

## Supplementary material

# Oxygenated cembrene diterpenes from *Sarcophyton convolutum*: cytotoxic sarcoconvolutum A-E

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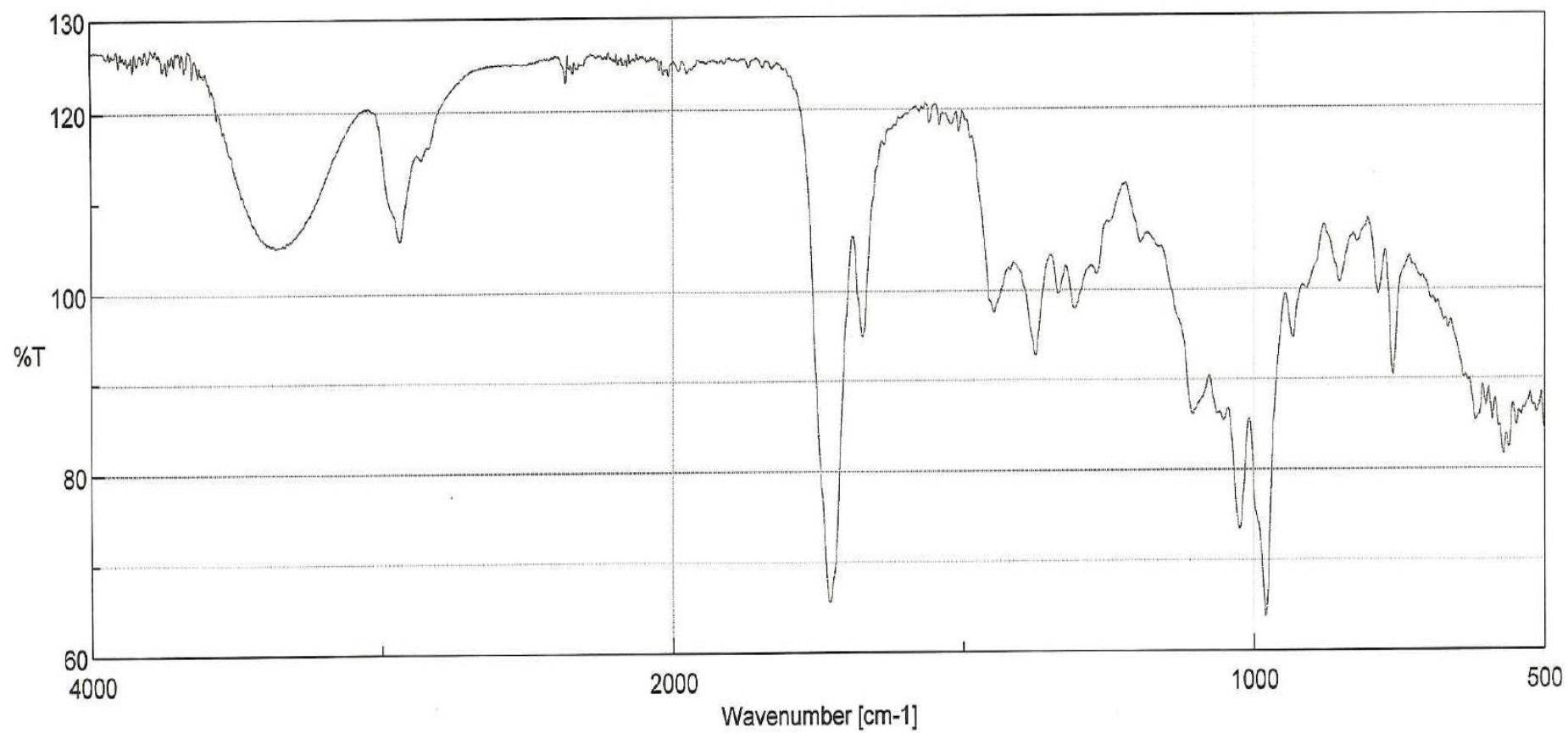
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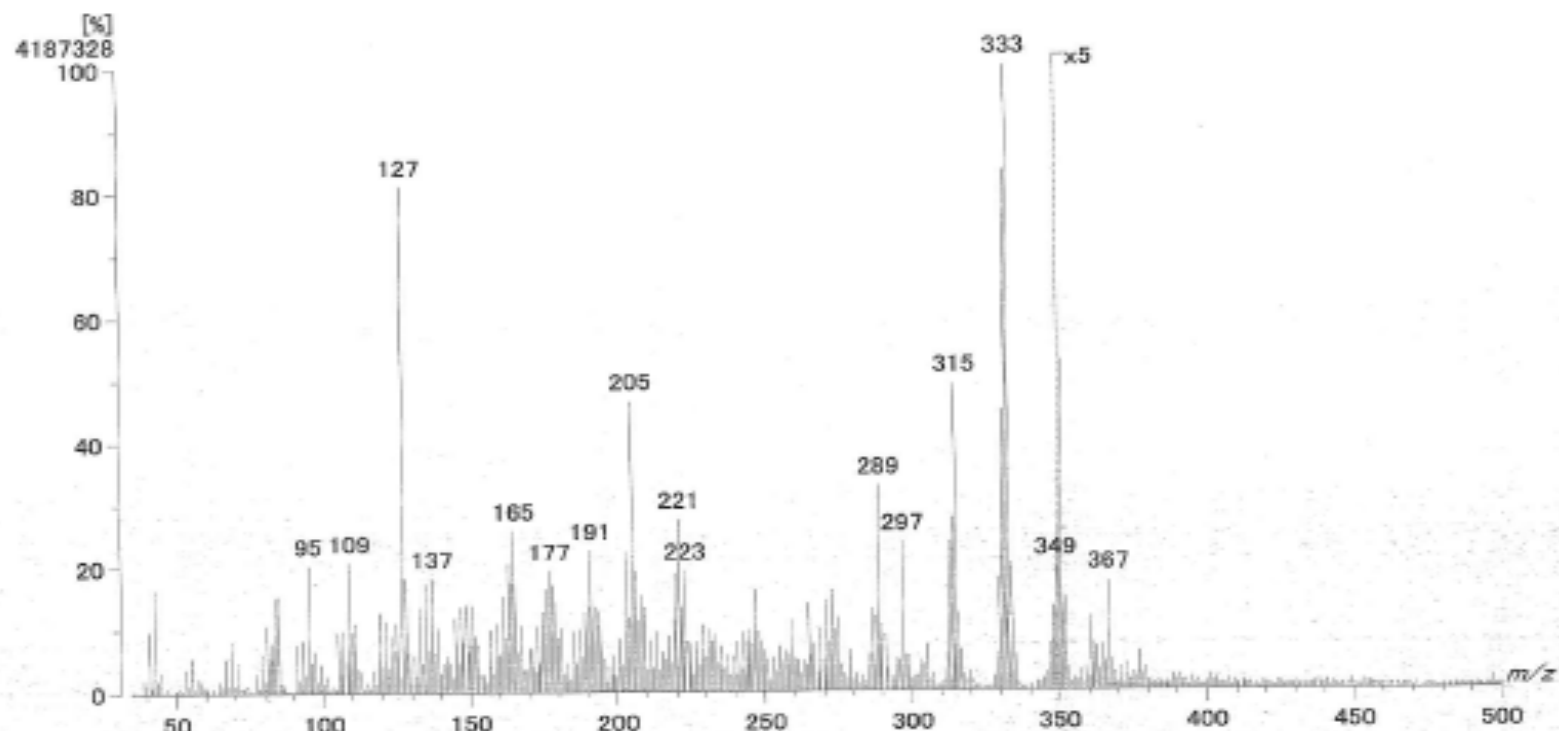
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S1: FTIR of **1**

Inlet : Direct Ion Mode : C<sup>+</sup>  
Spectrum Type : Normal Ion [MF-Linear]  
RT : 2.24 min Scan# : 83  
BP : m/z 333 Int. : 399.34 (4187328)  
Output m/z range : 35 to 500 Cut Level : 0.00 %



S2: LRCIMS of 1

Note : MStation

Inlet : Direct      Ion Mode : CI+

RT : 1.92 min      Scan# : 49

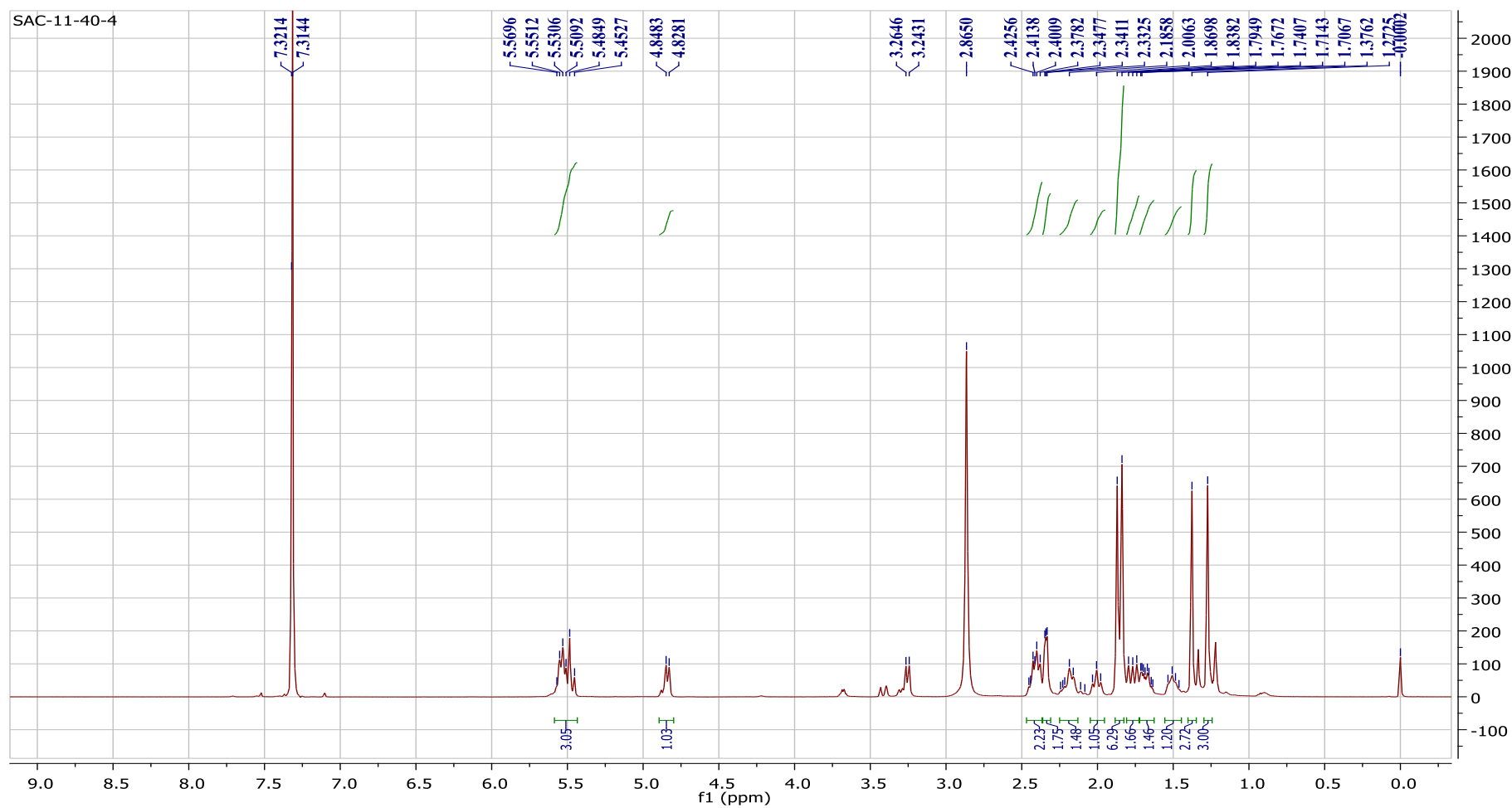
Elements : C 150/0, H 250/0, O 50/0

Mass Tolerance : 5mmu

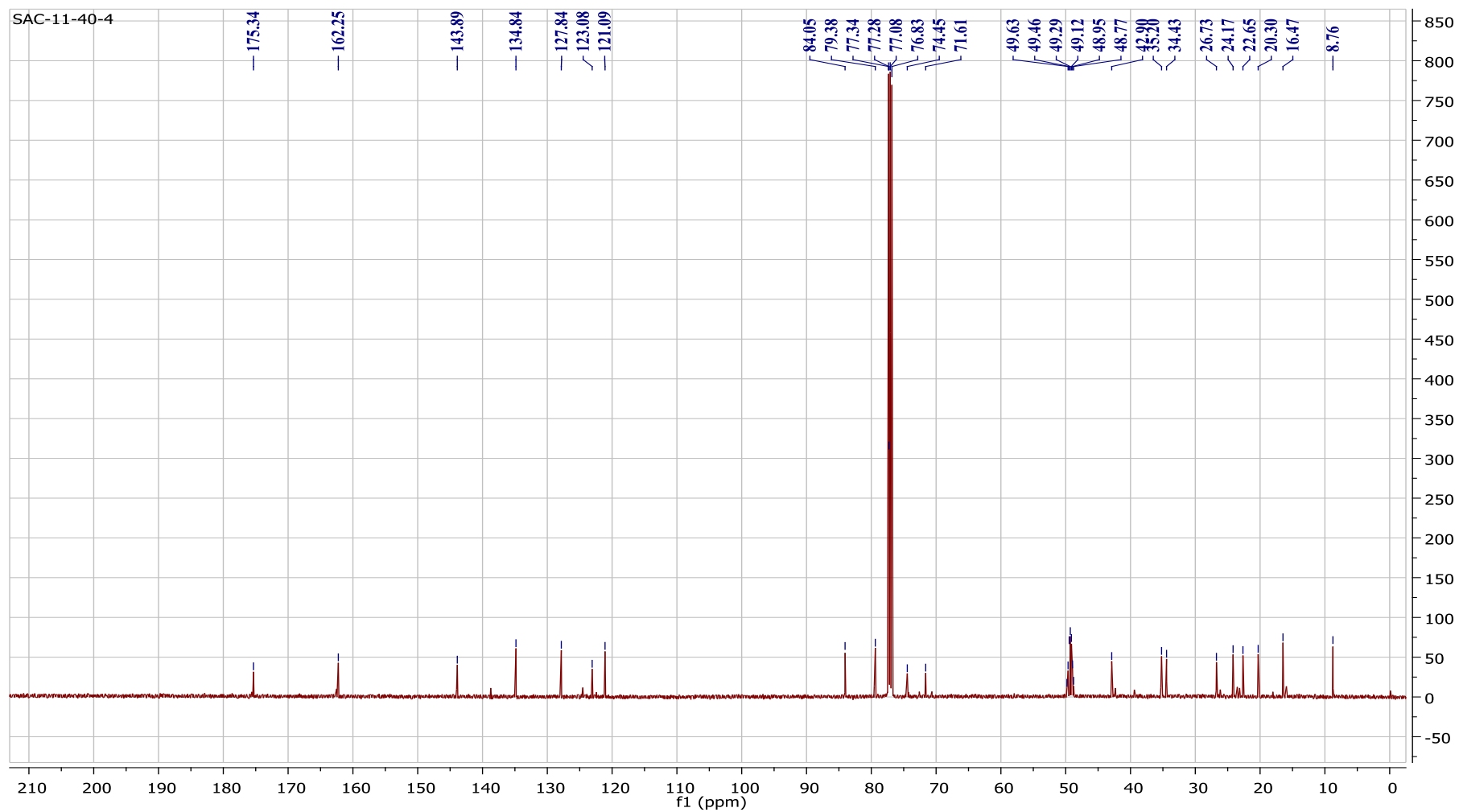
Unsaturation (U.S.) : 0.0 – 15.0

	Observed m/z	Int%	Err [ppm / mmu]	U.S. Composition
1	367.2122	4.78	+0.4 / +0.1	5.5 C20 H31 O6

S3: HRCIMS of 1

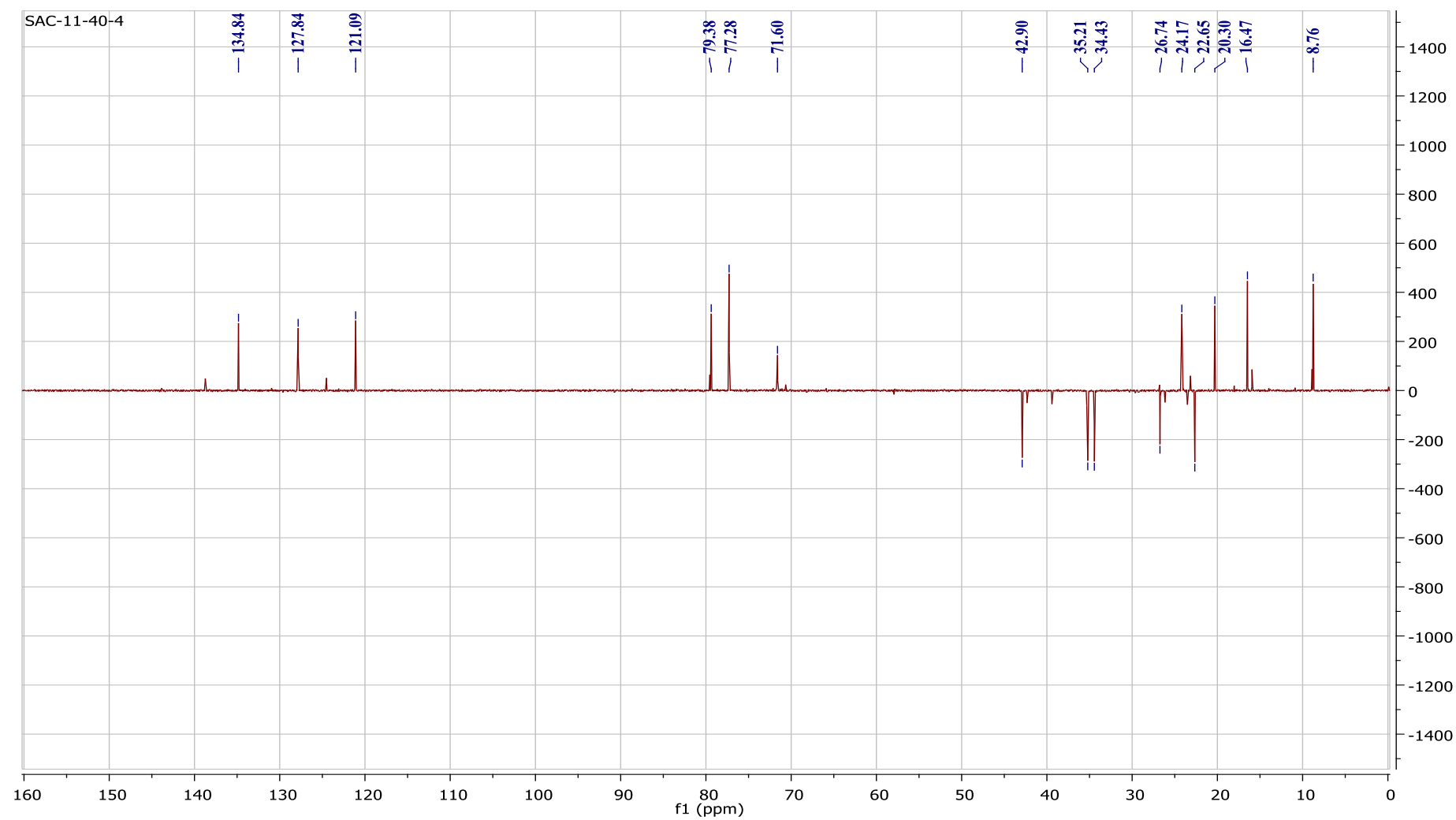


S4:  $^1\text{H}$  NMR of **1** (500 Hz,  $\text{CDCl}_3$ )

