

Conformational Comparison of Cyclic $\alpha_3\beta$ Tetra- vs $\alpha_4\beta$ Pentapeptides

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Material and Methods

High resolution mass spectroscopic analysis was performed on a Bruker MicroTOF QII mass spectrometer in positive mode with an internal calibration. All ¹H, ¹³C COSY, HSQC, HMBC spectra were recorded on a 9.4T Bruker AVANCE III 600 MHz instrument at room temperature (298K) with DMSO as an internal standard using a sample concentration in the range of 20 (mg/ml) and all ROESY Experiments were performed at room temperature (298K). Samples were dissolved in 450 μ l of deuterated DMSO-d₆ and measured at 25 °C.. Processing and assignments were carried out using the Topspin 2.3 software from Bruker Karlsruhe.

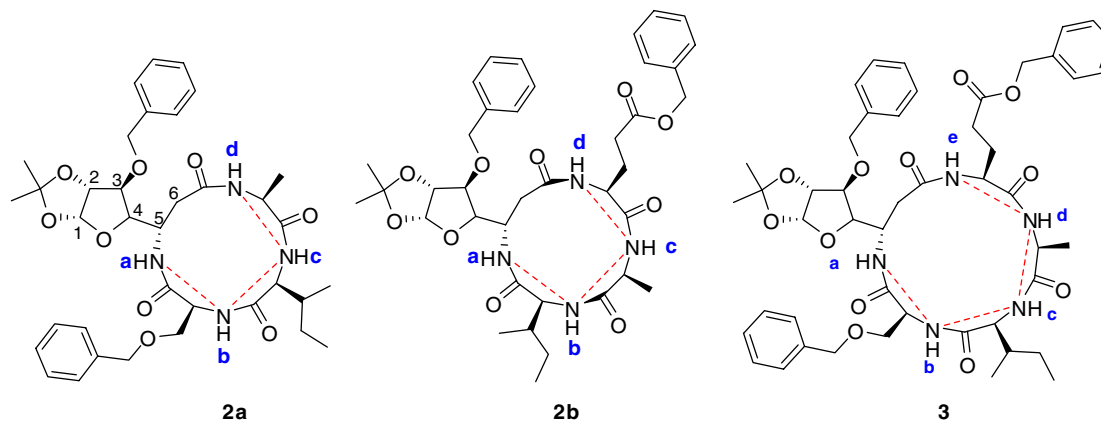


Table 1: ^1H NMR Chemical Shifts (δ , ppm) and Coupling Constants (J , Hz) of glycopeptides **2a**, **2b** and **3**

No.	2a	2b	3
Glycosyl			
1	5.81 (d, $J=3.8\text{Hz}$)	5.82 (d, $J=3.9\text{Hz}$)	5.80 (d, $J=3.0\text{Hz}$)
2	4.77 (d, $J=3.8\text{Hz}$)	4.77 (d, $J=3.9\text{Hz}$)	4.76 (d, $J=3.0\text{Hz}$)
3	4.00 (d, $J=2.7\text{Hz}$)	4.04 (d, $J=2.9\text{Hz}$)	3.88-3.90 (m)
4	4.69 (dd, Merged with H-8, $J=10.8, 2.7\text{Hz}$)	4.60 (dd, $J=10.0, 2.9\text{Hz}$)	4.41-4.49 (m)
5	4.27 (m)	4.21 (t, $J=7.8\text{Hz}$)	4.20-4.24 (m)
6a	2.35 (m)	2.25 (d, $J=15.1\text{Hz}$)	2.34 (d, $J=14.3\text{Hz}$)
6b	2.53 (dd, $J=6.1, 6.0\text{Hz}$)	2.57 (dd, $J=15.1, 2.58$)	2.58 (d, $J=14.3\text{Hz}$)
NH	6.85 (d, $J=9.7\text{Hz}$)	6.95 (d, $J=9.0\text{Hz}$)	7.75 (d, $J=8.8\text{Hz}$)

Glutamic			
NH		8.05 (d, $J=6.9\text{Hz}$)	7.96 (d, $J=5.1\text{Hz}$)
C_{αH}		3.98-4.02 (m)	3.98 (d, $J=7.9\text{Hz}$)
C_{βH}		1.85-1.88 (m)	1.81-1.92 (m)
C_{γH}		2.44 (t, $J=15.1\text{Hz}$)	2.41-2.42 (m)
Alanine			
NH	8.24 (m)	7.70 (d, $J=8.6\text{Hz}$)	8.10 (d, $J=7.9\text{Hz}$)
C_{αH}	3.93 (m)	4.21 (t, $J=7.8\text{Hz}$)	4.12 (t, $J=7.9\text{Hz}$)
C_{βH}	1.22 (d, $J=7.3\text{Hz}$)	1.23 (d, $J=7.3\text{Hz}$)	1.29 (d, $J=7.0\text{Hz}$)
Isoleucine			
NH	7.21 (d, $J=7.3\text{Hz}$)	7.66 (d, $J=9.0\text{Hz}$)	7.60 (d, $J=6.9\text{Hz}$)
C_{αH}	3.91 (m)	3.98-4.02 (m)	3.88-3.90 (m)
C_{βH}	1.62 (m)	1.85-1.88 (m)	1.82-1.91 (m)
C_{γH}	1.01 (m), 1.38 (m),	0.96 (m), 1.39 (m),	1.06-1.08 (m), 1.40 (m),
C_{δH}	0.82 (m)	0.80-0.83 (m)	0.75-0.80 (m)
Serine			
NH	8.23 (m)		8.19 (d, $J=6.4\text{Hz}$)
C_{αH}	4.29 (m)		4.20-4.24 (m)
C_{βH1}	3.56 (dd, $J=9.7, 5.8\text{Hz}$)		3.71-3.73 (m)
C_{βH2}	3.72 (dd, $J=9.7, 5.8\text{Hz}$)		

