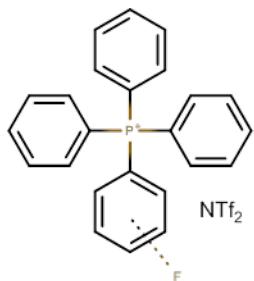


Tuning the Melting Point of Selected Ionic Liquids Through Adjustment of the Cation's Dipole Moment

General Information:

Commercial reagents were obtained from Aldrich Chemical and Oakwood Chemicals and used without further purification. ^1H , ^{19}F , ^{31}P and ^{13}C NMR was recorded on a 500 MHz JEOL spectrometer using CDCl_3 as a solvent at room temperature.

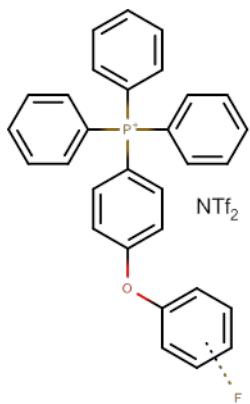
General Procedure for the Synthesis of Fluorophenyltriphenyl Phosphonium $[\text{NTf}_2^-]$ ILs



In a 50 ml heavy wall pressure vessels with an internal thread containing steering bar, Fluoro-iodobenzene (1.0 equiv), triphenyl phosphine (1.0 equiv), $\text{Pd}(\text{OAc})_2$ (1.5 mol%) and Xylene (15 ml) were added under nitrogen atmosphere and the reaction mixture stirred at 140 °C for 2 hours. The reaction mixture was then cooled down to the room temperature and filtered to yield pure fluorophenyltriphenyl phosphonium iodide as a pale-white solid. The $[\text{NTf}_2^-]$ salt was obtained by anion exchange of KNTf_2 (1.0 equiv) and phosphonium

salt (1.0 equiv) in water for 15 min at room temperature. The reaction mixture was then extracted three times with DCM and brine solution. The combined organic extracts were dried over anhydrous Na_2SO_4 and solvents were removed under reduced pressure to furnish desired fluorophenyltriphenyl phosphonium $[\text{NTf}_2^-]$ as a white solid.

General Procedure for the Synthesis of (fluorophenoxy)phenyltriphenylphosphonium $[\text{NTf}_2^-]$ ILs

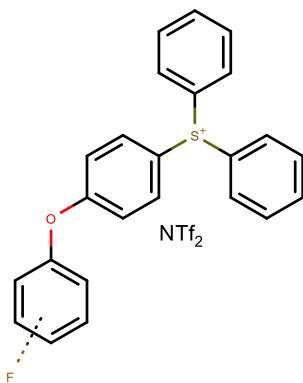


First 3-(fluorophenoxy)bromobenzene was prepared using following method: In a 100 ml heavy wall pressure vessels with an internal thread containing steering bar, 3-fluorophenylboronic acid (2.0 equiv, 10 g), 4-bromophenol (1.0 equiv), triethylamine (5.0 equiv), Copper(II) acetate (1.0 equiv), and MS 4Å(2.0 g) powder and the reaction mixture stirred for 5 hours at room temperature. After the completion of reaction the mixture was filtered through a filter paper and concentrated under reduced pressure. Further purification with column chromatography (Hexane/EtOAc) resulted pure 3-(fluorophenoxy)bromobenzene. Next step, In a 50 ml heavy wall pressure vessels with an internal thread containing steering bar, 3-(fluorophenoxy)bromobenzene (1.0 equiv), triphenyl phosphine (1.0 equiv), $\text{NiCl}_2 \cdot 6\text{H}_2\text{O}$ (10 mole%) ethylene glycol (10 ml) were added under nitrogen atmosphere and the reaction mixture stirred at 160 °C for 12 hours. The reaction mixture was then extracted three times with DCM and brine solution and concentrated under reduced pressure to yield pure 3-(fluorophenoxy)phenyltriphenylphosphonium bromide. The $[\text{NTf}_2^-]$ salt was obtained by anion exchange of KNTf_2 (1.0 equiv) and phosphonium salt (1.0 equiv) in water for 15 min at room temperature. The reaction mixture was then extracted three times with DCM and brine solution. The combined organic extracts were dried over anhydrous Na_2SO_4 and solvents were removed

The combined organic extracts were dried over anhydrous Na_2SO_4 and solvents were removed

under reduced pressure to furnish pure 3-(fluorophenoxy)phenyltriphenylphosphonium $[\text{NTf}_2^-]$ as a white solid. Same procedure was applied for the synthesis of 2-(fluorophenoxy)phenyltriphenylphosphonium $[\text{NTf}_2^-]$ and 4-(fluorophenoxy)phenyltriphenylphosphonium $[\text{NTf}_2^-]$.

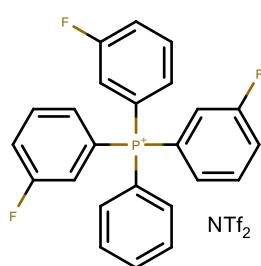
General Procedure for the Synthesis of triphenylsulfonium $[\text{NTf}_2^-]$ ILs



In a 100 ml round bottom flask containing a steering bar, Diphenyl Sulfoxide (1.0 equiv, 5g) and Fluorophenoxy benzene (1.1 equiv) in DCM (50 ml) was stirred for 10 minutes at room temperature. Triflic anhydride solution (1.05 equiv) in DCM was added dropwise to the reaction mixture at room temperature and reaction mixture stirred for overnight at room temperature. After completion of the reaction, solvent was removed under vacuum and washed with ether to remove the unreacted materials to furnish the desired sulfonium product. The $[\text{NTf}_2^-]$ salt was obtained by anion exchange of excess KNTf_2 (5 equiv) and sulfonium salt (1.0 equiv) in water-methanol for 30 min at room temperature.

The reaction mixture was then extracted three times with DCM and brine solution and the combined organic extracts were dried over anhydrous Na_2SO_4 . Then solvents were removed under reduced pressure to result desired triphenylsulfonium $[\text{NTf}_2^-]$.

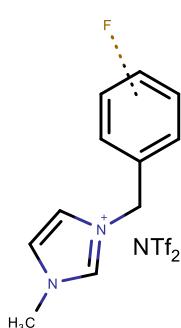
Synthesis of tris(3-fluorophenyl)(phenyl)phosphonium $[\text{NTf}_2^-]$ IL



In a 50 ml heavy wall pressure vessels with an internal thread containing steering bar, Bromobenzene (1.0 equiv), Tris(3-fluorophenyl)phosphine (1.0 equiv), $\text{NiCl}_2 \cdot 6\text{H}_2\text{O}$ (10 mole%) ethylene glycol (10 ml) were added under nitrogen atmosphere and the reaction mixture stirred at 180 °C for 15 hours. The reaction mixture was then cooled down to the room temperature and was extracted three times with DCM / brine solution and concentrated under reduced pressure to yield tris(3-fluorophenyl)(phenyl)phosphonium bromide. The $[\text{NTf}_2^-]$ salt was obtained by anion exchange of KNTf_2 (1.0 equiv) and phosphonium salt (1.0 equiv) in water for 15 min at room temperature.

The reaction mixture was then extracted three times with DCM and brine solution. The combined organic extracts were dried over anhydrous Na_2SO_4 and solvents were removed under reduced pressure to furnish pure of tris(3-fluorophenyl)(phenyl)phosphonium $[\text{NTf}_2^-]$. Same method was applied for the synthesis of tris(4-fluorophenyl)(phenyl)phosphonium $[\text{NTf}_2^-]$.

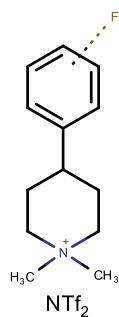
Synthesis of 3-benzyl-1-methyl-imidazolium $[\text{NTf}_2^-]$ ILs



In a 100 ml round bottom flask containing a steering bar, 1-methylimidazol (1.0 equiv, 5g) and Benzyl Bromide (1.0 equiv) in THF (50 ml) was stirred for overnight at 40 °C. Then the reaction mixture was filtered and white solid yielded as a pure product. Anion exchange with KNTf_2 (1.0 equiv) carried out in water and stirred for 30 min at room temperature. The reaction mixture was then extracted three times with DCM and brine solution and the combined organic extracts were dried

over anhydrous Na_2SO_4 . Then solvents were removed under reduced pressure to produce pure 3-benzyl-1-methyl-imidazolium $[\text{NTf}_2^-]$ as a colorless liquid.

1,1-dimethyl-4-phenylpiperidinium $[\text{NTf}_2^-]$ ILs



For the synthesis of 1,1-dimethyl-4-phenylpiperidin-1-ium $[\text{NTf}_2^-]$, first 1,1-dimethyl-4-phenylpiperidinium iodide salt was prepared using following method. In a 100 ml round bottom flask containing a steering bar, 4-phenylpiperidine (1.0 equiv, 2 g), iodomethane (5.0 equiv), and potassium bicarbonate (1.5 equiv) in methanol (50 ml) was stirred for 24 hours at room temperature. Then solvent was removed under vacuum and the mixture was dissolved in DCM and filtered through filter paper. Filtrate was concentrated under reduced pressure and pure product was obtained in high yield. Anion exchange with KNTf_2 in water furnished 1,1-dimethyl-4-phenylpiperidinium $[\text{NTf}_2^-]$. The reaction mixture was then extracted with DCM and the combined organic extracts were dried over anhydrous Na_2SO_4 . Then solvents were removed under reduced pressure to produce pure 1,1-dimethyl-4-phenylpiperidinium $[\text{NTf}_2^-]$ as a white solid.



```

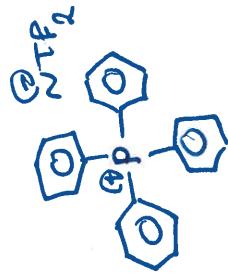
Filename = MS0299_PROTON-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0299
Solvent = CHLOROFORM-D
Changer_sample = 7
Creation_time = 9-JUN-2018 18:05:00
Revision_time = 9-JUN-2018 17:38:47
Current_time = 9-JUN-2018 17:38:47

Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JEOL-ECA500

Field_strength = 11.7473579 [T] (500 [MHz]
X_acq_duration = 1.74587904 [s]
X_domain = 1H
X_freq = 500.15991521 [MHz]
X_offset = 5.0 [ppm]
X_points = 16384
X_prescans = 1
X_resolution = 0.57277737 [Hz]
X_sweep = 9.38438438 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Tri_domain = 1H
Tri_freq = 500.15991521 [MHz]
Tri_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

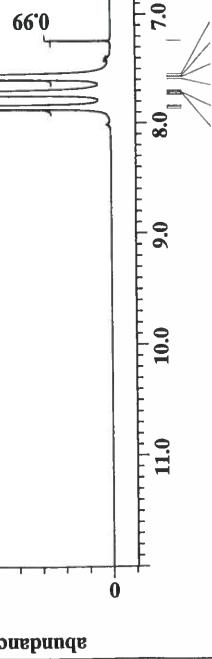
X_90_width = 12.4 [us]
X_acq_time = 1.74587904 [s]
X_angle = 45 [deg]
X_atn = 4 [dB]
X_pulse = 6.2 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Recv_gain = 28
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get = 21.1 [dC]

```



183.32

121.6



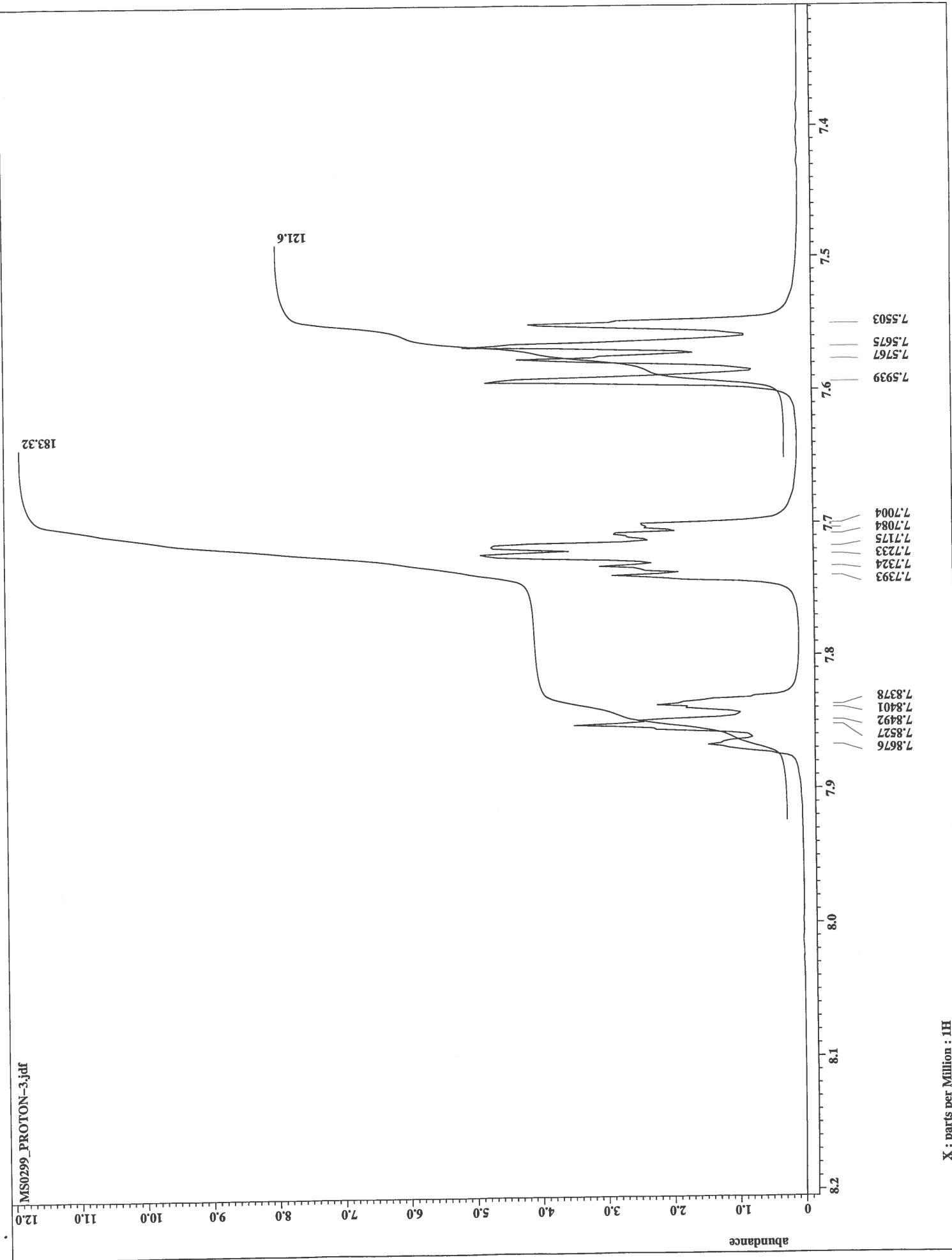
7.7233
7.7175
7.5939
7.5767
7.5503

MS0299_PROTON-2.jdf

5.0 4.0 3.0 2.0 1.0 0

11.0 10.0 9.0 8.0 7.0 6.0 5.0 4.0 3.0 2.0 1.0 0

abundance





```

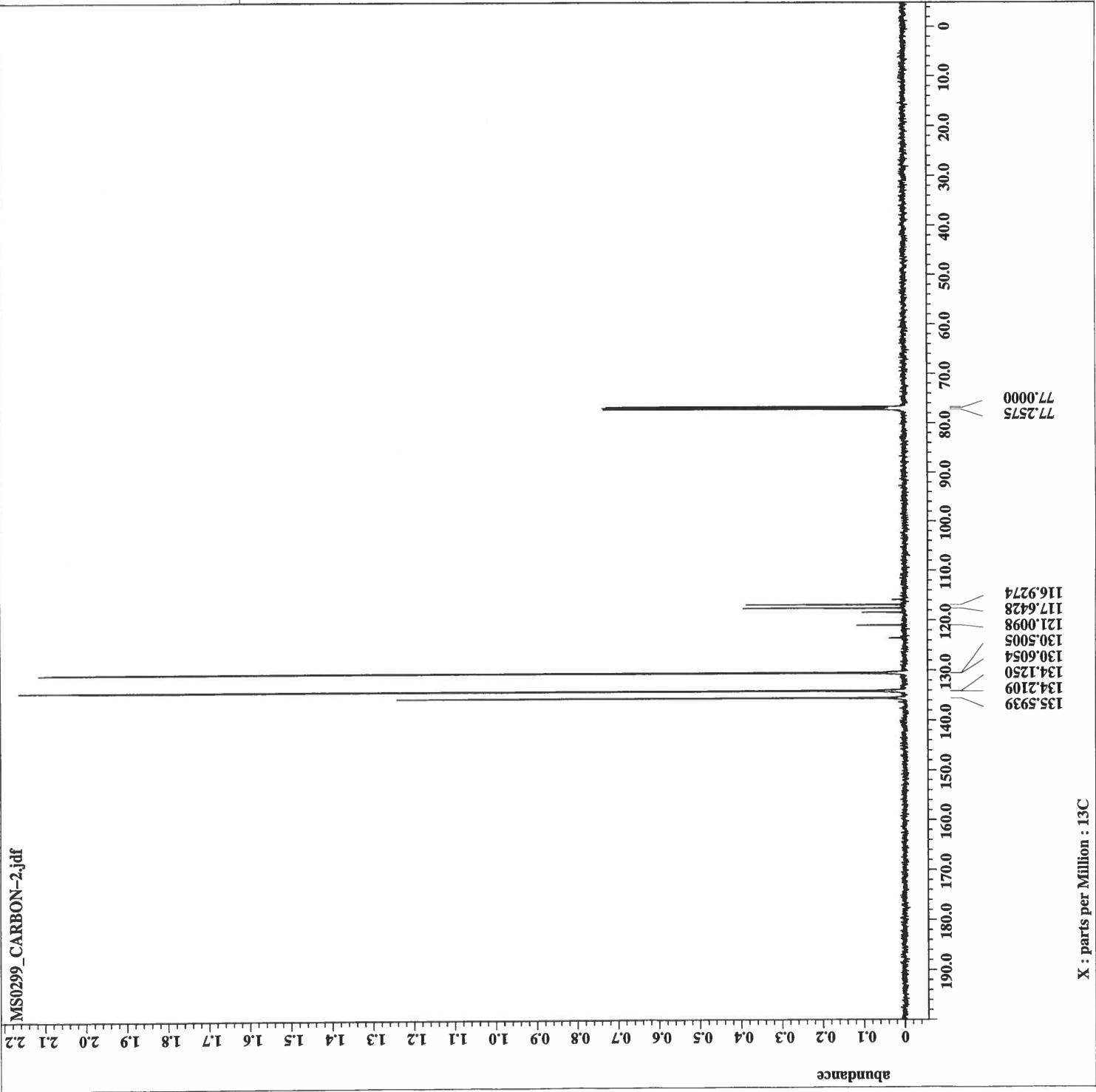
Filename = MS0299_CARBON-2.jdf
Author = Jim Davis
Experiment =
Sample_id = MS0299
Solvent = CHLOROFORM-D
Changer_sample = 7
Creation_time = 9-JUN-2018 18:19:22
Revision_time = 9-JUN-2018 17:53:10
Current_time = 9-JUN-2018 17:53:10

Data_format = 1D COMPLEX
Dim_size = 2614
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain = 13C
X_freq = 128.76539768 [MHz]
X_offset = 100.0 [ppm]
X_points = 32768
X_precans = 4
X_resolution = 1.19959034 [Hz]
X_sweep = 39.3082761 [Hz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 256
Total_scans = 256

X_90_width = 13.2 [us]
X_acq_time = 0.83361792 [s]
X_angle = 30 [deg]
X_atn = 6 [dB]
X_pulse = 4.4 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Regrv_gain = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get = 21.7 [dc]

```

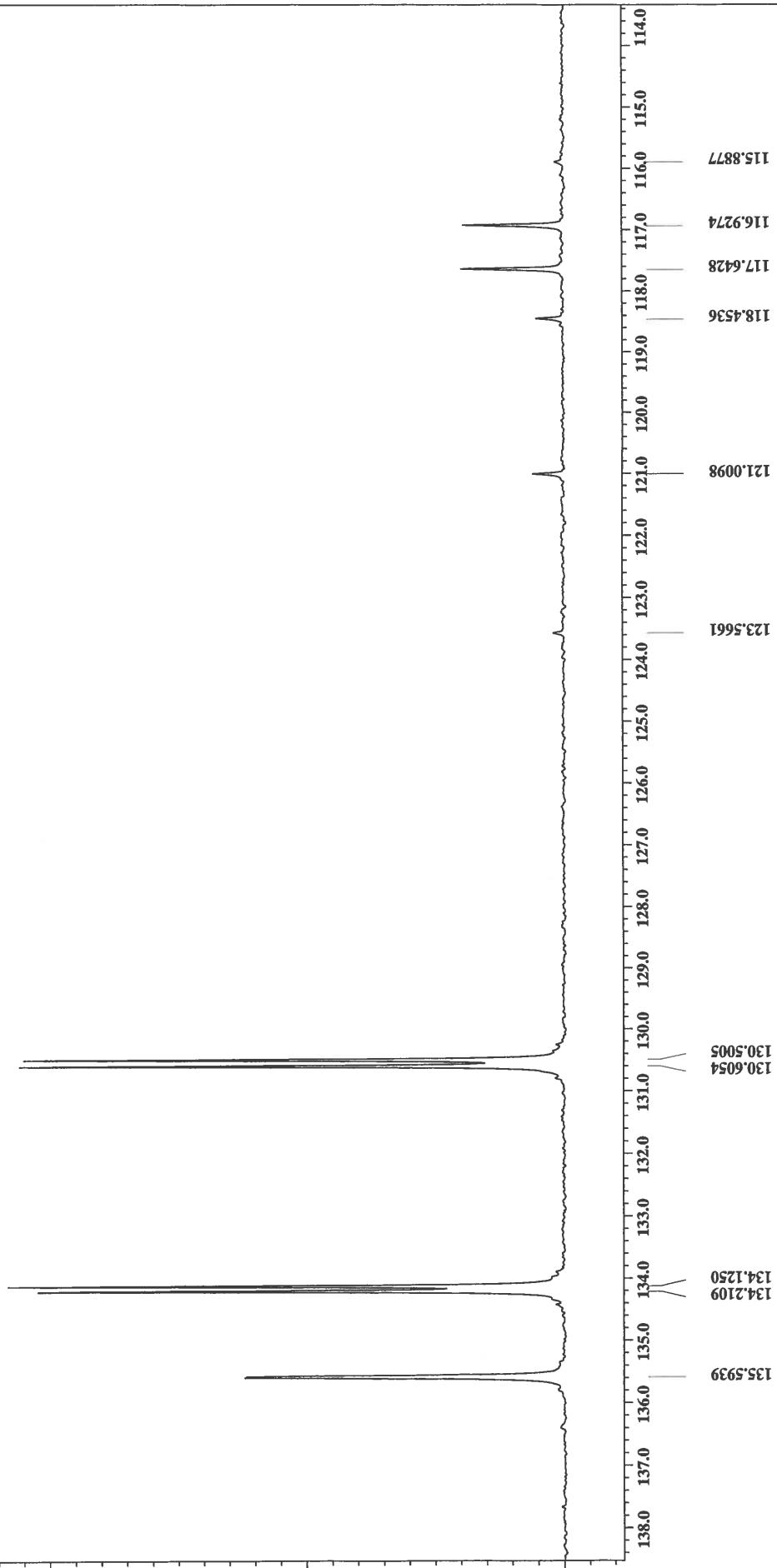


3.0

2.0

1.0

abundance

X : parts per Million : ^{13}C 



```

Filename = MS0299_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0299
Solvent = CHLOROFORM-D
Changer_sample = 7
Creation_time = 9-JUN-2018 18:22:19
Revision_time = 9-JUN-2018 17:56:06
Current_time = 9-JUN-2018 17:56:06

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574538 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 55536
X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [Hz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_pressat = FALSE
Initial_wait = 1 [s]
Regrv_gain = 38
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 21.4 [dc]


```





```

Filename      = MS0299_PHOSPHORUS-2.jdf
Author        = Jim Davis
Experiment    = single_pulse_dec
Sample_id     = MS0299
Solvent       = CHLOROFORM-D
Changer_sample = 7
Creation_time = 9-JUN-2018 18:25:53
Revision_time = 9-JUN-2018 17:59:40
Current_time  = 9-JUN-2018 17:59:40

Date_format   = 1D COMPLEX
Dim_size      = 26214
Dim_title     = 31P
Dim_units     = [ppm]
Dimensions    = X
Site          = ECA_500
Spectrometer  = JNM-ECA500

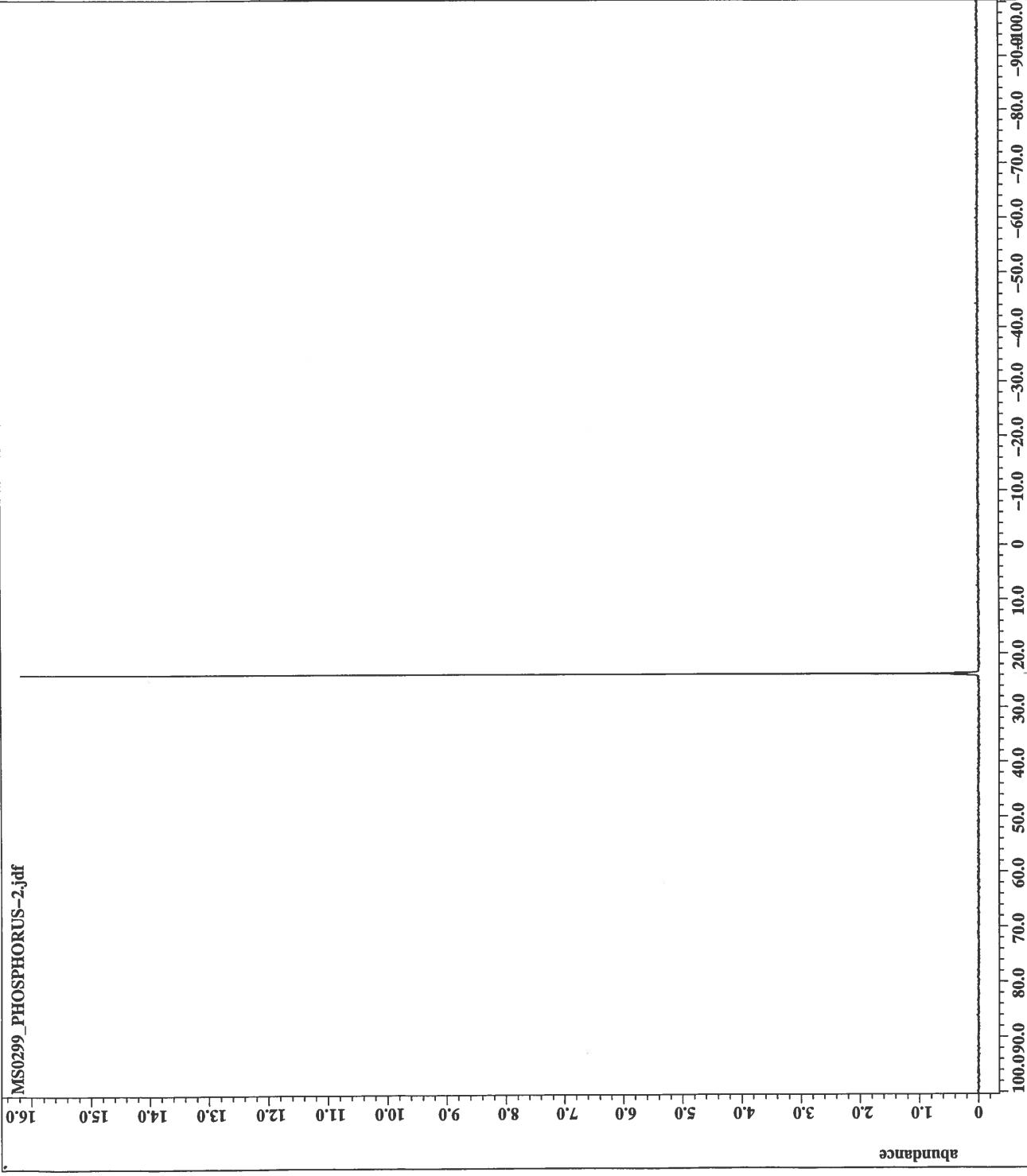
Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.64487424 [s]
X_domain      = 31P
X_freq         = 202.46831075 [MHz]
X_offset       = 0 [ppm]
X_points       = 32768
X_prescans    = 4
X_resolution  = 1.55068995 [Hz]
X_sweep        = 50.81300813 [kHz]
Irr_domain    = 1H
Irr_freq       = 500.15991521 [MHz]
Irr_offset     = 5.0 [ppm]
Clipped        = FALSE
Mod_return    = 1
Scans          = 25
Total_scans   = 25

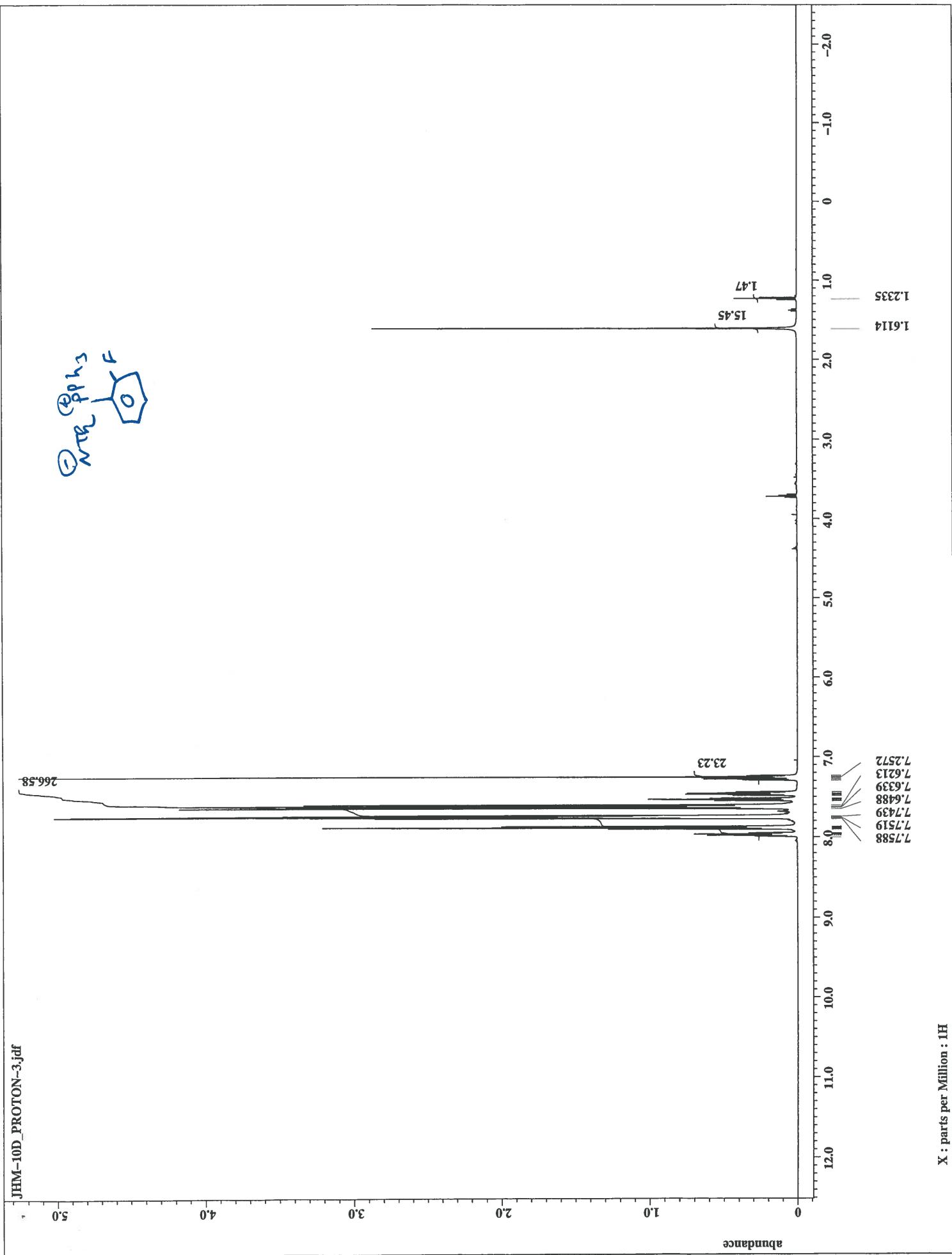
X_90_width    = 14.687 [us]
X_acq_time    = 0.64487424 [s]
X_angle        = 30 [deg]
X_atn         = 5 [dB]
X_pulse        = 4.89566667 [us]
Irr_atn_dec   = 20.7 [dB]
Irr_atn_noe   = 20.7 [dB]
Irr_noise     = WALTZ
Decoupling    = TRUE
Initial_wait   = 1 [s]
Noe           = TRUE
Noe_time      = 2 [s]
Regrv_gain    = 56
Relaxation_delay = 2 [s]
Repetition_time = 2.64487424 [s]
Temp_get      = 21.6 [dc]

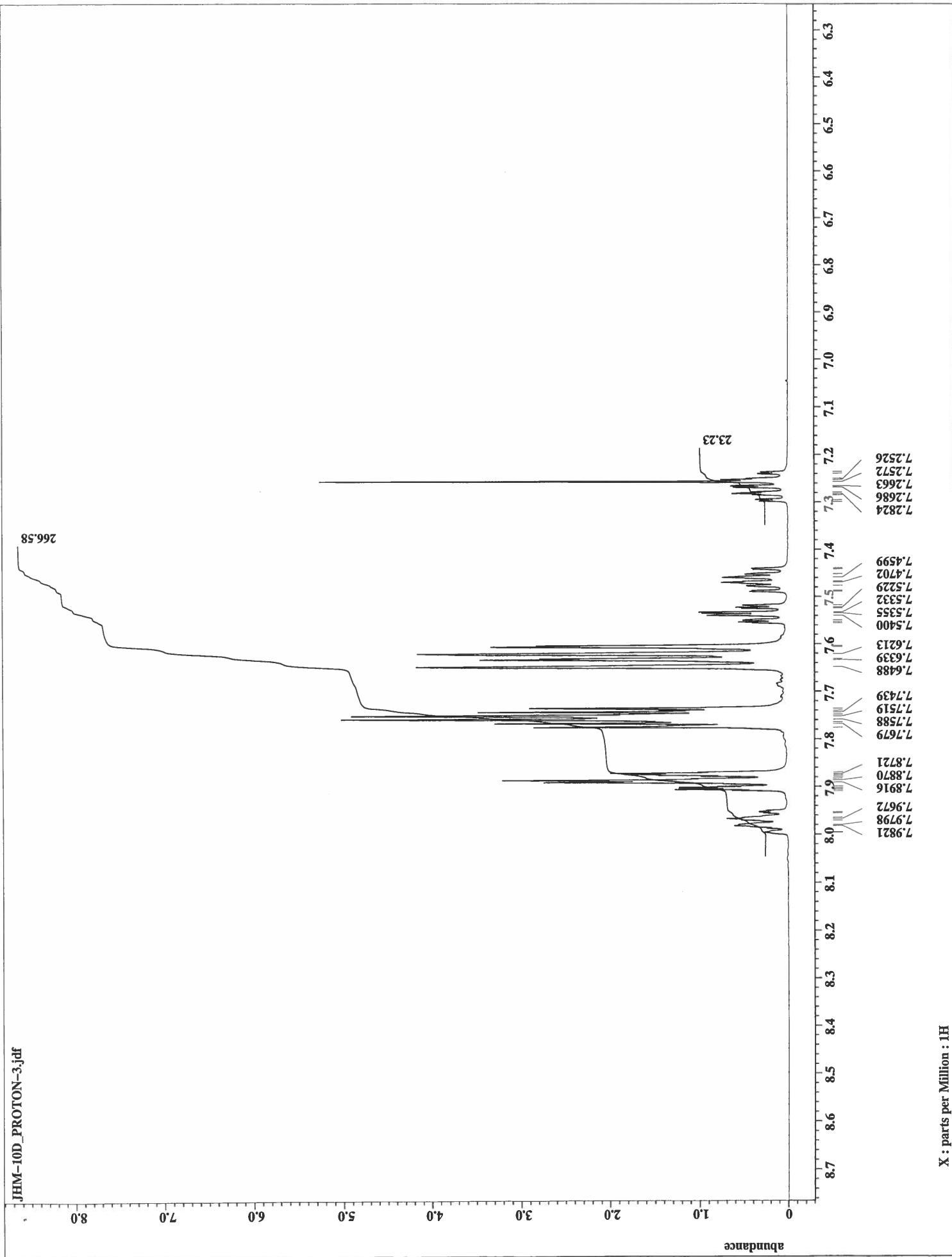
]

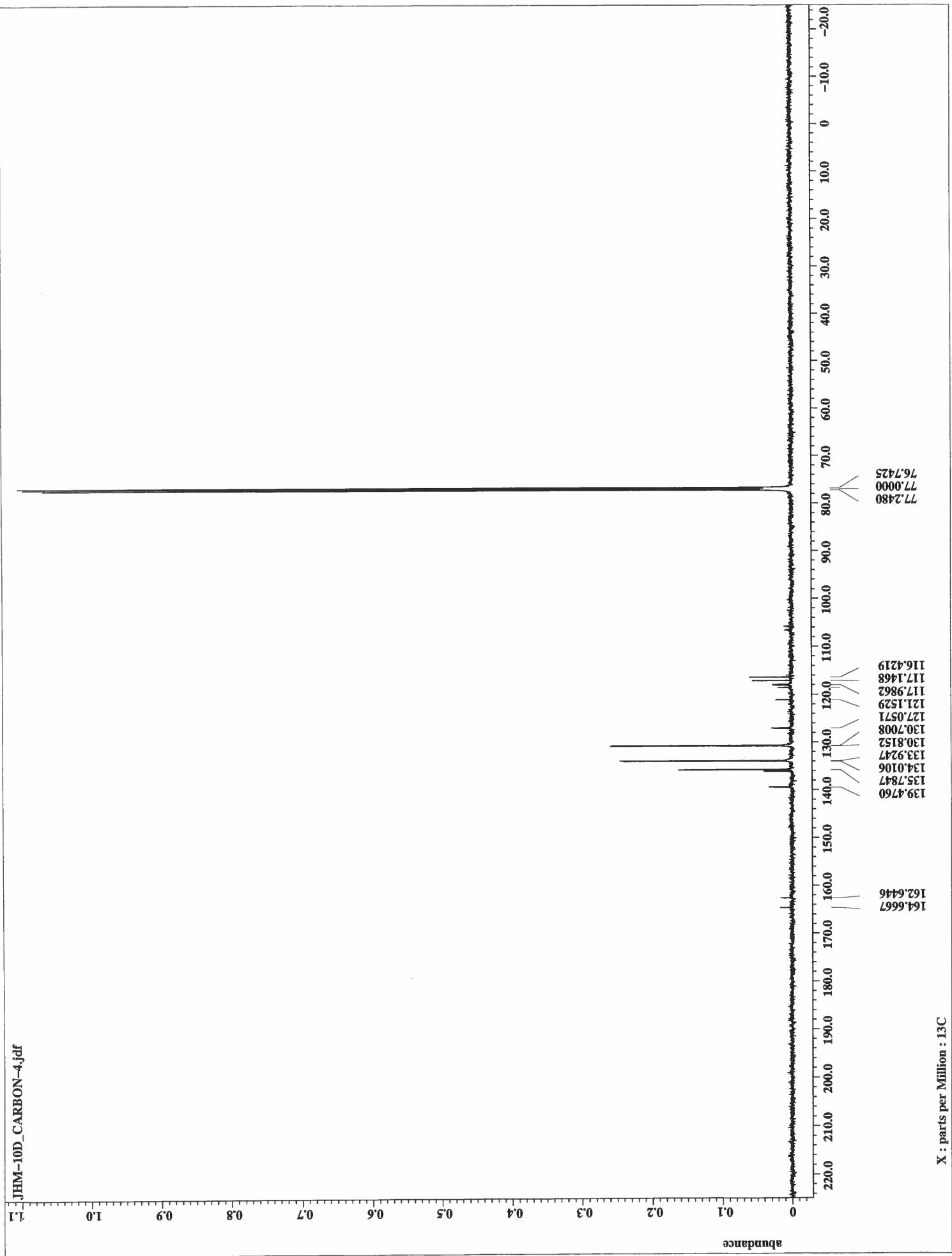
```

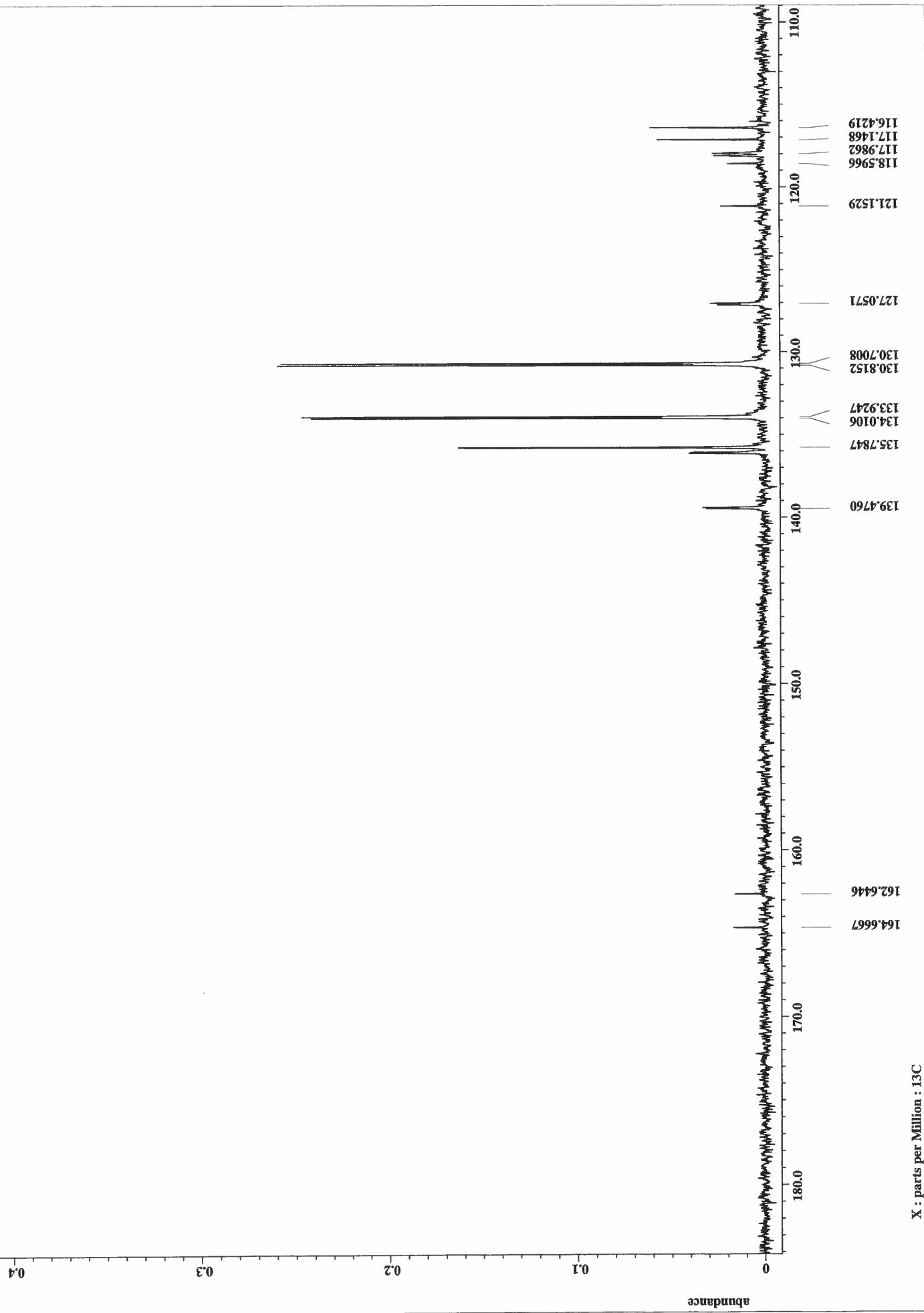
23.8193

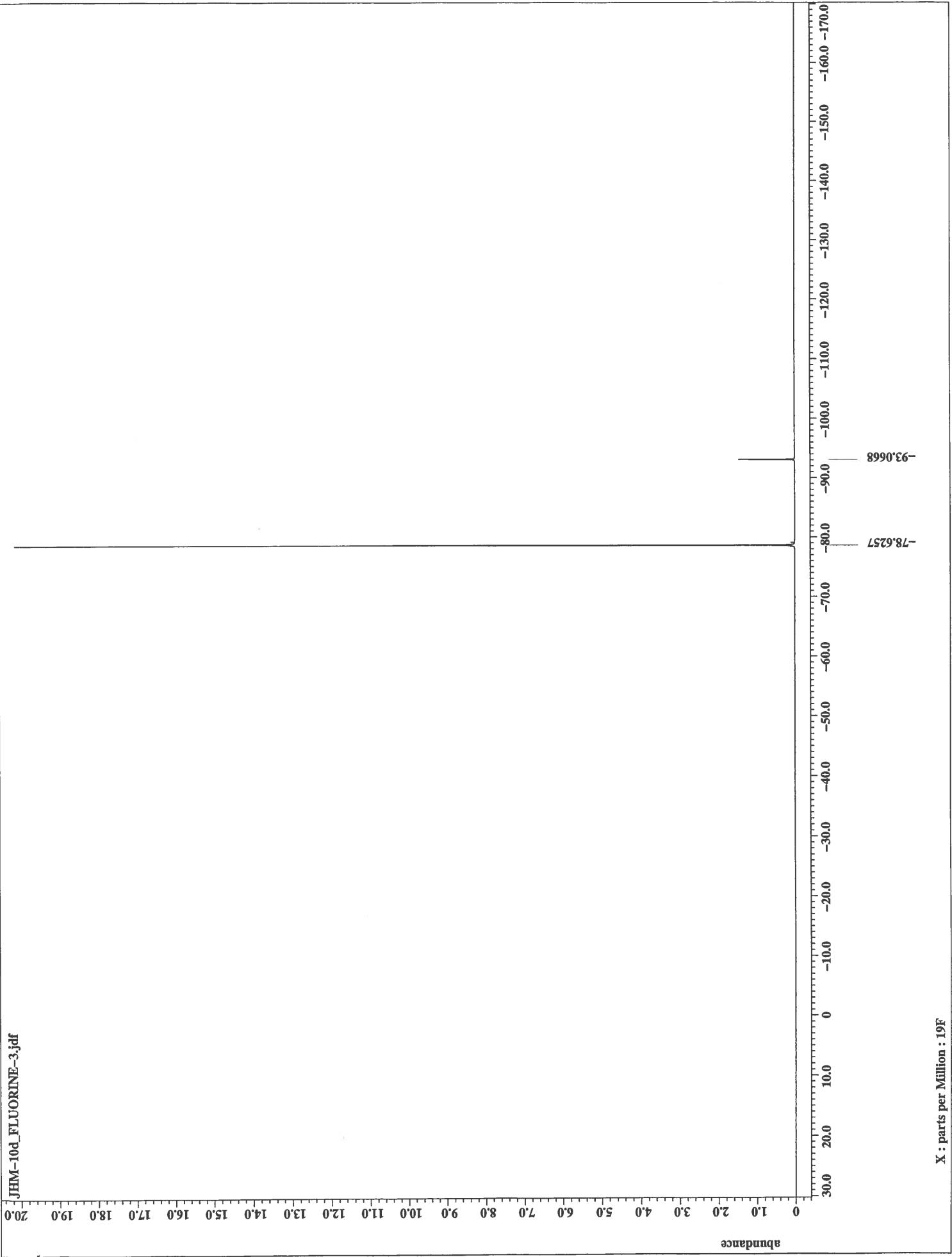












3.0

2.0

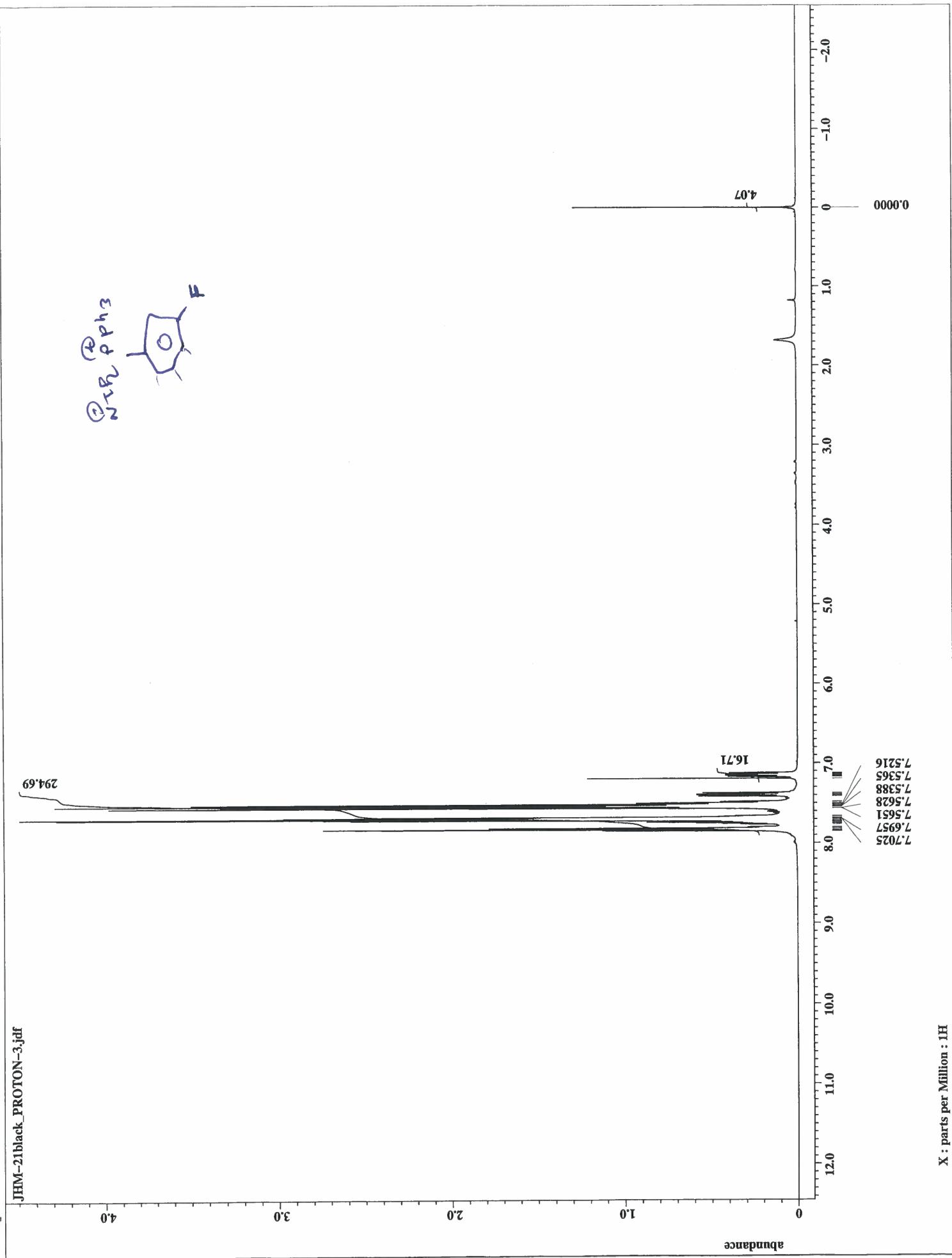
1.0

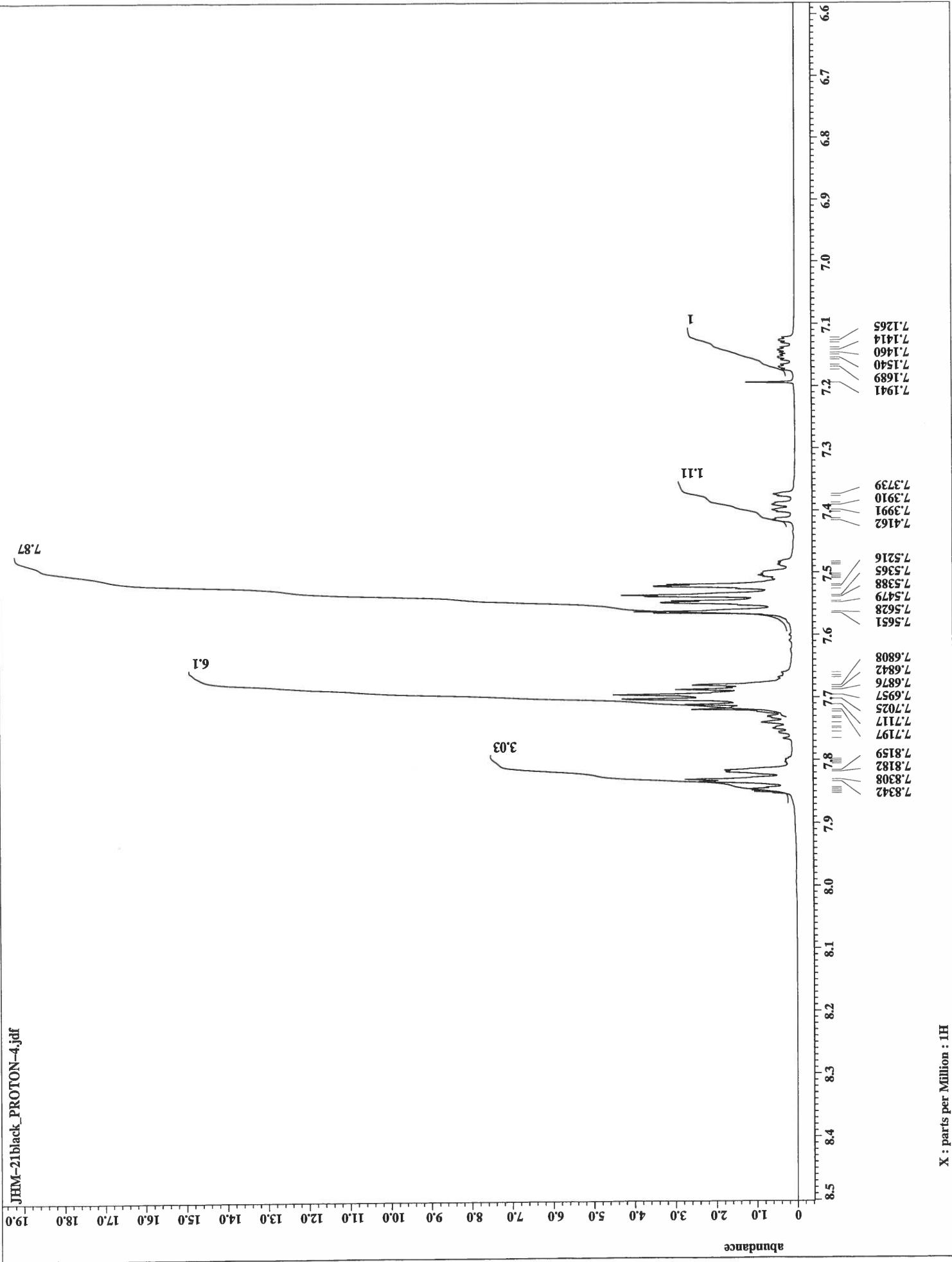
abundance

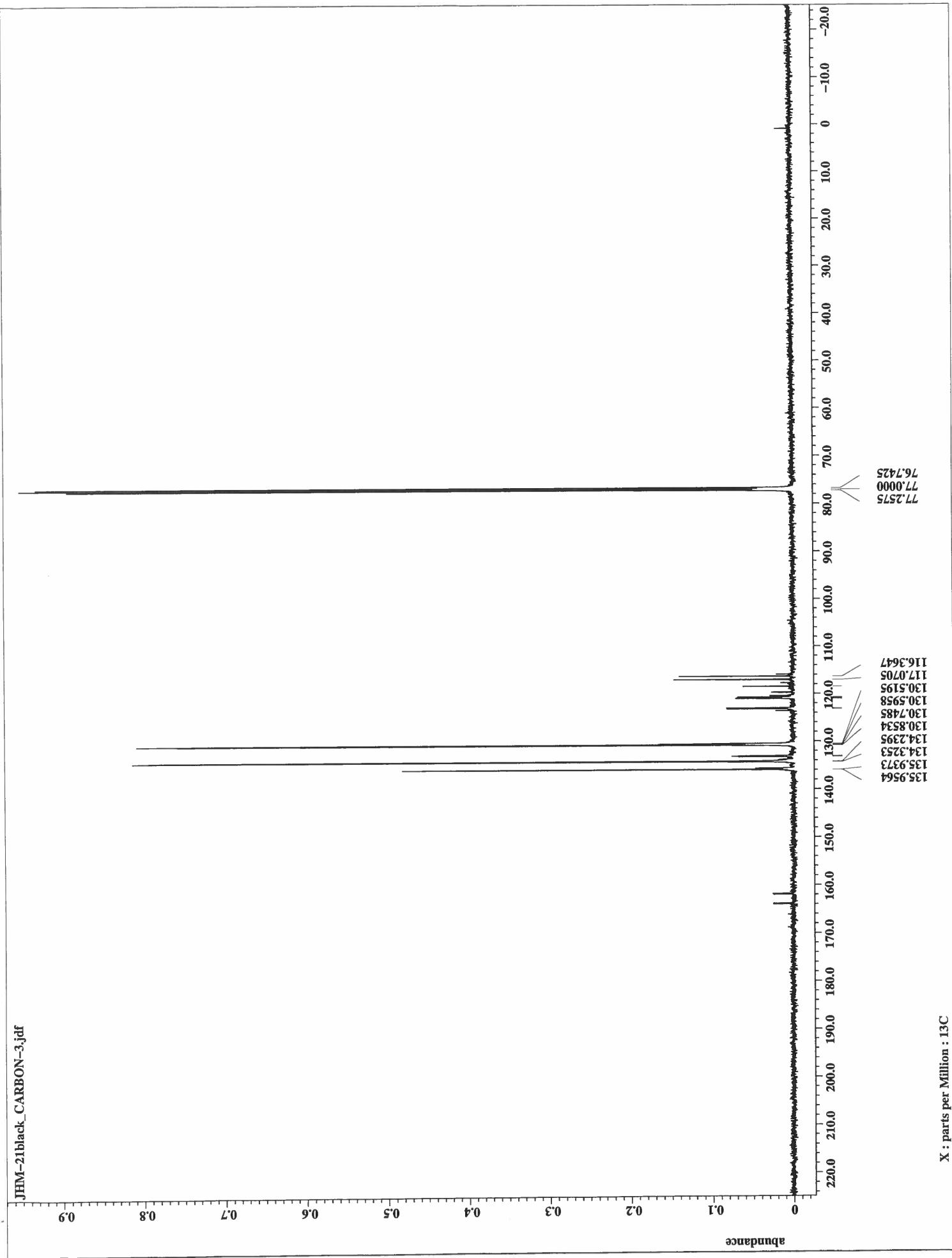
21.6365

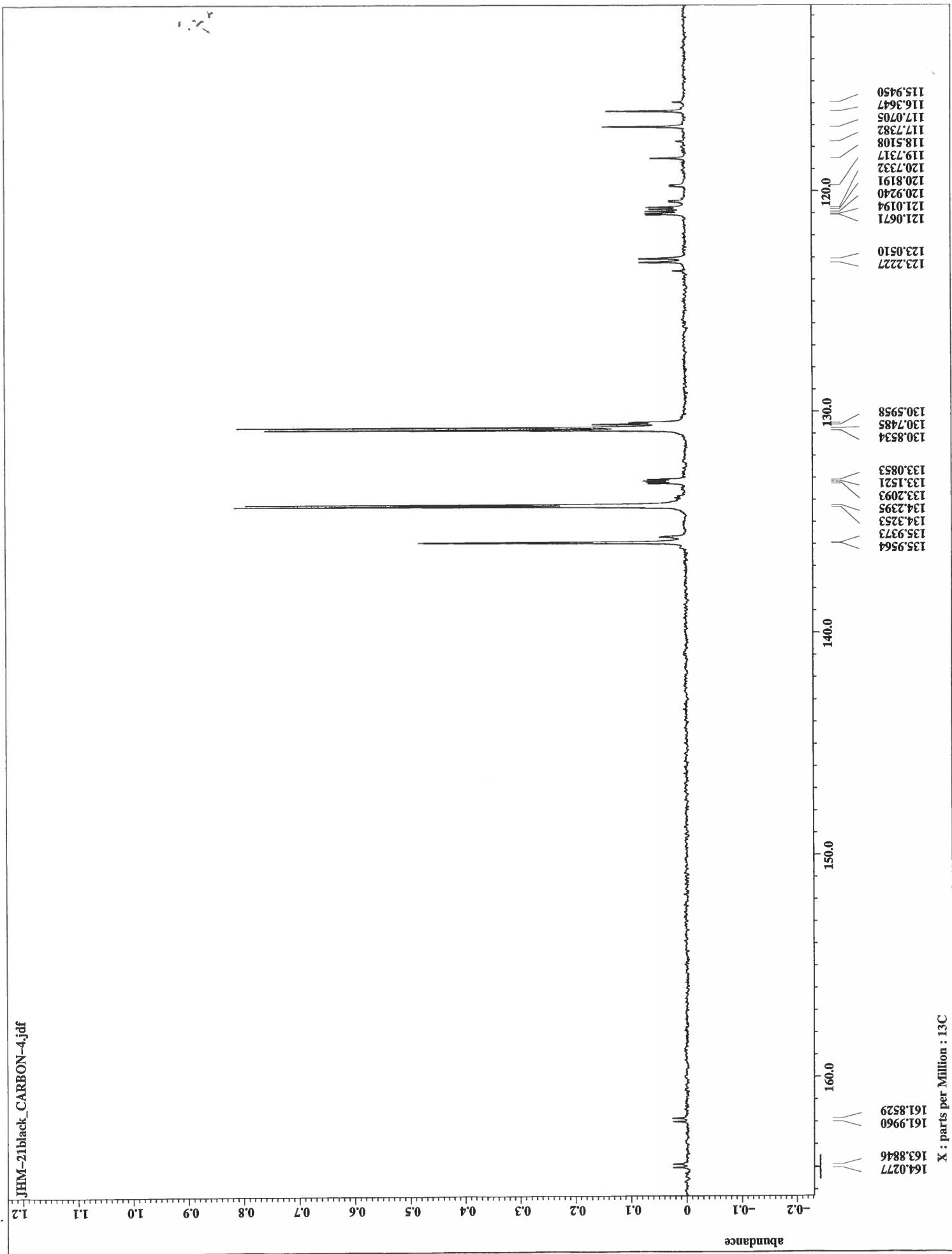
X : parts per Million : 31P

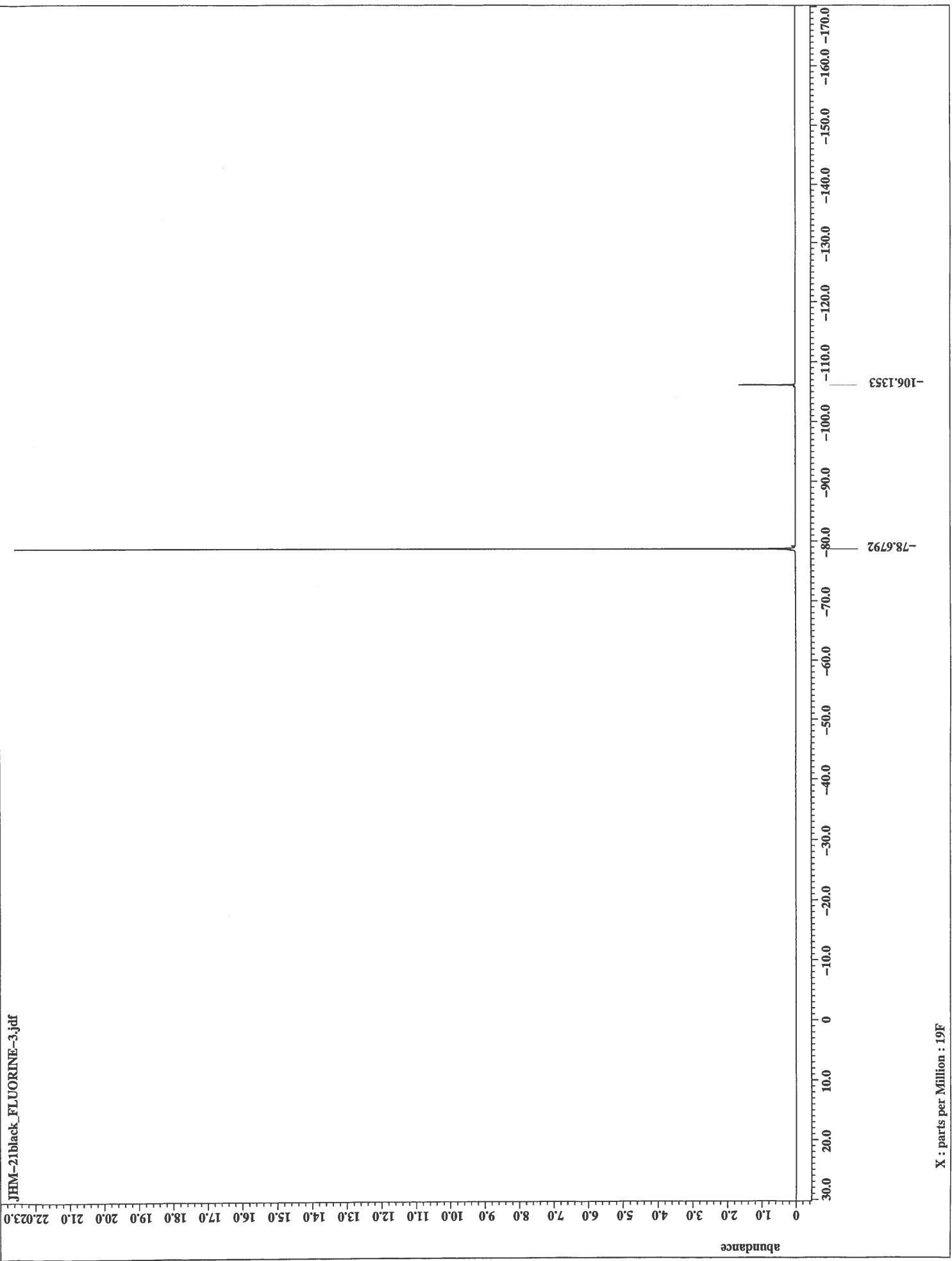
100.0 90.0 80.0 70.0 60.0 50.0 40.0 30.0 20.0 10.0 0 -10.0 -20.0 -30.0 -40.0 -50.0 -60.0 -70.0 -80.0 -90.0 -100.0

