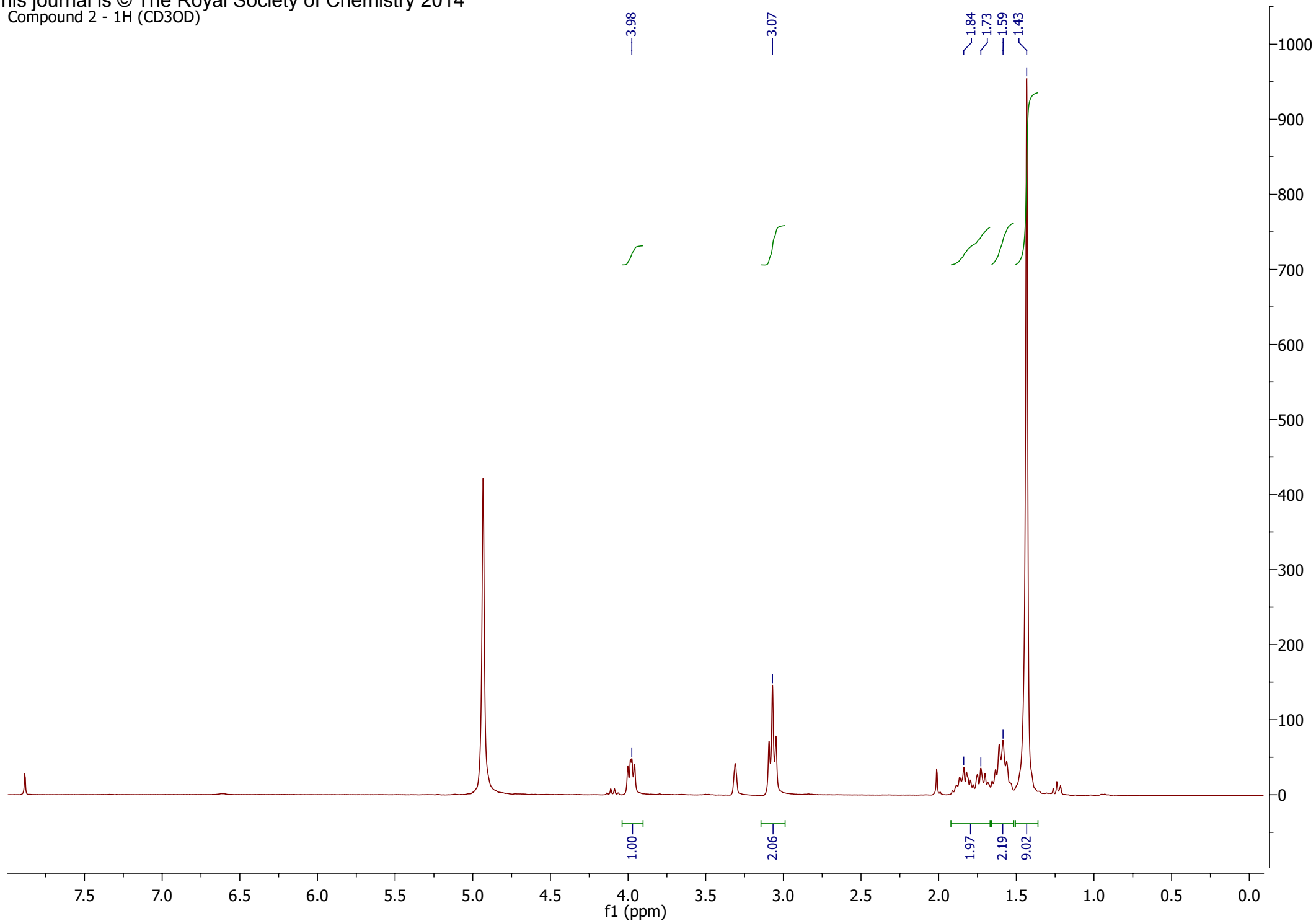
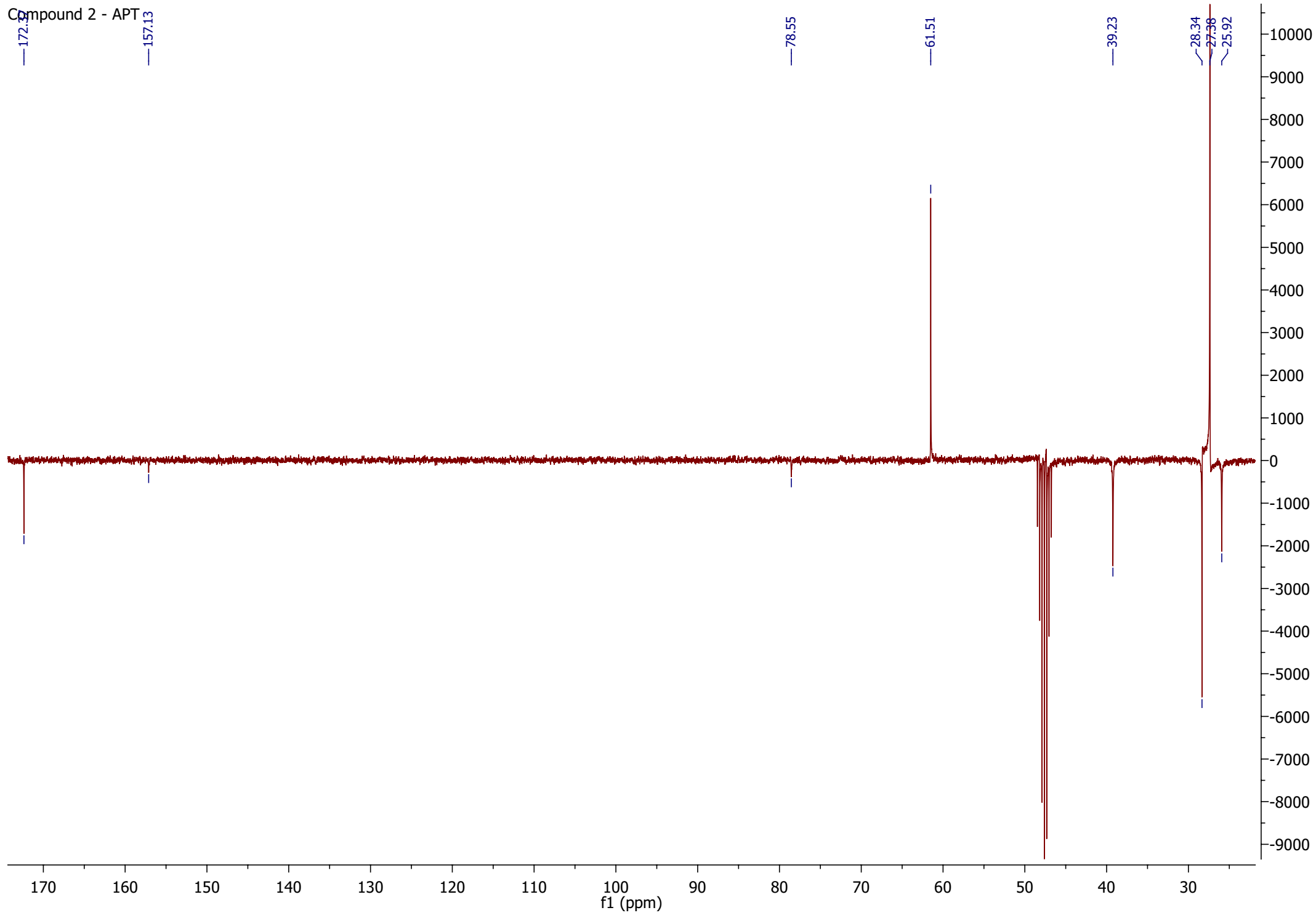
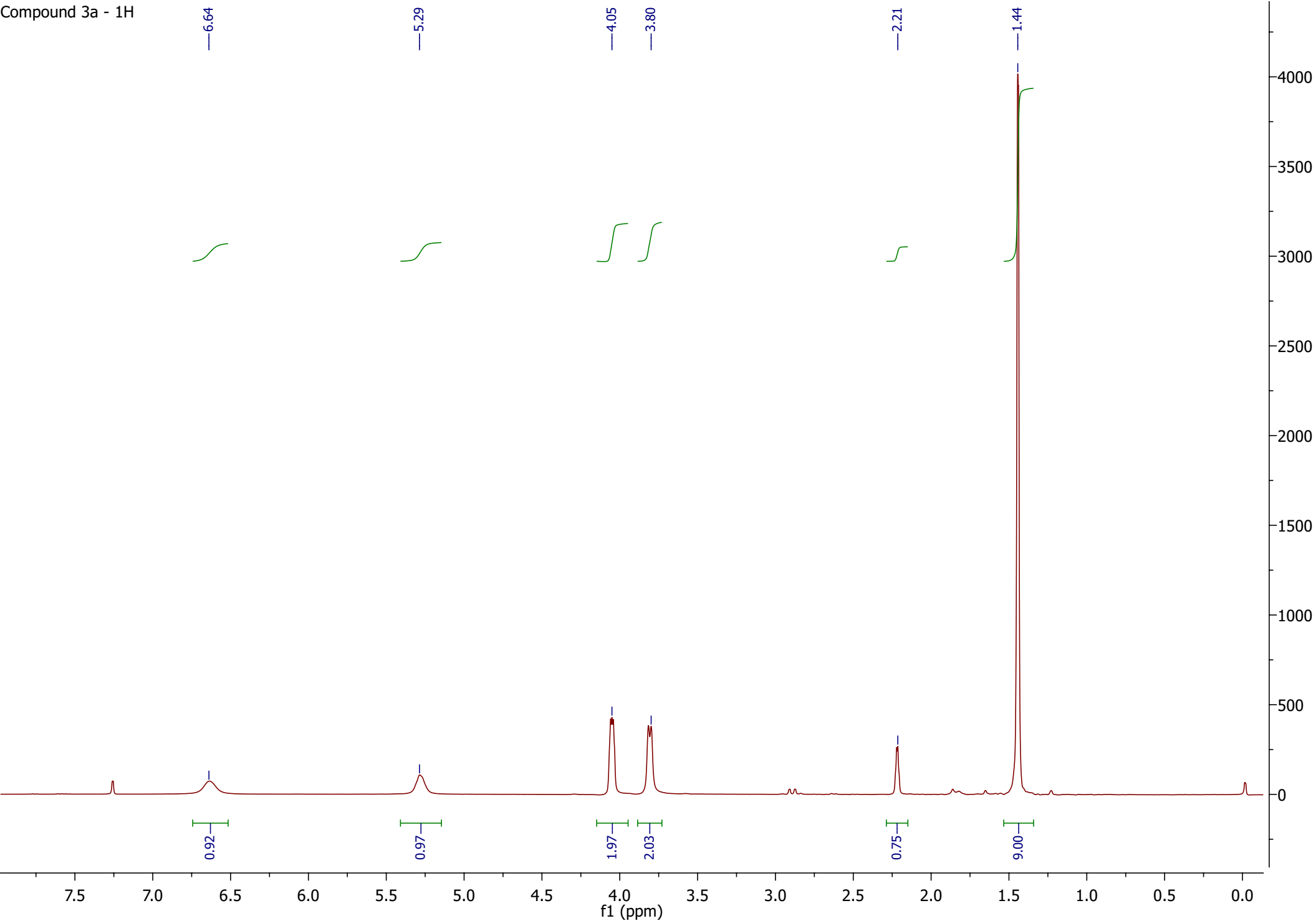