Figure 1 ^1H NMR spectrum of 2a

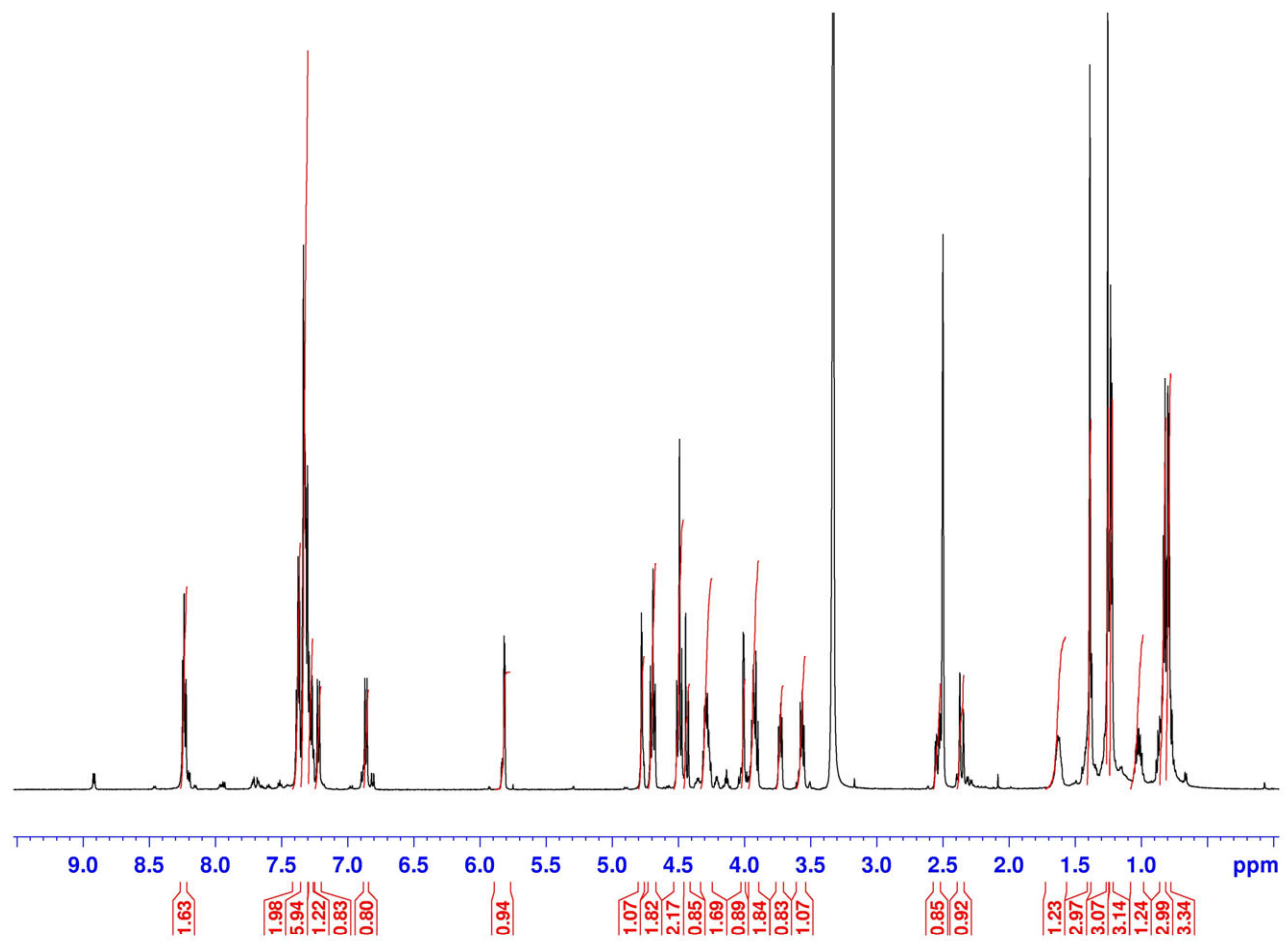


Figure 2 ^{13}C NMR spectrum of 2a

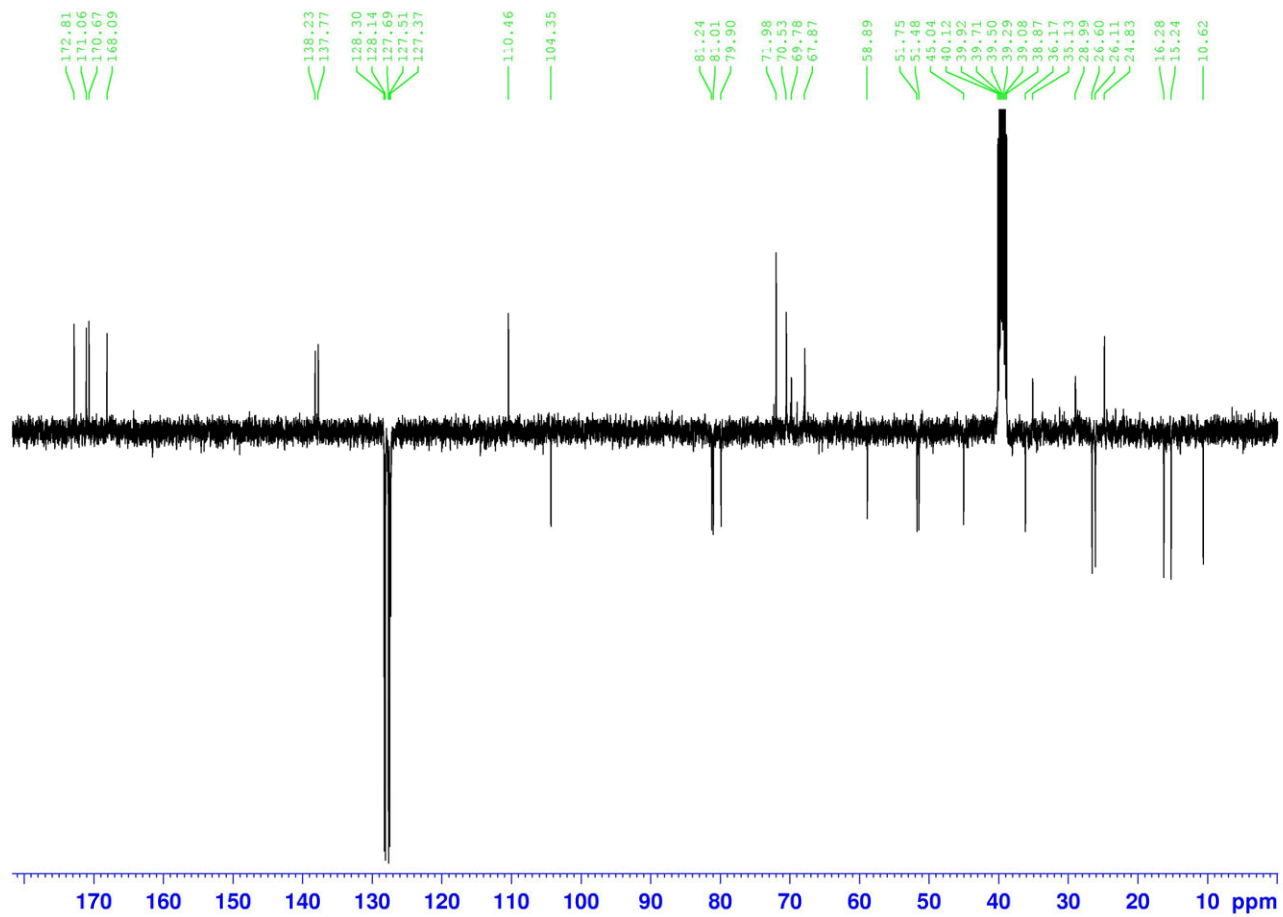


Figure 3 ROESY spectrum of 2a

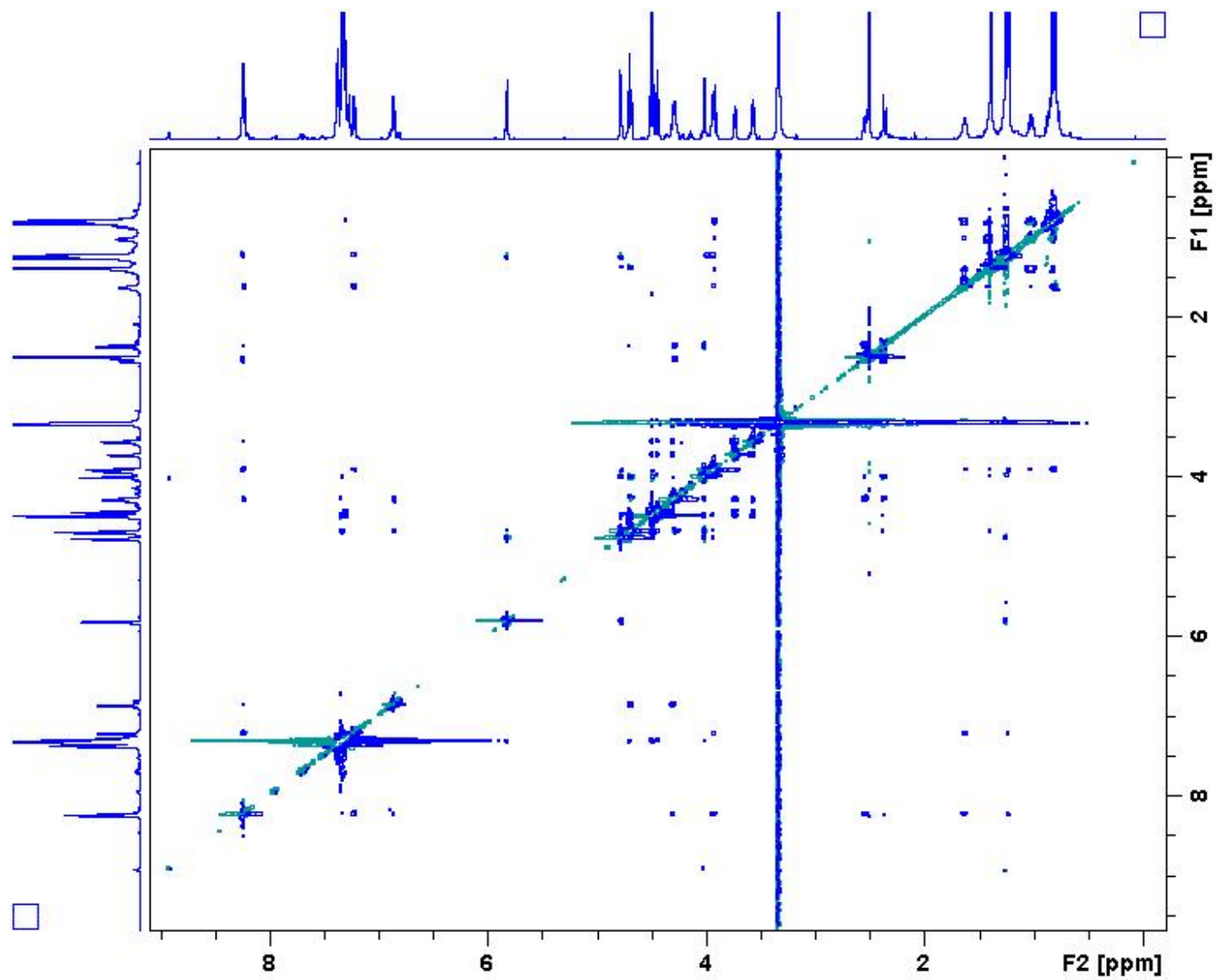


Figure 4 COSY spectrum of 2a

