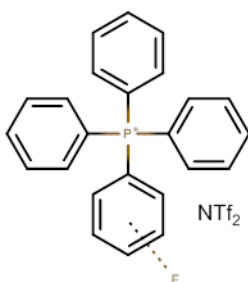


Tuning the Melting Point of Selected Ionic Liquids Through Adjustment of the Cation's Dipole Moment

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.

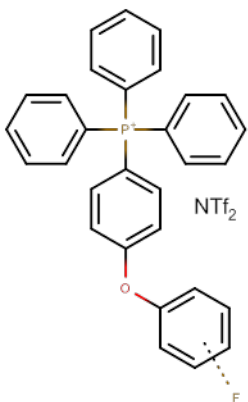
General Procedure for the Synthesis of Fluorophenyltriphenyl Phosphonium $[\text{NTf}_2]^-$ ILs



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

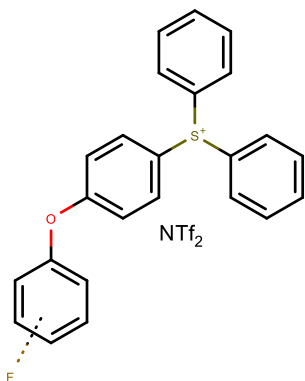


First 3-(fluorophenoxy)bromobenzene was prepared using following method: In a 100 ml heavy wall pressure vessels with an internal thread containing steering bar, 3-fluorophenylboronic acid (2.0 equiv, 10 g), 4-bromophenol (1.0 equiv), triethylamine (5.0 equiv), Copper(II) acetate (1.0 equiv), and MS 4\AA (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

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 $[\text{NTf}_2]^-$.

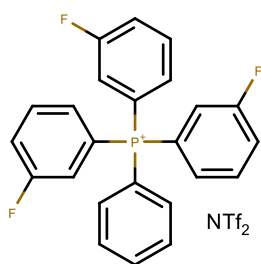
General Procedure for the Synthesis of triphenylsulfonium $[\text{NTf}_2]^-$ ILs



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

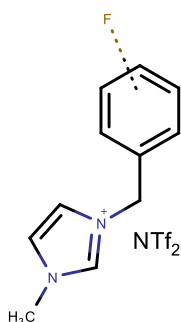


In a 50 ml heavy wall pressure vessels with an internal thread containing steering bar, Bromobenzene (1.0 equiv), Tris(3-fluorophenyl)phosphine (1.0 equiv), $\text{NiCl}_2 \cdot 6\text{H}_2\text{O}$ (10 mole%) ethylene glycol (10 ml) were added under nitrogen atmosphere and the reaction mixture stirred 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 $[\text{NTf}_2]^-$.

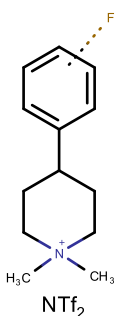
Synthesis of 3-benzyl-1-methyl-imidazolium $[\text{NTf}_2]^-$ ILs



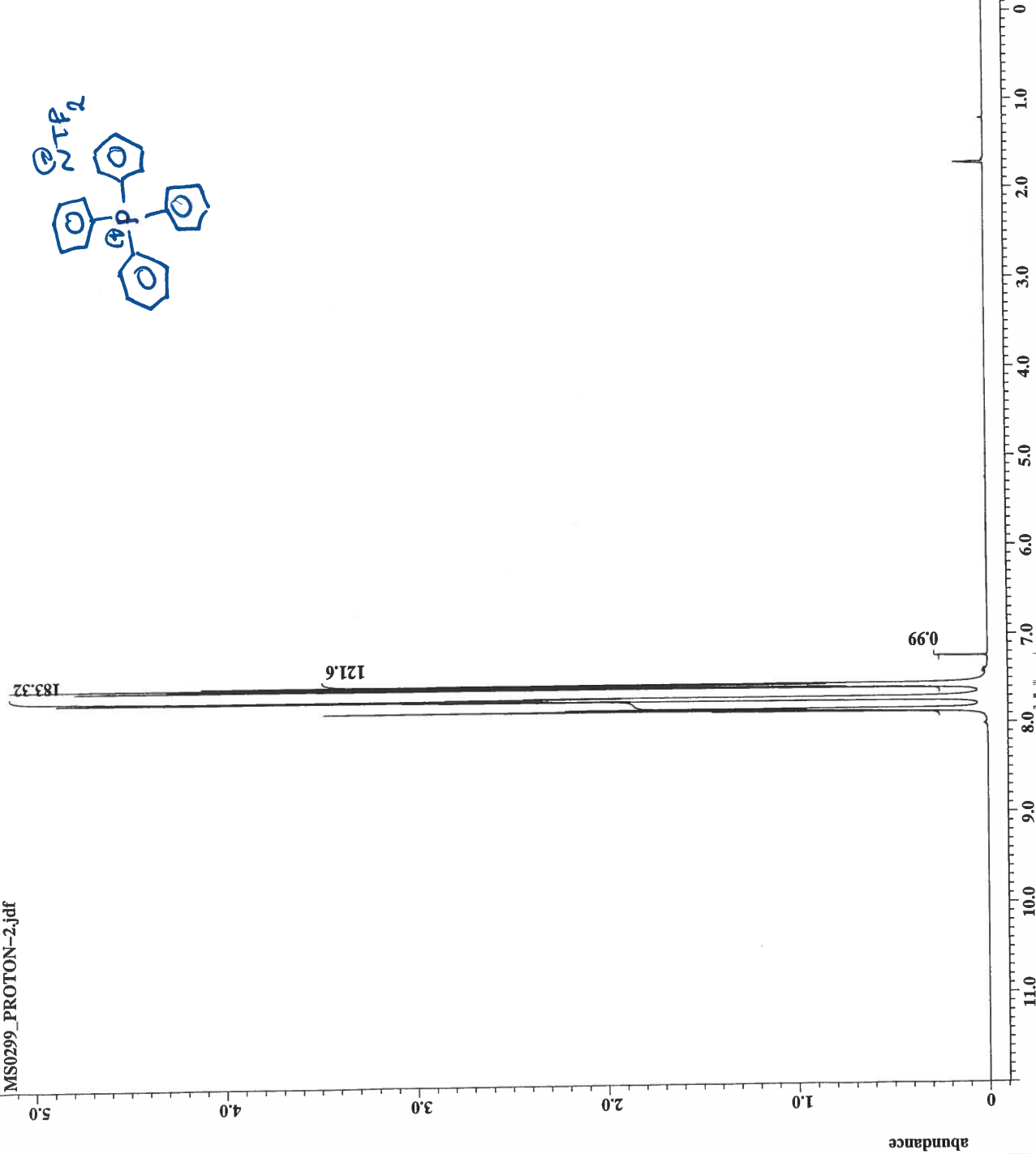
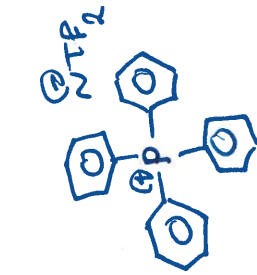
In a 100 ml round bottom flask containing a steering bar, 1-methylimidazol (1.0 equiv, 5g) and Benzyl Bromide (1.0 equiv) in THF (50 ml) was stirred for overnight at 40 °C. Then the reaction mixture was filtered and white solid yielded as a pure product. Anion exchange with KNTf_2 (1.0 equiv) carried 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



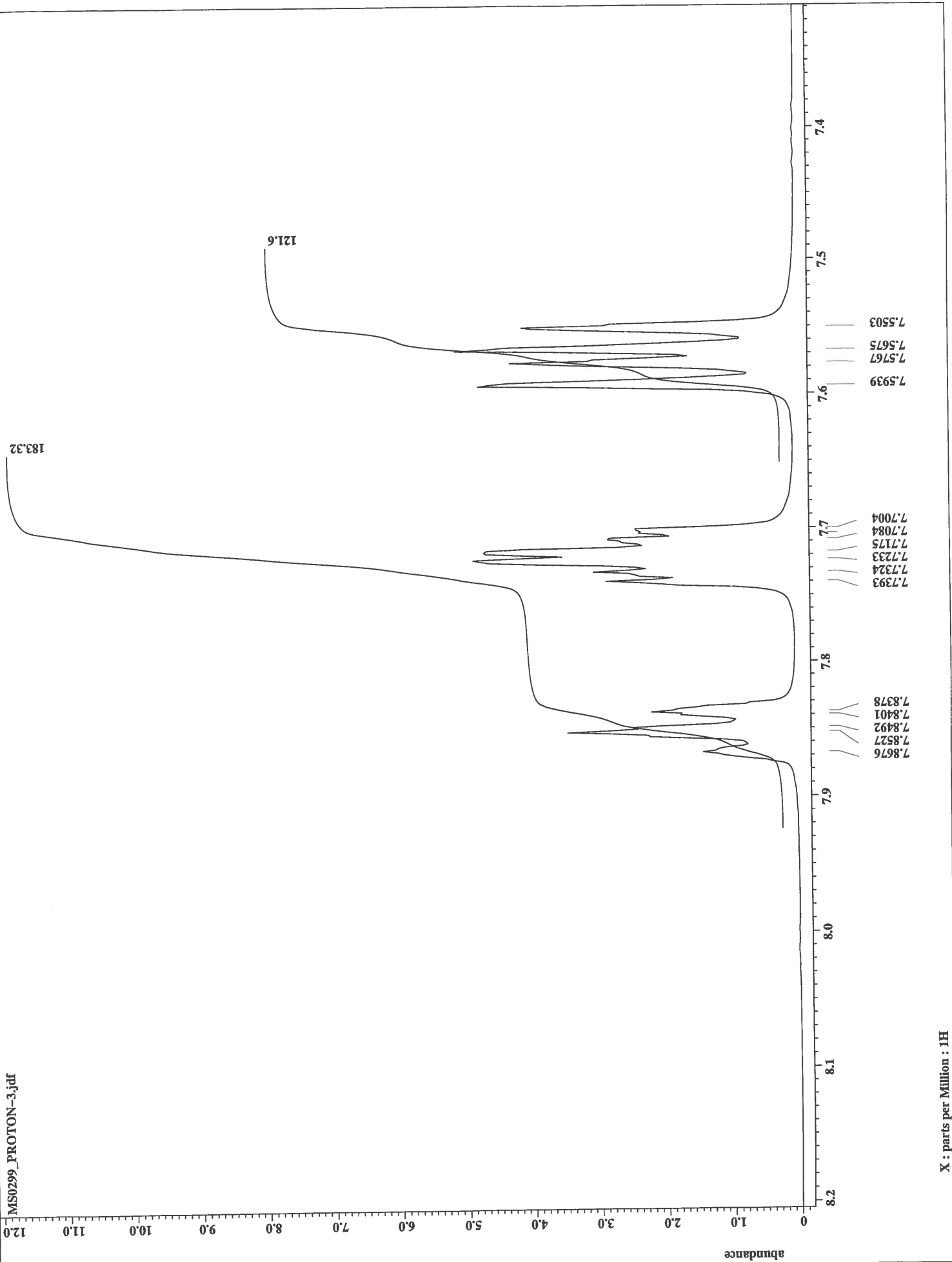
For the synthesis of 1,1-dimethyl-4-phenylpiperidinium $[\text{NTf}_2]^-$, first 1,1-dimethyl-4-phenylpiperidinium iodide salt was prepared using following method. In a 100 ml round bottom flask containing a stirring bar, 4-phenylpiperidine (1.0 equiv, 2 g), iodomethane (5.0 equiv), and potassium bicarbonate (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.



X : parts per Million : 1H



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 Irr_freq = 500.15991521[MHz]
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 Dante_presat = FALSE
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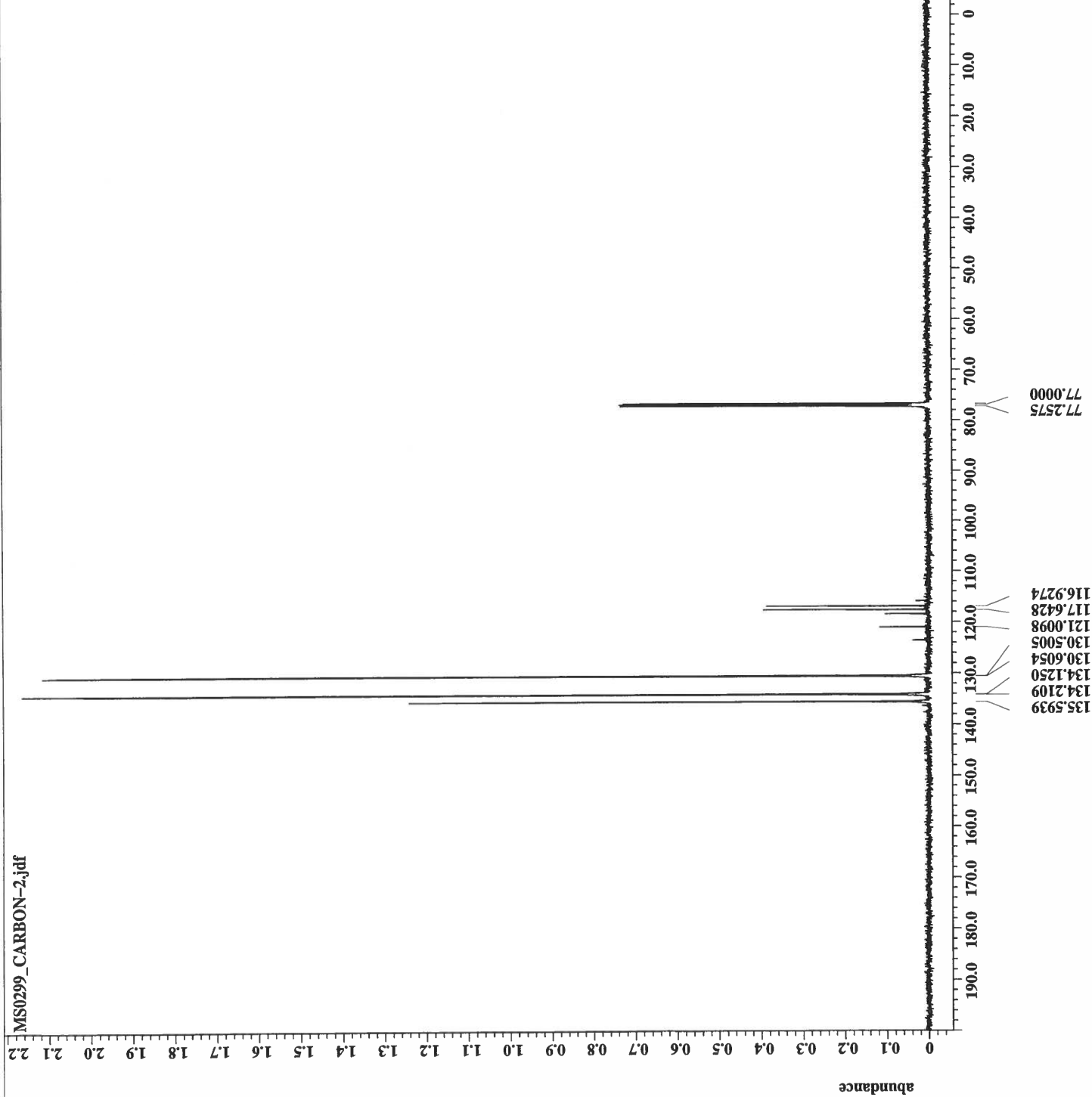
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Changer_sample
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Revision_time 9-JUN-2018 17:53:10
Current_time  9-JUN-2018 17:53:10

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X_pulse        = 4.4[us]
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X : parts per Million : 13C

3.0

2.0

1.0

abundance

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115.8877

116.9274

117.6428

118.4536

121.0098

123.5661

130.5005

130.6054

134.1250

134.2109

135.5939

X : parts per Million : 13C



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Filename      = MS0299_FLUORINE-2.jdf
Author       = Jim Davis
Experiment    = single_pulse.ex2
Sample_id    = MS0299
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Changer_sample_id = 7
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-78.6295

X : parts per Million : 19F

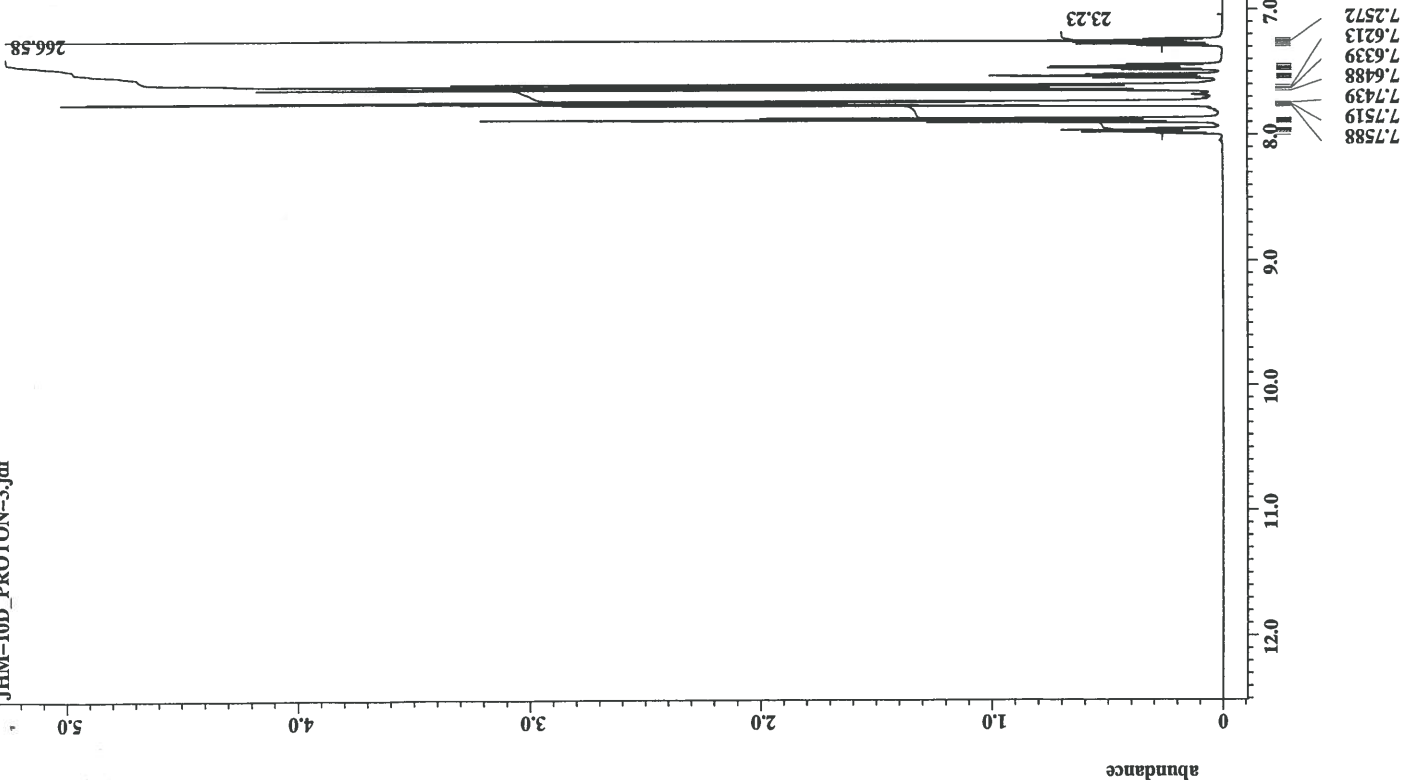
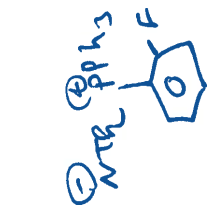
abundance

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Charger_sample = 7
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23.8193

X : parts per Million : 31P



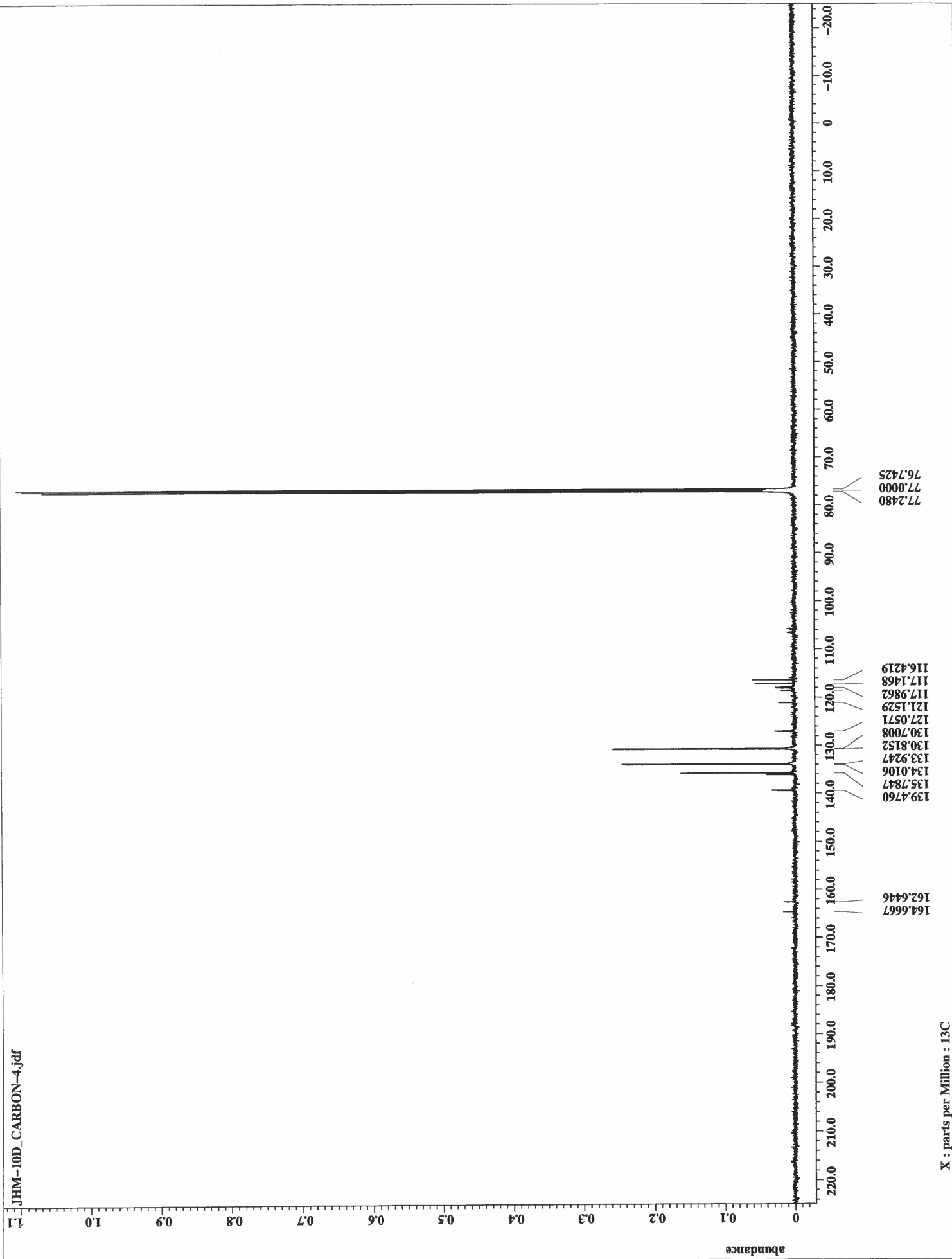
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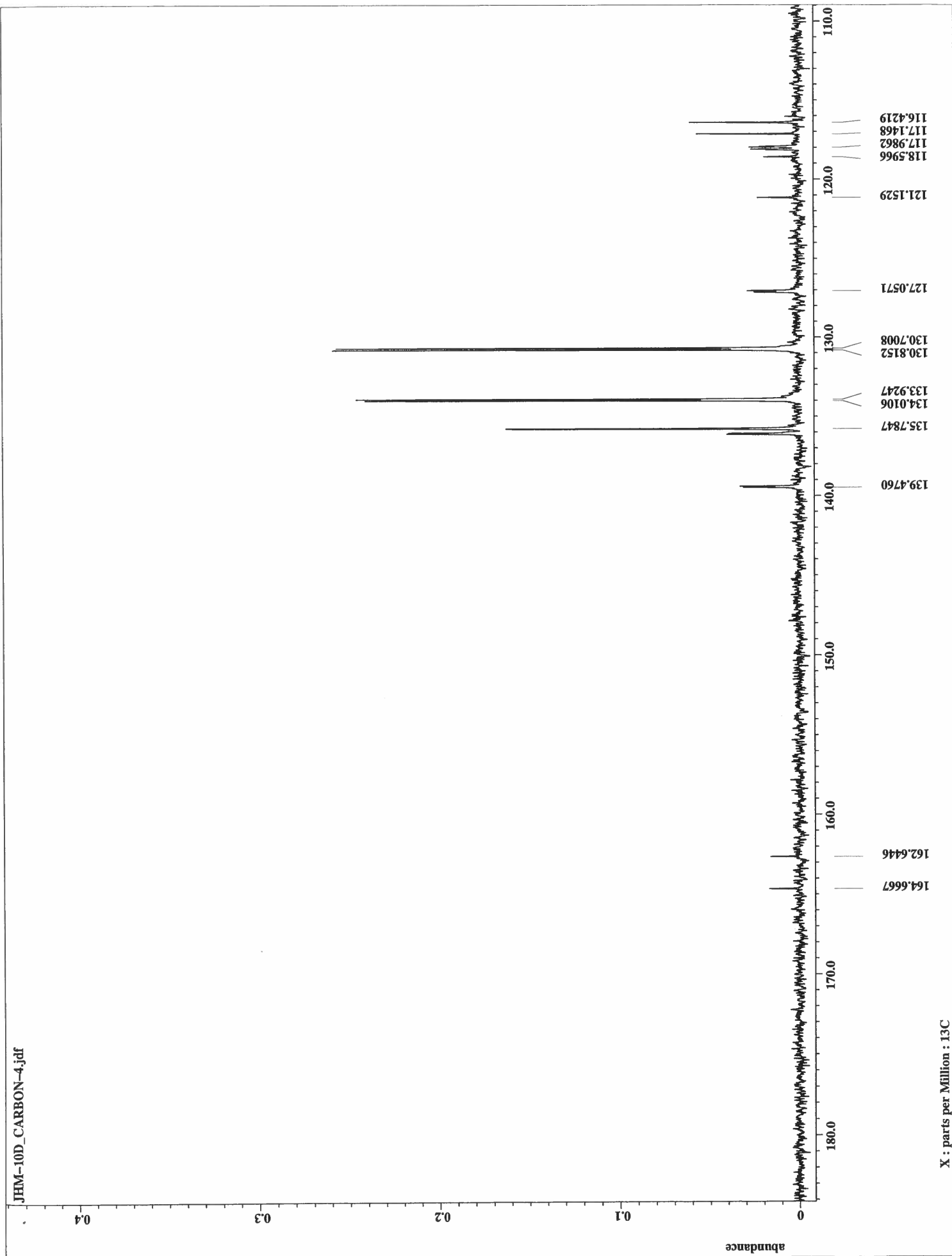
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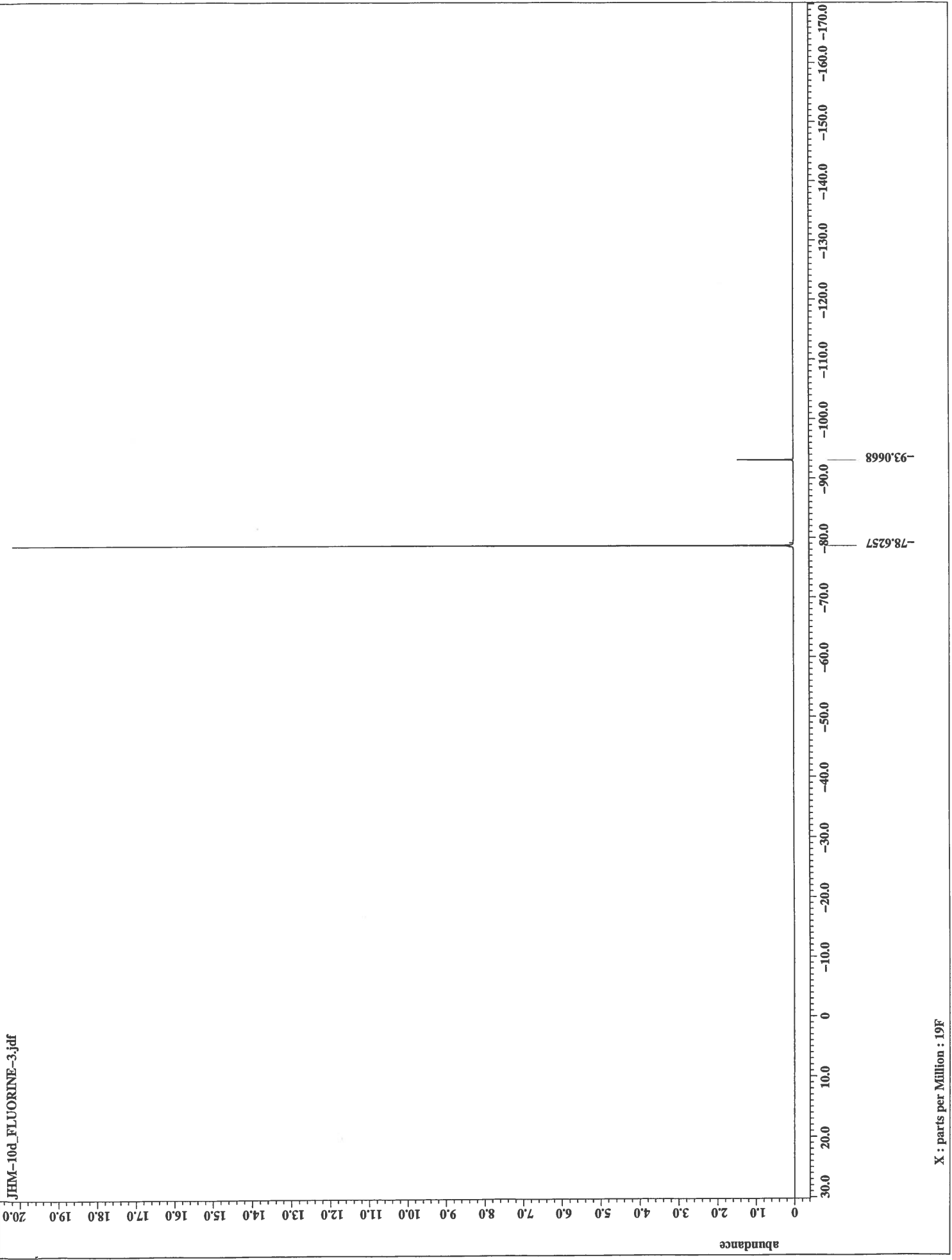
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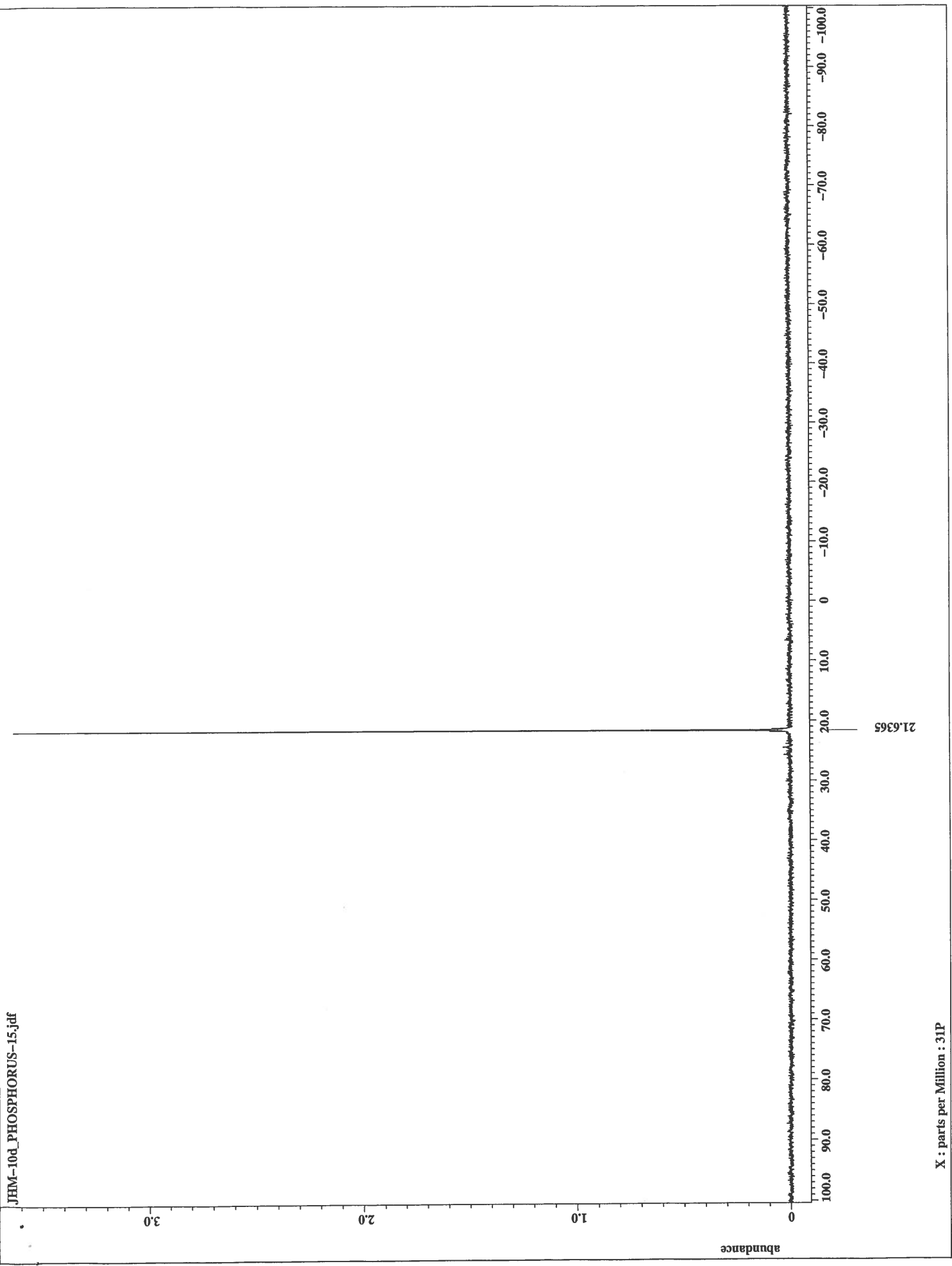
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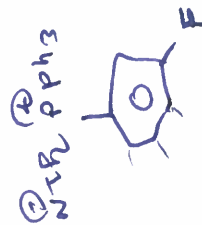
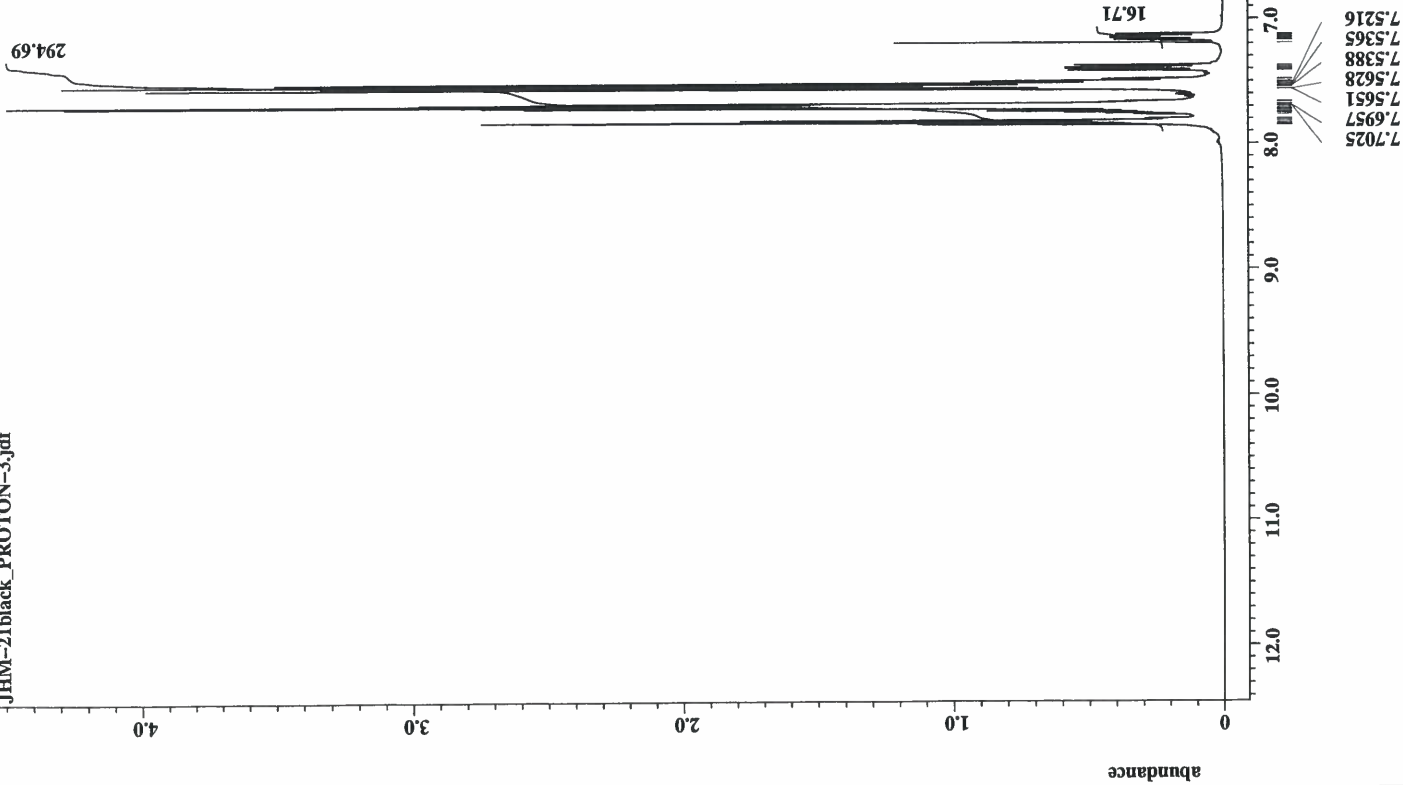
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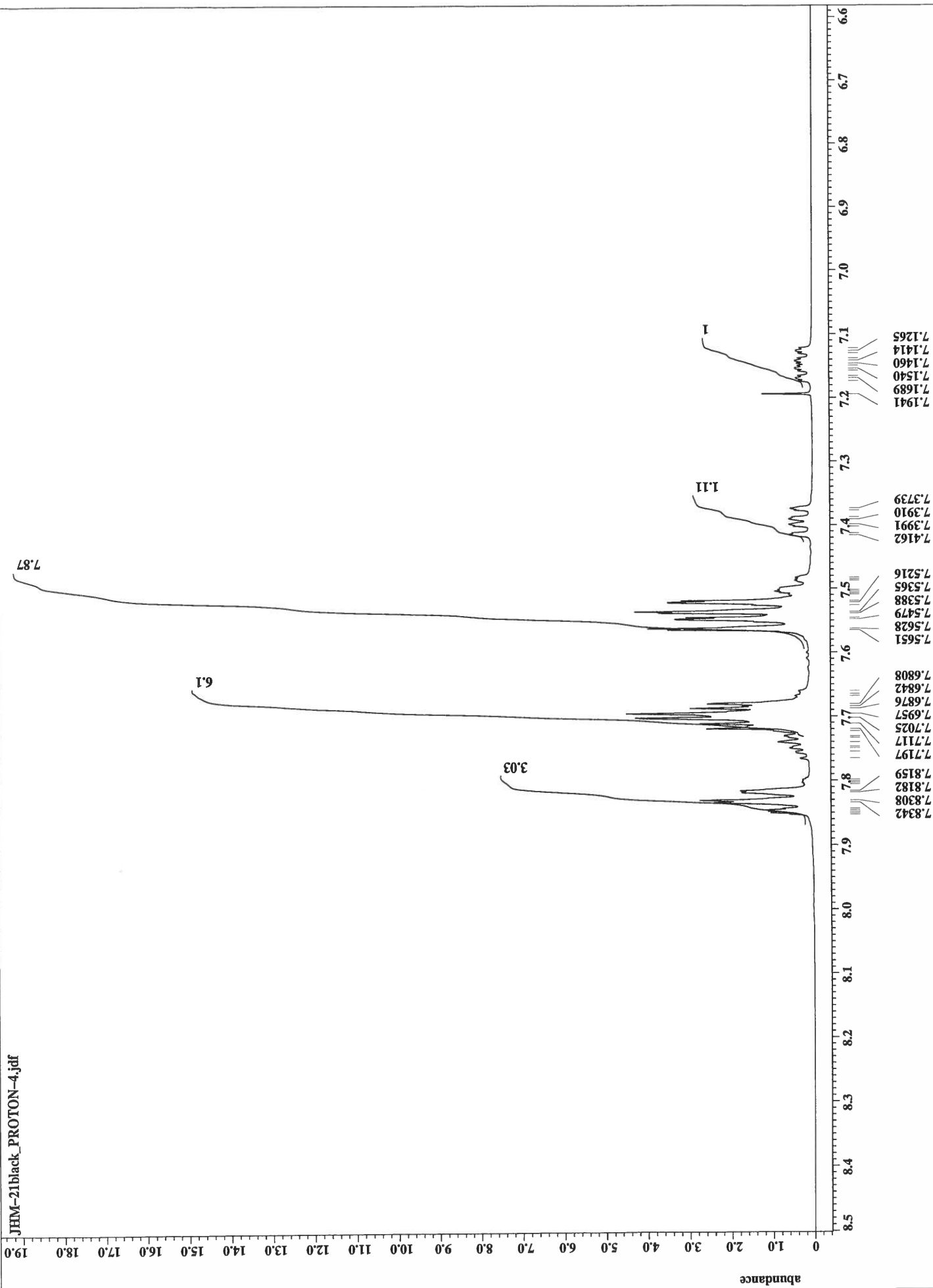