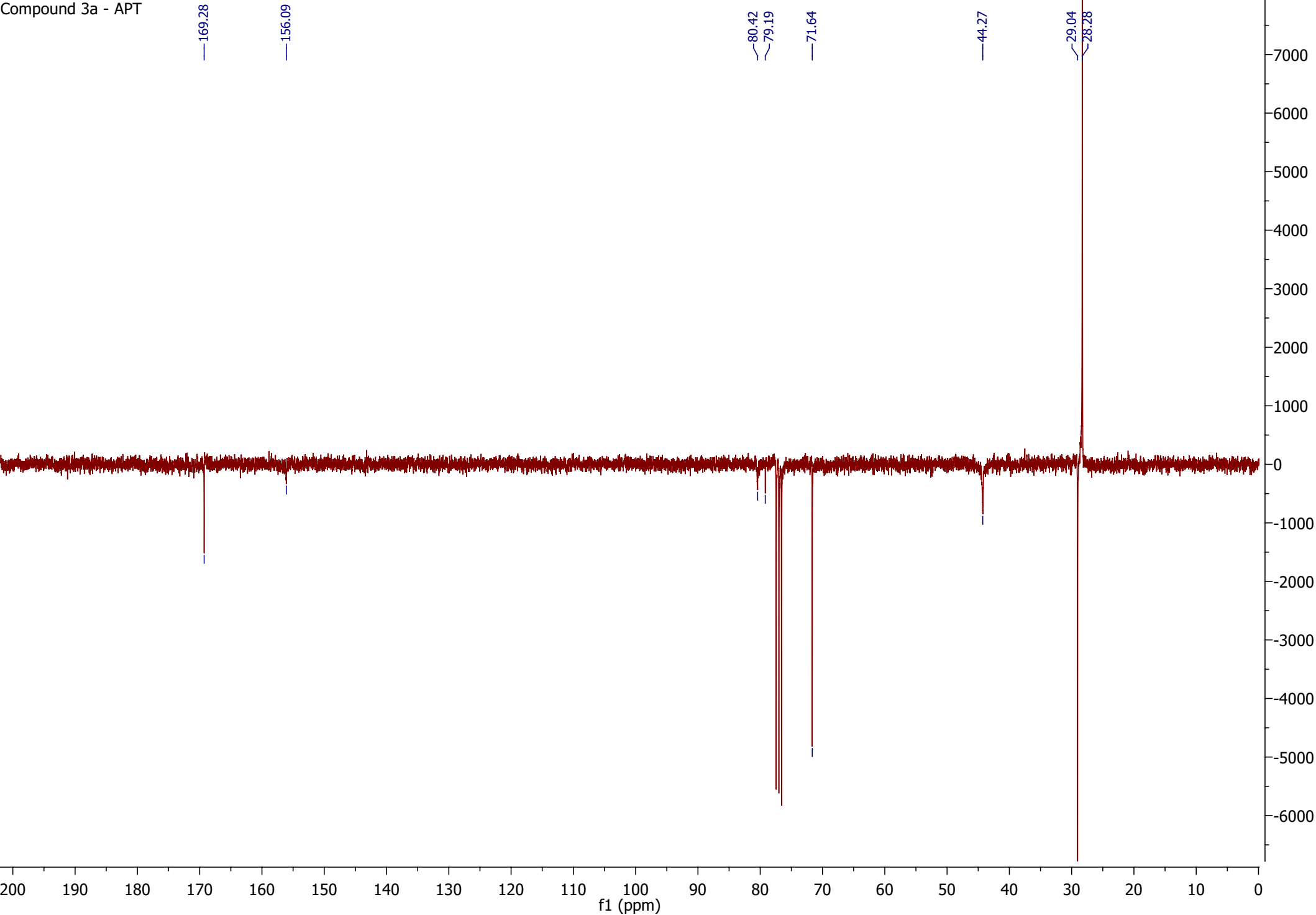
Compound 2 - APT



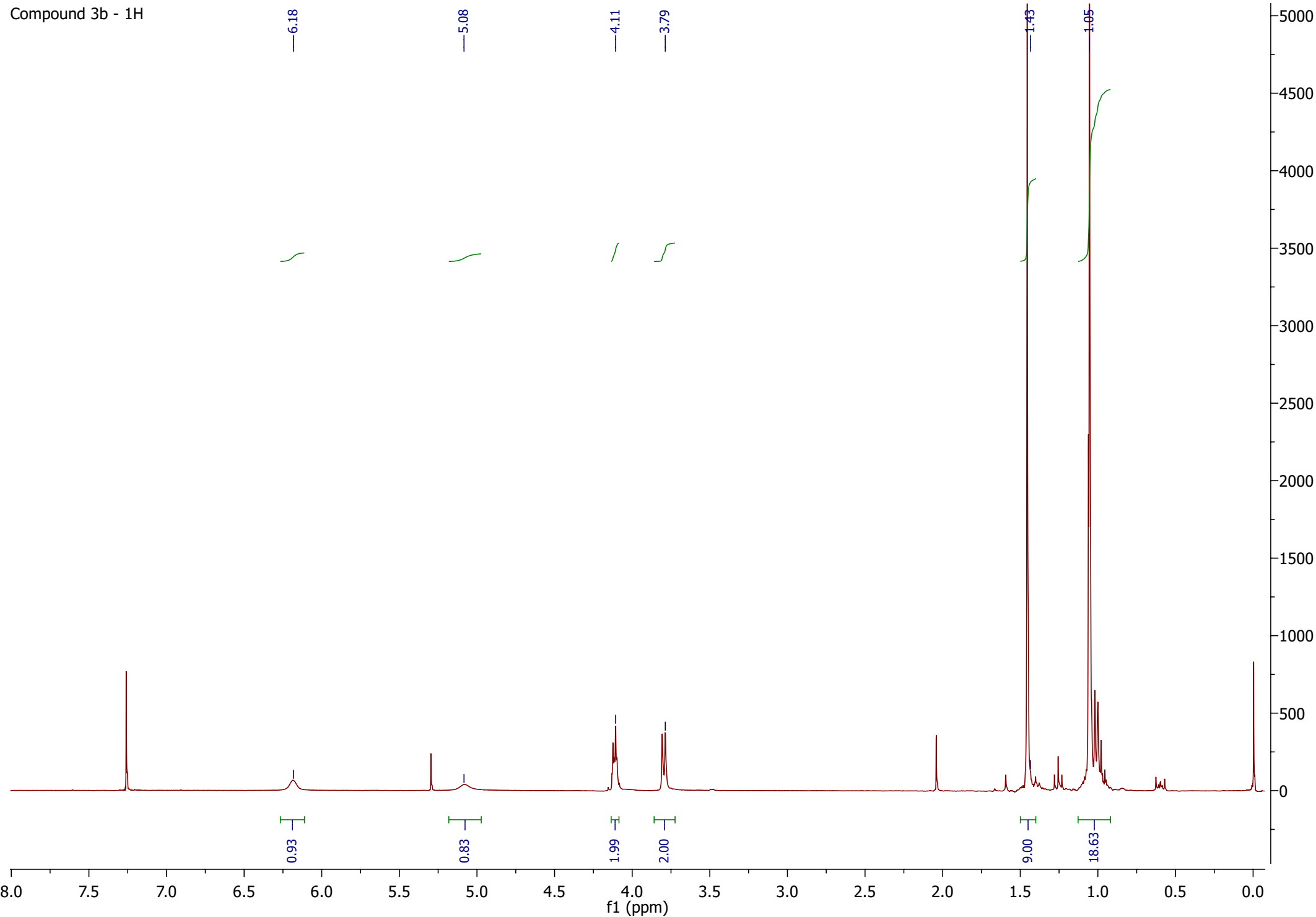
Compound 3a - 1H



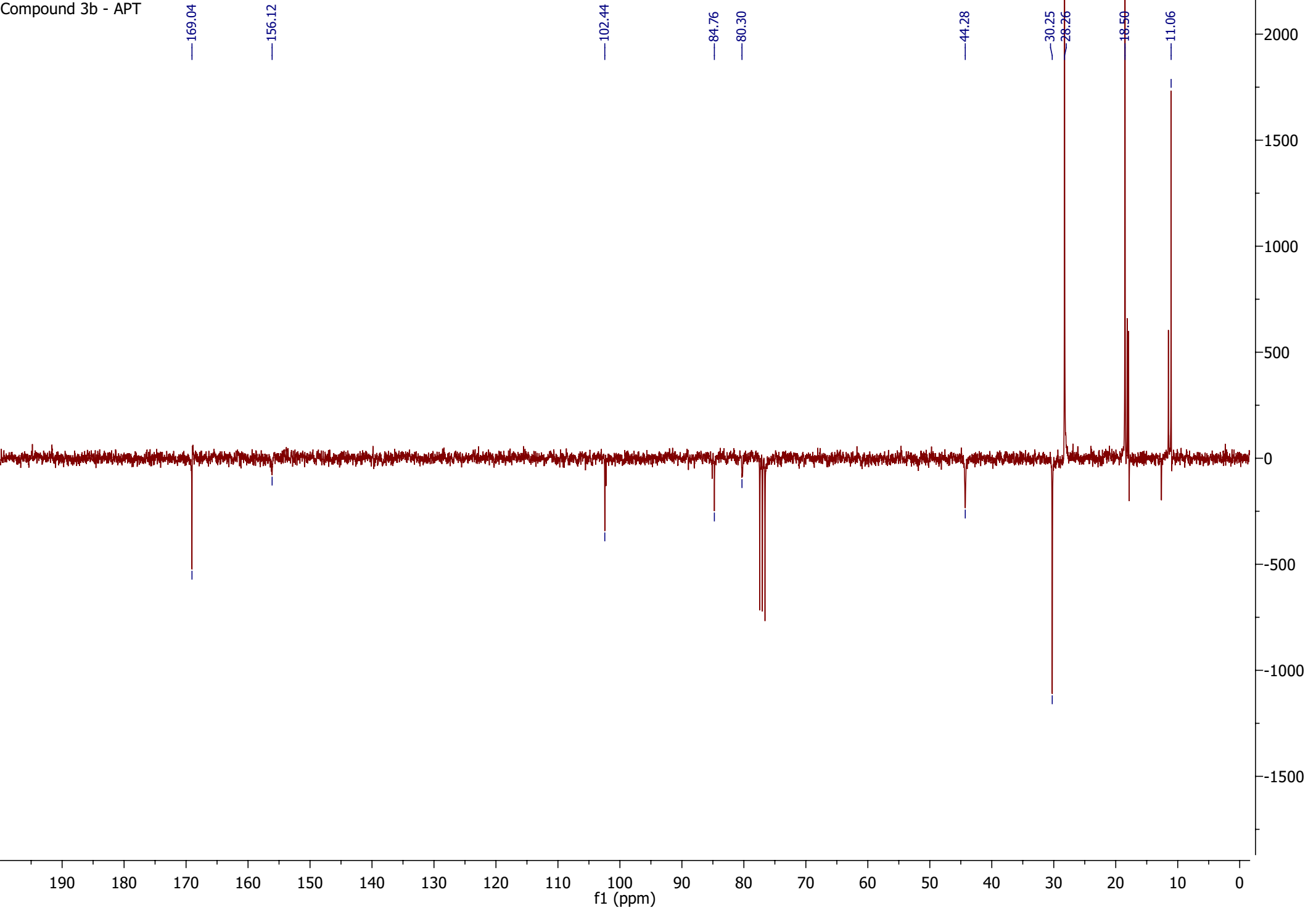
Compound 3a - APT



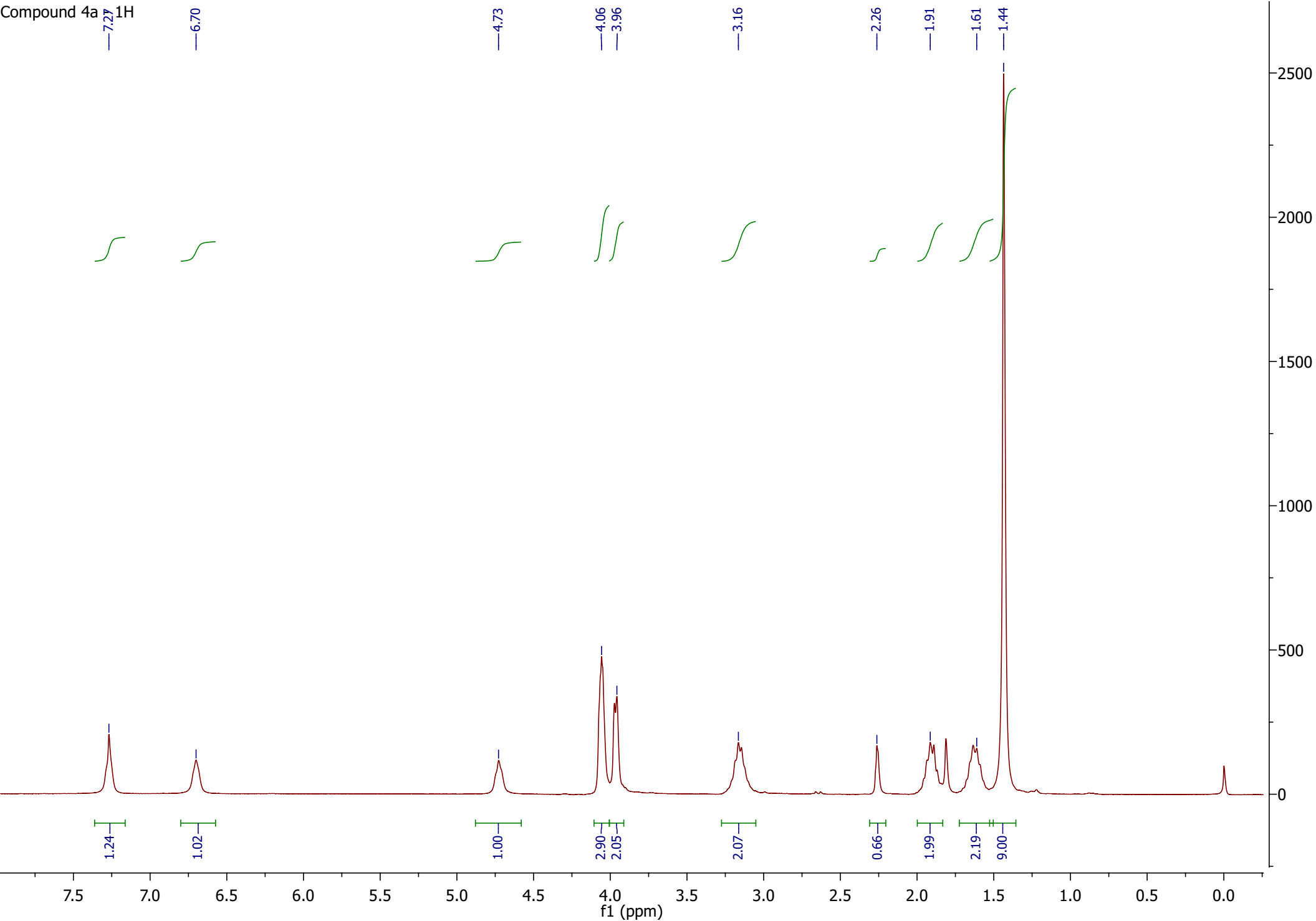
Compound 3b - 1H