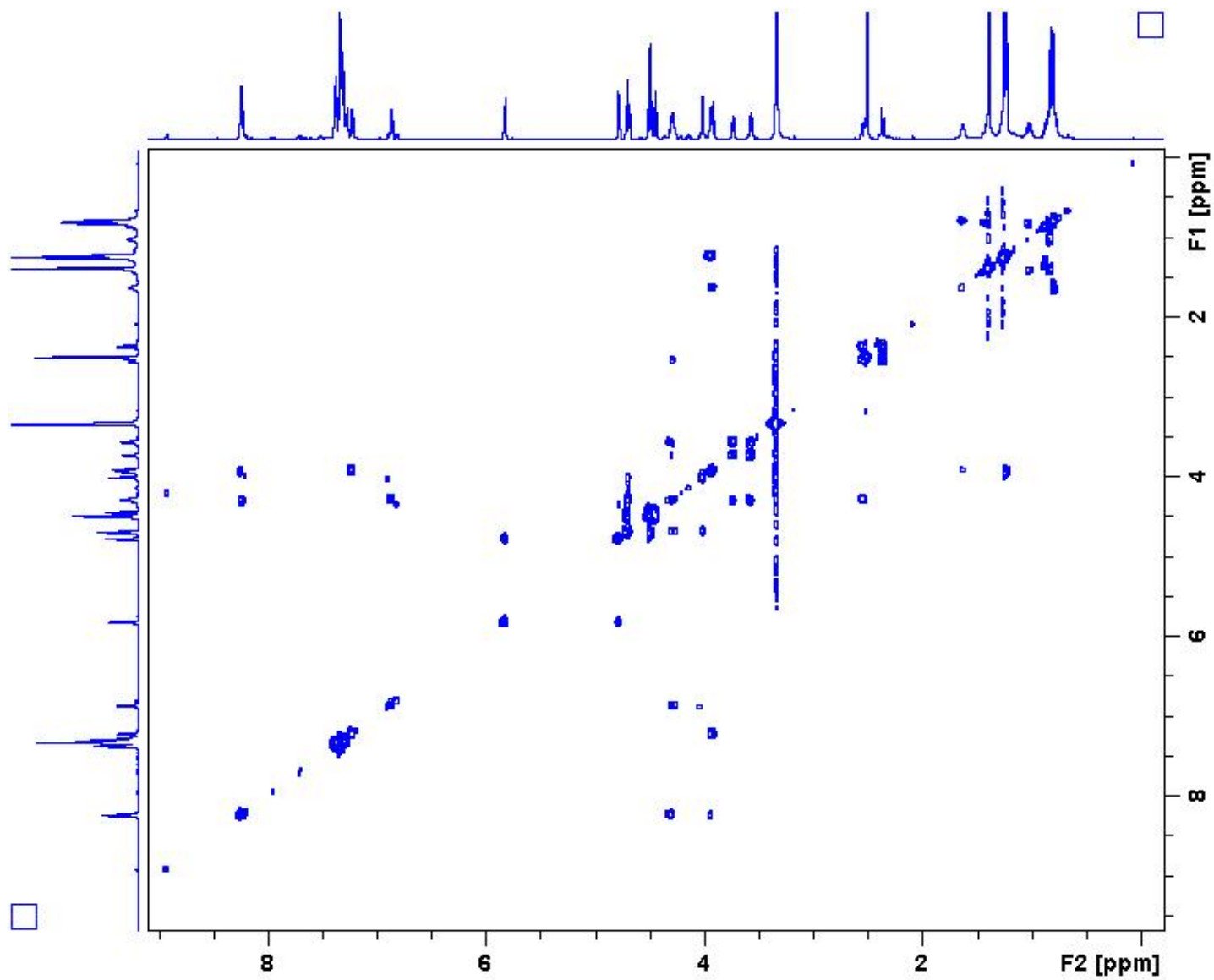


Figure 5 HSQC spectrum of 2a

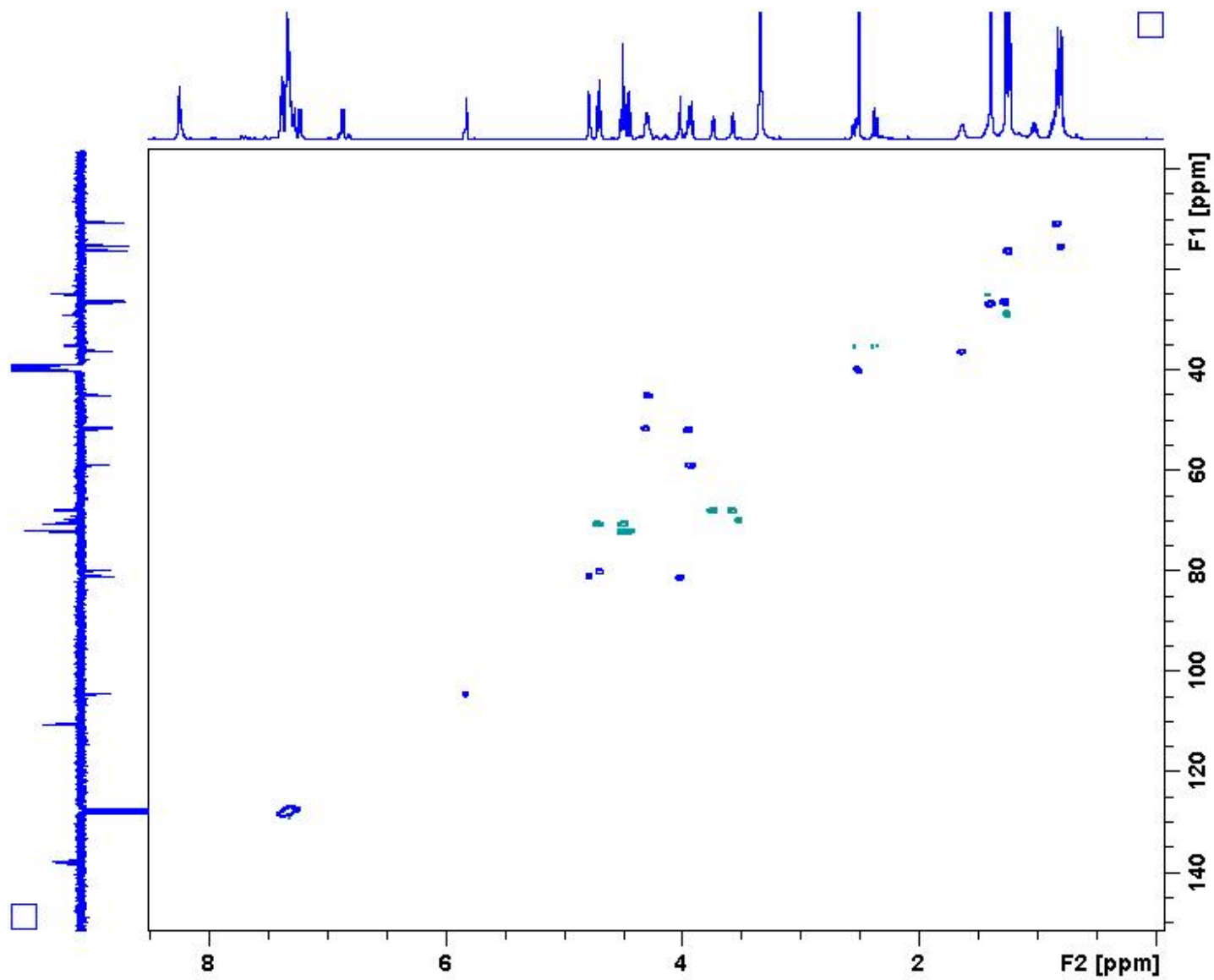


Figure 6 HMBC spectrum of 2a

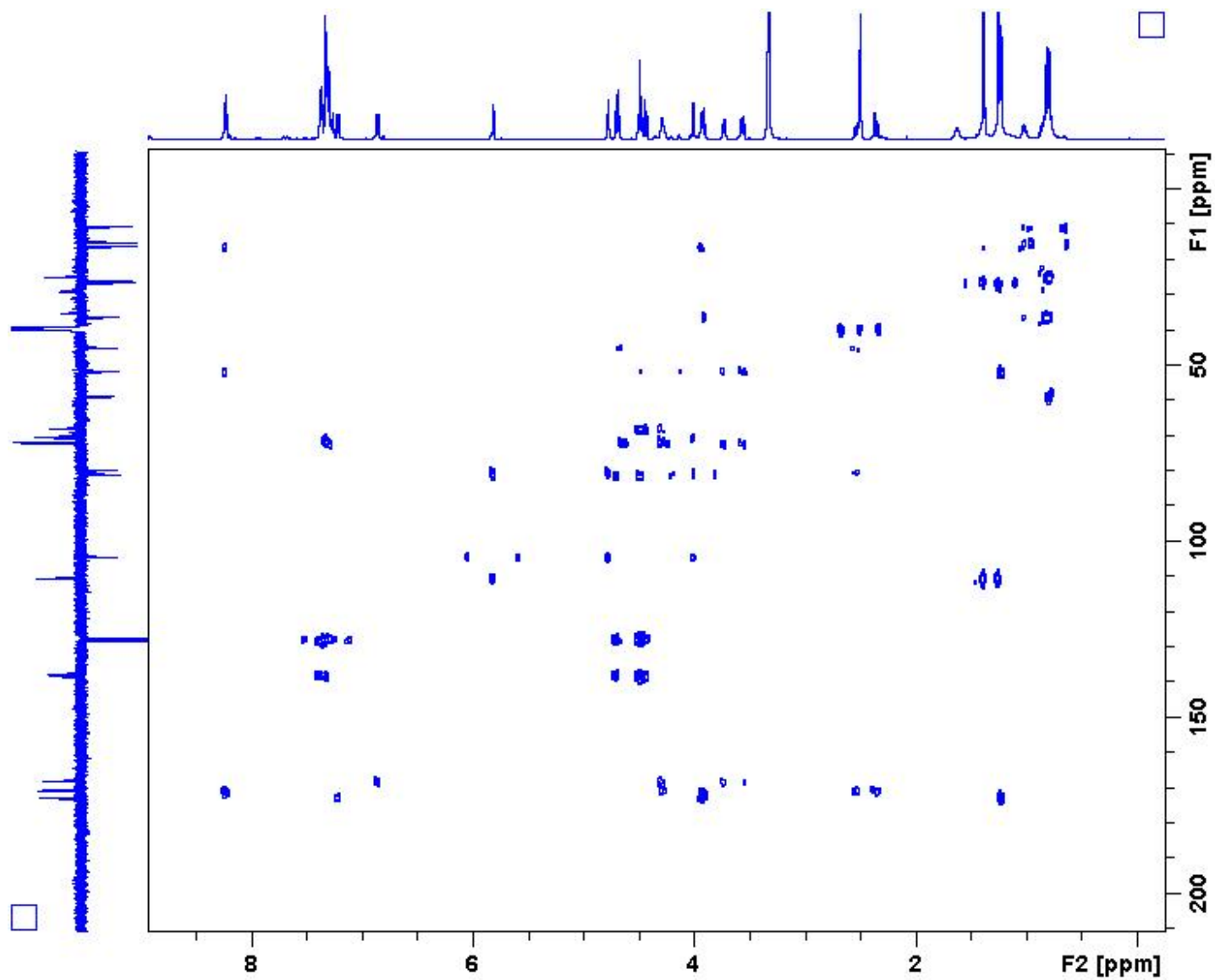


Figure 7 ^1H NMR spectrum of **2b**

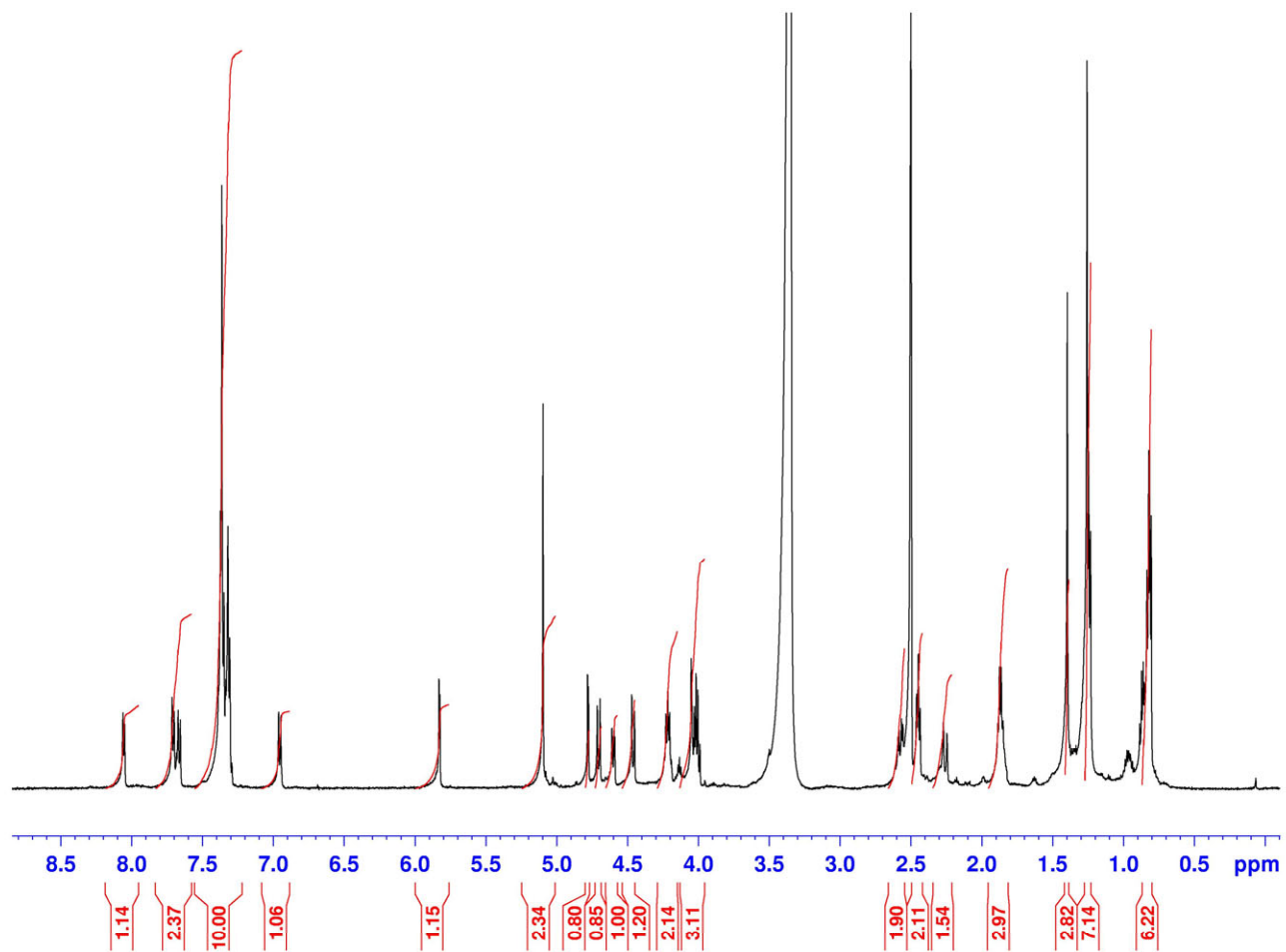


Figure 8 ^{13}C NMR spectrum of 2b