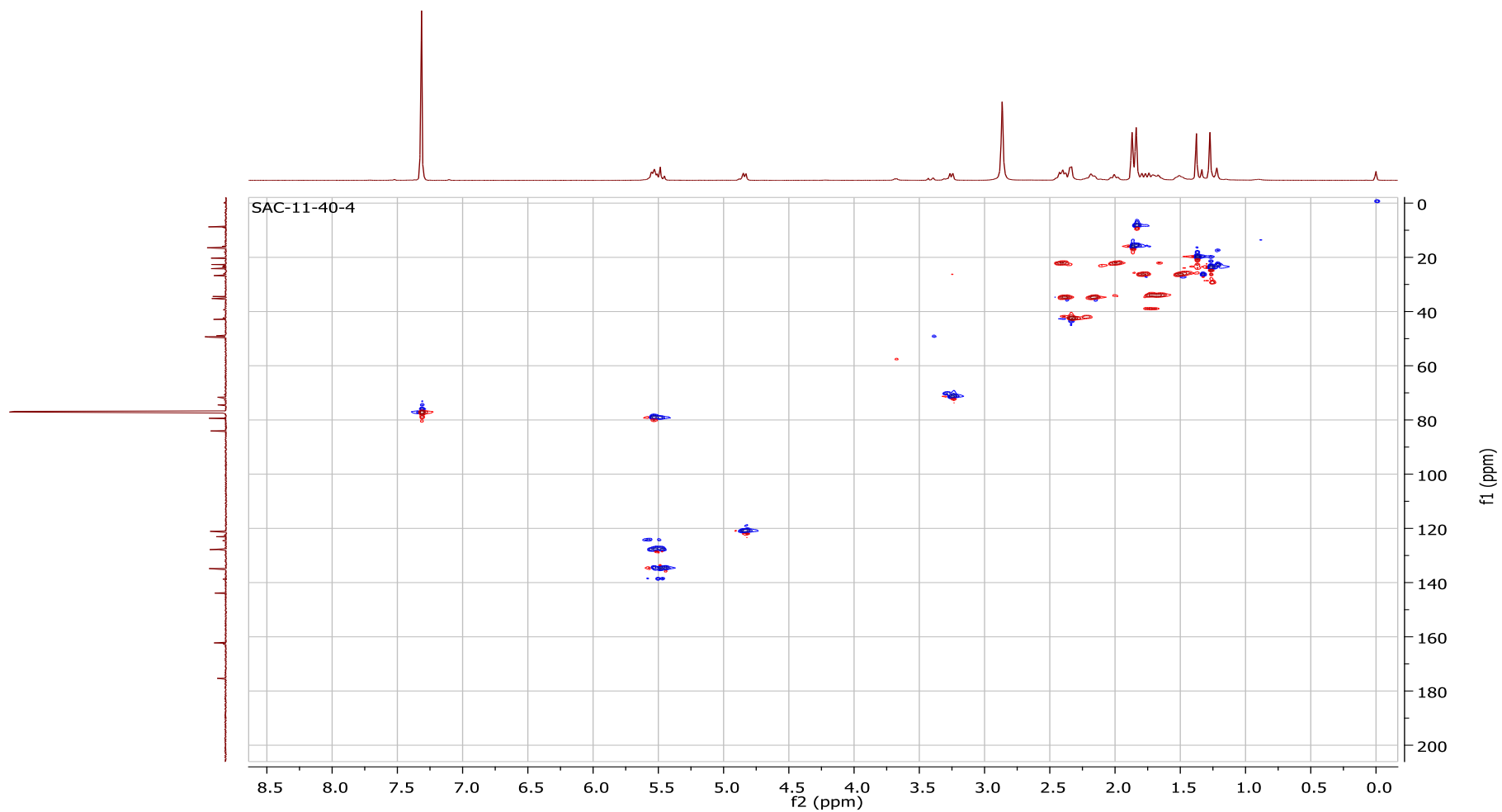


S5:  $^{13}\text{C}$  NMR of **1** (125 Hz,  $\text{CDCl}_3$ )

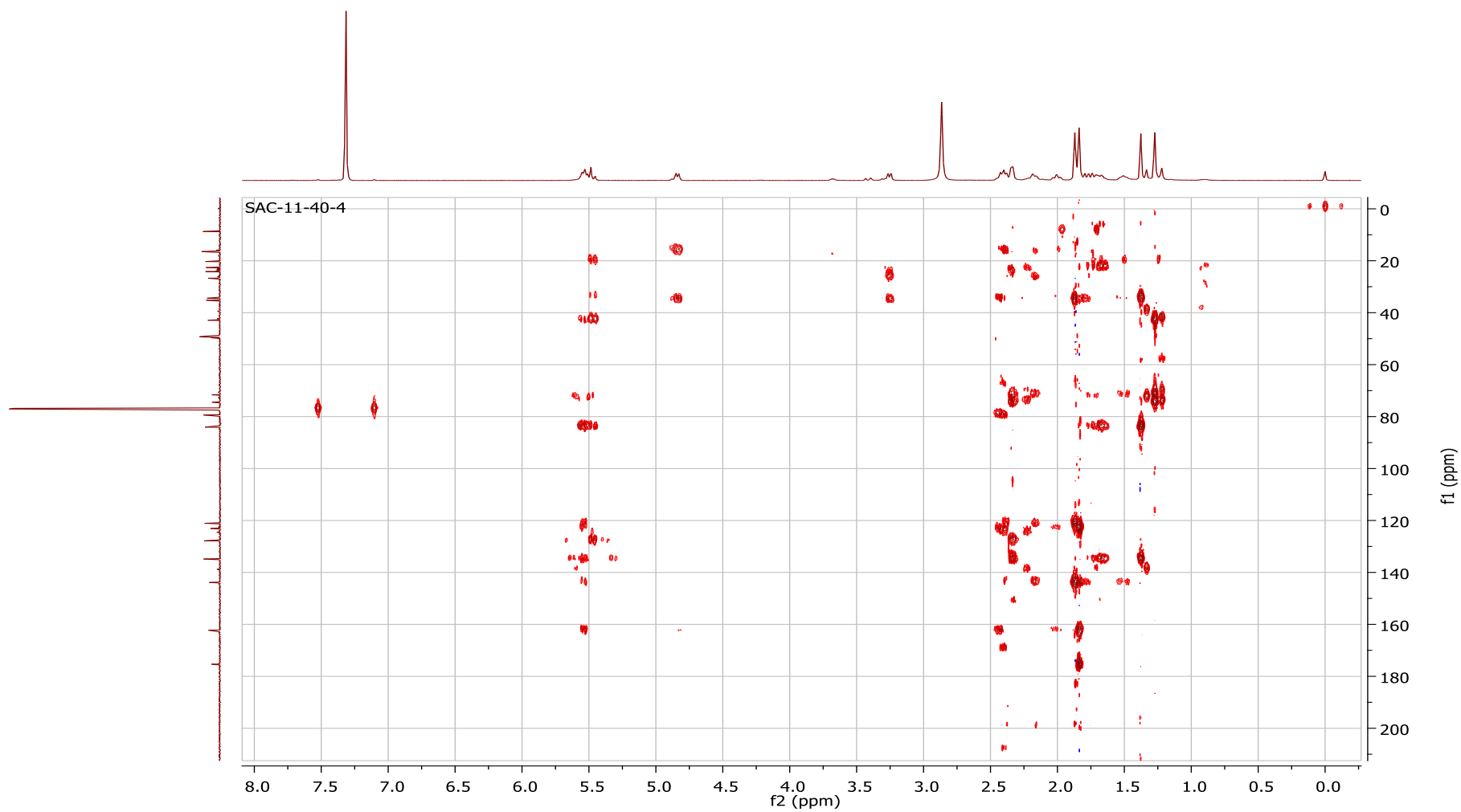




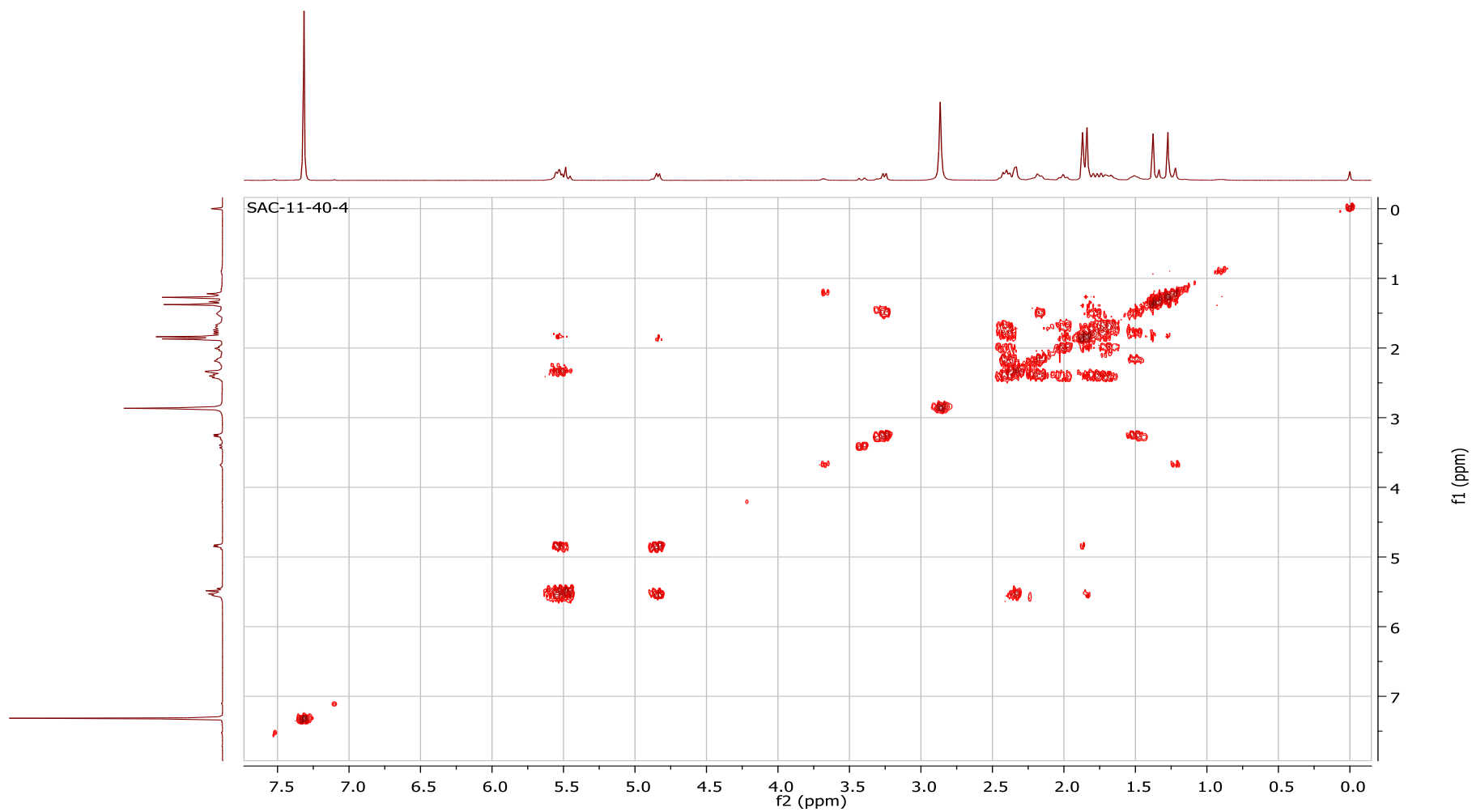
S6: DEPT of **1**



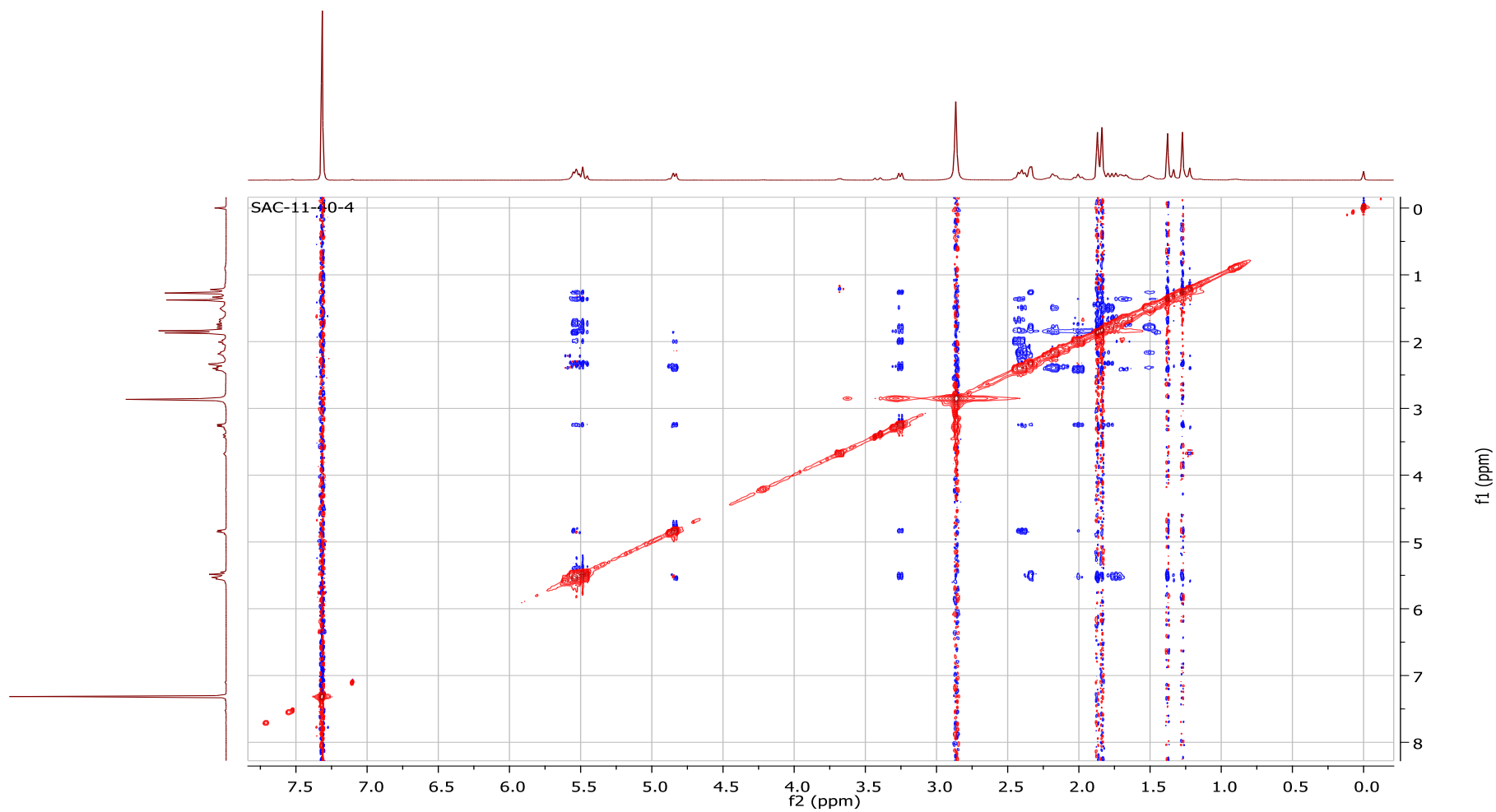
S7: HSQC of **1**



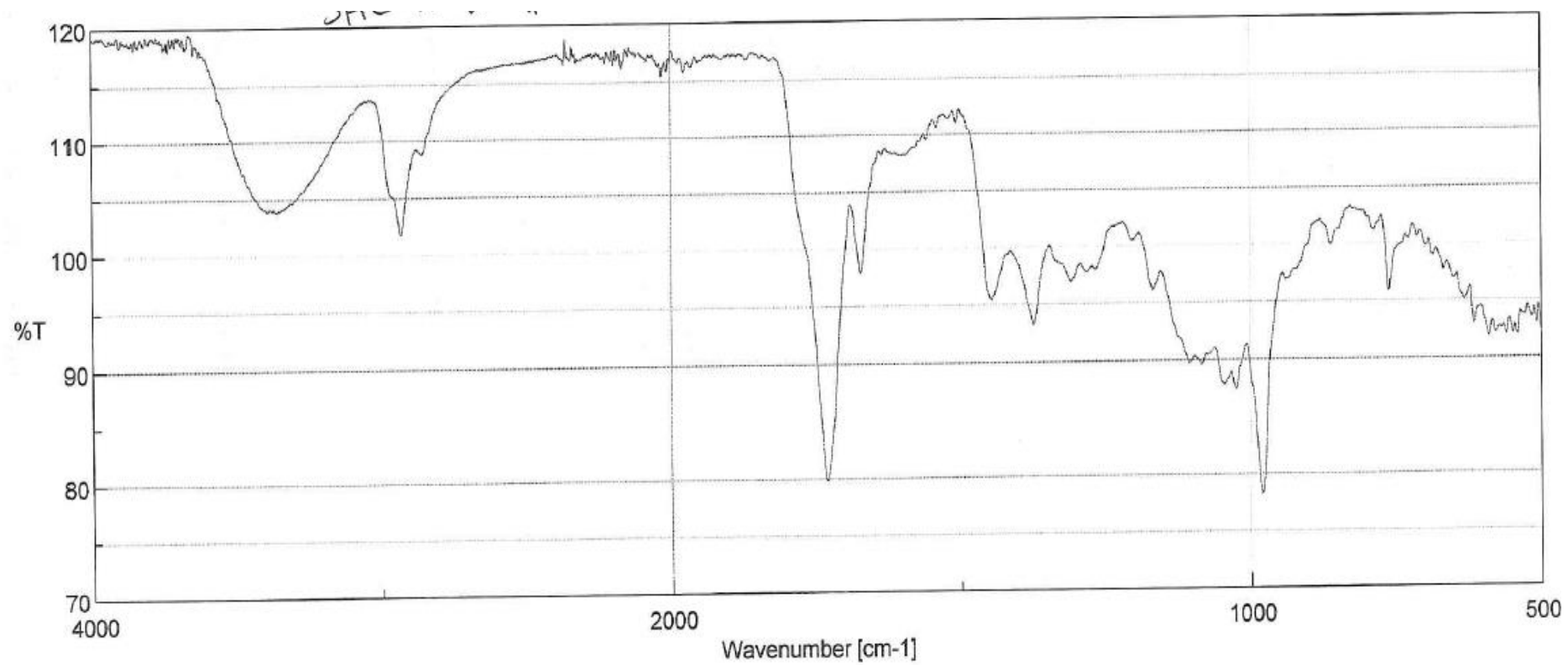
S8: HMBC of **1**



S9:  $^1\text{H}$   $^1\text{H}$  COSY of 1

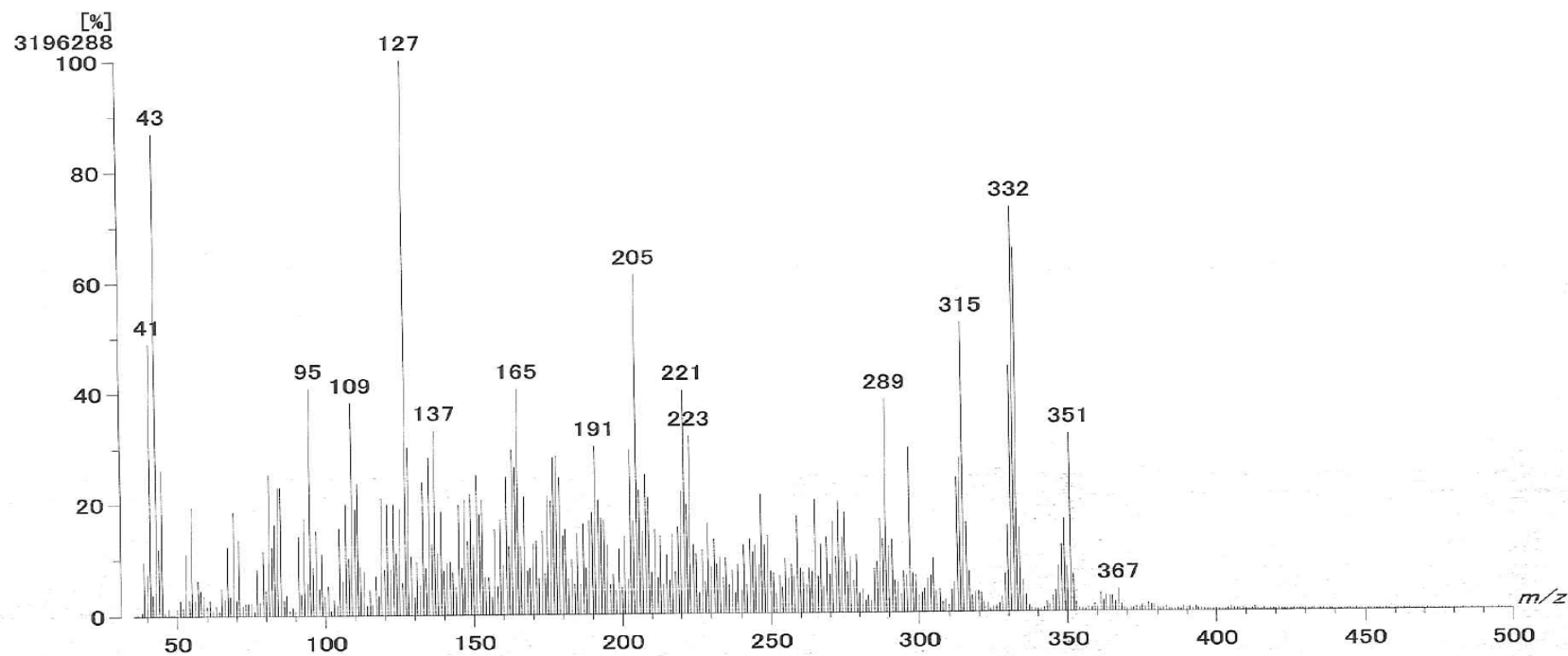


S10: NOESY of **1**



S11: FTIR of 2

Inlet : Direct      Ion Mode : CI+  
Spectrum Type : Normal Ion [MF-Linear]  
RT : 1.78 min      Scan# : 66  
BP : m/z 127      Int. : 304.82 (3196288)  
Output m/z range : 35 to 500      Cut Level : 0.00 %