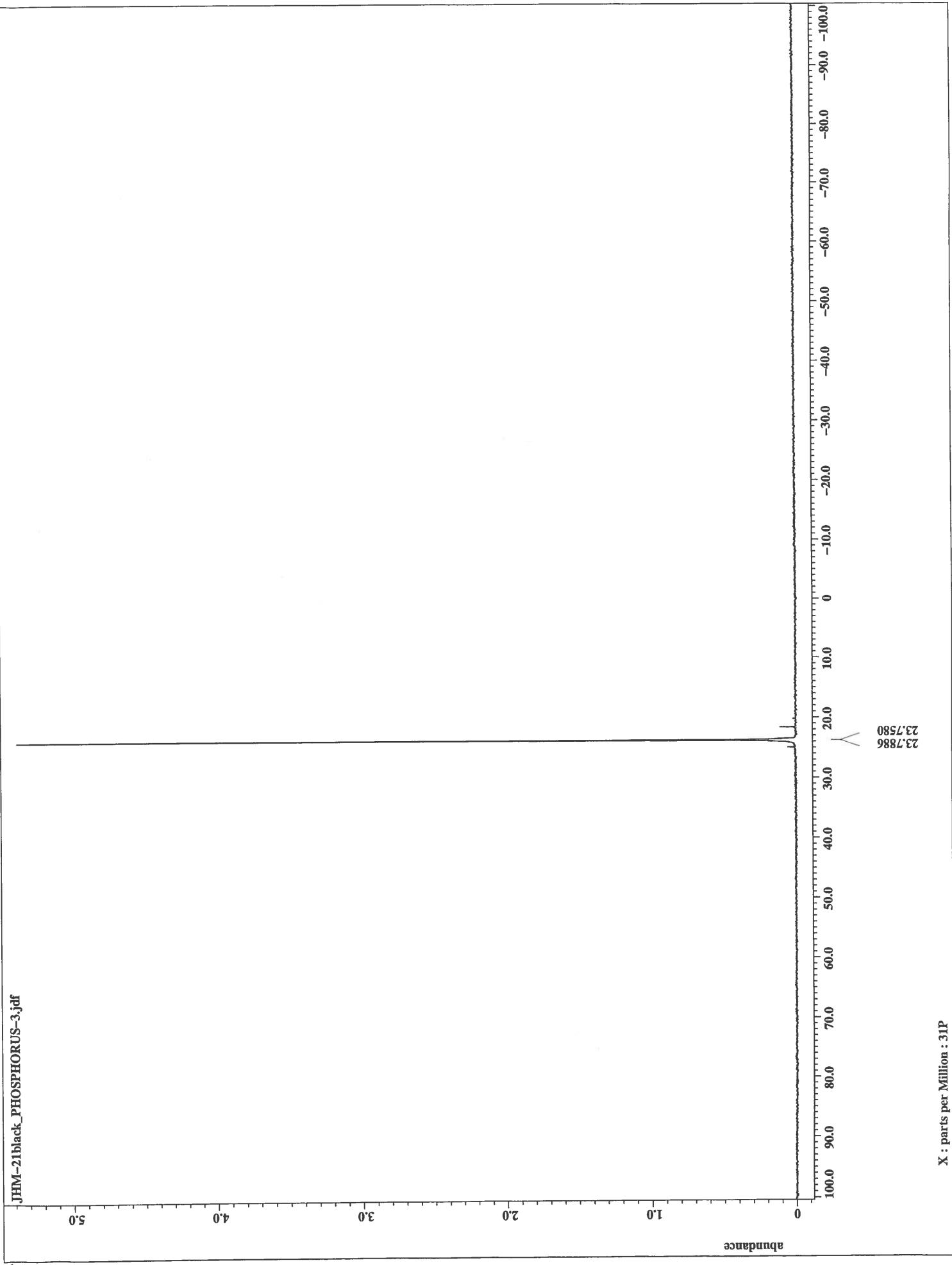


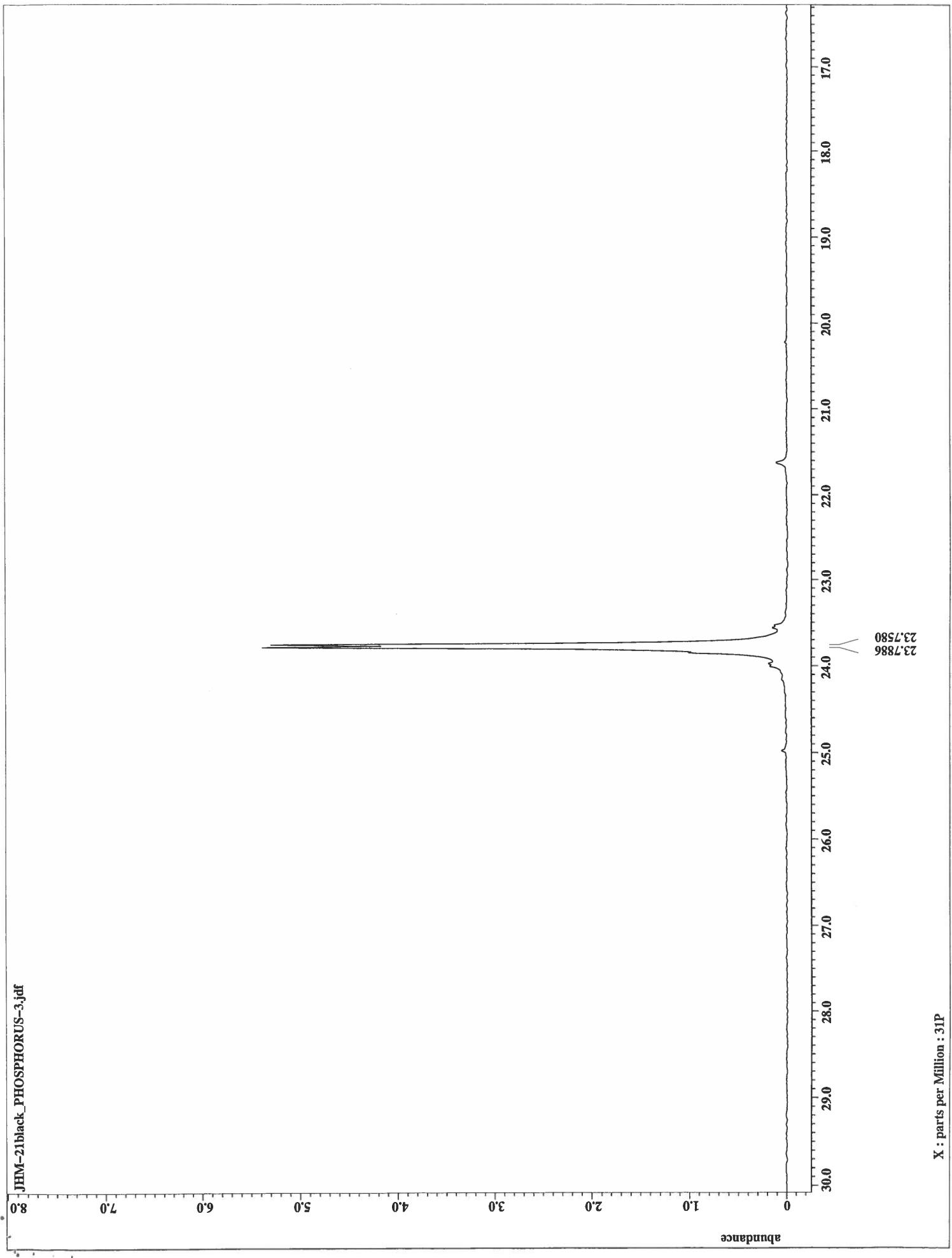


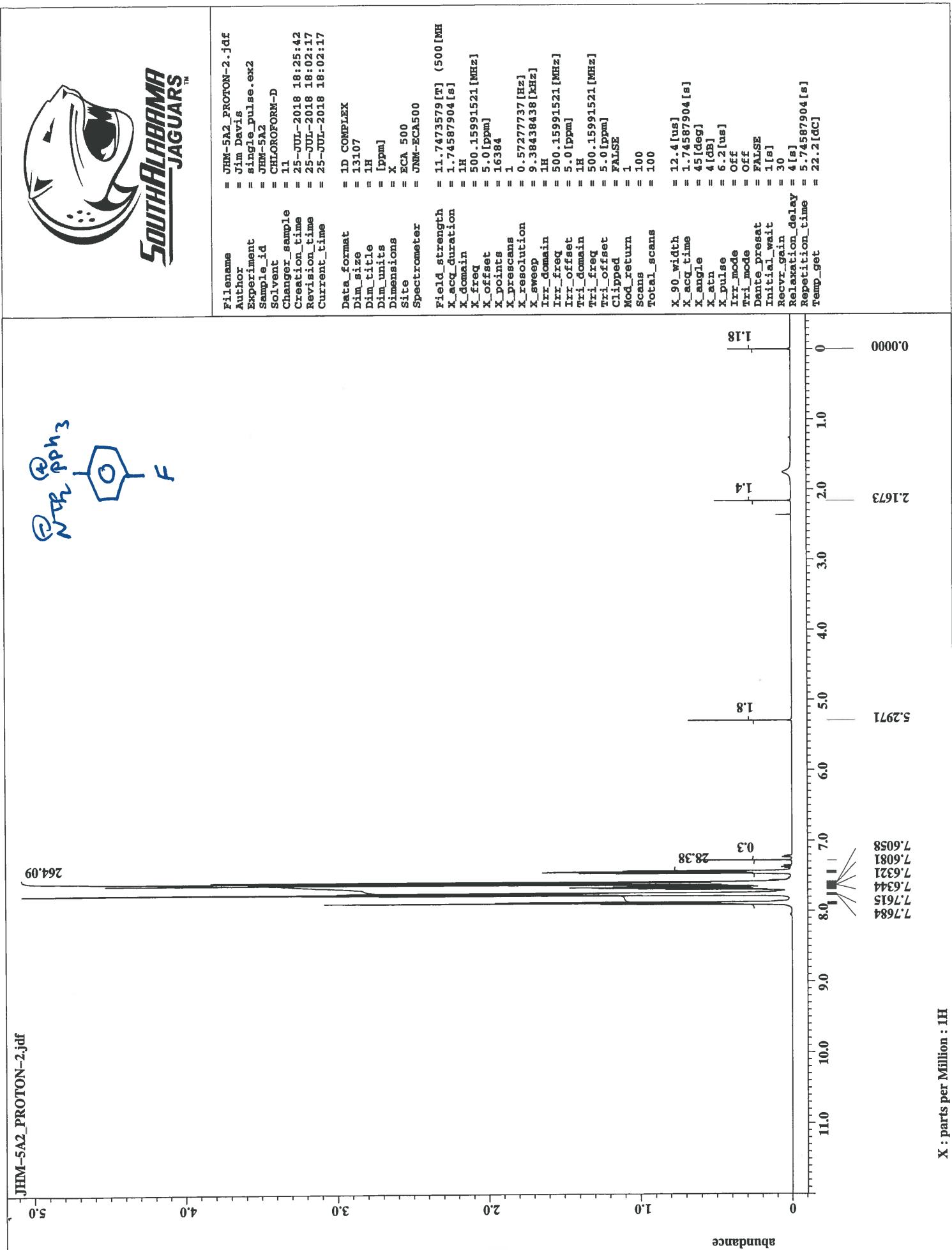


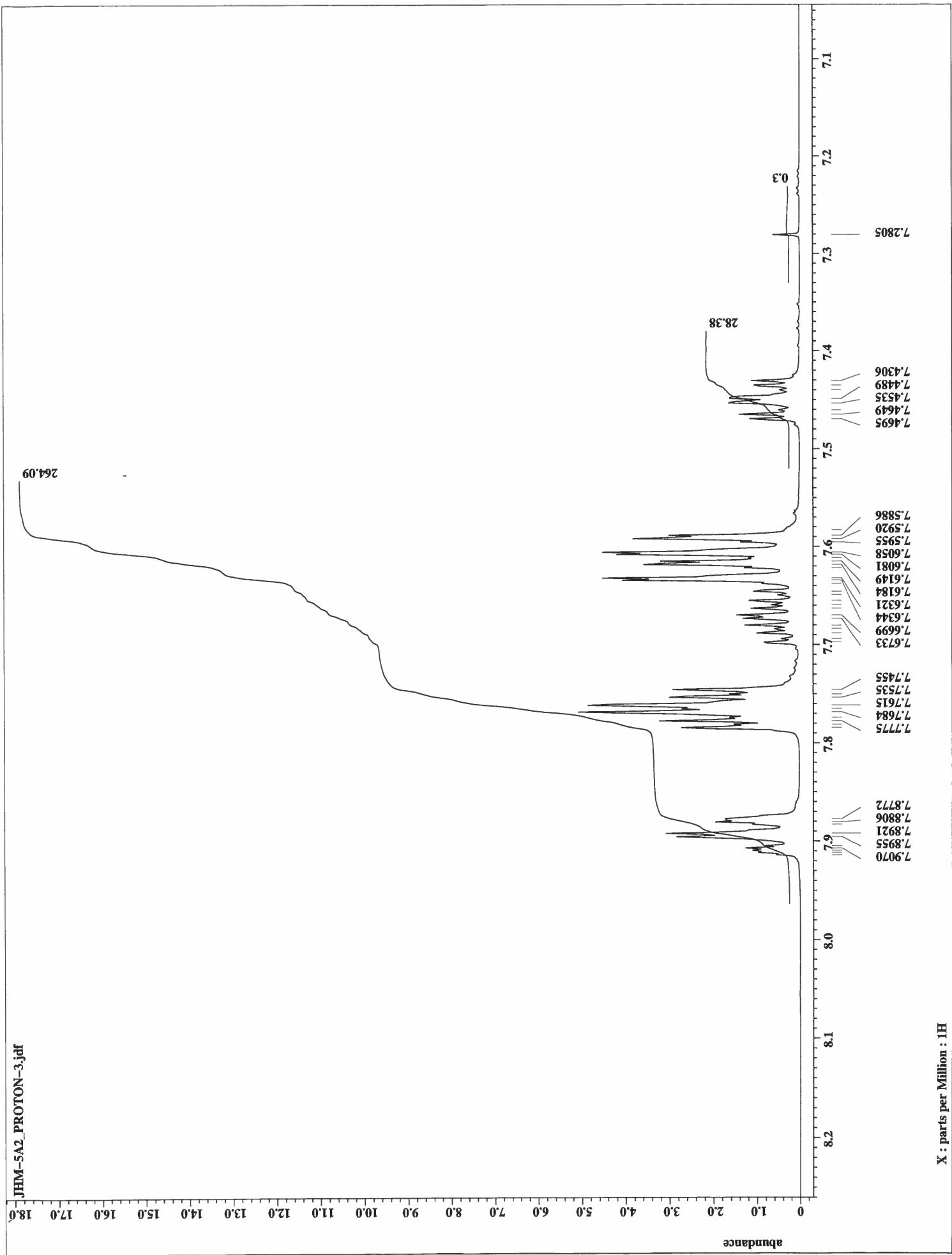














```

filename = JHM-5A2_CARBON-2.jdf
author = Jim Davis
experiment = single_pulse_dec
sample_id = JHM-5A2
solvent = CHLOROFORM-D
changer_sample = 11
creation_time = 25-JUL-2018 19:05:09
revision_time = 25-JUL-2018 18:45:43
current_time = 25-JUL-2018 18:45:43

data_format = 1D COMPLEX
dim_size = 2614
dim_title = 13C
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = JNM-ECA500

field_strength = 11.7473379 [T] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain = 13C
X_freq = 128.76529768 [MHz]
X_offset = 100.0 [ppm]
X_points = 32768
X_prescans = 4
X_resolution = 1.19959034 [Hz]
X_sweep = 39.3081761 [Hz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 300
Total_scans = 300

X_90_width = 13.2 [us]
X_acq_time = 0.83361792 [s]
X_angle = 30 [deg]
X_stn = 6 [dB]
X_pulse = 4.4 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Recur_grain = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get = 22.9 [dc]

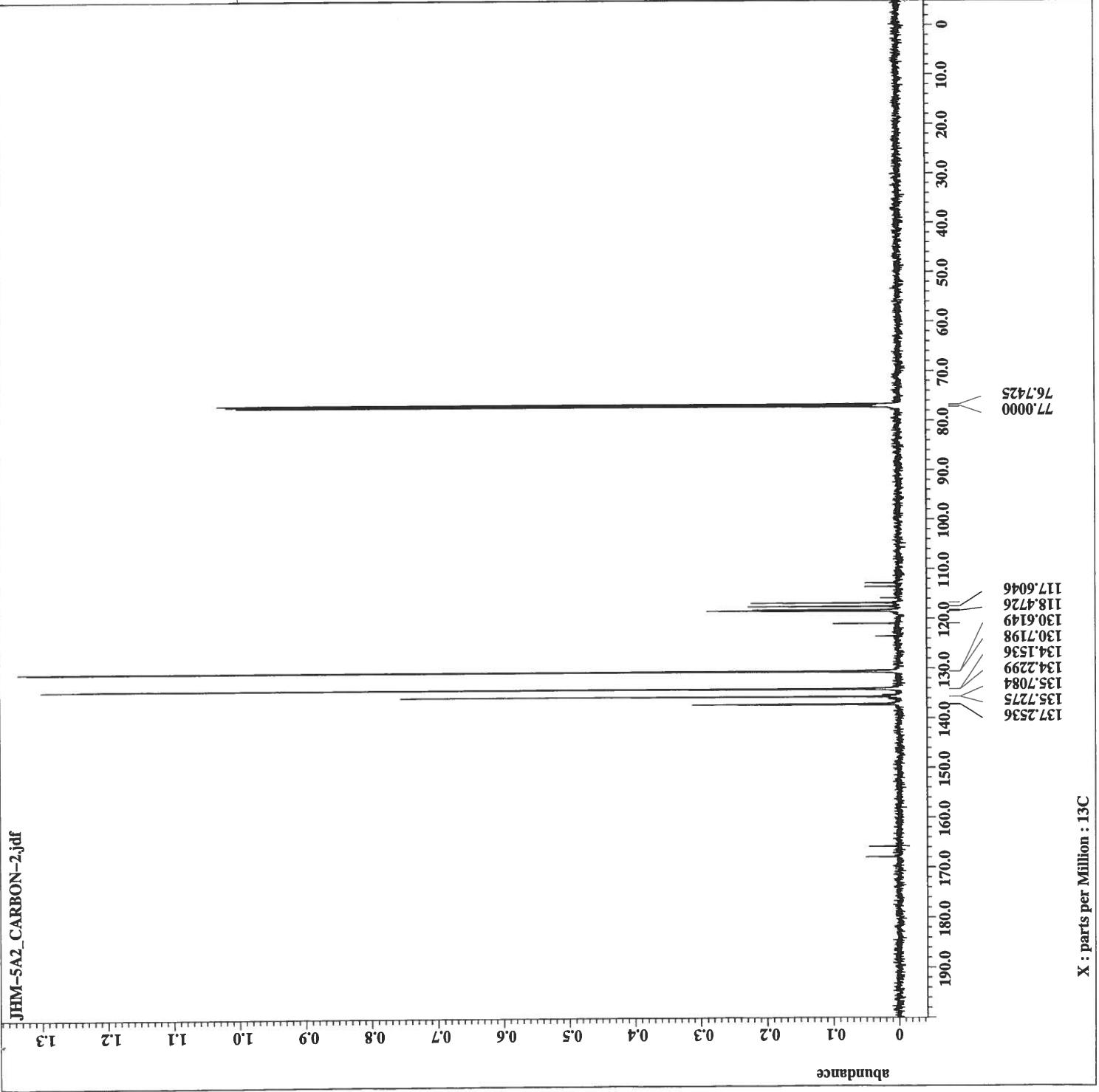
```

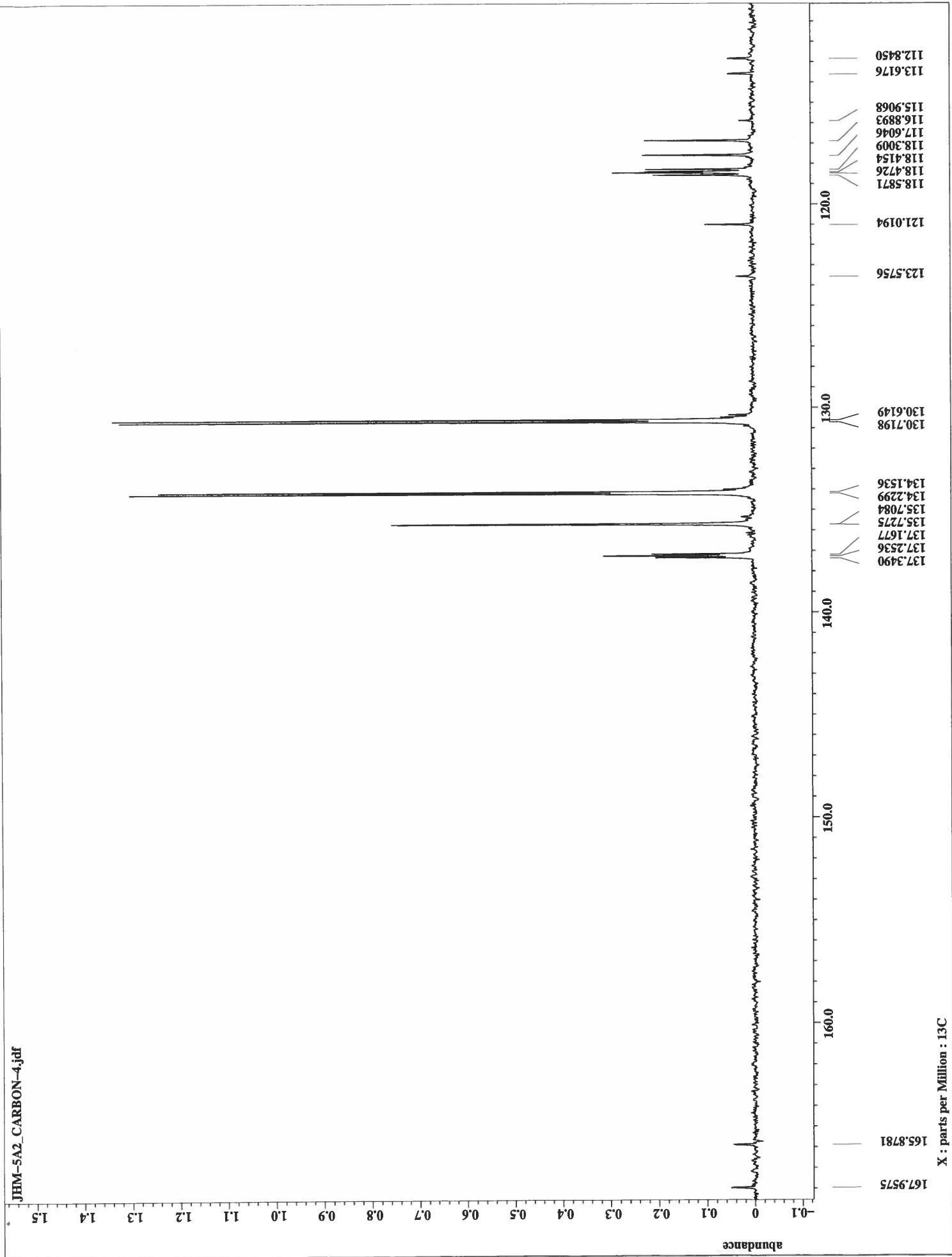
77.0000

76.7425

137.2536
 135.7275
 135.7084
 134.2299
 134.1536
 130.6149
 118.4726
 117.6066

X : parts per Million : 13C







```

Filename = JHM-5A2_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = JHM-5A2
Solvent = CHLOROFORM-D
Changer_sample = 11
Creation_time = 25-JUL-2018 18:14:32
Revision_time = 25-JUL-2018 17:51:07
Current_time = 25-JUL-2018 17:51:07

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JHM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [kHz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Racvr_gain = 38
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.5 [dc]

30.0 20.0 10.0 0 -10.0 -20.0 -30.0 -40.0 -50.0 -60.0 -70.0 -80.0 -90.0 -100.0 -110.0 -120.0 -130.0 -140.0 -150.0 -160.0 -170.0 -180.0 -190.0 -200.0 -210.0 -220.0 -230.0 -240.0 -250.0 -260.0 -270.0 -280.0 -290.0 -300.0

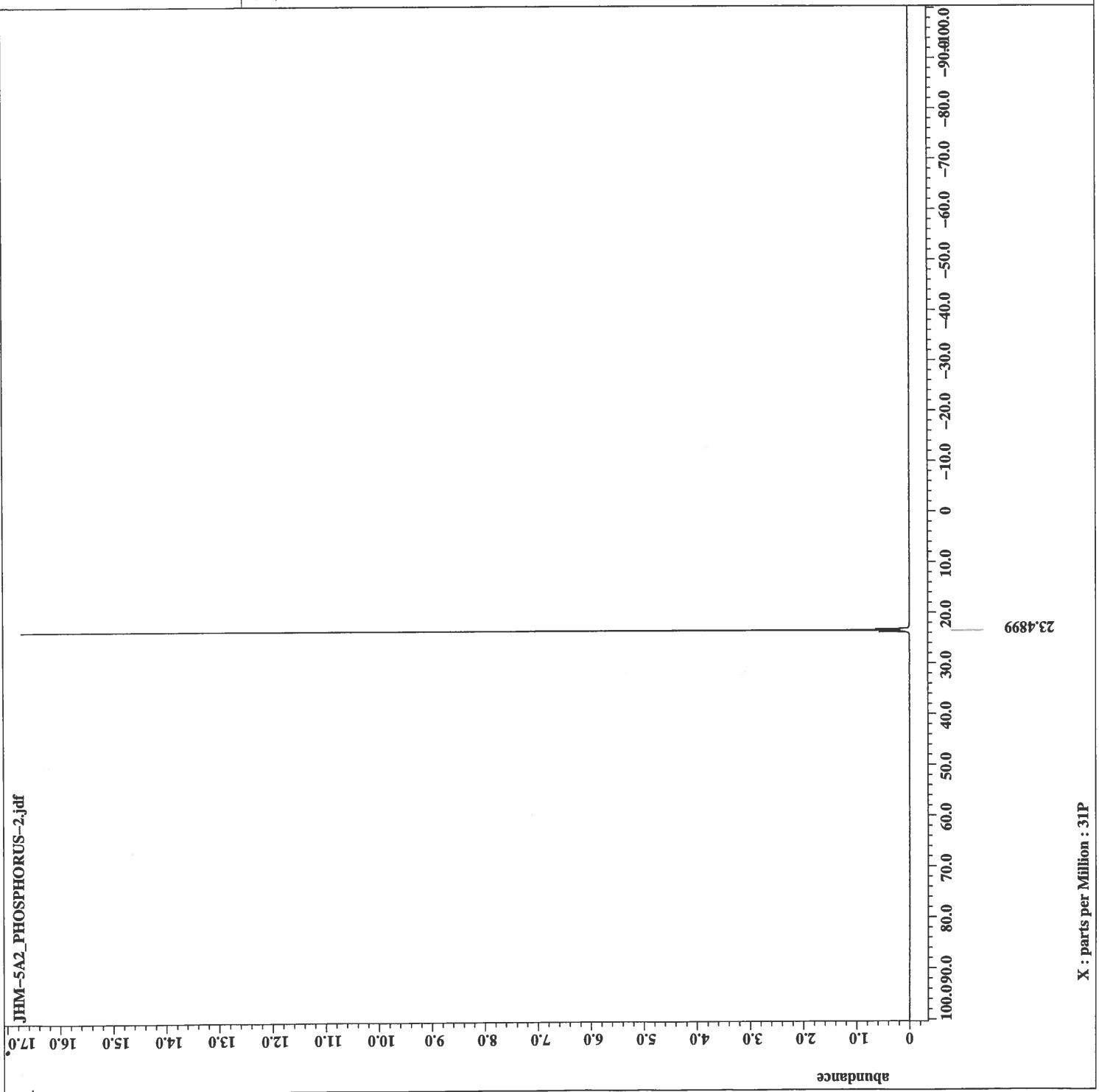
```

abundance

-98.8975

-78.9989

X : parts per Million : 19F



JHM-5A2_PHOSPHORUS-2.jdf



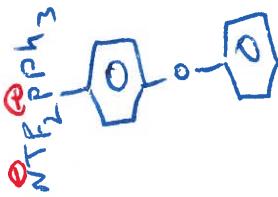
```

Filename = JHM-5A2_PHOSPHORUS-2.
Author = Jim Davis
Experiment =
Sample_id = Single_pulse_dec
Solvent = JHM-5A2
Solvant = CHLOROFORM-D
Solventer = JHM-ECA500
Sample = 11
Creation_time = 25-JUL-2018 18:38:23
Revision_time = 25-JUL-2018 18:14:57
Current_time = 25-JUL-2018 18:14:57
Data_format = 1D COMPLEX
Dim_size = 6214
Dim_title = 31P
Dim_units = [Dppm]
Dimensions = X
Site = ECA 500
Spectrometer = JHM-ECA500
Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.64487424 [s]
X_domain = 31P
X_fred = 202.46831075 [MHz]
X_offset = 0 [ppm]
X_points = 32768
X_precsans = 4
X_resolution = 1.55068995 [Hz]
X_sweep = 50.81300813 [Hz]
X_swep = 1H
Irr_domain = 0.15991521 [MHz]
Irr_freq = 50.0 [ppm]
Irr_offset = FALSE
Clipped = 1
Mod_return = 1
Scans = 234
Total_scans = 234
X_90_width = 14.687 [us]
X_acq_time = 0.64487424 [s]
X_angle = 30 [deg]
X_atn = 5 [dB]
X_pulse = 4.89566667 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Pecur.Gain = 56
Relaxation_delay = 2 [s]
Repetition_time = 2.64487424 [s]
Item_Set = 2,8,14,1

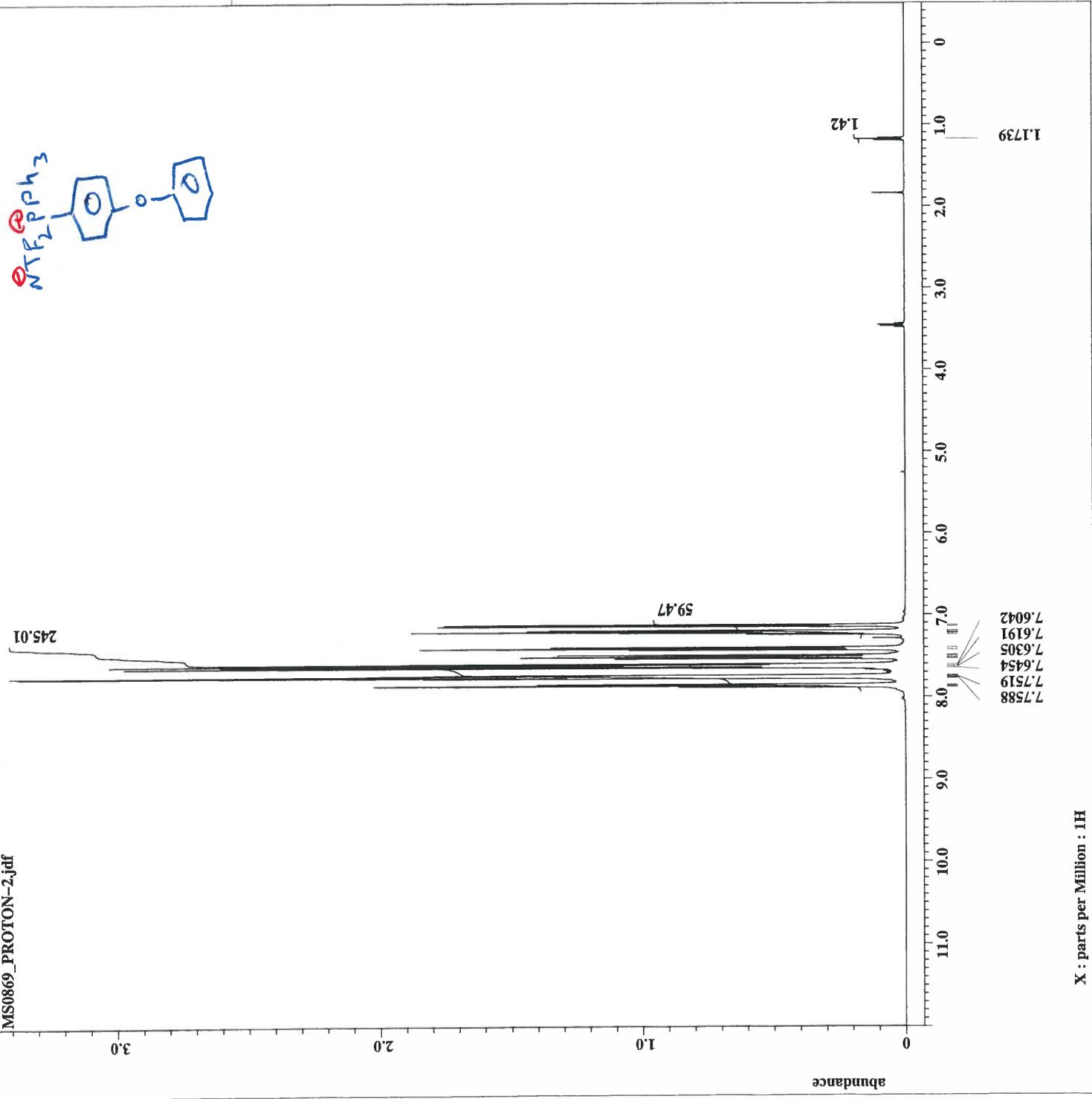
```

23.4899

X : parts per Million : 31P



245.01



```

Filename = MS0869_PROTON-2.jdf
Author = Jim Davis
Experiment = single_pulse_ex2
Sample_id = MS0869
Solvent = CHLOROFORM-D
Creation_time = 26-NOV-2019 11:19:03
Revision_time = 26-NOV-2019 10:49:59
Current_time = 26-NOV-2019 10:49:59

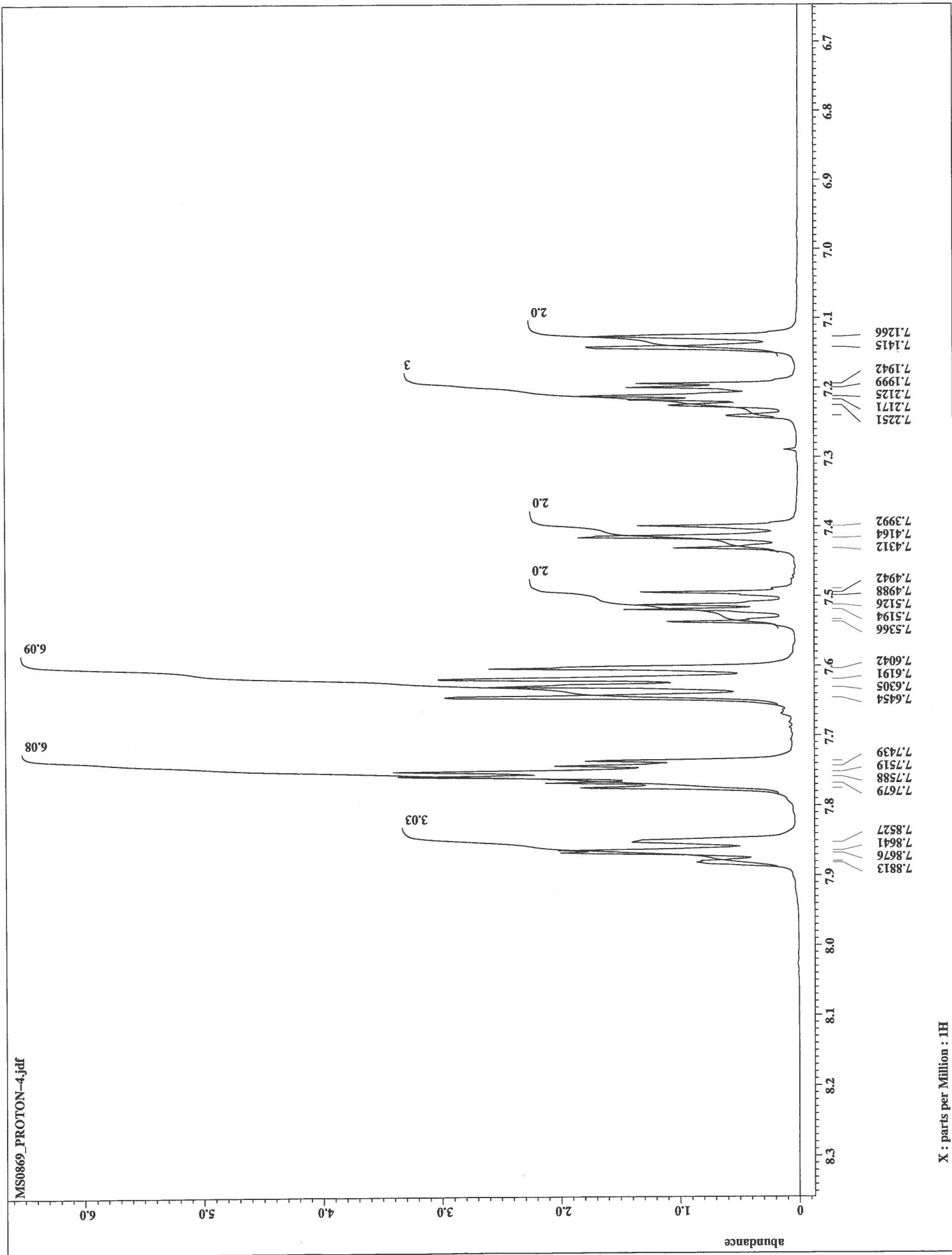
Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 1.74587904 [s]
X_domain = 1H
X_freq = 500.15991521 [MHz]
X_offset = 5.0 [ppm]
X_points = 16384
X_prescans = 1
X_resolution = 0.57277737 [Hz]
X_sweep = 9.38438438 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
total_scans = 16

X_90_width = 11.3 [us]
X_acq_time = 1.74587904 [s]
X_angle = 45 [deg]
X_atn = 4 [dB]
X_pulse = 5.65 [us]
Irr_mode = Off
Tri_mode = Off
Dante_presat = FALSE
Initial_wait = 1 [s]
Recv_gain = 20
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get = 19.3 [dc]

```

7.7588
7.6942
7.6391
7.6191
7.6541
7.7519





```

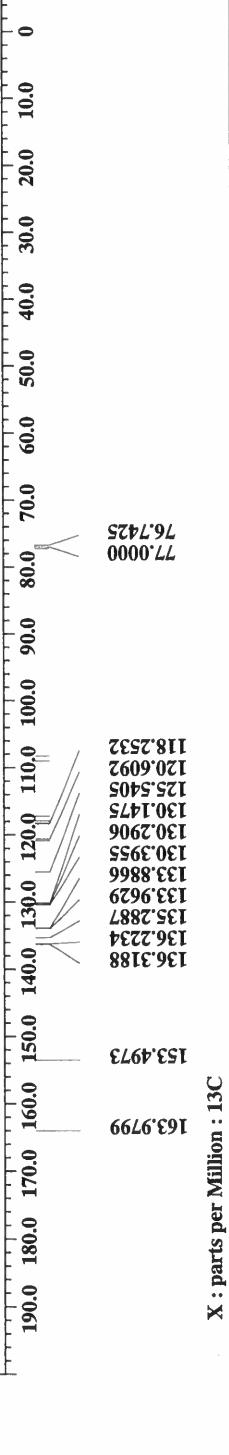
Filename = MS0869_CARBON-2.jdf
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0869
Solvent = CHLOROFORM-D
Creation_time = 26-NOV-2019 11:56:57
Revision_time = 26-NOV-2019 11:27:52
Current_time = 26-NOV-2019 11:27:53

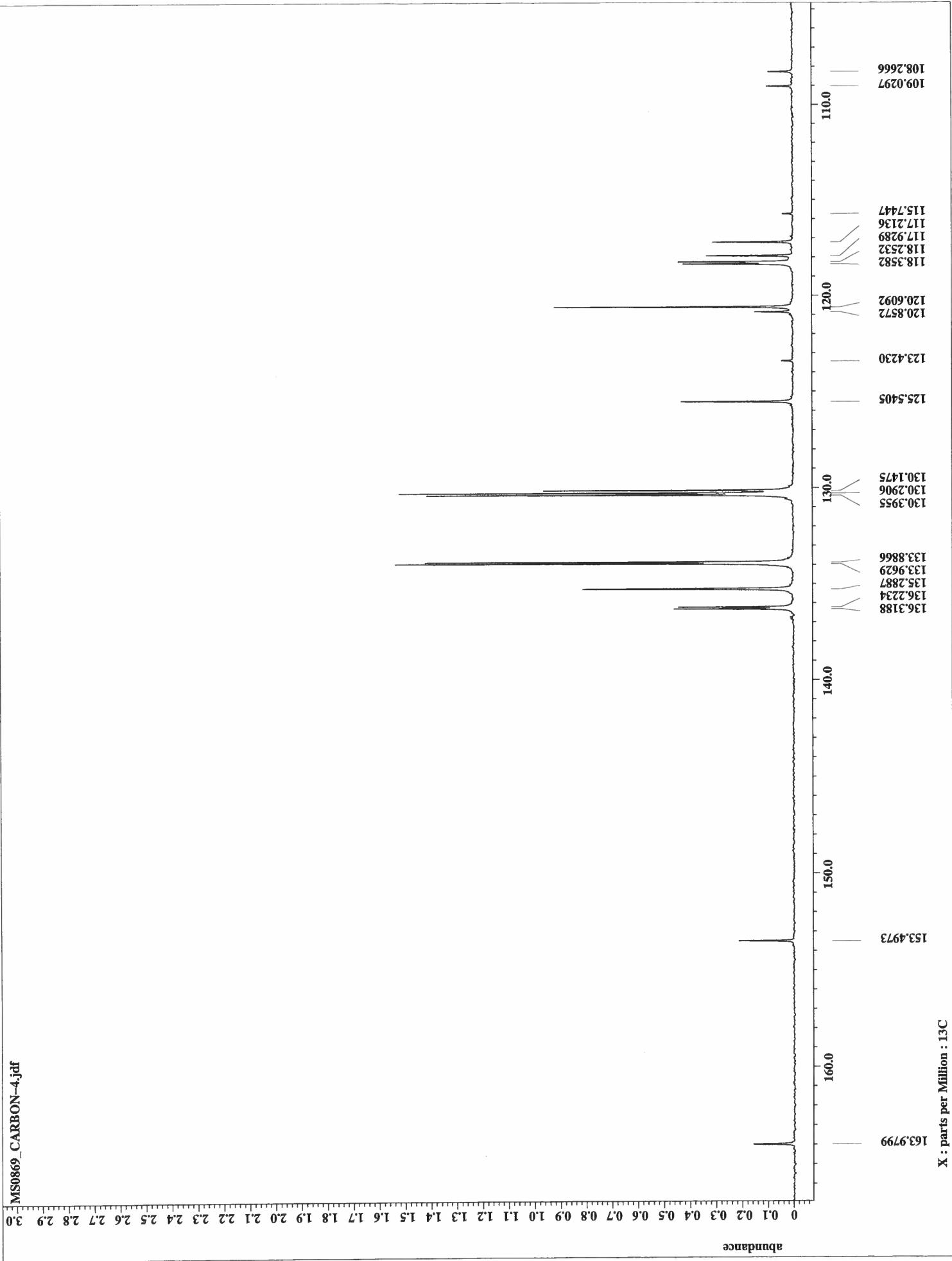
Data_format = 1D COMPLEX
Dim_size = 26214
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

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 = 39.3081751 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 449
Total_scans = 449

X_90_width = 13 [us]
X_acq_time = 0.83361792 [s]
X_angle = 30 [deg]
X_atn = 6 [dB]
X_pulse = 4.33333333 [us]
Irr_atn_dec = 21.2 [dB]
Irr_atn_noe = 21.2 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Recv_gain = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get = 18.8 [dC]

```





30.0

20.0

10.0

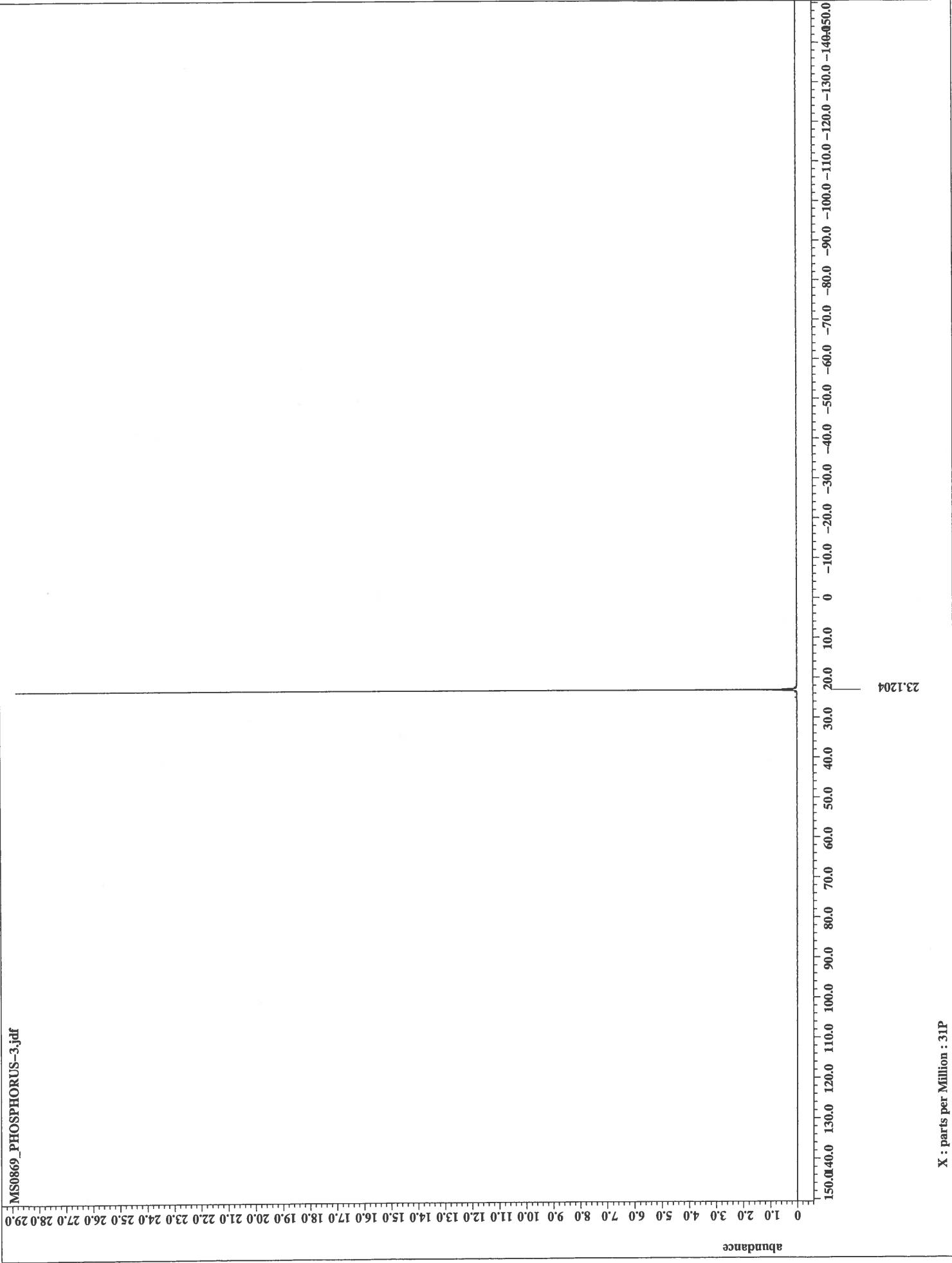
0

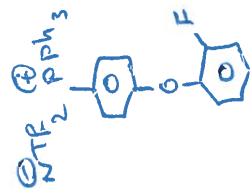
abundance

X : parts per Million : 19F

-78.8181

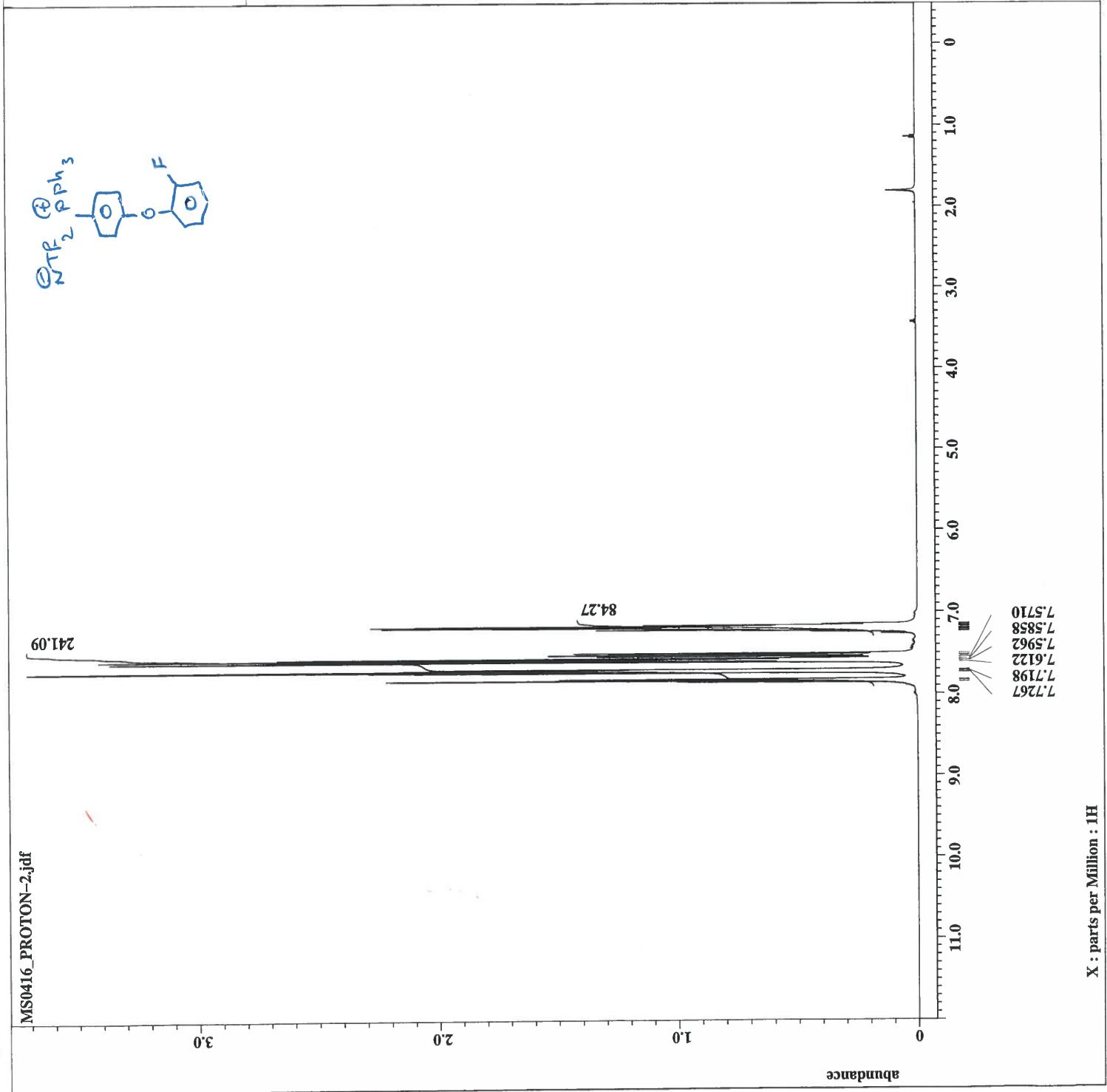
50.0 40.0 30.0 20.0 10.0 0 -10.0 -20.0 -30.0 -40.0 -50.0 -60.0 -70.0 -80.0 -90.0 -100.0 -110.0 -120.0 -130.0 -140.0 -150.0 -160.0 -170.0 -180.0 -190.0 -200.0 -210.0 -220.0 -230.0 -240.0 250.0

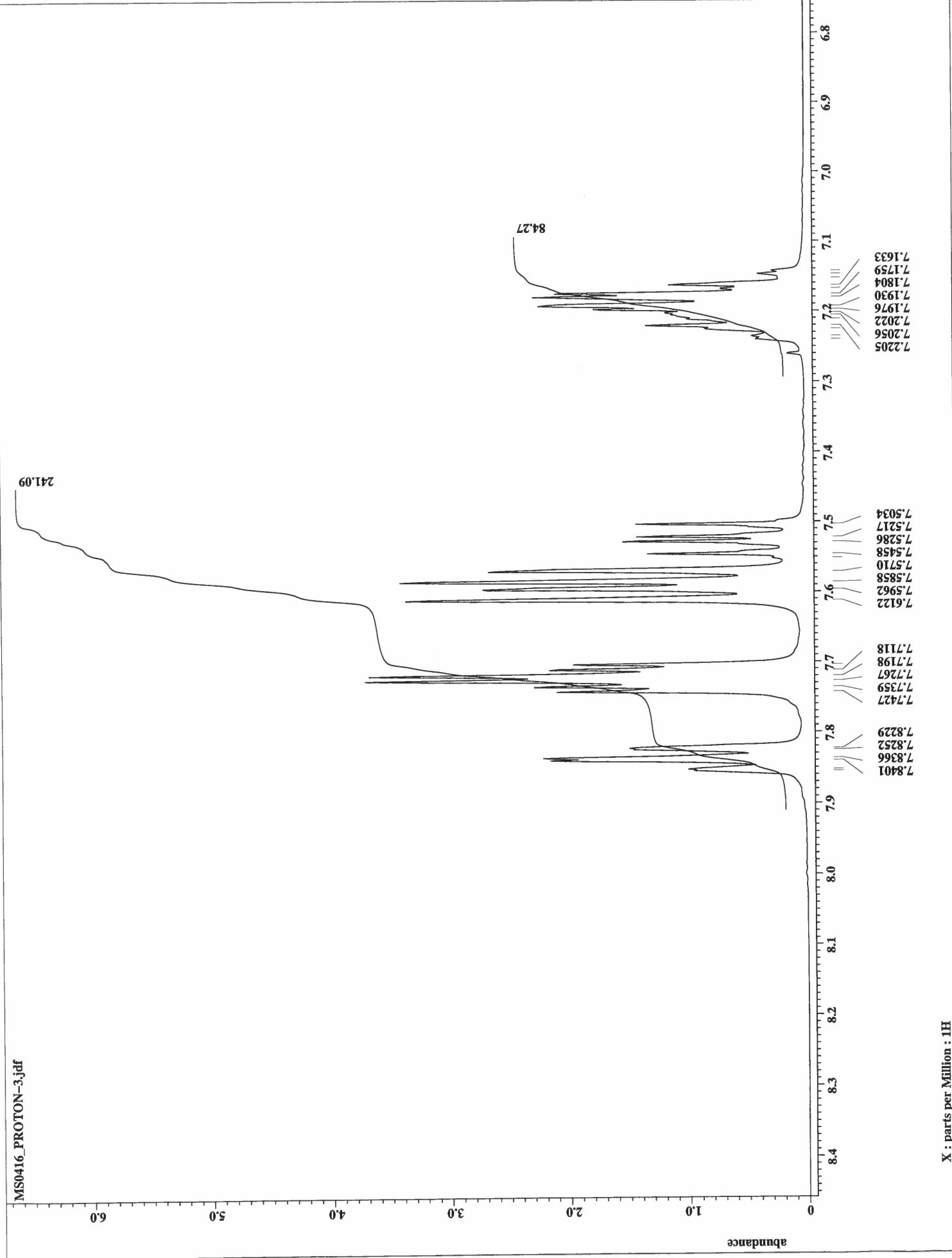




241.09

84.27







```

Filename          = MS0416_CARBON-2.jdf
Author           = Jim Davis
Experiment       = single_pulse_dec
Sample_id        = MS0416
Solvent          = CHLOROFORM-D
Changer_sample  = 10
Creation_time   = 14-APR-2018 11:47:07
Revision_time   = 14-APR-2018 11:45:15
Current_time    = 14-APR-2018 11:25:15

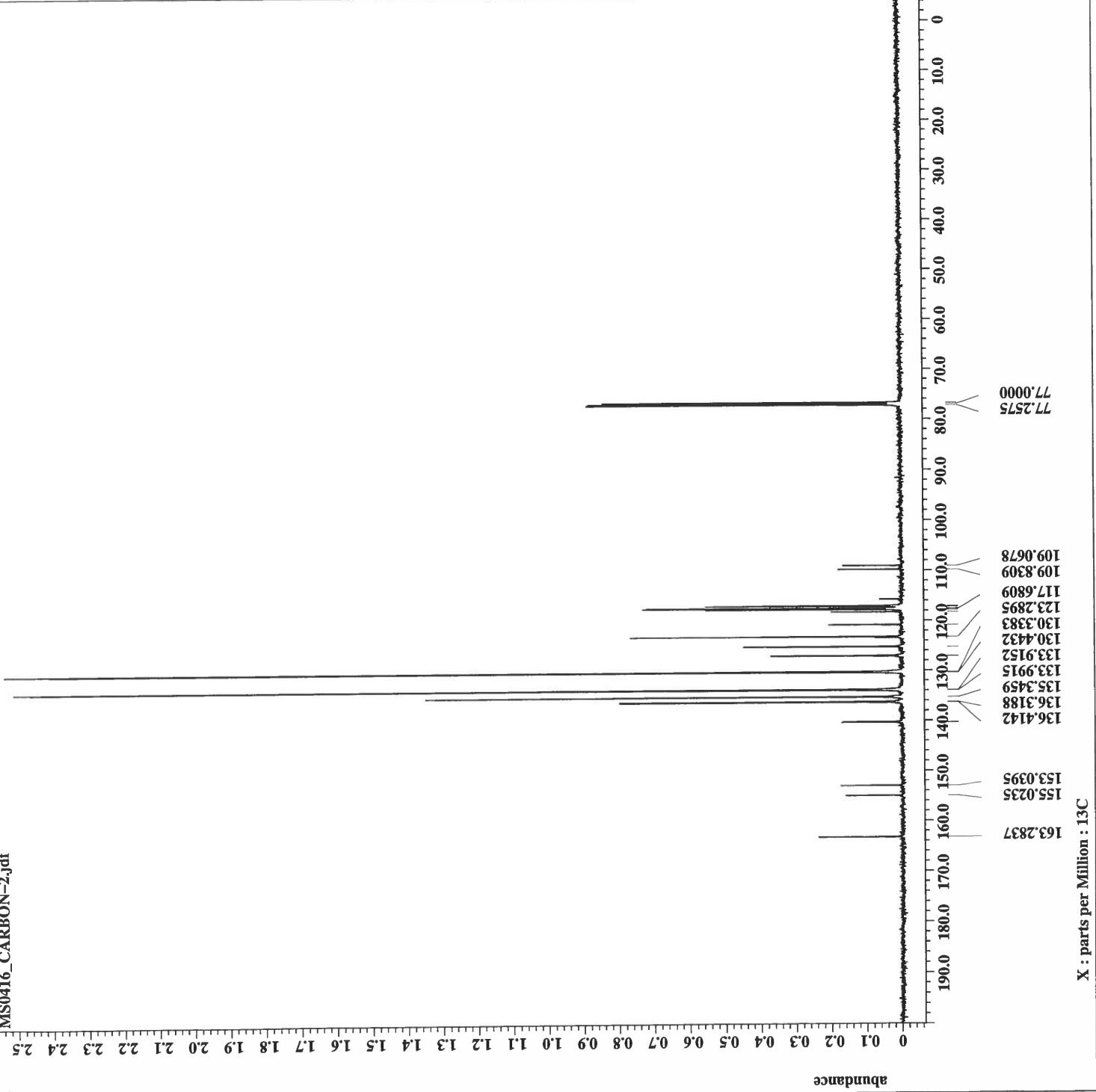
Data_format     = 1D COMPLEX
Dim_size        = 26214
Dim_title       = 13C
Dim_units       = [ppm]
Dimensions      = X
Site            = ECA 500
Spectrometer    = JNM-ECA500

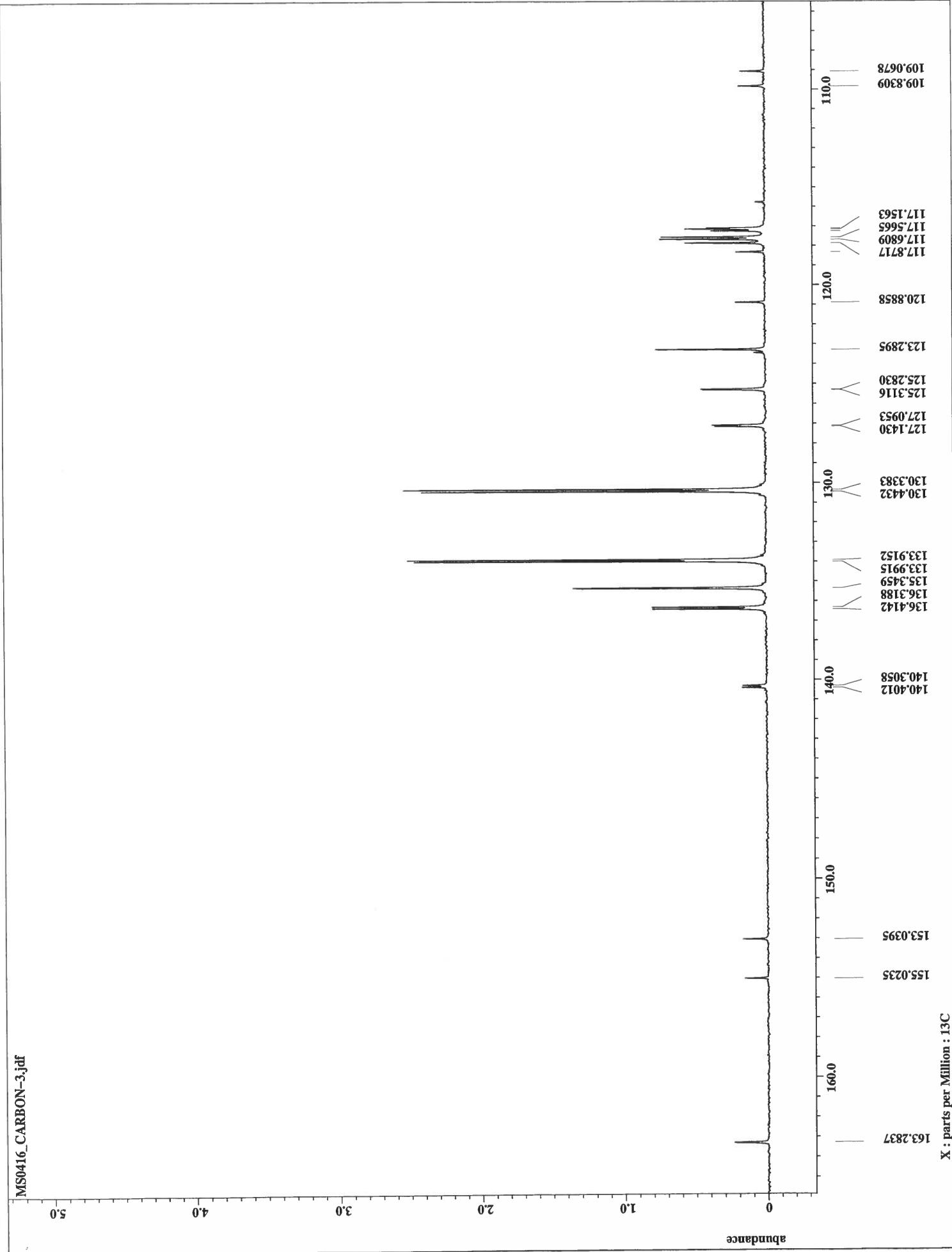
Field_strength  = 11.7473579 [Hz] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain        = 13C
X_freq          = 128.76529768 [MHz]
X_offset        = 100 [ppm]
X_points        = 32768
X_prescans     = 4
X_resolution   = 1.19959034 [Hz]
X_sweep         = 39.3082761 [Hz]
Irr_domain     = 1H
Irr_freq        = 500.15991521 [MHz]
Irr_offset      = 5.0 [ppm]
Clipped         = FALSE
Mod_return     = 1
Scans           = 256
total_scans    = 256

X_90_width     = 13.2 [us]
X_acq_time     = 0.83361792 [s]
X_angle         = 30 [deg]
X_atn          = 6 [dB]
X_pulse         = 4.4 [us]
Irr_atn_dec    = 20.7 [dB]
Irr_atn_noe    = 20.7 [dB]
Irr_noise       = WALTZ
Decoupling     = TRUE
Initial_wait   = 1 [s]
Noe            = TRUE
Noe_time        = 2 [s]
Regrv_gain     = 60
Relaxation_delay = 2 [s]
Relaxation_time = 2.83361792 [s]
Temp_get        = 23.21 [dc]

X : parts per Million : 13C