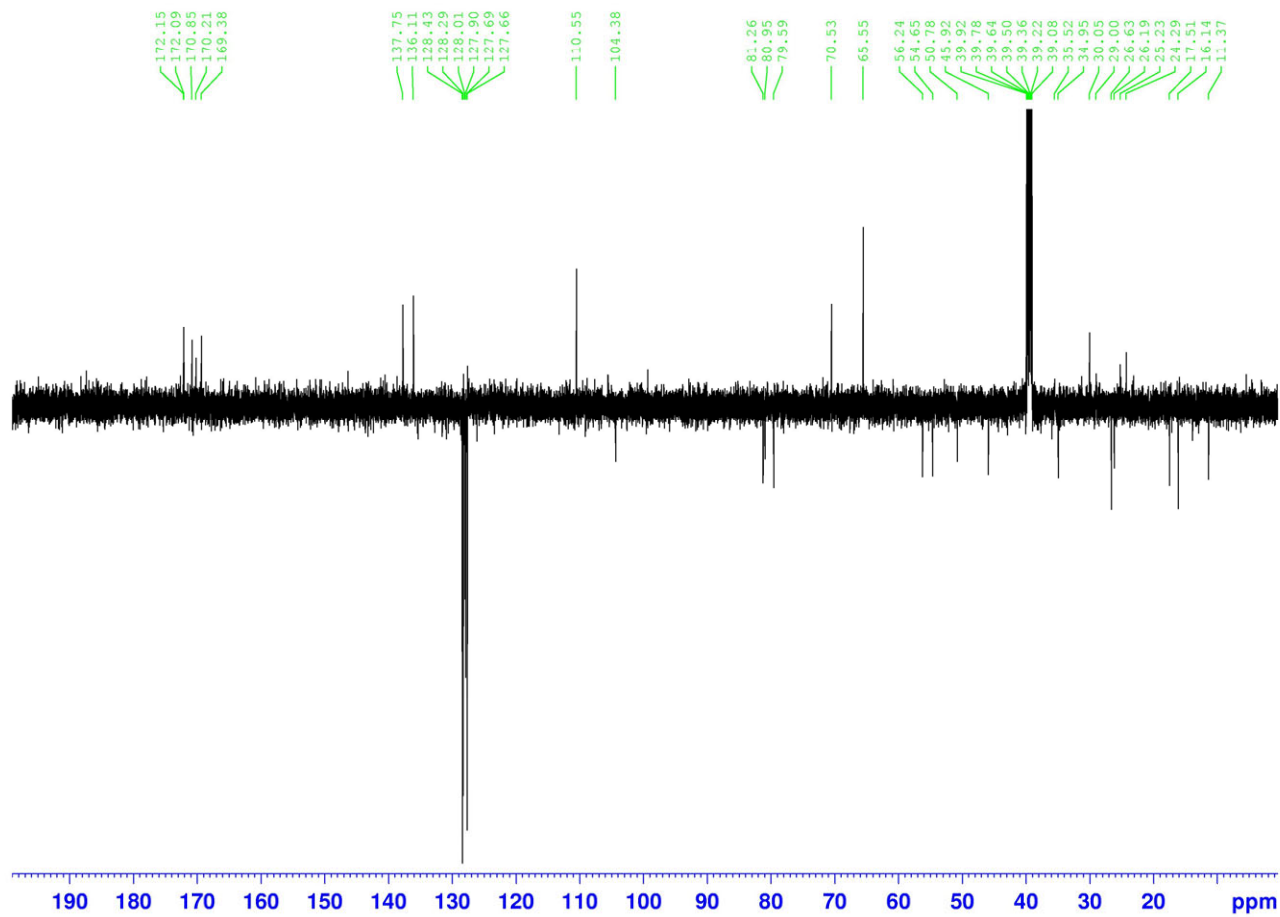


Figure 9 ROESY spectrum of 2b

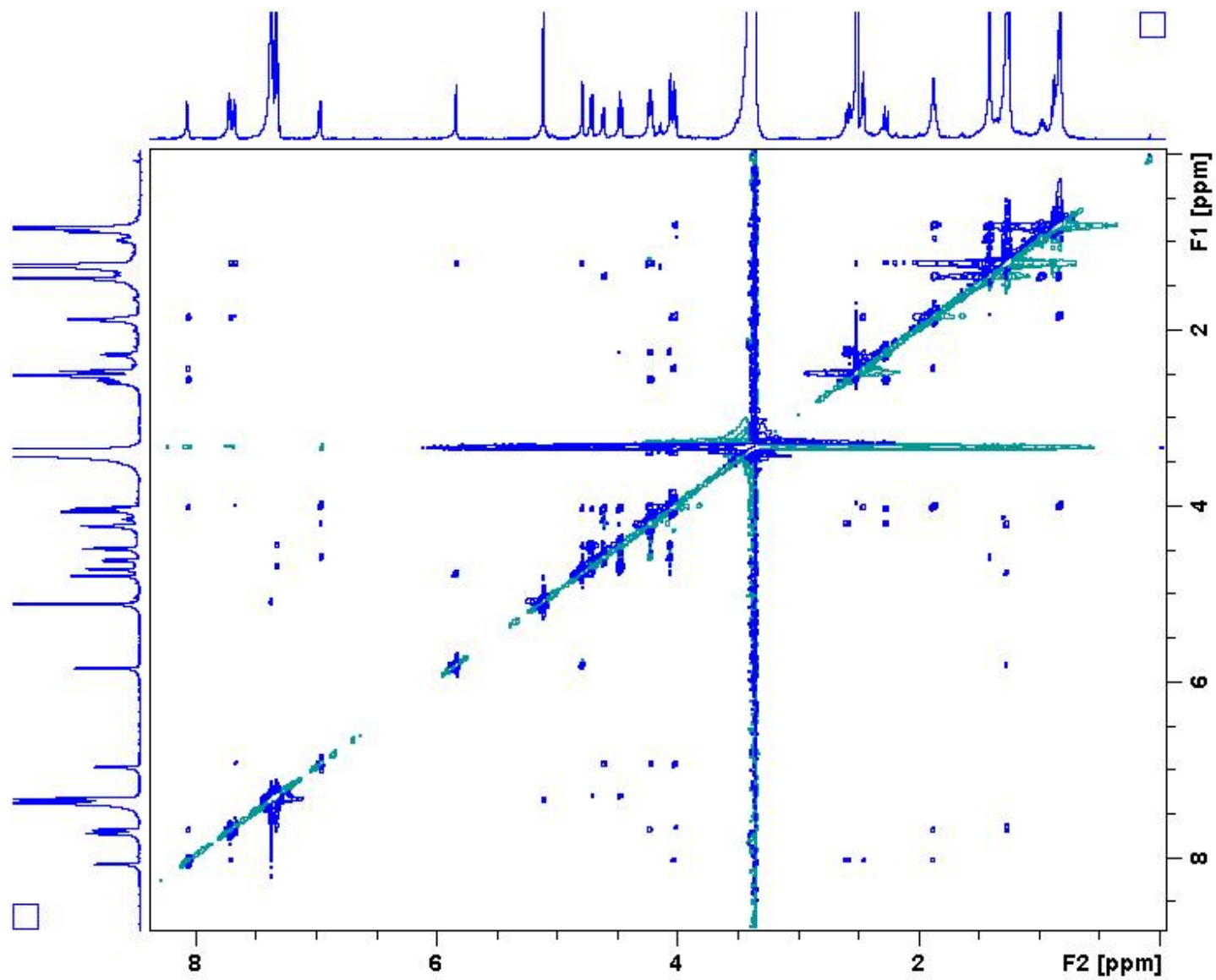


Figure 10 COSY spectrum of 2b

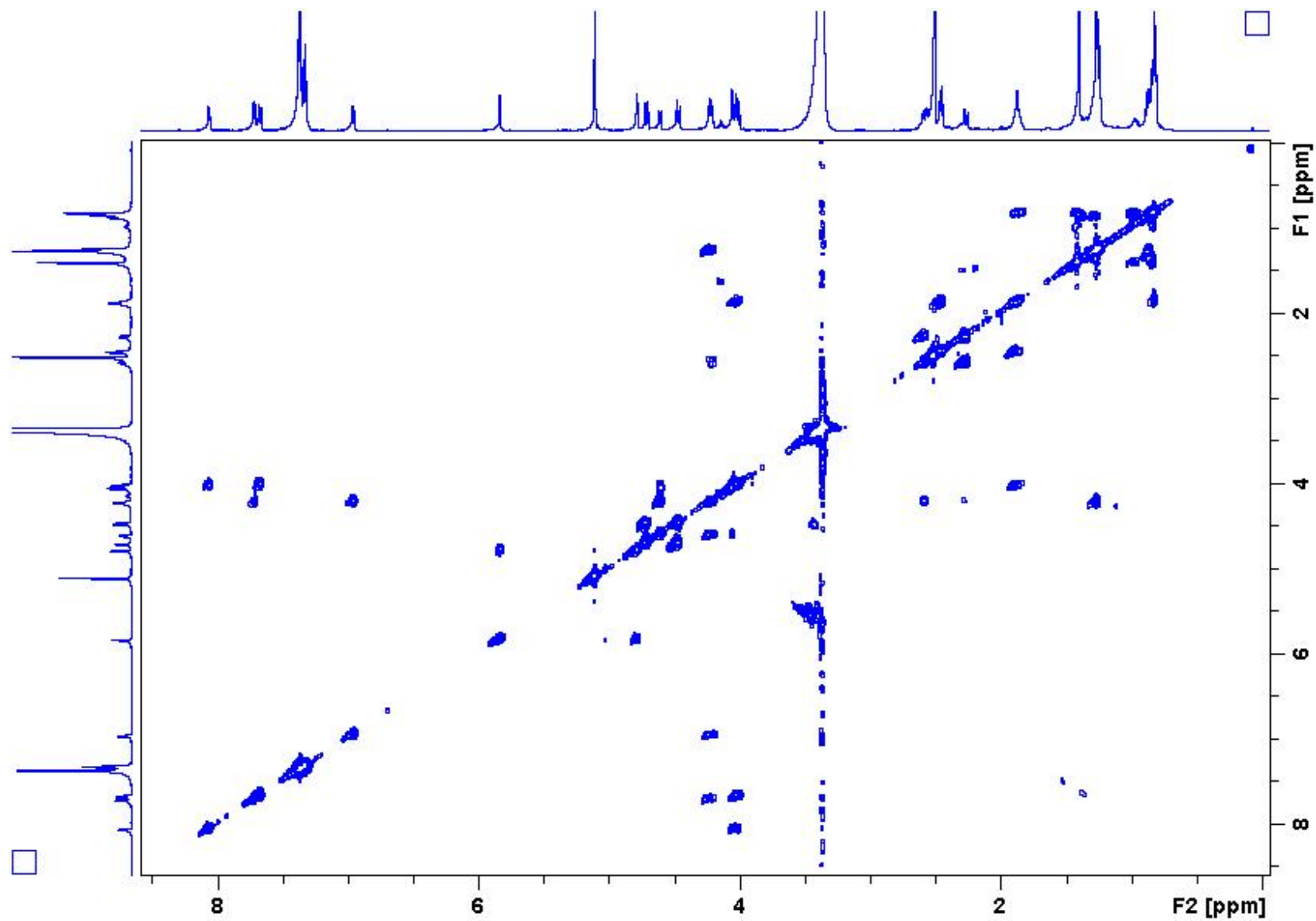


Figure 11 HSQC spectrum of 2b

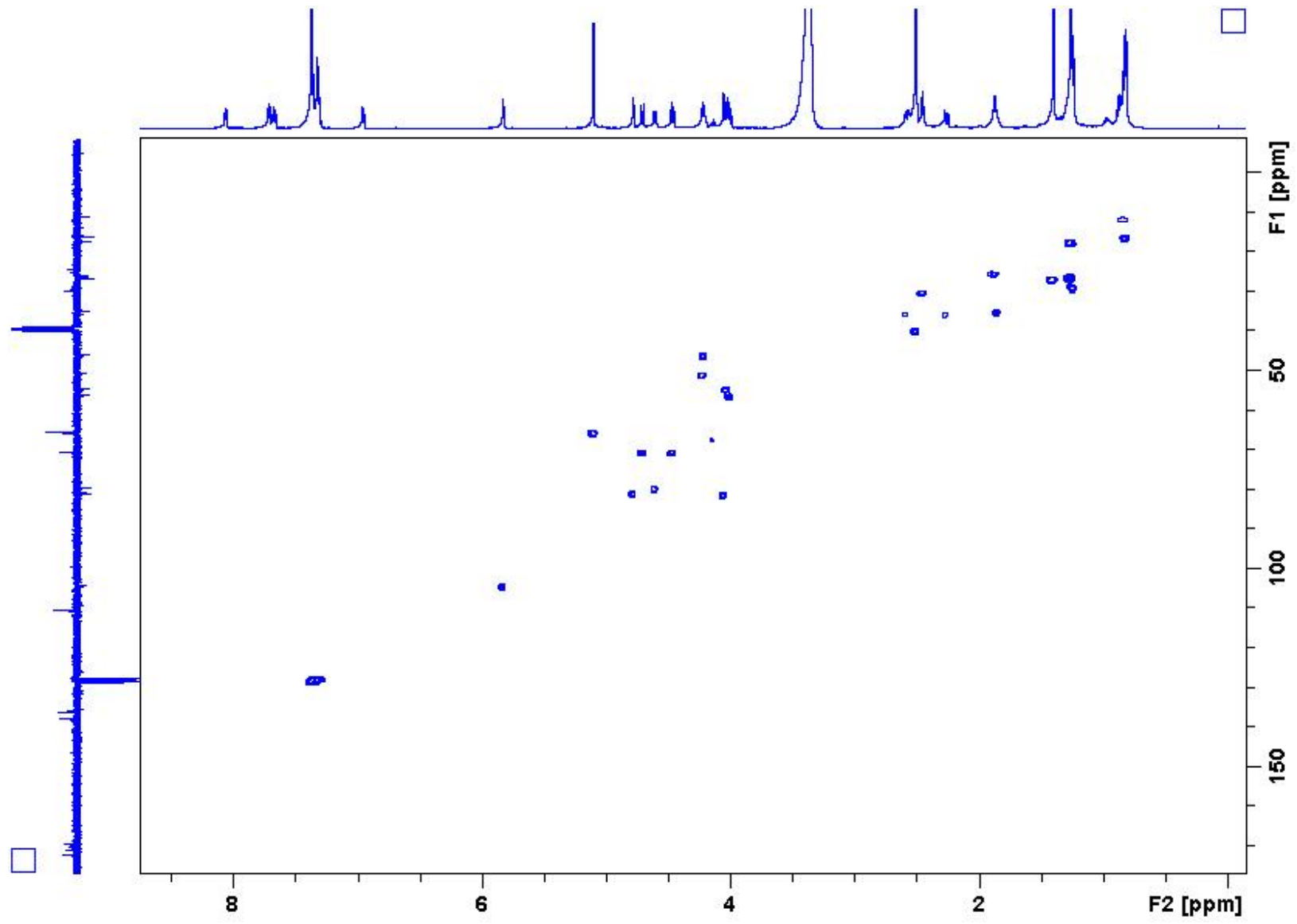