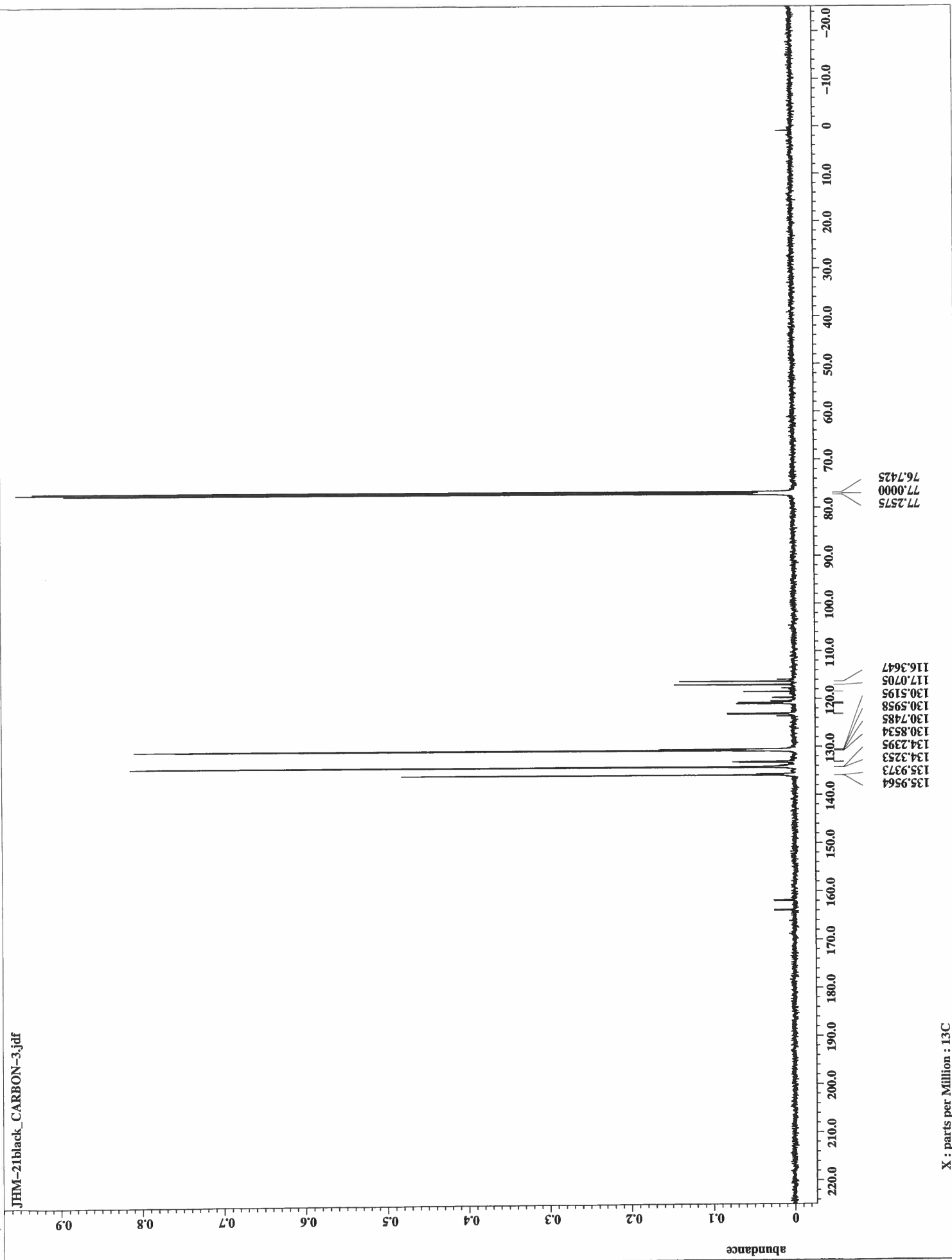


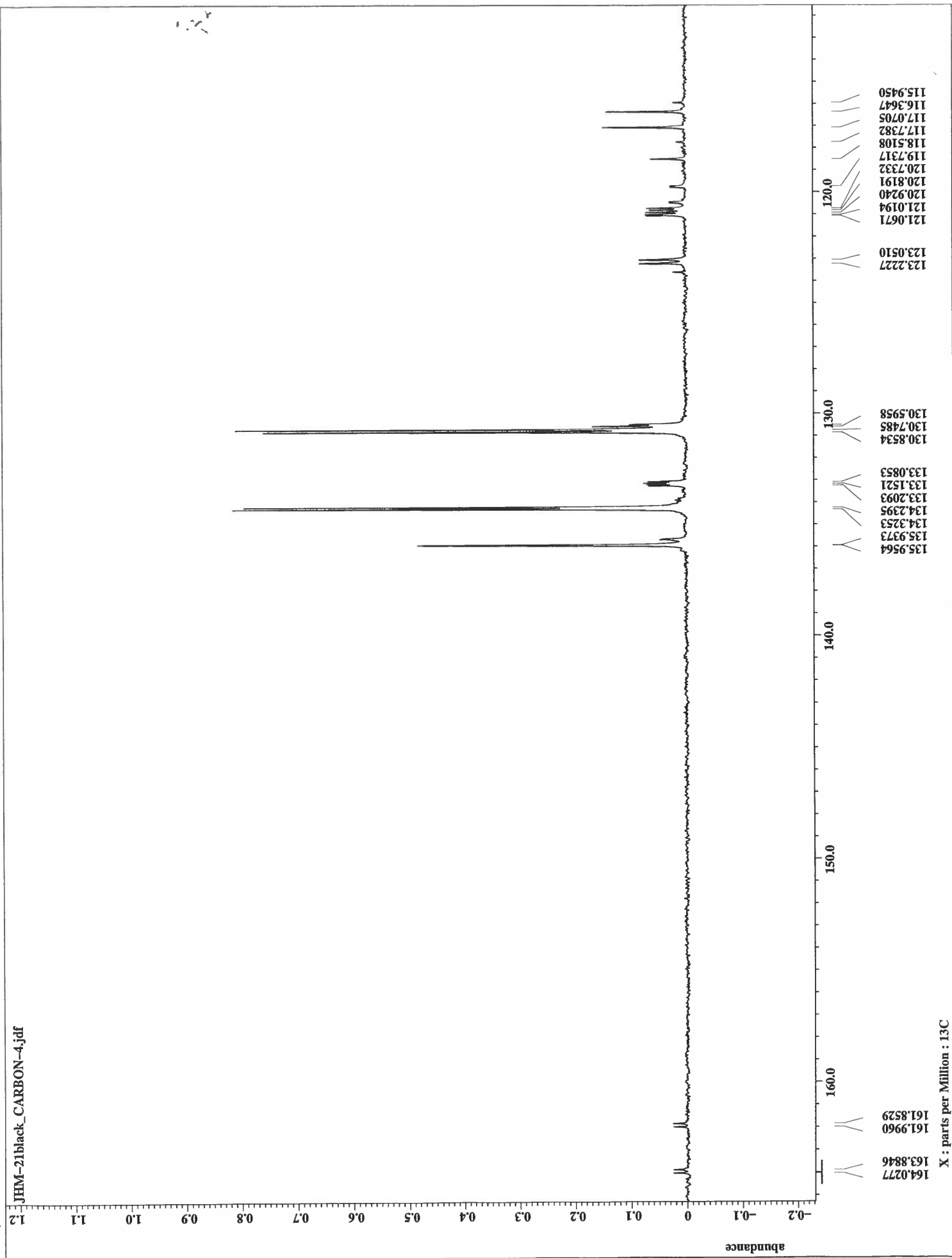


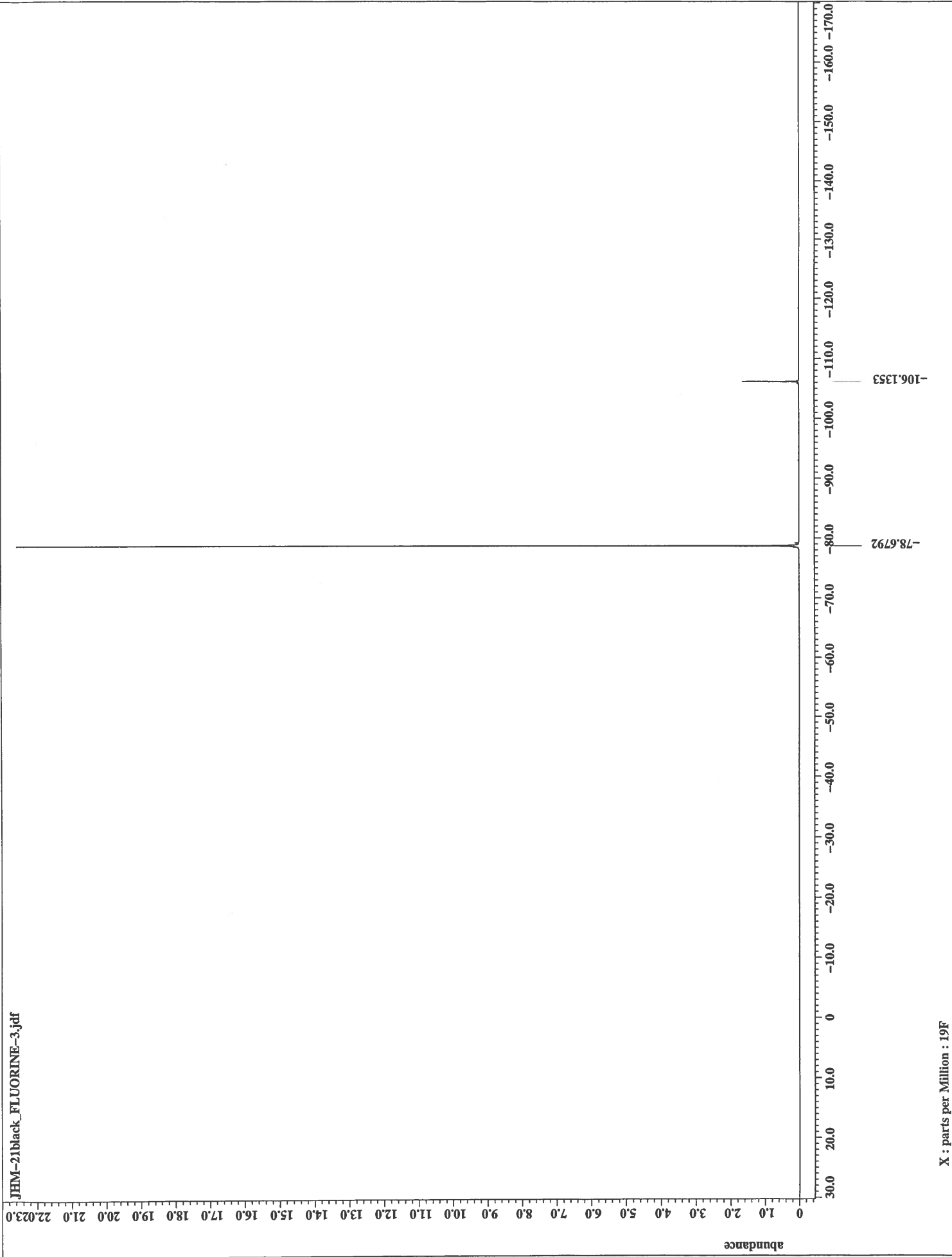


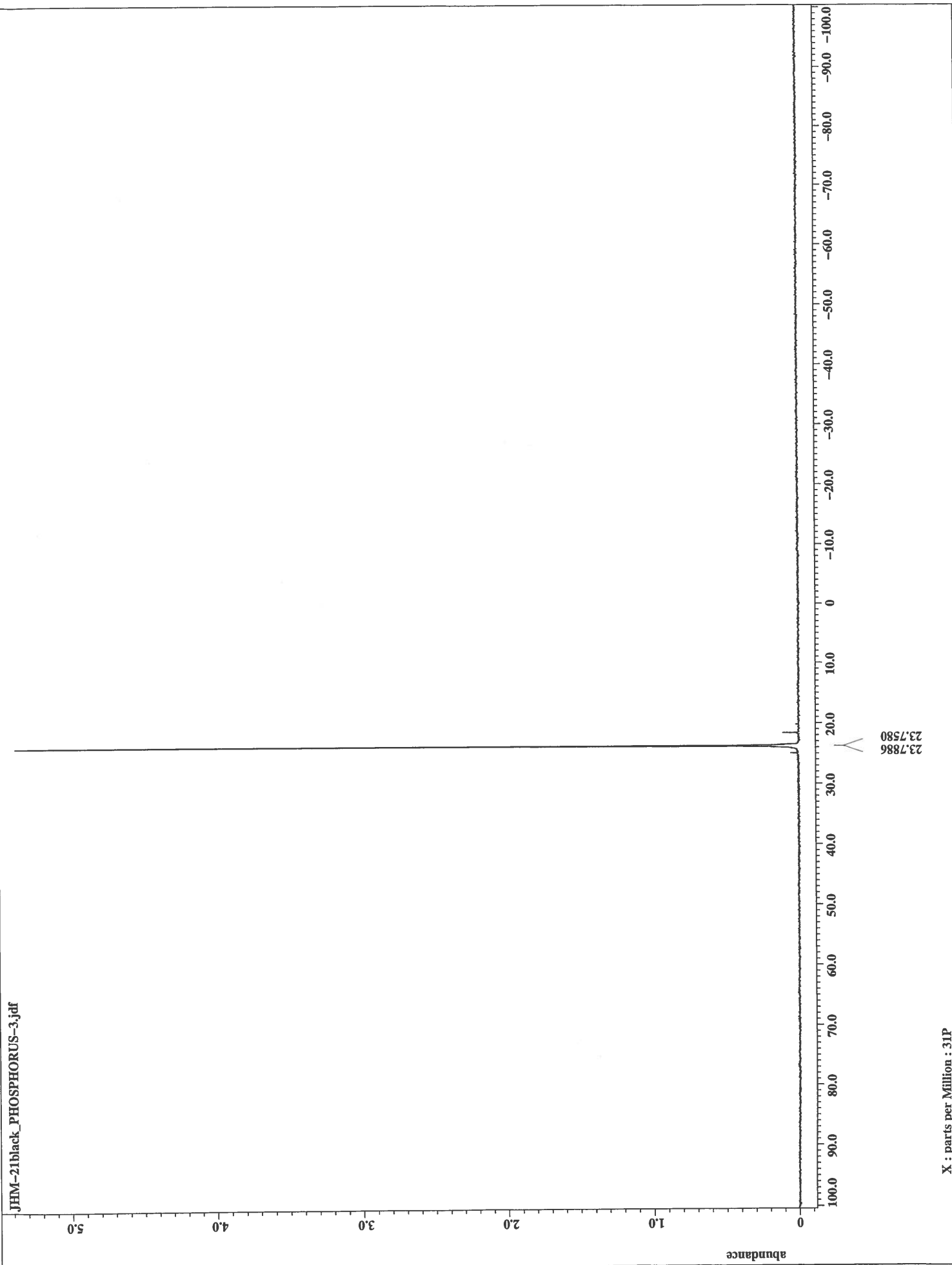




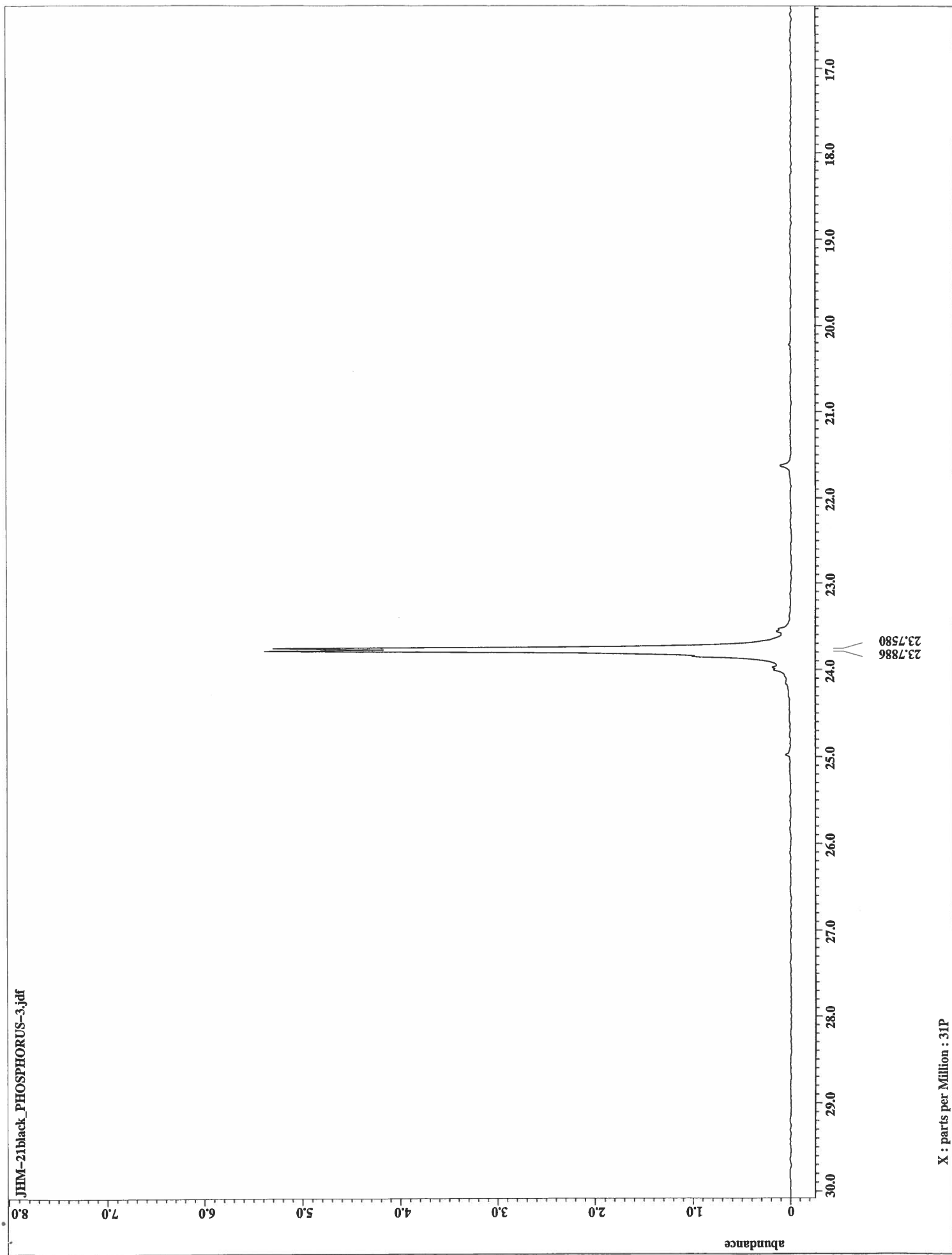


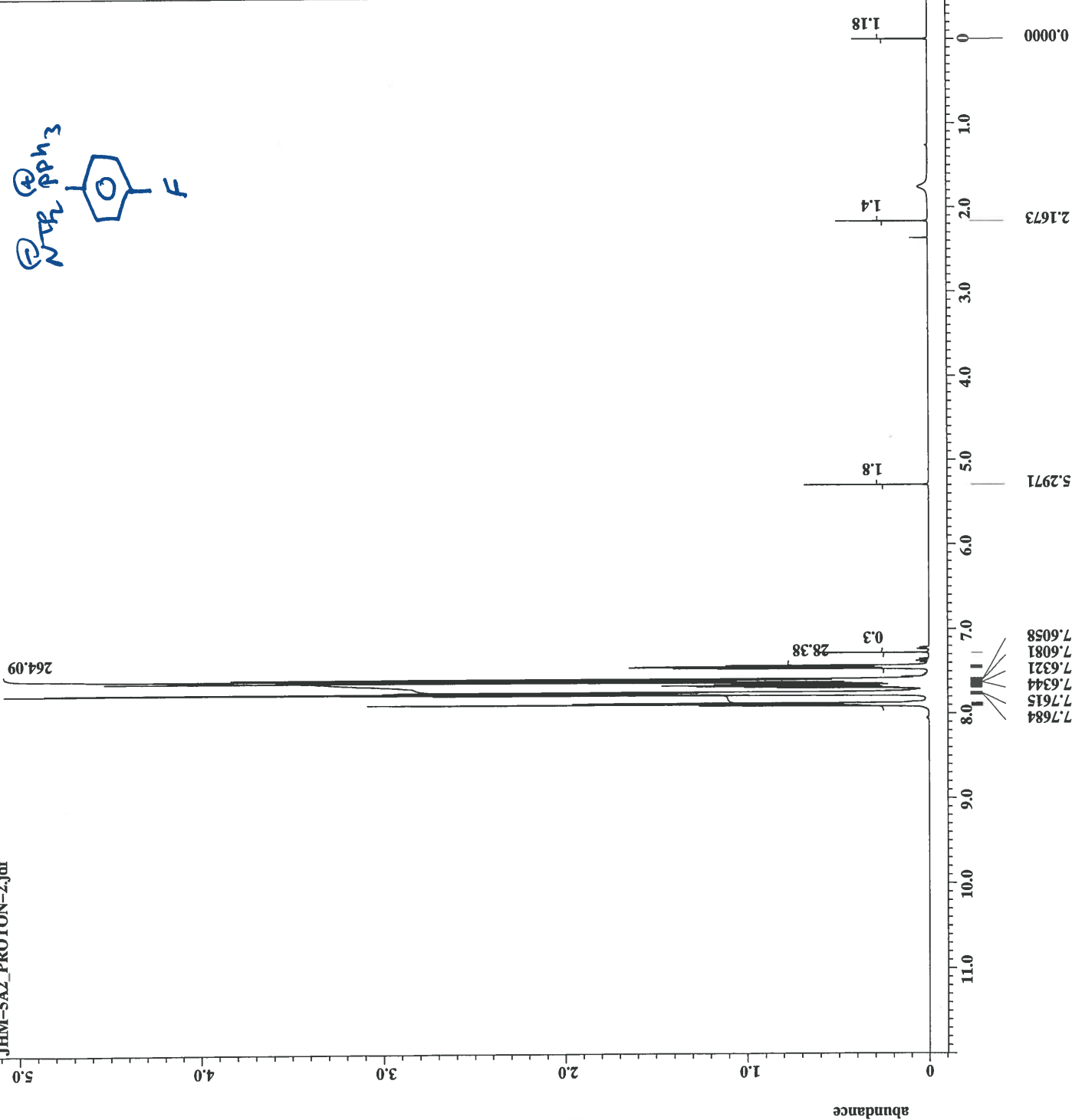
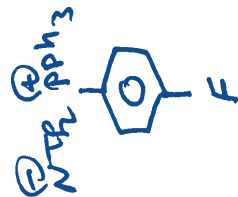






23.7886
23.7580

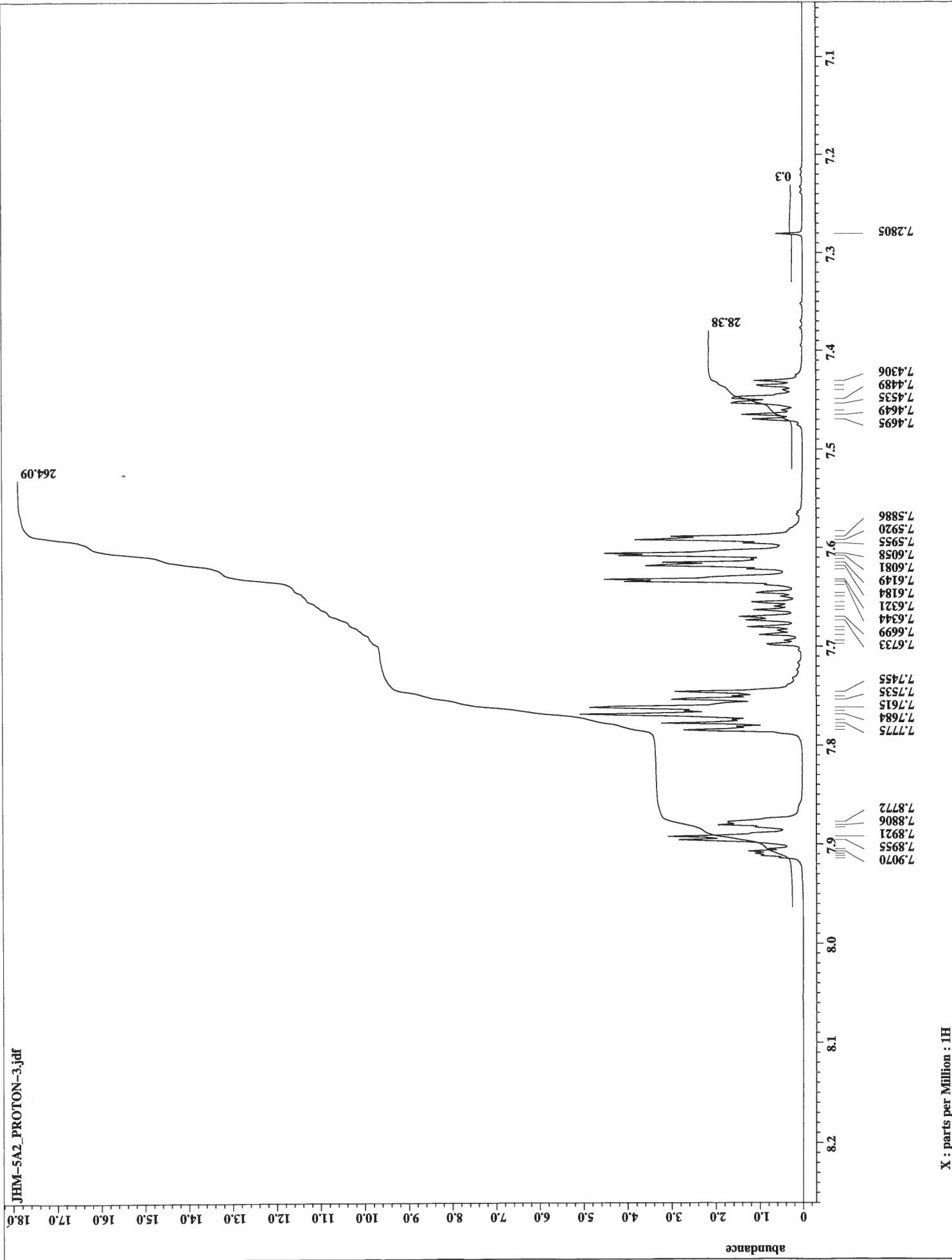




X : parts per Million : 1H



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 Irr_freq = 500.15991521[MHz]
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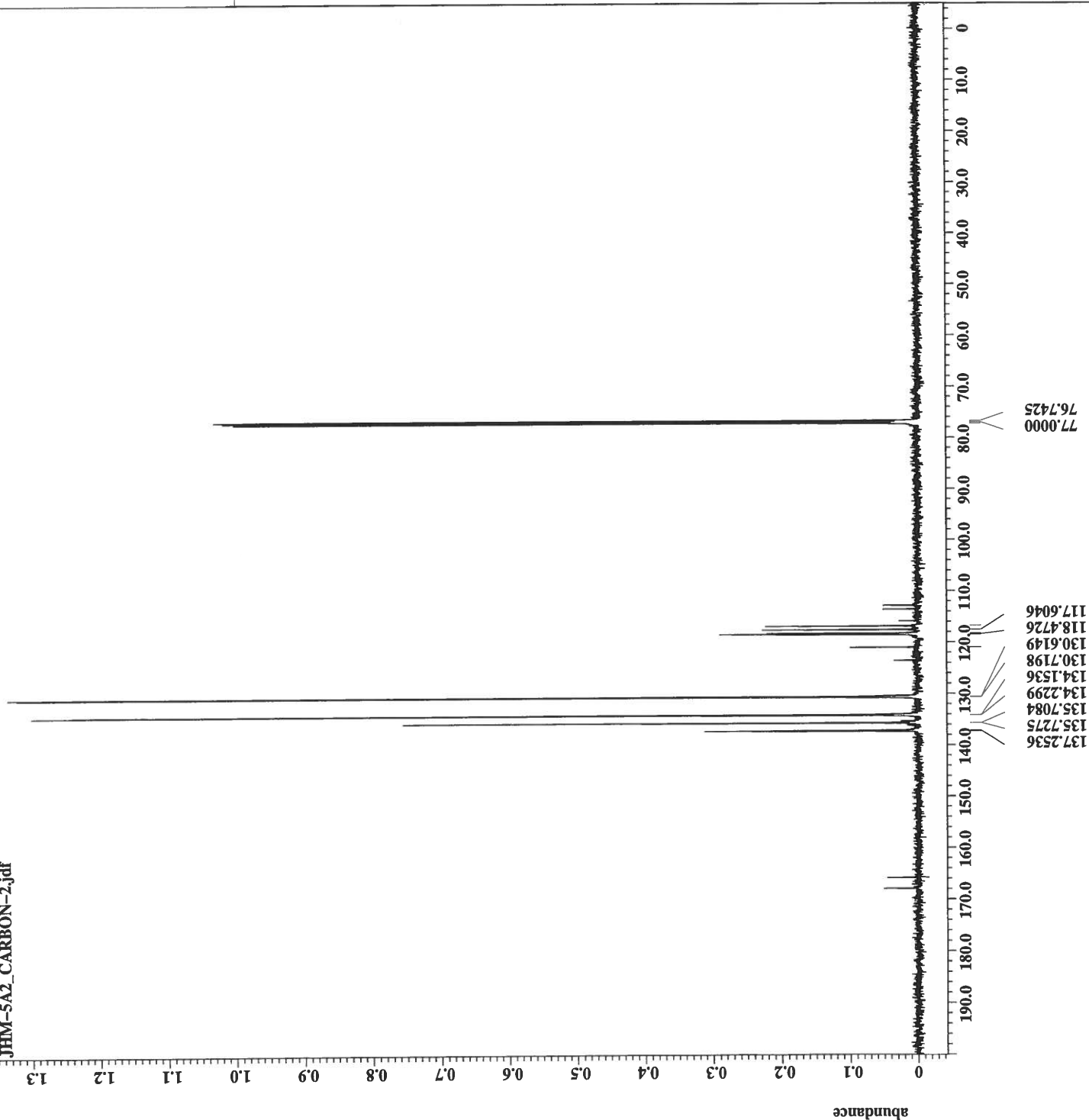
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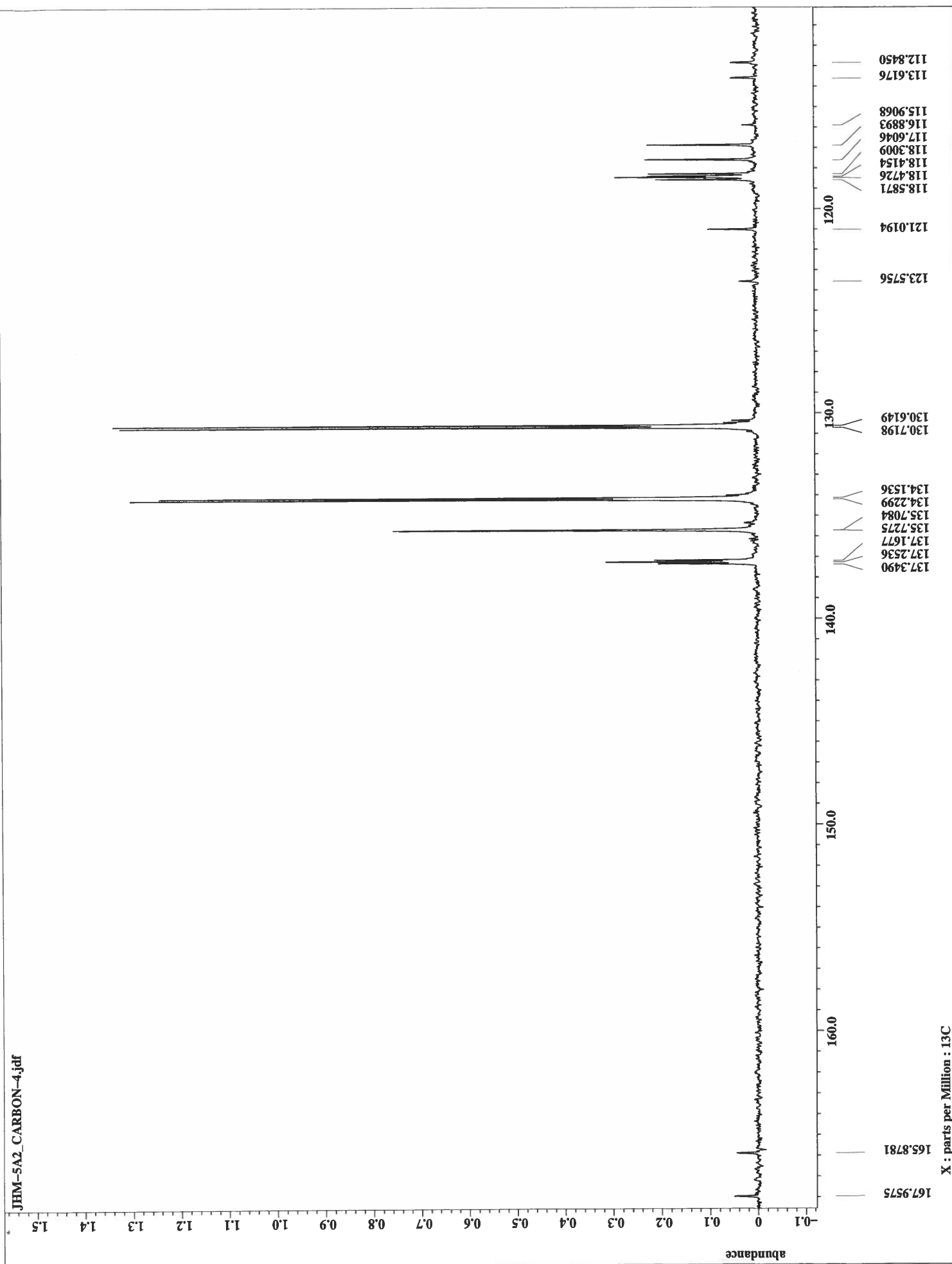
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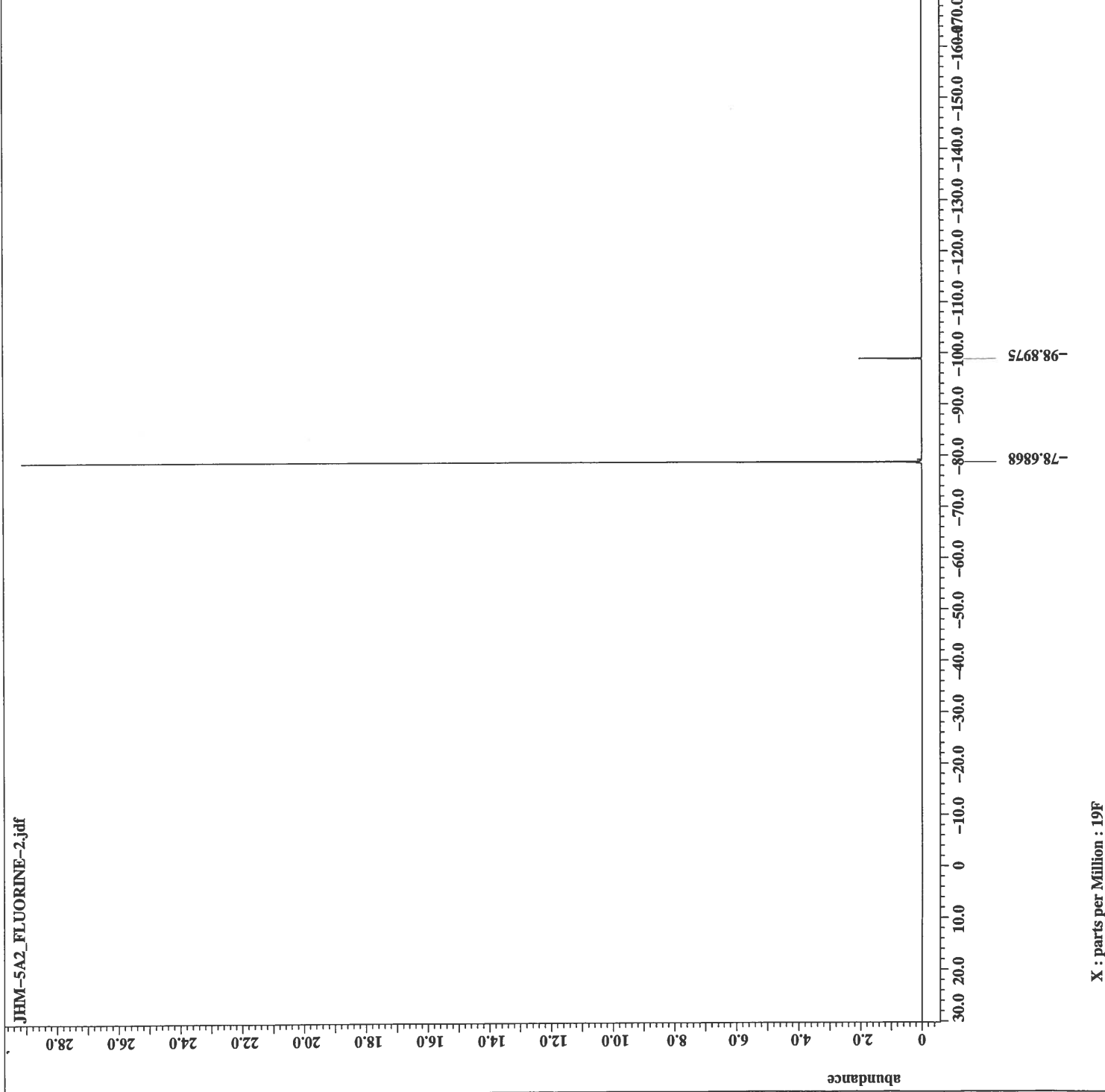
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X : parts per Million : 13C





X : parts per Million : 19F

**SOUTH ALABAMA**
JAGUARS

Filename = JHM-5A2_FLUORINE-2.jd
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 = JNM-ECA500
Field_strength = 11.7473579[T] (500[MH
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_pulse = Off
Irr_mode = Off
Dante_preset = FALSE
Initial_wait = 1[s]
Recvr_gain = 38
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get = 22.5[dc]

abundance

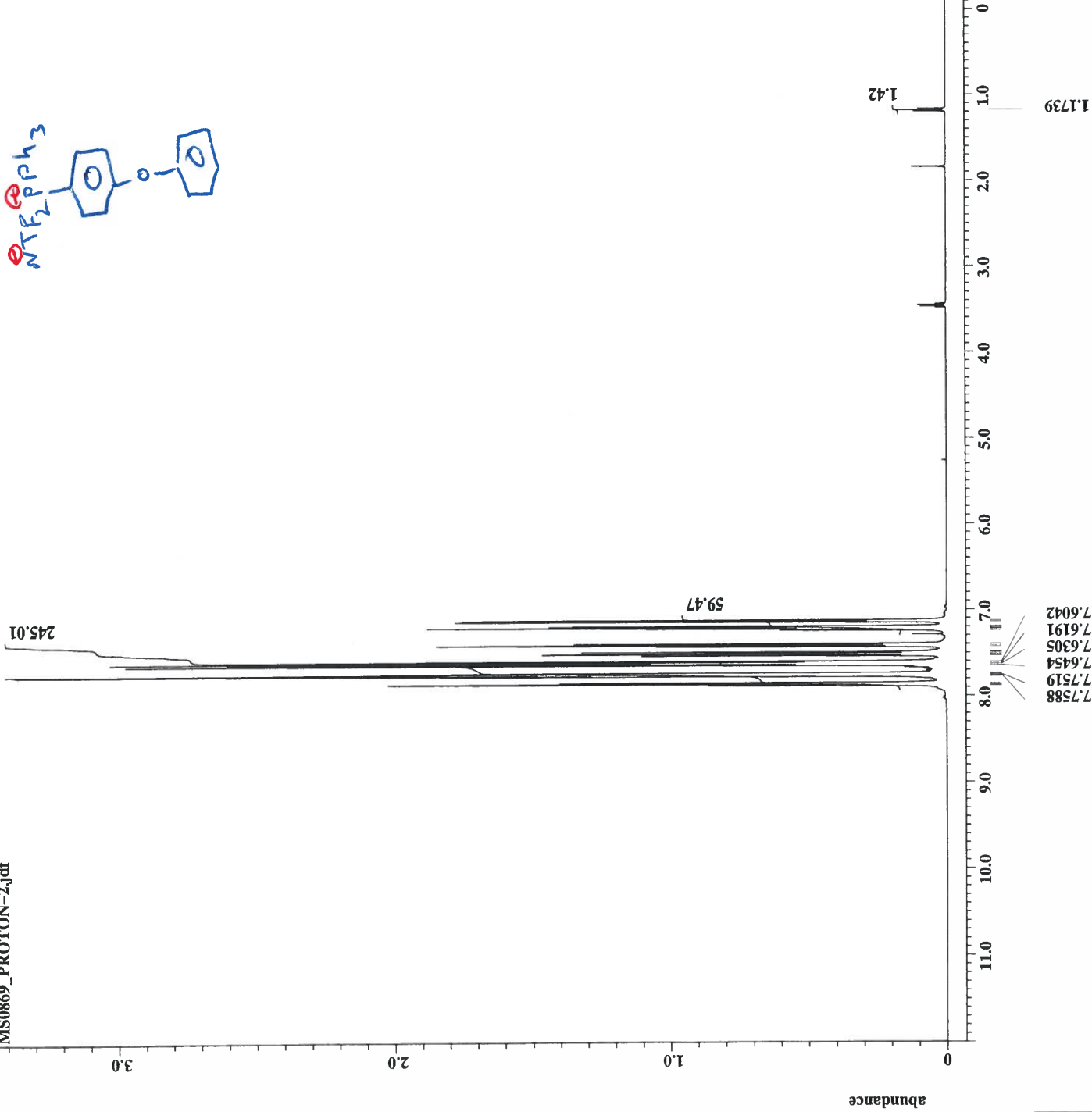
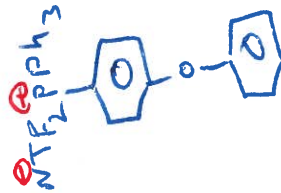


23.4899

X : parts per Million : 31P



Filename = JHM-5A2_PHOSPHORUS-2.
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = JHM-5A2
Solvent = CHLOROFORM-D
Changer_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 = 26214
Dim_title = 31P
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength = 11.7473579[T] (500[MH
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[MHz]
Irr_domain = 1H
Irr_freq = 500.15991521[MHz]
Irr_offset = 5.0[ppm]
Clipped = FALSE
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]
Recvr_gain = 56
Relaxation_delay = 2[s]
Repetition_time = 2.64487424[s]
Temp_set = 22.8[dc]



X : parts per Million : 1H



SOUTH ALABAMA
JAGUARS

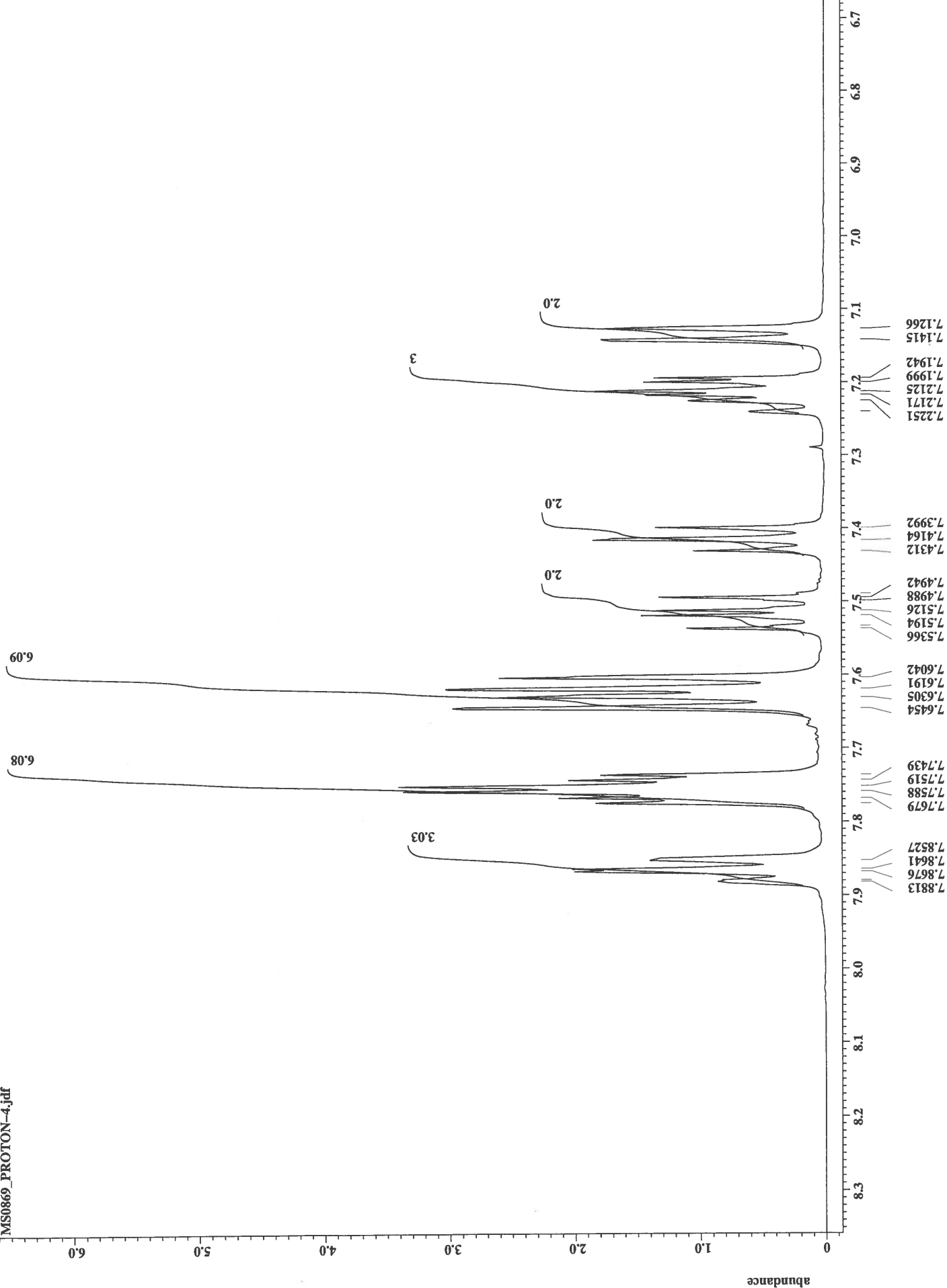
```

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[MH
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       = 9.38438438[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_atn         = 4[db]
X_pulse       = 5.65[us]
Irr_mode      = Off
Tri_mode      = Off
Dante_preset  = FALSE
Initial_wait  = 1[s]
Recvr_gain    = 20
Relaxation_delay = 4[s]
Repetition_time = 5.74587904[s]
Temp_get      = 19.3[dc]
  
```





SOUTH ALABAMA
JAGUARS™

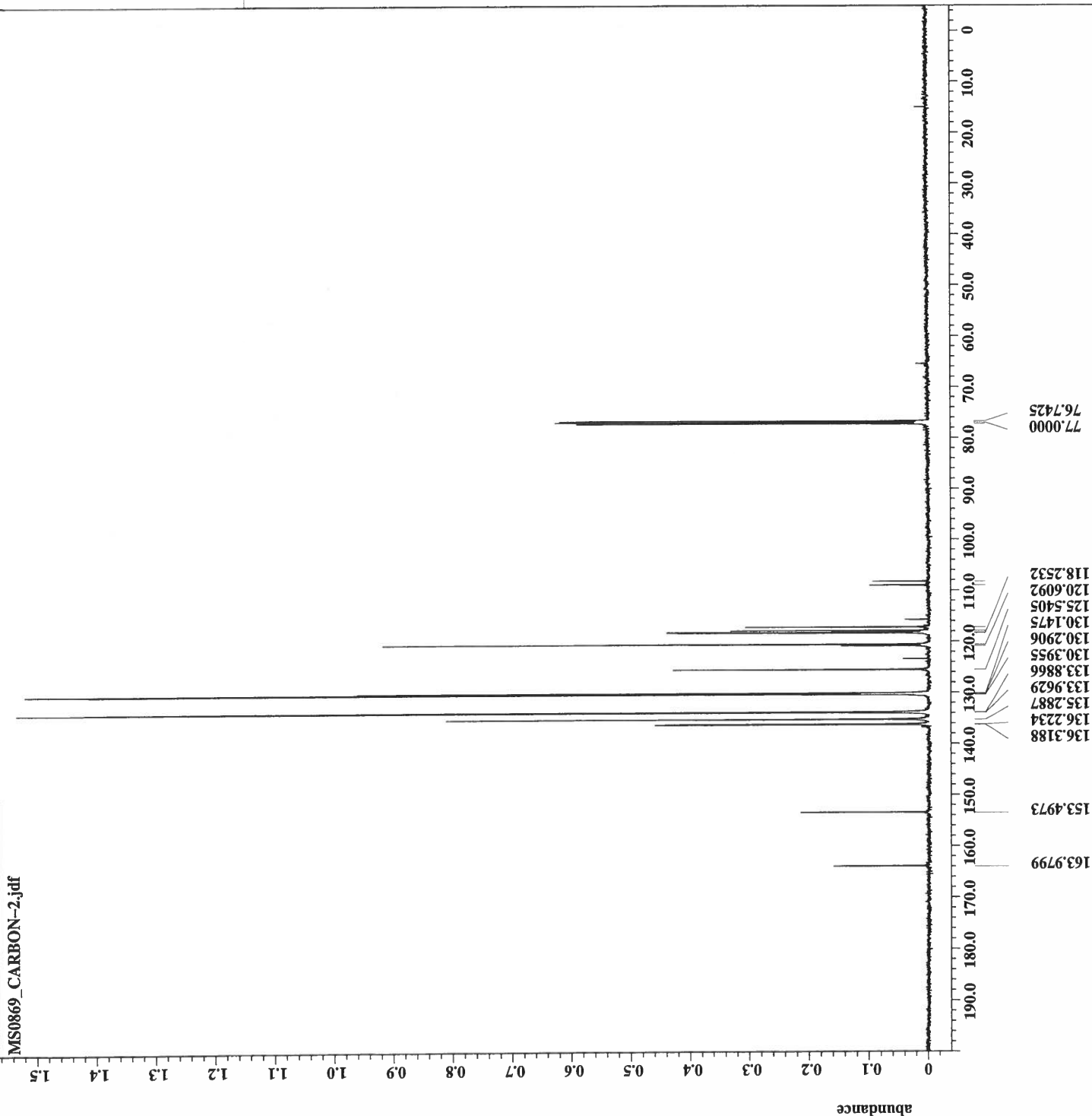
```

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

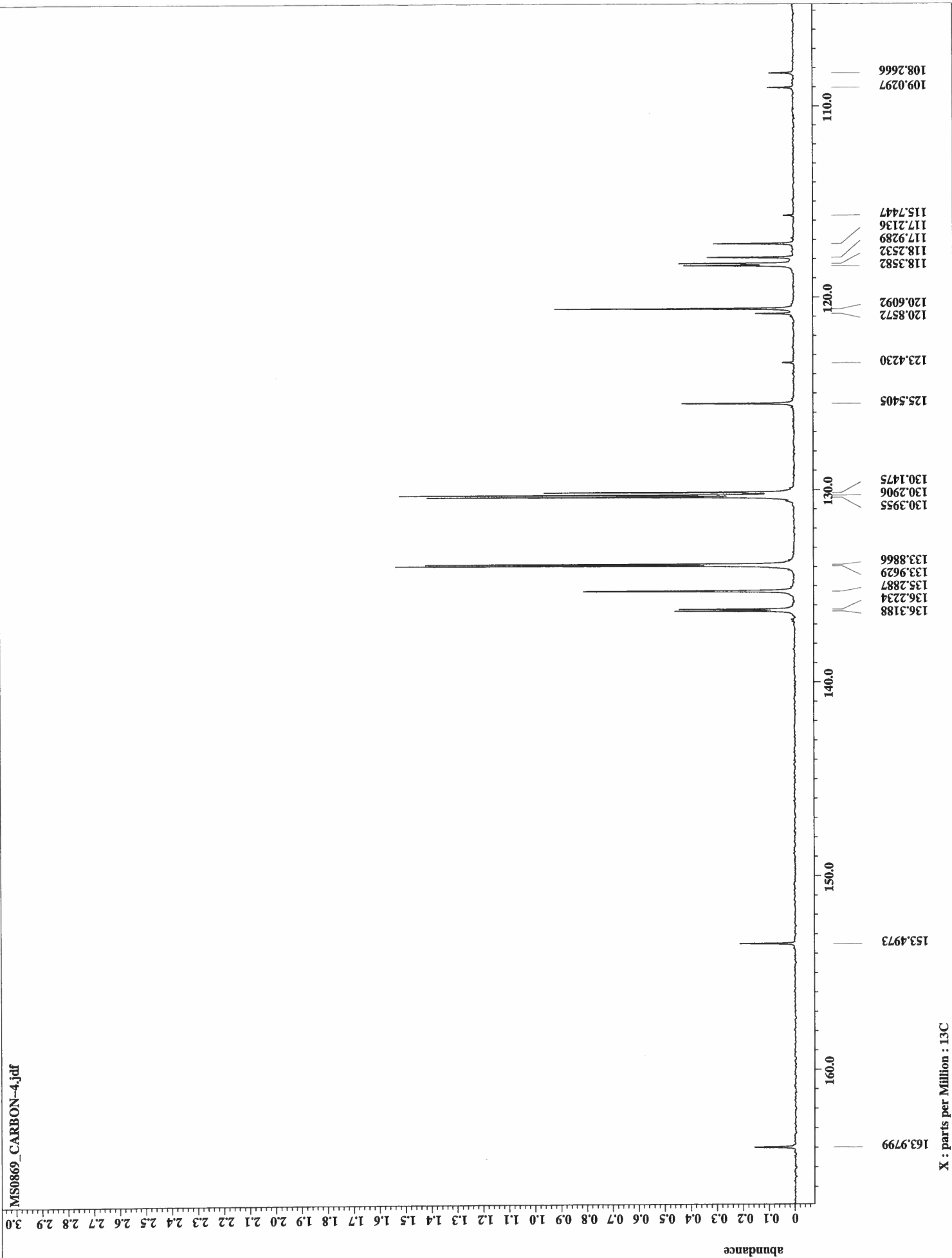
Data_format = 1D COMPLEX
Dim_size    = 26214
Dim_title   = 13C
Dim_units   = [ppm]
Dimensions  = X
Site        = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579[T] (500[MH
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          = 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        = 4.333333333[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]
Recvr_gain     = 60
Relaxation_delay = 2[s]
Repetition_time = 2.83361792[s]
Temp_get       = 18.8[dc]
  
```



X : parts per Million : 13C



30.0

20.0

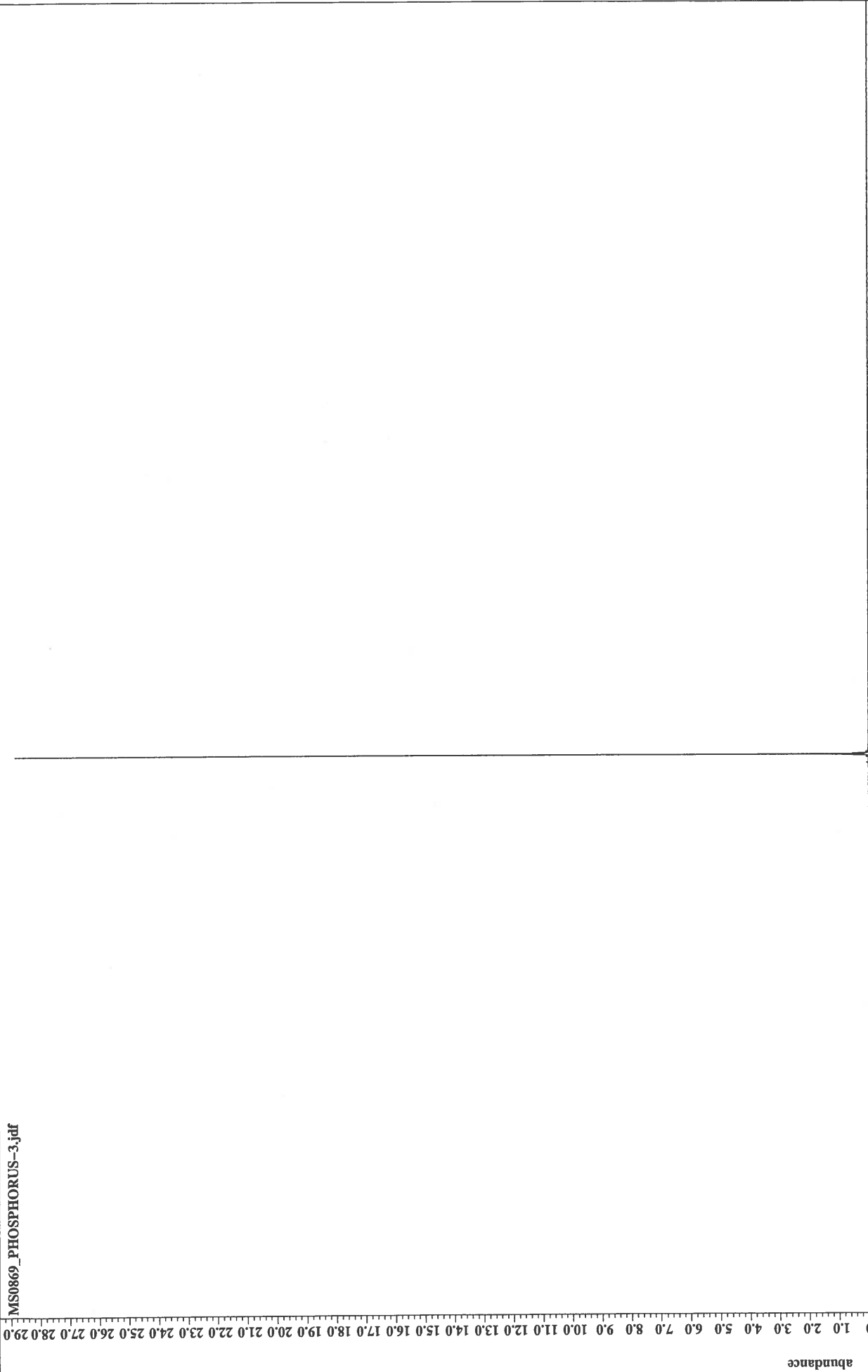
10.0

0

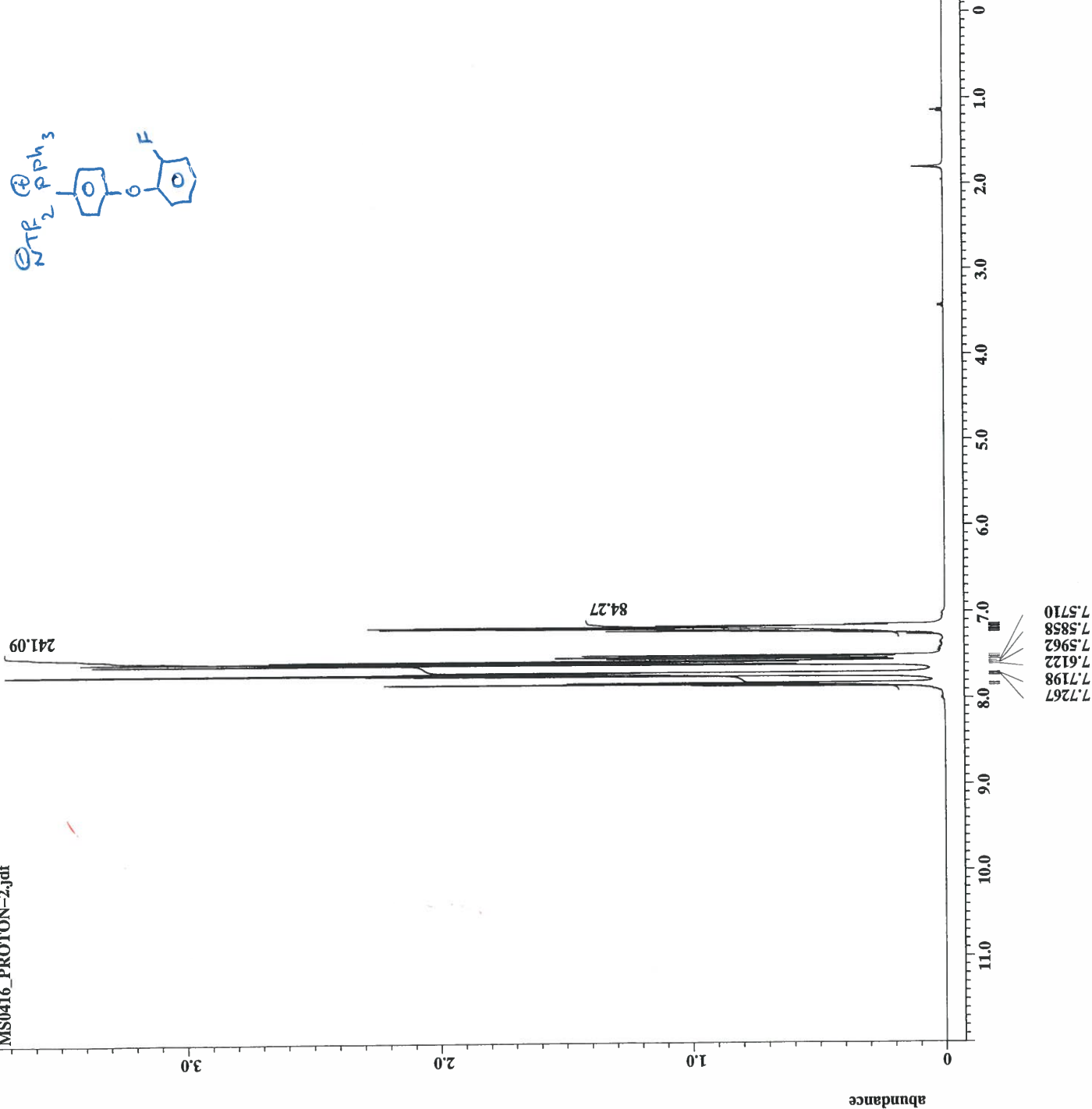
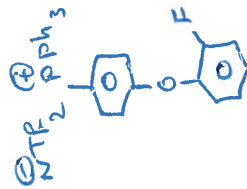
abundance

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

-78.8181



23.1204



X : parts per Million : 1H



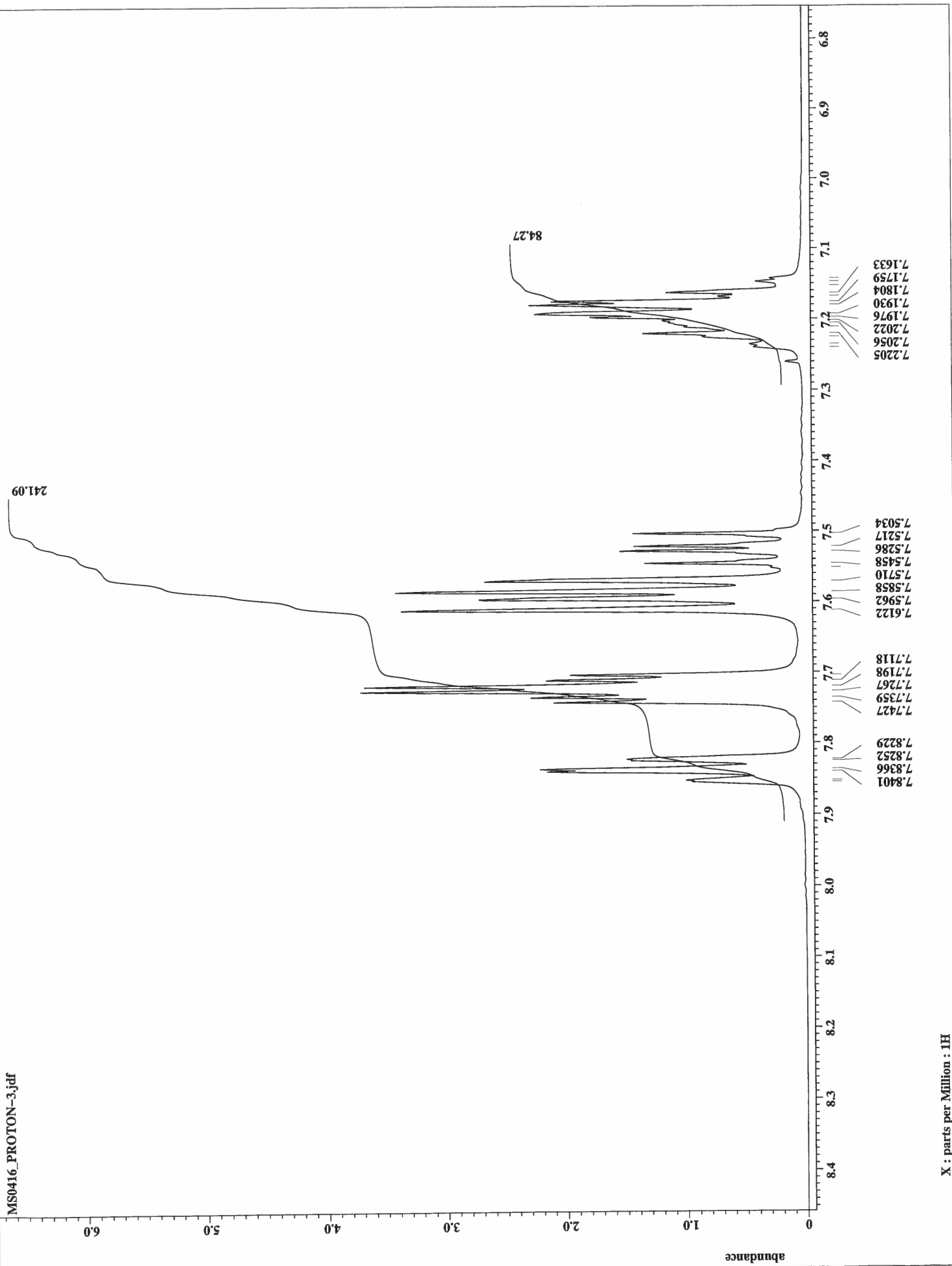
```

Filename      = MS0416_PROTON-2.jdf
Author       = Jim Davis
Experiment   = single_pulse.ex2
Sample_id    = MS0416
Solvent      = CHLOROFORM-D
Changer_sample
Creation_time = 14-APR-2018 11:32:37
Revision_time = 14-APR-2018 11:10:44
Current_time  = 14-APR-2018 11:10:44

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[MH
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]
Recvr_gain      = 24
Relaxation_delay = 4[s]
Repetition_time = 5.74587904[s]
Temp_get        = 22.8[dc]
  