S12: LRCIMS of 2

Note : MStation

Inlet : Direct Ion Mode : CI+

RT : 1.92 min Scan# : 49

Elements : C 150/0, H 250/0, O 50/0

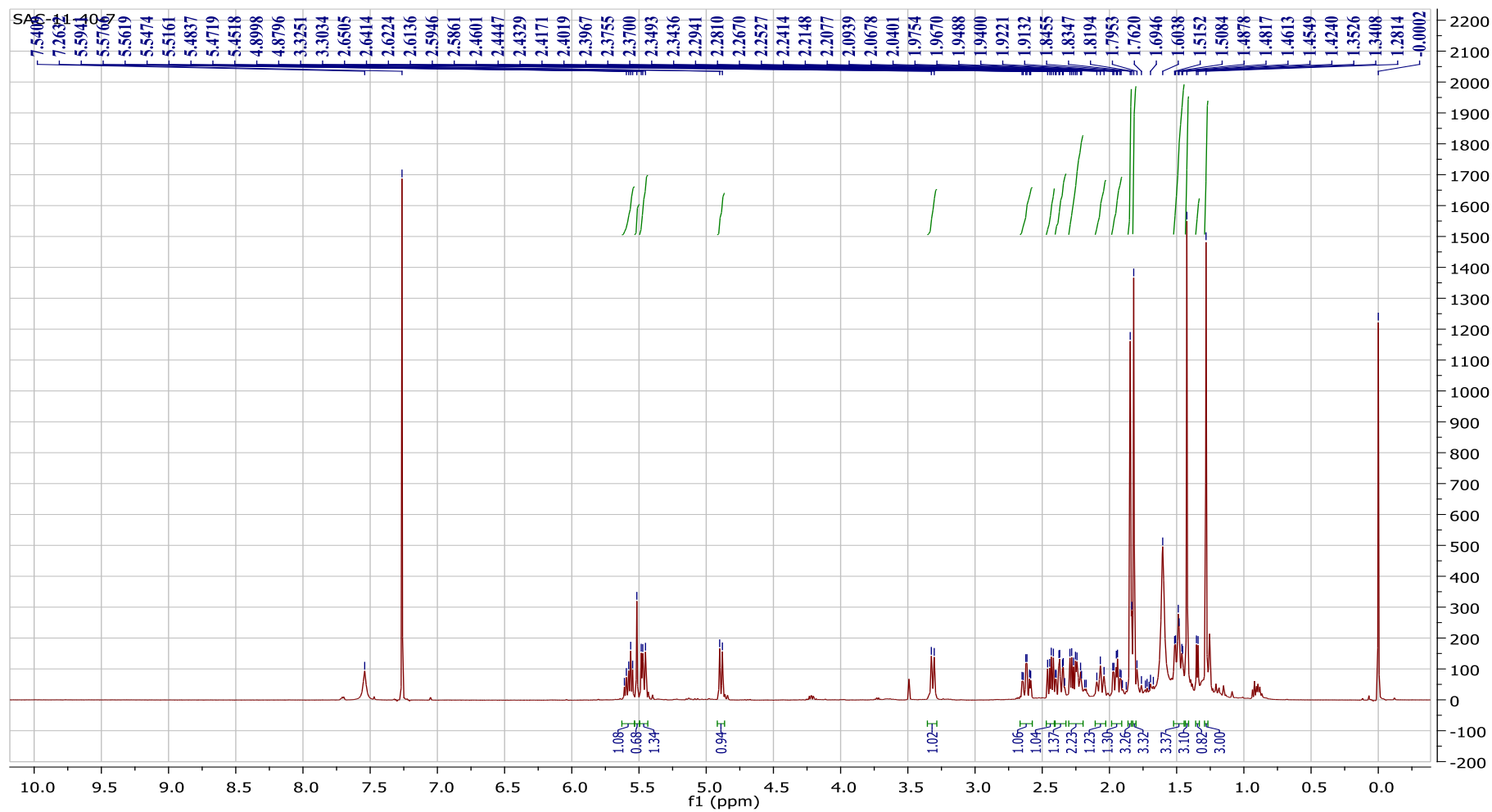
Mass Tolerance : 5mmu

Unsaturation (U.S.) : 0.0 – 15.0

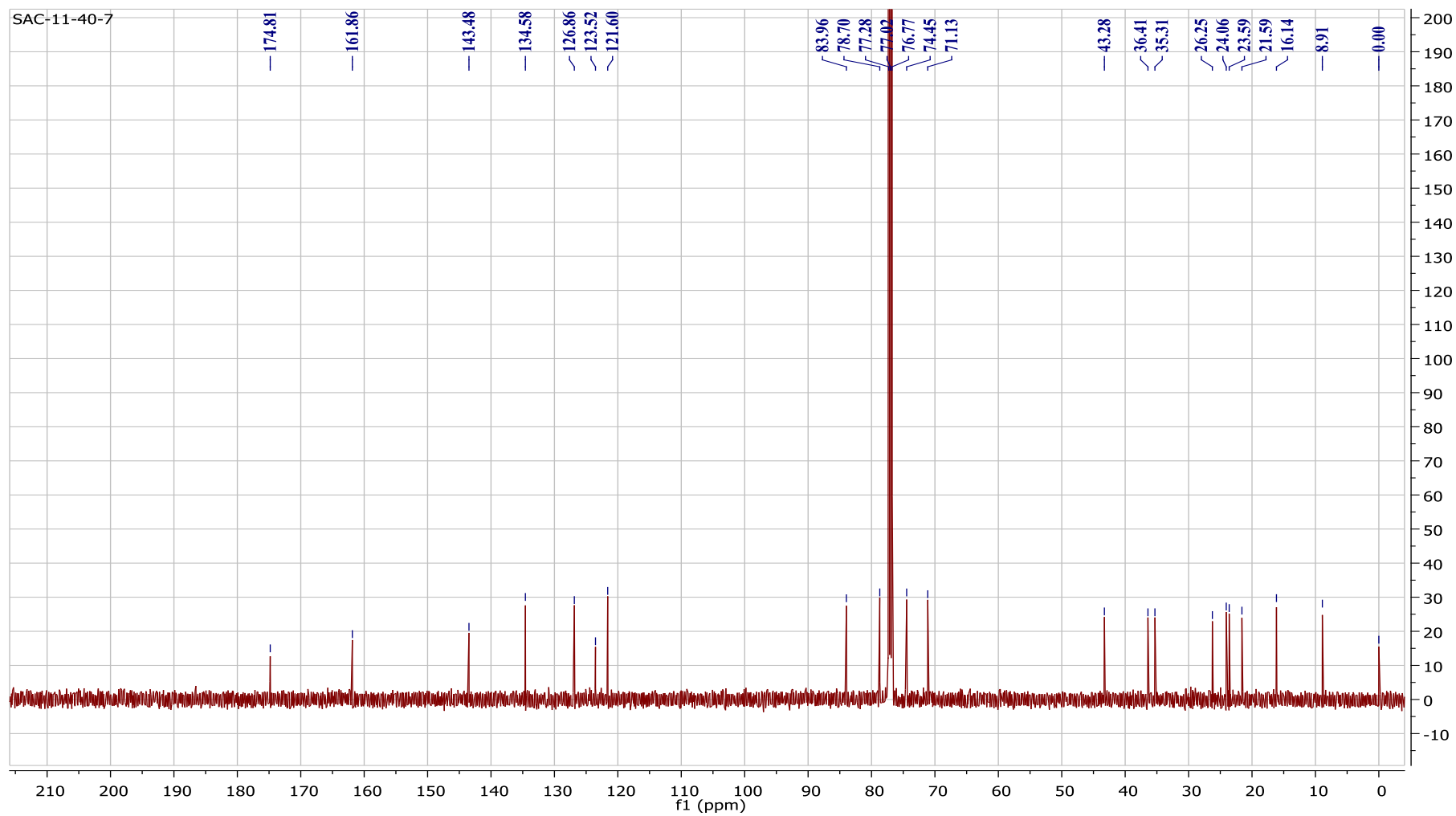
	Observed m/z	Int%	Err [ppm / mmu]	U.S. Composition
1	367.2122	4.78	+0.4 / +0.1	5.5 C20 H31 O6