```







```

Filename = MS0416_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0416
Solvent = CHLOROFORM-D
Changer_sample = 10
Creation_time = 14-APR-2018 11:25:31
Revision_time = 14-APR-2018 11:03:39
Current_time = 14-APR-2018 11:03:39

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X.acq_duration = 0.55574528 [s]
X.domain = 19F
X.freq = 470.62046084 [MHz]
X.offset = -70 [ppm]
X.points = 65536
X.precsans =
X.resolution =
X.sweep =
Irr_domain = 117.9245283 [kHz]
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
total_scans = 16

X.90_width = 13.1 [us]
X.acq_time = 0.55574528 [s]
X.angle = 45 [deg]
X.atn =
X.pulse =
Irr_mode = Off
Dante_presat = FALSE
Initial_wait = 1 [s]
Reacr_grain = 34
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.6 [dc]

```

-129.3244

-78.6218

X : parts per Million : 19F

30.0

20.0

10.0

0

abundance



```

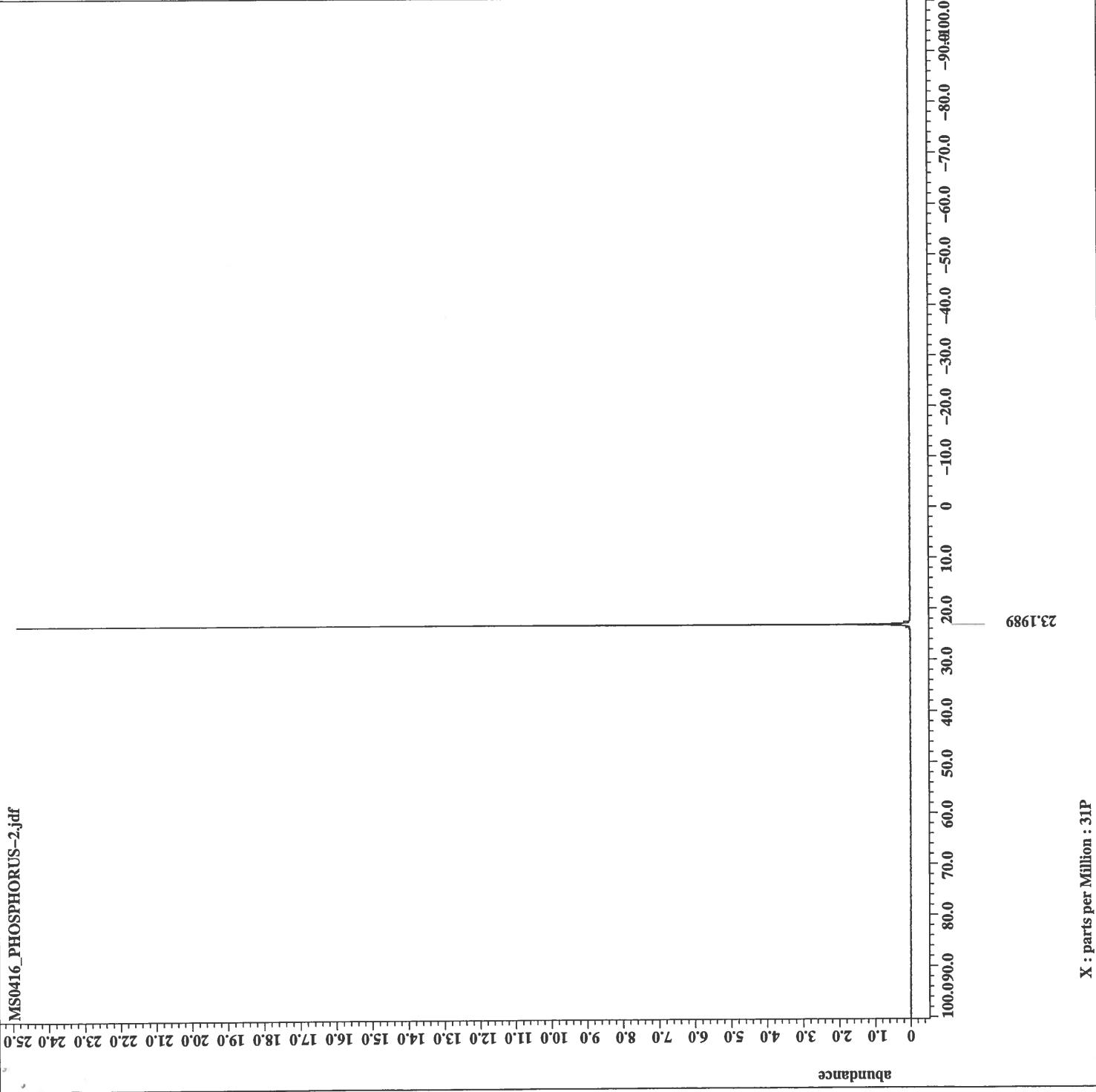
Filename = MS0416_PHOSPHORUS-2.j
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0416
Solvent = CHLOROFORM-D
Changer_sample = 10
Creation_sample = 14-APR-2018 11:29:07
Revision_time = 14-APR-2018 11:07:15
Current_time = 14-APR-2018 11:07:15

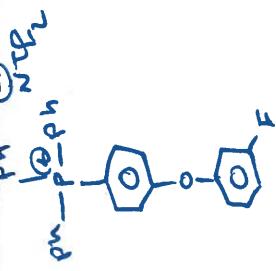
Data_format = 1D COMPLEX
Dim_size = 2614
Dim_title = 31P
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JEOL-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.64487424 [s]
X_domain = 31P
X_freq = 202.46831075 [MHz]
X_offset = 0 [ppm]
X_points = 32768
X_prescans = 4
X_resolution = 1.55068995 [Hz]
X_sweep = 50.81300813 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 25
Total_scans = 25

X_90_width = 14.687 [us]
X_acq_time = 0.64487424 [s]
X_angle = 30 [deg]
X_atn = 5 [dB]
X_pulse = 4.99566667 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Revr_gain = 54
Relaxation_delay = 2 [s]
Repetition_time = 2.64487424 [s]
Temp_get = 23 [dc]

```





SOUTH ALABAMA JAGUARS™

```

Filename      = MS0446_PROTON-2.jdf
Author        = Jim Davis
Experiment   = single_pulse.ex2
Sample_id    = MS0446
Solvent       = CHLOROFORM-D
Changer_sample = 1
Creation_time = 14-MAY-2018 12:32:44
Revision_time = 14-MAY-2018 12:08:31
Current_time  = 14-MAY-2018 12:08:31

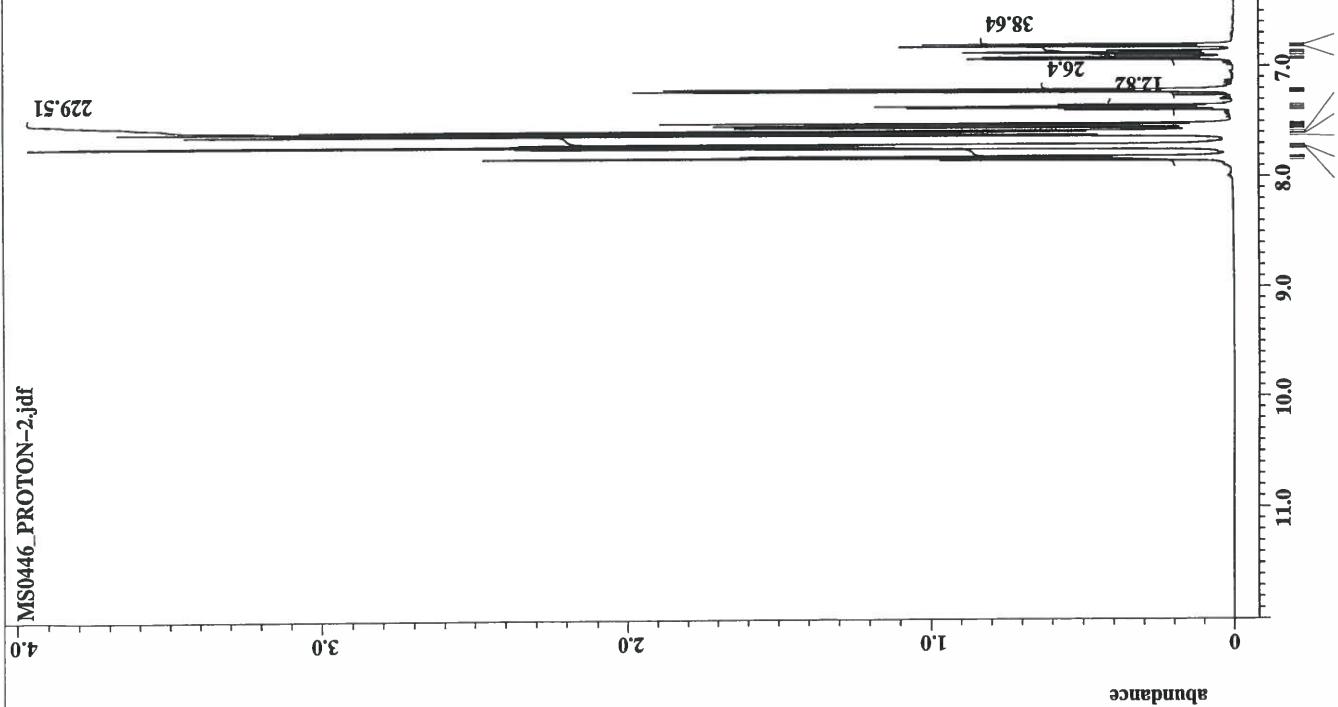
Data_format   = 1D COMPLEX
Dim_size      = 13107
Dim_title    = 1H
Dim_units    = [ppm]
Dimensions   = X
Site          = ECA 500
Spectrometer = JNM-ECA500

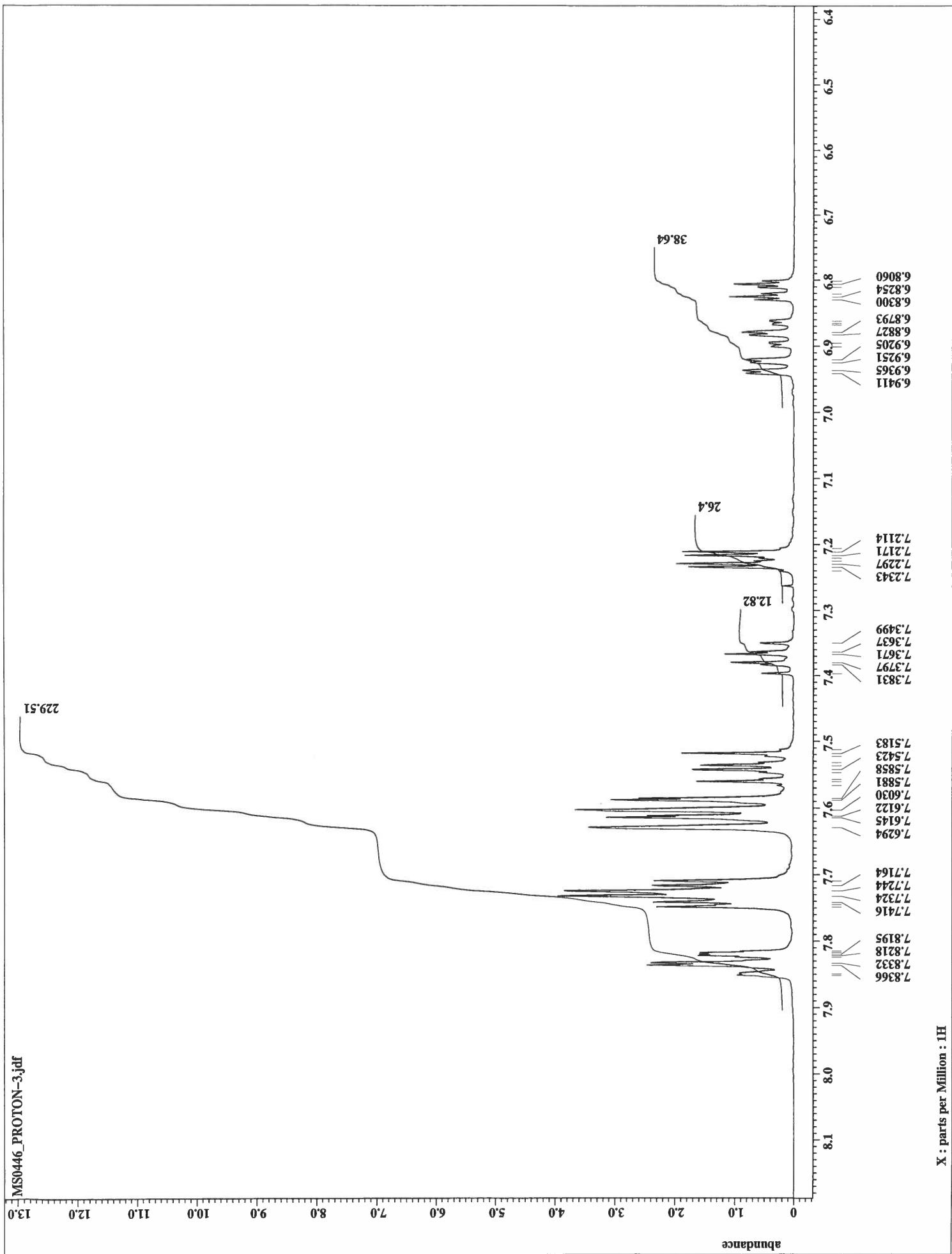
Field_strength = 11.7473579 [T] (500 [MHz])
X.acq_duration = 1.74587904 [s]
X.domain     = 1H
X.freq        = 500.15991521 [MHz]
X.offset      = 5.0 [ppm]
X.points      = 16384
X.precsans   = 1
X.resolution = 0.57277737 [Hz]
X.sweep       = 9.38438438 [Hz]
Irr_domain   = 1H
Irr_freq      = 500.15991521 [MHz]
Irr_offset    = 5.0 [ppm]
Irr_domain   = 1H
Irr_freq      = 500.15991521 [MHz]
Irr_offset    = 5.0 [ppm]
Clipped      = FALSE
Mod_return   = 1
Scans        = 16
total_scans  = 16

X.90_width   = 12.4 [us]
X.acq_time   = 1.74587904 [s]
X.angle       = 45 [deg]
X_tn          = 4 [dB]
X.pulse       = 6.2 [us]
Irr_mode     = Off
Tri_mode     = Off
Dante_presat = FALSE
Initial_wait = 1 [s]
Revr.gain     = 20
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get     = 22.8 [dC]

```

229.51







```

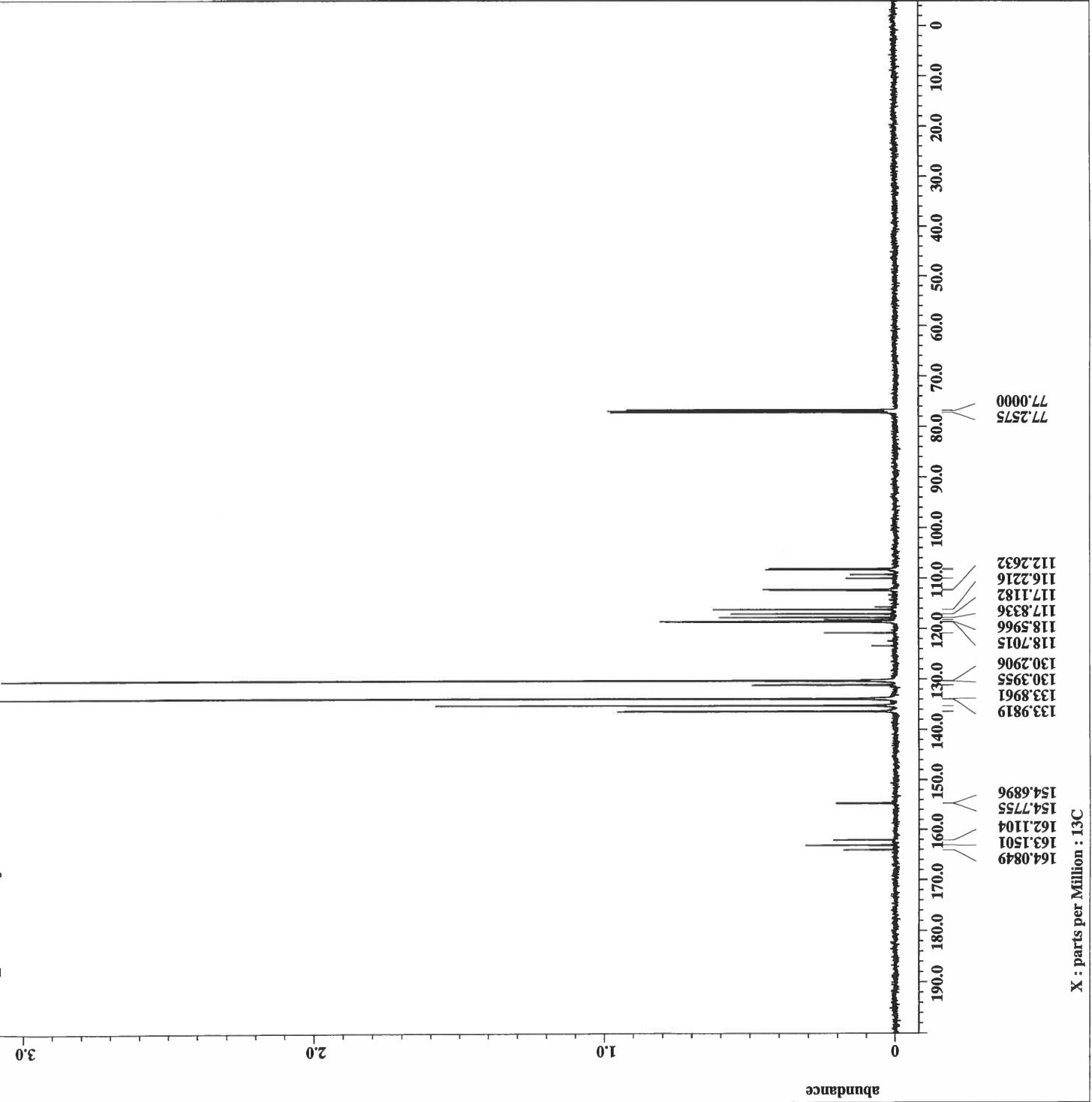
Filename      = MS0446_CARBON-2.jdf
Author        = Jim Davis
Experiment    = single_pulse_dec
Sample_id     = MS0446
Solvent       = CHLOROFORM-D
Changer_sample = 1
Creation_time = 14-MAY-2018 12:41:05
Revision_time = 14-MAY-2018 12:16:51
Current_time  = 14-MAY-2018 12:16:51

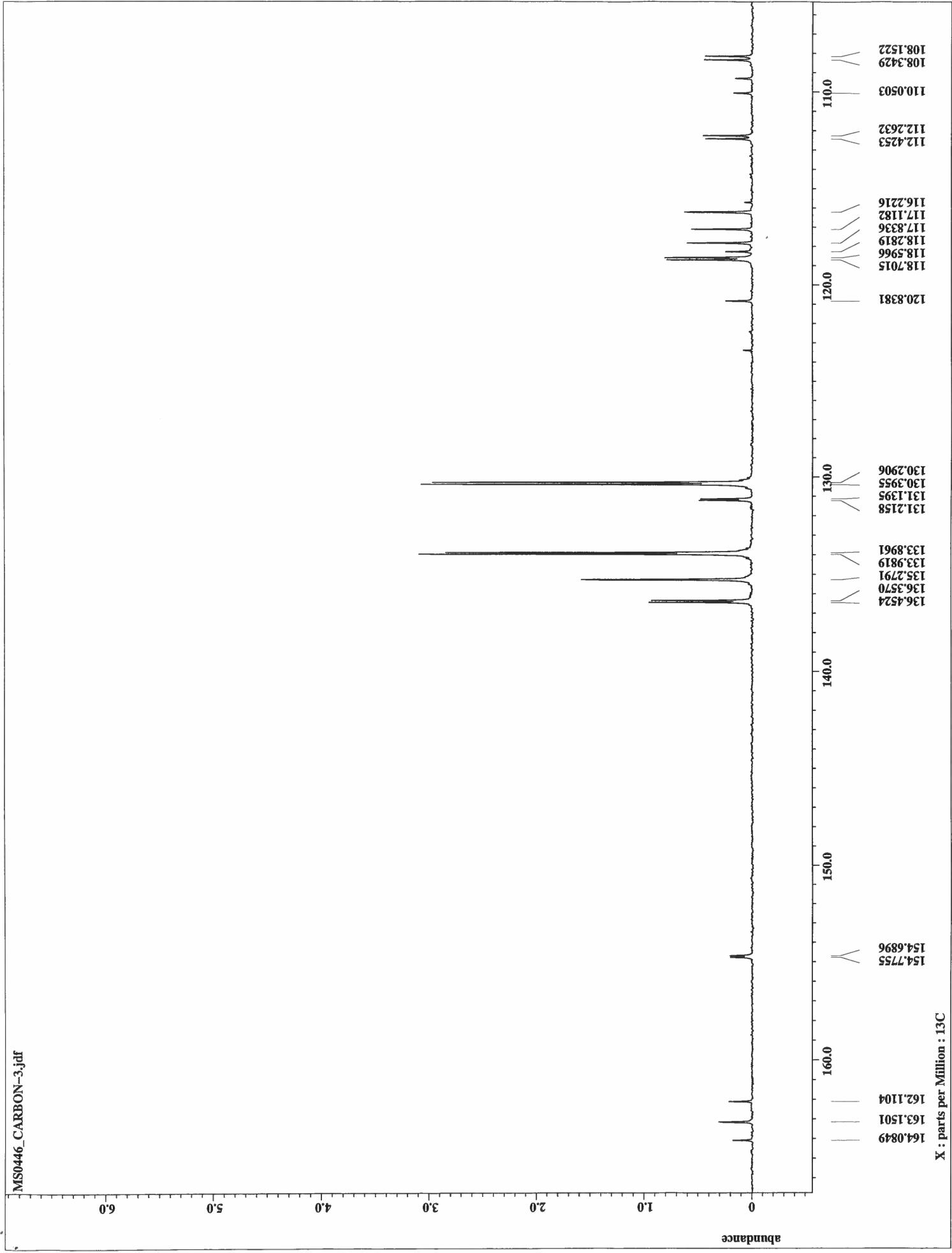
Data_format   = 1D COMPLEX
Dim_size      = 26214
Dim_title     = 13C
Dim_units     = [ppm]
Dimensions    = X
Site          = ECA 500
Spectrometer  = JNM-ECX500

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.precsans    = 4
X.resolution  = 1.19959034 [Hz]
X.sweep        = 39.3081761 [kHz]
Irr_domain    = 1H
Irr_freq       = 500.15991521 [MHz]
Irr_offset     = 5.0 [ppm]
Clipped        = FALSE
Mod_return    = 1
Scans          = 128
Total_scans   = 128

X_90_width   = 13.2 [us]
X.acq_time   = 0.83361792 [s]
X.angle       = 30 [deg]
X.atn         = 6 [dB]
X.pulse       = 4.4 [us]
Irr_atn_dec  = 20.7 [dB]
Irr_atn_noe  = 20.7 [dB]
Irr_noise     = WALTZ
Decoupling    = TRUE
Initial_wait  = 1 [s]
Noe           = TRUE
Noe_time      = 2 [s]
60
Recurr_grain = 2 [s]
Relaxation_delay = 2.33361792 [s]
Repetition_time = 22.8 [dC]
Temp_Set      = 22.8 [dC]

```







```

Filename      = MS0446_FLUORINE-2.jdf
Author        = Jim Davis
Experiment    = single_pulse.ex2
Sample_id     = MS0446
Solvent       = CHLOROFORM-D
Changer_sample = 1
Creation_time = 14-MAY-2018 12:25:44
Revision_time = 14-MAY-2018 12:01:32
Current_time  = 14-MAY-2018 12:01:32

Data_format   = 1D COMPLEX
Dim_size      = 52428
Dim_title     = 19F
Dim_units     = [ppm]
Dimensions    = X
Site          = ECA 500
Spectrometer  = JNM-ECX500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain      = 19F
X_freq         = 470.62046084 [MHz]
X_offset       = -70 [ppm]
X_points       = 65536
X_prescans    = 1
X_resolution  = 1.7993855 [Hz]
X_sweep        = 111.9245283 [Hz]
Irr_domain    = 19F
Irr_freq       = 470.62046084 [MHz]
Irr_offset     = 5 [ppm]
Irr_domain2   = 19F
Irr_freq2     = 470.62046084 [MHz]
Tri_offset    = 5 [ppm]
Clipped        = FALSE
Mod_return    = 1
scans         = 16
total_scans   = 16

X_90_width    = 13.1 [us]
X_acq_time    = 0.55574528 [s]
X_angle        = 45 [deg]
X_atn          = 2.5 [dB]
X_pulse        = 6.55 [us]
Irr_mode      = Off
Tri_mode      = Off
Dante_presat  = FALSE
Initial_wait   = 1 [s]
Revr_gain      = 30
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get      = 22.7 [dC]

```

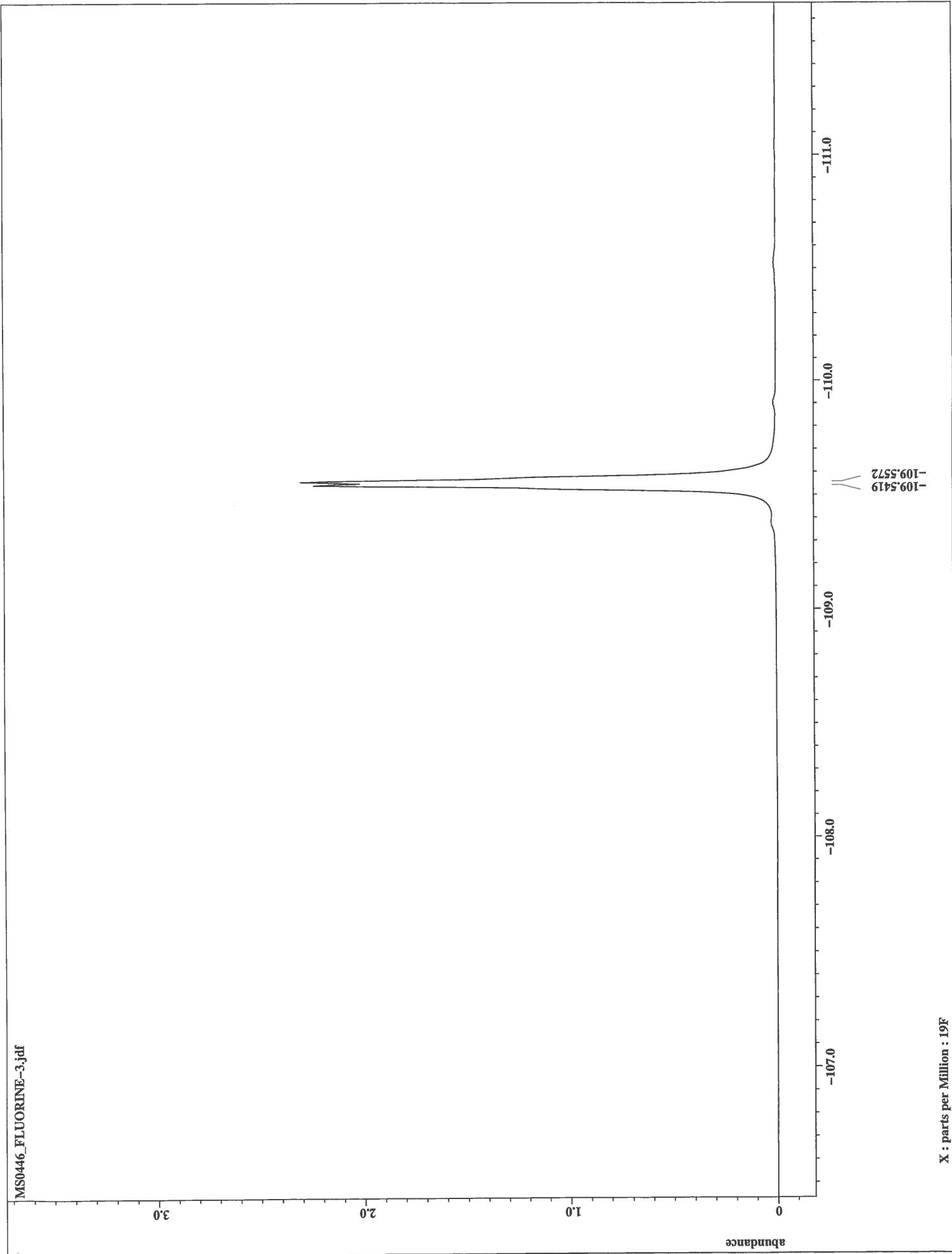
-109.5419 -109.5572

-78.6218

X : parts per Million : 19F



abundance





```

Filename = MS0446_PHOSPHORUS-2.j
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0446
Solvent = CHLOROFORM-D
Changer_sample = 1
Creation_time = 14-MAY-2018 12:29:19
Revision_time = 14-MAY-2018 12:05:05
Current_time = 14-MAY-2018 12:05:05

Data_format = 1D COMPLEX
Dim_size = 26214
Dim_title = 31P
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.644487424 [s]
X_domain = 31P
X_freq = 202.46831075 [MHz]
X_offset = 0 [ppm]
X_points = 32768
X_precsans = 4
X_resolution = 1.55068995 [Hz]
X_sweep = 50.8130081 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 25
Total_scans = 25

X_90_width = 14.687 [us]
X_acq_time = 0.644487424 [s]
X_angle = 30 [deg]
X_atn = 5 [dB]
X_pulse = 4.895666667 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Recvr_gain = 54
Relaxation_delay = 2 [s]
Repetition_time = 2.644487424 [s]
Temp_get = 23 [dC]

```

0 1.0 2.0 3.0 4.0 5.0 6.0 7.0 8.0 9.0 10.0 11.0 12.0 13.0 14.0 15.0 16.0 17.0 18.0 19.0 20.0 21.0 22.0 23.0 24.0 25.0 26.0 27.0

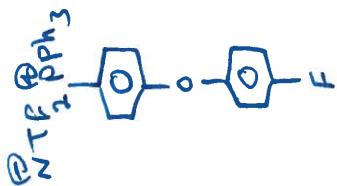
abundance

23.2219

X : parts per Million : 31P

MS0402 PROTON-5.jdf

215.53



```

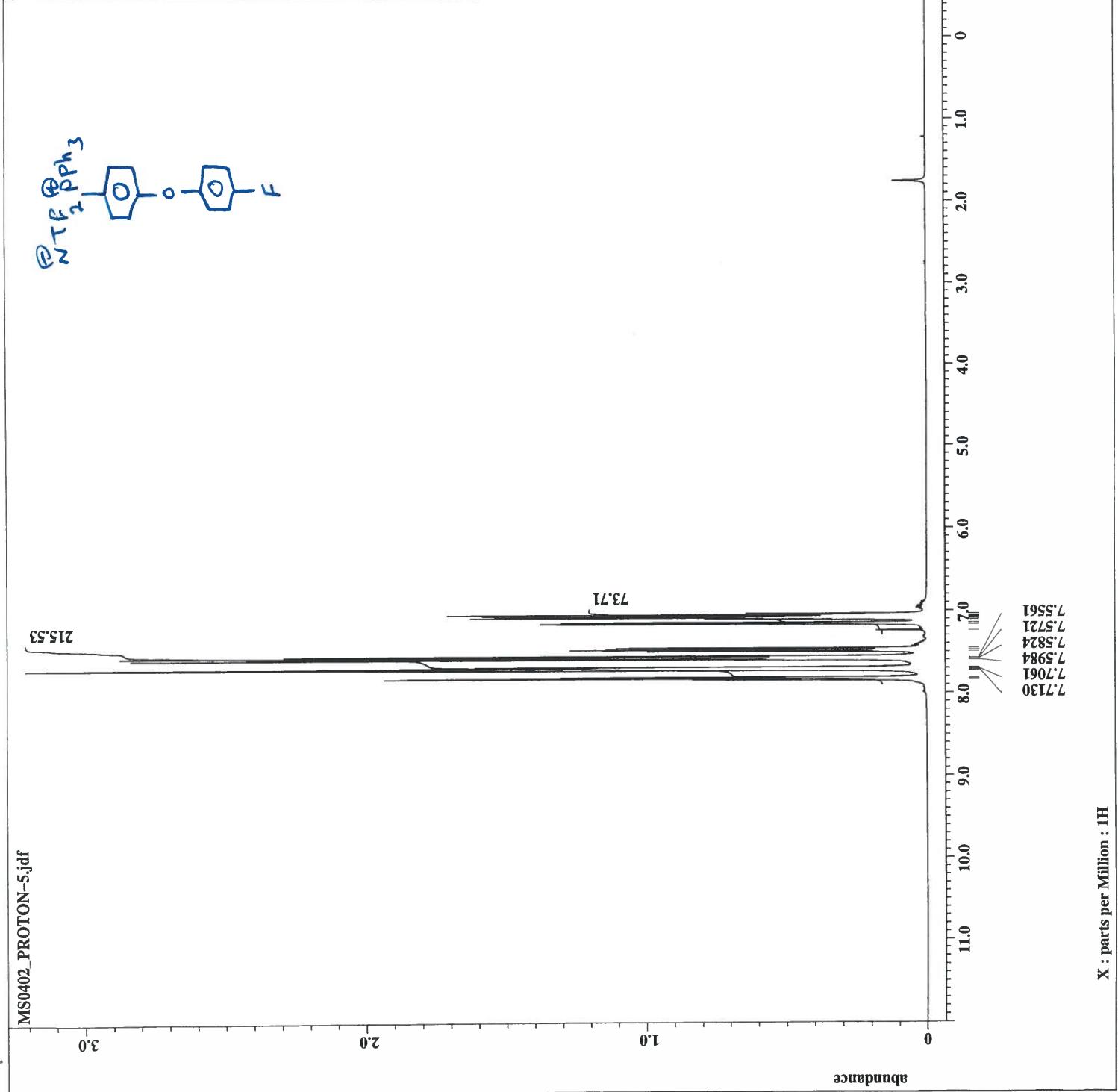
Filename      = MS0402_PROPON-5.jdf
Author        = Jim Davis
Experiment   = single_pulse_ex2
Sample_id    = MS0402
Solvent      = CHLOROFORM-D
Changer_sample = 6
Creation_time = 18-MAY-2018 16:54:40
Revision_time = 18-MAY-2018 16:30:05
Current_time = 18-MAY-2018 16:30:05

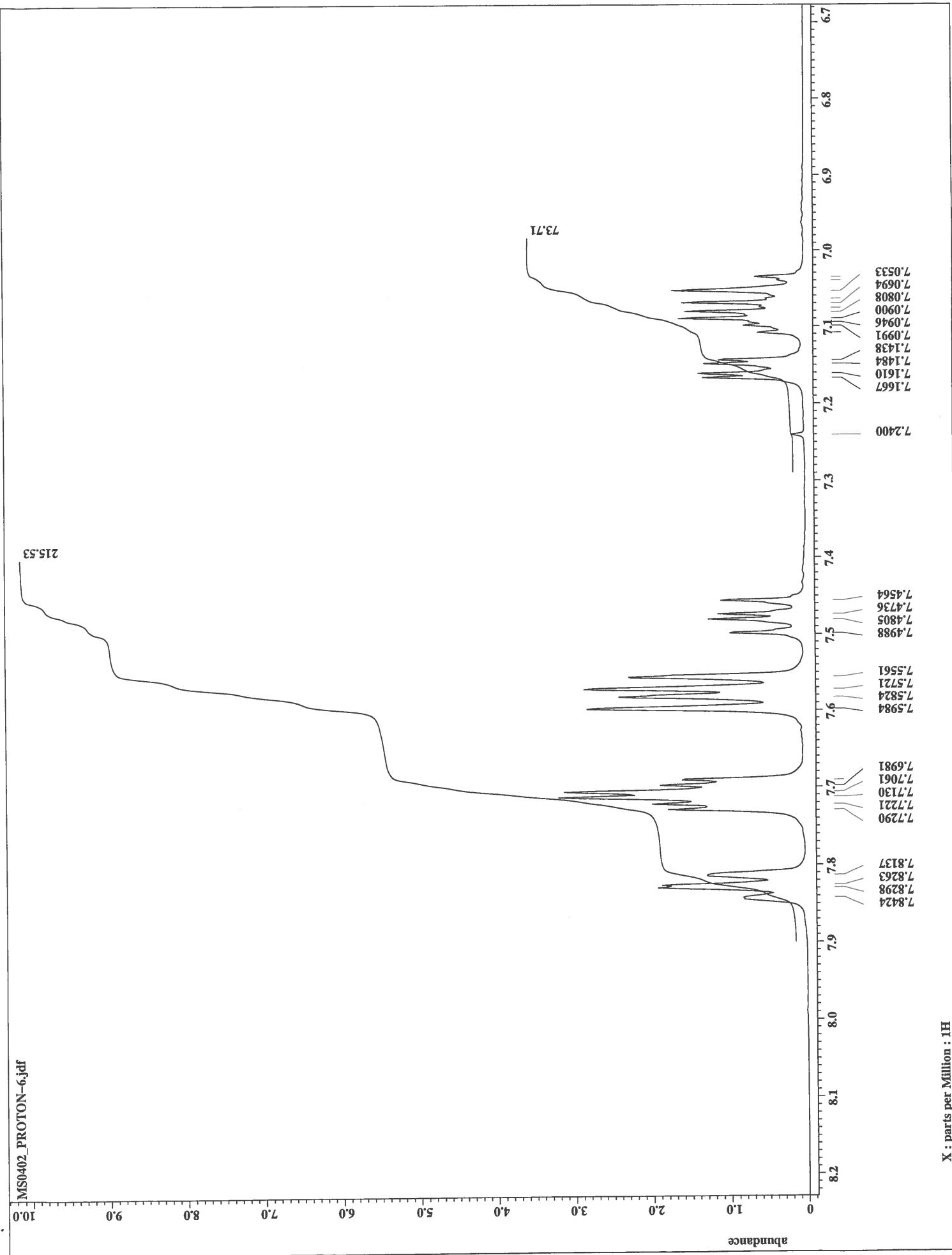
Data_format   = 1D COMPLEX
Dim_size      = 13107
Dim_title     = 1H
Dim_units     = [ppm]
Dimensions    = X
Site          = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 1.74587904 [s]
X_domain      = 1H
X_freq         = 500.15991521 [MHz]
X_offset       = 5.0 [ppm]
X_points       = 16384
X_prescans    = 1
X_resolution  = 0.5727773 [Hz]
X_sweep        = 9.38438438 [kHz]
Irr_domain    = 1H
Irr_freq       = 500.15991521 [MHz]
Irr_offset    = 5.0 [ppm]
Iri_domain   = 1H
Iri_freq      = 500.15991521 [MHz]
Tri_offset    = 5.0 [ppm]
Clipped       = FALSE
Mod_return    = 1
Scans         = 16
Total_scans   = 16

X_90_width    = 12.4 [us]
X_acq_time   = 1.74587904 [s]
X_angle       = 45 [deg]
X_atn         = 4 [dB]
X_pulse       = 6.2 [us]
Irr_mode     = Off
Tri_mode      = Off
Dante_preset = FALSE
Initial_wait  = 1 [s]
Recvz_gain    = 24
Relaxation_delay = 4 [s]
Temp_get      = 5.74587904 [s]
Temp_set      = 22.8 [dc]

```







```

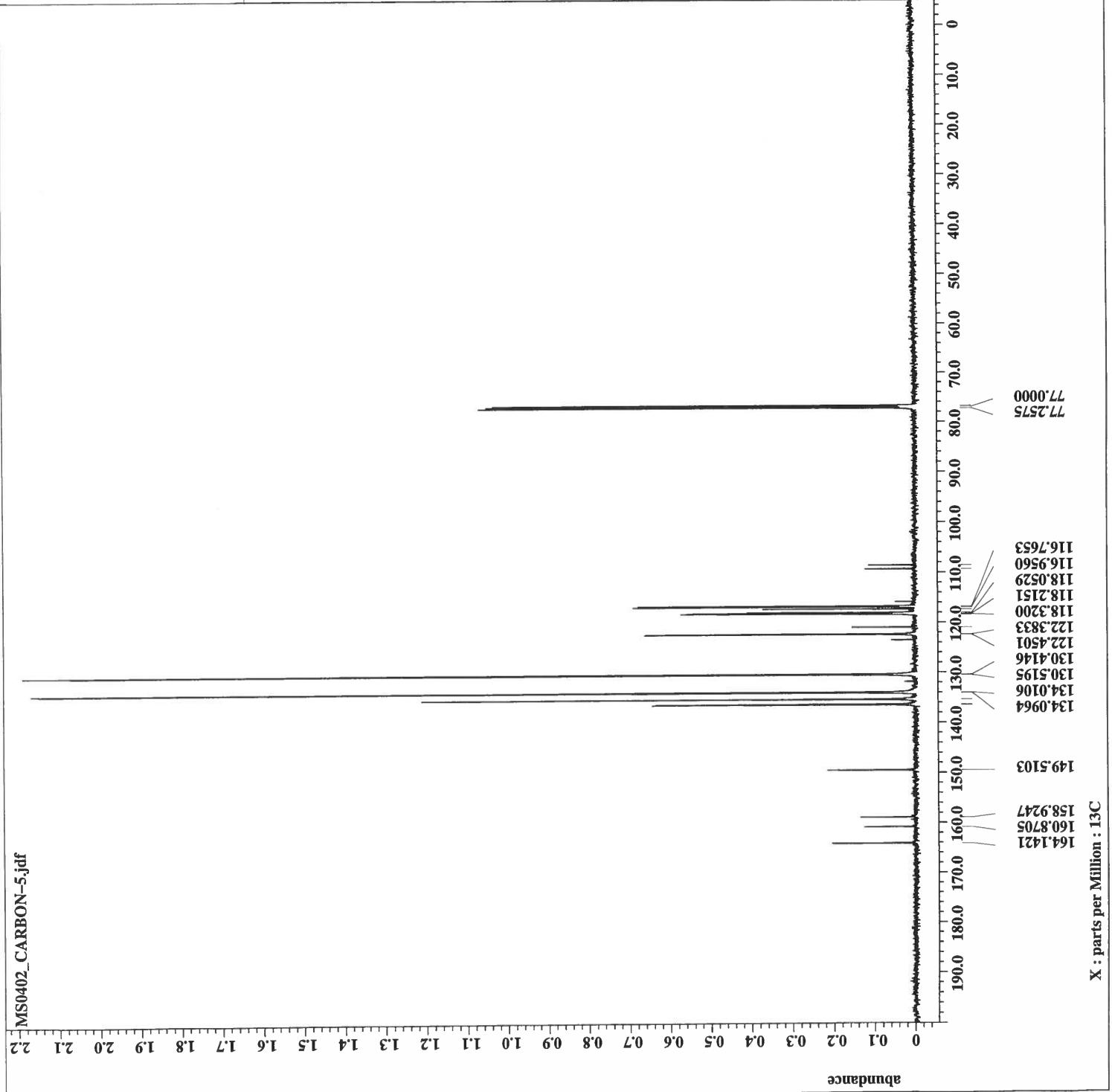
Filename = MS0402_CARBON-5.jdf
Author = Jim Davis
Experiment =
Sample_id =
Solvent =
Changer_sample =
Creation_time = 18-MAY-2018 17:08:56
Revision_time = 18-MAY-2018 16:44:20
Current_time = 18-MAY-2018 16:44:20

Data_format = 1D COMPLEX
Dim_size = 26214
Dim_title =
Dim_units = [ppm]
Dimensions =
X =
ECA 500
Site = JNM-ECX500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain =
X_freq = 125.76529768 [MHz]
X_offset = 100 [ppm]
X_points = 32768
X_Drscans =
X_resolution =
X_sweep =
Irr_domain =
Irr_freq = 500.15991521 [MHz]
Irr_offset =
Clipped =
Mod_return = FALSE
Scans = 1
Total_scans = 256

X_90_width =
X_acq_time = 0.83361792 [s]
X_angle =
X_atn =
X_pulse =
Irr_atn_dec =
Irr_atn_noe =
Irr_noise =
Decoupling =
Initial_wait =
Noe =
Noe_time =
Recv_grain =
Relaxation_delay =
Relaxation_time =
Temp_get =

```



3.0

2.0

1.0

0

abundance

X : parts per Million : ^{13}C 164.1421
160.8705
158.9247
149.5103136.4905
136.4047
135.4127
134.0964
134.0106
130.5195
130.4146
123.5375
122.4501
122.3833
120.9812
118.3200
118.2151
118.0529
117.3376
116.9560
116.7653
109.3349
108.5718



```

Filename          = MS0402_FLUORINE-5.jdf
Author           = Jim Davis
Experiment       = single_pulse.ex2
Sample_id        = MS0402
Solvent          = CHLOROFORM-D
Changer_sample  = 6
Creation_time   = 18-MAY-2018 17:11:54
Revision_time   = 18-MAY-2018 16:47:18
Current_time    = 18-MAY-2018 16:47:18

Data_format     = 1D COMPLEX
Dim_size        = 52428
Dim_title       = 19F
Dim_units       = [ppm]
Dimensions      = X
Site            = ECA 500
Spectrometer    = JNM-ECA500

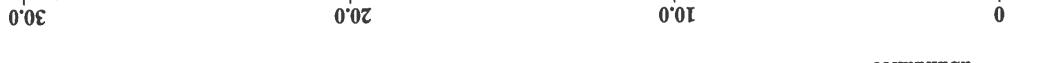
Field_strength  = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain        = 19F
X_freq          = 470.62046084 [MHz]
X_offset        = -70 [ppm]
X_points        = 65536
X_precsans     = 1
X_resolution   = 1.7993855 [Hz]
X_sweep         = 117.9245281 [kHz]
Irr_domain     = 19F
Irr_freq        = 470.62046084 [MHz]
Irr_offset      = 5 [ppm]
Tri_domain     = 19F
Tri_freq        = 470.62046084 [MHz]
Tri_offset      = 5 [ppm]
Clipped         = FALSE
Mod_return      = 1
Scans           = 16
Total_scans    = 16

X_90_width     = 13.1 [us]
X_acq_time     = 0.55574528 [s]
X_angle         = 45 [deg]
X_atn          = 2.5 [dB]
X_pulse         = 6.55 [us]
Irr_mode        = Off
Tri_mode        = Off
Dante_preset   = FALSE
Initial_wait    = 1 [s]
Recvr_gain      = 36
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get        = 22.6 [dc]

```

-116.4623

-78.6142





```

filename = MS0402_PHOSPHORUS-5.j
author = Jim Davis
Experiment = single_pulse_dec
sample_id = MS0402
solvent = CHLOROFORM-D
Changer_sample = 6
Creation_time = 18-MAY-2018 17:15:26
Revision_time = 18-MAY-2018 16:50:51
Current_time = 18-MAY-2018 16:50:51

data_format = 1D COMPLEX
dim_size = 26214
dim_title = 31P
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = JNM-ECA500

field_strength = 11.7473579 [T] (500 [MHz])
x_acq_duration = 0.64487424 [s]
x_domain = 31P
x_freq = 201.46831075 [MHz]
x_offset = 0 [ppm]
x_points = 32768
x_prescans = 4
x_resolution = 1.550688995 [Hz]
x_sweep = 50.-81.30081.3 [Hz]
irr_domain = 1H
irr_freq = 500.15991521 [MHz]
irr_offset = 5.0 [ppm]
clipped = FALSE
mod_return = 1
scans = 25
total_scans = 25

x_90_width = 14.687 [us]
x_acq_time = 0.64487424 [s]
x_angle = 30 [deg]
x_atn = 5 [dB]
x_pulse = 4.89566667 [us]
irr_atn_dec = 20.7 [dB]
irr_atn_noe = 20.7 [dB]
irr_noise = 56
decoupling = WALTZ
decoupling_true = TRUE
initial_wait = 1 [s]
noe_time = 2 [s]
recvr_gain = 56
relaxation_delay = 2 [s]
repetition_time = 2.64487424 [s]
temp_get = 22.9 [dC]

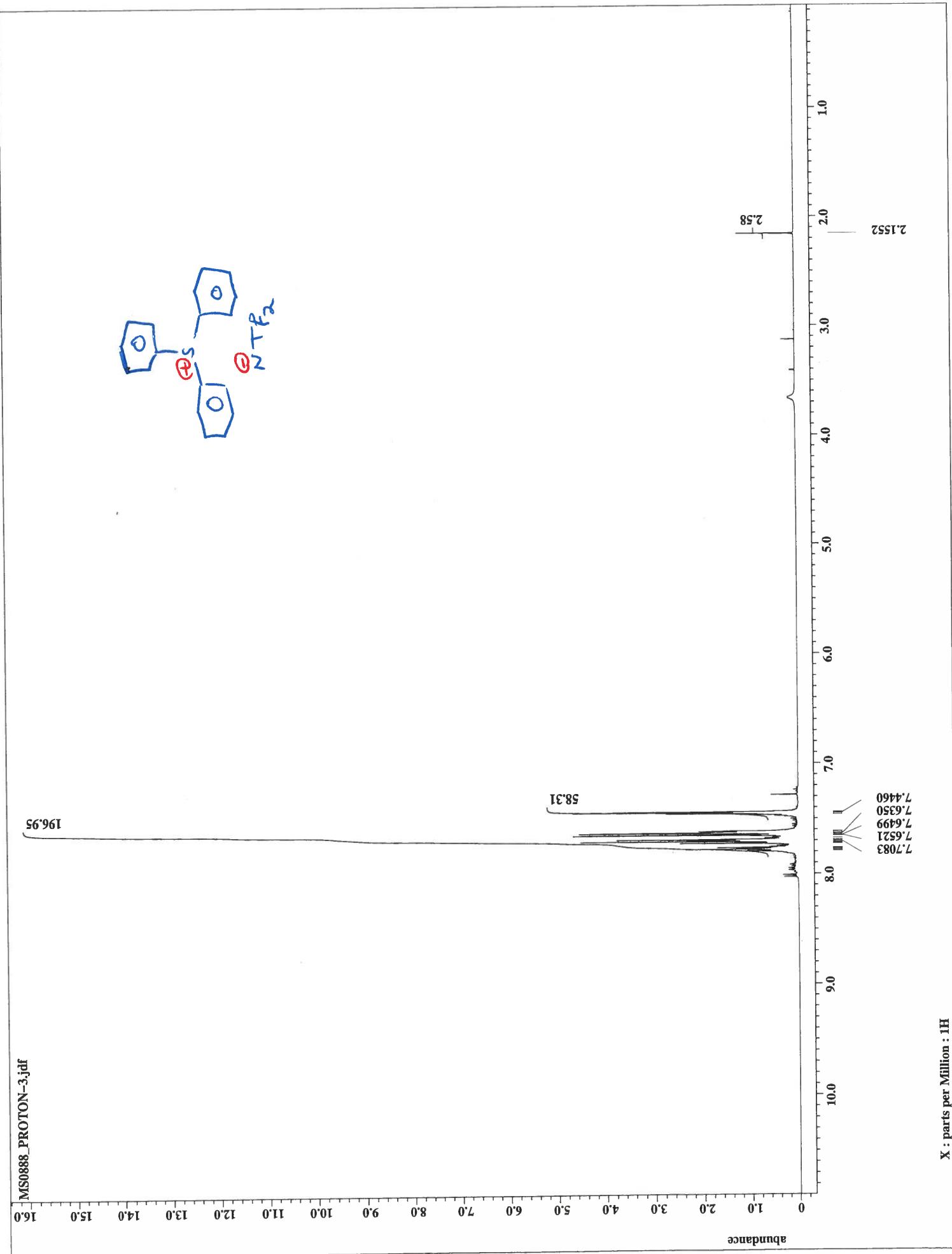
0.090.0 80.0 70.0 60.0 50.0 40.0 30.0 20.0 10.0 0 -10.0 -20.0 -30.0 -40.0 -50.0 -60.0 -70.0 -80.0 -90.0 100.0

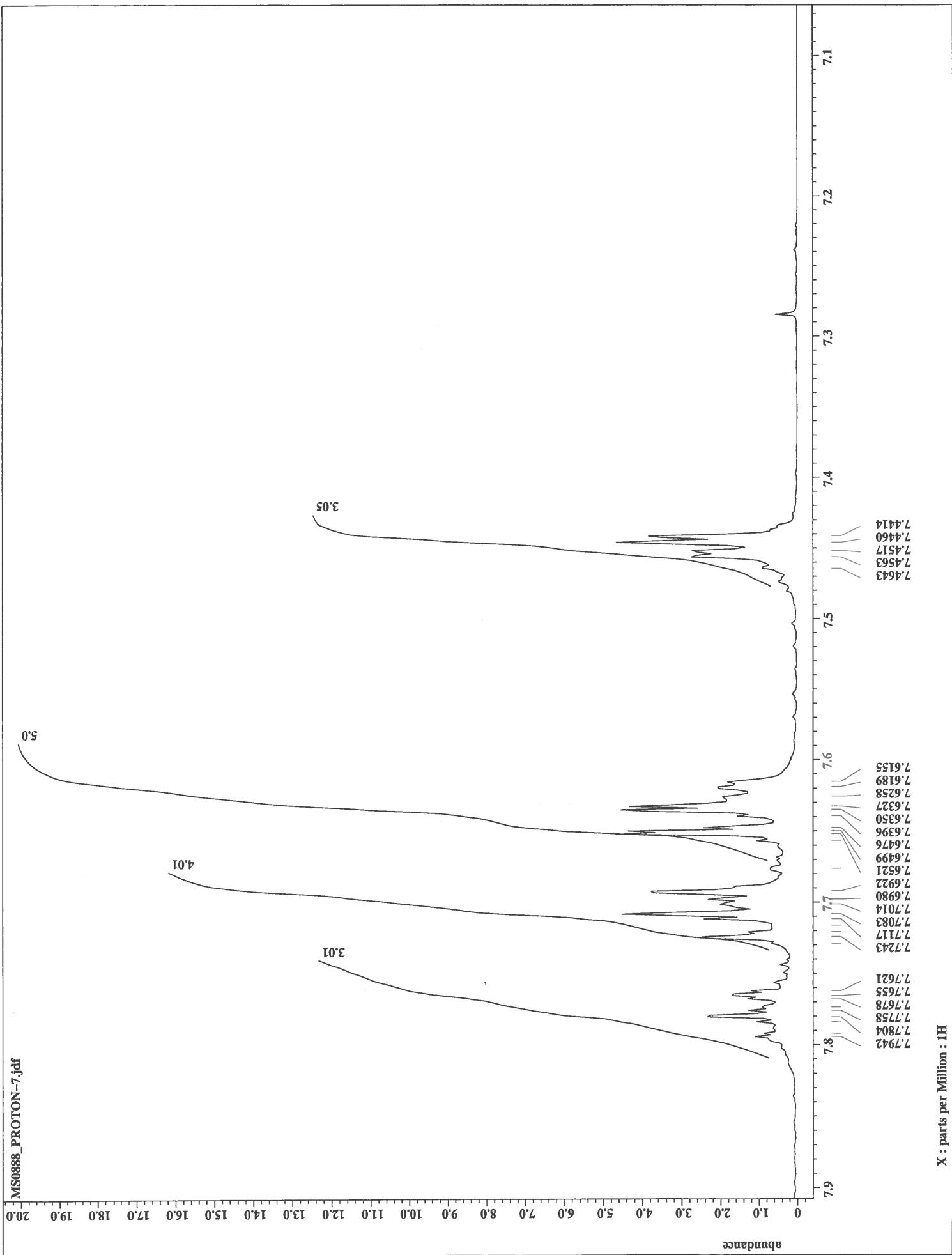
```

23.2295

X : parts per Million : 31P

abundance





4.0

3.0

2.0

1.0

0

abundance

X : parts per Million : 13C

144.9348

134.9291

132.1535
131.9522
131.9150
131.8006
131.3496
131.1424
130.8944
129.4351124.7804
123.8742
121.1081
118.5518
115.9956

113.0

115.0

117.0

119.0

121.0

123.0

125.0

127.0

129.0

131.0

133.0

135.0

137.0

139.0

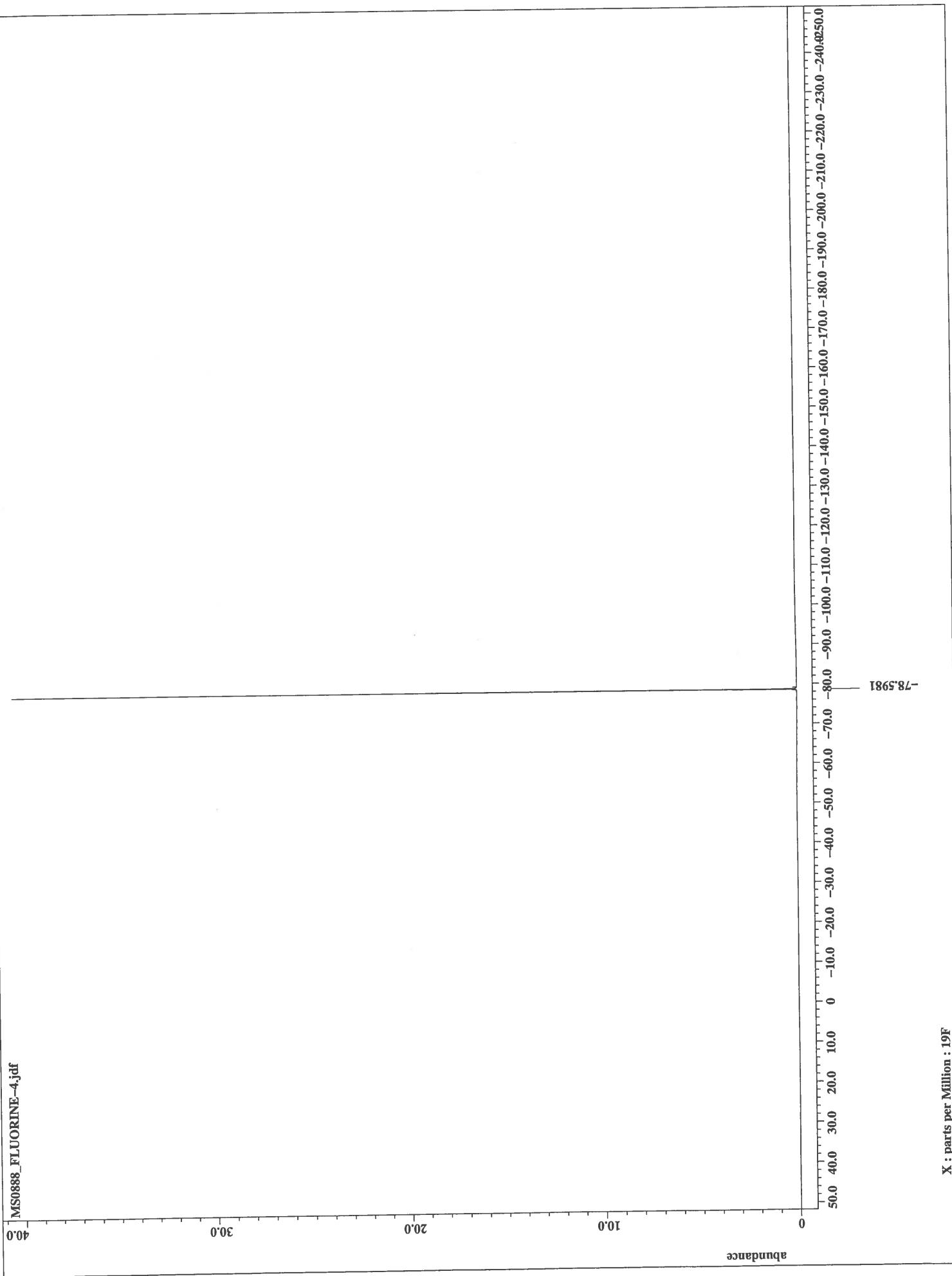
141.0

143.0

145.0

147.0

149.0





```

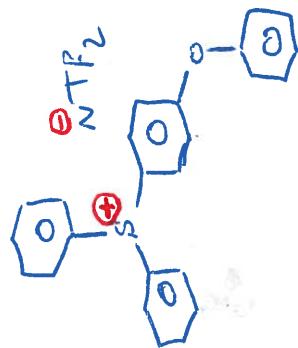
= MS0886_PROTON-2.jdf
= Jim Davis
= single_pulse.ex2
Sample_id = MS0886
Solvent = CHLOROFORM-D
Creation_time = 29-NOV-2019 12:51:31
Revision_time = 29-NOV-2019 12:23:15
Current_time = 29-NOV-2019 12:22:15

Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = RCA 500
Spectrometer = JNM-ECX500

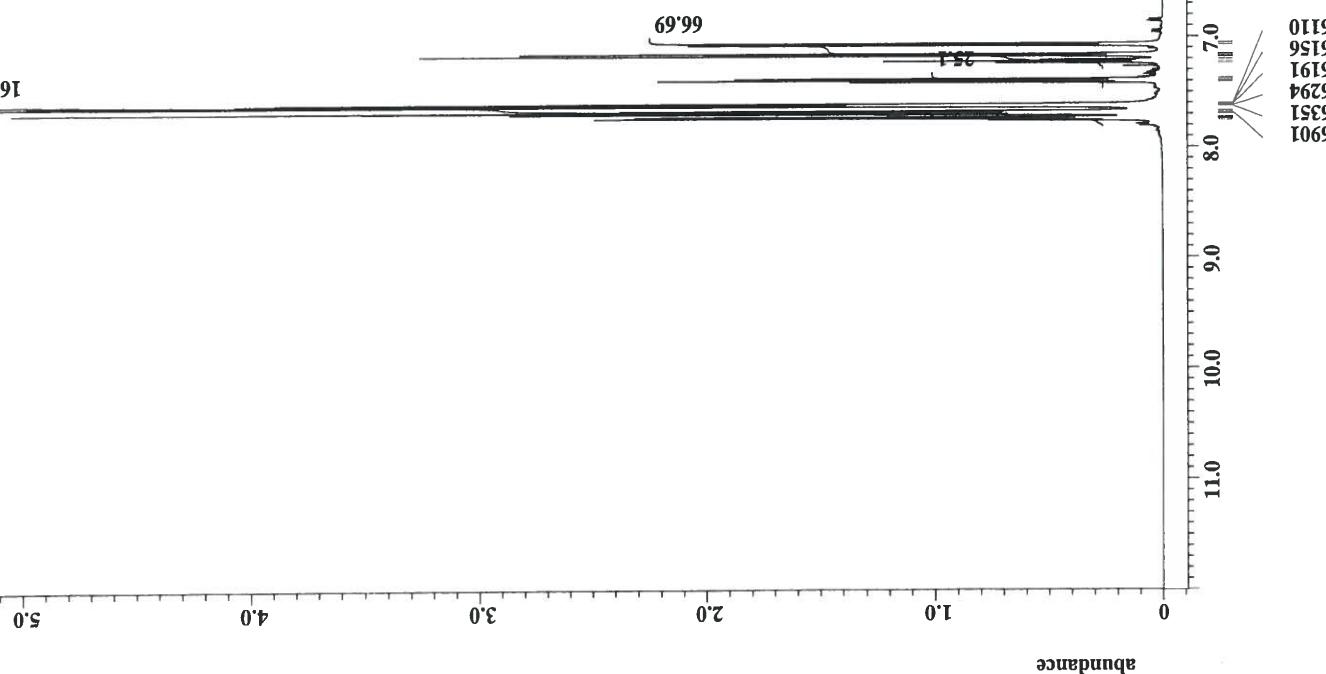
Field_strength = 11.7473579 [T] (500 [MHz]
X_acq_duration = 1.74587904 [s]
X_domain = 1H
X_freq = 500.15991521 [MHz]
X_offset = 5.0 [ppm]
X_points = 16384
X_prescans = 1
X_resolution = 0.57277737 [Hz]
X_sweep = 9.38438438 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Tri_domain = 1H
Tri_freq = 500.15991521 [MHz]
Tri_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

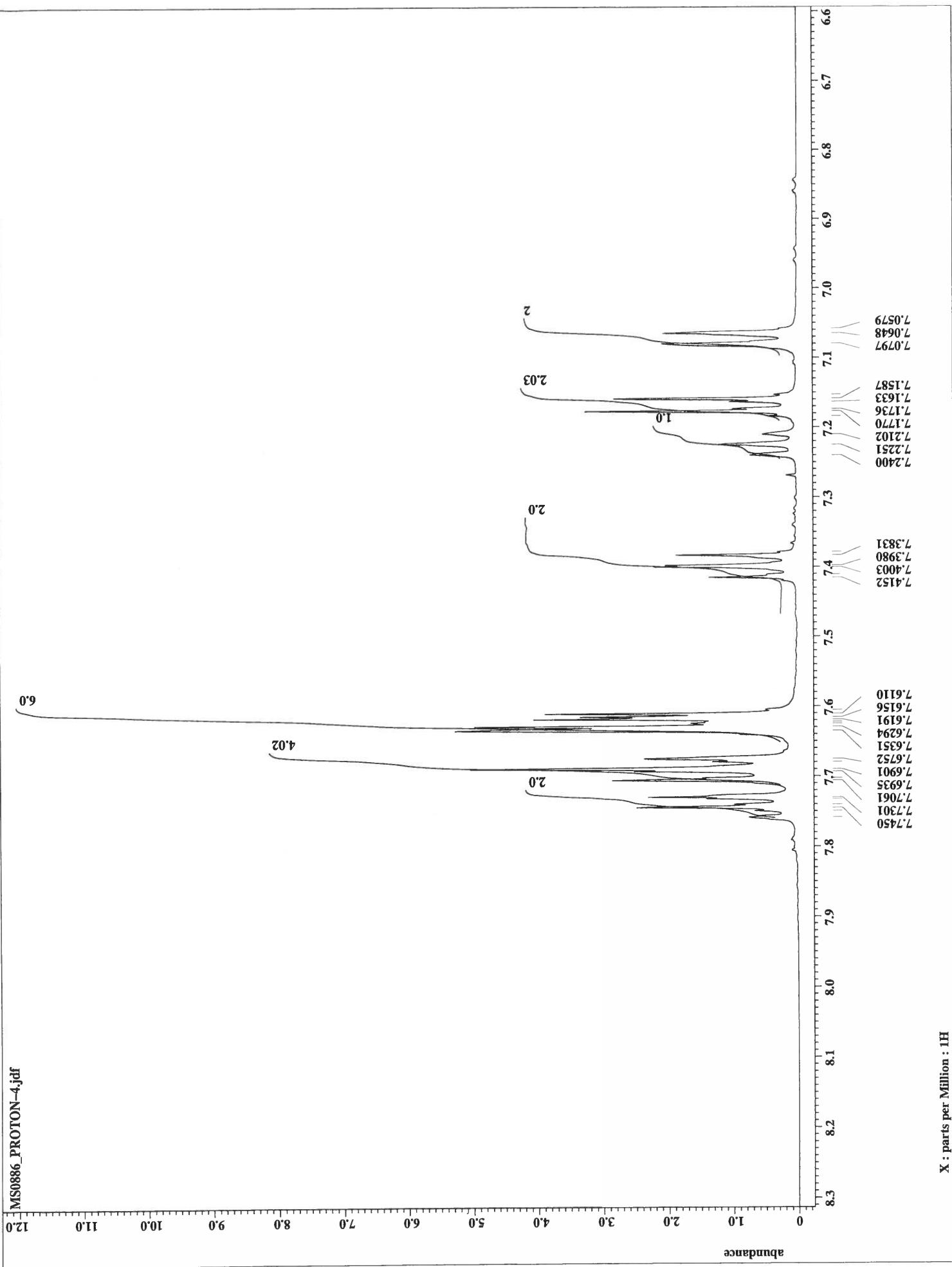
X_90_width = 11.3 [us]
X_acq_time = 1.74587904 [s]
X_angle = 45 [deg]
X_atm = 4 [dB]
X_pulse = 5.65 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Recv_gain = 22
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get = 19 [dc]

