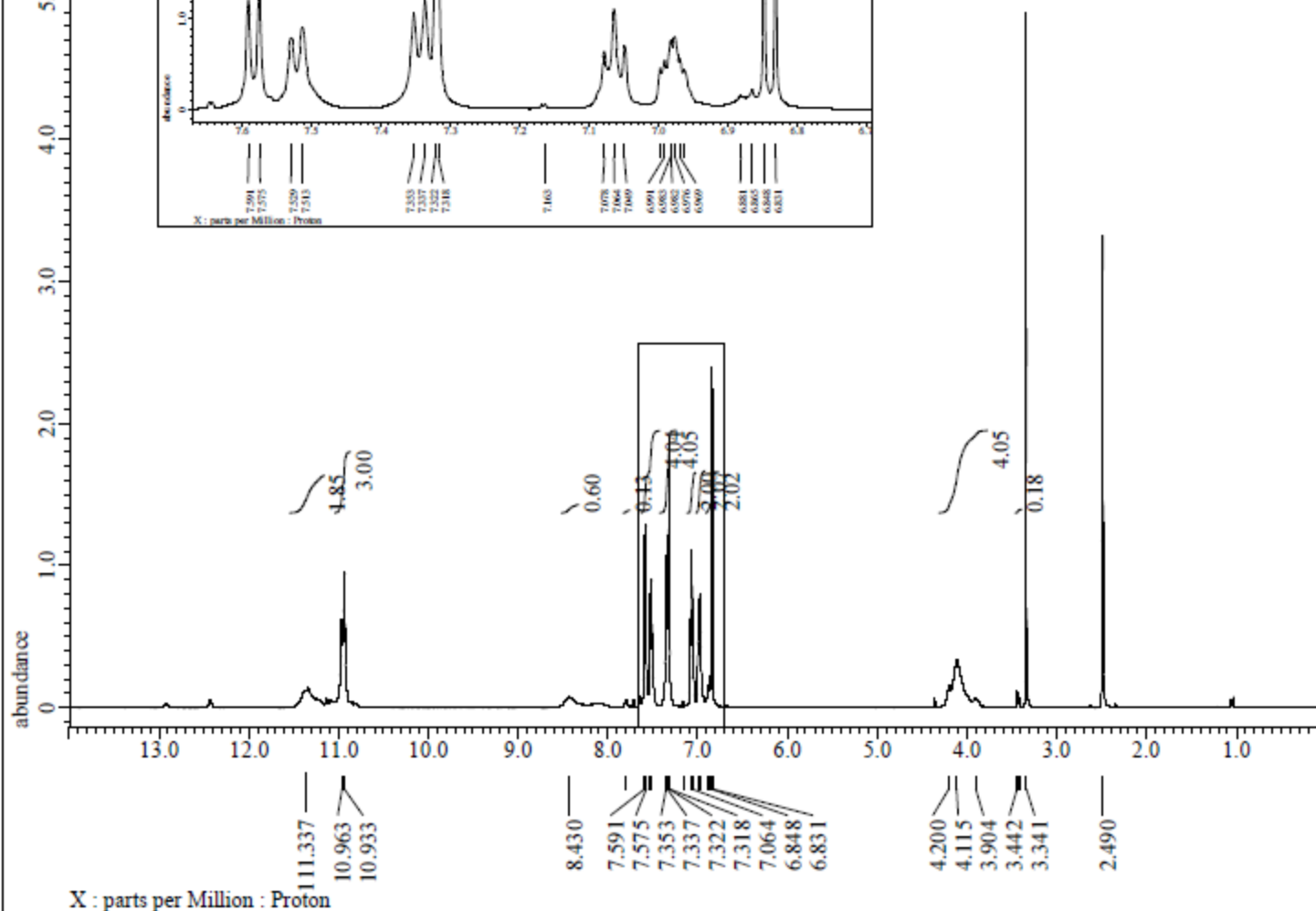
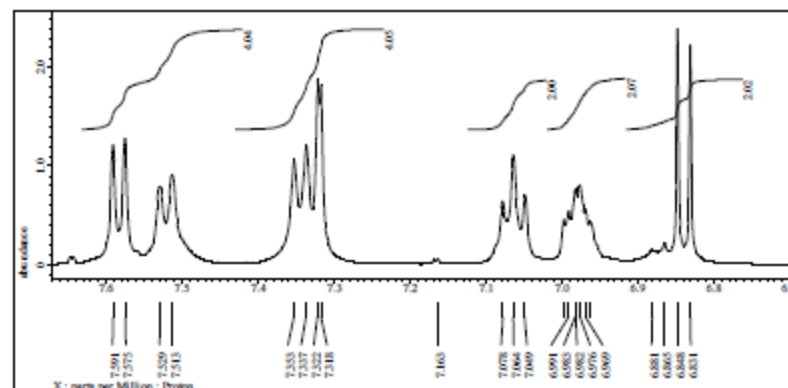
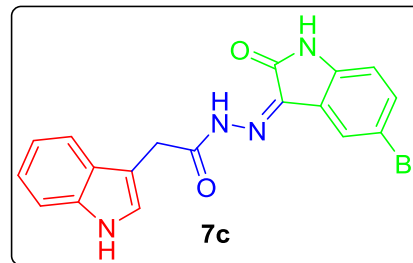




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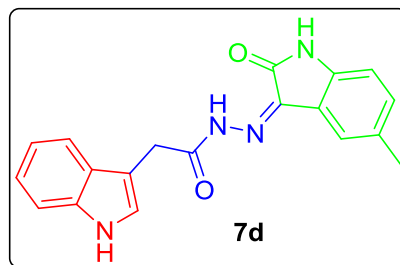


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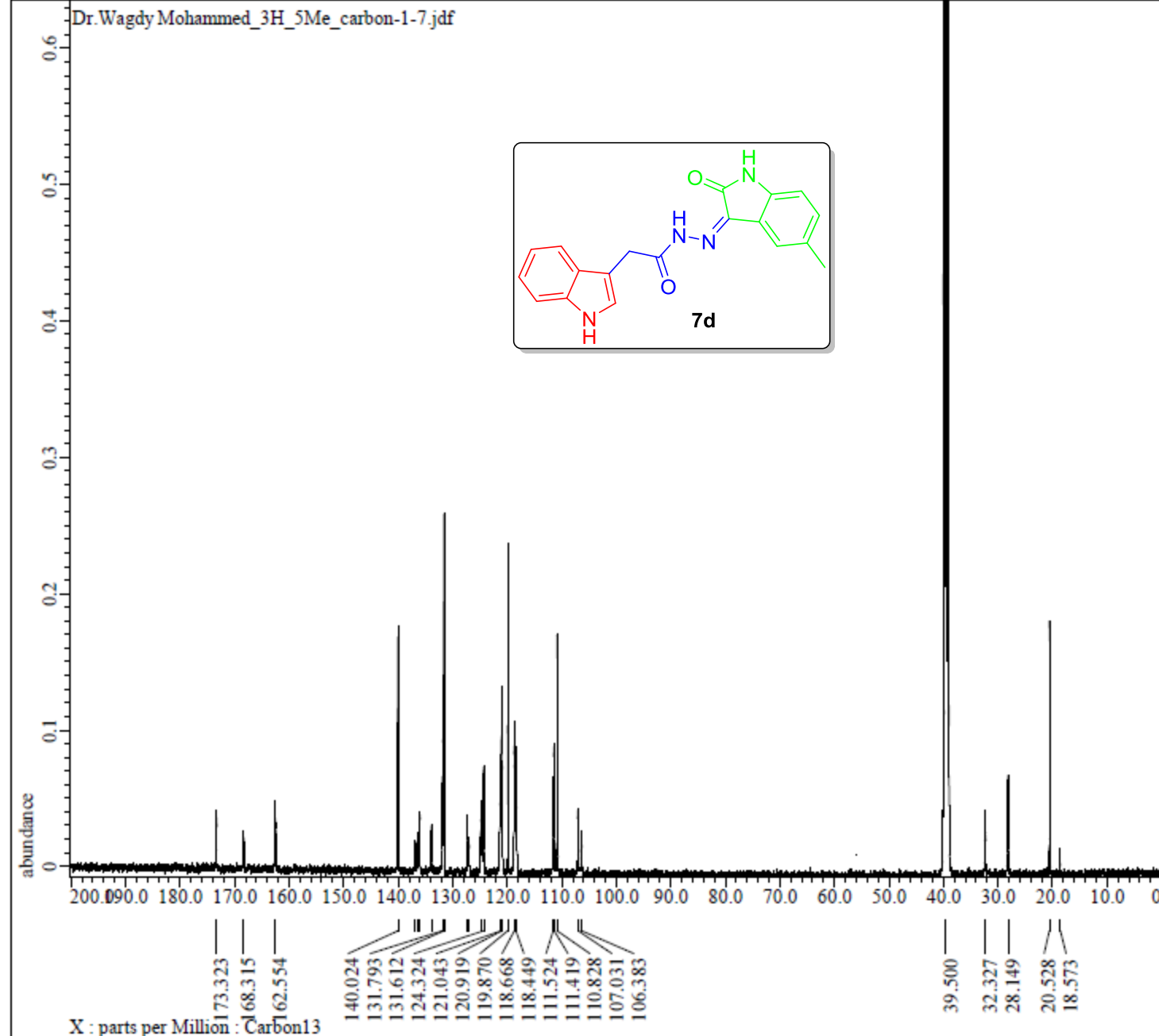


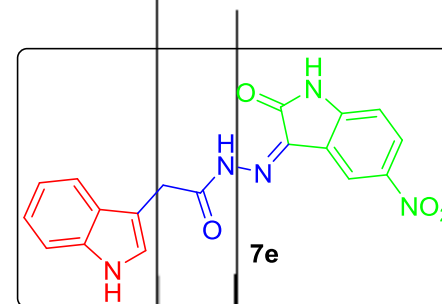
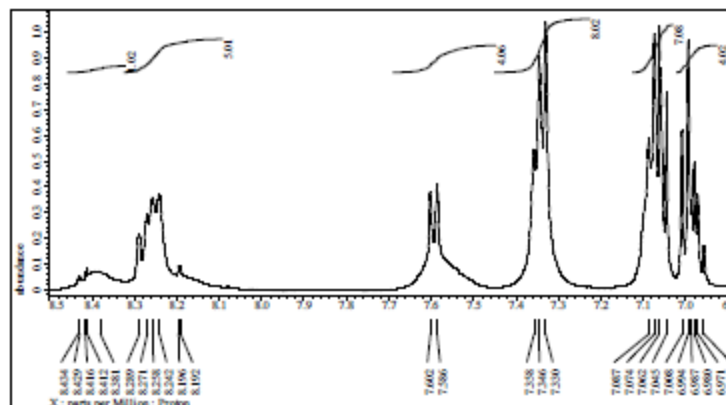
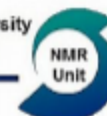
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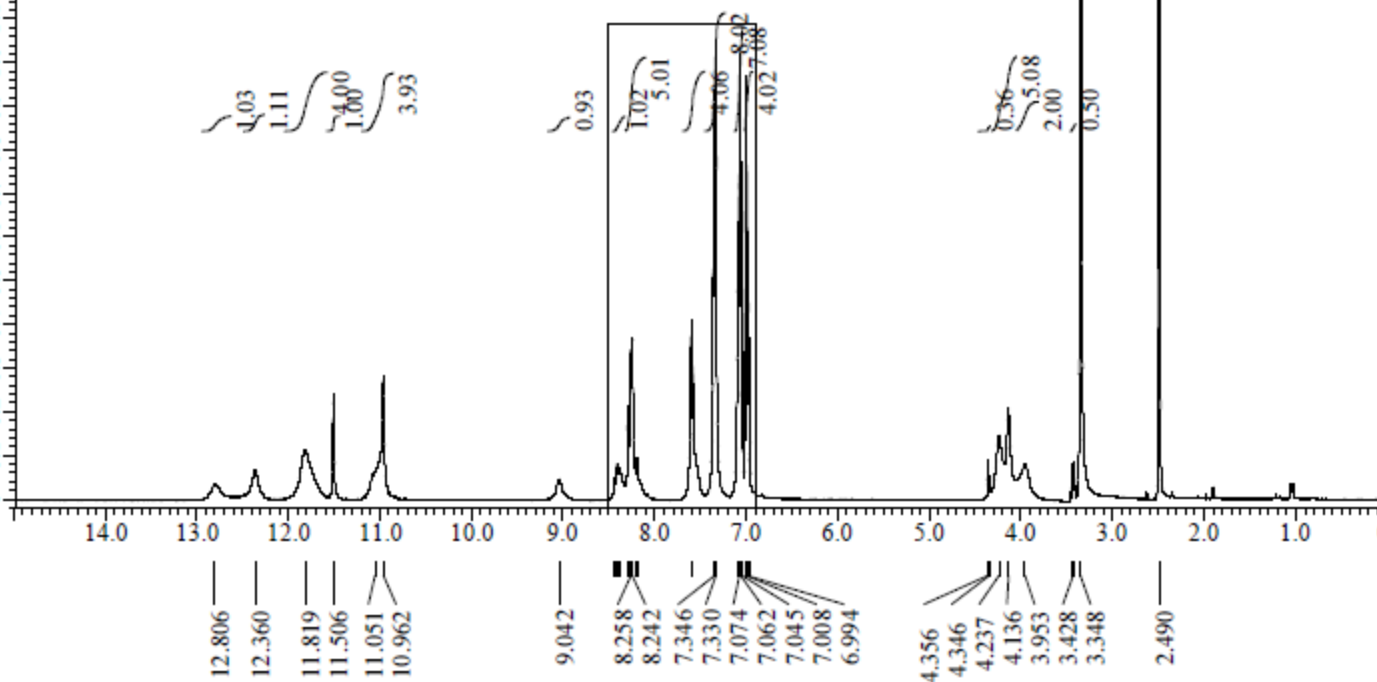
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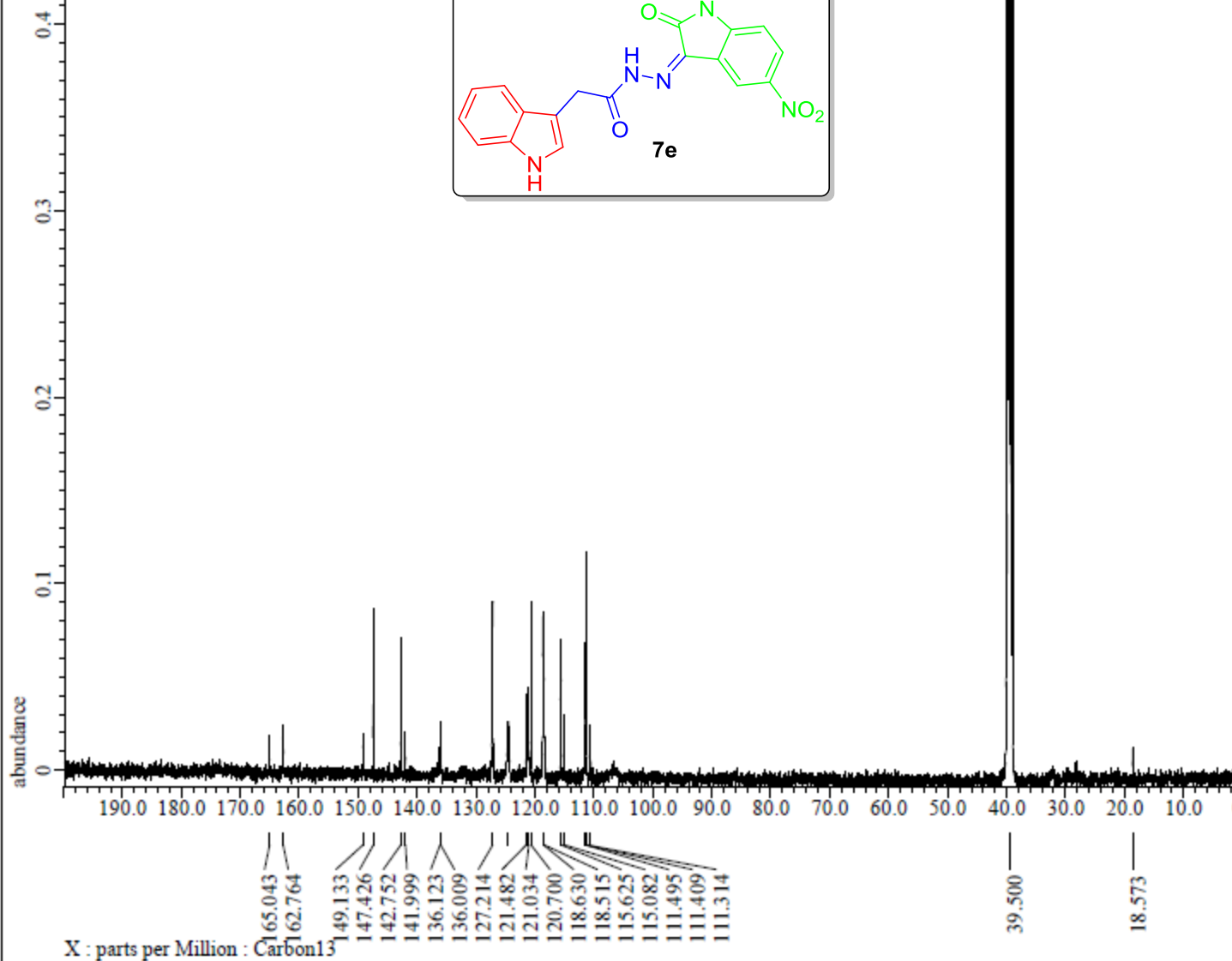
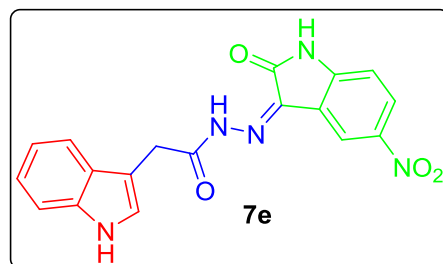
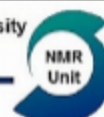
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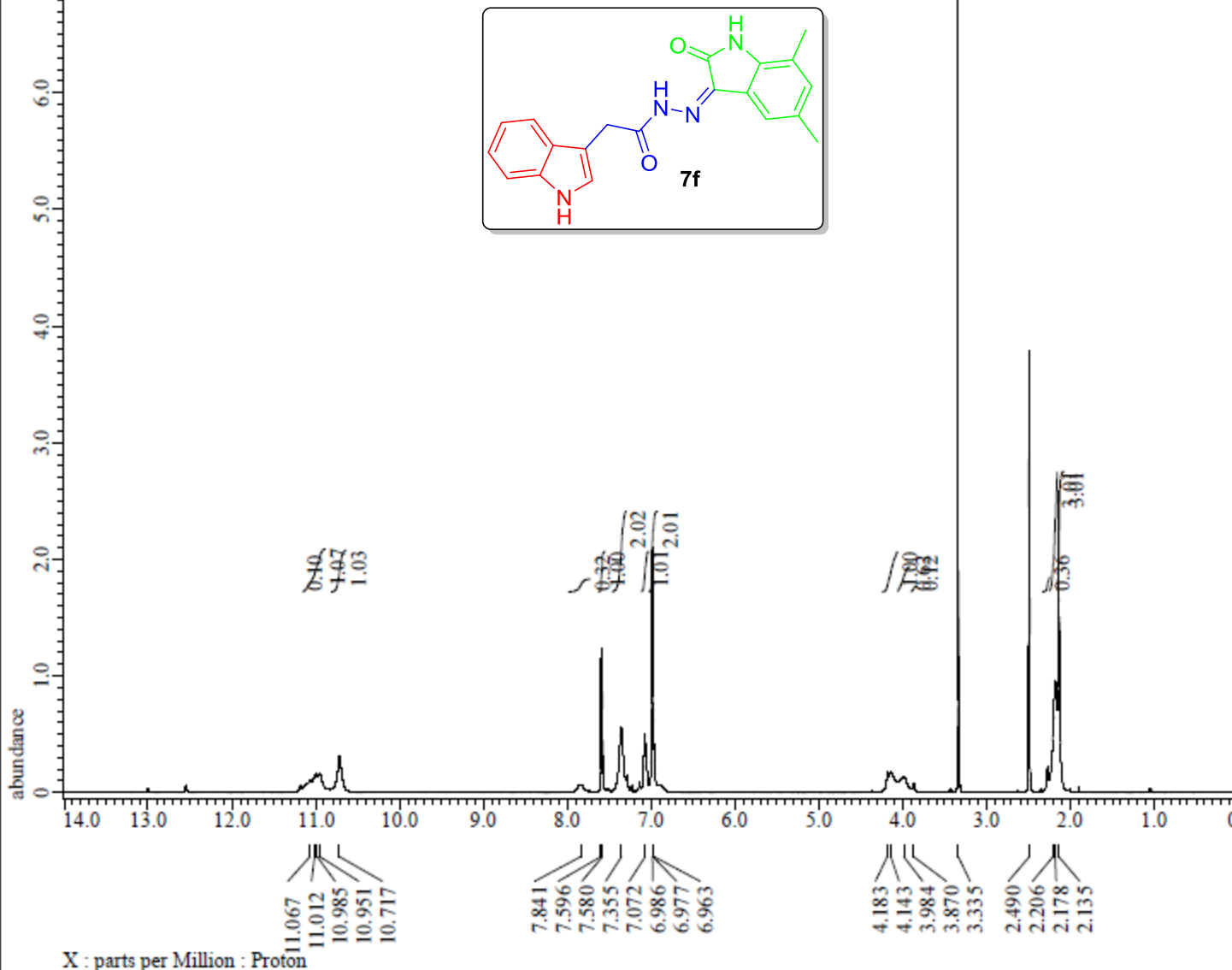
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X : parts per Million : Proton



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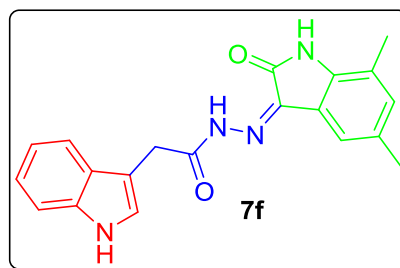
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Recvr_Gain         = 46
Temp_Get           = 19.9[dc]
X_90_Width         = 14.5[us]
X_Acq_Time         = 1.4548992[s]
X_Angle            = 45[deg]
X_Atn              = 4.9[dB]
X_Pulse            = 7.25[us]
Irr_Mode           = Off
Tri_Mode           = Off
Dante_Preset       = FALSE
Initial_Wait       = 1[s]
Repetition_Time    = 6.4548992[s]

```

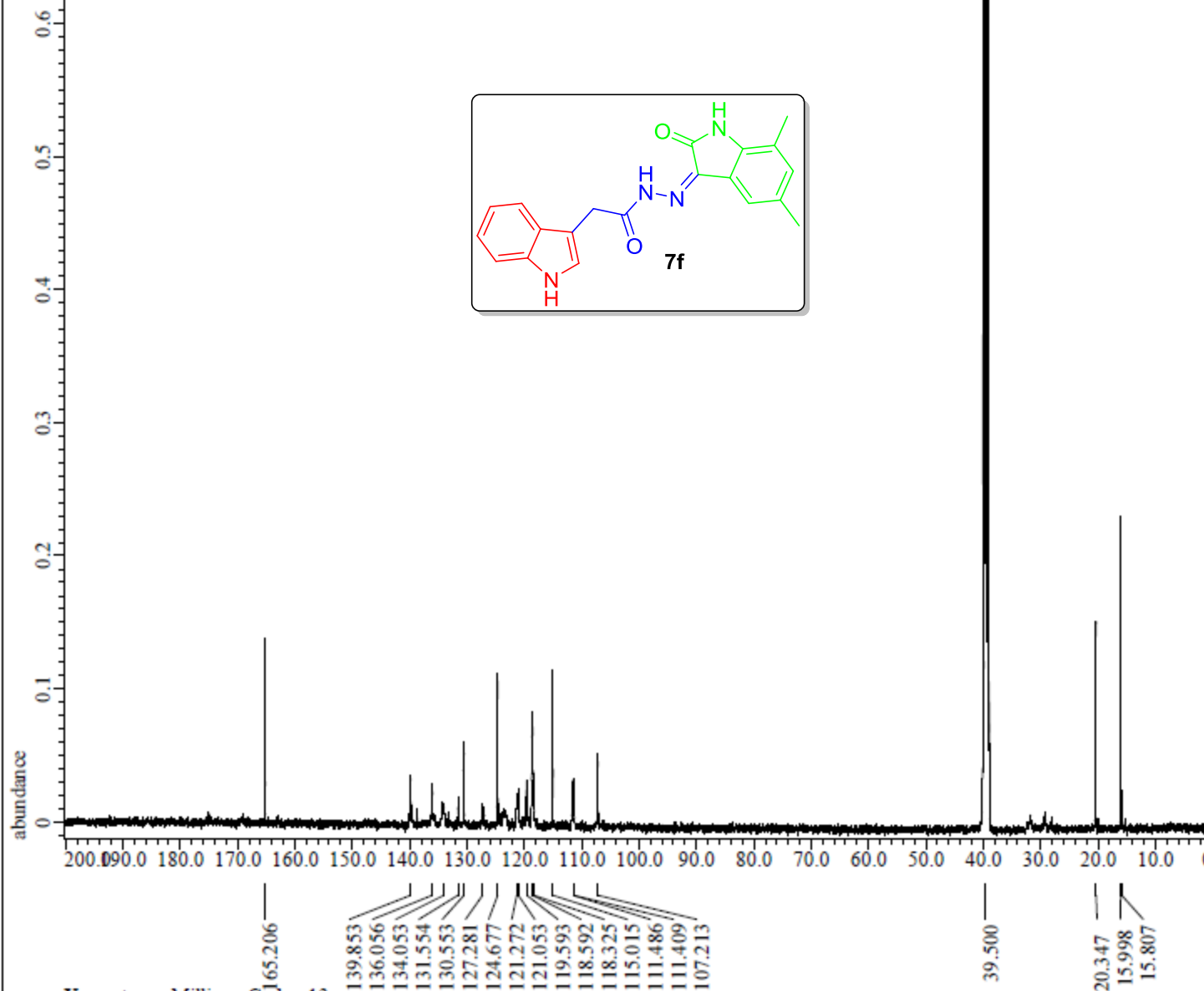


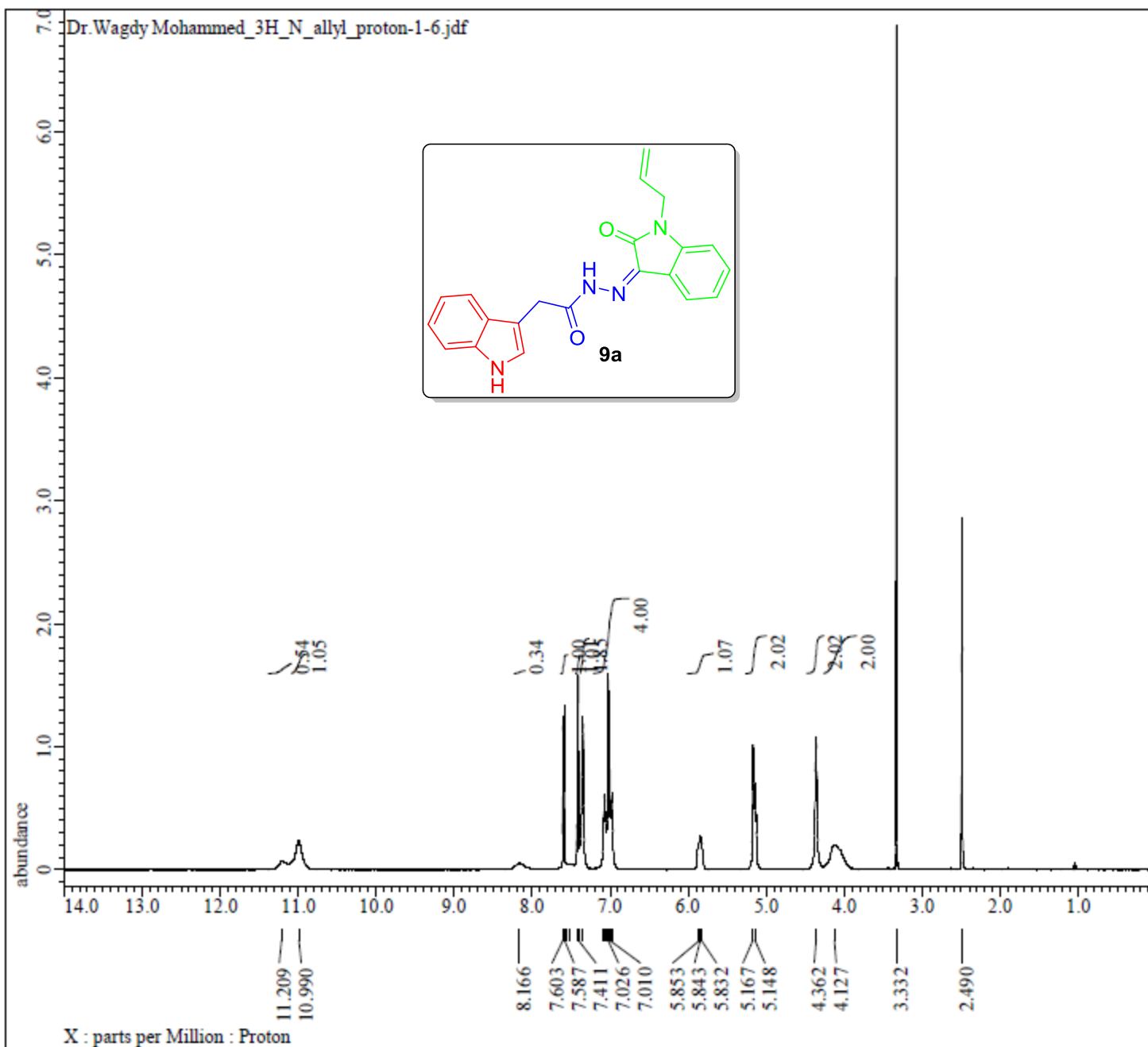
Filename = Dr.Wagdy Mohammed_3H_5,7_carbon-1-7.jdf
 Author = delta
 Experiment = carbon.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_5,7_carbon-1-7.jdf