Figure 12 HMBC spectrum of 2b

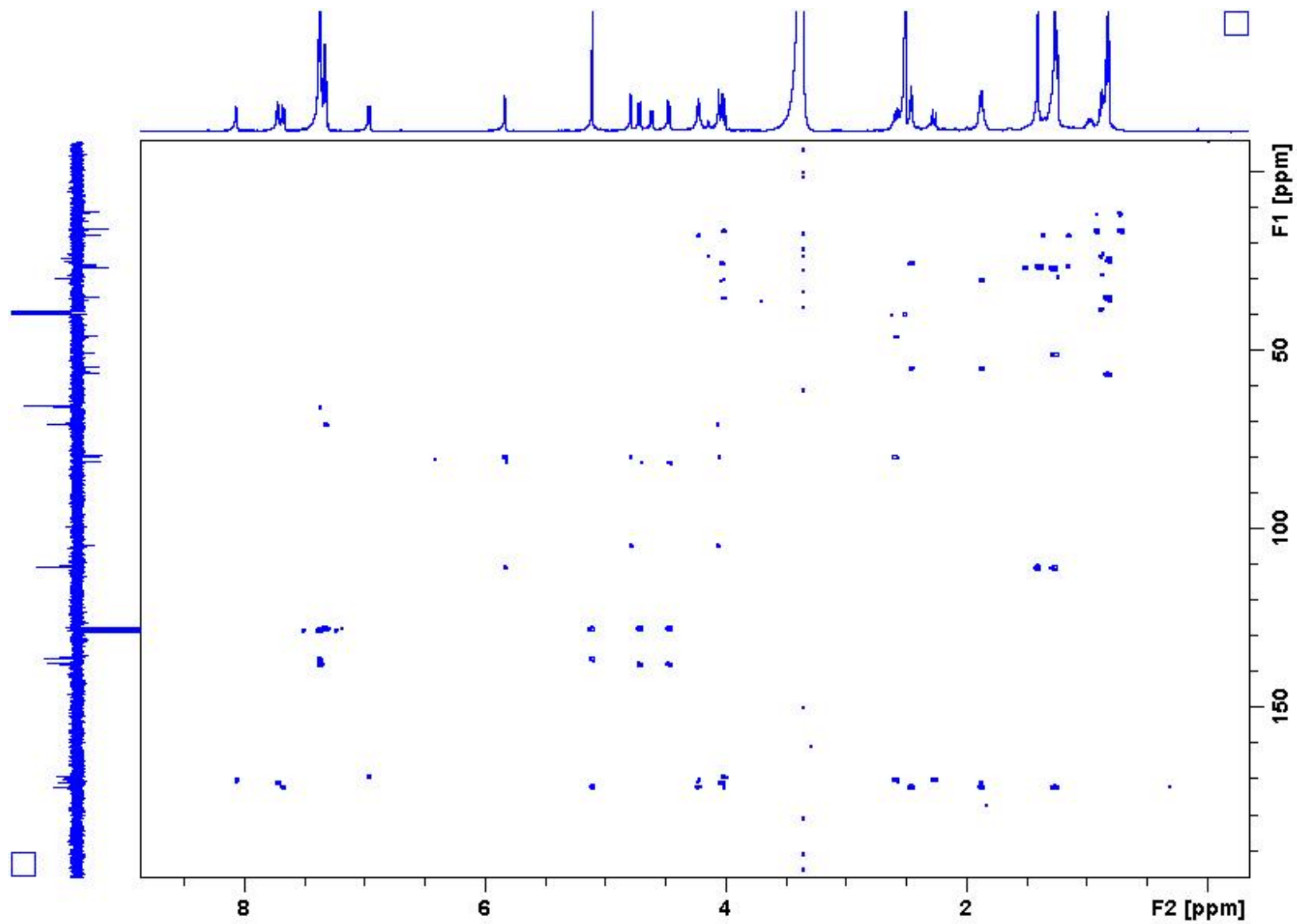


Figure 23 ^1H NMR spectrum of **3**

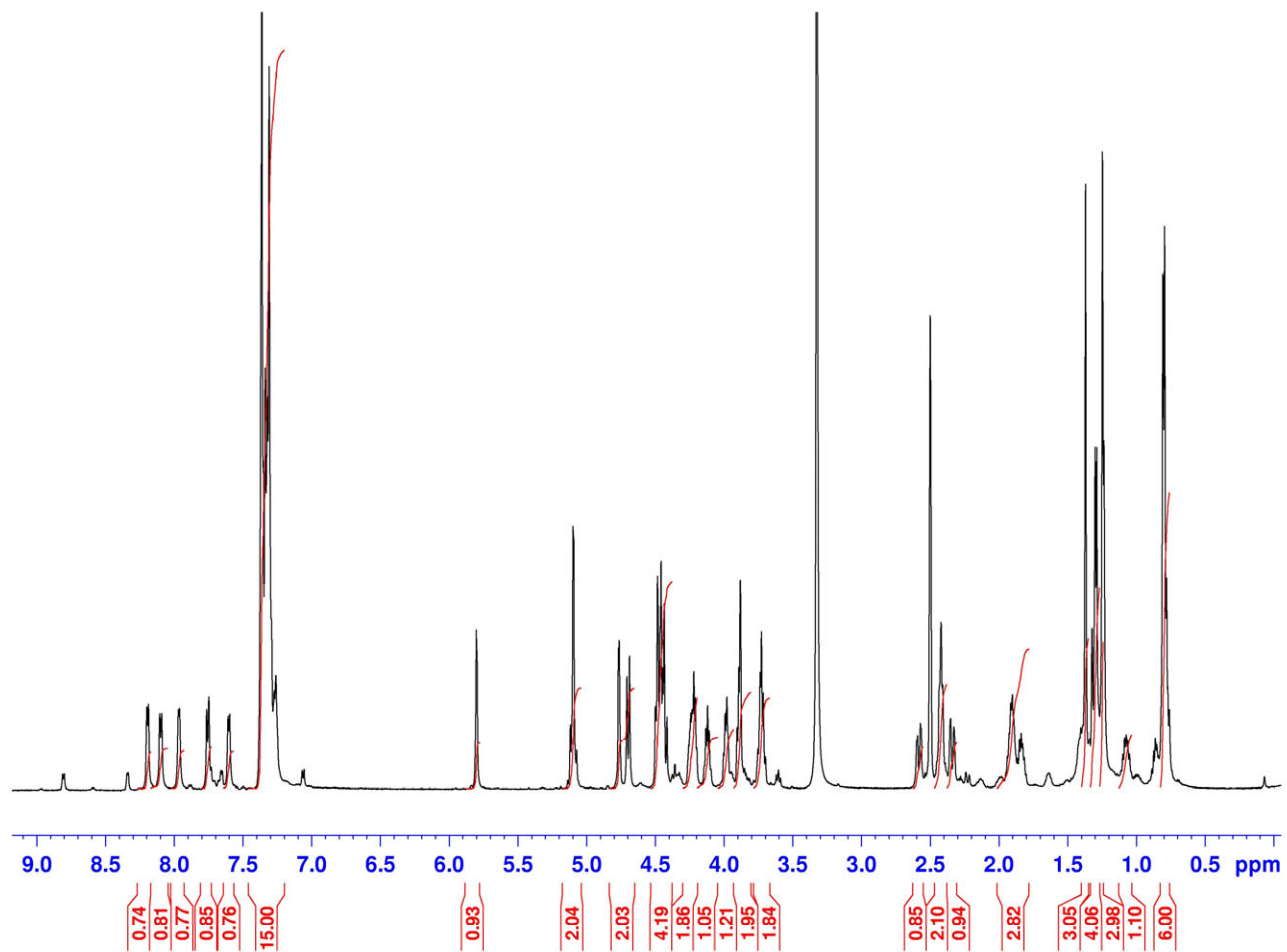


Figure 14 ^{13}C NMR spectrum of **3**

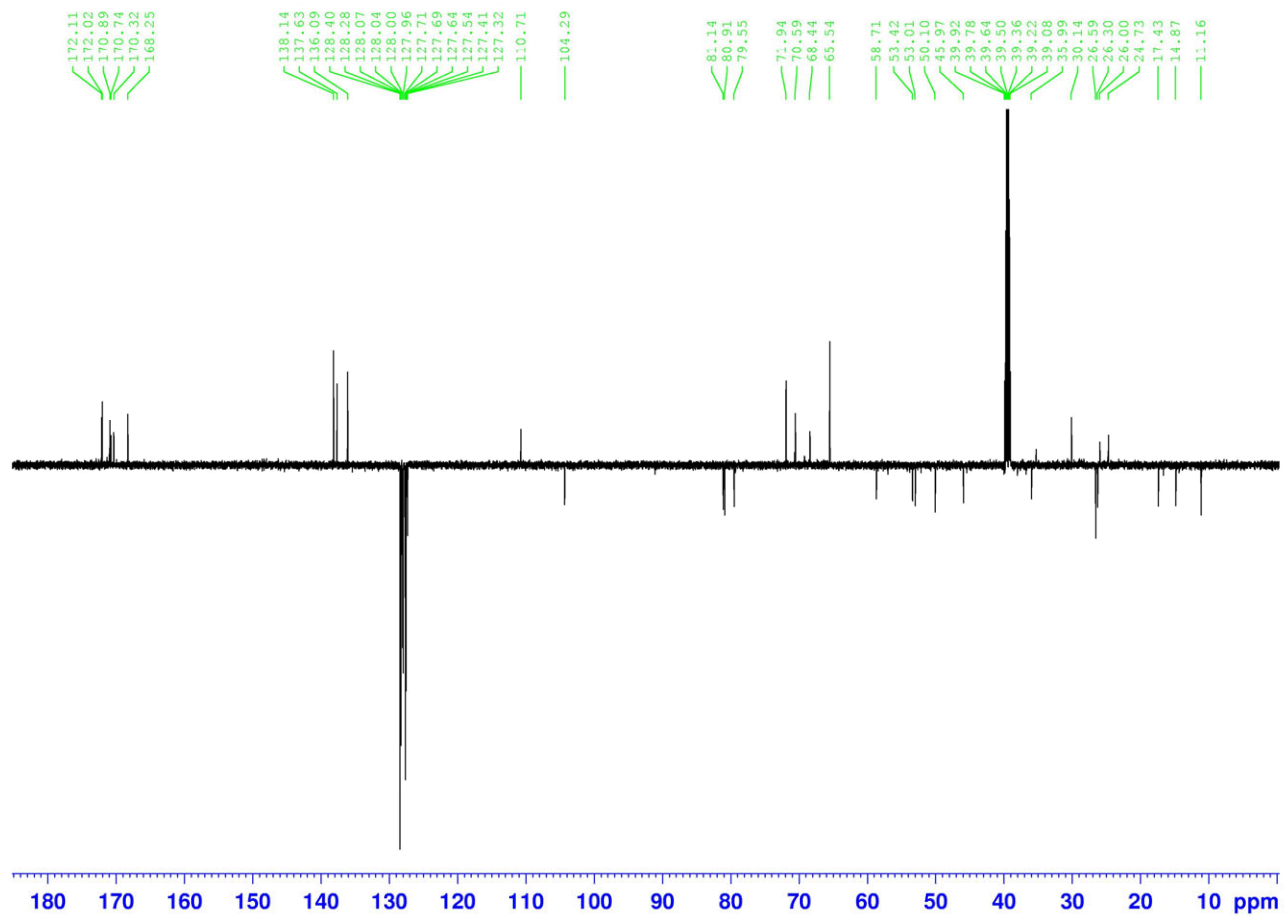


Figure 15 ROESY spectrum of 3

