

Supporting Information

Merging Norrish Type I Reaction and Transition Metal Catalysis: Photo- and Rh-promoted Borylation of C–C σ -Bonds of Arylketones

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General methods

All operations were performed under an argon atmosphere. ^1H and ^{13}C NMR spectra were recorded on a JEOL ECX-500 or a JEOL ECZ-500 (500 MHz for ^1H and 125 MHz for ^{13}C) spectrometer in CDCl_3 or C_6D_6 . Chemical shifts are expressed in parts per million (ppm) downfield from tetramethylsilane ($\delta_{\text{H}} = 0.00$, $\delta_{\text{C}} = 0.00$) and are referenced to residual solvents (δ_{H} 7.26 and δ_{C} 77.0 for CDCl_3 and δ_{H} 7.15 and δ_{C} 128.1 for C_6D_6). IR spectra were recorded on an IRTracer-100 (SHIMADZU Co., Ltd.) with an ATR QATR10 accessory. High resolution mass spectra (HRMS) were recorded on a BRUKER micrOTOF II and a JEOL JMS-T100 spectrometer. Gas chromatography (GC) analyses were conducted on a SHIMADZU GC-2010 equipped with an Agilent DB-1 column. EI-MS analyses (EI^+) were performed on a Shimadzu GCMS-QP2020. Silica Gel 60 (Kanto Chemical Co., Inc.) was used for flash column chromatography. Photoirradiation was carried out using a LED (365 nm, HLDEL-120U6-NWPSC/PSCC-60048(A), CCS). During the photoirradiation, the reaction temperature was kept at a constant temperature using a water bath equipped with a low temperature circulator (CTP-1000, EYELA). Merck Kieselgel 60 F254 (0.25 mm thickness, coated on glass 20x20 cm^2) plate was used for analytical thin layer chromatography (TLC). THF, Et_2O and hexane were purified by solvent purification system of Glass-Contour. Benzene- d_6 was purchased from Kanto chemicals and dried and degassed by benzophenone ketyl. All commercially available reagents were obtained from chemical suppliers and used after proper purification if necessary.

General procedures for the preparation of pivalophenone derivatives

Pivalophenone derivatives **1a-c**, **1e-g**, **1l**, **1n**, **1p-1q**, **1t** and **1u-1w** were synthesized by the general procedure A. **1h-k** and **1s** were synthesized by the general procedure B. **1d**, **1m**, **1o** and **1r** were synthesized according to a literature procedure.^{S1-S3} All analytical data of **1h**, **1j**, and **1m** are listed below. Other ketones are known compounds in literatures as referenced, and spectral data were in good agreement with literature values.

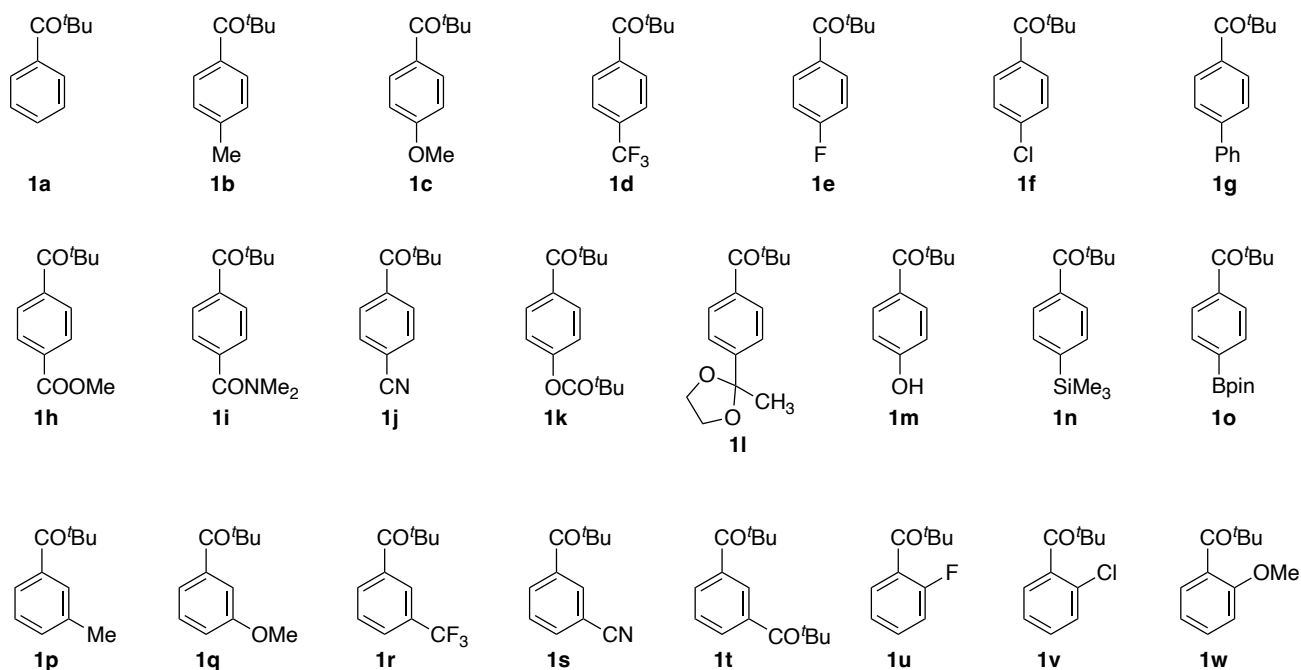


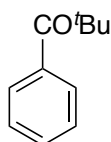
Figure S1. List of pivalophenone derivatives

General procedure A

Under a nitrogen atmosphere, a THF solution of Grignard reagent (ArMgBr, 20 mmol), which was prepared from Mg (20 mmol) and ArBr (21 mmol) in THF (40 mL), was added slowly to a solution of pivaloyl chloride (6.0 mL, 49 mmol) in THF (40 mL) at 0 °C. After stirring for 3 hours, the mixture was diluted with saturated aq. NH₄Cl (30 mL) and extracted with ethyl acetate three times. The combined organic layers were washed with saturated aq. Na₂CO₃ (30 mL) and brine (30 mL) and dried over anhydrous Na₂SO₄. After the solvent was removed under reduced pressure, the residue was purified by flash column chromatography followed by distillation to give pivalophenone derivatives in 22-72% yields.

General procedure B

Under a nitrogen atmosphere, a solution of $i\text{PrMgCl}\cdot\text{LiCl}$ (1.3 M in THF, 15.4 mL, 20 mmol) was added slowly to a solution of aryl iodide (20 mmol) in THF (10 mL) at $-40\text{ }^{\circ}\text{C}$. The solution of Grignard reagent (ArMgX) was added slowly to a solution of pivaloyl chloride (6.0 mL, 20 mmol) in THF (40 mL) at $-40\text{ }^{\circ}\text{C}$. After stirring for 3 hours, the mixture was diluted with saturated aq. NH_4Cl (30 mL) and extracted with ethyl acetate three times. The combined organic layers were washed with saturated aq. Na_2CO_3 (30 mL) and brine (30 mL) and dried over anhydrous Na_2SO_4 . After the solvent was removed under reduced pressure, the residue was purified by flash column chromatography followed by distillation to give pivalophenone derivatives in 36-70% yields.

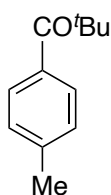


1a (colorless oil, 3.97 g, 24.5 mmol, 49%)

Synthesized according to the procedure **A** using 50 mmol of commercially available PhMgBr .

^1H NMR (500 MHz, CDCl_3) δ = 7.69 (d, J = 7.0 Hz, 2H), 7.46 (t, J = 7.5 Hz, 1H), 7.40 (t, J = 7.5 Hz, 2H), 1.35 (s, 9H).

Spectral data were in good agreement with literature values.^{S1}

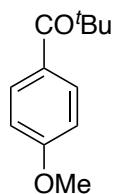


1b (colorless oil, 1.05 g, 5.94 mmol, 56%)

Synthesized according to the procedure **A** using 10 mmol of Mg.

^1H NMR (500 MHz, CDCl_3) δ = 7.66 (d, J = 8.5 Hz, 2H), 7.20 (d, J = 8.5 Hz, 2H), 2.39 (s, 3H), 1.35 (s, 9H).

Spectral data were in good agreement with literature values.^{S5}

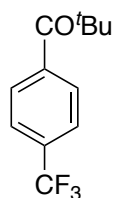


1c (colorless oil, 1.63 g, 8.50 mmol, 43%)

Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.86 (d, J = 8.0 Hz, 2H), 6.90 (d, J = 8.0 Hz, 2H), 3.86 (s, 3H), 1.37 (s, 9H) .

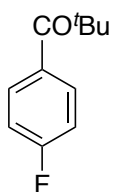
Spectral data were in good agreement with literature values.^{S1}



1d (colorless oil, 3.33 g, 14.5 mmol, 72%)

¹H NMR (500 MHz, CDCl₃) δ = 7.72 (d, J = 8.0 Hz, 2H), 7.67 (d, J = 8.5 Hz, 2H), 1.34 (s, 9H).

Spectral data were in good agreement with literature values.^{S1}

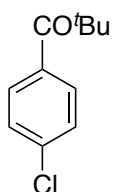


1e (colorless oil, 1.04 g, 5.78 mmol, 58%)

Synthesized according to the procedure **A** using 10 mmol of Mg.

¹H NMR (500 MHz, CDCl₃) δ = 7.82-7.76 (m, 2H), 7.08 (t, J = 8.5 Hz, 2H), 1.36 (s, 9H).

Spectral data were in good agreement with literature values.^{S6}

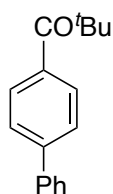


1f (colorless oil, 1.04 g, 5.21 mmol, 52%)

Synthesized according to the procedure **A** using 10 mmol of Mg.

¹H NMR (500 MHz, CDCl₃) δ = 7.67 (d, J = 9.0 Hz, 2H), 7.38 (d, J = 8.5 Hz, 2H), 1.34 (s, 9H).

Spectral data were in good agreement with literature values.^{S6}

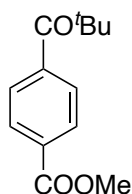


1g (white solids, 1.03 g, 4.33 mmol, 22%)

Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.82 (d, J = 9.0 Hz, 2H), 7.65-7.60 (m, 4H), 7.49-7.44 (m, 2H), 7.39 (t, J = 7.5 Hz, 1H), 1.40 (s, 9H).

Spectral data were in good agreement with literature values.^{S7}

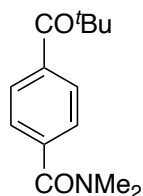


1h (white solids, 0.863 g, 3.91 mmol, 39%)

Synthesized according to the procedure **B** using 10 mmol of methyl 4-iodobenzoate.

¹H NMR (500 MHz, CDCl₃) δ = 8.07 (d, J = 8.5 Hz, 2H), 7.66 (d, J = 8.5 Hz, 2H), 3.94 (s, 3H), 1.33 (s, 9H).

Spectral data were in good agreement with literature values.^{S1}



1i (white solids, 1.68 g, 7.21 mmol, 36%)

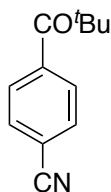
Synthesized according to the procedure **B**.

¹H NMR (500 MHz, CDCl₃) δ = 7.69 (d, J = 8.5 Hz, 2H), 7.45 (d, J = 8.5 Hz, 2H), 3.13 (s, 3H), 2.97 (s, 3H), 1.34 (s, 9H).

¹³C NMR (125 MHz, CDCl₃) δ = 209.0, 170.7, 139.7, 138.5, 127.8, 126.8, 44.3, 39.4, 35.3, 27.9.

IR (ATR) 2933.7, 1669.4, 1621.2, 1508.3, 1392.6, 1262.4, 1168.9.

HRMS (FD) Calcd for C₁₄H₁₉NO₂ [M]⁺:233.1416; Found: 233.1410

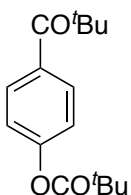


1j (pale yellow oil, 1.32 g, 7.03 mmol, 70%)

Synthesized according to the procedure **B**.

¹H NMR (500 MHz, CDCl₃) δ = 7.71 (s, 4H), 1.33 (s, 9H).

Spectral data were in good agreement with literature values.^{S5}

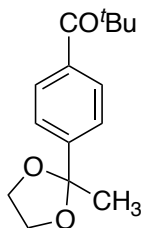


1k (white solids, 4.29 g, 16.3 mmol, 82%)

Synthesized according to the procedure **B**.

¹H NMR (500 MHz, CDCl₃) δ = 7.79 (dd, J = 8.5, 2.5 Hz, 2H), 7.10 (dd, J = 9.5, 2.3 Hz, 2H), 1.37 (s, 9H). 1.36 (s, 9H).

Spectral data were in good agreement with literature values.^{S8}



1l (white solid, 1.28 g, 5.14 mmol, 26%)

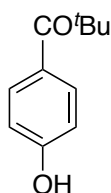
Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.68 (d, J = 8.0 Hz, 2H), 7.51 (d, J = 8.0 Hz, 2H), 4.00-4.09 (m, 2H), 3.81-3.75 (m, 2H), 1.66 (s, 3H), 1.35 (s, 9H).

¹³C NMR (125 MHz, CDCl₃) δ = 208.9, 146.1, 138.1, 128.0, 125.0, 108.5, 64.5, 44.2, 28.0, 27.4.

IR (ATR) 2971.3, 2935.7, 2883.6, 1669.4, 1605.7, 1568.1, 1465.9, 1401.3, 1369.5, 1273.0, 1242.2, 1220.0, 1191.0, 1175.6, 1142.8.

HRMS (ESI) Calcd for C₁₅H₂₀NaO₃ [M+Na]⁺: 271.1305; Found: 271.1317

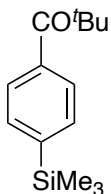


1m (white solids, 1.87 g, 10.5 mmol, 52%)

Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.71 (dt, J = 8.5 Hz, 2H), 6.82 (d, J = 8.5 Hz, 2H), 4.86 (s, 1H), 1.33 (s, 9H).

Spectral data were in good agreement with literature values.^{S4}

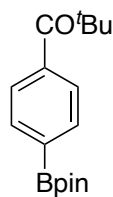


1n (colorless oil, 1.27 g, 5.42 mmol, 27%)

Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.66 (d, J = 8.5 Hz, 2H), 7.55 (d, J = 8.0 Hz, 2H), 1.35 (s, 9H), 0.28 (s, 9H).

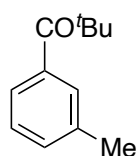
Spectral data were in good agreement with literature values.^{S9}



1o (pale yellow solids, 0.820 g, 2.85 mmol, 14%)

¹H NMR (500 MHz, CDCl₃) δ = 7.83 (d, J = 8.0 Hz, 2H), 7.61 (d, J = 9.0 Hz, 2H), 1.35 (s, 12H), 1.33 (s, 9H).

Spectral data were in good agreement with literature values.^{S10}

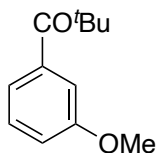


1p (colorless oil, 0.654 g, 3.71 mmol, 37%)

Synthesized according to the procedure **A** using 10 mmol of Mg.

¹H NMR (500 MHz, CDCl₃) δ = 7.50-7.45 (m, 2H), 7.30-7.25 (m, 2H), 2.39 (s, 3H), 1.34 (s, 9H).

Spectral data were in good agreement with literature values.^{S11}

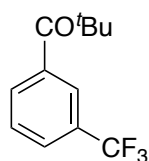


1q (colorless oil, 2.85 g, 14.8 mmol, 74%)

Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.33-7.26 (m, 2H), 7.19 (s, 1H), 7.00 (d, J = 8.0 Hz, 1H), 3.84 (s, 3H), 1.35 (s, 9H).

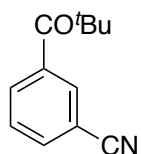
Spectral data were in good agreement with literature values.^{S6}



1r (colorless oil, 1.28 g, 5.56 mmol, 56%)

¹H NMR (500 MHz, CDCl₃) δ = 7.93 (s, 1H), 7.87 (d, J = 8.0 Hz, 1H), 7.73 (d, J = 8.0 Hz, 1H), 7.55 (t, J = 8.0, 1H), 1.36 (s, 9H).

Spectral data were in good agreement with literature values.^{S12}



1s (pale yellow oil, 2.82 g, 15.0 mmol, 69%)

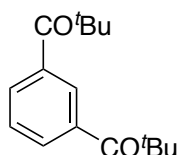
Synthesized according to the procedure **B** using 21.8 mmol of 3-iodobenzonitrile.

¹H NMR (500 MHz, CDCl₃) δ = 7.95 (s, 1H), 7.90 (d, J = 8.0 Hz, 1H), 7.75 (d, J = 8.0 Hz, 1H), 7.55 (t, J = 8.0, 1H), 1.35 (s, 9H).

¹³C NMR (125 MHz, CDCl₃) δ = 207.1, 139.5, 134.0, 131.9, 131.4, 129.1, 118.1, 112.6, 44.4, 27.8.

IR (ATR) 2972.3, 2873.9, 2231.6, 1680.0, 1477.5, 1367.5, 1276.9, 1215.2, 1149.6

The satisfactory data for HRMS analysis was not obtained by ESI- and FD-MS.

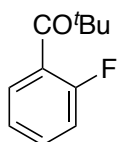


1t (white solids, 1.35 g, 15.0 mmol, 69%)

Synthesized according to the procedure **A** using 40 mmol of Mg and 10 mL of pivaloyl chloride.

¹H NMR (500 MHz, CDCl₃) δ = 8.03 (s, 1H), 7.79 (dd, J = 8.0, 2.0 Hz, 2H), 7.46 (t, J = 8.0 Hz, 1H), 1.37 (s, 18H).

Spectral data were in good agreement with literature values.^{S13}

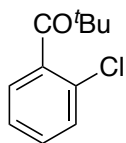


1u (colorless oil, 1.48 g, 8.22 mmol, 41%)

Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.40-7.34 (m, 1H), 7.21-7.13 (m, 2H), 7.09 (t, J = 9.0 Hz, 1H), 1.23 (s, 9H).

Spectral data were in good agreement with literature values.^{S14}

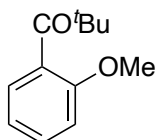


1v (colorless oil, 2.10 g, 10.7 mmol, 54%)

Synthesized according to the procedure **B**.

¹H NMR (500 MHz, CDCl₃) δ = 7.39 (dd, J = 7.8, 1.4 Hz, 1H), 7.31 (td, J = 7.9, 2.0 Hz, 1H), 7.27 (td, J = 7.6, 1.5 Hz, 1H), 7.15 (dd, J = 7.5, 1.7 Hz, 1H), 1.27 (s, 9H).

Spectral data were in good agreement with literature values.^{S15}



1w (colorless oil, 2.10 g, 10.7 mmol, 54%)

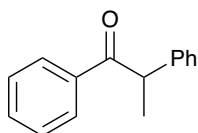
Synthesized according to the procedure **A**.

¹H NMR (500 MHz, CDCl₃) δ = 7.31 (t, J = 8.0 Hz, 1H), 7.02 (d, J = 7.5 Hz, 1H), 6.94 (t, J = 7.5 Hz, 1H), 6.90 (d, J = 8.5 Hz, 1H), 3.78 (s, 3H), 1.21 (s, 9H).

Spectral data were in good agreement with literature values.^{S11}

General procedures for the preparation of alkyl phenyl ketones

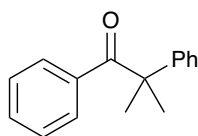
Alkyl phenyl ketones **3a**, **3b**, **3f** and **3g** were obtained from chemical suppliers and used after proper recrystallization or distillation. **3c-e** and **3h** were synthesized according to literature procedures. All ketones are known compounds in literatures as referenced, and spectral data were in good agreement with literature values.



3c (pale yellow solids, 1.38 g, 6.56 mmol, 33%)^{S16}

¹H NMR (500 MHz, CDCl₃) δ = 7.95 (d, J = 7.5 Hz, 2H), 7.47 (t, J = 7.5 Hz, 1H), 7.38 (t, J = 8.0 Hz, 2H), 7.32-7.27 (m, 4H), 7.23-7.17 (m, 1H), 4.69 (q, J = 7.0 Hz, 1H), 1.53 (d, J = 7.0 Hz, 3H).

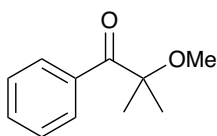
Spectral data were in good agreement with literature values.^{S16}



3d (white solids, 1.52 g, 6.78 mmol, 33%)^{S17}

¹H NMR (500 MHz, CDCl₃) δ = 7.48 (d, J = 7.5 Hz, 2H), 7.38-7.30 (m, 5H), 7.29-7.25 (m, 1H), 7.22 (t, J = 7.5 Hz, 2H), 1.61 (s, 6H).

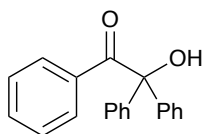
Spectral data were in good agreement with literature values.^{S17}



3e (white solids, 3.27 g, 18.3 mmol, 80%)^{S18}

¹H NMR (500 MHz, CDCl₃) δ = 8.28 (dd, J = 8.5, 1.0 Hz, 2H), 7.55 (t, J = 7.5 Hz, 1H), 7.44 (t, J = 7.5 Hz, 2H), 3.20 (s, 3H), 1.52 (s, 6H).

Spectral data were in good agreement with literature values.^{S18}

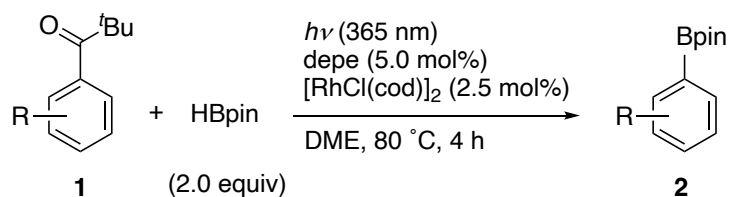


3h (white solids, 3.91 g, 13.6 mmol, 36%)^{S19}

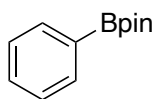
¹H NMR (500 MHz, CDCl₃) δ = 7.71 (dd, J = 8.5, 1.0 Hz, 2H), 7.47-7.38 (m, 5H), 7.37-7.27 (m, 8H), 4.98 (s, 1H).

Spectral data were in good agreement with literature values.^{S19}

A general procedure for the borylation of pivalophenone derivatives (Table 2)



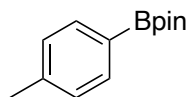
A solution of $[\text{RhCl(cod)}]_2$ (6.2 mg, 0.0125 mmol), depe (5.9 μL , 0.025 mmol), a pivalophenone derivative (0.5 mmol) and HBpin (145.0 μL , 1.0 mmol) in DME (5 mL) was placed in a sealed glass tube. The solution was photoirradiated at 365 nm with stirring at 80 $^\circ\text{C}$. After 4 h, the solvent was removed under reduced pressure to give a crude mixture, in which the yields of arylboronates were determined by ^1H NMR using tetrachloroethane as an internal standard or GC analysis using dodecane as an internal standard. The crude product was purified by silica gel column chromatography or PTLC to afford arylboronates **2a-w**.



2a (white solids, 59.2 mg, 0.29 mmol, 58%) (NMR yield = 68%)

^1H NMR (500 MHz, CDCl_3) δ = 7.81 (d, J = 6.9 Hz, 2H), 7.46 (t, J = 7.4 Hz, 1H), 7.37 (t, J = 7.4 Hz, 2H), 1.35 (s, 12H).

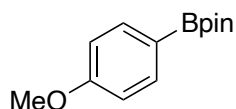
Spectral data were in good agreement with literature values.^{S20}



2b (white solids, 67.6 mg, 0.31 mmol, 62%) (NMR yield = 83%)

^1H NMR (500 MHz, CDCl_3) δ = 7.70 (d, J = 7.5 Hz, 2H), 7.18 (d, J = 8.0 Hz, 2H), 2.36 (s, 3H), 1.33 (s, 12H).

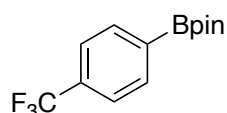
Spectral data were in good agreement with literature values.^{S20}



2c (white solids, 82.7 mg, 0.35 mmol, 71%) (NMR yield = 81%)

^1H NMR (500 MHz, CDCl_3) δ = 7.75 (d, J = 8.0 Hz, 2H), 6.90 (d, J = 8.5 Hz, 2H), 3.83 (s, 3H), 1.33 (s, 12H).

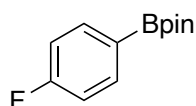
Spectral data were in good agreement with literature values.^{S20}



2d (white solids, 73.2 mg, 0.27 mmol, 55%) (NMR yield = 72%)

^1H NMR (500 MHz, CDCl_3) δ = 7.91 (d, J = 8.0 Hz, 2H), 7.61 (d, J = 8.0 Hz, 2H), 1.36 (s, 12H).

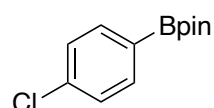
Spectral data were in good agreement with literature values.^{S20}



2e (colorless oil, 69.7 mg, 0.31 mmol, 63%) (NMR yield = 84%)

^1H NMR (500 MHz, CDCl_3) δ = 7.80 (dd, J = 8.0, 6.0 Hz, 2H), 7.04 (t, J = 9.0 Hz, 2H), 1.33 (s, 12H).

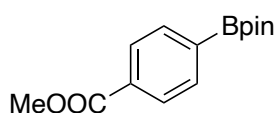
Spectral data were in good agreement with literature values.^{S20}



2f (white solids, 65.9 mg, 0.28 mmol, 55%) (NMR yield = 71%)

^1H NMR (500 MHz, CDCl_3) δ = 7.73 (d, J = 8.5 Hz, 2H), 7.33 (d, J = 8.5 Hz, 2H), 1.33 (s, 12H).

Spectral data were in good agreement with literature values.^{S20}

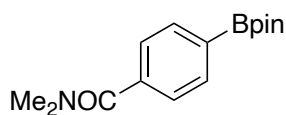


2h

2h (white solids, 59.4 mg, 0.23 mmol, 45%) (NMR yield = 66%)

^1H NMR (500 MHz, CDCl_3) δ = 8.02 (d, J = 8.0 Hz, 2H), 7.87 (d, J = 8.5 Hz, 2H), 3.92 (s, 3H), 1.36 (s, 12H).

Spectral data were in good agreement with literature values.^{S20}

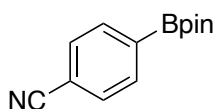


2i

2i (white solids, 110.4 mg, 0.4012 mmol, 80%) (NMR yield = 89%)

^1H NMR (500 MHz, CDCl_3) δ = 7.84 (d, J = 8.0 Hz, 2H), 7.40 (d, J = 8.0 Hz, 2H), 3.11 (s, 3H), 2.95 (s, 3H), 1.35 (s, 12H).

Spectral data were in good agreement with literature values.^{S23}

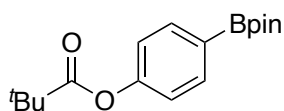


2j

2j was obtained as an inseparable mixture with **1j** (52.7 mg). The amount of **2j** was calculated to be 0.209 mmol (42%) from the ^1H NMR. (NMR yield = 58%)

^1H NMR (500 MHz, CDCl_3) δ = 7.86 (d, J = 8.5 Hz, 2H), 7.62 (d, J = 9.0 Hz, 2H), 1.33 (s, 12H).

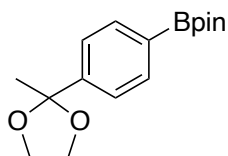
Spectral data were in good agreement with literature values.^{S20}



2k (white solids, 76.0 mg, 0.250 mmol, 50%) (NMR yield = 53%)

^1H NMR (500 MHz, CDCl_3) δ = 7.83 (d, J = 8.5 Hz, 2H), 7.06 (d, J = 8.5 Hz, 2H), 1.35 (s, 9H), 1.34 (s, 12H).

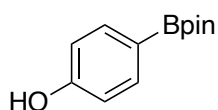
Spectral data were in good agreement with literature values.^{S8}



2l (white solids, 71.7 mg, 0.25 mmol, 50%) (NMR yield = 76%)

^1H NMR (500 MHz, CDCl_3) δ = 7.80 (d, J = 8.0 Hz, 2H), 7.49 (d, J = 8.0 Hz, 2H), 4.05-4.02 (m, 2H), 3.78-3.72 (m, 2H), 1.64 (s, 3H), 1.34 (s, 12H).

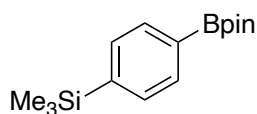
Spectral data were in good agreement with literature values.^{S24}



2m (white solids, 94.9 mg, 0.43 mmol, 86%) (NMR yield = 76%)

^1H NMR (500 MHz, CDCl_3) δ = 7.69 (d, J = 8.5 Hz, 2H), 6.82 (d, J = 8.5 Hz, 2H), 6.48 (s, 1H), 1.34 (s, 12H).

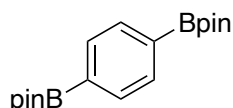
Spectral data were in good agreement with literature values.^{S20}



2n (white solids, 77.6 mg, 0.28 mmol, 56%) (NMR yield = 53%)

^1H NMR (500 MHz, CDCl_3) δ = 7.83 (d, J = 8.0 Hz, 2H), 7.57 (d, J = 8.0 Hz, 2H), 1.37 (s, 12H), 0.30 (s, 9H).

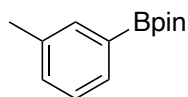
Spectral data were in good agreement with literature values.^{S22}



2o (white solids, 108.8 mg, 0.33 mmol, 66%) (NMR yield = quant.)

^1H NMR (500 MHz, CDCl_3) δ = 7.80 (s, 4H), 1.35 (s, 24H).

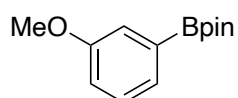
Spectral data were in good agreement with literature values.^{S20}



2p (white solids, 68.8 mg, 0.32 mmol, 63%) (NMR yield = 78%)

^1H NMR (500 MHz, CDCl_3) δ = 7.64 (s, 1H), 7.65-7.60 (m, 1H), 7.27–7.25 (m, 2H), 2.35 (s, 3H), 1.34 (s, 12H).

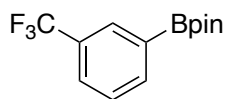
Spectral data were in good agreement with literature values.^{S20}



2q (white solids, 73.6 mg, 0.31 mmol, 63%) (NMR yield = 76%)

^1H NMR (500 MHz, CDCl_3) δ = 7.40 (d, J = 7.0 Hz, 1H), 7.33 (d, J = 2.5 Hz, 1H), 7.29 (t, J = 8.0 Hz, 1H), 8.01 (ddd, J = 8.0, 3.0, 1.0 Hz, 1H), 3.83 (s, 3H), 1.35 (s, 12H).

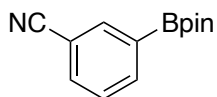
Spectral data were in good agreement with literature values.^{S22}



2r (white solids, 110.2 mg, 0.41 mmol, 81%) (NMR yield = 76%)

^1H NMR (500 MHz, CDCl_3) δ = 8.06 (s, 1H), 7.97 (d, J = 7.0 Hz, 1H), 7.70 (d, J = 8.0 Hz, 1H), 7.49 (t, J = 7.5 Hz, 1H), 1.36 (s, 12H).

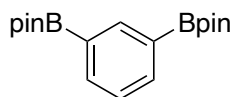
Spectral data were in good agreement with literature values.^{S24}



2s (white solids, 52.4 mg, 0.229 mmol, 46%) (NMR yield = 57%)

^1H NMR (500 MHz, CDCl_3) δ = 8.10 (s, 1H), 8.01 (d, J = 7.5 Hz, 1H), 7.72 (d, J = 7.5 Hz, 1H), 7.47 (t, J = 8.0 Hz, 1H), 1.36 (s, 12H).

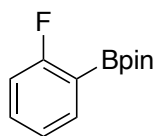
Spectral data were in good agreement with literature values.^{S20}



2t (white solids, 35.7 mg, 0.108 mmol, 22%) (NMR yield = 49%)

^1H NMR (500 MHz, CDCl_3) δ = 8.29 (s, 1H), 7.90 (dd, J = 7.5, 2.0 Hz, 2H), 7.38 (t, J = 7.5 Hz, 1H), 1.34 (s, 24H).

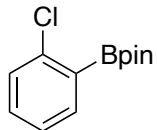
Spectral data were in good agreement with literature values.^{S20}



2u (white solids, 71.3 mg, 0.321 mmol, 64%) (NMR yield = 79%)

^1H NMR (500 MHz, CDCl_3) δ = 7.76–7.72 (m, 1H), 7.46–7.41 (m, 1H), 7.14 (t, J = 7.5 Hz, 1H), 7.03 (t, J = 8.5 Hz, 1H), 1.37 (s, 12H).

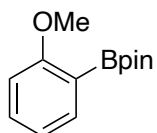
Spectral data were in good agreement with literature values.^{S20}



2v (white solids, 74.5 mg, 0.312 mmol, 62%) (NMR yield = 80%)

^1H NMR (500 MHz, CDCl_3) δ = 7.68 (d, J = 8.0 Hz, 1H), 7.40–7.29 (m, 2H), 7.24–7.19 (m, 1H), 1.36 (s, 12H).

Spectral data were in good agreement with literature values.^{S25}

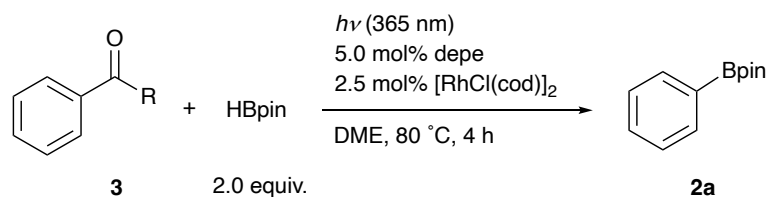


2w (white solids, 65.1 mg, 0.28 mmol, 56%) (NMR yield = 80%)

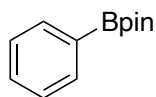
^1H NMR (500 MHz, CDCl_3) δ = 7.68 (dd, J = 9.0, 1.7 Hz, 1H), 7.41–7.37 (m, 1H), 6.94 (t, J = 8.0 Hz, 1H), 6.85 (dd, J = 8.0, 1 H), 3.82 (s, 3H), 1.35 (s, 12H).

Spectral data were in good agreement with literature values.^{S20}

A general procedure for the borylation of alkyl phenyl ketones (Table 3)



A solution of $[\text{RhCl}(\text{cod})]_2$ (6.2 mg, 0.0125 mmol), depe (5.9 μL , 0.025 mmol), an alkyl phenyl ketone (0.50 mmol) and HBpin (145.0 μL , 1.0 mmol) in DME (5 mL) was placed in a sealed glass tube. The solution was photoirradiated at 365 nm with stirring at 80 $^\circ\text{C}$. After 4 h, the solvent was removed under reduced pressure to give a crude mixture, in which the yield of phenylboronic acid pinacol ester **2a** was determined by ^1H NMR using tetrachloroethane as an internal standard. The crude product was purified by silica gel column chromatography to afford phenylboronic acid pinacol ester **2a**. For the reaction of **3g** and **3h**, separation of **2a** from **3** was difficult. Therefore, the crude mixture was treated with KHF_2 to convert **2a** to PhBF_3K **2a'**, which was isolated by washing with hexanes and a small amount of cold MeOH (ca. -20 $^\circ\text{C}$).



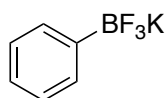
2a from **3b** (white solids, 16.3 mg, 0.080 mmol, 16%) (NMR yield = 30%)

2a from **3c** (white solids, 48.6 mg, 0.238 mmol, 48%) (NMR yield = 45%)

2a from **3d** (white solids, 47.0 mg, 0.230 mmol, 46%) (NMR yield = 59%)

2a from **3e** (white solids, 37.6 mg, 0.184 mmol, 37%) (NMR yield = 43%)

2a from **3f** (white solids, 10.2 mg, 0.050 mmol, 10%) (NMR yield = 17%)



2a' from **3g** (white solids, 5.8 mg, 0.032 mmol, 6%) (NMR yield = 16%)

2a' from **3h** (white solids, 13.8 mg, 0.075 mmol, 15%) (NMR yield = 26%)

^1H NMR (500 MHz, CDCl_3) δ = 7.81 (d, J = 6.9 Hz, 2H), 7.46 (t, J = 7.4 Hz, 1H), 7.37 (t, J = 7.4 Hz, 2H), 1.35 (s, 12H).

Spectral data were in good agreement with literature values.^{S21}

Optimization of reaction conditions

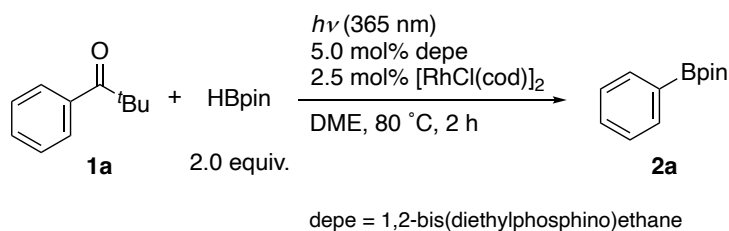


Table S1. Optimization of reaction conditions.

entry	variation from the standard conditions	yield of 2a ^a
1	—	68% (58%) ^b
2	dmpe instead of depe	19%
3	dippe instead of depe	12%
4	dtbpe instead of depe	12%
5	dppp instead of depe	54%
6	dppbz instead of depe	19%
7	depbz instead of depe	19%
8	DMA instead of DME	41%
9	toluene instead of DME	61%
10	[Rh(cod)I] ₂ instead of [Rh(cod)Cl] ₂	58%
11	[Rh(ethylene) ₂ Cl] ₂ instead of [Rh(cod)Cl] ₂	53%
12	[Rh(coe) ₂ Cl] ₂ instead of [Rh(cod)Cl] ₂	50%

^a Determined by NMR using 1,1,2,2-tetrachloroethane as an internal standard. ^b Isolated yield.

dmpe = 1,2-bis(dimethylphosphino)ethane

dippe = 1,2-bis(di-*i*-propylphosphino)ethane

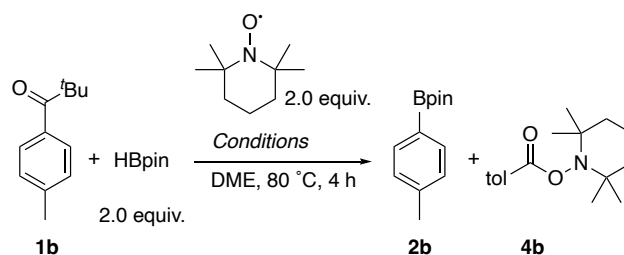
dtbpe = 1,2-bis(di-*tert*-butylphosphino)ethane

dppe = 1,2-bis(diphenylphosphino)ethane

dppbz = 1,2-bis(diphenylphosphino)benzene

depbz = 1,2-bis(diethylphosphino)benzene

Procedures for radical trapping experiments (Table 4)



For the experiment in entry 1, a solution of $[\text{RhCl}(\text{cod})]_2$ (2.5 mg, 0.0050 mmol), depe (2.4 μL , 0.01 mmol), *tert*-butyl *p*-tolyl ketone **1b** (35.3 mg, 0.20 mmol), TEMPO (62.5 mg, 0.40 mmol) and HBpin (58.0 μL , 0.4 mmol) in DME (2.0 mL) was placed in a sealed glass tube. The solution was photoirradiated at 365 nm with stirring at 80 °C for 4 hours. After removal of the solvent and volatile compounds, the crude mixture was analyzed by ^1H NMR in CDCl_3 to determine the yield of **2b** and **4b** using dibromomethane (14.0 μL , 0.20 mmol) as an internal standard. Spectral data were in good agreement with literature values.^{S20, S25} Other radical trapping experiments in the absence of the Rh catalyst (entry 2) or without photoirradiation (entry 3) were also conducted according to the same procedure.