 Solvent = DMSO-D6
 Creation_Time = 24-SEP-2019 13:28:13
 Revision_Time = 24-SEP-2019 15:44:03
 Current_Time = 24-SEP-2019 15:44:11

Comment = single pulse decoupled
 Data_Format = 1D COMPLEX
 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 = FALSE
 Scans = 3300
 Total_Scans = 3300

Relaxation_Delay = 2[s]
 Recvr_Gain = 58
 Temp_Get = 19.8[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 = WALTZ
 Irr_Pwidth = 92[us]
 Decoupling = TRUE
 Initial_Wait = 1[s]
 Noe = TRUE
 Noe_Time = 2[s]
 Repetition_Time = 2.83361792[s]





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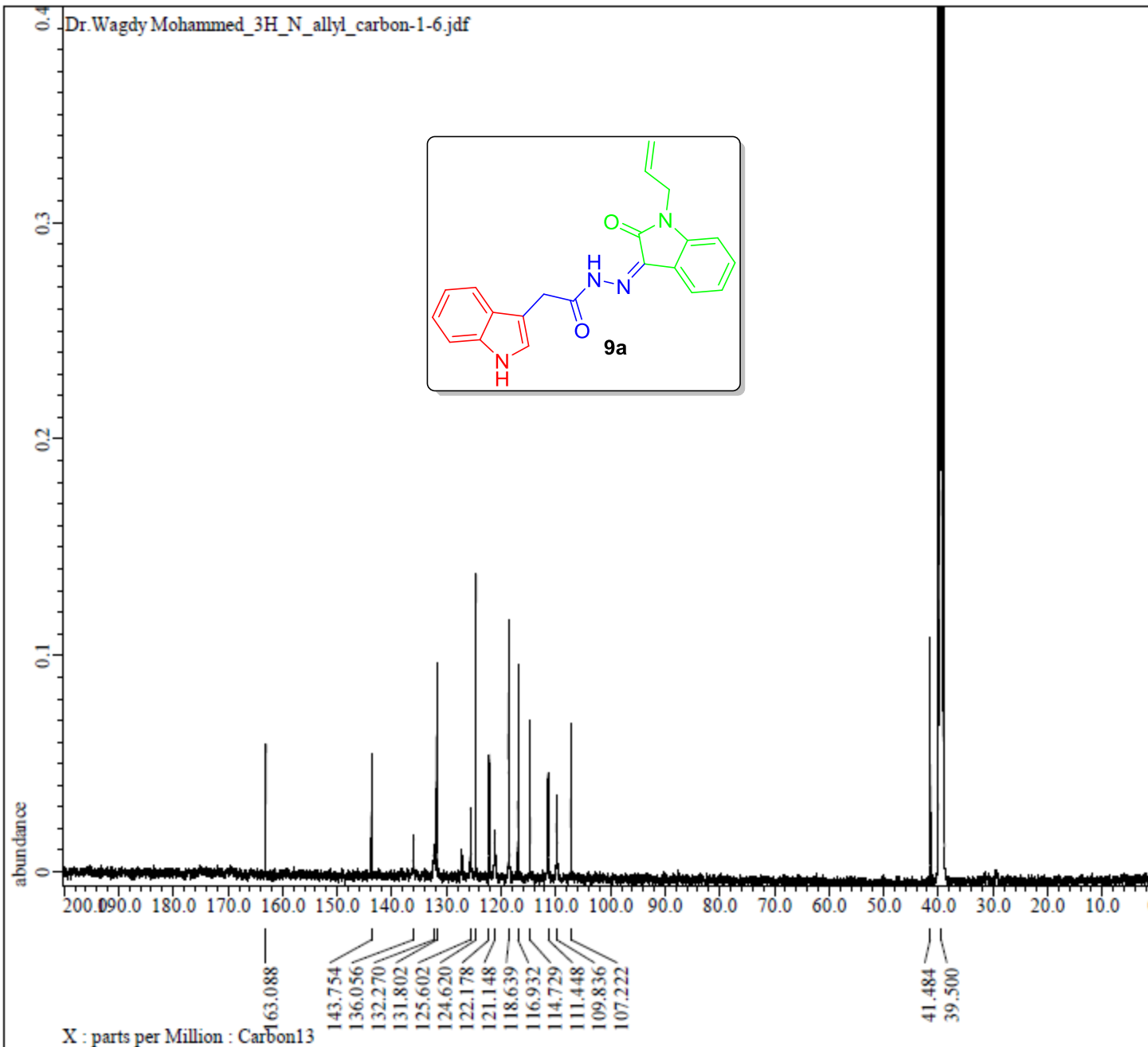
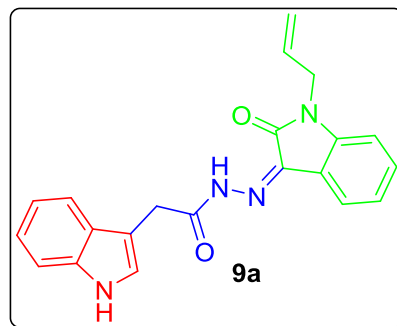


Filename = Dr.Wagdy Mohammed_3H_N_
 Author = delta
 Experiment = proton.jxp
 Sample Id = Dr.Wagdy Mohammed_3H_N_
 Solvent = DMSO-D6
 Creation Time = 23-SEP-2019 11:39:35
 Revision Time = 26-SEP-2019 12:56:00
 Current Time = 26-SEP-2019 12:56:28

Comment = single_pulse
 Data Format = 1D COMPLEX
 Dim Size = 13107
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2_MMR

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 = 16384
 X_Prescans = 1
 X_Resolution = 0.68733284[Hz]
 X_Sweep = 11.26126126[kHz]
 X_Sweep_Clipped = 9.00900901[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]
 Recvr_Gain = 44
 Temp_Get = 21.1[dC]
 X_90_Width = 14.5[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
 Dante_Presat = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 6.4548992[s]



Filename = Dr.Wagdy Mohammed_3H_N_...
 Author = delta