S13: HRCIMS of 2

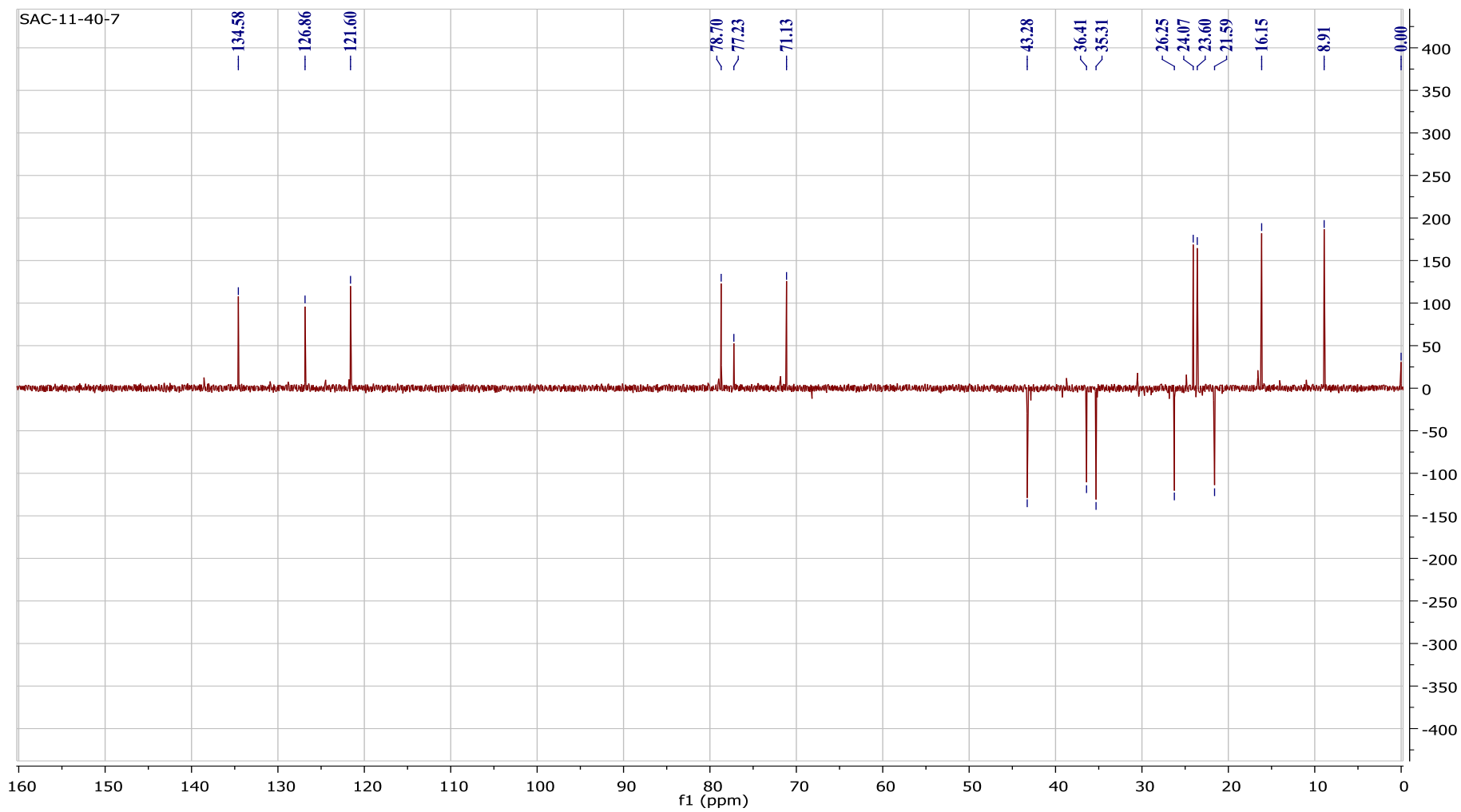




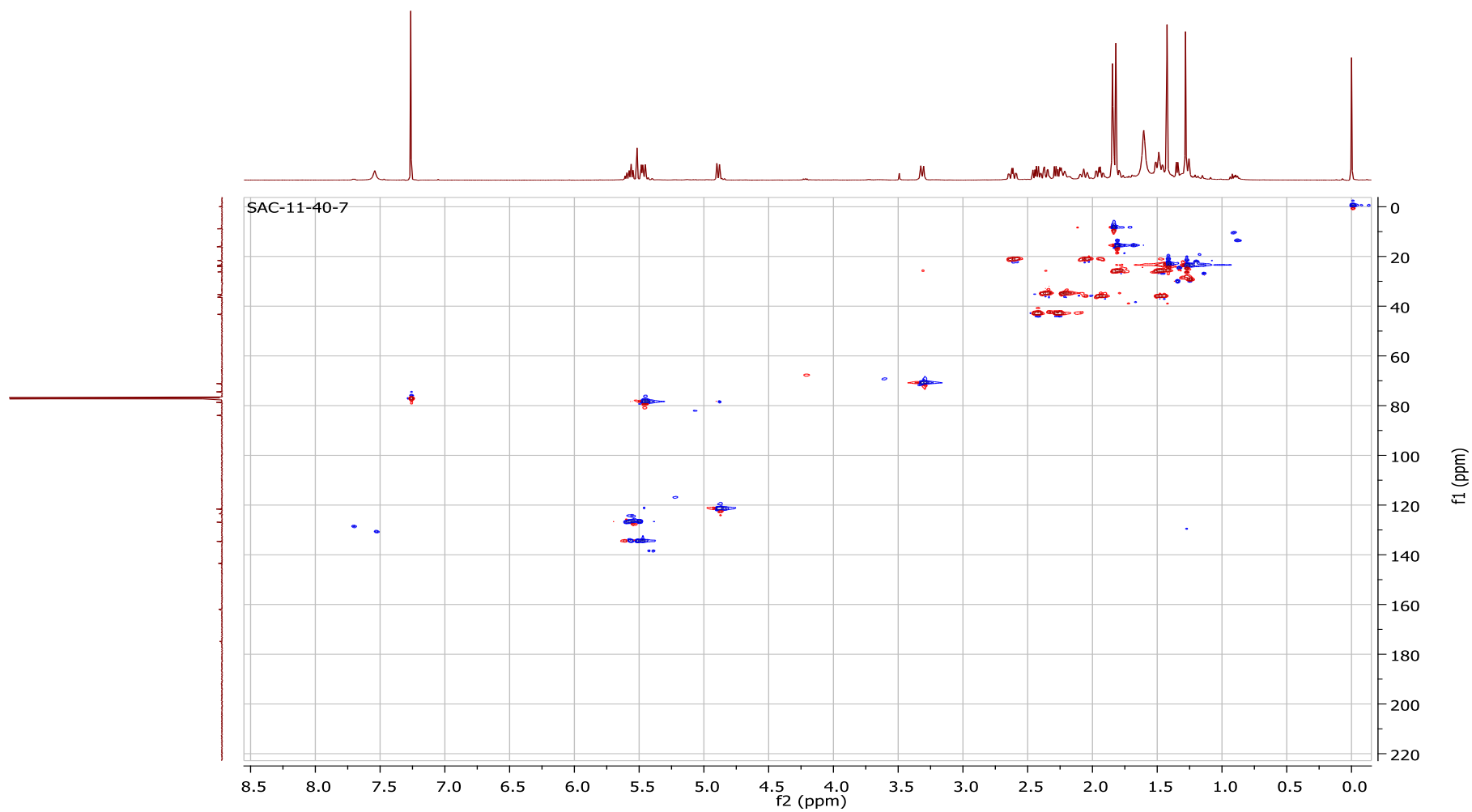
S14:  $^1\text{H}$  NMR of **2**



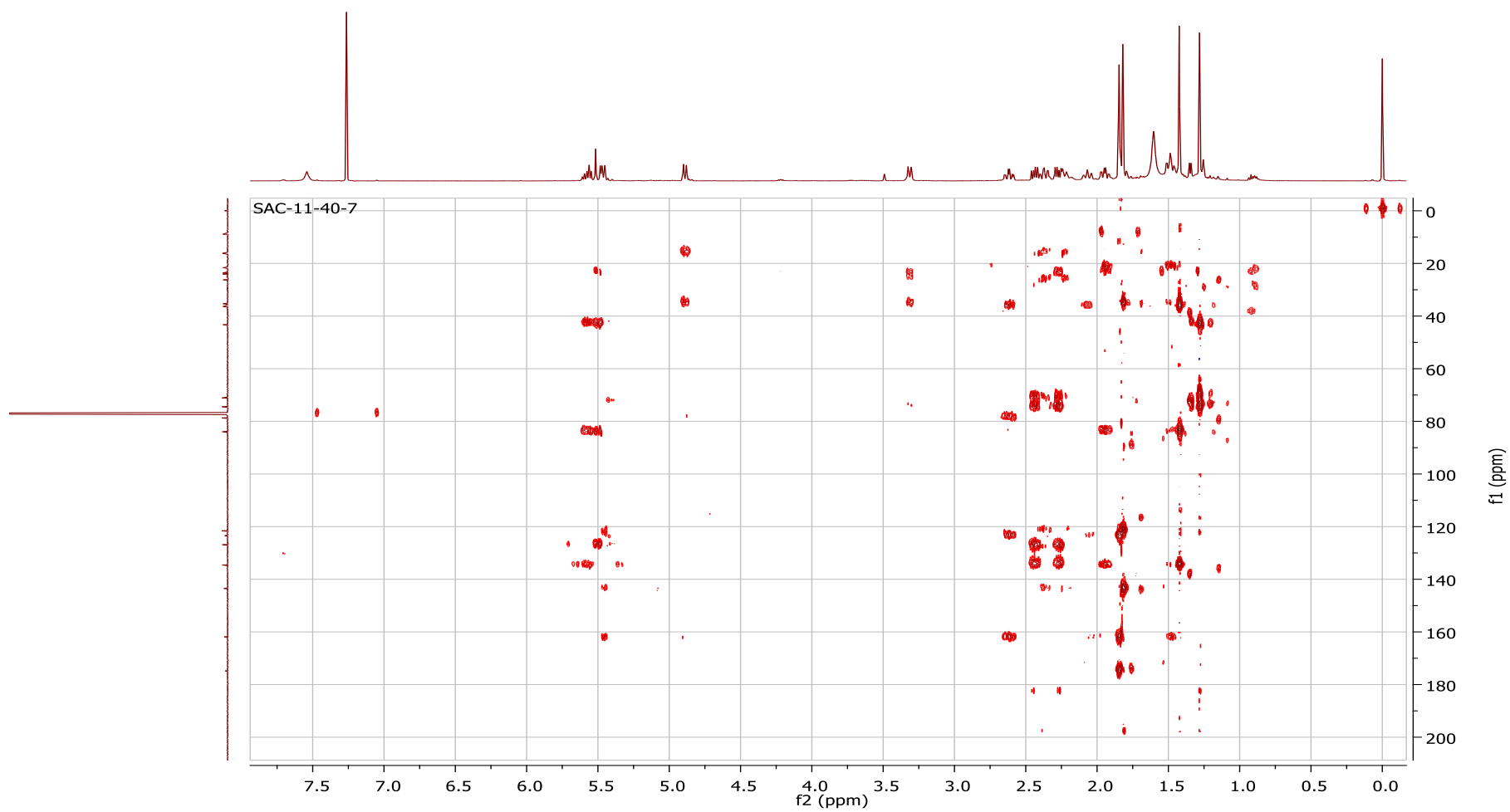
S15:  $^{13}\text{C}$  NMR of **2**



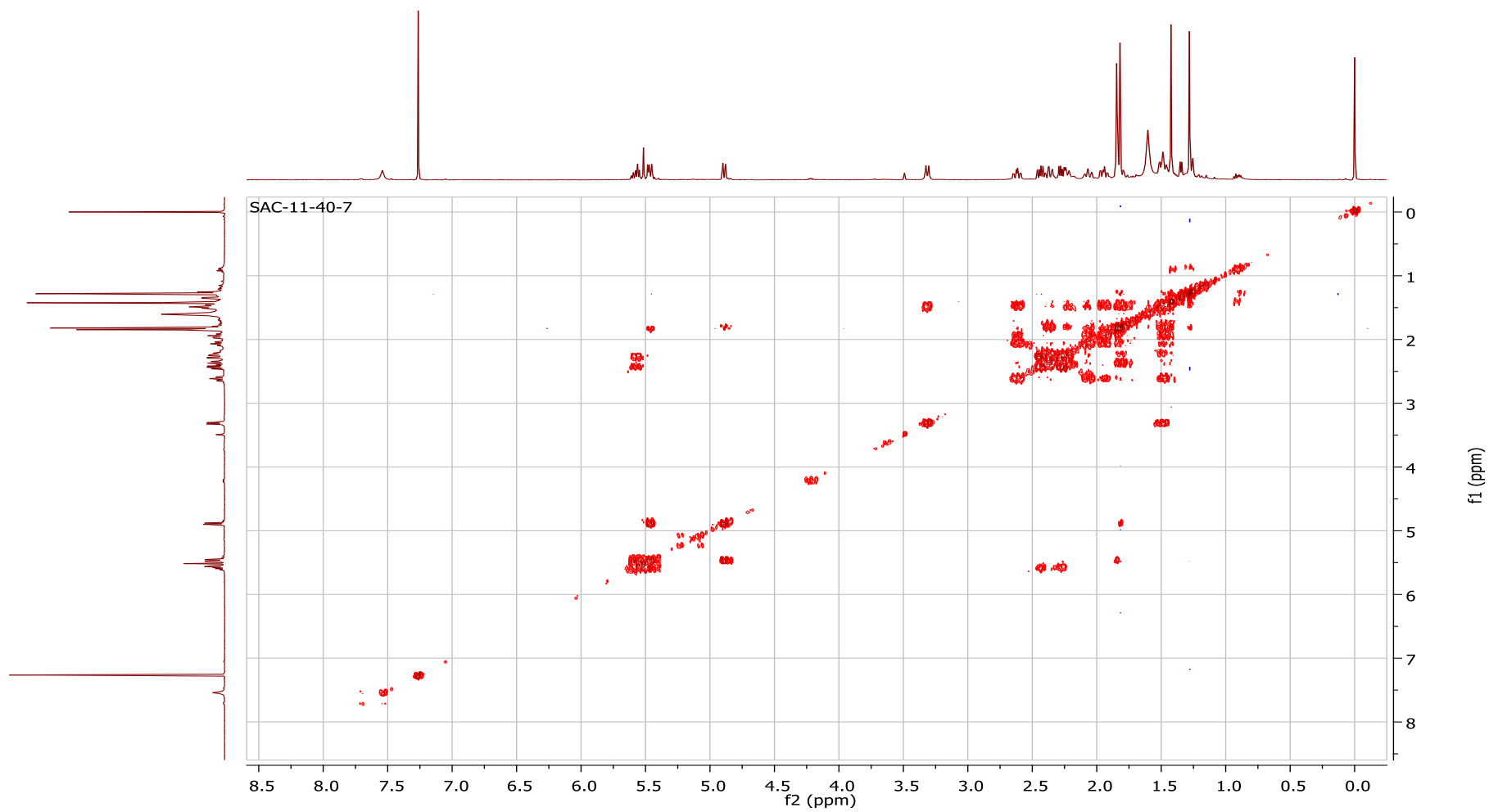
S16: DEPT of 2



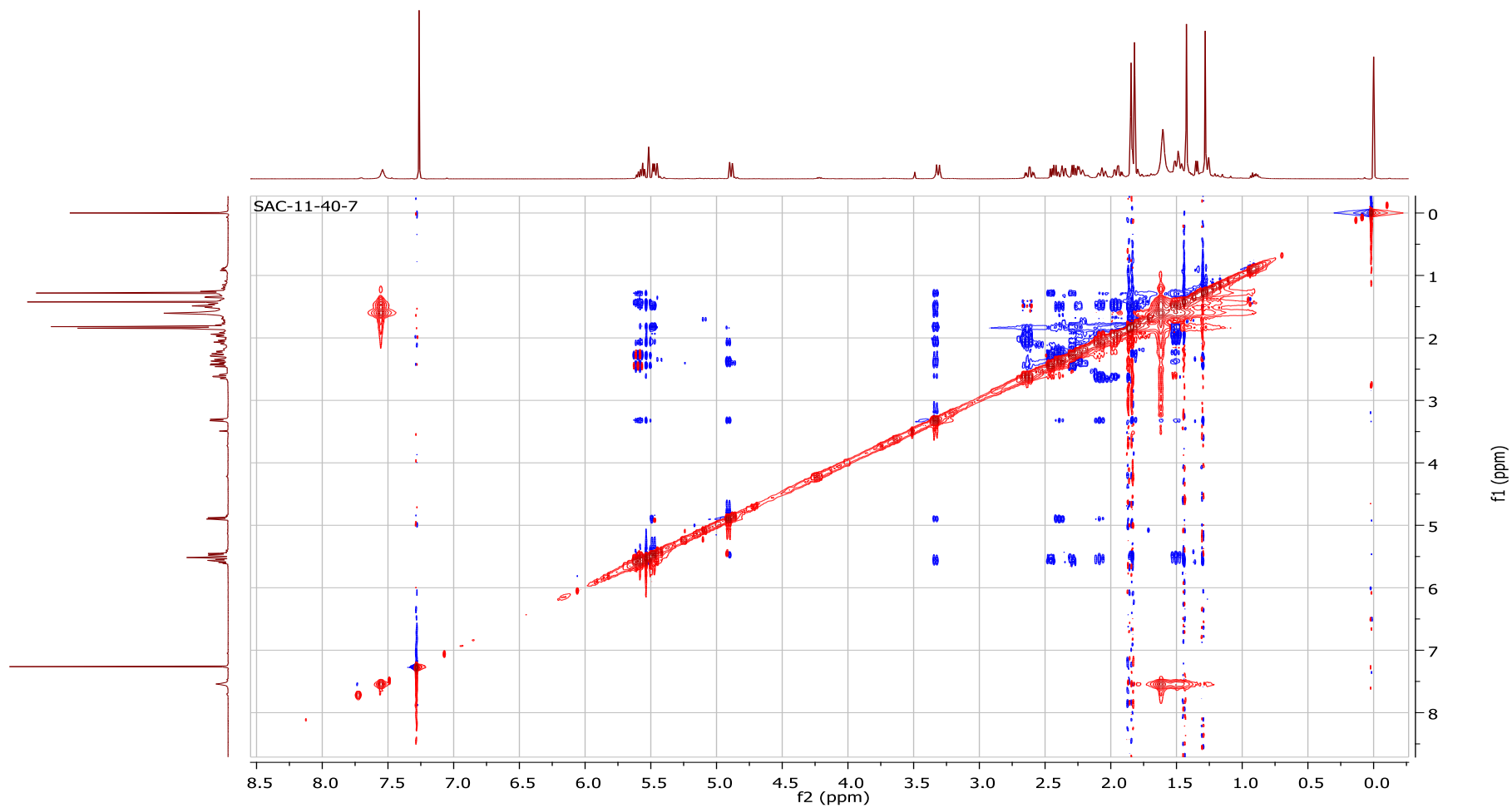
S17: HSQC of 2



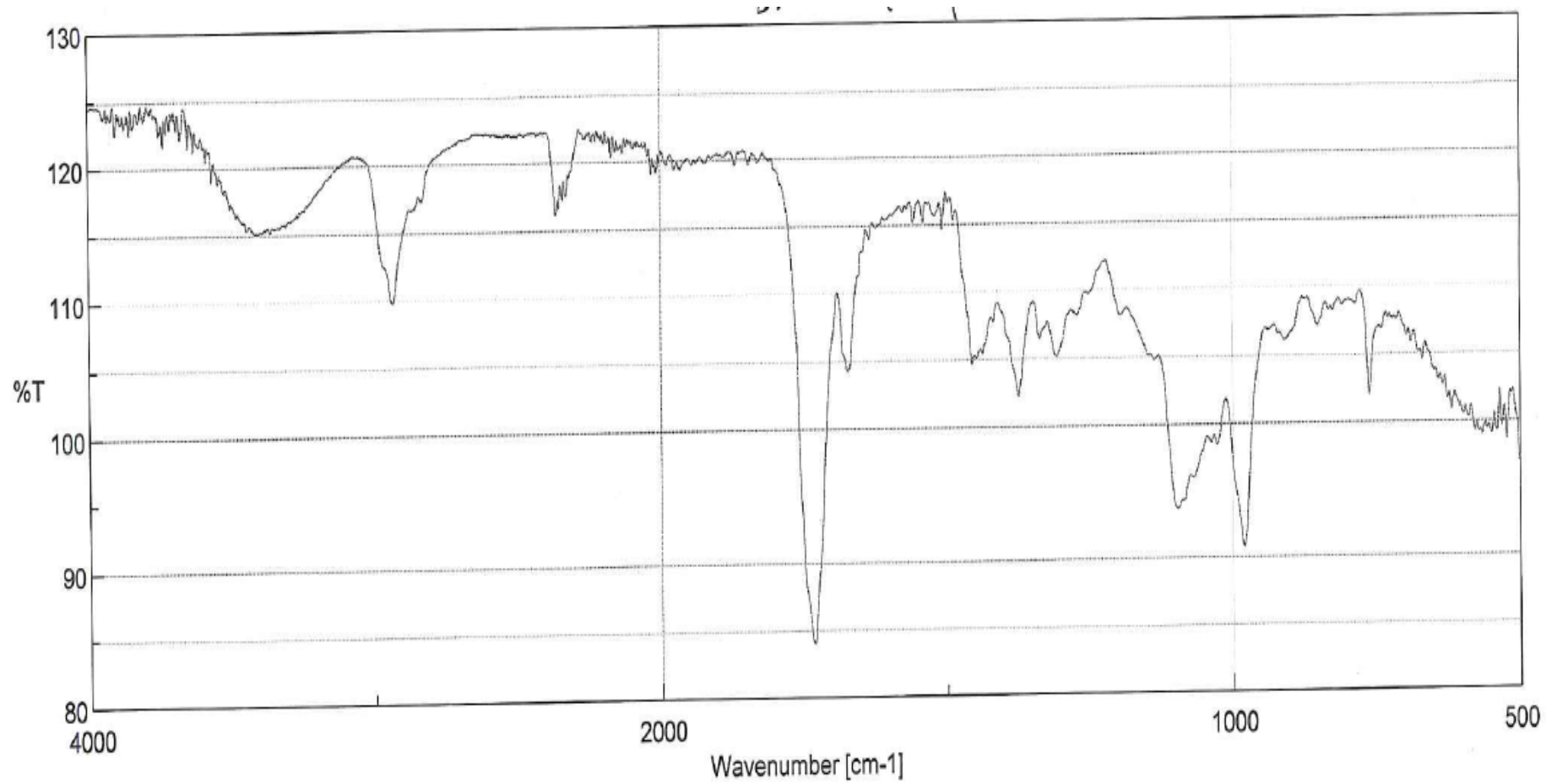
S18: HMBC of **2**



S19:  $^1\text{H}$   $^1\text{H}$  COSY of **2**



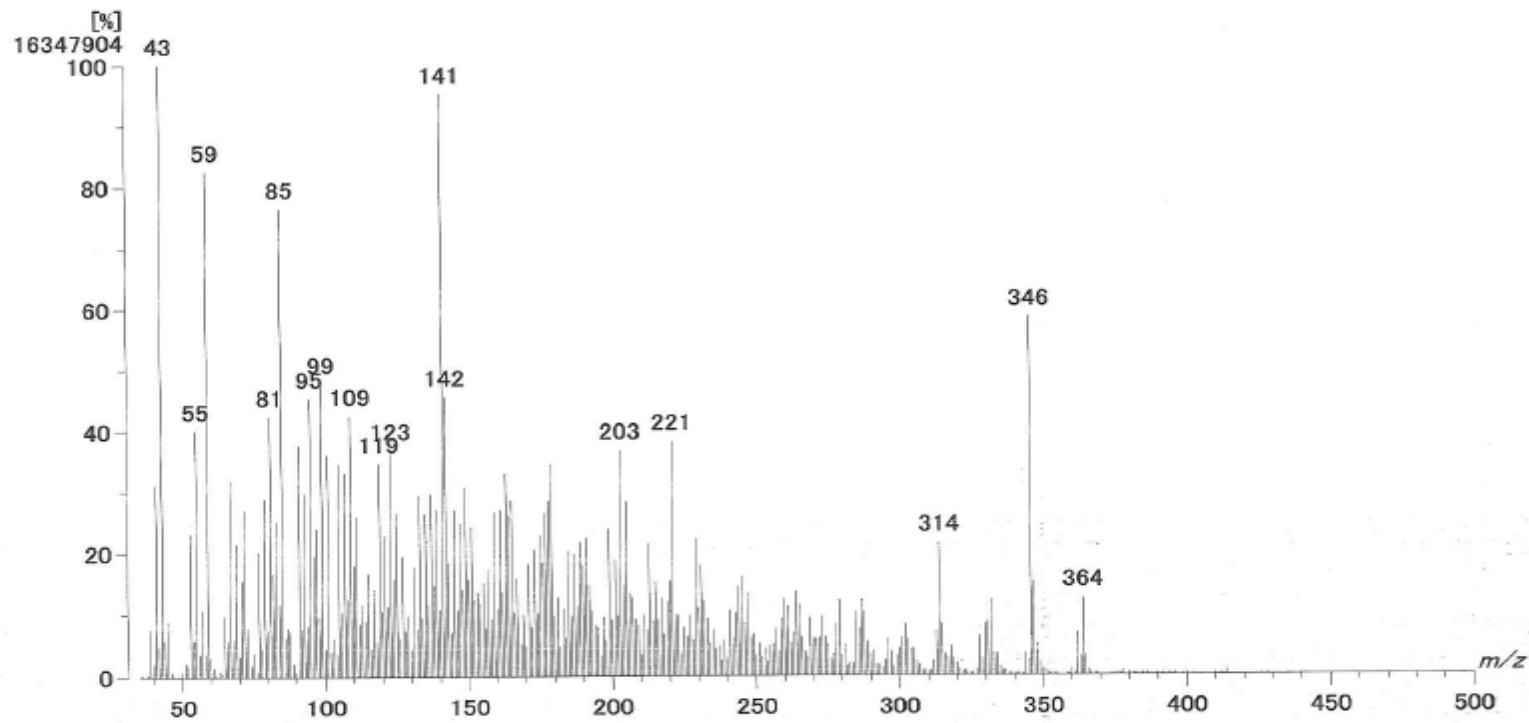
S20: NOESY of 2



S21: LRCIMS of 3



Inlet : Direct     Ion Mode : EI+  
Spectrum Type : Normal Ion [MF-Linear]  
RT : 1.07 min     Scan# : 33  
BP : m/z 43     Int. : 1559.06 (16347904)  
Output m/z range : 35 to 500     Cut Level : 0.00 %

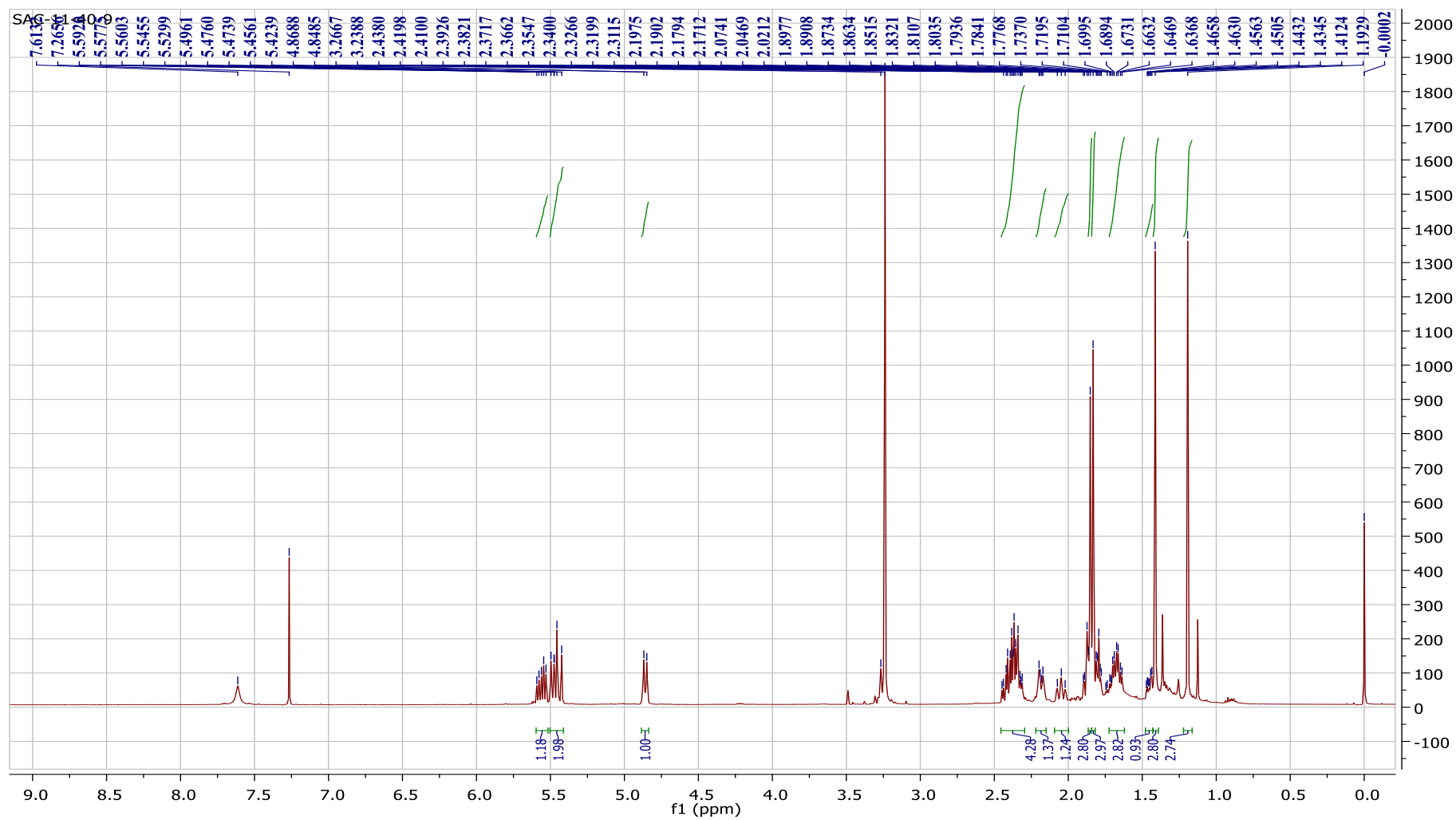


S22: LREIMS of 3

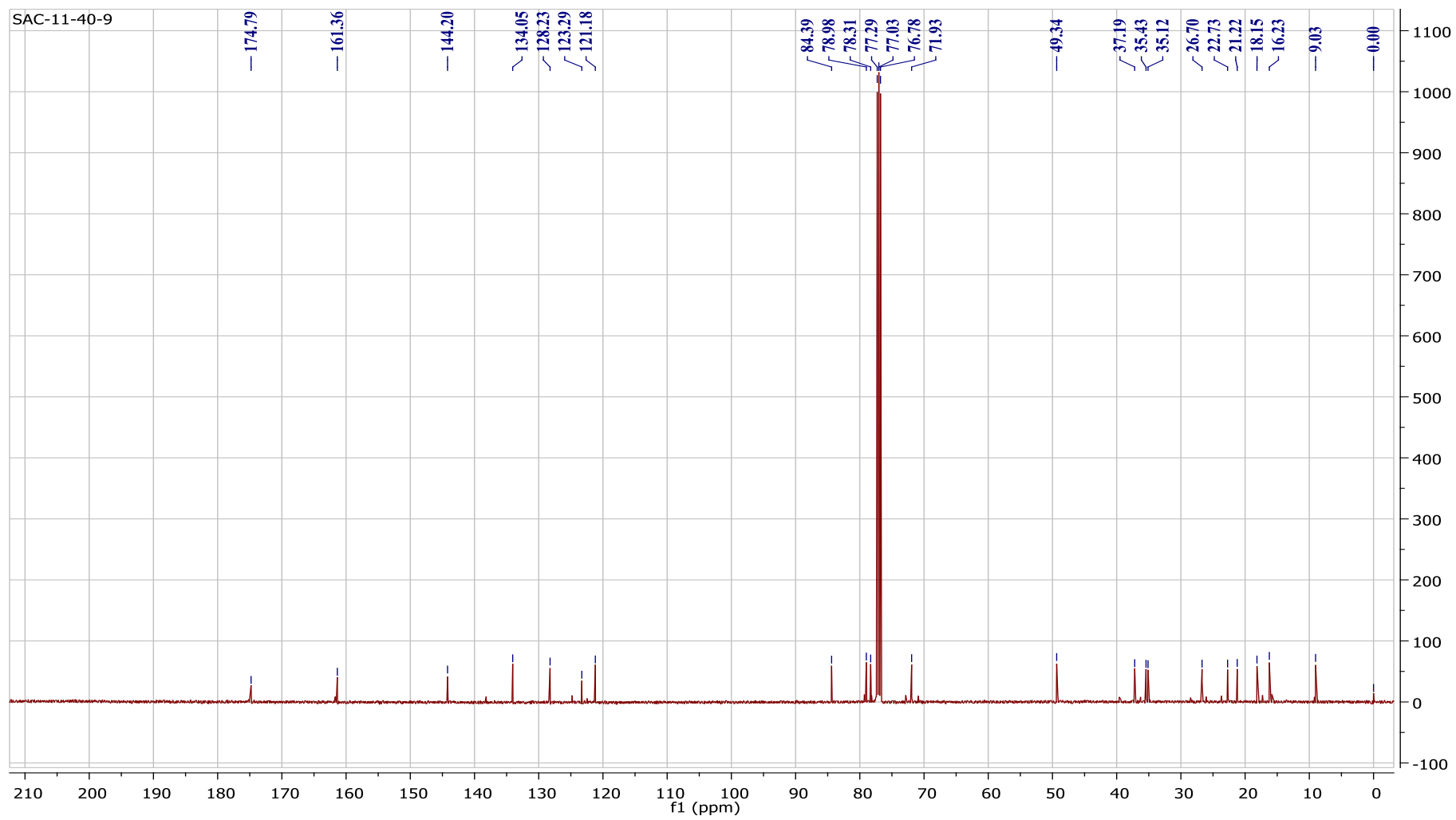
Inlet : Direct      Ion Mode : EI+  
RT : 1.48 min      Scan# : 38  
Elements : C 150/0, H 250/0, O 50/0  
Mass Tolerance : 5mmu  
Unsaturation (U.S.) : 0.0 – 20.0

	Observed m/z	Int%	Err [ppm / mmu]	U.S. Composition
1	364.2258	21.08	+2.3 / +0.8	6.0 C21 H32 O5