```



168.72







```

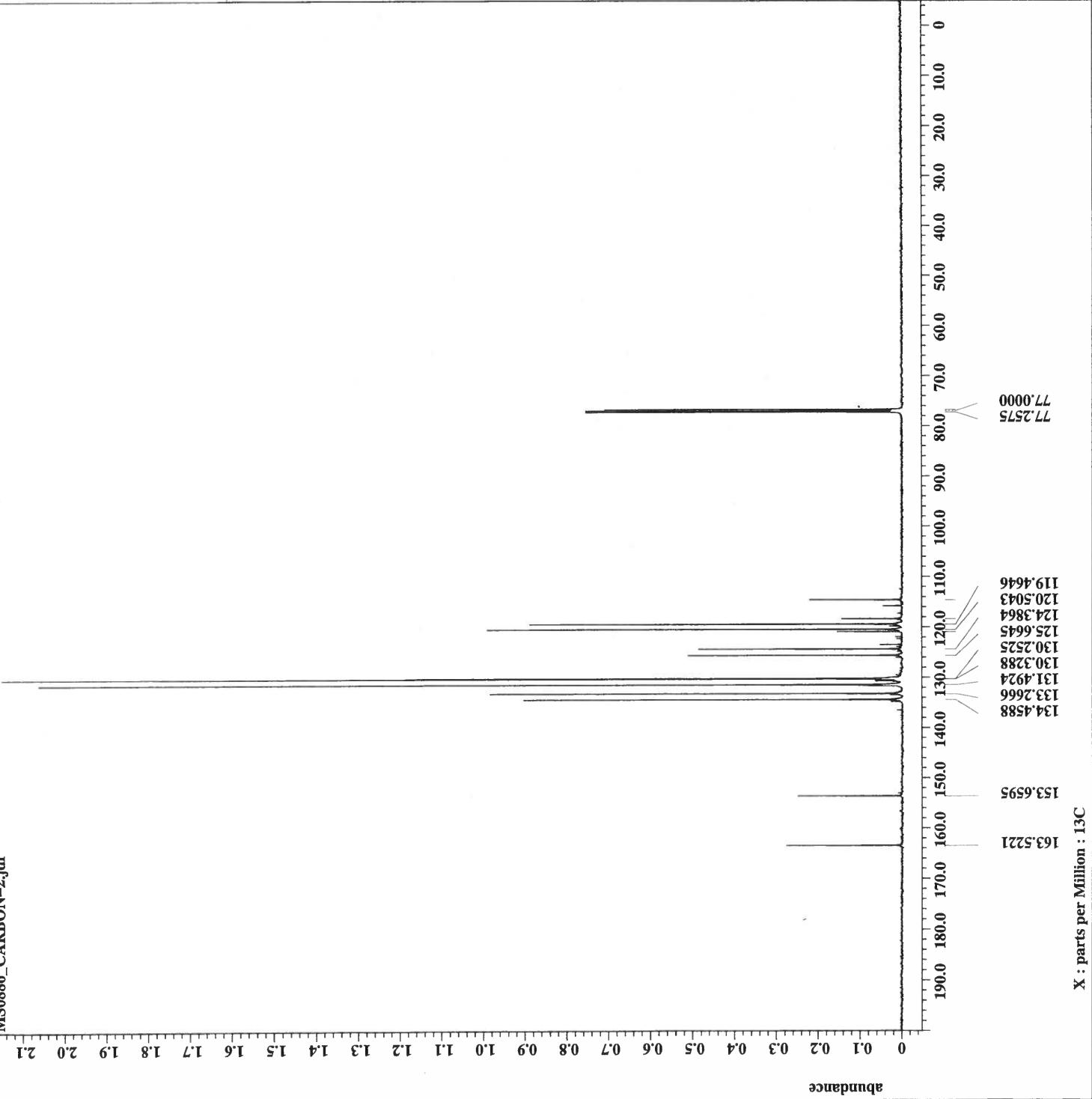
Filename      = MS0886_CARBON-2.jdf
Author        = Jim Davis
Experiment   = single_pulse_dec
Sample_id    = MS0886
Solvent      = CHLOROFORM-D
Creation_time = 29-NOV-2019 13:40:22
Revision_time = 29-NOV-2019 13:11:06
Current_time  = 29-NOV-2019 13:11:06

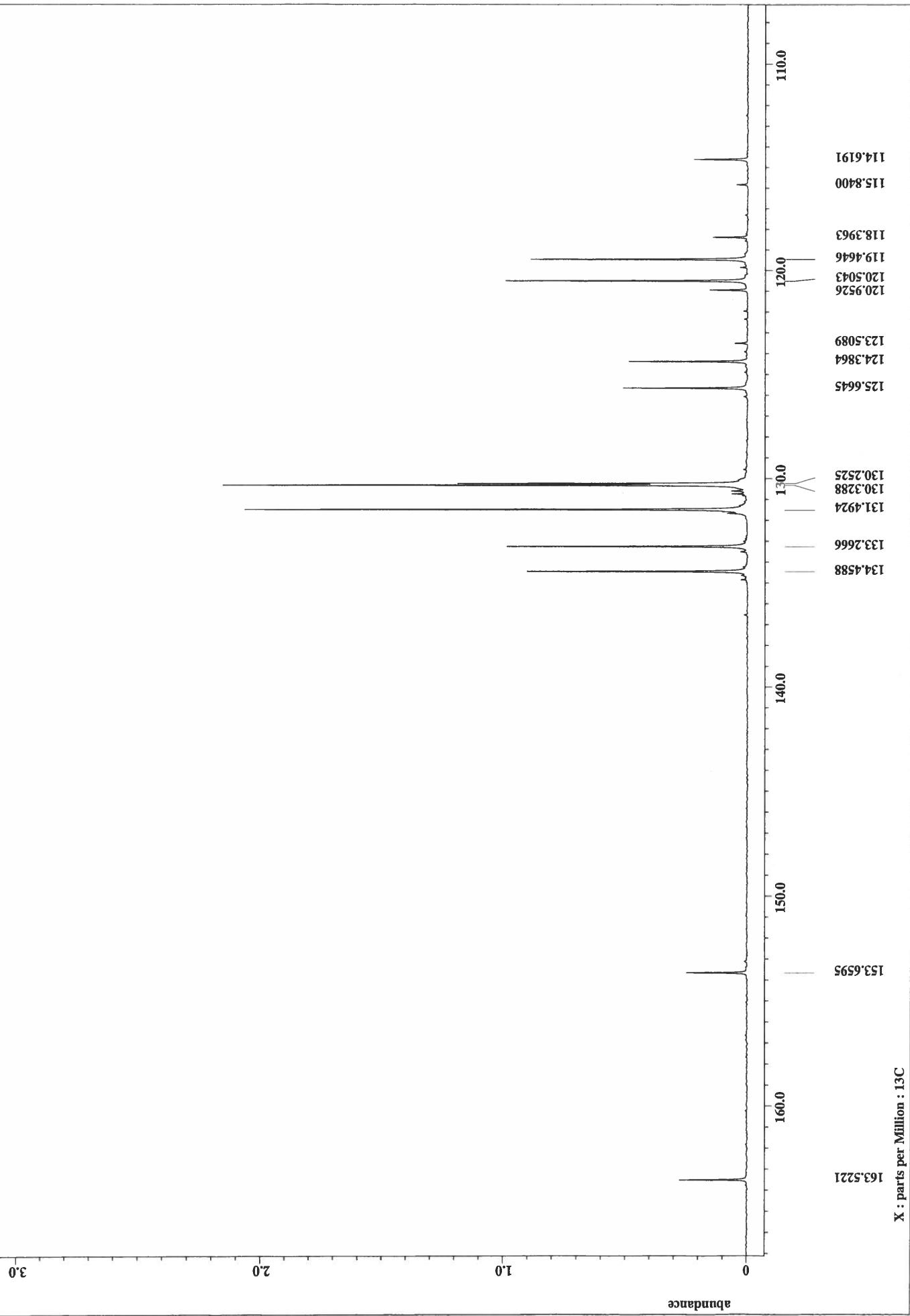
Data_format  = 1D COMPLEX
Dim_size     = 26214
Dim_title   = 13C
Dim_units   = [ppm]
Dimensions  = X
Site         = ECA 500
Spectrometer = JNM-ECA500

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]
Irr_domain   = 1H
Irr_freq      = 500.15991521 [MHz]
Irr_offset    = 5.0 [ppm]
Clipped      = FALSE
Mod_return   = 1
Scans         = 1024
Total_scans  = 1024

X_90_width   = 13 [us]
X_acq_time   = 0.83361792 [s]
X_angle       = 30 [deg]
X_atn        = 6 [dB]
X_pulse      = 4.33333333 [us]
Irr_atn_dec  = 21.2 [dB]
Irr_atn_noe  = 21.2 [dB]
Irr_noise    = WALETZ
Decoupling   = TRUE
Initial_wait = 1 [s]
Noe          = TRUE
Noe_time     = 2 [s]
Recvr_gain   = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get     = 20 [ac]

```





30.0

20.0

10.0

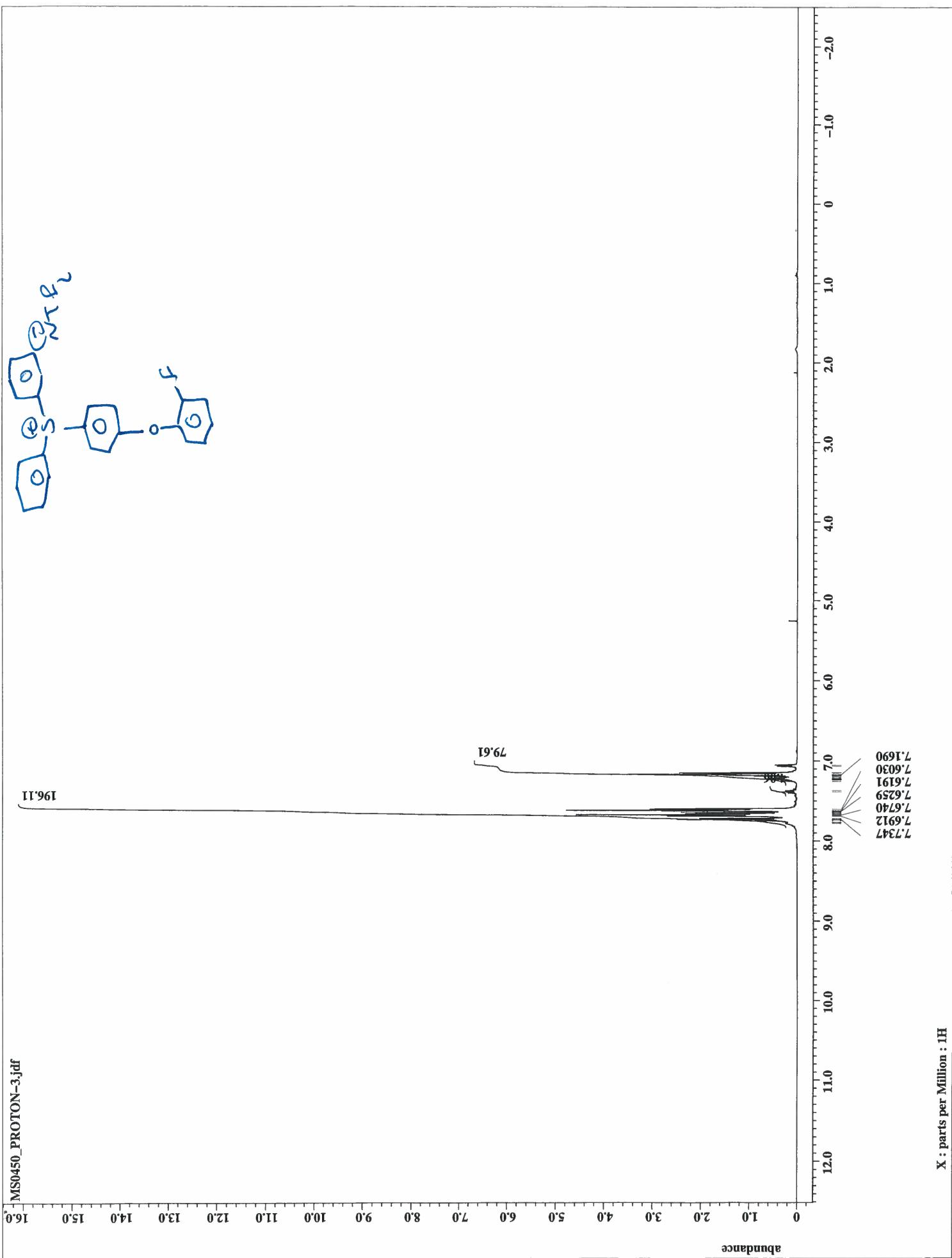
0

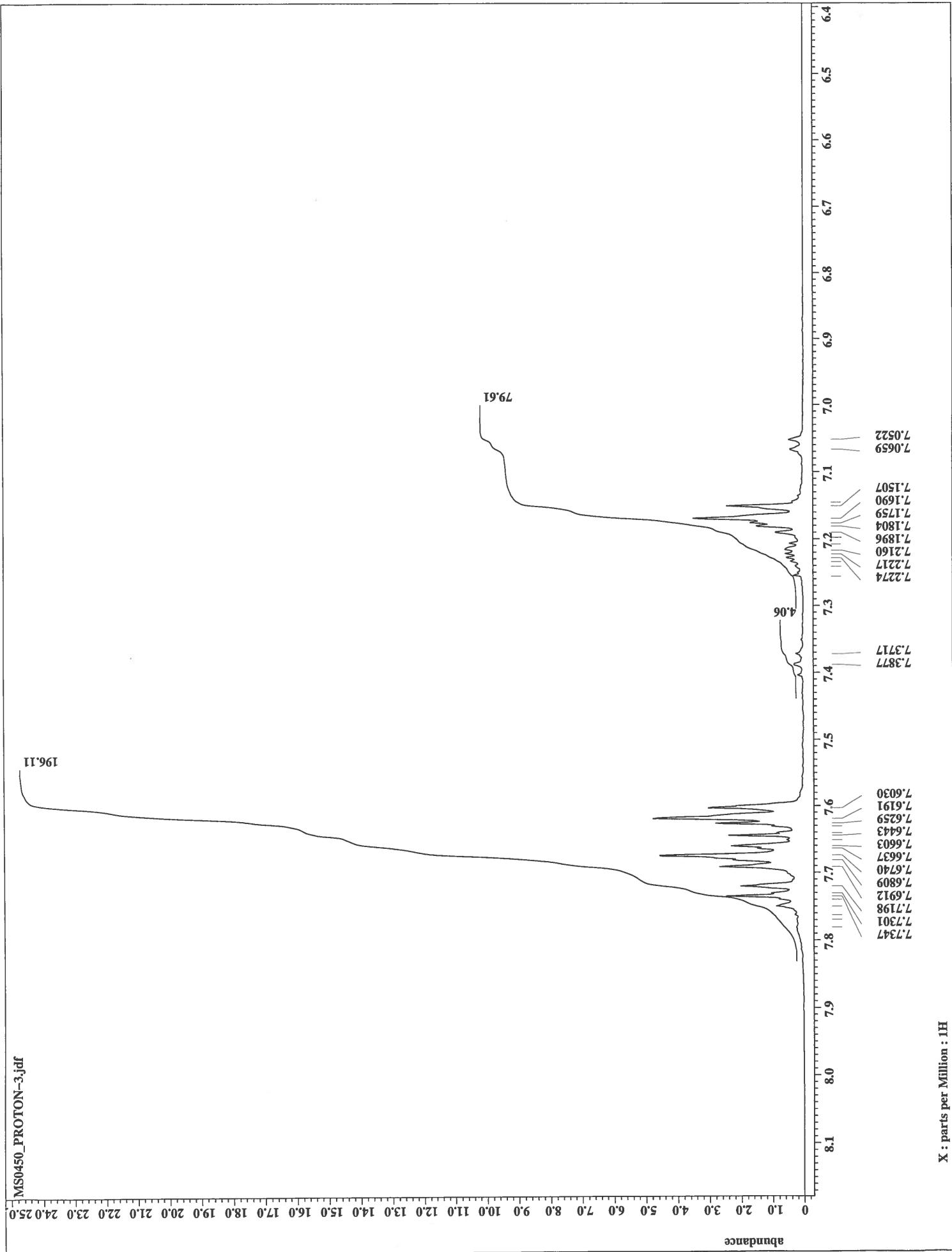
abundance

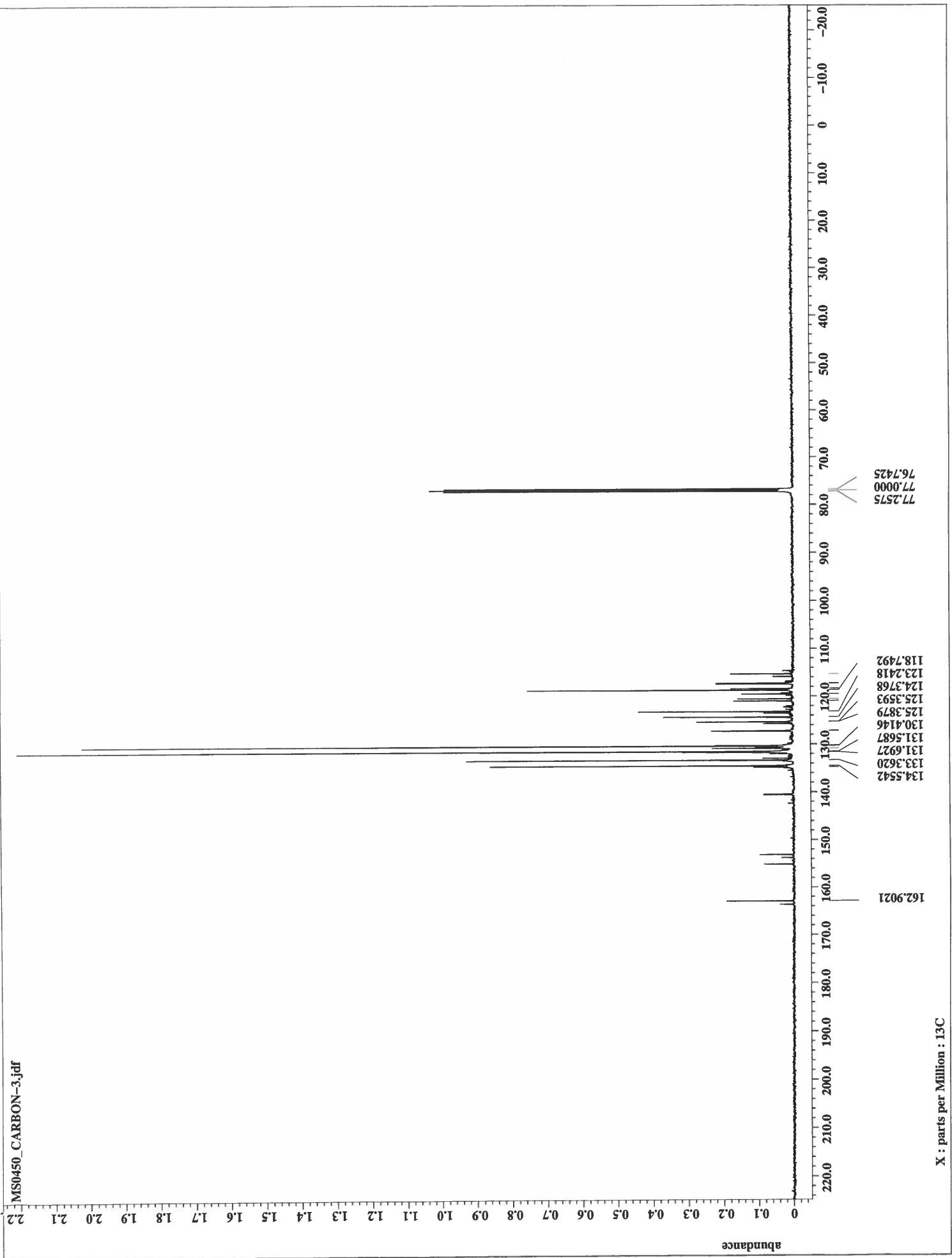
X : parts per Million : 19F

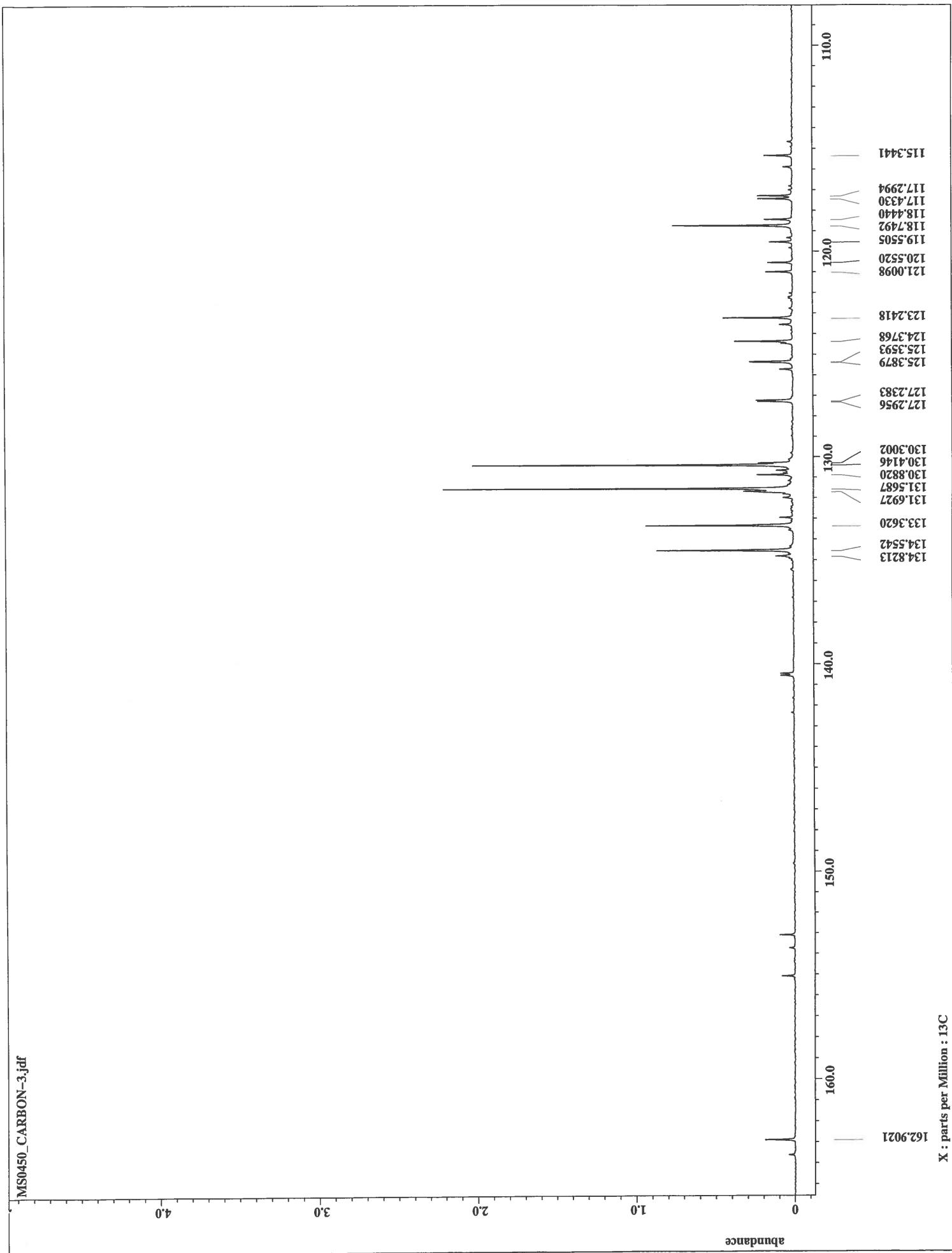
-78.6010

50.0 40.0 30.0 20.0 10.0 0 -10.0 -20.0 -30.0 -40.0 -50.0 -60.0 -70.0 -80.0 -90.0 -100.0 -110.0 -120.0 -130.0 -140.0 -150.0 -160.0 -170.0 -180.0 -190.0 -200.0 -210.0 -220.0 -230.0 -240.0 -250.0









30.0

20.0

10.0

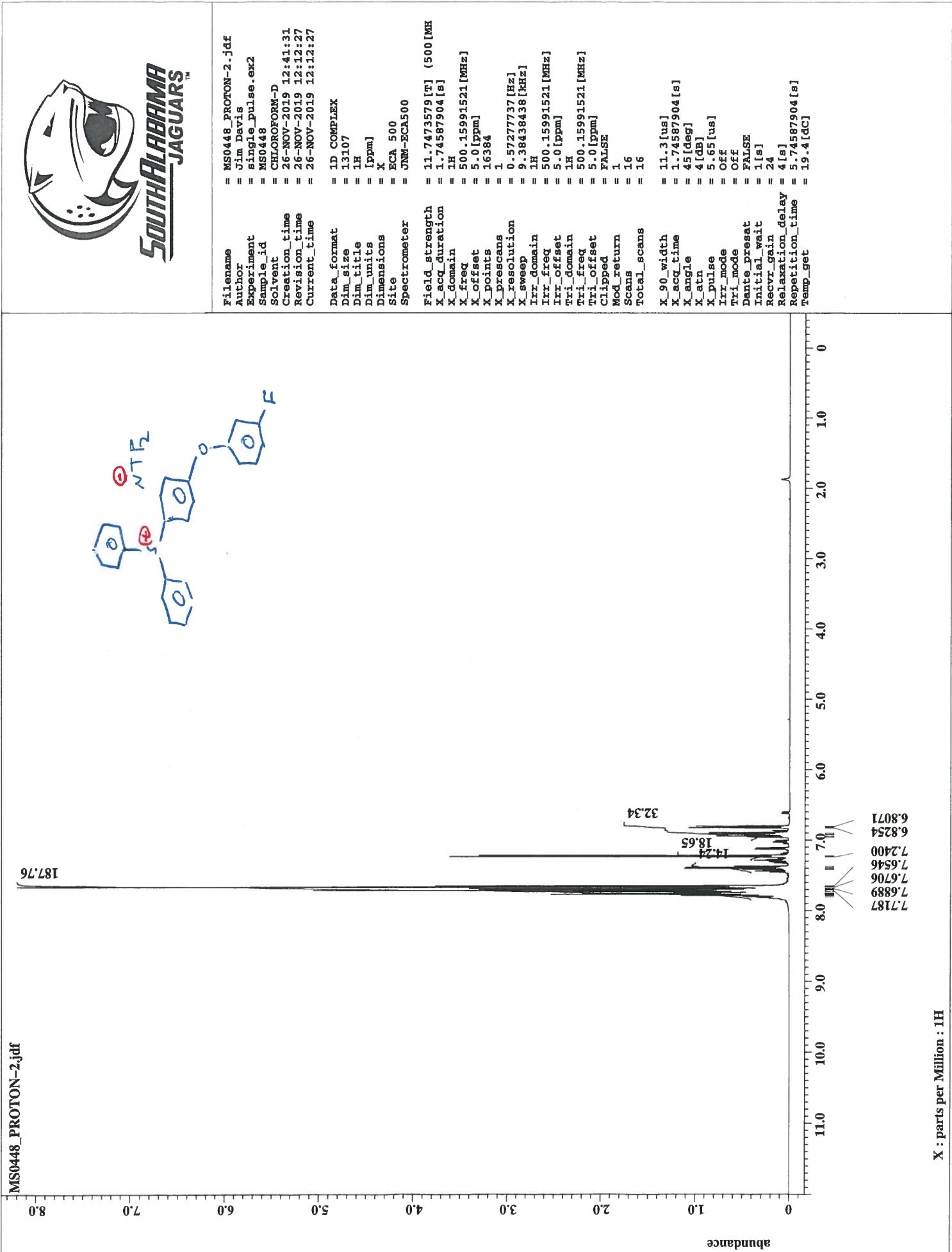
0

abundance

X : parts per Million : 19F

-78.6142

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





SOUTH ALABAMA
JAGUARS™

```

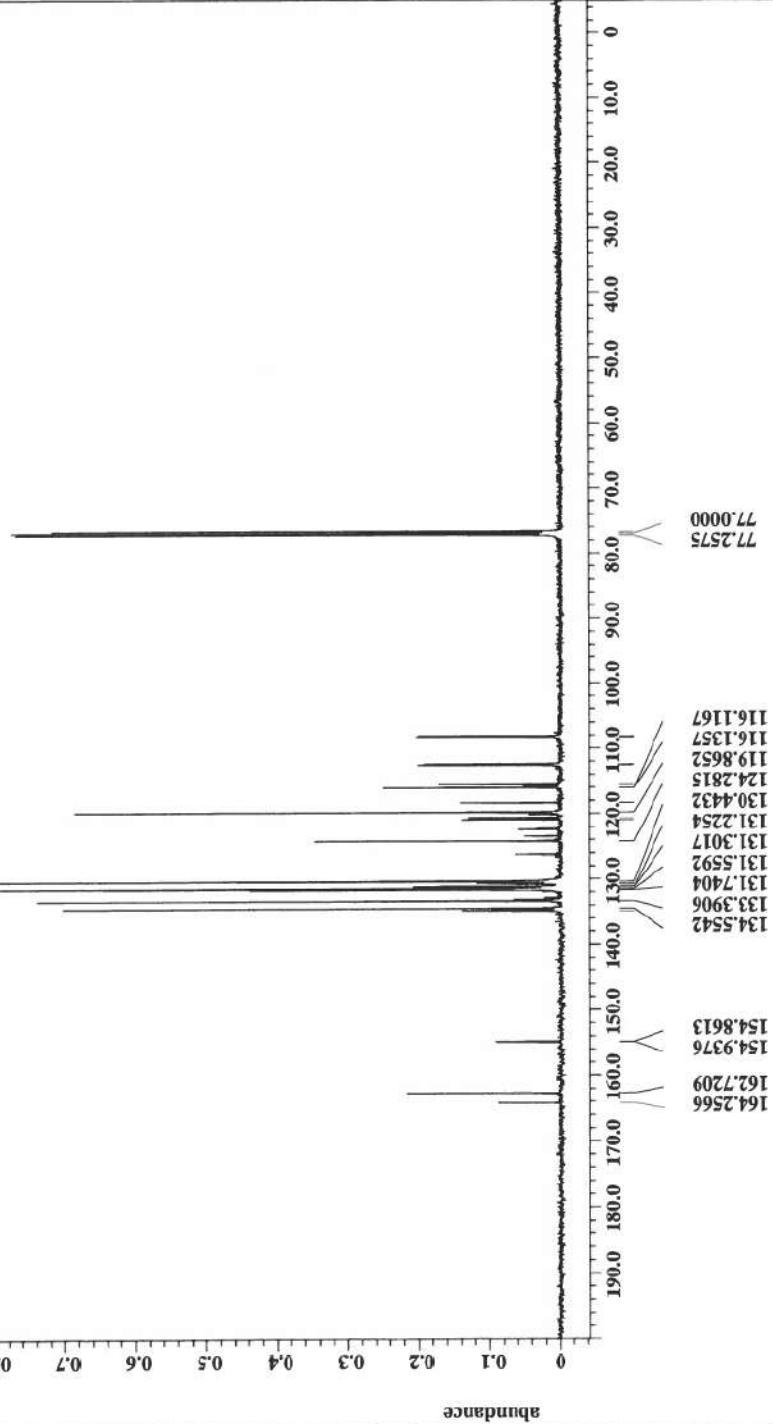
Filename = MS0448 CARBON-2.jdf
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0448
Solvent = CHLOROFORM-D
Creation_time = 26-NOV-2019 13:01:16
Revision_time = 26-NOV-2019 12:32:12
Current_time = 26-NOV-2019 12:32:12

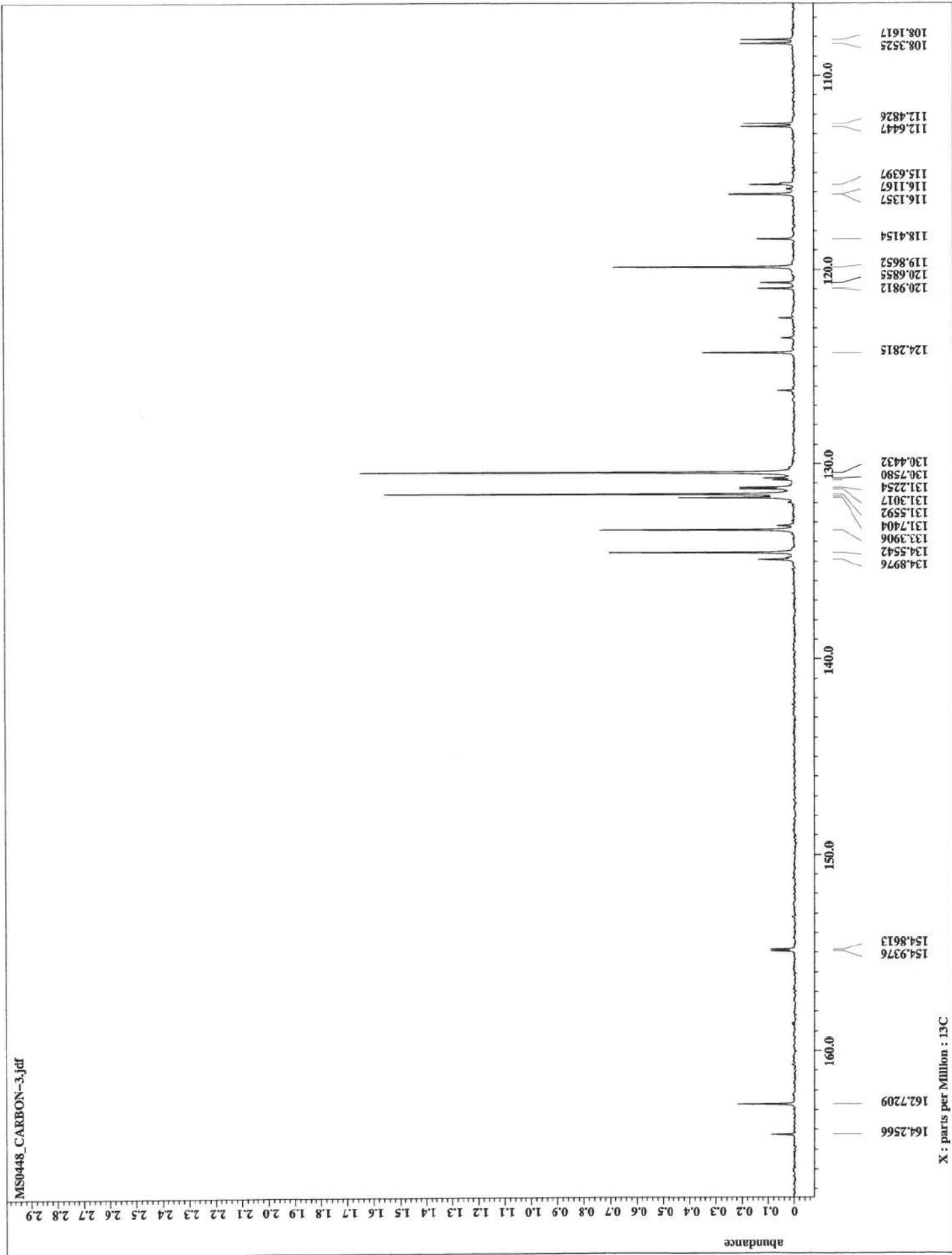
Data_format = 1D COMPLEX
Dim_size = 26214
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
spectrometer = URW-ECR500

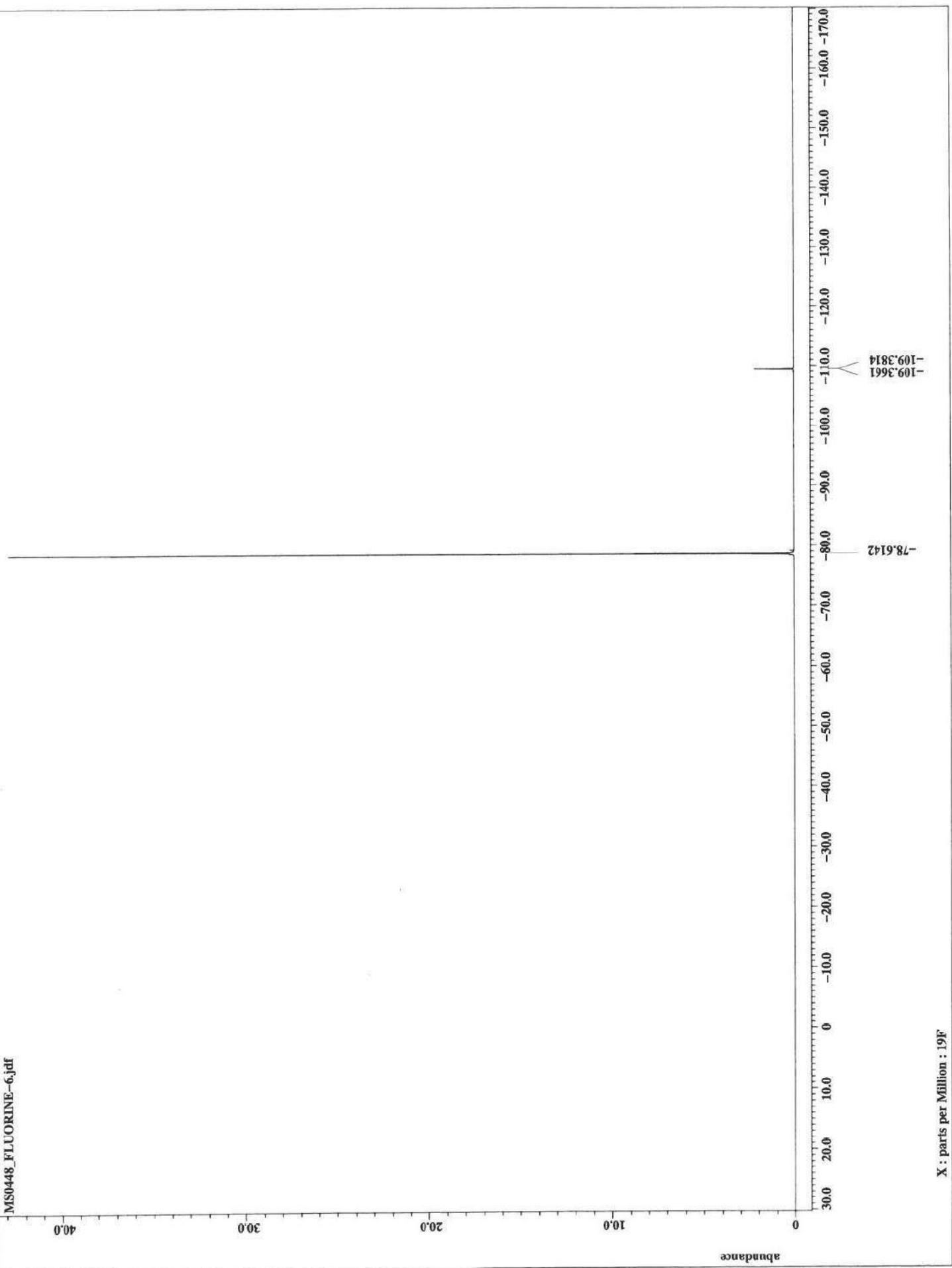
Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.833361792 [s]
X_domain = 13C
X_freq = 125.76529768 [MHz]
X_offset = 100 [ppm]
X_points = 32768
X_prescans = 4
X_resolution = 1.19959034 [Hz]
X_sweep = 39.3083761 [Hz]
Irr_domain = 1H
Irr_freq = 500.15591521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 400
Total_scans = 400

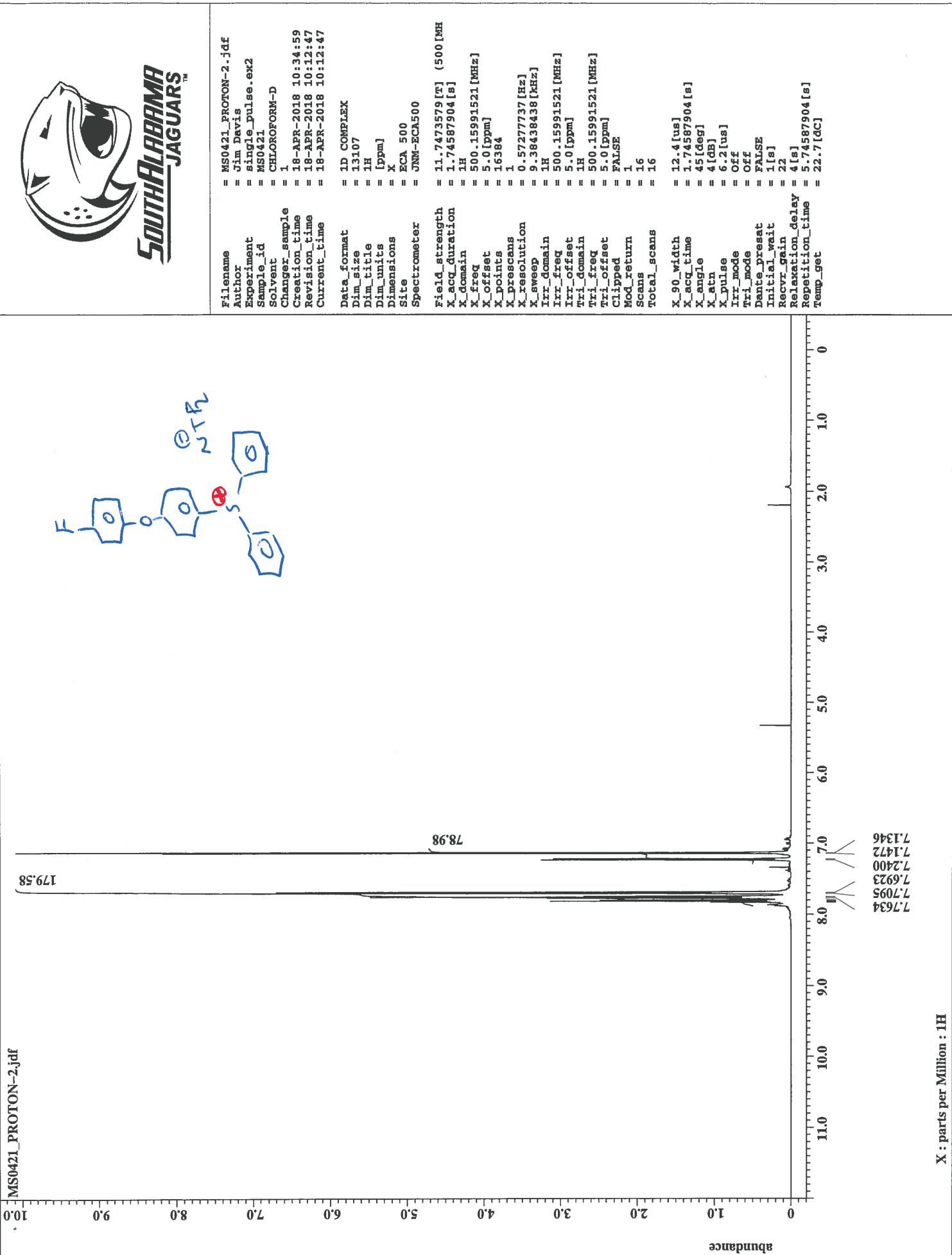
X_90_width = 13 [us]
X_acq_time = 0.833361792 [s]
X_angle = 30 [deg]
X_attn = 6 [dB]
X_pulse = 4.33333333 [us]
Irr_attn_dec = 21.2 [dB]
Irr_attn_noe = 21.2 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Recv_grain = 60
Relaxation_delay = 2.833361792 [s]
Repetition_time = 2.833361792 [s]
Temp_get = 19.6 [dc]

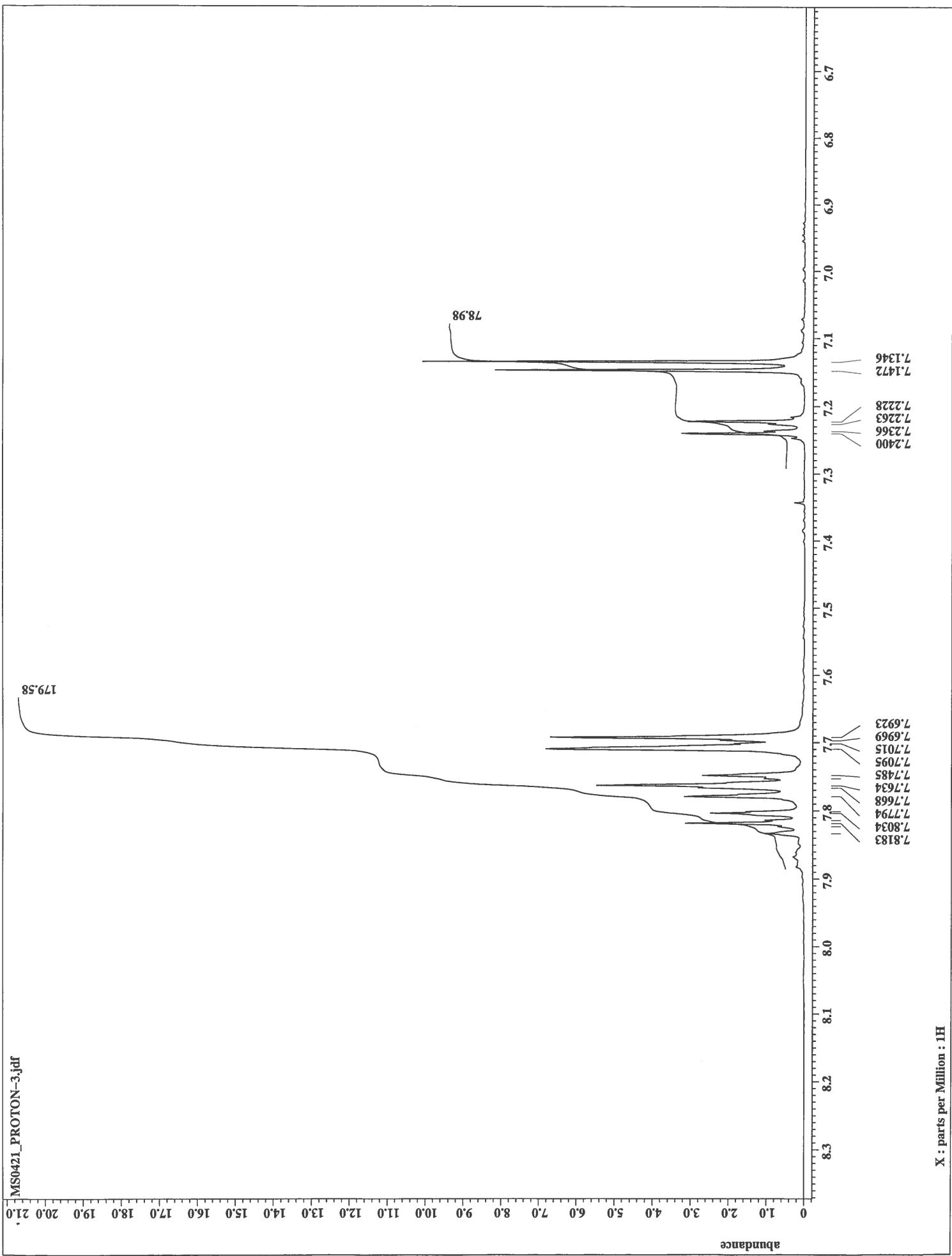
```













```

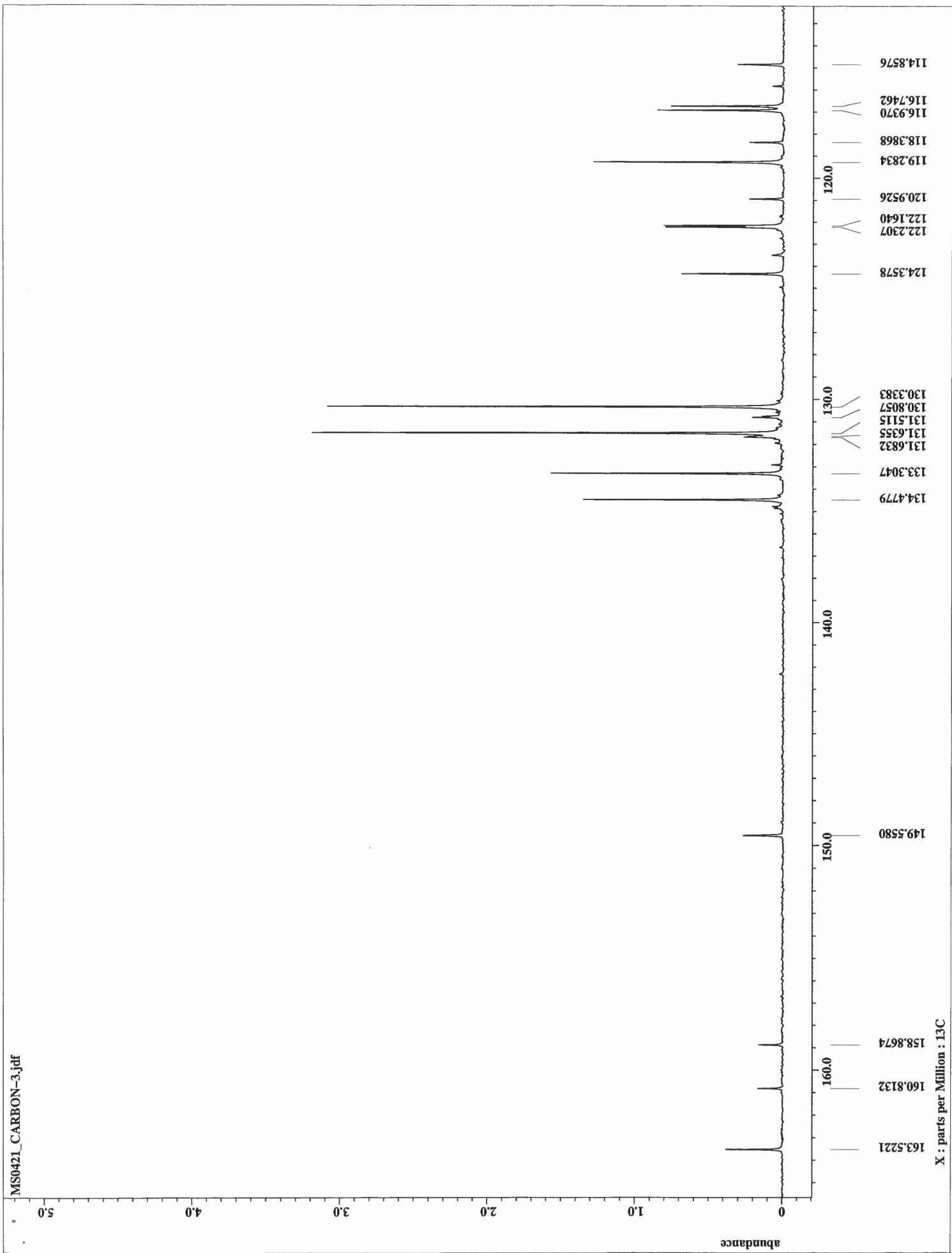
Filename      = MS0421_CARBON-2.jdf
Author        = Jim Davis
Experiment    = single_pulse_dec
Sample_id     = MS0421
Solvent       = CHLOROFORM-D
Changer sample = 1
Creation_time = 18-APR-2018 10:49:35
Revision_time = 18-APR-2018 10:27:22
Current_time  = 18-APR-2018 10:27:22

Data_format   = 1D COMPLEX
Dim_size      = 26214
Dim_title     = 13C
Dim_units     = [ppm]
Dimensions    = X
Site          = ECA 500
Spectrometer  = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X.acq_duration = 0.83361792 [s]
X.domain      = 13C
X.freq         = 125.765229768 [MHz]
X.offset       = 100 [ppm]
X.points       = 32768
X.precans     = 4
X.resolution  = 1.19959034 [Hz]
X.sweep        = 35.3084761 [kHz]
Irr_domain    = 1H
Irr_freq       = 500.15991521 [MHz]
Irr_offset     = 5.0 [ppm]
Clipped        = FALSE
Mod_return    = 1
Scans          = 256
Total_scans   = 256

X.90_width    = 13.2 [us]
X.acq_time    = 0.83361792 [s]
X.angle        = 30 [deg]
X.attn        = 6 [dB]
X.pulse        = 4.4 [us]
Irr_atr_dec   = 20.7 [dB]
Irr_atr_noe   = 20.7 [dB]
Irr_noise     = WALTZ
Decoupling    = TRUE
Initial_wait   = 1 [s]
Noe           = TRUE
Noe_time      = 2 [s]
Recvr.gain    = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get      = 23.4 [DC]
  
```







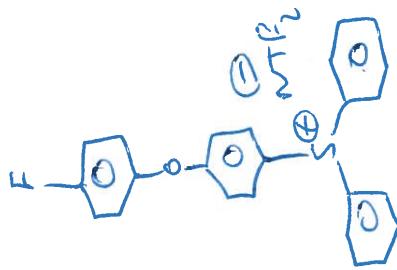
```

Filename = MS0421_FLUORINE-2.jdf:
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0421
Solvent = CHLOROFORM-D
Changer_sample = 1
Creation_time = 18-APR-2018 10:31:25
Revision_time = 18-APR-2018 10:09:13
Current_time = 18-APR-2018 10:09:13

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [kHz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

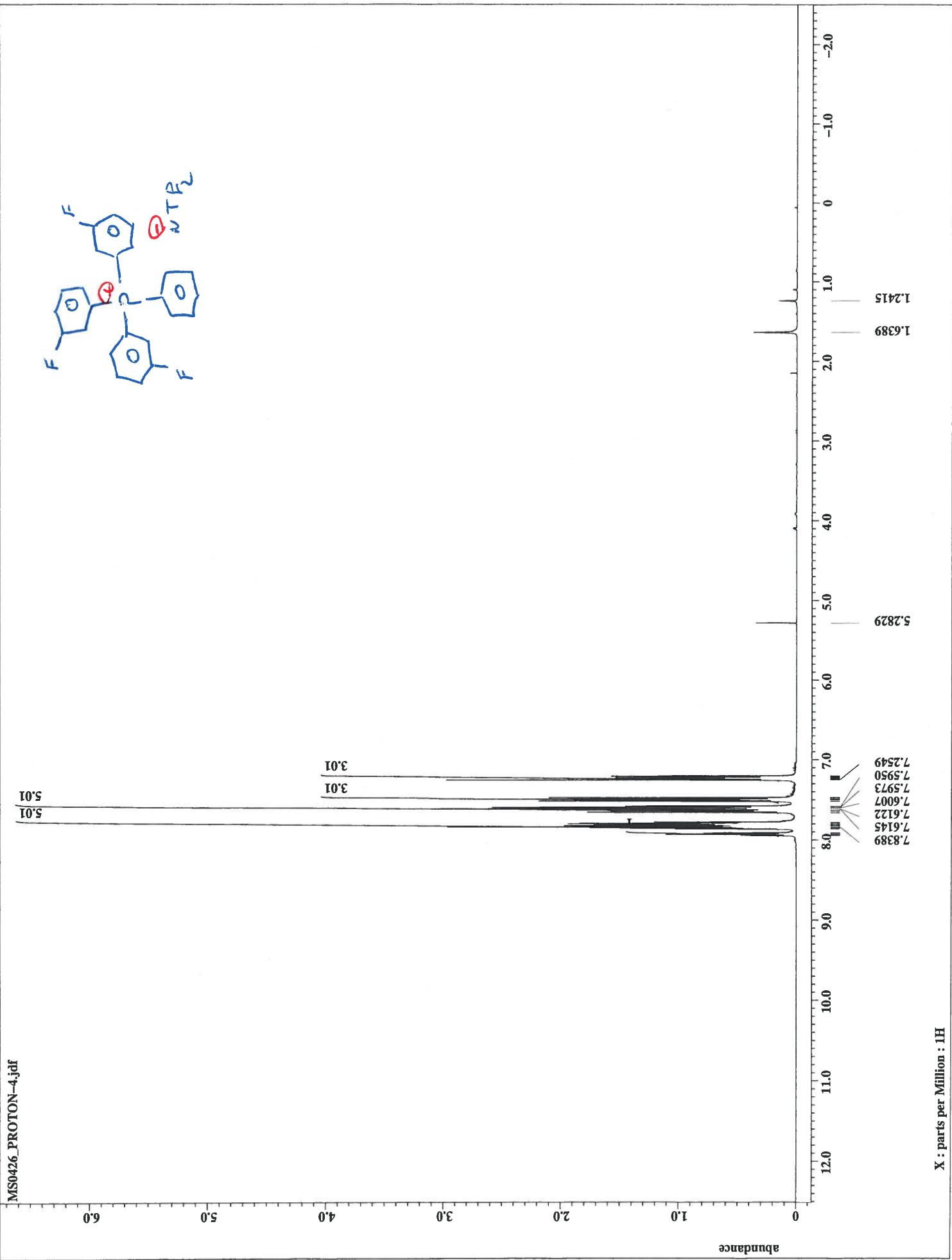
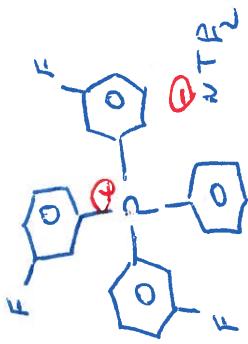
X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Irr_time = 0
Dante_presat = FALSE
Initial_wait = 1 [s]
Recvr_gain = 32
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_set = 22.6 [°C]
  