 Experiment = carbon.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_N_...
 Solvent = DMSO-D6
 Creation_Time = 24-SEP-2019 00:07:16
 Revision_Time = 24-SEP-2019 12:47:45
 Current_Time = 24-SEP-2019 12:48:40
 Comment = single pulse decoupled
 Data_Format = 1D COMPLEX
 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 = FALSE
 Scans = 3700
 Total_Scans = 3700
 Relaxation_Delay = 2 [s]
 Recvr_Gain = 56
 Temp_Get = 18.6 [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_Hoe = 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]

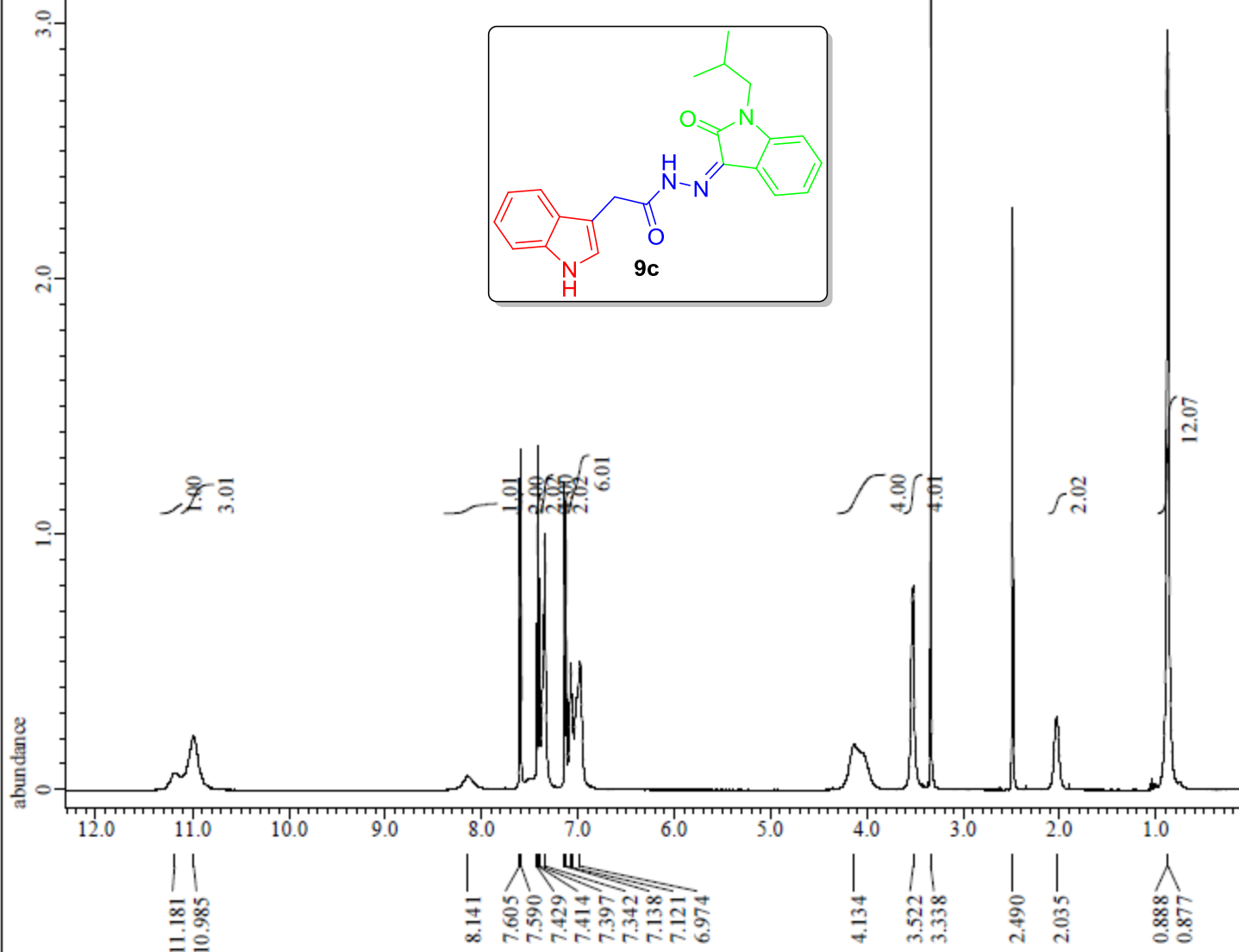
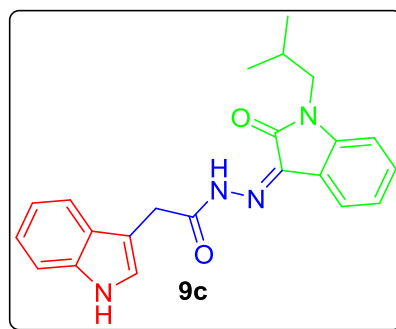


Filename = Dr.Wagdy Mohammed_3H_iso
 Author = delta
 Experiment = proton.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_iso
 Solvent = DMSO-D6
 Creation_Time = 23-SEP-2019 13:21:15
 Revision_Time = 24-SEP-2019 15:51:53
 Current_Time = 24-SEP-2019 15:52:30

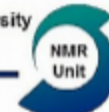
Comment = single_pulse
 Data_Format = 1D_COMPLEX
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2_NMR

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 = 16384
 X_Prescans = 1
 X_Resolution = 0.68733284[Hz]
 X_Sweep = 11.26126126[kHz]
 X_Sweep_Clipped = 9.00900901[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]
 Recvr_Gain = 42
 Temp_Get = 20.3[deg]
 X_90_Width = 14.5[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
 Dante_Presat = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 6.4548992[s]



X : parts per Million : Proton

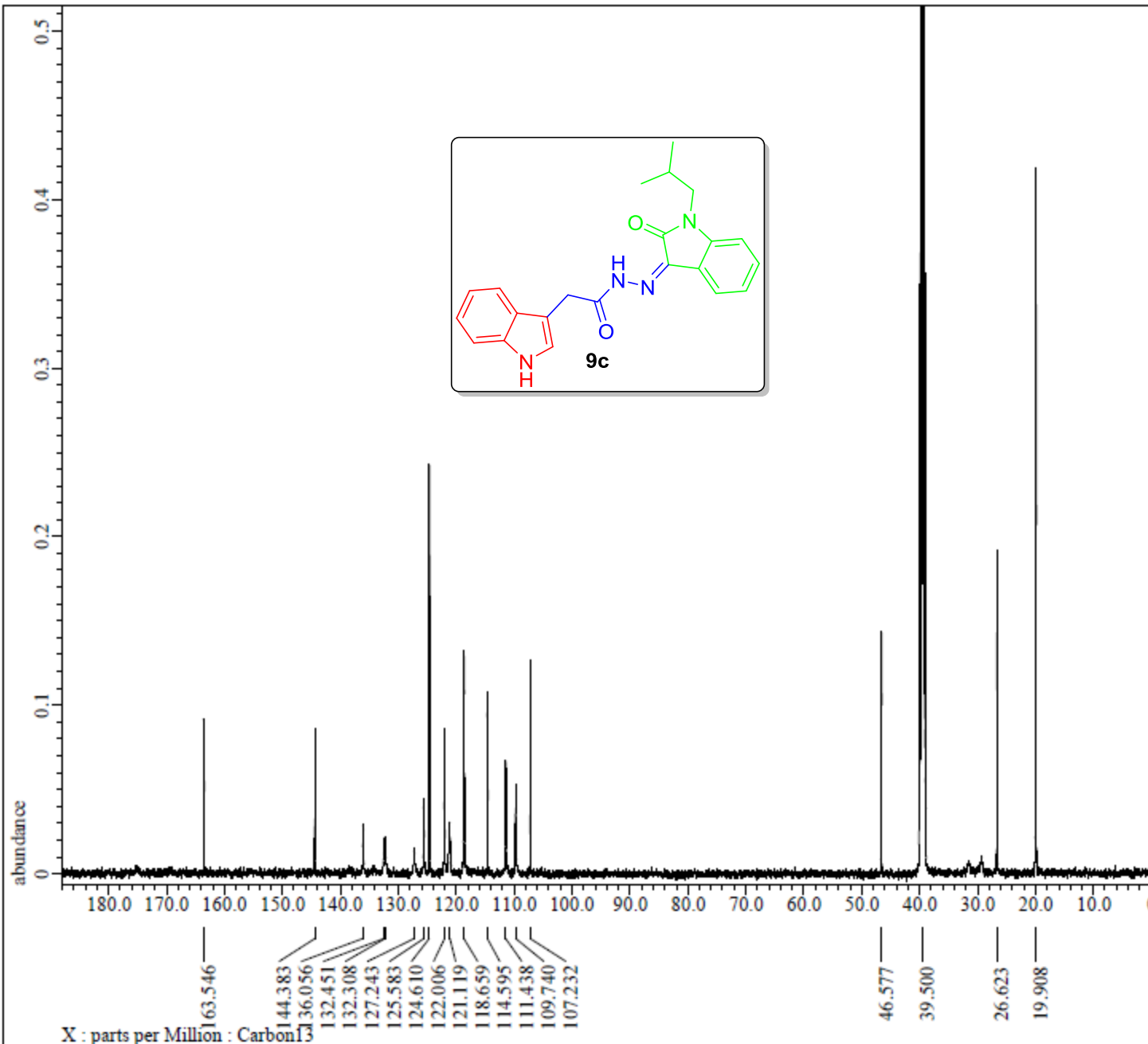
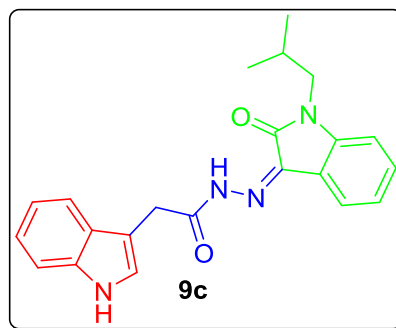


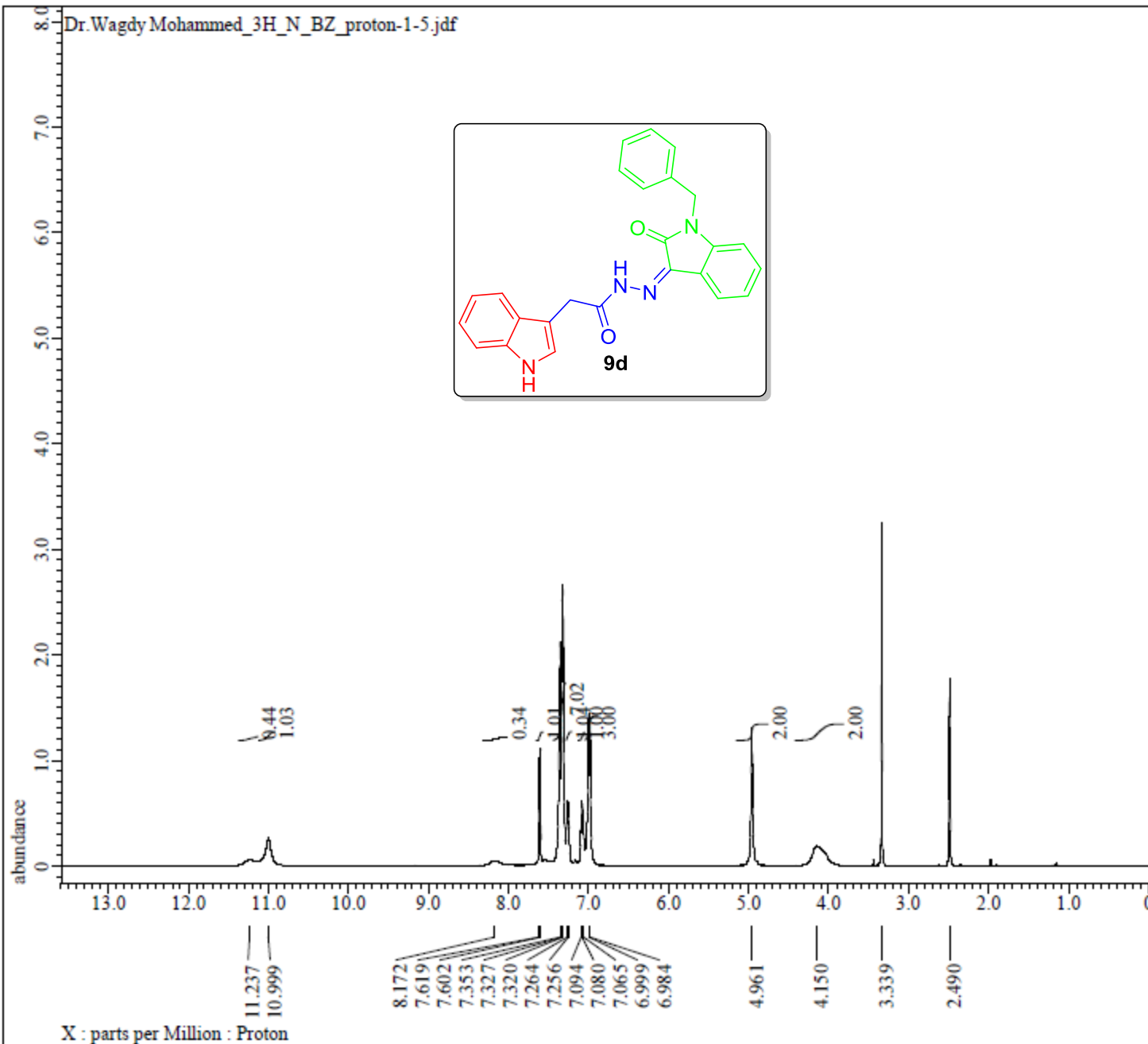