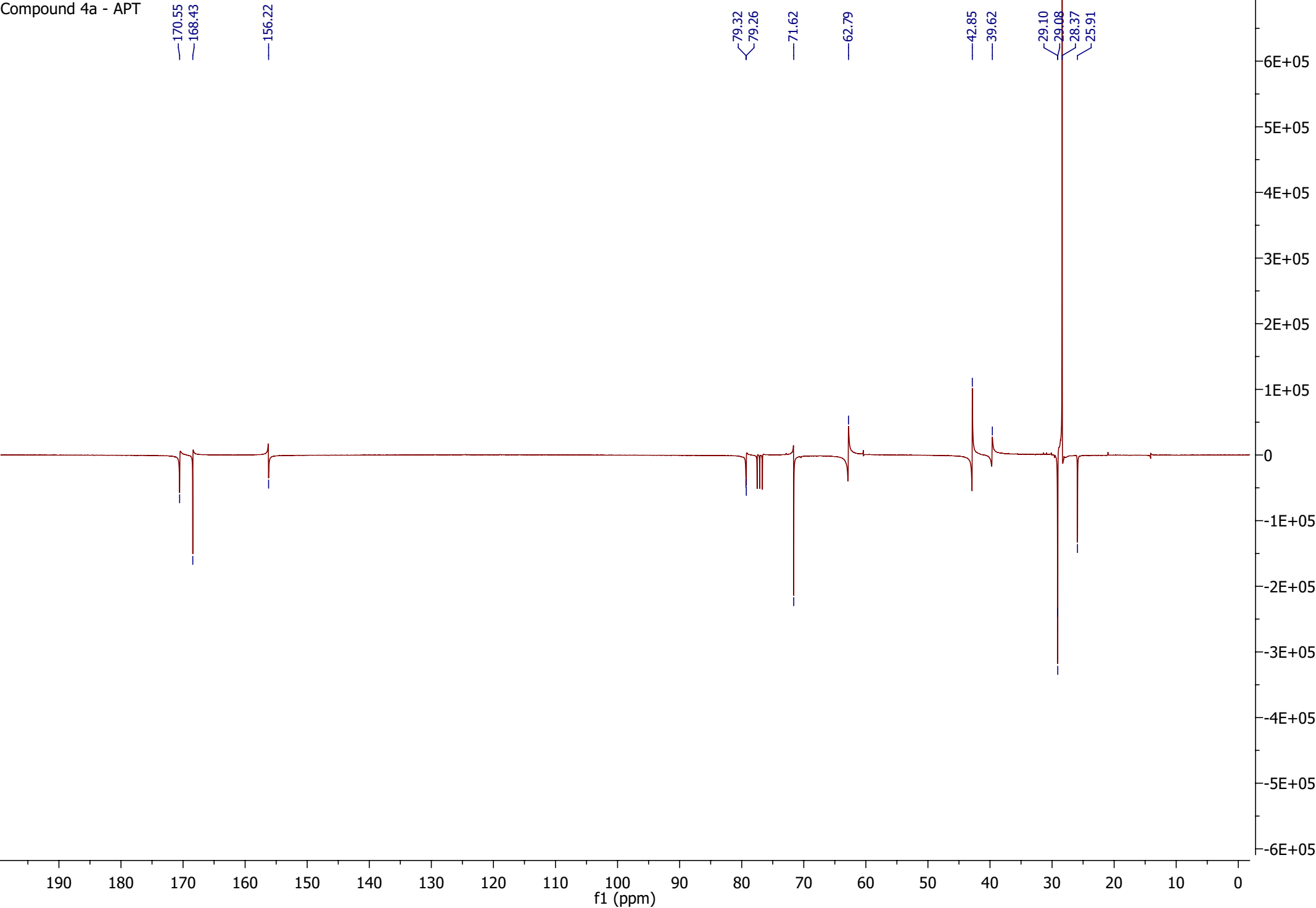
Compound 3b - APT



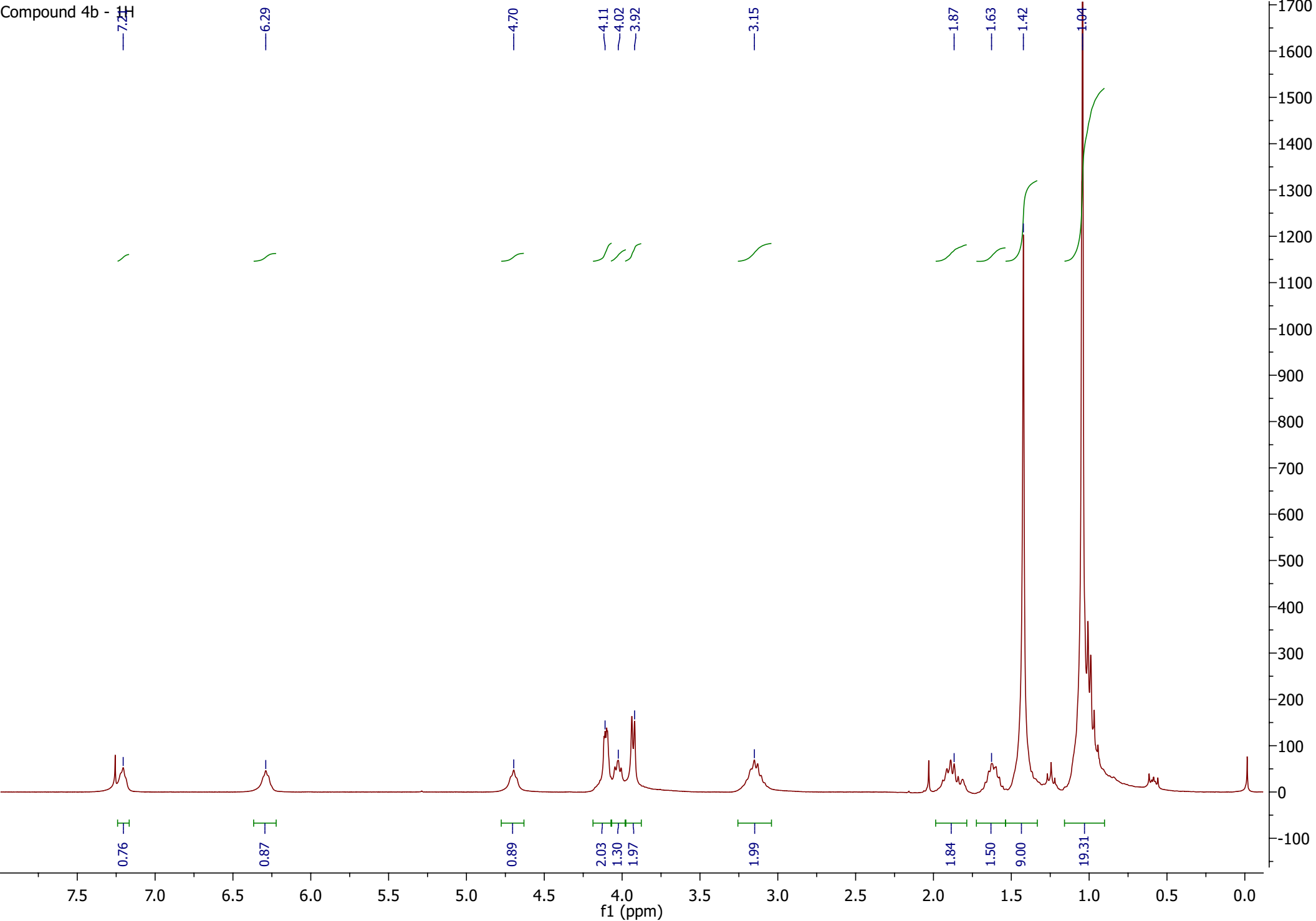
Compound 4a 1H



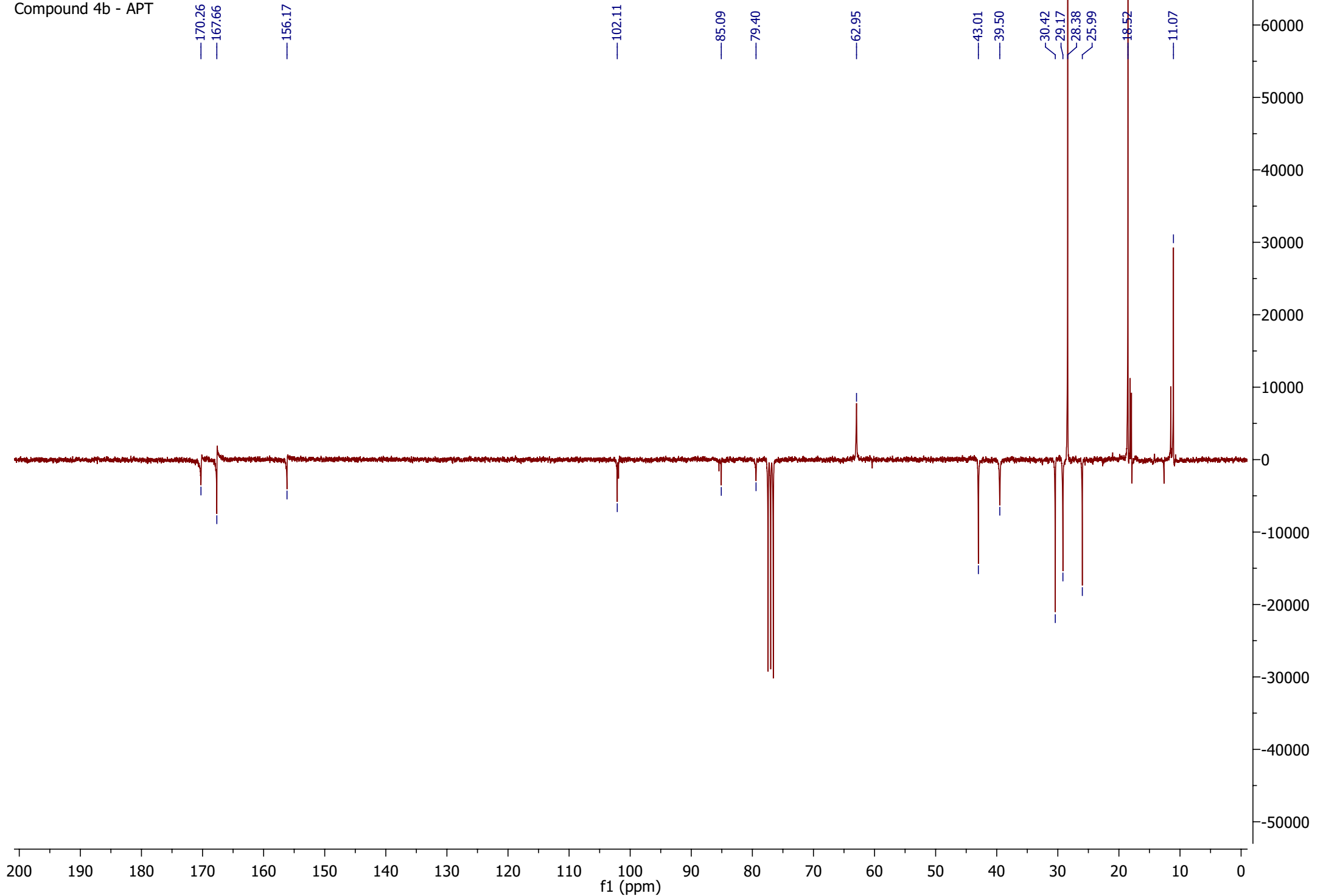
Compound 4a - APT



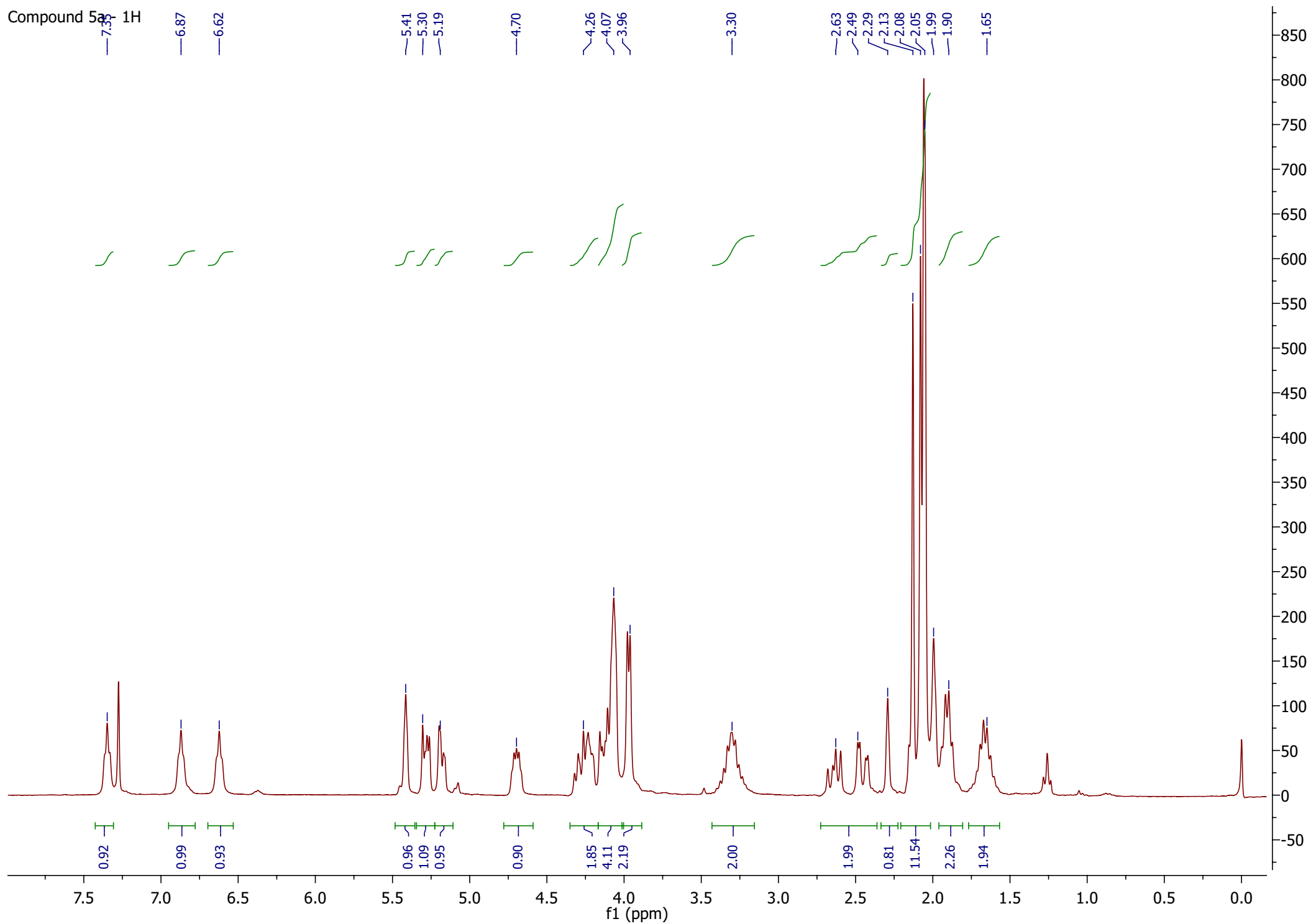
Compound 4b - ¹H



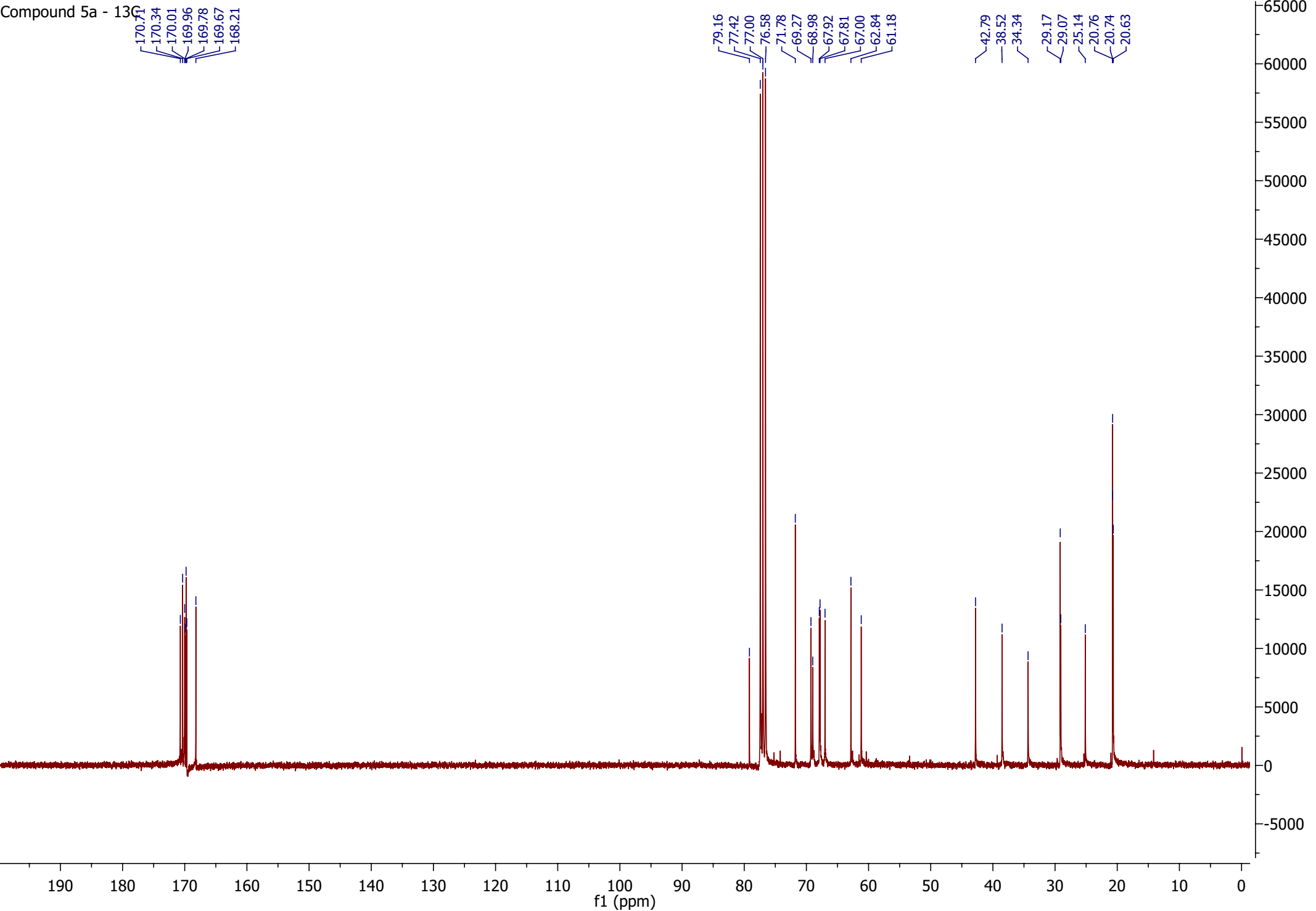