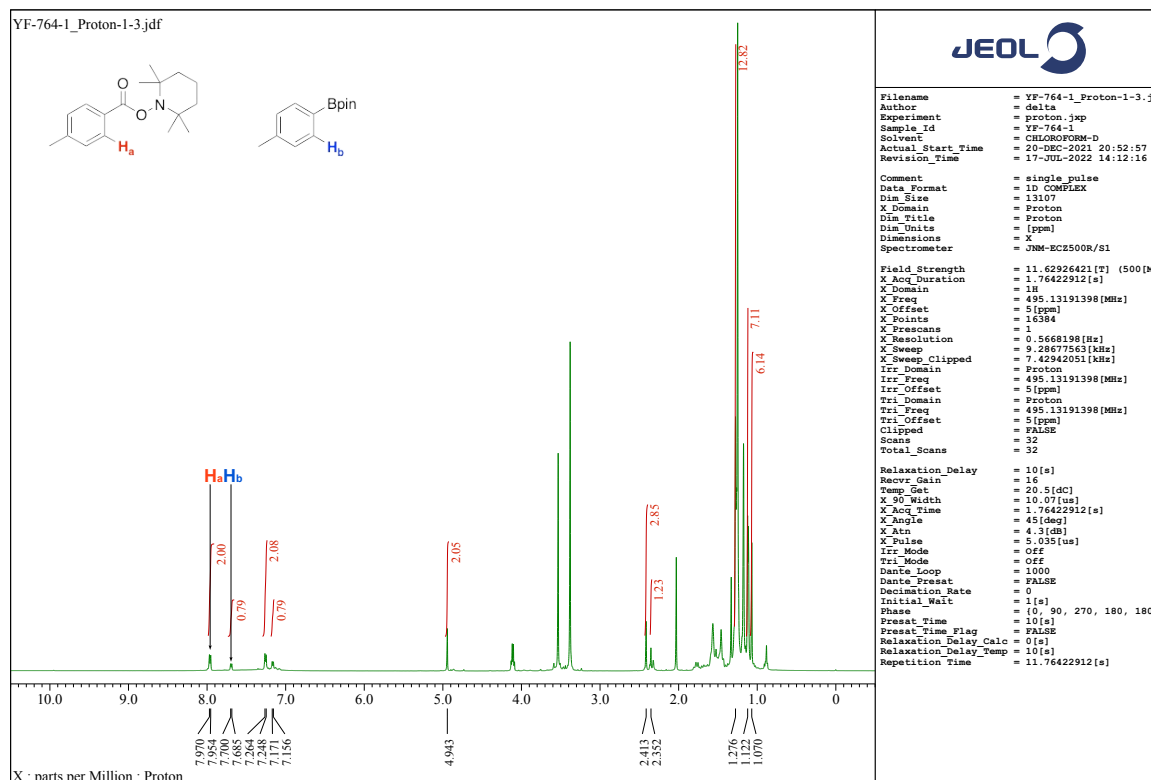


Figure S2. The radical trapping experiment under the standard conditions (entry 1).

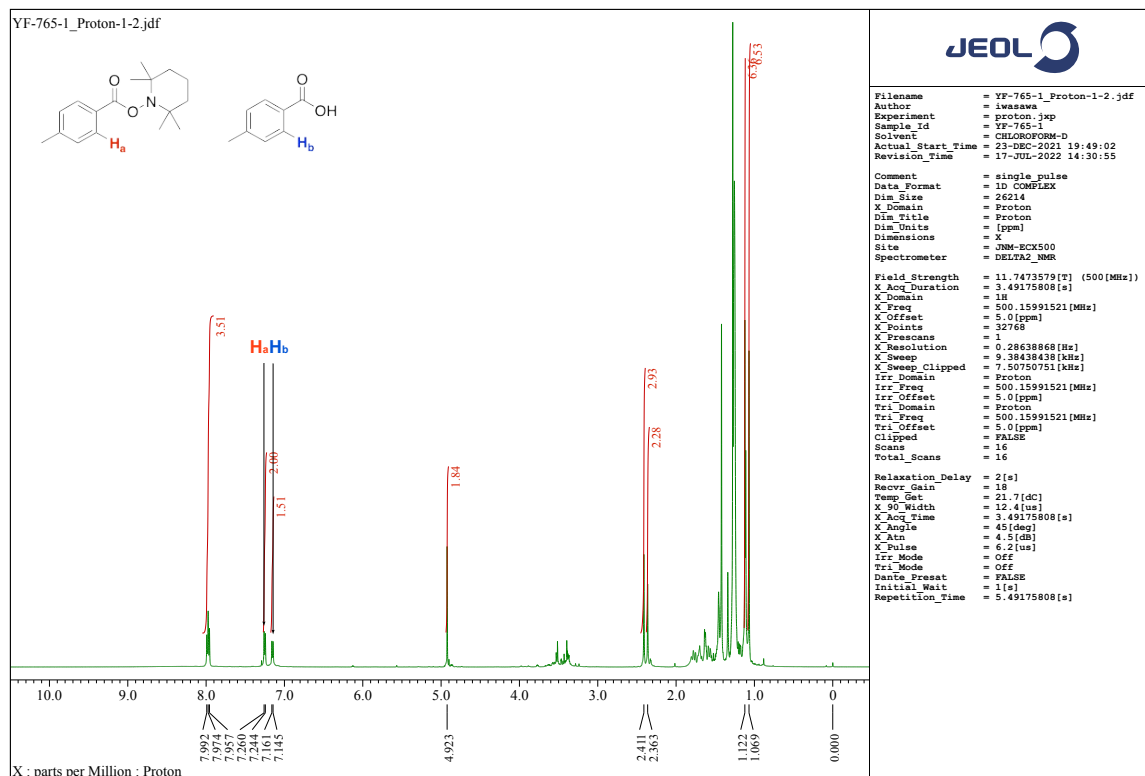


Figure S3. The radical trapping experiment without $[\text{RhCl}(\text{cod})]_2$ and depe (entry 2).^a

^a We confirmed that **4b** further reacted with HBpin to generate *p*-methylbenzoic acid under the reaction conditions. Therefore, the yield of the radical-TEMPO adduct was calculated to be the combined yield of **4b** and *p*-methylbenzoic acid.

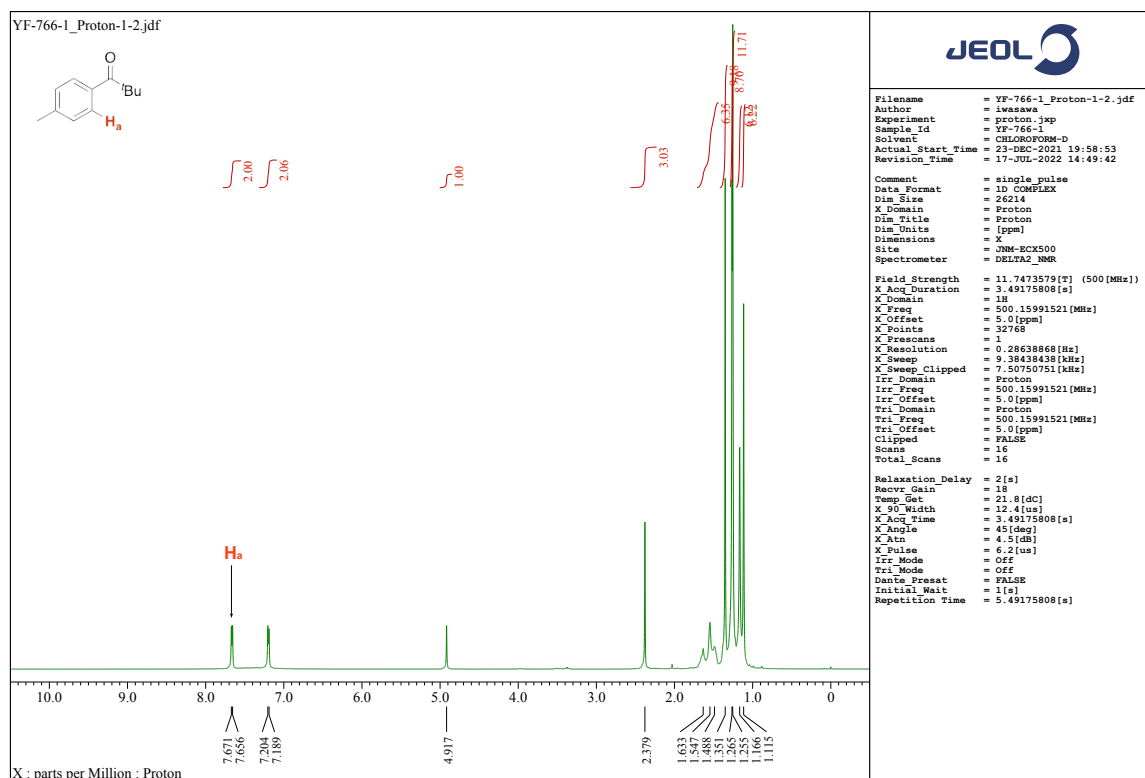
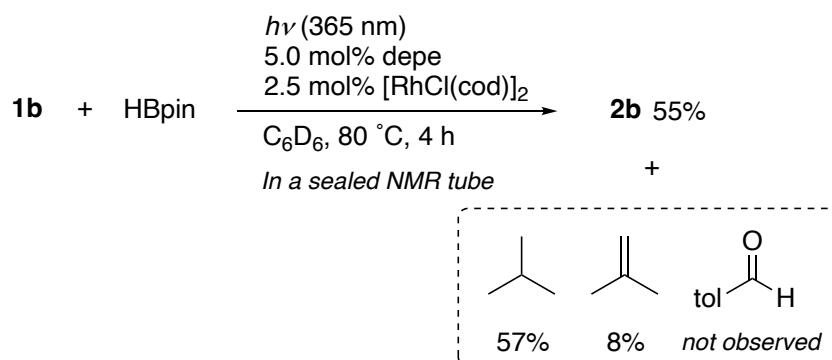


Figure S4. The radical trapping experiment without $h\nu$ (entry 3).

An NMR experiment in a sealed NMR tube (Scheme 1)



A solution of [RhCl(cod)]₂ (1.2 mg, 0.0025 mmol), depe (1.2 μ L, 0.005 mmol), pivalophenone **1a** (16.7 mg, 0.10 mmol), and HBpin (58 μ L, 0.20 mmol) in C₆D₆ (0.5 mL) was placed in a sealed NMR tube. The solution was photoirradiated at 365 nm without stirring at 80 °C for 4 hours. The solution was analyzed by ¹H NMR, demonstrating the formation of 2-methylpropane and 2-methylpropene as major by-products. The yields were determined after adding 1,1,2,2-tetrachloroethane quickly into the NMR tube as an internal standard.

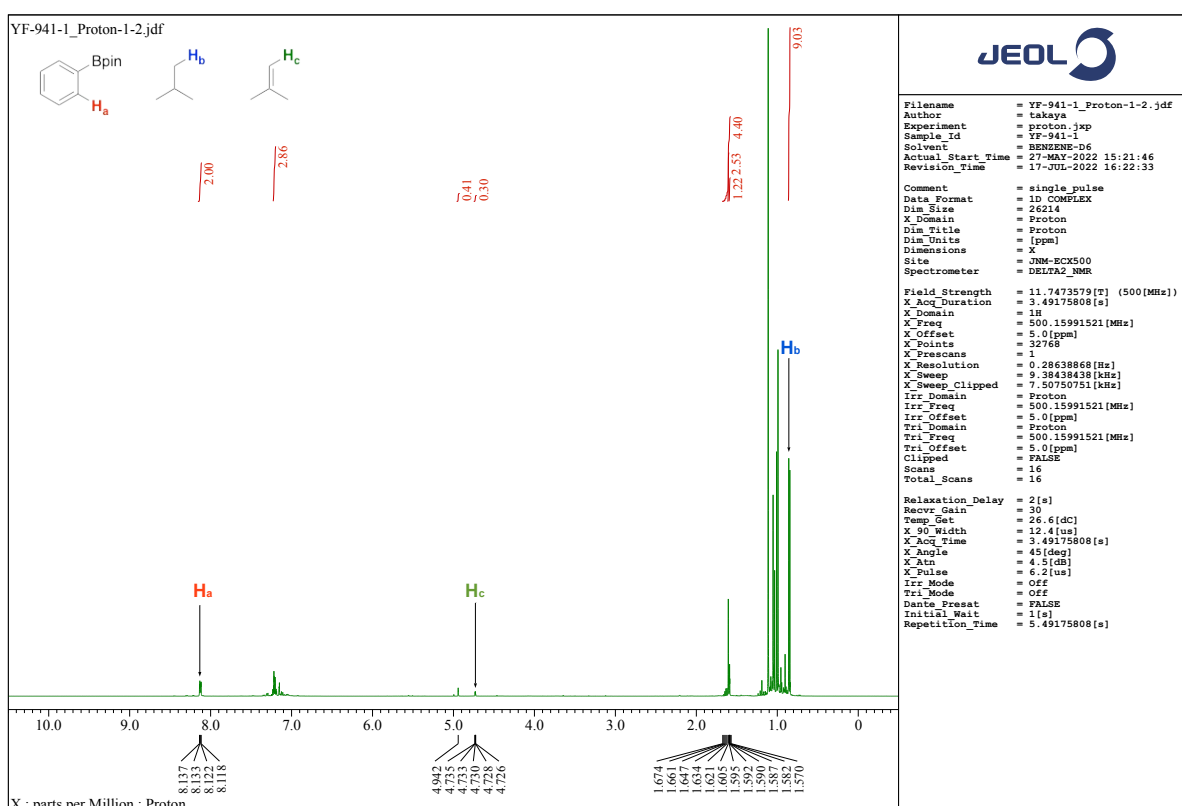
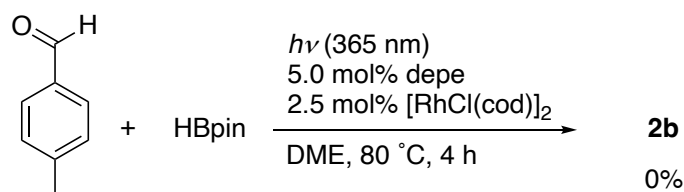


Figure S5. ¹H NMR after the reaction (NMR experiment in C₆D₆)

A reaction of aldehyde (Scheme 2)



A solution of [RhCl(cod)]₂ (6.2 mg, 0.012 mmol), depe (2.4 μL, 0.025 mmol), *p*-tolualdehyde (59.6 mg, 0.496 mmol) and HBpin (145 μL, 1.0 mmol) in DME (5.0 mL) was placed in a sealed glass tube. The solution was photoirradiated at 365 nm with stirring at 80 °C for 4 hours. After removal of the solvent and volatile compounds, the crude mixture was analyzed by ¹H NMR with 1,1,2,2-tetrachloroethane as an internal standard (50.0 μL, 0.472 mmol), supporting no formation of *p*-tolylboronate **2b** and formation of hydroboration product in 71% yield.

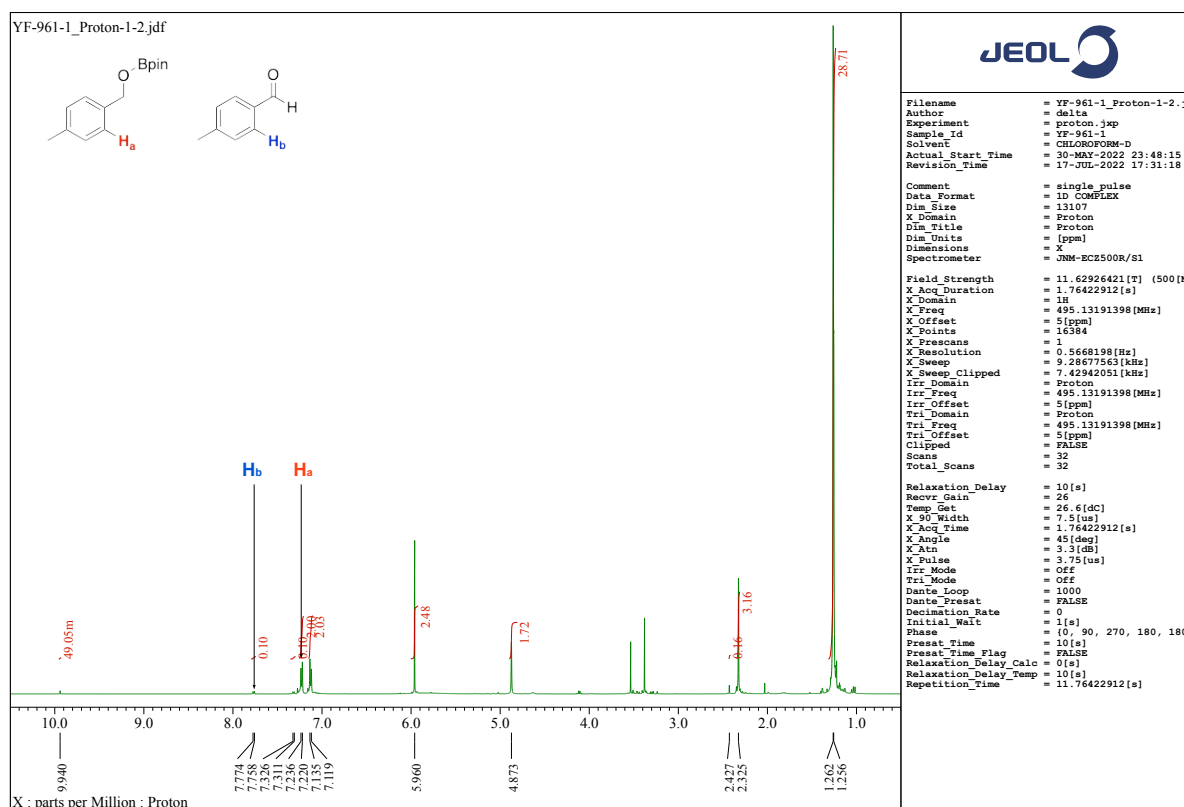
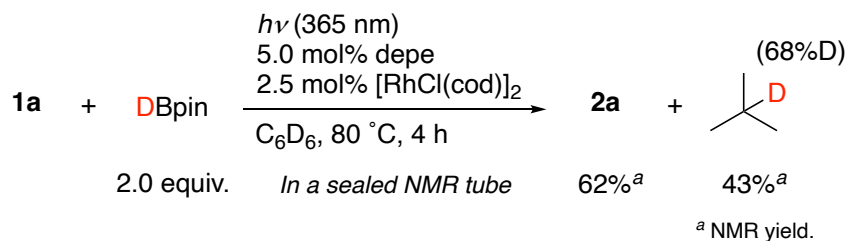


Figure S6. ^1H NMR of the reaction of *p*-tolylaldehyde under the standard conditions

A deuterium labeling experiment (Scheme 3)



A solution of [RhCl(cod)]₂ (1.2 mg, 0.0025 mmol), depe (1.2 μL, 0.0050 mmol), pivalophenone **1a** (16.7 mg, 0.10 mmol), and DBpin (51.6 mg, 0.20 mmol) in C₆D₆ (0.5 mL) was placed in a sealed NMR tube. The solution was photoirradiated at 365 nm without stirring at 80 °C for 4 hours. The solution was analyzed by ¹H NMR after adding 1,1,2,2-tetrachloroethane quickly, demonstrating the formation of **2a** in 62% yield and 2-methylpropane in 43% yield. The incorporation of a deuterium at the methine carbon was confirmed by ¹H NMR. The D-content of 2-methylpropane was determined to be 68% by EI-MS, in which the relative abundance of non-labeled 2-methylpropane (m/z 58) to D-labeled one (m/z 59, including a natural isotope of non-labeled 2-methylpropane) was 126628:268735.

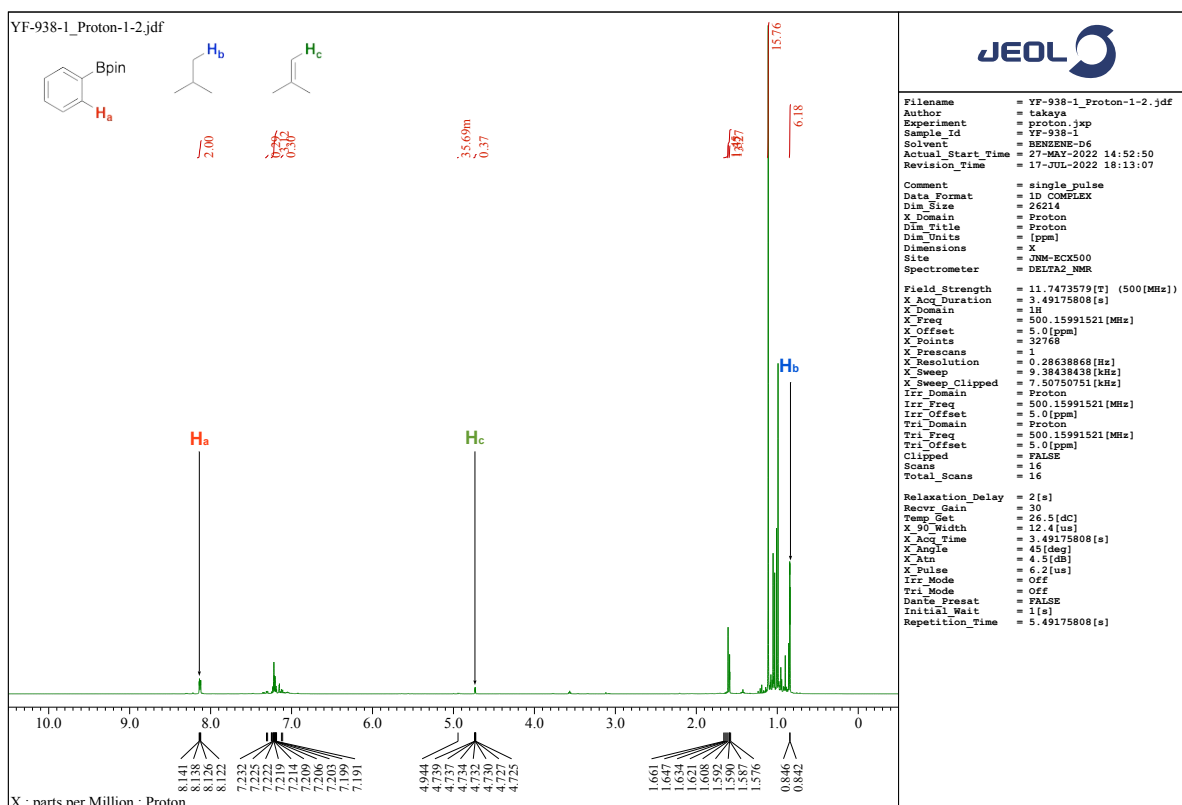
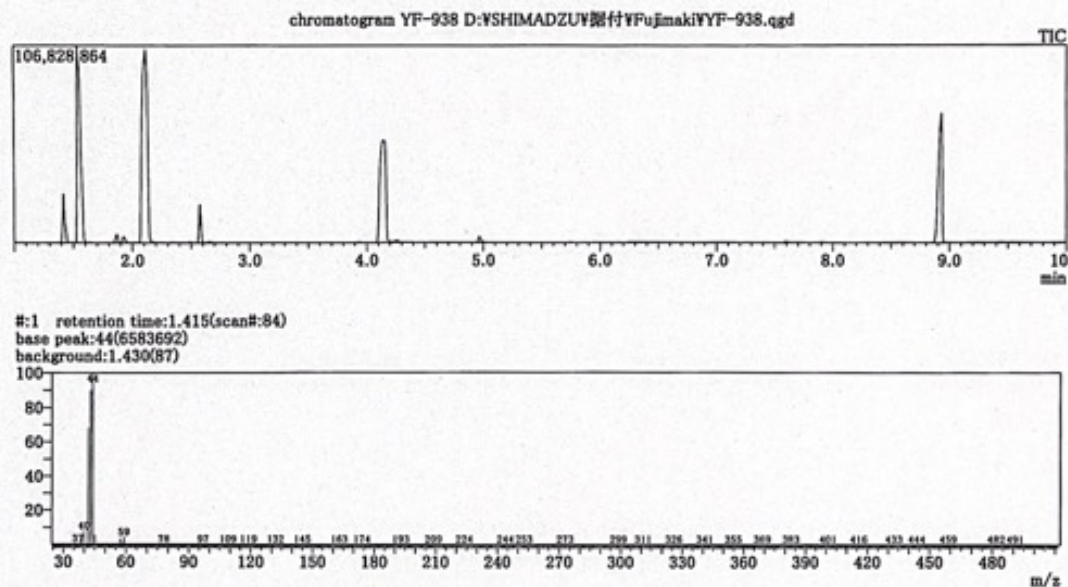


Figure S7. ^1H NMR of the deuterium labeling experiment before adding an internal standard

date : 2022/05/27 18:04:25
sample name : YF-938
date file : D:\SHIMADZUY\YF\YFujimaki\YF-938.qgd
method file : D:\SHIMADZUY\YF\YFujimaki\method_line1_standard_分子量500_1min-.qgm



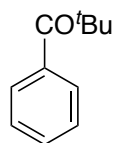
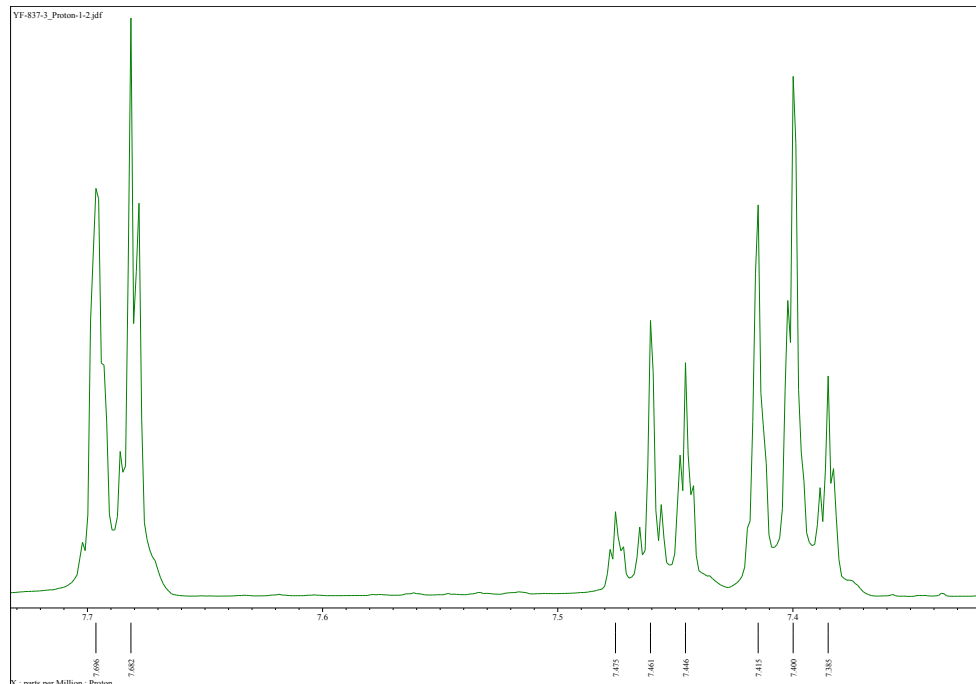
#	m/z	absolute intensity	relative intensity
1	37.05	23928	0.36
2	38.05	66514	1.01
3	39.05	242660	3.69
4	40.05	512871	7.79
5	41.05	44144	0.67
6	42.05	4483647	68.10
7	43.05	5923539	89.97
8	44.10	6583692	100.00
9	45.10	364925	5.54
10	46.10	4304	0.07
11	57.10	205296	3.12
12	58.10	126628	1.92
13	59.10	274307	4.17
14	60.10	11807	0.18
15	61.10	73	0.00

Figure S8. EI-MS spectra

References

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**1a**

X : parts per Million : Proton



----- PROCESSING PARAMETERS -----
sexp(0.2[Hz], 0.0[s])
trapezoid(0[%], 0[%], 80[%], 100[%])
zerofill(1, TRUE)
fft(1, TRUE, TRUE)
machinephase
ppm

以下に由来 : YF-837-3_Proton-1-1.jdf

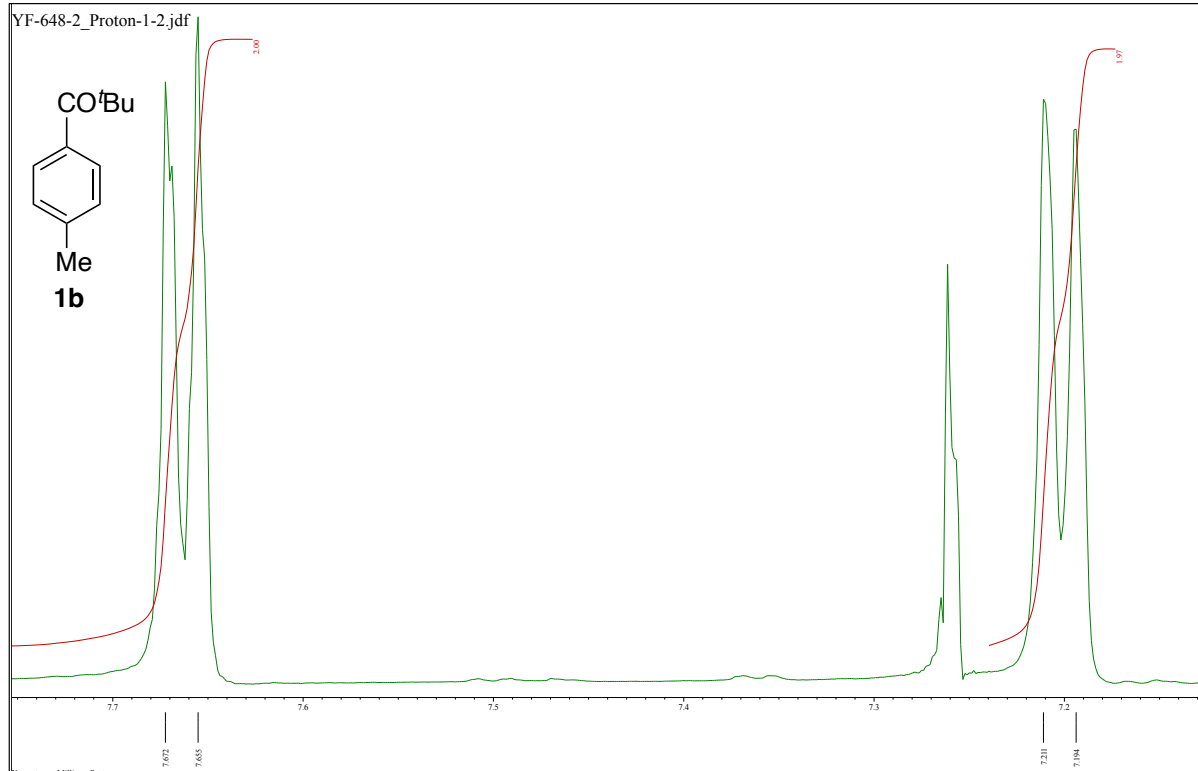
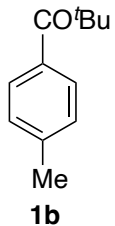
Filename = YF-837-3_Proton-1-2.j
Author = delta
Experiment = proton.jxp
Sample_Id = YF-837-3
Solvent = CHLOROFORM-D
Actual_Start_Time = 9-MAR-2022 22:18:57
Revision_Time = 5-JUL-2022 17:46:22

Comment = single_pulse
Data Format = 1D COMPLEX
Dim_Size = 13107
X_Domain = Proton
Dim Title = Proton
Dim Units = [ppm]
Dimensions = X
Spectrometer = JNM-ECZ500R/S1

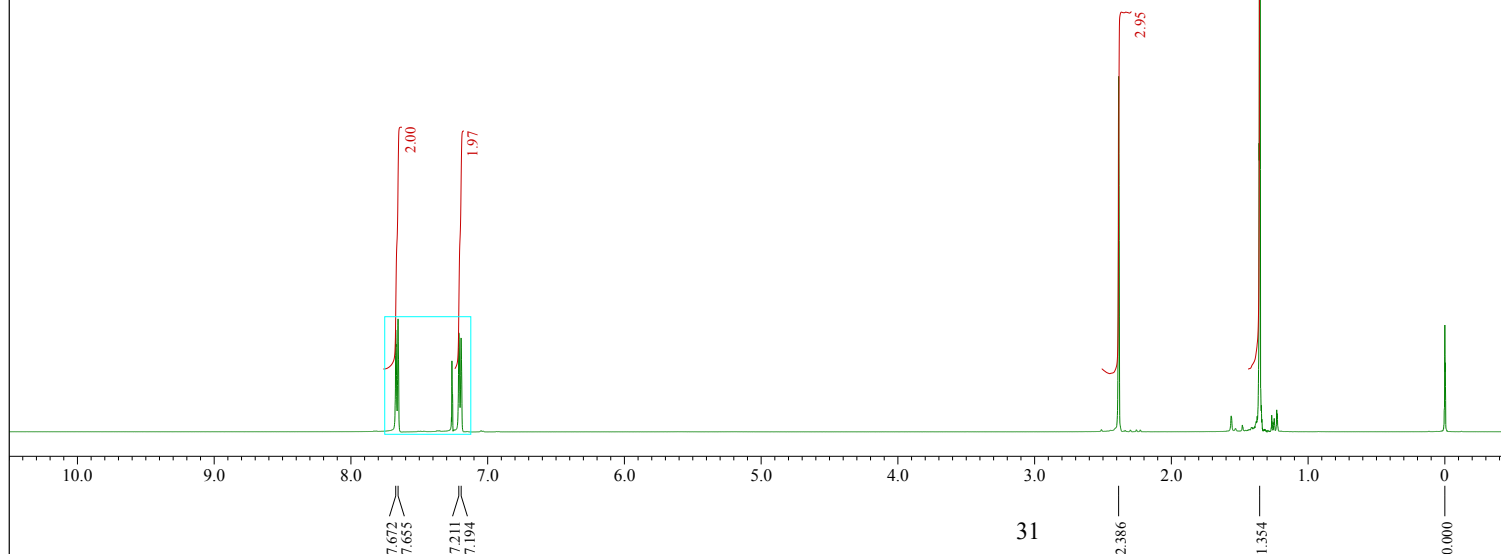
Field_Strength = 11.62926421[T] (500[M]
X_Acq_Duration = 1.76422912[s]
X_Domain = 1H
X_Freq = 495.13191398[MHz]
X_Offset = 5[ppm]
X Points = 16384
X_Prescans = 1
X_Resolution = 0.5668198[Hz]
X_Sweep = 9.28677563[kHz]
X_Sweep_Clippped = 7.42942051[kHz]
Irr_Domain = Proton
Irr_Freq = 495.13191398[MHz]
Irr_Offset = 5[ppm]
Tri_Domain = Proton
Tri_Freq = 495.13191398[MHz]
Tri_Offset = 5[ppm]
Clipped = FALSE
Scans = 32
Total_Scans = 32

Relaxation_Delay = 5[s]
Recvr Gain = 36
Temp_Get = 20.6[dC]
X_90_Width = 9.47[us]
X_Acq_Time = 1.76422912[s]
X_Angle = 45[deg]
X_Atn = 4.3[dB]
X_Pulse = 4.735[us]
Irr_Mode = Off
Tri_Mode = Off
Dante_Loop = 500
Dante_Presat = FALSE
Decimation_Rate = 0
Initial_Wait = 1[s]
Phase = {0, 90, 270, 180, 180
Presat_Time = 5[s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 5[s]
Repetition_Time = 6.76422912[s]

YF-648-2_Proton-1-2.jdf



X : parts per Million : Proton



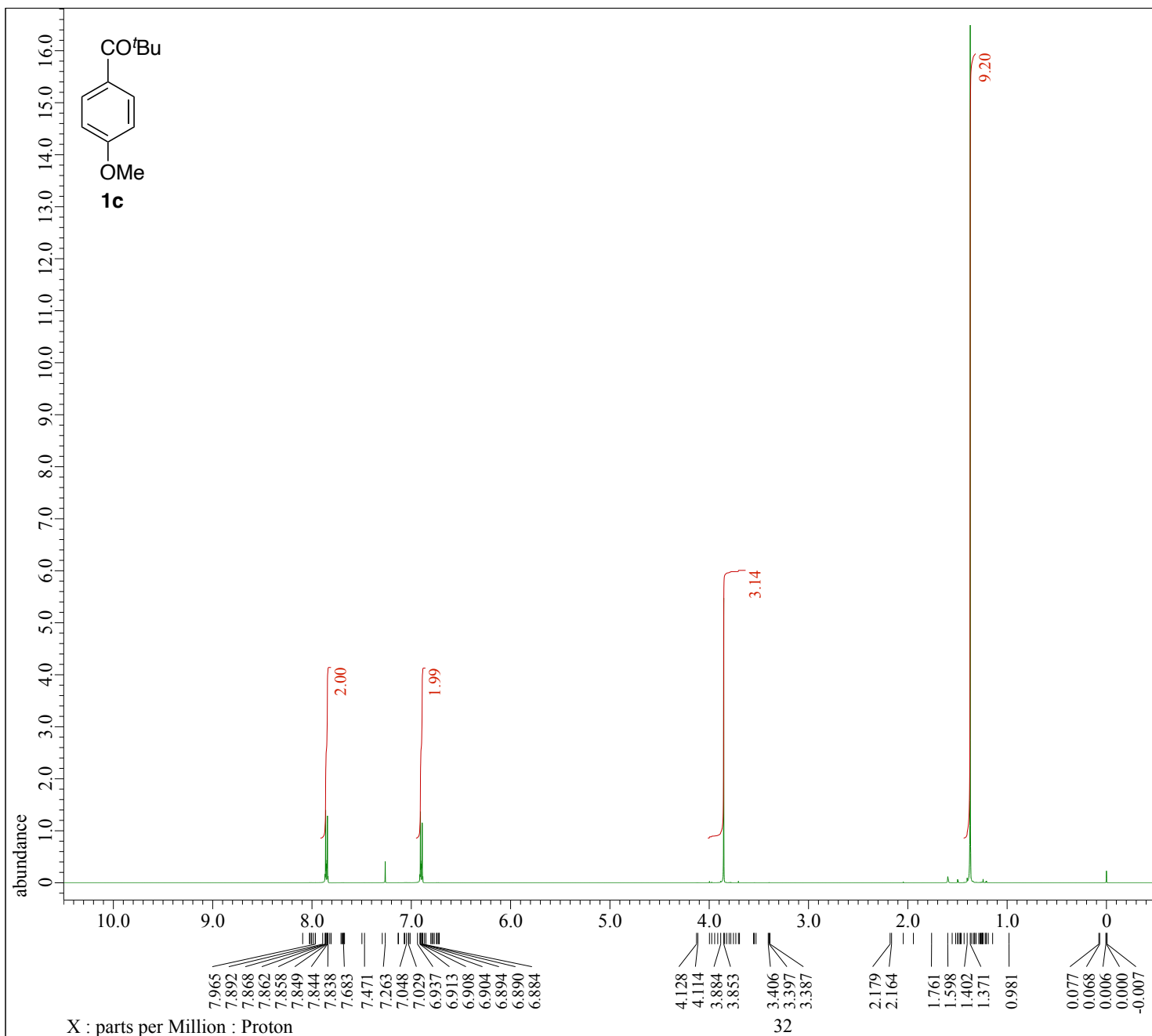
X : parts per Million : Proton



---- PROCESSING PARAMETERS ----
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm

以下に由来 : YF-648-2_Proton-1-1.jdf

Filename = YF-648-2_Proton-1-2.j
 Author = delta
 Experiment = proton.jpg
 Sample_Id = YF-648-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 20-OCT-2021 20:30:43
 Revision_Time = 5-JUL-2022 20:43:18
 Comment = single_pulse
 Data_Format = 1D COMPLEX
 Dim_Size = 13107
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-EZ500R/S1
 Field_Strength = 11.62926421[T] (500[M
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clipped = 7.42942051[kHz]
 IFR_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 32
 Total_Scans = 32
 Relaxation_Delay = 5[s]
 Recvr_Gain = 46
 Temp_Get = 22.3[dC]
 X_90_Width = 10.07[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[dB]
 X_Pulse = 5.035[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

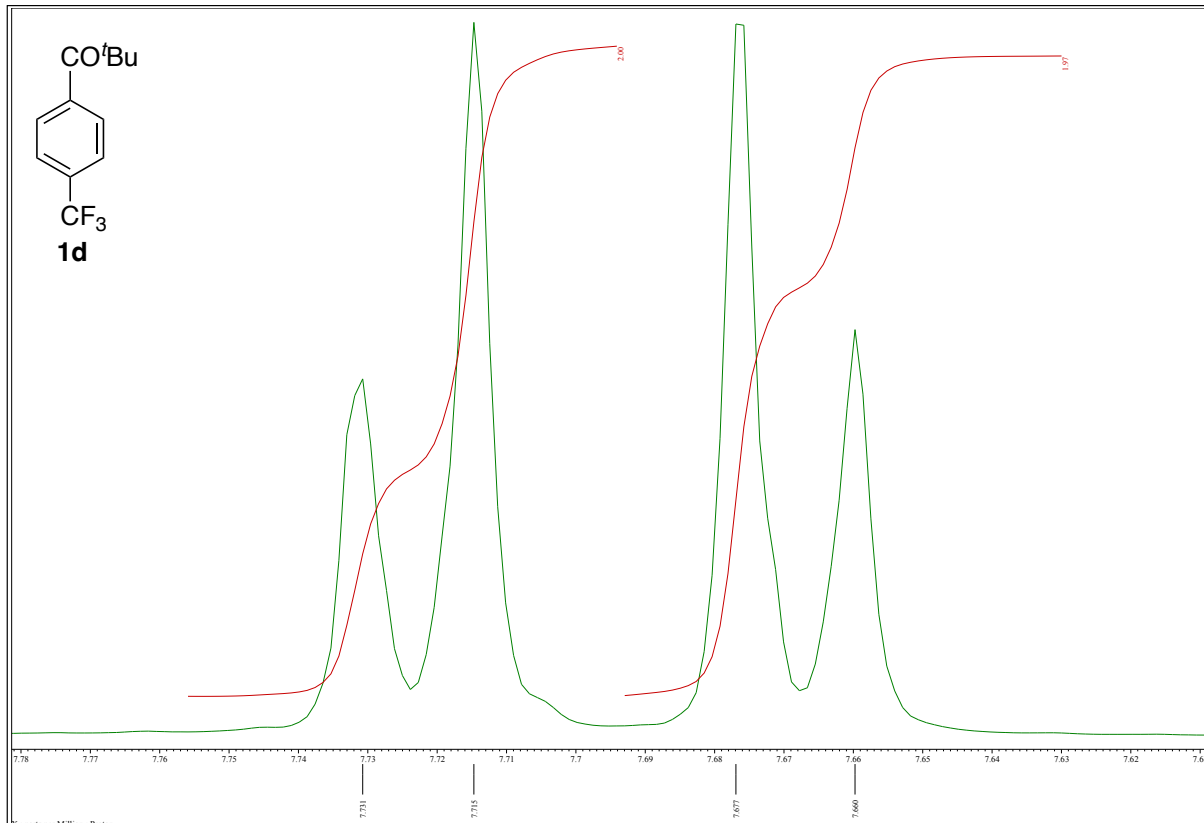
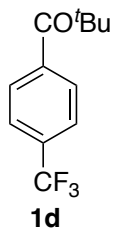


Filename = YF-809-2_Proton-1-2.j
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-809-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 18-FEB-2022 22:24:01
 Revision_Time = 19-FEB-2022 09:11:38

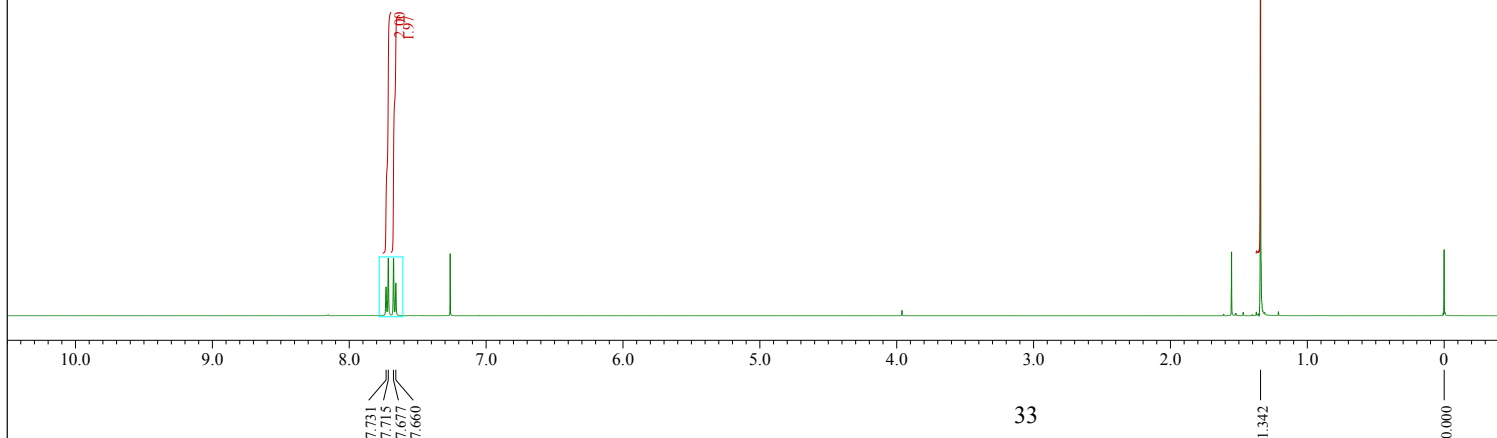
Comment = single_pulse
 Data_Format = 1D REAL
 Dim_Size = 13107
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1

Field_Strength = 11.62926421[T] (500[M]
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398 [MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198 [Hz]
 X_Sweep = 9.28677563 [kHz]
 X_Sweep_Clippped = 7.42942051 [kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398 [MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398 [MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 32
 Total_Scans = 32

Relaxation_Delay = 5[s]
 Recvr_Gain = 56
 Temp_Get = 21.8 [dC]
 X_90_Width = 7.33 [us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 3.3 [dB]
 X_Pulse = 3.665 [us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]



X : parts per Million : Proton



X : parts per Million : Proton



```

---- PROCESSING PARAMETERS ----
sexp( 0.2[Hz], 0.0[s] )
trapezoid( 0[%], 0[%], 80[%], 100[%] )
zerofill( 1, TRUE )
fft( 1, TRUE, TRUE )
machinephase
ppm

```

以下に由来 : YF-1008-3_Proton-1-1.jdf

```

Filename      = YF-1008-3_Proton-1-2.
Author       = delta
Experiment    = proton.jxp
Sample_Id     = YF-1008-3
Solvent       = Chloroform-D
Actual_Start_Time = 17-JUN-2022 10:51:29
Revision_Time = 5-JUL-2022 22:16:38

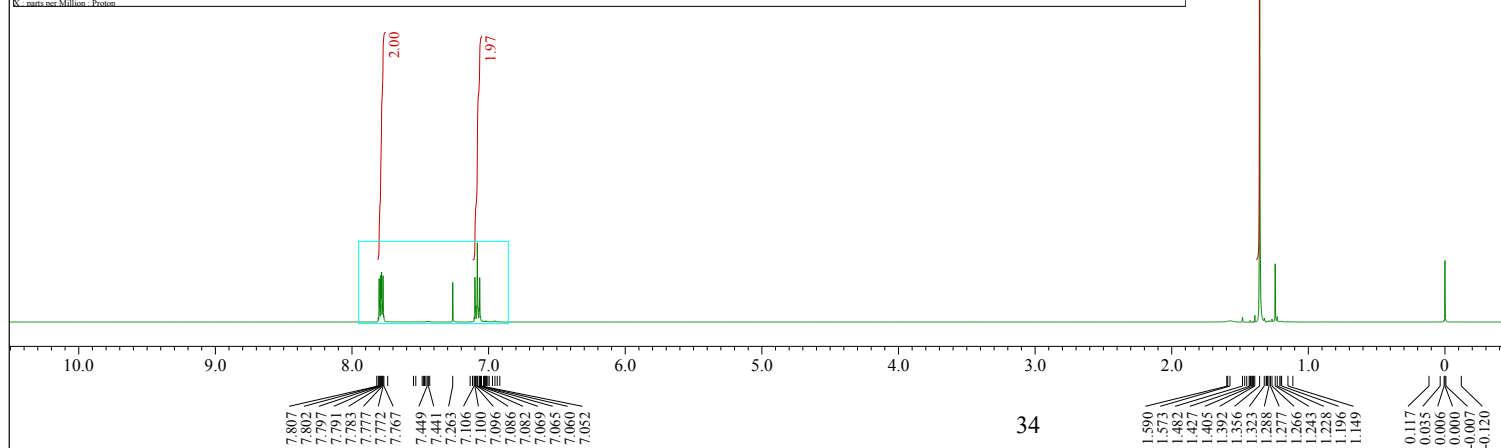
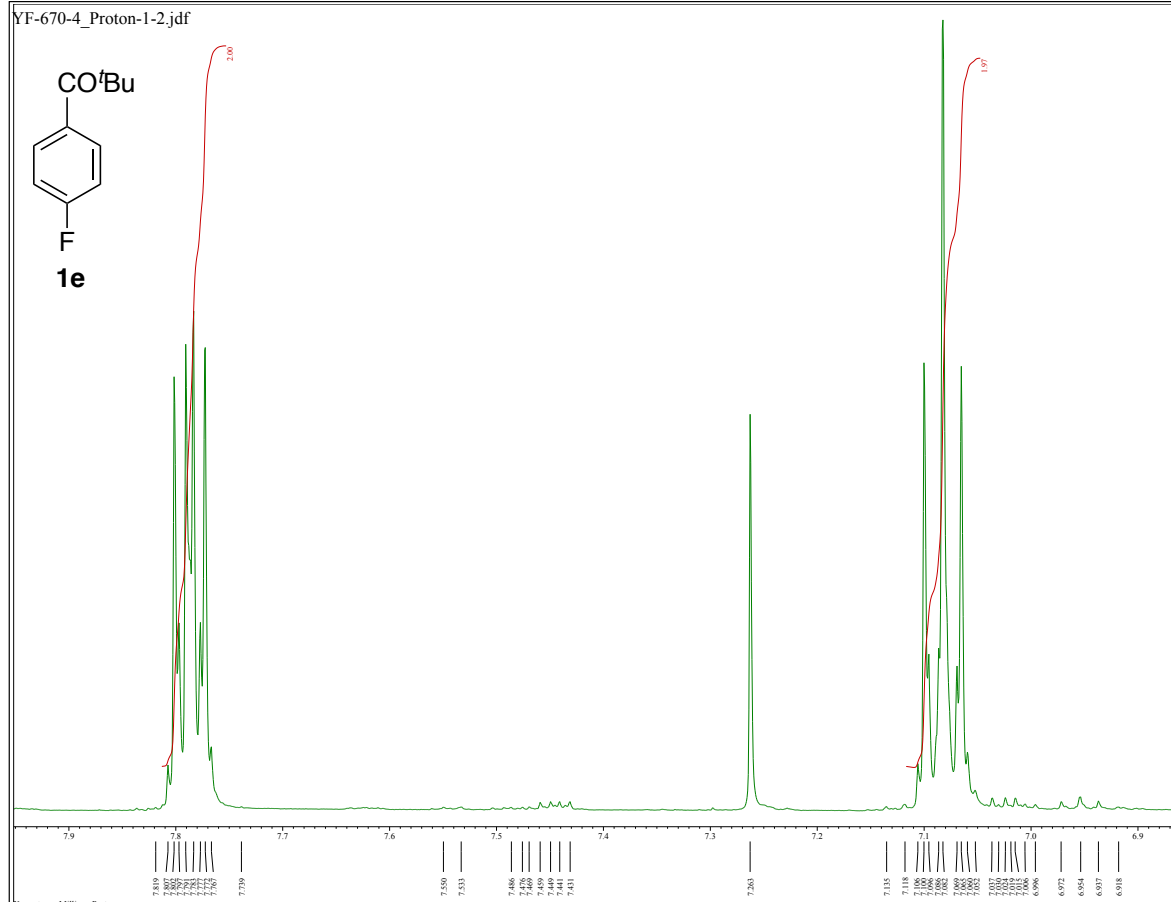
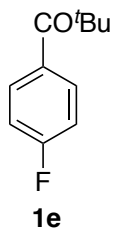
Comment       = single_pulse
Data Format    = 1D COMPLEX
Dim_Size      = 13107
X_Domain      = Proton
Dim_Title     = Proton
Dim_Units     = [ppm]
Dimensions    = X
Spectrometer  = JNM-ECZ500R/S1

Field_Strength = 11.62926421[T] (500[M]
X_Acq_Duration = 1.764222912[s]
X_Domain       = 1H
X_Freq         = 495.13191398[MHz]
X_Offset       = 5[ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198[Hz]
X_Sweep        = 9.28677563[kHz]
X_Sweep_Clipped = 7.42942051[kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398[MHz]
Irr_Offset     = 5[ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398[MHz]
Tri_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 32
Total_Scans    = 32

Relaxation_Delay = 5[s]
Recvr_Gain       = 46
Temp_Get         = 24.1[dC]
X_90_Width       = 9.49[us]
X_Acq_Time       = 1.764222912[s]
X_Angle          = 45[deg]
X_Atn            = 4.3[dB]
X_Pulse          = 4.745[us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 500
Dante_Preset     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1[s]
Phase            = {0, 90, 270, 180, 180}
Preset_Time      = 5[s]
Preset_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 5[s]
Repetition_Time  = 6.764222912[s]

```

YF-670-4_Proton-1-2.jdf



X : parts per Million : Proton



```

---- PROCESSING PARAMETERS ----
dc_balance(0, FALSE)
sexp( 0.2[Hz], 0.0[s] )
trapezoid( 0[%], 0[%], 80[%], 100[%] )
zerofill( 1, TRUE )
fft( 1, TRUE, TRUE )
machinephase
ppm
base_correct( Akima, 5, 0, FALSE, 3, None, FA

```

```

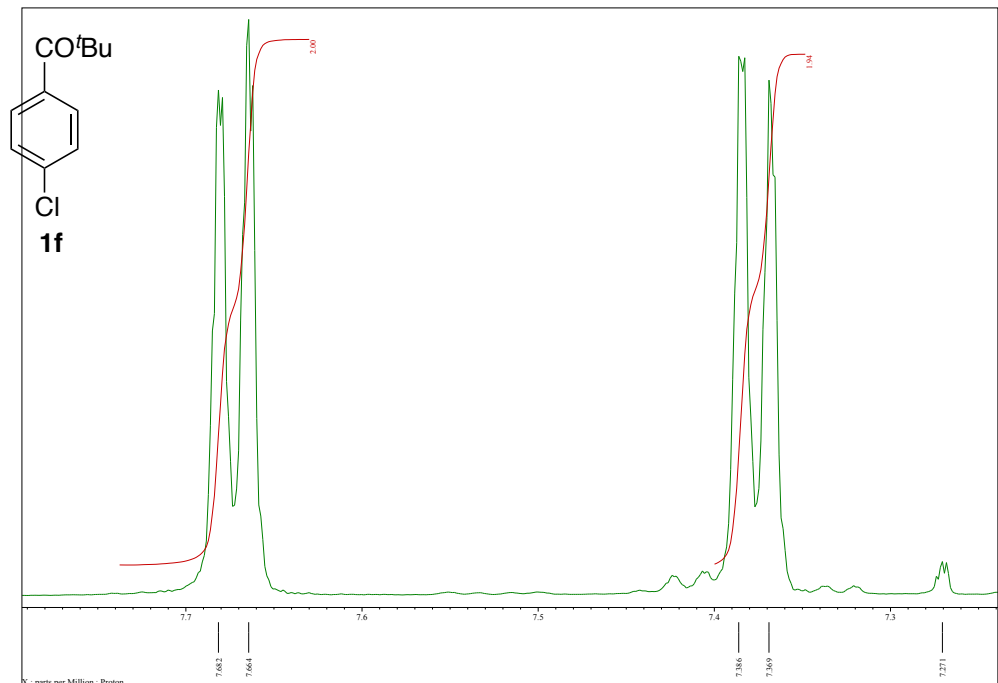
Filename      = YF-670-4_Proton-1-2_jdf
Author        = iwawasa
Experiment     = proton_jp
Sample Id     = YF-670-4
Solvent       = DMSO-D6FORM-D
Actual Start Time = 26-NOV-2021 20:04:12
Revision Time  = 6-JUL-2022 11:51:55

Comment
Data Format    = single pulse
Dim Size      = 1D REAL
X Domain      = 26214
X Min         = Proton
X Max         = Proton
X Title       =
X Units       = [ppm]
Dimensions    = X
Site          = JNM-ECK500
Spectrometer  = DELTA2 NMR

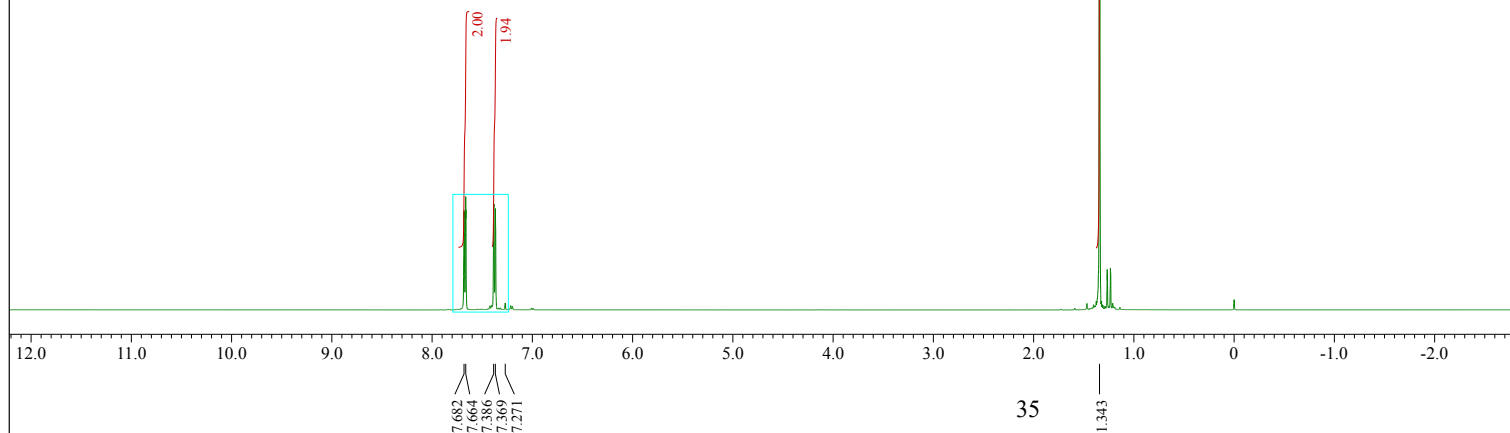
Field Strength = 174.73579 [T] (500 [MHz])
X Acquisition  = 3.49175808 [s]
X Domain      = 1H
X Freq        = 500.15991521 [MHz]
X Offset      = 5.0 [ppm]
X Points      = 6
X Prescans    = 1
X Resolution  = 0.26838668 [Hz]
X Sweep      = 3.68438438 [kHz]
X Sweep Clipped = 7.50750751 [kHz]
Irr Domain    = Proton
Irr Freq     = 500.15991521 [MHz]
Irr Offset   = 5 [ppm]
Tri Domain   = Proton
Tri Freq     = 500.15991521 [MHz]
Tri Offset   = 5.0 [ppm]
Clipped     = FALSE
Scans        = 16
Total Scans  = 16

Relaxation Delay = 2 [s]
Recvr Gain       = 44
Temp Ctg        = 22.2 [dC]
X X 90 Width     = 12.4 [us]
X Acq Time       = 3.49175808 [s]
X Angle         = 45 [deg]
X Atn           = 45 [dB]
X Pulse         = 6 [us]
Irr Mode        = off
Tri Mode        = off
Dante Presat    = FALSE
Initial Wait    = 1 [s]
Repetition Time = 5.49175808 [s]

```



X : parts per Million : Proton



X : parts per Million : Proton



```

---- PROCESSING PARAMETERS ----
sexp( 0.2[Hz], 0.0[s] )
trapezoid( 0[%], 0[%], 80[%], 100[%] )
zerofill( 1, TRUE )
fft( 1, TRUE, TRUE )
machinephase
ppm

```

以下に由来 : YF-743-3_Proton-1-1.jdf

```

Filename      = YF-743-3_Proton-1-2.j
Author        = delta
Experiment     = proton.jpg
Sample_Id     = YF-743-3
Solvent       = CHLOROFORM-D
Actual_Start_Time = 3-DEC-2021 22:23:42
Revision_Time  = 8-JUL-2022 10:04:11

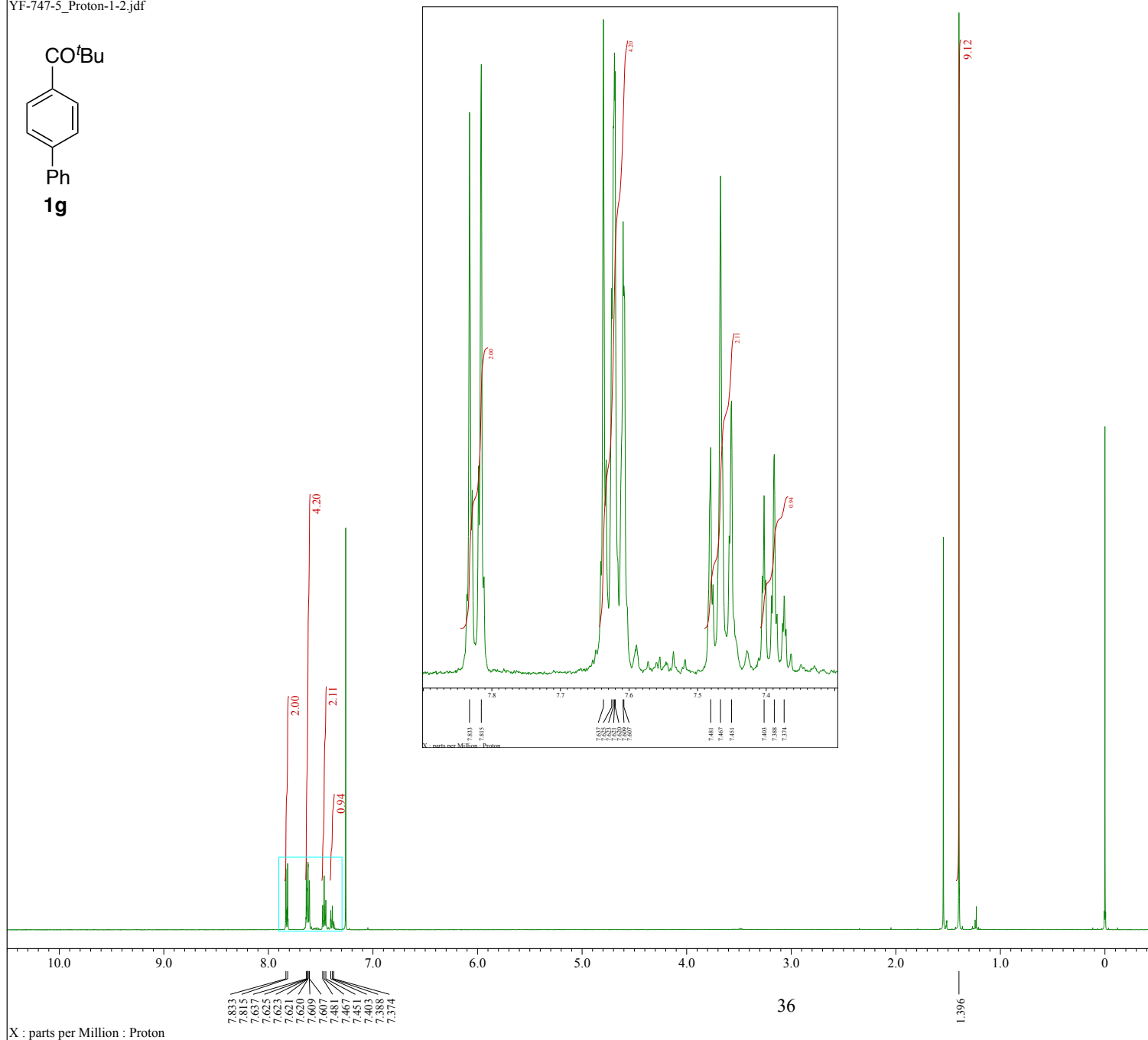
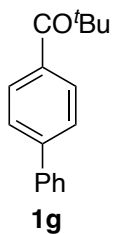
Comment       = single_pulse
Data Format    = 1D COMPLEX
Dim_Size      = 13107
X_Domain      = Proton
Dim_Title     = Proton
Dim_Units     = [ppm]
Dimensions    = X
Spectrometer  = JNM-EZ500R/S1

Field_Strength = 11.62926421[T] (500[M
X_Acq_Duration = 1.76422912[s]
X_Domain       = 1H
X_Freq         = 495.13191398[MHz]
X_Offset       = 5[ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198[Hz]
X_Sweep        = 9.28677563[kHz]
X_Sweep_Clippped = 7.42942051[kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398[MHz]
Irr_Offset     = 5[ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398[MHz]
Tri_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 32
Total_Scans    = 32

Relaxation_Delay = 5[s]
Recvr_Gain       = 26
Temp_Get         = 21[dC]
X_90_Width       = 10.07[us]
X_Acq_Time       = 1.76422912[s]
X_Angle          = 45[deg]
X_Atn            = 4.3[dB]
X_Pulse          = 5.035[us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 500
Dante_Preset     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1[s]
Phase            = {0, 90, 270, 180, 180}
Preset_Time      = 5[s]
Preset_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 5[s]
Repetition_Time  = 6.76422912[s]

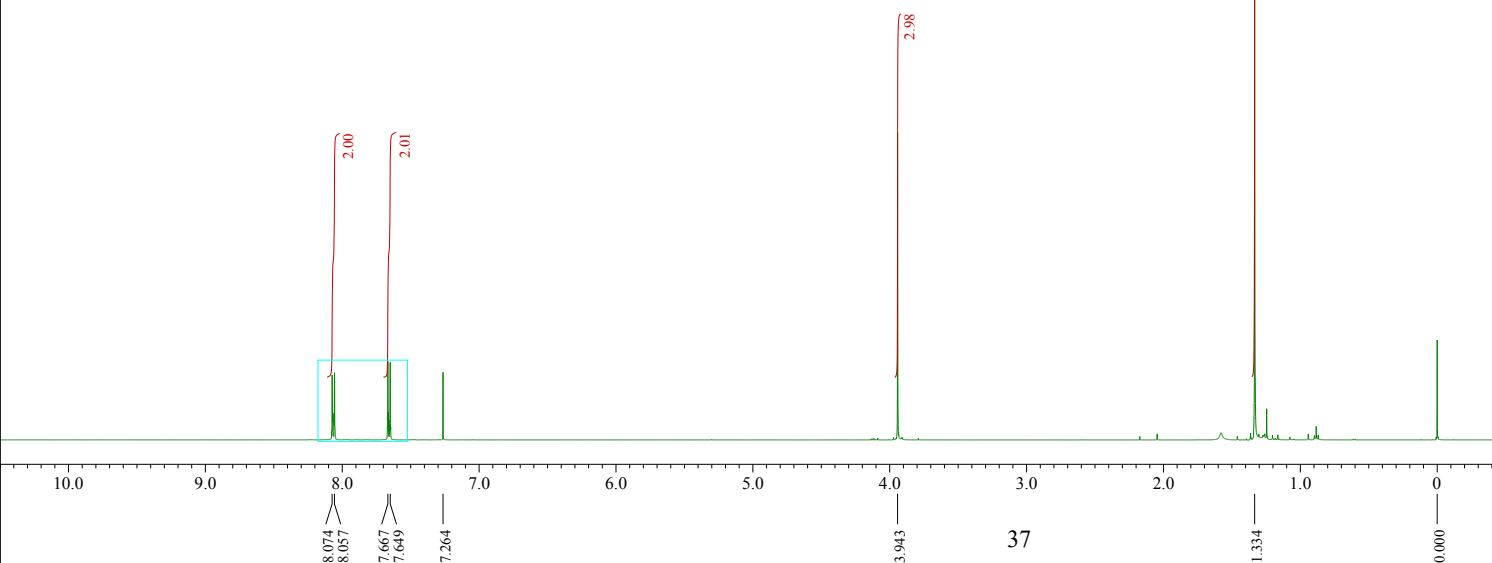
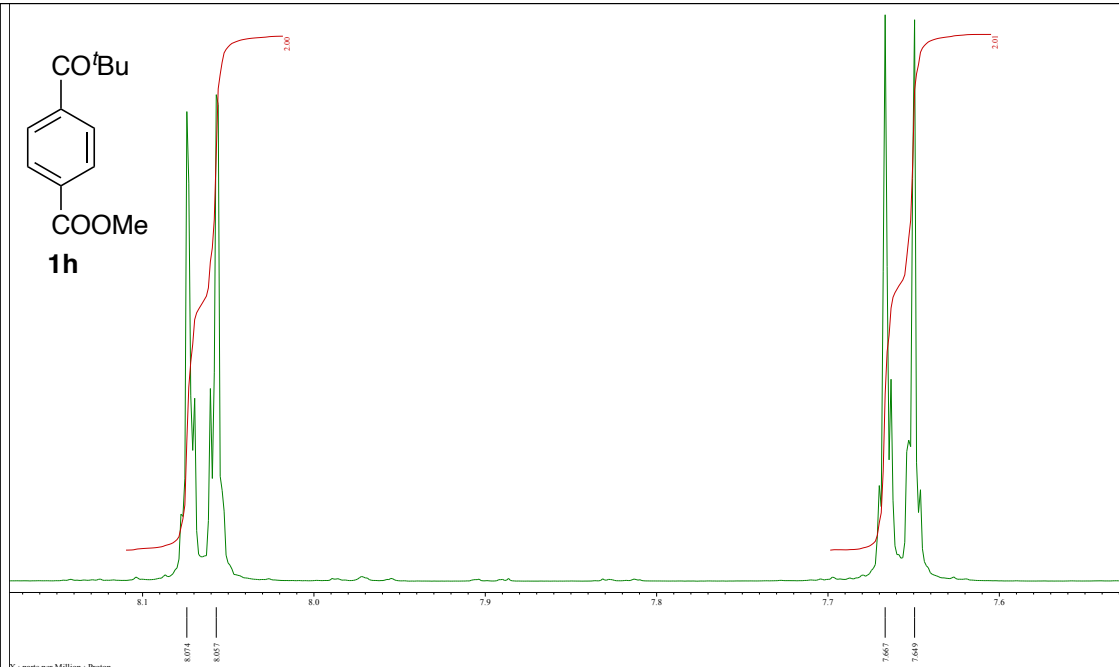
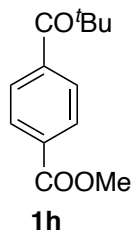
```

YF-747-5_Proton-1-2.jdf



---- PROCESSING PARAMETERS ----
 dc balance(0, FALSE)
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm
 以下に由来 : YF-747-5_Proton-1-1.jdf

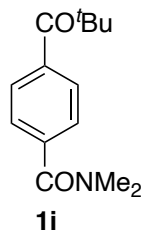
Filename = YF-747-5_Proton-1-2.jdf
 Author = iwasawa
 Experiment = proton.jxp
 Sample_Id = YF-747-5
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 9-DEC-2021 19:28:43
 Revision_Time = 7-JUL-2022 22:19:39
 Comment = single_pulse
 Data Format = 1D COMPLEX
 Dim_Size = 26214
 X_Domain = Proton
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Site = JNM-ECX500
 Spectrometer = DELTA2_NMR
 Field_Strength = 11.7473579[T] (500[MHz])
 X_Acq_Duration = 3.49175808[s]
 X_Domain = 1H
 X_Freq = 500.15991521[MHz]
 X_Offset = 5.0[ppm]
 X_Points = 32768
 X_Prescans = 1
 X_Resolution = 0.28638868[Hz]
 X_Sweep = 9.38438438[kHz]
 X_Sweep_Clippped = 7.50750751[kHz]
 Irr_Domain = Proton
 Irr_Freq = 500.15991521[MHz]
 Irr_Offset = 5.0[ppm]
 Tri_Domain = Proton
 Tri_Freq = 500.15991521[MHz]
 Tri_Offset = 5.0[ppm]
 Clipped = FALSE
 Scans = 16
 Total_Scans = 16
 Relaxation_Delay = 2[s]
 Recvr_Gain = 54
 Temp_Get = 22.2[dC]
 X_90_Width = 12.4[us]
 X_Acq_Time = 3.49175808[s]
 X_Angle = 45[deg]
 X_Atn = 4.5[dB]
 X_Pulse = 6.2[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Preset = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 5.49175808[s]



---- PROCESSING PARAMETERS ----
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm
 以下に由来 : YF-1018-4_Proton-1-1.jdf

Filename = YF-1018-4_Proton-1-2.
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-1018-4
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 18-JUN-2022 17:29:47
 Revision_Time = 6-JUL-2022 14:33:31
 Comment = single_pulse
 Data Format = 1D COMPLEX
 Dim_Size = 13107
 X_Domain = Proton
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1
 Field_Strength = 11.62926421[T] (500[M
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X Points = 16384
 X_Prescans = 1
 X Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clippped = 7.42942051[kHz]
 IFR_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8
 Relaxation_Delay = 5[s]
 Recvr Gain = 46
 Temp Get = 24.1[dC]
 X_90_Width = 9.49[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[dB]
 X_Pulse = 4.745[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat Time = 5[s]
 Presat Time Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

YF-0-2_Proton-1-2.jdf



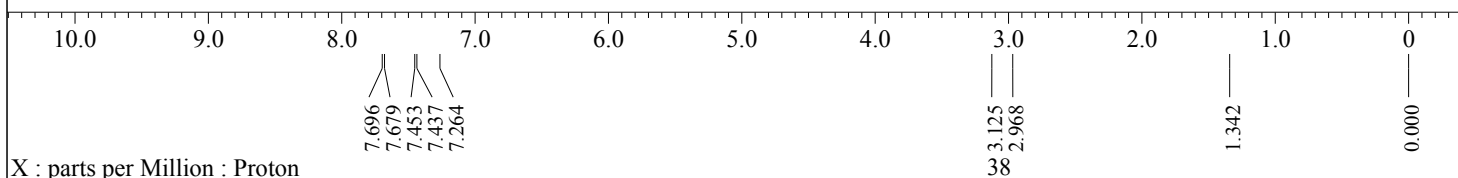
```

Filename      = YF-0-2_Proton-1-2.jdf
Author       = delta
Experiment    = proton.jxp
Sample Id    = YF-0-2
Solvent      = CHLOROFORM-D
Actual_Start_Time = 21-MAY-2022 11:28:04
Revision_Time  = 23-MAY-2022 15:43:07

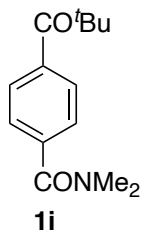
Comment      = single_pulse
Data Format   = 1D COMPLEX
Dim Size     = 13107
X_Domain     = Proton
Dim Title    = Proton
Dim Units    = [ppm]
Dimensions   = X
Spectrometer = JNM-ECZ500R/S1

Field Strength = 11.62926421[T] (500[M]
X_Acq_Duration = 1.76422912[s]
X_Domain       = 1H
X_Freq         = 495.13191398 [MHz]
X_Offset       = 5[ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198 [Hz]
X_Sweep        = 9.28677563 [kHz]
X_Sweep_Clippped = 7.42942051 [kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398 [MHz]
Irr_Offset     = 5[ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398 [MHz]
Tri_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 32
Total_Scans    = 32

Relaxation_Delay = 5[s]
Recvr_Gain       = 56
Temp_Get         = 24.9 [dC]
X_90_Width       = 7.5 [us]
X_Acq_Time       = 1.76422912[s]
X_Angle          = 45[deg]
X_Atn            = 3.3 [dB]
X_Pulse          = 3.75 [us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 500
Dante_Presat     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1[s]
Phase            = {0, 90, 270, 180, 180}
Presat_Time      = 5[s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 5[s]
Repetition_Time  = 6.76422912[s]
  