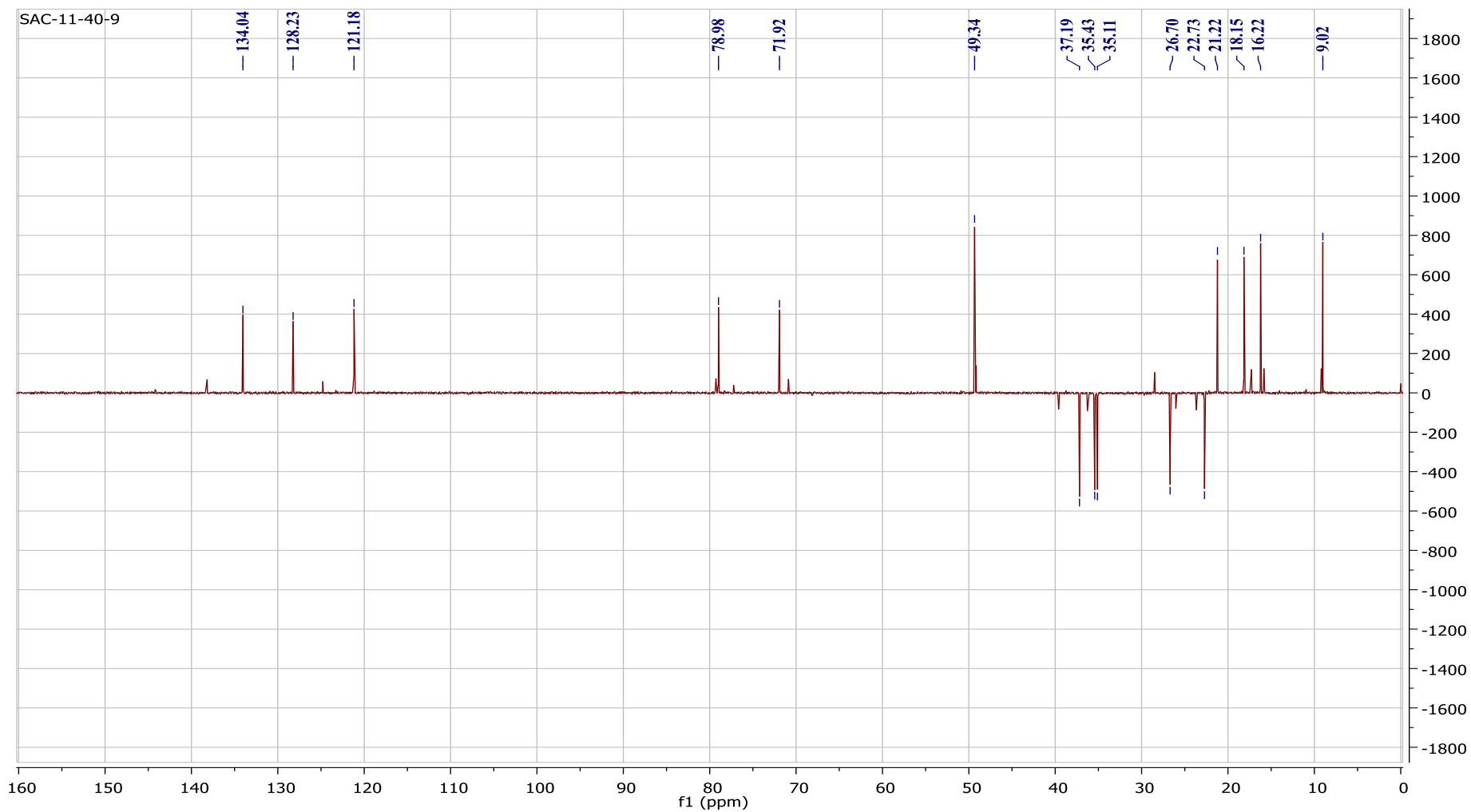
S23: HREIMS of 3



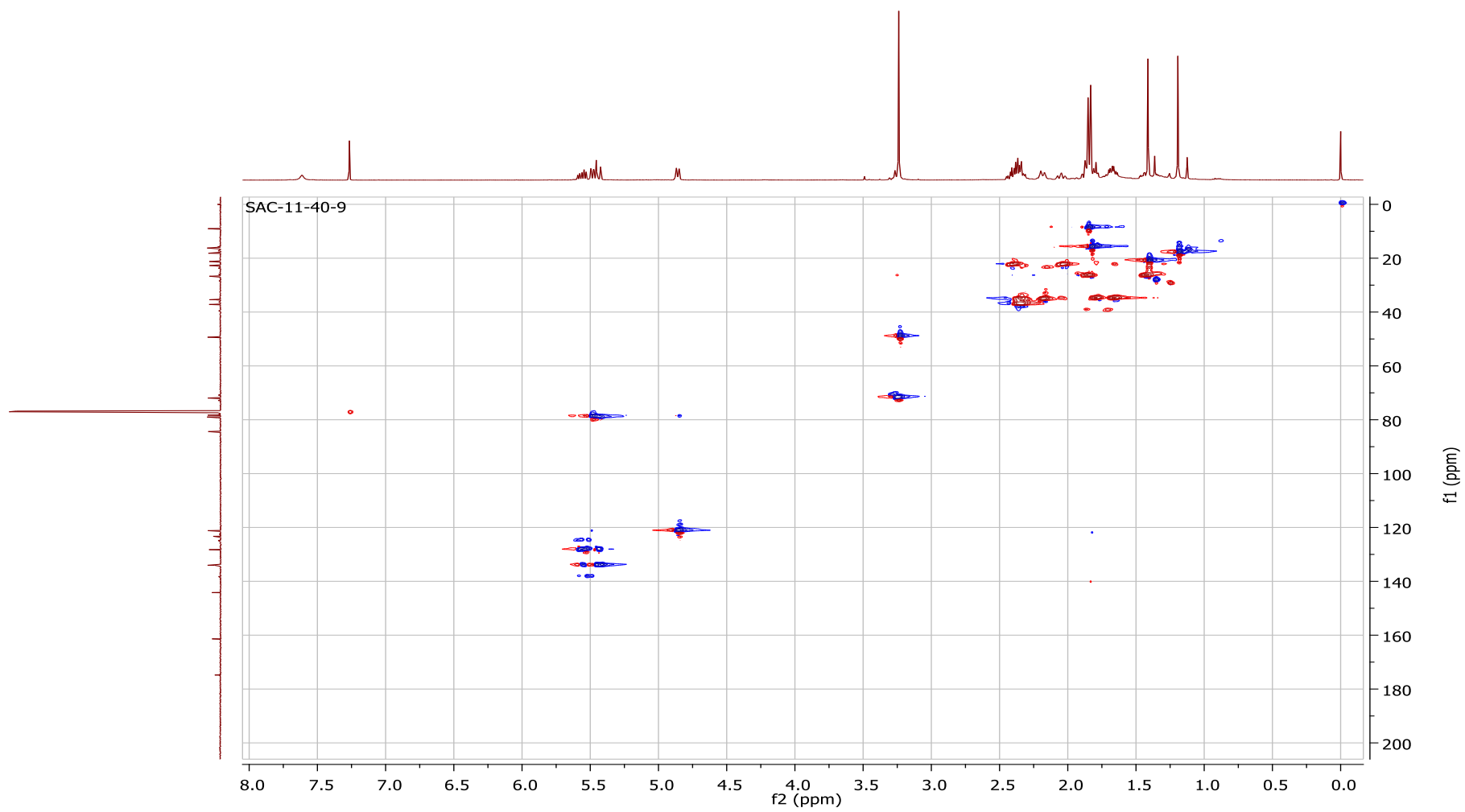
S24:  $^1\text{H}$  NMR of **3**



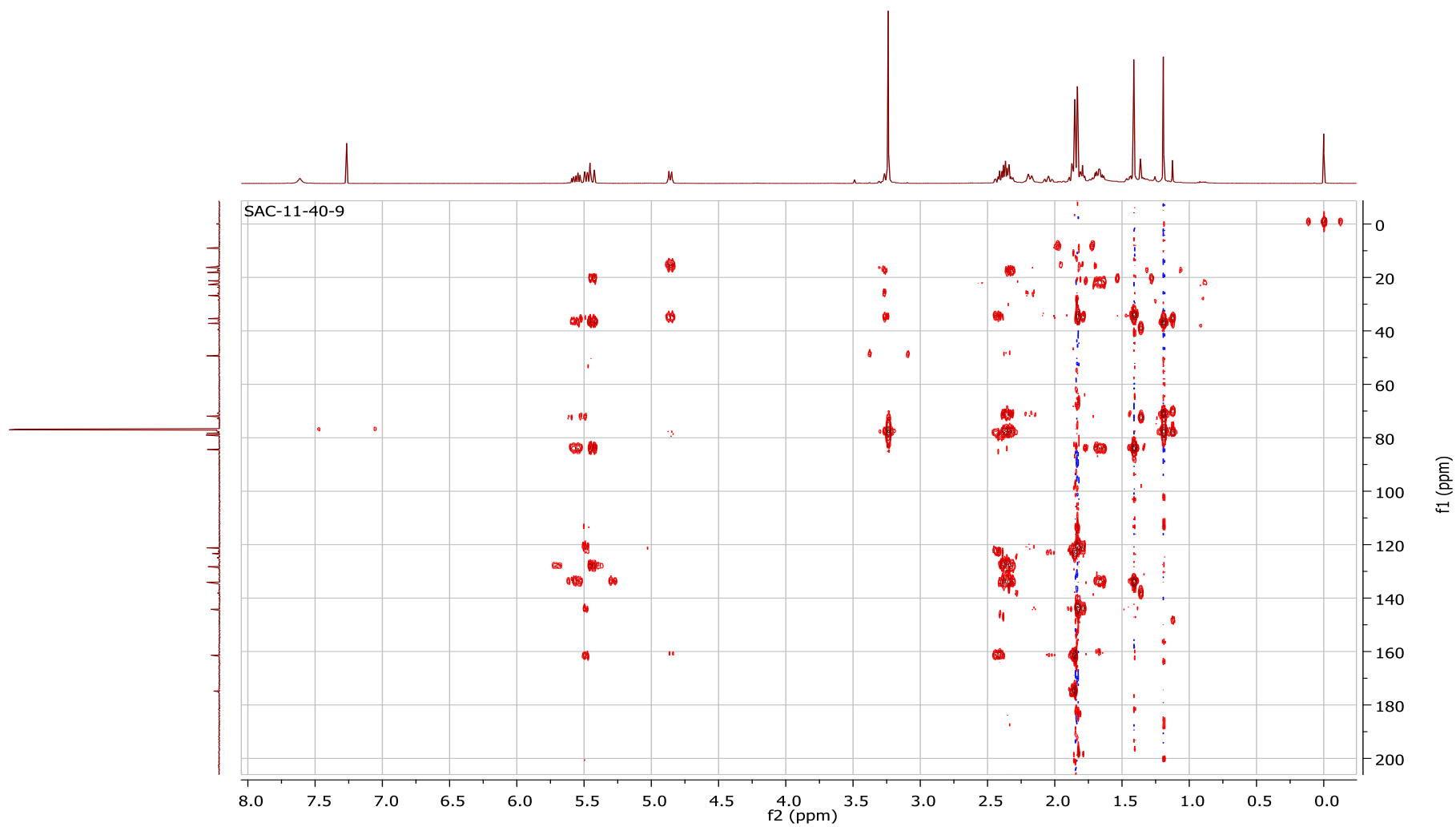
S25:  $^{13}\text{C}$  NMR of **3**



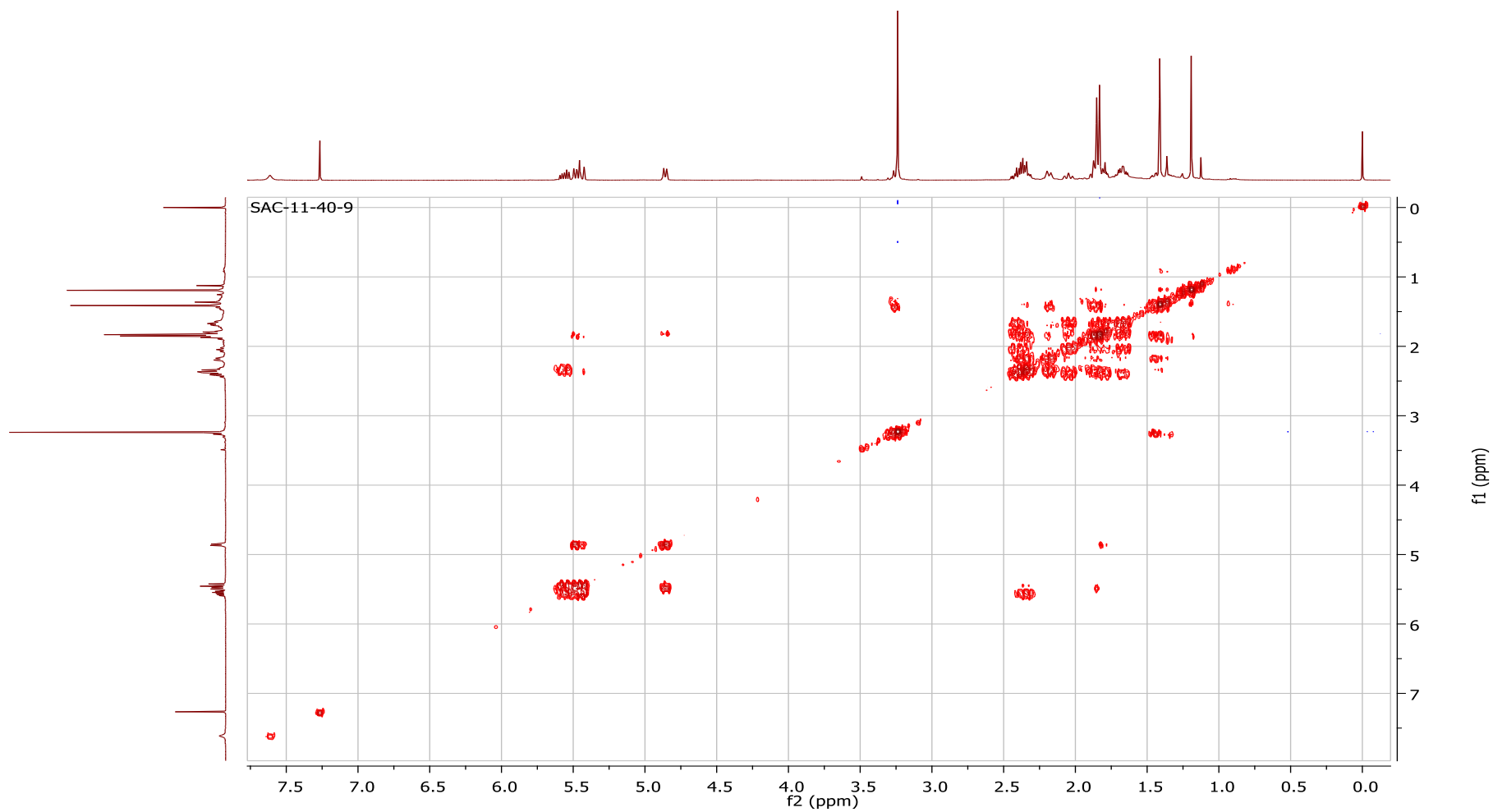
S26: DEPT of 3



S27: HSQC of **3**

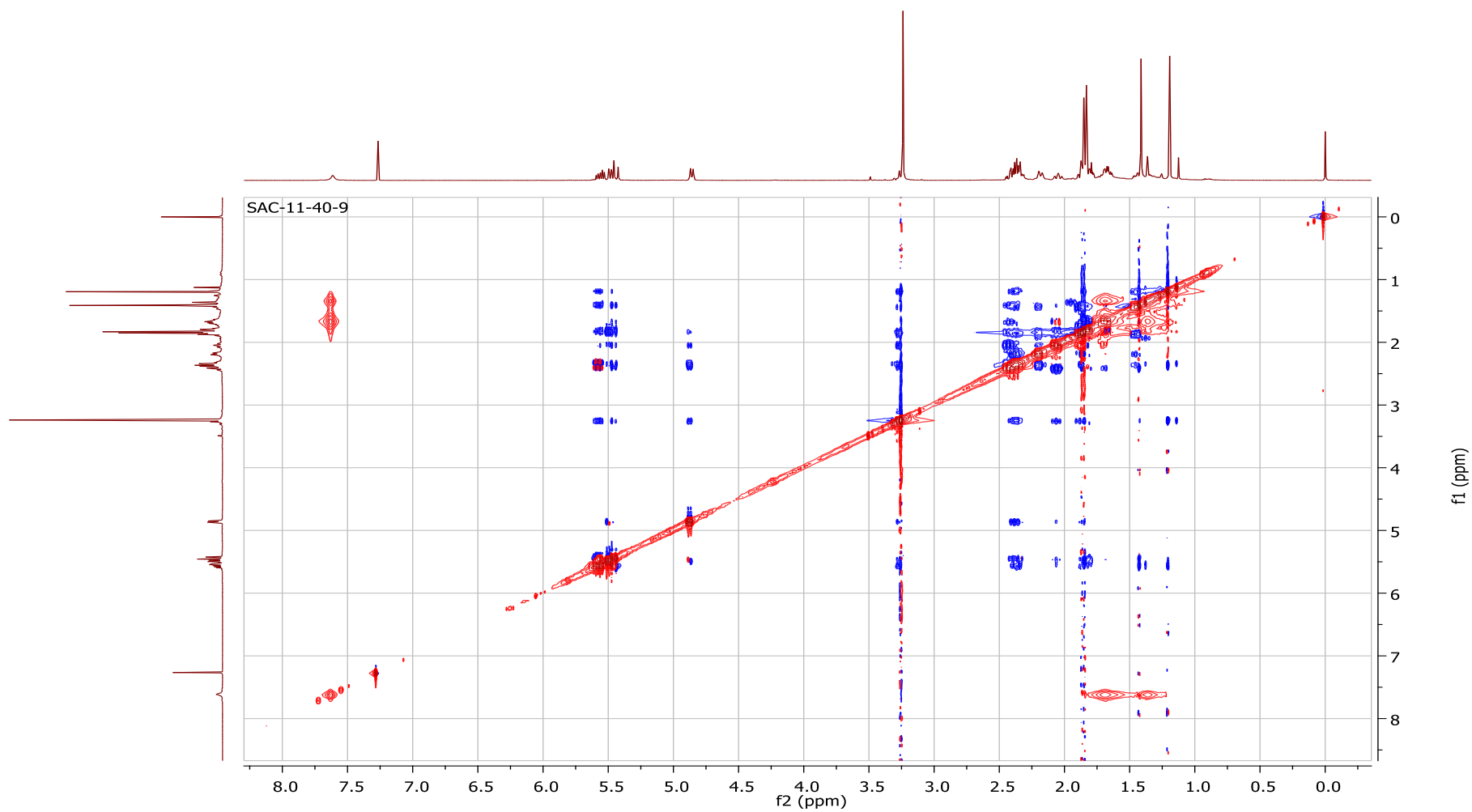


S28: HMBC of **3**

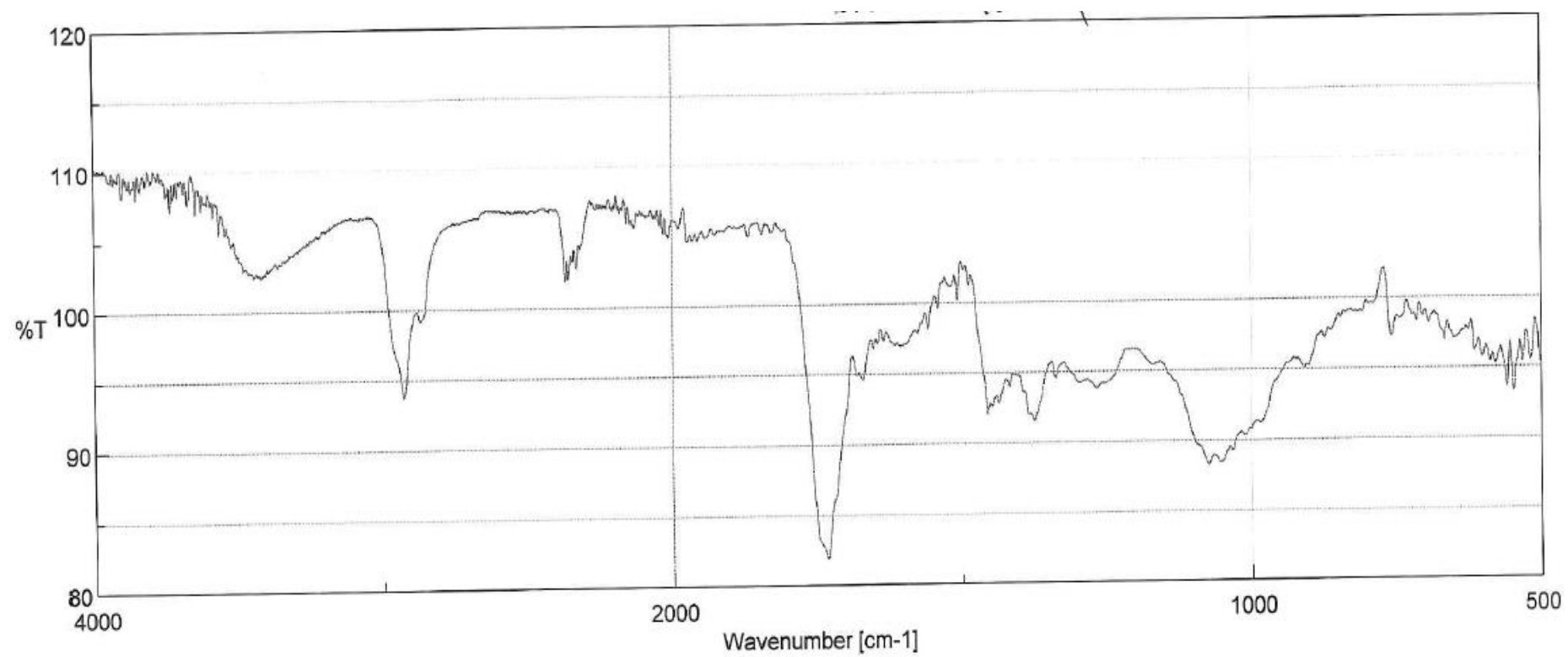


S29:  $^1\text{H}$   $^1\text{H}$  COSY of **3**



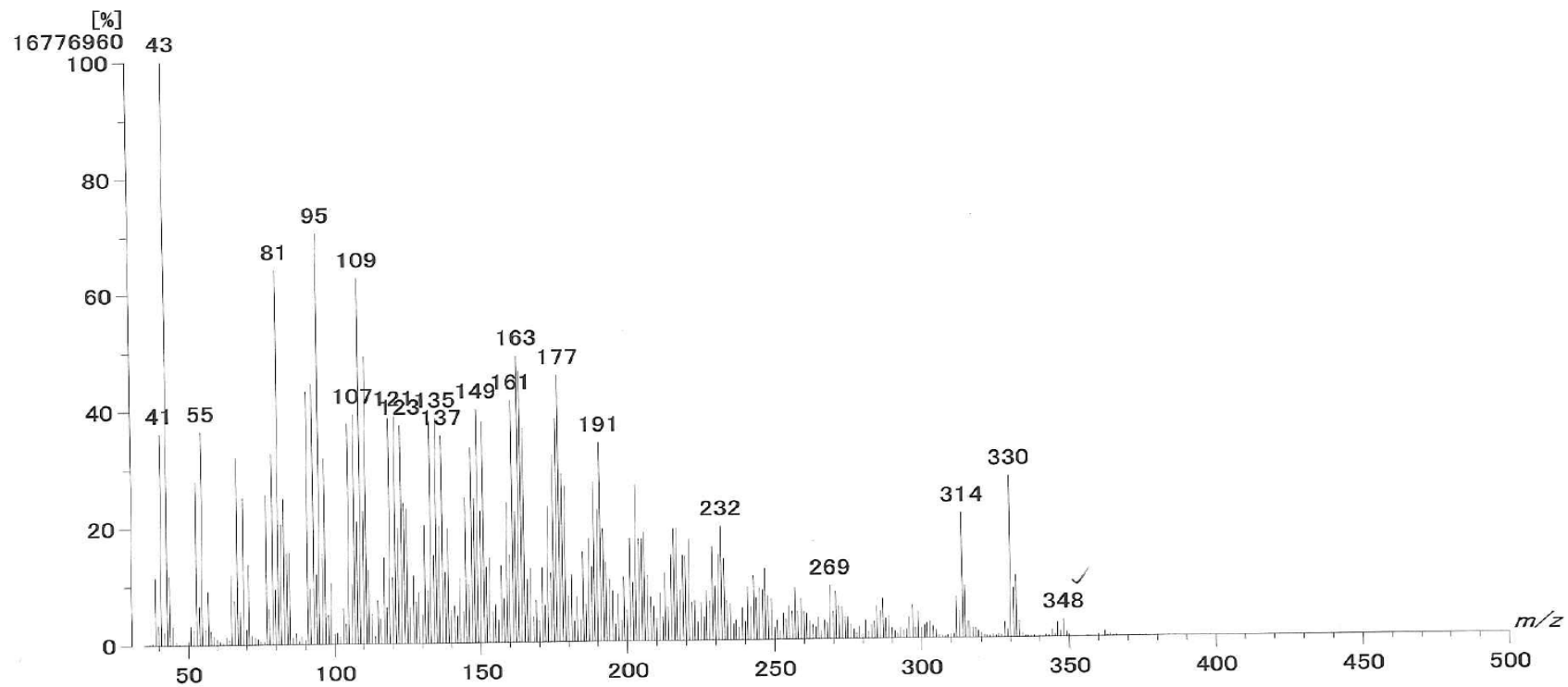


S30: NOESY of **3**



S31: FTIR of **4**

Inlet : Direct      Ion Mode : EI+  
Spectrum Type : Normal Ion [MF-Linear]  
RT : 1.44 min      Scan# : 44  
BP : m/z 43      Int. : 1599.98 (16776960)  
Output m/z range : 35 to 500      Cut Level : 0.00 %