Compound 4b - APT



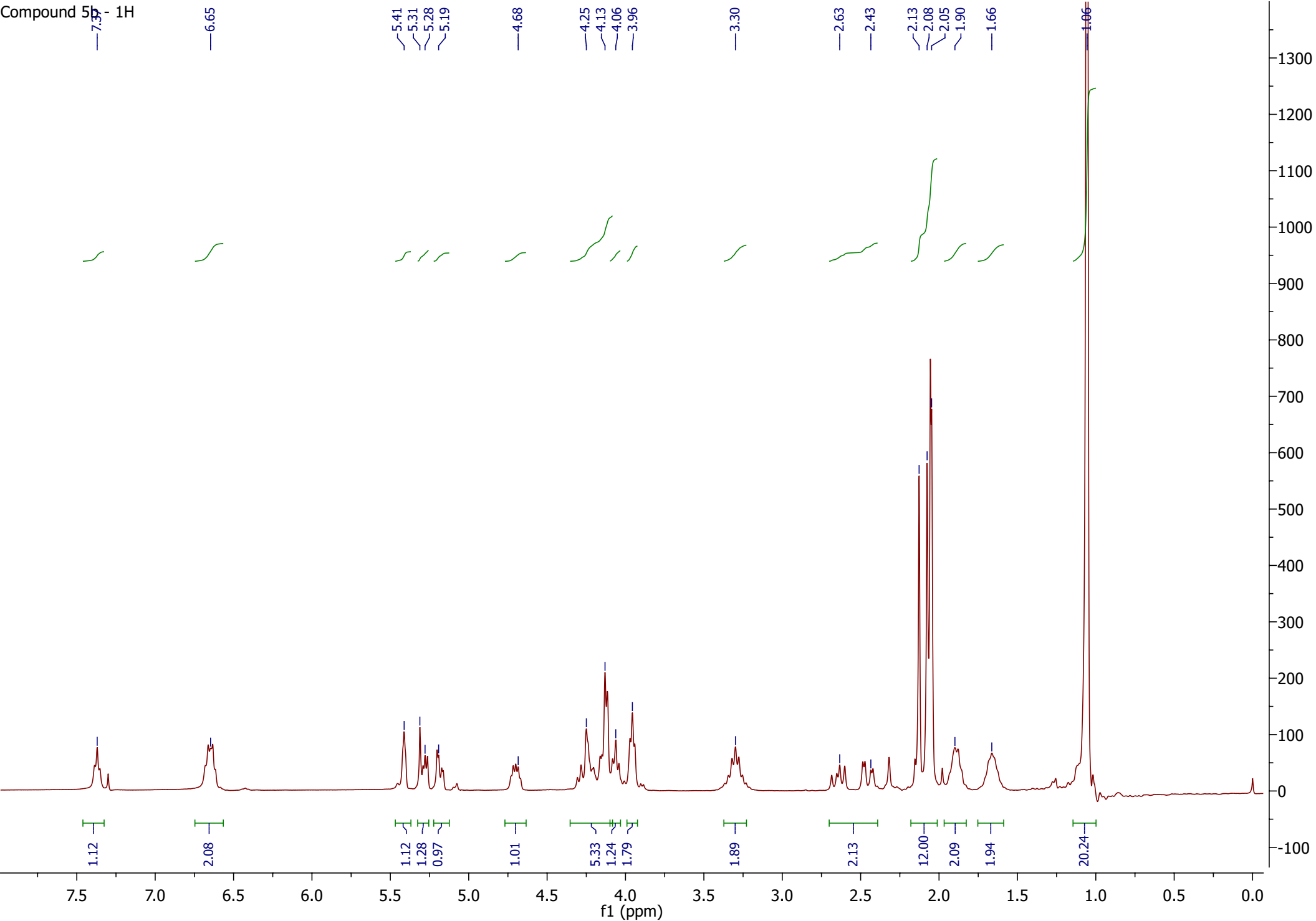
Compound 5a- 1H



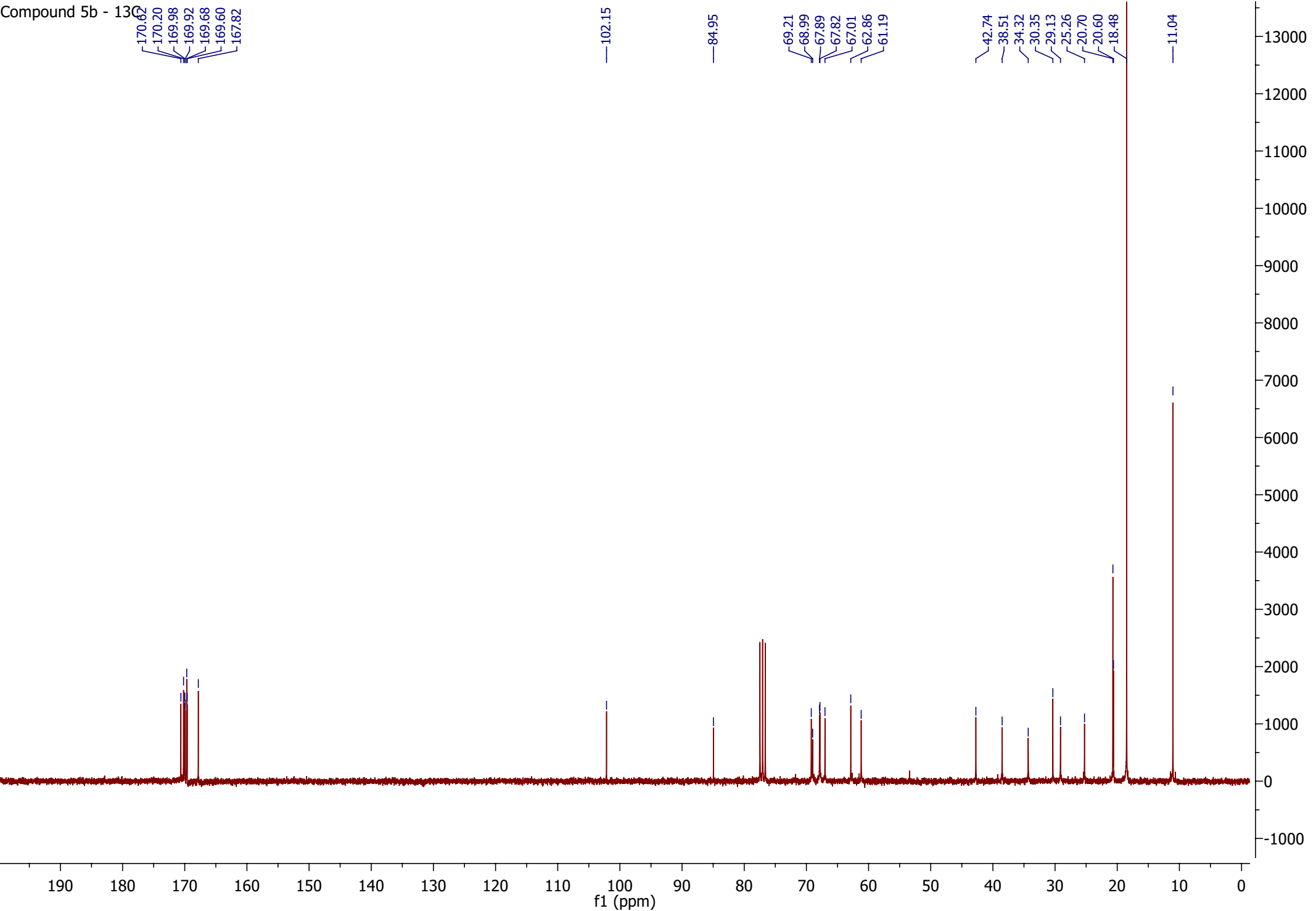
Compound 5a - 13C



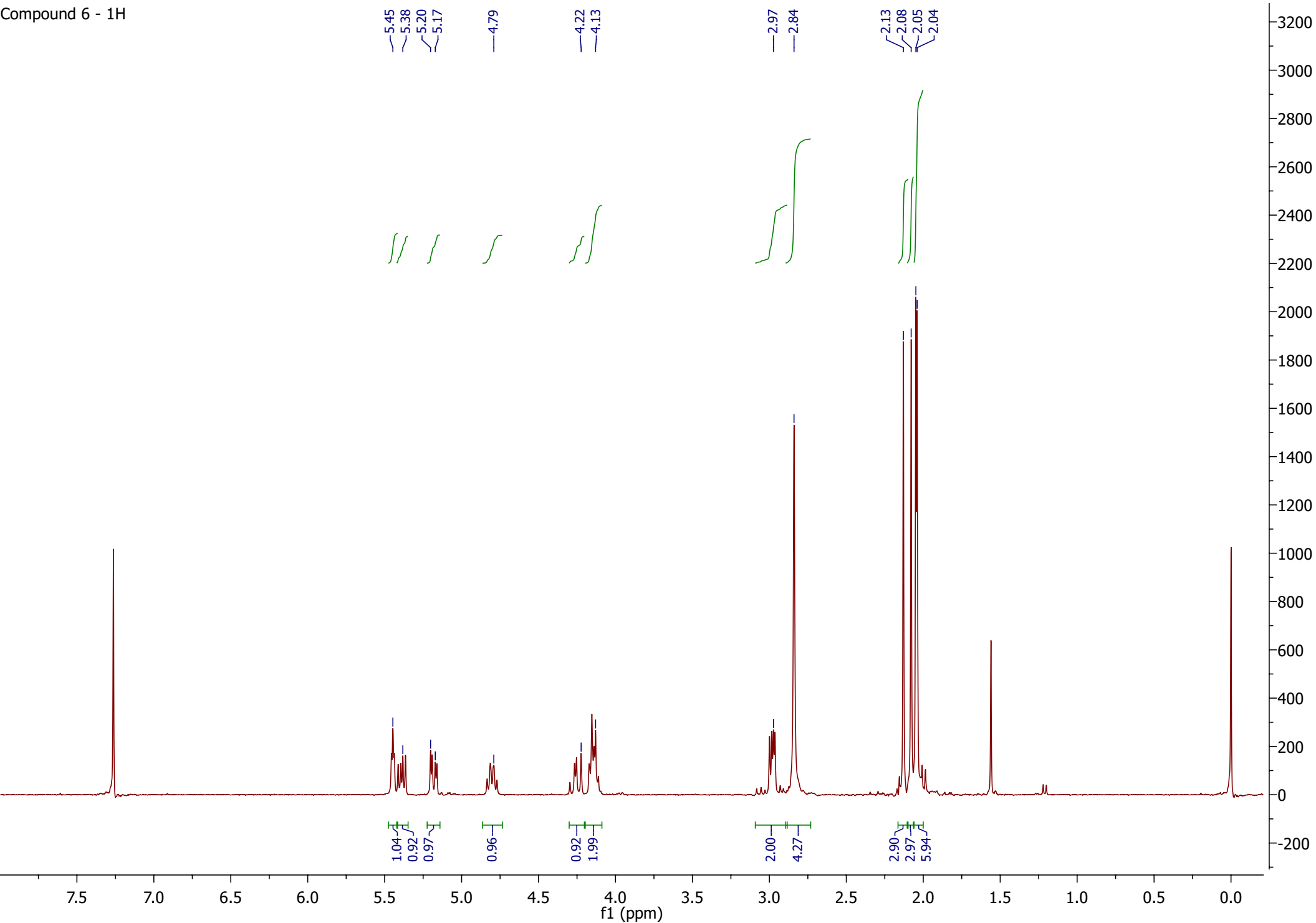
Compound 5b - 1H



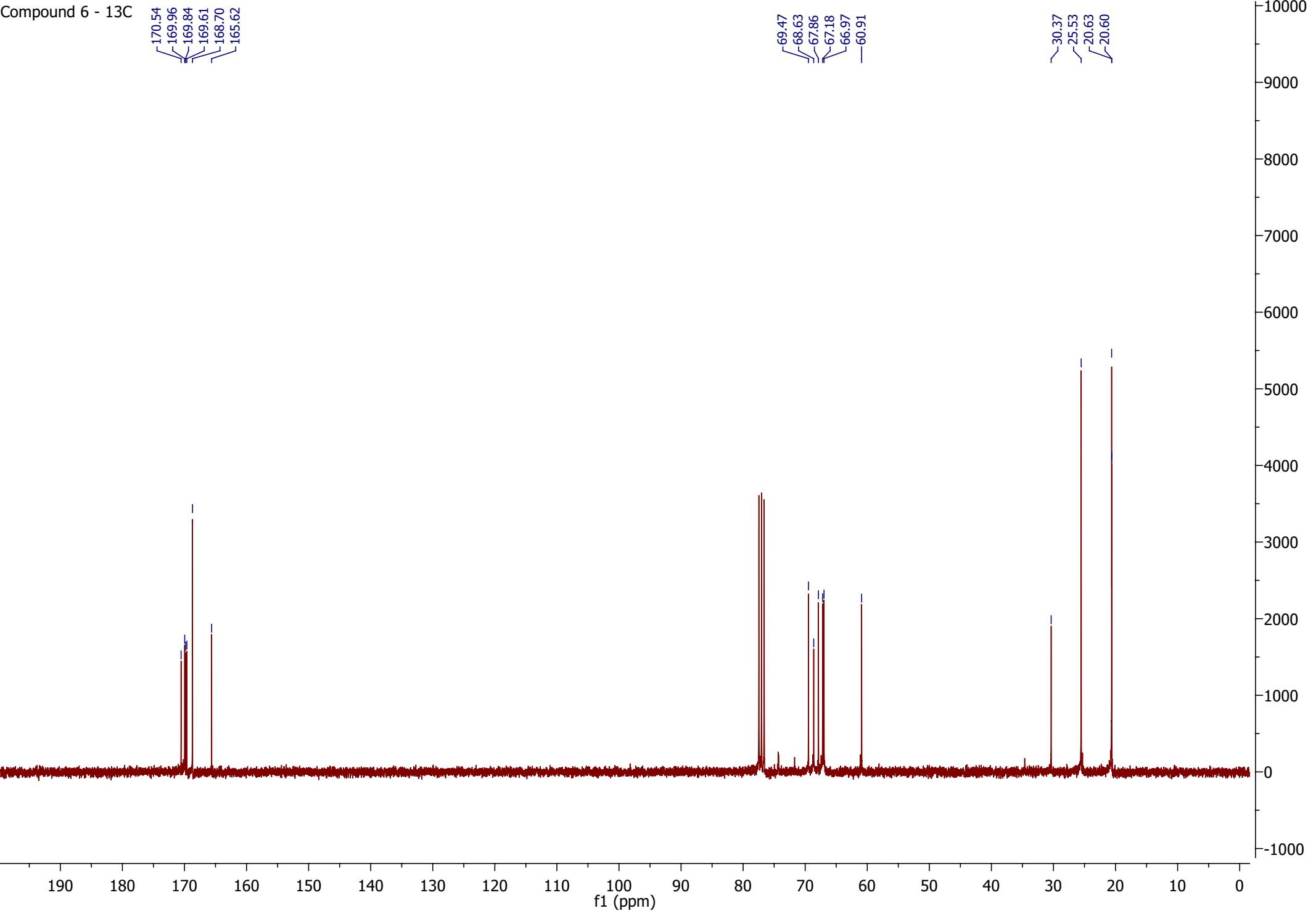
Compound 5b - 13C