```

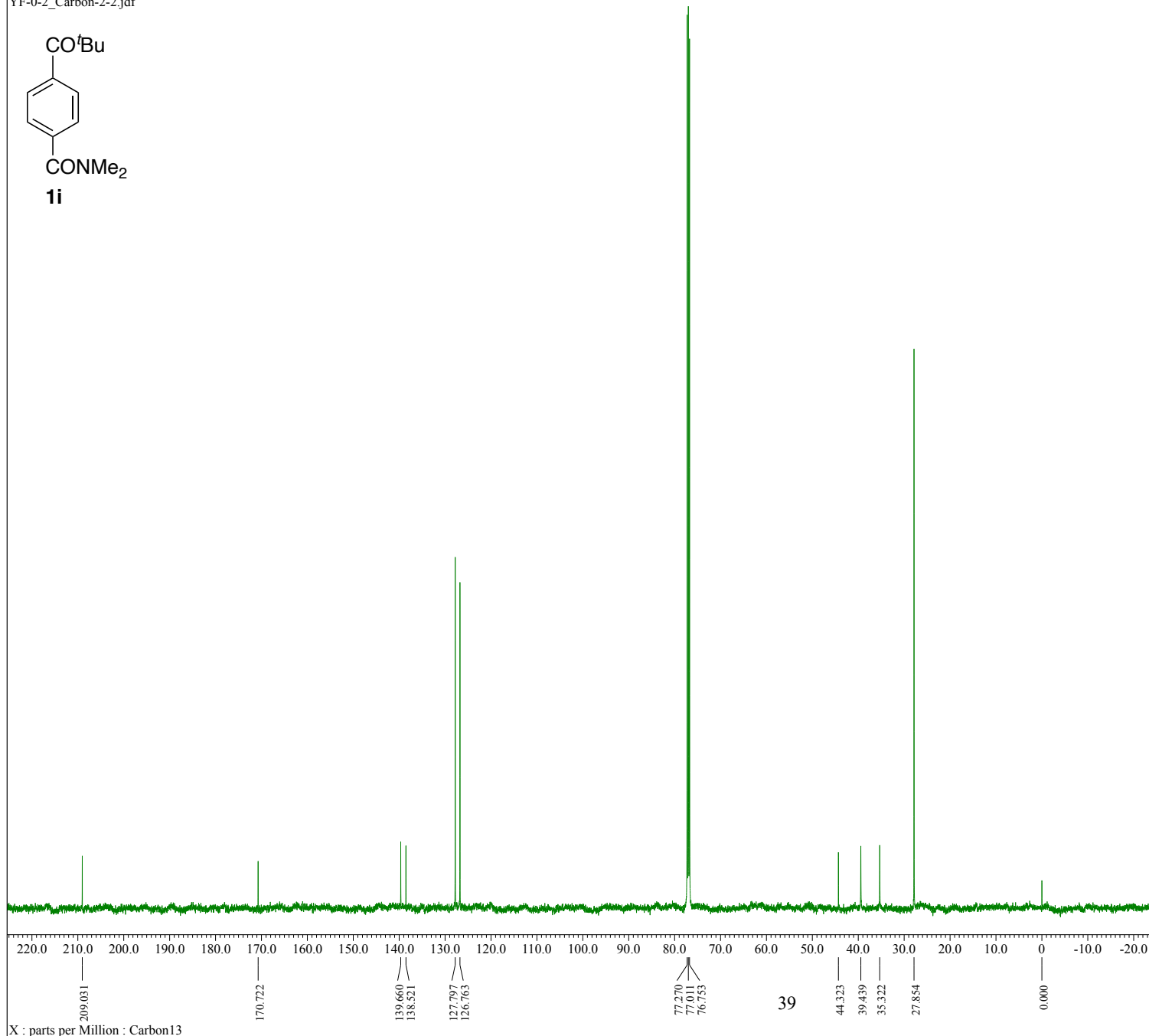


YF-0-2_Carbon-2-2.jdf



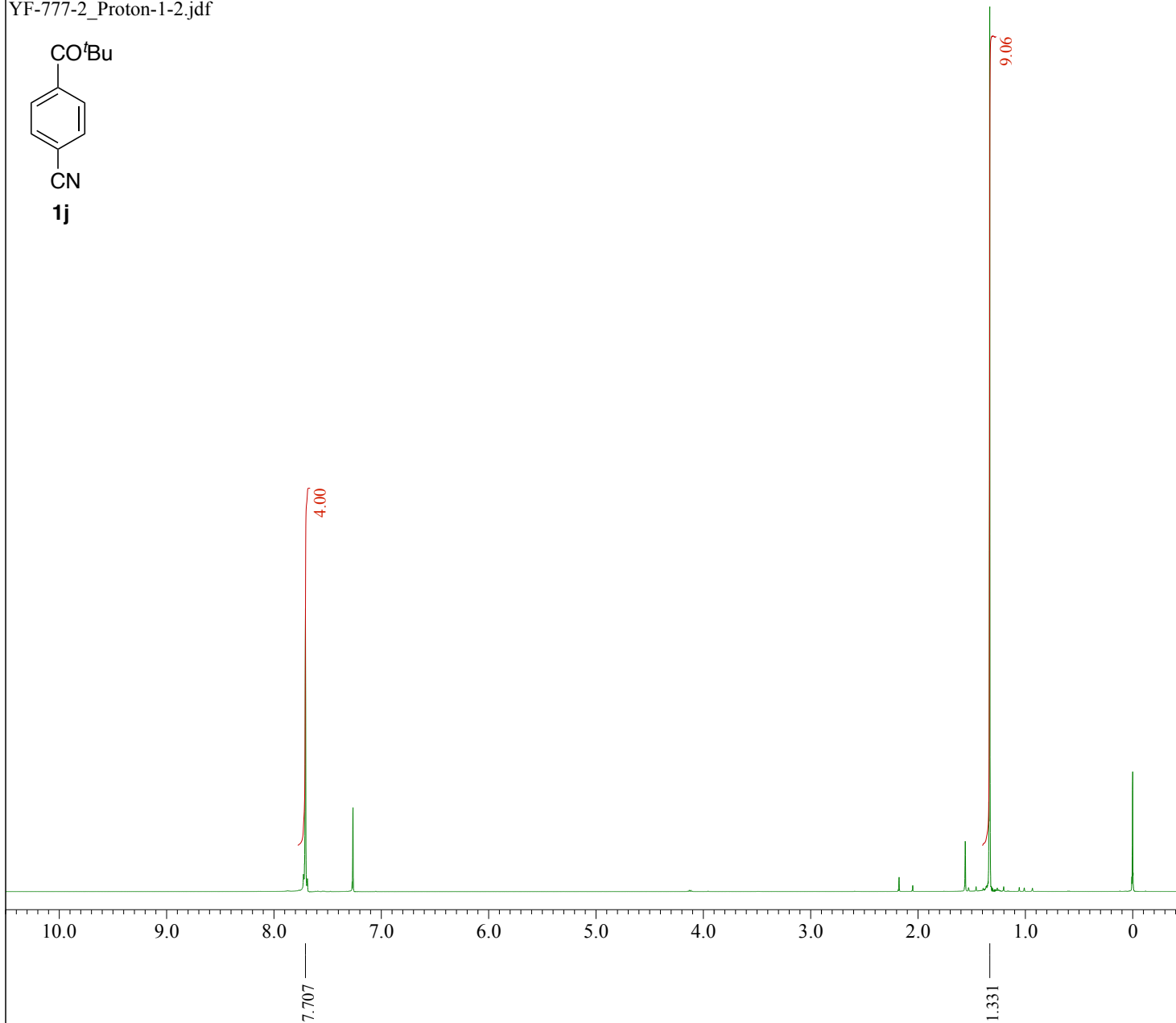
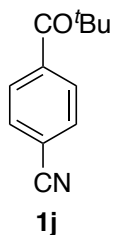
---- PROCESSING PARAMETERS ----
sexp(2.0[Hz], 0.0[s])
trapezoid(0[%], 0[%], 80[%], 100[%])
zerofill(1, TRUE)
fft(1, TRUE, TRUE)
machinephase
ppm
phase(-1.03703, 0, 59.13478[%])
base_correct(Akima, 5, 0, FALSE, 3, None, FA
以下に由来: YF-0-2_Carbon-2-1.jdf

Filename = YF-0-2_Carbon-2-2.
Author = delta
Experiment = carbon.jxp
Sample_Id = YF-0-2
Solvent = CHLOROFORM-D
Actual_Start_Time = 22-MAY-2022 03:24:
Revision_Time = 9-JUL-2022 12:28:
Comment = single pulse decou
Data Format = 1D REAL
Dim_Size = 26214
X_Domain = Carbon13
Dim_Title = Carbon13
Dim_Units = [ppm]
Dimensions = X
Spectrometer = JNM-ECZ500R/S1
Field_Strength = 11.62926421[T] (50
X_Acq_Duration = 0.8388608[s]
X_Domain = 13C
X_Freq = 124.5010059[MHz]
X_Offset = 100[ppm]
X_Points = 32768
X_Prescans = 4
X_Resolution = 1.1920929[Hz]
X_Sweep = 39.0625[kHz]
X_Sweep_Clippped = 31.25[kHz]
Irr_Domain = Proton
Irr_Freq = 495.13191398[MHz]
Irr_Offset = 5[ppm]
Clipped = FALSE
Scans = 4096
Total_Scans = 4096
Relaxation_Delay = 2[s]
Recvr_Gain = 56
Temp_Get = 24.9[dc]
X_90_Width = 14[us]
X_Acq_Time = 0.8388608[s]
X_Angle = 30[deg]
X_Atn = 11[db]
X_Pulse = 4.66666667[us]
Irr_Atn_Dec = 25.8[db]
Irr_Atn_Dec_Calc = 25.8[db]
Irr_Atn_Dec_Default_Calc = 25.8[db]
Irr_Atn_No = 25.8[db]
Irr_Dec_Bandwidth_Hz = 5.97826087[kHz]
Irr_Dec_Bandwidth_Ppm = 12.07407703[ppm]
Irr_Dec_Freq = 495.13191398[MHz]
Irr_Dec_Merit_Factor = 2.2
Irr_Decoupling = TRUE
Irr_No = TRUE
Irr_Noise = WALTZ
Irr_Offset_Default = 5[ppm]
Irr_Pwidth = 92[us]
Irr_Pwidth_Default = 92[us]
Irr_Pwidth_Default_Calc = 92[us]
Irr_Pwidth_Templ = 92[us]
Irr_Wurst = FALSE
Decimation_Rate = 0
Initial_Wait = 1[s]
Noe_Time = 2[s]
Noe_Time_Flag = FALSE
Relaxation_Delay_Calc = 2[s]
Relaxation_Delay_Temp = 2[s]
Repetition_Time = 2.8388608[s]



X : parts per Million : Carbon13

YF-777-2_Proton-1-2.jdf



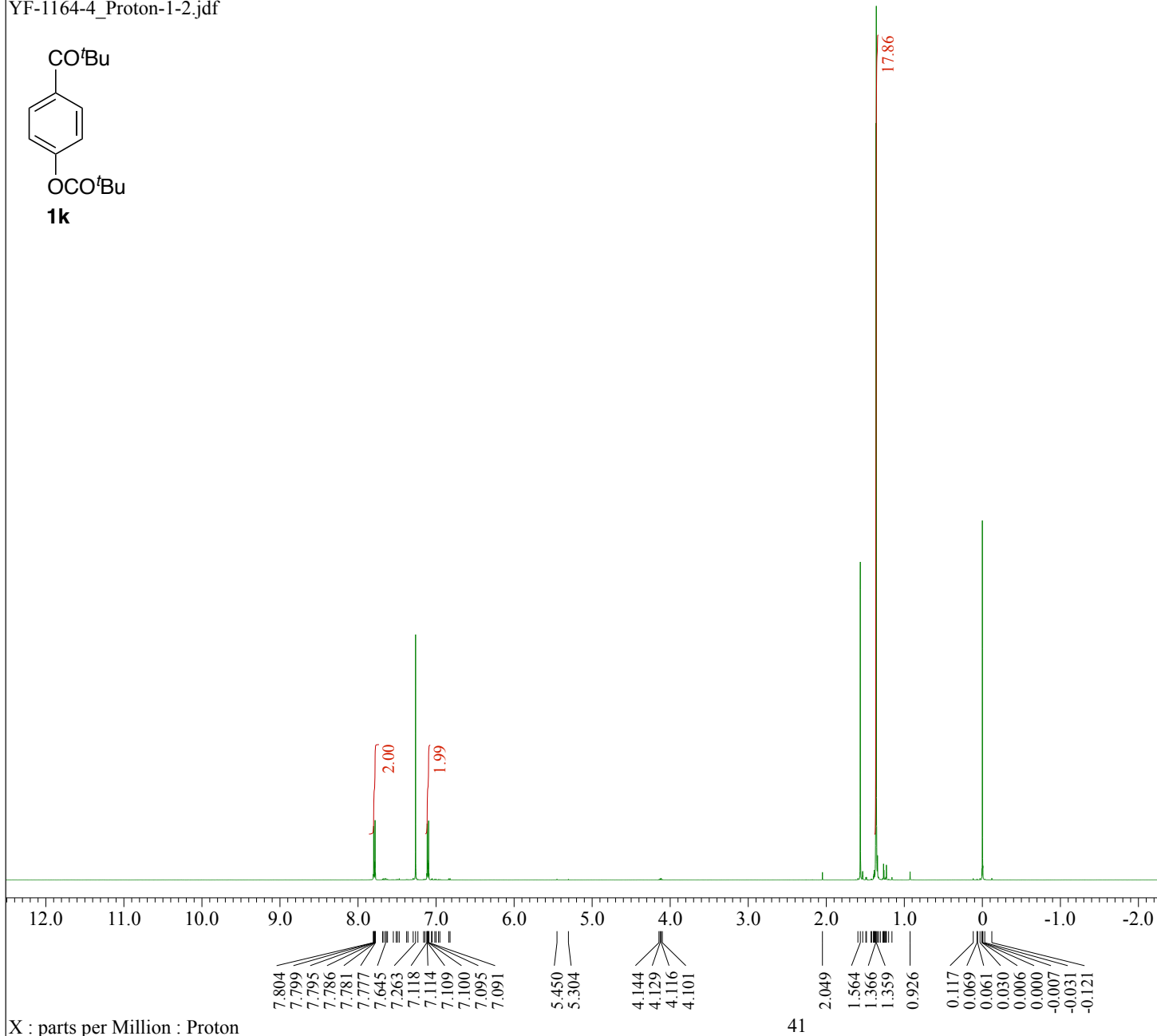
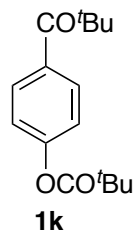
X : parts per Million : Proton

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Filename	= YF-777-2_Proton-1-2.j
Author	= delta
Experiment	= proton.jxp
Sample Id	= YF-777-2
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 8-JAN-2022 20:48:23
Revision_Time	= 18-JUL-2022 20:08:52
Comment	= single_pulse
Data Format	= 1D COMPLEX
Dim Size	= 13107
X Domain	= Proton
Dim Title	= Proton
Dim Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ500R/S1
Field Strength	= 11.62926421[T] (500[M
X Acq_Duration	= 1.76422912[s]
X Domain	= 1H
X_Freq	= 495.13191398 [MHz]
X_Offset	= 5[ppm]
X Points	= 16384
X_Prescans	= 1
X Resolution	= 0.5668198 [Hz]
X Sweep	= 9.28677563 [kHz]
X Sweep_Clippped	= 7.42942051 [kHz]
Irr Domain	= Proton
Irr_Freq	= 495.13191398 [MHz]
Irr_Offset	= 5[ppm]
Tri Domain	= Proton
Tri_Freq	= 495.13191398 [MHz]
Tri_Offset	= 5[ppm]
Clipped	= TRUE
Scans	= 32
Total_Scans	= 32
Relaxation_Delay	= 5[s]
Recvr Gain	= 56
Temp_Get	= 20.5 [dC]
X 90_Width	= 9.38 [us]
X Acq_Time	= 1.76422912[s]
X_Angle	= 45[deg]
X_Atn	= 4.3[dB]
X_Pulse	= 4.69 [us]
Irr_Mode	= Off
Tri_Mode	= Off
Dante_Loop	= 500
Dante_Presat	= FALSE
Decimation_Rate	= 0
Initial_Wait	= 1[s]
Phase	= {0, 90, 270, 180, 180
Presat_Time	= 5[s]
Presat_Time_Flag	= FALSE
Relaxation_Delay_Calc	= 0[s]
Relaxation_Delay_Temp	= 5[s]
Repetition_Time	= 6.76422912[s]

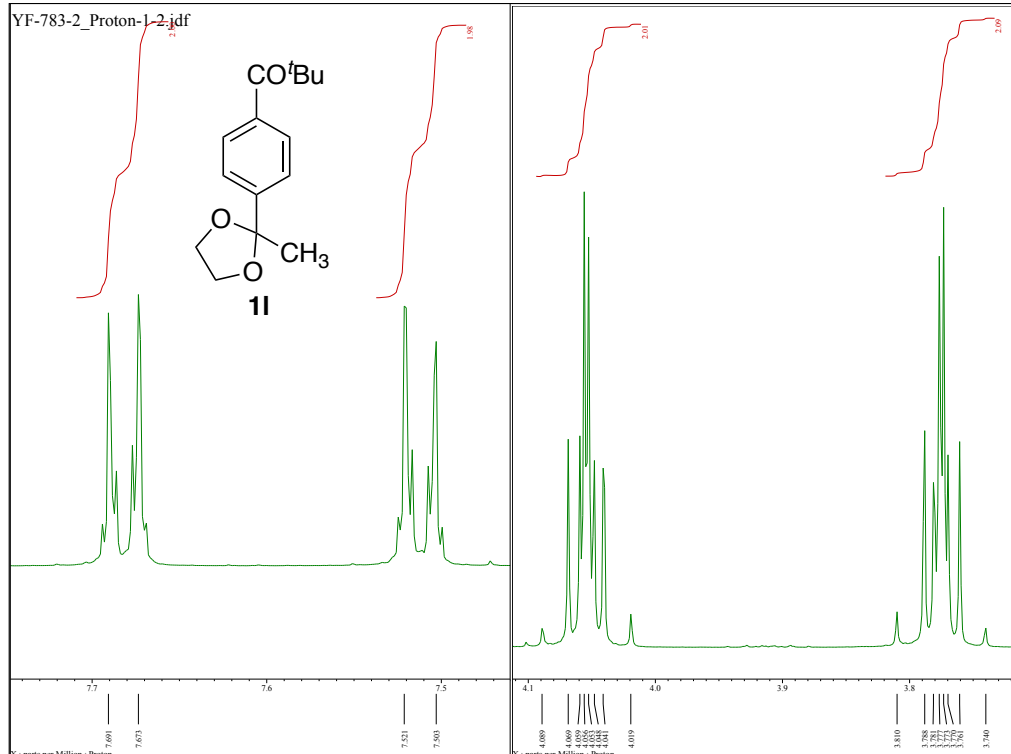
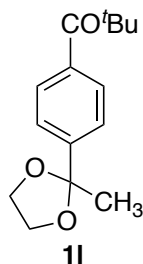
YF-1164-4_Proton-1-2.jdf



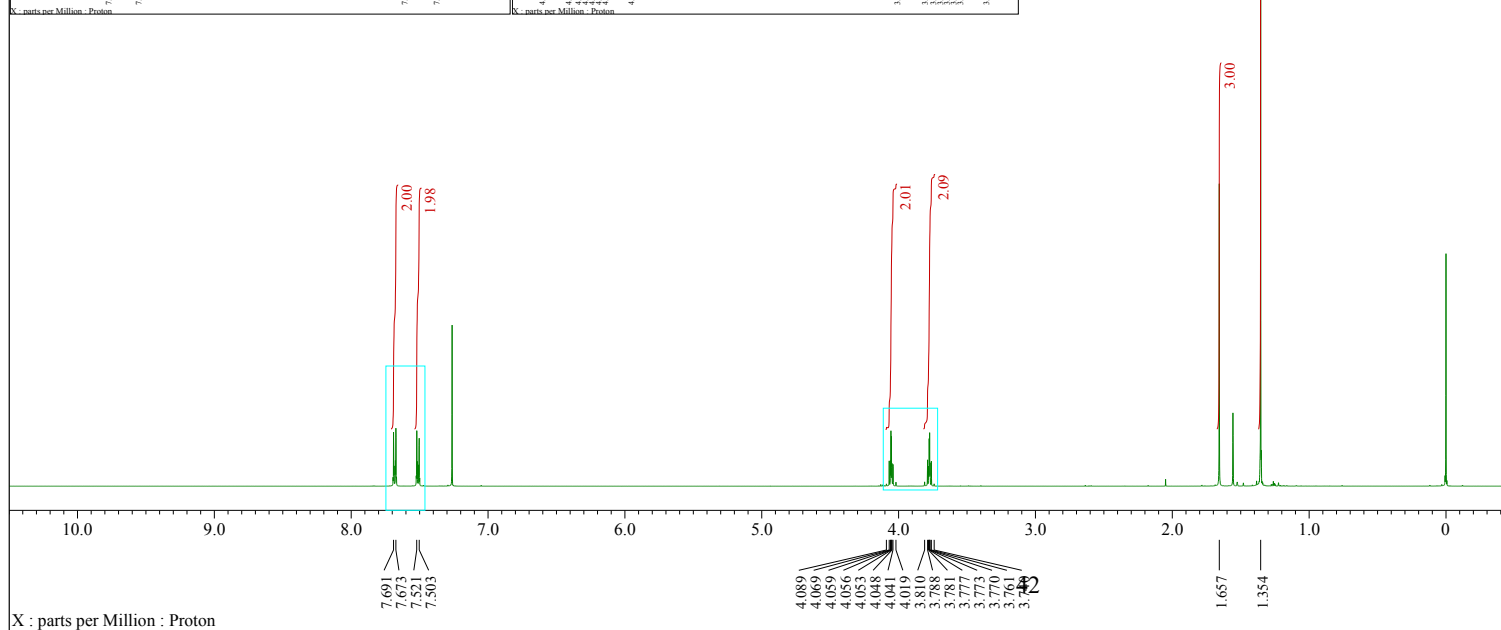
Filename = YF-1164-4_Proton-1-2.
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-1164-4
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 17-OCT-2022 22:51:41
 Revision_Time = 18-OCT-2022 13:37:37
 Comment = single_pulse
 Data_Format = 1D REAL
 Dim_Size = 13107
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1
 Field_Strength = 11.62926421[T] (500[M]
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398 [MHz]
 X_Offset = 5 [ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198 [Hz]
 X_Sweep = 9.28677563 [kHz]
 X_Sweep_Clippped = 7.42942051 [kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398 [MHz]
 Irr_Offset = 5 [ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398 [MHz]
 Tri_Offset = 5 [ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8
 Relaxation_Delay = 5 [s]
 Recvr_Gain = 56
 Temp_Get = 19.2 [dC]
 X_90_Width = 8.7 [us]
 X_Acq_Time = 1.76422912 [s]
 X_Angle = 45 [deg]
 X_Atn = 4.3 [dB]
 X_Pulse = 4.35 [us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1 [s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5 [s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0 [s]
 Relaxation_Delay_Temp = 5 [s]
 Repetition_Time = 6.76422912 [s]

X : parts per Million : Proton

YF-783-2_Proton-1-2.jdf



X : parts per Million : Proton



X : parts per Million : Proton

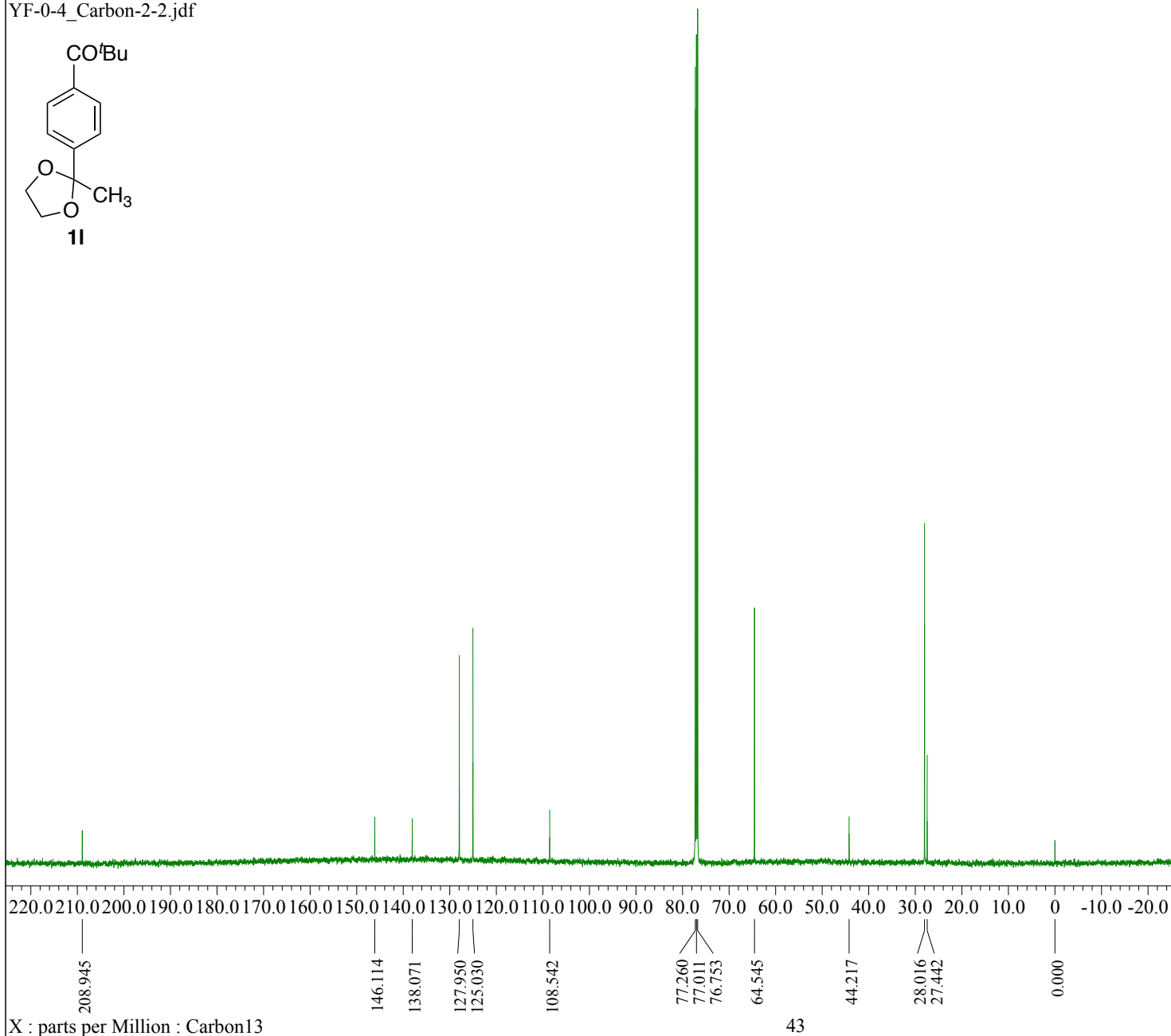
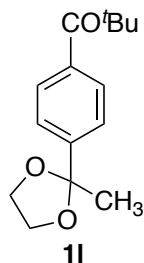


----- PROCESSING PARAMETERS -----
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm

以下に由来 : YF-783-2_Proton-1-1.jdf

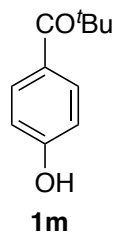
Filename = YF-783-2_Proton-1-2.j
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-783-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 12-JAN-2022 21:44:03
 Revision_Time = 7-JUL-2022 21:34:34
 Comment = single_pulse
 Data_Format = 1D COMPLEX
 Dim_Size = 13107
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-EZ500R/S1
 Field_Strength = 11.62926421[T] (500[M]
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clipped = 7.42942051[kHz]
 IFR_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = TRUE
 Scans = 32
 Total_Scans = 32
 Relaxation_Delay = 5[s]
 Recvr_Gain = 56
 Temp_Get = 20.8[dC]
 X_90_Width = 9.38[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[dB]
 X_Pulse = 4.69[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Preset = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

YF-0-4_Carbon-2-2.jdf

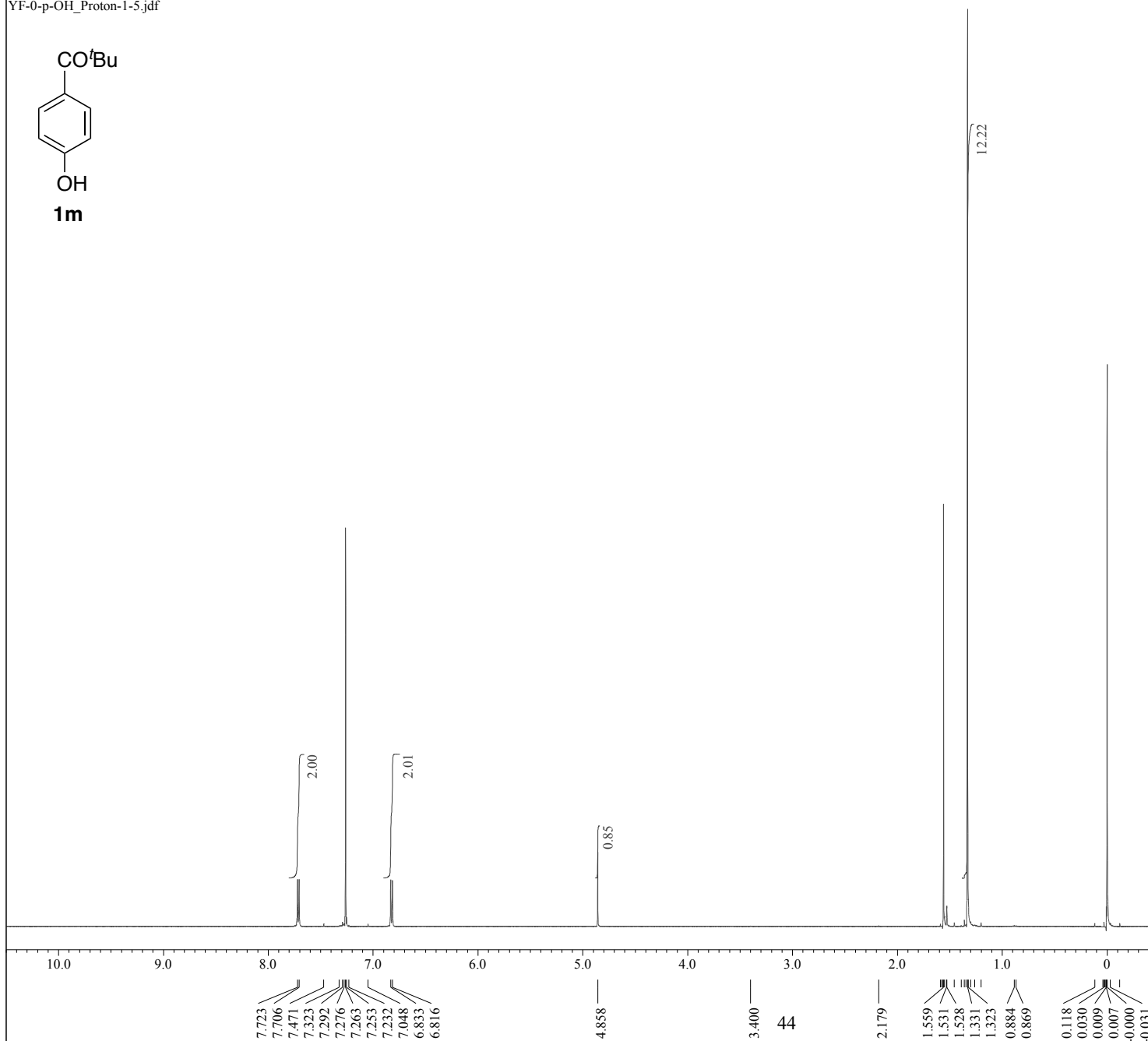


Filename	= YF-0-4_Carbon-2-2.
Author	= delta
Experiment	= carbon.jxp
Sample Id	= YF-0-4
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 22-MAY-2022 06:44:
Revision_Time	= 23-MAY-2022 15:57:
Comment	= single pulse decou
Data Format	= 1D COMPLEX
Dim Size	= 26214
X Domain	= Carbon13
Dim Title	= Carbon13
Dim Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ500R/S1
Field Strength	= 11.62926421[T] (50
X Acq Duration	= 0.8388608[s]
X Domain	= 13C
X Freq	= 124.5010059[MHz]
X Offset	= 100[ppm]
X Points	= 32768
X Prescans	= 4
X Resolution	= 1.1920929[Hz]
X Sweep	= 39.0625[kHz]
X Sweep_Clip	= 31.25[kHz]
Irr Domain	= Proton
Irr Freq	= 495.13191398[MHz]
Irr Offset	= 5[ppm]
Clipped	= FALSE
Scans	= 4096
Total_Scans	= 4096
Relaxation_Delay	= 2[s]
Recvr Gain	= 56
Temp Get	= 24.8[dC]
X 90_Width	= 14[us]
X Acq Time	= 0.8388608[s]
X Angle	= 30[deg]
X Atn	= 11[dB]
X Pulse	= 4.66666667[us]
Irr_Atn_Dec	= 25.8[dB]
Irr_Atn_Dec_Calc	= 25.8[dB]
Irr_Atn_Dec_Default_Calc	= 25.8[dB]
Irr_Atn_No	= 25.8[dB]
Irr_Dec_Bandwidth_Hz	= 5.97826087[kHz]
Irr_Dec_Bandwidth_Ppm	= 12.07407703[ppm]
Irr_Dec_Freq	= 495.13191398[MHz]
Irr_Dec_Merit_Factor	= 2.2
Irr_Decoupling	= TRUE
Irr Noe	= TRUE
Irr Noise	= WALTZ
Irr Offset Default	= 5[ppm]
Irr Pwidth	= 92[us]
Irr_Pwidth Default	= 92[us]
Irr_Pwidth Default_Calc	= 92[us]
Irr_Pwidth_Templ	= 92[us]
Irr_Wurst	= FALSE
Decimation_Rate	= 0
Initial_Wait	= 1[s]
Noe_Time	= 2[s]
Noe_Time_Flag	= FALSE
Relaxation_Delay_Calc	= 0[s]
Relaxation_Delay_Temp	= 2[s]
Repetition_Time	= 2.8388608[s]

YF-0-p-OH_Proton-1-5.jdf

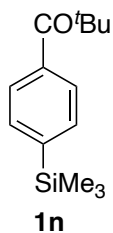


Filename = YF-0-p-OH_Proton-1-5.
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-0-p-OH
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 11-NOV-2022 19:34:07
 Revision_Time = 11-NOV-2022 18:35:33
 Comment = single_pulse
 Data_Format = 1D REAL
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1
 Field_Strength = 11.62926421 [T] (500 [M])
 X_Acq_Duration = 1.76422912 [s]
 X_Domain = 1H
 X_Freq = 495.13191398 [MHz]
 X_Offset = 5 [ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198 [Hz]
 X_Sweep = 9.28677563 [kHz]
 X_Sweep_Clipped = 7.42942051 [kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398 [MHz]
 Irr_Offset = 5 [ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398 [MHz]
 Tri_Offset = 5 [ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8
 Relaxation_Delay = 5 [s]
 Recvr_Gain = 56
 Temp_Get = 19.9 [dC]
 X_90_Width = 8.7 [us]
 X_Acq_Time = 1.76422912 [s]
 X_Angle = 45 [deg]
 X_Atn = 4.3 [dB]
 X_Pulse = 4.35 [us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1 [s]
 Phase = {0, 90, 270, 180, 180}
 Presat_Time = 5 [s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0 [s]
 Relaxation_Delay_Temp = 5 [s]
 Repetition_Time = 6.76422912 [s]



X : parts per Million : Proton

YF-1183-4_Proton-1-4.jdf



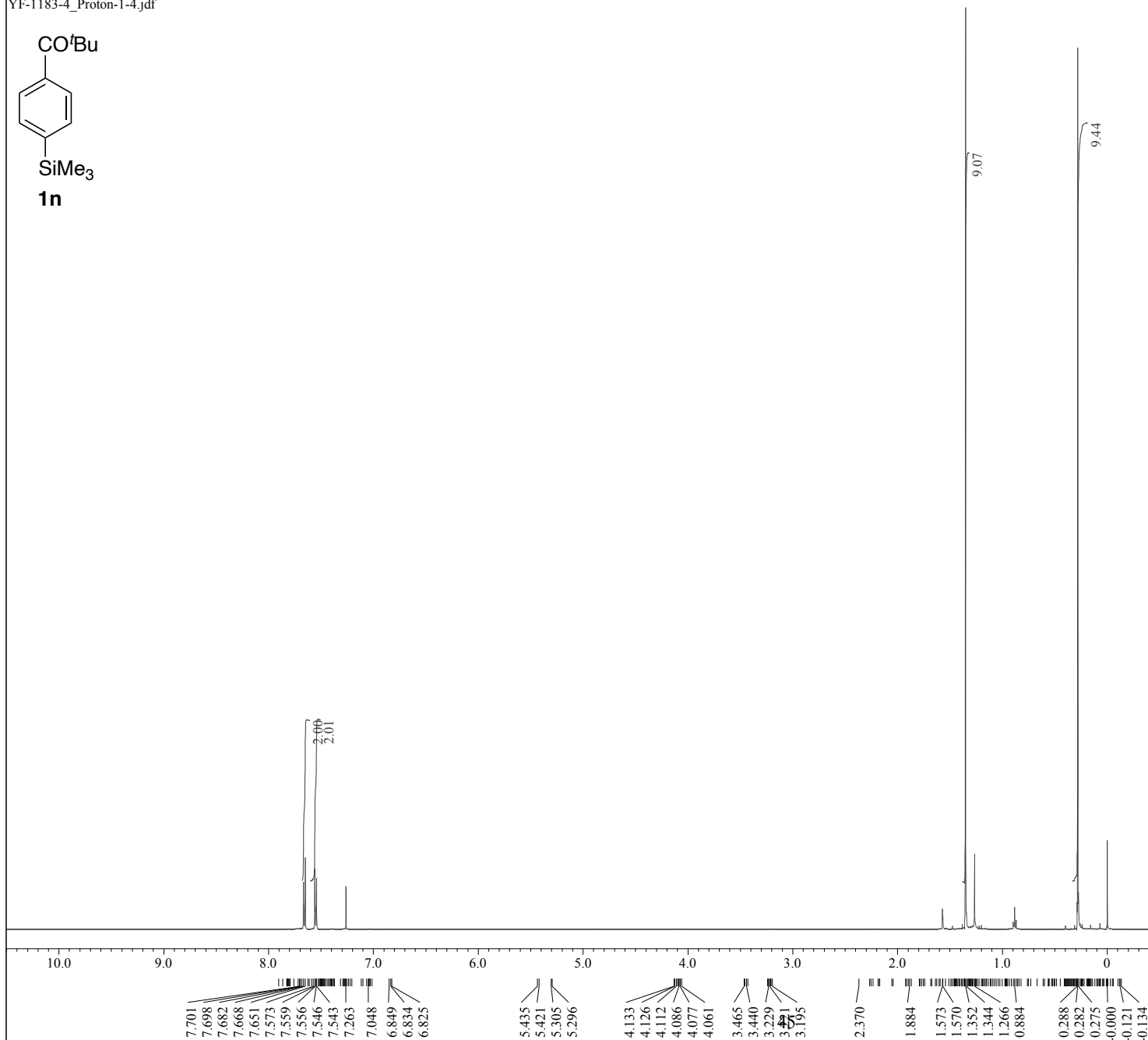
```

Filename      = YF-1183-4_Proton-1-4.
Author       = delta
Experiment    = proton.jxp
Sample_Id     = YF-1183-4
Solvent       = CHLOROFORM-D
Actual_Start_Time = 26-OCT-2022 21:49:09
Revision_Time  = 26-OCT-2022 20:51:44

Comment      = single_pulse
Data_Format   = 1D REAL
Dim_Size      = 13107
Dim_Title     = Proton
Dim_Units     = [ppm]
Dimensions    = X
Spectrometer  = JNM-ECZ500R/S1

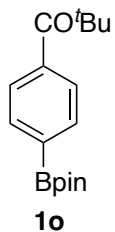
Field_Strength = 11.62926421 [T] (500 [M]
X_Acq_Duration = 1.76422912 [s]
X_Domain       = 18
X_Freq         = 495.13191398 [MHz]
X_Offset       = 5 [ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198 [Hz]
X_Sweep        = 9.28677563 [kHz]
X_Sweep_Clipped = 7.42942051 [kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398 [MHz]
Irr_Offset     = 5 [ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398 [MHz]
Tri_Offset     = 5 [ppm]
Clipped        = FALSE
Scans          = 8
Total_Scans    = 8

Relaxation_Delay = 5 [s]
Recvr_Gain       = 46
Temp_Get         = 19.8 [dC]
X_90_Width       = 8.7 [us]
X_Acq_Time       = 1.76422912 [s]
X_Angle          = 45 [deg]
X_Atn            = 4.3 [dB]
X_Pulse          = 4.35 [us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 500
Dante_Presat     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1 [s]
Phase            = {0, 90, 270, 180, 180}
Presat_Time      = 5 [s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0 [s]
Relaxation_Delay_Temp = 5 [s]
Repetition_Time  = 6.76422912 [s]
    
```