```

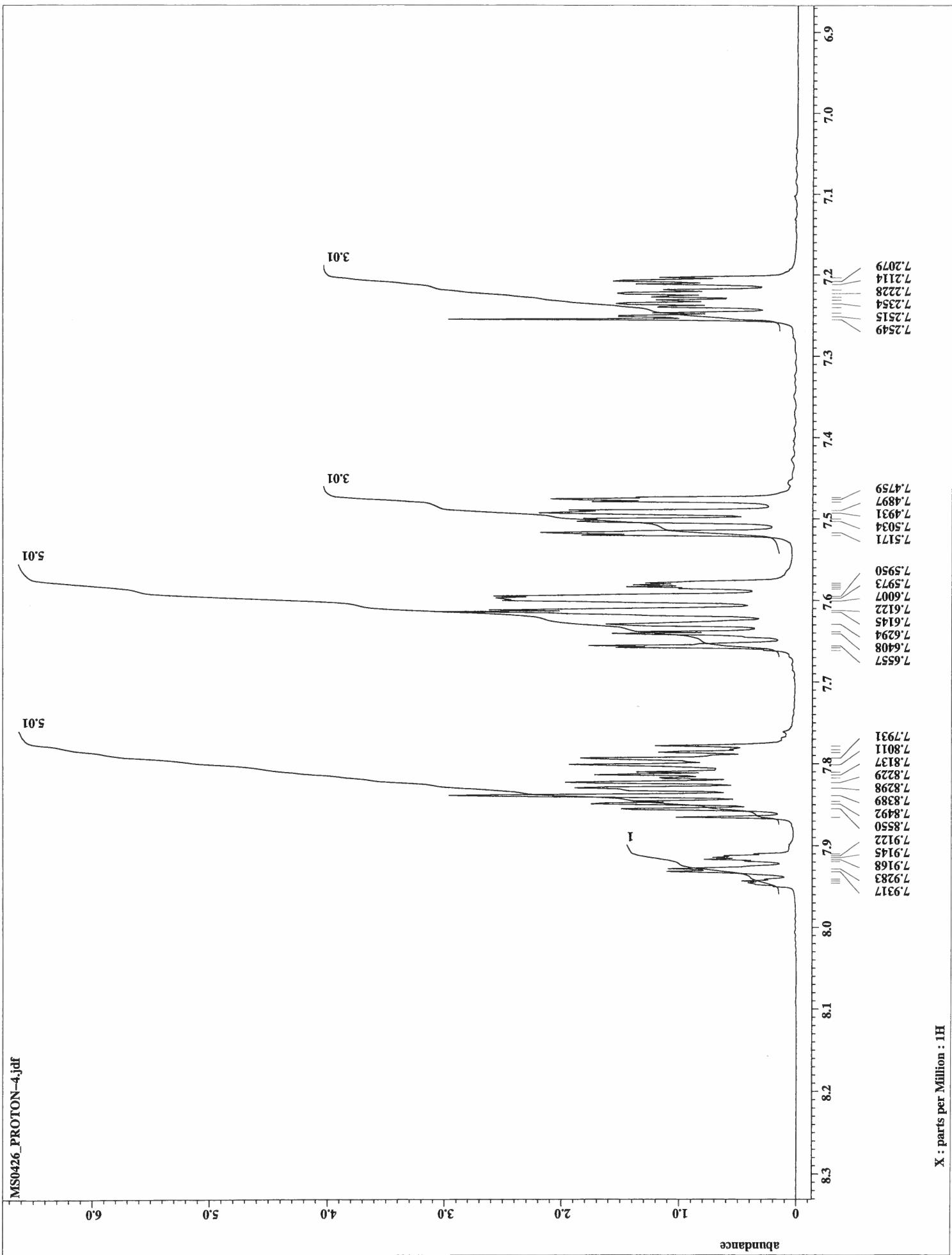


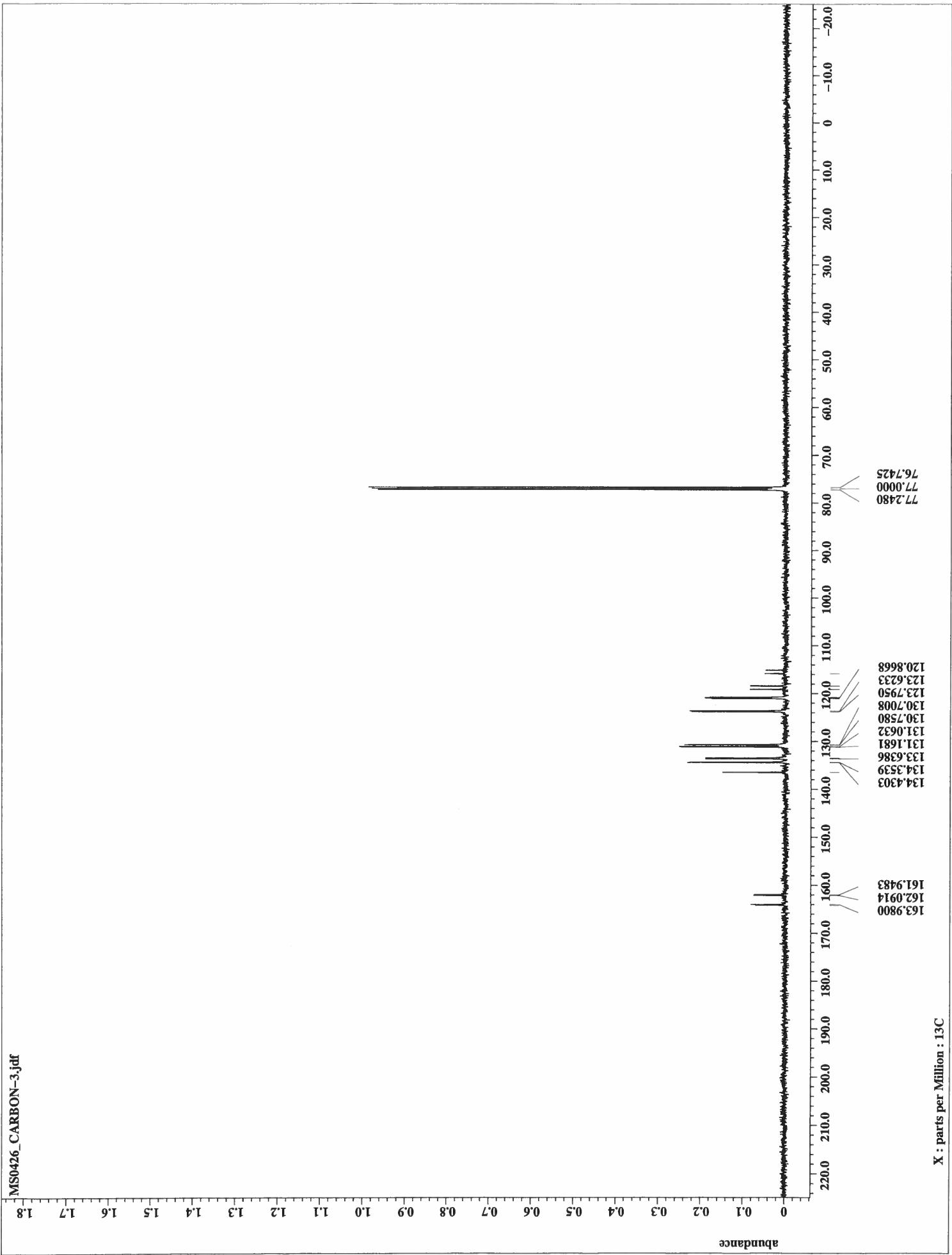
X : parts per Million : 19F

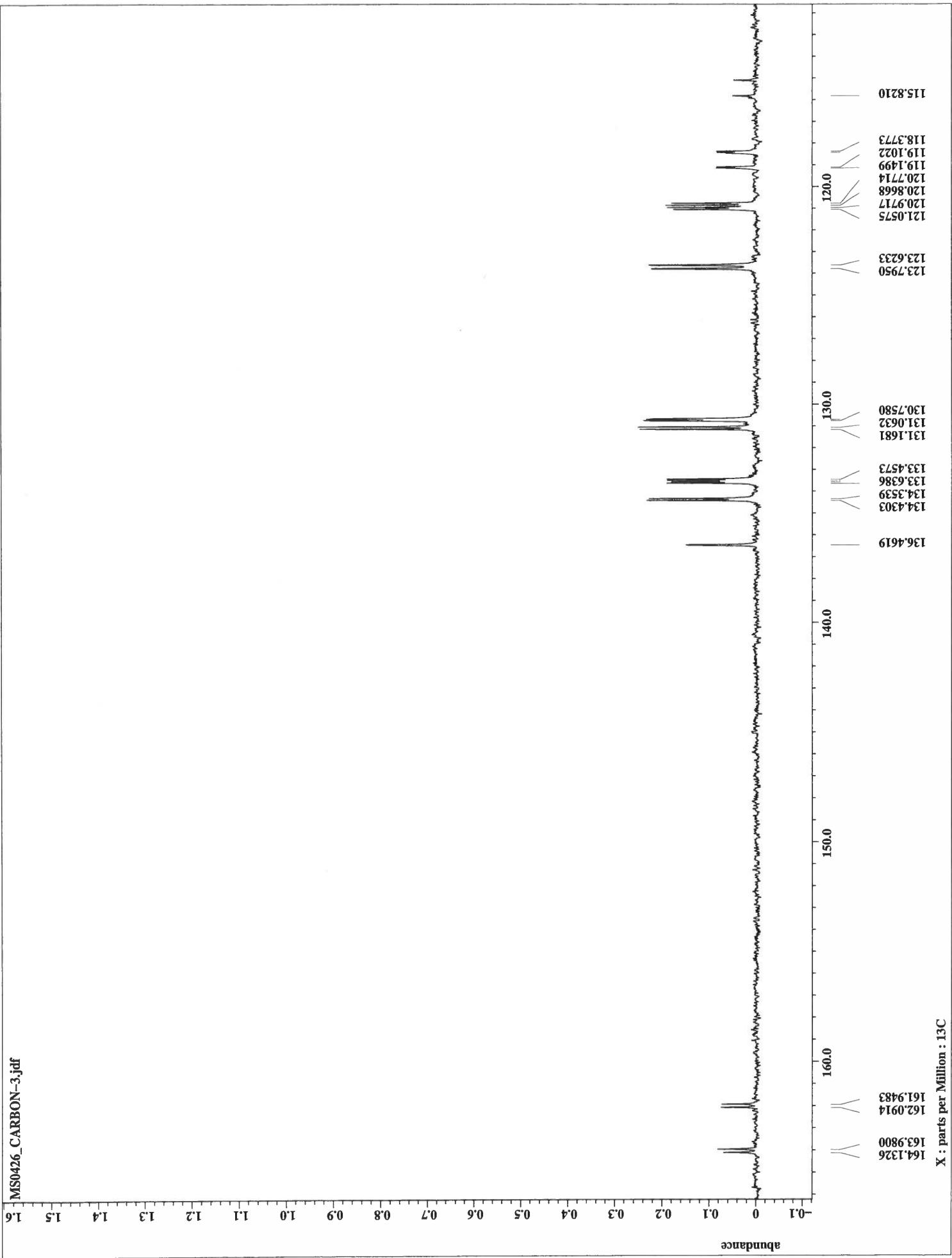
-78.6142

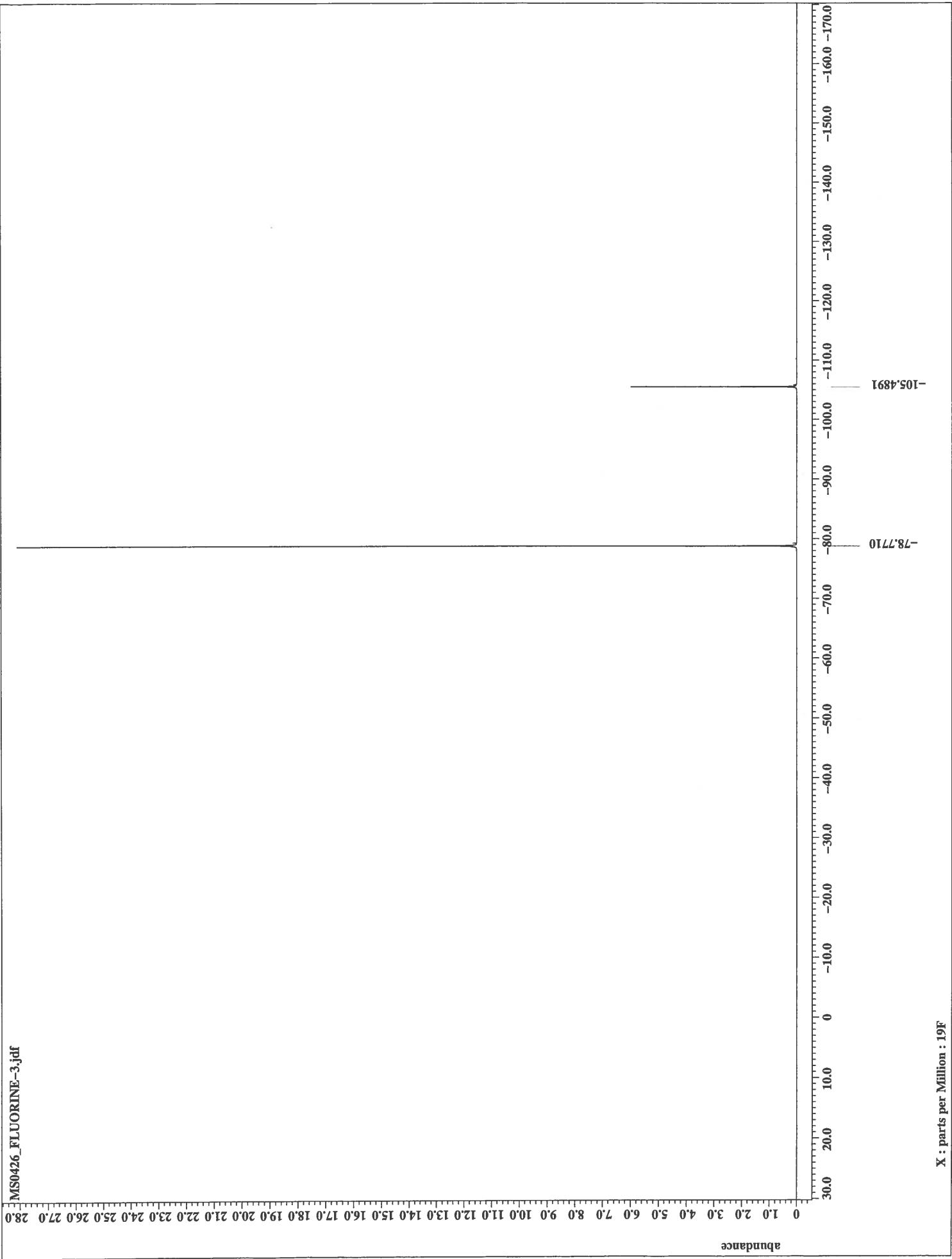
-116.3821











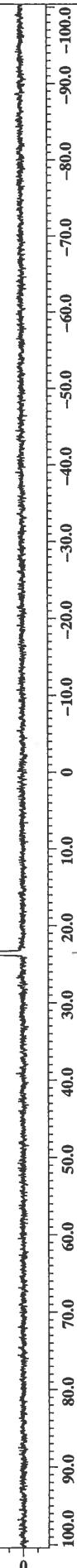
6.0 5.0 4.0 3.0 2.0 1.0 0

X : parts per Million : 31P

abundance

23.5895

23.5589





```

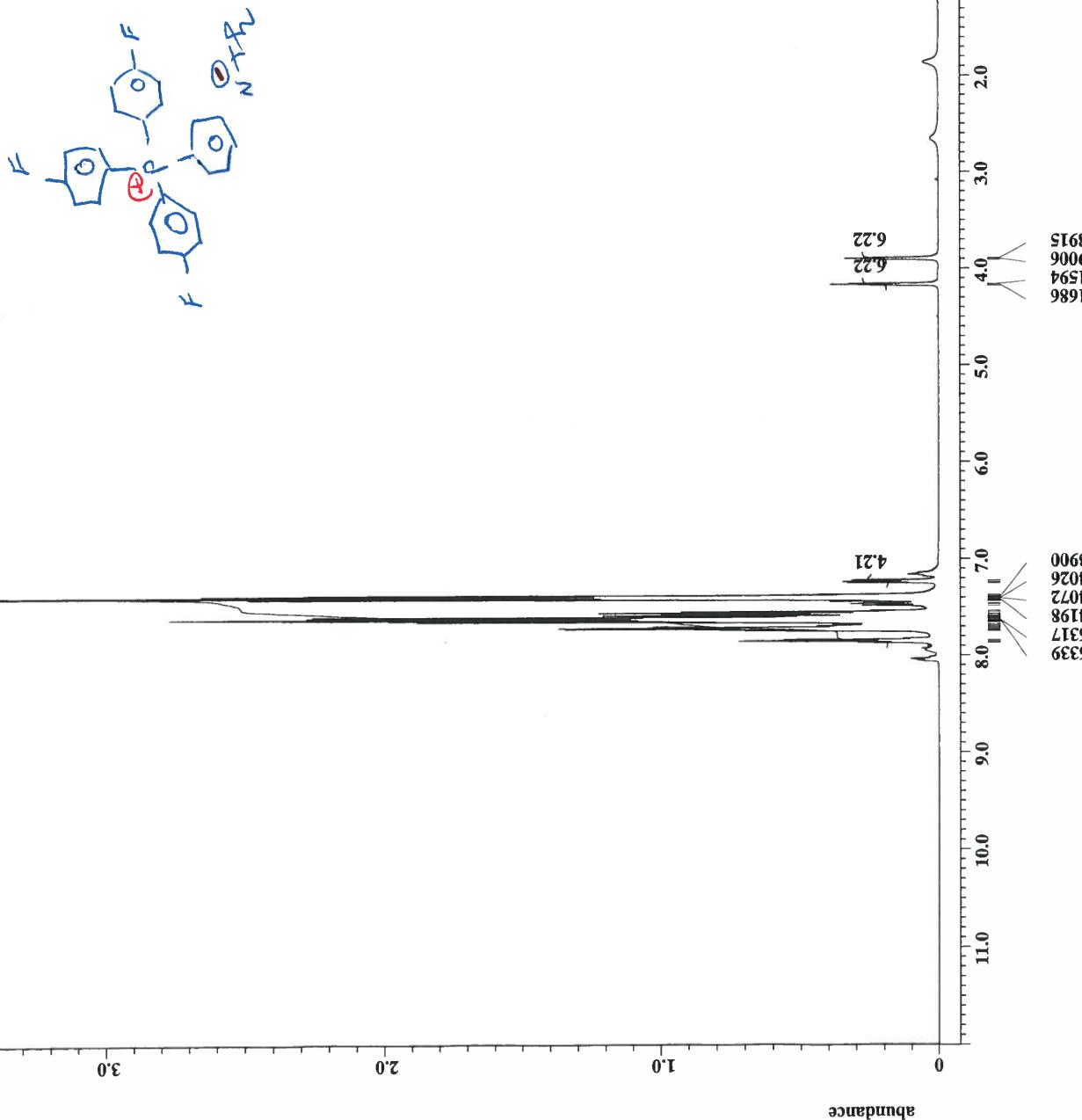
Filename = MS0430_PROTON-2.jdf
Author = Jim Davis
Experiment = single_pulse_ex2
Sample_id = MS0430
Solvent = CHLOROFORM-D
Creation_time = 26-NOV-2019 12:03:37
Revision_time = 26-NOV-2019 11:34:33
Current_time = 26-NOV-2019 11:34:33

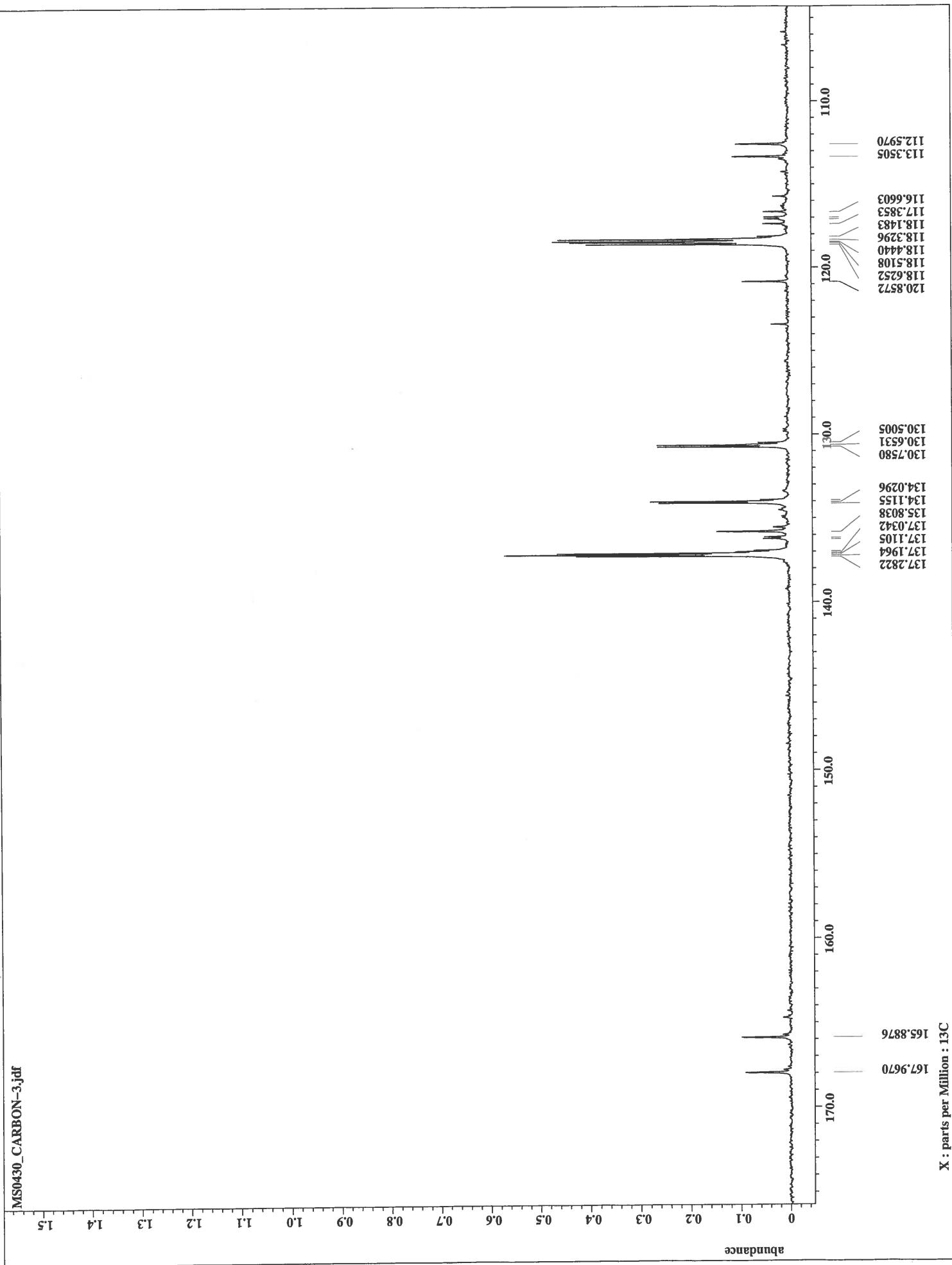
Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

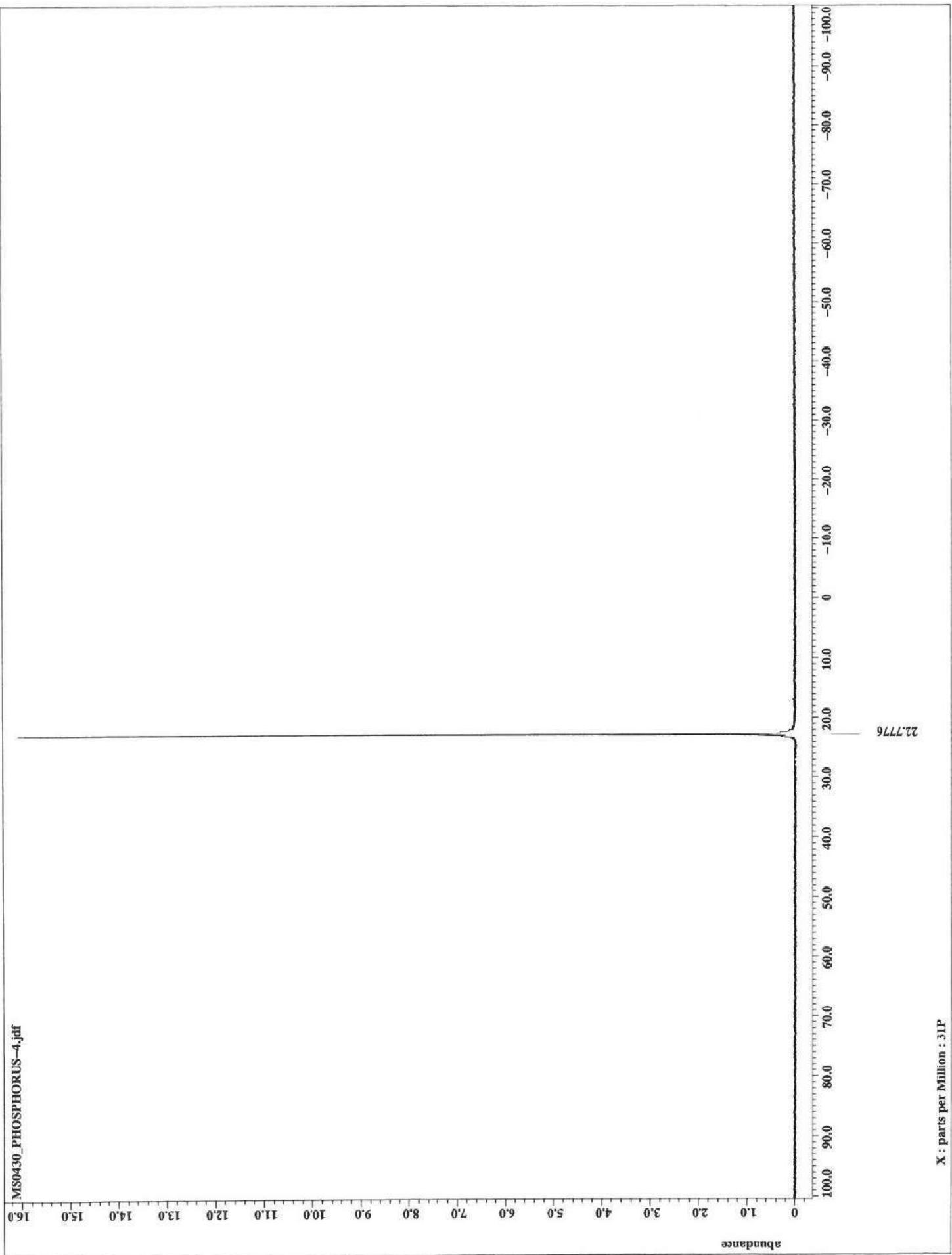
Field_strength = 11.7473579 [MHz] (500 MHz)
X_acq_duration = 1.74587904 [s]
X_domain = 1H
X_freq = 500.15991521 [MHz]
X_offset = 5.0 [ppm]
X_points = 16384
X_precsabs = 1
X_resolution = 0.57277737 [Hz]
X_sweep = 9.38438438 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Tri_domain = 1H
Tri_freq = 500.15991521 [MHz]
Tri_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
total_scans = 16

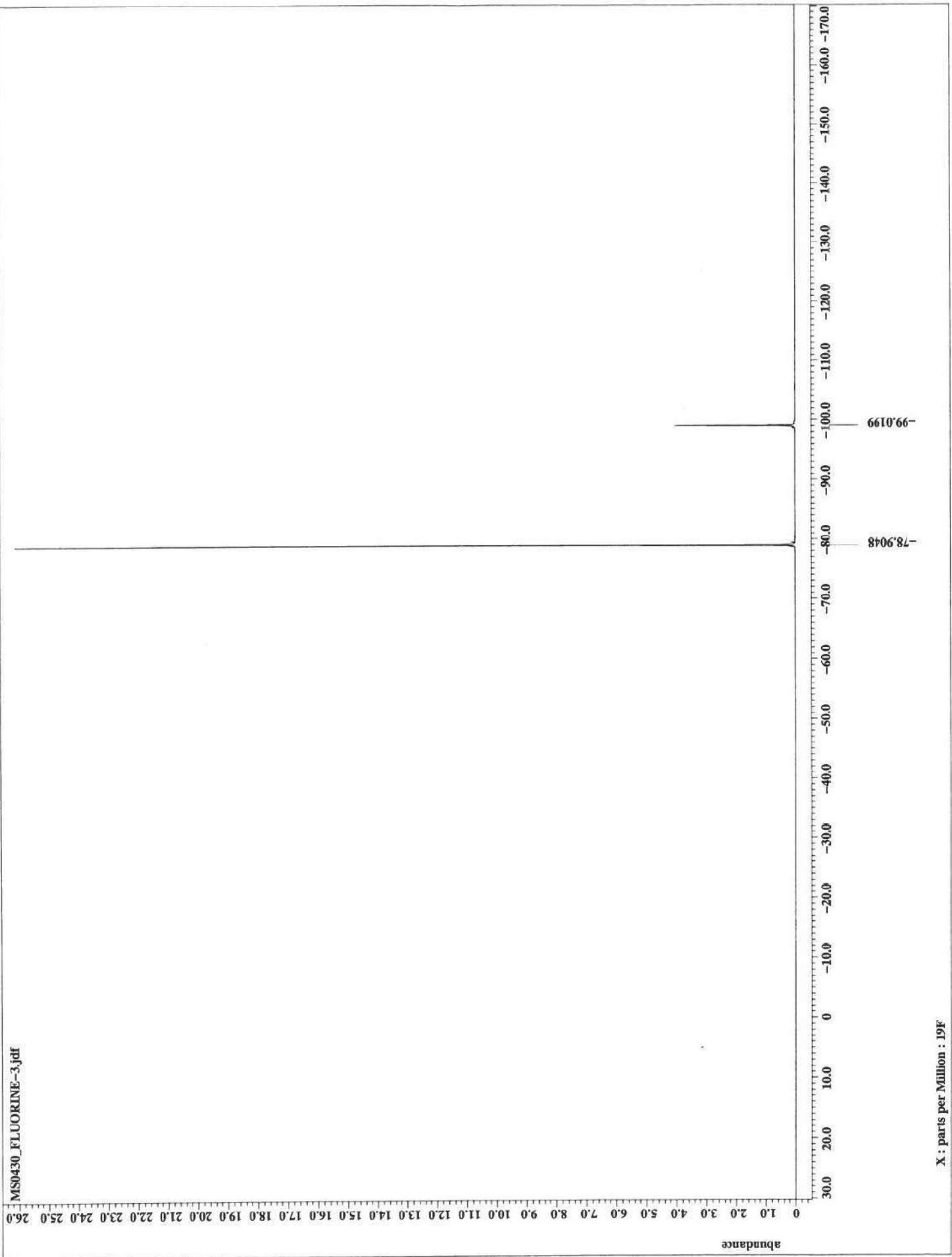
X_90_width = 11.3 [us]
X_acq_time = 1.74587904 [s]
X_angle = 45 [deg]
X_atm = 4 [dB]
X_pulse = 5.65 [us]
Irr_mode = Off
Dante_presat = FALSE
Initial_wait = 1 [s]
Recv_gain = 26
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get = 19.1 [dC]

```











```

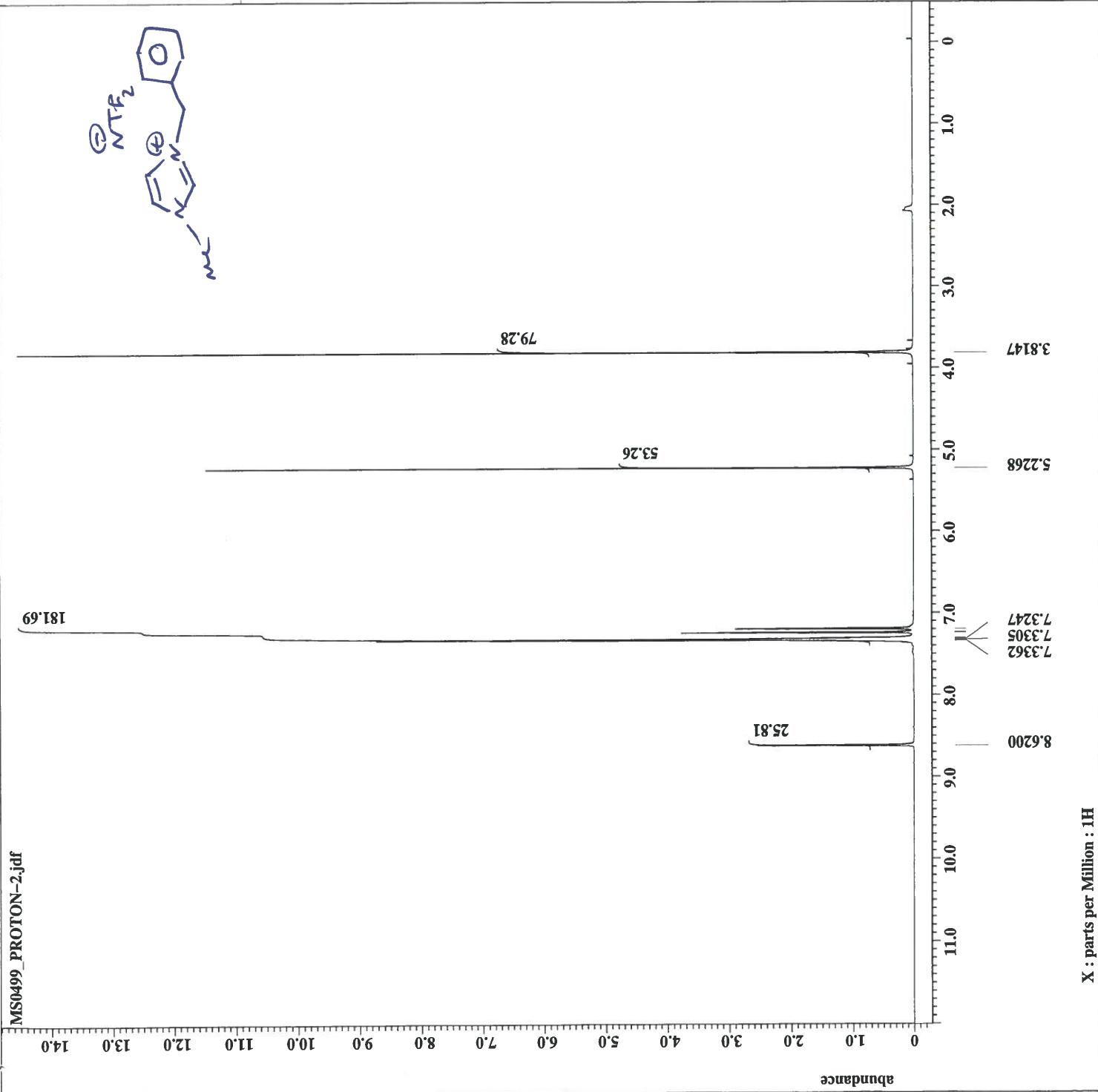
Filename == MS0499_PROTON-2.jdf
Author == Jim Davis
Experiment == single_pulse.ex2
Sample_id == MS0499
Solvent == CHLOROFORM-D
Changer_sample == 9
Creation_time == 12-JUL-2018 18:21:04
Revision_time == 12-JUL-2018 17:58:39
Current_time == 12-JUL-2018 17:58:39

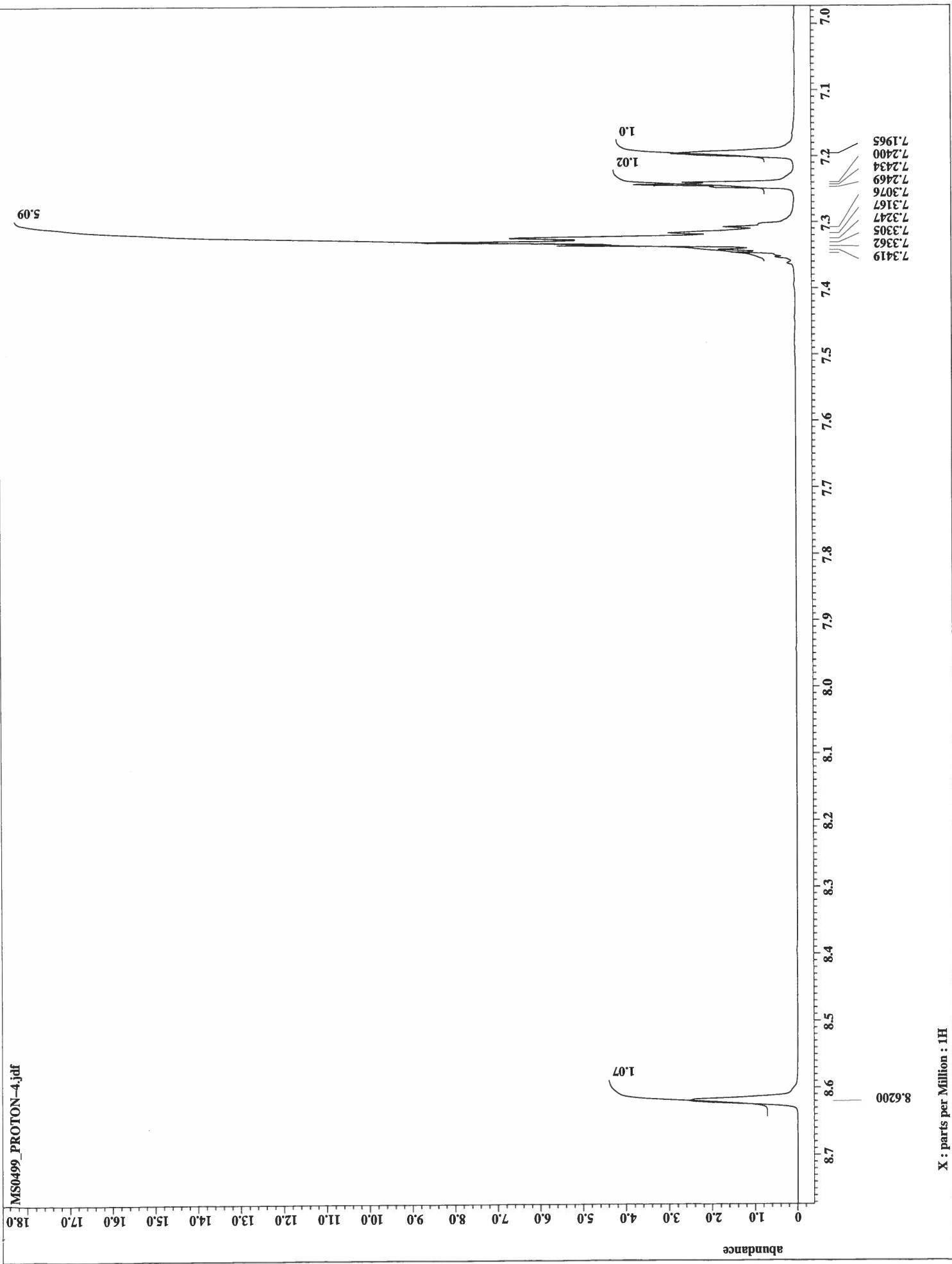
Data_format == 1D COMPLEX
Dim_size == 13107
Dim_title == 1H
Dim_units == [ppm]
Dimensions == X
Site == ECA 500
Spectrometer == JNM-ECA500

Field_strength == 11.7473579 [T] (500 [MHz])
X_acq_duration == 1.7458790 [s]
X_domain == 1H
X_freq == 500.15991521 [MHz]
X_offset == 5.0 [ppm]
X_points == 16384
X_prescans == 1
X_resolution == 0.57277737 [Hz]
X_sweep == 9.38438438 [kHz]
Irr_domain == 1H
Irr_freq == 500.15991521 [MHz]
Irr_offset == 5.0 [ppm]
Tri_domain == 1H
Tri_freq == 500.15991521 [MHz]
Tri_offset == 5.0 [ppm]
Clipped == FALSE
Mod_return == 1
Scans == 16
total_scans == 16

X_90_width == 12.4 [us]
X_acq_time == 1.74587904 [s]
X_angle == 45 [deg]
X_atn == 4 [dB]
X_pulse == 6.2 [us]
Irr_mode == Off
Tri_mode == Off
Dante_preset == FALSE
Initial_preset == 1 [s]
Recv_gain == 24
Relaxation_delay == 4 [s]
Relaxation_time == 5.74587904 [s]
Temp_get == 22.7 [dc]

```







```

Filename      = MS0499_CARBON-2.jdf
Author        = Jim Davis
Experiment   = single_pulse_dec
Sample_id    = MS0499
Solvent       = CHLOROFORM-D
Changer_sample = 9
Creation_time = 12-JUL-2018 18:33:38
Revision_time = 12-JUL-2018 18:11:15
Current_time  = 12-JUL-2018 18:11:15

Date_format   = 1D COMPLEX
Dim_size      = 26214
Dim_title     = 13C
Dim_units     = [ppm]
Dimensions    = X
Site          = ECA_500
Spectrometer  = JNM-ECA500

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_precsams    = 4
X_resolution  = 1.19959034 [Hz]
X_sweep        = 39.3081761 [kHz]
Irr_domain.n = 1H
Irr_freq       = 500.15991521 [MHz]
Irr_offset     = 5.0 [ppm]
Clipped        = FALSE
Mod_return    = 1
Scans          = 256
Total_scans   = 256

X_90_width    = 13.2 [us]
X_acq_time    = 0.83361792 [s]
X_angle        = 30 [deg]
X_attn        = 6 [dB]
X_pulse        = 4.4 [us]
Irr_attn_dec  = 20.7 [dB]
Irr_attn_noe  = 20.7 [dB]
Irr_noise      = WALTZ
Decoupling     = TRUE
Initial_wait   = 1 [s]
Noe           = TRUE
Noe_time       = 2 [s]
Recvr_gain    = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get      = 23.2 [dc]

```



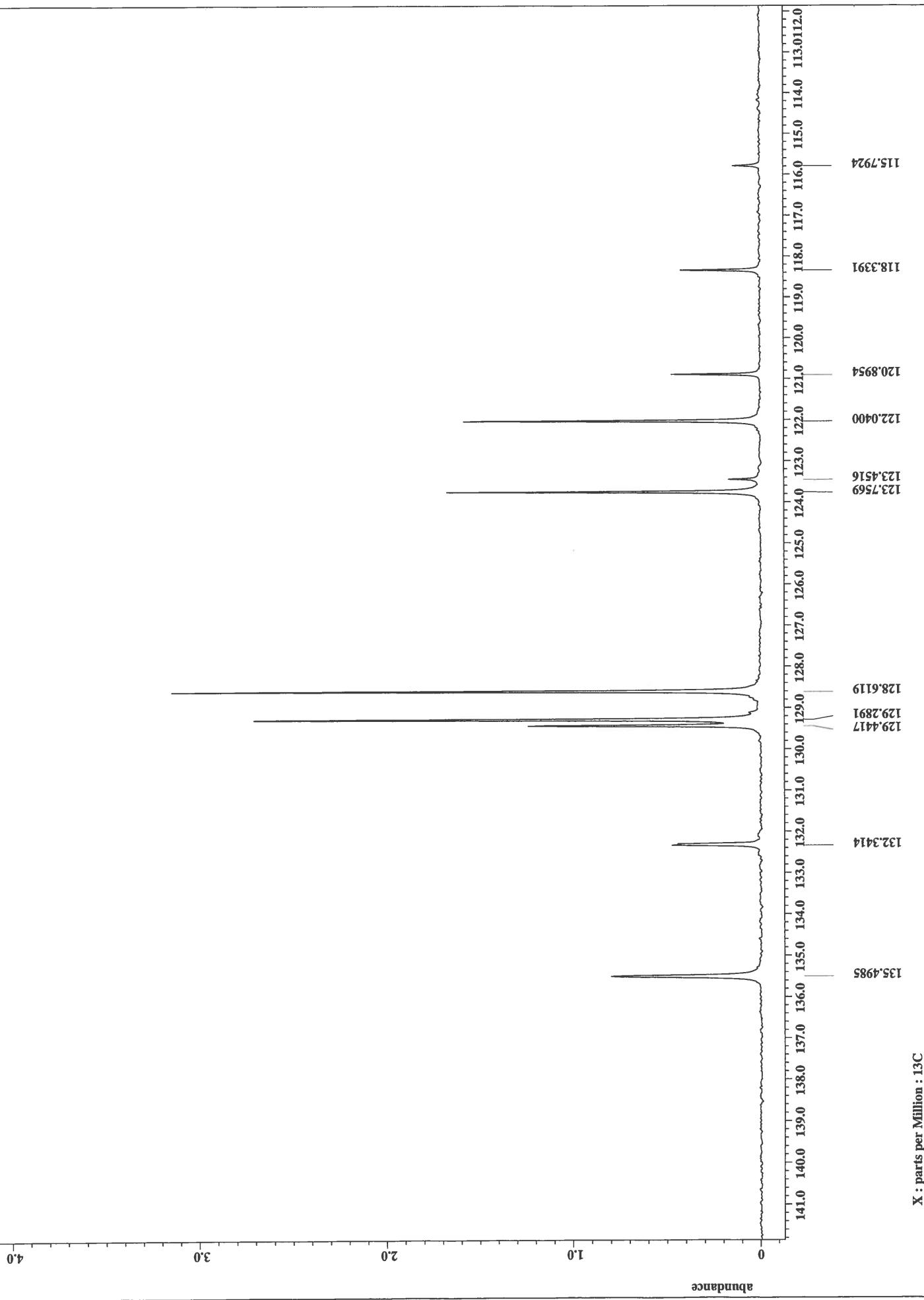
X : parts per Million : 13C

135.4985
 132.3414
 129.4417
 129.2891
 128.6119
 123.7569
 122.0400
 120.9554

77.0000
 77.2575

35.9948

53.1923





MS0499 FLUORINE-2.jdf

```

Filename = MS0499_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0499
Solvent = CHLOROFORM-D
Changer_sample = 9
Creation_time = 12-JUL-2018 16:38:27
Revision_time = 12-JUL-2018 16:16:04
Current_time = 12-JUL-2018 16:16:04

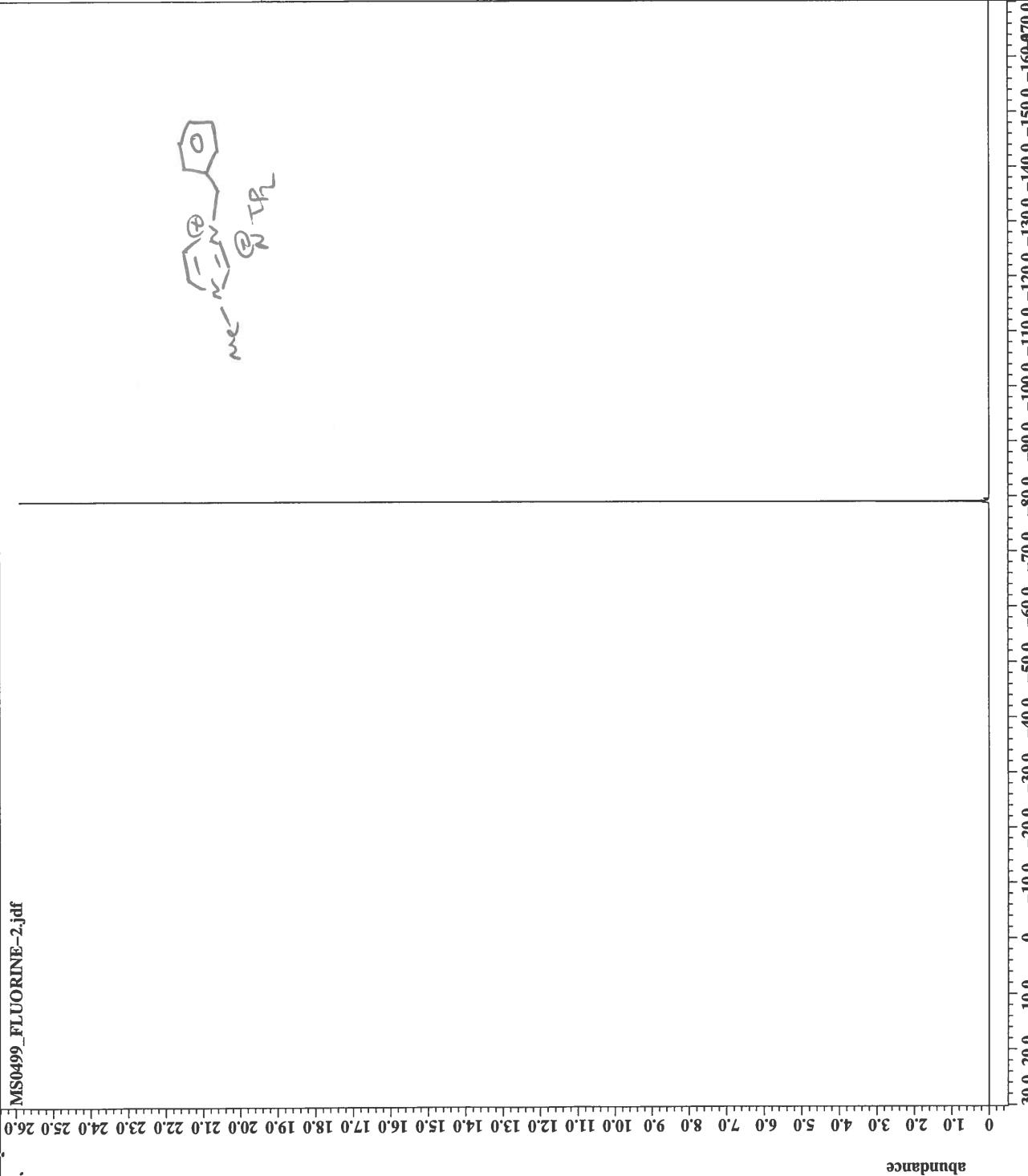
Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = DPPM
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_Precans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [Hz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Recv_gain = 30

Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.5 [dc]

```



X : parts per Million : 19F

-79.0271



```

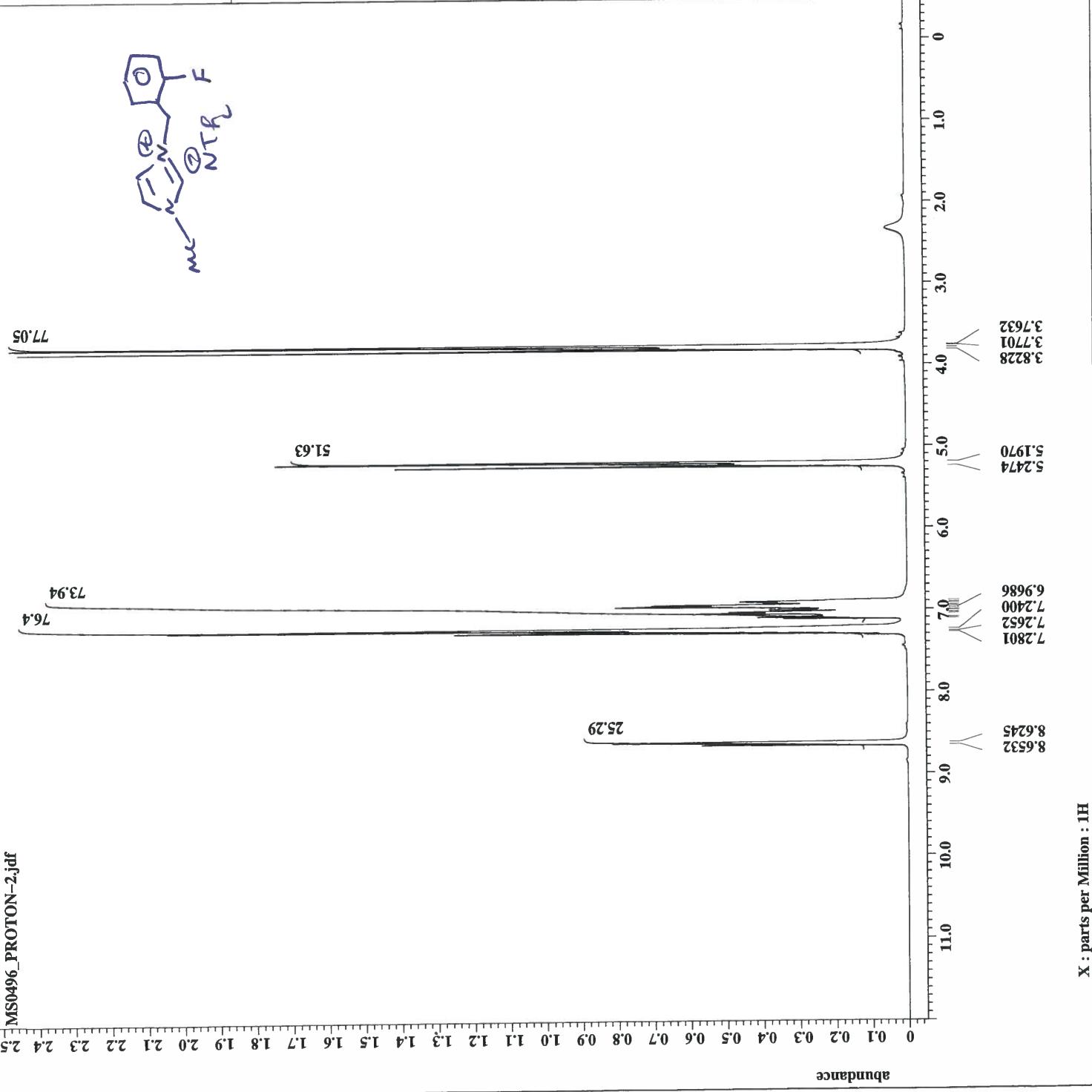
Filename = MS0496_PROTON-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0496
Solvent = CHLOROFORM-D
Changer_sample = 6
Creation_time = 12-JUL-2018 17:39:41
Revision_time = 12-JUL-2018 17:17:17
Current_time = 12-JUL-2018 17:17:17

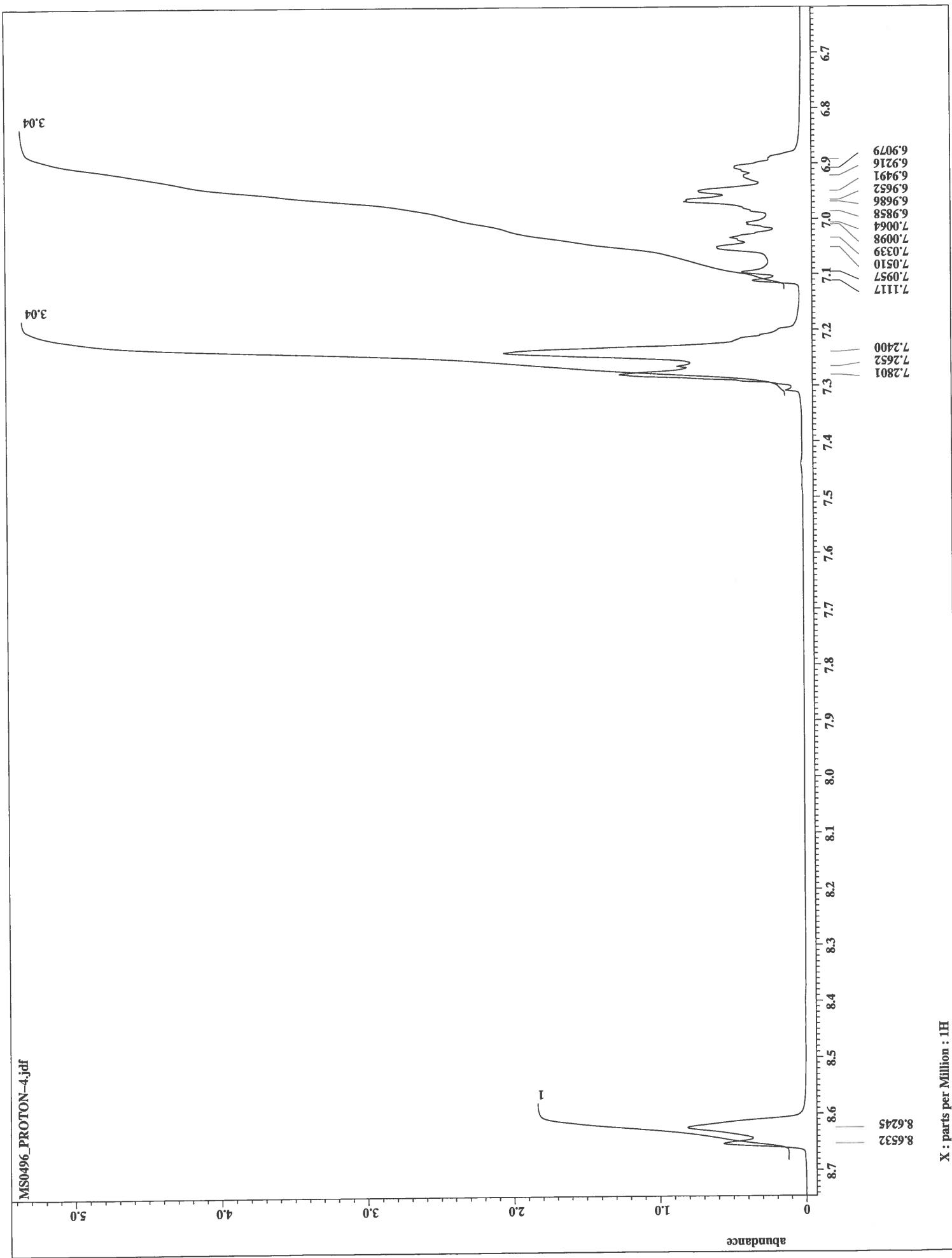
Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 1.74587904 [s]
X_domain = 1H
X_freq = 500.15991521 [MHz]
X_offset = 5.0 [ppm]
X_points = 16384
X_prescans = 1
X_resolution = 0.57277737 [Hz]
X_sweep = 9.38438438 [Hz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Tri_domain = 1H
Tri_freq = 500.15991521 [MHz]
Tri_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 12.4 [us]
X_acq_time = 1.74587904 [s]
X_angle = 45 [deg]
X_atn = 4 [dB]
X_pulse = 6.2 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Recv_gain = 22
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get = 22.5 [degC]

```







```

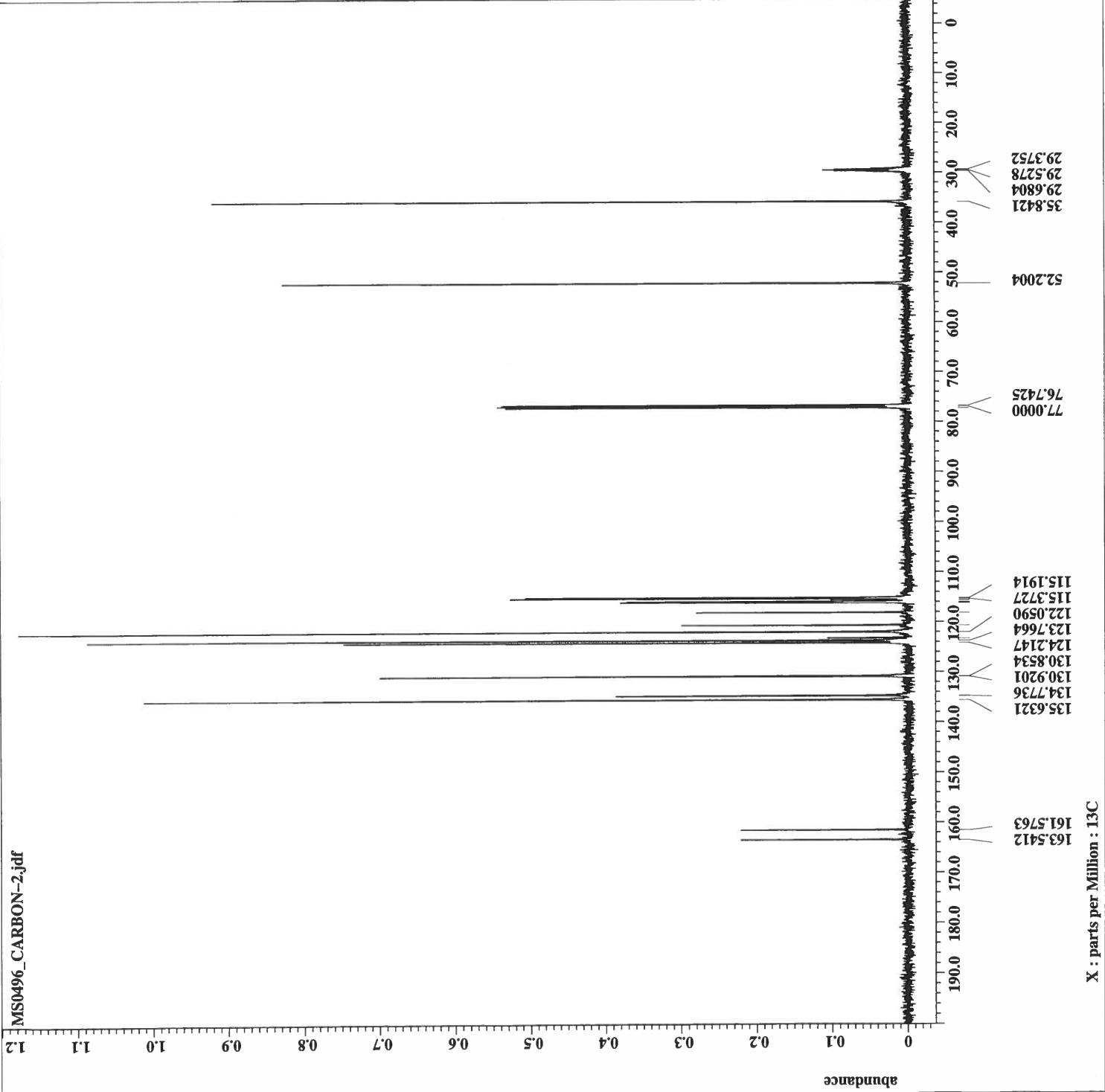
filename = MS0496_CARBON-2.jdf
author = Jim Davis
experiment = single_pulse_dec
sample_id = MS0496
solvent = CHLOROFORM-D
changer_sample = 6
creation_time = 12-JUL-2018 17:54:02
revision_time = 12-JUL-2018 17:31:38
current_time = 12-JUL-2018 17:31:38

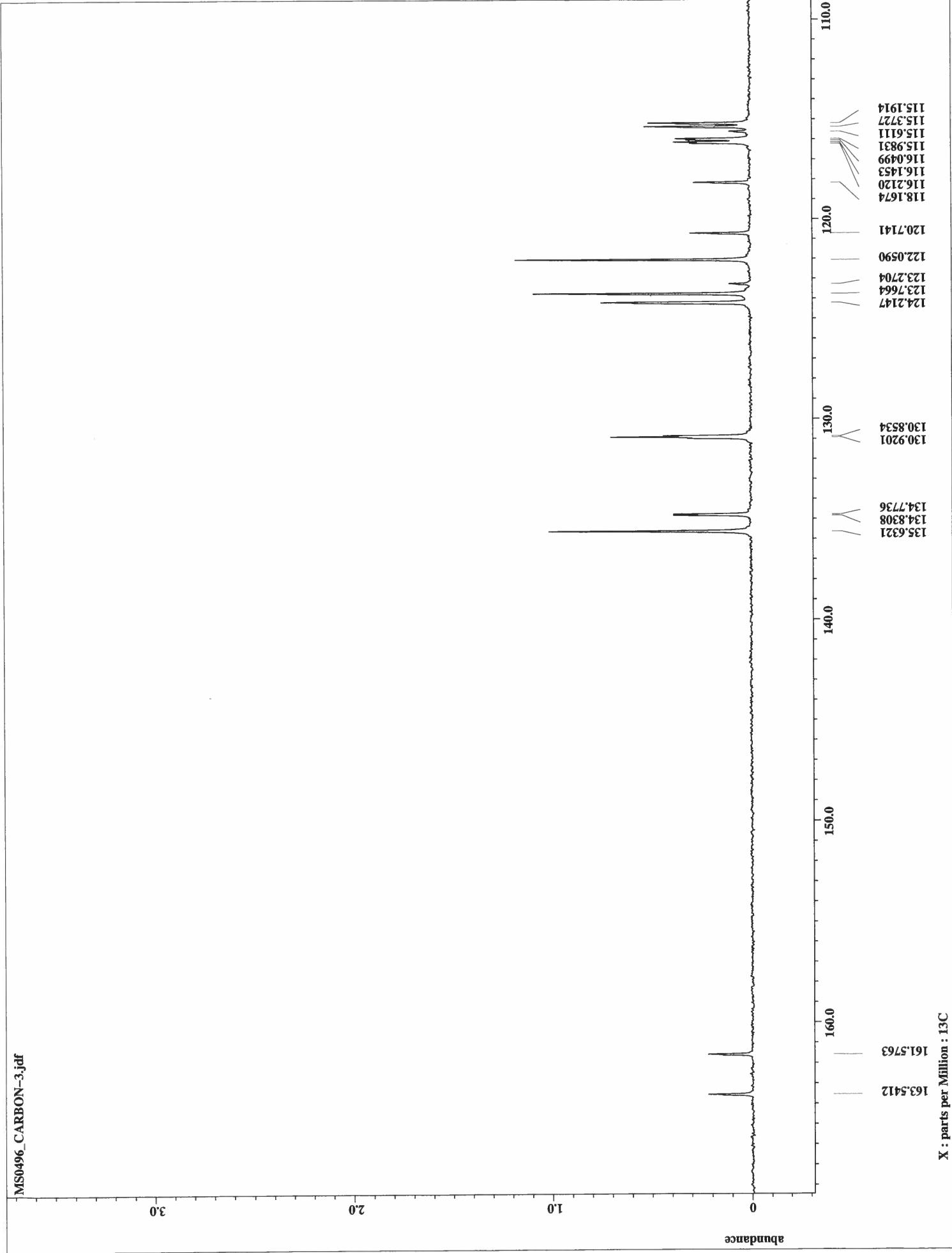
data_format = 1D COMPLEX
dim_size = 26214
dim_title = 13C
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = JEOL-ECA500

field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain = 13C
X_freq = 128.76529768 [MHz]
X_offset = 100.0 [ppm]
X_points = 32768
X_prescans = 4
X_resolution = 1.19959034 [Hz]
X_sweep = 39.3082761 [Hz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 256
total_scans = 256

X_90_width = 13.2 [us]
X_acq_time = 0.83361792 [s]
X_angle = 30 [deg]
X_atn = 6 [dB]
X_pulse = 4.4 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
Nose = TRUE
Nose_time = 2 [s]
Regrv_grain = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get = 23 [DC]

```







```

filename = MS0496_FLUORINE-2.jdf
author = Jim Davis
experiment = single_pulse-ex2
sample_id = MS0496
solvent = CHLOROFORM-D
changer_sample = 6
creation_time = 12-JUL-2018 16:18:09
revision_time = 12-JUL-2018 15:55:46
current_time = 12-JUL-2018 15:55:46

data_format = 1D COMPLEX
dim_size = 52428
dim_title = 19F
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = JNM-ECA500

field_strength = 11.7473579 [T] (500 [MHz])
x_acq_duration = 0.55574528 [s]
x_domain = 19F
x_freq = 470.62046684 [MHz]
x_offset = -70 [ppm]
x_points = 65536
x_prescans = 1
x_resolution = 1.7993855 [Hz]
x_sweep = 117.924523 [kHz]
irr_domain = 19F
irr_freq = 470.62046684 [MHz]
irr_offset = 5 [ppm]
tri_domain = 19F
tri_freq = 470.62046684 [MHz]
tri_offset = 5 [ppm]
clipped = FALSE
mod_return = 1
scans = 16
total_scans = 16

x_90_width = 13.1 [us]
x_acq_time = 0.55574528 [s]
x_angle = 45 [deg]
x_atn = 2.5 [dB]
x_pulse = 6.55 [us]
irr_mode = Off
tri_mode = Off
dante_preset = FALSE
initial_wait = 1 [s]
recvr_gain = 26
relaxation_delay = 4 [s]
repetition_time = 4.55574528 [s]
temp_get = 22.5 [DC]