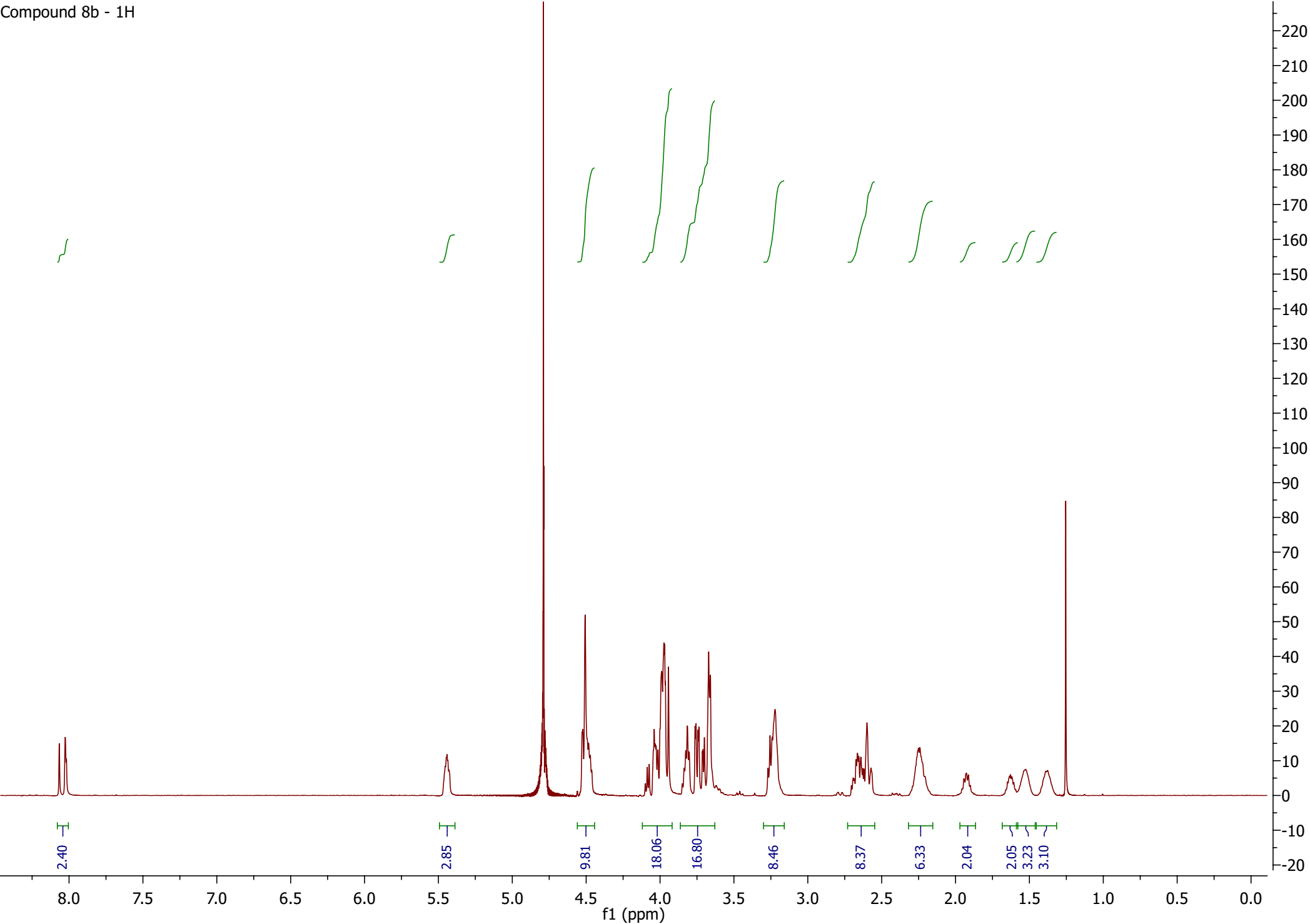
Compound 6 - 1H

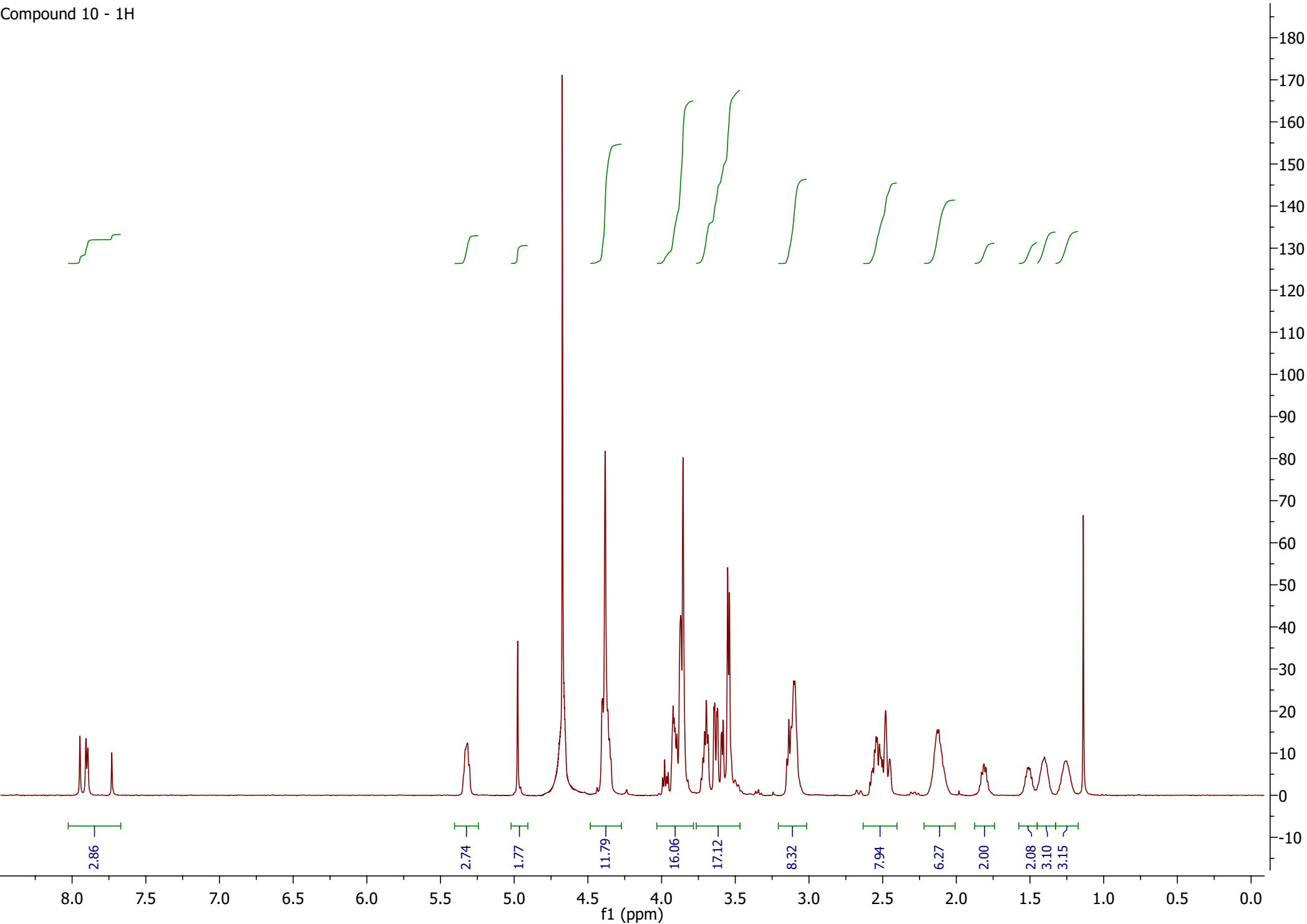


Compound 6 - ¹³C

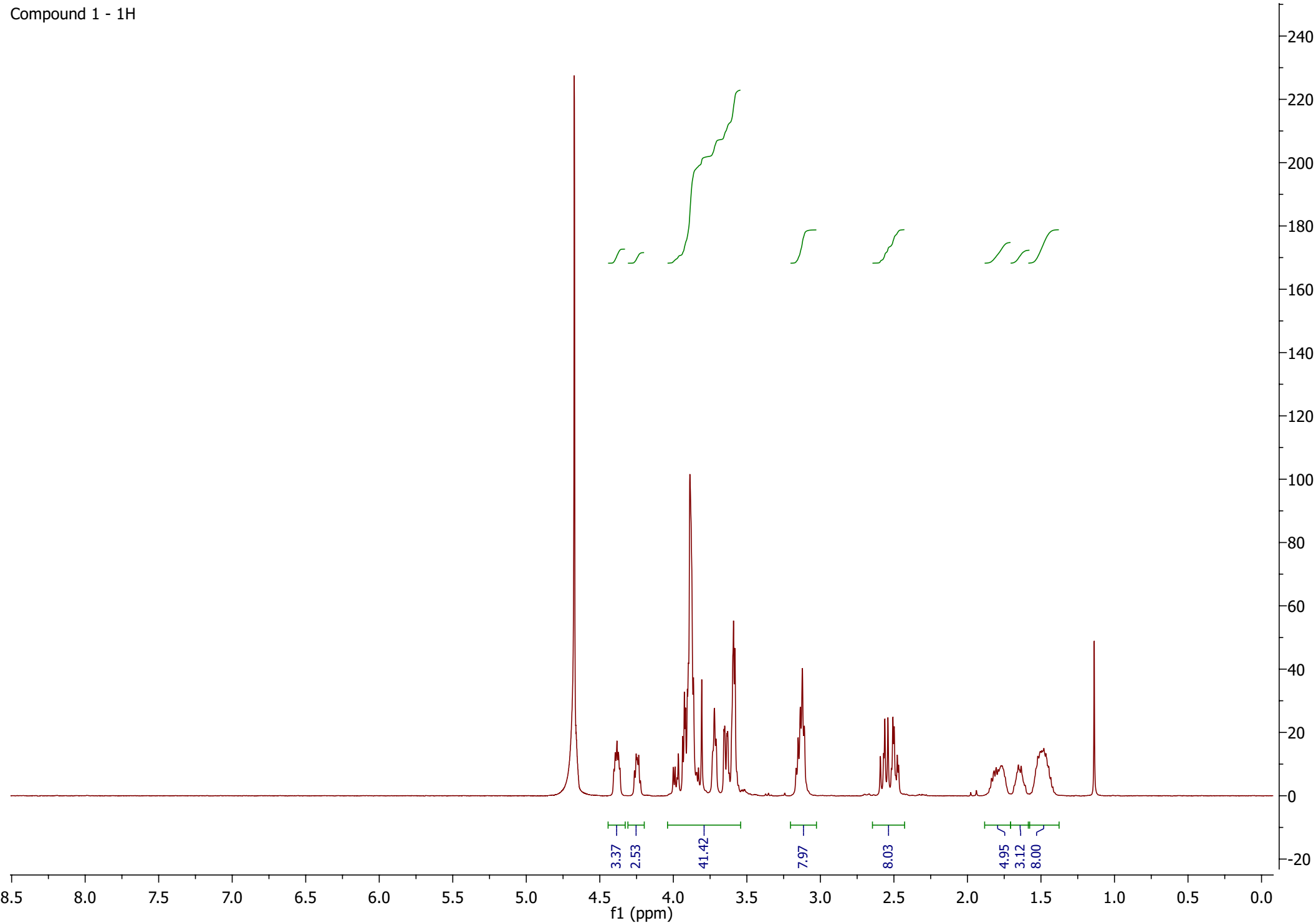


Compound 8b - 1H

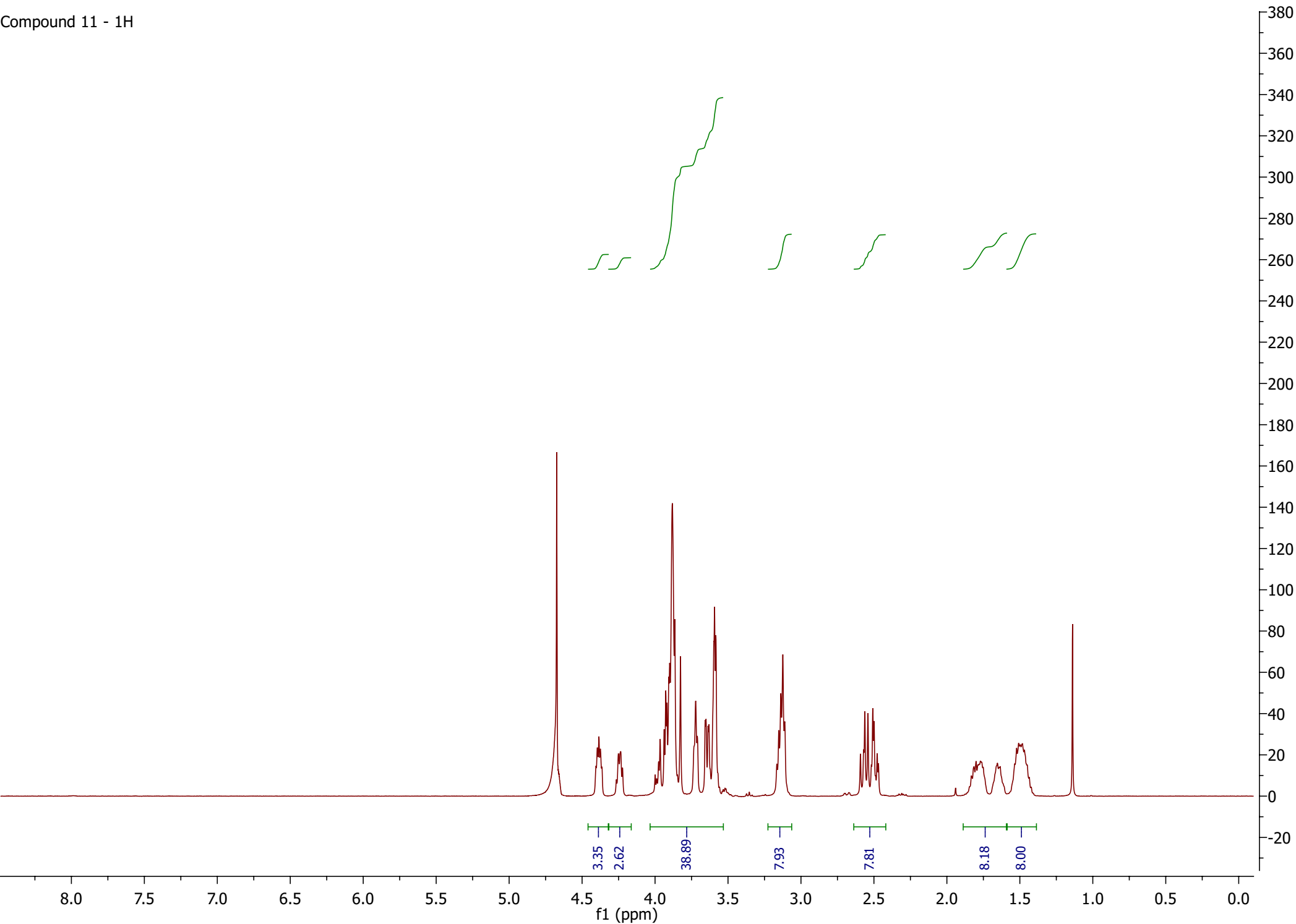




Compound 1 - 1H



Compound 11 - 1H