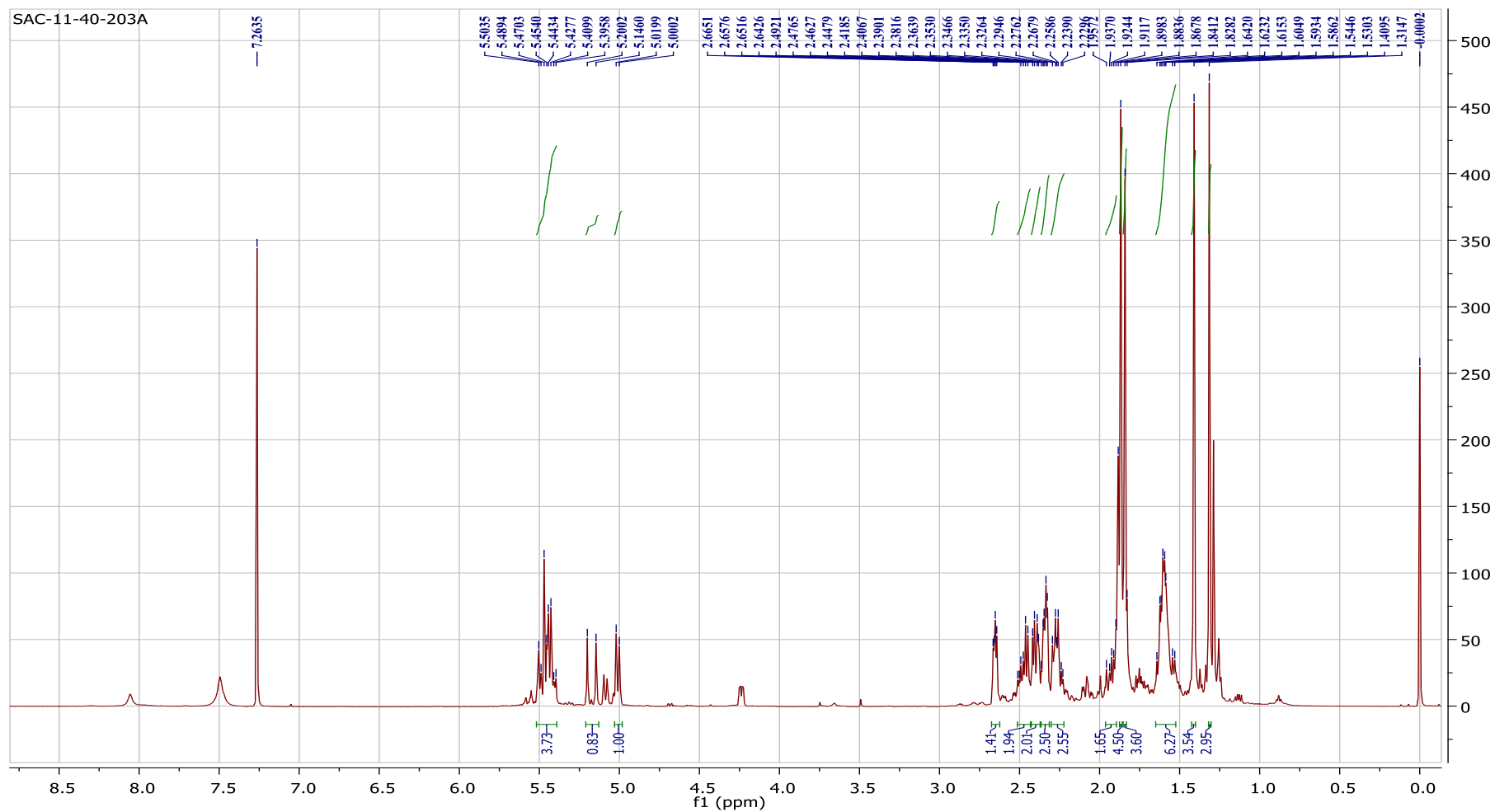


S32: LREIMS of **4**

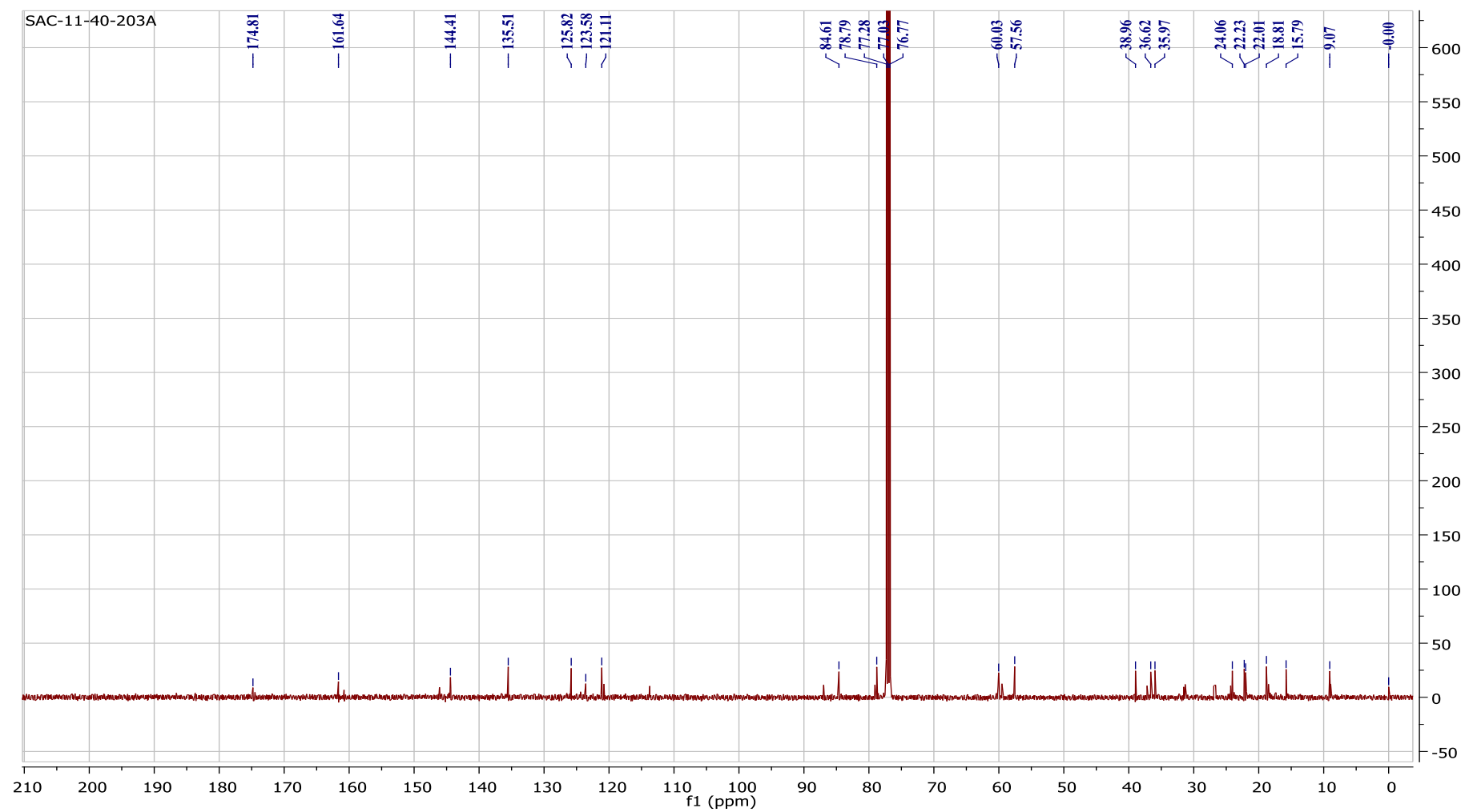
Inlet : Direct      Ion Mode : EI+  
RT : 2.08 min      Scan# : 53  
Elements : C 150/0, H 250/0, O 50/0  
Mass Tolerance : 5mmu  
Unsaturation (U.S.) : 0.0 – 15.0

	Observed m/z	Int%	Err [ppm / mmu]	U.S. Composition
1	348.1939	5.37	+0.6 / +0.2	7.0 C20 H28 O5