```





SOUTH ALABAMA
JAGUARS

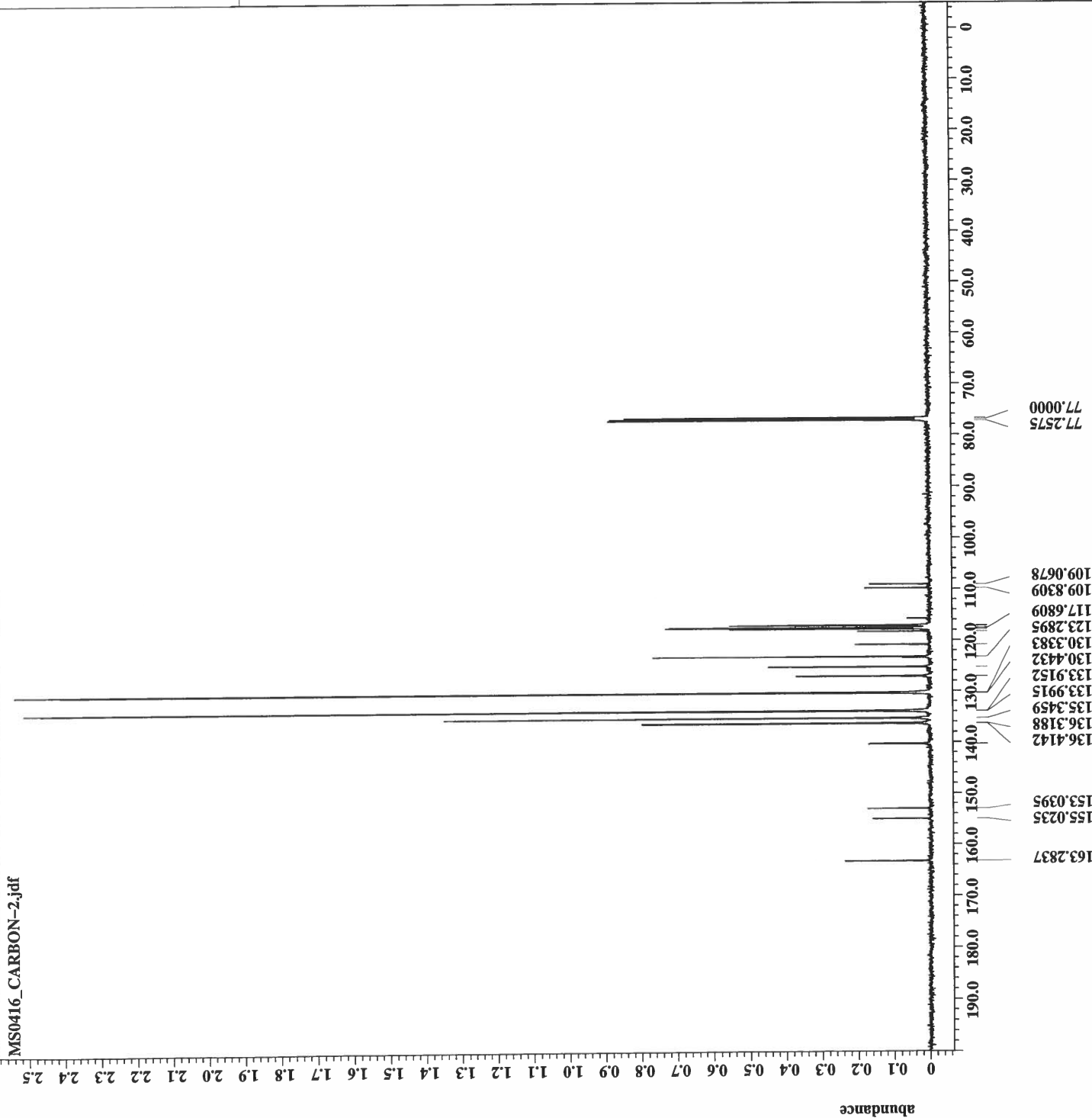
```

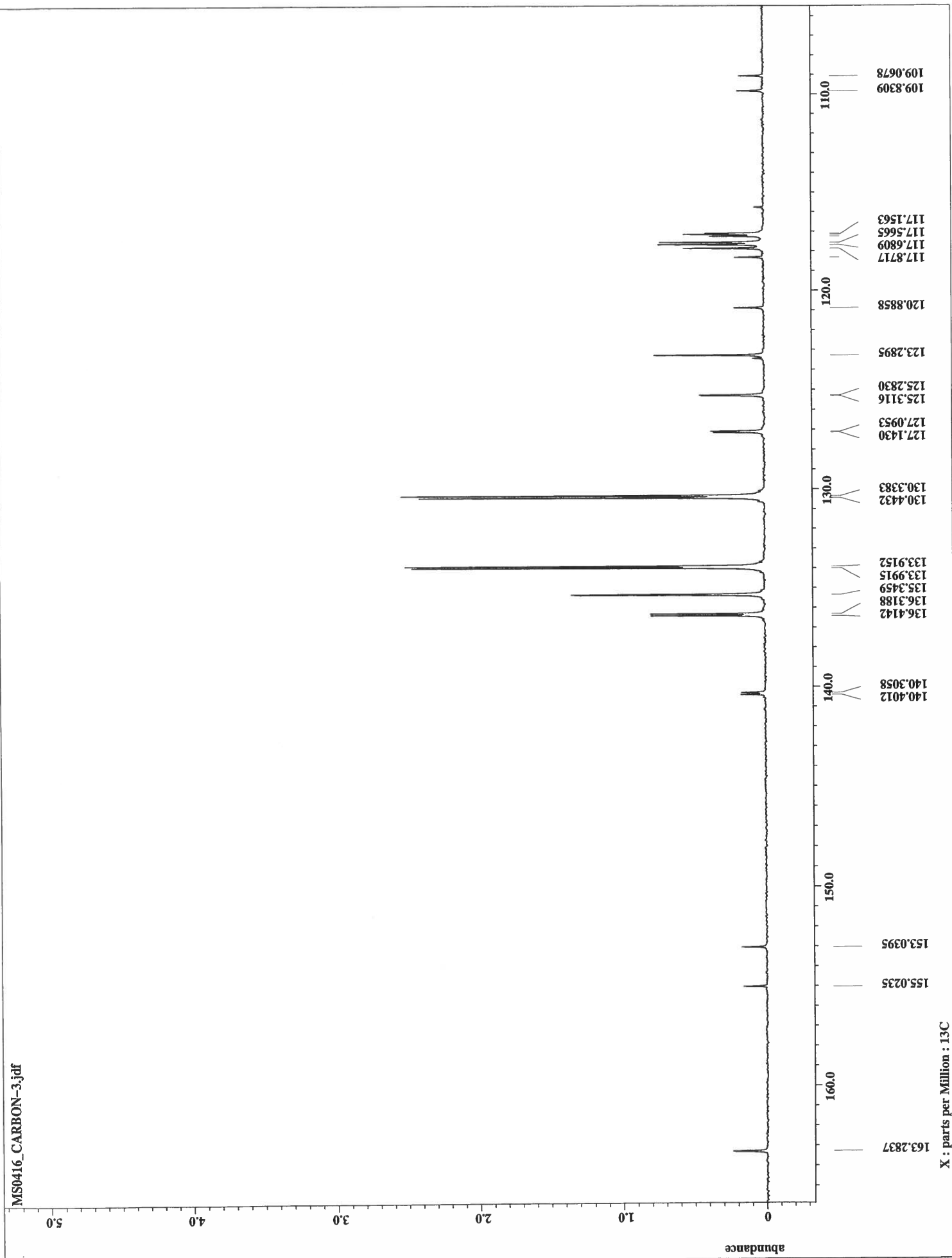
Filename      = MS0416_CARBON-2.jdf
Author       = Jim Davis
Experiment   = single_pulse_dec
Sample_id    = MS0416
Solvent      = CHLOROFORM-D
Changer_sample
Creation_time = 14-APR-2018 11:47:07
Revision_time = 14-APR-2018 11:25:15
Current_time  = 14-APR-2018 11:25:15

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[T] (500[MH
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

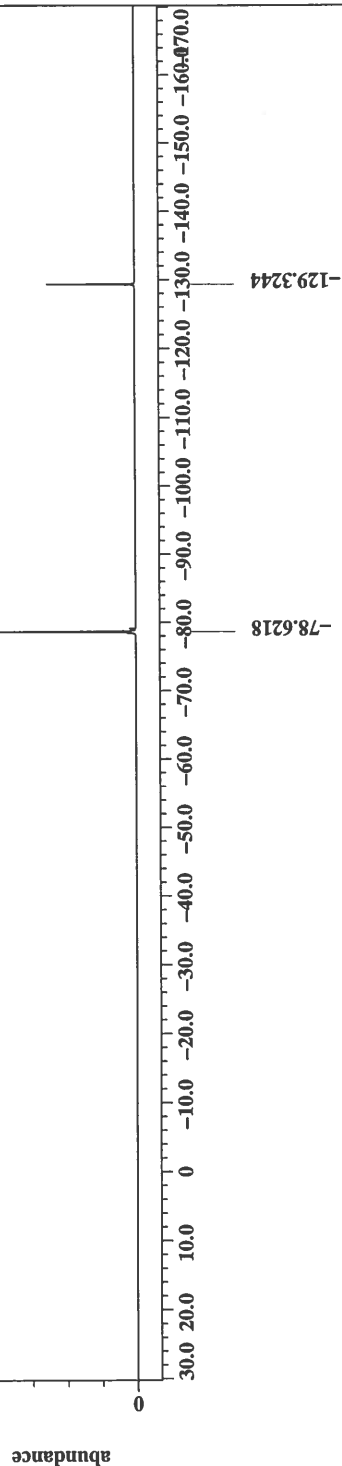
X_90_width     = 13.2[us]
X_acq_time     = 0.83361792[s]
X_angle        = 30[deg]
X_atn          = 6[dB]
X_pulse        = 4.4[us]
Irr_atn_dec    = 20.7[dB]
Irr_atn_noe    = 20.7[dB]
Irr_noise      = WALTZ
Decoupling     = TRUE
Initial_wait   = 1[s]
Noe            = TRUE
Noe_time       = 2[s]
Recvr_gain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

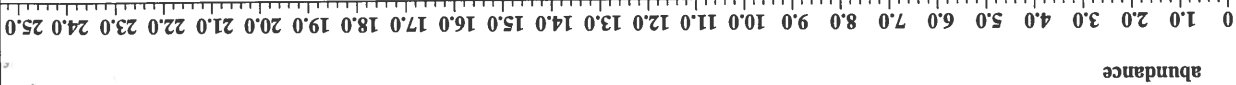




**SOUTH ALABAMA**
JAGUARS

Filename = MS0416_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0416
Solvent = CHLOROFORM-D
Charger_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[MH
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_pulse = Off
Irr_mode = Off
Tri_mode = Off
Dante_preset = FALSE
Initial_wait = 1[s]
Recvr_gain = 34
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get = 22.6[degC]



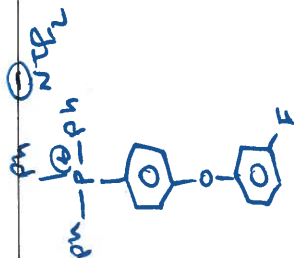


23.1989

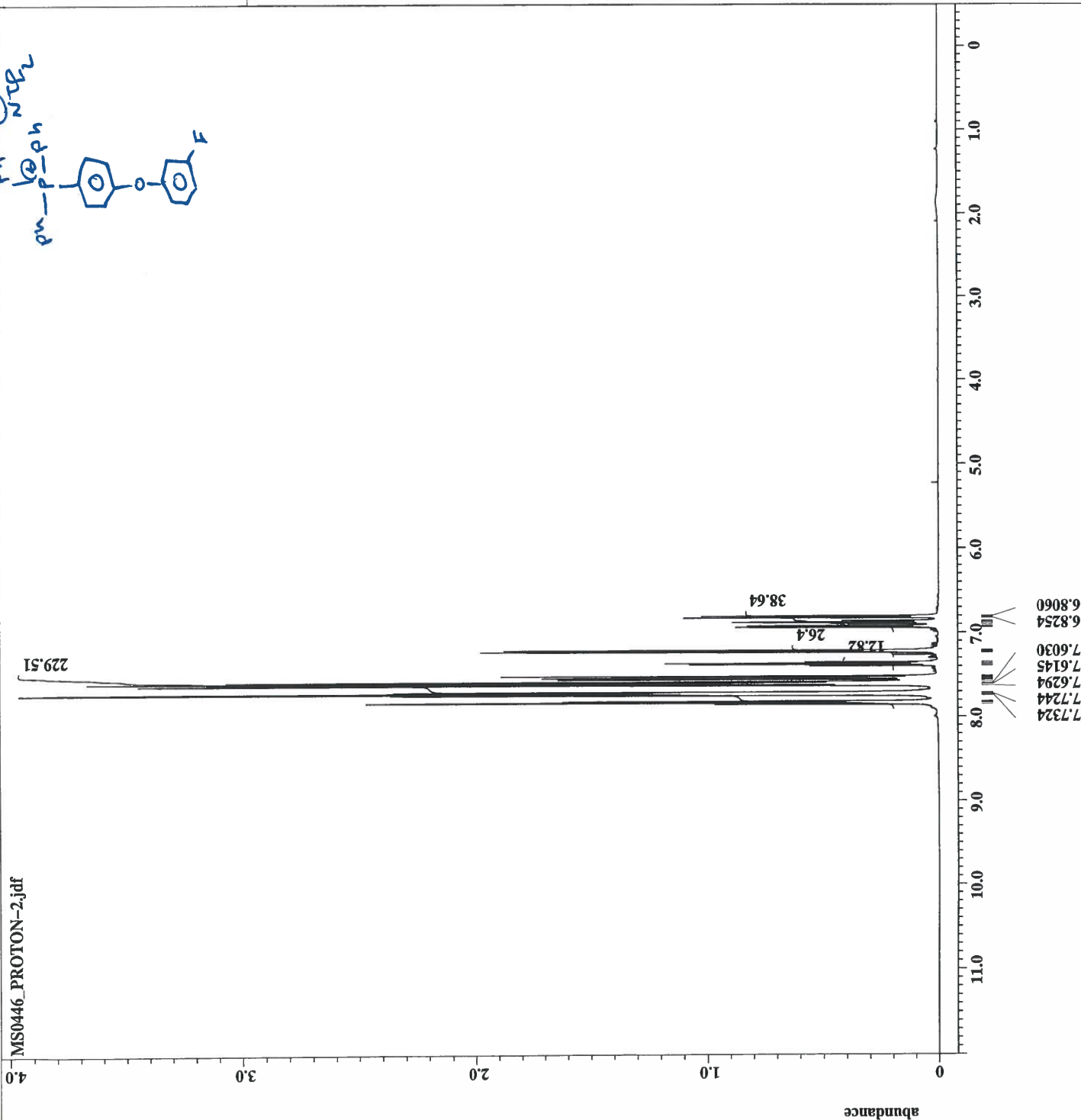
X : parts per Million : 31P



MS0416_PHOSPHORUS-2.j
= Jim Davis
= single_pulse_dec
= MS0416
= CHLOROFORM-D
= 10
= 14-APR-2018 11:29:07
= 14-APR-2018 11:07:15
= 14-APR-2018 11:07:15
= 1D COMPLEX
= 26214
= 31P
= [ppm]
= X
= ECA 500
= JNM-ECA500
Field_strength = 11.7473579[T] (500[MH
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]
Relaxr_gain = 54
Relaxation_delay = 2[s]
Repetition_time = 2.64487424[s]
Temp_get = 23[dc]



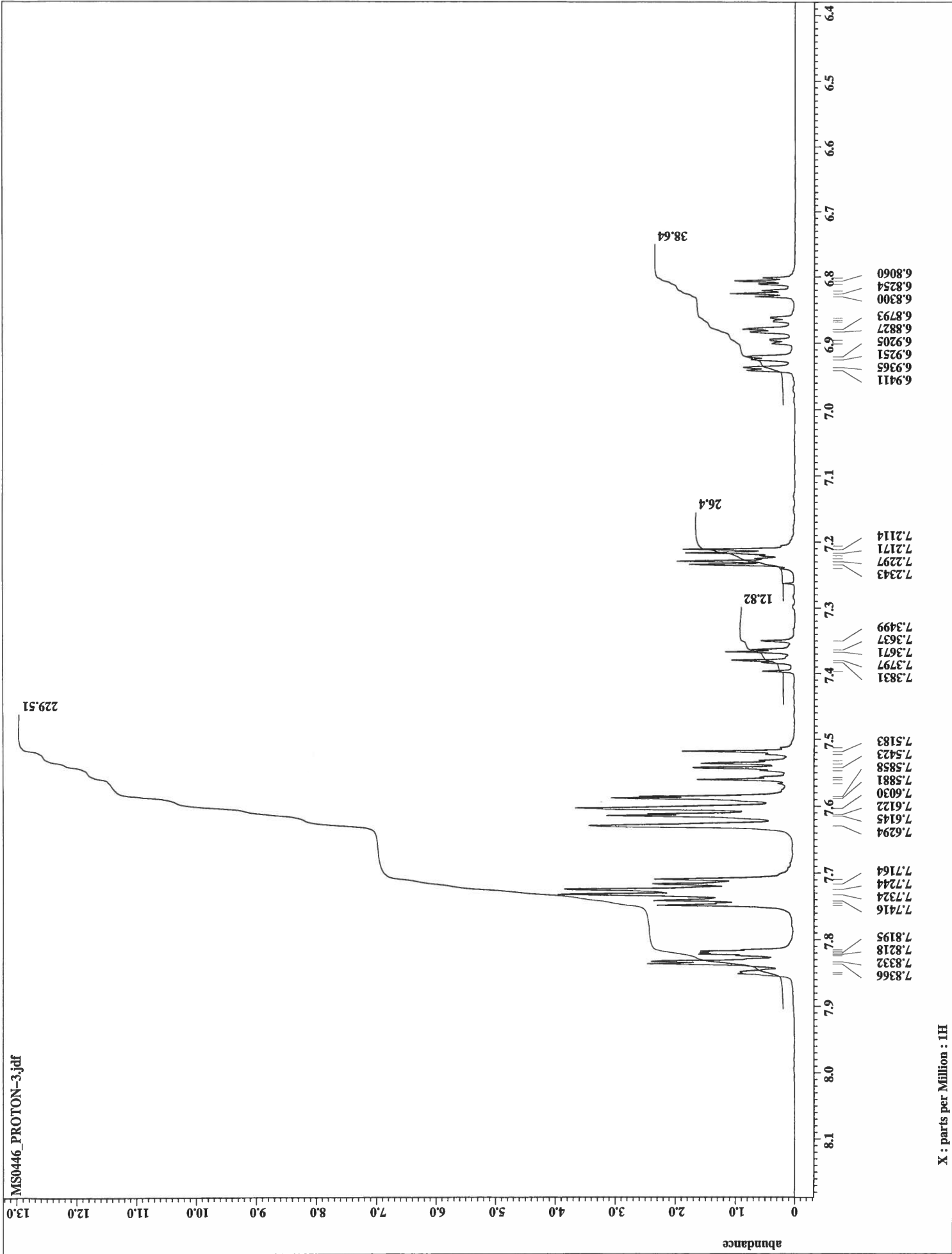
MS0446_PROTON-2.jdf



X : parts per Million : 1H



Filename = MS0446_PROTON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0446
 Solvent = CHLOROFORM-D
 Changen_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[MH
 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]
 Recvr_gain = 20
 Relaxation_delay = 4[s]
 Repetition_time = 5.74587904[s]
 Temp_get = 22.8[dc]





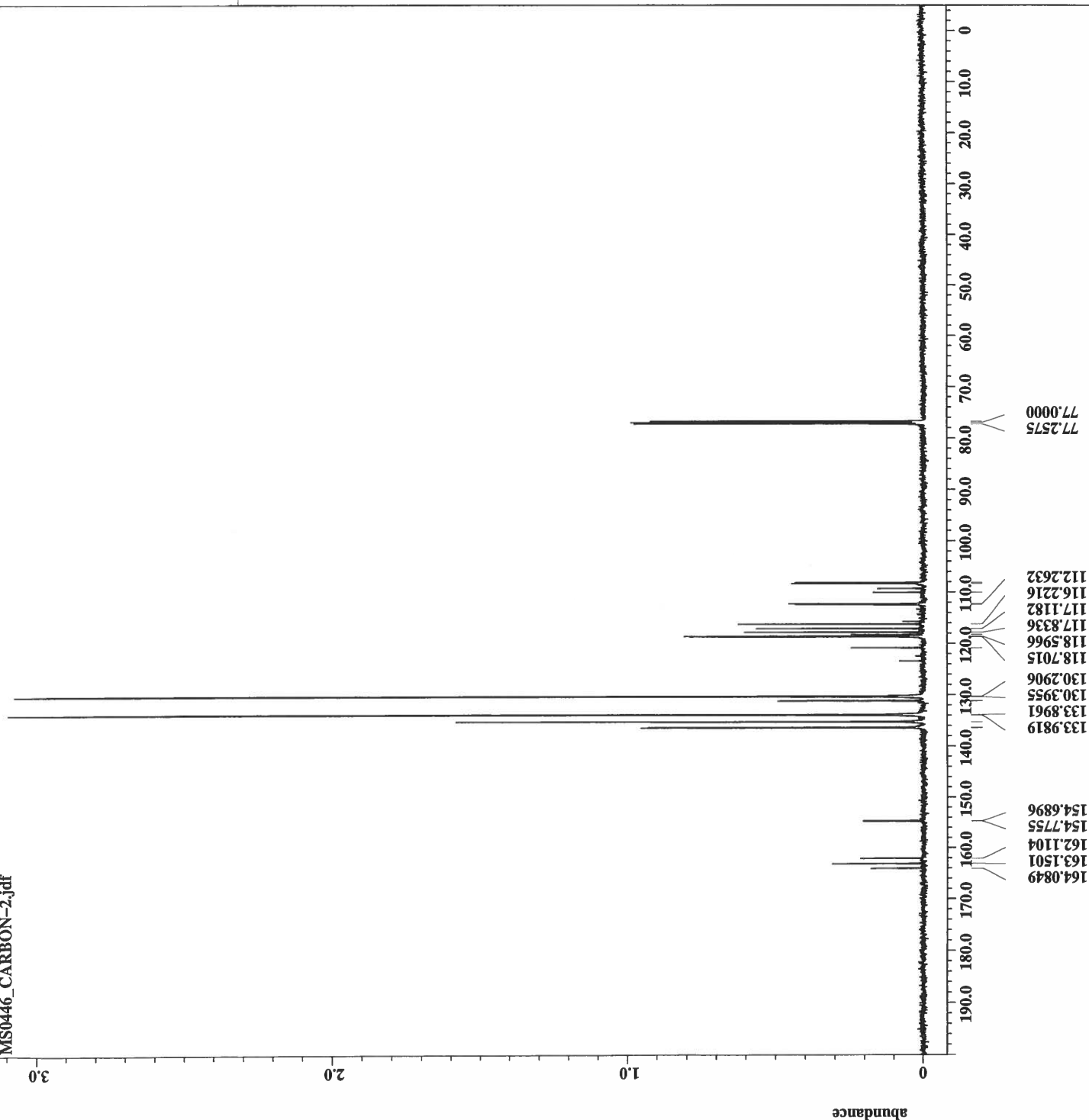
SOUTH ALABAMA
JAGUARS

```

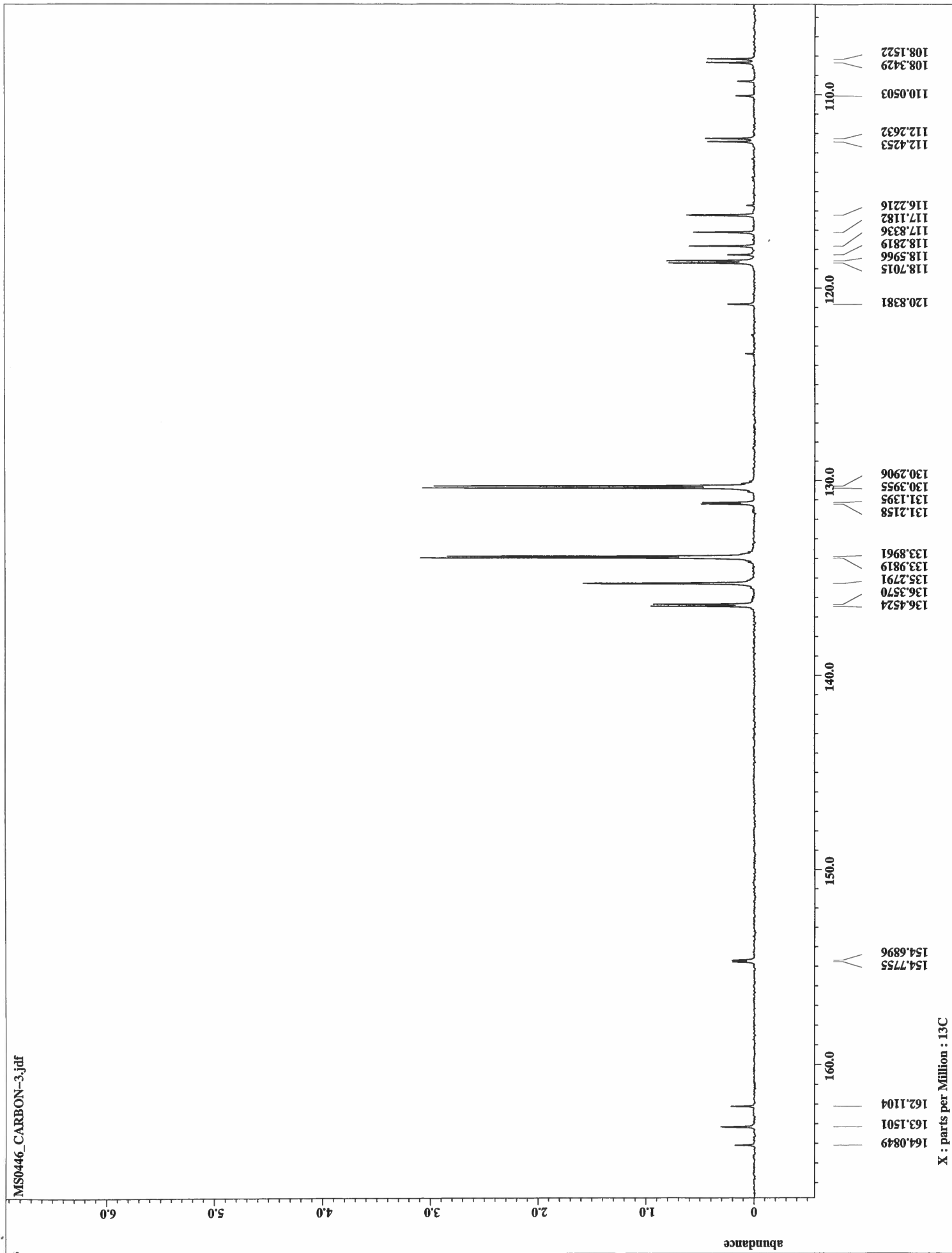
= MS0446_CARBON-2.jdf
= Jim Davis
= single_pulse_dec
= MS0446
= CHLOROFORM-D
= 1
= 14-MAY-2018 12:41:05
= 14-MAY-2018 12:16:51
= 14-MAY-2018 12:16:51
= 1D COMPLEX
= 26214
= 13C
= [ppm]
= X
= ECA 500
= JNM-ECA500
= 11.7473579 [T] (500 [MH
= 0.83361792 [s]
= 13C
= 125.76529768 [MHz]
= 100 [ppm]
= 32768
= 4
= 1.19959034 [Hz]
= 1H
= 39.3081761 [kHz]
= 500.15991521 [MHz]
= 5.0 [ppm]
= FALSE
= 1
= 128
= 128
= 13.2 [us]
= 0.83361792 [s]
= 30 [deg]
= 6 [dB]
= 4.4 [us]
= 20.7 [dB]
= 20.7 [dB]
= WALTZ
= TRUE
= 1 [s]
= TRUE
= 2 [s]
= 60
= 2 [s]
= 2.83361792 [s]
= 22.8 [dC]
  
```

```

Filename
Author
Experiment
Sample_id
Solvent
Changer_sample
Creation_time
Revision_time
Current_time
Data_format
Dim_size
Dim_title
Dim_units
Dimensions
Site
Spectrometer
Field_strength
X_acq_duration
X_domain
X_freq
X_offset
X_points
X_prescans
X_resolution
X_sweep
Irr_domain
Irr_freq
Irr_offset
Clipped
Mod_return
Scans
Total_scans
X_90_width
X_acq_time
X_angle
X_atn
X_pulse
Irr_atn_dec
Irr_atn_noe
Irr_noise
Decoupling
Initial_wait
Noe
Noe_time
Recvr_gain
Relaxation_delay
Repetition_time
Temp_get
  
```



X : parts per Million : 13C





SOUTH ALABAMA
JAGUARS

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-ECA500
Field_strength = 11.7473579[T] (500[MH
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]
Recvr_gain = 30
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get = 22.7[dc]

abundance

30.0

20.0

10.0

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 -160-070.0

-78.6218

-109.5572
-109.5419

X : parts per Million : 19F

3.0

2.0

1.0

0

abundance

-107.0

-108.0

-109.0

-109.5419
-109.5572

-110.0

-111.0

X : parts per Million : 19F

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

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

23.2219

X : parts per Million : 31P

**SOUTH ALABAMA**
JAGUARS™

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[MH
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 = TRUZ
Initial_wait = 1[s]
Noe = TRUE
Noe_time = 2[s]
Recvr_gain = 54
Relaxation_delay = 2[s]
Repetition_time = 2.64487424[s]
Temp_get = 23[degC]



215.53

73.71

7.7130
7.7061
7.5984
7.5824
7.5721
7.5561

X : parts per Million : 1H



SOUTH ALABAMA
JAGUARS

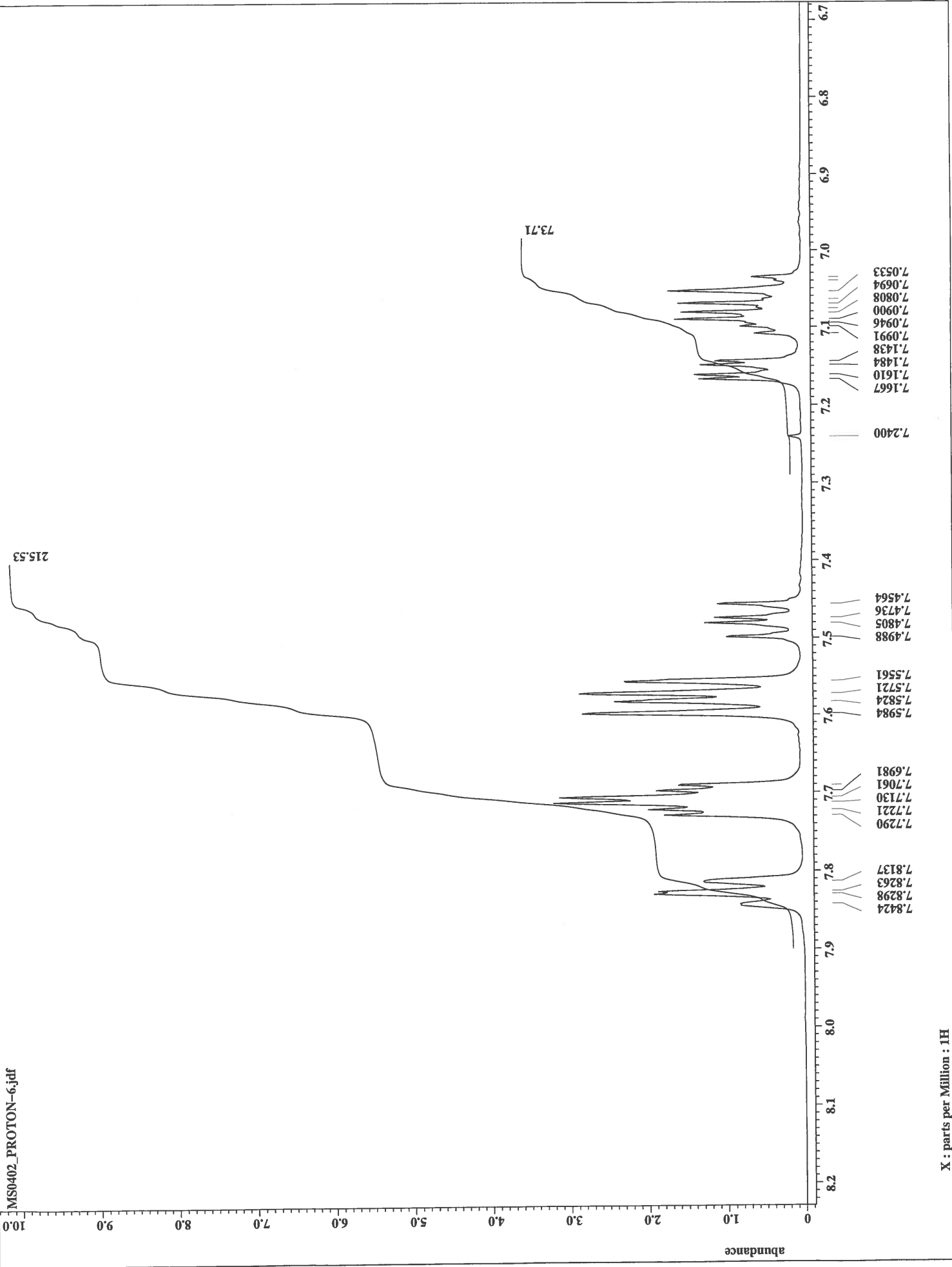
```

Filename      = MS0402_PROTON-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[MH
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_pulse      = Off
Irr_mode       = Off
Dante_preset   = FALSE
Initial_wait   = 1[s]
Recvr_gain     = 24
Relaxation_delay = 4[s]
Repetition_time = 5.74587904[s]
Temp_get       = 22.8[dc]
  
```



SOUTH ALABAMA
JAGUARS

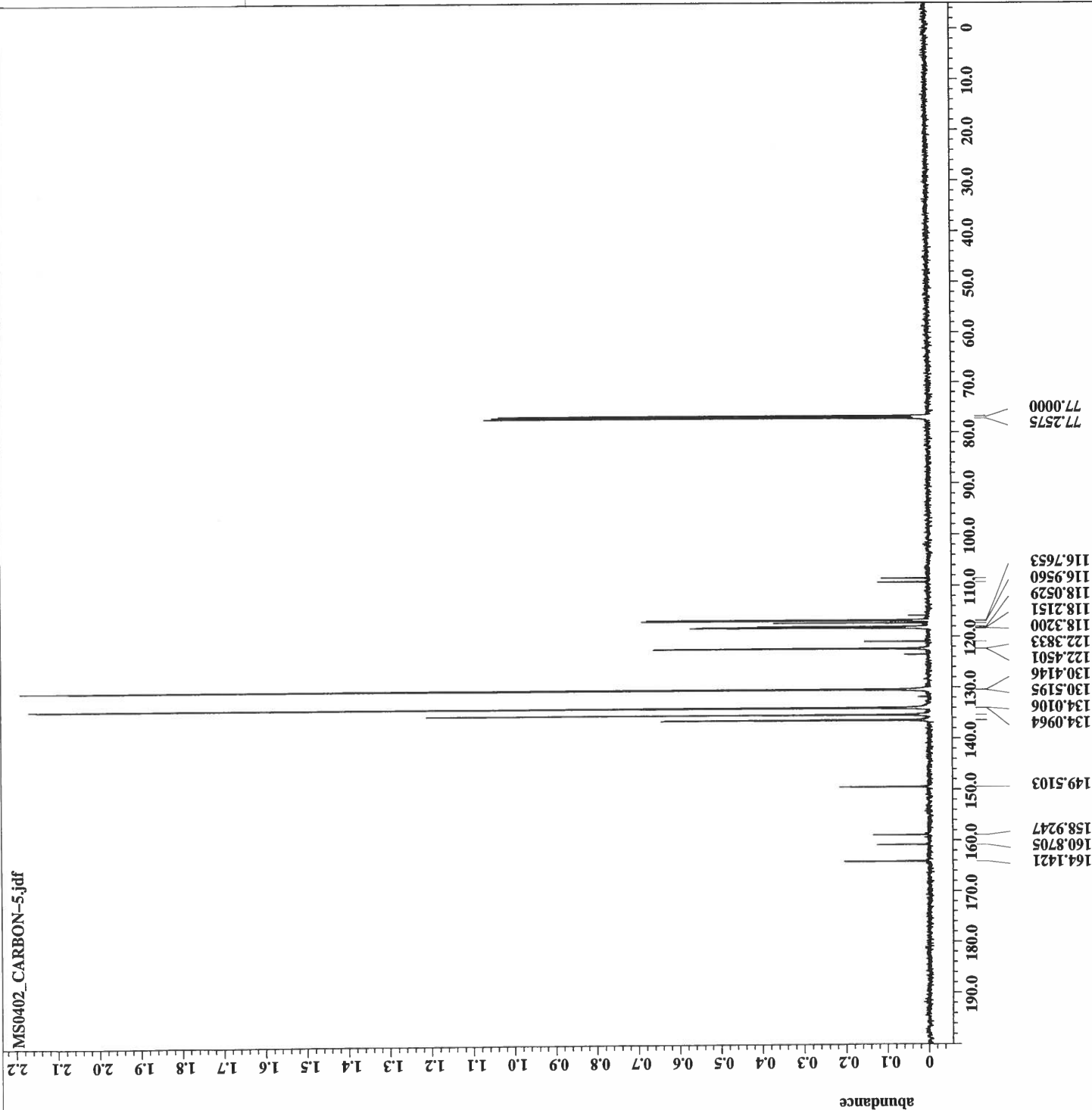
```

Filename      = MS0402_CARBON-5.jdf
Author       = Jim Davis
Experiment    = single_pulse_dec
Sample_id    = MS0402
Solvent      = CHLOROFORM-D
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    = 13C
Dim_units    = [ppm]
Dimensions   = x
Site         = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579[T] (500[MH
X_acq_duration = 0.83361792[s]
X_domain       = 13C
X_freq         = 125.76529768[MHz]
X_offset       = 100[ppm]
X_points       = 32768
X_prescans     = 4
X_resolution   = 1.19959034[Hz]
X_sweep        = 39.3081761[kHz]
X_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      = WAITZ
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       = 22.9[dc]
  
```



X : parts per Million : 13C

3.0

2.0

1.0

abundance

164.1421

160.8705

158.9247

149.5103

160.0

150.0

140.0

136.4905

136.4047

135.4127

134.0964

134.0106

130.5195

130.4146

130.0

123.5375

122.4501

122.3833

120.9812

120.0

118.3200

118.2151

118.0529

117.3376

116.9560

116.7653

109.3349

108.5718

110.0

X : parts per Million : 13C



SOUTH ALABAMA
JAGUARS

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[MH
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_pulse = Off
Irr_mode = Off
Dante_presat = FALSE
Initial_wait = 1[s]
Recvr_gain = 36
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get = 22.6[degC]

30.0

20.0

10.0

0

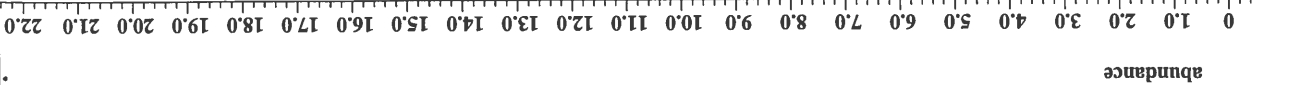
abundance

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

-78.6142

-116.4623

X : parts per Million : 19F



23.2295

X : parts per Million : 31P



SOUTH ALABAMA
JAGUARS

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[MH
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 = 1.5068995[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]
Recvr_gain = 56
Relaxation_delay = 2[s]
Repetition_time = 2.64487424[s]
Temp_get = 22.9[dc]

abundance

196.95

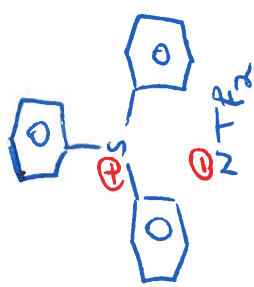
58.31

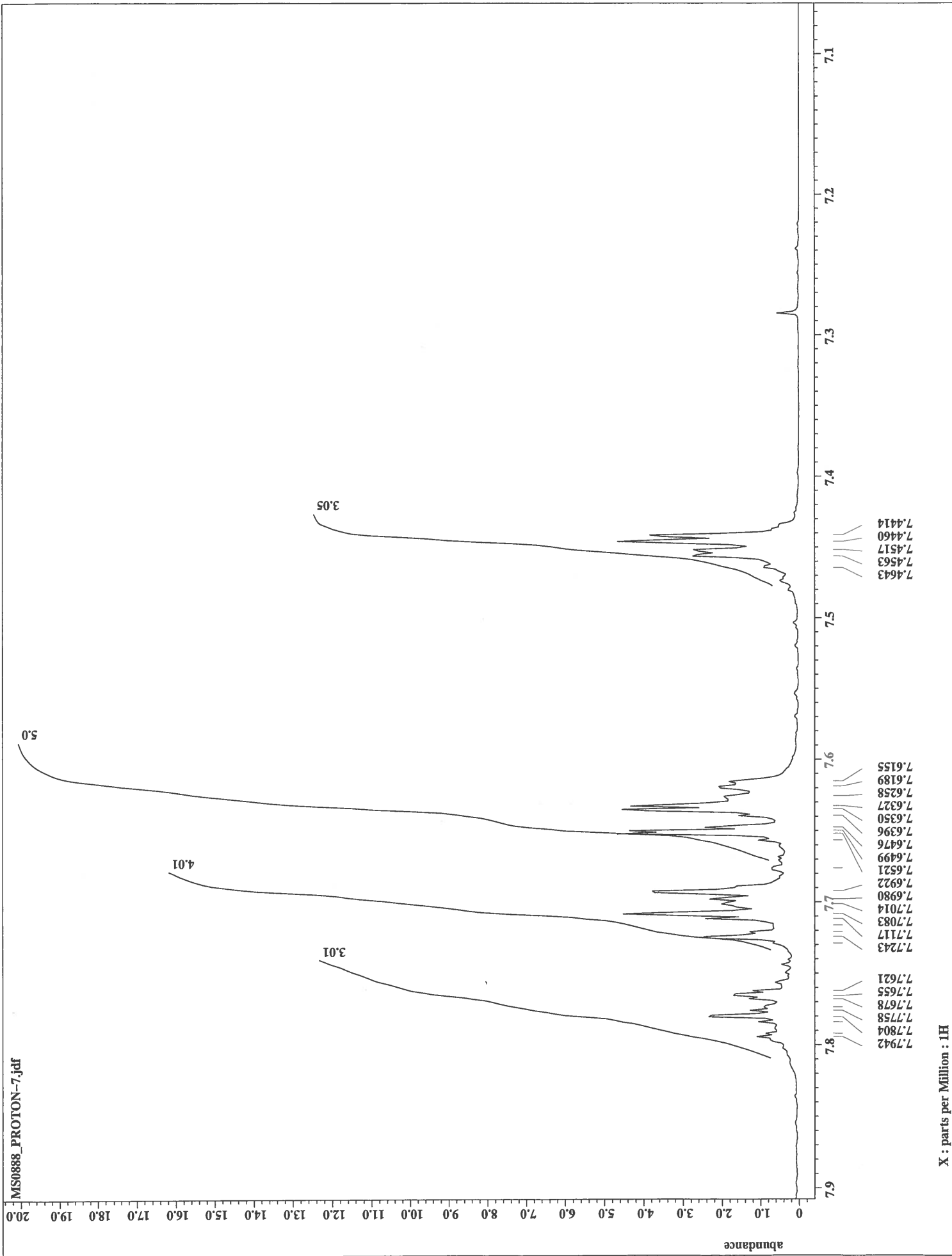
7.7083
7.6521
7.6499
7.6350
7.4460

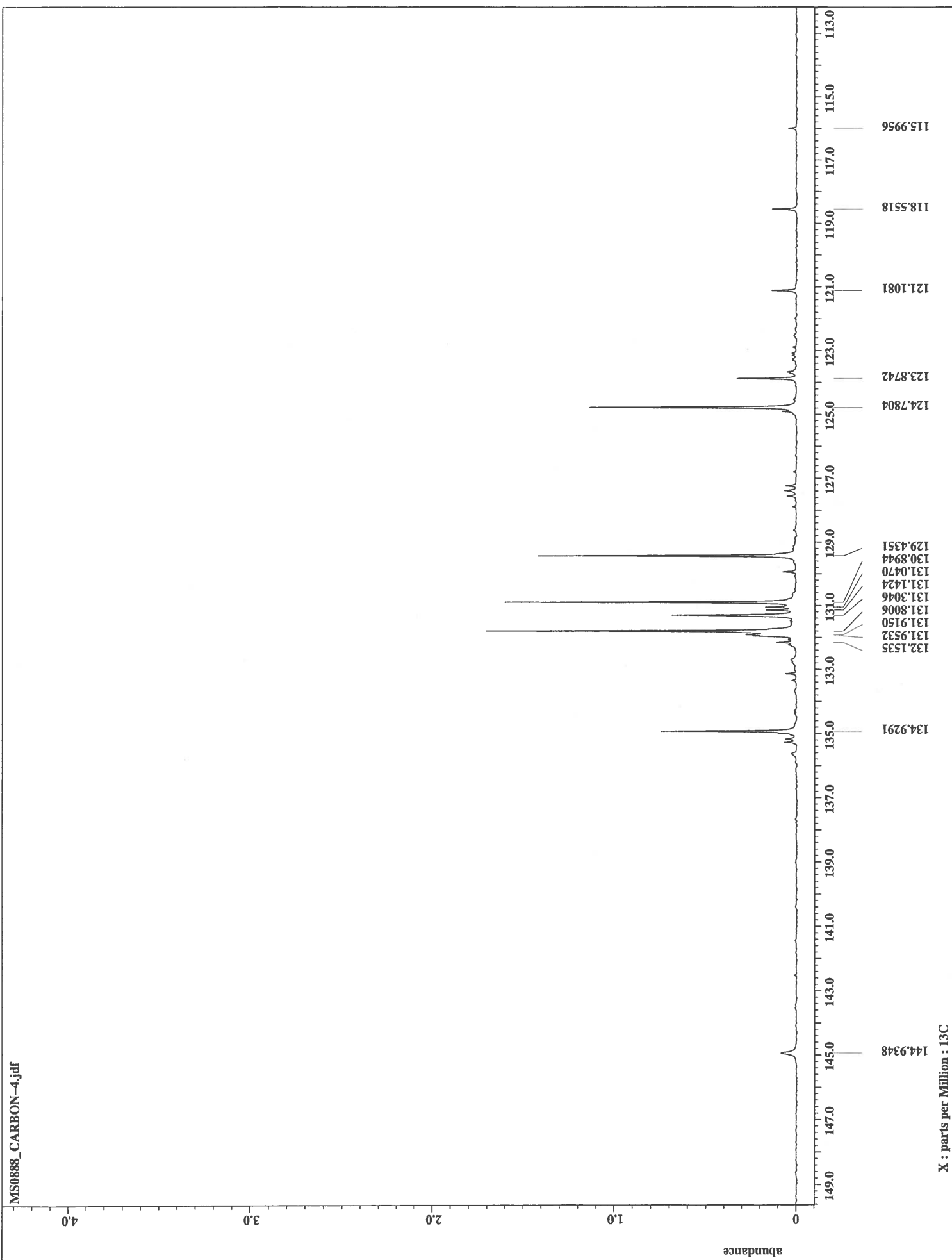
10.0 9.0 8.0 7.0 6.0 5.0 4.0 3.0 2.0 1.0

2.1552

2.58







MS0888_FLUORINE-4.jdf

40.0

30.0

20.0

10.0

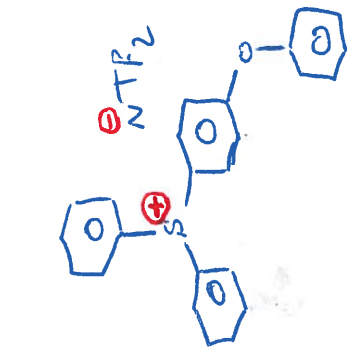
0

abundance

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

-78.5981

X : parts per Million : 19F



168.72

66.69

25.1

abundance

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

7.6901
7.6351
7.6294
7.6191
7.6156
7.6110

X : parts per Million : 1H

SOUTH ALABAMA
JAGUARSTM

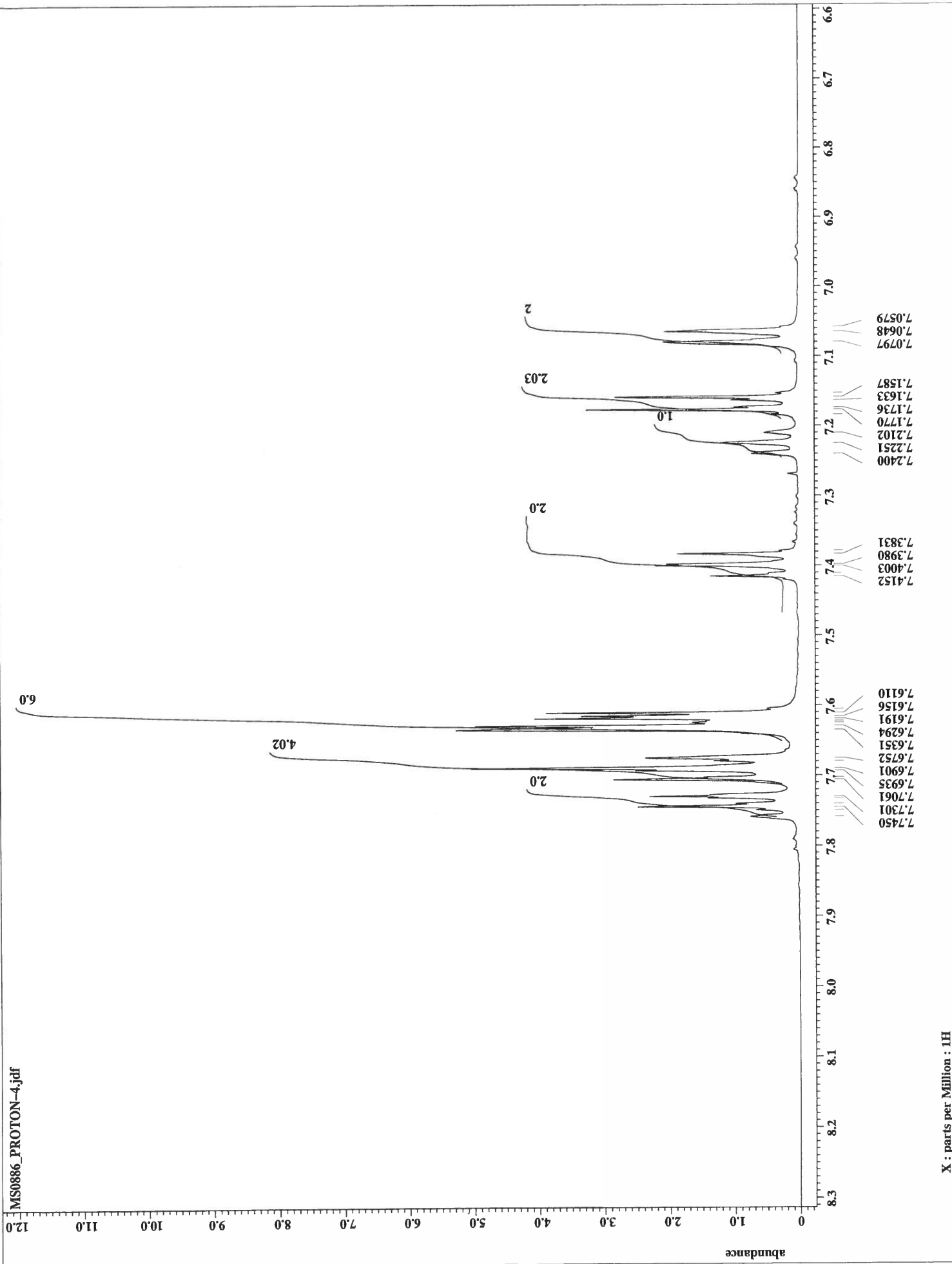
```

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

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

Field_strength = 11.7473579[T] (500[MH
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_atn         = 4[dB]
X_pulse       = 5.65[us]
Irr_mode      = Off
Tri_mode      = Off
Dante_preset  = FALSE
Initial_wait  = 1[s]
Recvr_gain    = 22
Relaxation_delay = 4[s]
Repetition_time = 5.74587904[s]
Temp_get      = 19[dc]
  
```

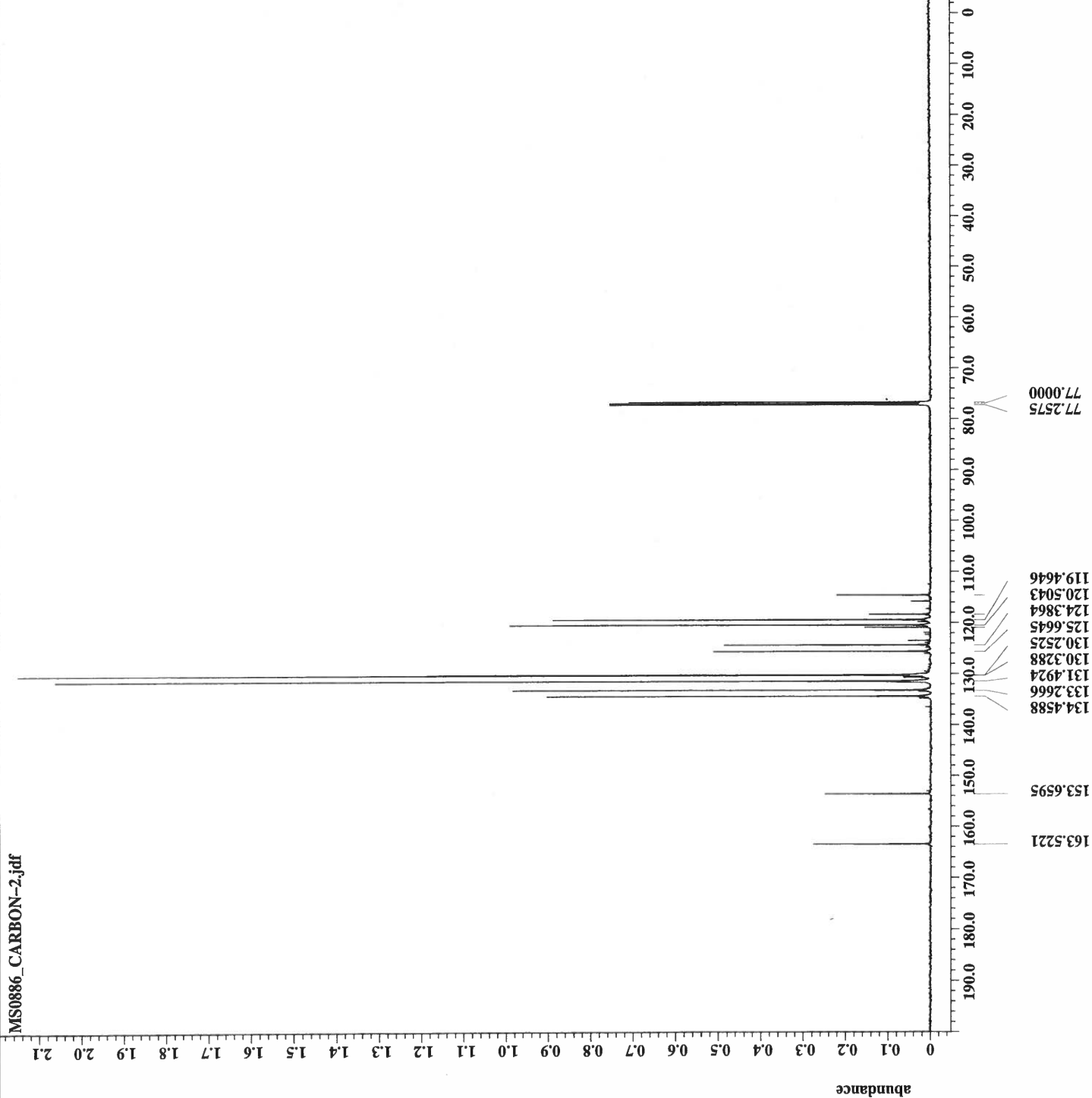




SOUTH ALABAMA
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```

= MS0886_CARBON-2.jdf
= Jim Davis
= single_pulse_dec
= MS0886
= CHLOROFORM-D
= 29-NOV-2019 13:40:22
= 29-NOV-2019 13:11:06
= 29-NOV-2019 13:11:06
= 1D COMPLEX
= 26214
= 13C
= [ppm]
= X
= ECA 500
= JNM-ECA500
= 11.7473579[T] (500[MH
= 0.83361792[s]
= 13C
= 125.76529768[MHz]
= 100[ppm]
= 32768
= 4
= 1.19959034[Hz]
= 39.3081761[KHz]
= 1H
= 500.15991521[MHz]
= 5.0[ppm]
= FALSE
= 1
= 1024
= 1024
= 13[us]
= 0.83361792[s]
= 30[deg]
= 6[db]
= 4.33333333[us]
= 21.2[db]
= 21.2[db]
= WAITZ
= TRUE
= 1[s]
= TRUE
= 2[s]
= 60
= 2[s]
= 2.83361792[s]
= 20[dc]
  