Sample Information Item

Date and Time : 7/25/2013 13:06:45 PM

User : Steffen

Type : Unknown

Inj. Volume : 20.000

Data Name : E:\Users\Steffen\Important files\SWalk_prep.qld

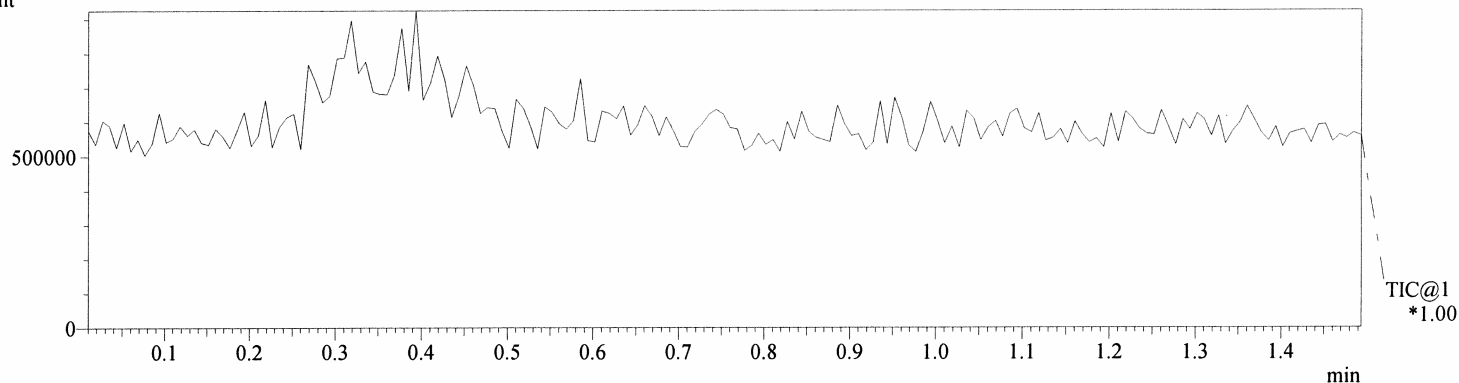
Method Name : E:\Mass files\Method\Positive mode\200-2000 0.3ml-min flow.qlm