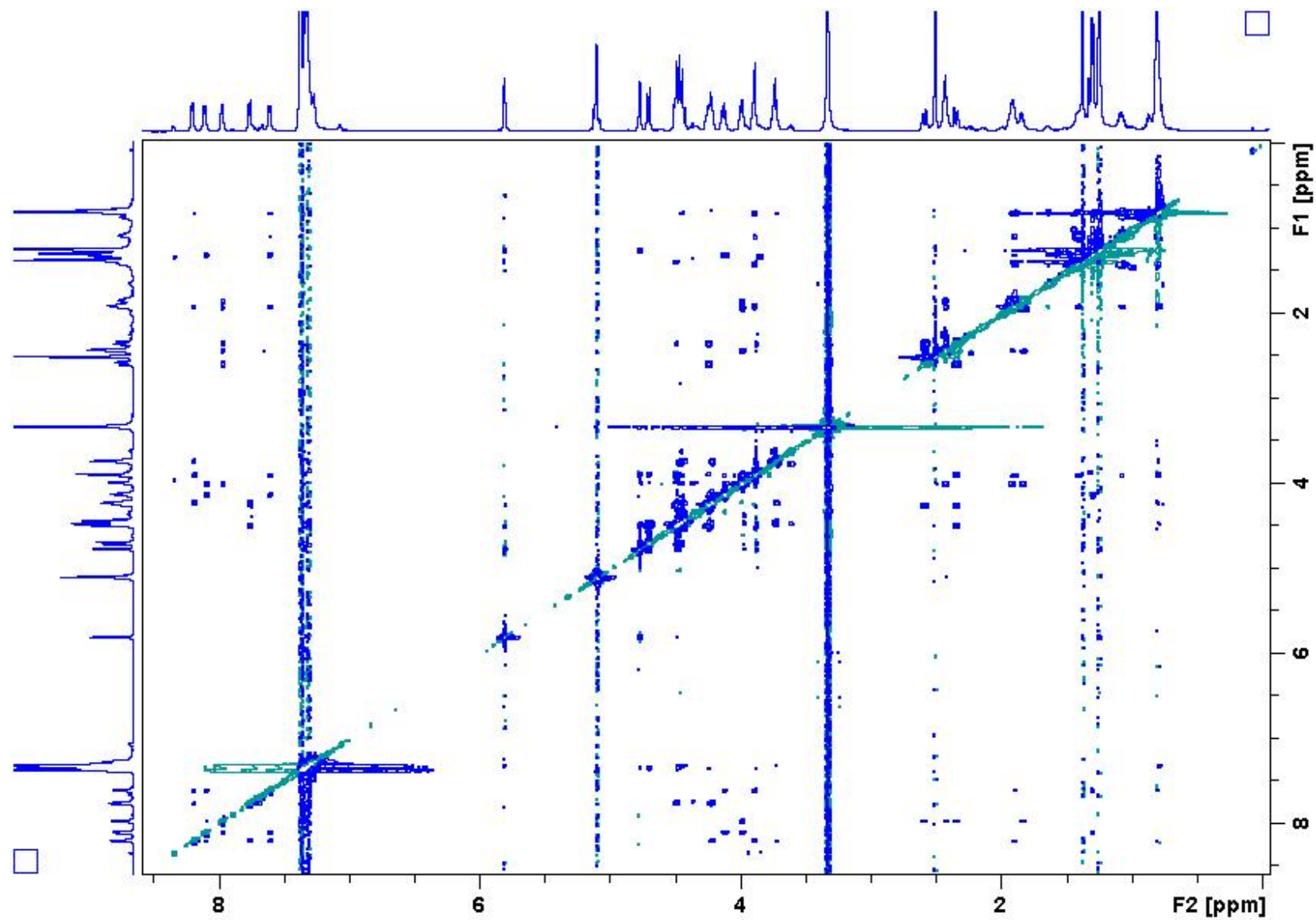


Figure 16 COSY spectrum of 3

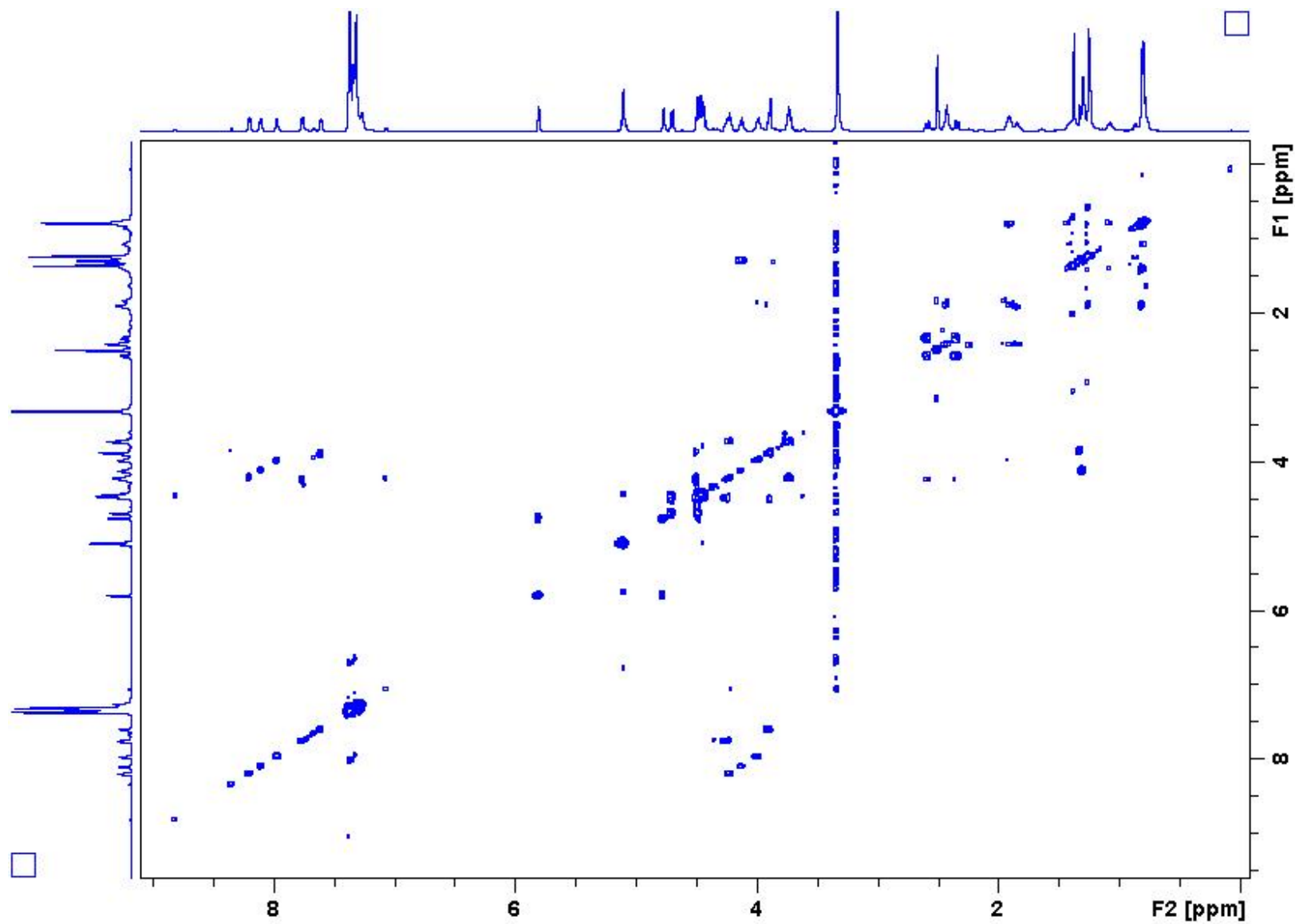


Figure 17 HSQC spectrum of 3

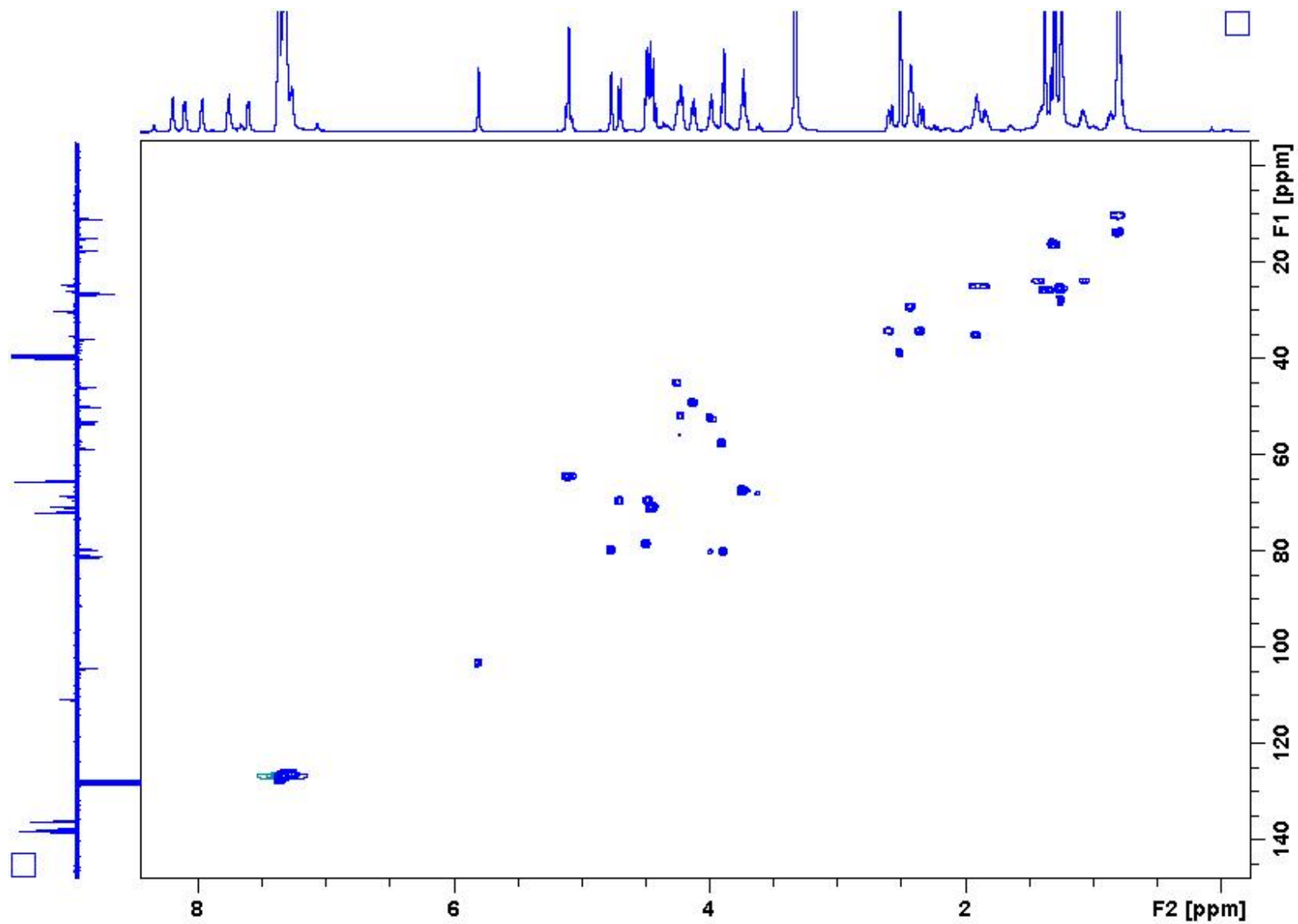


Figure 18 HMBC spectrum of 3

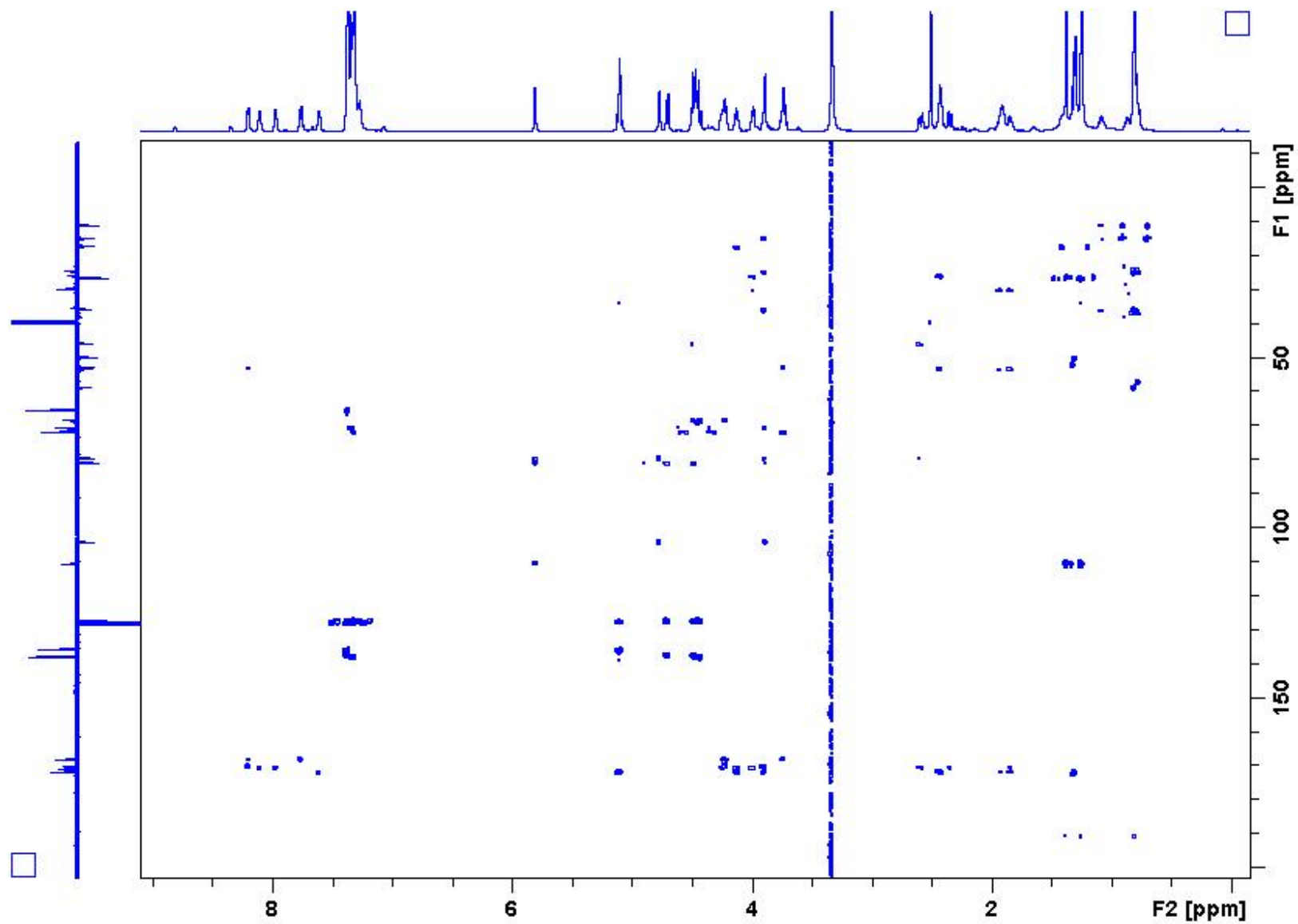


Figure 19 ^1H NMR spectrum of 4