```



X : parts per Million : 13C

3.0

2.0

1.0

abundance

0

160.0

150.0

140.0

130.0

120.0

110.0

163,5221

153,6595

134,4588

133,2666

131,4924

130,3288

130,2525

125,6645

124,3864

123,5089

120,9526

120,5043

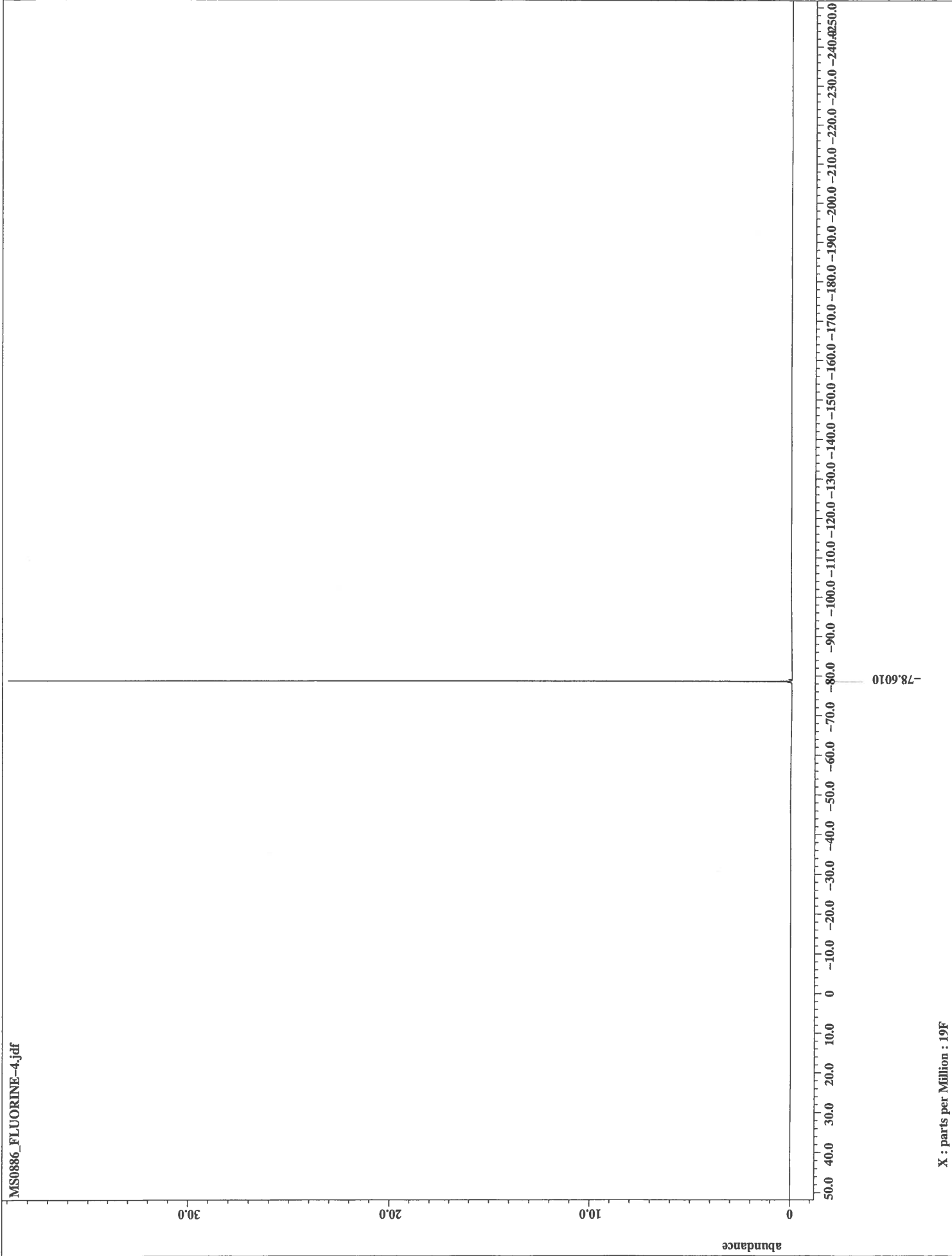
119,4646

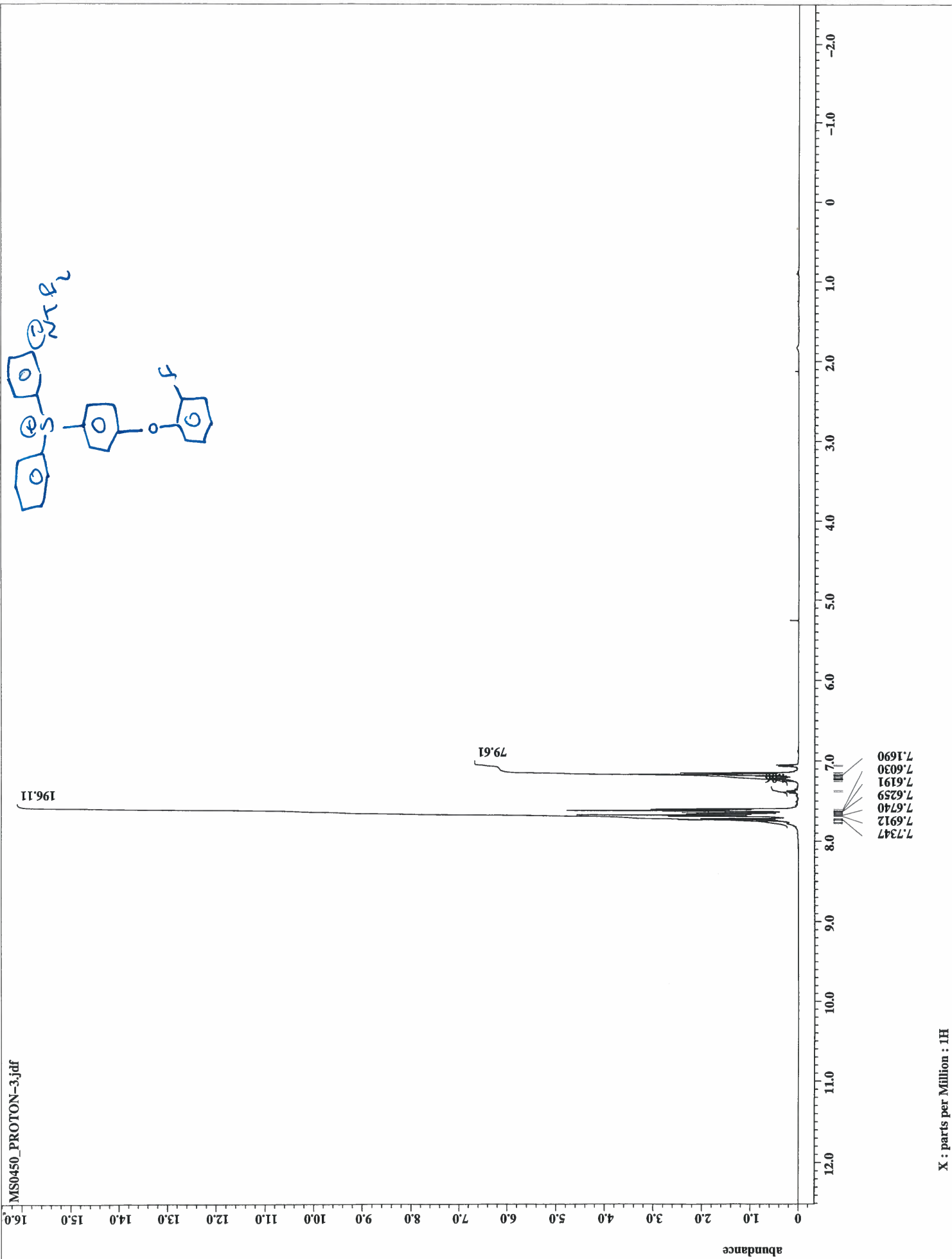
118,3963

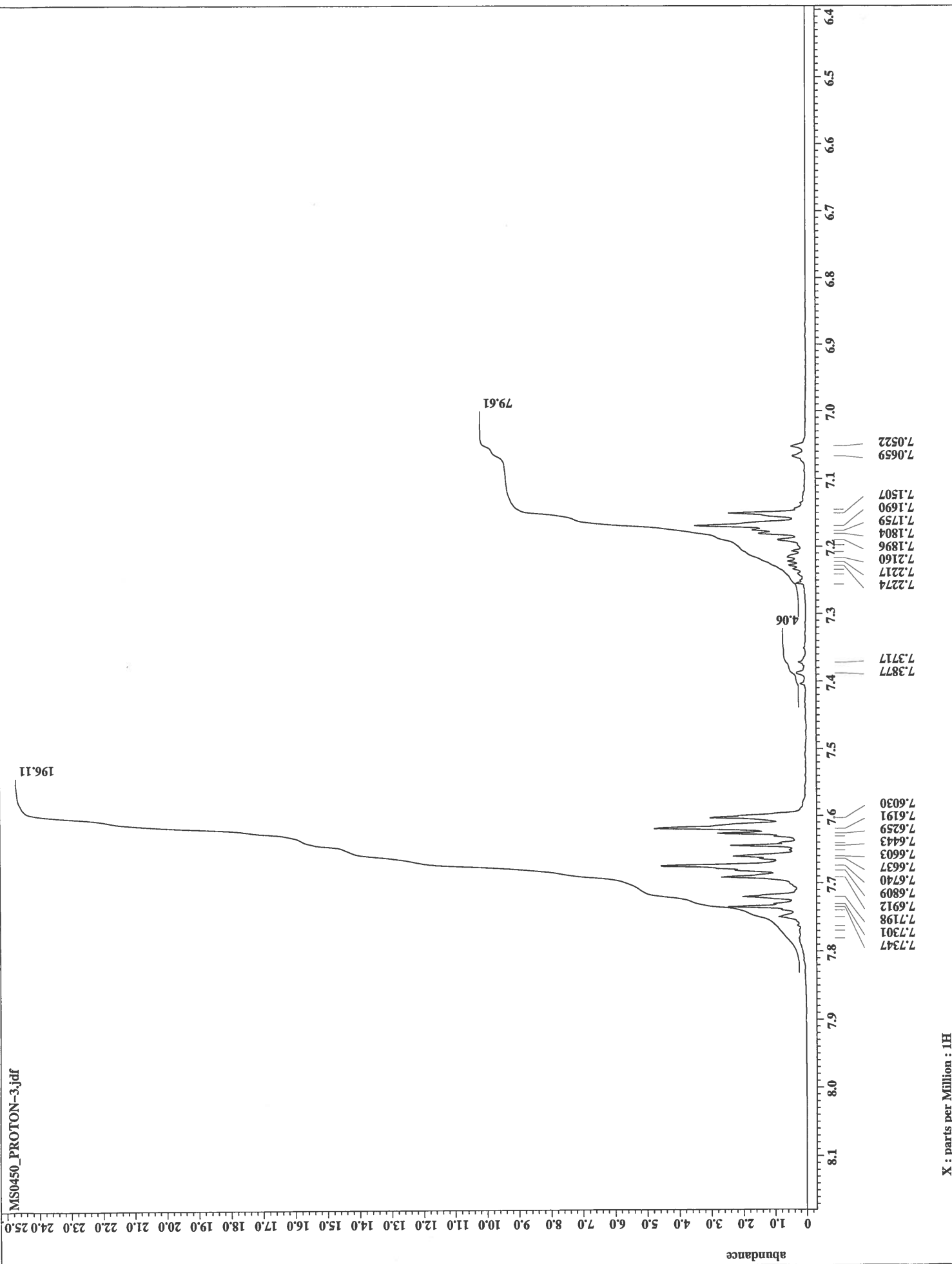
115,8400

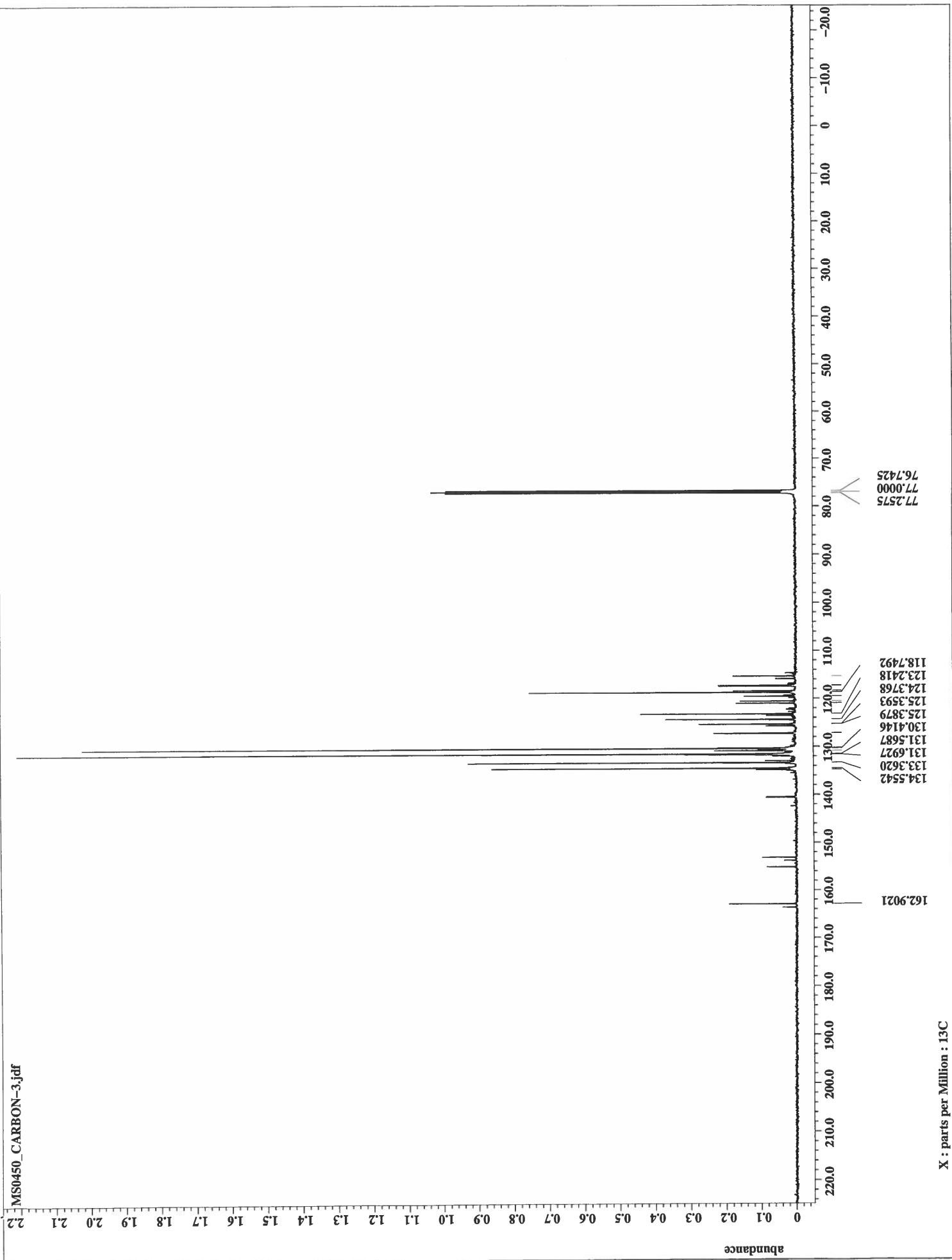
114,6191

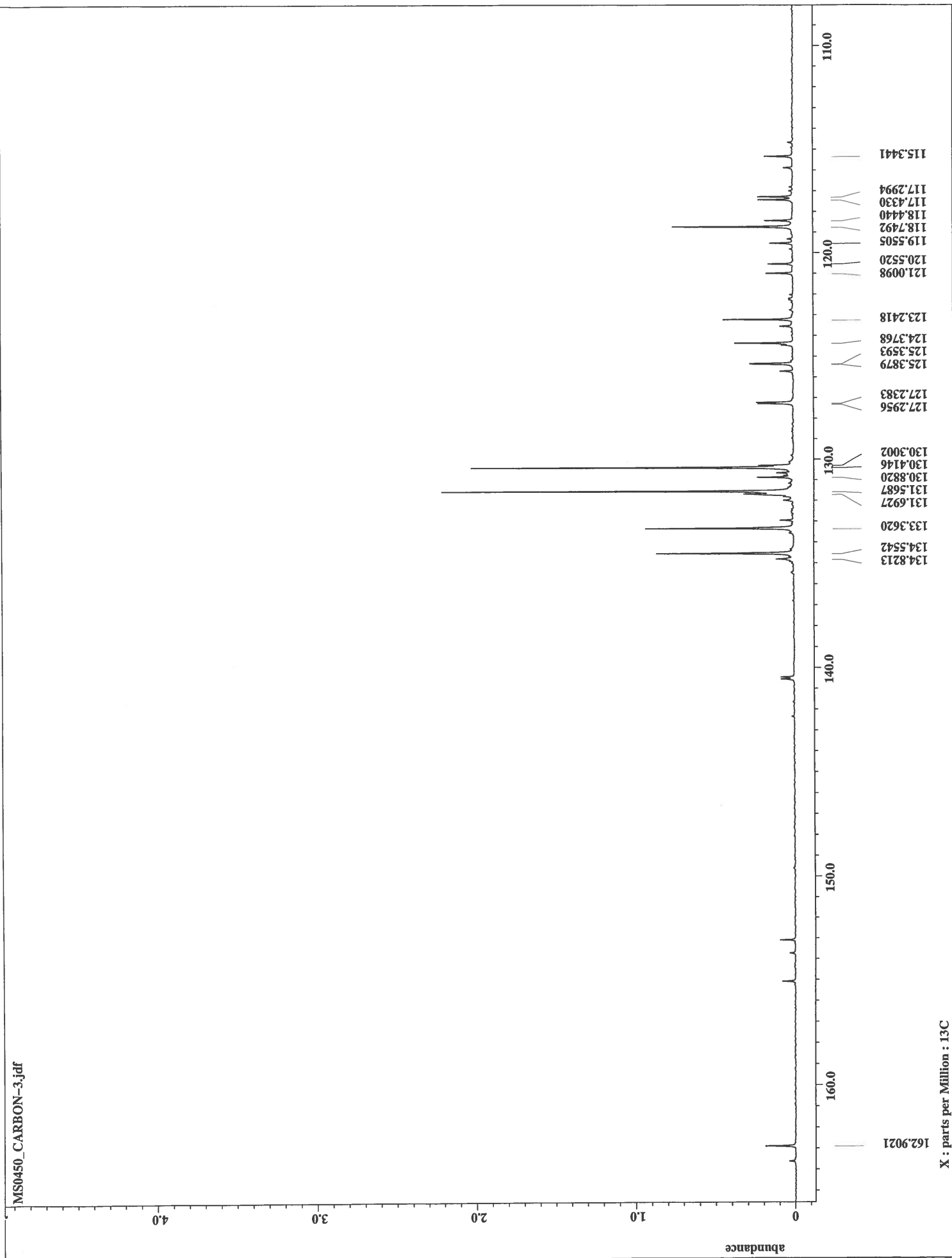
X : parts per Million : 13C











30.0

20.0

10.0

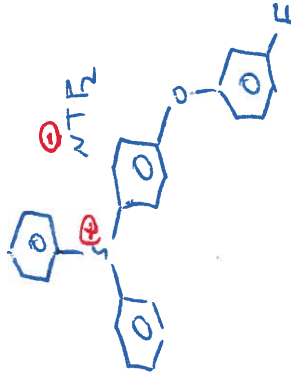
0

abundance

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

-78.6142

187.76



32.34

18.65

14.24

7.7187
7.6889
7.6706
7.6546
7.2400
6.8254
6.8071

X : parts per Million : 1H



```

Filename      = MS0448_PROTON-2.jdf
Author       = Jim Davis
Experiment    = single_pulse.ex2
Sample_id    = MS0448
Solvent      = CHLOROFORM-D
Creation_time = 26-NOV-2019 12:41:31
Revision_time = 26-NOV-2019 12:12:27
Current_time  = 26-NOV-2019 12:12:27

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[MH
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_atn          = 4[db]
X_pulse        = 5.65[us]
Irr_mode       = Off
Tri_mode       = Off
Dante_preset   = FALSE
Initial_wait   = 1[s]
Recvr_gain     = 24
Relaxation_delay = 4[s]
Repetition_time = 5.74587904[s]
Temp_get       = 19.4[dc]
  
```



SOUTH ALABAMA
JAGUARS

```

= MS0448_CARBON-2.jdf
= Jim Davis
= single_pulse_dec
= MS0448
= CHLOROFORM-D
= 26-NOV-2019 13:01:16
= 26-NOV-2019 12:32:12
= 26-NOV-2019 12:32:12
= 1D COMPLEX
= 26214
= 13C
= [ppm]
= X
= ECA 500
= JNM-ECA500
= 11.7473579[T] (500[MHz]
= 0.83361792[s]
= 13C
= 125.76529768[MHz]
= 100[ppm]
= 32768
= 4
= 1.19959034[Hz]
= 39.3081761[MHz]
= 1H
= 500.15991521[MHz]
= 5.0[ppm]
= FALSE
= 1
= 400
= 400
= 13[us]
= 0.83361792[s]
= 30[deg]
= 6[db]
= 4.33333333[us]
= 21.2[db]
= 21.2[db]
= WALTZ
= TRUE
= 1[s]
= TRUE
= 2[s]
= 60
= 2[s]
= 2.83361792[s]
= 19.6[dc]
  
```

```

Filename
Author
Experiment
Sample_id
Solvent
Creation_time
Revision_time
Current_time
  
```

```

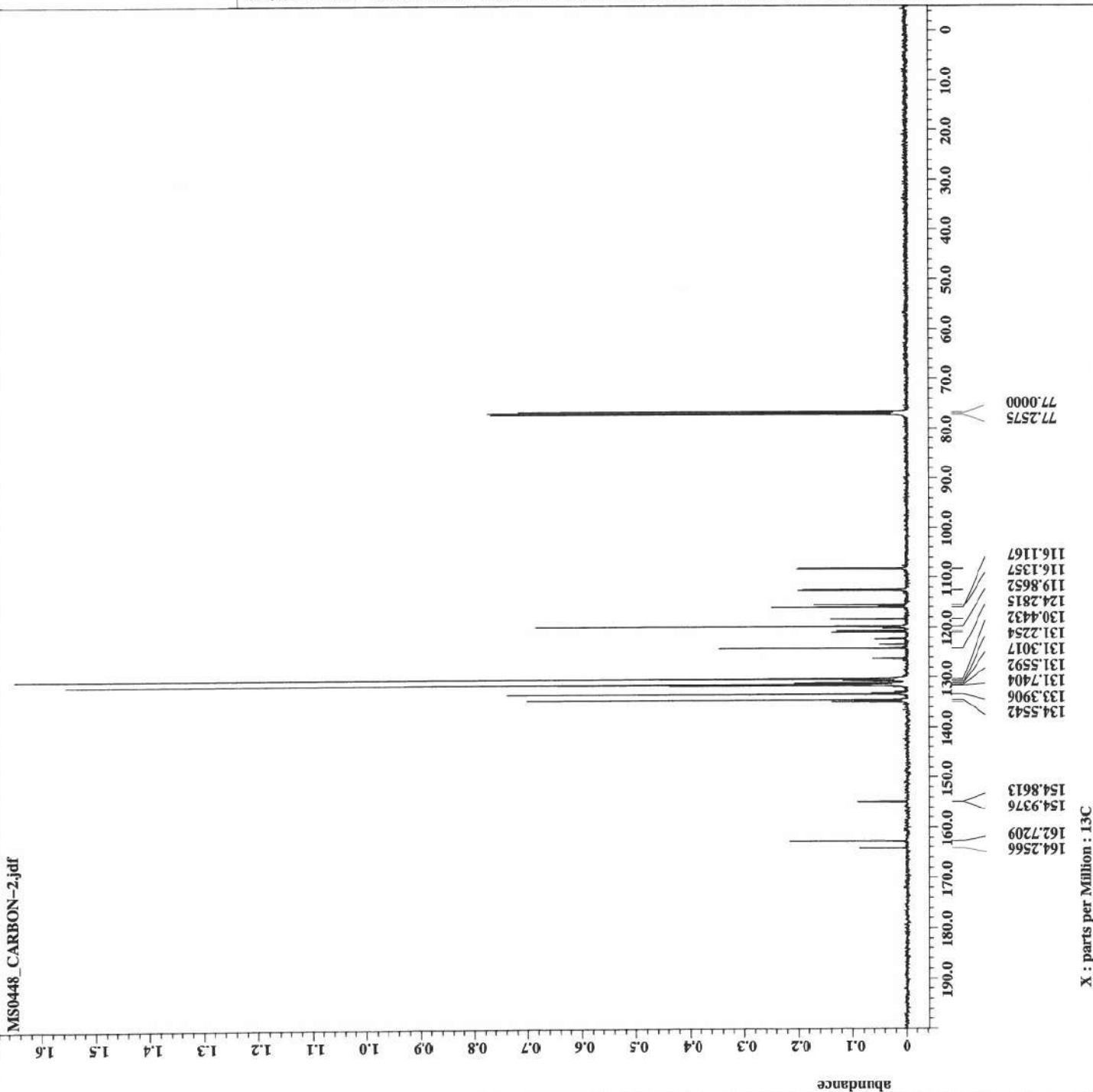
Data_format
Dim_size
Dim_title
Dim_units
Dimensions
Site
Spectrometer
  
```

```

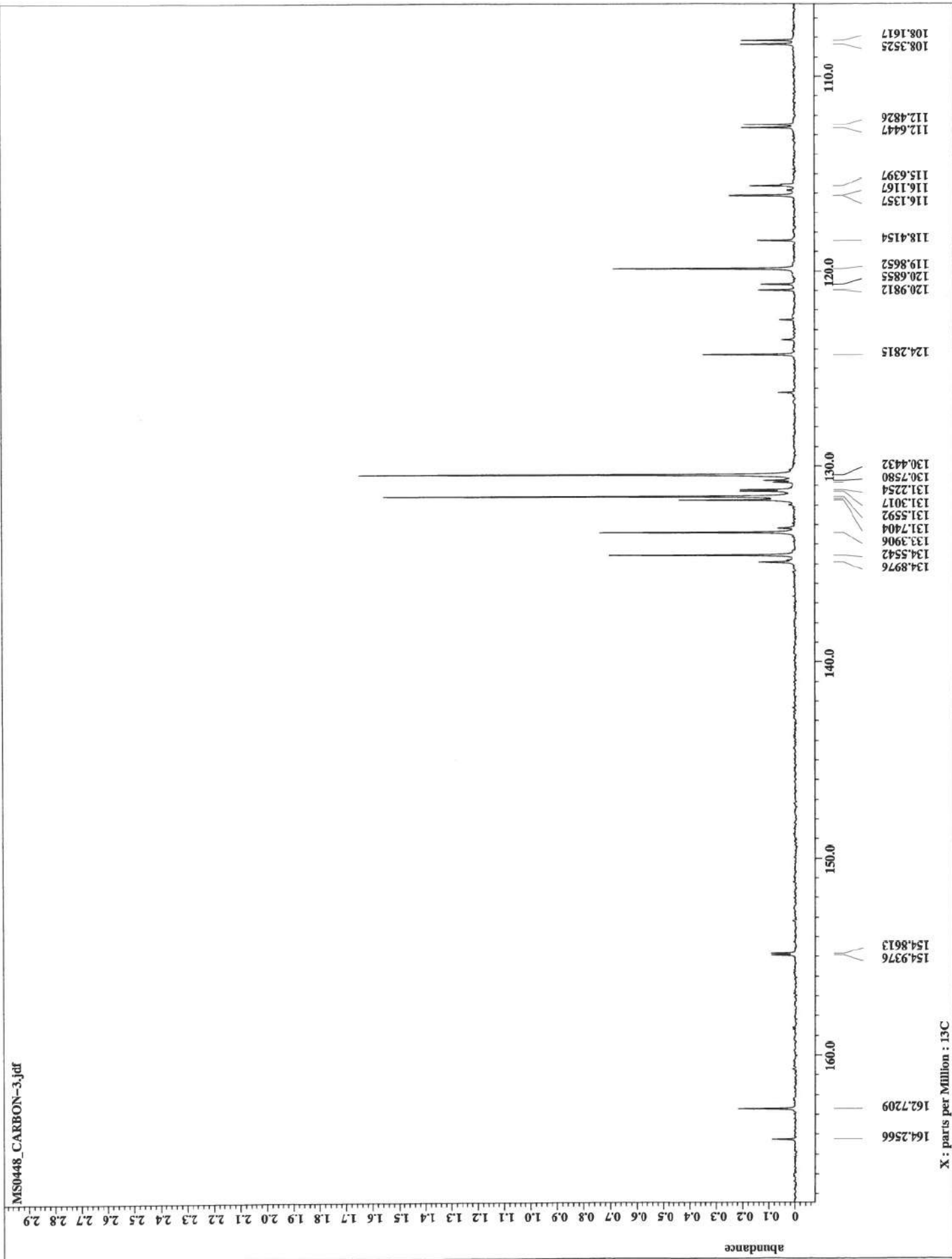
Field_strength
X_acq_duration
X_domain
X_freq
X_offset
X_points
X_prescans
X_resolution
X_sweep
Irr_domain
Irr_freq
Irr_offset
Clipped
Mod_return
Scans
Total_scans
  
```

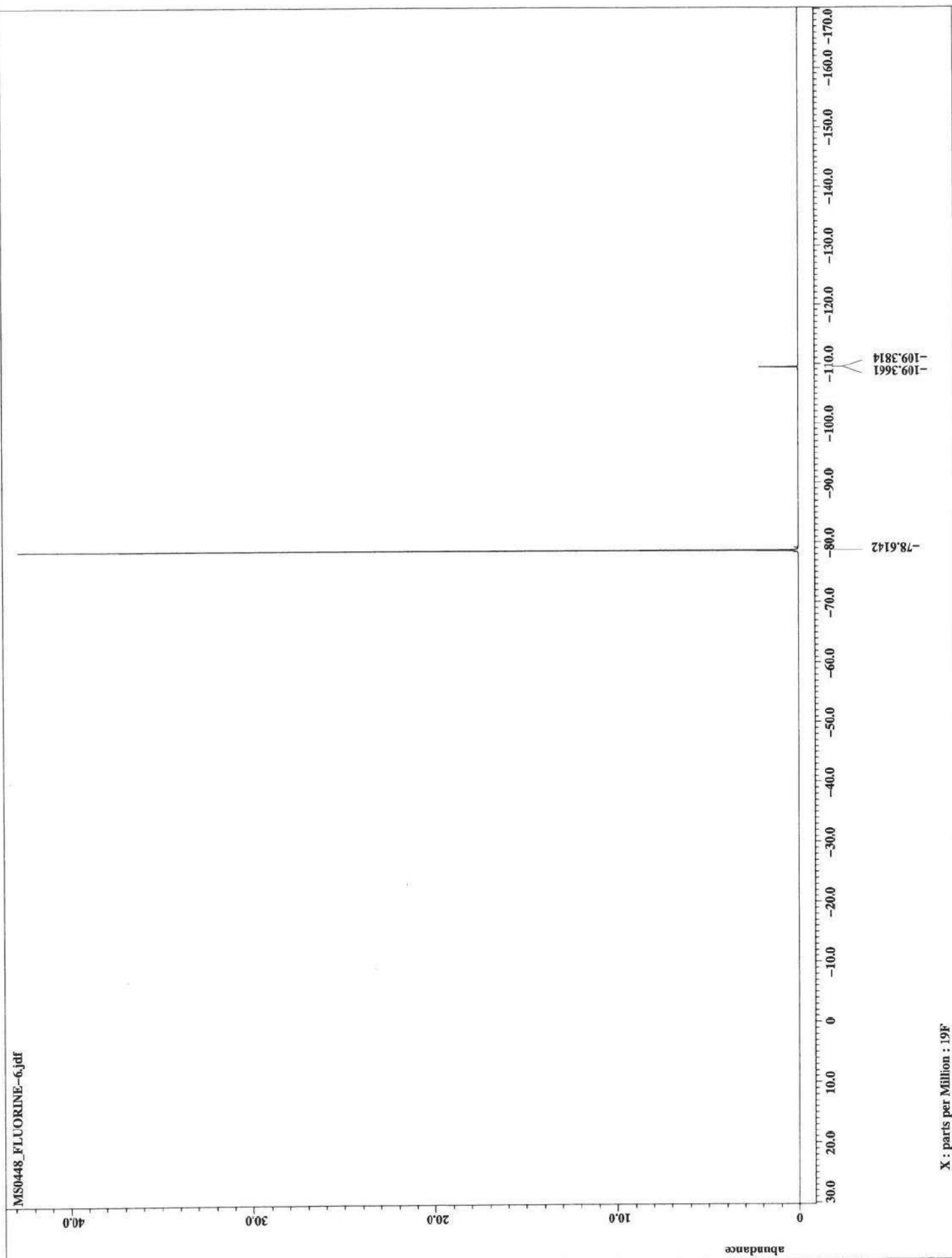
```

X_90_width
X_acq_time
X_angle
X_atn
X_pulse
Irr_atn_dec
Irr_atn_noe
Irr_noise
Decoupling
Initial_wait
Noe
Noe_time
Recvr_gain
Relaxation_delay
Repetition_time
Temp_get
  
```



X : parts per Million : 13C





179.58

78.98

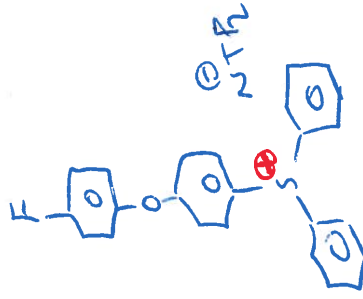
7.7634
7.7095
7.6923
7.2400
7.1472
7.1346

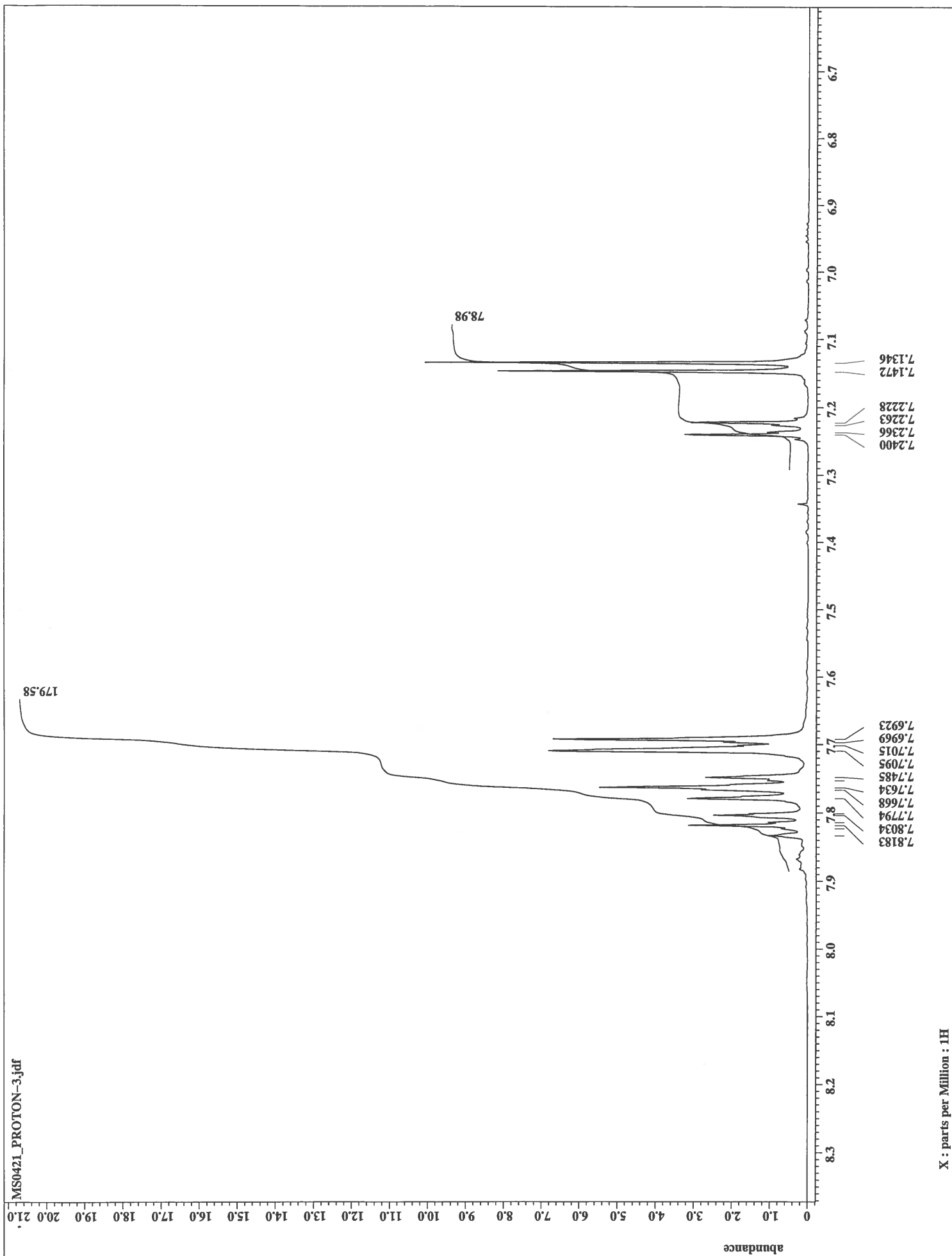
X : parts per Million : 1H



SOUTH ALABAMA
JAGUARS

Filename = MS0421_PROTON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0421
 Solvent = CHLOROFORM-D
 Changer_sample = 1
 Creation_time = 18-APR-2018 10:34:59
 Revision_time = 18-APR-2018 10:12:47
 Current_time = 18-APR-2018 10:12:47
 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 [MH
 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_presat = FALSE
 Initial_wait = 1 [s]
 Recvr_gain = 22
 Relaxation_delay = 4 [s]
 Repetition_time = 5.74587904 [s]
 Temp_get = 22.7 [dC]







SOUTH ALABAMA
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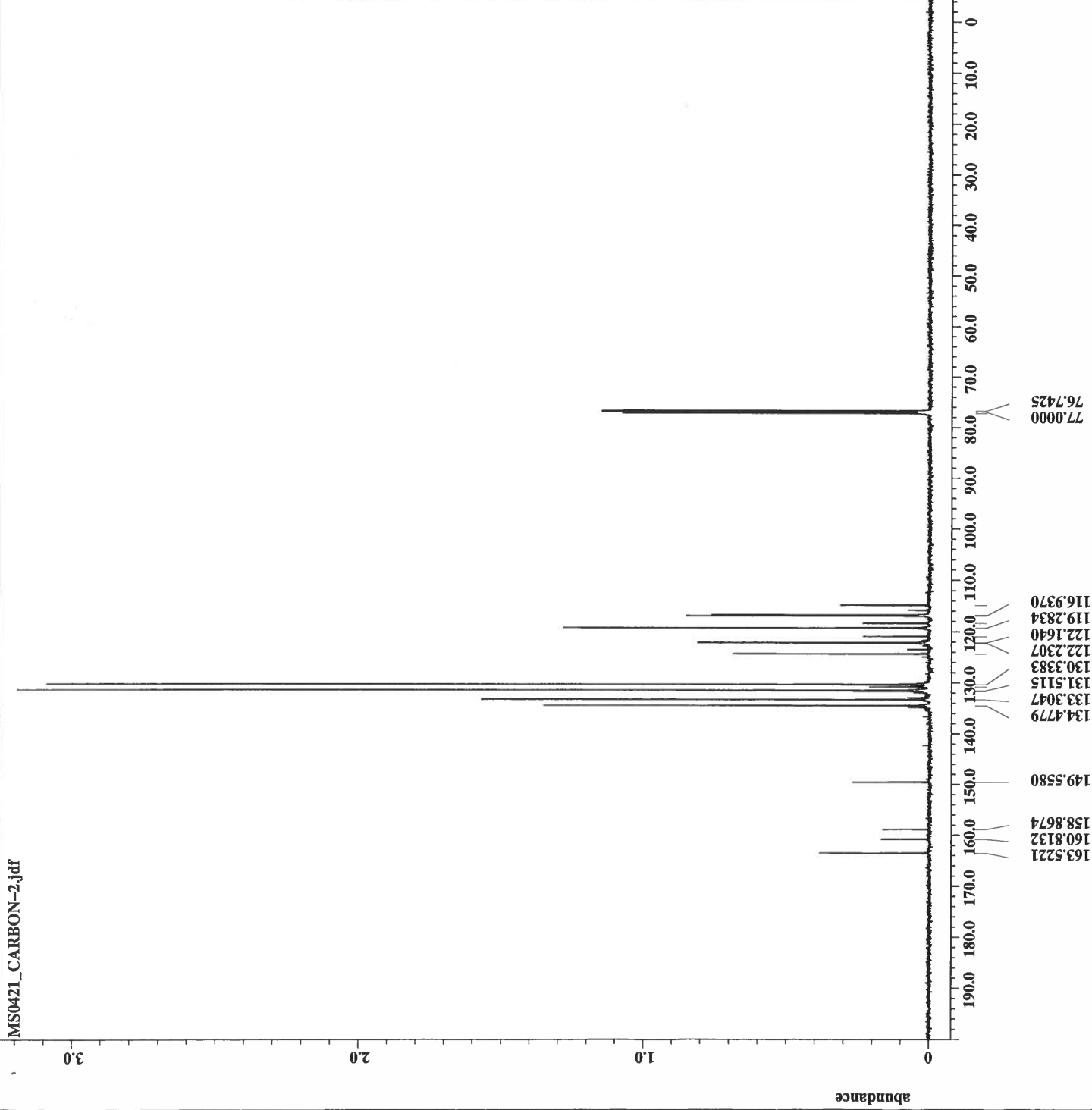
```

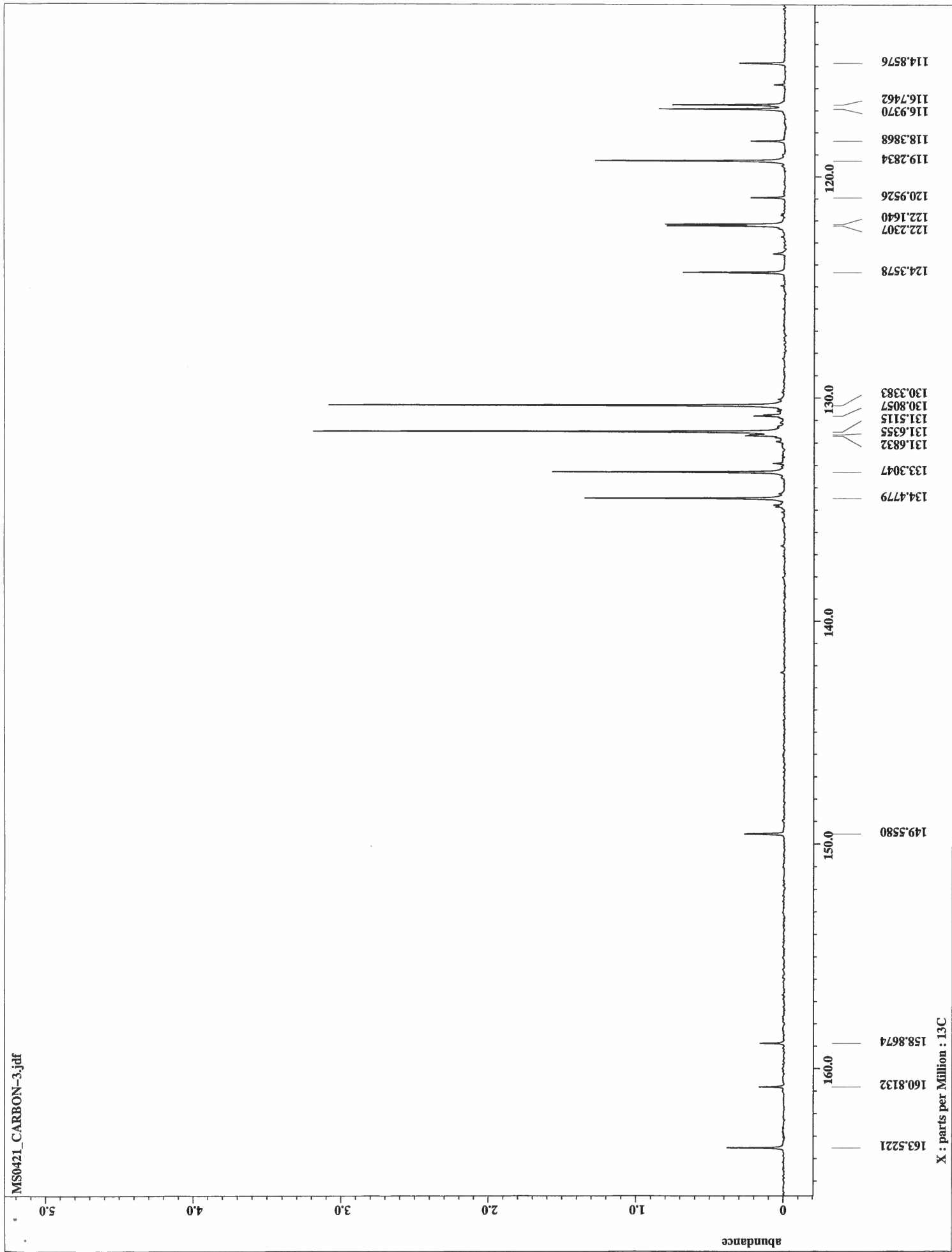
Filename      = MS0421_CARBON-2.jdf
Author       = Jim Davis
Experiment   = single_pulse_dec
Sample_id    = MS0421
Solvent      = CHLOROFORM-D
Charger_sample = 1
Creation_time   = 18-APR-2018 10:49:35
Revision_time  = 18-APR-2018 10:27:22
Current_time   = 18-APR-2018 10:27:22

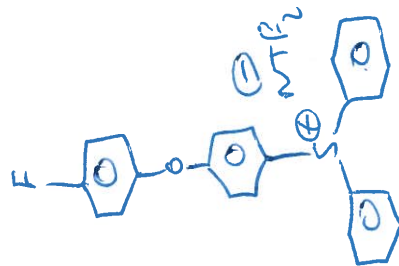
Data_format  = 1D COMPLEX
Dim_size     = 26214
Dim_title    = 13C
Dim_units    = [ppm]
Dimensions   = X
Site         = ECA 500
Spectrometer = JNM-ECA500

Field_strength = 11.7473579 [T] (500 [MH
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       = 23.4 [dC]
  
```





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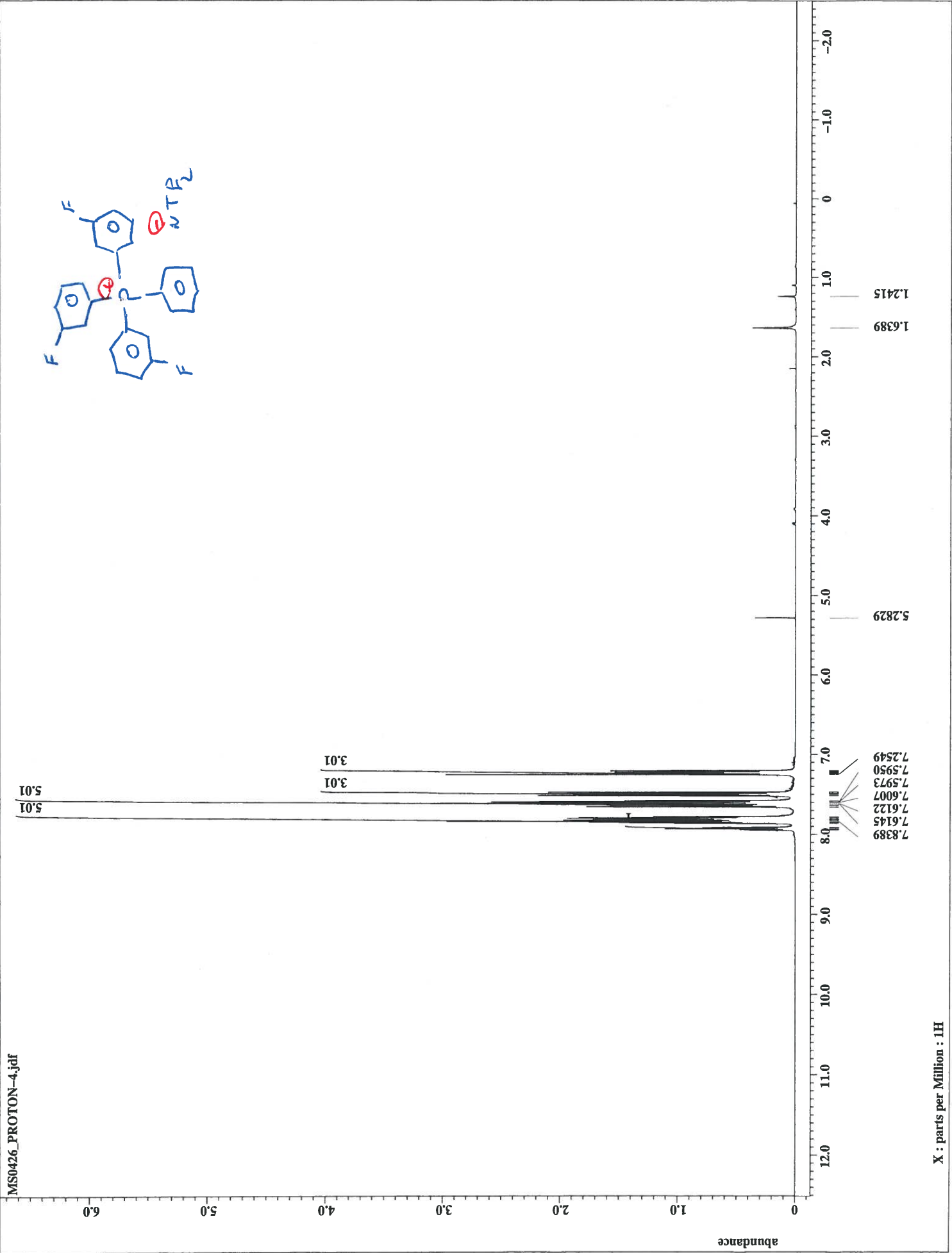
Filename = MS0421_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0421
Solvent = CHLOROFORM-D
Charger_sample = 1
Creation_time = 18-APR-2018 10:31:25
Revision_time = 18-APR-2018 10:09:13
Current_time = 18-APR-2018 10:09:13
Data_format = 1D COMPLEX
Dim_size = 52428
Dim_title = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength = 11.7473579 [T] (500 [MH
X_acq_duration = 0.55574528 [s]
X_domain = 19F
X_freq = 470.62046084 [MHz]
X_offset = -70 [ppm]
X_points = 65536
X_resolution = 1
X_prescans = 1
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_presat = FALSE
Initial_wait = 1 [s]
Recvr_gain = 32
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.6 [dC]