```





```

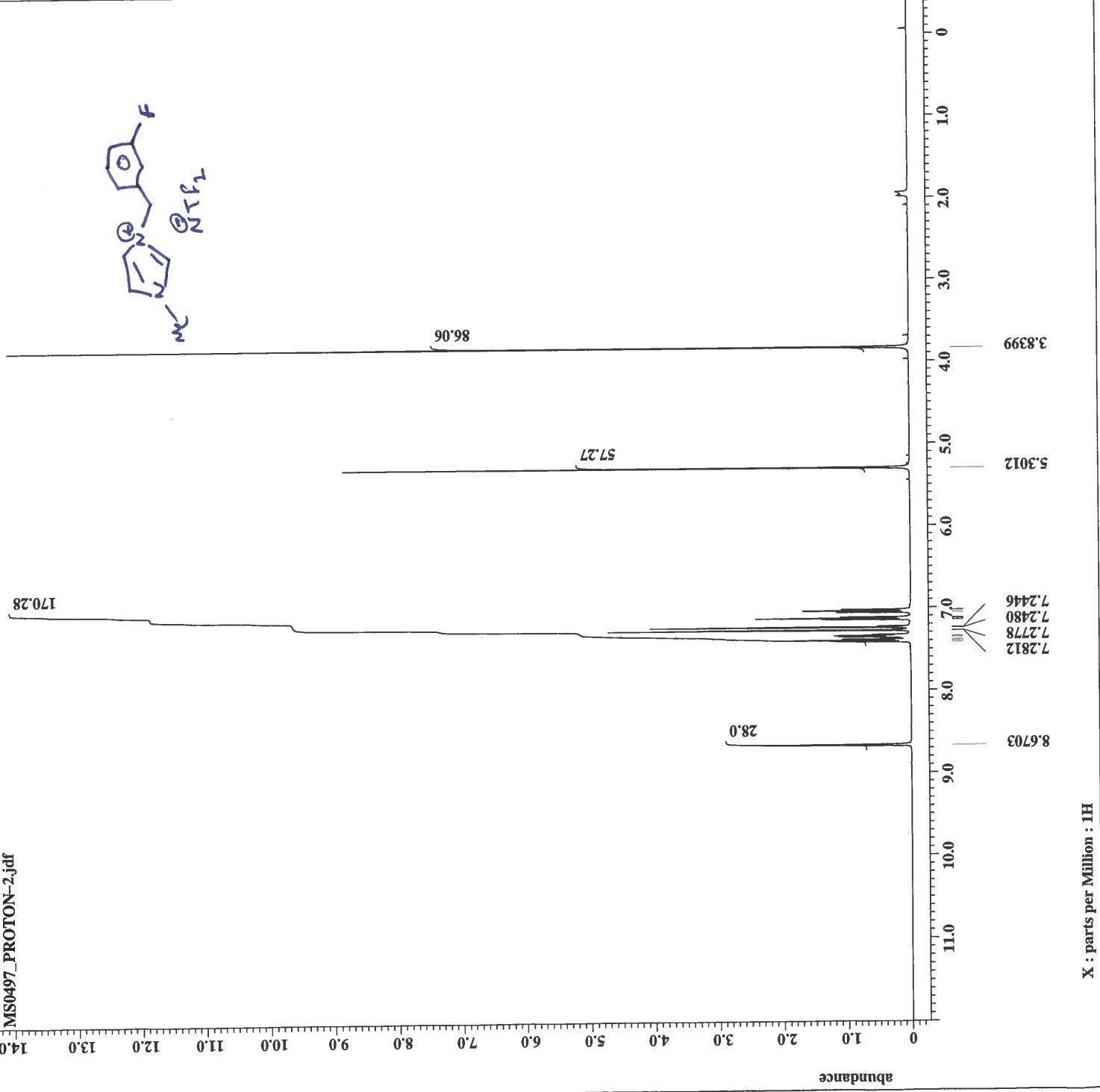
= MS0497_PROTON-2.jdf
= Jim Davis
= single_pulse.ex2
Experiment      = MS0497
Sample_id       = CHLOROFORM-D
Solvent         = 7
Changer_sample = 12-JUL-2018 18:01:19
Creation_time   = 12-JUL-2018 17:38:55
Revision_time   = 12-JUL-2018 17:38:55
Current_time    = 12-JUL-2018 17:38:55

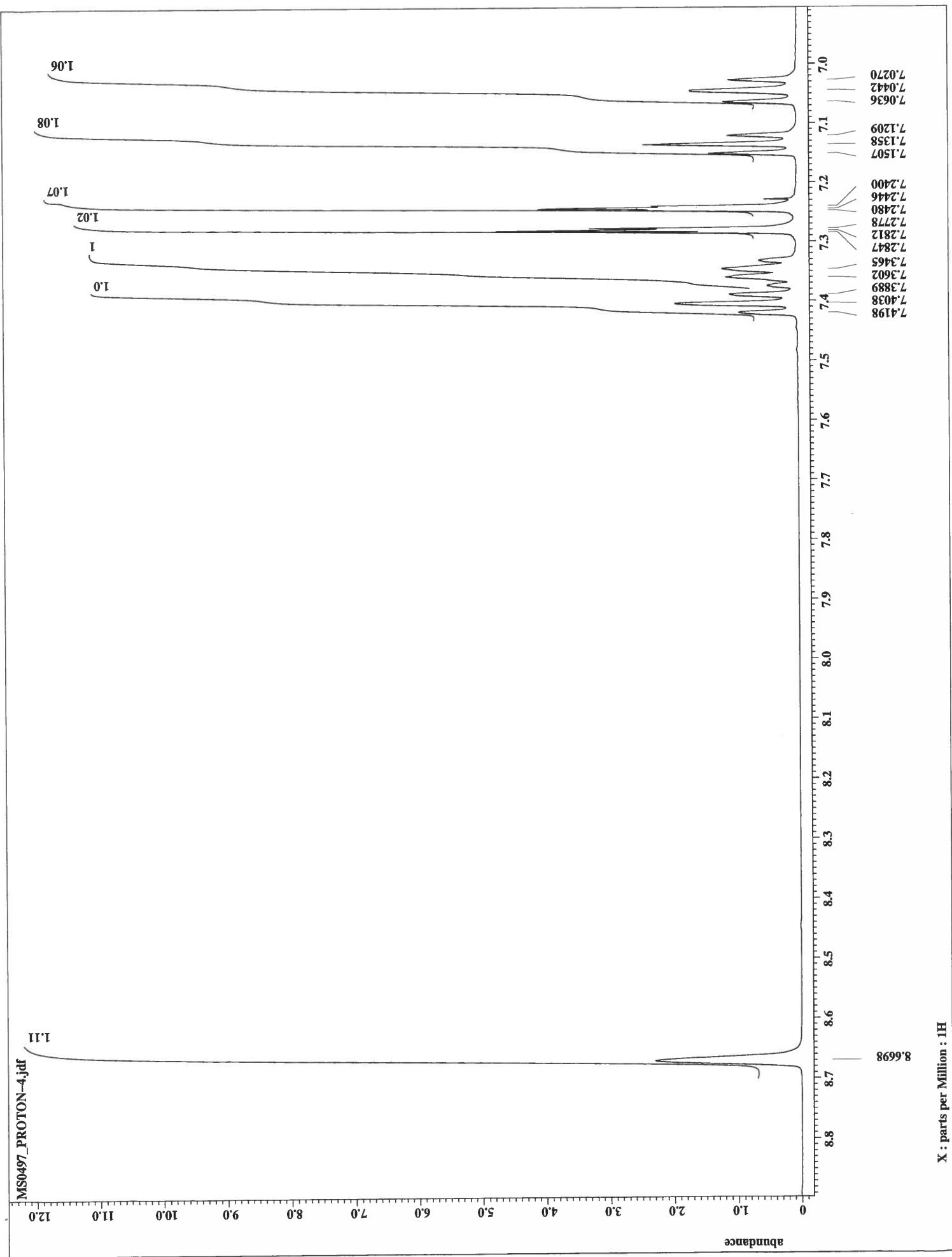
Data_format     = 1D COMPLEX
Dim_size        = 13107
Dim_title       = 1H
Dim_units       = [ppm]
Dimensions      = X
Site            = ECA 500
Spectrometer    = JNM-ECA500

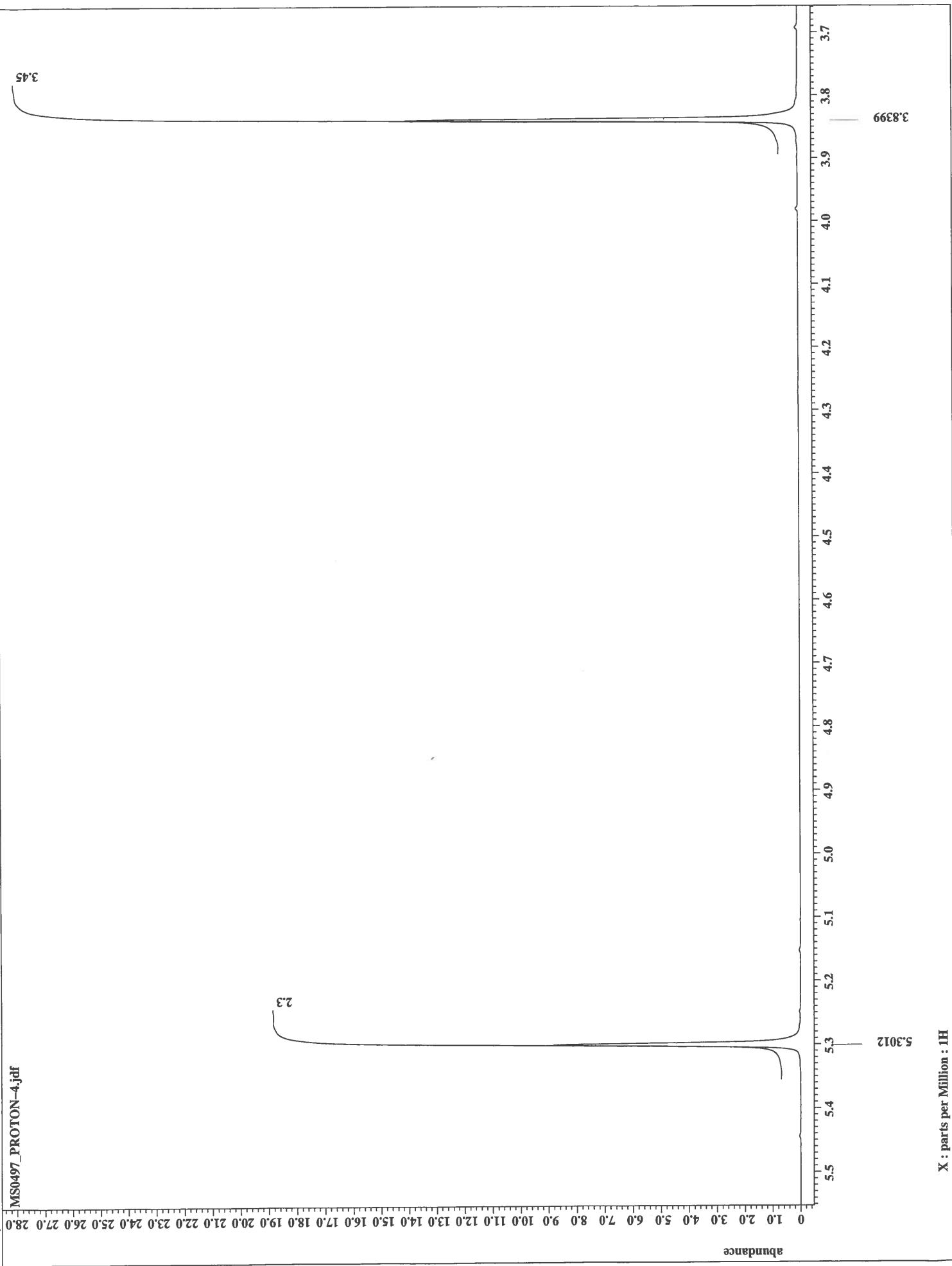
Field_strength  = 11.7473579 [T] (500 [MHz])
X_acq_duration = 1.74587904 [s]
X_domain        = 1H
X_freq          = 500.15991521 [MHz]
X_offset         = 5.0 [ppm]
X_points         = 16384
X_prescans      = 1
X_resolution    = 0.57277737 [Hz]
X_sweep          = 9.3843838 [Hz]
Irr_domain      = 1H
Irr_freq         = 500.15991521 [MHz]
Irr_offset       = 5.0 [ppm]
Tri_domain      = 1H
Tri_freq         = 500.15991521 [MHz]
Tri_offset       = 5.0 [ppm]
Clipped          = FALSE
Mod_return      = 1
Scans            = 16
Total_scans     = 16

X_90_width      = 12.4 [us]
X_acq_time      = 1.74587904 [s]
X_angle          = 45 [deg]
X_atn            = 4 [dB]
X_pulse          = 6.2 [us]
Irr_mode         = Off
Tri_mode         = Off
Dante_pressat   = FALSE
Initial_wait     = 1 [s]
Recv_Gain        = 28
Relaxation_delay = 4 [s]
Repetition_time  = 5.74587904 [s]
Temp_Get         = 22.7 [idc]

```









```

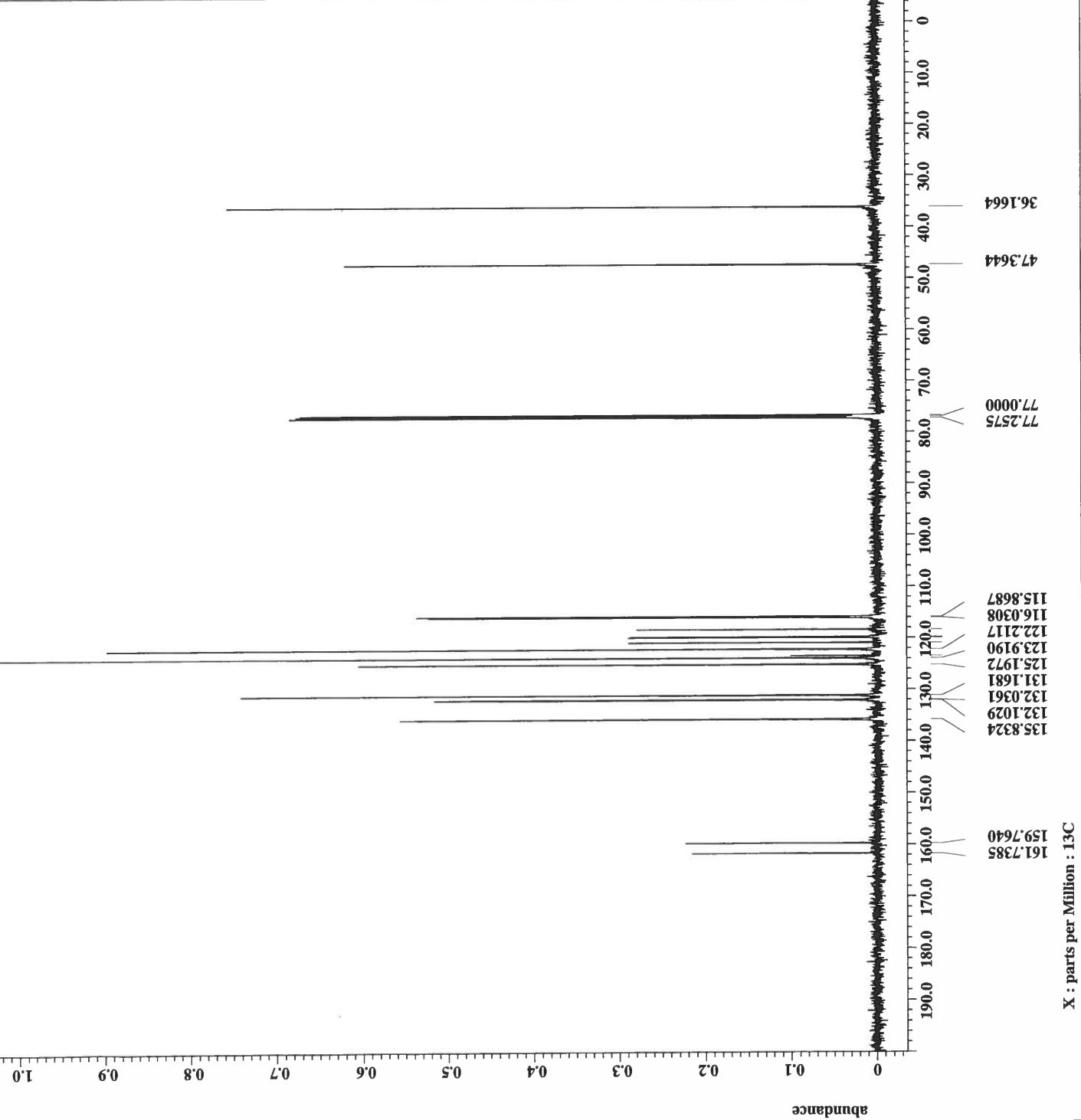
Filename           = MS0497_CARBON-2.jdf
Author            = Jim Davis
Experiment        = single_pulse_dec
Sample_id         = MS0497
Solvent           = CHLOROFORM-D
Changer_sample   = 7
Creation_time    = 12-JUL-2018 18:13:54
Revision_time    = 12-JUL-2018 17:51:29
Current_time     = 12-JUL-2018 17:51:29

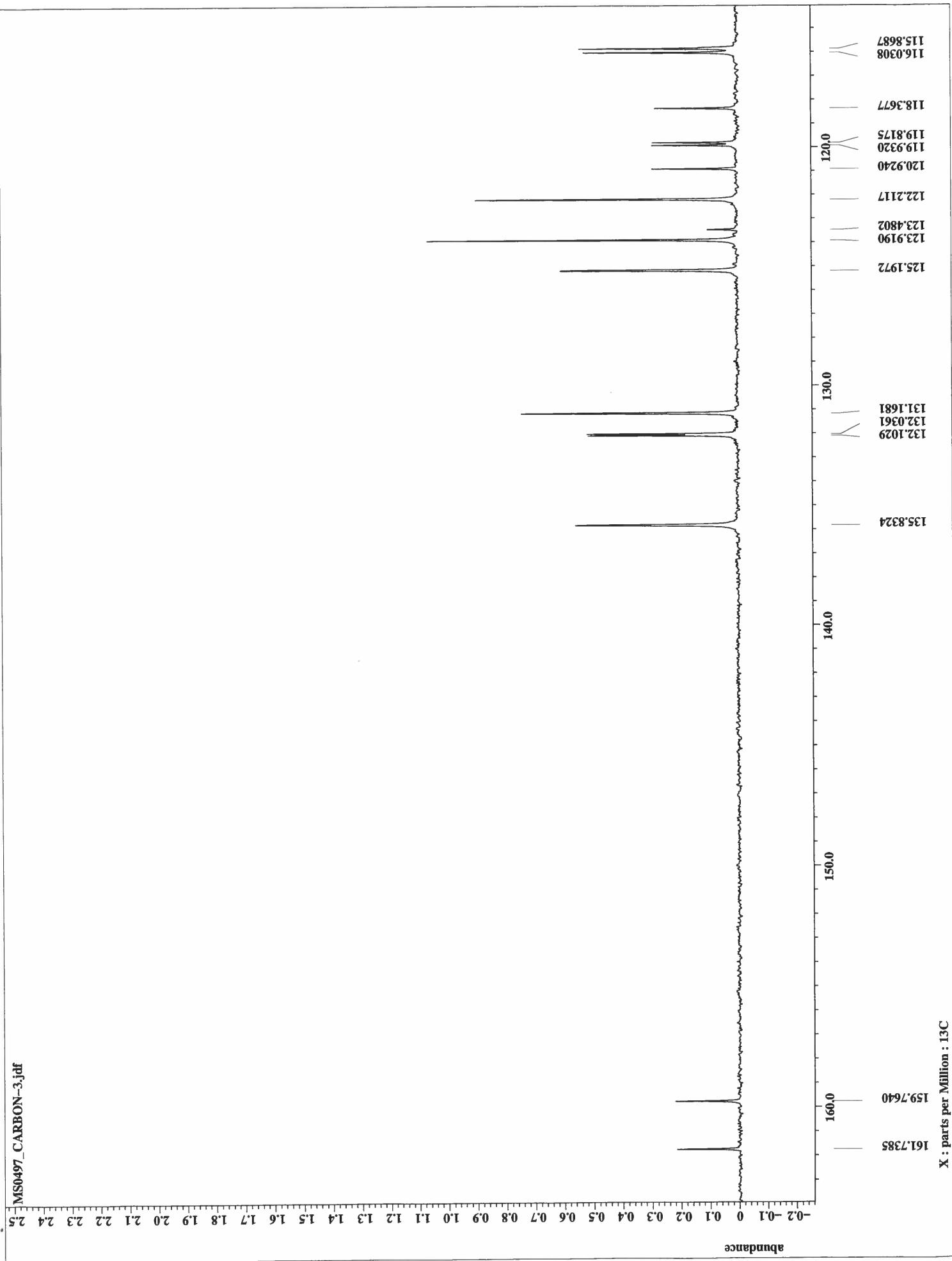
Data_format       = 1D COMPLEX
Dim_size          = 26214
Dim_title         = 13C
Dim_units         = [ppm]
Dimensions        = X
Site              = ECA 500
Spectrometer      = JNM-ECAS500

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]
Irr_domain       = 1H
Irr_freq          = 500.15991521 [MHz]
Irr_offset         = 5.0 [ppm]
Clipped           = FALSE
Mod_return        = 1
Scans             = 256
Total_scans       = 256

X_90_width       = 13.2 [us]
X_acq_time       = 0.83361792 [s]
X_angle           = 30 [deg]
X_atn             = 6 [dB]
X_pulse           = 4.4 [us]
Irr_atn_dec       = 20.7 [dB]
Irr_atn_noe       = 20.7 [dB]
Irr_noise          = WALTZ
Decoupling        = TRUE
Initial_wait      = 1 [s]
Noe               = TRUE
Noe_time          = 2 [s]
Recvr_gain        = 60
Relaxation_delay = 2 [s]
Repetition_time   = 2.83361792 [s]
Temp_get          = 23.1 [dc]

```







```

Filename = MS0497_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0497
Solvent = CHLOROFORM-D
Changer_sample = 7
Creation_time = 12-JUL-2018 16:25:02
Revision_time = 12-JUL-2018 16:02:38
Current_time = 12-JUL-2018 16:02:38

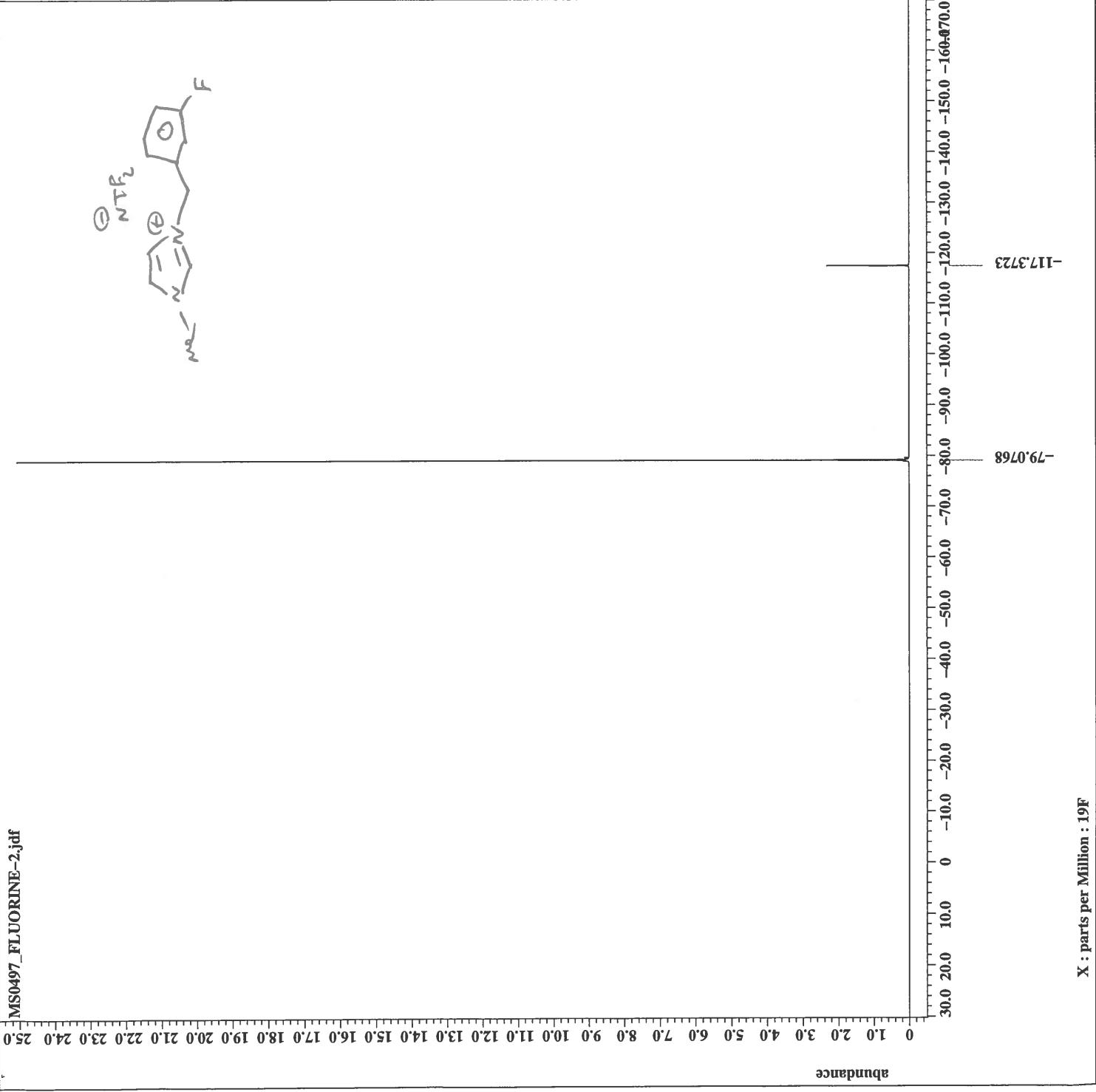
Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acc_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536

X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.924523 [kHz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acc_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Recvrv_gain = 30
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.4 [DC]

```





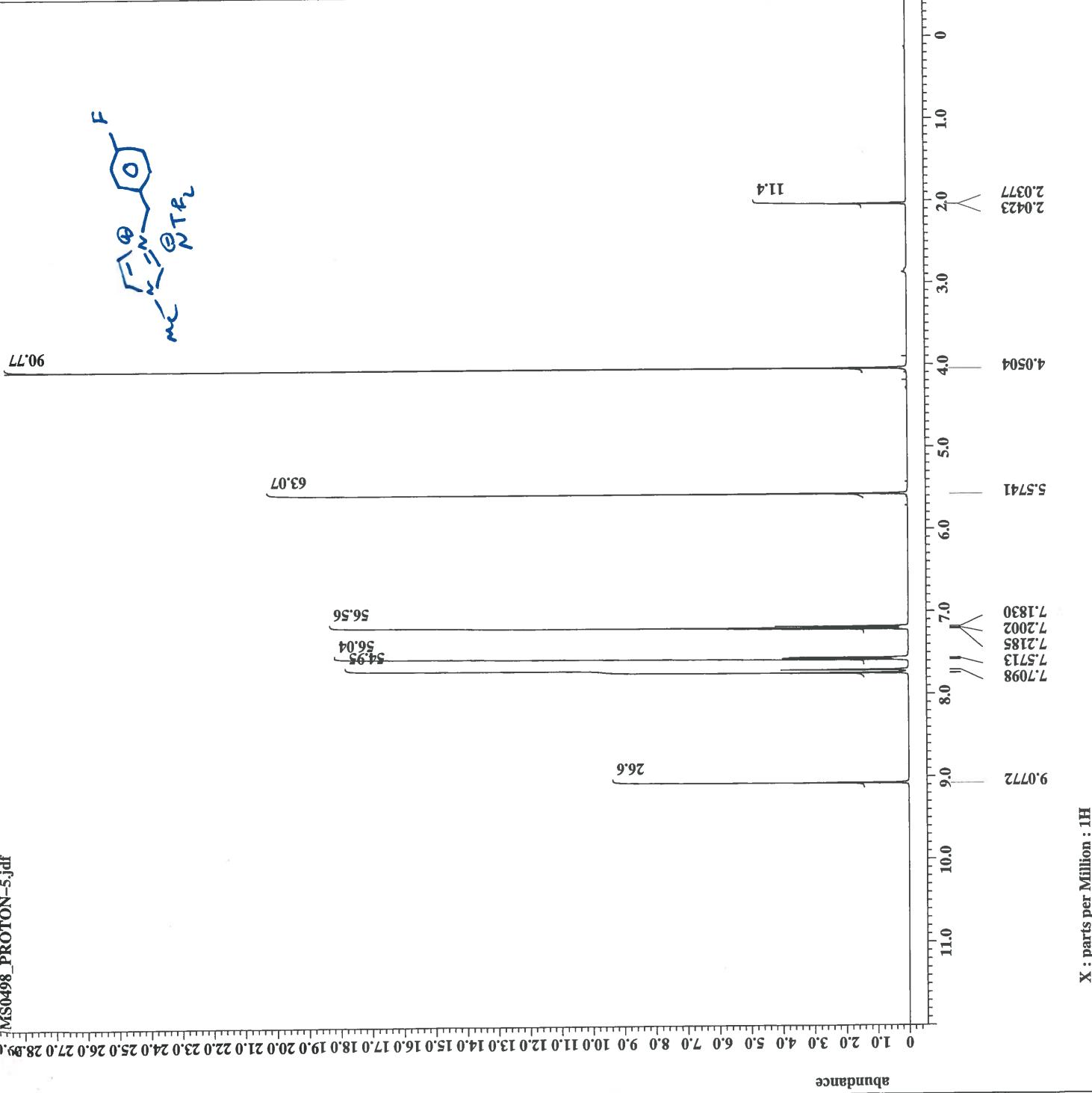
```

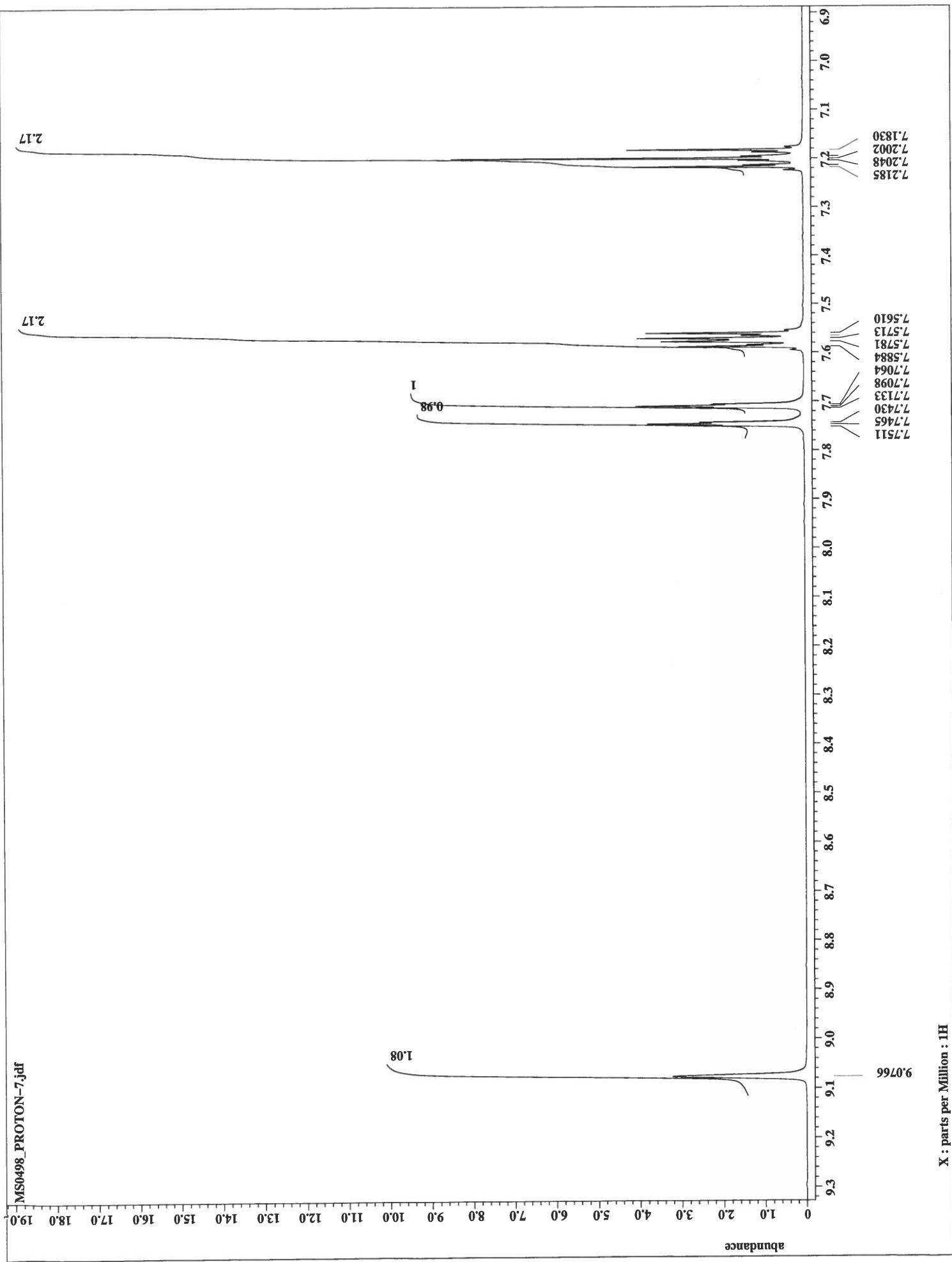
Filename = MS0498_PROTON-5.jdf
Author = Jim Davis
Experiment =
Sample_id =
Solvent =
Changer_sample =
Creation_time = 13-JUL-2018 09:18:13
Revision_time = 13-JUL-2018 08:55:45
Current_time = 13-JUL-2018 08:55:45
Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500.0 [MHz])
X.acq_duration = 1.74587904 [s]
X.domain = 1H
X.freq = 500.15991521 [MHz]
X.offset = 5.0 [ppm]
X.points = 16384
X.prsccans = 1
X.resolution =
X.sweep =
Irr_domain =
Irr.freq = 500.15991521 [MHz]
Irr.offset = 5.0 [ppm]
Tri_domain =
Tri.freq = 500.15991521 [MHz]
Tri.offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 12.4 [us]
X_acq_time = 1.74587904 [s]
X_angle =
X_atn =
X_pulse =
Irr_mode = 4 [deg]
4 [dB] =
6.2 [us]
Off =
Off =
FALSE =
1 [s]
34 =
4 [s]
5.74587904 [s]
22.5 [dB]
Temp_get =

```







```

filename = MS0498_CARBON-2.rdf
author = Jim Davis
experiment = single_pulse_dec
sample_id = MS0498
solvent = CHLOROFORM-D
changer_sample = 8
creation_time = 12-JUL-2018 17:31:59
revision_time = 12-JUL-2018 17:09:35
current_time = 12-JUL-2018 17:09:35

data_format = 1D COMPLEX
dim_size = 2614
dim_title = 13C
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = JNM-ECA500

field_strength = 11.7473579 [T] (500 [MHz])
x_acq_duration = 0.83361792 [s]
x_domain = 13C
x_freq = 125.76529768 [MHz]
x_offset = 100.0 [ppm]
x_points = 32768
x_prescans = 4
x_resolution = 1.19959034 [Hz]
x_sweep = 39.3081761 [PPB]
irr_domain = 1H
irr_freq = 500.15991521 [MHz]
irr_offset = 5.0 [ppm]
clipped = FALSE
mod_return = 1
scans = 128
total_scans = 128

x90_width = 13.2 [us]
x_acq_time = 0.83361792 [s]
x_angle = 30 [deg]
x_atn = 6 [dB]
x_pulse = 4.4 [us]
irr_atn_dec = 20.7 [dB]
irr_atn_noe = 60
irr_noise = 100
decoupling = 100
initial_wait = 1 [s]
noe_time = 2 [s]
recvr_gain = 60
relaxation_delay = 2 [s]
repetition_time = 2.83361792 [s]
temp_get = 23 [IC]

```

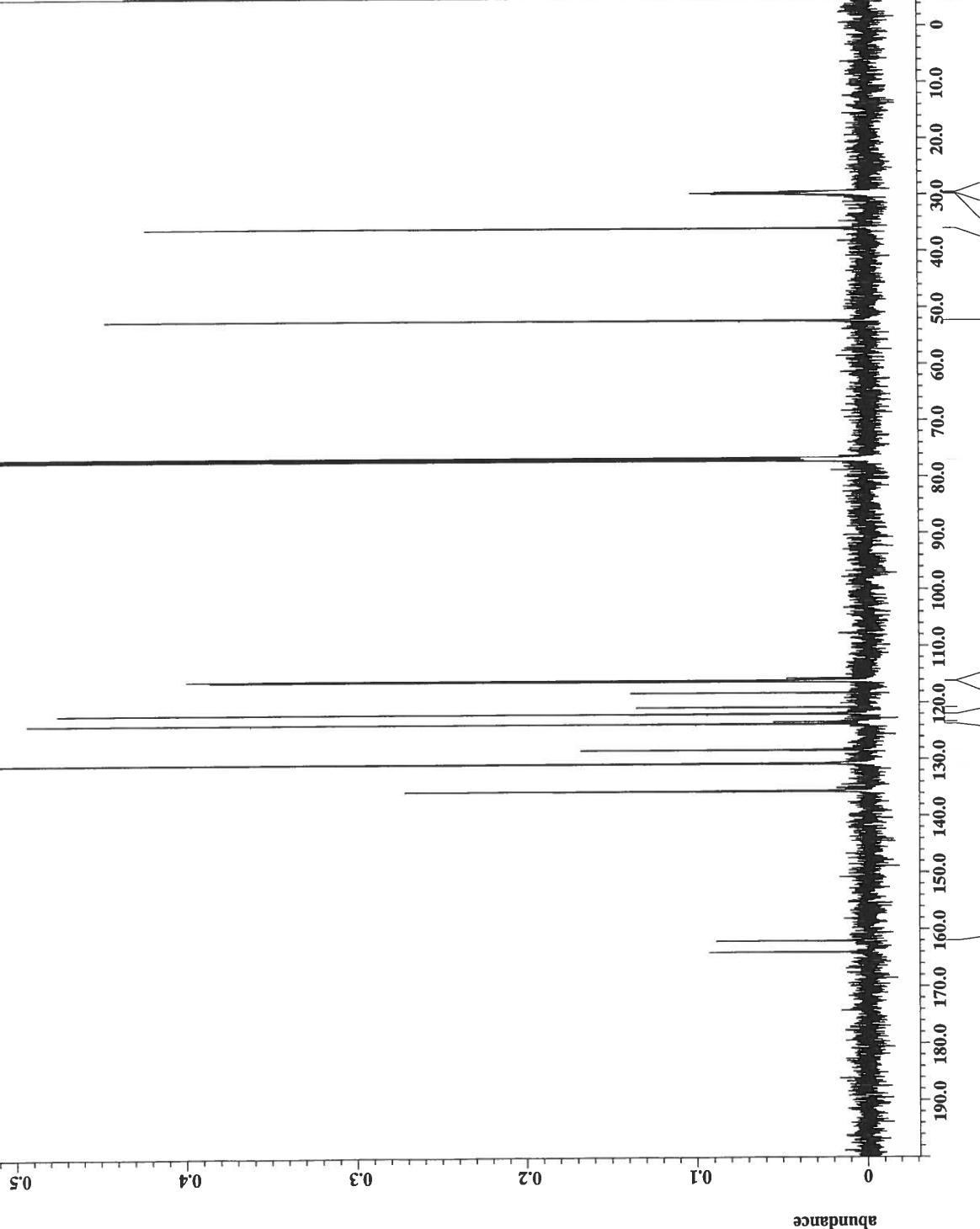
36.0520
29.9665
29.8139
29.6613

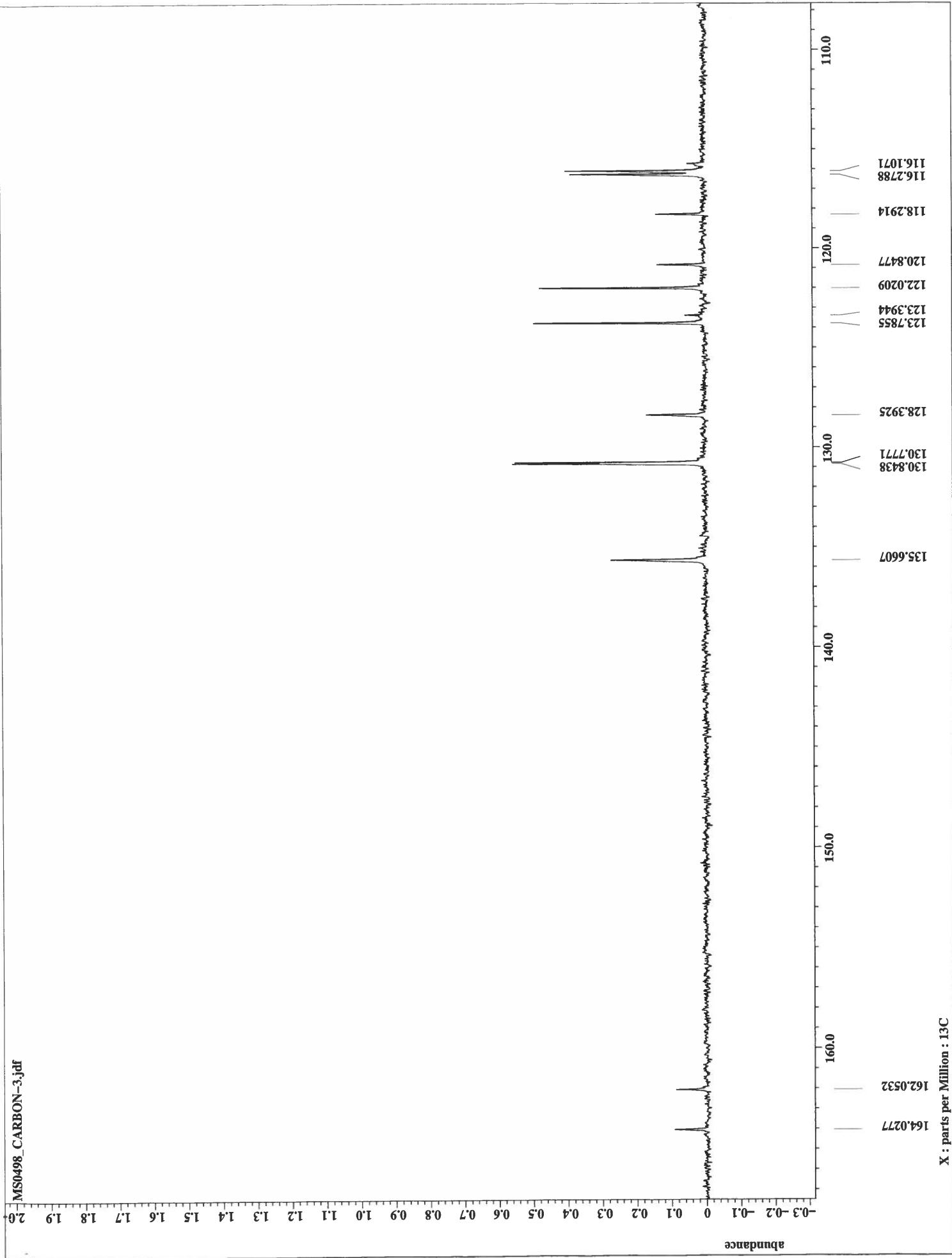
52.3816

77.2575
77.0000

135.6607
130.8438
128.3777
123.7855
122.0209
116.1071

164.0277
162.0532
X : parts per Million : 13C







```

Filename = MS0498_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse_ex2
Sample_id = MS0498
Solvent = CHLOROFORM-D
Changer_sample = 8
Creation_time = 12-JUL-2018 16:31:45
Revision_time = 12-JUL-2018 16:09:22
Current_time = 12-JUL-2018 16:09:23

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [MHz] (500 [MHz]
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [Hz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.05 [us]
Irr_mode = Off
Tri_mode = Off
Dante_presat = FALSE
Initial_wait = 1 [s]
Revr_gain = 36
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.4 [dC]

```

-111.5760

-79.1265

X : parts per Million : 19F

Abundance





```

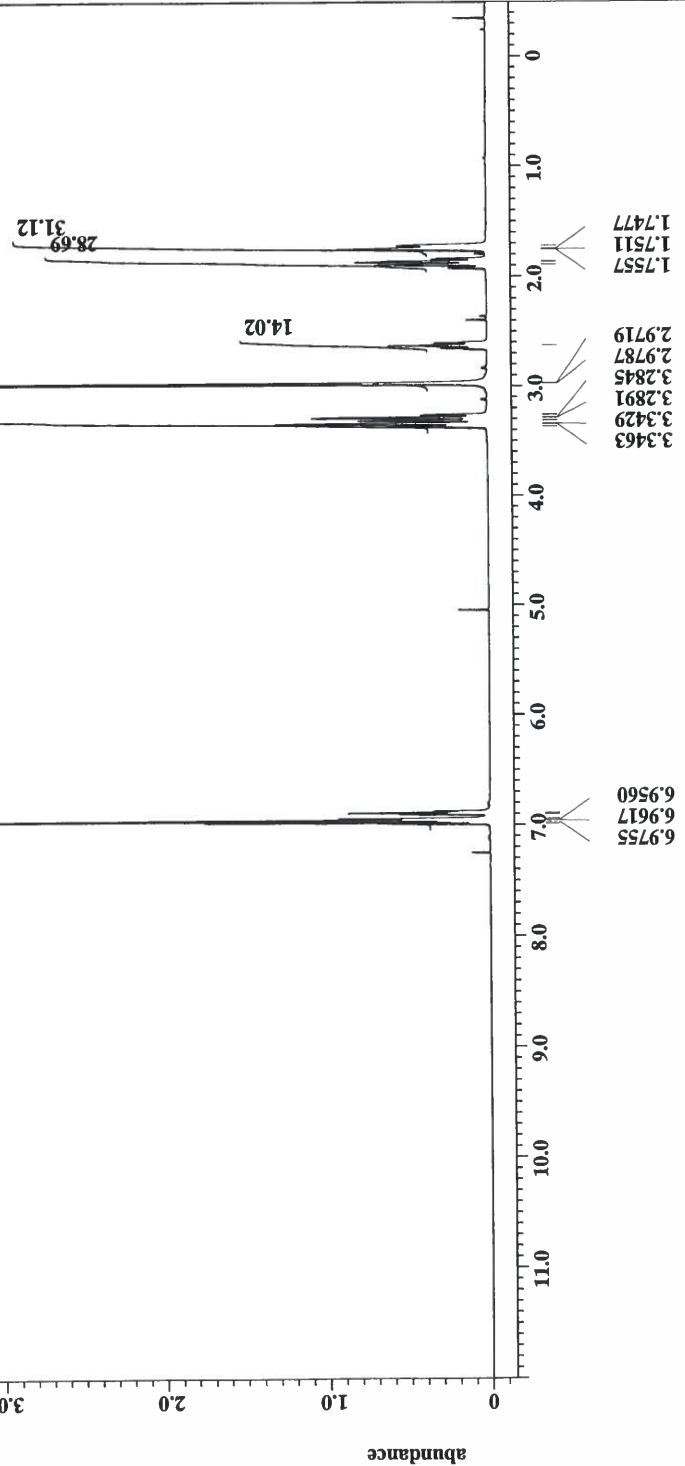
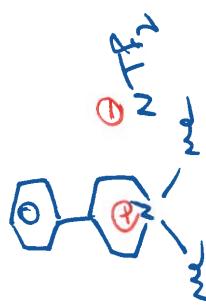
= MS0501.PROTON-2.jdf
= Vilm Davis
= single_pulse.ex2
Sample_id = MS0501
Solvent = CHLOROFORM-D
Changer_sample = 1.1
Creation_time = 5-JUL-2018 16:43:37
Revision_time = 5-JUL-2018 16:20:08
Current_time = 5-JUL-2018 16:20:08

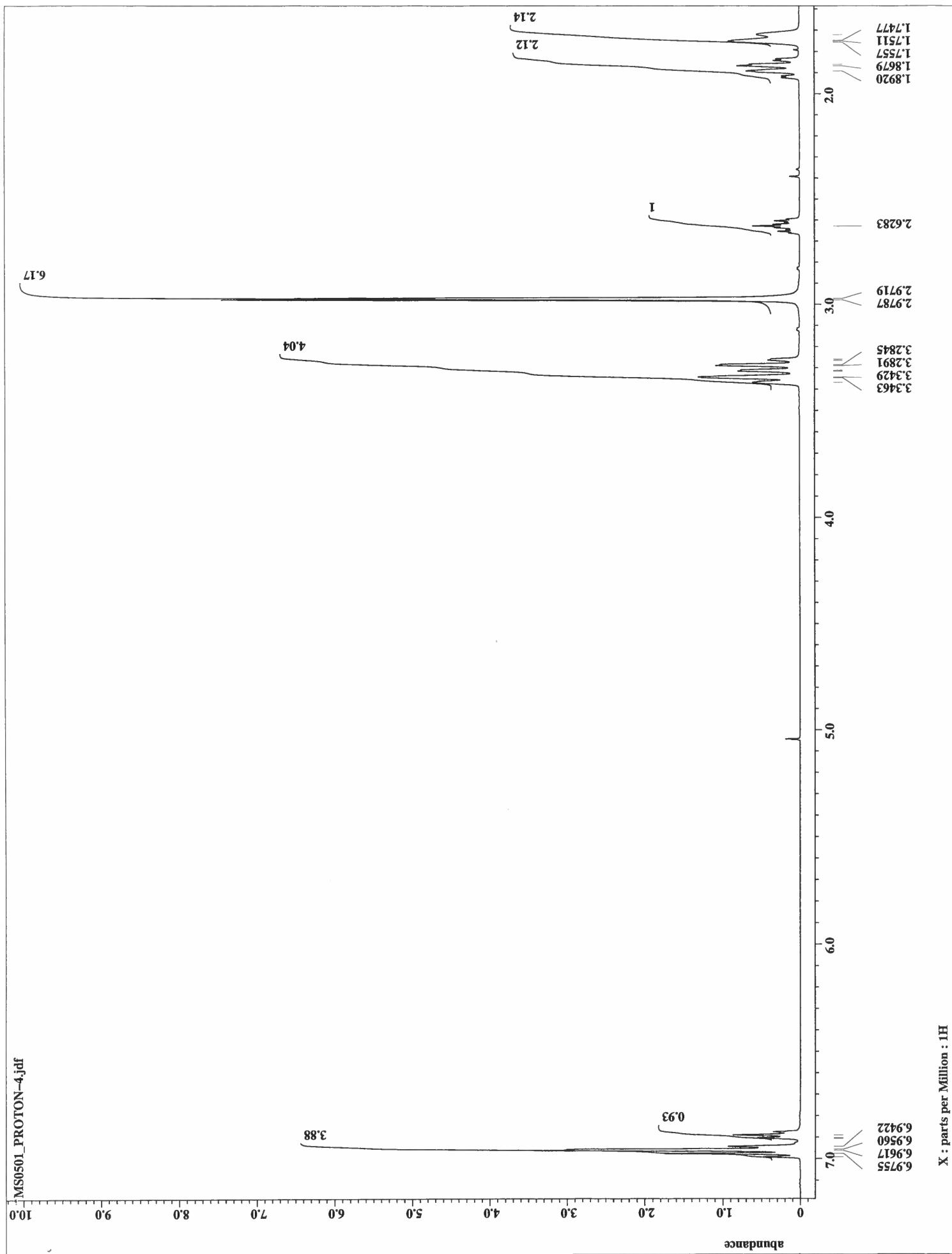
Data_format = 1D COMPLEX
Dim_size = 13107
Dim_title = 1H
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JEOL-ECA500

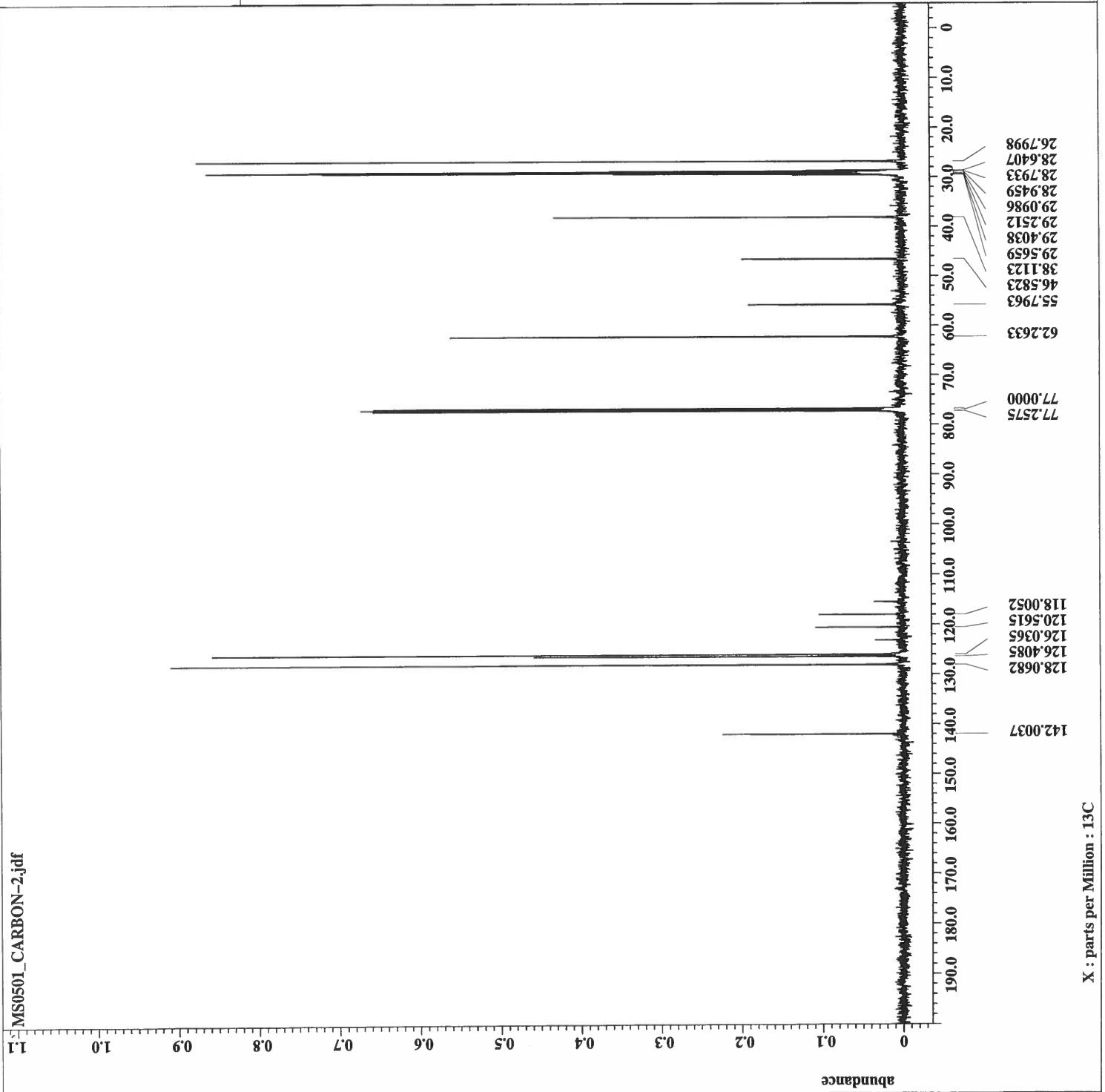
Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 1.74587904 [s]
X_domain = 1H
X_freq = 500.15991521 [MHz]
X_offset = 5.0 [ppm]
X_points = 16384
X_prescans = 1
X_resolution = 0.5727737 [Hz]
X_sweep = 9.38439438 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Tri_domain = 1H
Tri_freq = 500.15991521 [MHz]
Tri_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 12.4 [us]
X_acq_time = 1.74587904 [s]
X_angle = 45 [deg]
X_stn = 4 [dB]
X_pulse = 6.2 [us]
Irr_mode = Off
Tri_mode = Off
Dante_pressat = FALSE
Initial_wait = 1 [s]
Recvr_gain = 30
Relaxation_delay = 4 [s]
Repetition_time = 5.74587904 [s]
Temp_get = 22.7 [dc]

```







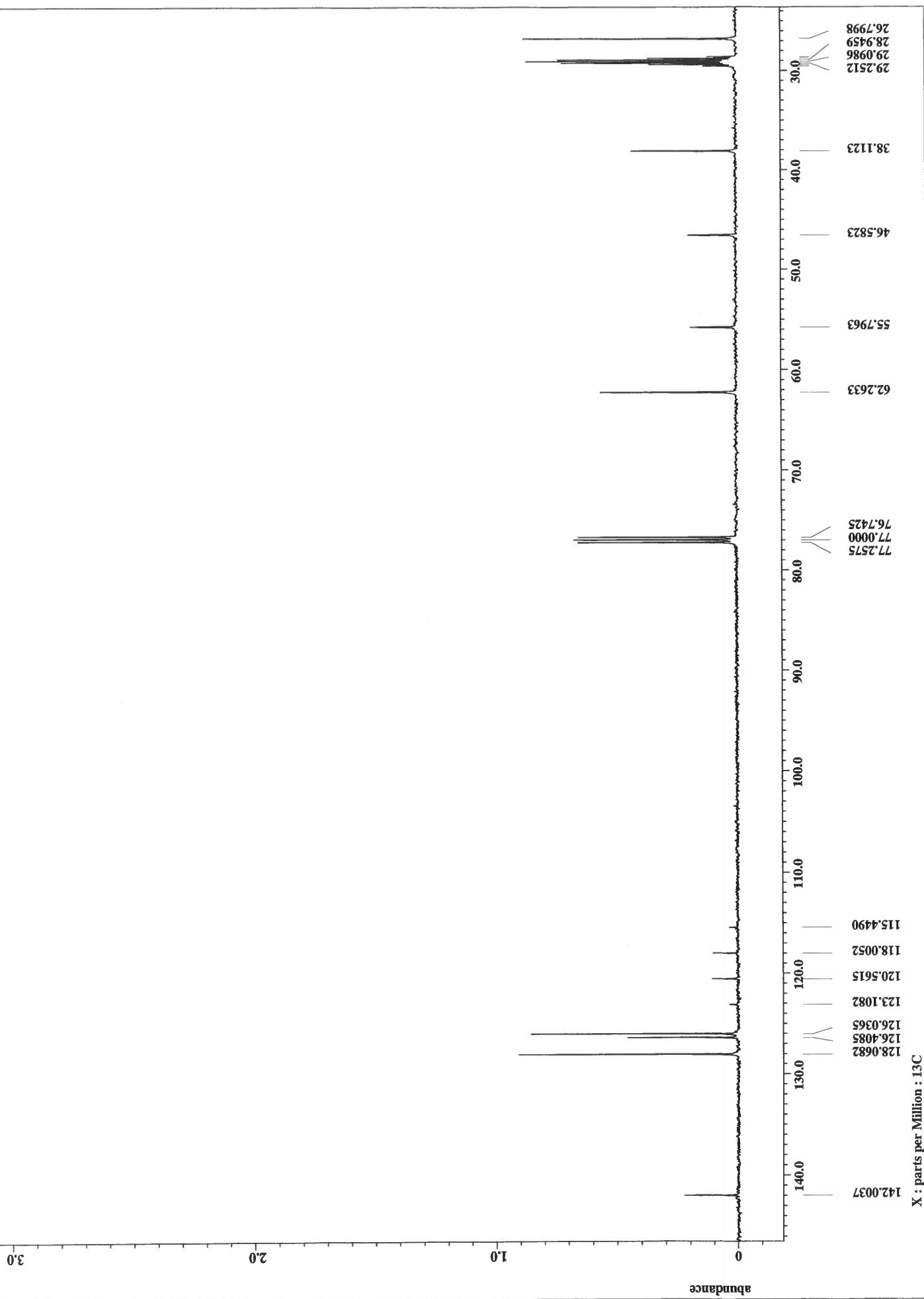
MS0501 CARBON-2.jdf

The logo for the South Alabama Jaguars. It features a stylized jaguar head in profile, facing right, with its mouth open as if roaring. The head is rendered in a dark, textured style. To the right of the head, the word "SOUTH" is stacked above "ALABAMA", and "JAGUARS" is stacked below "ALABAMA". The entire wordmark is enclosed in a thick, dark, horizontal bar. The "S" in "SOUTH" and the "J" in "JAGUARS" are partially cut off by the bar's edge.

```

Filename = MS0501_CARBON-2.jdf
Author = Jim Davis
Experiment =
Sample_id =
Solvent =
Chloroform-D
Changer_sample =
Creation_time = 5-JUL-2018 16:57:55
Revision_time = 5-JUL-2018 16:44:26
Current_time = 5-JUL-2018 16:34:26
Data_format = 1D COMPLEX
Dim_size = 2614
Dim_title =
Dim_units [ppm]
Dimensions =
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength =
X_accg_duration =
X_domain =
X_freq =
X_offset =
X_prestabs =
X_resolution =
X_sweep =
Irr_domain =
Irr_freq =
Irr_offset =
Clipped =
Mod_return =
Scans = 256
Total_scans = 256
X_90_width =
X_acc_time =
X_angle =
X_atn =
X_pulse =
Irr_stn_dec =
Irr_atn_nce =
Irr_noise =
Decoupling =
Initial_wait =
Noe =
Noe_time =
Recvr_gain =
Relaxation_delay =
Repetition_time =
Temp_get =
WALTZ =
TRUE =
1[s] =
TRUE =
2[s] =
23.2[dc]

```





```

Filename = MS0501_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0501
Solvent = CHLOROFORM-D
Changer_sample = 11
Creation_time = 5-JUL-2018 17:00:44
Revision_time = 5-JUL-2018 16:37:16
Current_time = 5-JUL-2018 16:37:16

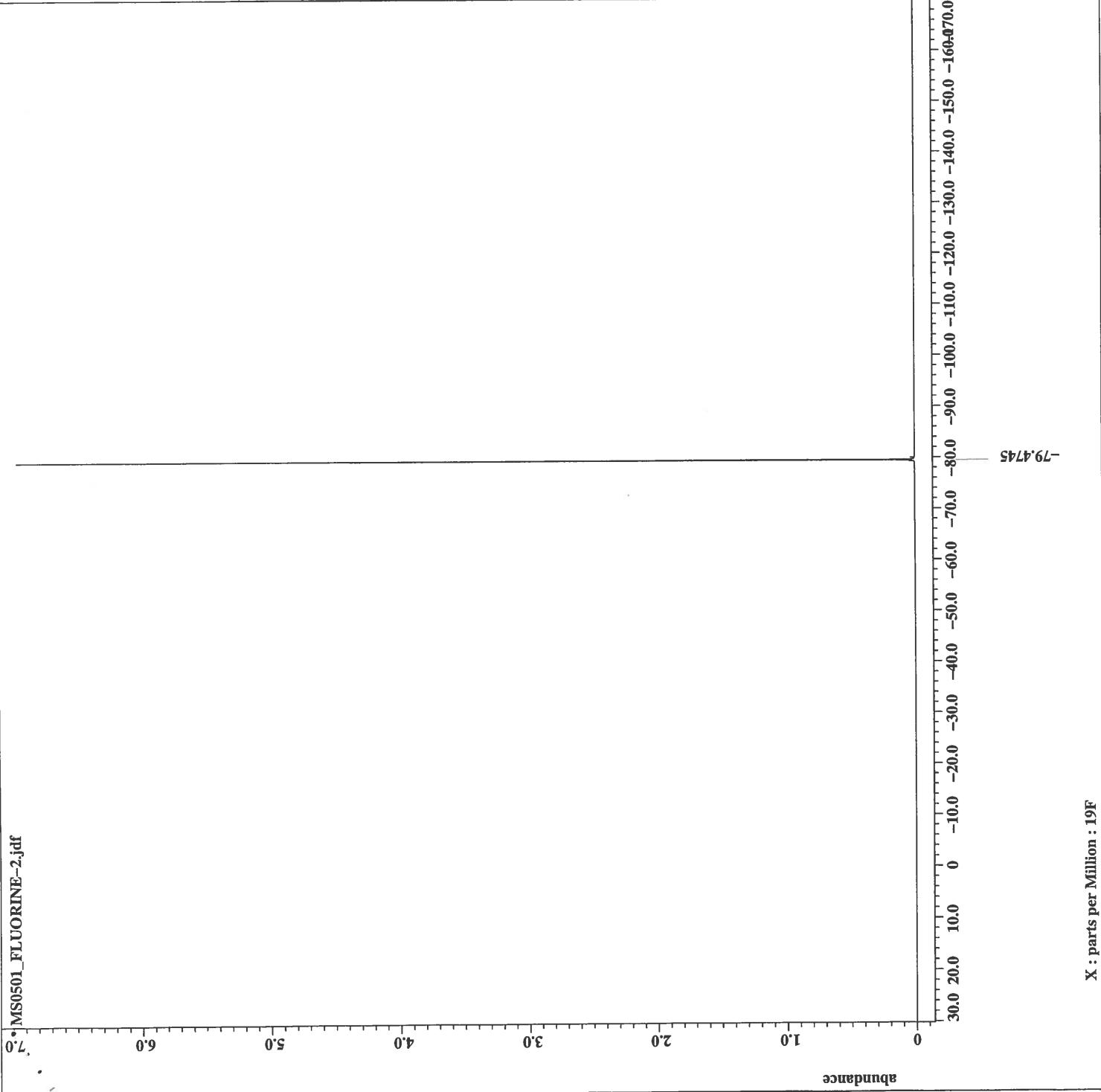
Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [Hz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_pressat = FALSE
Initial_wait = 1 [s]
Recv_rgrain = 38
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.0 [idC]

