

Supplementary data

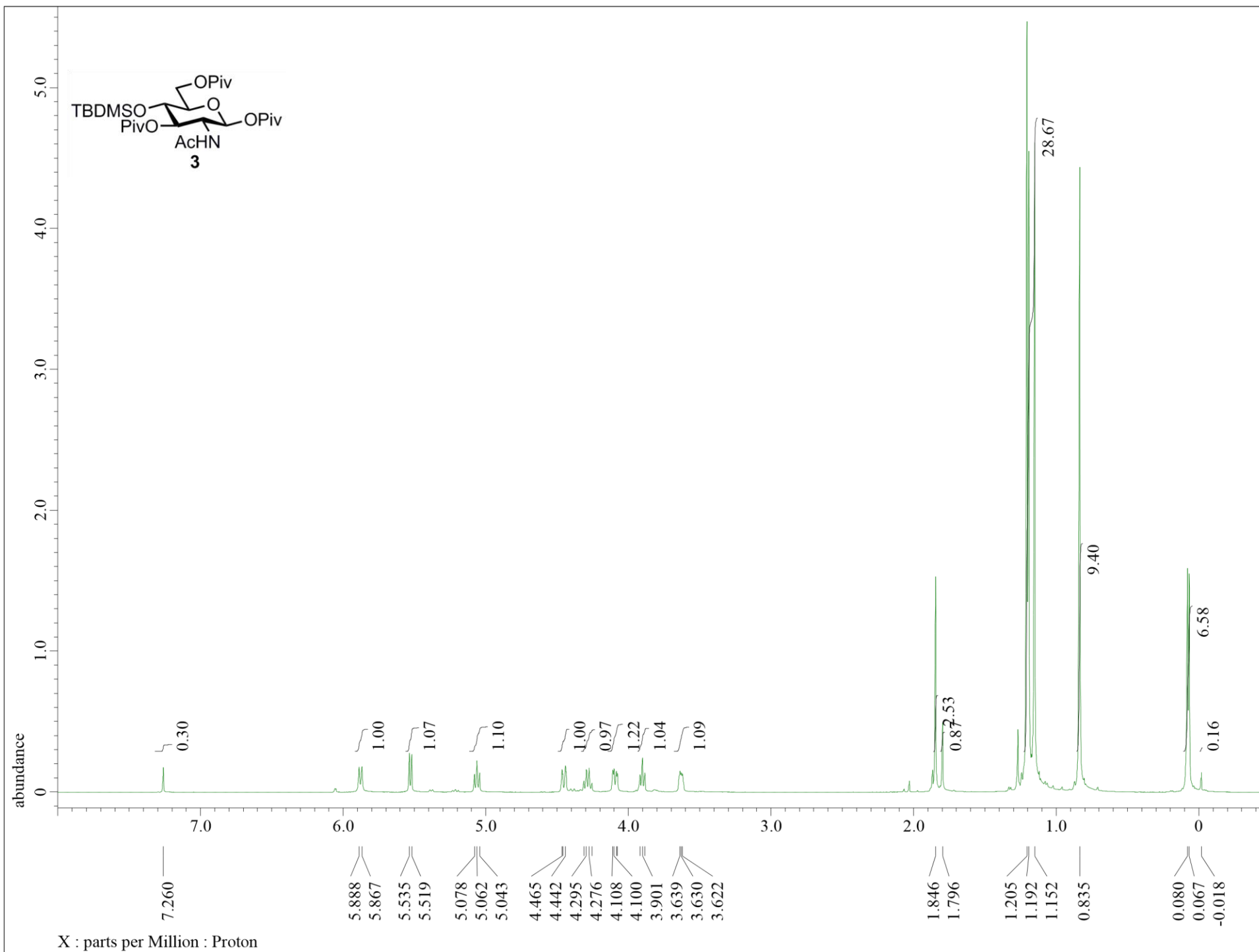
for

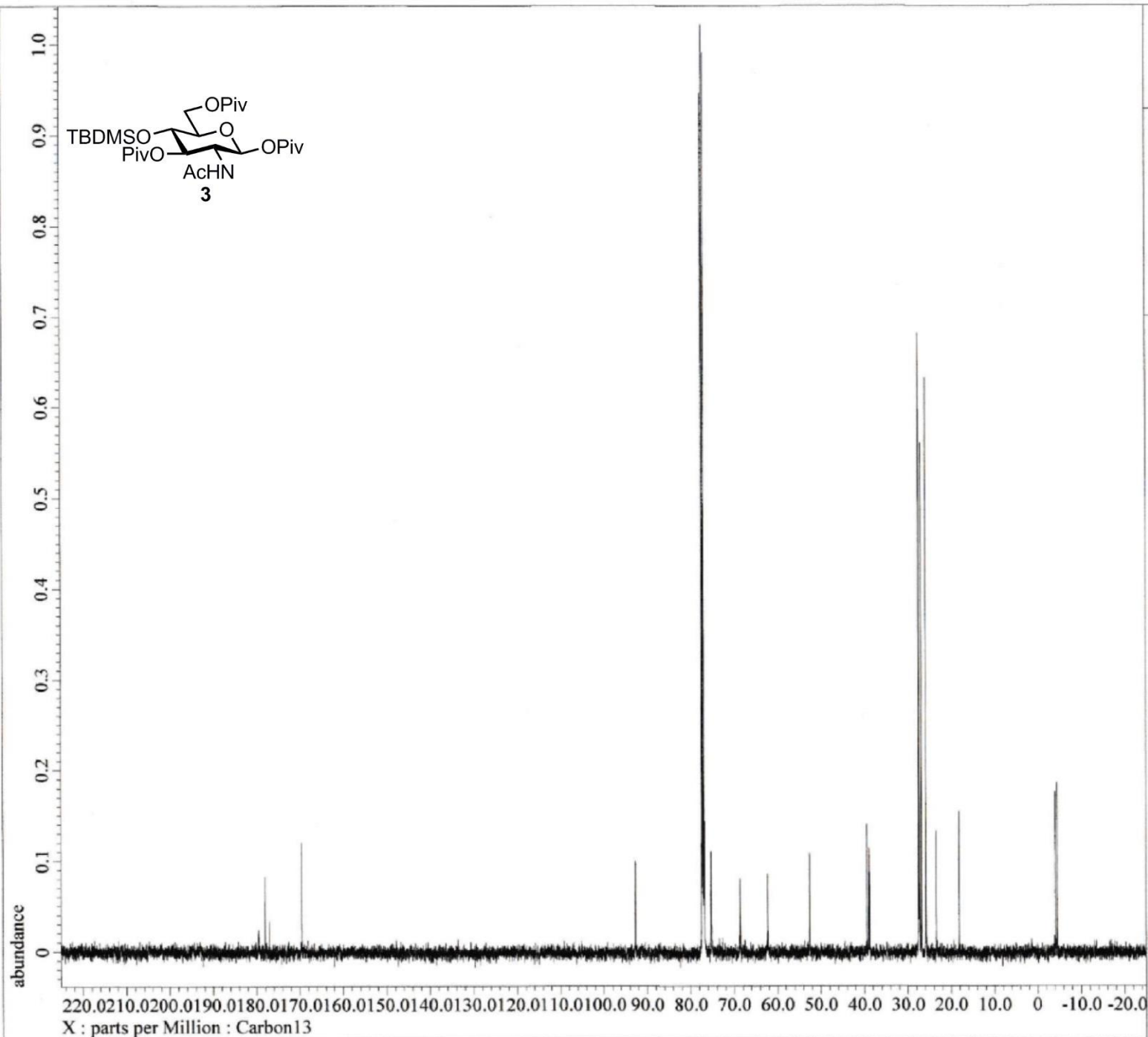
A direct method for β -glycosylation with an *N*-acetylglucosamine aimed by
a 4-*O*-TBDMS protecting group

Contents

NMR spectra of compound 3	-----	p. 3–4
NMR spectra of compound 6	-----	p. 5–6
NMR spectra of compound 7	-----	p. 7–8