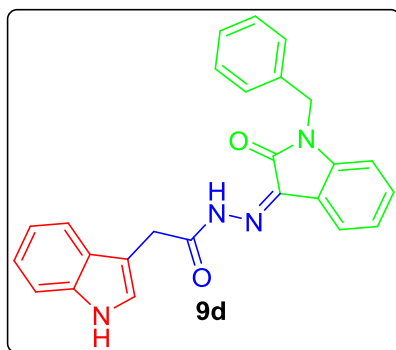
Filename = Dr.Wagdy Mohammed_3H_isc
Author = delta
Experiment = carbon.jxp
Sample_Id = Dr.Wagdy Mohammed_3H_isc
Solvent = DMSO-D6
Creation_Time = 25-SEP-2019 02:48:45
Revision_Time = 25-SEP-2019 09:25:27
Current_Time = 25-SEP-2019 09:28:42

Comment = single pulse decoupled g
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 = FALSE
Scans = 3700
Total_Scans = 3700

Relaxation_Delay = 2[s]
Recvr_Gain = 56
Temp_Get = 18.8[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_Noce = 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]



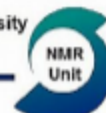


Filename = Dr.Wagdy Mohammed_3H_N
 Author = delta
 Experiment = proton.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_N
 Solvent = DMSO-D6
 Creation_Time = 23-SEP-2019 11:28:22
 Revision_Time = 25-SEP-2019 10:35:45
 Current_Time = 25-SEP-2019 10:35:51

Comment = single_pulse
 Data_Format = 1D_COMPLEX
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2_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 = 16384
 X_Prescans = 1
 X_Resolution = 0.68733284[Hz]
 X_Sweep = 11.26126126[kHz]
 X_Sweep_Clipped = 9.00900901[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]
 Recvr_Gain = 42
 Temp_Get = 21.5[dC]
 X_90_Width = 14.5[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
 Dante_Presat = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 6.4548992[s]

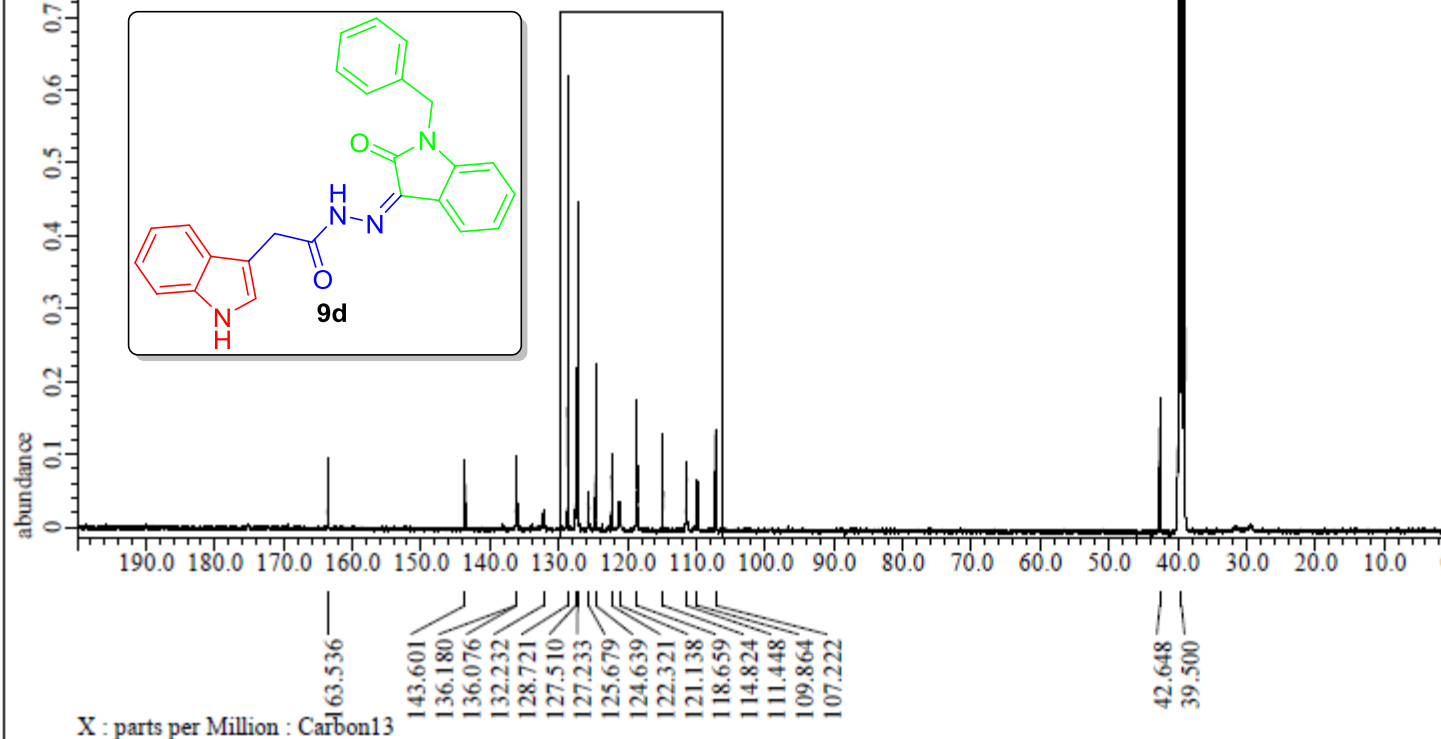
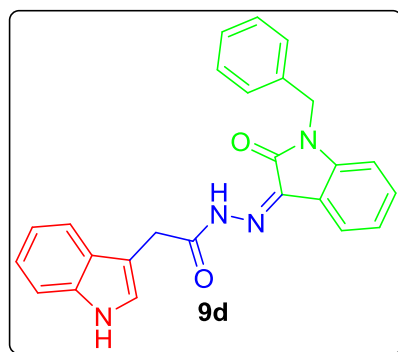
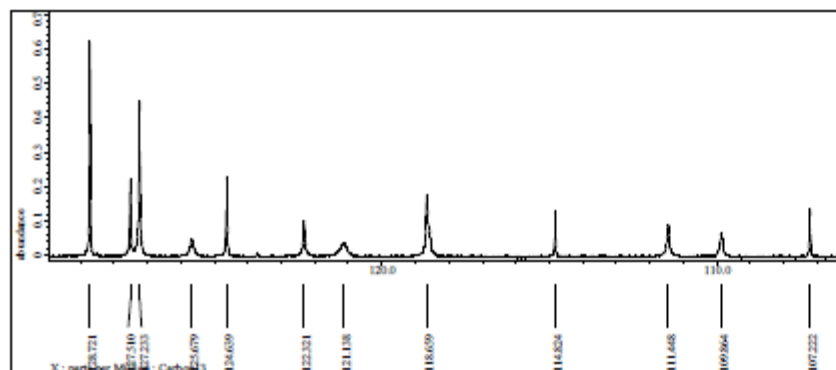


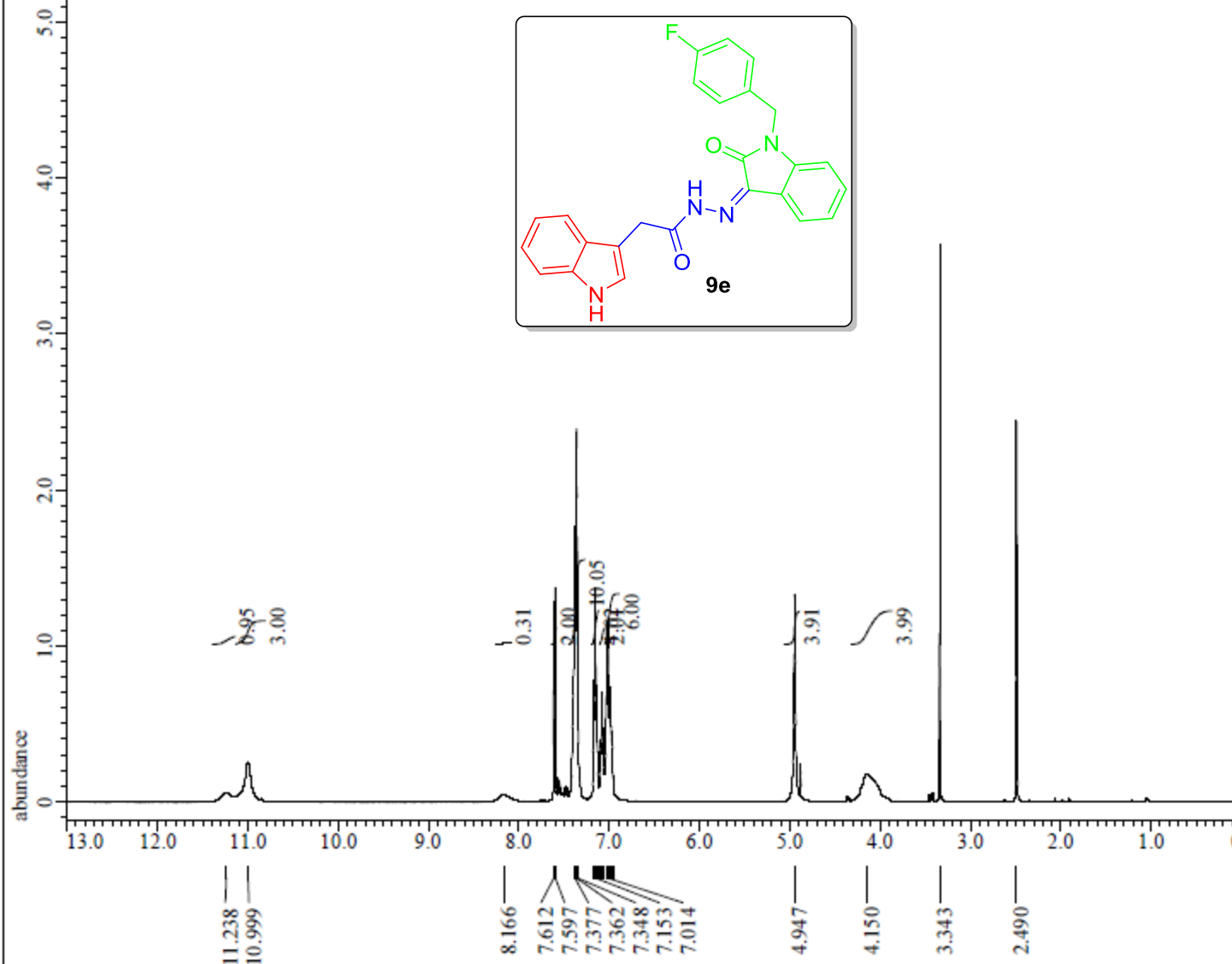