```

-79.4745





```

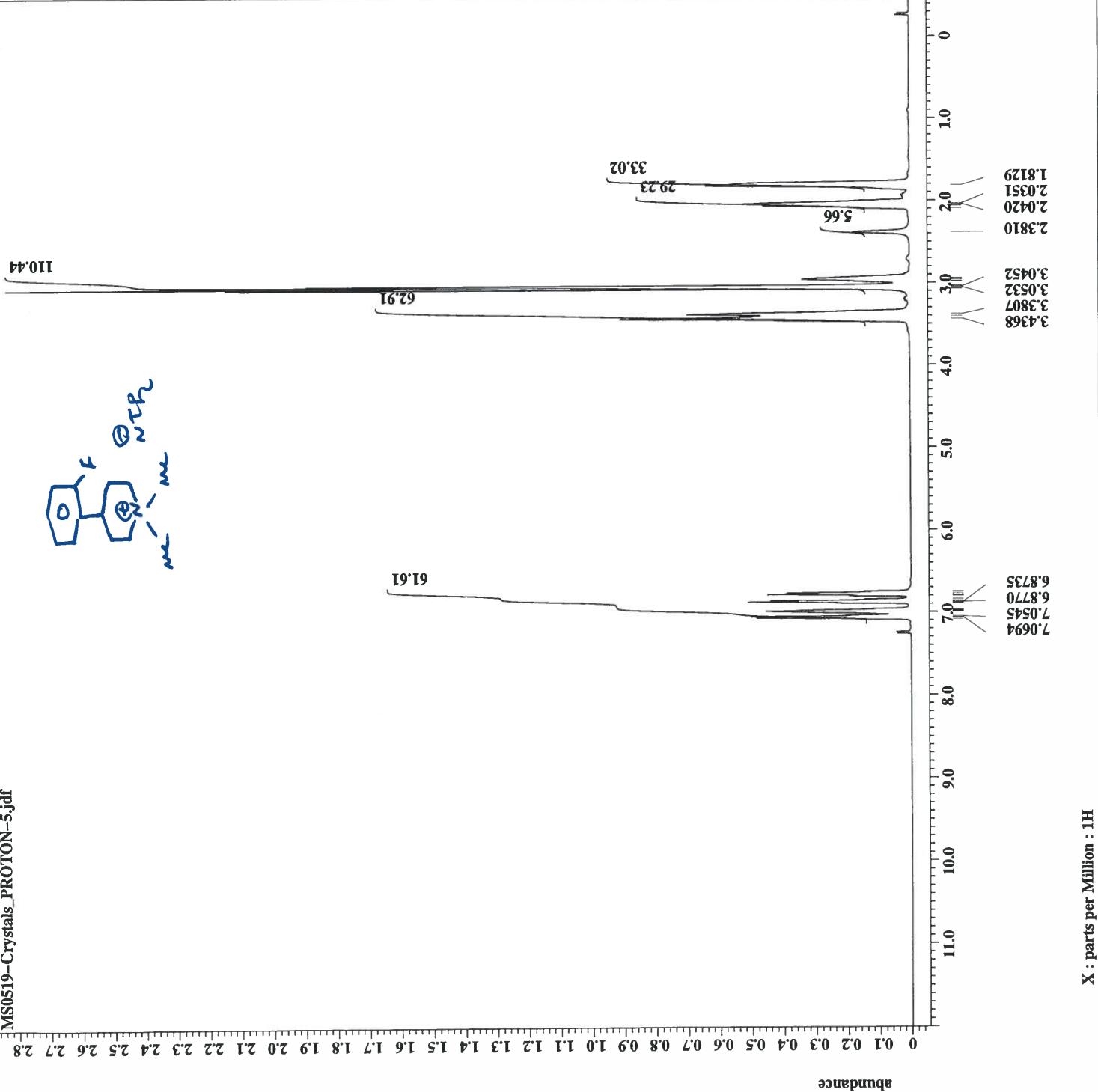
filename      = MS0519-Crystals.PROTON
author        = Jim Davis
experiment   = single_pulse-ex2
sample_id    = MS0519-Crystals
solvent       = CHLOROFORM-D
changer_sample = 6
creation_time = 25-JUL-2018 09:12:03
revision_time = 25-JUL-2018 08:48:40
current_time  = 25-JUL-2018 08:48:40

data_format   = 1D COMPLEX
dim_size      = 13107
dim_title     = 1H
dim_units     = [ppm]
dimensions   = X
site          = ECA 500
spectrometer  = JNM-ECA500

field_strength = 11.7473579 [T] (500 [MHz])
x_acq_duration = 1.74587904 [s]
x_domain      = 1H
x_freq         = 500.15991521 [MHz]
x_offset       = 5.0 [ppm]
x_points       = 16384
x_prescans    = 1
x_resolution  = 0.57277737 [Hz]
x_sweep        = 9.38458438 [kHz]
irr_domain    = 1H
irr_freq       = 500.15991521 [MHz]
irr_offset     = 5.0 [ppm]
tri_domain    = 1H
tri_freq       = 500.15991521 [MHz]
tri_offset    = 5.0 [ppm]
clipped       = FALSE
mod_return    = 1
scans          = 50
total_scans   = 50

x_90_width    = 12.4 [us]
x_acq_time    = 1.74587904 [s]
x_angle        = 45 [deg]
x_atn          = 4 [dB]
x_pulse        = 6.2 [us]
irr_mode       = Off
tri_mode       = Off
dante_preset  = FALSE
initial_wait   = 1 [s]
recv_gain      = 30
relaxation_delay = 4 [s]
repetition_time = 5.74587904 [s]
temp_get       = 22.4 [dc]

```





```

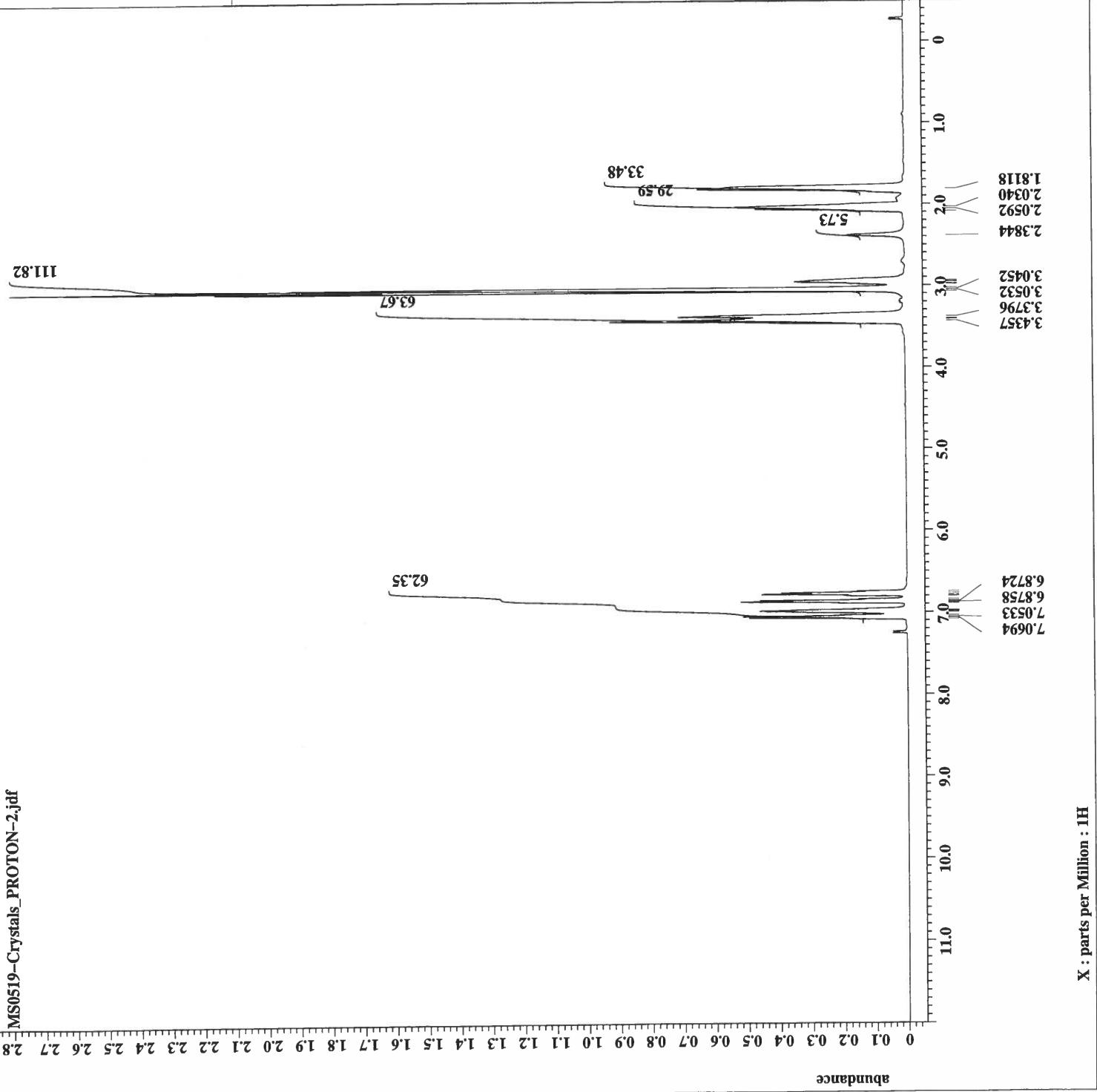
filename = MS0519-Crystals_PROTO
author = Jim Davis
Experiment = Single_pulse-ex2
sample_id = MS0519-Crystals
solvent = CHLOROFORM-D
Changer sample = 6
Creation_time = 25-JUL-2018 08:46:11
Revision_time = 25-JUL-2018 08:22:50
Current_time = 25-JUL-2018 08:22:50

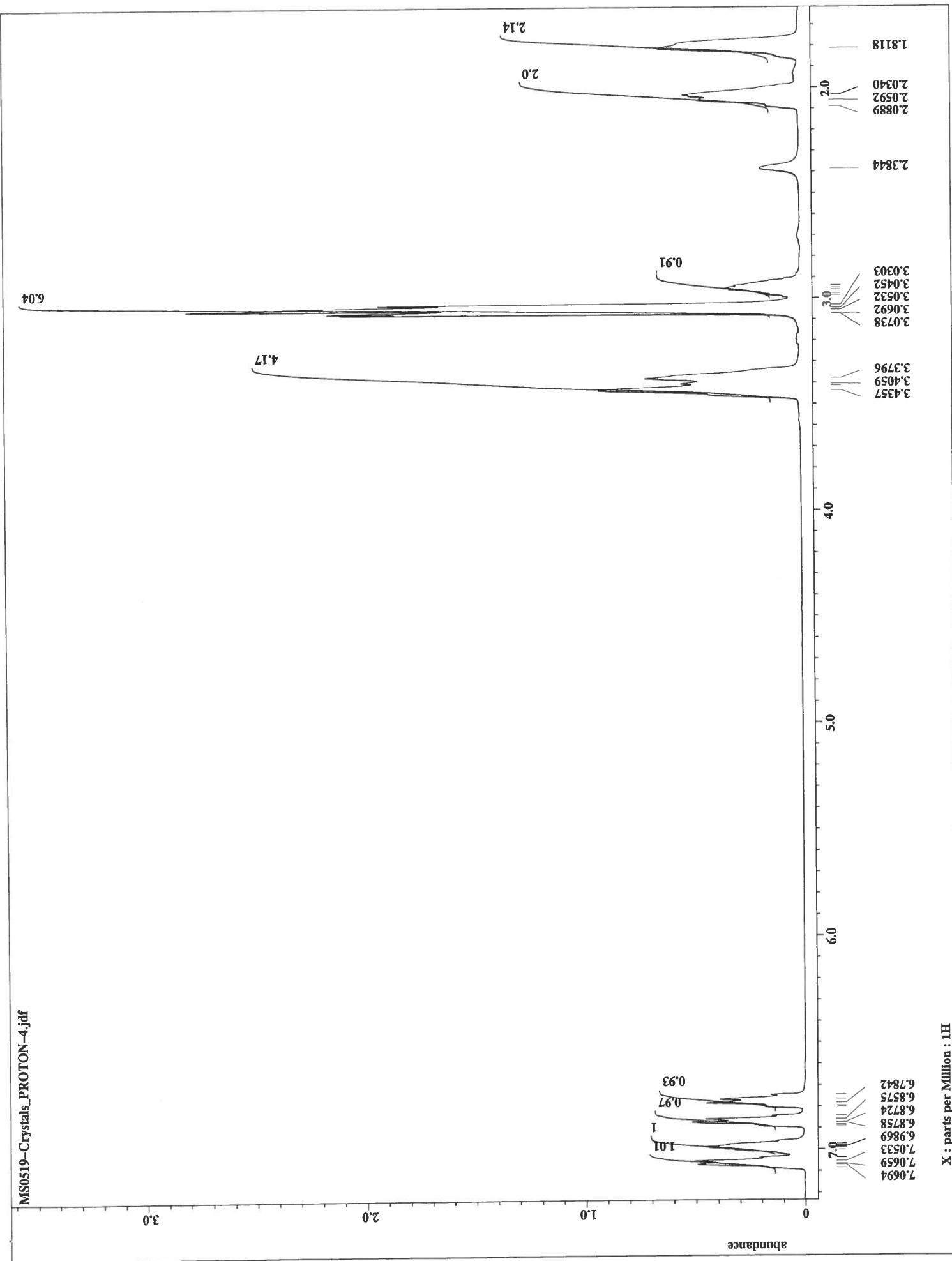
data_format = 1D COMPLEX
dim_size = 13107
dim_title = 1H
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = QNM-ECA500

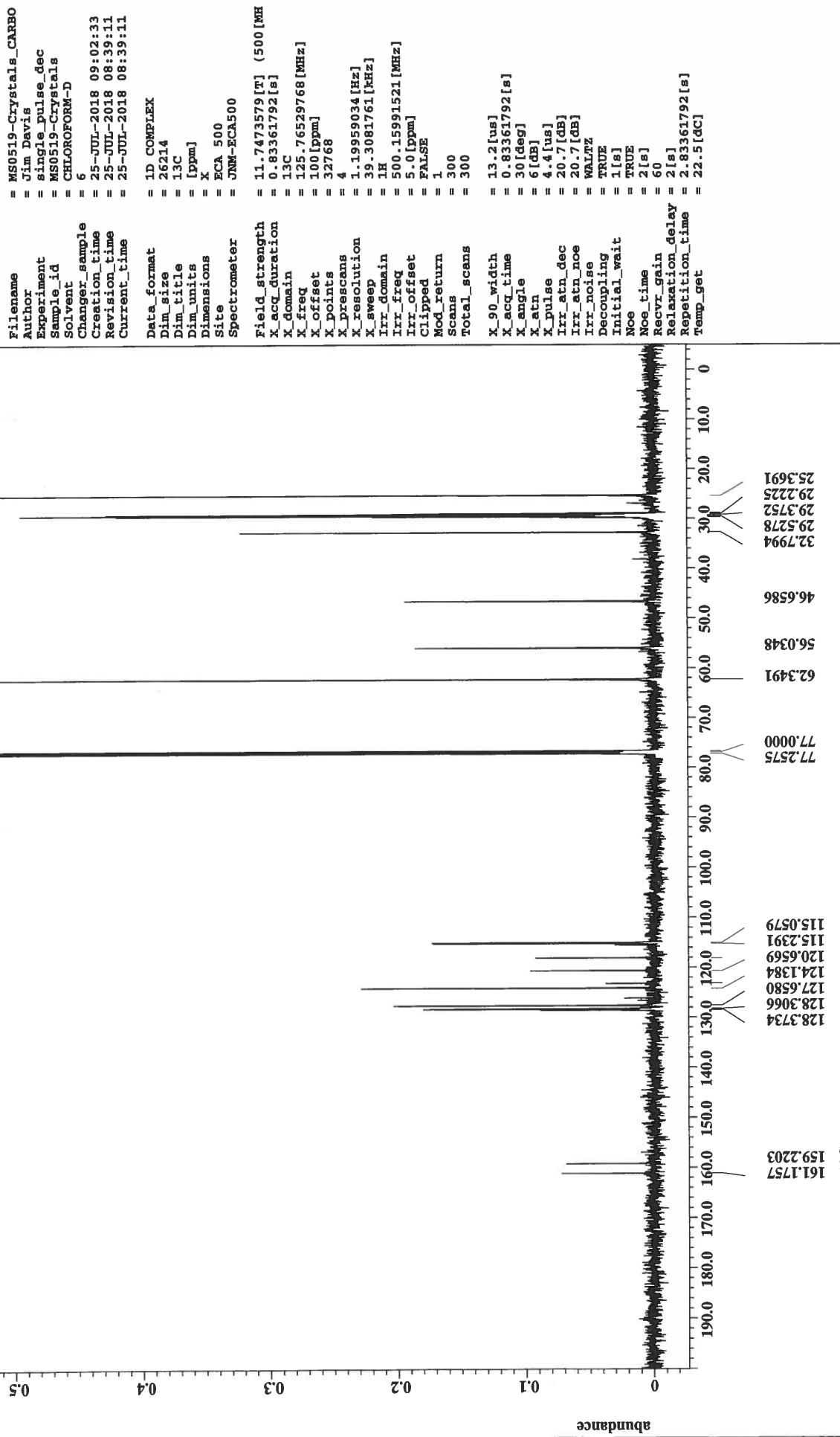
field_strength = 11.7473579 [T] (500 [MHz])
x_acq_duration = 1.74587904 [s]
x_domain = 1H
x_freq = 500.15991521 [MHz]
x_offset = 5.0 [ppm]
x_points = 16384
x_prescans = 1
x_resolution = 0.57277737 [Hz]
x_sweep = 9.38458438 [kHz]
irr_domain = 1H
irr_freq = 500.15991521 [MHz]
irr_offset = 5.0 [ppm]
tri_domain = 1H
tri_freq = 500.15991521 [MHz]
tri_offset = 5.0 [ppm]
clipped = FALSE
mod_return = 1
scans = 16
total_scans = 16

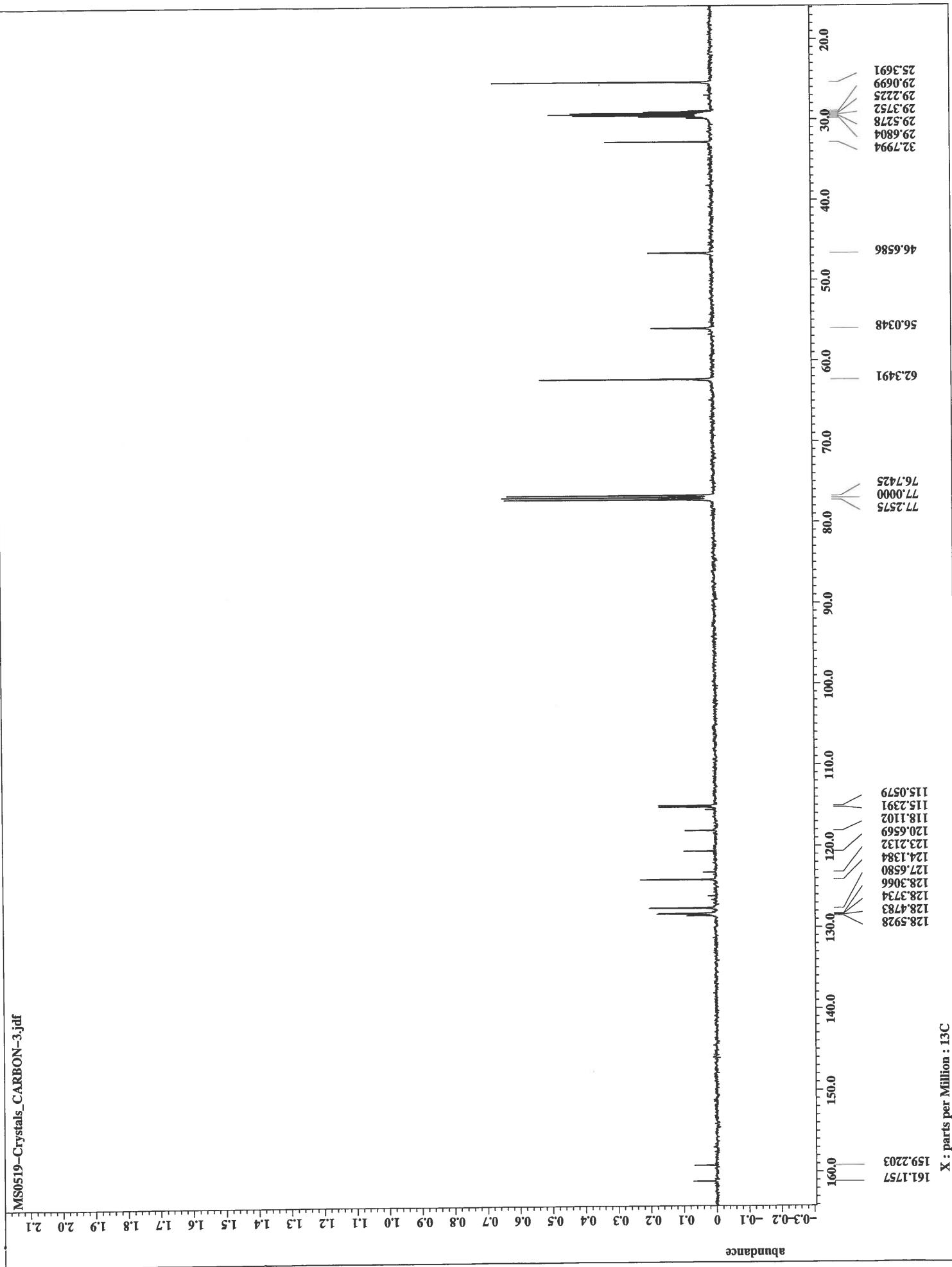
x90_width = 12.4 [us]
x_acq_time = 1.74587904 [s]
x_angle = 45 [deg]
x_atn = 4 [dB]
x_pulse = 6.2 [us]
irr_mode = OFF
tr1_mode = OFF
dante_preset = FALSE
initial_wait = 1 [s]
recvr_gain = 30
relaxation_delay = 4 [s]
relaxation_time = 5.74587904 [s]
temp_get = 22 [dc]


```











```

Filename = MS0519-Crystals_FLUOR
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0519-Crystals
Solvent = CHLOROFORM-D
Changer_sample = 6
Creation_time = 25-JUL-2018 09:05:38
Revision_time = 25-JUL-2018 08:42:15
Current_time = 25-JUL-2018 08:42:15

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 1.9F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.555745281 [s]
X_domain = 1.9F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_precans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [kHz]
Irr_domain = 1.9F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 1.9F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.555745281 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Recvr_gain = 36
Relaxation_delay = 4 [s]
Repetition_time = 4.555745281 [s]
Temp_get = 22.4 [dc]

```

-118.8367

-79.3292





```

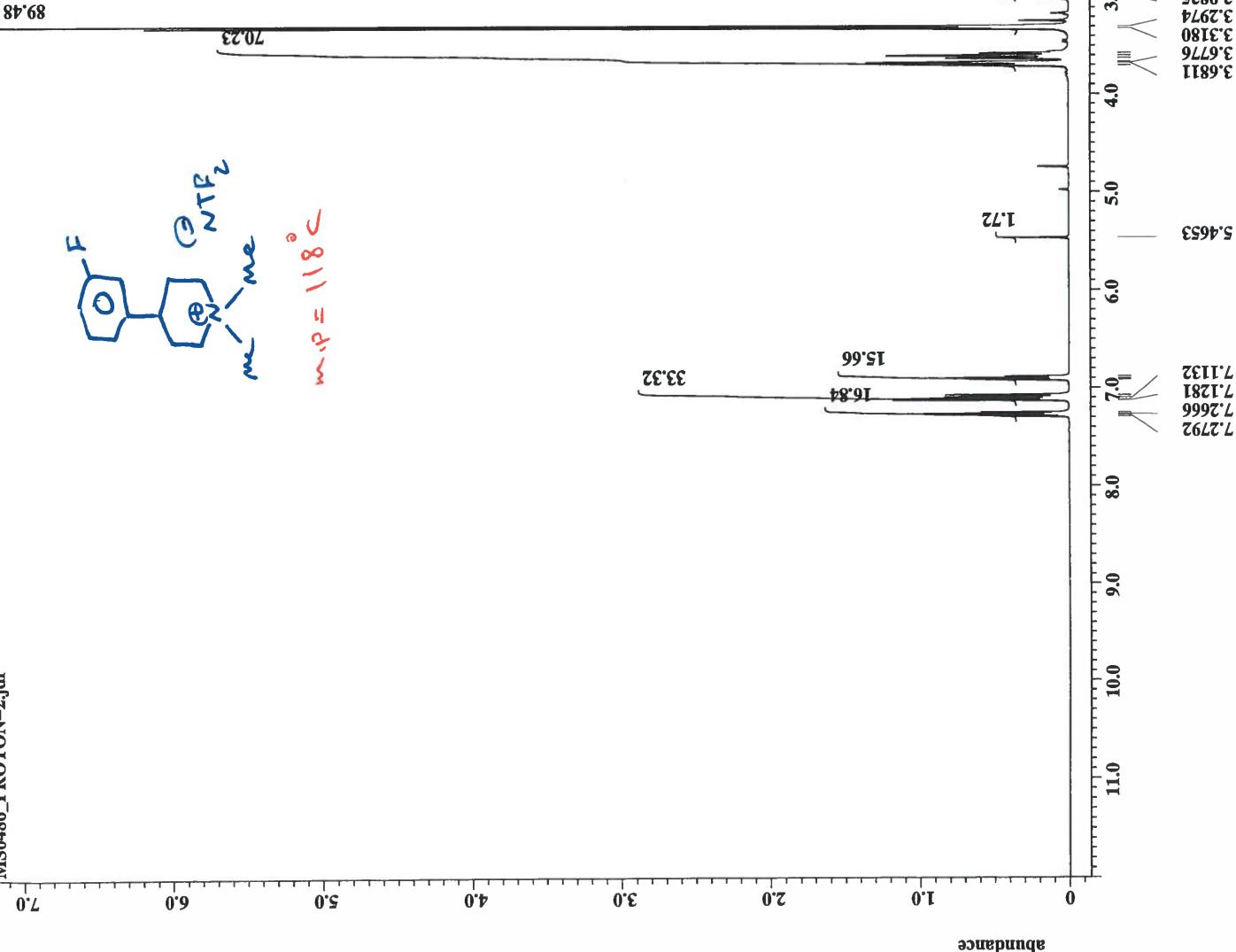
filename = MS0486_PROTON-2.jdf
author = Jim Davis
experiment = single_pulse.ex2
sample_id = MS0486
solvent = ACETONE-D6
changer_sample = 11
creation_time = 27-JUN-2018 10:42:54
revision_time = 27-JUN-2018 10:20:02
current_time = 27-JUN-2018 10:20:02

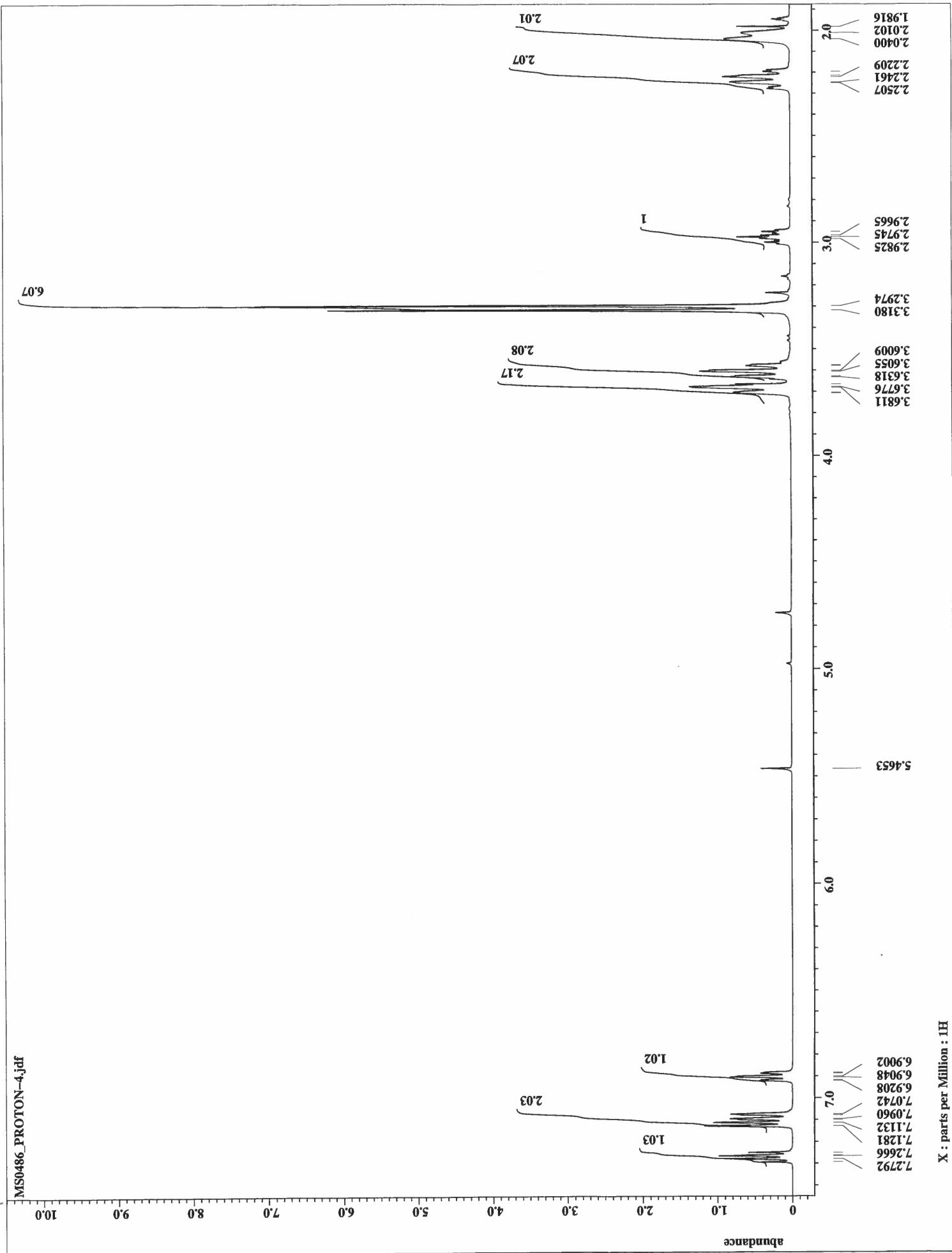
data_format = 1D COMPLEX
dim_size = 13107
dim_title = 1H
dim_units = [ppm]
dimensions = X
site = ECA 500
spectrometer = JEOL-ECA500

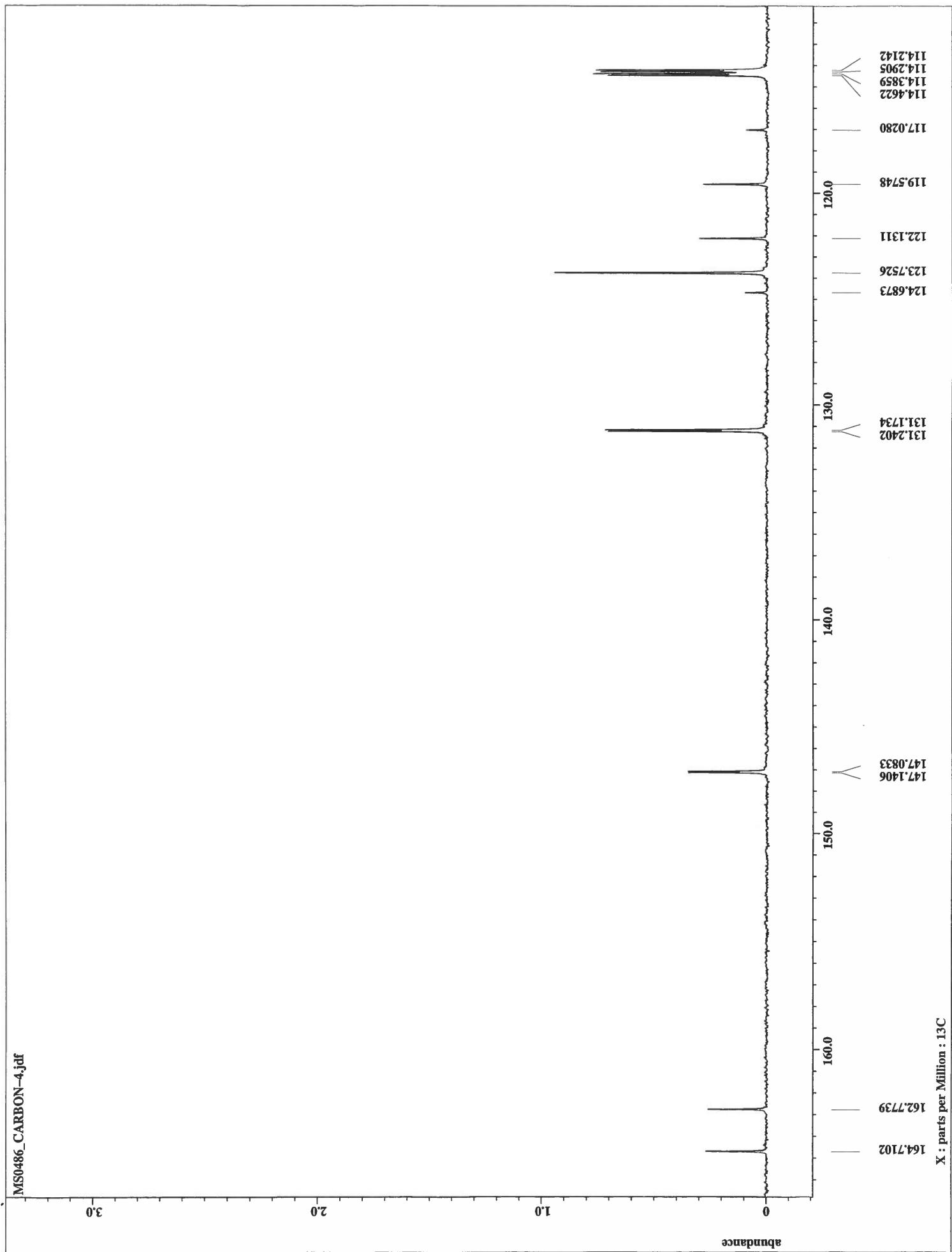
field_strength = 11.7473579 [T] (500 [MHz])
x_acq_duration = 1.74587904 [s]
x_domain = 1H
x_freq = 500.15991521 [MHz]
x_offset = 5.0 [ppm]
x_points = 16384
x_prescans = 1
x_resolution = 0.57277737 [Hz]
x_sweep = 9.38438438 [kHz]
irr_domain = 1H
irr_freq = 500.15991521 [MHz]
irr_offset = 5.0 [ppm]
tri_domain = 1H
tri_freq = 500.15991521 [MHz]
tri_offset = 5.0 [ppm]
clipped = FALSE
mod_return = 1
scans = 16
total_scans = 16

x90_width = 12.4 [us]
x_acq_time = 1.74587904 [s]
x_angle = 45 [deg]
x_atn = 4 [dB]
x_pulse = 6.2 [us]
irr_mode = Off
off_time = 0
dante_preset = FALSE
initial_wait = 1 [s]
recvr_gain = 22
relaxation_delay = 4 [s]
relaxation_time = 5.74587904 [s]
temp_get = 22.3 [dc]

```









```

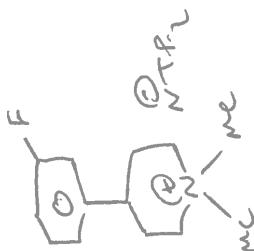
Filename = MS0486_CARBON-2.jdf
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0486
Solvent = ACETONE-D6
Changer_sample = 11
Creation_time = 27-JUN-2018 10:57:27
Revision_time = 27-JUN-2018 10:34:33
Current_time = 27-JUN-2018 10:34:33

Data_format = 1D_COMPLEX
Dim_size = 2614
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECA_500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain = 13C
X_freq = 128.76529768 [MHz]
X_offset = 100.0 [ppm]
X_points = 32768
X_prescans = 4
X_resolution = 1.19959034 [Hz]
X_sweep = 39.3082761 [Hz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 256
Total_scans = 256

X_90_width = 13.2 [us]
X_acq_time = 0.83361792 [s]
X_angle = 30 [deg]
X_atn = 6 [dB]
X_pulse = 4.1 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = 5000
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 2 [s]
Regrv_gain = 60
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get = 22.8 [adc]

```





```

Filename = MS0486_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0486
Solvent = ACETONE-D6
Changer_sample = 11
Creation_time = 27-JUN-2018 10:39:16
Revision_time = 27-JUN-2018 10:16:23
Current_time = 27-JUN-2018 10:16:24

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

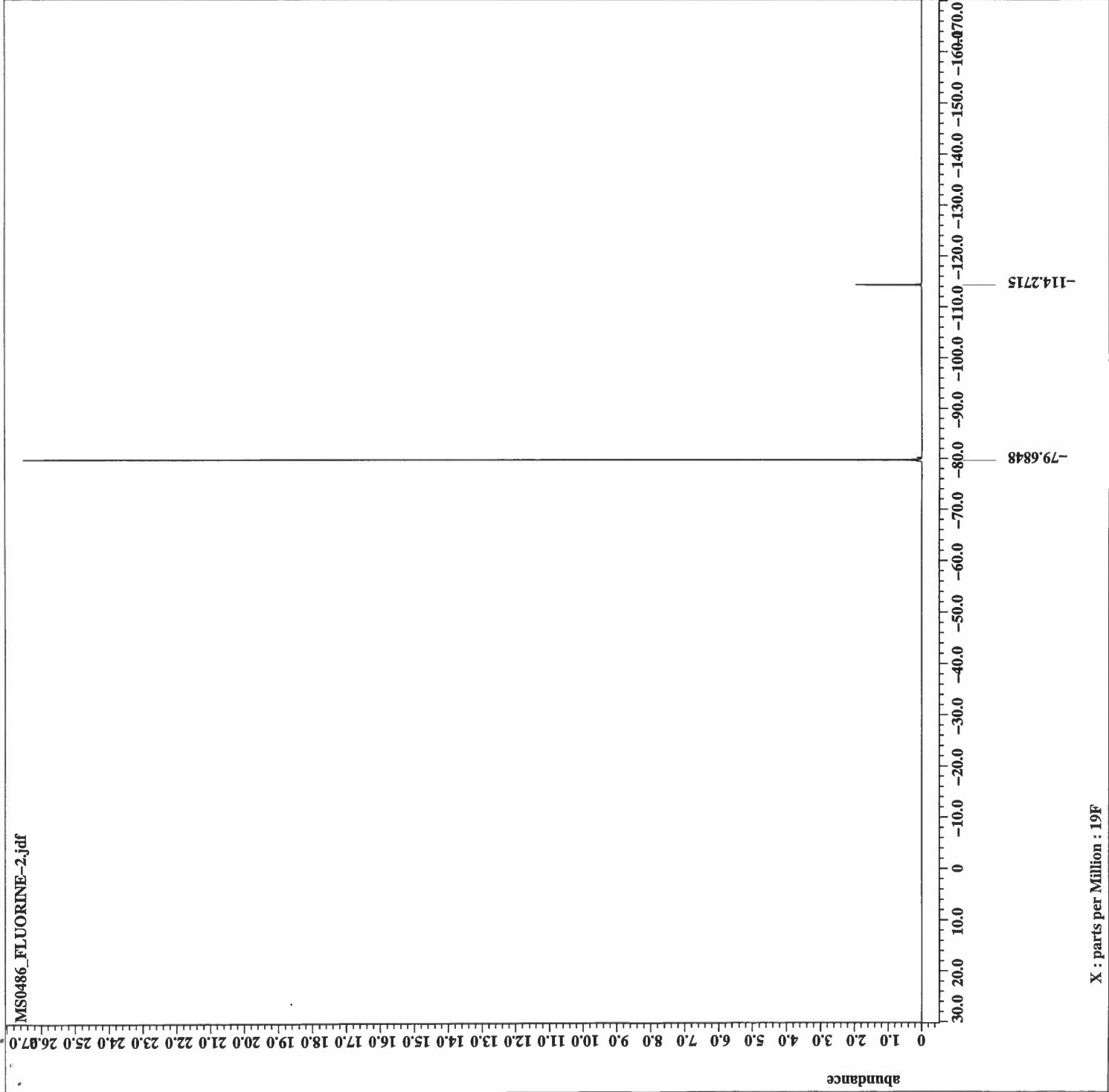
Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_prescans = 1
X_resolution = 1.7993655 [Hz]
X_sweep = 117.9245283 [kHz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1 [s]
Revr_gain = 30
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.3 [dc]

```

-114.2715

-79.6848





```

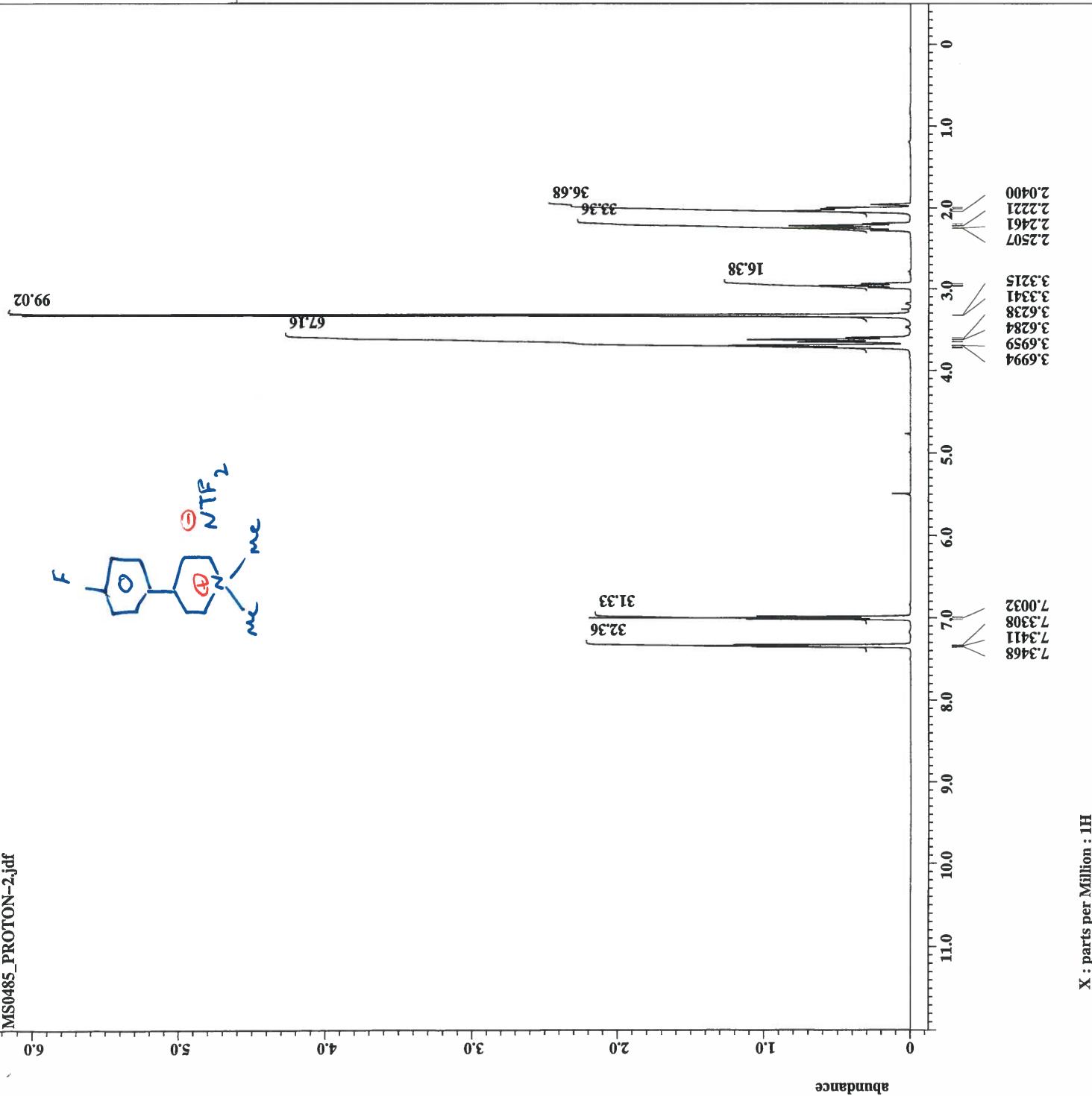
Filename           = MS0485.PROTON-2.jdf
Author            = Jim Davis
Experiment        = single_pulse.ex2
Sample_id         = MS0485
Solvent           = ACETONE-D6
Changer_sample   = 12
Creation_time    = 27-JUN-2018 11:38:17
Revision_time    = 27-JUN-2018 11:15:23
Current_time     = 27-JUN-2018 11:15:23

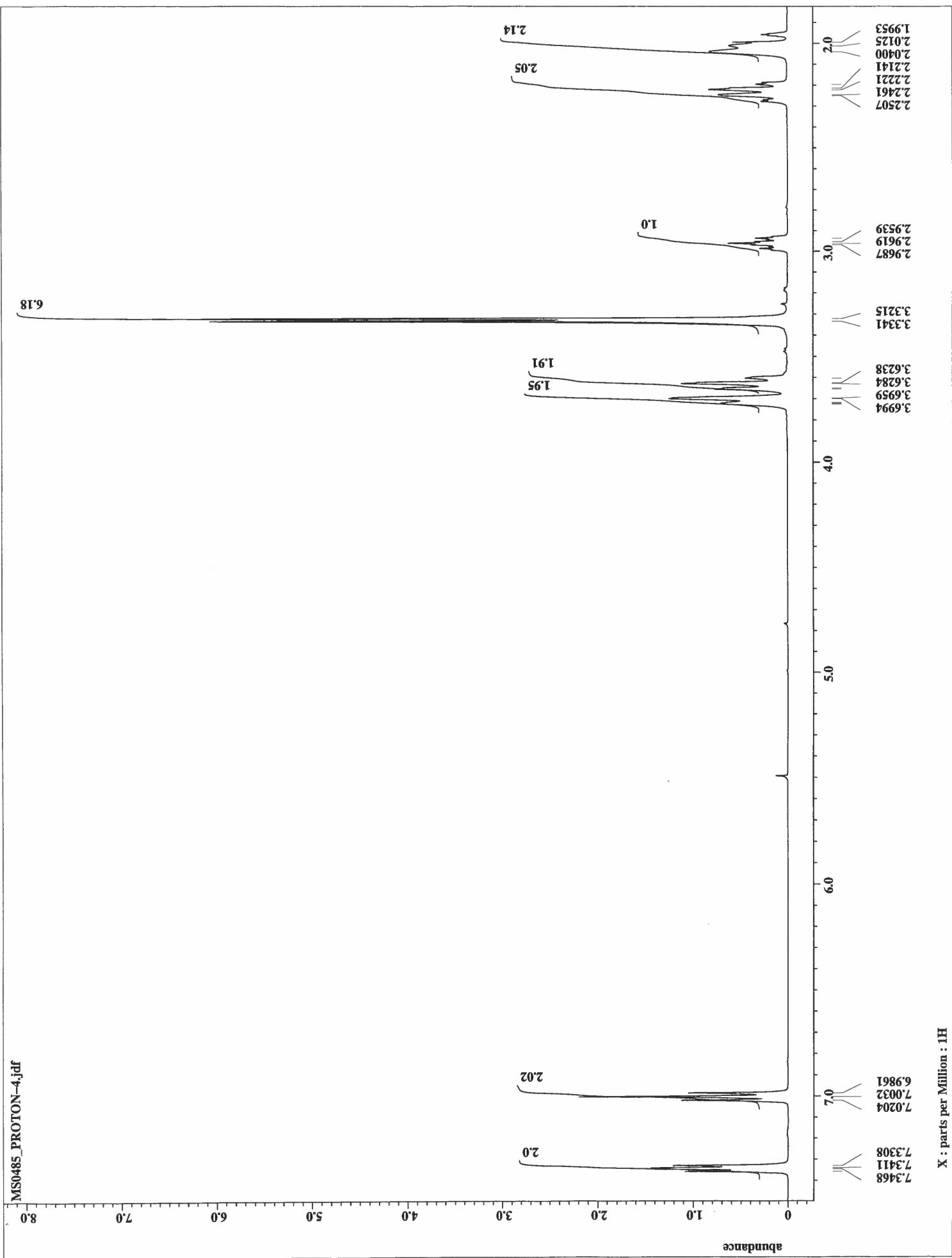
Data_format      = 1D COMPLEX
Dim_size          = 13107
Dim_title         = 1H
Dim_units         = [ppm]
Dimensions        = X
Site              = ECA 500
Spectrometer      = JNM-ECA500

Field_strength   = 11.7475579 [T] (500 [MHz])
X_acc_duration   = 1.74587904 [s]
X_domain          = 1H
X_freq            = 500.15591521 [MHz]
X_offset          = 5.0 [ppm]
X_points          = 16384
X_prescans        = 1
X_resolution      = 0.57277737 [Hz]
X_sweep           = 9.38438438 [kHz]
Irr_domain        = 1H
Irr_freq          = 500.15591521 [MHz]
Irr_offset        = 5.0 [ppm]
Iri_domain        = 1H
Iri_freq          = 500.15591521 [MHz]
Iri_offset        = 5.0 [ppm]
Clipped           = FALSE
Mod_return        = 1
Scans             = 16
Total_scans       = 16

X_90_width        = 12.4 [us]
X_acc_time         = 1.74587904 [s]
X_angle            = 45 [deg]
X_attn             = 4 [dB]
X_pulse            = 6.2 [us]
Irr_mode           = Off
Tri_mode           = Off
Dante_preset       = FALSE
Initial_wait       = 1 [s]
Recv_gain          = 24
Relaxation_delay   = 4 [s]
Repetition_time    = 5.74587904 [s]
Temp_get           = 22.3 [dc]

```







```

Filename = MS0485_CARBON-2.jdf
Author = Jim Davis
Experiment =
Sample_id = MS0485
Solvent =
Changer_sample = 12
Creation_time = 27-JUN-2018 11:59:25
Revision_time = 27-JUN-2018 11:36:31
Current_time = 27-JUN-2018 11:36:31

Data_format = 1D COMPLEX
Dim_size = 2614
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.83361792 [s]
X_domain = 13C
X_freq = 123.76529768 [MHz]
X_offset = 100.0 [ppm]
X_points = 32768
X_prescans = 4
X_resolution = 1.19959034 [Hz]
X_sweep = 39.3082761 [kHz]
Irr_domain = 1H
Irr_freq = 500.15991521 [MHz]
Irr_offset = 5.0 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 400
Total_scans = 400

X_90_width = 13.2 [us]
X_acq_time = 0.83361792 [s]
X_angle = 30 [deg]
X_atn = 6 [dB]
X_pulse = 4.4 [us]
Irr_atn_dec = 20.7 [dB]
Irr_atn_noe = 20.7 [dB]
Irr_noise = 0.0172
Decoupling = TRUE
Initial_wait = 1 [s]
Noe = TRUE
Noe_time = 60
Regrv_gain = 2 [s]
Relaxation_delay = 2 [s]
Repetition_time = 2.83361792 [s]
Temp_get = 22.9 [dc]

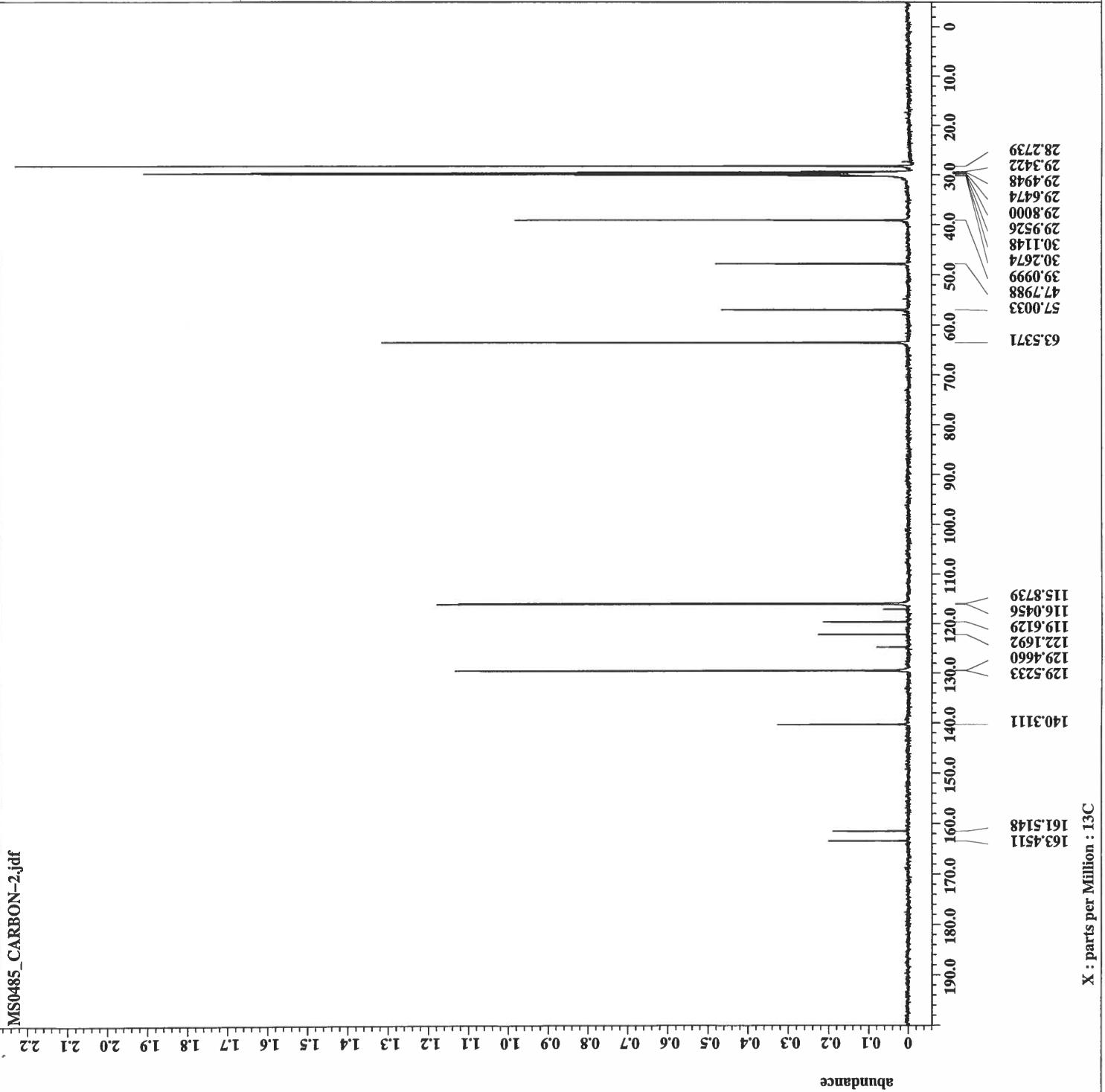
```

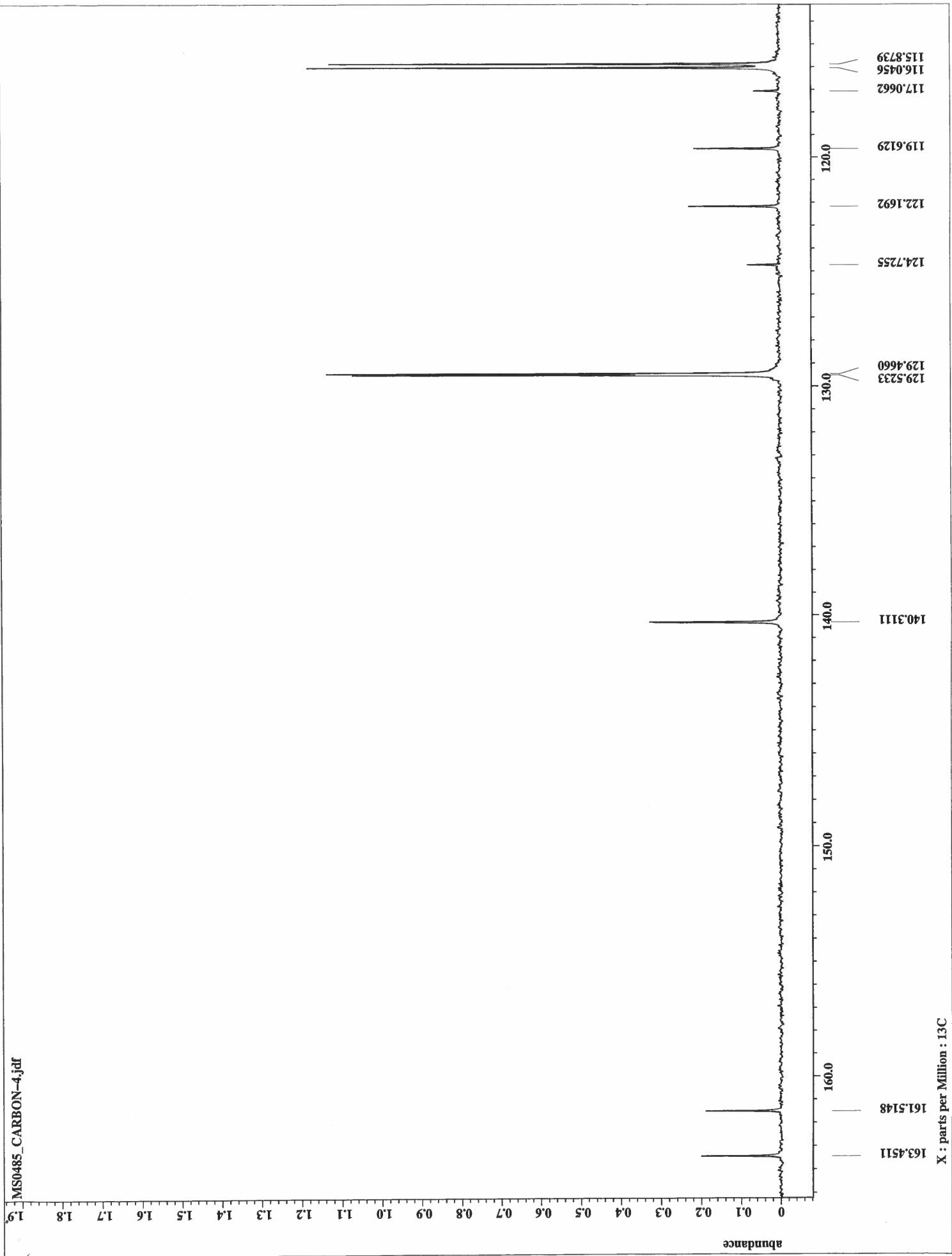
63.5371
 57.0033
 47.7988
 30.2674
 39.0999
 30.1148
 29.9526
 29.8000
 29.6474
 29.4948
 29.3422
 28.2739

140.3111
 129.5233
 122.1692
 119.6129
 116.0456

163.4511
 161.5148

X : parts per Million : 13C







```

filename = MS0485_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0485
Solvent = ACETONE-D6
Changer_sample = 12
Creation_time = 27-JUN-2018 11:34:58
Revision_time = 27-JUN-2018 11:12:05
Current_time = 27-JUN-2018 11:12:05

Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MHz])
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_prescans = 1
X_resolution = 1.7993855 [Hz]
X_sweep = 117.9245283 [Hz]
Irr_domain = 19F
Irr_freq = 470.62046084 [MHz]
Irr_offset = 5 [ppm]
Tri_domain = 19F
Tri_freq = 470.62046084 [MHz]
Tri_offset = 5 [ppm]
Clipped = FALSE
Mod_return = 1
Scans = 16
Total_scans = 16

X_90_width = 13.1 [us]
X_acq_time = 0.55574528 [s]
X_angle = 45 [deg]
X_atn = 2.5 [dB]
X_pulse = 6.55 [us]
Irr_mode = Off
Tri_mode = Off
Dante_pressat = FALSE
Initial_wait = 1 [s]
Recv_gain = 34
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.3 [dc]


```

-117.4832

-79.7153



abundance

X : parts per Million : 19F

checkCIF (basic structural check) running

Checking for embedded fcf data in CIF ...
 Found embedded fcf data in CIF. Extracting fcf data from uploaded CIF, please wait

checkCIF/PLATON (basic structural check)

Structure factors have been supplied for datablock(s) JD_POP_0m

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

No syntax errors found.
 Please wait while processing

[CIF dictionary](#)
[Interpreting this report](#)

Structure factor report

Datablock: JD_POP_0m

Bond precision:	C-C = 0.0046 Å	Wavelength=1.54178	
Cell:	a=18.1884 (8) alpha=90	b=9.1975 (4) beta=90	c=18.6788 (7) gamma=90
Temperature:	150 K		
	Calculated	Reported	
Volume	3124.7 (2)	3124.7 (2)	
Space group	P c a 21	P c a 21	
Hall group	P 2c -2ac	P 2c -2ac	
Moiety formula	C30 H24 O P, C2 F6 N O4 S2	C30 H24 O P, C2 F6 N O4 S2	
Sum formula	C32 H24 F6 N O5 P S2	C32 H24 F6 N O5 P S2	
Mr	711.61	711.61	
Dx, g cm-3	1.513	1.513	
Z	4	4	
Mu (mm-1)	2.739	2.739	
F000	1456.0	1456.0	
F000'	1464.89		
h,k,lmax	23,11,23	23,11,23	
Nref	6873 [3547]	5537	
Tmin, Tmax	0.663, 0.826	0.535, 0.754	
Tmin'	0.536		
Correction method= # Reported T Limits: Tmin=0.535 Tmax=0.754			
AbsCorr = MULTI-SCAN			
Data completeness=	1.56/0.81	Theta(max)=	80.721
R(reflections)=	0.0303 (5354)	wR2(reflections)=	0.0813 (5537)
S =	1.092	Npar=	561

The following ALERTS were generated. Each ALERT has the format

[test-name_ALERT_alert-type_alert-level](#).

Click on the hyperlinks for more details of the test.

● Alert level C

[PLAT089_ALERT_3_C](#) Poor Data / Parameter Ratio (Zmax < 18) 6.24 Note
[PLAT340_ALERT_3_C](#) Low Bond Precision on C-C Bonds 0.0046 Ang.
[PLAT911_ALERT_3_C](#) Missing FCF Refl Between Thmin & STh/L= 0.600 10 Report
[PLAT913_ALERT_3_C](#) Missing # of Very Strong Reflections in FCF 5 Note
[PLAT915_ALERT_3_C](#) No Flack x Check Done: Low Friedel Pair Coverage 61 %

● Alert level G

PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	30	Note
PLAT003_ALERT_2_G	Number of Uiso or Uij Restrained non-H Atoms ...	30	Report
PLAT175_ALERT_4_G	The CIF-Embedded .res File Contains SAME Records	1	Report
PLAT178_ALERT_4_G	The CIF-Embedded .res File Contains SIMU Records	1	Report
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 2)	100%	Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 3)	100%	Note
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 2	11.74	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 3	3.25	Check
PLAT432_ALERT_2_G	Short Inter X...Y Contact O4B ..C3	2.87	Ang.
	x,y,z = 1_555		Check
PLAT811_ALERT_5_G	No ADDSYM Analysis: Too Many Excluded Atoms		! Info
PLAT860_ALERT_3_G	Number of Least-Squares Restraints	532	Note
PLAT910_ALERT_3_G	Missing # of FCF Reflection(s) Below Theta(Min).	2	Note
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	33	Note
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.	7	Info

0 **ALERT level A** = Most likely a serious problem - resolve or explain

0 **ALERT level B** = A potentially serious problem, consider carefully

5 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight

14 **ALERT level G** = General information/check it is not something unexpected

0 ALERT type 1 CIF construction/syntax error, inconsistent or missing data

4 ALERT type 2 Indicator that the structure model may be wrong or deficient

7 ALERT type 3 Indicator that the structure quality may be low

7 ALERT type 4 Improvement, methodology, query or suggestion

1 ALERT type 5 Informative message, check

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

Publication of your CIF in IUCr journals

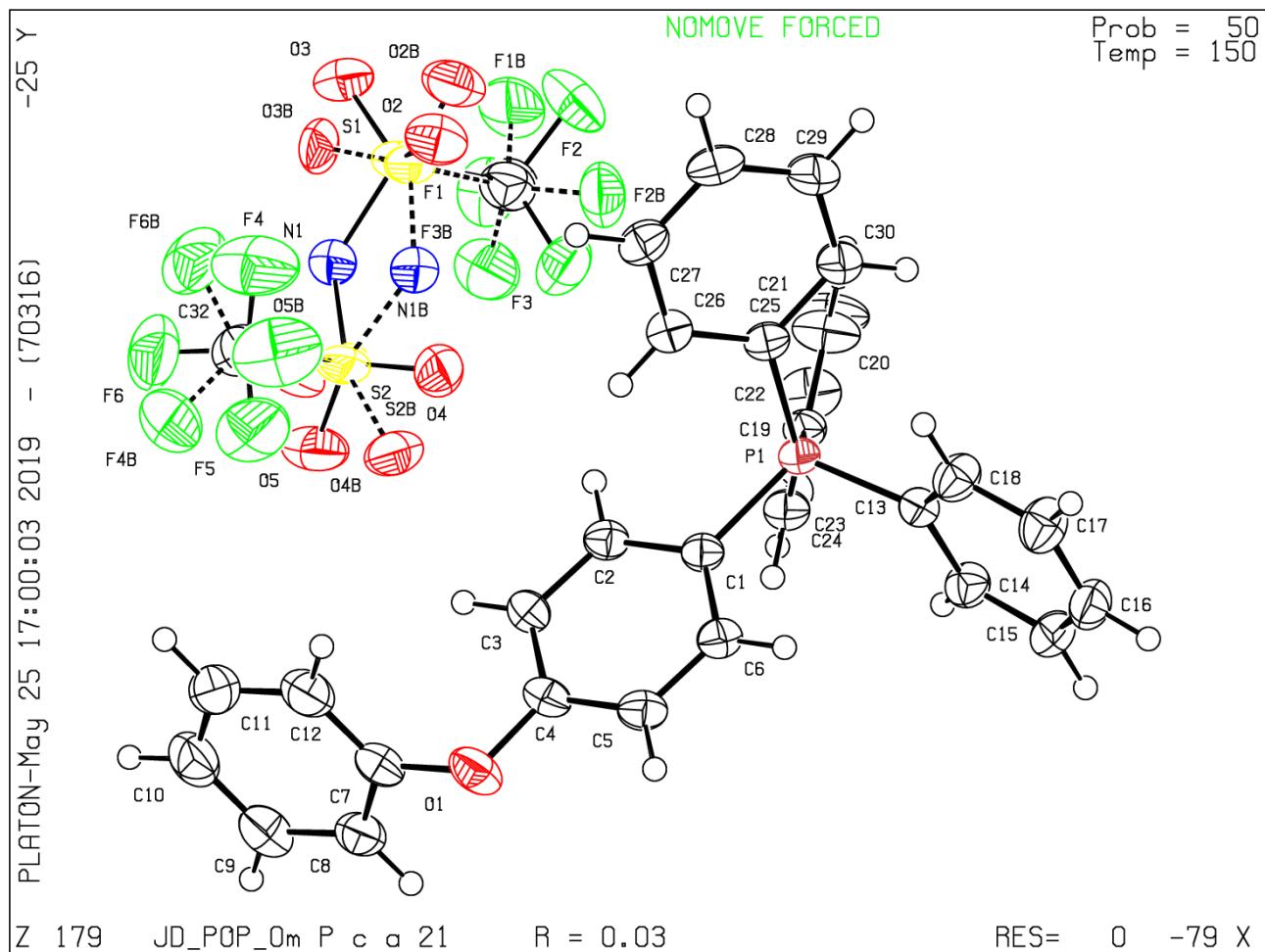
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that **full publication checks** are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 03/05/2019; check.def file version of 29/04/2019

Datablock JD_POP_0m - ellipsoid plot



[Download CIF editor \(publCIF\) from the IUCr](#)

[Download CIF editor \(enCIFer\) from the CCDC](#)

[Test a new CIF entry](#)

checkCIF (basic structural check) running

Checking for embedded fcif data in CIF ...