X : parts per Million : Proton

YF-1187-2_Proton-1-2.jdf



```

---- PROCESSING PARAMETERS ----
sexf( 0.2[Hz], 0.0[s] )
trapezoid( 0[%], 0[%], 80[%], 100[%] )
sexfill( 1, TRUE )
fft( 1, TRUE, TRUE )
machinphase
ppm
base_correct( Akima, 5, 0, FALSE, 3, None, FA
以下に由来 : YF-1187-2_Proton-1-1.jdf
  
```

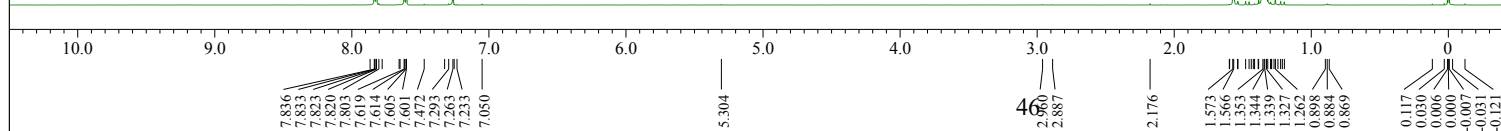
```

Filename      = YF-1187-2_Proton-1-2.
Author        = delta
Experiment     = proton.jpg
Sample_Id      = YF-1187-2
Solvent        = CHLOROFORM-D
Actual_Start_Time = 28-OCT-2022 20:02:12
RevisiOn_Time  = 29-OCT-2022 15:18:43

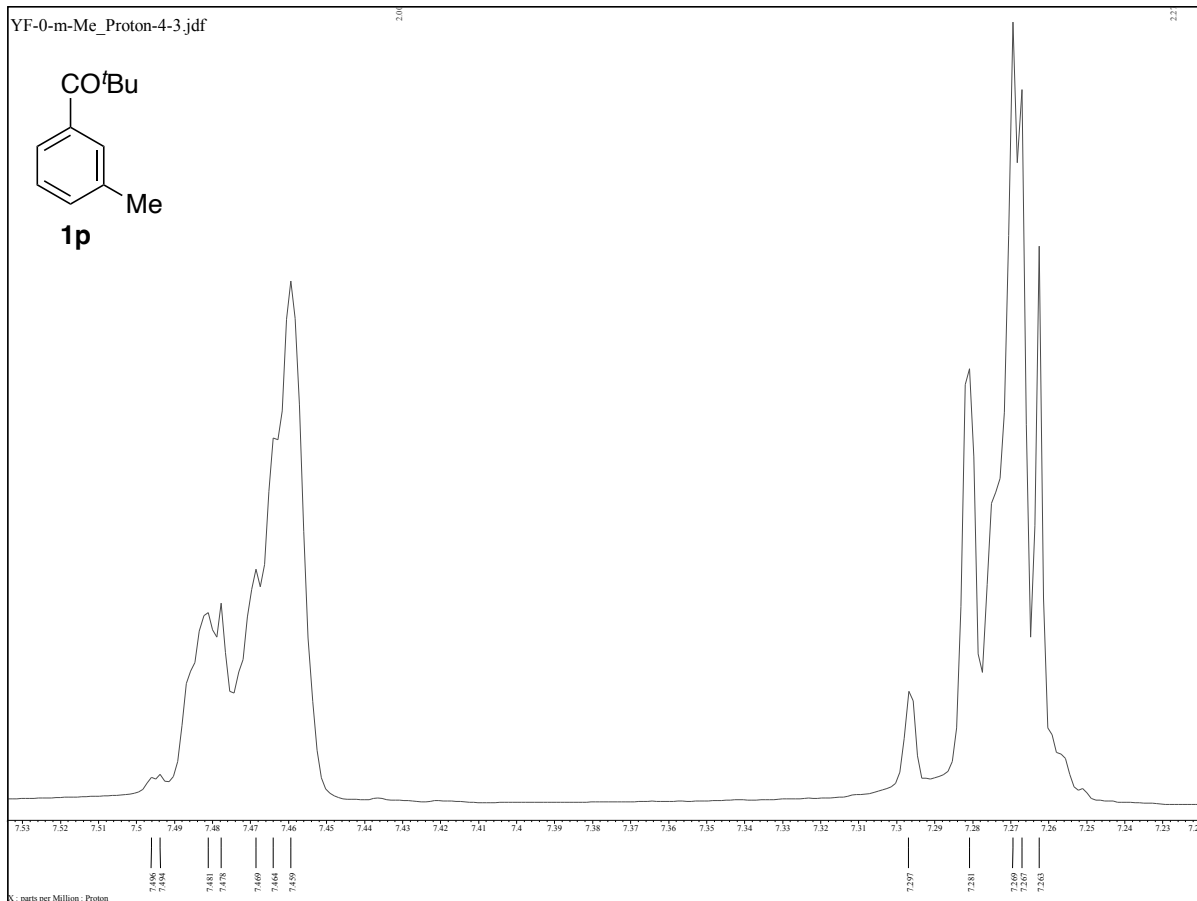
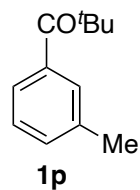
Comment       = single_pulse
Data_Format   = 1D_REAL
Dim_Size      = 13107
X_Domain      = Proton
Dim_Title     = Proton
Dim_Units     = [ppm]
Dimensions    = X
Spectrometer  = JNM-ECZ500R/S1

Field_Strength = 11.62926421[T] (500[M
X_Acq_Duration = 1.76422912[s]
X_Domain       = 1H
X_Freq         = 495.13191398[MHz]
X_Offset       = 5[ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198[Hz]
X_Sweep        = 9.28677563[kHz]
X_Sweep_Clippped = 7.42942051[kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398[MHz]
Irr_Offset     = 5[ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398[MHz]
Tri_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 8
Total_Scans    = 8

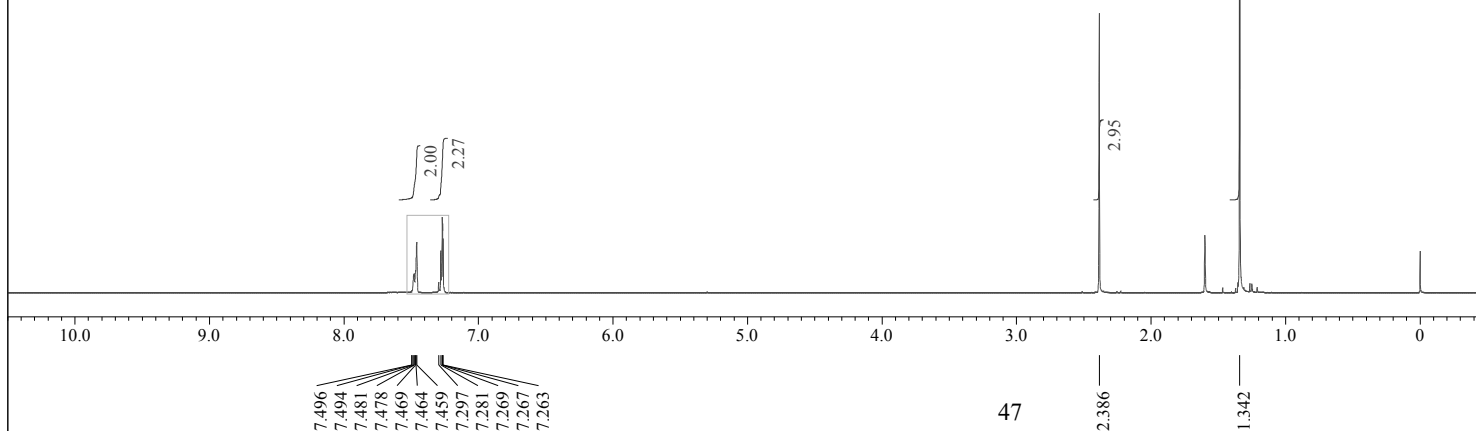
Relaxation_Delay = 5[s]
Recvr_Gain       = 56
Temp_Get         = 19.1[dC]
X_90_Width       = 8.7[us]
X_Acq_Time       = 1.76422912[s]
X_Angle          = 45[deg]
X_Atn            = 4.3[dB]
X_Pulse         = 4.35[us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 500
Dante_Presat     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1[s]
Phase           = {0, 90, 270, 180, 180
Presat_Time      = 5[s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 5[s]
Repetition_Time  = 6.76422912[s]
  
```



X : parts per Million : Proton



X : parts per Million : Proton



X : parts per Million : Proton



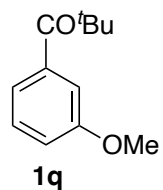
Filename = YF-0-m-Me_Proton-4-3.
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-0-m-Me
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 8-JUL-2022 15:19:07
 Revision_Time = 8-JUL-2022 14:38:13

Comment = single_pulse
 Data_Format = 1D COMPLEX
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1

Field_Strength = 11.62926421 [T] (500[M]
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clipped = 7.42942051[kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 32
 Total_Scans = 32

Relaxation_Delay = 5[s]
 Recvr_Gain = 36
 Temp_Get = 20.2[dc]
 X_90_Width = 9.49[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[db]
 X_Pulse = 4.745[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Preset = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

YF-0-m-OMe_Proton-2-3.jdf

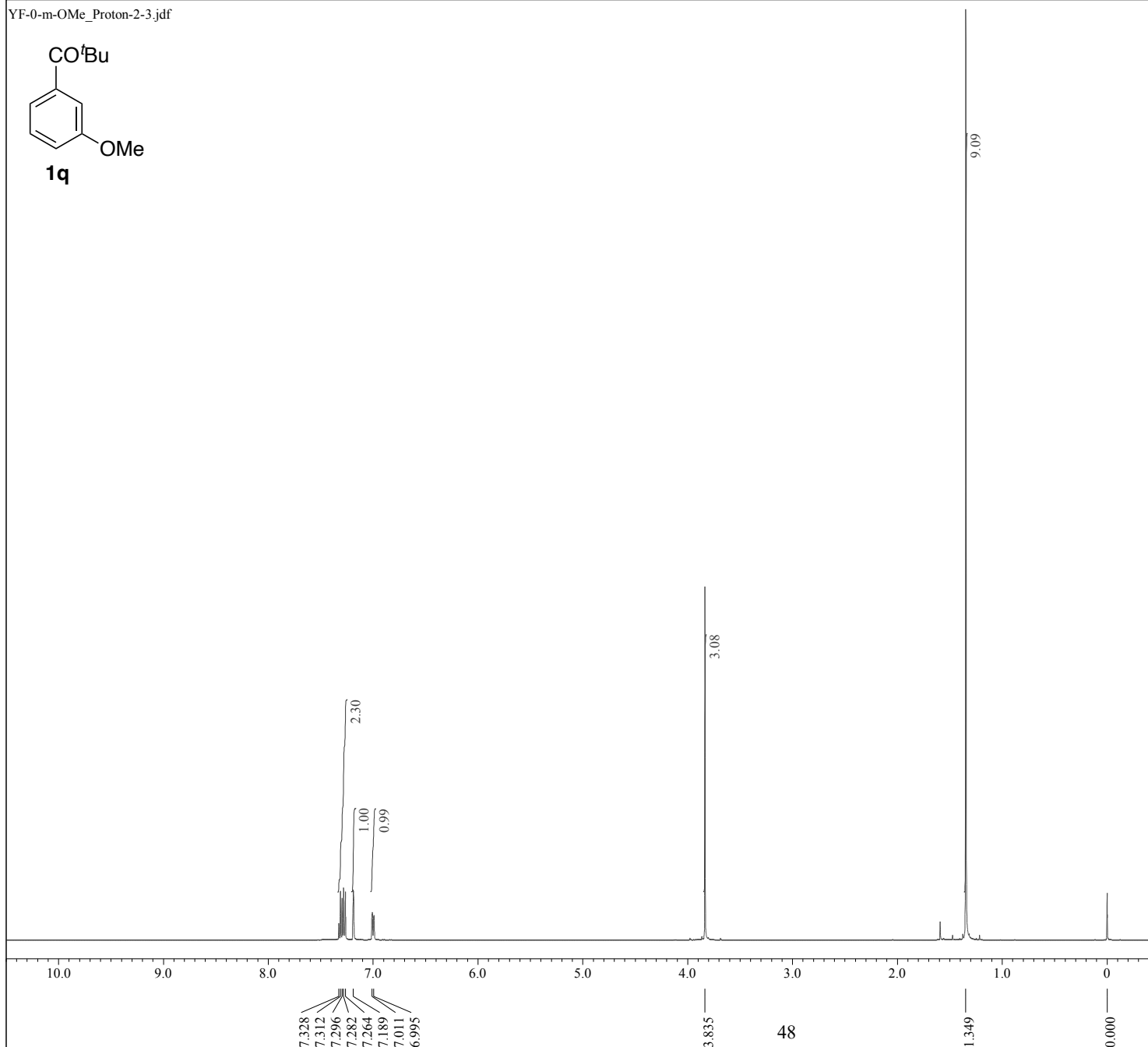


Filename = YF-0-m-OMe_Proton-2-2
 Author = delta
 Experiment = proton.jxp
 Sample_Id = YF-0-m-OMe
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 23-NOV-2022 17:48:44
 Revision_Time = 23-NOV-2022 16:55:42

Comment = single_pulse
 Data_Format = 1D COMPLEX
 Dim_Size = 13107
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1

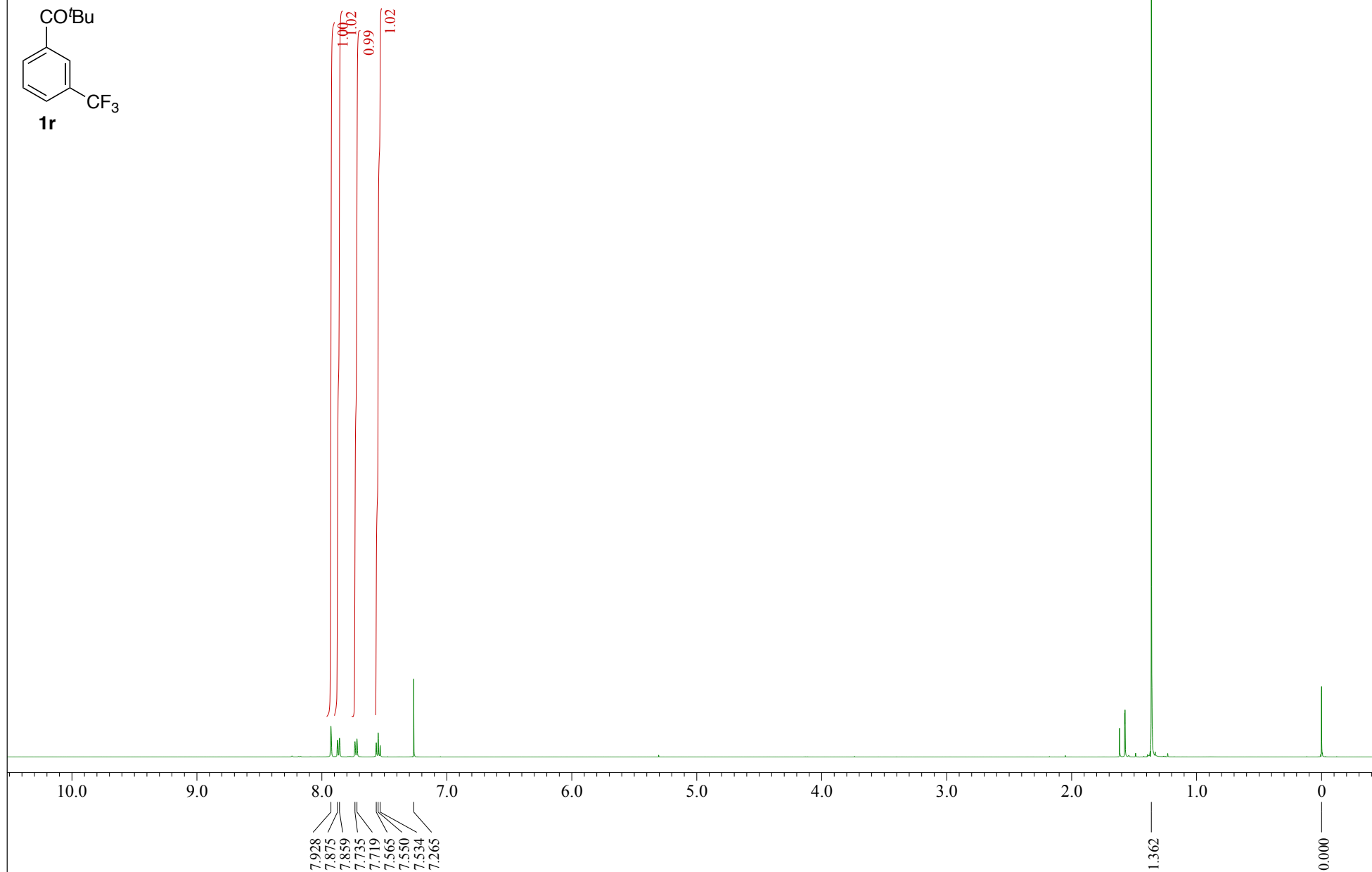
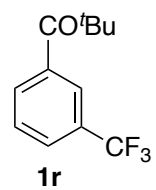
Field_Strength = 11.62926421 [T] (500 [M])
 X_Acq_Duration = 1.76422912 [s]
 X_Domain = 1H
 X_Freq = 495.13191398 [MHz]
 X_Offset = 5 [ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198 [Hz]
 X_Sweep = 9.28677563 [kHz]
 X_Sweep_Clipped = 7.42942051 [kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398 [MHz]
 Irr_Offset = 5 [ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398 [MHz]
 Tri_Offset = 5 [ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8

Relaxation_Delay = 5 [s]
 Recvr_Gain = 36
 Temp_Get = 19.6 [dC]
 X_90_Width = 8.7 [us]
 X_Acq_Time = 1.76422912 [s]
 X_Angle = 45 [deg]
 X_Atn = 4.3 [dB]
 X_Pulse = 4.35 [us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1 [s]
 Phase = {0, 90, 270, 180, 180}
 Presat_Time = 5 [s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0 [s]
 Relaxation_Delay_Temp = 5 [s]
 Repetition_Time = 6.76422912 [s]



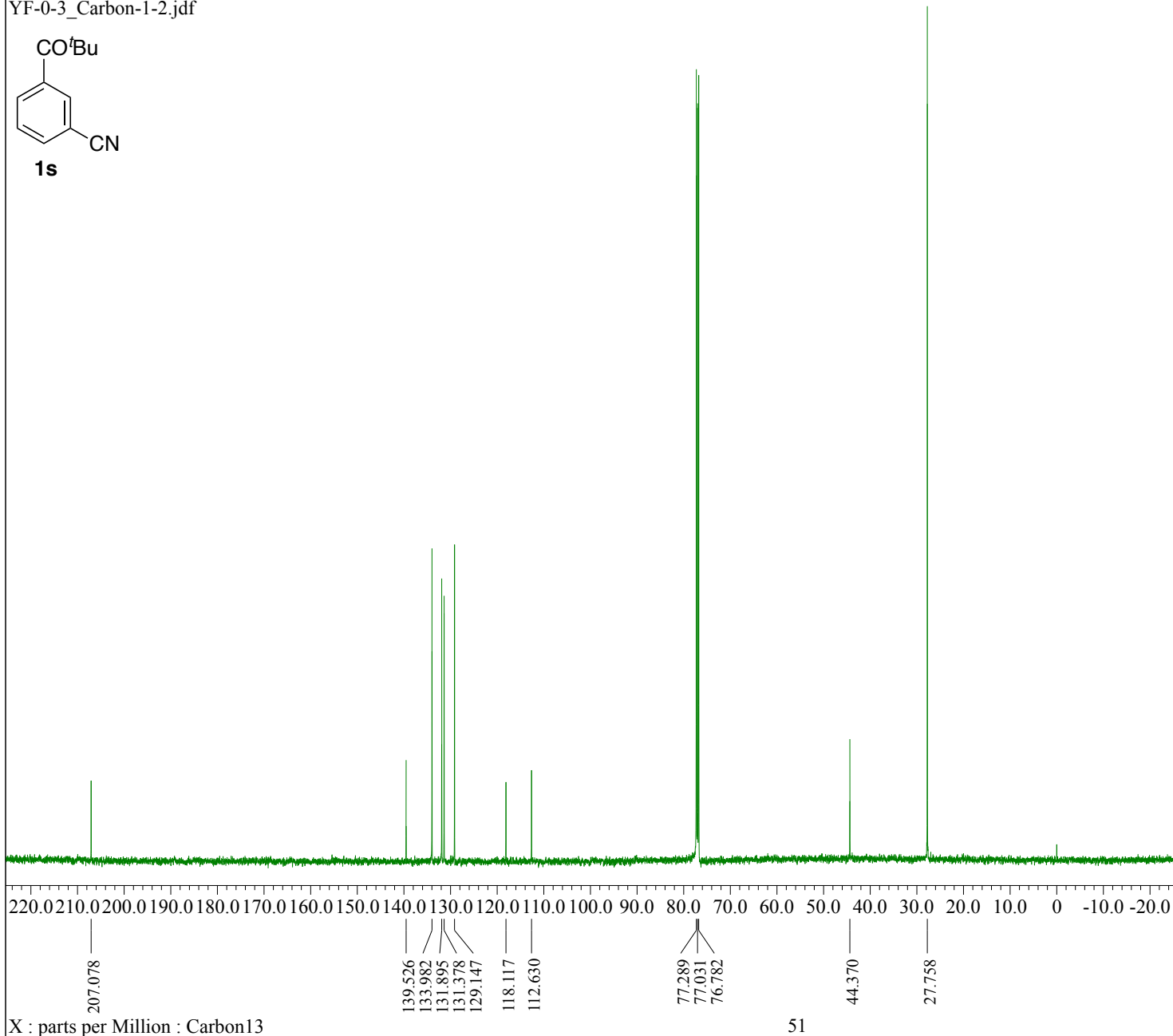
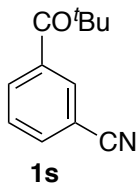
X : parts per Million : Proton

YF-1138-3_Proton-1-2.jdf



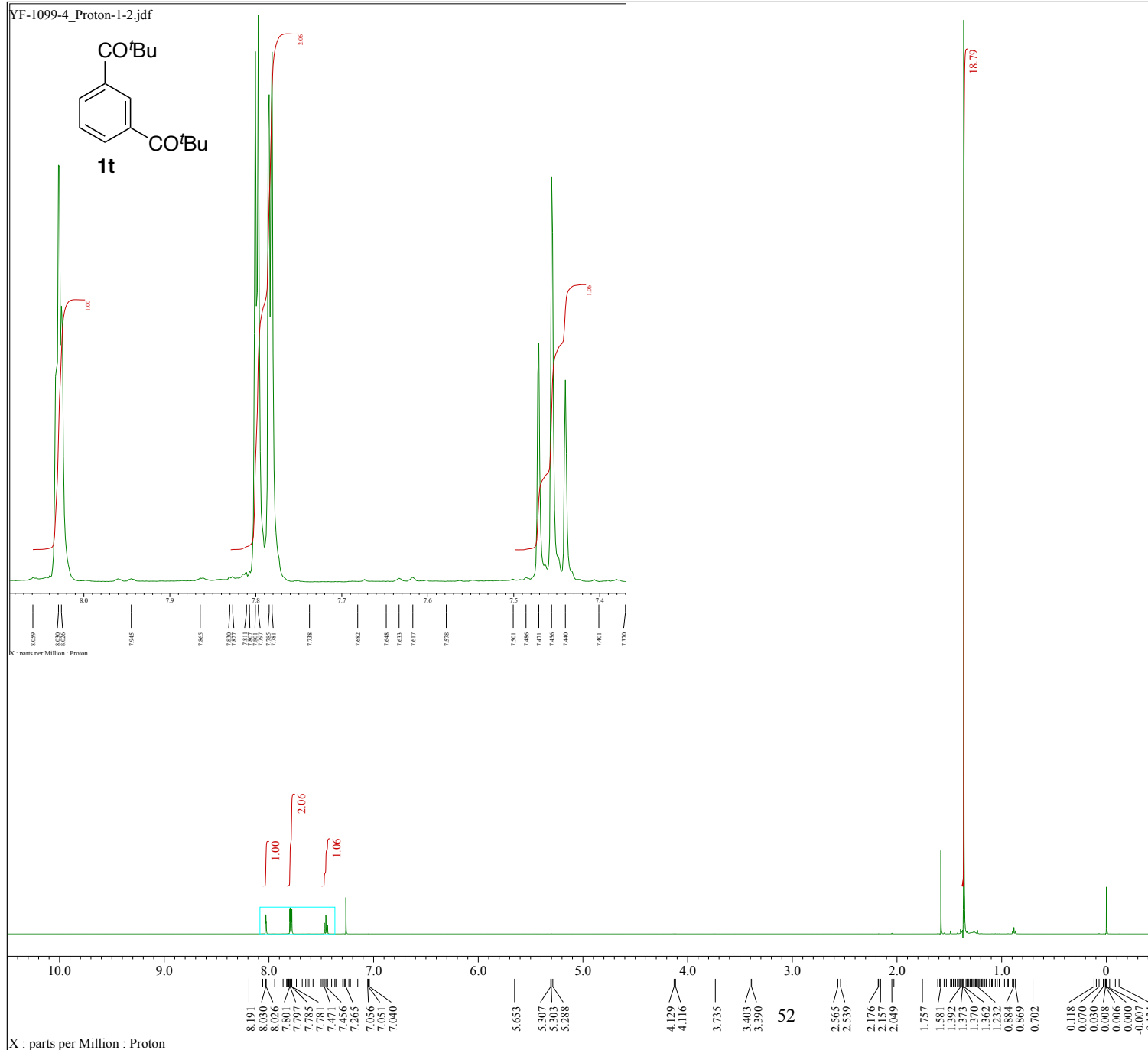
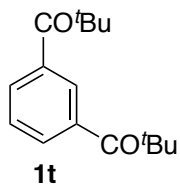
X : parts per Million : Proton

YF-0-3_Carbon-1-2.jdf



Filename	= YF-0-3_Carbon-1-2.
Author	= delta
Experiment	= carbon.jxp
Sample_Id	= YF-0-3
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 21-MAY-2022 03:58:
Revision_Time	= 23-MAY-2022 15:48:
Comment	= single pulse decou
Data_Format	= 1D COMPLEX
Dim_Size	= 26214
X_Domain	= Carbon13
Dim_Title	= Carbon13
Dim_Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ500R/S1
Field_Strength	= 11.62926421[T] (50
X_Acq_Duration	= 0.8388608[s]
X_Domain	= 13C
X_Freq	= 124.5010059[MHz]
X_Offset	= 100[ppm]
X_Points	= 32768
X_Prescans	= 4
X_Resolution	= 1.1920929[Hz]
X_Sweep	= 39.0625[kHz]
X_Sweep_Clippped	= 31.25[kHz]
Irr_Domain	= Proton
Irr_Freq	= 495.13191398[MHz]
Irr_Offset	= 5[ppm]
Clipped	= FALSE
Scans	= 1024
Total_Scans	= 1024
Relaxation_Delay	= 2[s]
Recvr_Gain	= 56
Temp_Get	= 25.3[dC]
X_90_Width	= 14[us]
X_Acq_Time	= 0.8388608[s]
X_Angle	= 30[deg]
X_Atn	= 11[dB]
X_Pulse	= 4.66666667[us]
Irr_Atn_Dec	= 25.8[dB]
Irr_Atn_Dec_Calc	= 25.8[dB]
Irr_Atn_Dec_Default_Calc	= 25.8[dB]
Irr_Atn_No	= 25.8[dB]
Irr_Dec_Bandwidth_Hz	= 5.97826087[kHz]
Irr_Dec_Bandwidth_Ppm	= 12.07407703[ppm]
Irr_Dec_Freq	= 495.13191398[MHz]
Irr_Dec_Merit_Factor	= 2.2
Irr_Decoupling	= TRUE
Irr_No	= TRUE
Irr_Noise	= WALTZ
Irr_Offset_Default	= 5[ppm]
Irr_Pwidth	= 92[us]
Irr_Pwidth_Default	= 92[us]
Irr_Pwidth_Default_Calc	= 92[us]
Irr_Pwidth_Templ	= 92[us]
Irr_Wurst	= FALSE
Decimation_Rate	= 0
Initial_Wait	= 1[s]
Noe_Time	= 2[s]
Noe_Time_Flag	= FALSE
Relaxation_Delay_Calc	= 0[s]
Relaxation_Delay_Temp	= 2[s]
Repetition_Time	= 2.8388608[s]

YF-1099-4_Proton-1-2.jdf

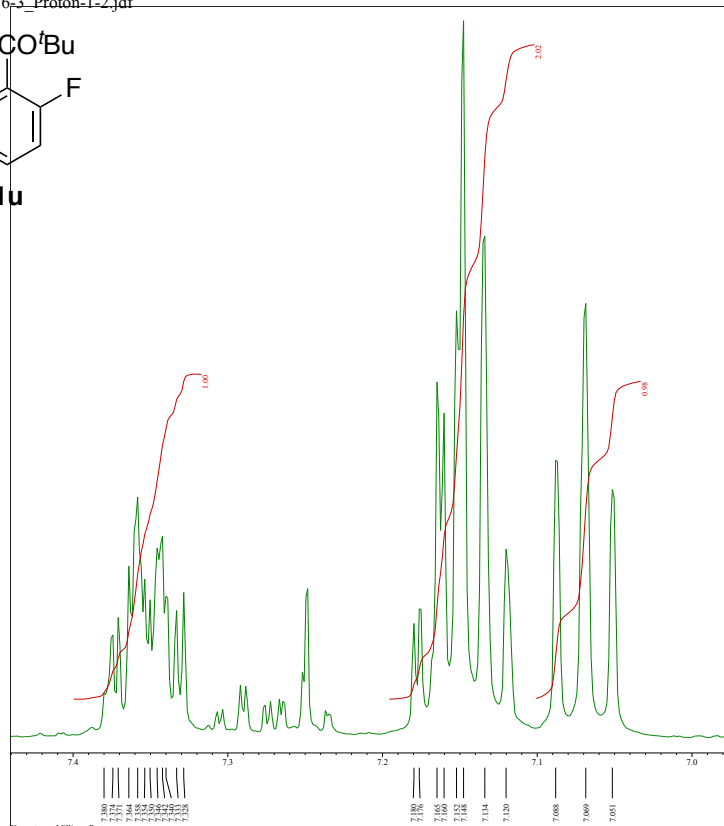
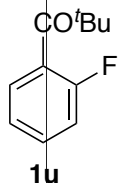


----- PROCESSING PARAMETERS -----
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm
 base_correct(Akima, 5, 0, FALSE, 3, None, FA
 以下に由来 : YF-1099-4_Proton-1-1.jdf

Filename = YF-1099-4_Proton-1-2.
 Author = delta
 Experiment = proton.jpg
 Sample_Id = YF-1099-4
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 11-OCT-2022 15:10:30
 Revision_Time = 11-OCT-2022 20:28:03
 Comment = single_pulse
 Data_Format = 1D_REAL
 Dim_Size = 13107
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1
 Field_Strength = 11.62926421[T] (500[M
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clippped = 7.42942051[kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8
 Relaxation_Delay = 5[s]
 Recvz_Gain = 46
 Temp_Get = 20.1[dC]
 X_90_Width = 8.7[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[dB]
 X_Pulse = 4.35[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Preset = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

X : parts per Million : Proton

YF-816-3_Proton-1-2.jdf



7.364
7.358
7.354
7.346
7.342
7.340
7.328
7.165
7.152
7.148
7.134
7.120
7.088
7.069
7.051

X : parts per Million : Proton

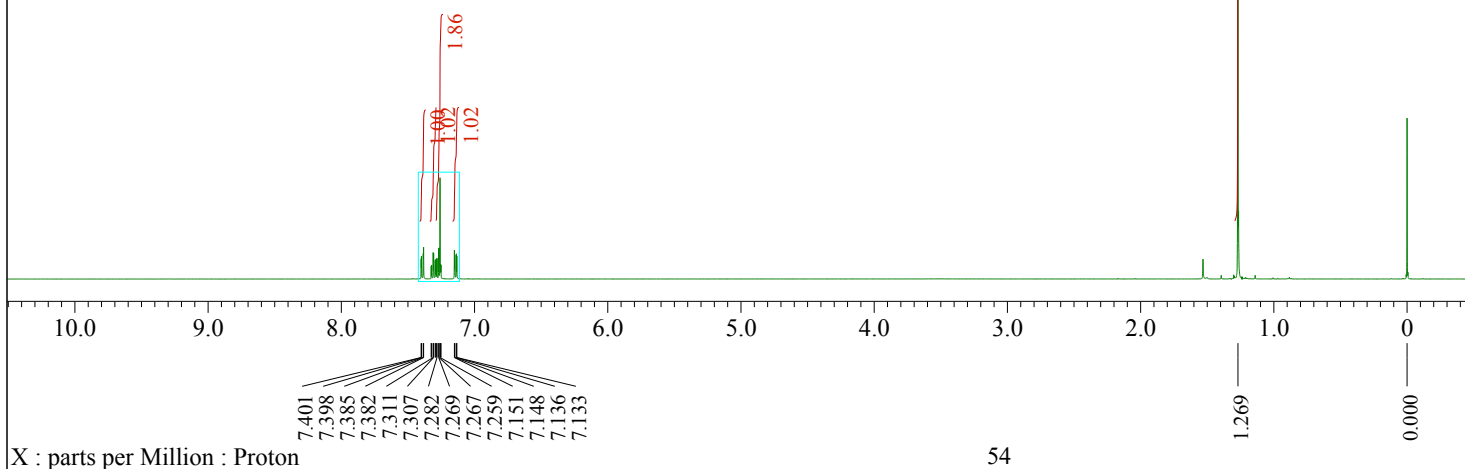
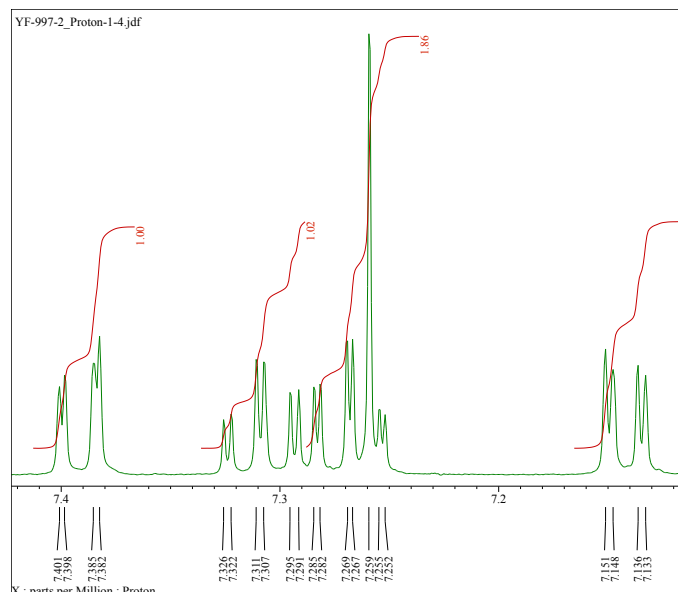
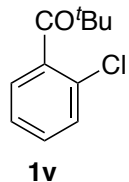
53



---- PROCESSING PARAMETERS ----
sexp(0.2[Hz], 0.0[s])
trapezoid(0[%], 0[%], 80[%], 100[%])
zerofill(1, TRUE)
fft(1, TRUE, TRUE)
machinephase
ppm
以下に由来 : YF-816-3_Proton-1-1.jdf