Filename = Dr.Wagdy Mohammed_3H_N_BZ_carbon-1-6.jdf
 Author = delta
 Experiment = carbon.jxp
 Sample Id = Dr.Wagdy Mohammed_3H_N_BZ_carbon-1-6.jdf
 Solvent = DMSO-D6
 Creation Time = 23-SEP-2019 21:03:23
 Revision Time = 24-SEP-2019 12:45:31
 Current Time = 24-SEP-2019 12:46:17

Comment = single pulse decoupled
 Data Format = 1D COMPLEX
 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 = FALSE
 Scans = 3700
 Total_Scans = 3700

Relaxation_Delay = 2 [s]
 Recvr_Gain = 58
 Temp_Get = 18.6 [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]





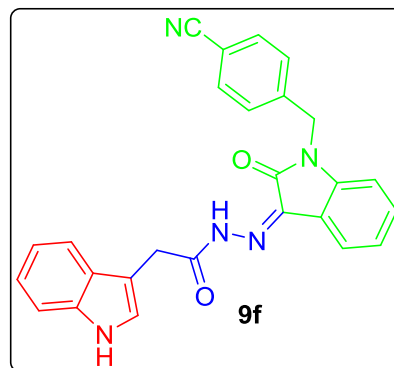
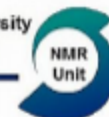
X : parts per Million : Proton

Filename = Dr. Wagdy Mohamd_3H-4FBz
 Author = delta
 Experiment = proton.jxp
 Sample_Id = Dr. Wagdy Mohamd_3H-4FBz
 Solvent = DMSO-D6
 Creation_Time = 17-SEP-2019 14:55:04
 Revision_Time = 17-SEP-2019 14:37:29
 Current_Time = 17-SEP-2019 14:37:55

Comment = single_pulse
 Data_Format = 1D_COMPLEX
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2_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 = 16384
 X_Prescans = 1
 X_Resolution = 0.68733284[Hz]
 X_Sweep = 11.26126126[kHz]
 X_Sweep_Clipped = 9.00900901[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]
 Recvr_Gain = 42
 Temp_Get = 20[dC]
 X_90_Width = 14.5[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
 Dante_Preset = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 6.4548992[s]

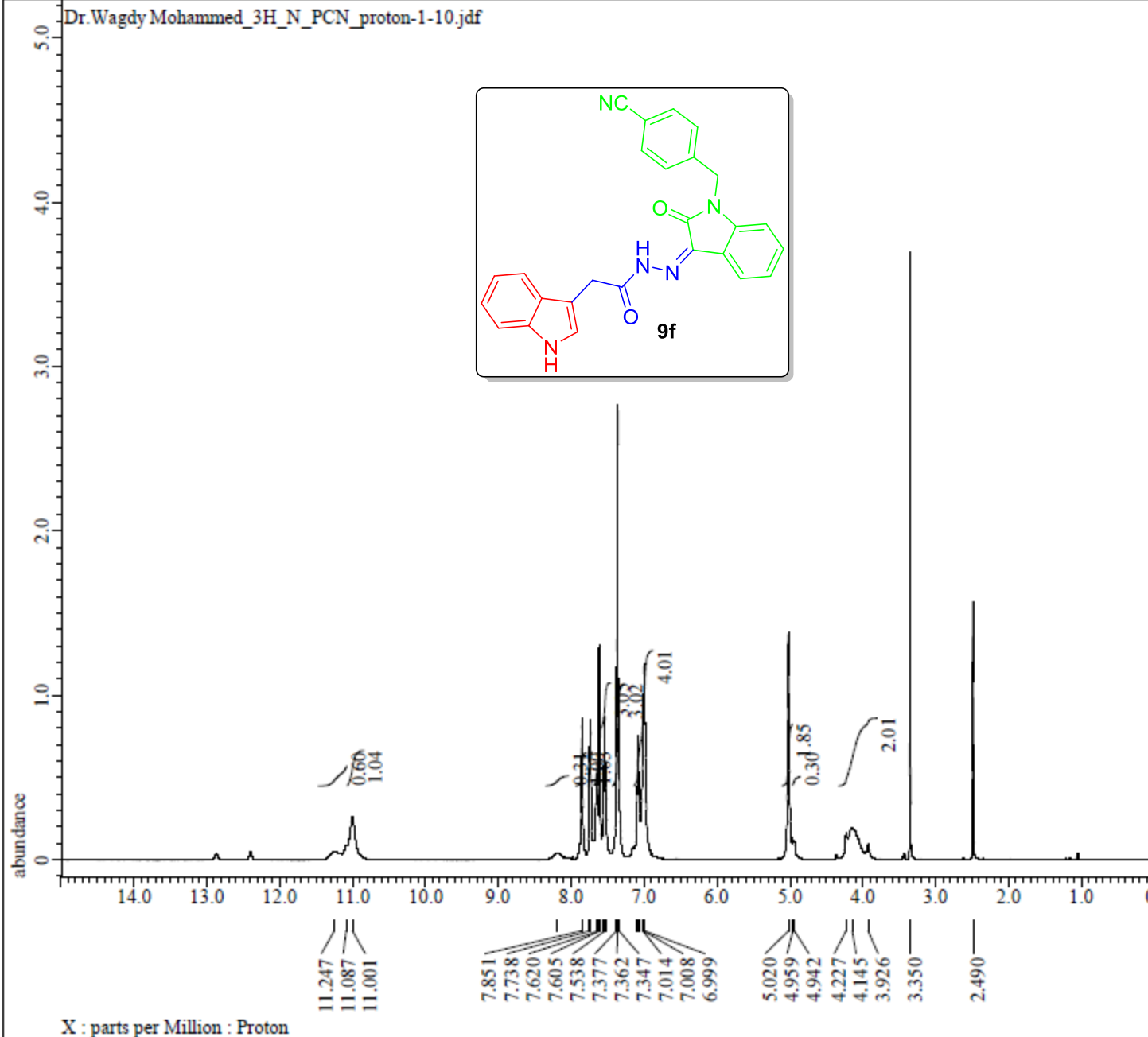


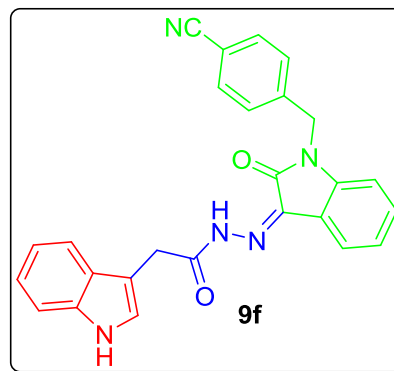
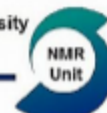
Filename = Dr.Wagdy Mohammed_3H_N_
 Author = delta