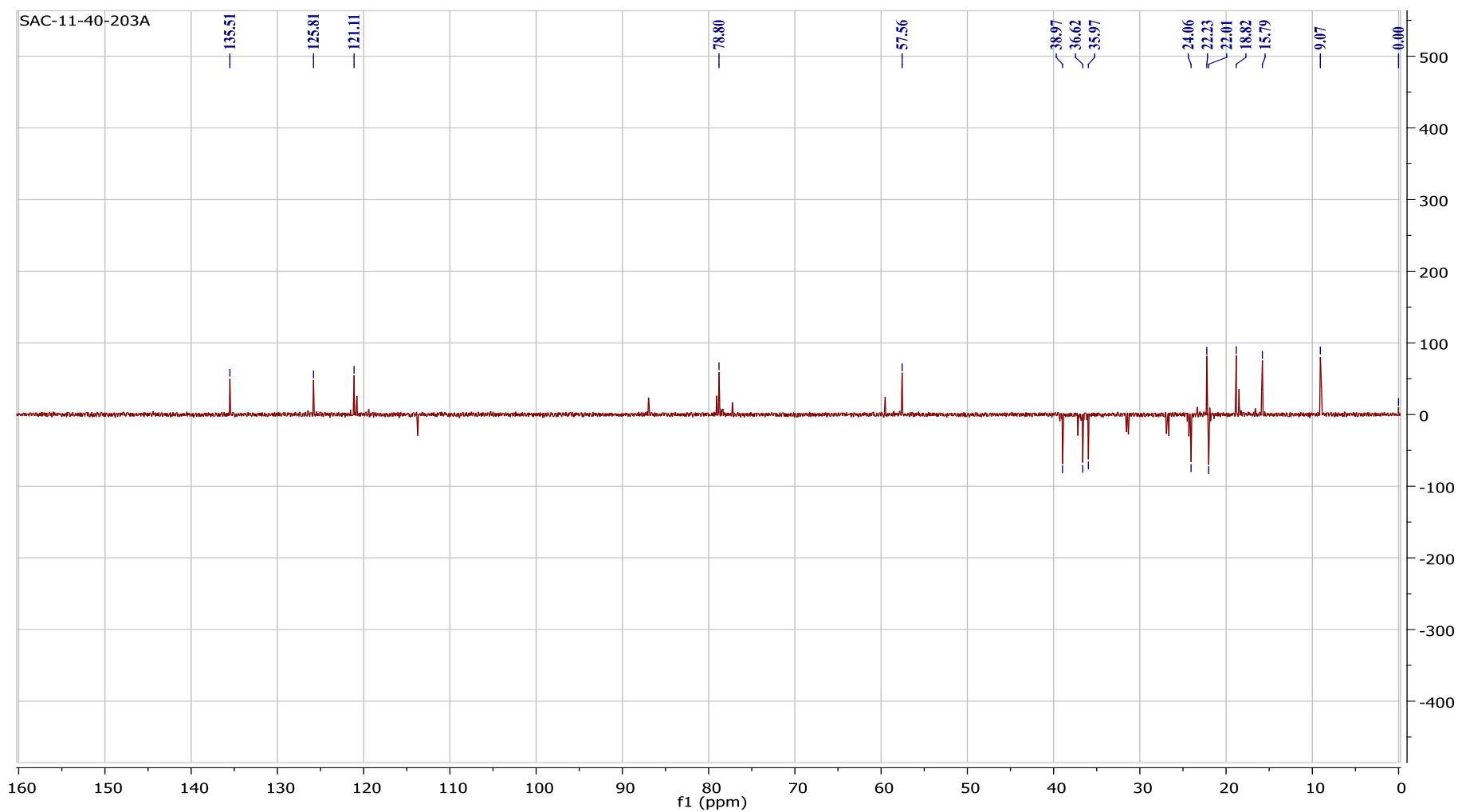
S33: HRCIMS of 4



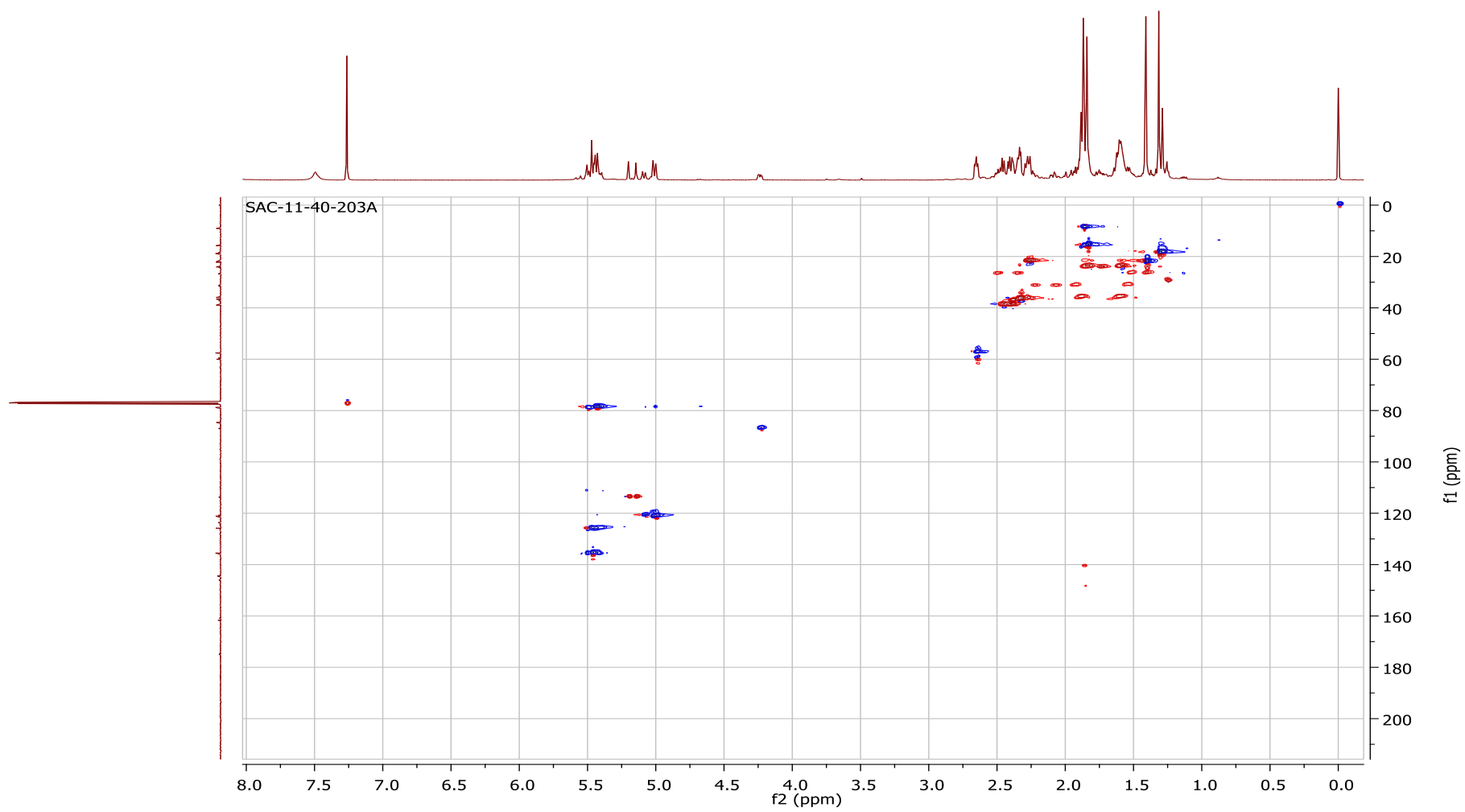
S34:  $^1\text{H}$  NMR of **4**



S35:  $^{13}\text{C}$  NMR of **4**

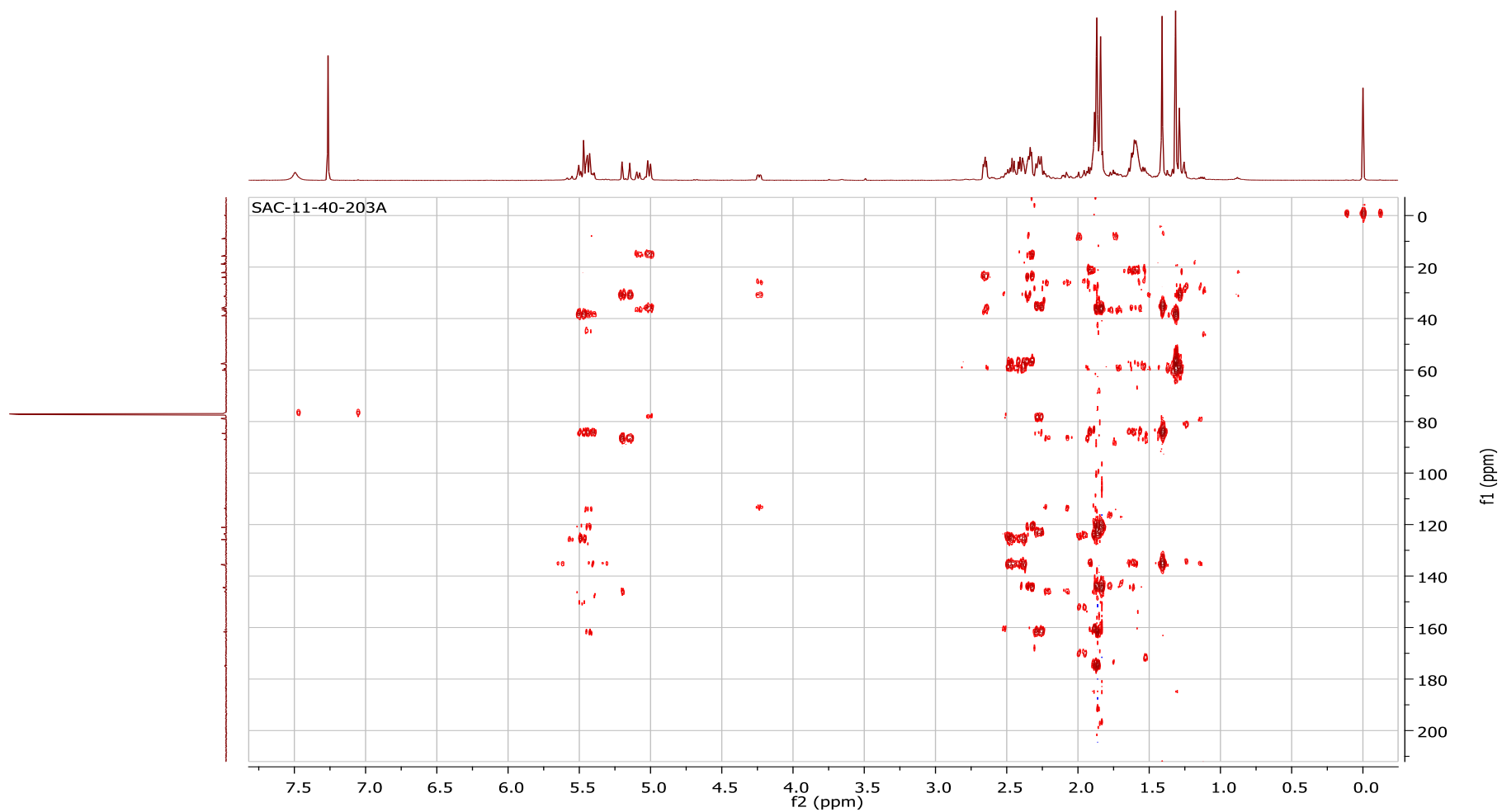


S36: DEPT of 4

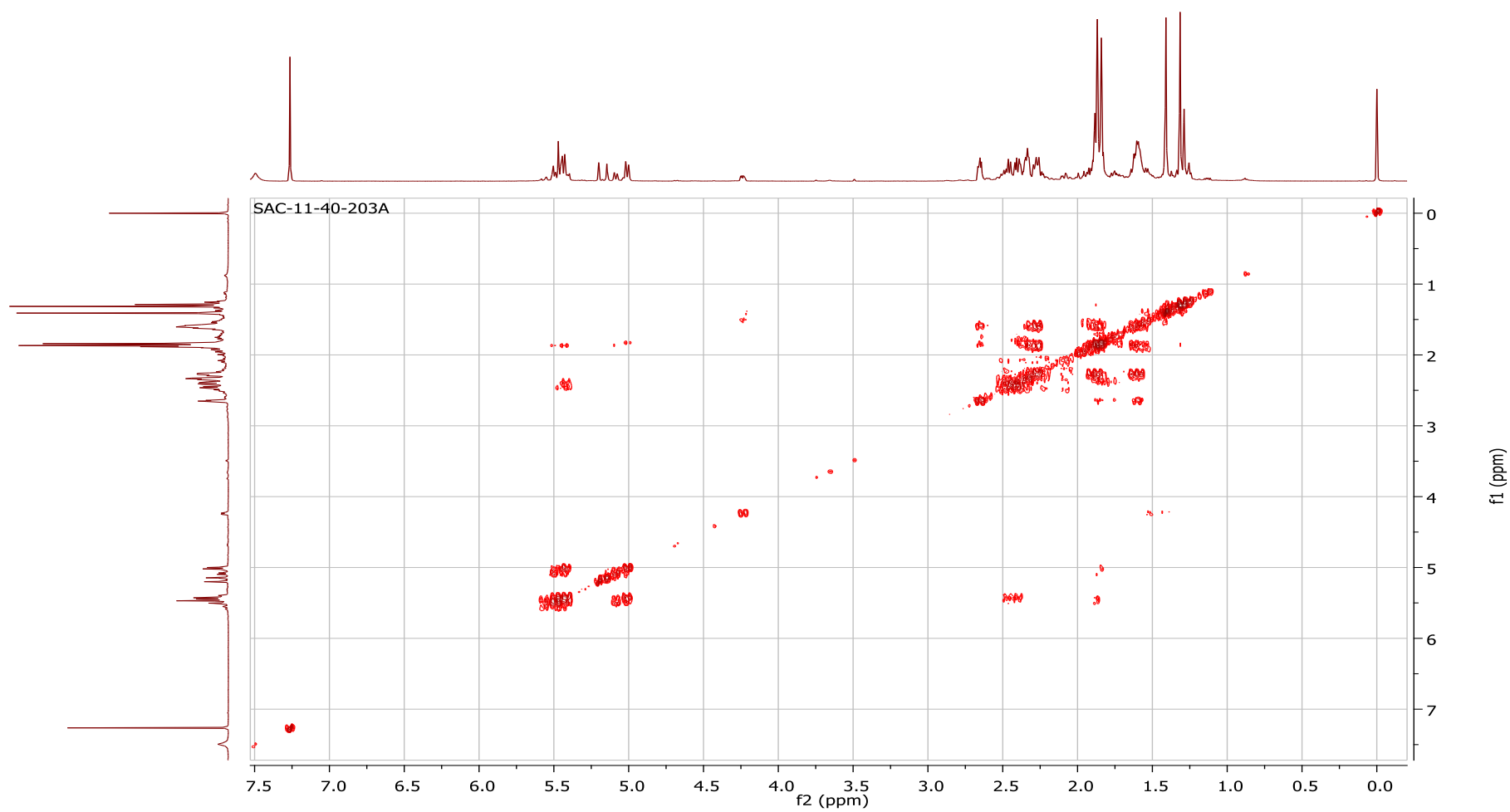


S37: HSQC of **4**

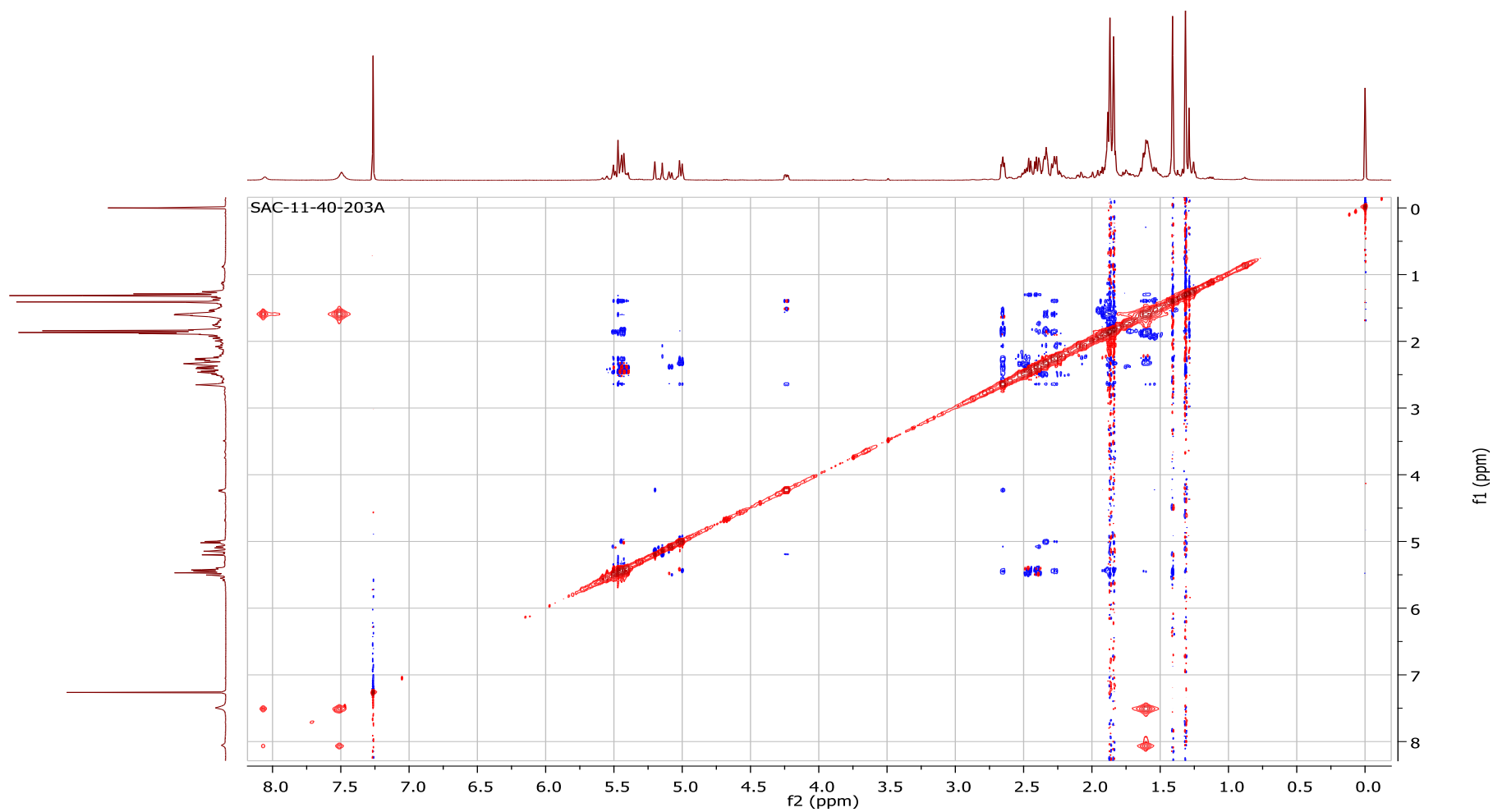




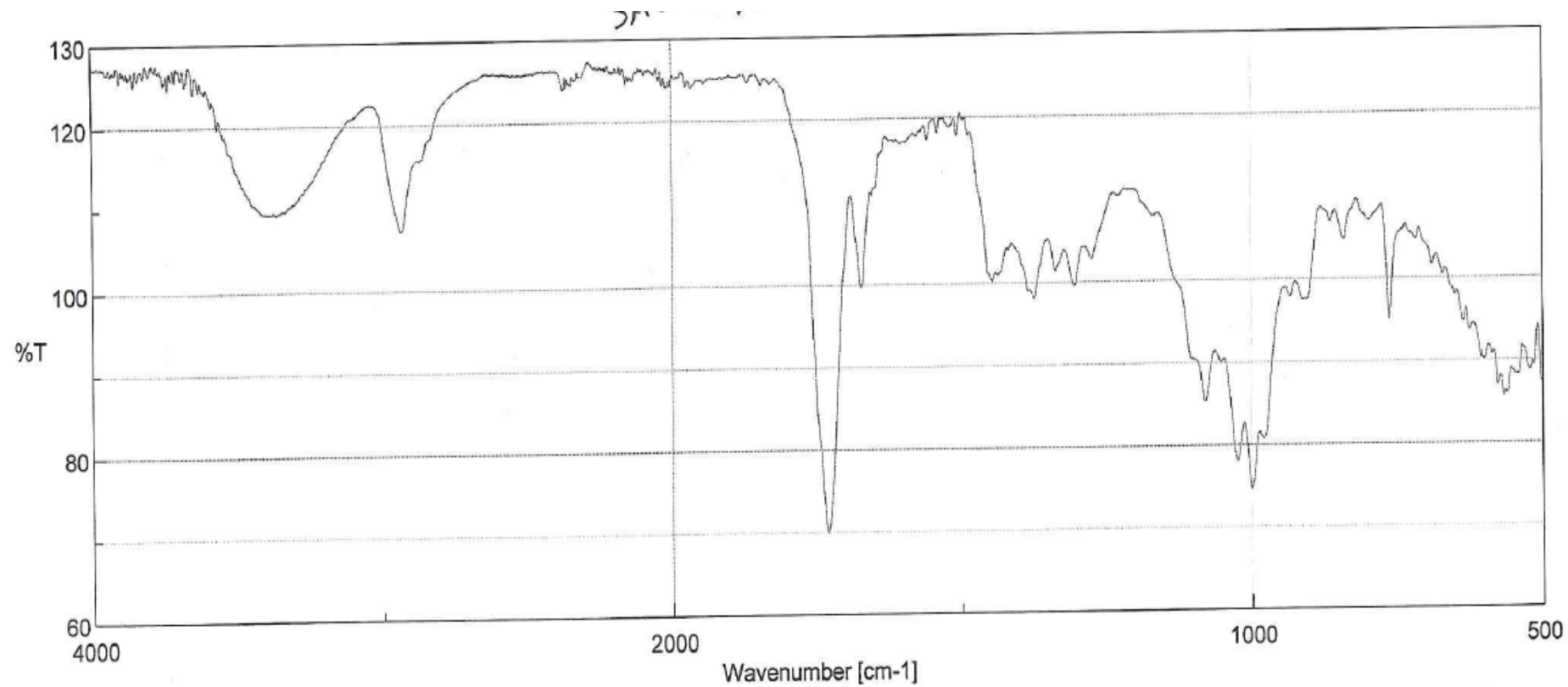
S38: HMBC of **4**



S39:  $^1\text{H}$   $^1\text{H}$  COSY of **4**

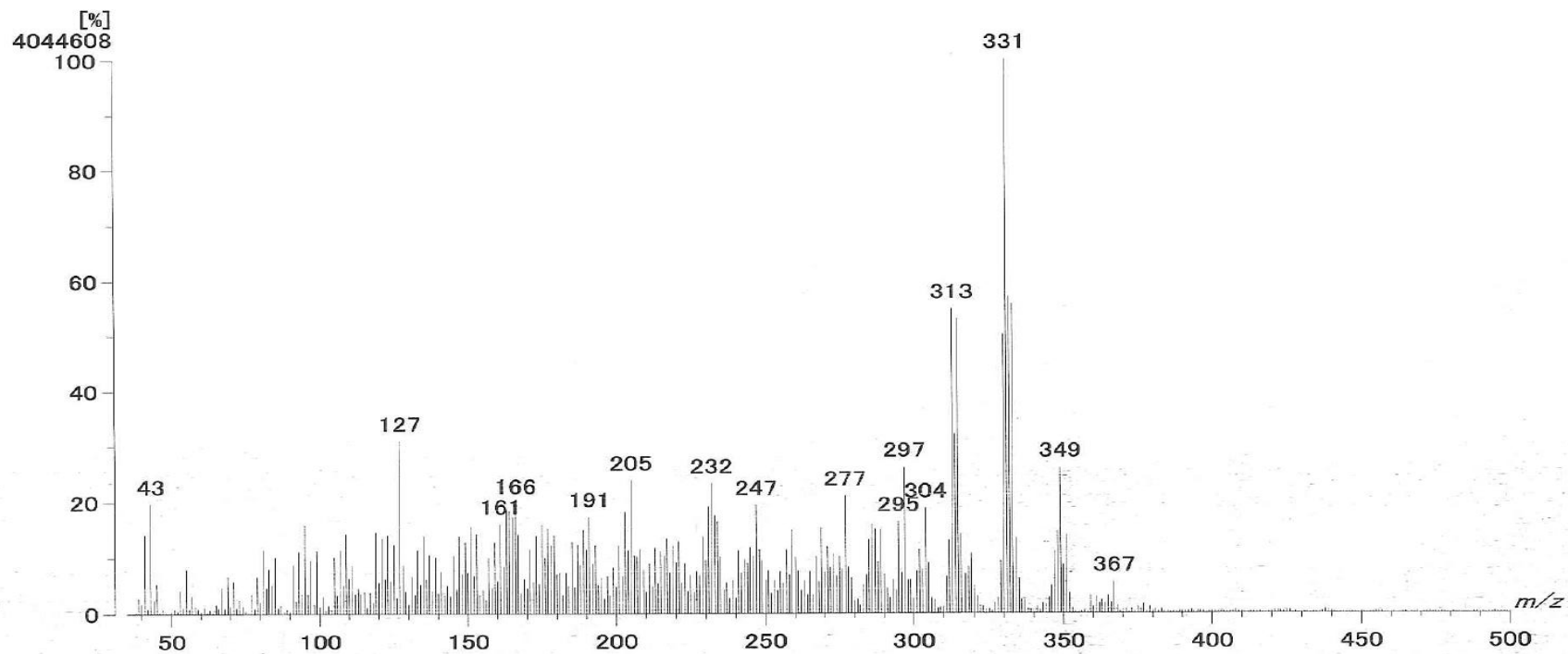


S40: NOESY of **4**



S41: FTIR of 5

Inlet : Direct      Ion Mode : CI+  
Spectrum Type : Normal Ion [MF-Linear]  
RT : 1.99 min      Scan# : 74  
BP : m/z 331      Int. : 385.72 (4044608)  
Output m/z range : 35 to 500      Cut Level : 0.00 %

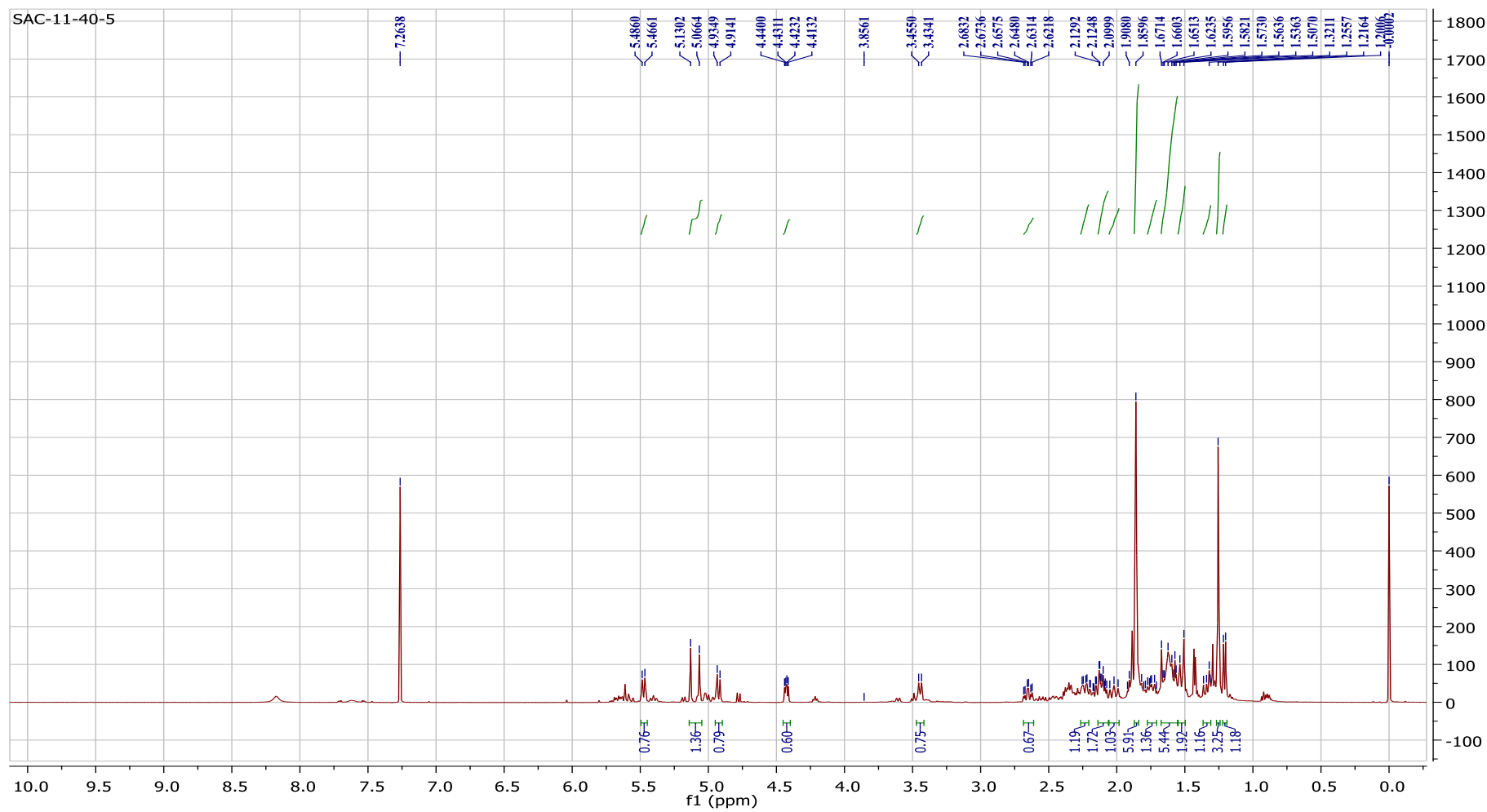


S42: LRCIMS of 5

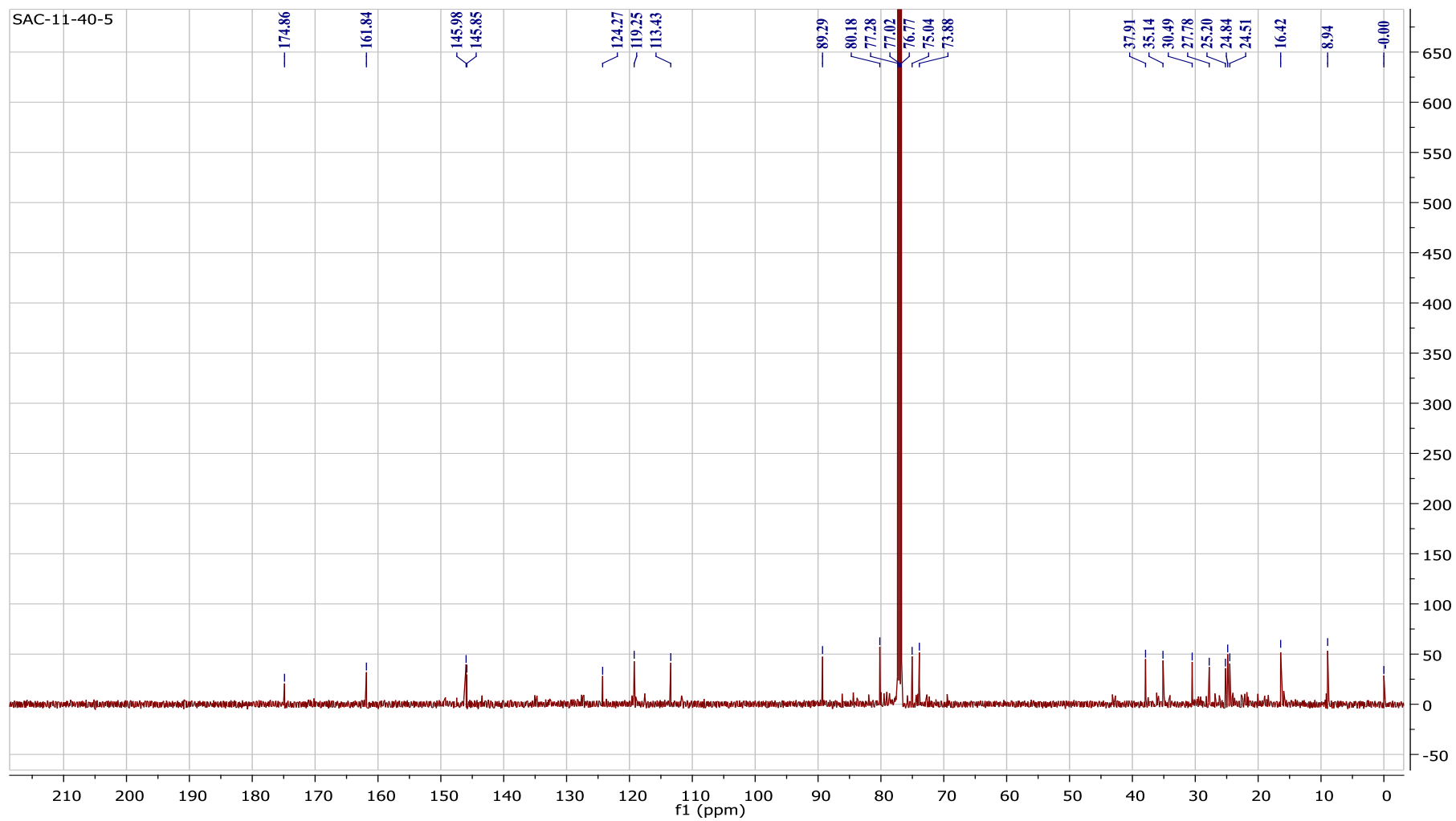
Inlet : Direct      Ion Mode : CI+  
 RT : 1.64 min      Scan# : 42  
 Elements : C 150/0, H 250/0, O 50/0  
 Mass Tolerance : 5mmu  
 Unsaturation (U.S.) : 0.0 – 15.0