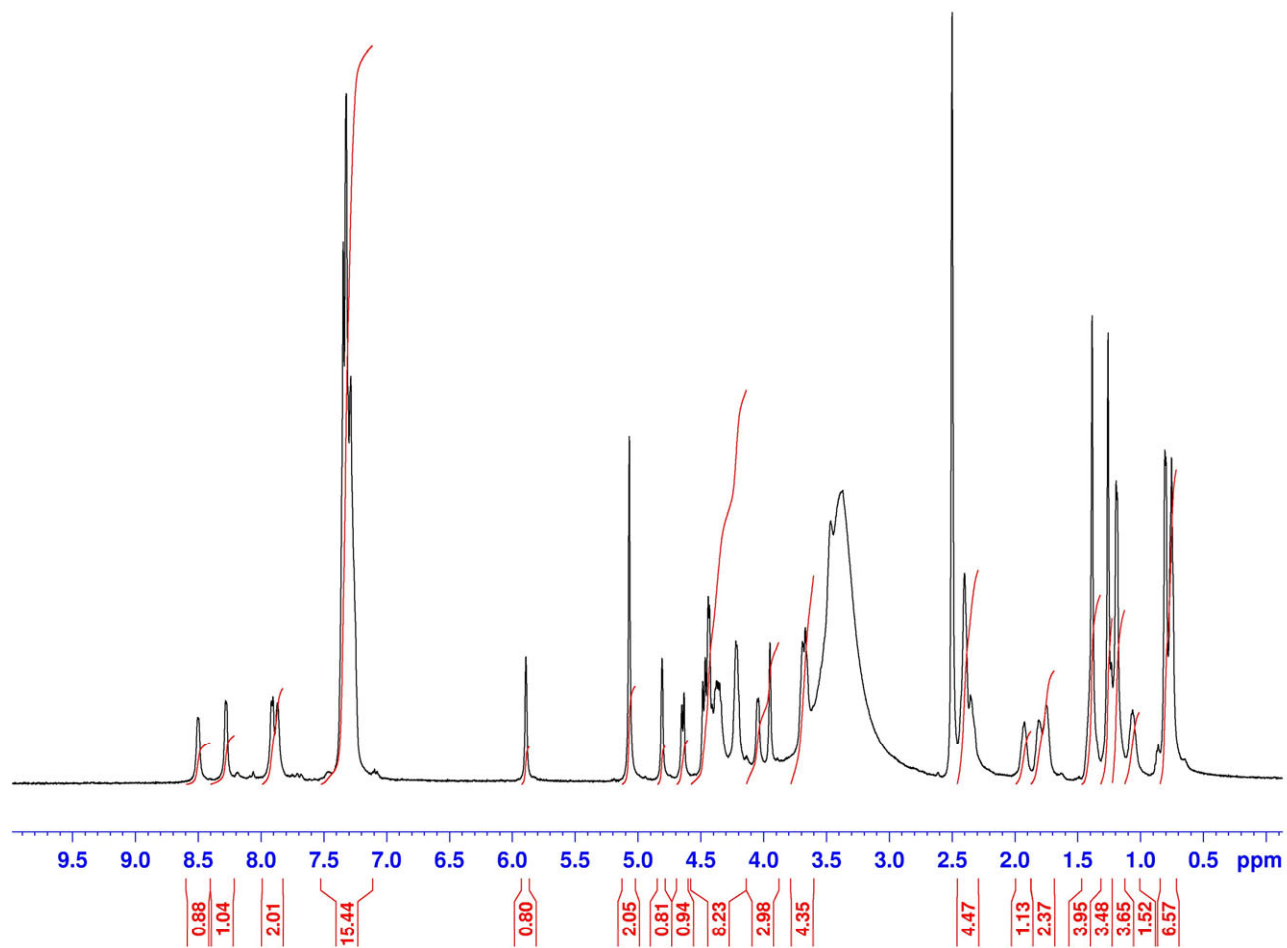


Figure 20 ^{13}C NMR spectrum of 4

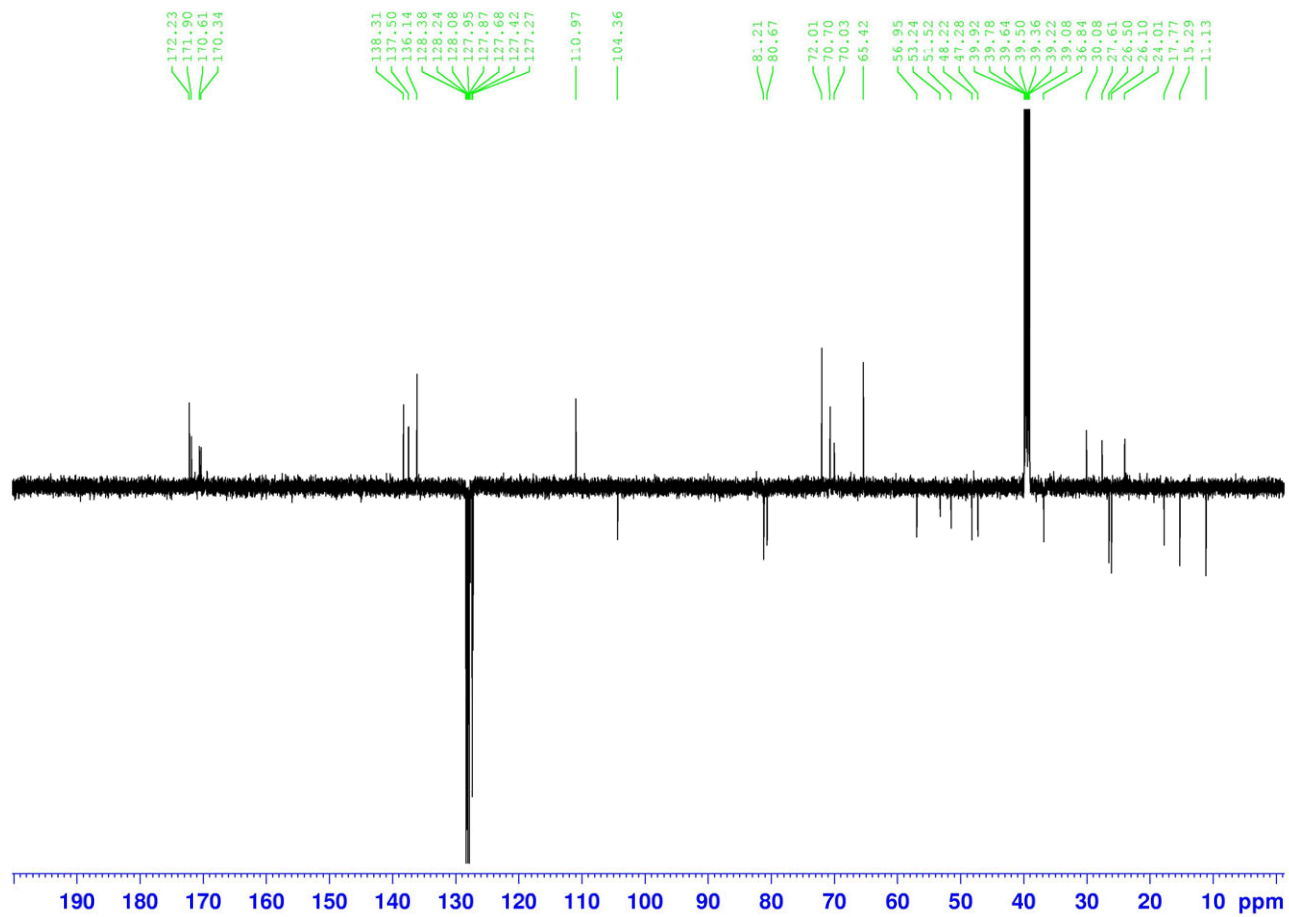


Figure 21 HRMS spectrum of 3

Analysis Info

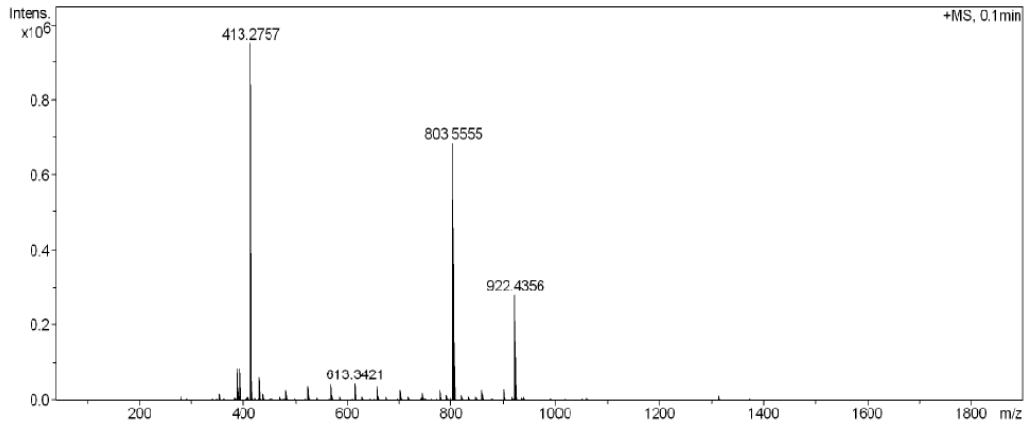
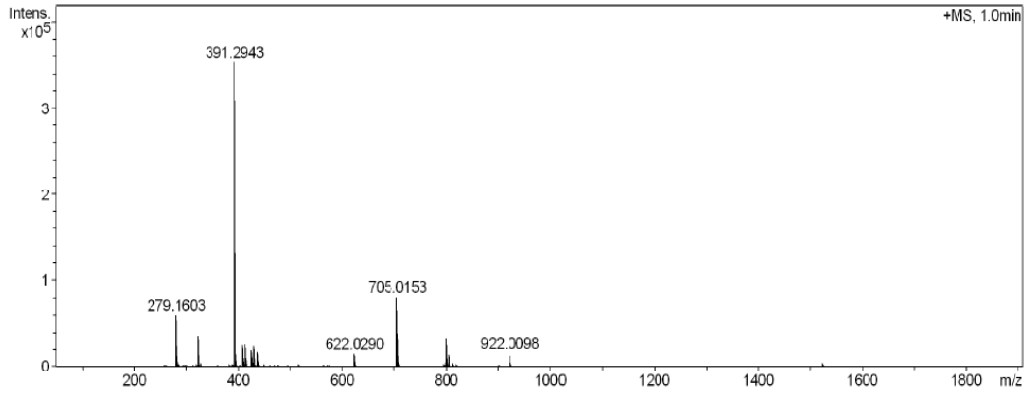
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Comment