 Experiment = proton.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_N_
 Solvent = DMSO-D6
 Creation_Time = 23-SEP-2019 12:02:28
 Revision_Time = 26-SEP-2019 13:50:57
 Current_Time = 26-SEP-2019 13:51:02

Comment = single pulse
 Data_Format = 1D COMPLEX
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2_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 = 16384
 X_Prescans = 1
 X_Resolution = 0.68733284 [Hz]
 X_Sweep = 11.26126126 [kHz]
 X_Sweep_Clippped = 9.00900901 [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]
 Recvr_Gain = 40
 Temp_Get = 21 [dC]
 X_90_Width = 14.5 [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
 Dante_Presat = FALSE
 Initial_Wait = 1 [s]
 Repetition_Time = 6.4548992 [s]





Filename = Dr.Wagdy Mohammed_3H_N_1
 Author = delta
 Experiment = carbon.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_N_1
 Solvent = DMSO-D6
 Creation_Time = 24-SEP-2019 06:14:19
 Revision_Time = 24-SEP-2019 12:59:14
 Current_Time = 24-SEP-2019 13:00:02

Comment = single pulse decoupled
 Data Format = 1D COMPLEX
 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_Clippped = 31.44654088[kHz]
 Irr_Domain = Proton
 Irr_Freq = 500.15991521[MHz]
 Irr_Offset = 5.0[ppm]
 Clipped = TRUE
 Scans = 3700
 Total_Scans = 3700

Relaxation_Delay = 2[s]
 Recvr_Gain = 58
 Temp_Get = 19.7[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]
 Irr_Atn_Dec = 20.948[dB]
 Irr_Atn_No = 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]

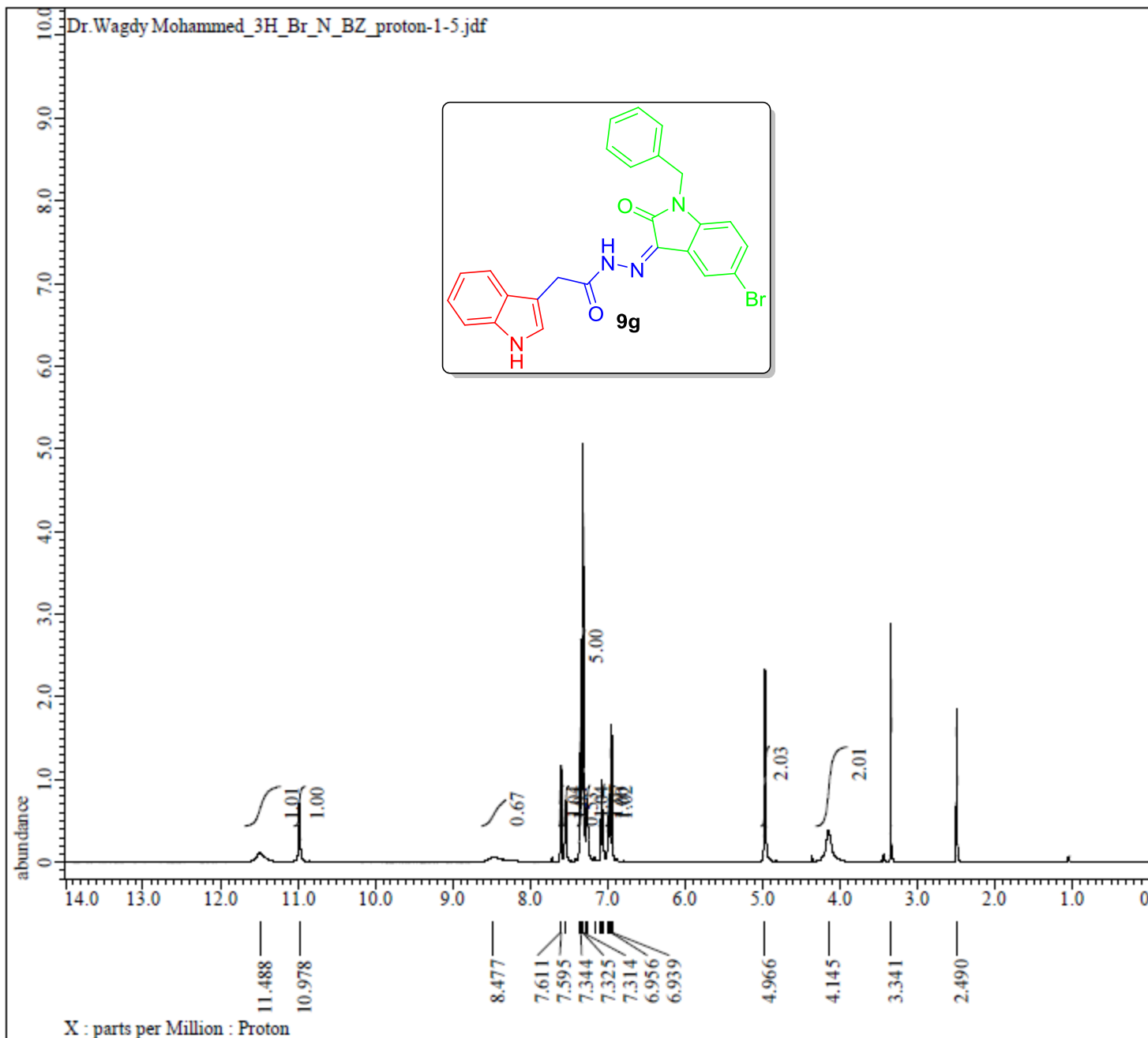
abundance

200.0 190.0 180.0 170.0 160.0 150.0 140.0 130.0 120.0 110.0 100.0 90.0 80.0 70.0 60.0 50.0 40.0 30.0 20.0 10.0 0