	Observed m/z	Int%	Err [ppm / mmu]	U.S. Composition
1	367.2130	3.63	+2.5 / +0.9	5.5 C20 H31 O6

S43: HRCIMS of 5

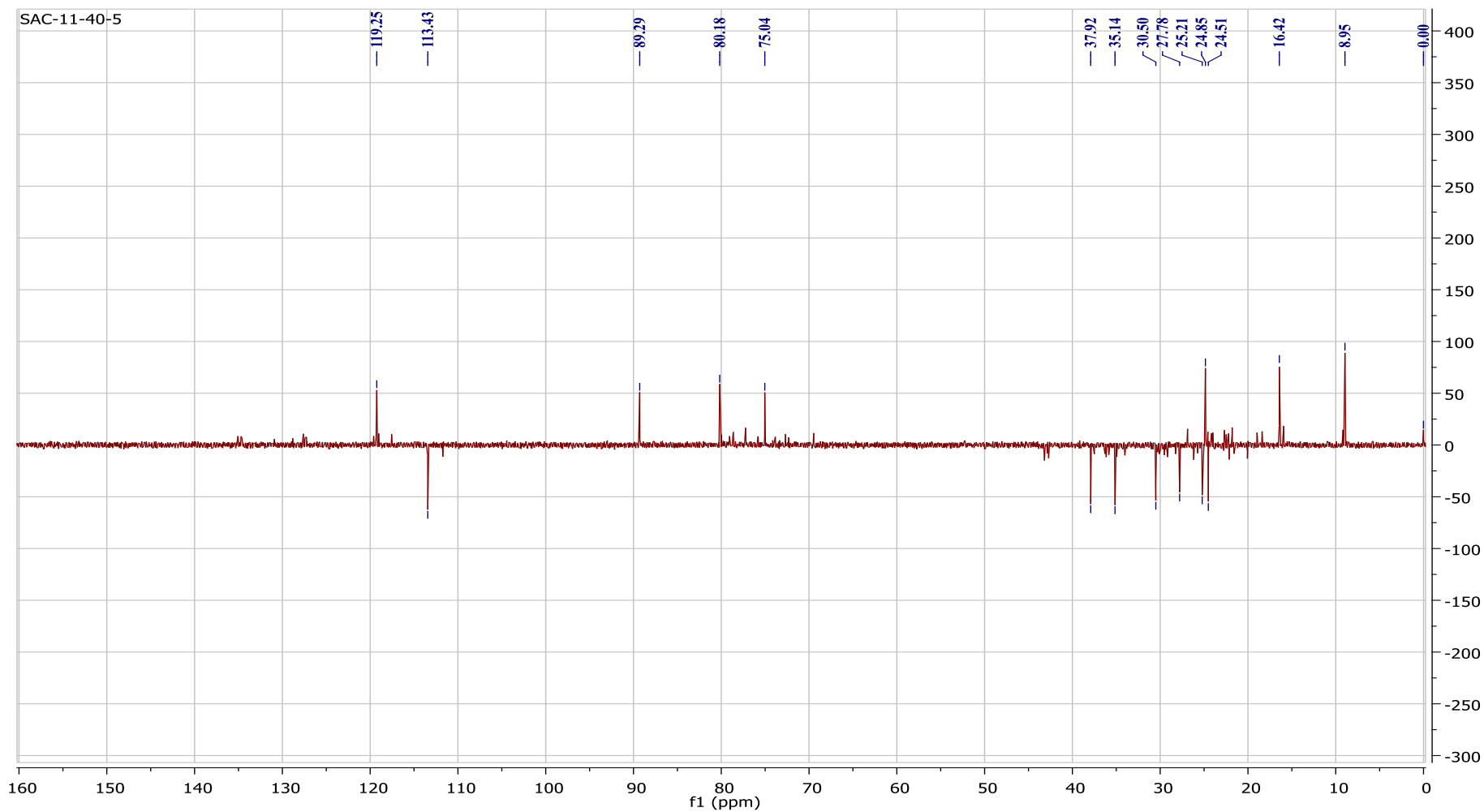


S44:  $^1\text{H}$  NMR of **5**

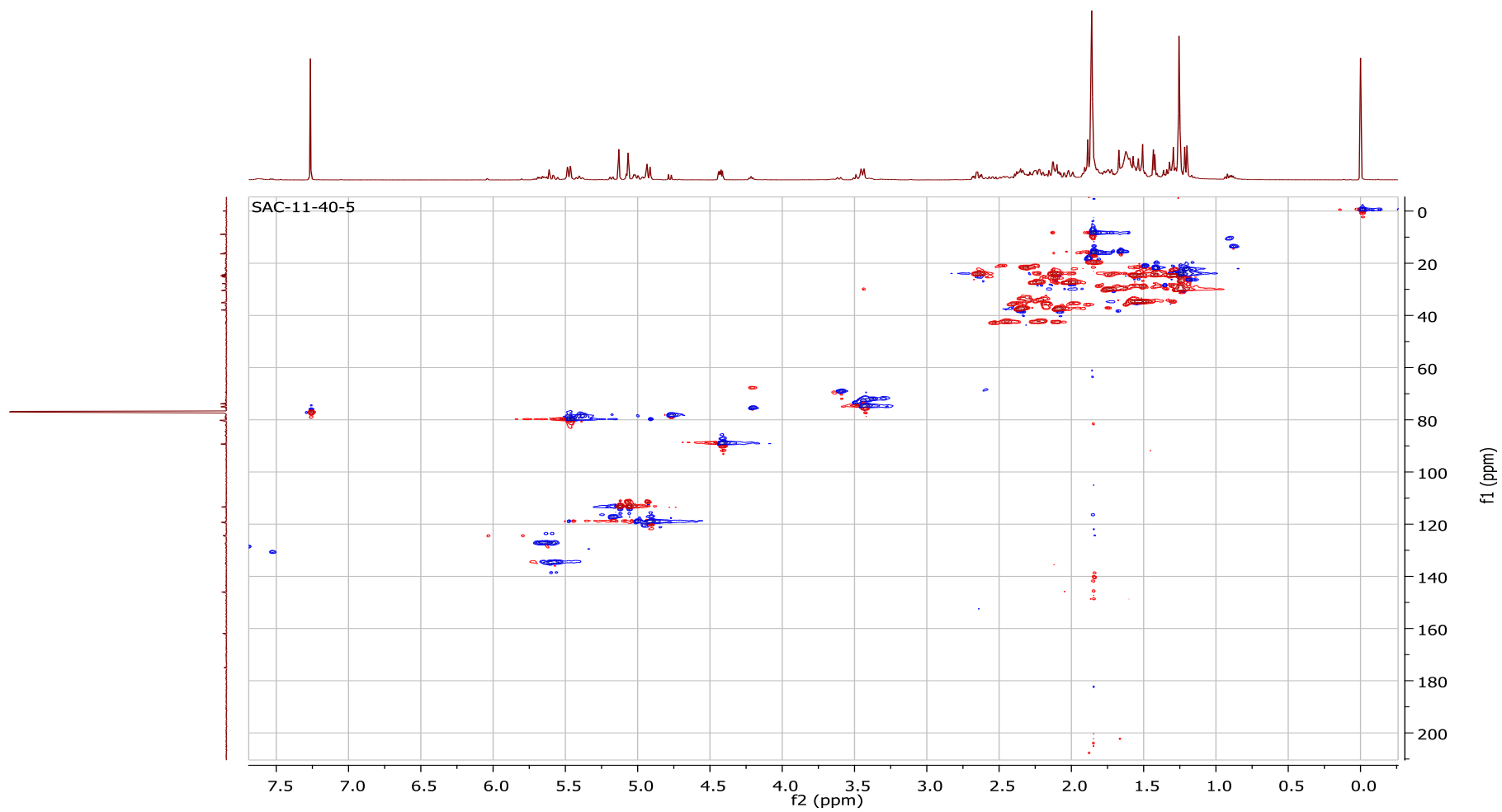


S45:  $^{13}\text{C}$  NMR of **5**

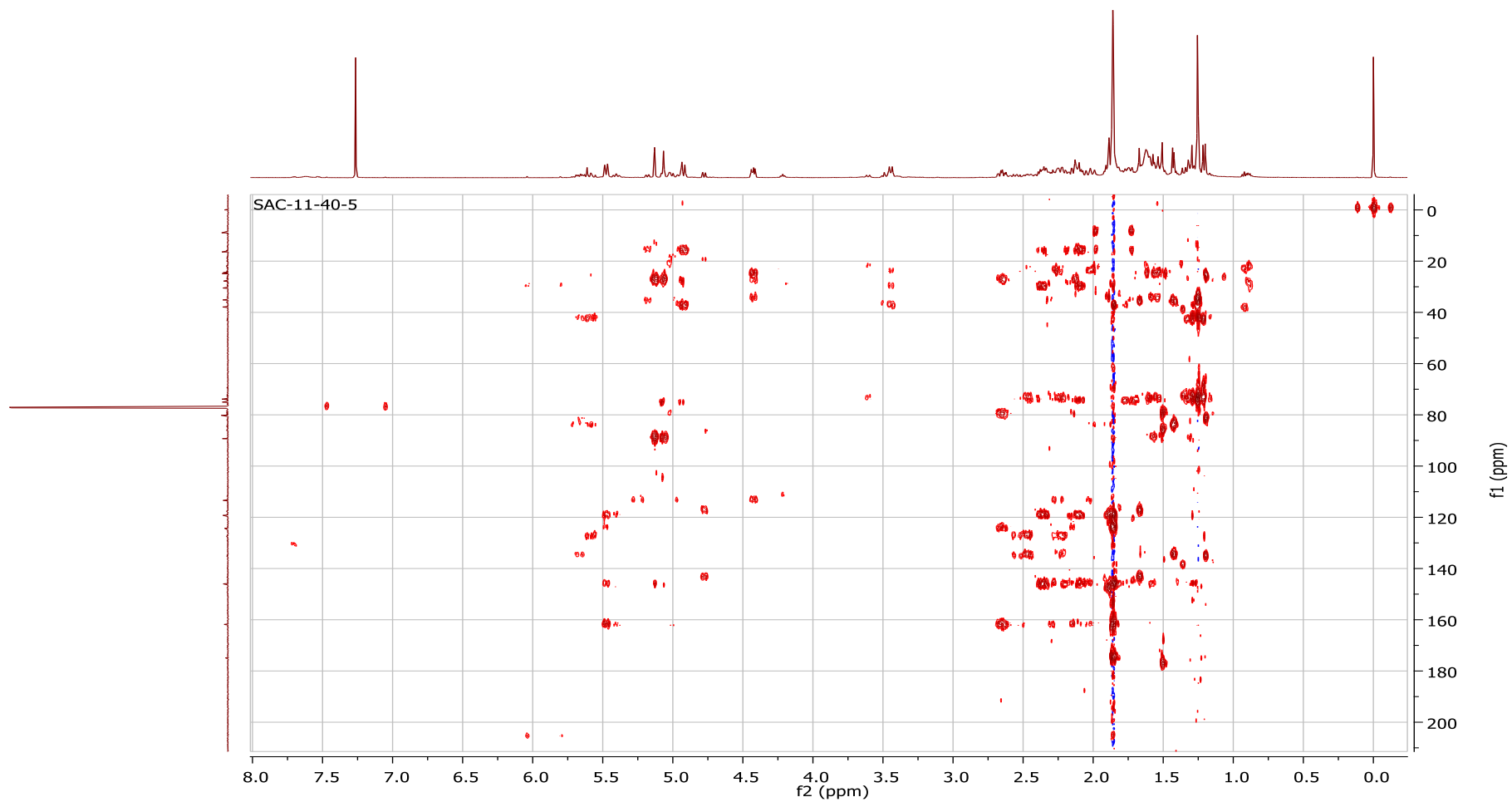




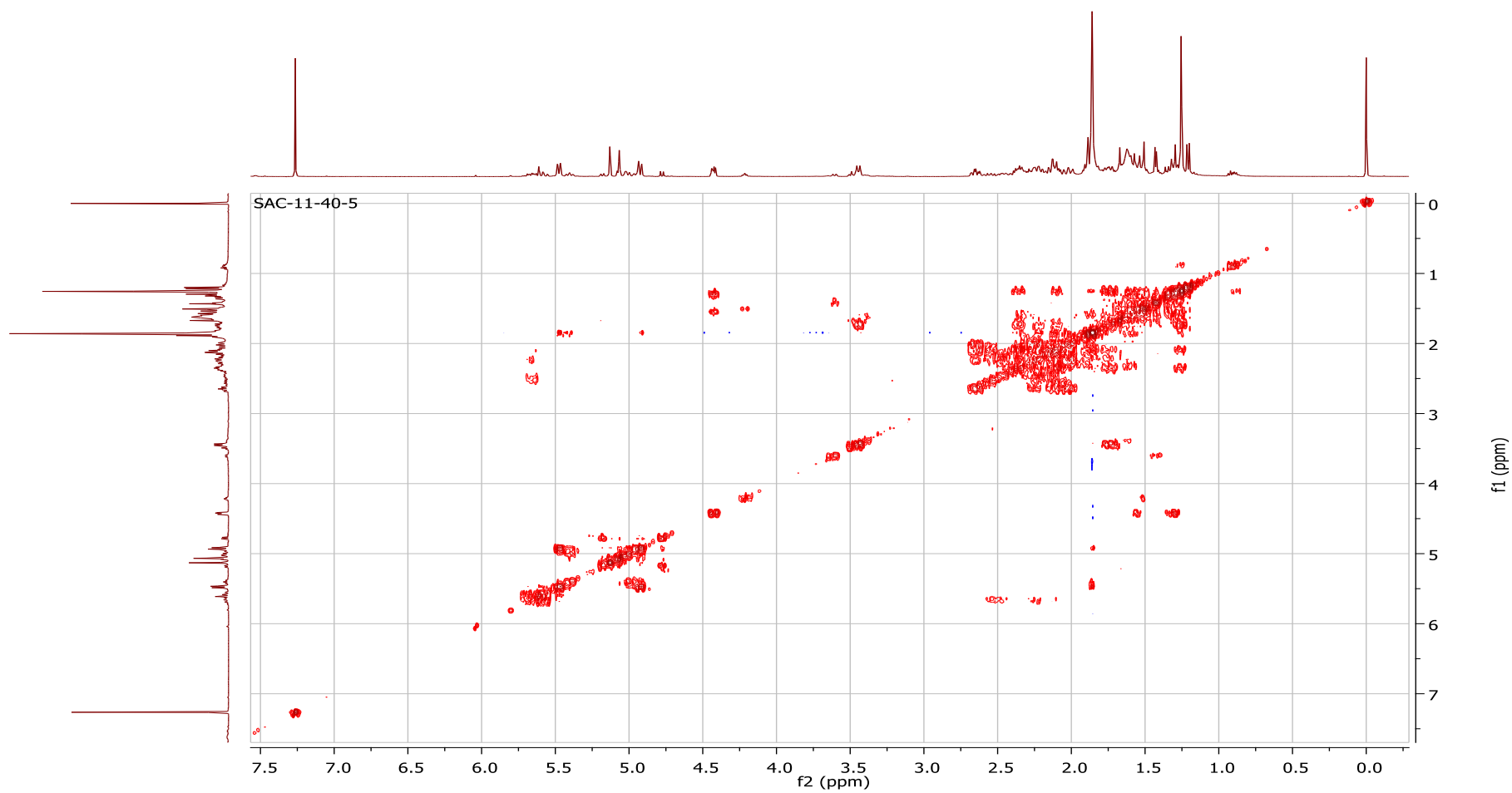
S46: DEPT of 5



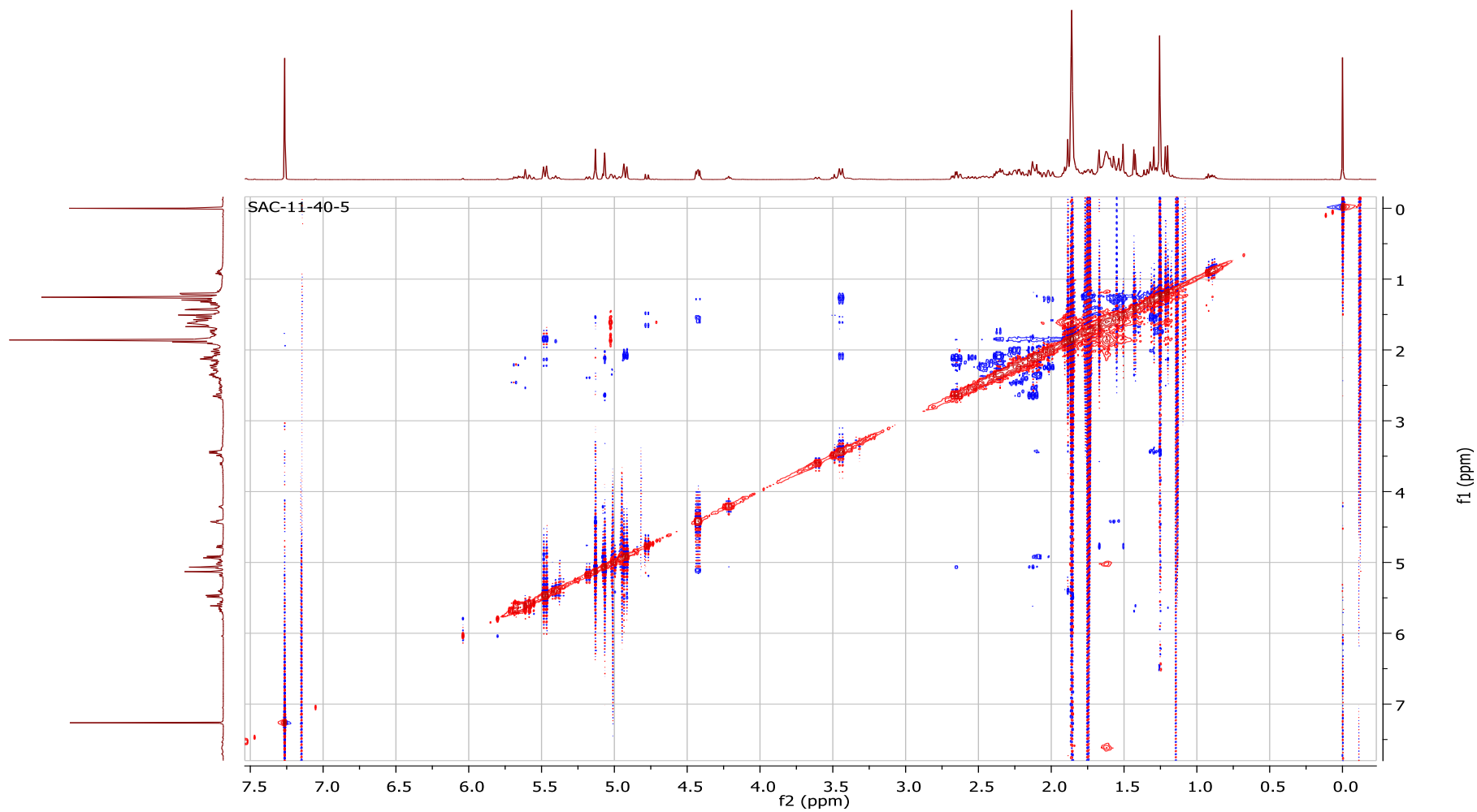
S47: HSQC of **5**



S48: HMBC of **5**



S49:  $^1\text{H}$   $^1\text{H}$  COSY of **5**



S50: NOESY of **5**