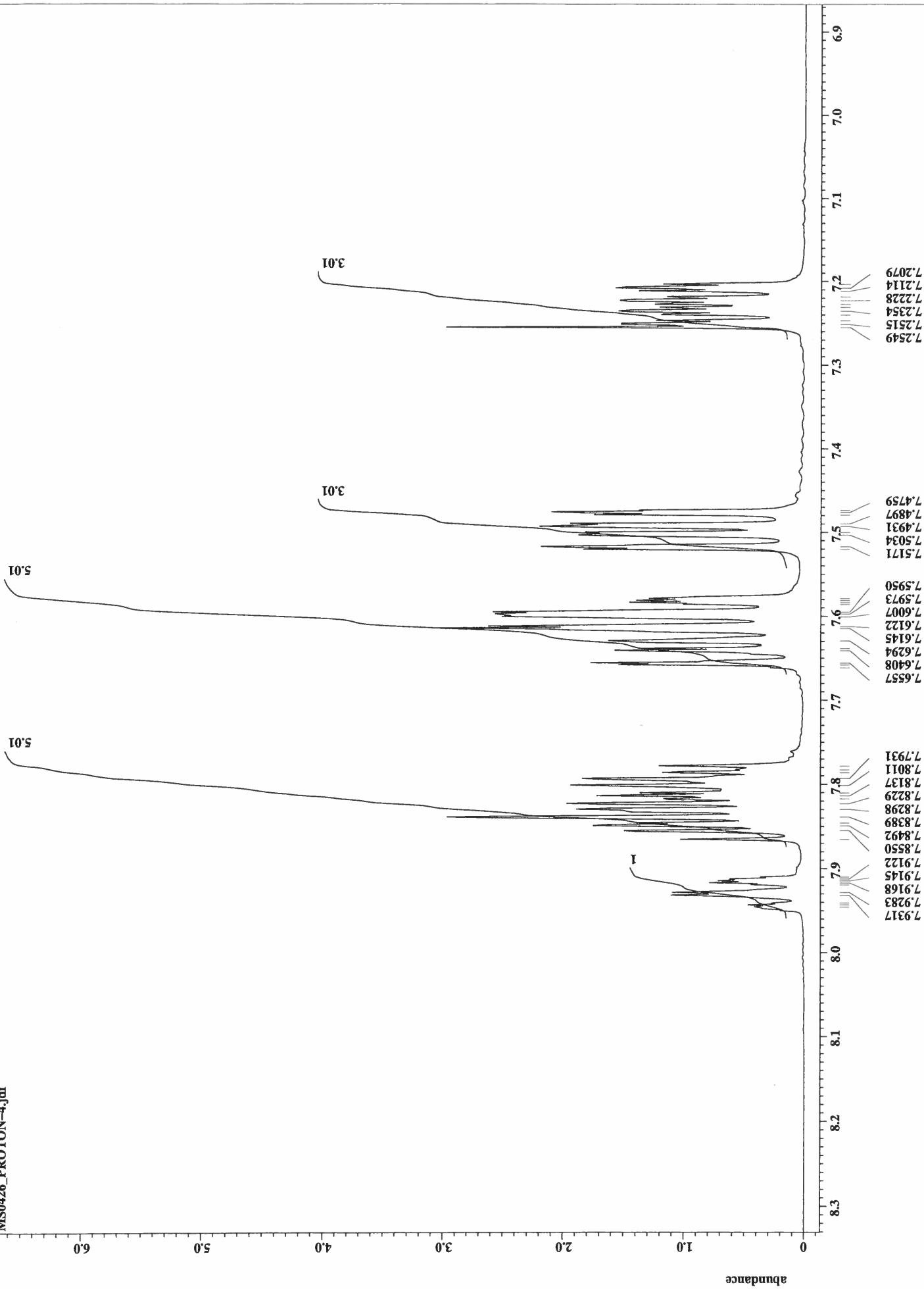
abundance

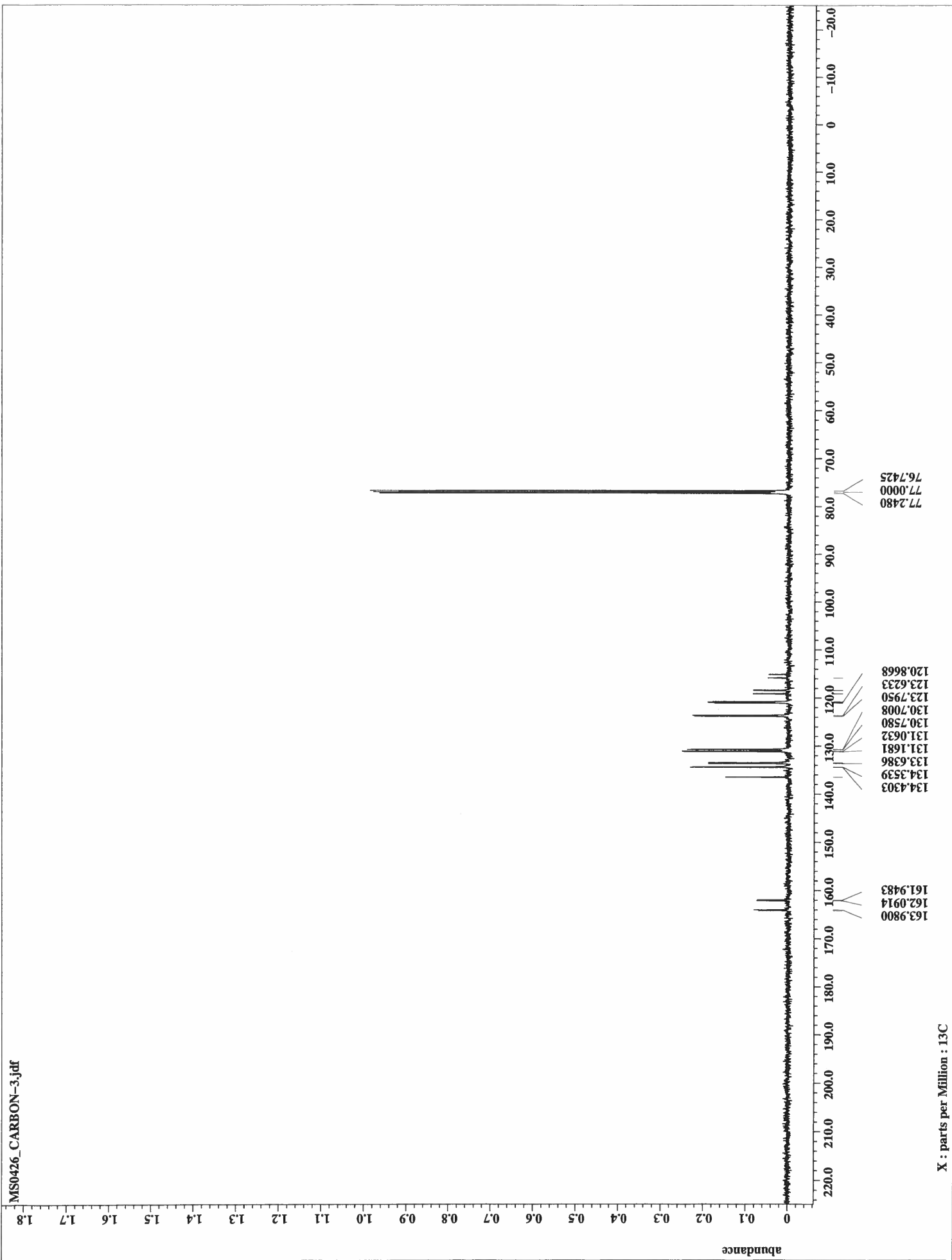
30.0 20.0 10.0 0 -10.0 -20.0 -30.0 -40.0 -50.0 -60.0 -70.0 -80.0 -90.0 -100.0 -110.0 -120.0 -130.0 -140.0 -150.0 -160-170.0

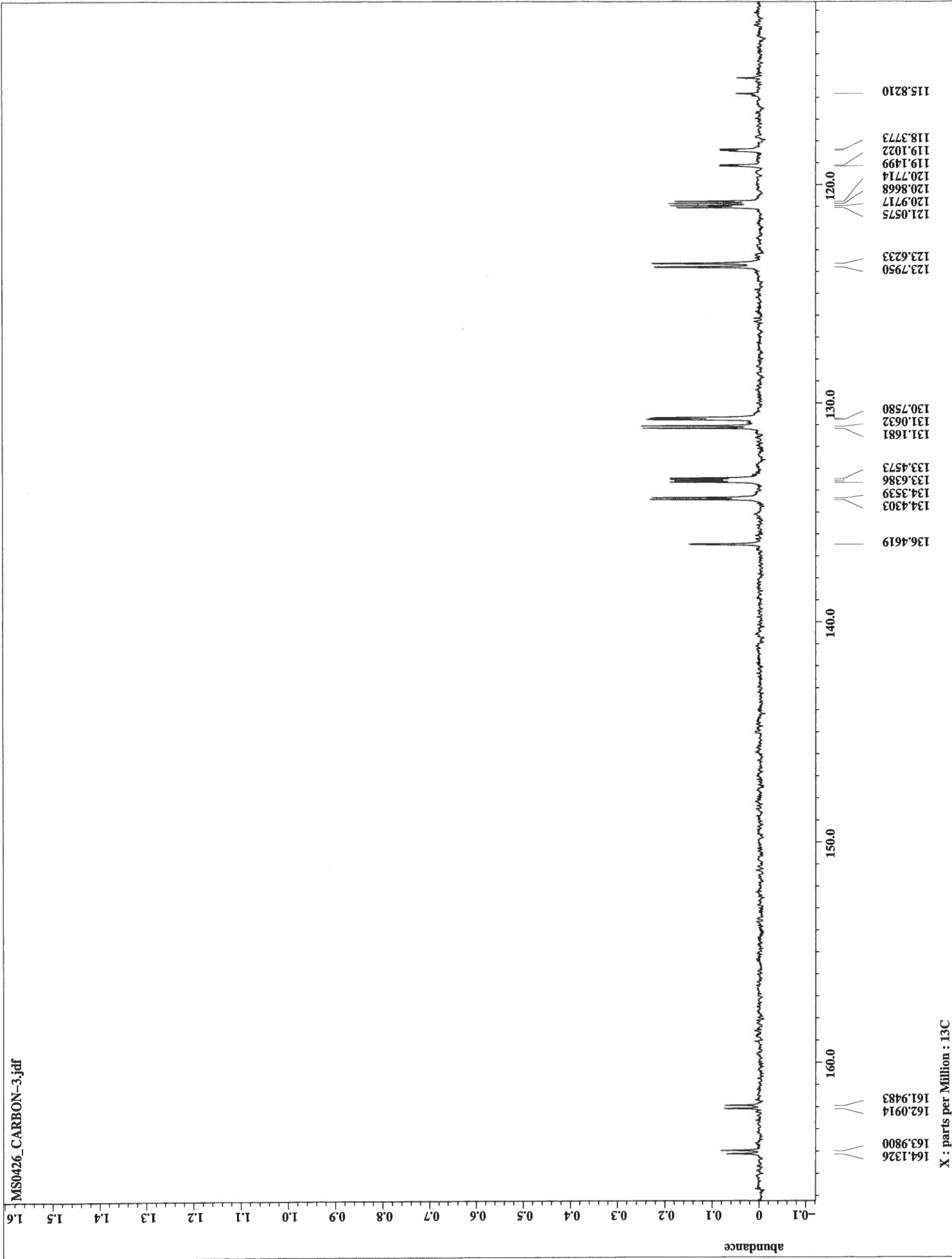
-78.6142

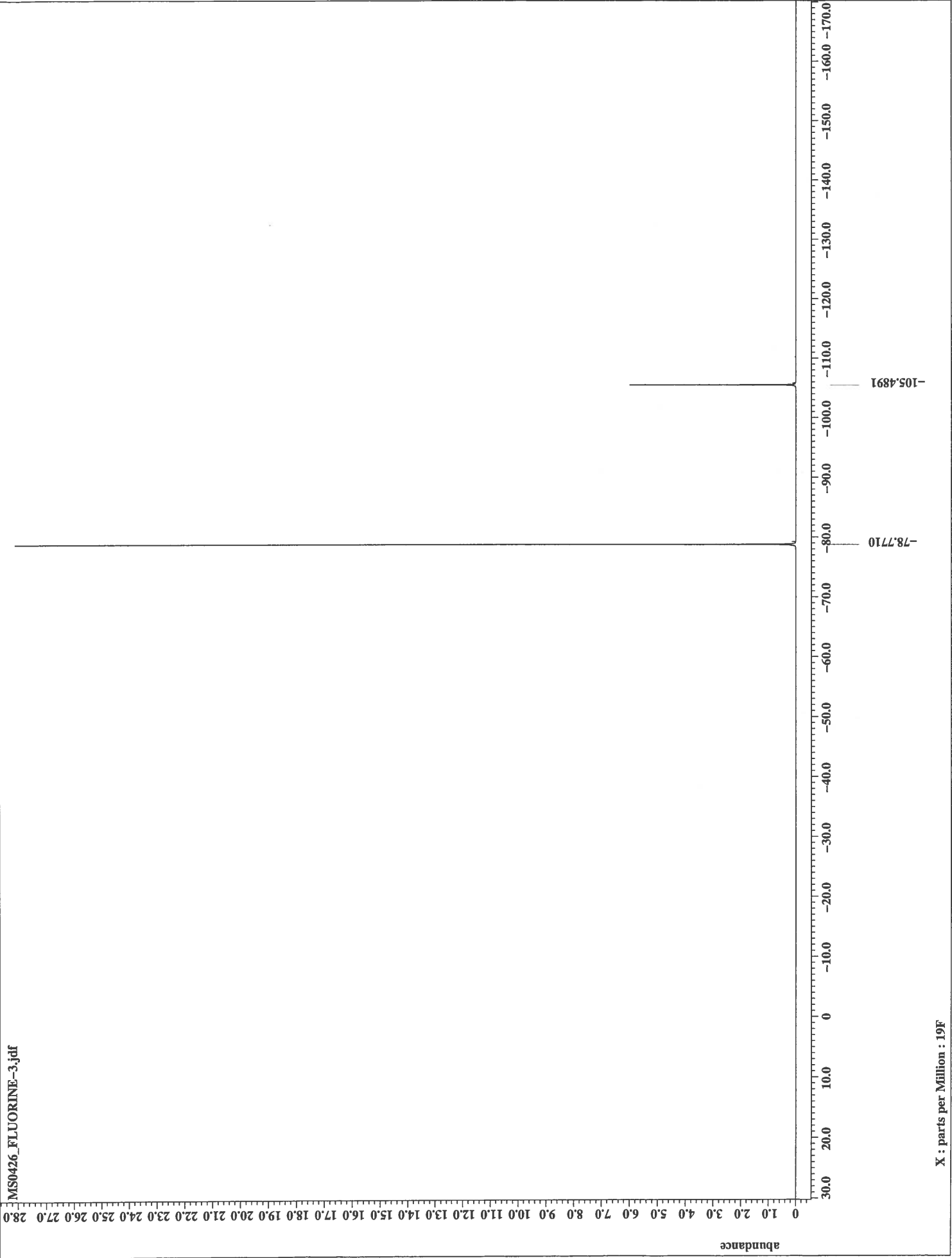
-116.3821

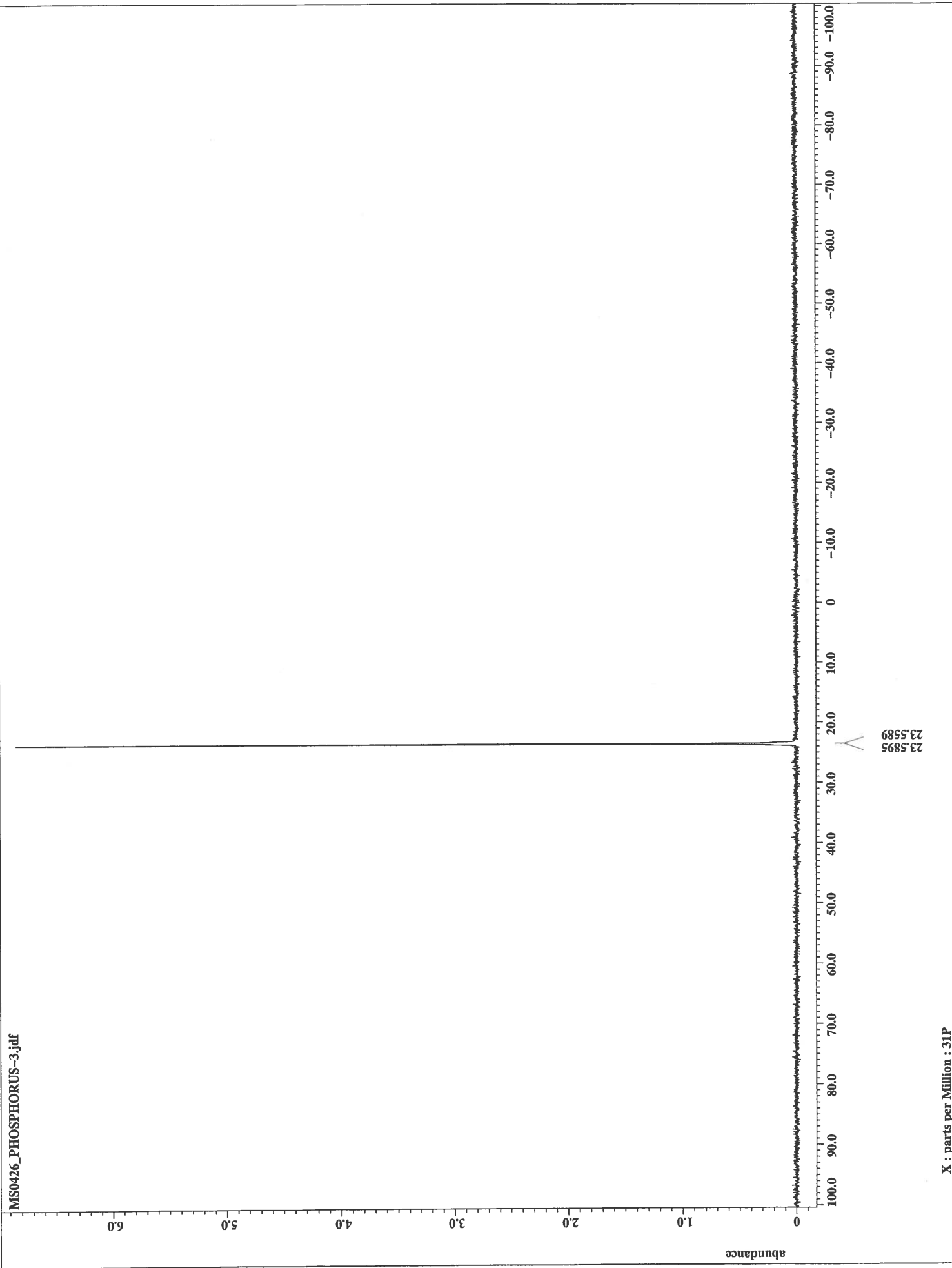




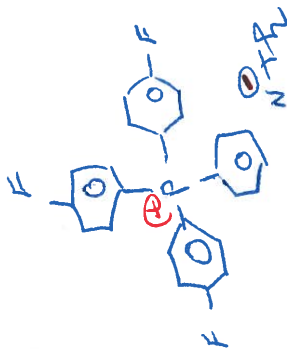








264.57



3.0

2.0

1.0

abundance

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

7.6339
7.6317
7.4198
7.4072
7.4026
7.39004.1686
4.1594
3.9006
3.8915

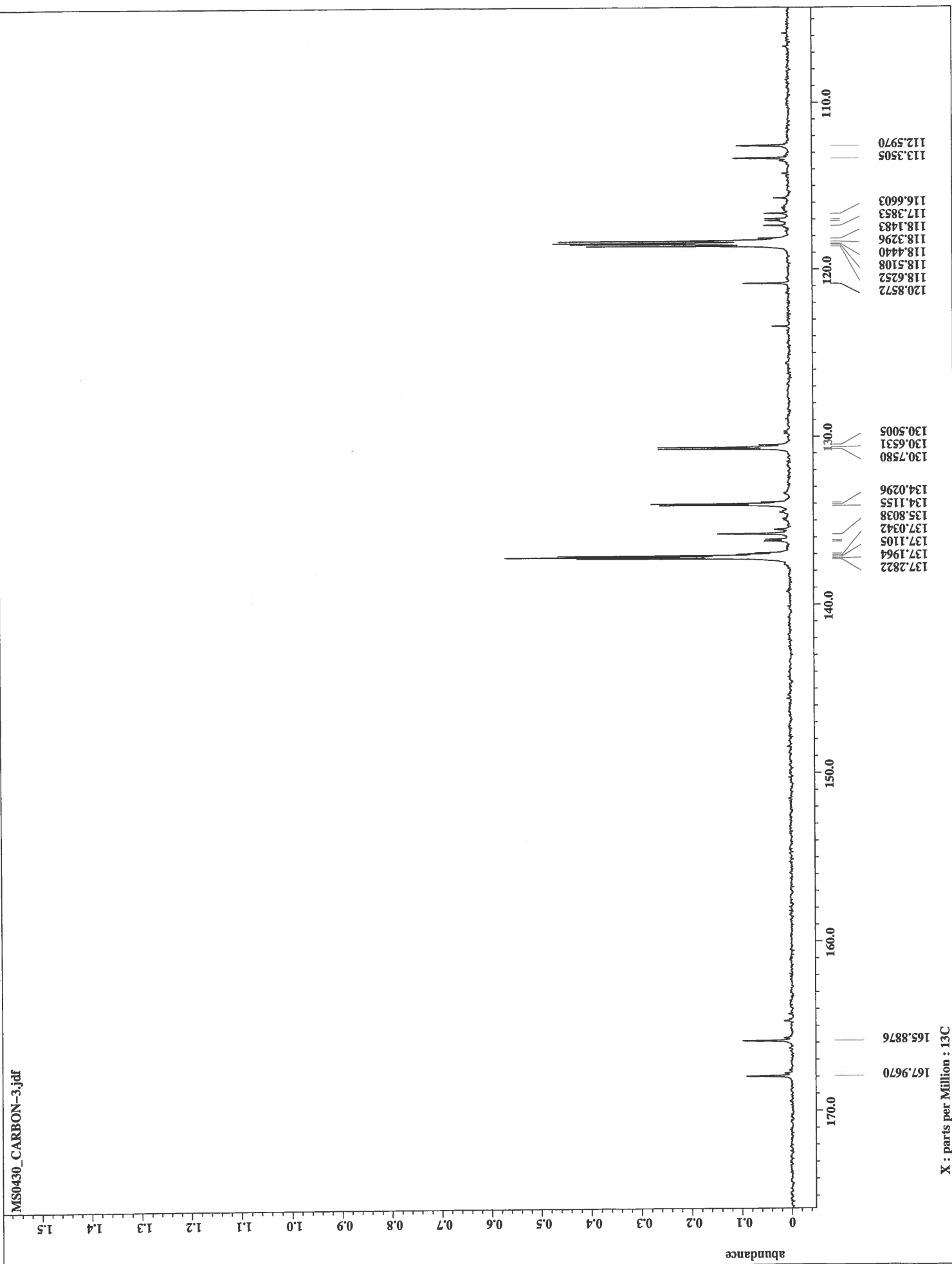
4.21

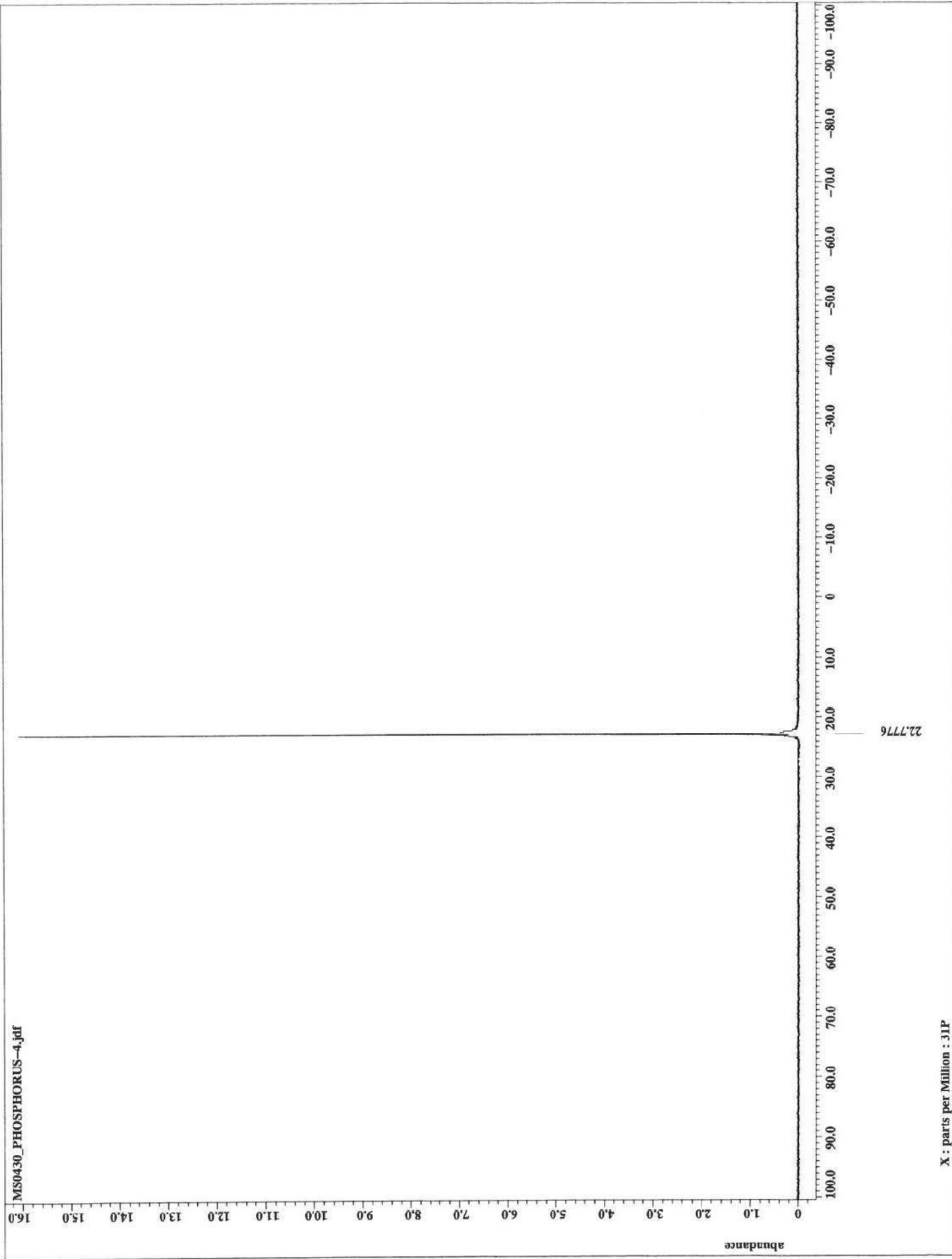
6.22

6.22

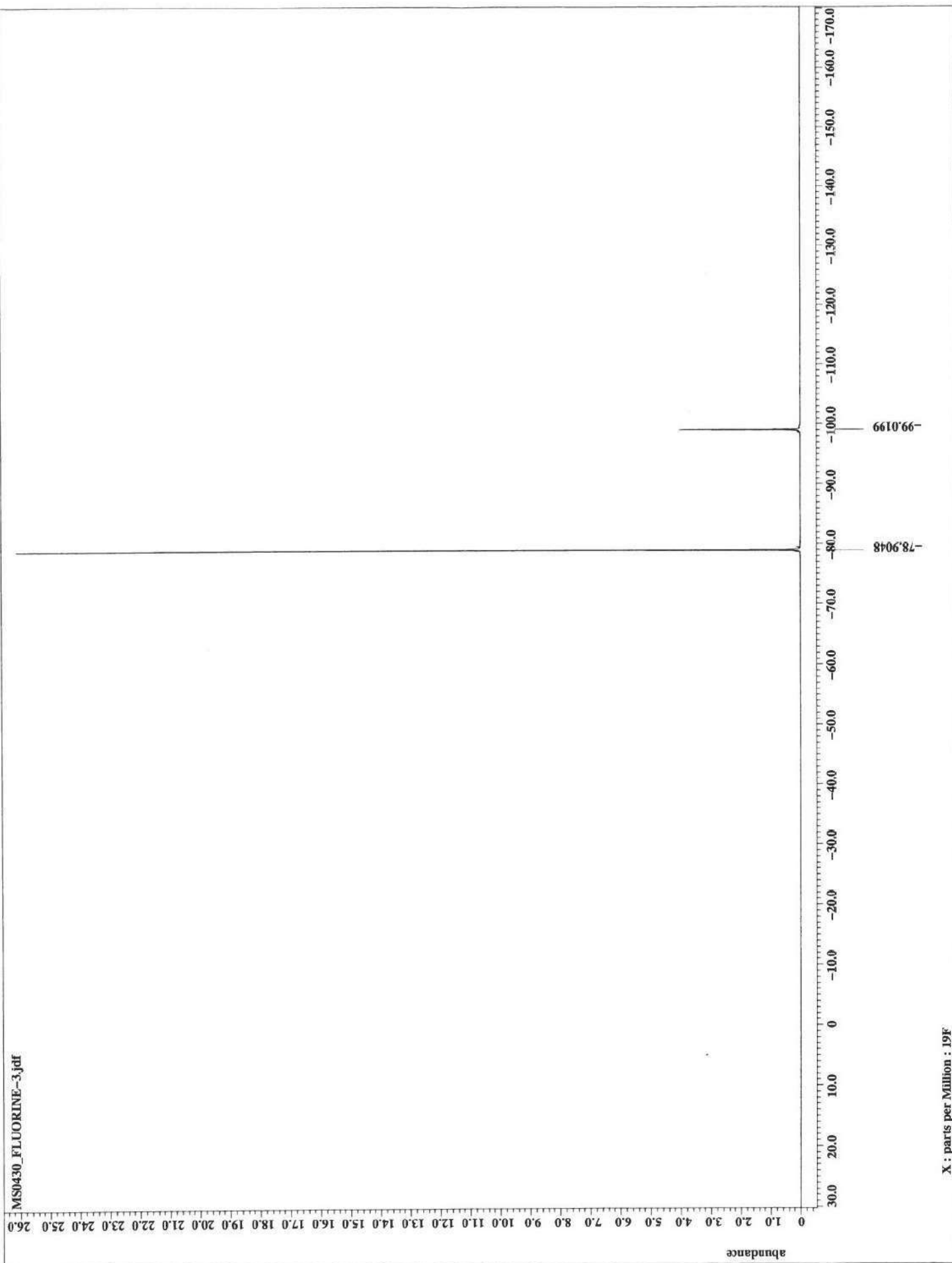


Filename = MS0430_PROTON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0430
 Solvent = CHLOROFORM-D
 Creation_time = 26-NOV-2019 12:03:37
 Revision_time = 26-NOV-2019 11:34:33
 Current_time = 26-NOV-2019 11:34:33
 Data_format = 1D COMPLEX
 Dim_size = 13107
 Dim_title = 1H
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579 [T] (500 [MH
 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_atn = 4 [dB]
 X_pulse = 5.65 [us]
 Irr_mode = Off
 Tri_mode = Off
 Dante_preset = FALSE
 Initial_wait = 1 [s]
 Recvr_gain = 26
 Relaxation_delay = 4 [s]
 Repetition_time = 5.74587904 [s]
 Temp_get = 19.1 [dC]





MS0430_FLUORINE-3.jdf





181.69

79.28

53.26

25.81

3.8147

5.2268

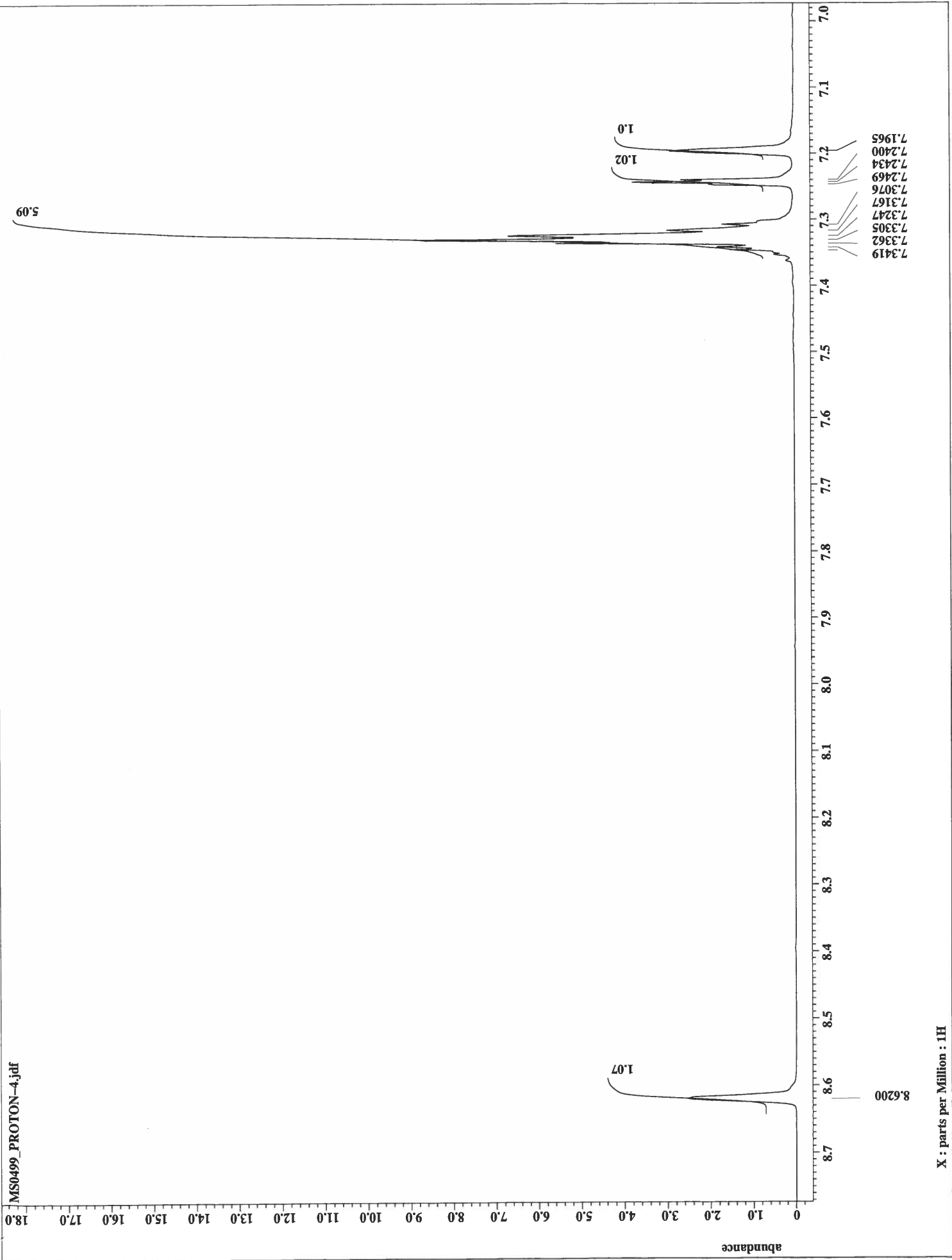
7.3362
7.3305
7.3247

8.6200

X : parts per Million : 1H



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[MH
 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[MHz]
 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_presat = FALSE
 Initial_wait = 1[s]
 Recvr_gain = 24
 Relaxation_delay = 4[s]
 Repetition_time = 5.74587904[s]
 Temp_get = 22.7[dc]




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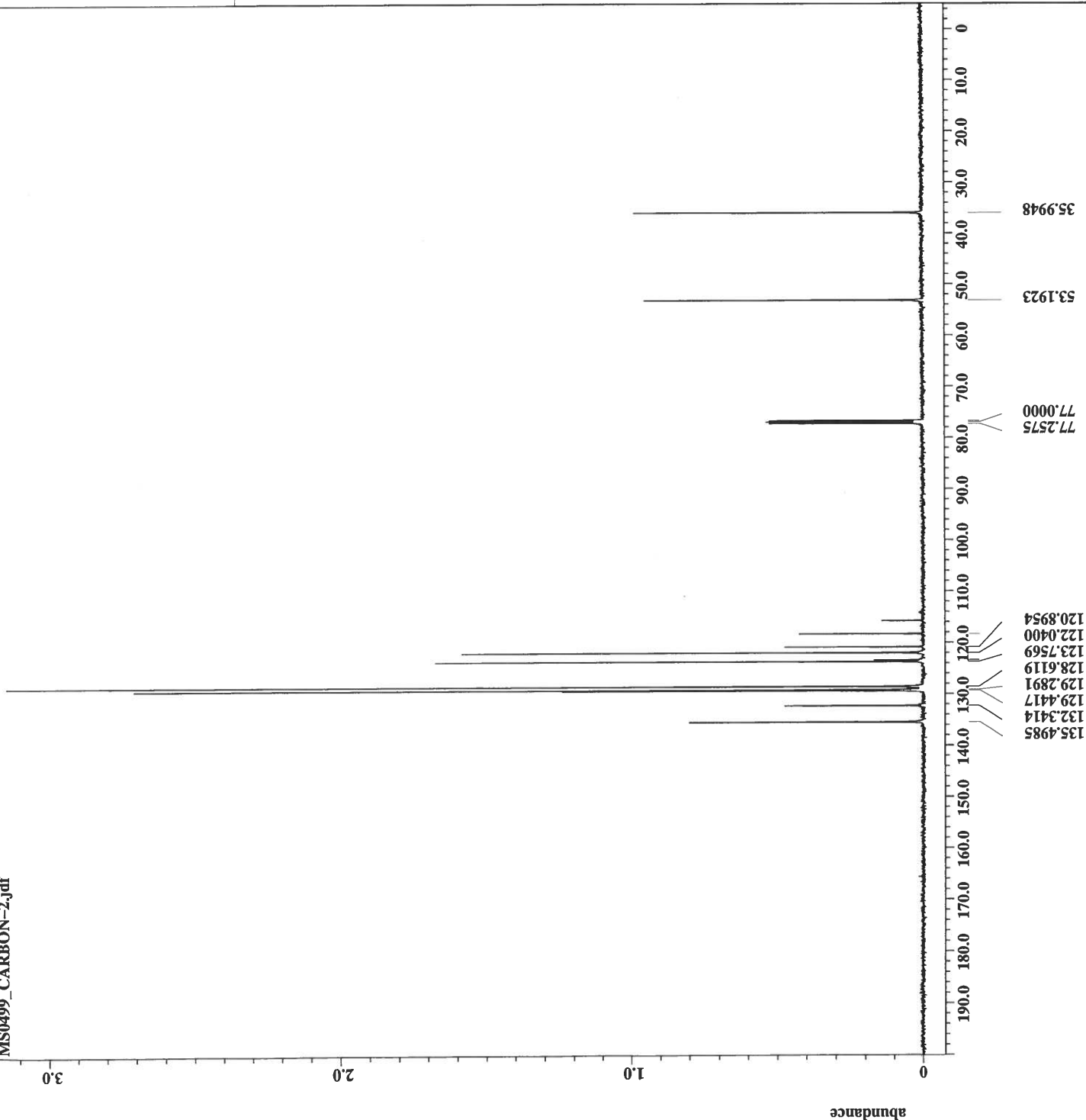
```

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

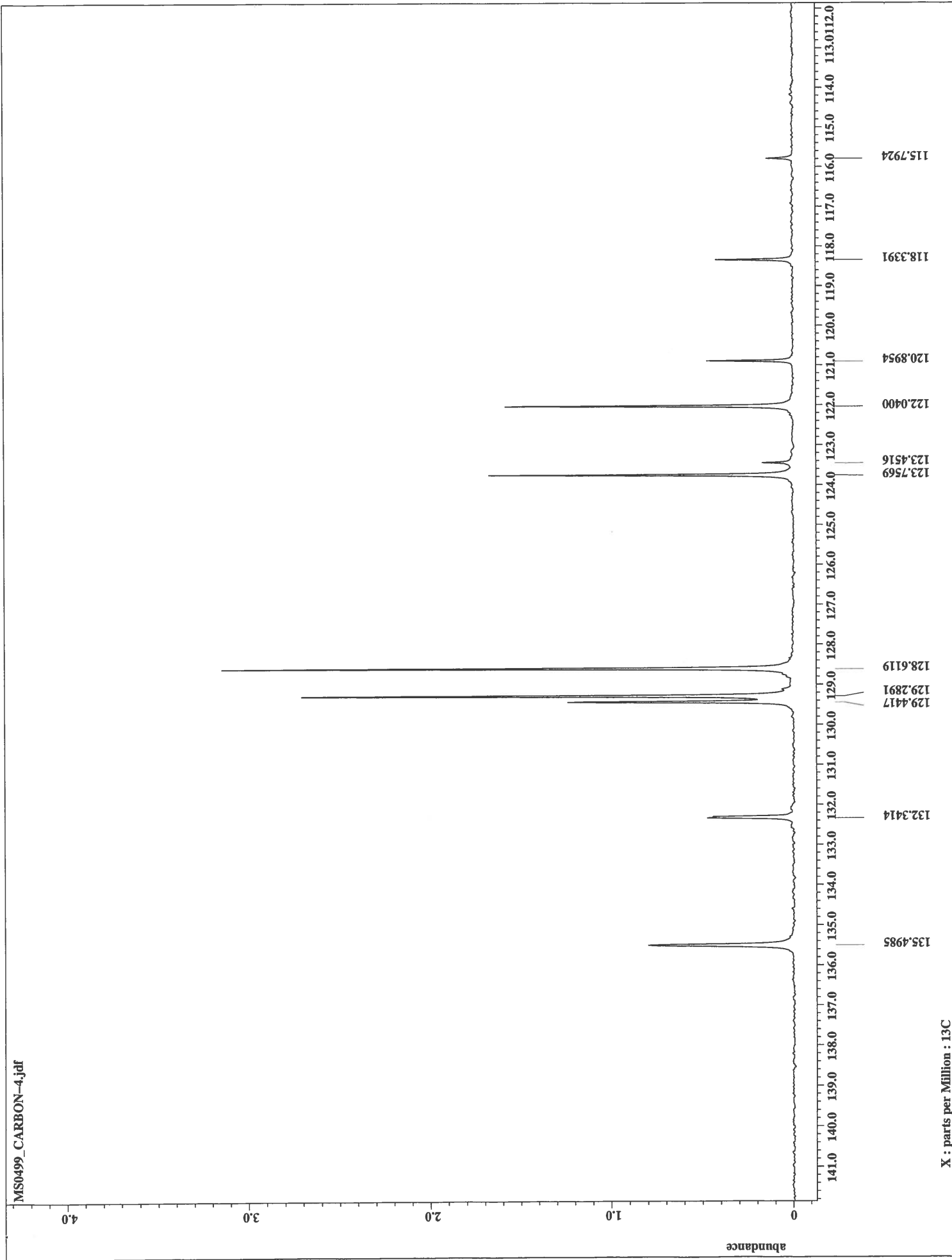
Data_format  = 1D COMPLEX
Dim_size     = 26214
Dim_title    = 13C
Dim_units    = [ppm]
Dimensions   = X
Site         = ECA 500
Spectrometer = JNM-ECA500

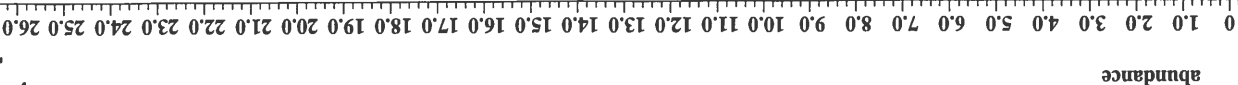
Field_strength = 11.7473579[T] (500[MH
X_acq_duration = 0.83361792[s]
X_domain       = 13C
X_freq         = 125.76529768[MHz]
X_offset       = 100[ppm]
X_points       = 32768
X_prescans     = 4
X_resolution   = 1.19959034[Hz]
X_sweep        = 39.3081761[KHz]
Irr_domain     = 1H
Irr_freq       = 500.15991521[MHz]
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       = 23.2[dc]
  
```



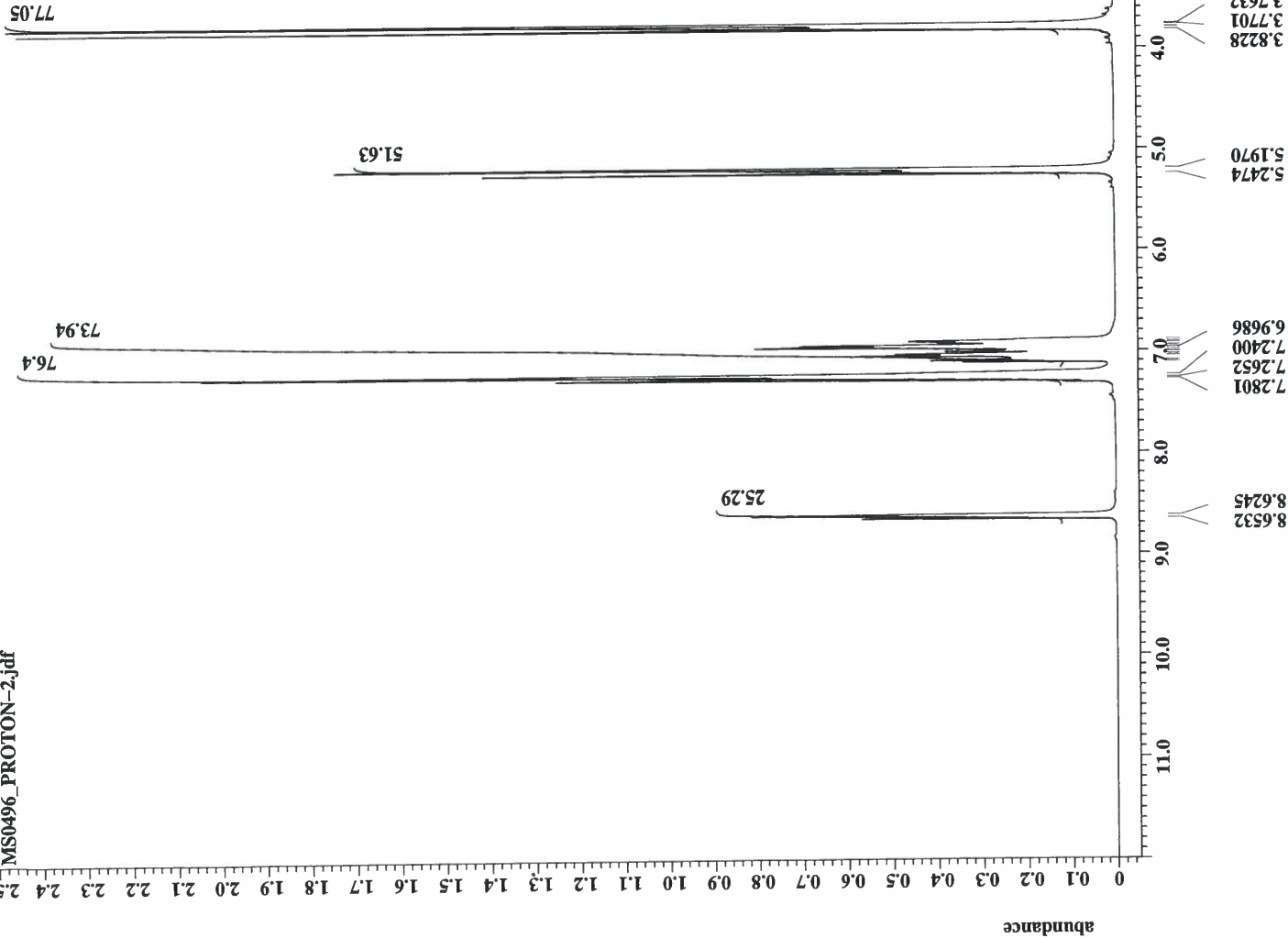
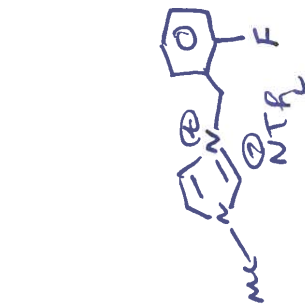
X : parts per Million : 13C





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 = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength = 11.7473579 [T] (500 [MH
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]
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]
Recvr_gain = 30
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.5 [dC]

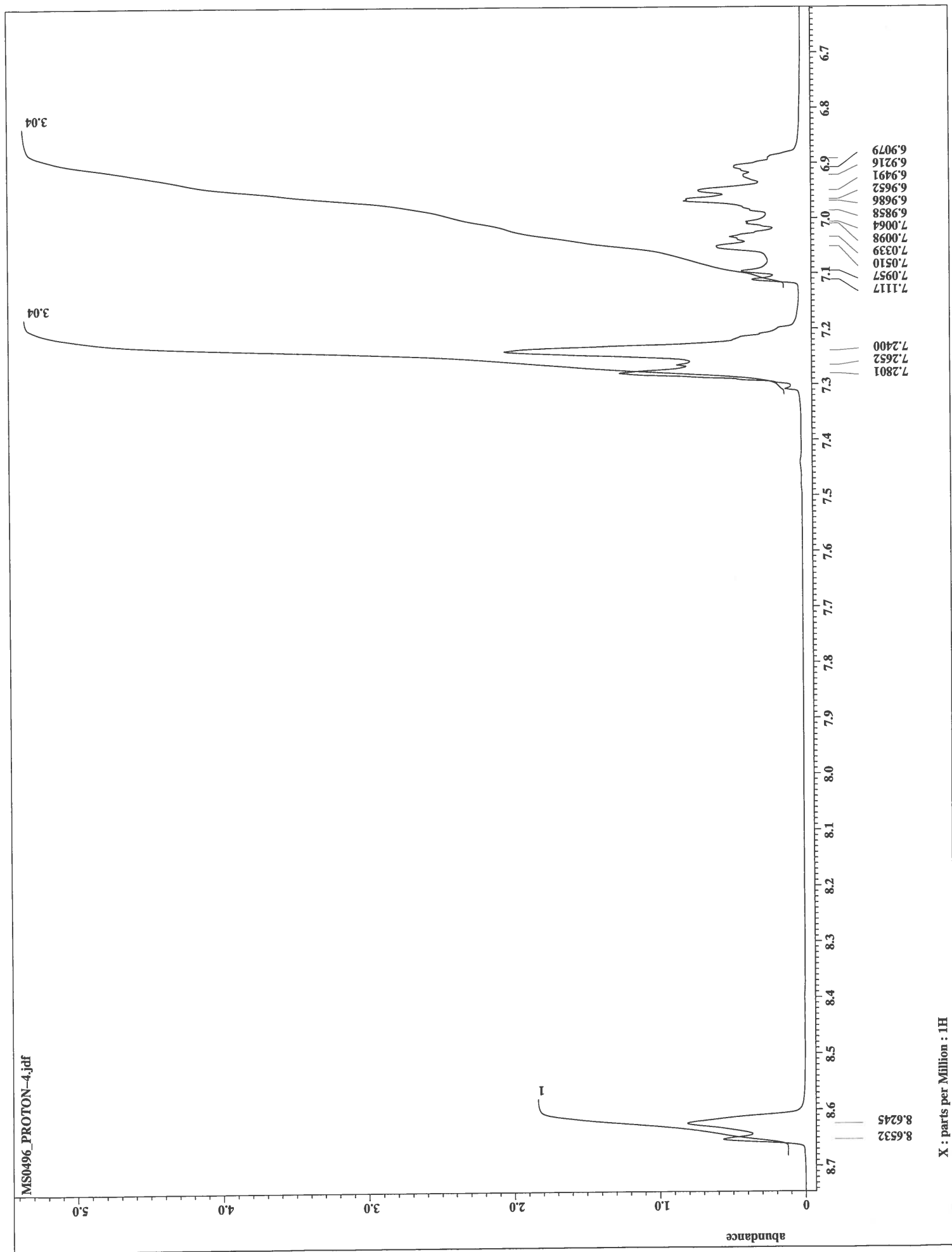
-79.0271



X : parts per Million : 1H



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 [MH
 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
 Pulse_presat = FALSE
 Initial_wait = 1 [s]
 Recvr_gain = 22
 Relaxation_delay = 4 [s]
 Repetition_time = 5.74587904 [s]
 Temp_get = 22.5 [dC]

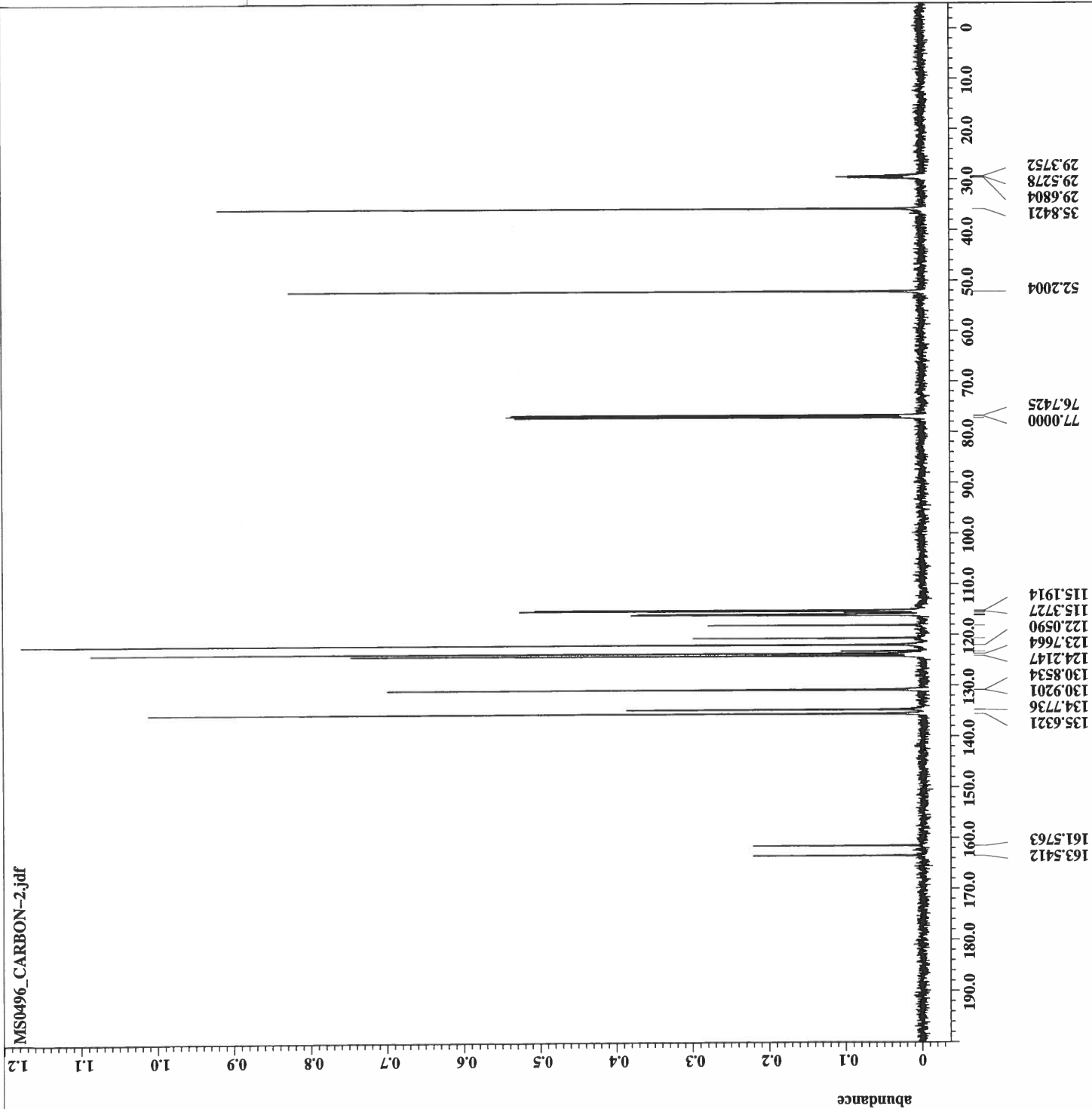


X : parts per Million : 1H

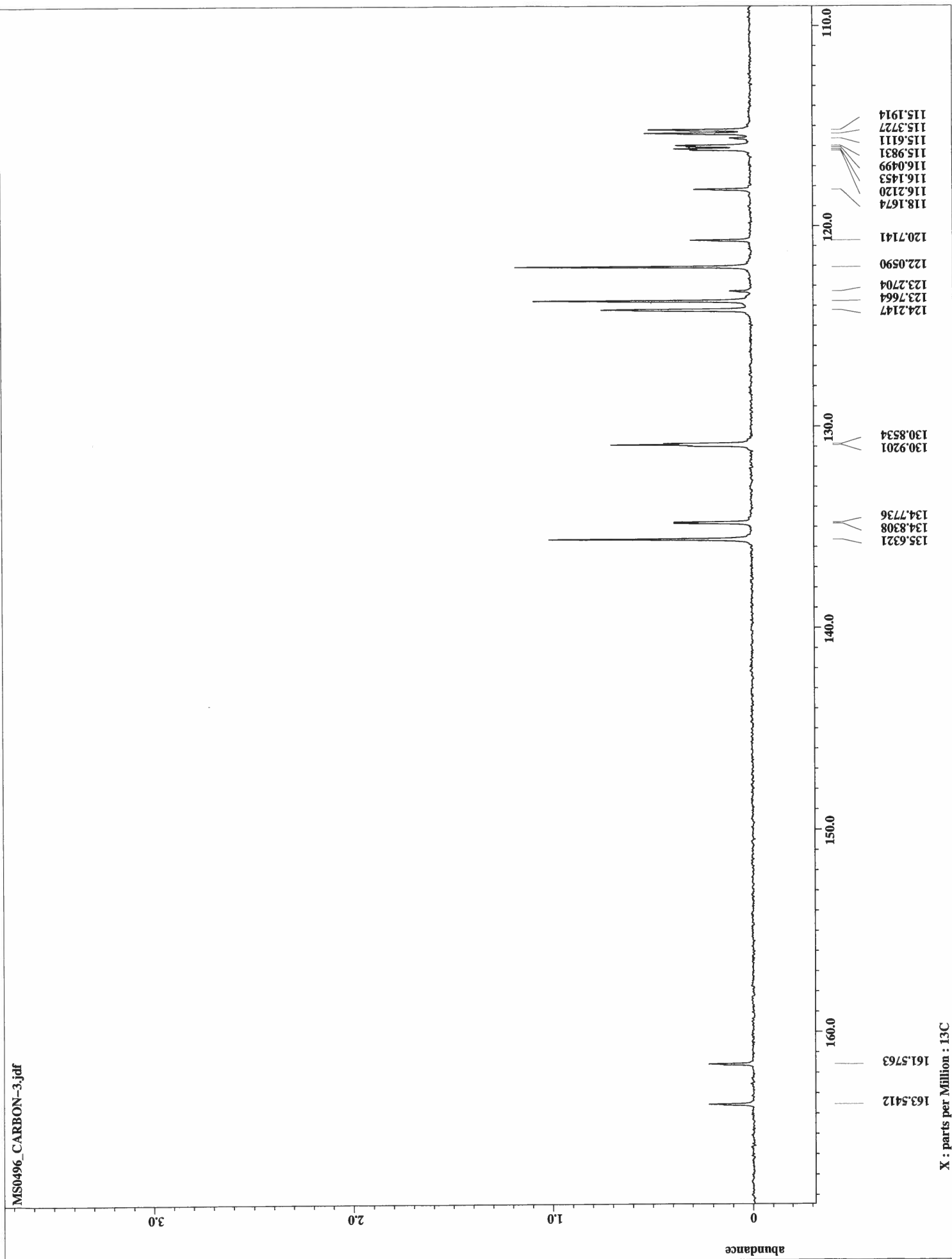


SOUTH ALABAMA
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Filename = MS0496_CARBON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse_dec
 Sample_id = MS0496
 Solvent = CHLOROFORM-D
 Changer_sample = 6
 Creation_time = 12-JUL-2018 17:54:02
 Revision_time = 12-JUL-2018 17:31:38
 Current_time = 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 = JNM-ECA500
 Field_strength = 11.7473579[T] (500[MH
 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_time = TRUE
 Recvr_gain = 2[s]
 Relaxation_delay = 60
 Repetition_time = 2.83361792[s]
 Temp_get = 23[dc]