Filename = YF-816-3_Proton-1-2.j
Author = delta
Experiment = proton.jpg
Sample_Id = YF-816-3
Solvent = CHLOROFORM-D
Actual_Start_Time = 7-MAR-2022 22:07:59
Revision_Time = 9-JUL-2022 11:54:30
Comment = single_pulse
Data_Format = 1D COMPLEX
Dim_Size = 13107
X_Domain = Proton
Dim_Title = Proton
Dim_Units = [ppm]
Dimensions = X
Spectrometer = JNM-EZ500R/S1
Field_Strength = 11.62926421[T] (500[M
X_Acq_Duration = 1.76422912[s]
X_Domain = 1H
X_Freq = 495.13191398[MHz]
X_Offset = 5[ppm]
X_Points = 16384
X_Prescans = 1
X_Resolution = 0.5668198[Hz]
X_Sweep = 9.28677563[kHz]
X_Sweep_Clipped = 7.42942051[kHz]
Irr_Domain = Proton
Irr_Freq = 495.13191398[MHz]
Irr_Offset = 5[ppm]
Tri_Domain = Proton
Tri_Freq = 495.13191398[MHz]
Tri_Offset = 5[ppm]
Clipped = FALSE
Scans = 32
Total_Scans = 32
Relaxation_Delay = 5[s]
Recvr_Gain = 26
Temp_Get = 20.4[dC]
X_90_Width = 9.47[us]
X_Acq_Time = 1.76422912[s]
X_Angle = 45[deg]
X_Atn = 4.3[dB]
X_Pulse = 4.735[us]
Irr_Mode = Off
Tri_Mode = Off
Dante_Loop = 500
Dante_Preset = FALSE
Decimation_Rate = 0
Initial_Wait = 1[s]
Phase = {0, 90, 270, 180, 180
Preset_Time = 5[s]
Preset_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 5[s]
Repetition_Time = 6.76422912[s]

YF-997-2_Proton-1-4.jdf



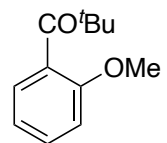
Filename = YF-997-2_Proton-1-4.jdf
 Author = takaya
 Experiment = proton.jxp
 Sample_Id = YF-997-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 13-JUN-2022 20:41:57
 Revision_Time = 19-JUL-2022 22:32:42

Comment = single pulse
 Data_Format = 1D COMPLEX
 Dim_Size = 26214
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Site = JNM-ECX500
 Spectrometer = DELTA2_NMR

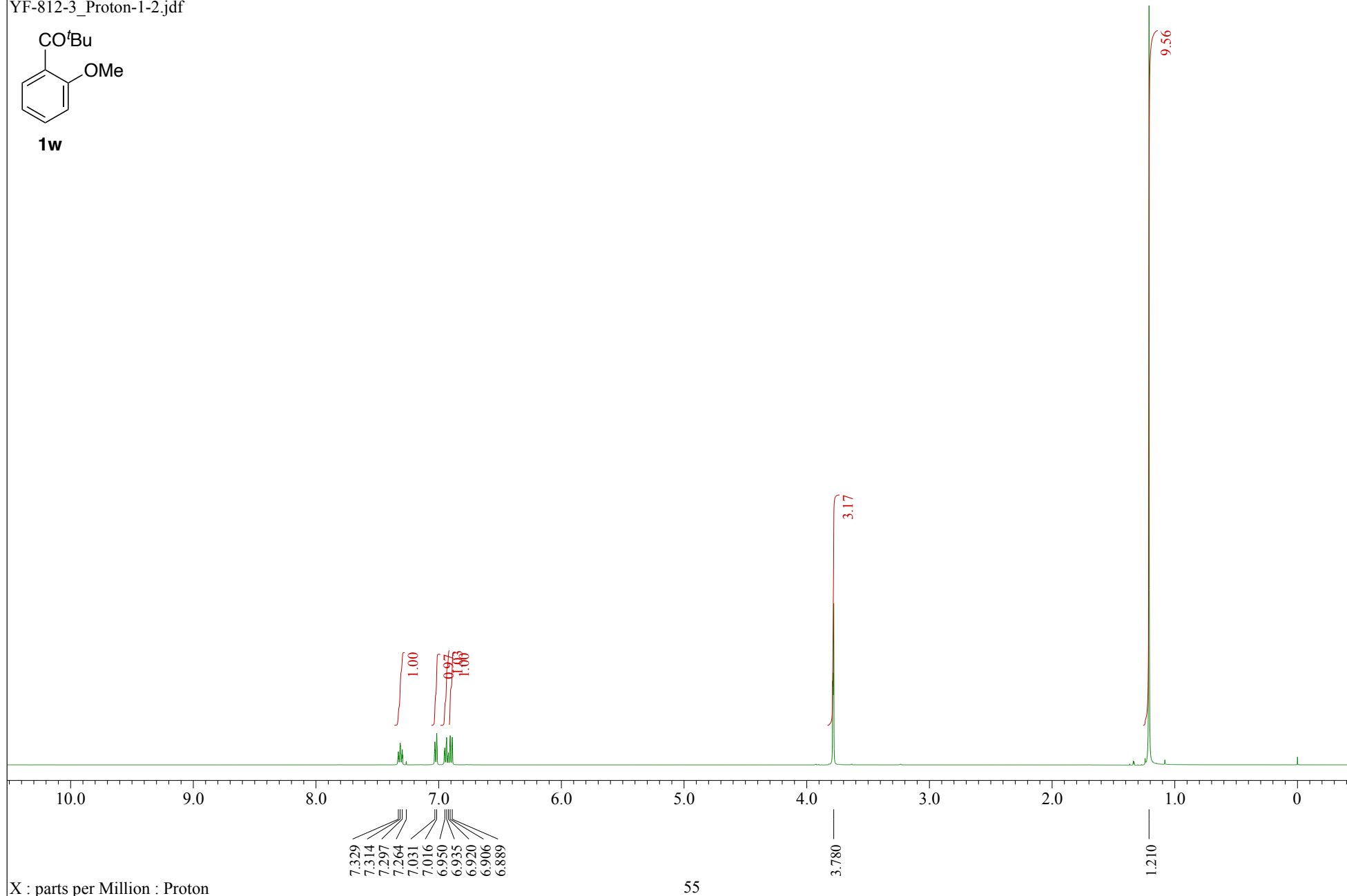
Field_Strength = 11.7473579[T] (500[MHz])
 X_Acq_Duration = 3.49175808[s]
 X_Domain = 1H
 X_Freq = 500.15991521[MHz]
 X_Offset = 5.0[ppm]
 X_Points = 32768
 X_Prescans = 1
 X_Resolution = 0.28638868[Hz]
 X_Sweep = 9.38438438[kHz]
 X_Sweep_Clippped = 7.50750751[kHz]
 Irr_Domain = Proton
 Irr_Freq = 500.15991521[MHz]
 Irr_Offset = 5.0[ppm]
 Tri_Domain = Proton
 Tri_Freq = 500.15991521[MHz]
 Tri_Offset = 5.0[ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8

Relaxation_Delay = 2[s]
 Recvr_Gain = 58
 Temp_Get = 26.5[dC]
 X_90_Width = 12.4[us]
 X_Acq_Time = 3.49175808[s]
 X_Angle = 45[deg]
 X_Atn = 4.5[dB]
 X_Pulse = 6.2[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Presat = FALSE
 Initial_Wait = 1[s]
 Repetition_Time = 5.49175808[s]

YF-812-3_Proton-1-2.jdf



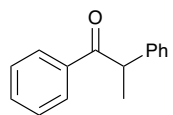
1w



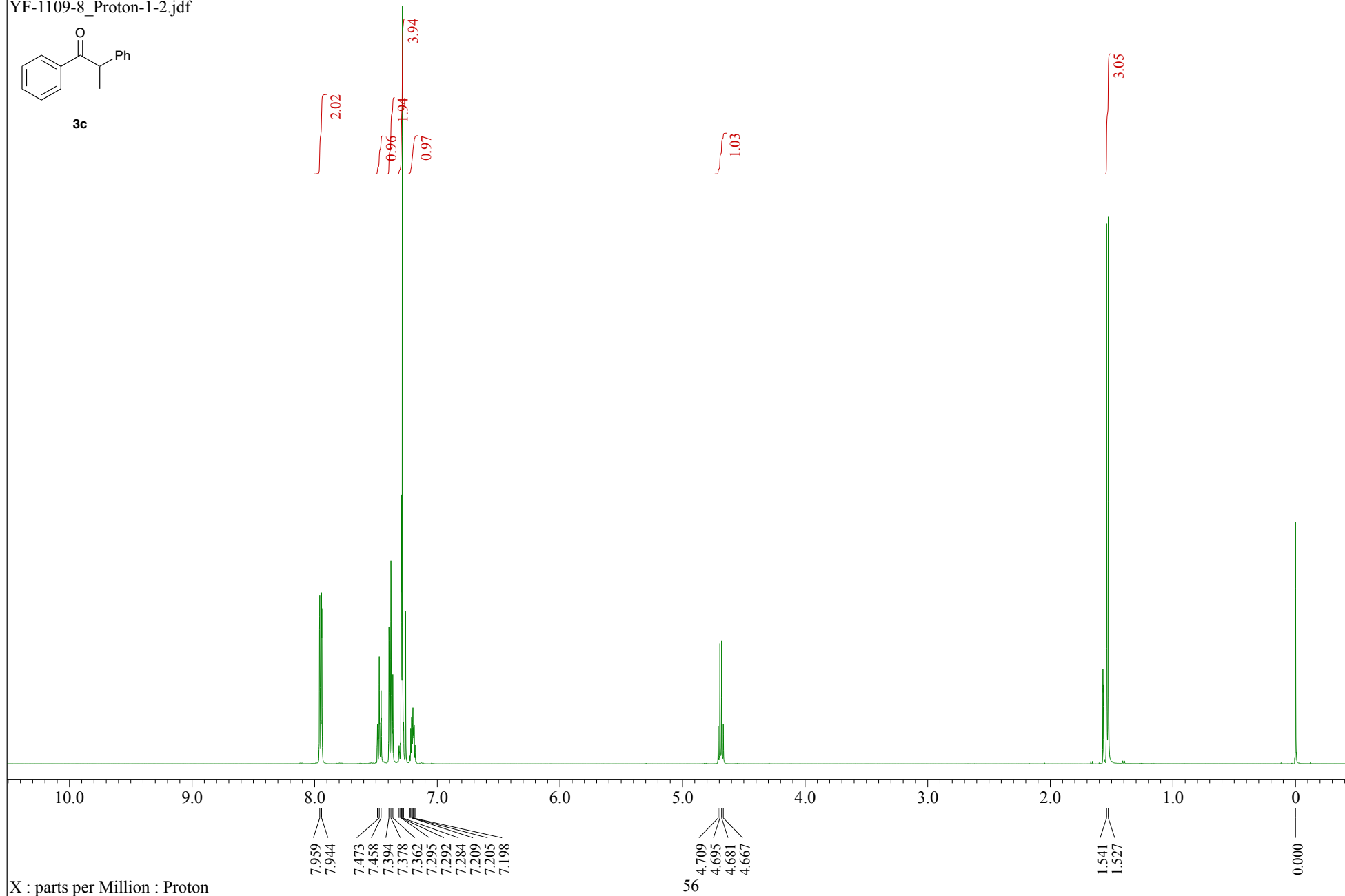
X : parts per Million : Proton

55

YF-1109-8_Proton-1-2.jdf

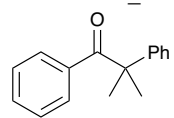


3c



X : parts per Million : Proton

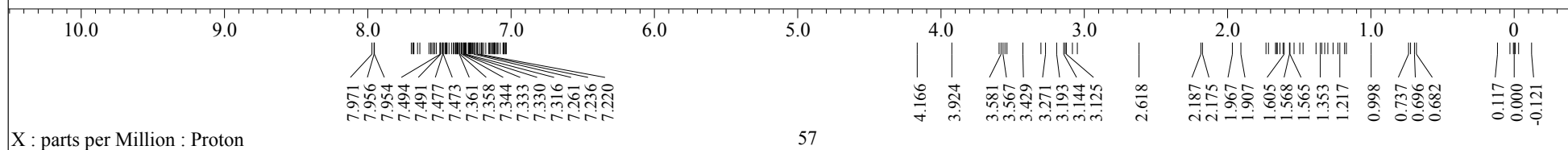
YF-0-Me2Ph_Proton-1-2.jdf



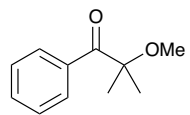
3d

2.00
1.98
5.07

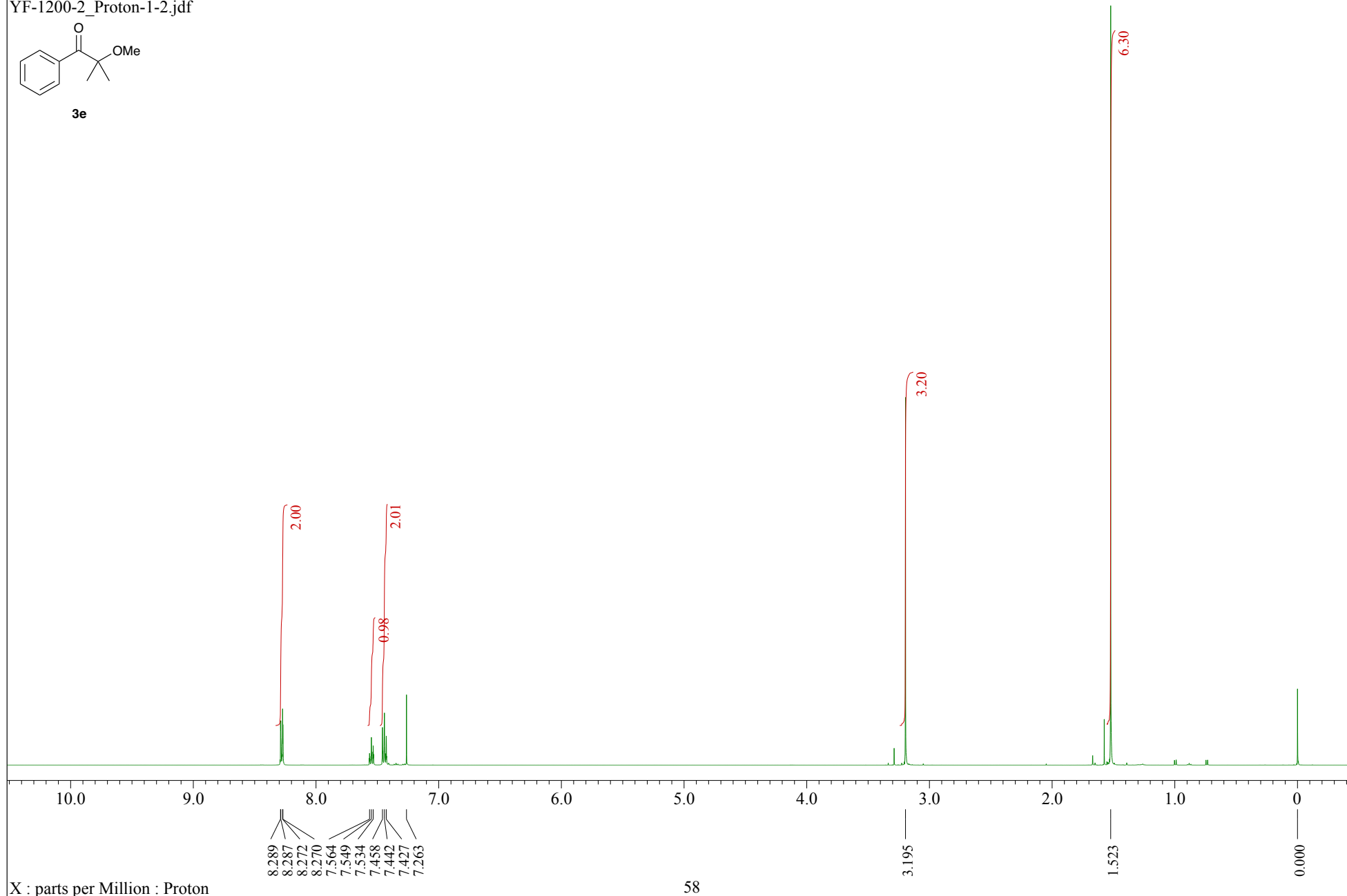
6.0



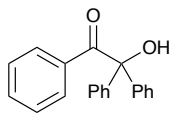
YF-1200-2_Proton-1-2.jdf



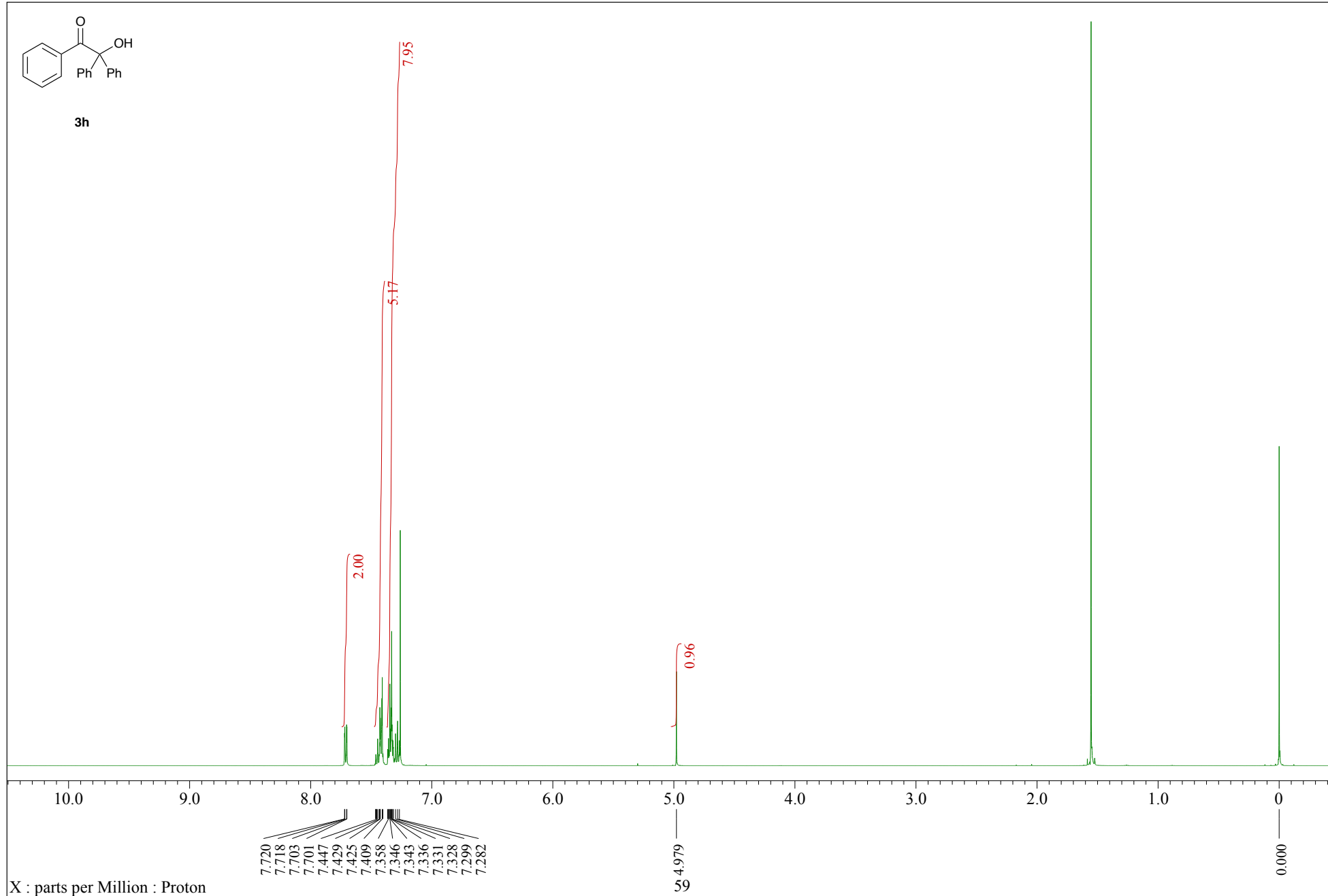
3e



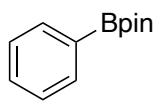
X : parts per Million : Proton



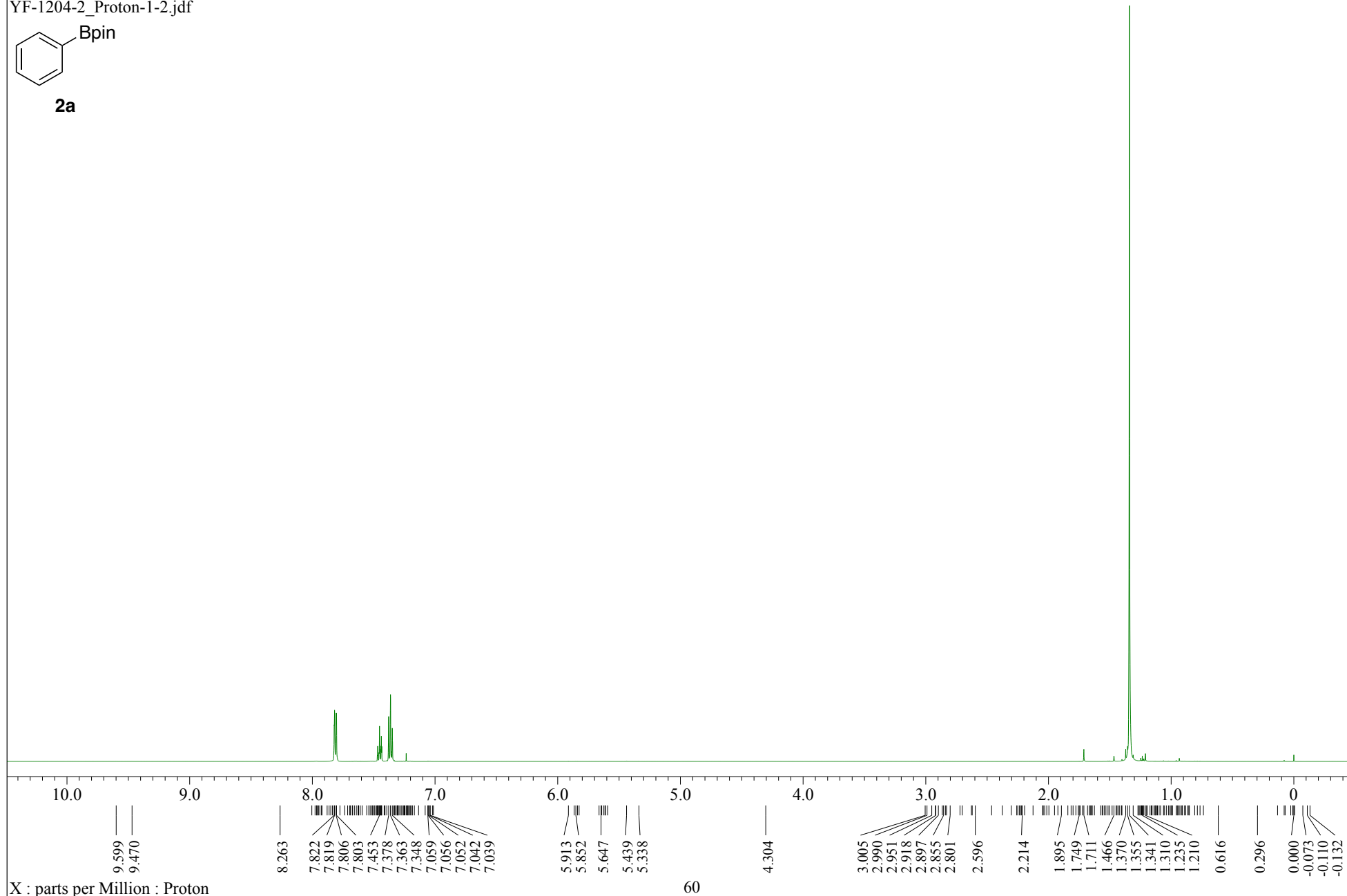
3h



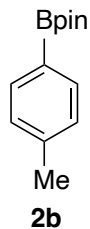
YF-1204-2_Proton-1-2.jdf



2a

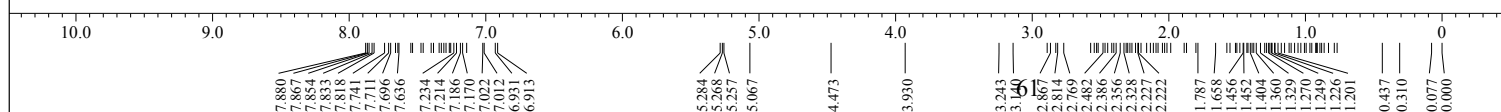


YF-965-2_Proton-2-2.jdf



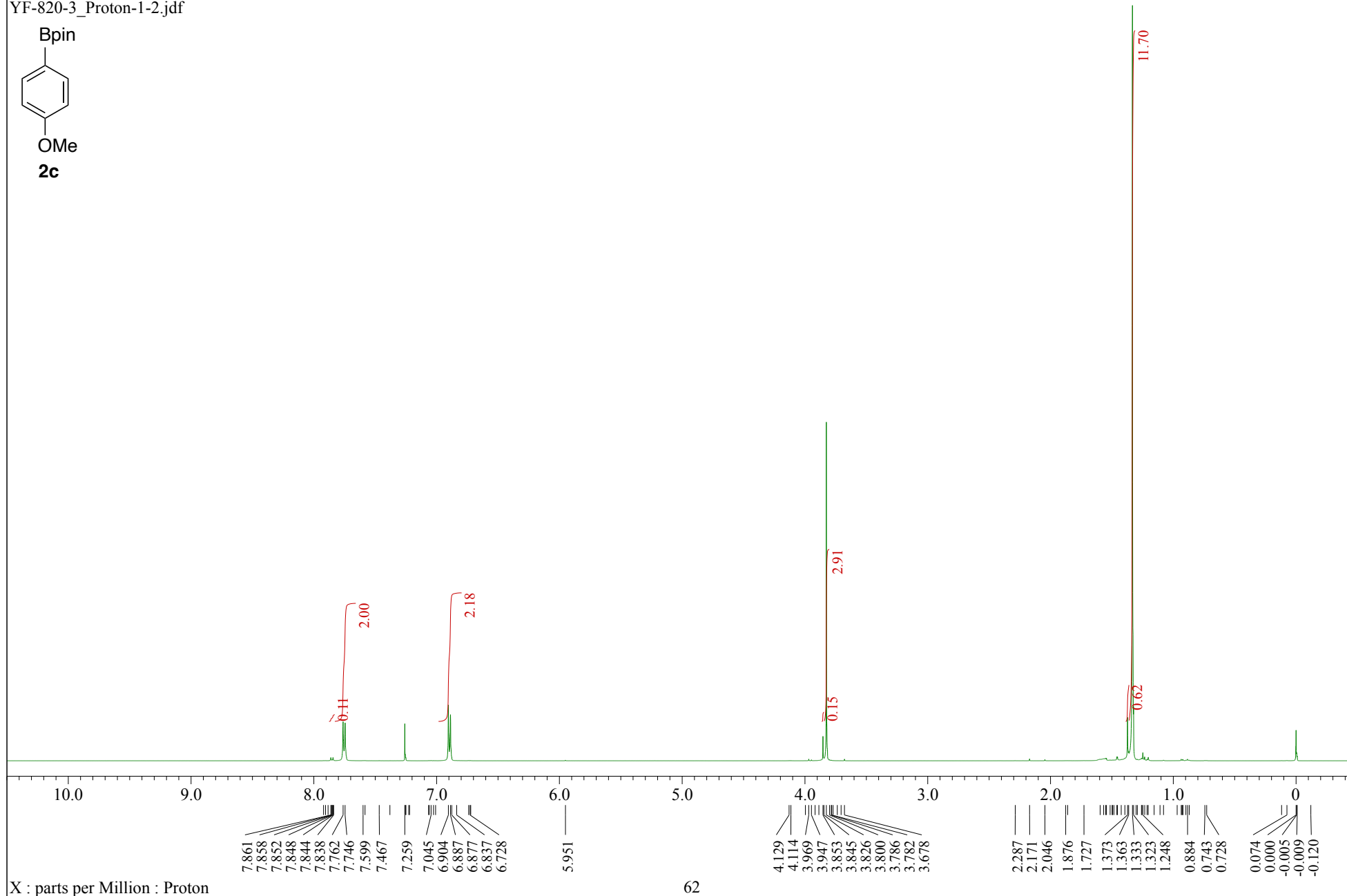
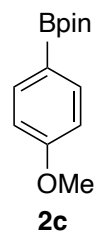
---- PROCESSING PARAMETERS ----
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm
 base_correct(Akima, 5, 0, FALSE, 3, None, FA
 以下に由来: YF-965-2_Proton-2-1.jdf

Filename = YF-965-2_Proton-2-2.j
 Author = delta
 Experiment = proton.jpg
 Sample_Id = YF-965-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 2-JUN-2022 17:10:48
 Revision_Time = 2-JUN-2022 19:57:33
 Comment = single_pulse
 Data Format = 1D REAL
 Dim_Size = 13107
 X_Domain = Proton
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECS500R/S1
 Field_Strength = 11.62926421[T] (500[M
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X Points = 16384
 X_Prescans = 1
 X Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clippped = 7.42942051[kHz]
 Iir Domain = Proton
 Iir_Freq = 495.13191398[MHz]
 Iir_Offset = 5[ppm]
 Tri Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 32
 Total_Scans = 32
 Relaxation_Delay = 10[s]
 Recvr Gain = 36
 Temp Get = 27[dC]
 X_90_Width = 7.5[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 3.3[dB]
 X_Pulse = 3.75[us]
 Iir Mode = Off
 Tri_Mode = Off
 Dante Loop = 1000
 Dante Presat = FALSE
 Decimation_Rate = 0
 Initial Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat Time = 10[s]
 Presat Time Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 10[s]
 Repetition_Time = 11.76422912[s]



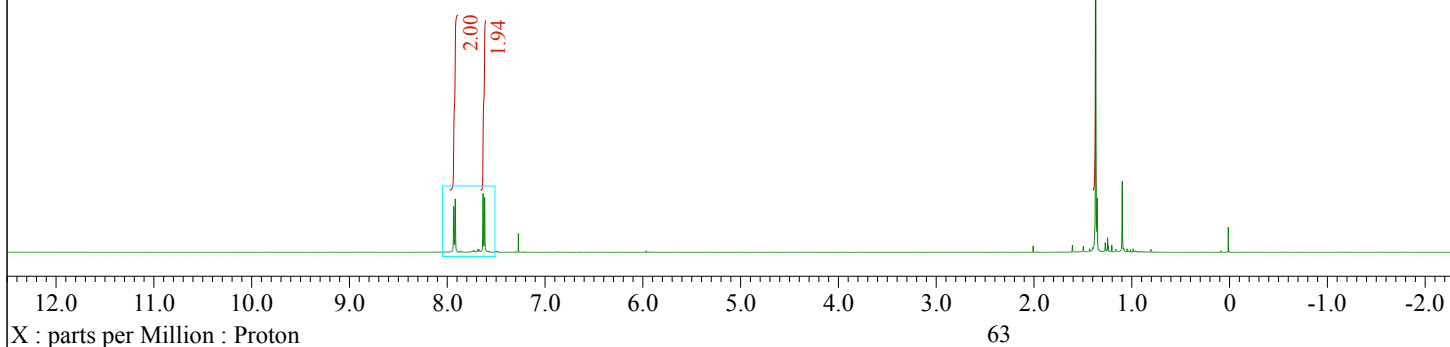
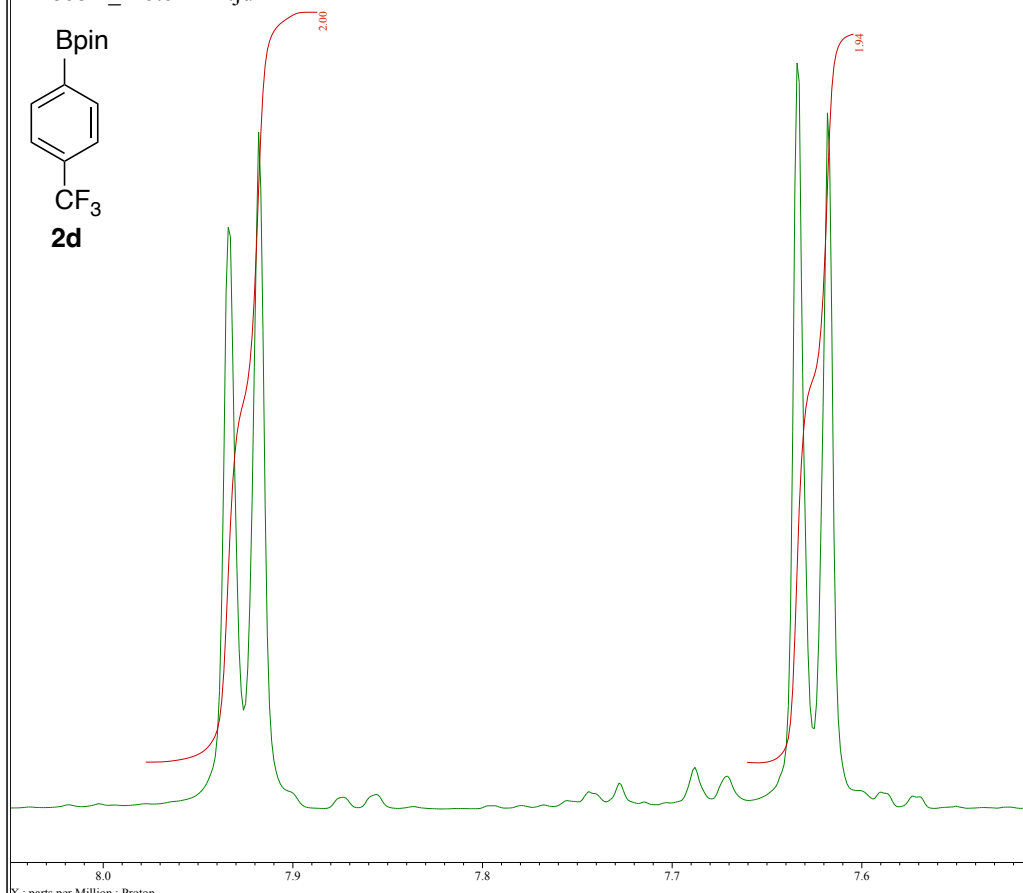
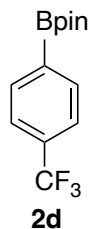
X : parts per Million : Proton

YF-820-3_Proton-1-2.jdf



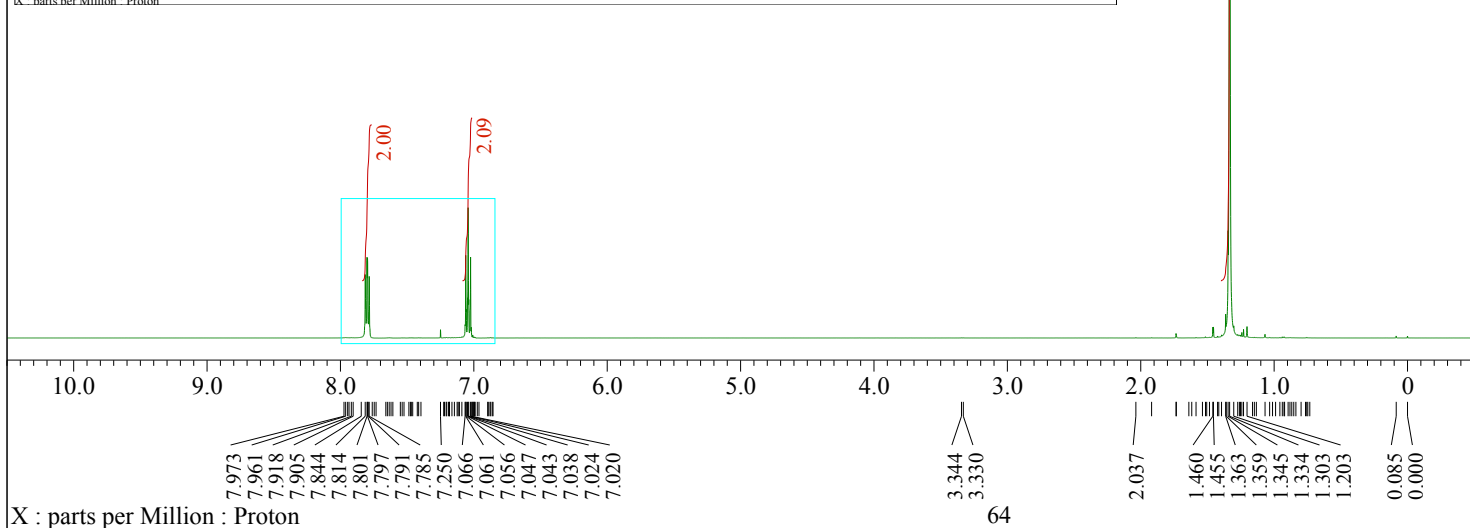
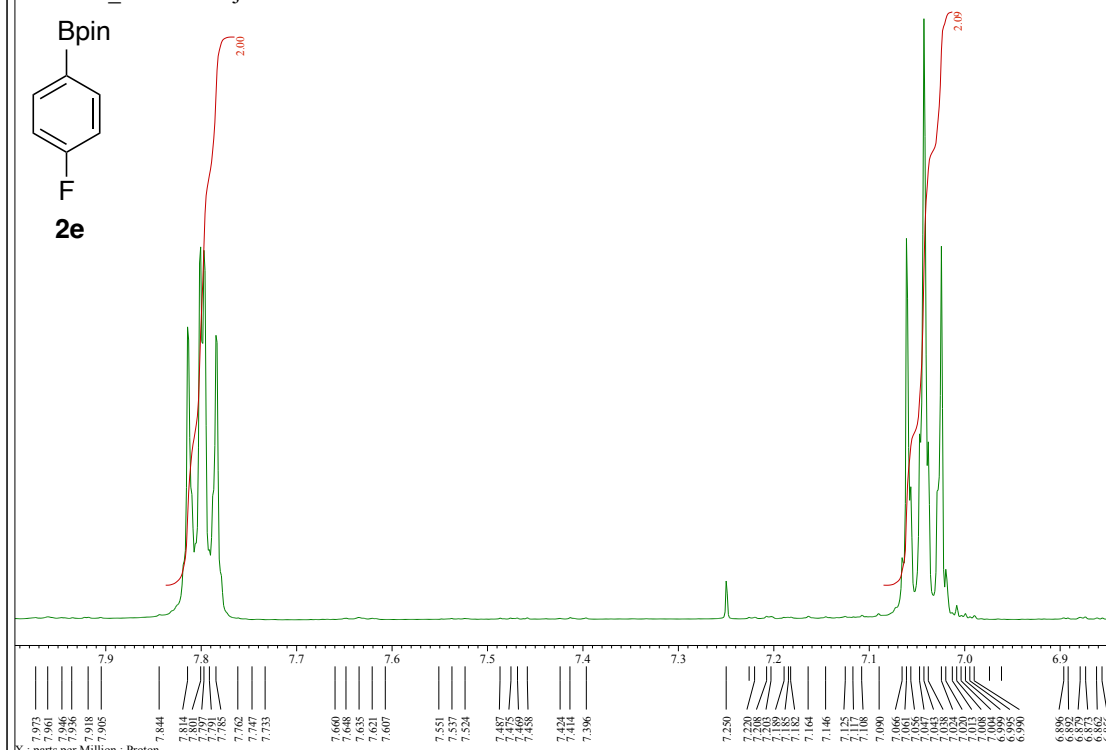
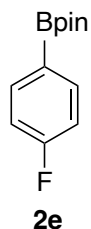
X : parts per Million : Proton