163.670
 160.789
 143.315
 142.323
 137.955
 132.098
 131.402
 130.906
 129.981
 129.895
 124.639
 123.275
 122.455
 118.630
 115.015
 111.619
 111.457
 109.683
 107.213

42.018
 41.761
 39.500

X : parts per Million : Carbon13



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ECA- 500 II

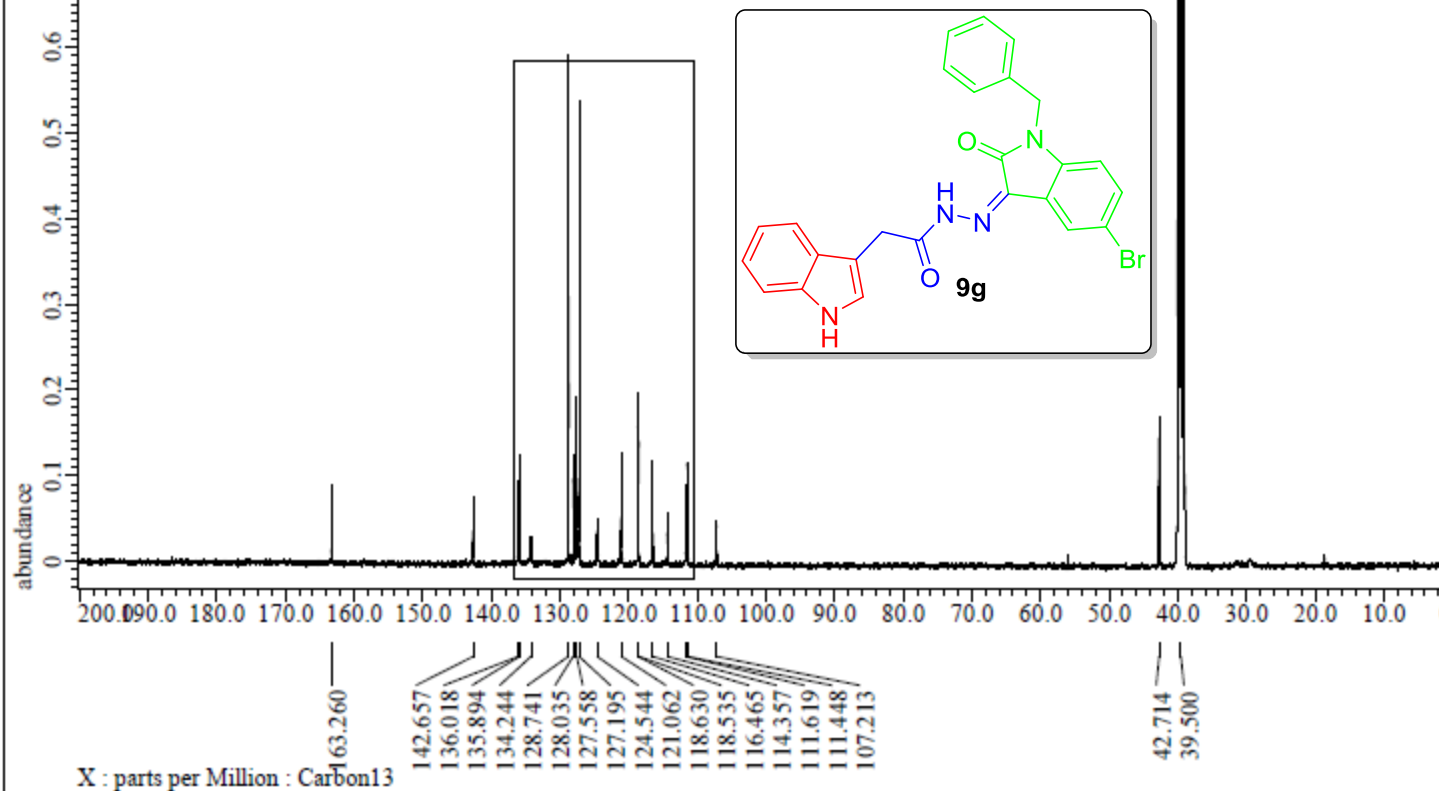
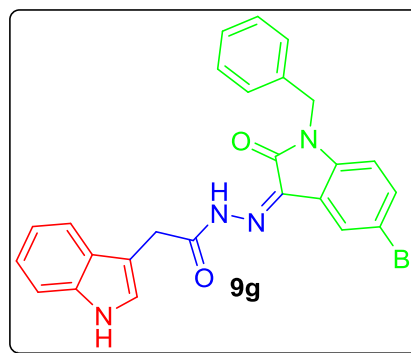
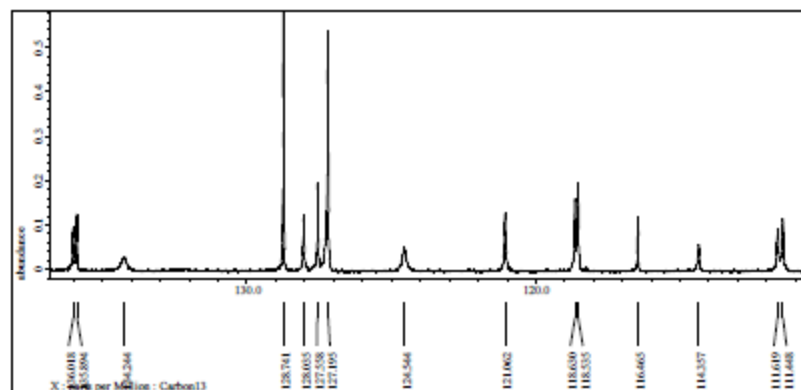


Filename = Dr.Wagdy Mohammed_3H_Br_N_BZ_proton-1-5.jdf
 Author = delta
 Experiment = proton.jxp
 Sample_Id = Dr.Wagdy Mohammed_3H_Br_N_BZ_proton-1-5.jdf
 Solvent = DMSO-D6
 Creation_Time = 23-SEP-2019 11:51:01
 Revision_Time = 17-OCT-2019 14:15:57
 Current_Time = 17-OCT-2019 14:16:04

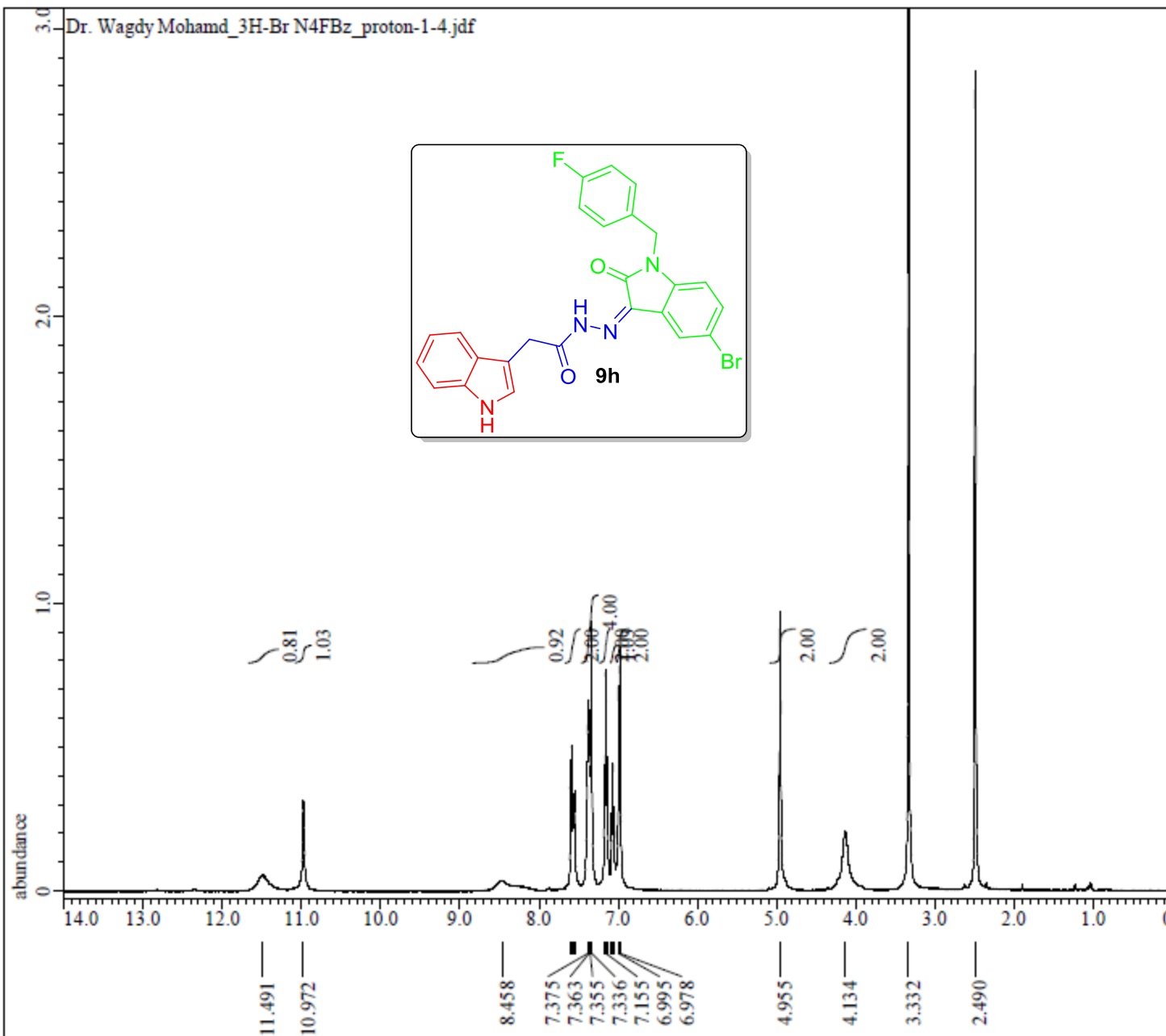
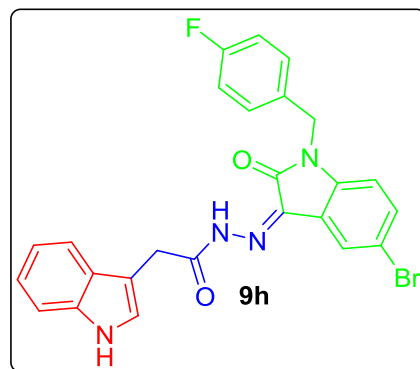
Comment = single_pulse
 Data_Format = 1D_COMPLEX
 Dim_Size = 26214
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2_NMR

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 = 16384
 X_Prescans = 1
 X_Resolution = 0.68733284[Hz]
 X_Sweep = 11.26126126[kHz]
 X_Sweep_Clipped = 9.00900901[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]
 Recvr_Gain = 42
 Temp_Get = 21[dC]
 X_90_Width = 14.5[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
 Dante_Presat = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 6.4548992[s]



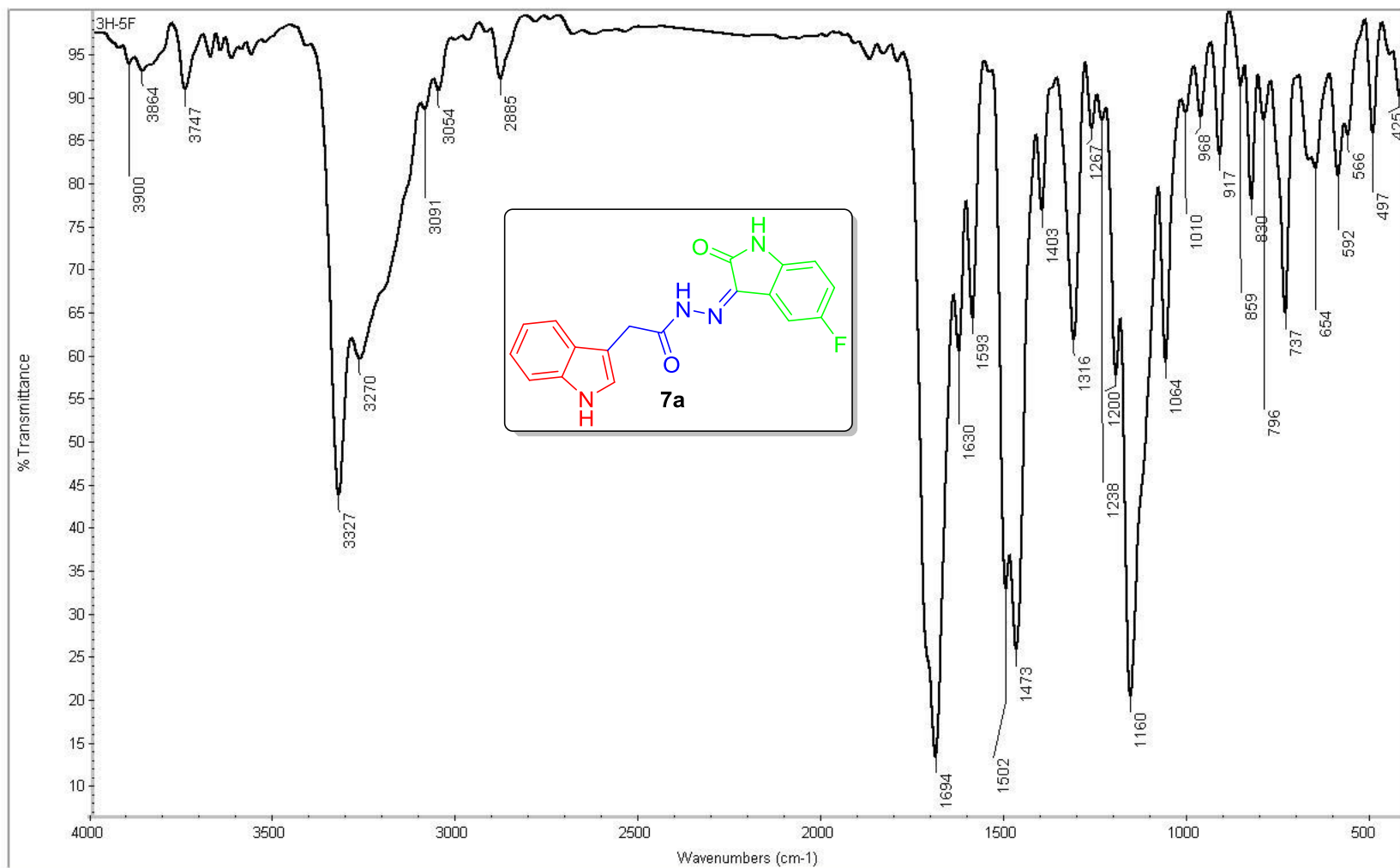
Filename = Dr.Wagdy Mohammed_3H_Br
 Author = delta
 Experiment = carbon.jxp
 Sample Id = Dr.Wagdy Mohammed_3H_Br
 Solvent = DMSO-D6
 Creation Time = 24-SEP-2019 03:09:35