NMR spectra of compound 8	p. 9–10
NMR spectra of compound 9	p. 11–12
NMR spectra of compound 17	p. 13–14
NMR spectra of compound 18	p. 15–16
NMR spectra of compound 19	p. 17–18
NMR spectra of compound 20	p. 19–20
NMR spectra of compound 21	p. 21–22
NMR spectra of compound 22	p. 23–24





```

---- PROCESSING PARAMETERS ----
dc balance( 0, FALSE )
sexp( 2.0[Hz], 0.0[s] )
trapezoid( 0[%], 0[%], 80[%], 100[%] )
zerofill( 1 )
fft( 1, TRUE, TRUE )
machinephase
ppm

```

以下に由来: NAc 4-Si carbon-2-1.jdf

```

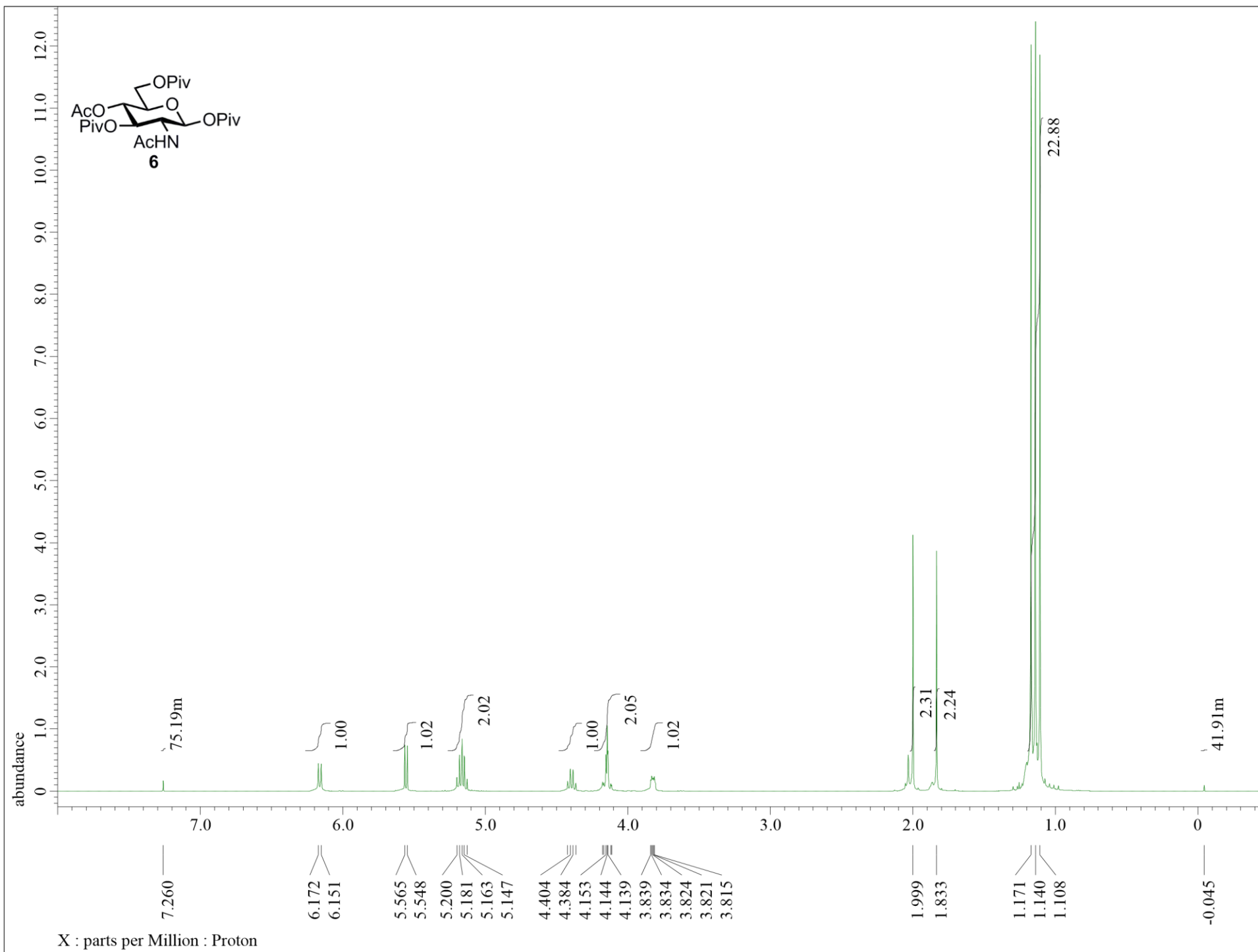
Filename           = NAC_4-Si_carbon-2-2.jdf
Author             = delta
Experiment         = carbon.jxp
Sample_Id         = NAC_4-Si
Solvent           = CHLOROFORM-D
Creation_Time      = 27-JUL-2015 10:51:47
Revision_Time     = 27-JUL-2015 11:03:06
Current_Time      = 27-JUL-2015 11:03:57

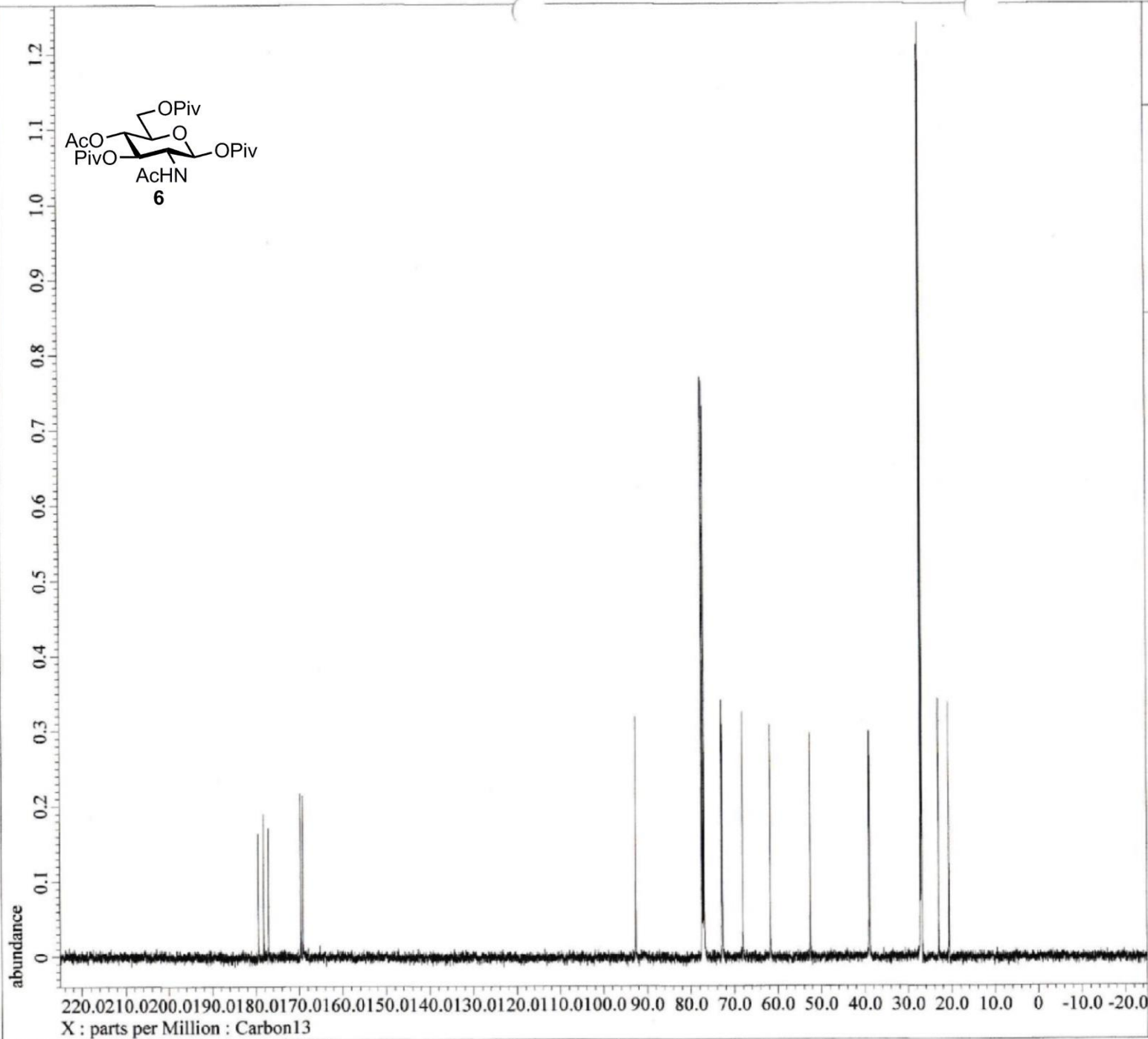
Comment           = single pulse decoupled gat
Data_Format       = 1D COMPLEX
Dim_Size          = 26214