YF-868-2_Proton-1-2.jdf



Filename	= YF-868-2_Proton-1-2.j
Author	= delta
Experiment	= proton.jxp
Sample Id	= YF-868-2
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 21-MAR-2022 18:02:13
Revision_Time	= 22-MAR-2022 10:46:54
Comment	= single_pulse
Data Format	= 1D COMPLEX
Dim Size	= 13107
X_Domain	= Proton
Dim Title	= Proton
Dim Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ500R/S1
Field Strength	= 11.62926421[T] (500[M
X_Acq_Duration	= 1.76422912[s]
X_Domain	= 1H
X_Freq	= 495.13191398[MHz]
X_Offset	= 5[ppm]
X_Points	= 16384
X_Prescans	= 1
X_Resolution	= 0.5668198[Hz]
X_Sweep	= 9.28677563[kHz]
X_Sweep_Clippped	= 7.42942051[kHz]
Irr_Domain	= Proton
Irr_Freq	= 495.13191398[MHz]
Irr_Offset	= 5[ppm]
Tri_Domain	= Proton
Tri_Freq	= 495.13191398[MHz]
Tri_Offset	= 5[ppm]
Clipped	= FALSE
Scans	= 32
Total_Scans	= 32
Relaxation_Delay	= 5[s]
Recvr_Gain	= 36
Temp_Get	= 20.8[dC]
X_90_Width	= 9.47[us]
X_Acq_Time	= 1.76422912[s]
X_Angle	= 45[deg]
X_Atn	= 4.3[dB]
X_Pulse	= 4.735[us]
Irr_Mode	= Off
Tri_Mode	= Off
Dante_Loop	= 500
Dante_Presat	= FALSE
Decimation_Rate	= 0
Initial_Wait	= 1[s]
Phase	= {0, 90, 270, 180, 180
Presat_Time	= 5[s]
Presat_Time_Flag	= FALSE
Relaxation_Delay_Calc	= 0[s]
Relaxation_Delay_Temp	= 5[s]
Repetition_Time	= 6.76422912[s]

YF-813-2_Proton-1-2.jdf



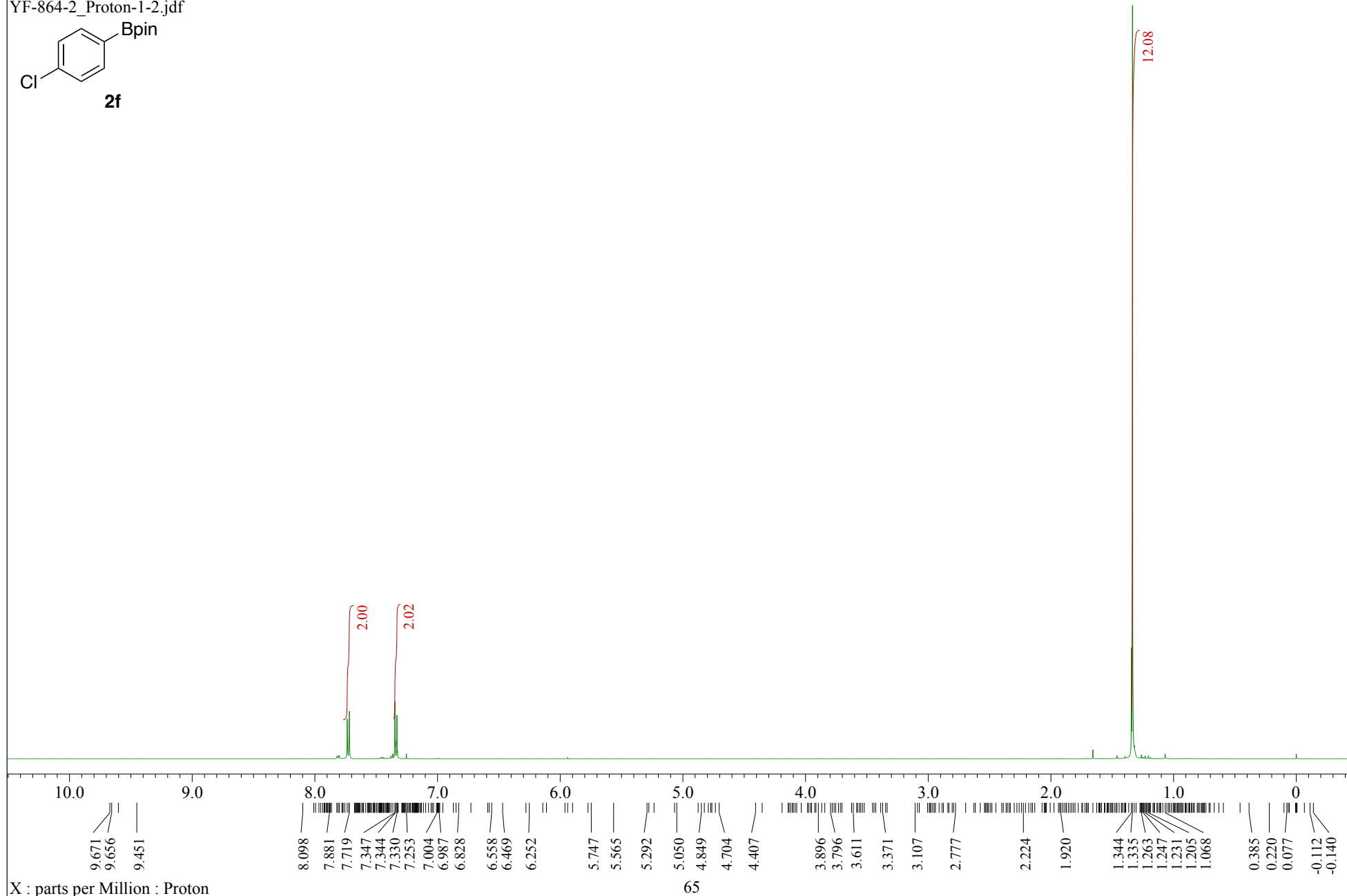
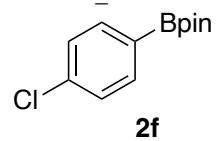
Filename = YF-813-2_Proton-1-2.j
 Author = delta
 Experiment = proton.jxp
 Sample Id = YF-813-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 21-FEB-2022 23:40:31
 Revision_Time = 22-FEB-2022 10:27:30

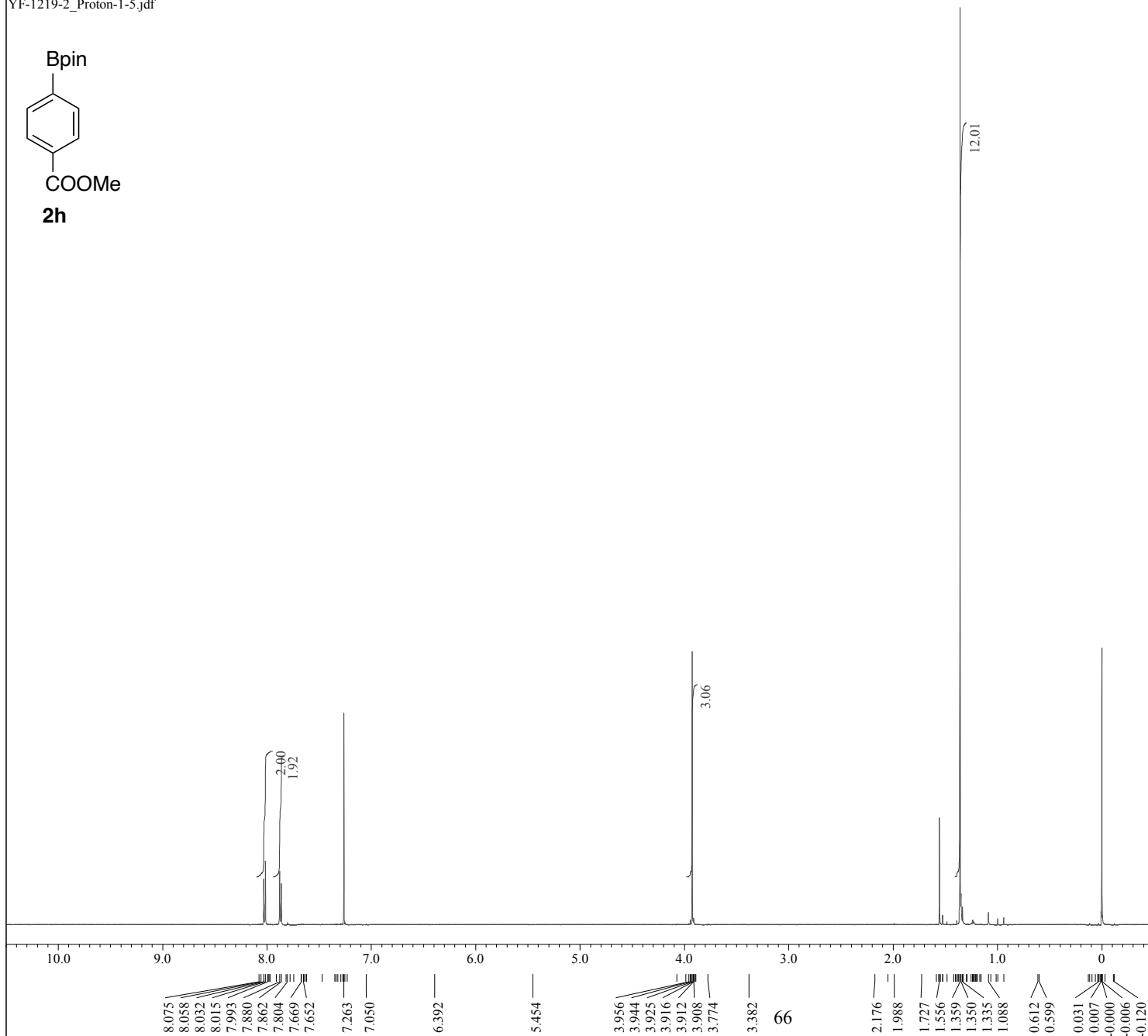
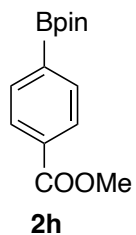
Comment = single_pulse
 Data Format = 1D REAL
 Dim Size = 13107
 X_Domain = Proton
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1

Field Strength = 11.62926421[T] (500[M]
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clippped = 7.42942051[kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 32
 Total_Scans = 32

Relaxation_Delay = 5[s]
 Recvr_Gain = 26
 Temp_Get = 22.1[dC]
 X_90_Width = 7.33[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 3.3[dB]
 X_Pulse = 3.665[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

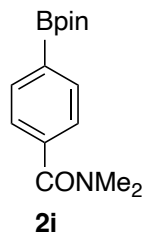
YF-864-2_Proton-1-2.jdf





Filename = YF-1219-2_Proton-1-5.
Author = delta
Experiment = proton.jxp
Sample_Id = YF-1219-2
Solvent = CHLOROFORM-D
Actual_Start_Time = 9-NOV-2022 14:05:18
Revision_Time = 9-NOV-2022 13:07:20
Comment = single_pulse
Data_Format = 1D REAL
Dim_Size = 13107
Dim_Title = Proton
Dim_Units = [ppm]
Dimensions = X
Spectrometer = JNM-ECZ500R/S1
Field_Strength = 11.62926421 [T] (500 [M])
X_Acq_Duration = 1.76422912 [s]
X_Domain = 18
X_Freq = 495.13191398 [MHz]
X_Offset = 5 [ppm]
X_Points = 16384
X_Prescans = 1
X_Resolution = 0.5668198 [Hz]
X_Sweep = 9.28677563 [kHz]
X_Sweep_Clippped = 7.42942051 [kHz]
Irr_Domain = Proton
Irr_Freq = 495.13191398 [MHz]
Irr_Offset = 5 [ppm]
Tri_Domain = Proton
Tri_Freq = 495.13191398 [MHz]
Tri_Offset = 5 [ppm]
Clipped = FALSE
Scans = 8
Total_Scans = 8
Relaxation_Delay = 5 [s]
Recvr_Gain = 56
Temp_Get = 20.1 [dC]
X_90_Width = 8.7 [us]
X_Acq_Time = 1.76422912 [s]
X_Angle = 45 [deg]
X_Atn = 4.3 [dB]
X_Pulse = 4.35 [us]
Irr_Mode = Off
Tri_Mode = Off
Dante_Loop = 500
Dante_Presat = FALSE
Decimation_Rate = 0
Initial_Wait = 1 [s]
Phase = {0, 90, 270, 180, 180}
Presat_Time = 5 [s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0 [s]
Relaxation_Delay_Temp = 5 [s]
Repetition_Time = 6.76422912 [s]

YF-1068-4_Proton-1-2.jdf

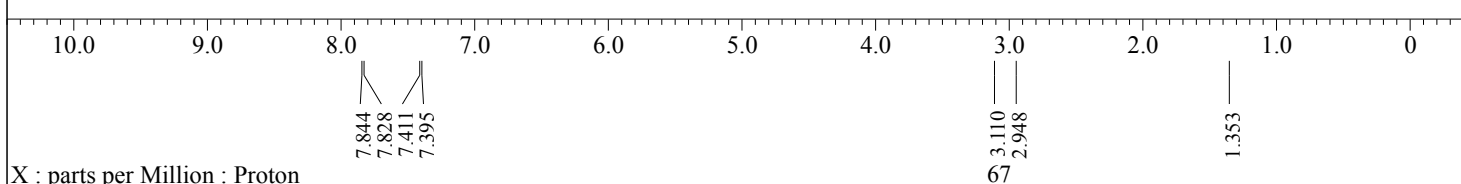


Filename = YF-1068-4_Proton-1-2.
 Author = delta
 Experiment = proton.jxp
 Sample Id = YF-1068-4
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 14-JUL-2022 00:35:08
 Revision_Time = 14-JUL-2022 14:55:05

Comment = single_pulse
 Data Format = 1D COMPLEX
 Dim Size = 13107
 X_Domain = Proton
 Dim Title = Proton
 Dim Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1

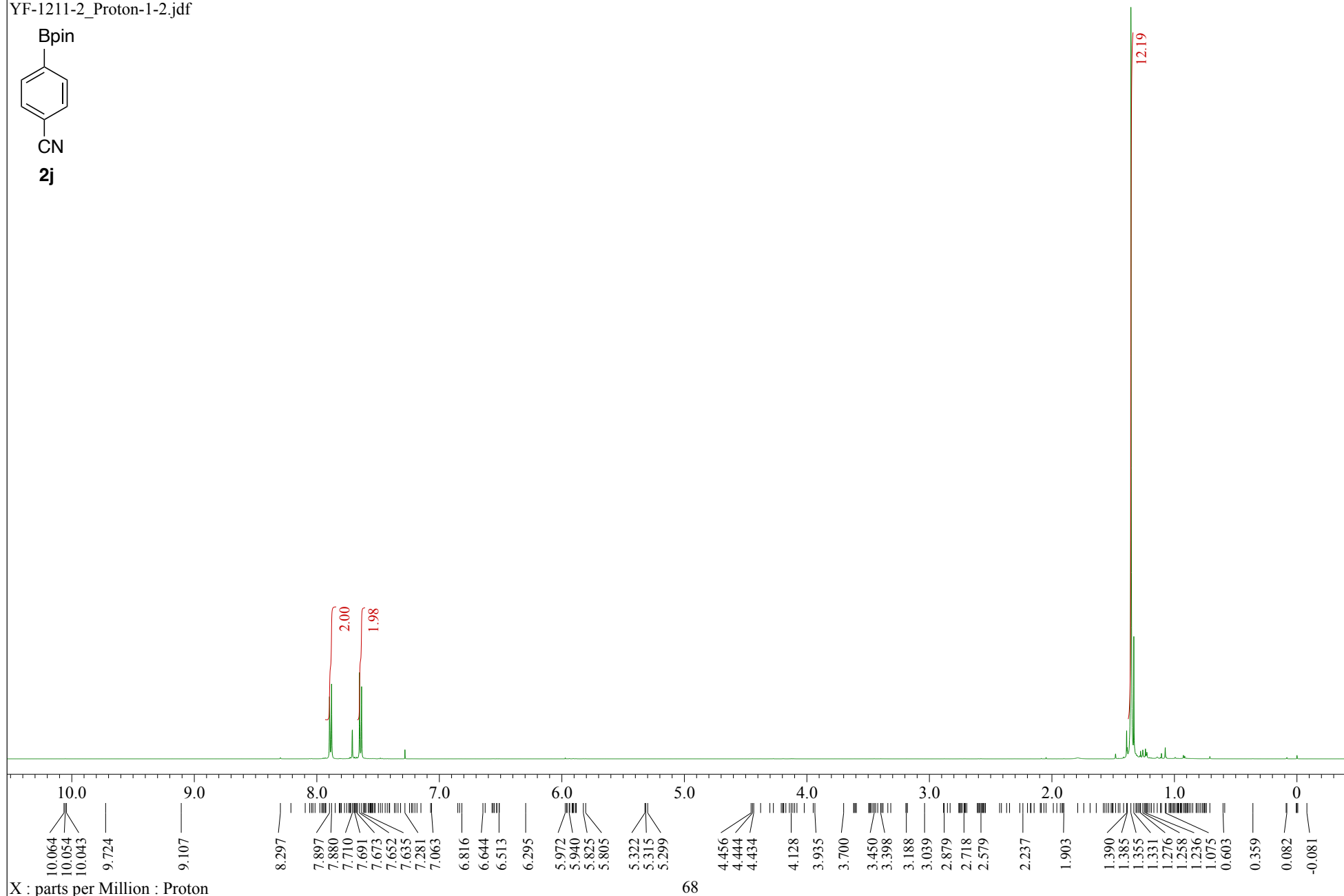
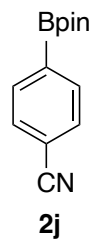
Field Strength = 11.62926421[T] (500[M]
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clippped = 7.42942051[kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 32
 Total_Scans = 32

Relaxation_Delay = 10[s]
 Recvr_Gain = 36
 Temp_Get = 18.9[dC]
 X_90_Width = 9.49[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[dB]
 X_Pulse = 4.745[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 1000
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 10[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 10[s]
 Repetition_Time = 11.76422912[s]



X : parts per Million : Proton

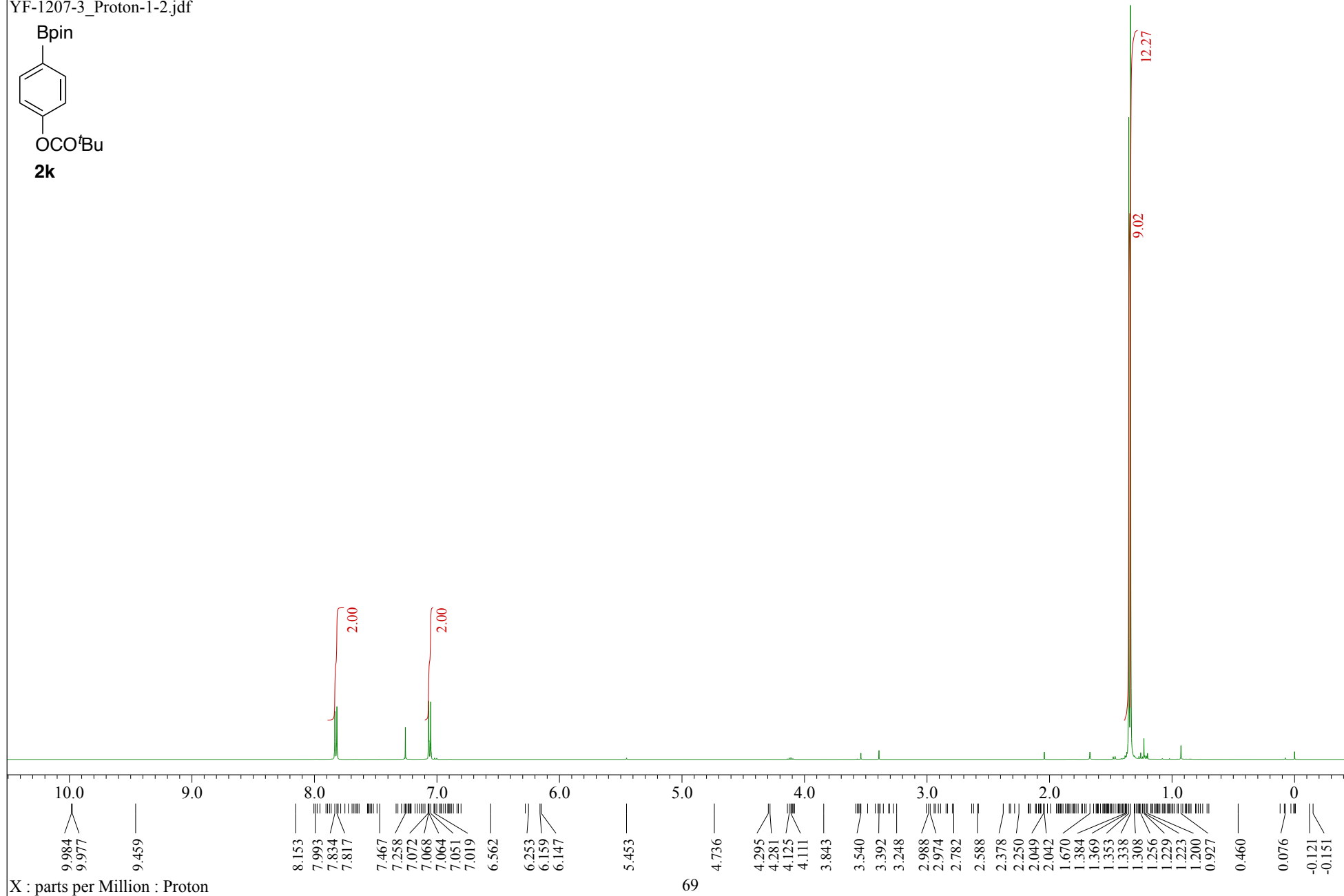
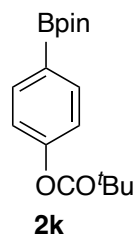
YF-1211-2_Proton-1-2.jdf



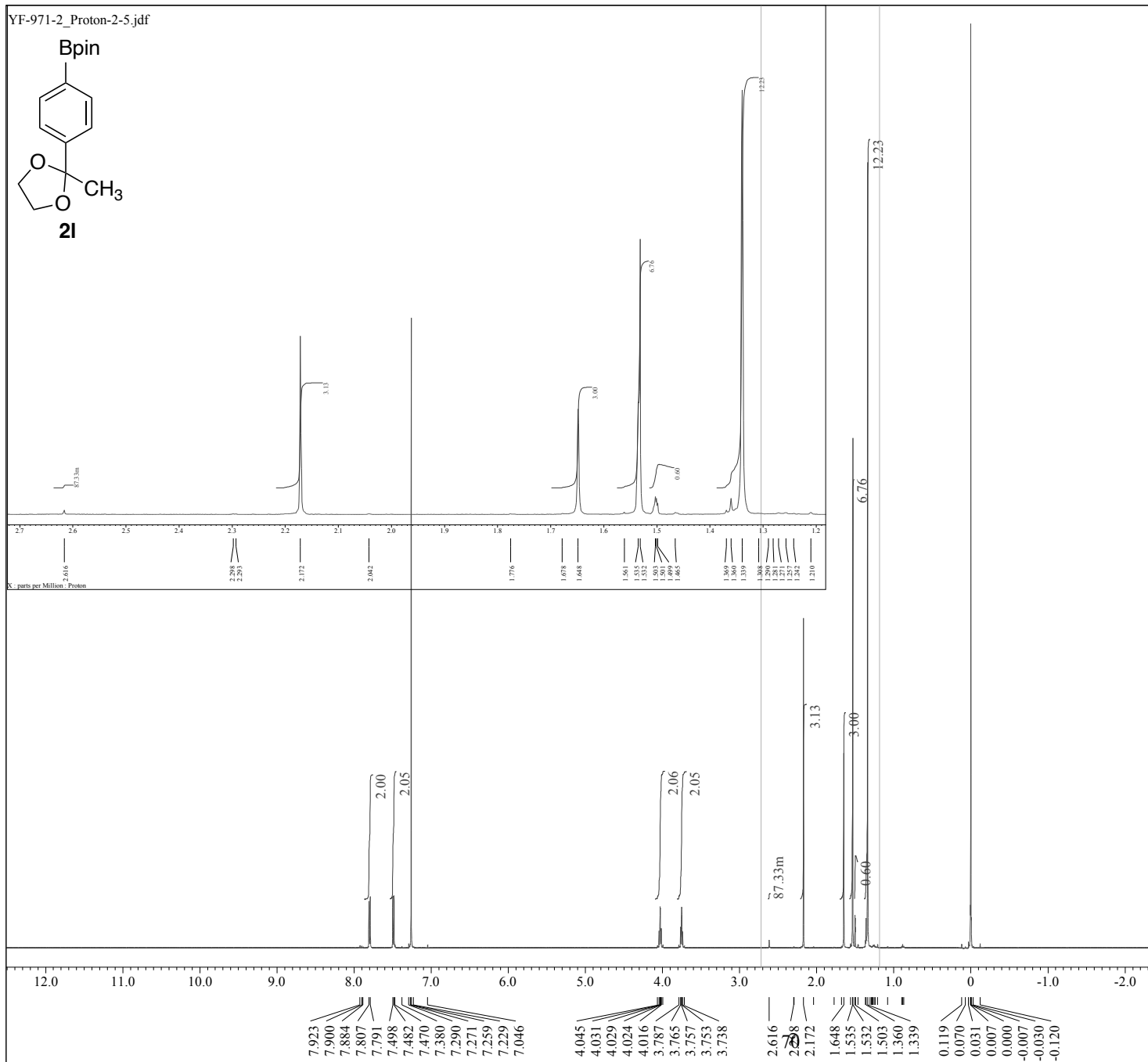
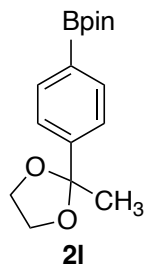
X : parts per Million : Proton

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YF-1207-3_Proton-1-2.jdf



YF-971-2_Proton-2-5.jdf



```

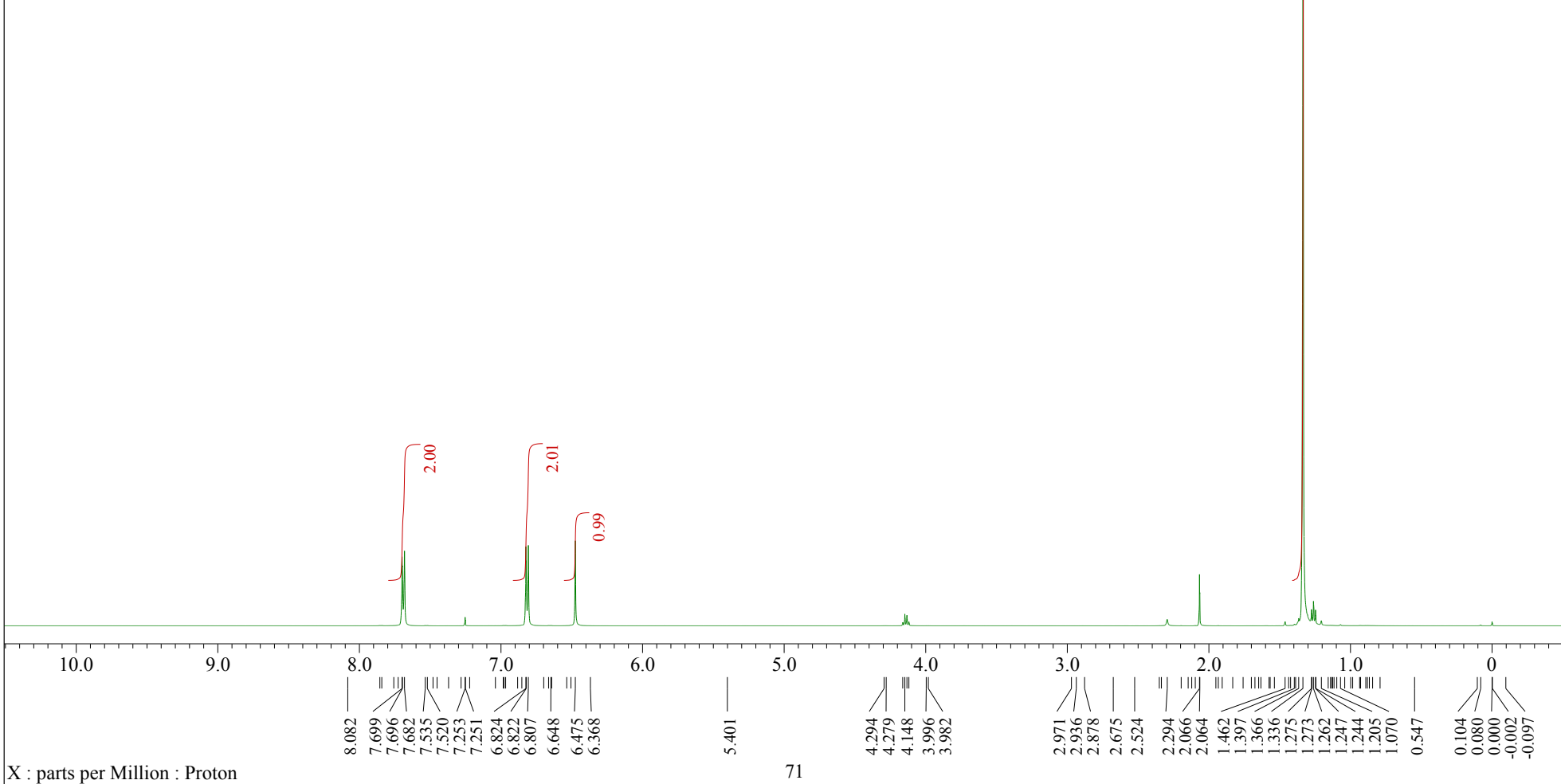
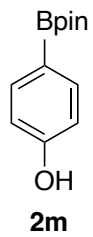
Filename      = YF-971-2_Proton-2-5.j
Author       = delta
Experiment    = proton.jxp
Sample_Id     = YF-971-2
Solvent       = CHLOROFORM-D
Actual_Start_Time = 3-JUN-2022 09:08:41
Revision_Time  = 3-JUN-2022 09:10:45

Comment      = single_pulse
Data_Format   = 1D_REAL
Dim_Size      = 13107
Dim_Title     = Proton
Dim_Units     = [ppm]
Dimensions    = X
Spectrometer  = JNM-ECZ500R/S1

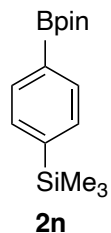
Field_Strength = 11.62926421 [T] (500[M]
X_Acq_Duration = 1.76422912[s]
X_Domain       = 1H
X_Freq         = 495.13191398[MHz]
X_Offset       = 5[ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198[Hz]
X_Sweep        = 9.28677563[kHz]
X_Sweep_Clipped = 7.42942051[kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398[MHz]
Irr_Offset     = 5[ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398[MHz]
Tri_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 32
Total_Scans    = 32

Relaxation_Delay = 10[s]
Recvr_Gain       = 66
Temp_Get         = 26.7[dc]
X_90_Width       = 7.5[us]
X_Acq_Time       = 1.76422912[s]
X_Angle          = 45[deg]
X_Atn            = 3.3[db]
X_Pulse          = 3.75[us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 1000
Dante_Preset     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1[s]
Phase            = {0, 90, 270, 180, 180}
Preset_Time      = 10[s]
Preset_Time_Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 10[s]
Repetition_Time  = 11.76422912[s]
  
```

YF-1226-2_Proton-1-2.jdf

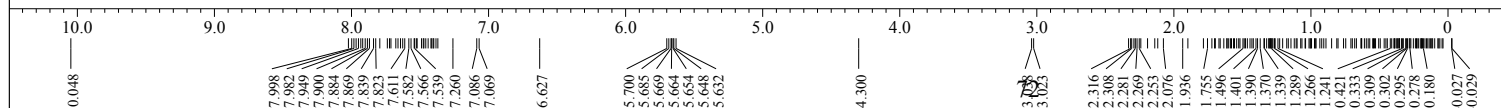


YF-1190-2_Proton-1-2.jdf



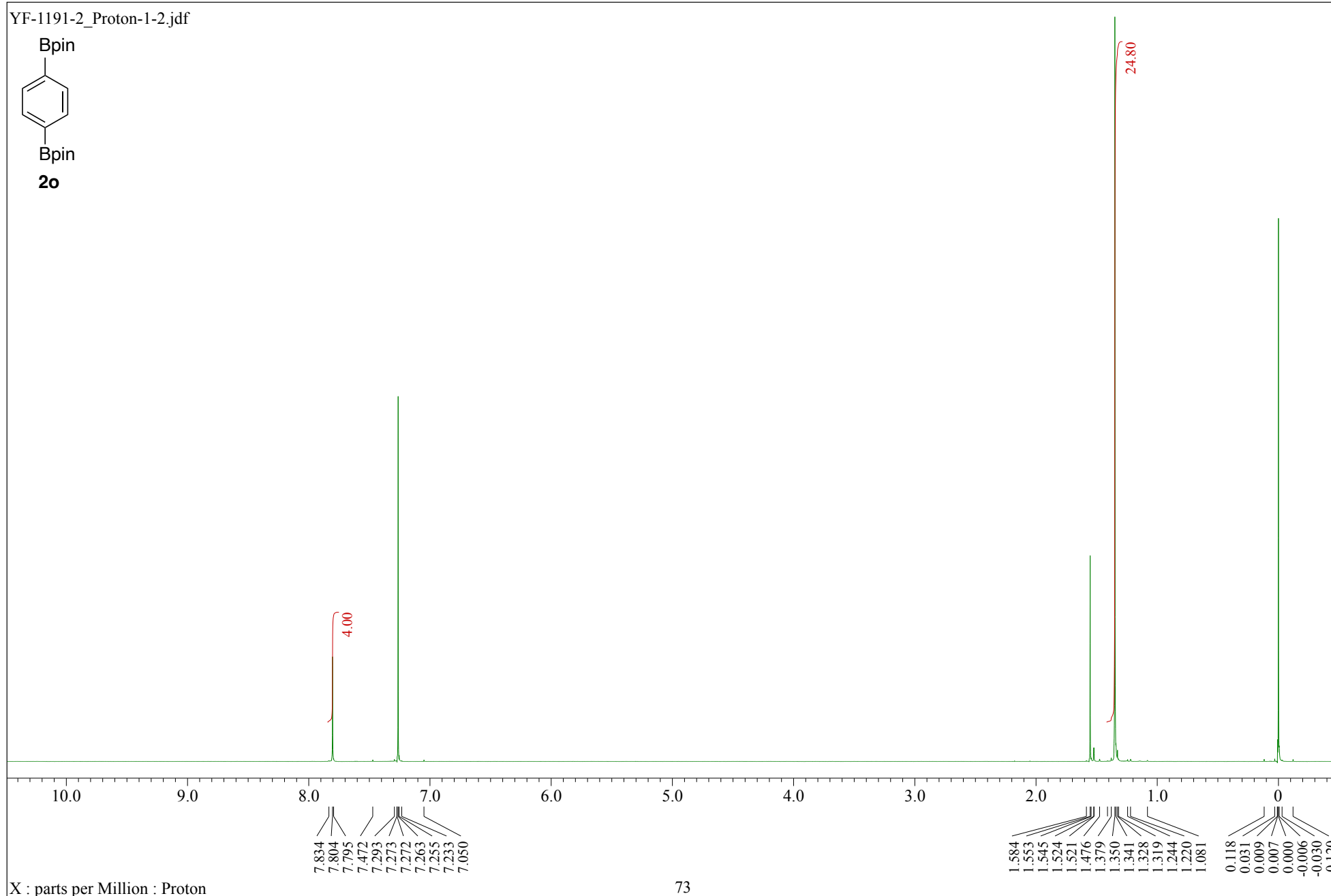
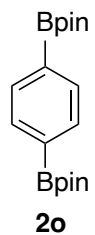
----- PROCESSING PARAMETERS -----
 sexp(0.2[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1, TRUE)
 fft(1, TRUE, TRUE)
 machinephase
 ppm
 base_correct(Akima, 5, 0, FALSE, 3, None, FA
 以下に由来 : YF-1190-2_Proton-1-1.jdf

Filename = YF-1190-2_Proton-1-2.
 Author = delta
 Experiment = proton.jpg
 Sample_Id = YF-1190-2
 Solvent = CHLOROFORM-D
 Actual_Start_Time = 28-OCT-2022 20:25:39
 RevisiOn_Time = 29-OCT-2022 15:11:32
 Comment = single_pulse
 Data_Format = 1D REAL
 Dim_Size = 13107
 X_Domain = Proton
 Dim_Title = Proton
 Dim_Units = [ppm]
 Dimensions = X
 Spectrometer = JNM-ECZ500R/S1
 Field_Strength = 11.62926421[T] (500[M
 X_Acq_Duration = 1.76422912[s]
 X_Domain = 1H
 X_Freq = 495.13191398[MHz]
 X_Offset = 5[ppm]
 X_Points = 16384
 X_Prescans = 1
 X_Resolution = 0.5668198[Hz]
 X_Sweep = 9.28677563[kHz]
 X_Sweep_Clipped = 7.42942051[kHz]
 Irr_Domain = Proton
 Irr_Freq = 495.13191398[MHz]
 Irr_Offset = 5[ppm]
 Tri_Domain = Proton
 Tri_Freq = 495.13191398[MHz]
 Tri_Offset = 5[ppm]
 Clipped = FALSE
 Scans = 8
 Total_Scans = 8
 Relaxation_Delay = 5[s]
 Recvr_Gain = 26
 Temp_Get = 19.9[dC]
 X_90_Width = 8.7[us]
 X_Acq_Time = 1.76422912[s]
 X_Angle = 45[deg]
 X_Atn = 4.3[dB]
 X_Pulse = 4.35[us]
 Irr_Mode = Off
 Tri_Mode = Off
 Dante_Loop = 500
 Dante_Presat = FALSE
 Decimation_Rate = 0
 Initial_Wait = 1[s]
 Phase = {0, 90, 270, 180, 180
 Presat_Time = 5[s]
 Presat_Time_Flag = FALSE
 Relaxation_Delay_Calc = 0[s]
 Relaxation_Delay_Temp = 5[s]
 Repetition_Time = 6.76422912[s]