 Revision Time = 24-SEP-2019 12:51:49
 Current Time = 24-SEP-2019 12:52:22
 Comment = single pulse decoupled
 Data Format = 1D COMPLEX
 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 = 3700
 Total_Scans = 3700
 Relaxation_Delay = 2 [s]
 Recvr_Gain = 58
 Temp_Get = 18.9 [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_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]



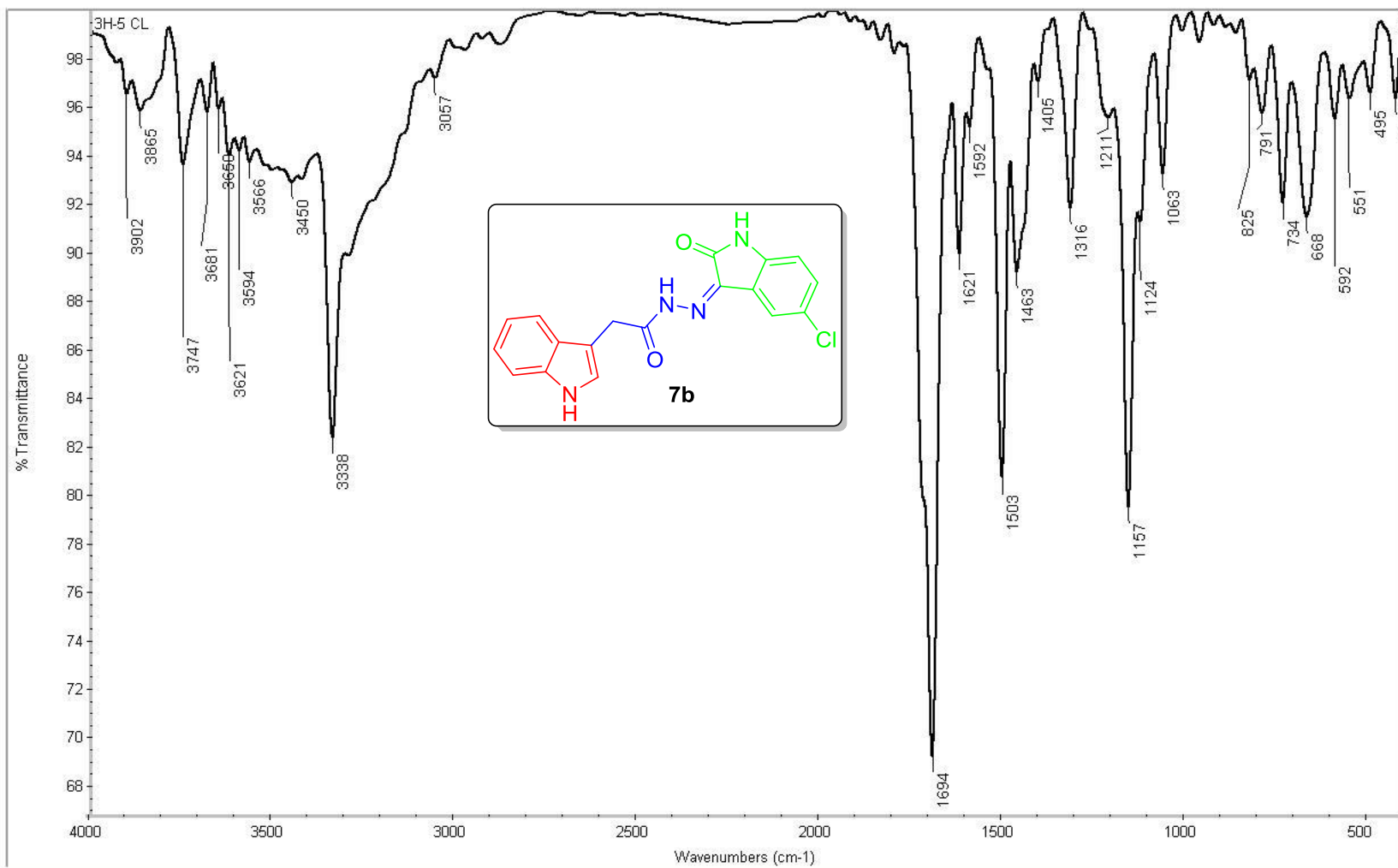
X : parts per Million : Proton

Filename = Dr. Wagdy Mohamd_3H-Br 1
 Author = delta
 Experiment = proton.jxp
 Sample Id = Dr. Wagdy Mohamd_3H-Br 1
 Solvent = DMSO-D6
 Creation Time = 17-SEP-2019 15:23:50
 Revision Time = 17-SEP-2019 15:55:40
 Current Time = 17-SEP-2019 15:55:46
 Comment = single_pulse
 Data Format = 1D COMPLEX
 Dim Size = 13107
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Site = JNM-ECA500II
 Spectrometer = DELTA2 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 = 16384
 X Prescans = 1
 X Resolution = 0.68733284 [Hz]
 X Sweep = 11.26126126 [kHz]
 X Sweep Clipped = 9.00900901 [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]
 Recvr Gain = 48
 Temp Get = 20 [dC]
 X 90 Width = 14.5 [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
 Dante Presat = FALSE
 Initial Wait = 1 [s]
 Repetition Time = 6.4548992 [s]

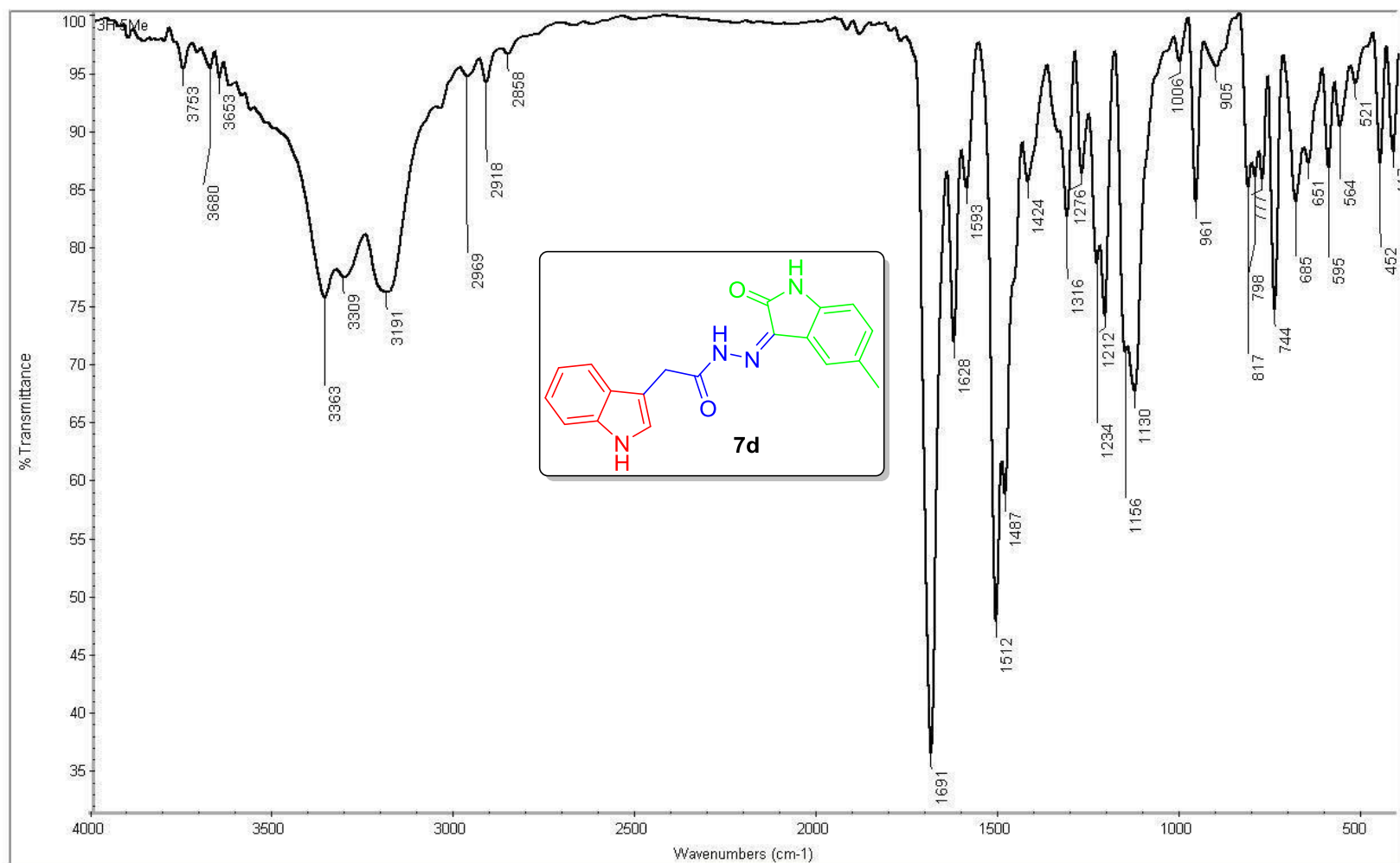
3H-5F



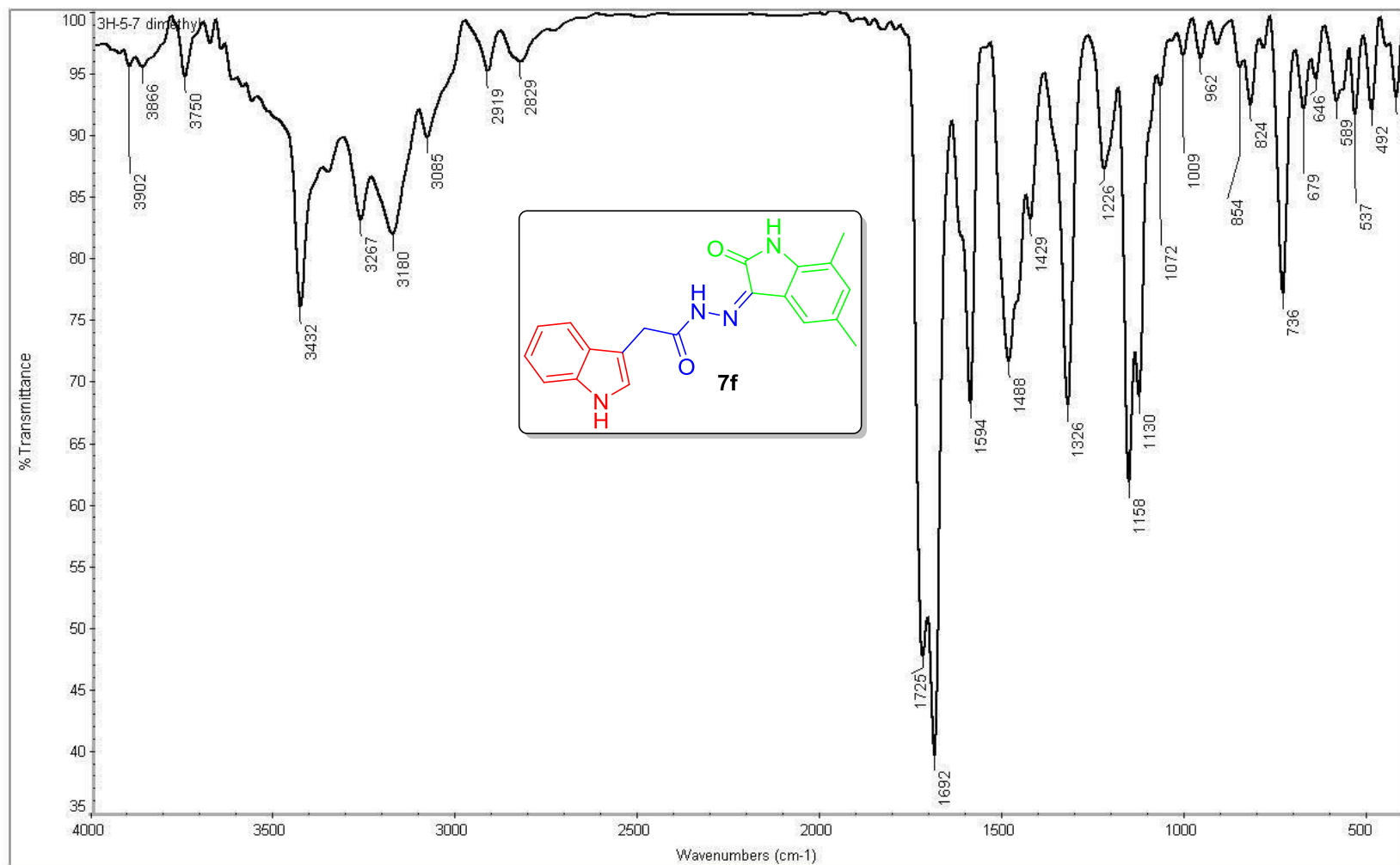
3H-5 CL



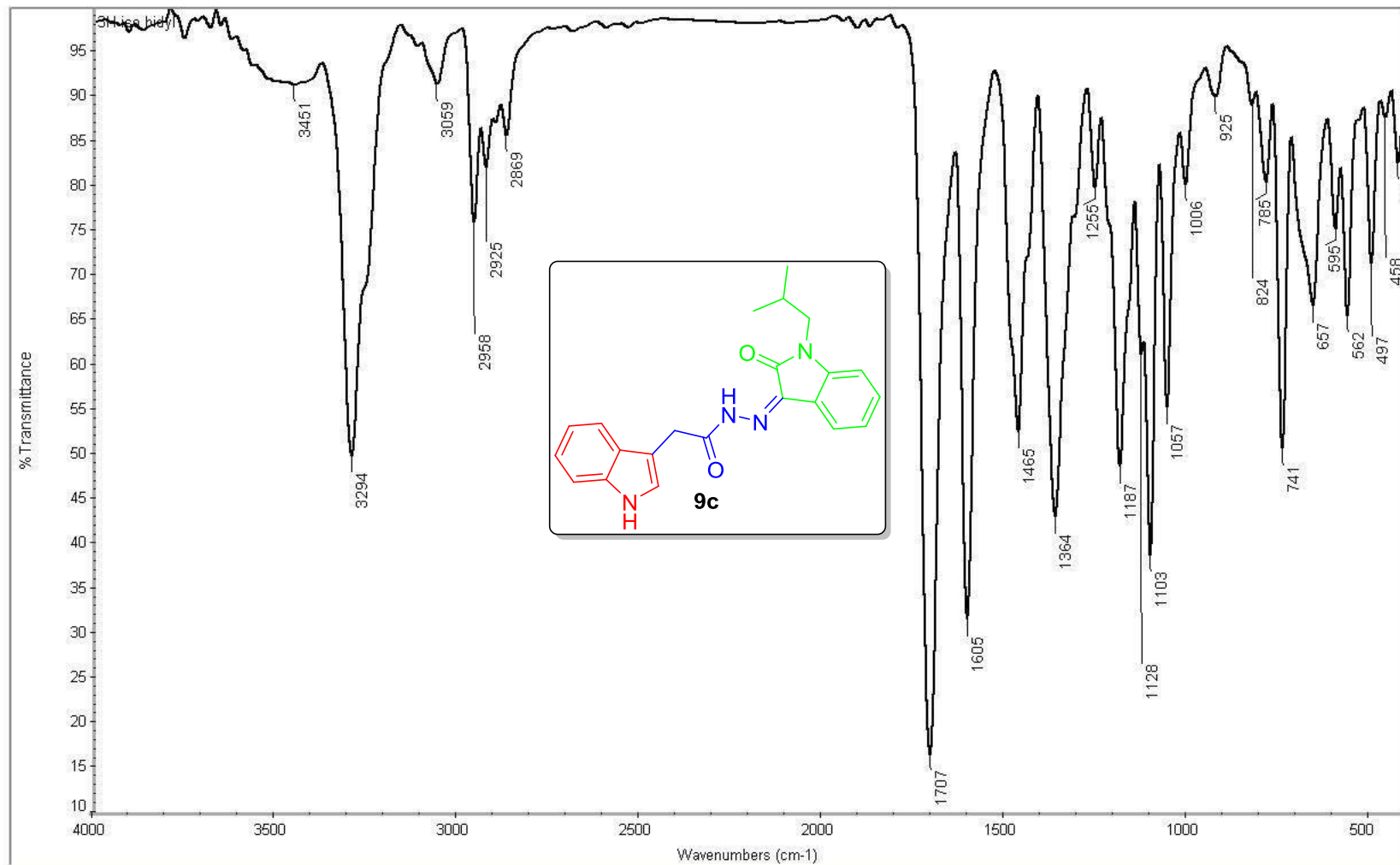
3H-5Me



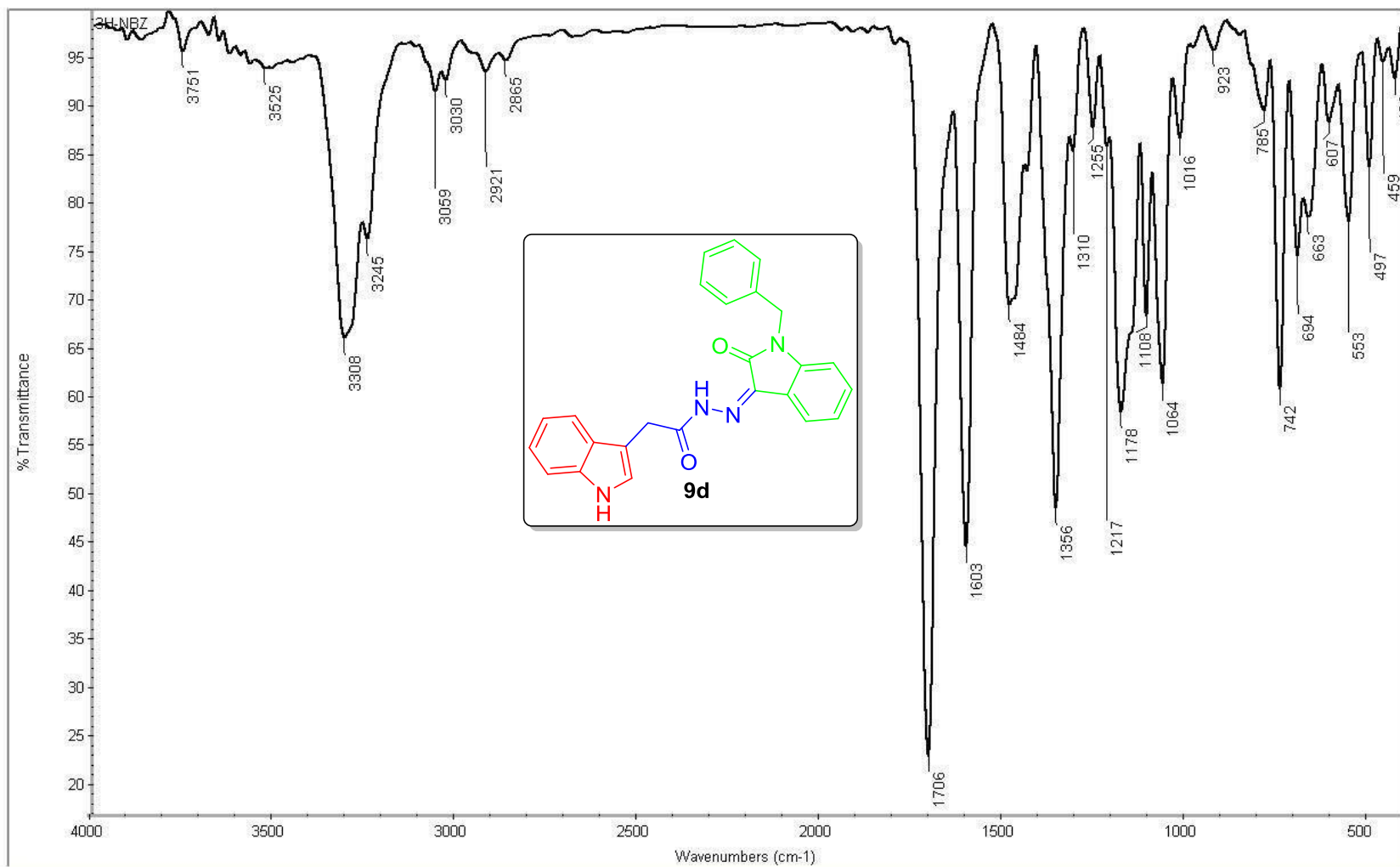
3H-5-7 dimethyl



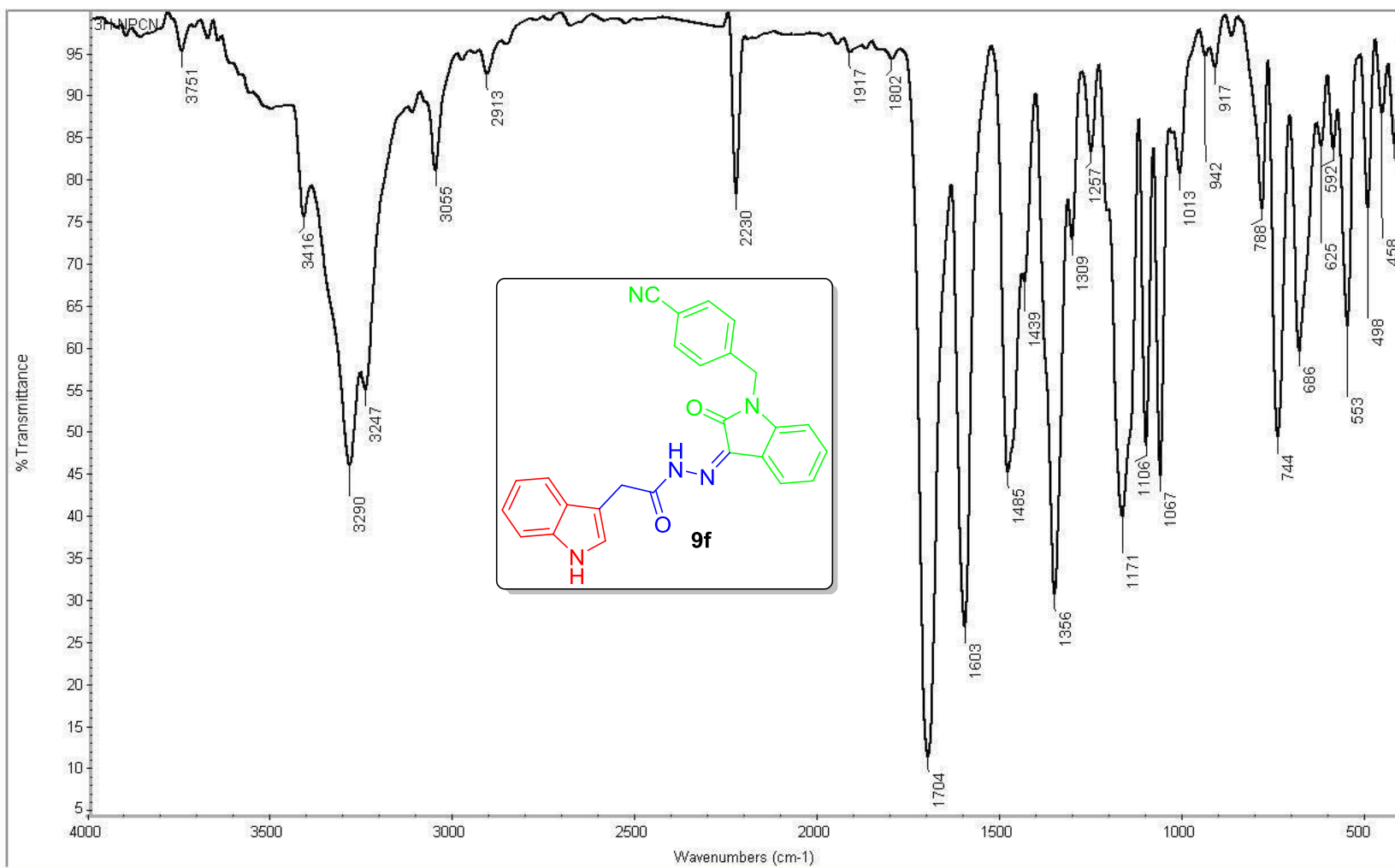
3H-iso bidyl



3H-NBZ



3H-NPCN



3H-Br NBZ