Dim_Title         = Carbon13
Dim_Units         = [ppm]
Dimensions        = X
Site              = JMN-ECA500
Spectrometer      = DELTA2_NMR

Field_Strength    = 11.7473579[T] (500[MHz])
X_Acq_Duration    = 0.83361792[s]
X_Domain          = 13C
X_Freq            = 125.76529768[MHz]
X_Offset          = 100[ppm]
X_Points          = 32768
X_Prescans        = 4
X_Resolution      = 1.19959034[Hz]
X_Sweep           = 39.3081761[kHz]
X_Sweep_Clippped = 31.44654088[kHz]
Irr_Domain        = Proton
Irr_Freq          = 500.15991521[MHz]
Irr_Offset        = 5.0[ppm]
Clipped           = FALSE
Scans             = 200
Total_Scans       = 200

Relaxation_Delay  = 2[s]
Recvr_Gain        = 56
Temp_Get          = 21.4[°C]
X_90_Width        = 8.17[us]
X_Acq_Time        = 0.83361792[s]
X_Angle           = 30[deg]
X_Atn             = 6.5[dB]
X_Pulse           = 2.72333333[us]
Irr_Atn_Dec       = 22.66[dB]
Irr_Atn_Noise     = 22.66[dB]
Irr_Noise         = WALTZ
Irr_Pwidth        = 92[us]
Decoupling        = TRUE