Tuning Name : E:\Mass files\Tuning\Autotune 20120620.QLT

Compound 8b

MS Chromatogram

height



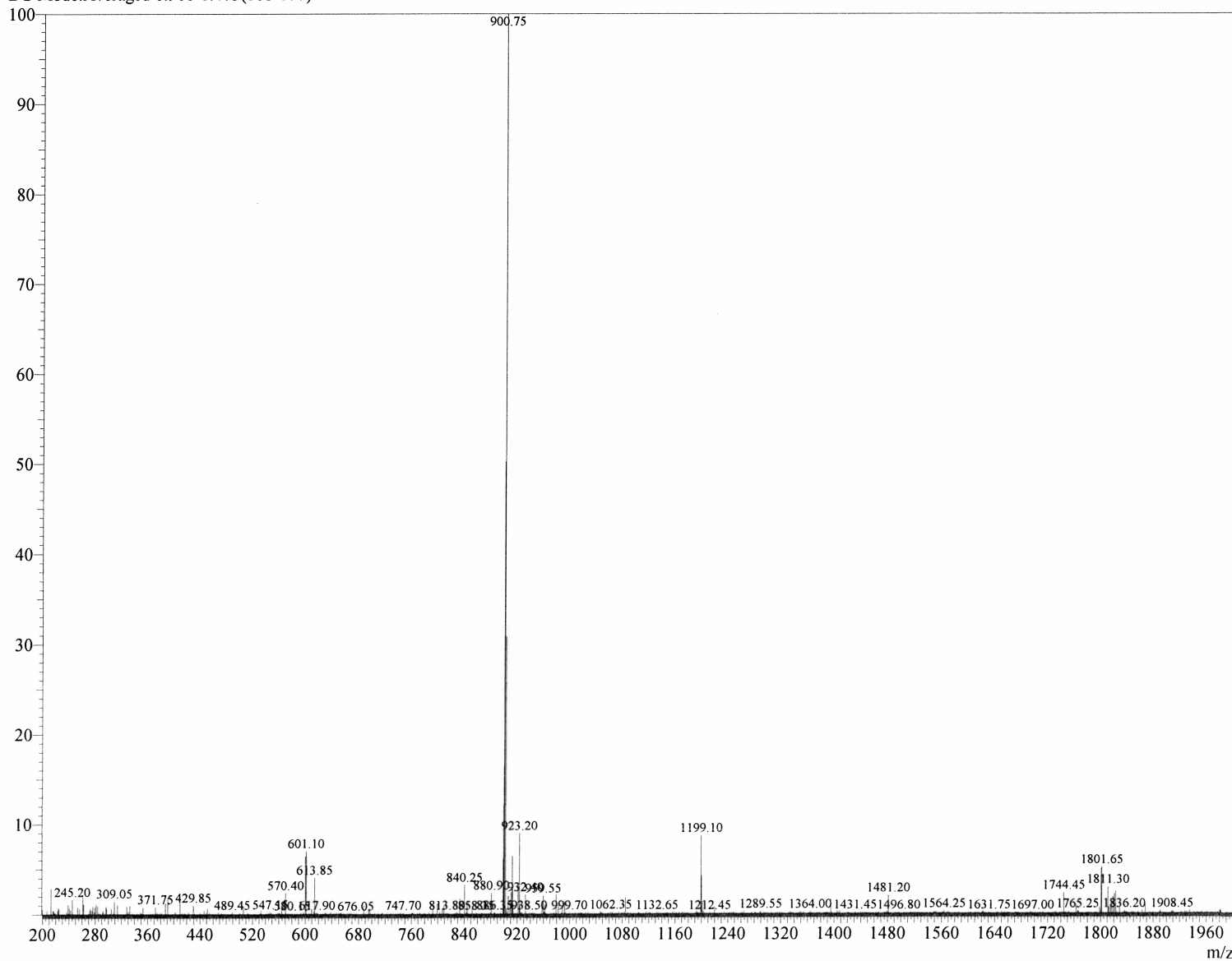
MS Spectrum

Line#:1 R.Time:0.393(Scan#:47) Positive

MassPeaks:879 BasePeak:900.75(40662)