X : parts per Million : 13C

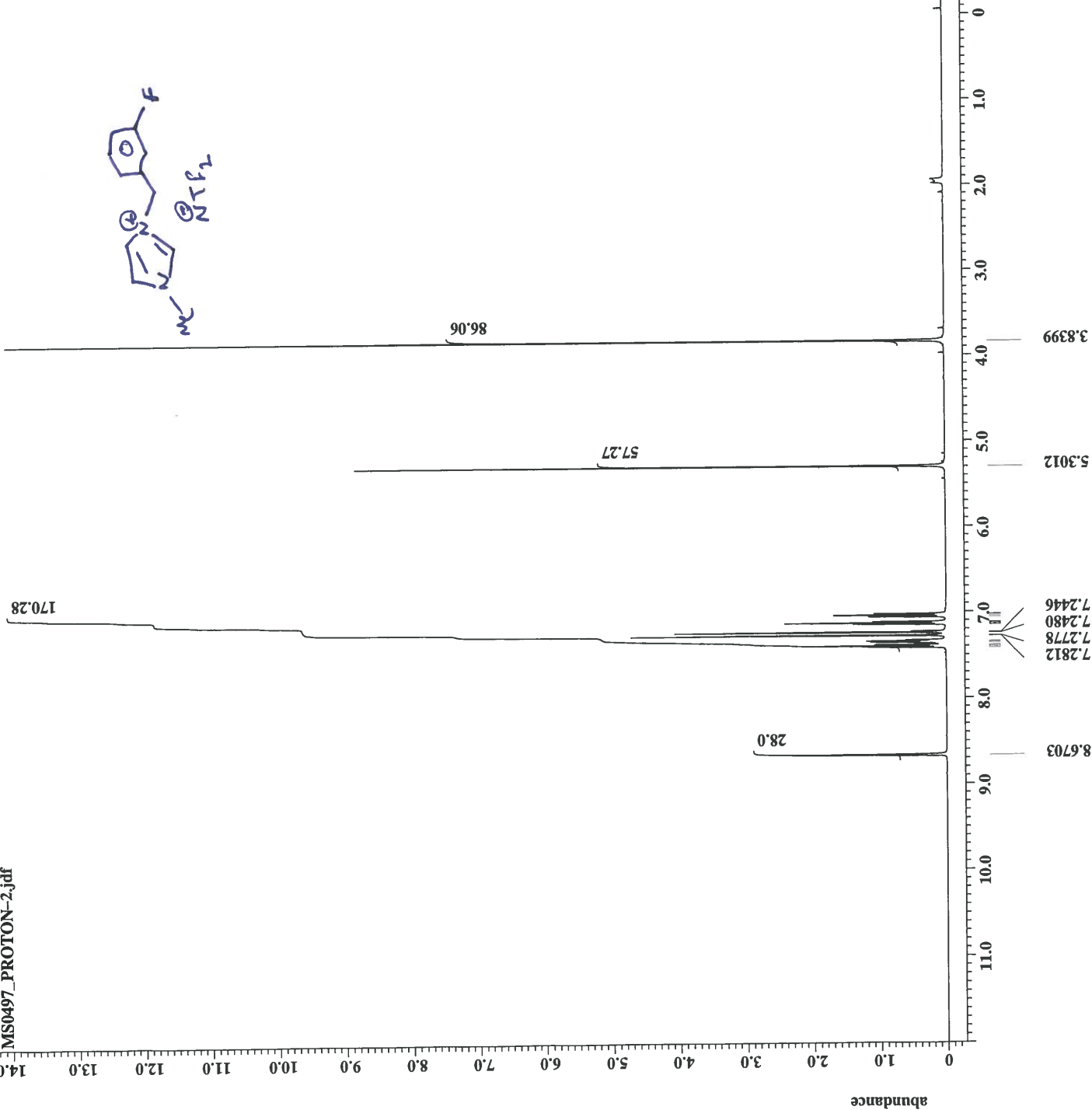
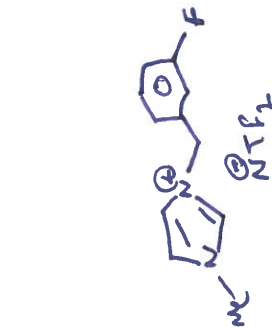




Filename = MS0496_FLUORINE-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0496
 Solvent = CHLOROFORM-D
 Changer_sample = 6
 Creation_time = 12-JUL-2018 16:18:09
 Revision_time = 12-JUL-2018 15:55:46
 Current_time = 12-JUL-2018 15:55:46
 Data_format = 1D COMPLEX
 Dim_size = 52428
 Dim_title = 19F
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579 [T] (500 [MH
 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_presat = FALSE
 Initial_wait = 1 [s]
 Recvr_gain = 26
 Relaxation_delay = 4 [s]
 Repetition_time = 4.55574528 [s]
 Temp_get = 22.5 [dC]

-111.7825

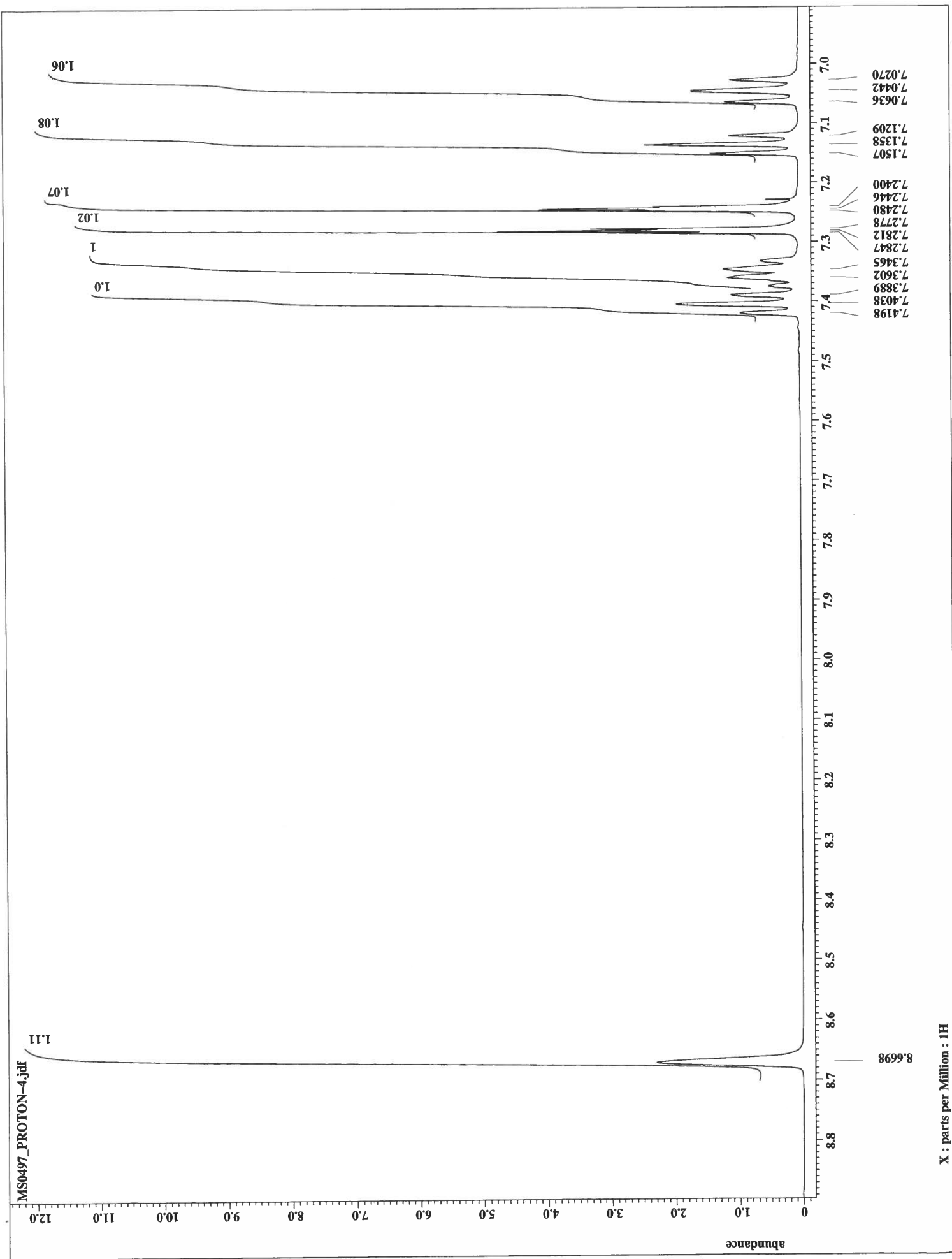
-79.2642
-79.2833

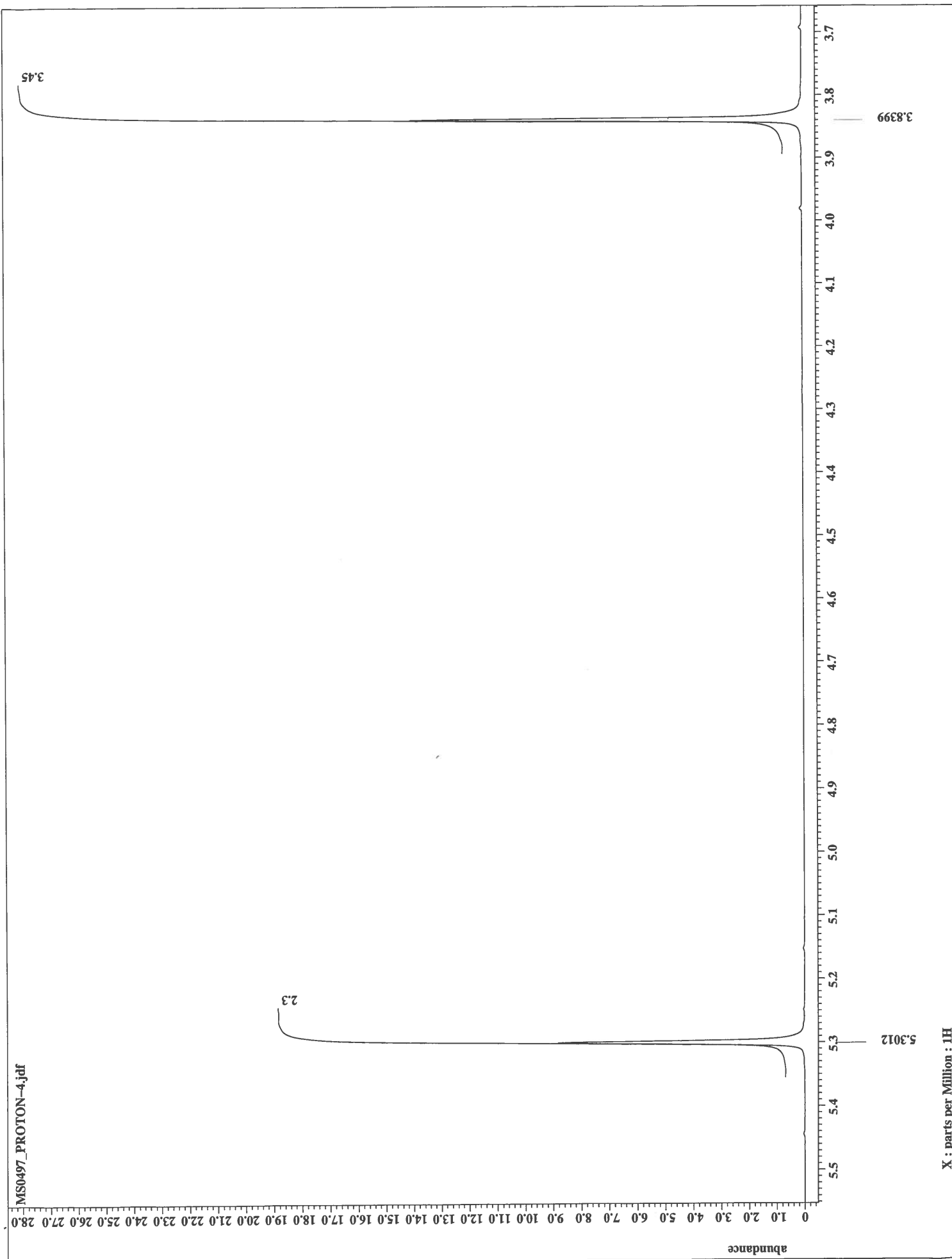


X : parts per Million : 1H



Filename = MS0497_PROTON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0497
 Solvent = CHLOROFORM-D
 Changer_sample = 7
 Creation_time = 12-JUL-2018 18:01:19
 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[MH
 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[MHz]
 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]
 Recvr_gain = 28
 Relaxation_delay = 4[s]
 Repetition_time = 5.74587904[s]
 Temp_get = 22.7[dc]





X : parts per Million : 1H

3.8399

5.3012

2.3

3.45

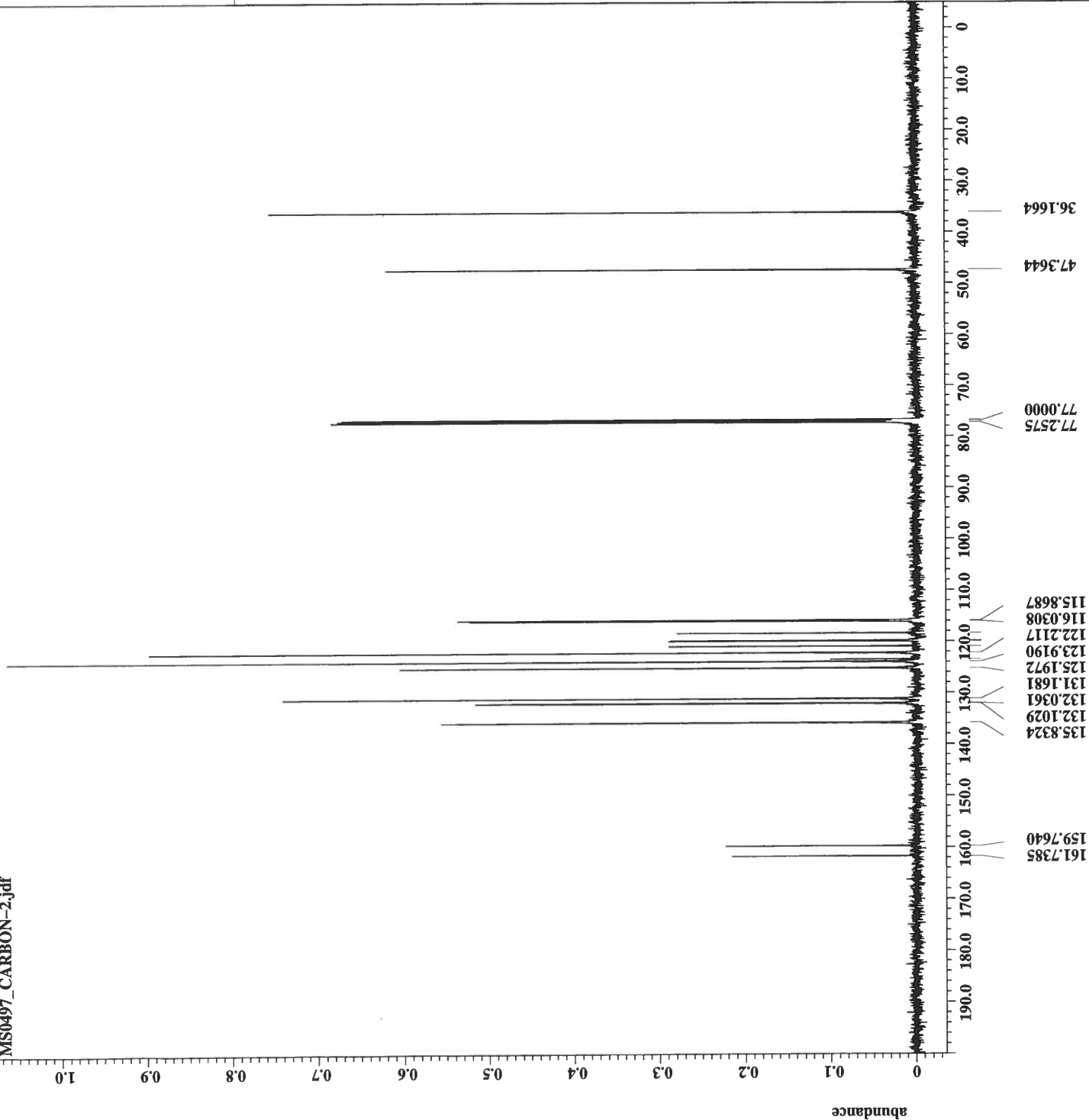
MS0497_PROTON-4.jdf

abundance

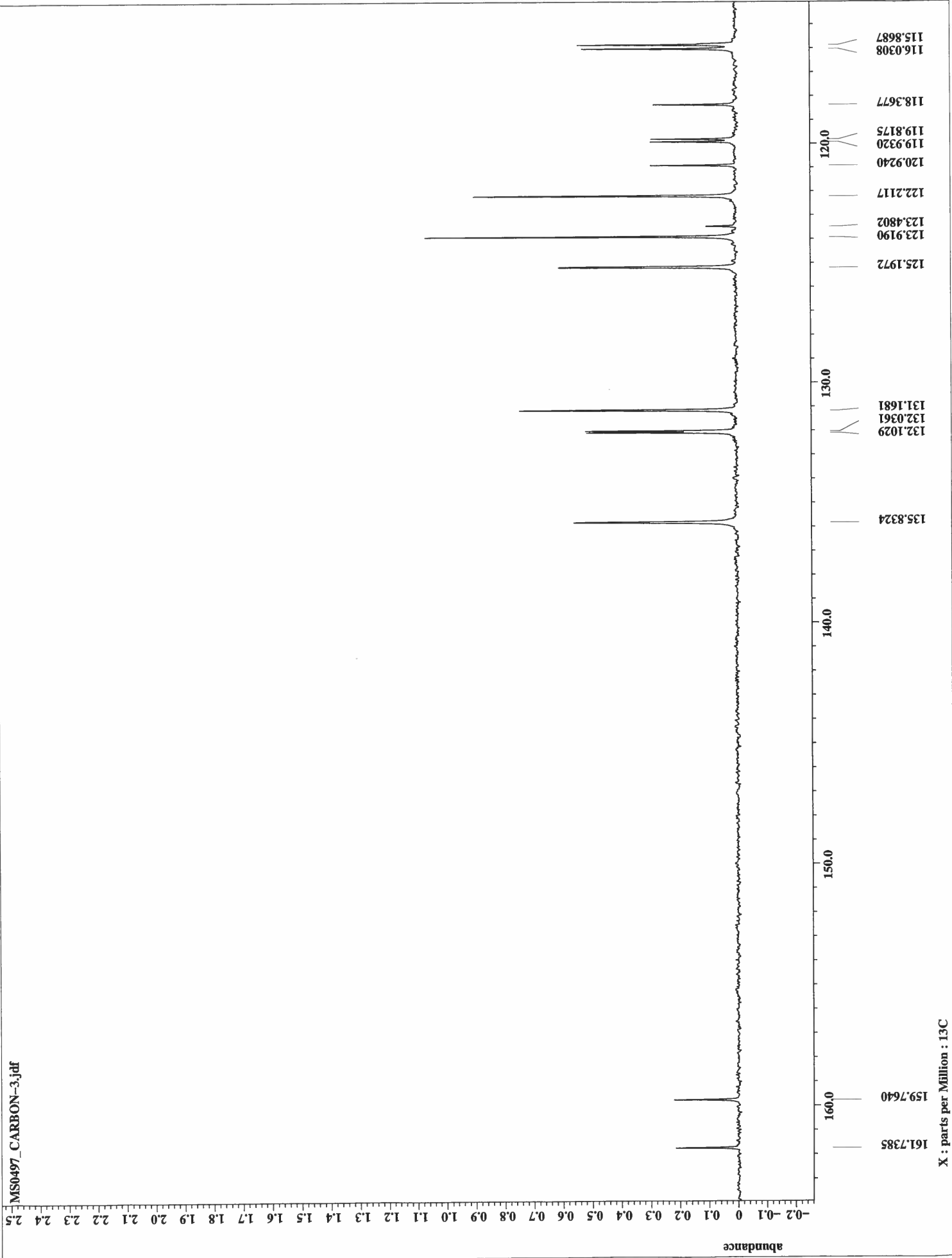


SOUTH ALABAMA
JAGUARS

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 = UNM-ECA500
 Field_strength = 11.7473579[T] (500[MH
 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_time = TRUE
 Noe = 2[s]
 Recvr_gain = 60
 Relaxation_delay = 2[s]
 Repetition_time = 2.83361792[s]
 Temp_get = 23.1[dc]

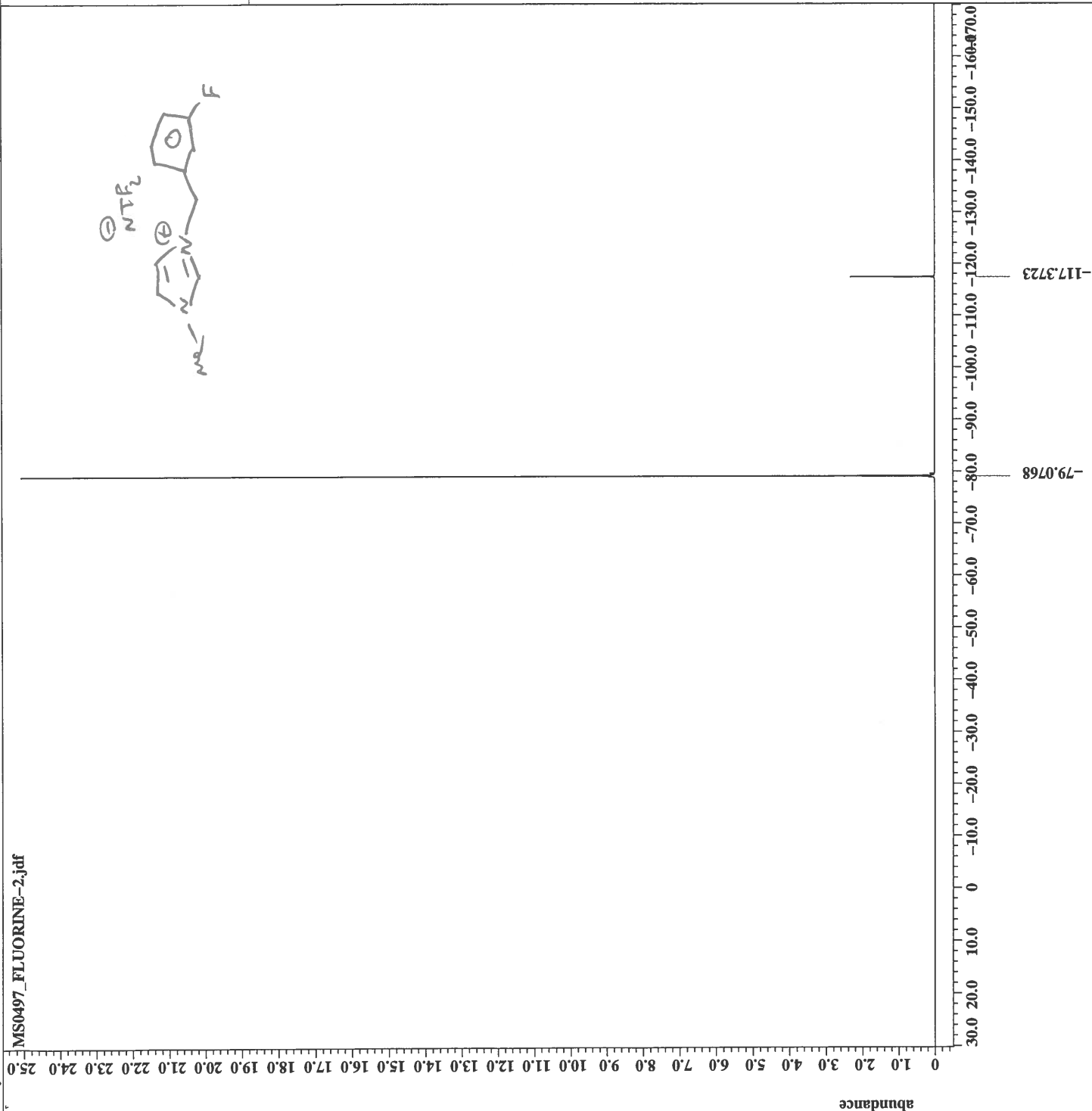


X : parts per Million : 13C





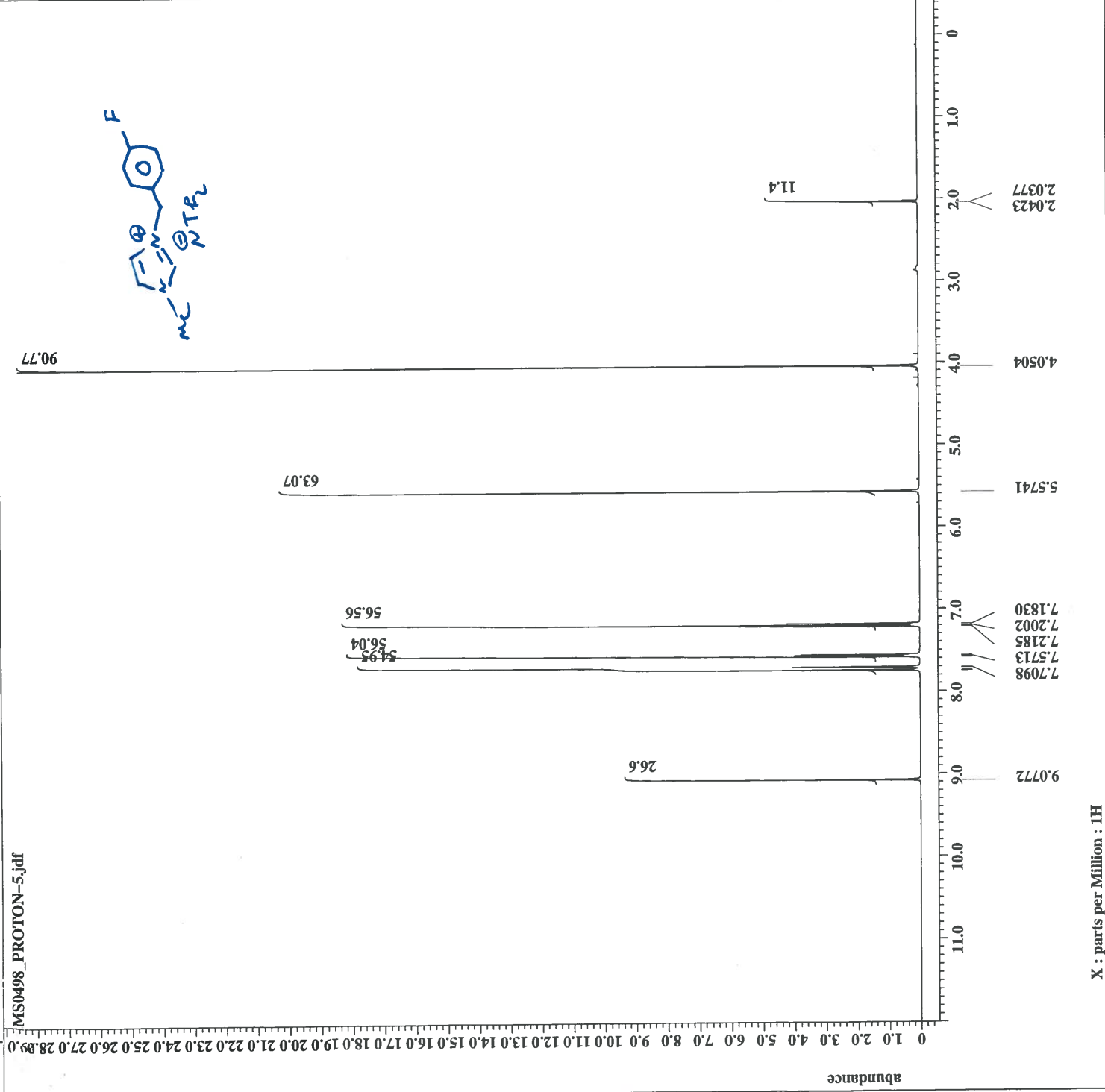
Filename = MS0497_FLUORINE-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0497
 Solvent = CHLOROFORM-D
 Changer_sample = 7
 Creation_time = 12-JUL-2018 16:25:02
 Revision_time = 12-JUL-2018 16:02:38
 Current_time = 12-JUL-2018 16:02:38
 Data_format = 1D COMPLEX
 Dim_size = 52428
 Dim_title = 19F
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579 [T] (500 [MH
 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_presat = FALSE
 Initial_wait = 1 [s]
 Recvr_gain = 30
 Relaxation_delay = 4 [s]
 Repetition_time = 4.55574528 [s]
 Temp_get = 22.4 [dC]

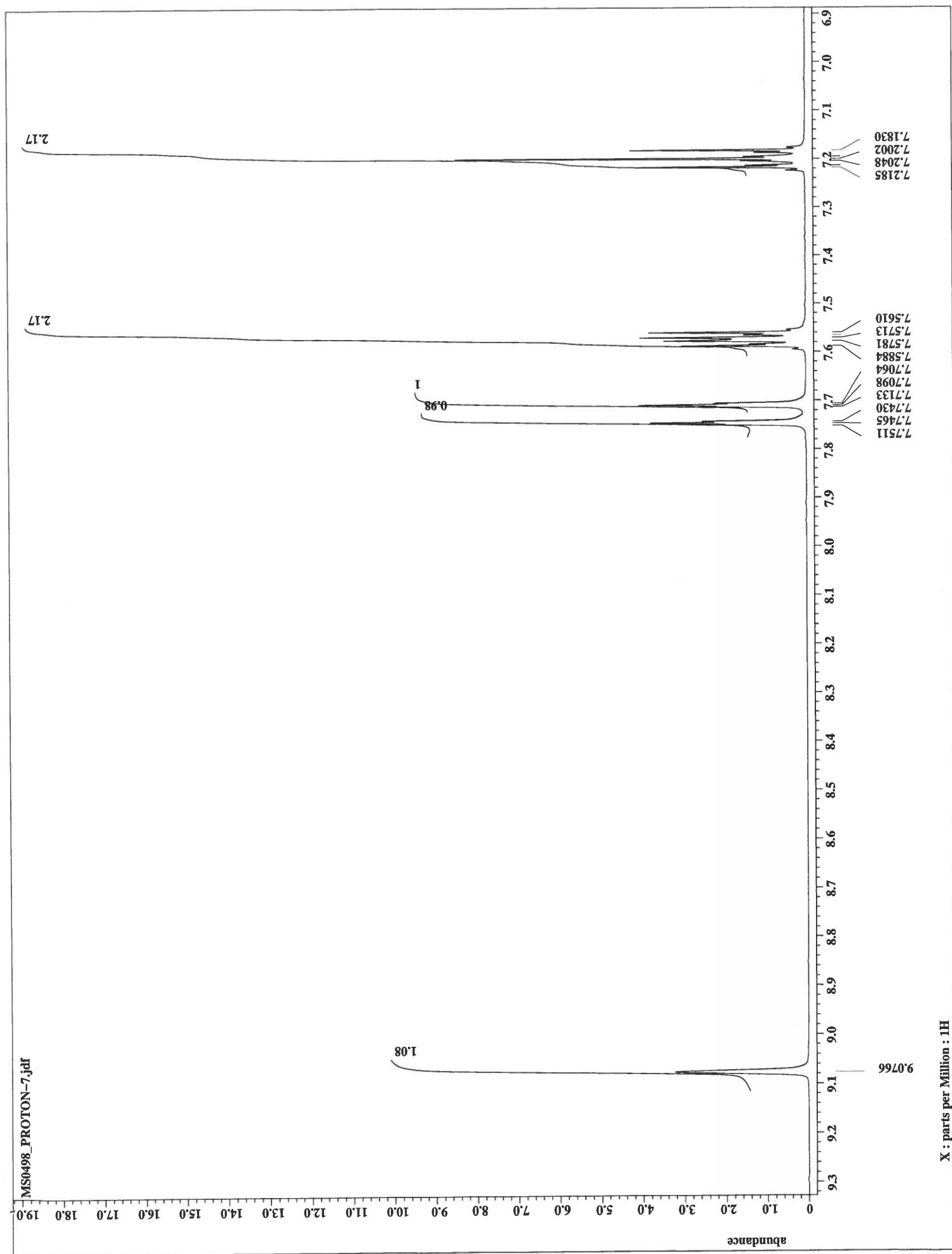


90.77



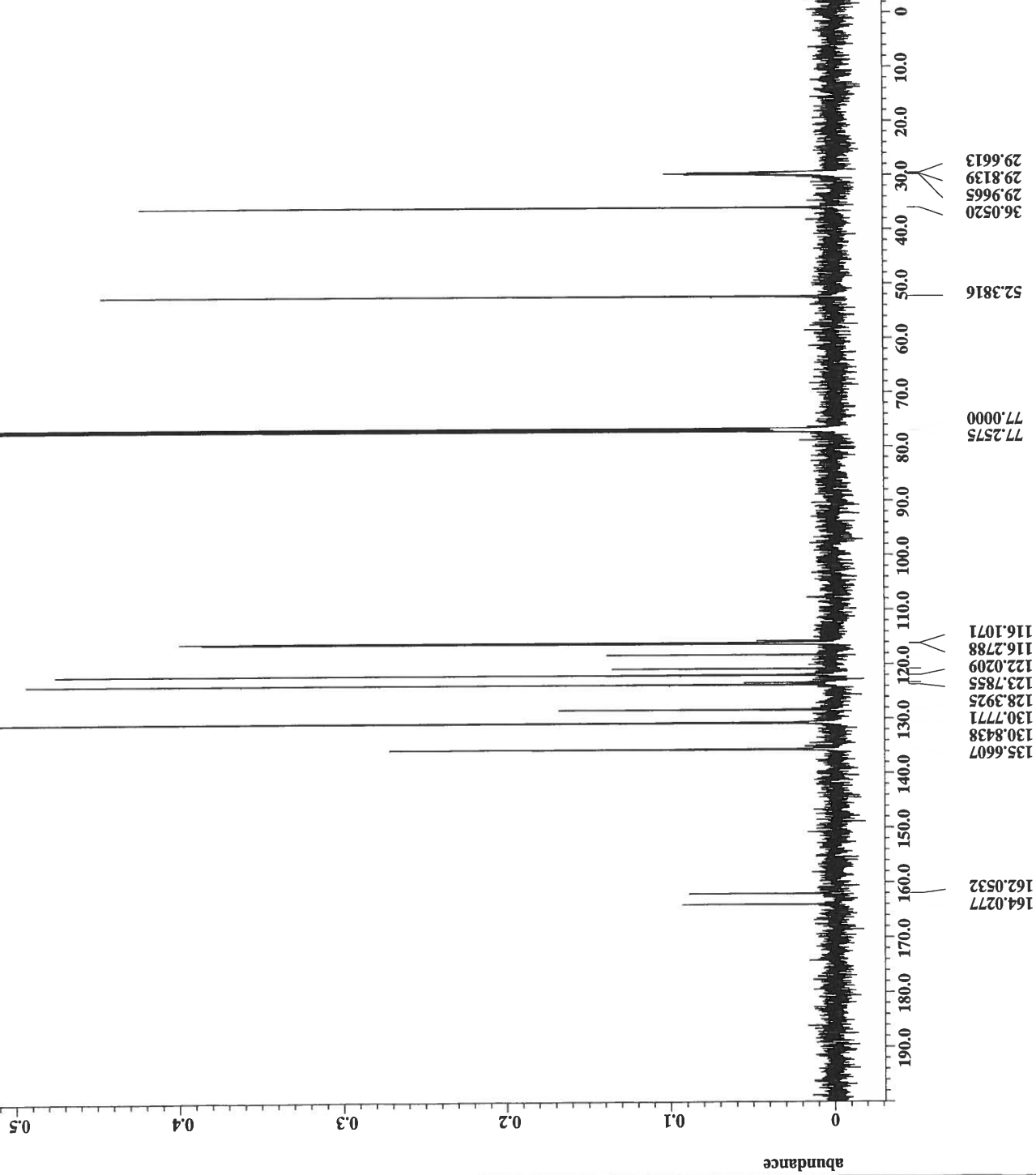
Filename = MS0498_PROTON-5.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0498
 Solvent = ACETONE-D6
 Changer_sample = 8
 Creation_time = 13-JUL-2018 09:18:13
 Revision_time = 13-JUL-2018 08:55:45
 Current_time = 13-JUL-2018 08:55:45
 Data_format = 1D COMPLEX
 Dim_size = 13107
 Dim_title = 1H
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579 [T] (500 [MH
 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]
 Recvr_gain = 34
 Relaxation_delay = 4 [s]
 Repetition_time = 5.74587904 [s]
 Temp_get = 22.5 [dC]



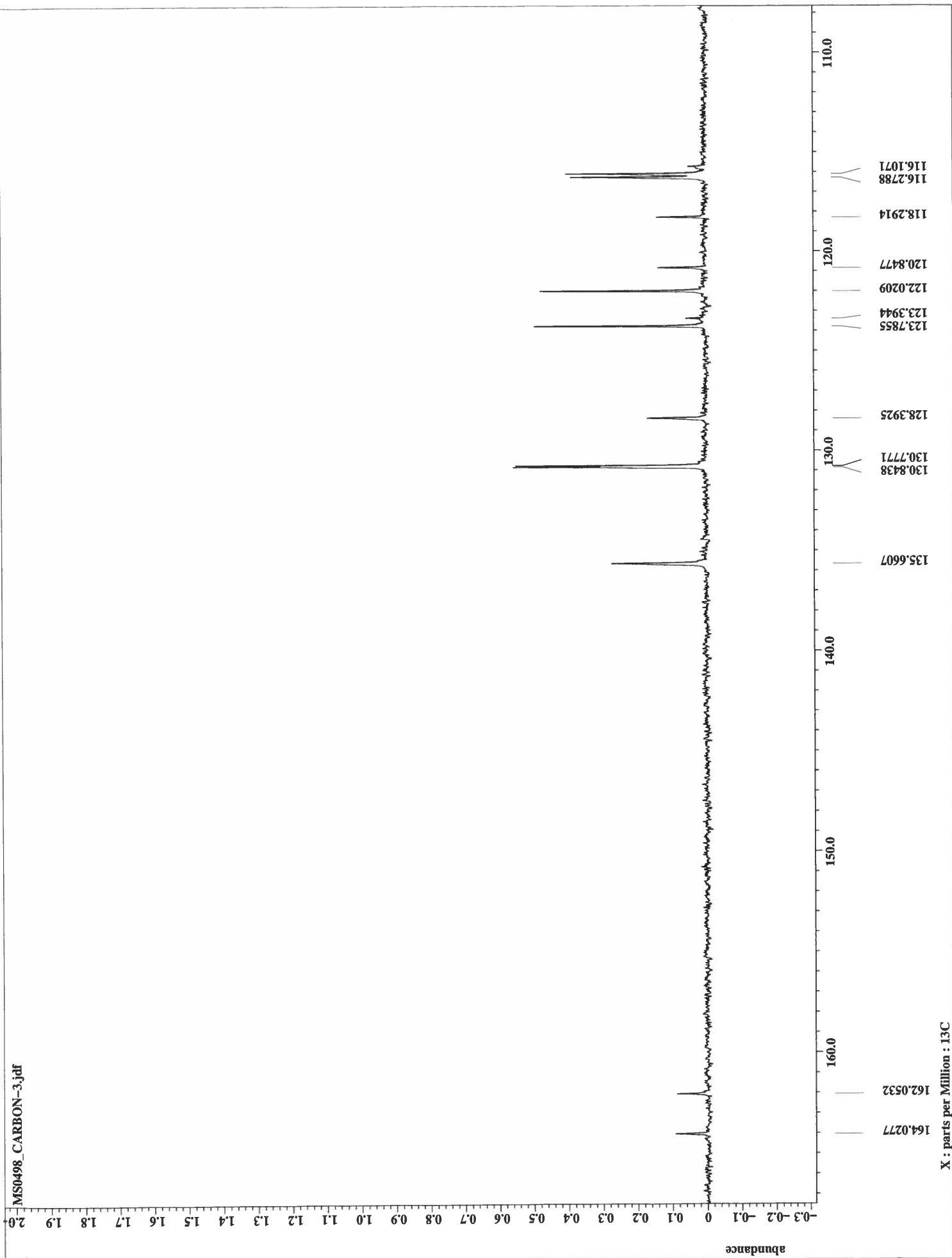


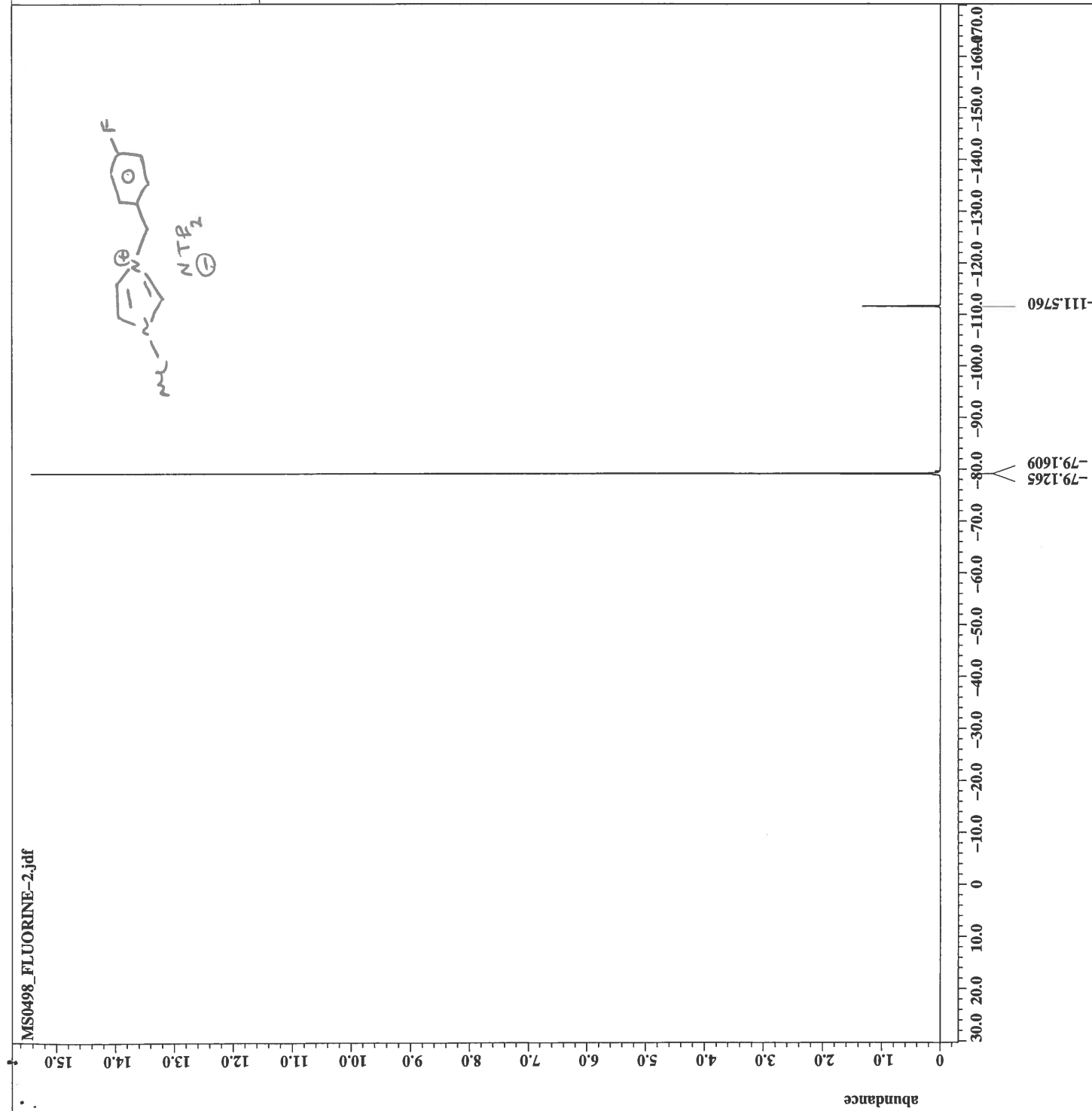
**SOUTH ALABAMA**
JAGUARS

Filename = MS0498_CARBON-2.jdf
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0498
Solvent = CHLOROFORM-D
Charger_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 = 26214
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength = 11.7473579[T] (500[MH
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 = 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]
Recvr_gain = 60
Relaxation_delay = 2[s]
Repetition_time = 2.83361792[s]
Temp_get = 23[dc]



X : parts per Million : 13C





X : parts per Million : 19F



```

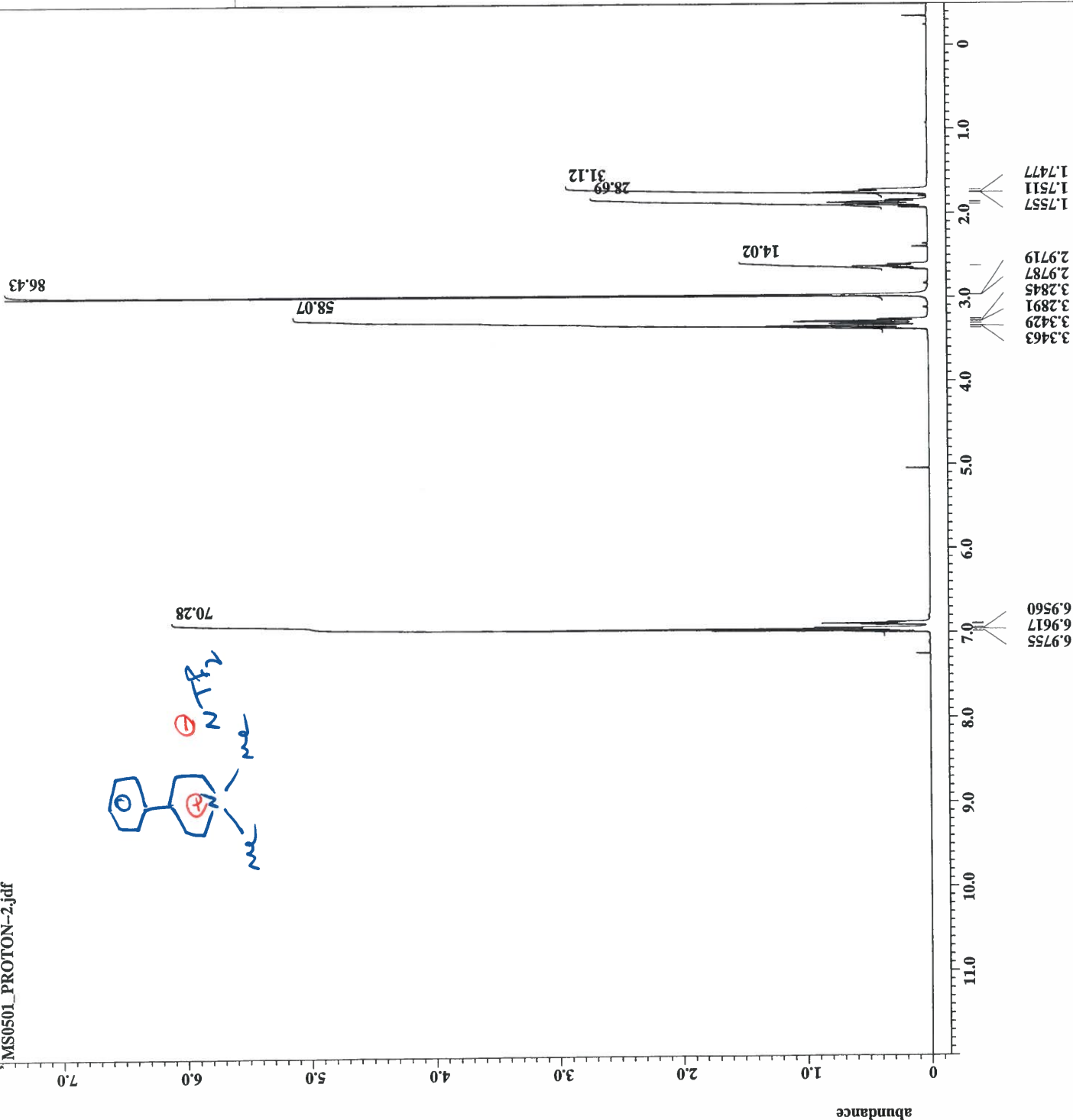
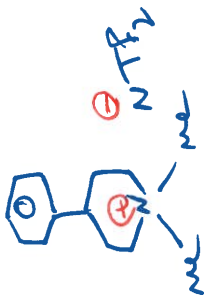
Filename = MS0498_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0498
Solvent = CHLOROFORM-D
Changer_sample = 8
Creation_time = 12-JUL-2018 16:31:45
Revision_time = 12-JUL-2018 16:09:22
Current_time = 12-JUL-2018 16:09:23

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[T] (500[MH
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]
Recvr_gain = 36
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get = 22.4[dc]

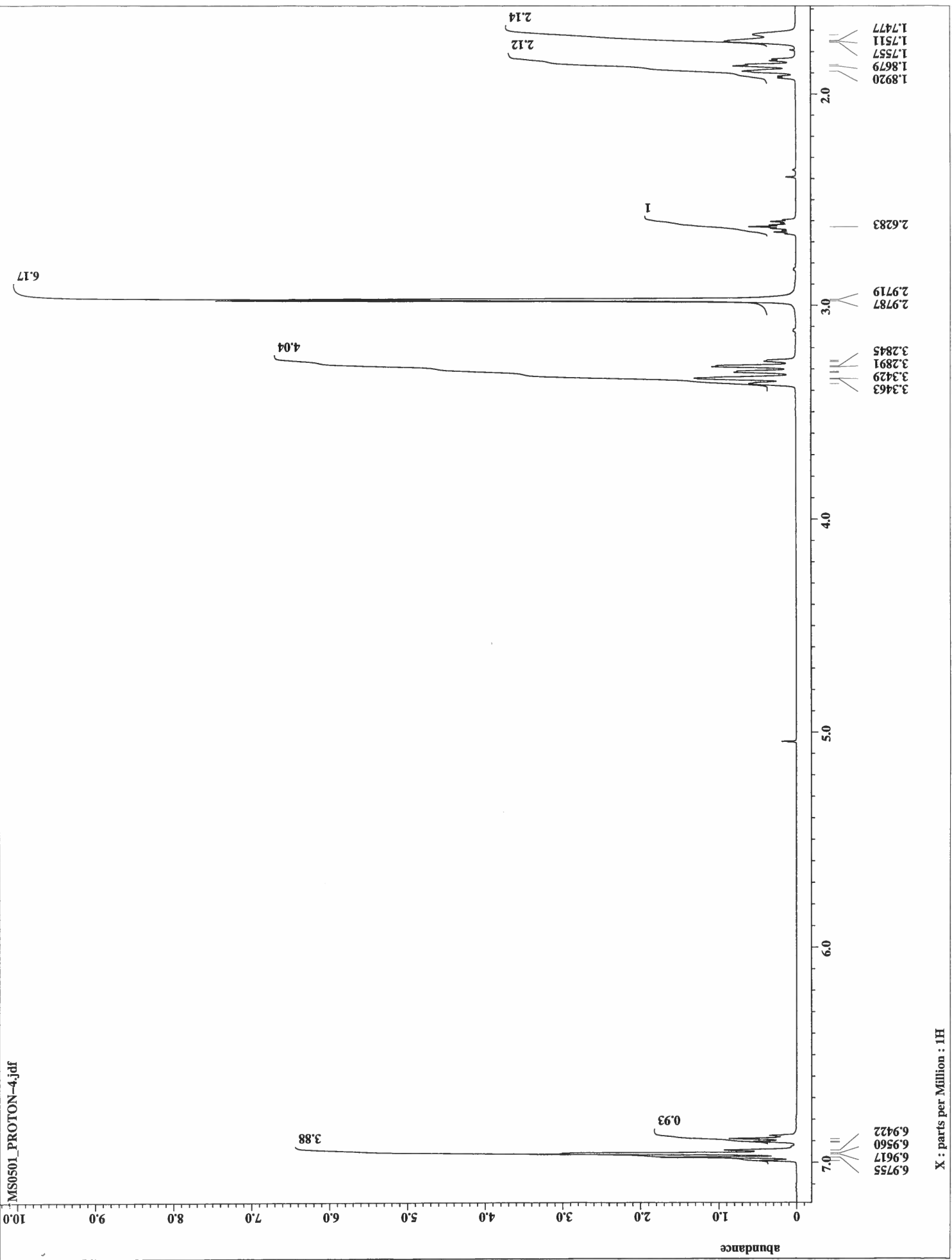
```