```





---- PROCESSING PARAMETERS ----
 dc_balance(0, FALSE)
 sexp(2.0[Hz], 0.0[s])
 trapezoid(0[%], 0[%], 80[%], 100[%])
 zerofill(1)
 fft(1, TRUE, TRUE)
 machinephase
 ppm

以下に由来: NAc_4Ac_carbon-1-1.jdf

Filename = NAc_4Ac_carbon-1-2.jdf
 Author = console
 Experiment = carbon.jxp
 Sample_Id = NAc_4Ac
 Solvent = CHLOROFORM-D
 Creation_Time = 17-JUL-2015 15:26:10
 Revision_Time = 17-JUL-2015 15:37:46
 Current_Time = 17-JUL-2015 15:40:02

Comment = single pulse decoupled gat
 Data Format = 1D COMPLEX
 Dim_Size = 26214
 Dim_Title = Carbon13
 Dim_Units = [ppm]
 Dimensions = X
 Site = JMN-ECA500
 Spectrometer = DELTA2_NMR

Field Strength = 11.7473579[T] (500[MHz])
 X_Acq_Duration = 0.83361792[s]
 X_Domain = 13C
 X_Freq = 125.76529768[MHz]
 X_Offset = 100[ppm]
 X_Points = 32768
 X_Prescans = 4
 X_Resolution = 1.19959034[Hz]
 X_Sweep = 39.3081761[kHz]
 X_Sweep_Clippped = 31.44654088[kHz]
 Irr_Domain = Proton
 Irr_Freq = 500.15991521[MHz]
 Irr_Offset = 5.0[ppm]
 Clipped = FALSE
 Scans = 200
 Total_Scans = 200

Relaxation_Delay = 2[s]
 Recvr_Gain = 56
 Temp_Get = 22.3[dC]
 X_90_Width = 8.17[us]
 X_Acq_Time = 0.83361792[s]
 X_Angle = 30[deg]
 X_Atn = 6.5[dB]
 X_Pulse = 2.7233333[us]
 Irr_Atn_Dec = 22.66[dB]
 Irr_Atn_Noe = 22.66[dB]
 Irr_Noise = WALTZ
 Irr_Pwidth = 92[us]
 Decoupling = TRUE