RawMode:Averaged 0.259-0.577(31-69)

BG Mode:Averaged 0.900-1.473(108-177)

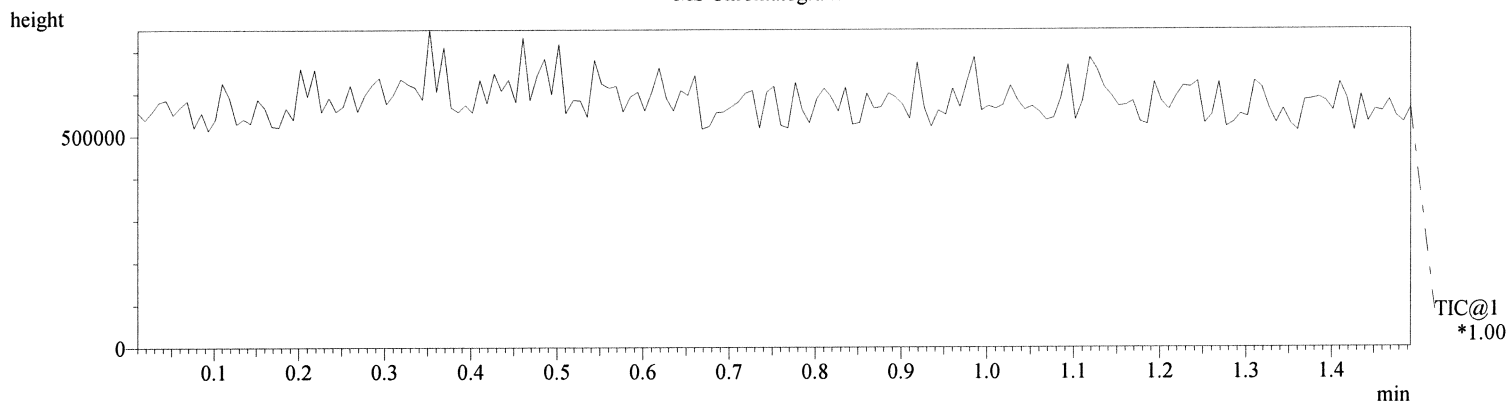


Sample Information Item

Date and Time : 7/25/2013 13:09:51 PM
User : Steffen
Type : Unknown
Inj. Volume : 20.000
Data Name : E:\Users\Steffen\Important files\SWacid_prep.qld
Method Name : E:\Mass files\Method\Positive mode\200-2000 0.3ml-min flow.qlm
Tuning Name : E:\Mass files\Tuning\Autotune 20120620.QLT

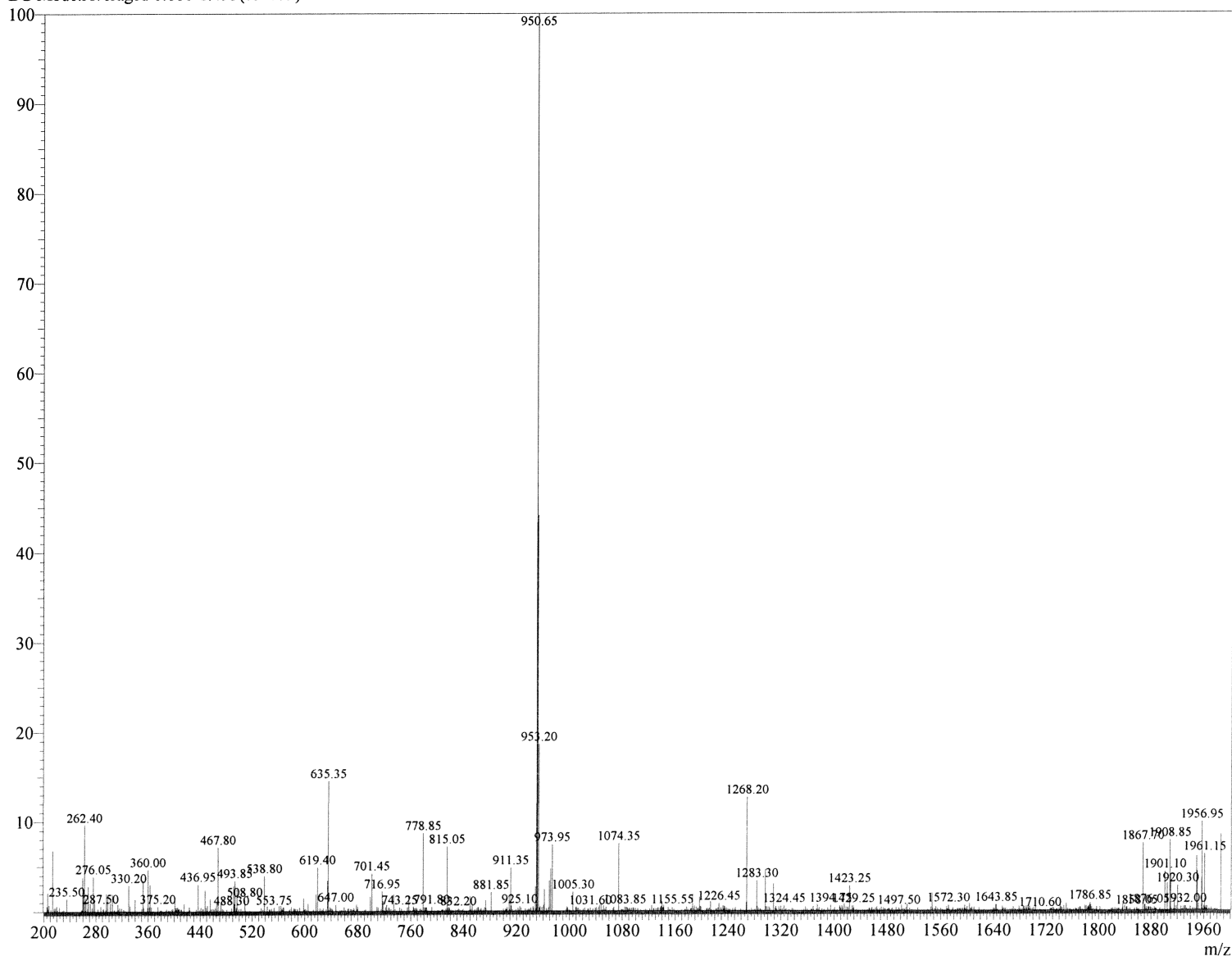
Compound 10

MS Chromatogram



MS Spectrum

Line#:1 R.Time:0.352(Scan#:42) Positive
MassPeaks:856 BasePeak:950.65(15885)
RawMode:Averaged 0.185-0.492(22-59)
BG Mode:Averaged 0.660-1.493(79-179)



Tuning Name : E:\Mass files\Tuning\Autotune 20120620.QLT

Mass spectrum of compound 10. The x-axis represents the mass-to-charge ratio (m/z) from 200 to 1960, and the y-axis represents the relative intensity in percent from 0 to 100. The base peak is at m/z 902.90. Other labeled peaks include:

m/z	Relative Intensity (%)
236.05	~1
293.00	~1
358.70	~1
418.20	~1
483.50	~1
491.20	~2
508.35	~1
577.75	~1
639.20	~1
696.30	~1
766.50	~1
831.50	~1
893.75	~1
902.90	100
914.70	~1
1007.50	~1
1081.85	~1
1159.05	~1
1227.70	~1
1298.95	~1
1370.90	~1
1441.05	~1
1510.80	~1
1581.10	~1
1600.35	~1
1640.05	~1
1683.85	~1
1695.70	~1
1773.85	~2
1788.58	~1
1805.20	~40
1806.60	~10
1822.40	~5
1841.65	~1
1914.55	~1

Sample Information Item

Date and Time : 3/12/2013 3:42:22 AM

User : Steffen

Type : Unknown

Inj. Volume : 20.000

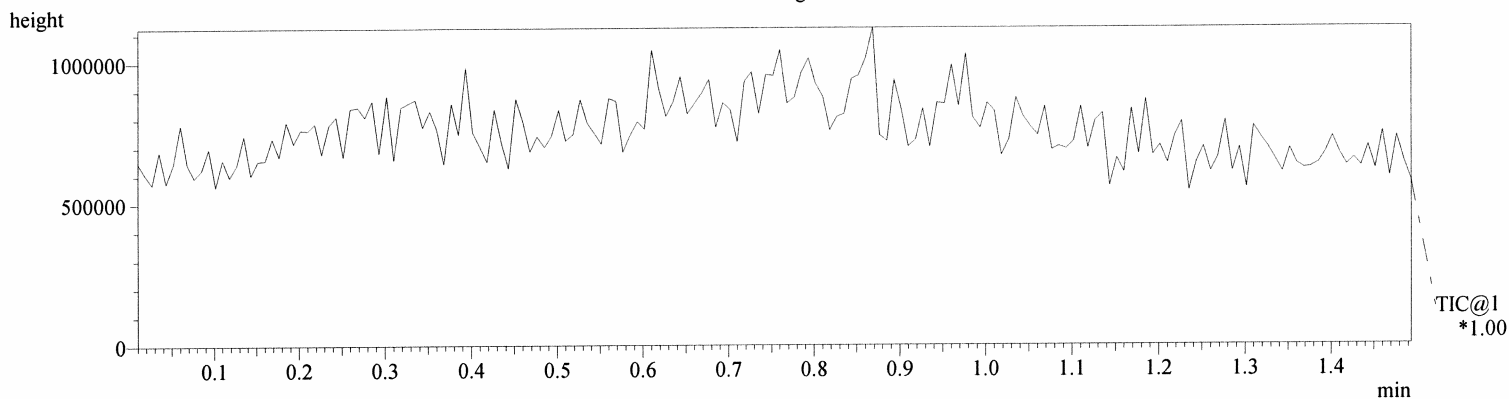
Data Name : E:\Users\Steffen\Important files\4peptidesben\amide-amide-deprotected.qld

Method Name : E:\Mass files\Method\Positive mode\200-2000 0.3ml-min flow.qlm

Tuning Name : E:\Mass files\Tuning\Autotune 20120620.QLT

Compound 11

MS Chromatogram



MS Spectrum

Line#:1 R.Time:0.393(Scan#:47) Positive

MassPeaks:821 BasePeak:874.35(36431)

RawMode:Averaged 0.200-0.560(24-67)

BG Mode:Averaged 0.936-1.449(112-174)