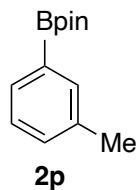


X : parts per Million : Proton

YF-1191-2_Proton-1-2.jdf

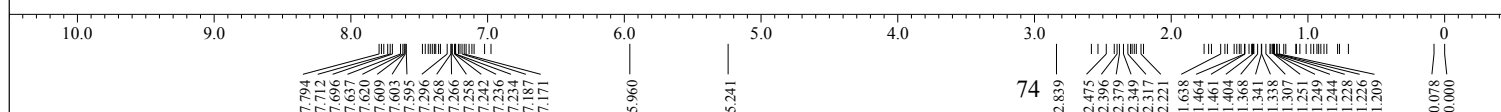


YF-964-2_Proton-1-2.jdf



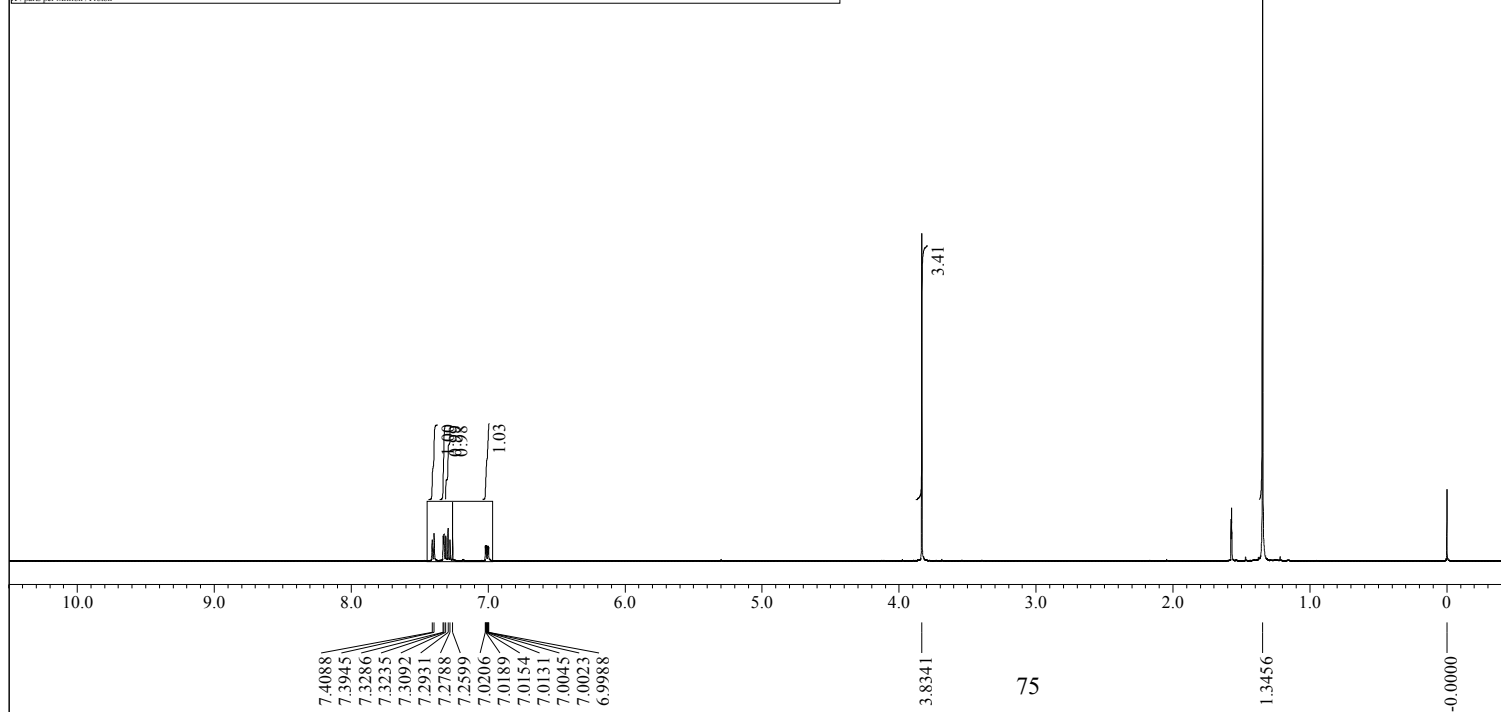
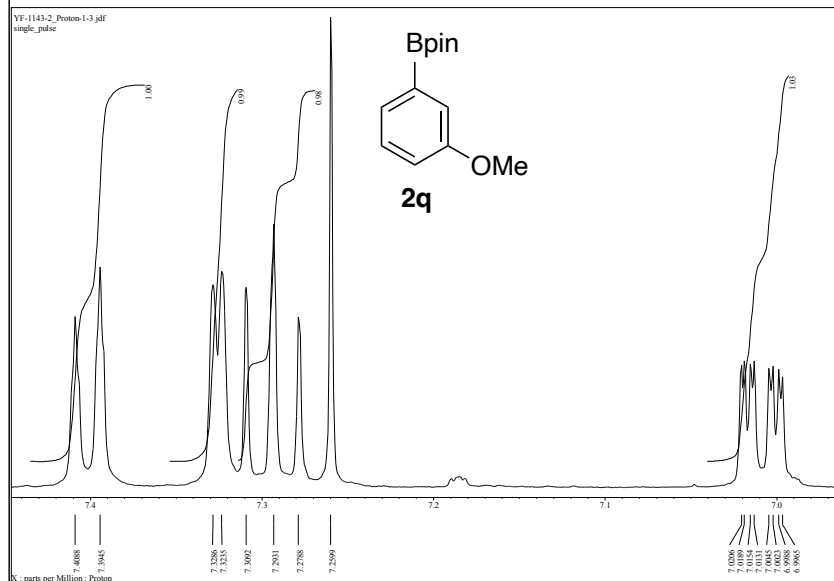
----- PROCESSING PARAMETERS -----
sexp(0.2[Hz], 0.0[s])
trapezoid(0[%], 0[%], 80[%], 100[%])
zerofill(1, TRUE)
fft(1, TRUE, TRUE)
machinephase
ppm
base_correct(Akima, 5, 0, FALSE, 3, None, FA
以下に由来: YF-964-2_Proton-1-1.jdf

Filename = YF-964-2_Proton-1-2.j
Author = delta
Experiment = proton.jpg
Sample_Id = YF-964-2
Solvent = CHLOROFORM-D
Actual_Start_Time = 2-JUN-2022 04:15:45
Revision_Time = 2-JUN-2022 11:35:10
Comment = single_pulse
Data Format = 1D REAL
Dim_Size = 13107
X_Domain = Proton
Dim Title = Proton
Dim Units = [ppm]
Dimensions = X
Spectrometer = JNM-ECP500R/S1
Field_Strength = 11.62926421[T] (500[M
X_Acq_Duration = 1.76422912[s]
X_Domain = 1H
X_Freq = 495.13191398[MHz]
X_Offset = 5[ppm]
X Points = 16384
X_Prescans = 1
X Resolution = 0.5668198[Hz]
X_Sweep = 9.28677563[kHz]
X_Sweep_Clippped = 7.42942051[kHz]
Irr_Domain = Proton
Irr_Freq = 495.13191398[MHz]
Irr_Offset = 5[ppm]
Tri_Domain = Proton
Tri_Freq = 495.13191398[MHz]
Tri_Offset = 5[ppm]
Clipped = FALSE
Scans = 32
Total_Scans = 32
Relaxation_Delay = 10[s]
Recvr Gain = 36
Temp Get = 26.7[dc]
X_90_Width = 7.5[us]
X_Acq_Time = 1.76422912[s]
X_Angle = 45[deg]
X_Atn = 3.3[dB]
X_Pulse = 3.75[us]
Irr_Mode = Off
Tri_Mode = Off
Dante_Loop = 1000
Dante_Presat = FALSE
Decimation_Rate = 0
Initial_Wait = 1[s]
Phase = {0, 90, 270, 180, 180
Presat Time = 10[s]
Presat Time Flag = FALSE
Relaxation_Delay_Calc = 0[s]
Relaxation_Delay_Temp = 10[s]
Repetition_Time = 11.76422912[s]



X : parts per Million : Proton

YF-1143-2_Proton-1-3.jdf
single_pulse



---- PROCESSING PARAMETERS ----
dc balance(0, FALSE)
sexp(0.2[Hz], 0.0[s])
trapezoid(0[%], 0[%], 80[%], 100[%])
zerofill(1)
fft(1, TRUE, TRUE)
machinphase
ppm

以下に由来: YF-1143-2_Proton-1-1.jdf

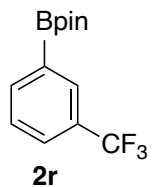
Filename = YF-1143-2_Proton-1-3.jdf
Author = takaya
Experiment = proton.jpg
Sample_Id = YF-1143-2
Solvent = CHLOROFORM-D
Creation_Time = 7-OCT-2022 20:36:08
Revision_Time = 7-OCT-2022 21:07:19
Current_Time = 7-OCT-2022 21:08:04

Comment = single_pulse
Data_Format = 1D COMPLEX
Dim_Size = 26214
Dim_Title = Proton
Dim_Units = [ppm]
Dimensions = X
Site = JNM-ECK500
Spectrometer = DELTA2_NMR

Field_Strength = 11.7473579[T] (500[MHz])
X_Acq_Duration = 3.49175808[s]
X_Domain = 1H
X_Freq = 500.15991521[MHz]
X_Offset = 5.0[ppm]
X_Points = 32768
X_Fscans = 1
X_Resolution = 0.28638868[Hz]
X_Sweep = 9.38438438[kHz]
X_Sweep_Clipped = 7.50750751[kHz]
Irr_Domain = Proton
Irr_Freq = 500.15991521[MHz]
Irr_Offset = 5.0[ppm]
Tri_Domain = Proton
Tri_Freq = 500.15991521[MHz]
Tri_Offset = 5.0[ppm]
Clipped = FALSE
Scans = 8
Total_Scans = 8

Relaxation_Delay = 2[s]
Recvr_Gain = 46
Temp_Get = 21.9[dc]
X_90_Width = 12.4[us]
X_Acq_Time = 3.49175808[s]
X_Angle = 45[deg]
X_Atn = 4.5[db]
X_Pulse = 6.2[us]
Irr_Mode = Off
Tri_Mode = Off
Dante_Presat = FALSE
Initial_Wait = 1[s]
Repetition_Time = 5.49175808[s]

YF-1163-2_Proton-1-6.jdf



0.99
0.99
1.01
1.00

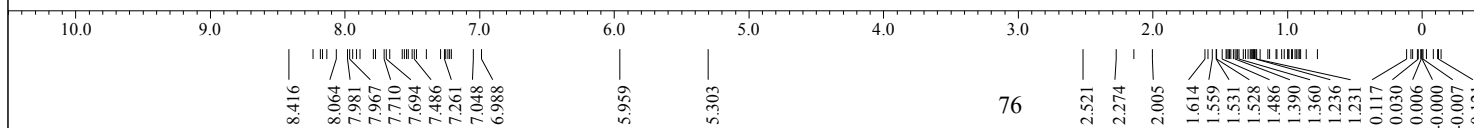


Filename = YF-1163-2_Proton-1-6.
Author = delta
Experiment = proton.jxp
Sample_Id = YF-1163-2
Solvent = CHLOROFORM-D
Actual_Start_Time = 14-OCT-2022 21:52:35
Revision_Time = 14-OCT-2022 20:56:09

Comment = single_pulse
Data_Format = 1D_REAL
Dim_Size = 13107
Dim_Title = Proton
Dim_Units = [ppm]
Dimensions = X
Spectrometer = JNM-ECZ500R/S1

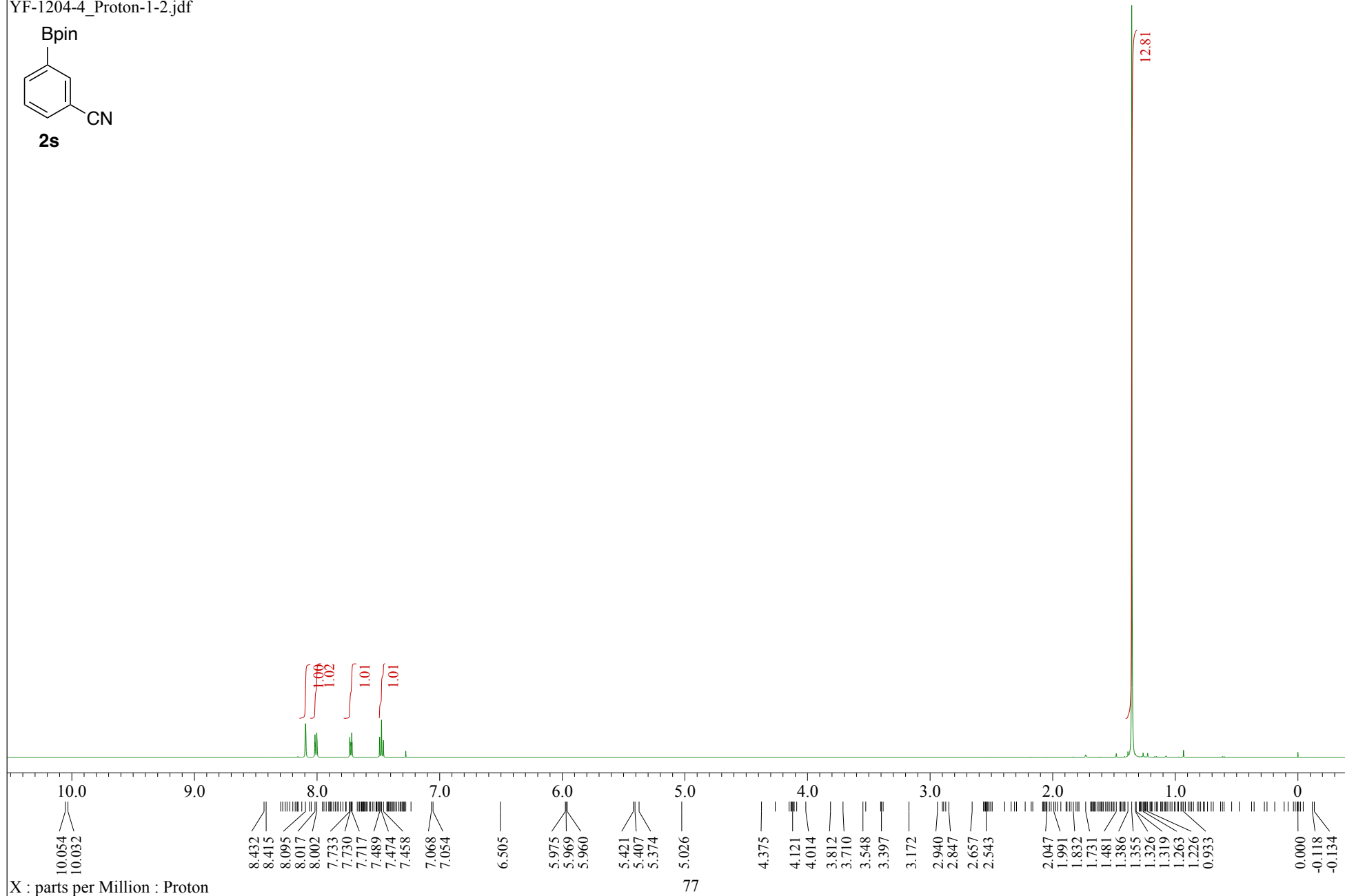
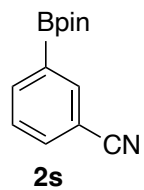
Field_Strength = 11.62926421 [T] (500 [M])
X_Acq_Duration = 1.76422912 [s]
X_Domain = 18
X_Freq = 495.13191398 [MHz]
X_Offset = 5 [ppm]
X_Points = 16384
X_Prescans = 1
X_Resolution = 0.5668198 [Hz]
X_Sweep = 9.28677563 [kHz]
X_Sweep_Clipped = 7.42942051 [kHz]
Irr_Domain = Proton
Irr_Freq = 495.13191398 [MHz]
Irr_Offset = 5 [ppm]
Tri_Domain = Proton
Tri_Freq = 495.13191398 [MHz]
Tri_Offset = 5 [ppm]
Clipped = FALSE
Scans = 8
Total_Scans = 8

Relaxation_Delay = 5 [s]
Recvr_Gain = 56
Temp_Get = 20 [dc]
X_90_Width = 8.7 [us]
X_Acq_Time = 1.76422912 [s]
X_Angle = 45 [deg]
X_Atn = 4.3 [dB]
X_Pulse = 4.35 [us]
Irr_Mode = Off
Tri_Mode = Off
Dante_Loop = 500
Dante_Presat = FALSE
Decimation_Rate = 0
Initial_Wait = 1 [s]
Phase = {0, 90, 270, 180, 180}
Presat_Time = 5 [s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0 [s]
Relaxation_Delay_Temp = 5 [s]
Repetition_Time = 6.76422912 [s]

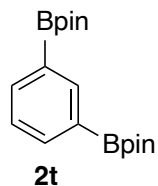


X : parts per Million : Proton

YF-1204-4_Proton-1-2.jdf



YF-1180-4_Proton-1-4.jdf



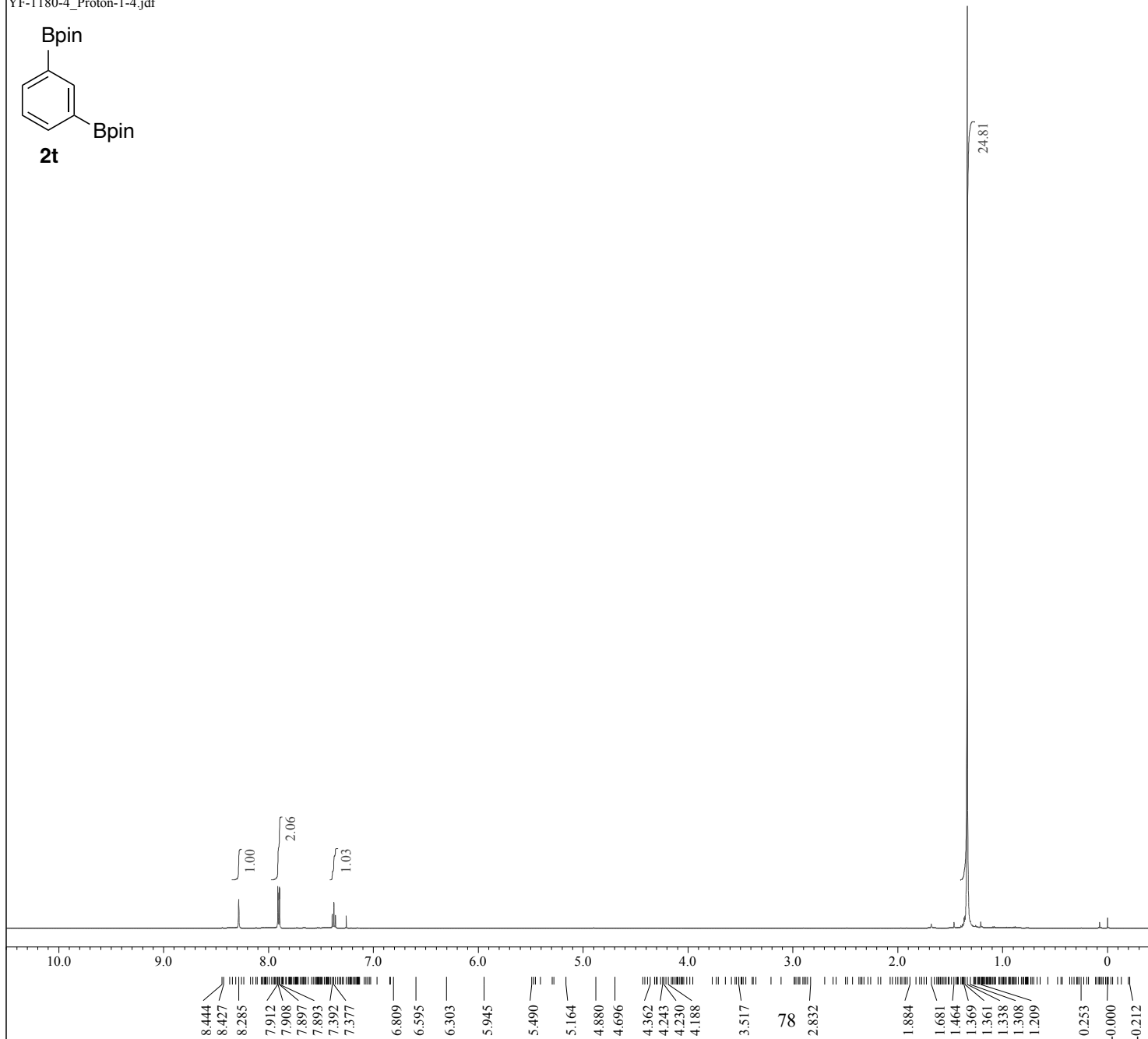
```

Filename      = YF-1180-4_Proton-1-4.
Author       = delta
Experiment    = proton.jxp
Sample_Id    = YF-1180-4
Solvent       = CHLOROFORM-D
Actual_Start_Time = 26-OCT-2022 21:44:12
Revision_Time  = 26-OCT-2022 20:46:45

Comment      = single_pulse
Data_Format   = 1D_REAL
Dim_Size      = 13107
Dim_Title     = Proton
Dim_Units     = [ppm]
Dimensions    = X
Spectrometer  = JNM-ECZ500R/S1

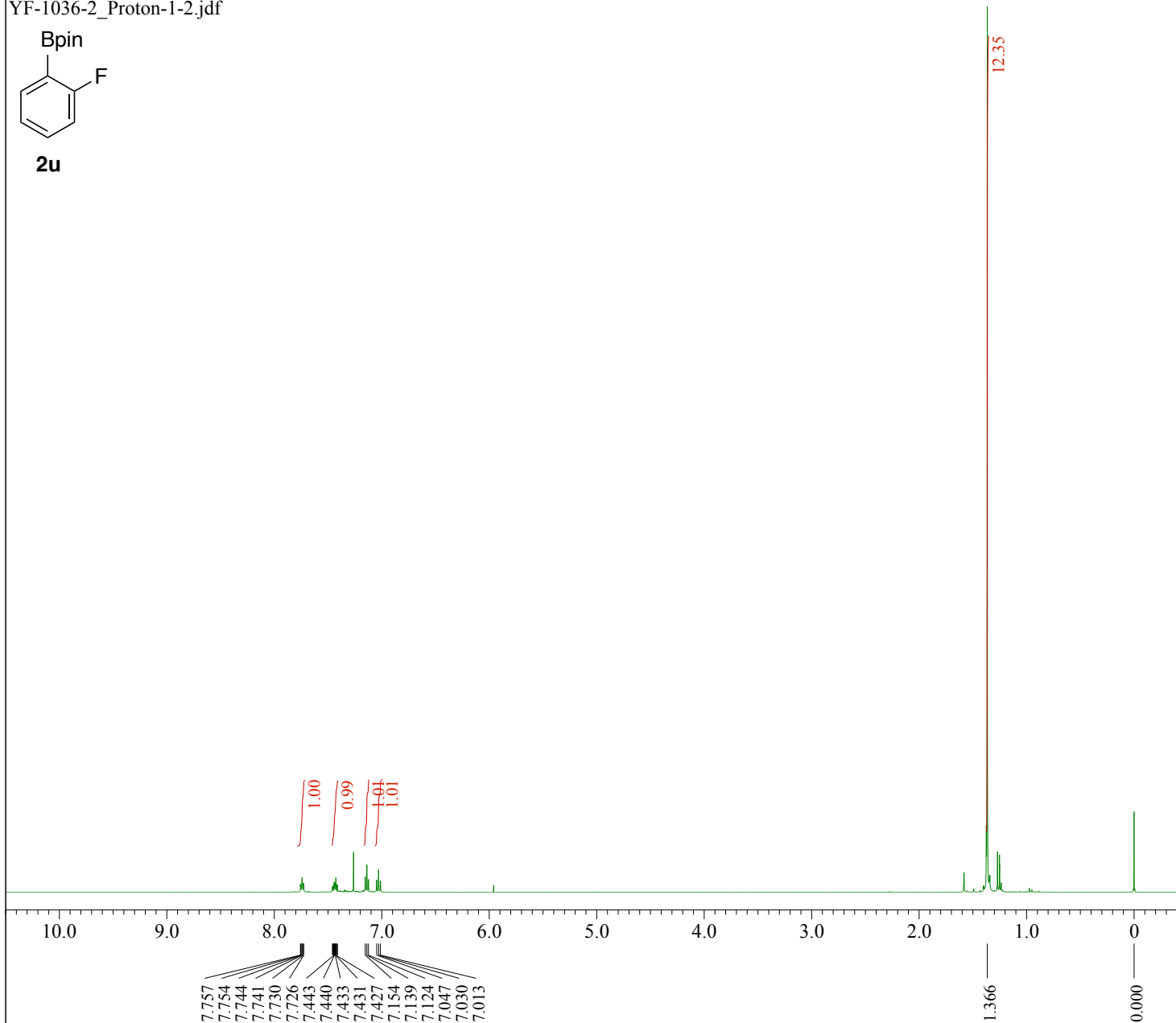
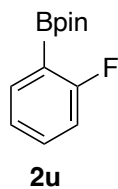
Field_Strength = 11.62926421 [T] (500 [M]
X_Acq_Duration = 1.76422912 [s]
X_Domain       = 1H
X_Freq         = 495.13191398 [MHz]
X_Offset       = 5 [ppm]
X_Points       = 16384
X_Prescans     = 1
X_Resolution   = 0.5668198 [Hz]
X_Sweep        = 9.28677563 [kHz]
X_Sweep_Clipped = 7.42942051 [kHz]
Irr_Domain     = Proton
Irr_Freq       = 495.13191398 [MHz]
Irr_Offset     = 5 [ppm]
Tri_Domain     = Proton
Tri_Freq       = 495.13191398 [MHz]
Tri_Offset     = 5 [ppm]
Clipped        = FALSE
Scans          = 8
Total_Scans    = 8

Relaxation_Delay = 5 [s]
Recvr_Gain       = 26
Temp_Get         = 19.8 [dC]
X_90_Width       = 8.7 [us]
X_Acq_Time       = 1.76422912 [s]
X_Angle          = 45 [deg]
X_Atn            = 4.3 [dB]
X_Pulse          = 4.35 [us]
Irr_Mode         = Off
Tri_Mode         = Off
Dante_Loop       = 500
Dante_Presat     = FALSE
Decimation_Rate  = 0
Initial_Wait     = 1 [s]
Phase            = {0, 90, 270, 180, 180}
Presat_Time      = 5 [s]
Presat_Time_Flag = FALSE
Relaxation_Delay_Calc = 0 [s]
Relaxation_Delay_Temp = 5 [s]
Repetition_Time  = 6.76422912 [s]
  
```



X : parts per Million : Proton

YF-1036-2_Proton-1-2.jdf



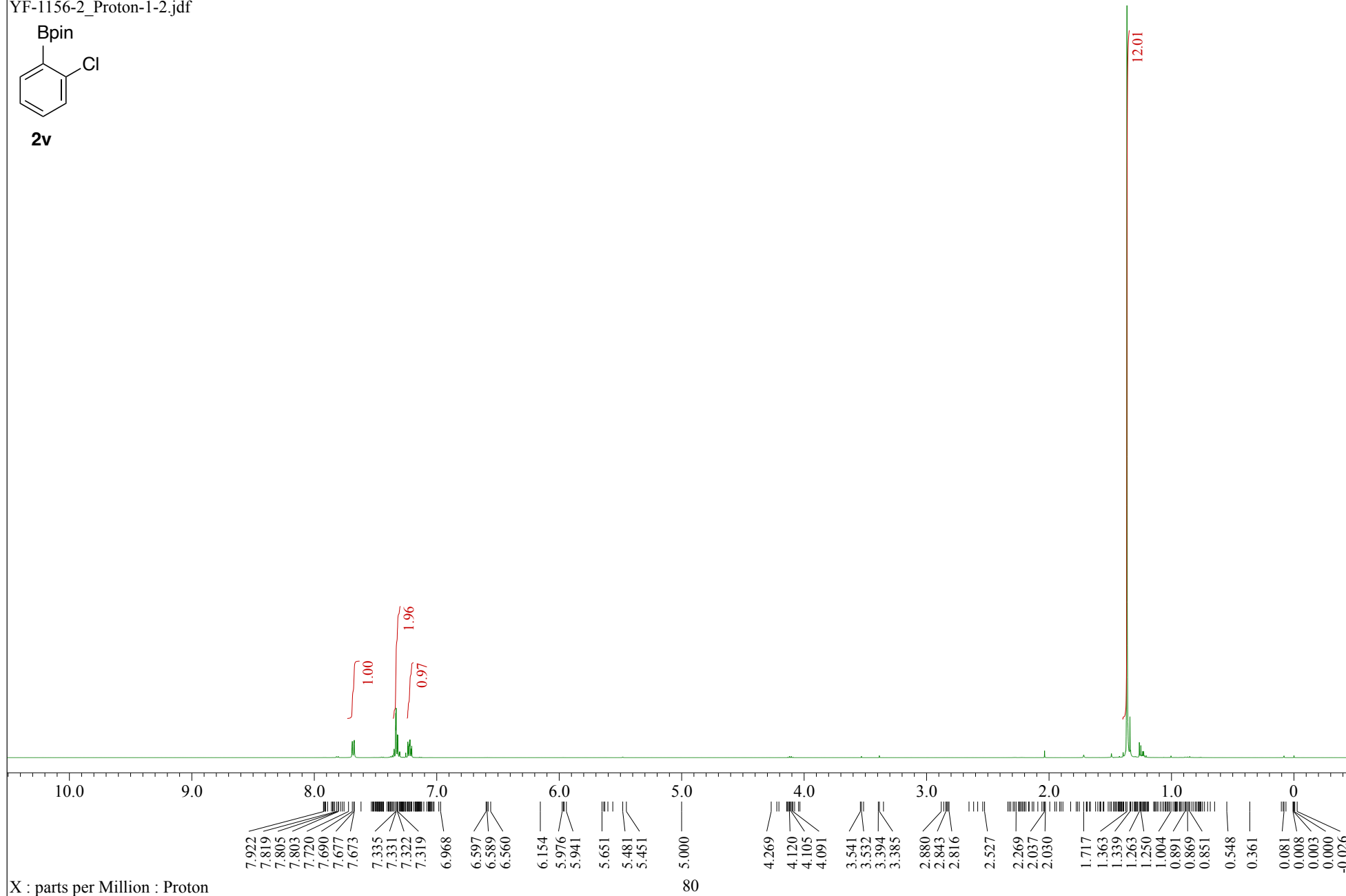
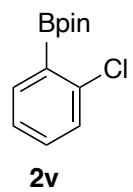
X : parts per Million : Proton

79

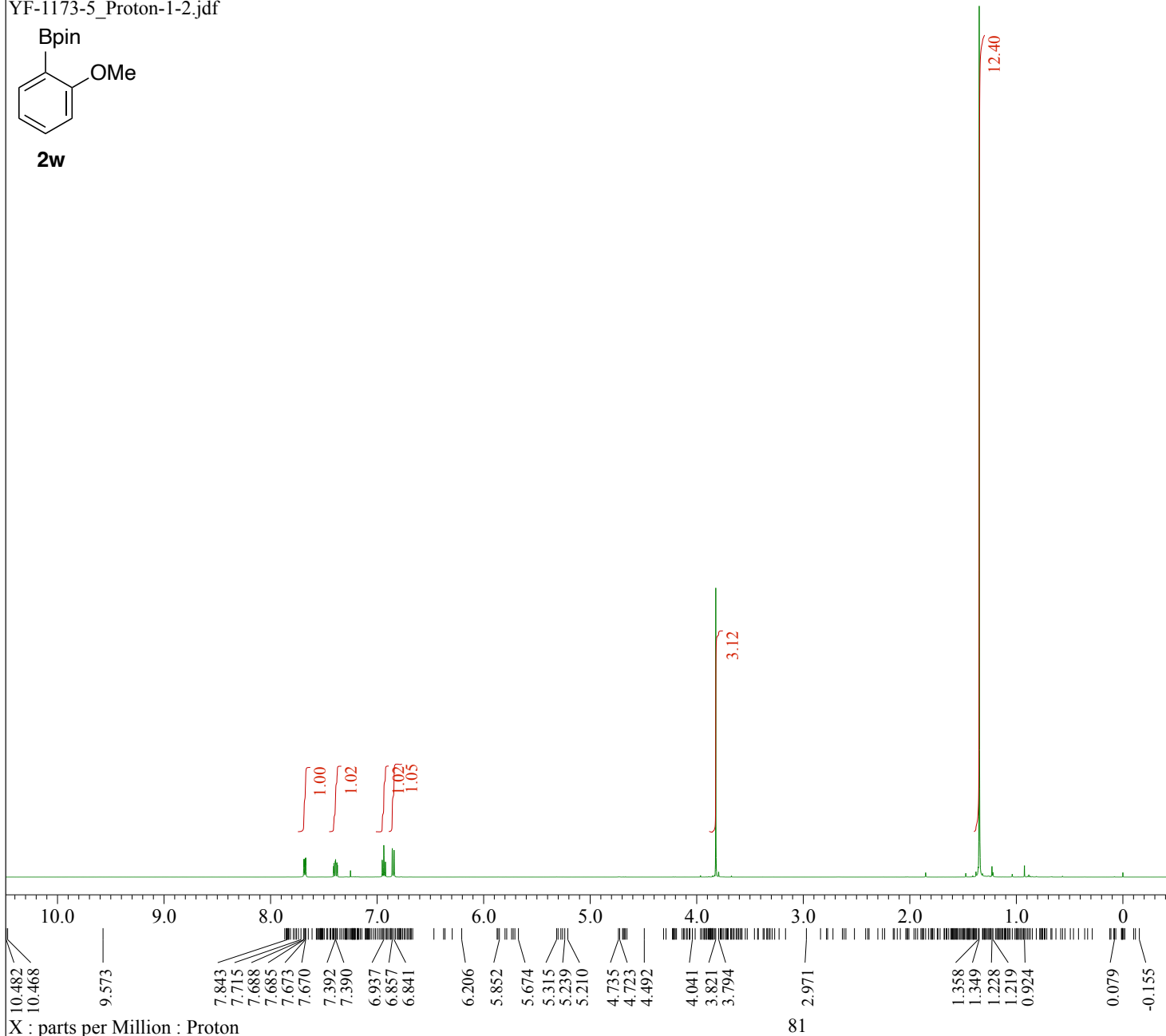
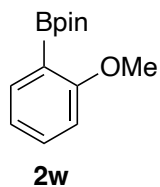


Filename	= YF-1036-2_Proton-1-2.
Author	= delta
Experiment	= proton.jxp
Sample_Id	= YF-1036-2
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 28-JUN-2022 19:23:37
Revision_Time	= 18-JUL-2022 19:35:21
Comment	= single_pulse
Data_Format	= 1D COMPLEX
Dim_Size	= 13107
X_Domain	= Proton
Dim_Title	= Proton
Dim_Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ500R/S1
Field_Strength	= 11.62926421[T] (500[M
X_Acq_Duration	= 1.76422912[s]
X_Domain	= 1H
X_Freq	= 495.13191398[MHz]
X_Offset	= 5[ppm]
X_Points	= 16384
X_Prescans	= 1
X_Resolution	= 0.5668198[Hz]
X_Sweep	= 9.28677563[kHz]
X_Sweep_Clippped	= 7.42942051[kHz]
Irr_Domain	= Proton
Irr_Freq	= 495.13191398[MHz]
Irr_Offset	= 5[ppm]
Tri_Domain	= Proton
Tri_Freq	= 495.13191398[MHz]
Tri_Offset	= 5[ppm]
Clipped	= FALSE
Scans	= 32
Total_Scans	= 32
Relaxation_Delay	= 5[s]
Recvr_Gain	= 46
Temp_Get	= 18.5[dC]
X_90_Width	= 9.49[us]
X_Acq_Time	= 1.76422912[s]
X_Angle	= 45[deg]
X_Atn	= 4.3[dB]
X_Pulse	= 4.745[us]
Irr_Mode	= Off
Tri_Mode	= Off
Dante_Loop	= 500
Dante_Presat	= FALSE
Decimation_Rate	= 0
Initial_Wait	= 1[s]
Phase	= {0, 90, 270, 180, 180
Presat_Time	= 5[s]
Presat_Time_Flag	= FALSE
Relaxation_Delay_Calc	= 0[s]
Relaxation_Delay_Temp	= 5[s]
Repetition_Time	= 6.76422912[s]

YF-1156-2_Proton-1-2.jdf



YF-1173-5_Proton-1-2.jdf



X : parts per Million : Proton

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Filename	= YF-1173-5_Proton-1-2.
Author	= delta
Experiment	= proton.jxp
Sample_Id	= YF-1173-5
Solvent	= CHLOROFORM-D
Actual_Start_Time	= 20-OCT-2022 15:53:42
Revision_Time	= 20-OCT-2022 19:53:36
Comment	= single_pulse
Data_Format	= 1D REAL
Dim_Size	= 13107
X_Domain	= Proton
Dim_Title	= Proton
Dim_Units	= [ppm]
Dimensions	= X
Spectrometer	= JNM-ECZ500R/S1
Field_Strength	= 11.62926421[T] (500[M
X_Acq_Duration	= 1.76422912[s]
X_Domain	= 1H
X_Freq	= 495.13191398[MHz]
X_Offset	= 5[ppm]
X_Points	= 16384
X_Prescans	= 1
X_Resolution	= 0.5668198[Hz]
X_Sweep	= 9.28677563[kHz]
X_Sweep_Clippped	= 7.42942051[kHz]
Irr_Domain	= Proton
Irr_Freq	= 495.13191398[MHz]
Irr_Offset	= 5[ppm]
Tri_Domain	= Proton
Tri_Freq	= 495.13191398[MHz]
Tri_Offset	= 5[ppm]
Clipped	= FALSE
Scans	= 8
Total_Scans	= 8
Relaxation_Delay	= 5[s]
Recvr_Gain	= 36
Temp_Get	= 20.5[dC]
X_90_Width	= 8.7[us]
X_Acq_Time	= 1.76422912[s]
X_Angle	= 45[deg]
X_Atn	= 4.3[dB]
X_Pulse	= 4.35[us]
Irr_Mode	= Off
Tri_Mode	= Off
Dante_Loop	= 500
Dante_Presat	= FALSE
Decimation_Rate	= 0
Initial_Wait	= 1[s]
Phase	= {0, 90, 270, 180, 180
Presat_Time	= 5[s]
Presat_Time_Flag	= FALSE
Relaxation_Delay_Calc	= 0[s]
Relaxation_Delay_Temp	= 5[s]
Repetition_Time	= 6.76422912[s]