Found embedded fcif data in CIF. Extracting fcif data from uploaded CIF, please wait

checkCIF/PLATON (basic structural check)

Structure factors have been supplied for datablock(s) JD_POP_MF_0m_5

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A
REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT
REPLACE THE EXPERTISE OF AN EXPERIENCED
CRYSTALLOGRAPHIC REFEREE.

No syntax errors found.
Please wait while processing

[CIF dictionary](#)
[Interpreting this report](#)

[Structure factor report](#)

Datablock: JD_POP_MF_0m_5

Bond precision: C-C = 0.0040 Å Wavelength=0.71073
 Cell: a=18.0534(13) b=19.0695(14) c=20.2712(15)
 alpha=90 beta=114.239(3) gamma=90
 Temperature: 150 K

	Calculated	Reported
Volume	6363.5(8)	6363.5(8)
Space group	P 21/n	P 21/n
Hall group	-P 2yn	-P 2yn
Moiety formula	C30 H23 F O P, C2 F6 N O4 S2	C30 H23 F O P, C2 F6 N O4 S2
Sum formula	C32 H23 F7 N O5 P S2	C32 H23 F7 N O5 P S2
Mr	729.60	729.60
Dx, g cm ⁻³	1.523	1.523
Z	8	8
Mu (mm ⁻¹)	0.302	0.302
F000	2976.0	2976.0
F000'	2980.76	
h, k, lmax	26, 28, 29	26, 28, 29
Nref	21348	20359
Tmin, Tmax	0.871, 0.962	0.400, 0.746
Tmin'	0.865	
Correction method=	# Reported T Limits: Tmin=0.400	
Tmax=	0.746	AbsCorr = MULTI-SCAN
Data completeness=	0.954	Theta(max) = 31.592
R(reflections)=	0.0534(11965)	wR2(reflections) = 0.1450(20359)
S =	1.021	Npar= 1152

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.

Alert level C

[PLAT241_ALERT_2_C](#) High 'MainMol' Ueq as Compared to Neighbors of
[PLAT910_ALERT_3_C](#) Missing # of FCF Reflection(s) Below Theta(Min).

02 Check
10 Note

[PLAT911_ALERT_3_C](#) Missing FCF Refl Between Thmin & STh/L= 0.600
[PLAT913_ALERT_3_C](#) Missing # of Very Strong Reflections in FCF

58 Report
7 Note

● **Alert level G**

PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	68 Note
PLAT003_ALERT_2_G	Number of Uiso or Uij Restrained non-H Atoms ...	45 Report
PLAT171_ALERT_4_G	The CIF-Embedded .res File Contains EADP Records	1 Report
PLAT175_ALERT_4_G	The CIF-Embedded .res File Contains SAME Records	3 Report
PLAT176_ALERT_4_G	The CIF-Embedded .res File Contains SADI Records	1 Report
PLAT178_ALERT_4_G	The CIF-Embedded .res File Contains SIMU Records	1 Report
PLAT230_ALERT_2_G	Hirshfeld Test Diff for F1B --C5 .	6.1 s.u.
PLAT242_ALERT_2_G	Low 'MainMol' Ueq as Compared to Neighbors of	C61 Check
PLAT242_ALERT_2_G	Low 'MainMol' Ueq as Compared to Neighbors of	C62 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of S3	Constrained at 0.5 Check

And 14 other PLAT300 Alerts

PLAT300_ALERT_4_G	Atom Site Occupancy of S4	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F9	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F10	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F11	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F12	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F13	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F14	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O7	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O8	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O9	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O10	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N2	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C63	Constrained at 0.5 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C64	Constrained at 0.5 Check

PLAT301_ALERT_3_G	Main Residue Disorder(Resd 1)	3% Note
PLAT301_ALERT_3_G	Main Residue Disorder(Resd 2)	3% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 4)	100% Note

And 2 other PLAT302 Alerts

PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 5)	100% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 6)	100% Note

[PLAT304_ALERT_4_G](#) Non-Integer Number of Atoms in Resd 4 7.50 Check

And 2 other PLAT304 Alerts

PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 5	4.80 Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 6	2.70 Check

PLAT432_ALERT_2_G	Short Inter X...Y Contact O12B ..C6 x,y,z = 1_555	2.90 Ang. Check
PLAT789_ALERT_4_G	Atoms with Negative _atom_site_disorder_group #	45 Check
PLAT860_ALERT_3_G	Number of Least-Squares Restraints	786 Note
PLAT870_ALERT_4_G	ALERTS Related to Twinning Effects Suppressed ..	! Info
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	1245 Note
PLAT931_ALERT_5_G	CIFcalcFCF Twin Law (0 0 1) Est.d BASF	0.16 Check
PLAT933_ALERT_2_G	Number of OMIT Records in Embedded .res File ...	3 Note

0 **ALERT level A** = Most likely a serious problem - resolve or explain
0 **ALERT level B** = A potentially serious problem, consider carefully
4 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
39 **ALERT level G** = General information/check it is not something unexpected

0 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
8 ALERT type 2 Indicator that the structure model may be wrong or deficient
6 ALERT type 3 Indicator that the structure quality may be low
28 ALERT type 4 Improvement, methodology, query or suggestion
1 ALERT type 5 Informative message, check

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should

normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

Publication of your CIF in IUCr journals

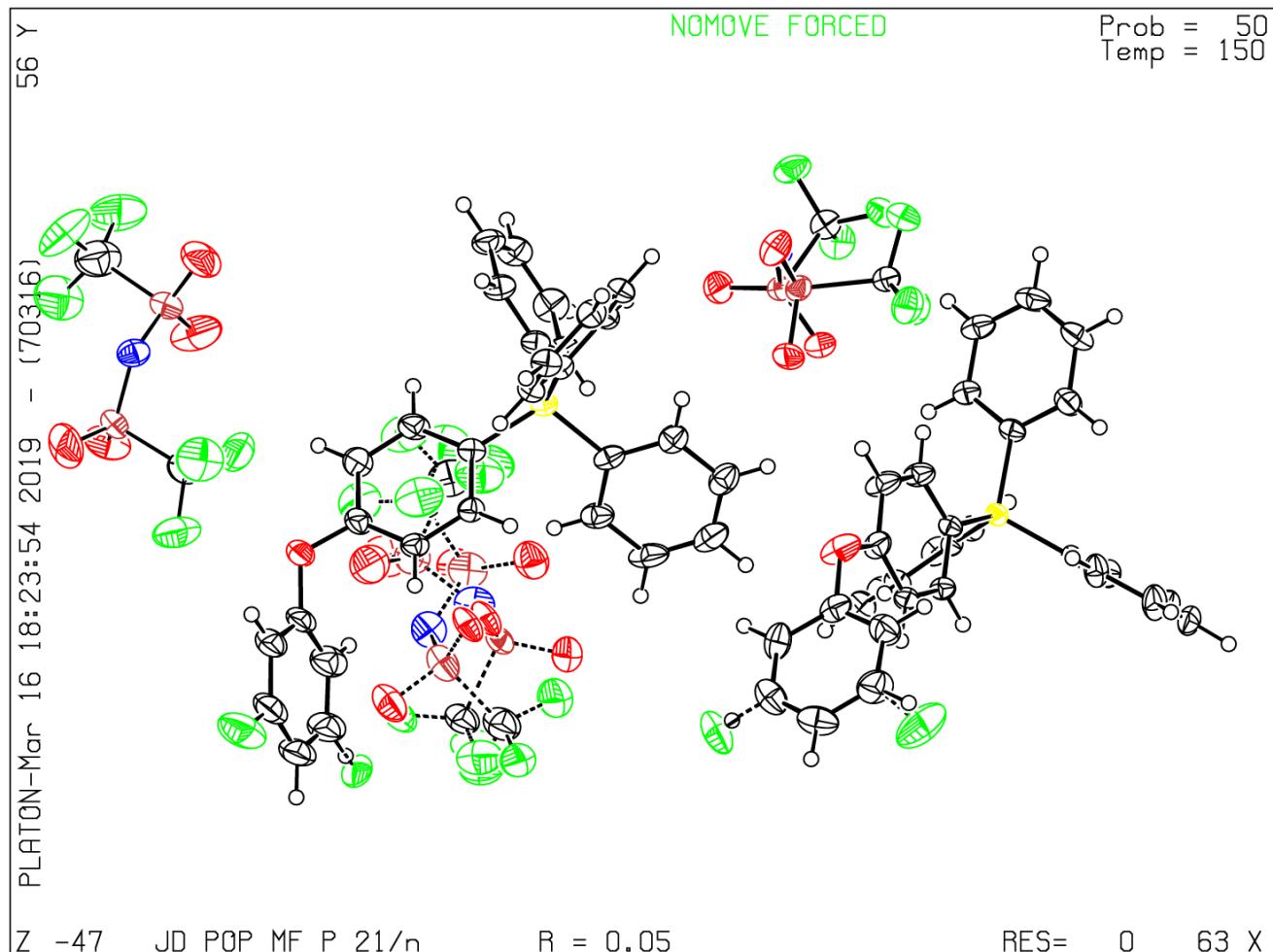
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that [full publication checks](#) are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 18/02/2019; check.def file version of 18/02/2019

Datablock JD_POP_MF_0m_5 - ellipsoid plot



[Download CIF editor \(publCIF\) from the IUCr](#)

[Download CIF editor \(enCIFer\) from the CCDC](#)

[Test a new CIF entry](#)

checkCIF (basic structural check) running

Checking for embedded fcif data in CIF ...

Found embedded fcif data in CIF. Extracting fcif data from uploaded CIF, please wait

checkCIF/PLATON (basic structural check)

Structure factors have been supplied for datablock(s) JD_POP_OF_0m

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A
REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT
REPLACE THE EXPERTISE OF AN EXPERIENCED
CRYSTALLOGRAPHIC REFEREE.

No syntax errors found.
Please wait while processing

[CIF dictionary](#)
[Interpreting this report](#)

[Structure factor report](#)

Datablock: JD_POP_OF_0m

Bond precision:	C-C = 0.0026 A	Wavelength=0.71073
Cell:	a=18.0015(14) b=18.7263(15) c=18.8837(16)	
	alpha=90 beta=90 gamma=90	
Temperature:	150 K	
	Calculated	Reported
Volume	6365.7(9)	6365.7(9)
Space group	P b c a	P b c a
Hall group	-P 2ac 2ab	-P 2ac 2ab
Moiety formula	C30 H23 F O P, C2 F6 N O4 S2	C30 H23 F O P, C2 F6 N O4 S2
Sum formula	C32 H23 F7 N O5 P S2	C32 H23 F7 N O5 P S2
Mr	729.60	729.60
Dx, g cm ⁻³	1.523	1.523
Z	8	8
Mu (mm ⁻¹)	0.301	0.301
F000	2976.0	2976.0
F000'	2980.76	
h, k, lmax	27, 28, 29	27, 28, 29
Nref	12223	12152
Tmin, Tmax	0.855, 0.939	0.689, 0.747
Tmin'	0.855	
Correction method=	# Reported T Limits: Tmin=0.689	
Tmax=0.747	AbsCorr = MULTI-SCAN	
Data completeness=	0.994	Theta(max) = 33.230
R(reflections)=	0.0579(8861)	wR2(reflections) = 0.1854(12152)
S =	1.046	Npar= 579

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.

● Alert level B

[PLAT430_ALERT_2_B](#) Short Inter D...A Contact 01 ..01 . 2.76 Ang.
 1-x,-y,1-z = 5_656 Check

● Alert level C

[PLAT905_ALERT_3_C](#) Negative K value in the Analysis of Variance ... -2.090 Report
[PLAT911_ALERT_3_C](#) Missing FCF Refl Between Thmin & STh/L= 0.600 4 Report

● Alert level G

PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	30	Note
PLAT003_ALERT_2_G	Number of Uiso or Uij Restrained non-H Atoms ...	30	Report
PLAT175_ALERT_4_G	The CIF-Embedded .res File Contains SAME Records	1	Report
PLAT178_ALERT_4_G	The CIF-Embedded .res File Contains SIMU Records	1	Report
PLAT230_ALERT_2_G	Hirshfeld Test Diff for F1 --C2 .	6.4	s.u.
PLAT301_ALERT_3_G	Main Residue Disorder(Resd 1)	3%	Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 2)	100%	Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 3)	100%	Note
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 2	12.68	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 3	2.32	Check
PLAT432_ALERT_2_G	Short Inter X...Y Contact 05B ..C16	2.95	Ang.
	1/2+x,y,1/2-z =	6_656	Check
PLAT811_ALERT_5_G	No ADDSYM Analysis: Too Many Excluded Atoms	! Info	
PLAT860_ALERT_3_G	Number of Least-Squares Restraints	513	Note
PLAT910_ALERT_3_G	Missing # of FCF Reflection(s) Below Theta(Min).	1	Note
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	67	Note
PLAT913_ALERT_3_G	Missing # of Very Strong Reflections in FCF	3	Note
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.	9	Info

0 **ALERT level A** = Most likely a serious problem - resolve or explain

1 **ALERT level B** = A potentially serious problem, consider carefully

2 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight

17 **ALERT level G** = General information/check it is not something unexpected

0 ALERT type 1 CIF construction/syntax error, inconsistent or missing data

6 ALERT type 2 Indicator that the structure model may be wrong or deficient

6 ALERT type 3 Indicator that the structure quality may be low

7 ALERT type 4 Improvement, methodology, query or suggestion

1 ALERT type 5 Informative message, check

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

Publication of your CIF in IUCr journals

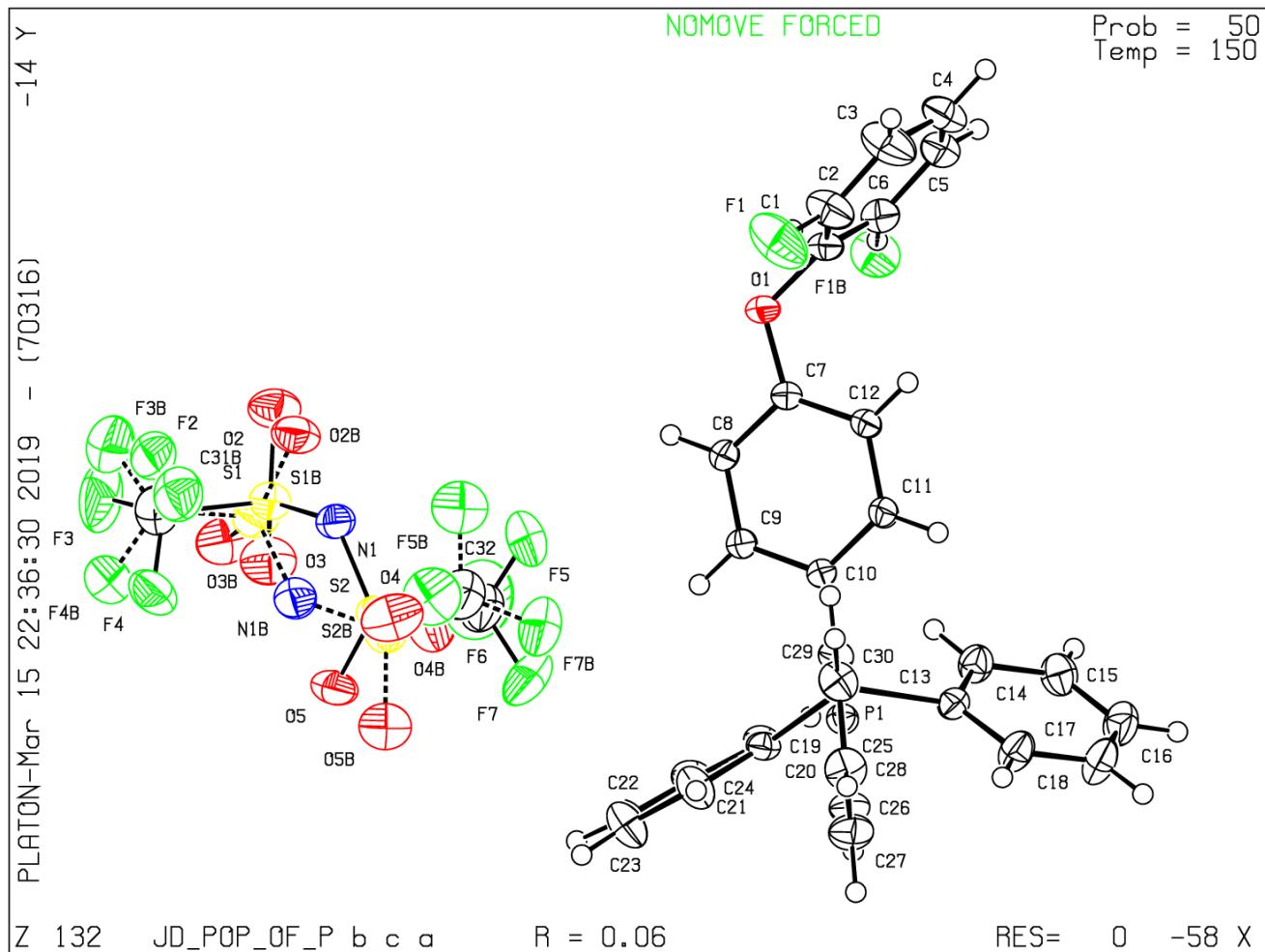
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that [full publication checks](#) are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 18/02/2019; check.def file version of 18/02/2019

Datablock JD_POP_OF_0m - ellipsoid plot



[Download CIF editor \(publCIF\) from the IUCr](#)

[Download CIF editor \(enCIFer\) from the CCDC](#)

[Test a new CIF entry](#)

checkCIF (basic structural check) running

Checking for embedded fcif data in CIF ...

Found embedded fcif data in CIF. Extracting fcif data from uploaded CIF, please wait

checkCIF/PLATON (basic structural check)

Structure factors have been supplied for datablock(s) JD_POP_PF_0m

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A
REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT
REPLACE THE EXPERTISE OF AN EXPERIENCED
CRYSTALLOGRAPHIC REFEREE.

No syntax errors found.
Please wait while processing

[CIF dictionary](#)
[Interpreting this report](#)

[Structure factor report](#)

Datablock: JD_POP_PF_0m

Bond precision: C-C = 0.0024 Å Wavelength=0.71073
Cell: a=9.5981(8) b=13.6324(11) c=22.2398(18)
alpha=102.938(3) beta=92.159(3) gamma=109.211(3)
Temperature: 150 K

	Calculated	Reported
Volume	2658.5(4)	2658.5(4)
Space group	P -1	P -1
Hall group	-P 1	-P 1
Moiety formula	C24 H19 F P, C2 F6 N O4 S2	C24 H19 F P, C2 F6 N O4 S2
Sum formula	C26 H19 F7 N O4 P S2	C26 H19 F7 N O4 P S2
Mr	637.50	637.51
Dx, g cm ⁻³	1.593	1.593
Z	4	4
μ (mm ⁻¹)	0.345	0.345
F000	1296.0	1296.0
F000'	1298.27	
h, k, lmax	13, 19, 32	13, 19, 32
Nref	17132	16986
Tmin, Tmax	0.921, 0.973	0.694, 0.746
Tmin'	0.921	
Correction method=	# Reported T Limits: Tmin=0.694	
Tmax=0.746	AbsCorr = MULTI-SCAN	
Data completeness=	0.991	Theta(max) = 31.139
R(reflections)=	0.0440(12700)	wR2(reflections)= 0.1276(16986)
S =	1.035	Npar= 1073

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.

Alert level C

[PLAT214_ALERT_2_C](#) Atom C51B (Anion/Solvent) ADP max/min Ratio 4.3 oblate
[PLAT906_ALERT_3_C](#) Large K Value in the Analysis of Variance 2.746 Check

[PLAT910_ALERT_3_C](#) Missing # of FCF Reflection(s) Below Theta(Min).
[PLAT911_ALERT_3_C](#) Missing FCF Refl Between Thmin & STh/L= 0.600

6 Note
36 Report

● Alert level G

PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	76 Note
PLAT003_ALERT_2_G	Number of Uiso or Uij Restrained non-H Atoms ...	60 Report
PLAT154_ALERT_1_G	The s.u.'s on the Cell Angles are Equal ..(Note)	0.003 Degree
PLAT175_ALERT_4_G	The CIF-Embedded .res File Contains SAME Records	3 Report
PLAT176_ALERT_4_G	The CIF-Embedded .res File Contains SADI Records	1 Report
PLAT178_ALERT_4_G	The CIF-Embedded .res File Contains SIMU Records	2 Report
PLAT230_ALERT_2_G	Hirshfeld Test Diff for F2 --C28 .	5.1 s.u.
PLAT230_ALERT_2_G	Hirshfeld Test Diff for S3 --C51 .	6.2 s.u.
PLAT301_ALERT_3_G	Main Residue Disorder(Resd 1)	4% Note
PLAT301_ALERT_3_G	Main Residue Disorder(Resd 2)	4% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 3)	100% Note

And 3 other PLAT302 Alerts

PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 4)	100% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 5)	100% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 6)	100% Note

[PLAT304_ALERT_4_G](#) Non-Integer Number of Atoms in Resd 3 14.25 Check

And 3 other PLAT304 Alerts

PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 4	13.05 Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 5	0.75 Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms in Resd 6	1.95 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O3 ..C39 3.01 Ang.
x,y,z = 1_555 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact F1C ..C3 2.93 Ang.
-1+x,y,z = 1_455 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact F5B ..C39 2.89 Ang.
2-x,2-y,1-z = 2_776 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O1B ..C11 2.82 Ang.
1-x,1-y,1-z = 2_666 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O1B ..C12 2.90 Ang.
1-x,1-y,1-z = 2_666 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O2B ..C9 2.91 Ang.
2-x,1-y,1-z = 2_766 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O5B ..C47 2.85 Ang.
1+x,y,z = 1_655 Check

[PLAT432_ALERT_2_G](#) Short Inter X...Y Contact O6B ..C26 2.89 Ang.
1-x,1-y,-z = 2_665 Check

[PLAT811_ALERT_5_G](#) No ADDSYM Analysis: Too Many Excluded Atoms ! Info

[PLAT860_ALERT_3_G](#) Number of Least-Squares Restraints 1242 Note

[PLAT912_ALERT_4_G](#) Missing # of FCF Reflections Above STh/L= 0.600 103 Note

[PLAT913_ALERT_3_G](#) Missing # of Very Strong Reflections in FCF 3 Note

[PLAT933_ALERT_2_G](#) Number of OMIT Records in Embedded .res File ... 3 Note

[PLAT978_ALERT_2_G](#) Number C-C Bonds with Positive Residual Density. 12 Info

0 **ALERT level A** = Most likely a serious problem - resolve or explain

0 **ALERT level B** = A potentially serious problem, consider carefully

4 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight

32 **ALERT level G** = General information/check it is not something unexpected

1 ALERT type 1 CIF construction/syntax error, inconsistent or missing data

15 ALERT type 2 Indicator that the structure model may be wrong or deficient

7 ALERT type 3 Indicator that the structure quality may be low

12 ALERT type 4 Improvement, methodology, query or suggestion

1 ALERT type 5 Informative message, check

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual

parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

Publication of your CIF in IUCr journals

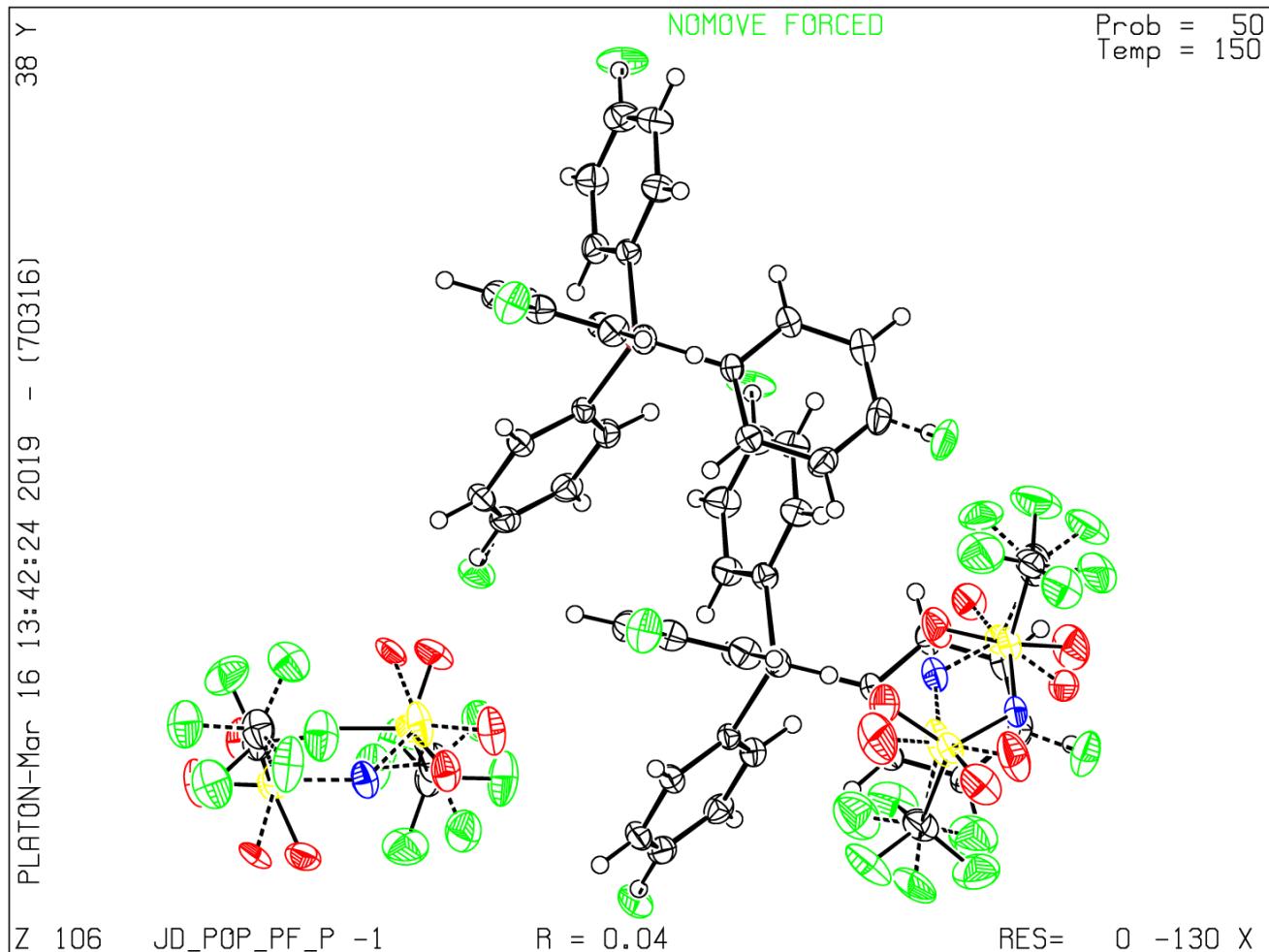
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that [full publication checks](#) are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

PLATON version of 18/02/2019; check.def file version of 18/02/2019

Datablock JD_POP_PF_0m - ellipsoid plot



[Test a new CIF entry](#)

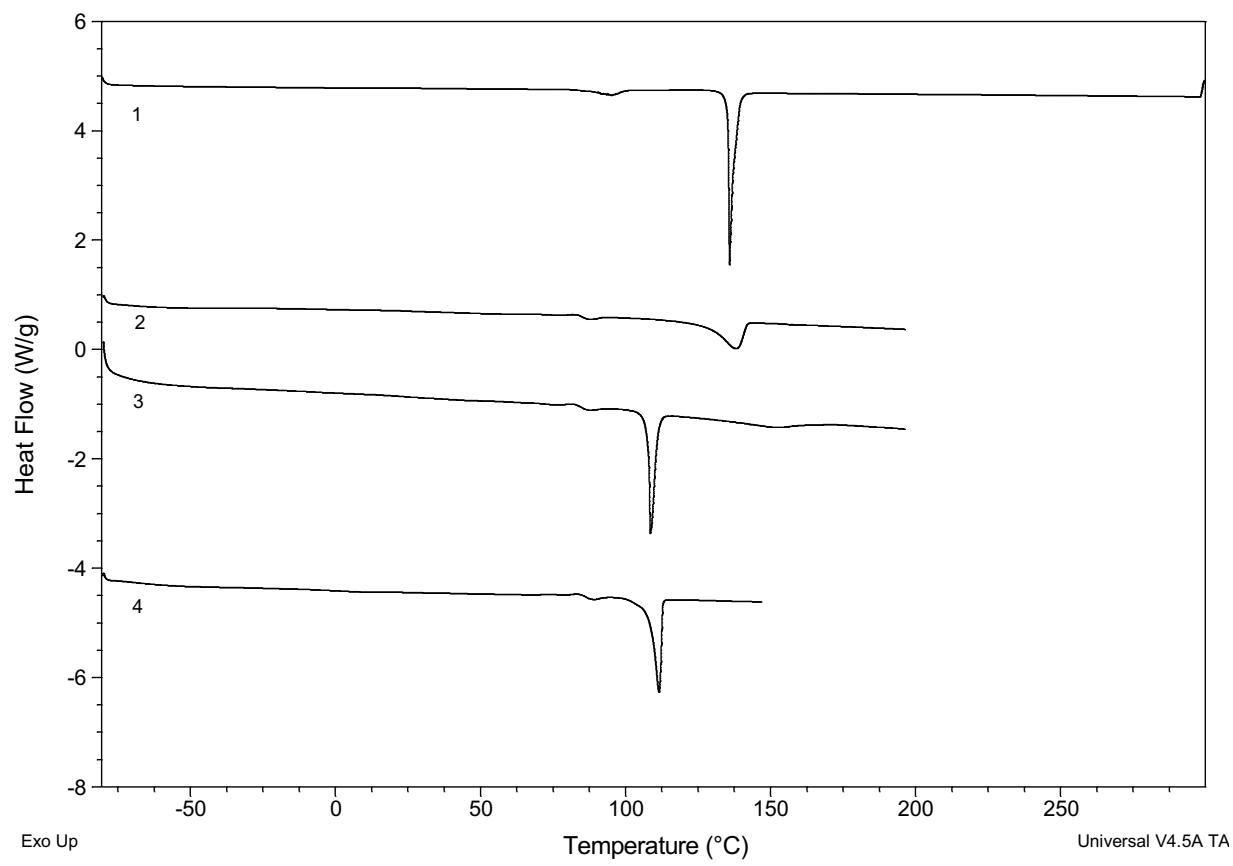


Figure S1: Representative DSC scans for compounds **1-4**.

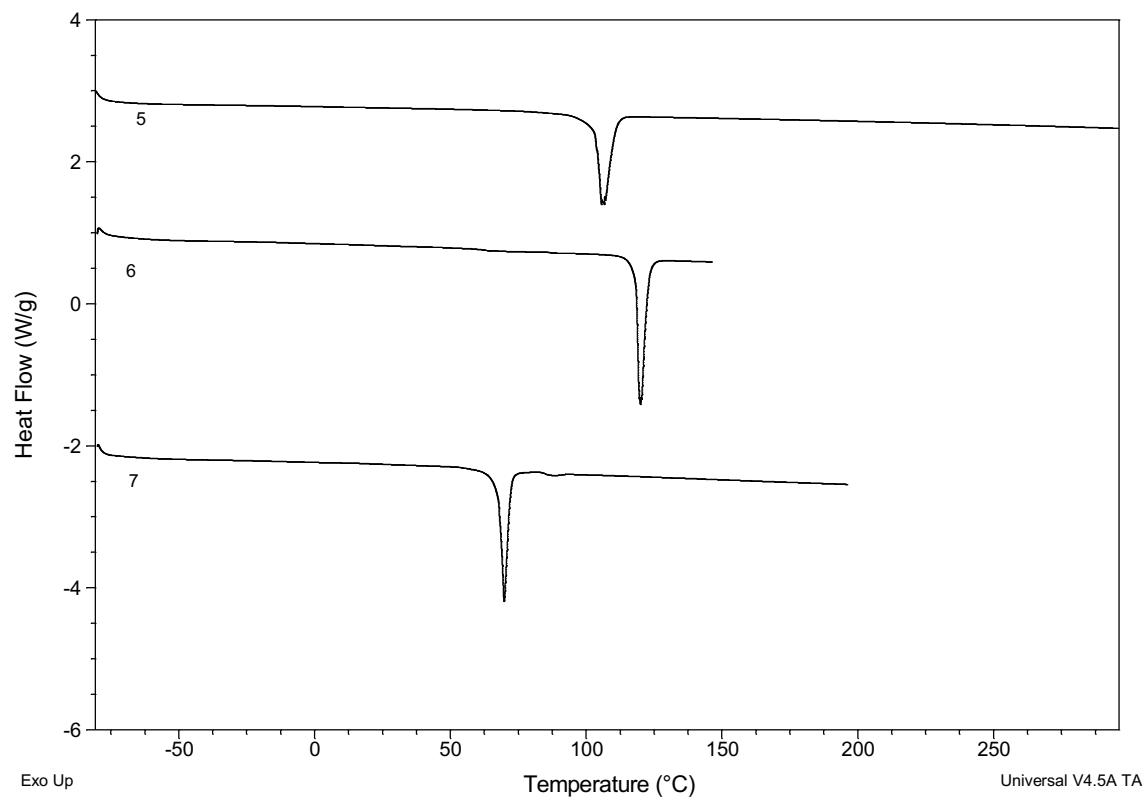


Figure S2: Representative DSC scans for compounds **5-7**.

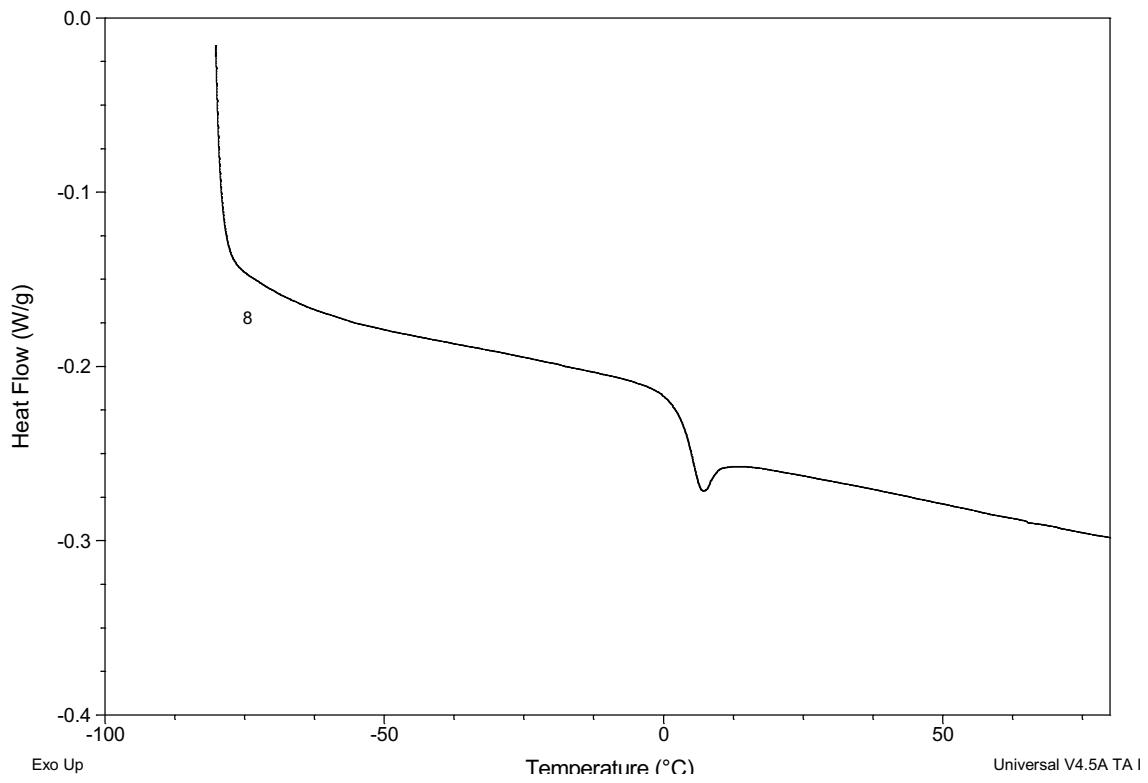


Figure S3: Representative DSC scan for compound **8**, which exhibits a weak glass transition.

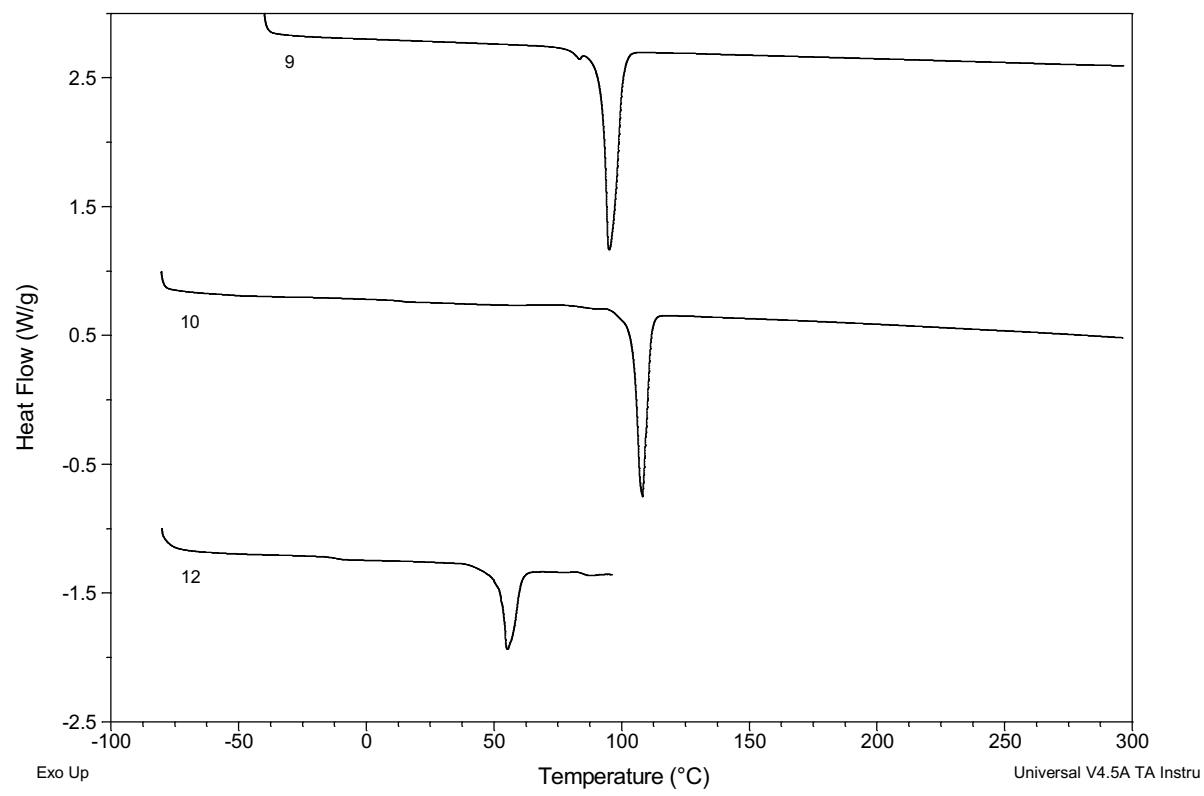


Figure S4: Representative DSC scans for compounds **9,10** and **12**.

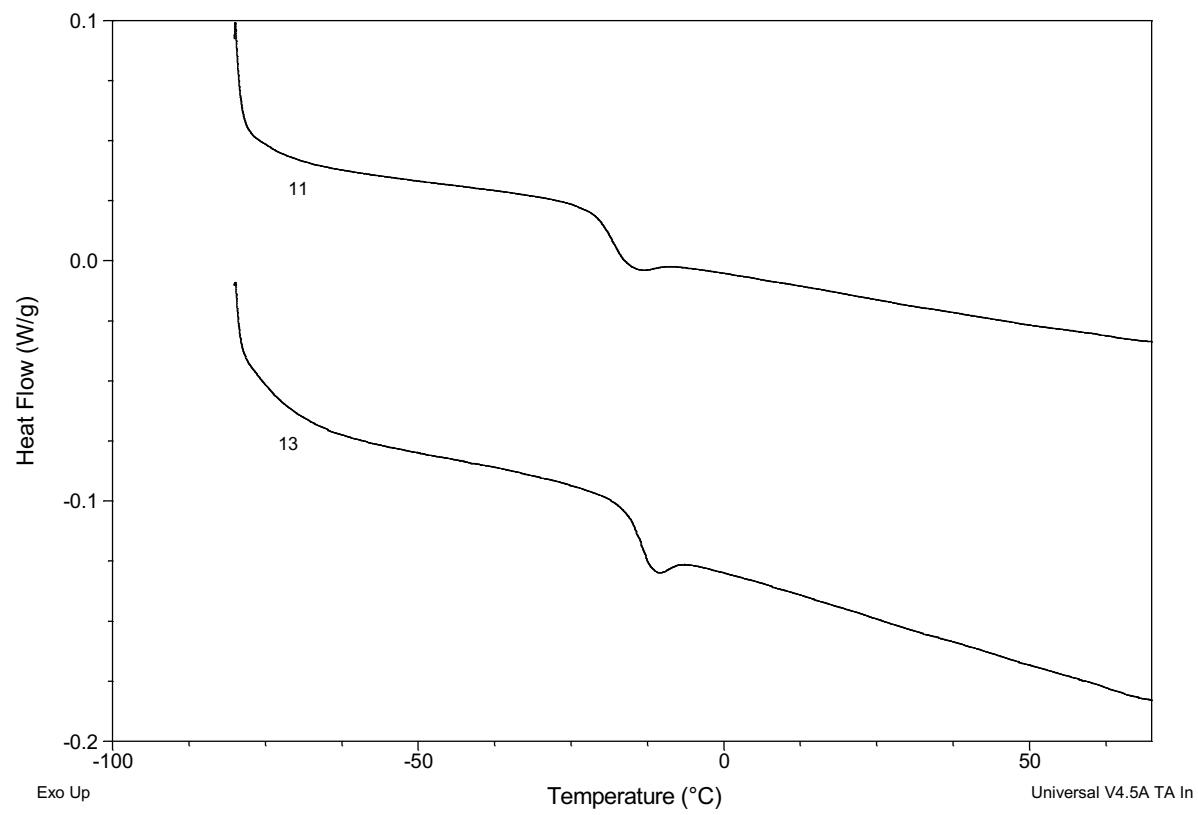


Figure S5: Representative DSC scans for compounds **11** and **13**, which exhibit weak glass transitions.

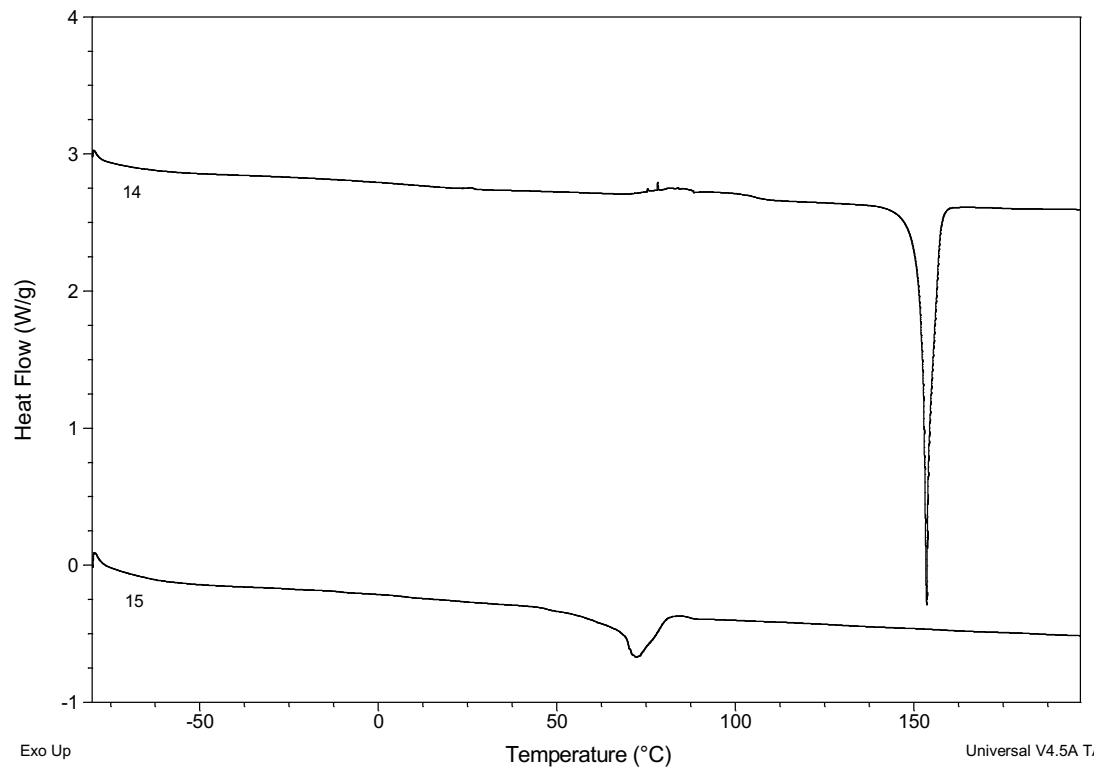


Figure S6: Representative DSC scans for compounds **14** and **15**.

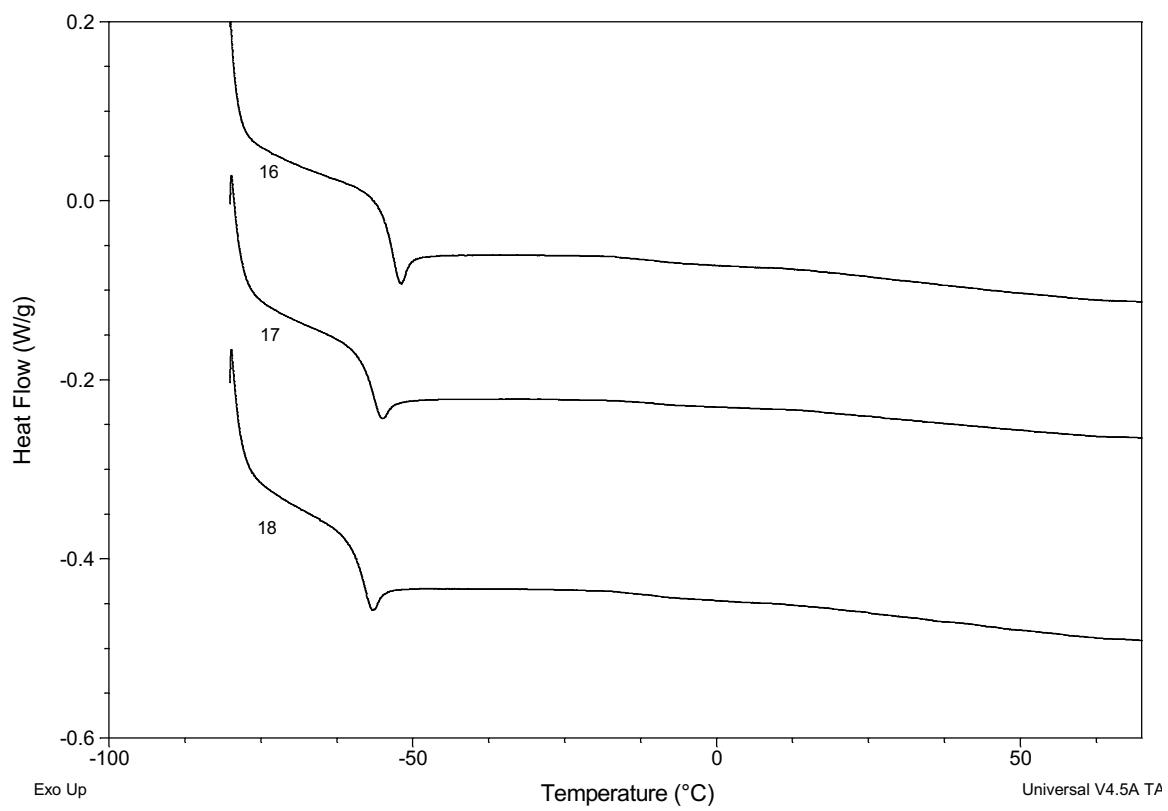


Figure S7: Representative DSC scans for compounds **16-18**, and **13**, which exhibit weak glass transitions.

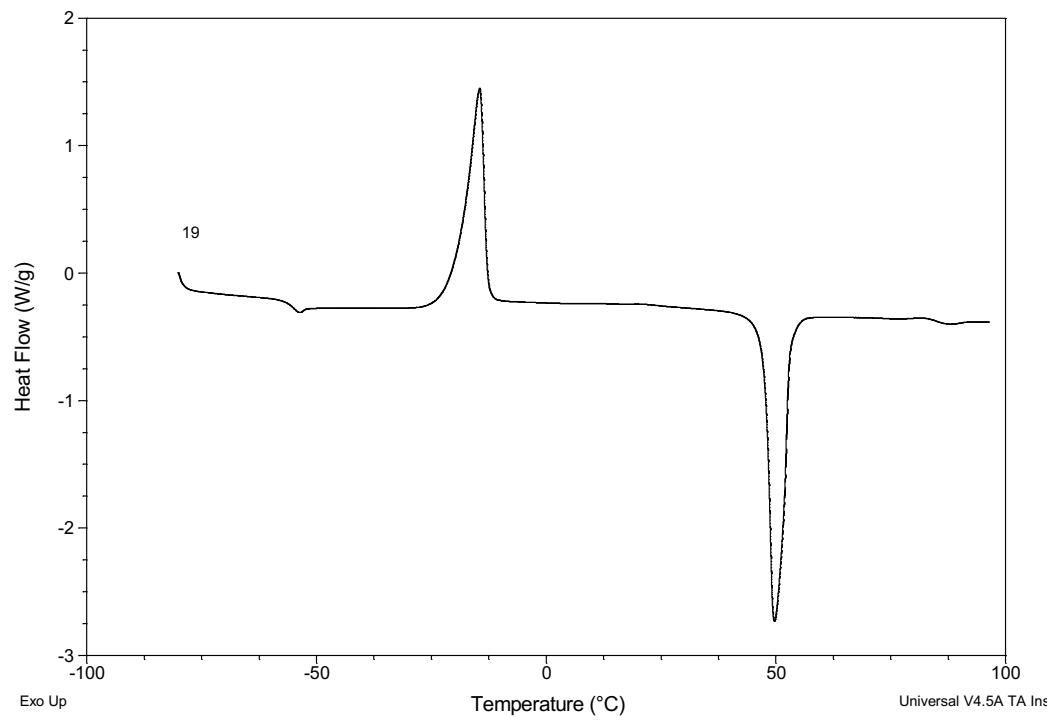


Figure S8: Representative DSC scan for compound **19**.

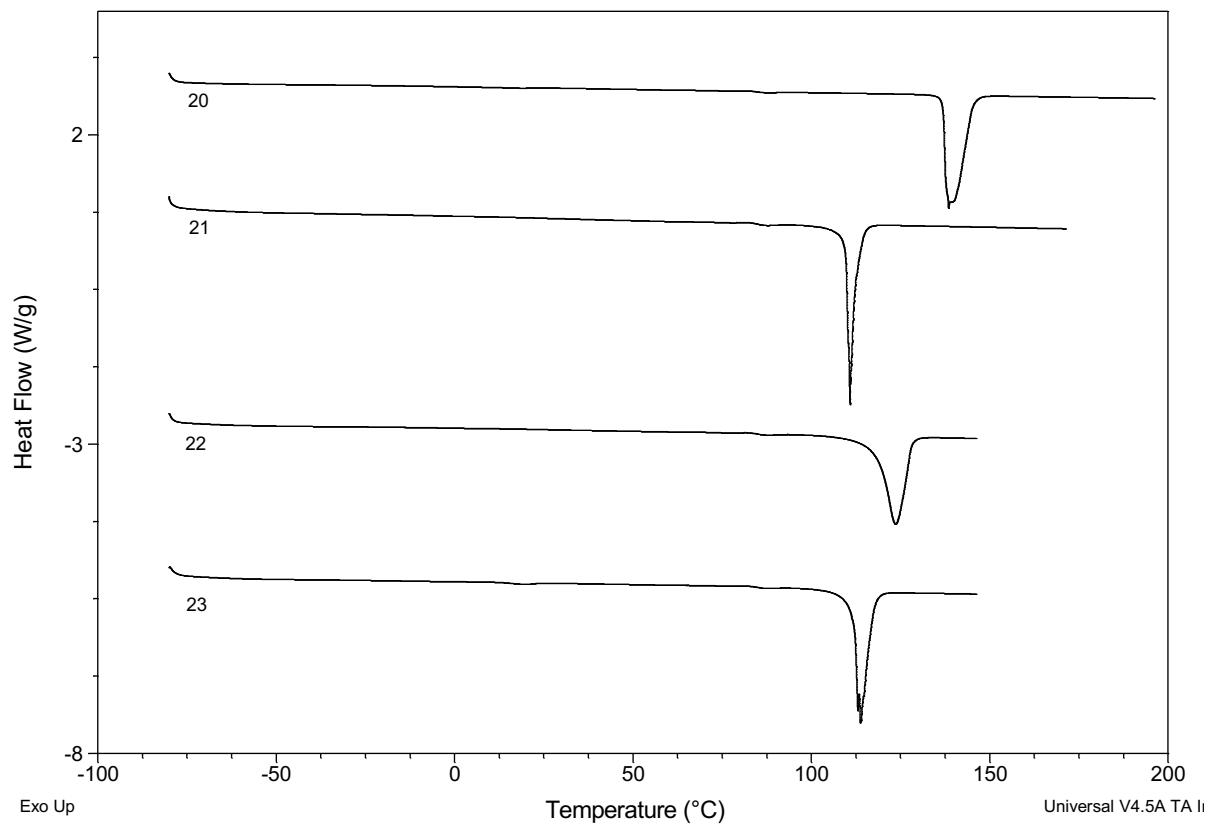


Figure S9: Representative DSC scans for compounds **20-23**.

Details of the Single Crystal X-Ray Diffraction Data Collection

CRYSTAL GROWTH:

Crystals of the salts were grown by slow evaporation from solutions in methanol. The crystals were collected by decantation, dried in vacuo, and transferred into a glass vial for storage.

SAMPLE PREPARATION:

Single crystals were taken out of the vial by either spatula (for dry samples) or via pasteur pipette (for crystals in mother liquor) and immersed in a few drops of Fomblin oil (a perfluoroether). Crystals were separated from each other with a MiTeGen micromesh mount (on a magnetic snap on mount), cut to size when required using triangular tipped a carpet cutting razor blade, and picked up on a MiTeGen micromesh mount from the Fomblin oil (using as little oil as possible). The MiTeGen micromesh mount on the magnetic snap on mount was transferred into the cold stream of the diffractometer and flash cooled.

INSTRUMENTATION:	Bruker AXS D8 Quest CMOS diffractometer
RADIATION:	Triumph curved graphite crystal
SOFTWARE PACKAGES:	Apex3 v2018.1-0 (Bruker, 2018) SAINT V8.38A (Bruker, 2016) SHELXS-97 (Sheldrick, 2008)
DATA COLLECTION:	omega and phi scans

Software and Refinement Methods

DATA COLLECTION:	Apex3 v2018.1-0 (Bruker, 2018)
CELL REFINEMENT:	SAINT V8.38A (Bruker, 2016)
DATA REDUCTION:	SAINT V8.38A (Bruker, 2016)
STRUCTURE SOLUTION:	SHELXS-97 (Sheldrick, 2008)

Special Details of Restraints and Methods

Cation 5, P(Ph)₃-p-POP⁺

```
_reflns_special_details
;
Reflections were merged by SHELXL according to the crystal class for the calculation of
statistics and refinement.
```

_reflns_Friedel_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences. Completeness statistics refer to single and composite reflections containing twin component 1 only.

```
;
```

```
_computing_data_collection      'Apex3 v2018.7-2 (Bruker, 2017)'
_computing_cell_refinement      'SAINT V8.38A (Bruker, 2016)'
```

```
_computing_data_reduction      'SAINT V8.38A (Bruker, 2016)'  
_computing_structure_solution  'SHELXS-97 (Sheldrick, 2008)'  
_computing_structure_refinement  
;  
SHELXL-2018/3 (Sheldrick, 2015, 2018),  
SHELXLE Rev946 (H\"ubschle et al., 2011)  
;  
_computing_molecular_graphics ?  
_computing_publication_material ?  
_refine_special_details  
;  
Refined as a 2-component inversion twin.
```

The anion is disordered by a pseudo-mirror operation. The two disordered moieties were restrained to have similar geometries. Uij components of ADPs for disordered atoms closer to each other than 2.0 Angstrom were restrained to be similar. Subject to these conditions the occupancy ratio refined to 0.783(3) to 0.217(3).

Cation 6, P(Ph)₃-p-(o-F-POP)⁺

```
_reflns_special_details  
;  
Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.  
  
_reflns_Friedel_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.  
;  
_computing_data_collection      'Apex3 v2018.1-0 (Bruker, 2018)'  
_computing_cell_refinement      'SAINT V8.38A (Bruker, 2016)'  
_computing_data_reduction      'SAINT V8.38A (Bruker, 2016)'  
_computing_structure_solution  'SHELXS-97 (Sheldrick, 2008)'  
_computing_structure_refinement  
;  
SHELXL-2018/3 (Sheldrick, 2015, 2018),  
SHELXLE Rev946 (H\"ubschle et al., 2011)  
;  
_computing_molecular_graphics ?  
_computing_publication_material ?  
_refine_special_details  
;
```

The fluorine atom of the fluorophenyl group is disordered over the two ortho positions. The occupancy ratio refined to 0.683(4) to 0.683(4). The anion is disordered over two orientations, with indication of additional disorder which was not sufficiently defined for unambiguous refinement. The two disordered moieties were restrained to have similar geometries. Uij components of ADPs for disordered atoms closer to each other than 2.0 Angstrom were restrained to be similar. Subject to these conditions the occupancy ratio refined to 0.8456(13) to 0.1544(13).

NOTE: The refined structure for this compound had a B-level alert for an O-O contact between the cations. This is due to the dipole-dipole interaction between the C-O bonds of neighboring cations, which overcomes the expected O-O repulsion and the alert regarded as a false alarm.

Cation 7, P(Ph)₃-p-(m-F-POP)⁺

```
_reflns_special_details
;
_reflns_Friedel_fraction is defined as the number of unique Friedel pairs measured divided by
the number that would be possible theoretically, ignoring centric projections and systematic
absences. Completeness statistics refer to single and composite reflections containing twin
component 1 only.
;

_computing_data_collection      'Apex3 v2018.1-0 (Bruker, 2018)'
_computing_cell_refinement      'SAINT V8.38A (Bruker, 2016)'
_computing_data_reduction       'SAINT V8.38A (Bruker, 2016)'
_computing_structure_solution   'SHELXS-97 (Sheldrick, 2008)'
_computing_structure_refinement
;
SHELXL-2018/3 (Sheldrick, 2015, 2018),
SHELXLE Rev946 (H\"ubschle et al., 2011)
;
_computing_molecular_graphics  ?
_computing_publication_material ?
_refine_special_details
;
The crystal under investigation was found to be non-merohedrally twinned. The orientation
matrices for the two components were identified using the program Cell_Now, with the two
components being related by a 180 degree rotation around the real a-axis. The two
components were integrated using Saint and corrected for absorption using twinabs, resulting
in the following statistics:
```

15502 data (4686 unique) involve domain 1 only, mean I/sigma 23.1
15148 data (4609 unique) involve domain 2 only, mean I/sigma 11.5
75199 data (19329 unique) involve 2 domains, mean I/sigma 14.4
19 data (19 unique) involve 3 domains, mean I/sigma 11.7

The exact twin matrix identified by the integration program was found to be:

```
1.00019 0.00032 0.00040
-0.00035 -1.00000 0.00029
-0.92160 -0.00042 -1.00019
```

The structure was solved using direct methods with only the non-overlapping reflections of component 1. The structure was refined using the hklf 5 routine with all reflections of component 1 (including the overlapping ones), resulting in a BASF value of 0.1967(6).

The Rint value given is for all reflections and is based on agreement between observed single and composite intensities and those calculated from refined unique intensities and twin fractions (TWINABS (Sheldrick, 2012)).

In both cations, the fluorine atom of the fluorophenyl group is disordered over the two meta positions. The C-F distances in the cations were restrained to be similar. Subject to these conditions the occupancy ratio refined to 0.835(6) to 0.165(6) in the cation of P1, and to 0.833(5) to 0.167(5) in the cation of P2.

Two anions are located on inversion centers and are disordered around these. Of these, one is in addition disordered over two alternative positions not related by inversion. The three disordered moieties were restrained to have similar geometries as the one not disordered anion in a general position. Uij components of ADPs for disordered atoms closer to each other than 2.0 Angstrom were restrained to be similar. Subject to these conditions the occupancy rates for the molecule of N3 and N3B refined to two times 0.3200(12) and two times 0.1800(12).

;

Cation 4, P(Ph)₃-p-F-POP⁺

_reflns_special_details

;

Reflections were merged by SHELXL according to the crystal class for the calculation of statistics and refinement.

_reflns_Friedel_fraction is defined as the number of unique Friedel pairs measured divided by the number that would be possible theoretically, ignoring centric projections and systematic absences.

;

_computing_data_collection 'Apex3 v2018.1-0 (Bruker, 2018)'

_computing_cell_refinement 'SAINT V8.38A (Bruker, 2016)'

_computing_data_reduction 'SAINT V8.38A (Bruker, 2016)'

_computing_structure_solution 'SHELXS-97 (Sheldrick, 2008)'

_computing_structure_refinement

;

SHELXL-2018/3 (Sheldrick, 2015, 2018),

SHELXLE Rev946 (H\"ubschle et al., 2011)

;

_computing_molecular_graphics ?

_computing_publication_material ?

_refine_special_details

;

In both p-F-Ph-PPh₃ cations the fluorine atom is disordered over all four possible para positions. All C-F bond lengths in the cations were restrained to be similar. The total occupancy was each constrained to one fluorine. Subject to these conditions the occupancy rates for the F atoms refined to 0.493(3), 0.080(3), 0.175(3) and 0.251(3) for the cation of P1, and to 0.497(3), 0.079(3), 0.248(3) and 0.176(3) for the cation of P2.

The two anions are disordered over each two orientations. The four disordered moieties were restrained to have similar geometries. Uij components of ADPs for disordered atoms closer to each other than 2.0 Angstrom were restrained to be similar. Subject to these conditions the occupancy ratio refined to 0.9503(12) to 0.0497(12) for the molecule of N1, and to 0.8698(16) to 0.1302(16) for the molecule of N2.

;