X : parts per Million : 1H



Filename	=	MS0501_PROTON-2.jdf
Author	=	Jim Davis
Experiment	=	single_pulse.ex2
Sample_id	=	MS0501
Solvent	=	CHLOROFORM-D
Changer_sample	=	11
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	=	JNM-ECA500
Field_strength	=	11.7473579[M] (500[MH
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[kHz]
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]
Recvr_gain	=	30
Relaxation_delay	=	4[s]
Repetition_time	=	5.74587904[s]
Temp_set	=	22.7[deg]





SOUTH ALABAMA
JAGUARS

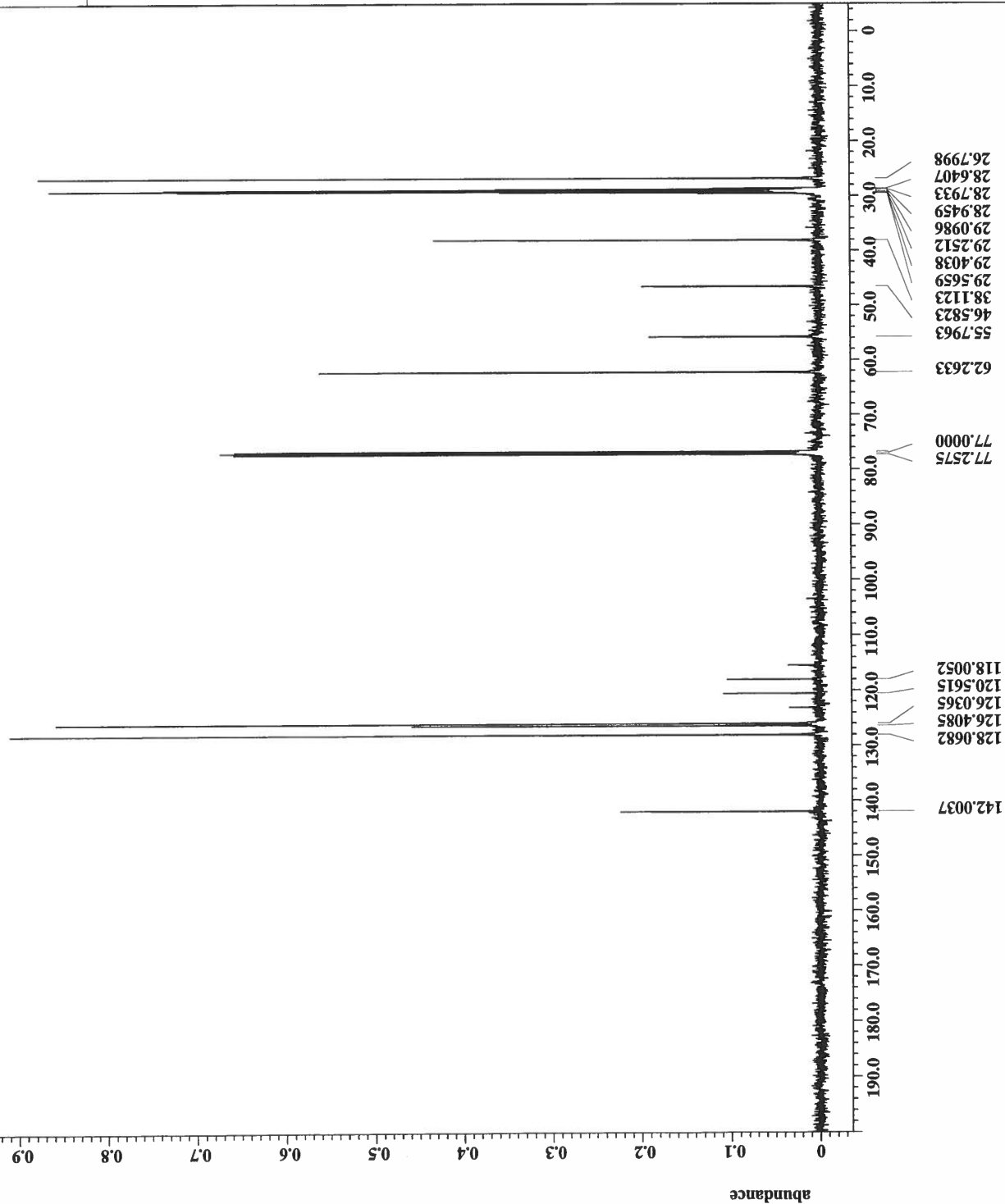
```

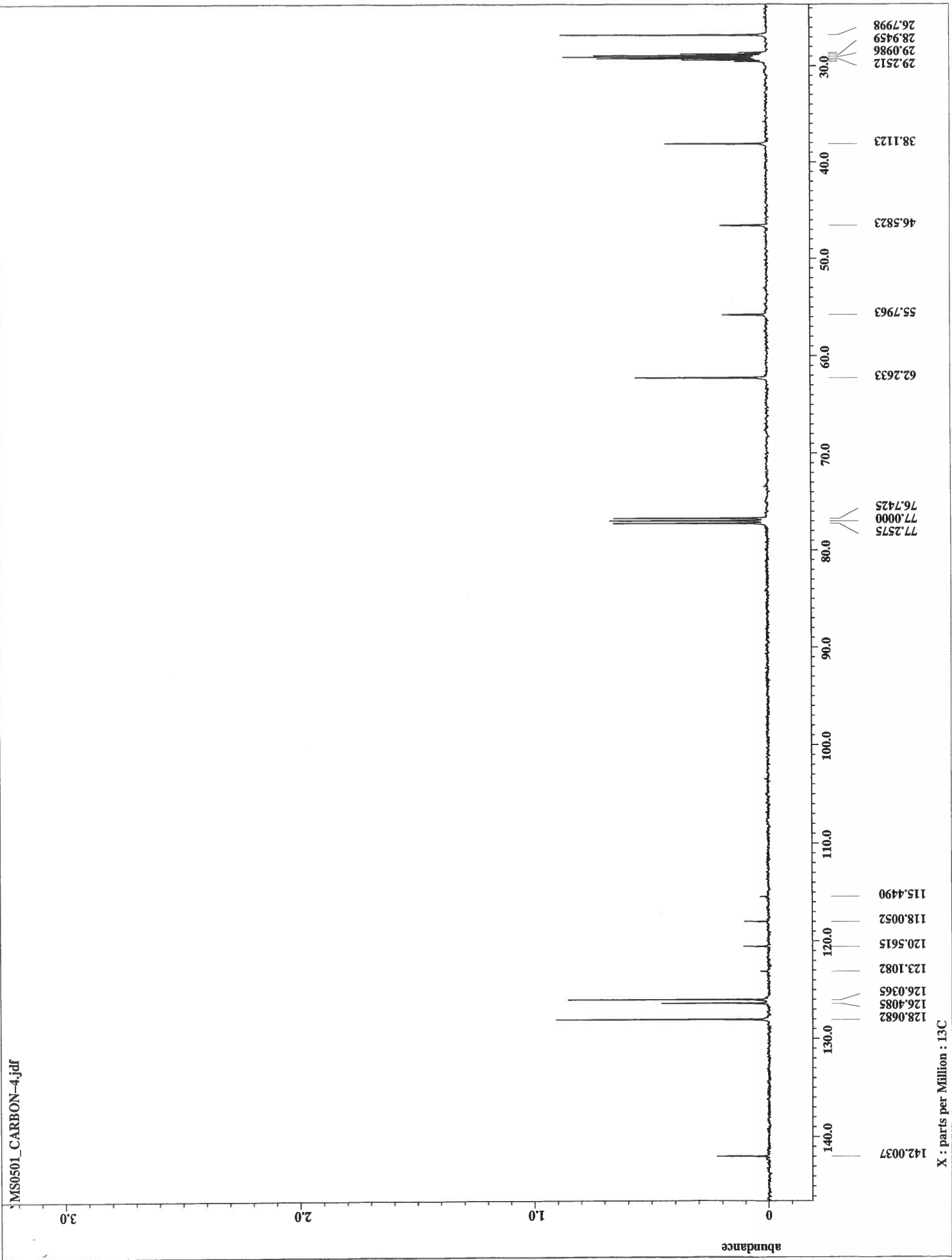
Filename      = MS0501_CARBON-2.jdf
Author       = Jim Davis
Experiment   = single_pulse_dec
Sample_id    = MS0501
Solvent      = CHLOROFORM-D
Changer_sample  = 11
Creation_time  = 5-JUL-2018 16:57:55
Revision_time  = 5-JUL-2018 16:34:26
Current_time   = 5-JUL-2018 16:34:26

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[MH
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       = 23.2[dc]
  
```

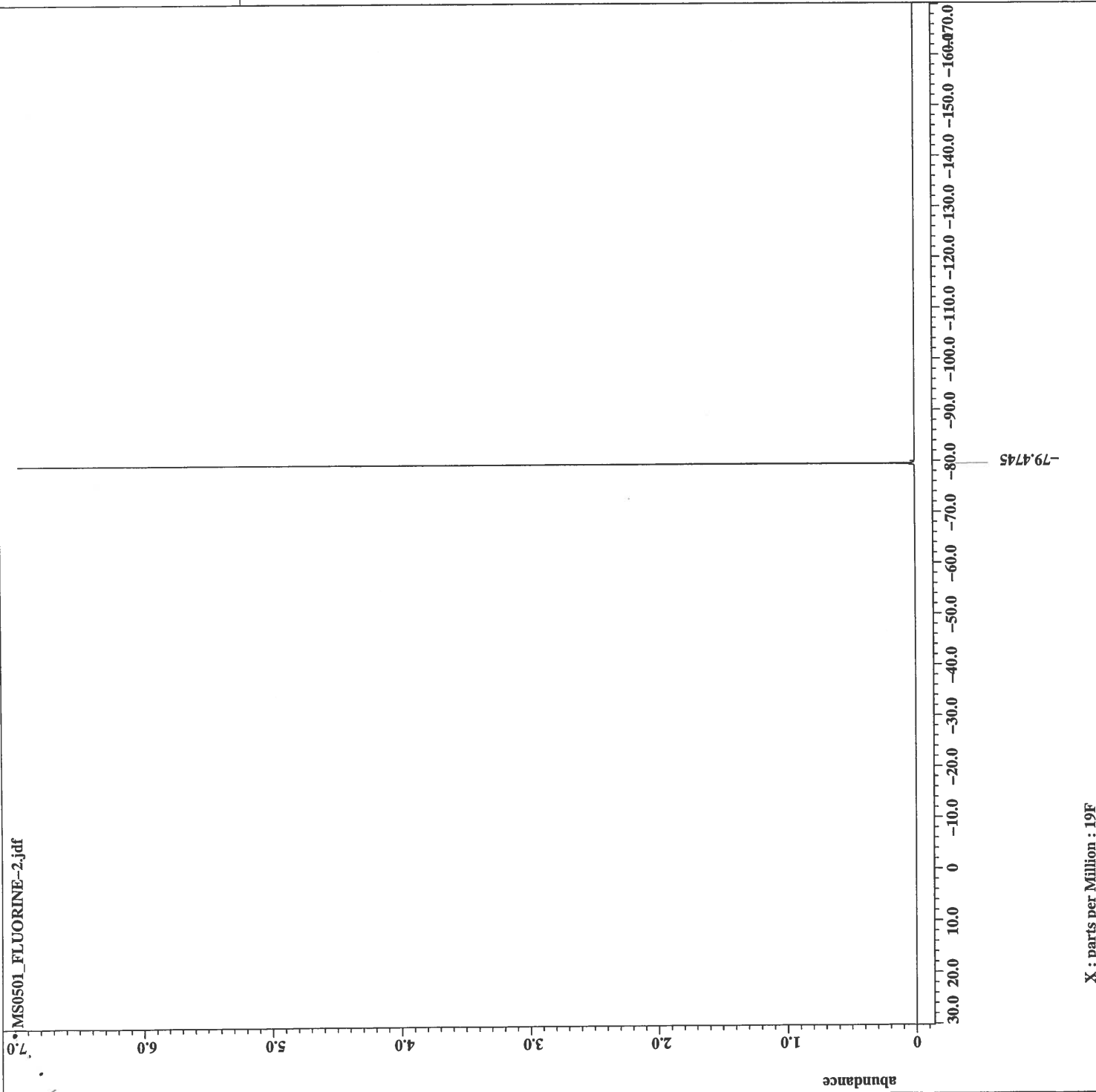




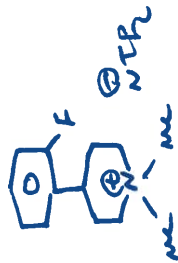
MS0501_CARBON-4.jdf

SOUTH ALABAMA
JAGUARS™

Filename = MS0501_FLUORINE-2.jdf
Author = Jim Davis
Experiment = single_pulse.ex2
Sample_id = MS0501
Solvent = CHLOROFORM-D
Charger_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[MH
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]
Recvr_gain = 38
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get = 22.8[dc]



110.44



61.61

62.91

29.23
33.02

abundance

7.0694
7.0545
6.8770
6.87353.4368
3.3807
3.0532
3.04522.3810
2.0420
2.0351
1.8129

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

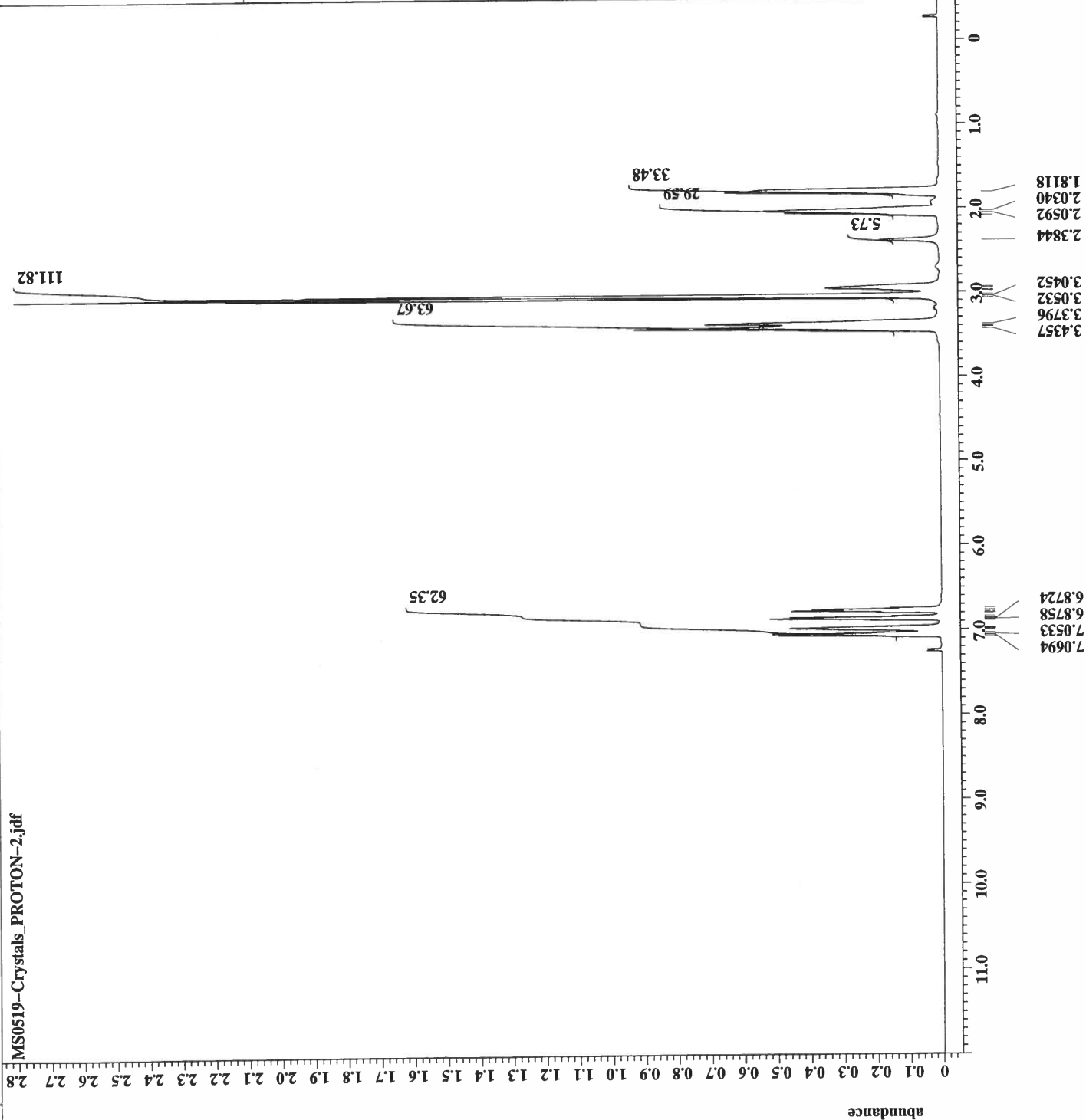
X : parts per Million : 1H

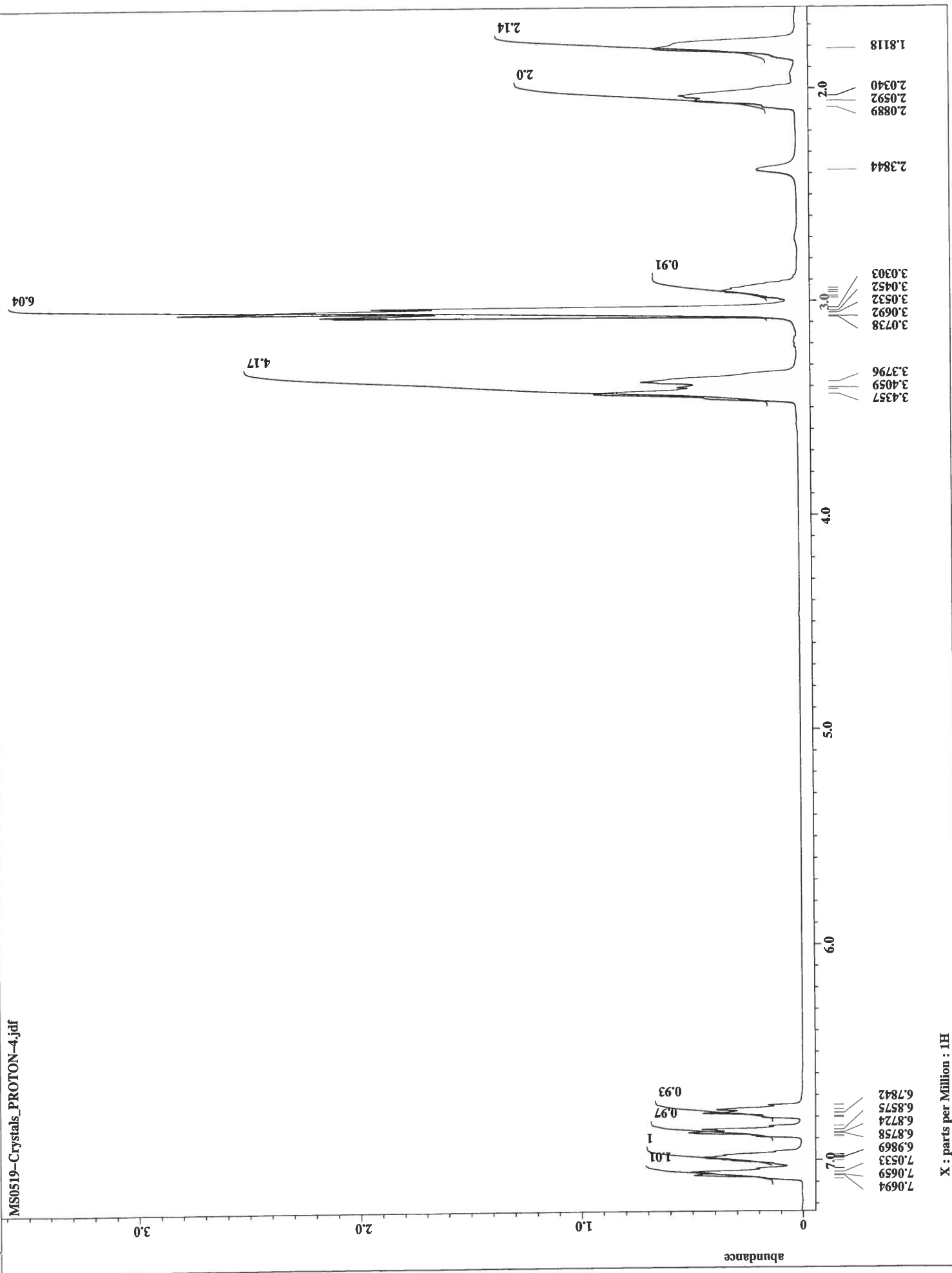


Filename = MS0519-Crystals_PROTO
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0519-Crystals
 Solvent = CHLOROFORM-D
 Changer_sample = 6
 Creation_time = 25-JUL-2018 09:12:03
 Revision_time = 25-JUL-2018 08:48:40
 Current_time = 25-JUL-2018 08:48:40
 Data_format = 1D COMPLEX
 Dim_size = 13107
 Dim_title = 1H
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579[T] (500[MH
 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 = 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]
 Recvr_gain = 30
 Relaxation_delay = 4[s]
 Repetition_time = 5.74587904[s]
 Temp_get = 22.4[dc]

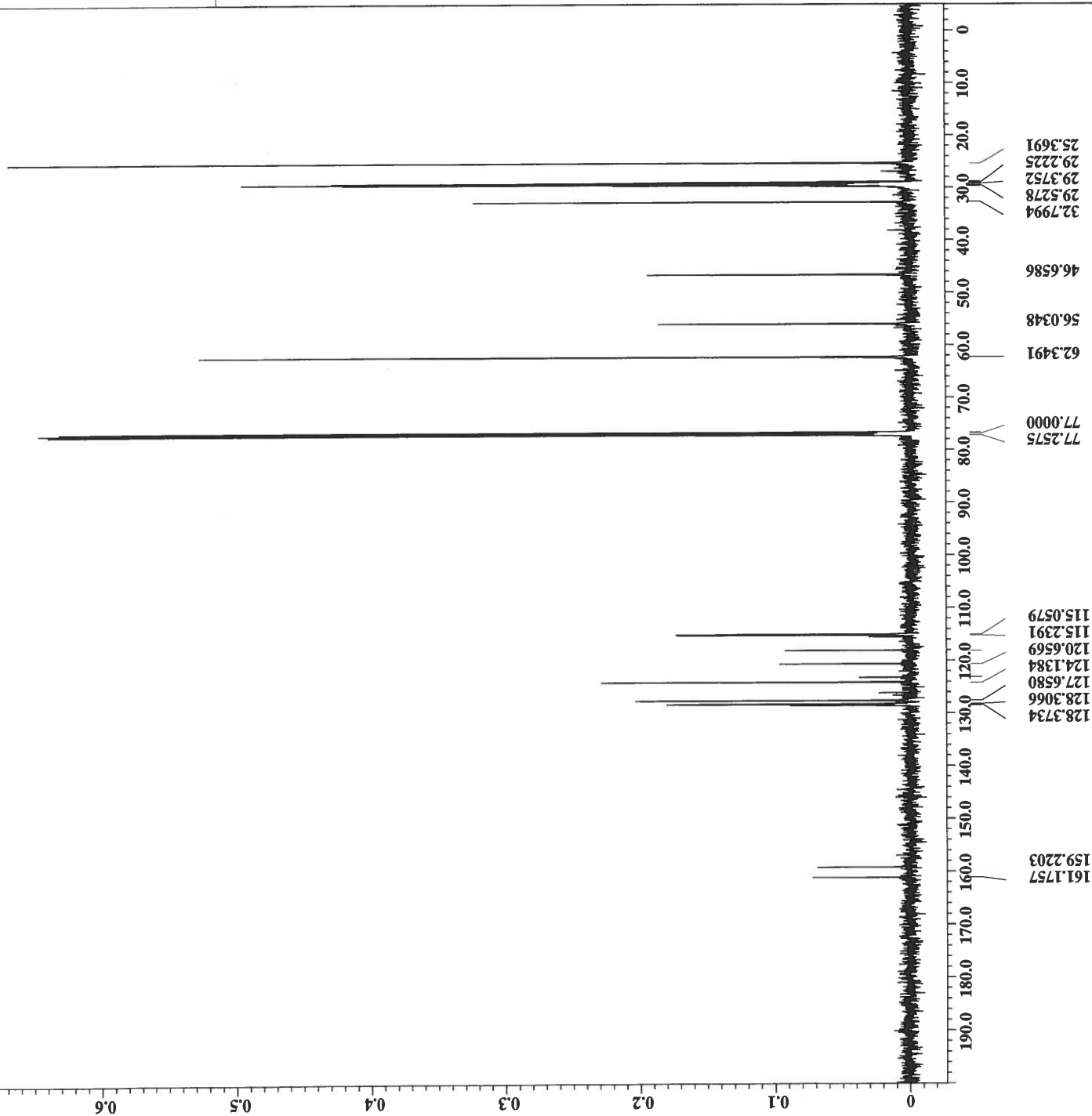


Filename = MS0519-Crystals_PROTO
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0519-Crystals
 Solvent = CHLOROFORM-D
 Changer_sample = 6
 Creation_time = 25-JUL-2018 08:46:11
 Revision_time = 25-JUL-2018 08:22:50
 Current_time = 25-JUL-2018 08:22:50
 Data_format = 1D COMPLEX
 Dim_size = 13107
 Dim_title = 1H
 Dim_units = [ppm]
 Dimensions = x
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579[T] (500 [MH
 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]
 Recvr_gain = 30
 Relaxation_delay = 4 [s]
 Repetition_time = 5.74587904 [s]
 Temp_get = 22 [C]





abundance

X : parts per Million : ¹³C
SOUTH ALABAMA
JAGUARS

```

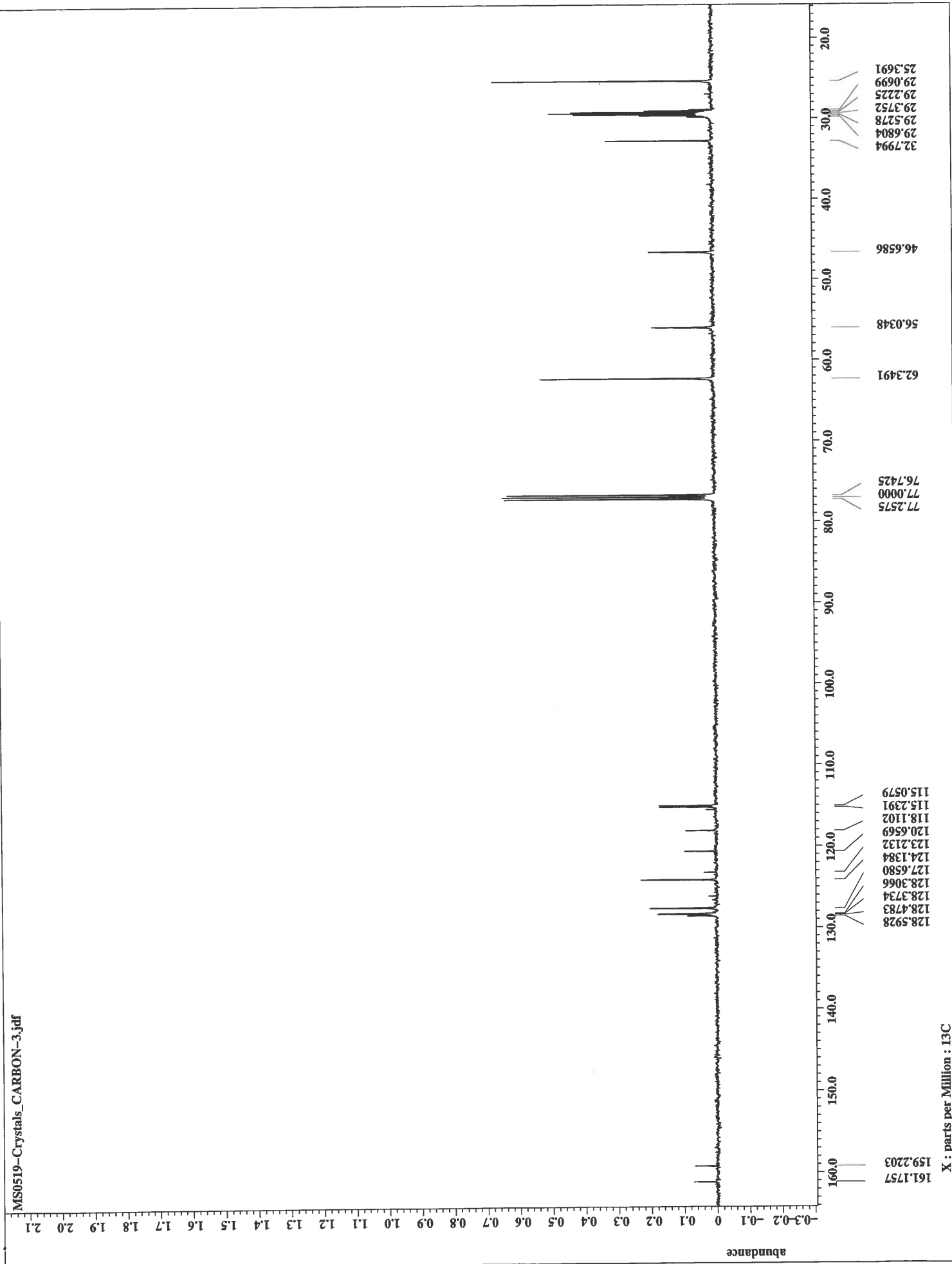
Filename = MS0519-Crystals_CARBO
Author = Jim Davis
Experiment = single_pulse_dec
Sample_id = MS0519-Crystals
Solvent = CHLOROFORM-D
Changer_sample = 6
Creation_time = 25-JUL-2018 09:02:33
Revision_time = 25-JUL-2018 08:39:11
Current_time = 25-JUL-2018 08:39:11

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[MH
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 = 300
Total_scans = 300

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 = 22.5[dc]

```





SOUTH ALABAMA
JAGUARS™

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 = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength = 11.7473579[T] (500[MH
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]
Recvr_gain = 36
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.4 [dC]



SOUTH ALABAMA
JAGUARS™

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 = 19F
Dim_units = [ppm]
Dimensions = X
Site = ECA 500
Spectrometer = JNM-ECA500
Field_strength = 11.7473579[T] (500[MH
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]
Recvr_gain = 36
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get = 22.4 [dC]

abundance

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 70.0

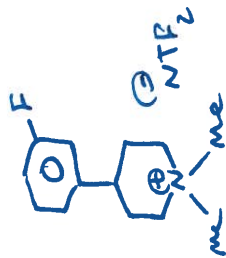
-79.3292

-118.8367

X : parts per Million : 19F



SOUTH ALABAMA
JAGUARS



m.p = 118°C

89.48

70.23

37.34

33.32

15.66

16.84

16.7

1.72

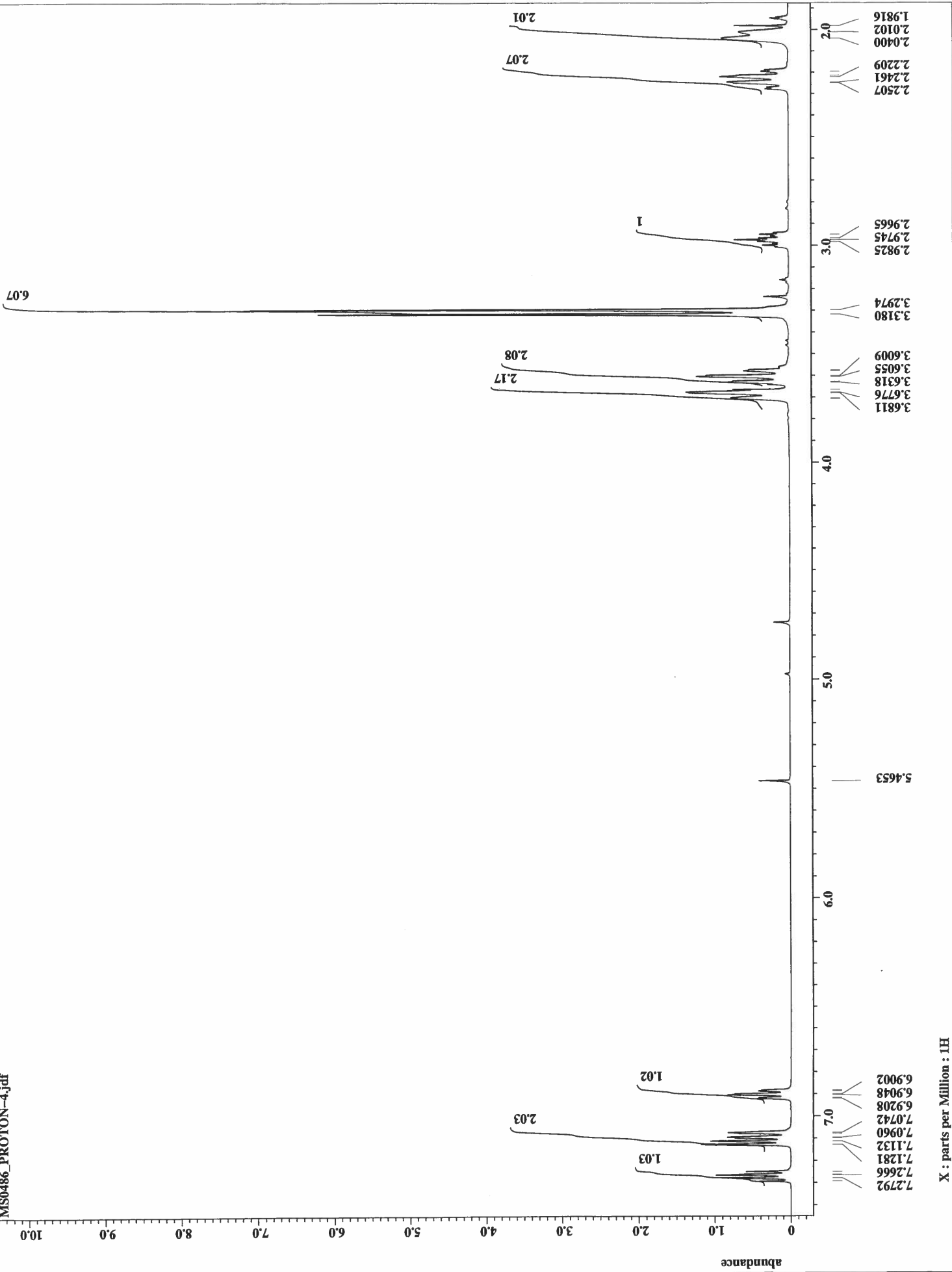
2.0400
2.2209
2.2461
2.2507
2.9745
2.9825
3.2974
3.3180
3.6776
3.6811

7.2792
7.2666
7.1281
7.1132

5.4653

X : parts per Million : 1H

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 = JNM-ECA500
 Field_strength = 11.7473579[T] (500[MH
 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]
 Recvr_gain = 22
 Relaxation_delay = 4[s]
 Repetition_time = 5.74587904[s]
 Temp_get = 22.3[dc]



3.0

2.0

1.0

abundance

164.7102

162.7739

160.0

147.1406
147.0833

150.0

140.0

131.2402
131.1734

130.0

124.6873

123.7526

122.1311

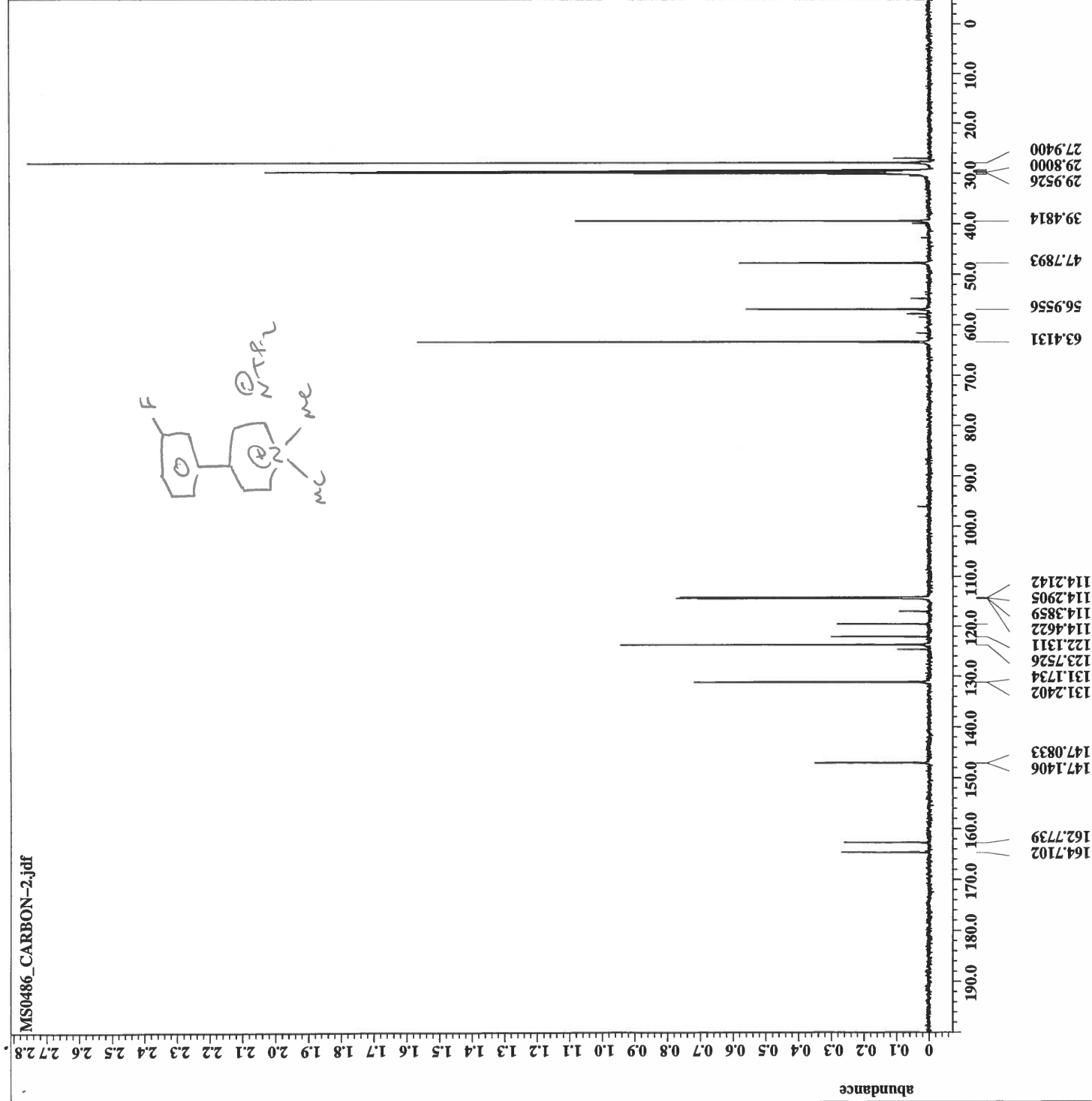
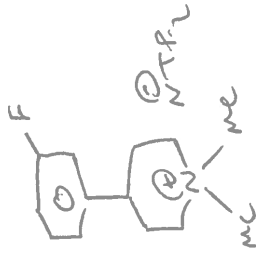
120.0

119.5748

117.0280

114.4622
114.3859
114.2905
114.2142

X : parts per Million : 13C





SOUTH ALABAMA
JAGUARS

```

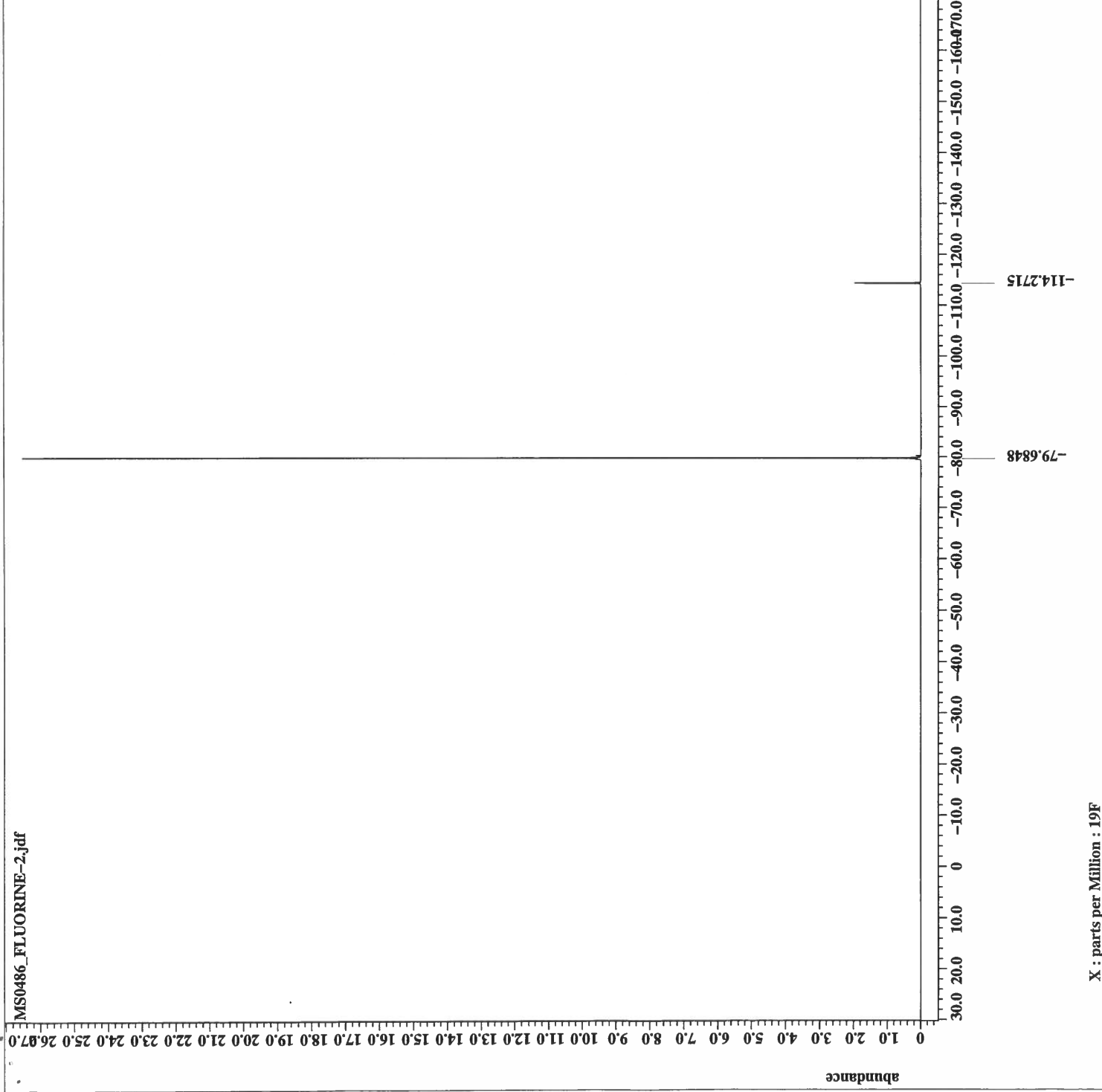
Filename      = MS0486_FLUORINE-2.jdf
Author       = Jim Davis
Experiment    = single_pulse.ex2
Sample_id    = MS0486
Solvent      = ACETONE-D6
Charger_sample 11
Creation_time 27-JUN-2018 10:39:16
Revision_time 27-JUN-2018 10:16:23
Current_time  27-JUN-2018 10:16:24

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[MH
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

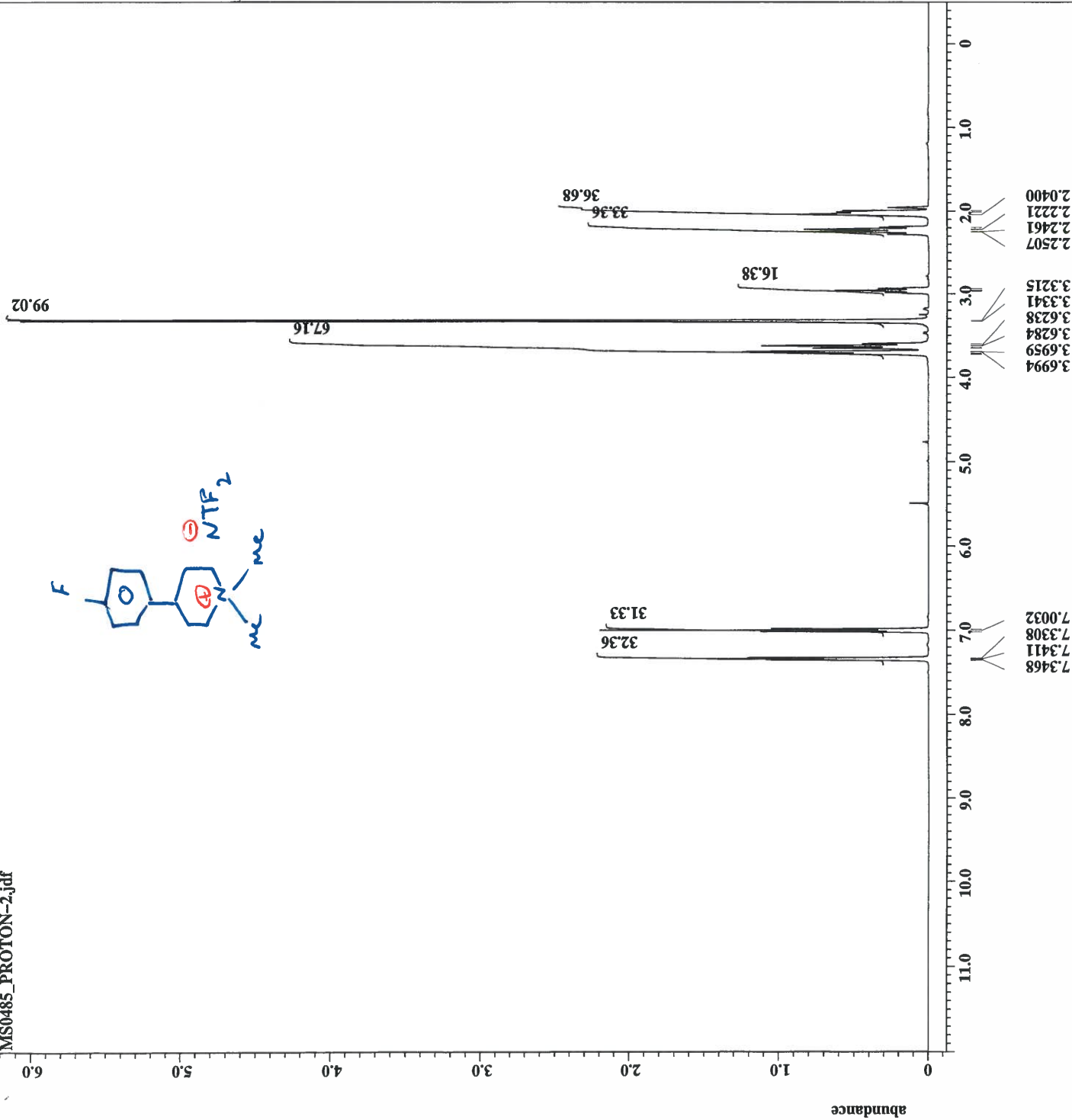
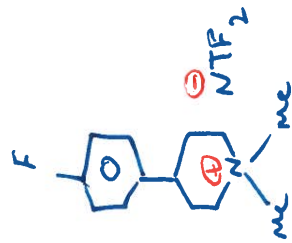
X_90_width    = 13.1[us]
X_acq_time    = 0.55574528[s]
X_angle       = 45[deg]
X_atn         = 2.5[dB]
X_pulse       = 6.55[us]
Irr_mode      = Off
Tri_mode      = Off
Dante_preset  = FALSE
Initial_wait  = 1[s]
Recvr_gain    = 30
Relaxation_delay = 4[s]
Repetition_time = 4.55574528[s]
Temp_get      =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

```



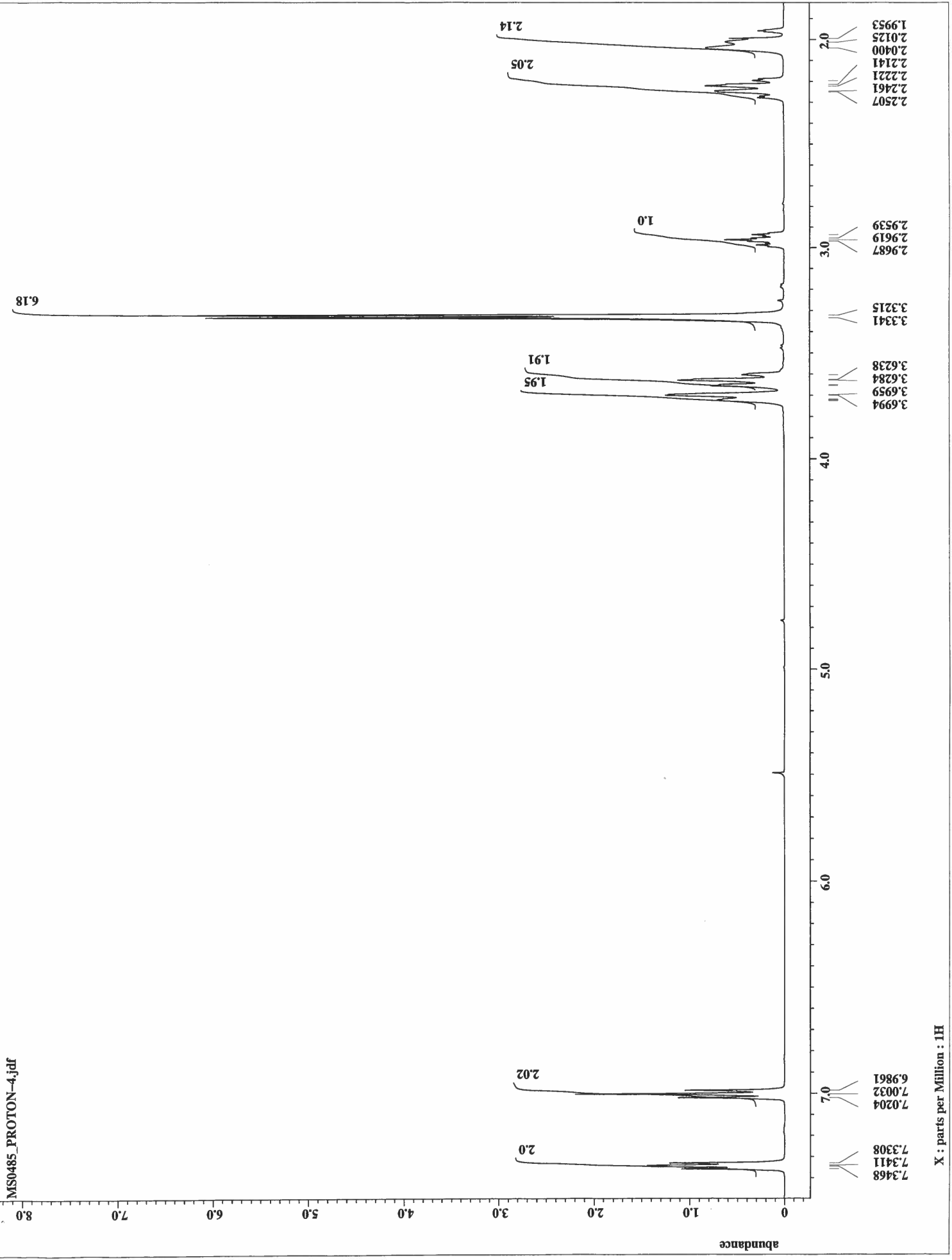


SOUTH ALABAMA
JAGUARS



X : parts per Million : 1H

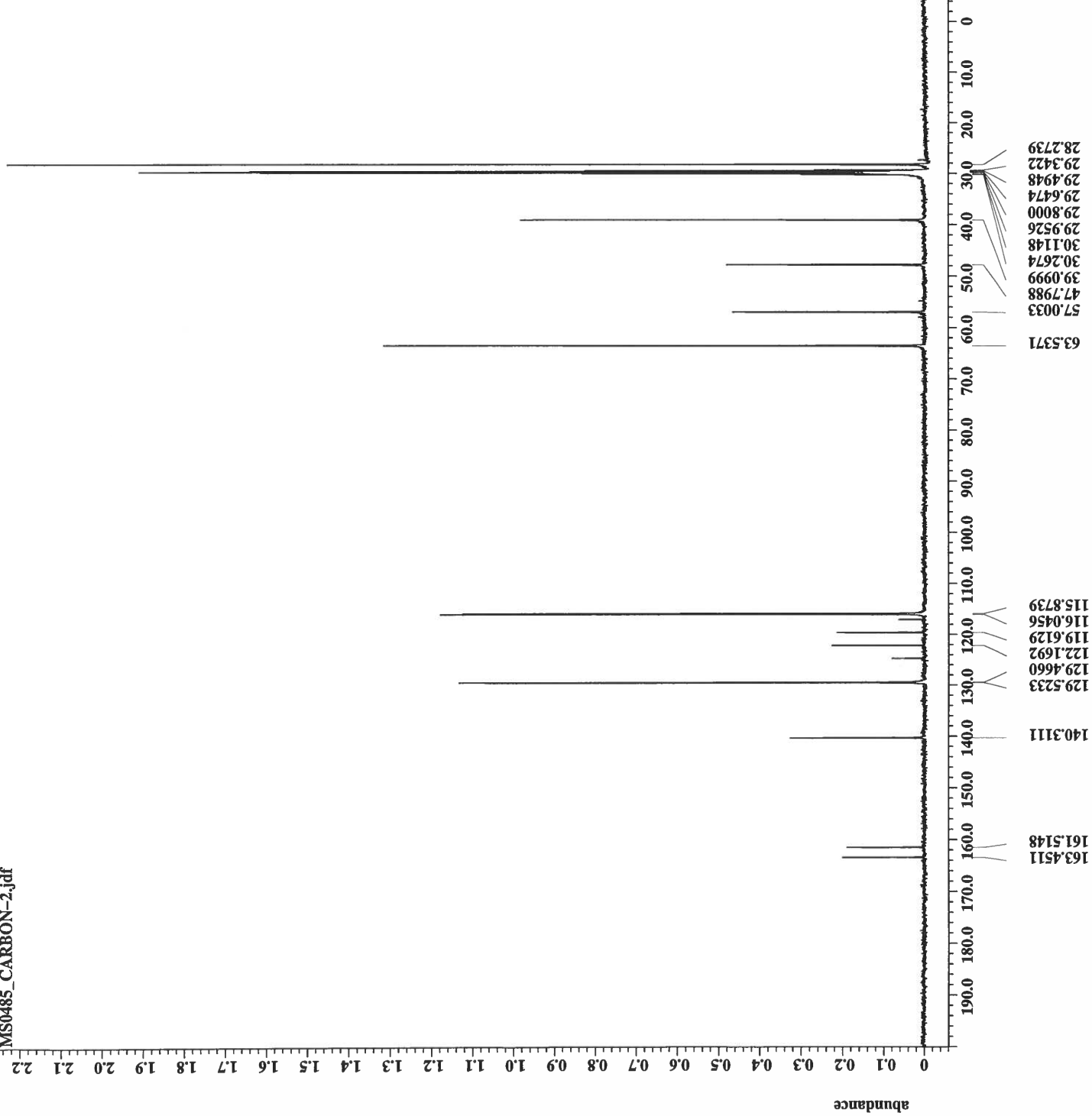
Filename = MS0485_PROTON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse.ex2
 Sample_id = MS0485
 Solvent = ACETONE-D6
 Changer_sample = 12
 Creation_time = 27-JUN-2018 11:38:17
 Revision_time = 27-JUN-2018 11:15:23
 Current_time = 27-JUN-2018 11:15:23
 Data_format = 1D COMPLEX
 Dim_size = 13107
 Dim_title = 1H
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579[T] (500[MH
 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]
 Recvr_gain = 24
 Relaxation_delay = 4[s]
 Repetition_time = 5.74587904[s]
 Temp_get = 22.3[dc]



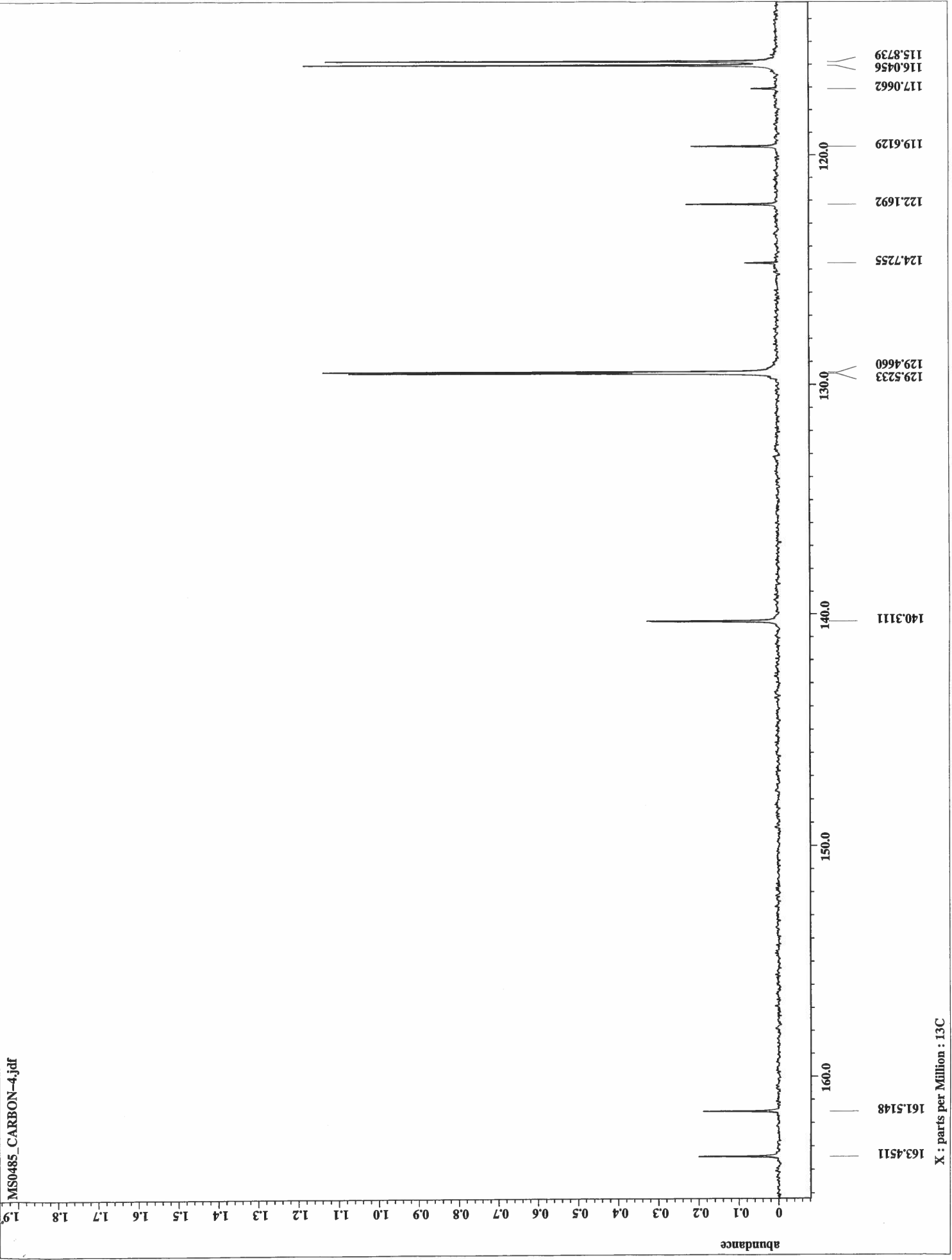


SOUTH ALABAMA
JAGUARS™

Filename = MS0485_CARBON-2.jdf
 Author = Jim Davis
 Experiment = single_pulse_dec
 Sample_id = MS0485
 Solvent = ACETONE-D6
 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 = 26214
 Dim_title = 13C
 Dim_units = [ppm]
 Dimensions = X
 Site = ECA 500
 Spectrometer = JNM-ECA500
 Field_strength = 11.7473579[T] (500[MH
 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 = 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 = 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 = 22.9 [dC]



X : parts per Million : 13C





SOUTH ALABAMA
JAGUARS

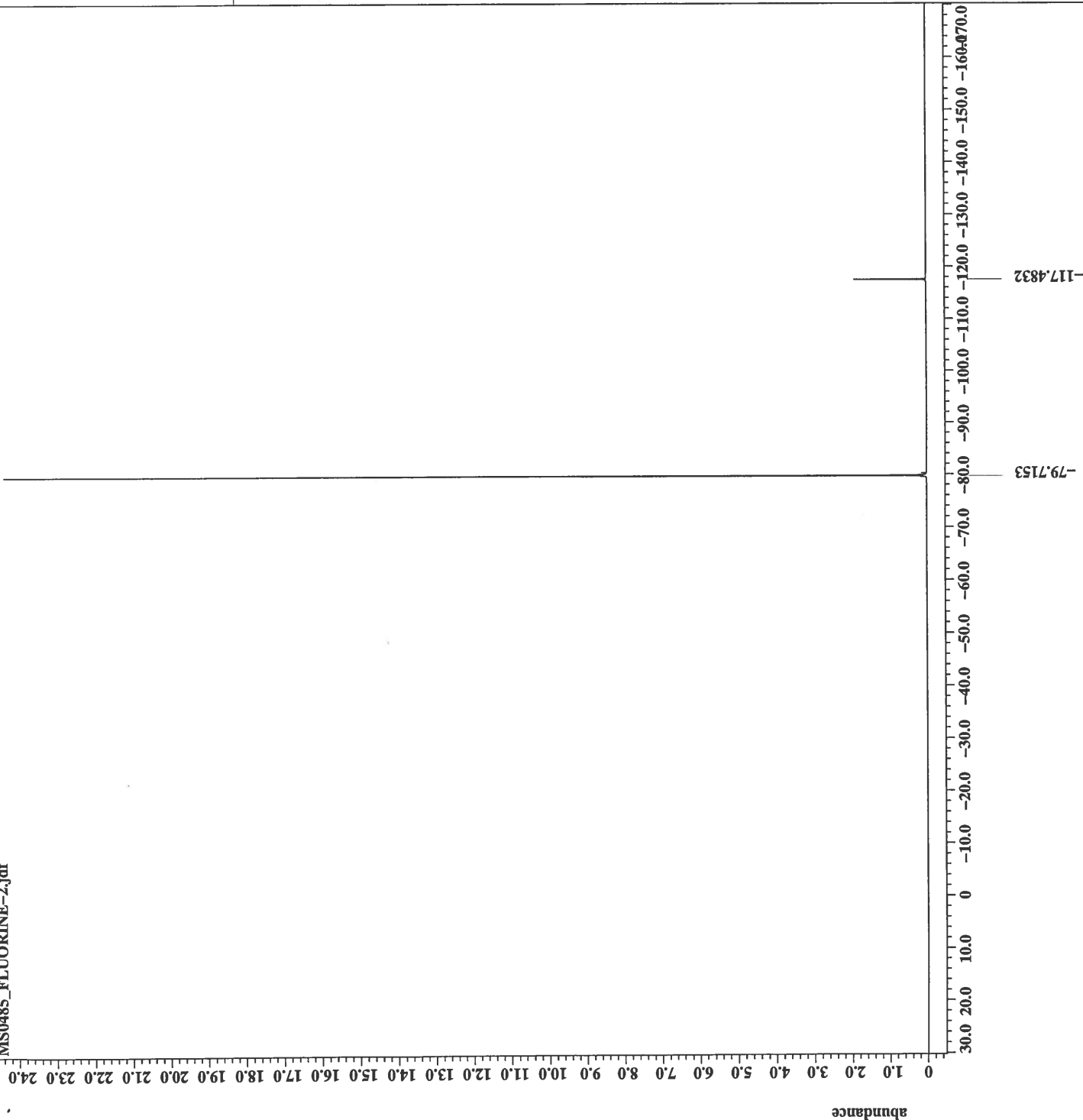
```

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[MH
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_presat   = FALSE
Initial_wait   = 1 [s]
Recvr_gain     = 34
Relaxation_delay = 4 [s]
Repetition_time = 4.55574528 [s]
Temp_get       = 22.3 [dC]
  
```



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) b=9.1975(4) c=18.6788(7)	
	alpha=90 beta=90 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

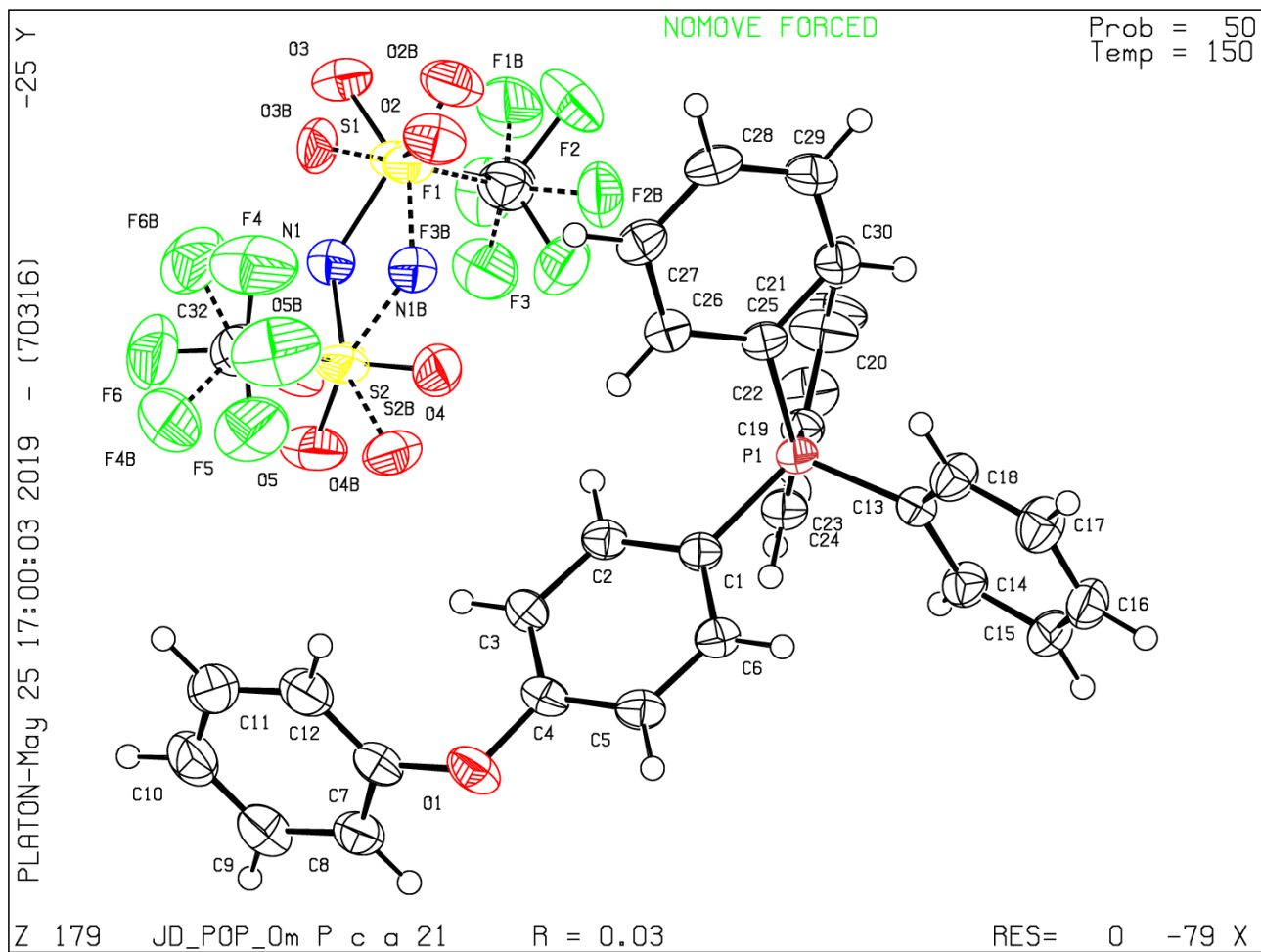
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section 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 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_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	58 Report
PLAT913_ALERT_3_C	Missing # of Very Strong Reflections in FCF		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	2.90 Ang.
	x,y,z =	1_555 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

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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

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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

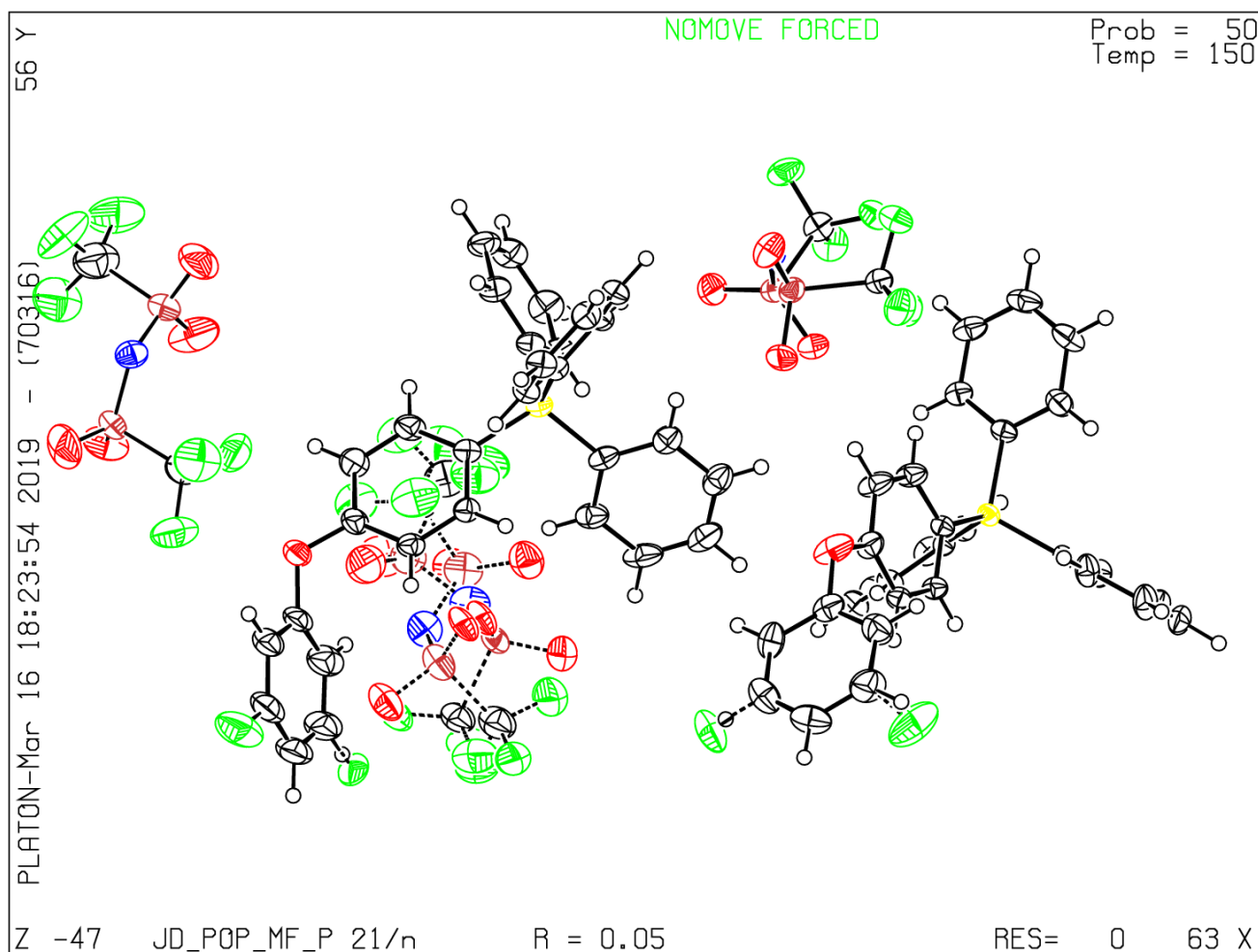
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section 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.

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Datablock JD_POP_MF_0m_5 - ellipsoid plot



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[Download CIF editor \(enCIFer\) from the CCDC](#)
[Test a new CIF entry](#)

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_OF_0m

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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 Å	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 O1 ..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 O5B ..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
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2 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
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- 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

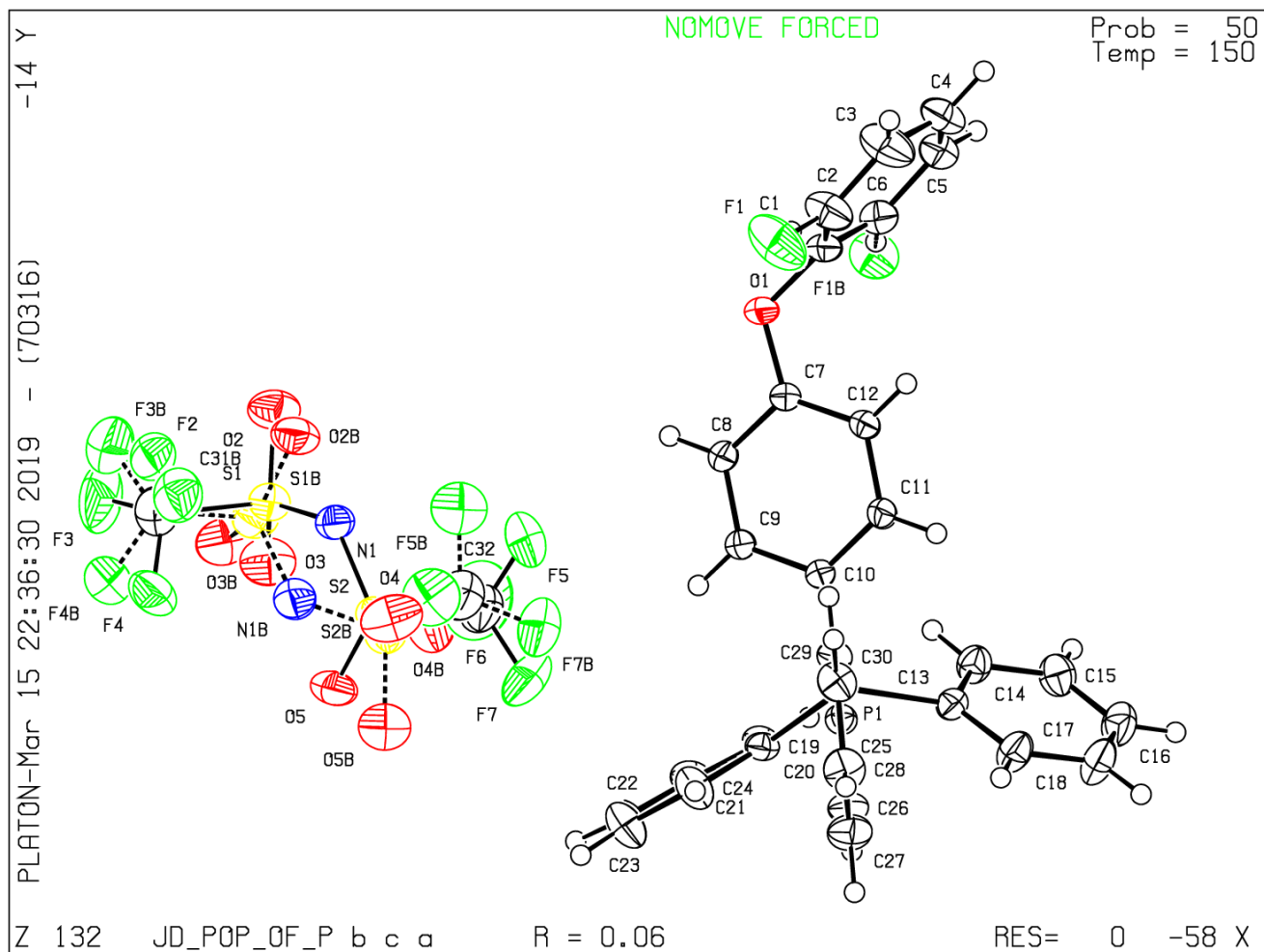
A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section 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.

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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 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_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
Mu (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).	6 Note
PLAT911_ALERT_3_C	Missing FCF Refl Between Thmin & STh/L= 0.600	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
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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

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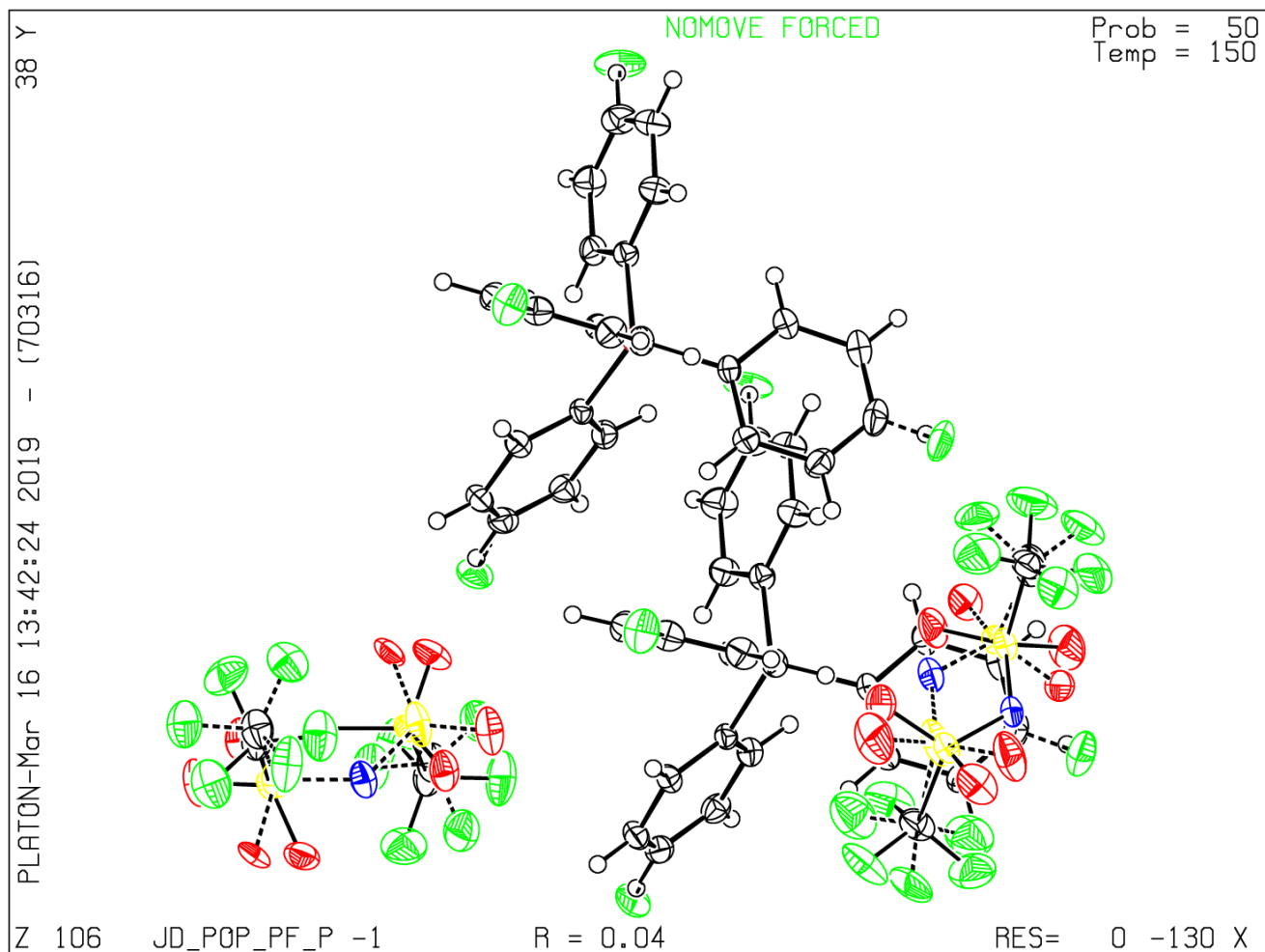
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PLATON version of 18/02/2019; check.def file version of 18/02/2019

Datablock JD_POP_PF_0m - ellipsoid plot



[Test a new CIF entry](#)

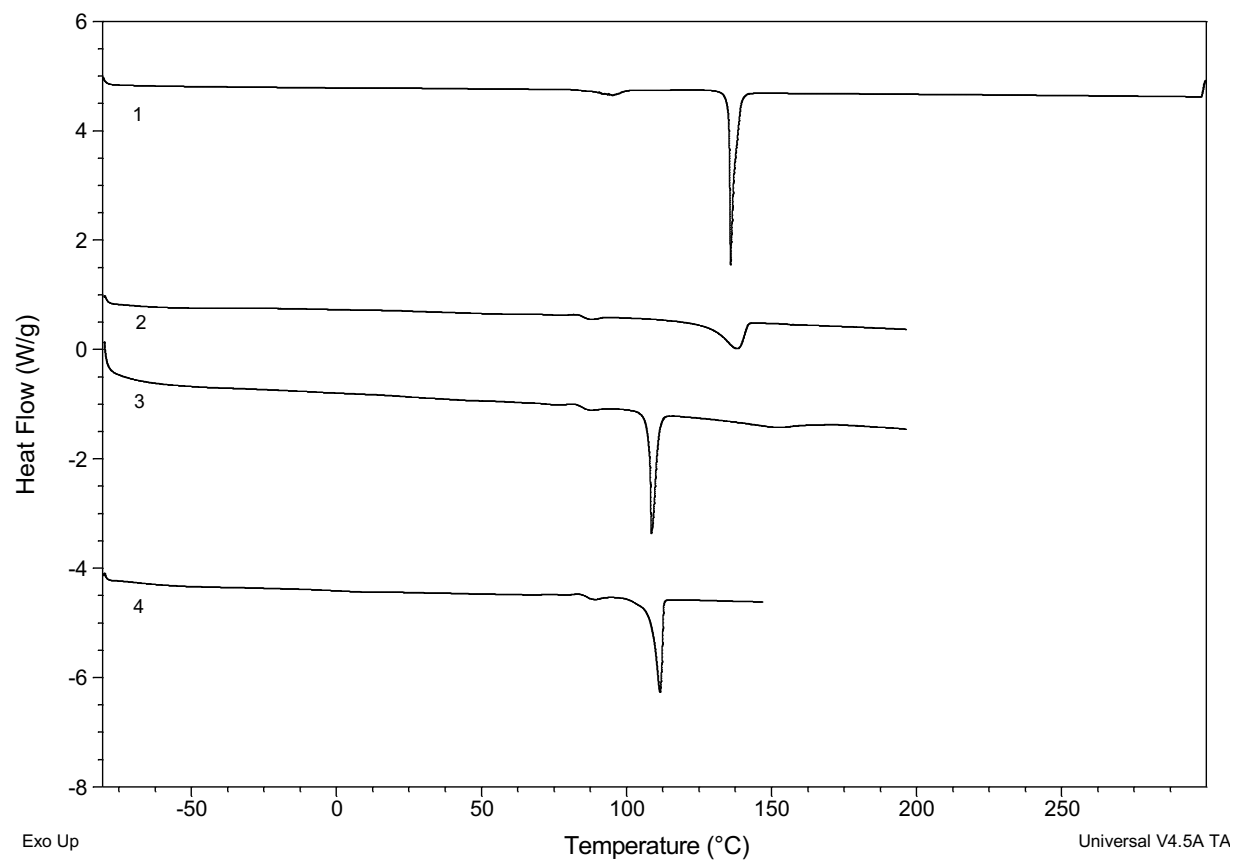


Figure S1: Representative DSC scans for compounds **1-4**.

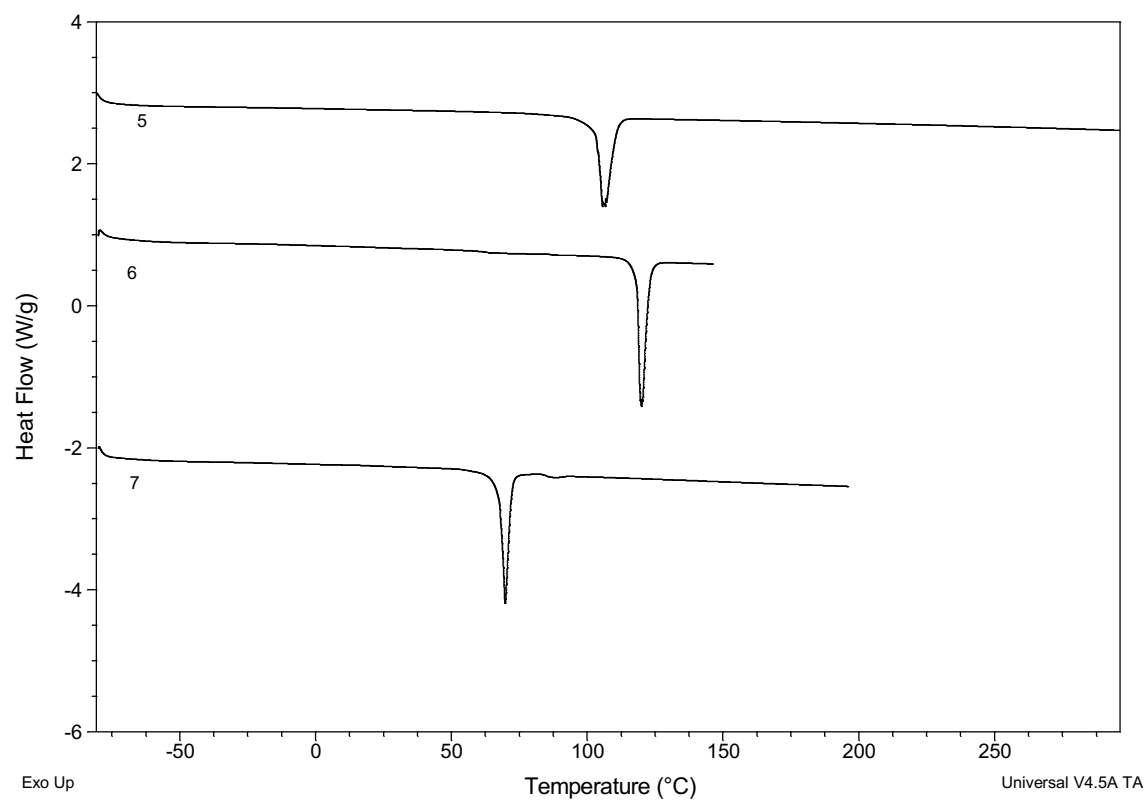


Figure S2: Representative DSC scans for compounds **5-7**.

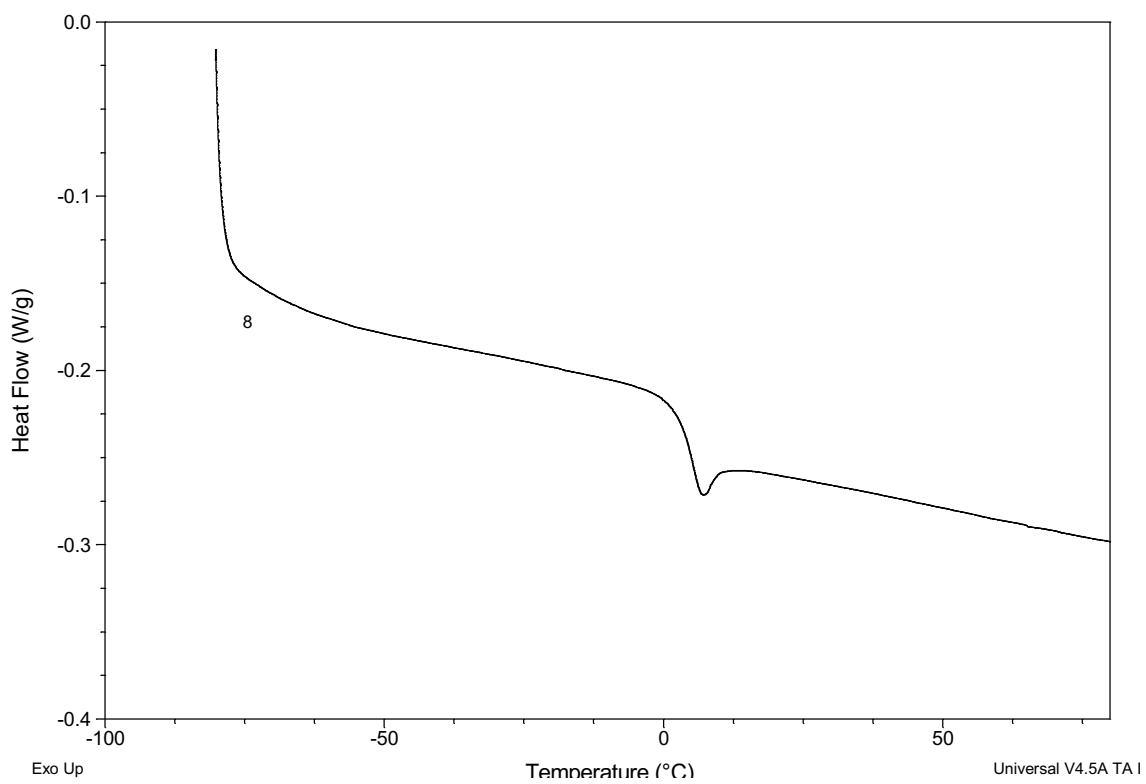


Figure S3: Representative DSC scan for compound **8**, which exhibits a weak glass transition.

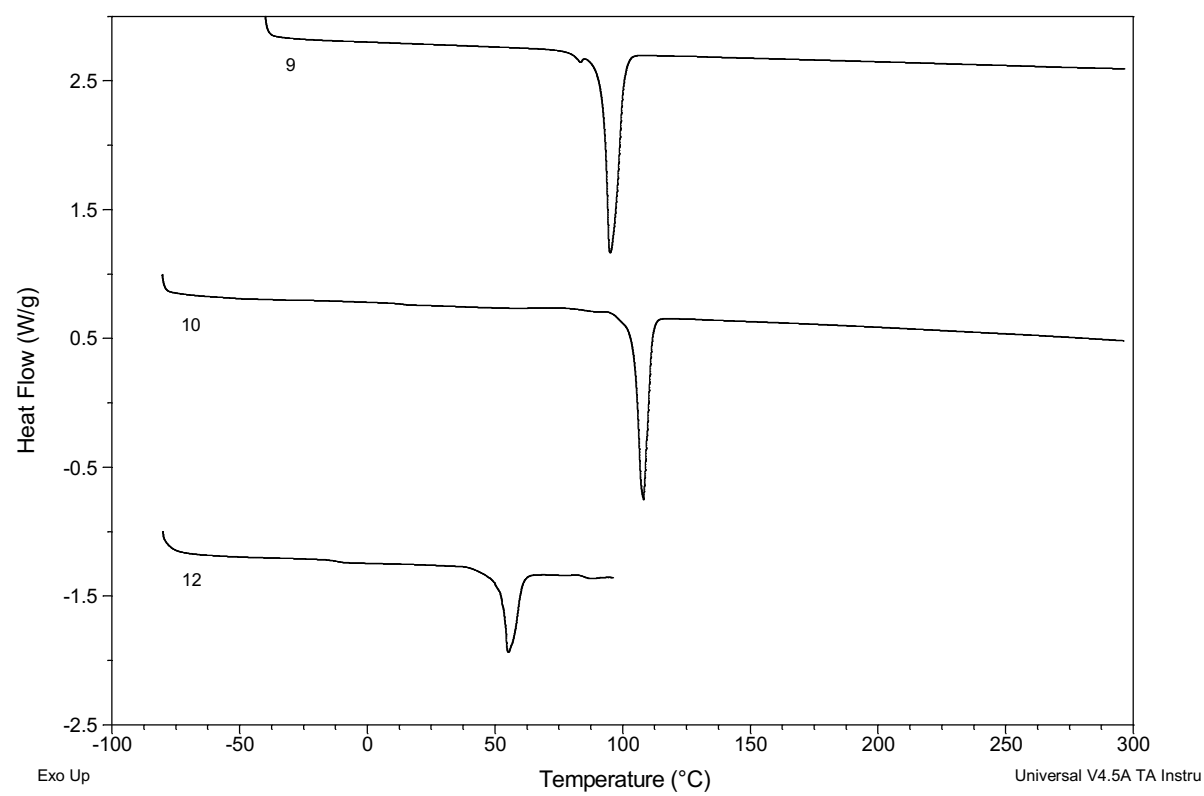


Figure S4: Representative DSC scans for compounds **9,10** and **12**.

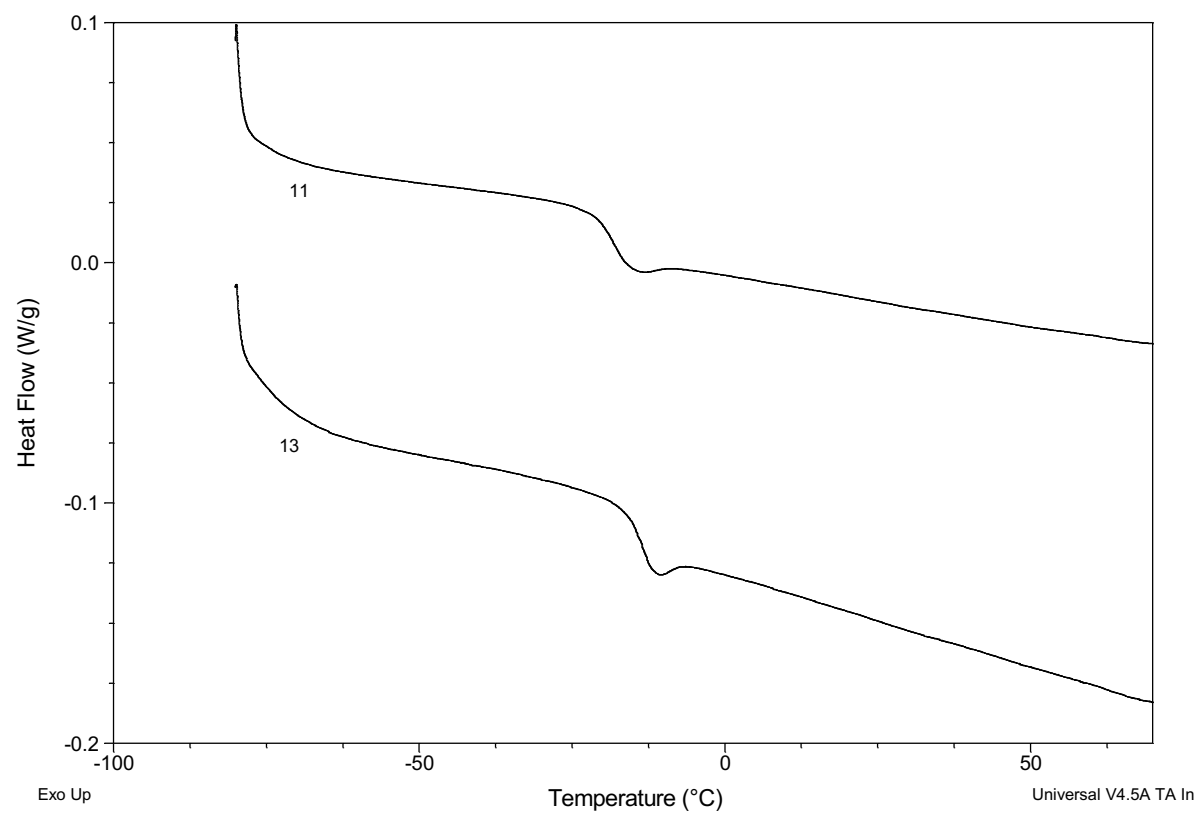


Figure S5: Representative DSC scans for compounds **11** and **13**, which exhibit weak glass transitions.

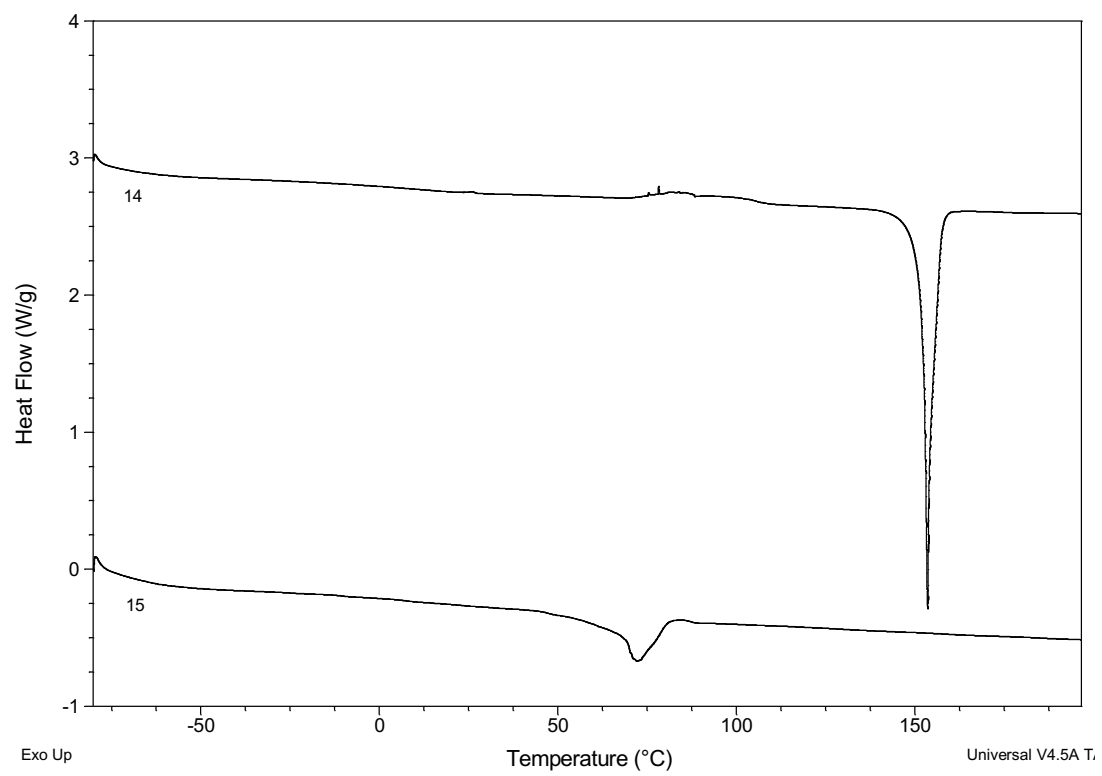


Figure S6: Representative DSC scans for compounds **14** and **15**.

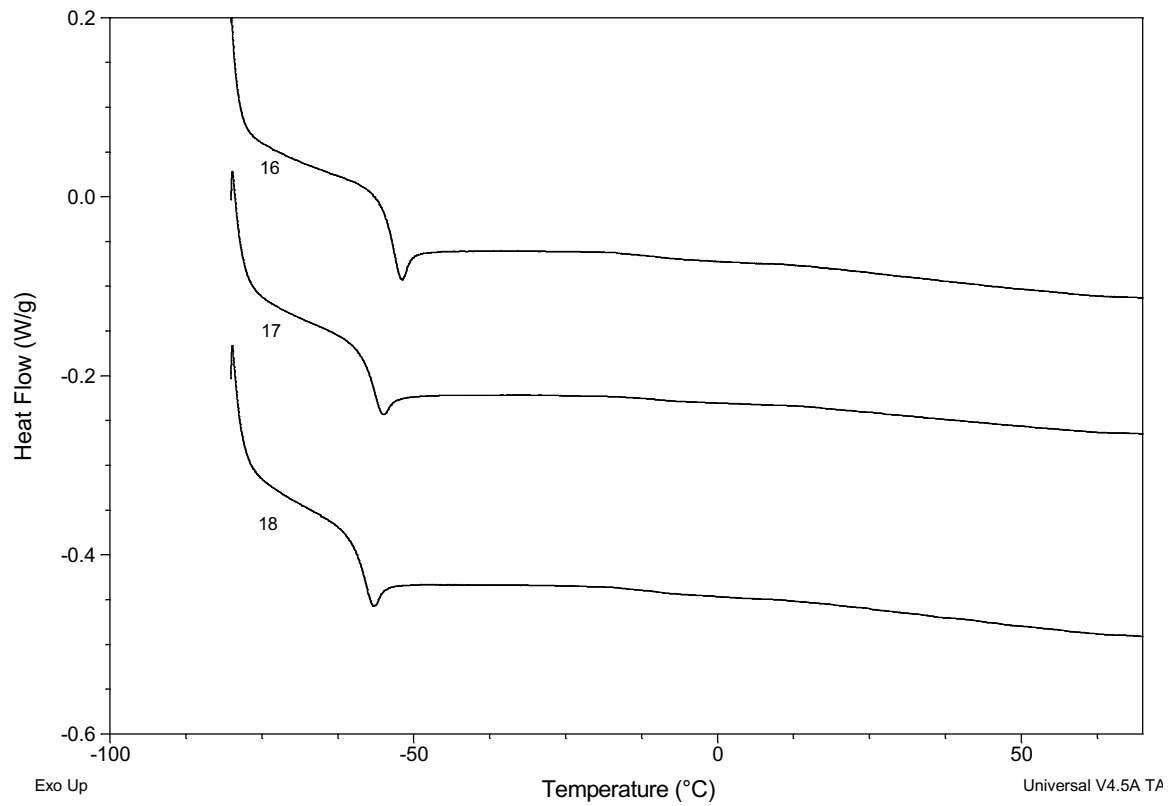


Figure S7: Representative DSC scans for compounds **16-18**, and **13**, which exhibit weak glass transitions.

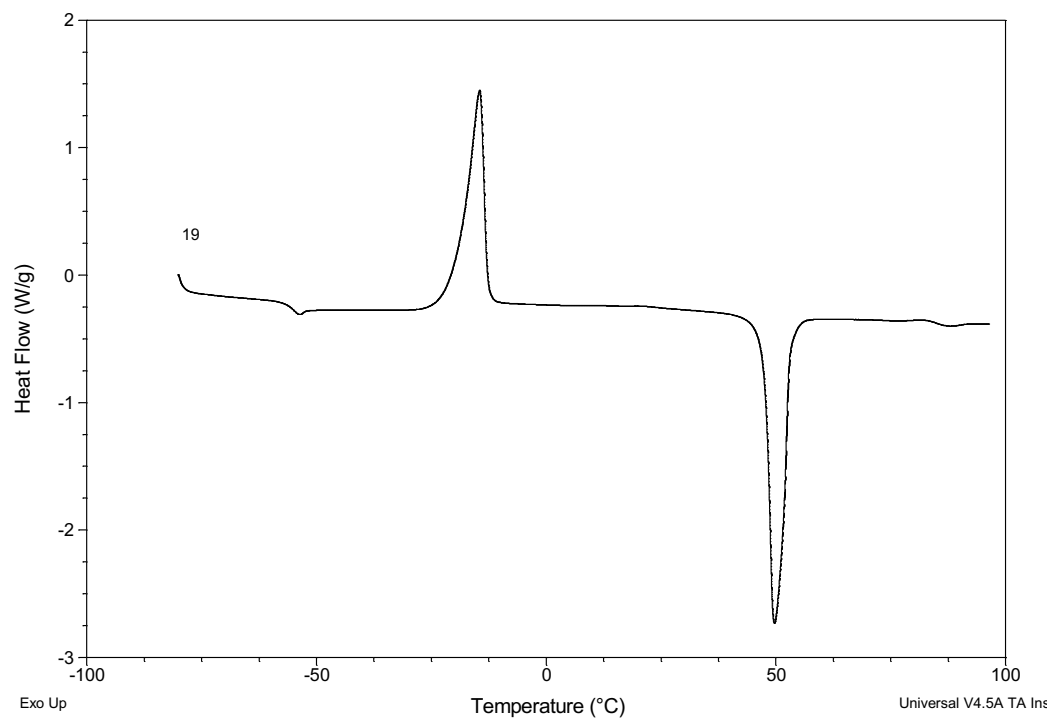


Figure S8: Representative DSC scan for compound **19**.

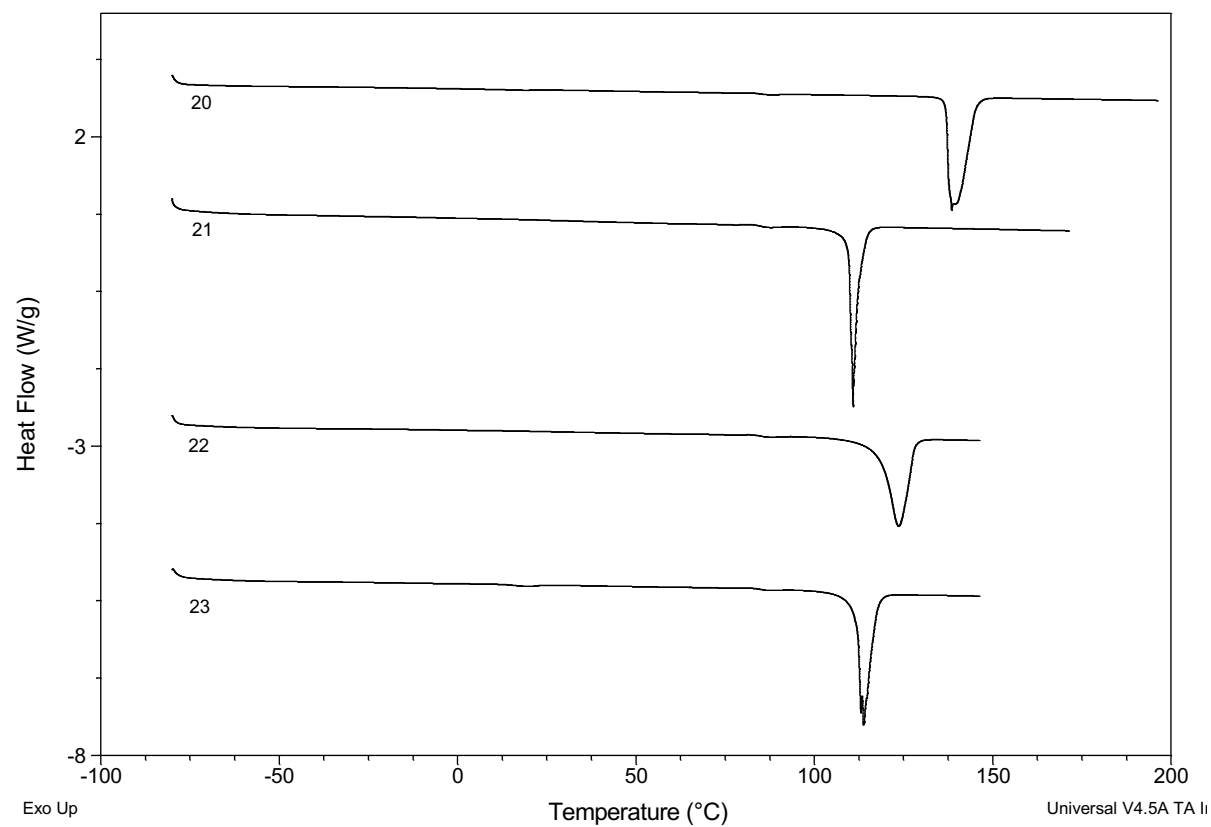


Figure S9: Representative DSC scans for compounds **20-23**.

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)'


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_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.

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_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 (Hübschle 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 (H²ubschle et al., 2011)

;

_computing_molecular_graphics ?

_computing_publication_material ?

_refine_special_details

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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.

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