Acquisition Date 5/11/2012 4:27:44 PM

Operator BDAL@DE
Instrument / Ser# micrOTOF Q 10139

Acquisition Parameter

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Figure 22 HRMS spectrum of 4

Analysis Info

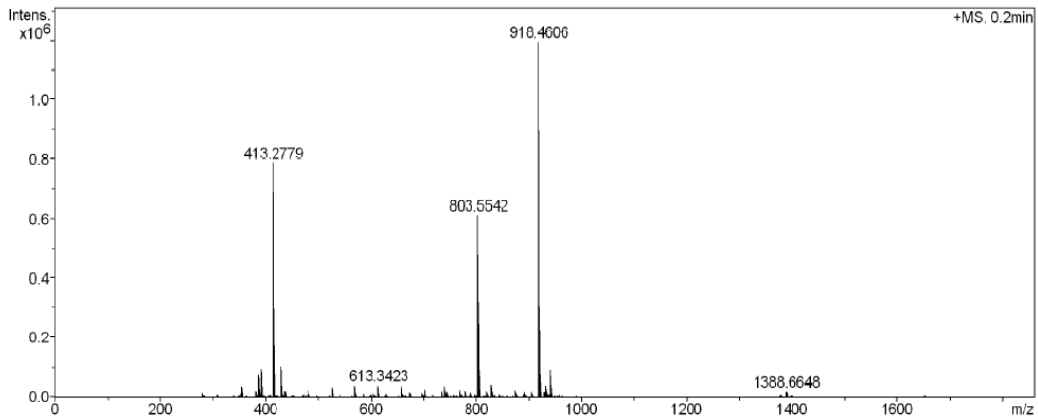
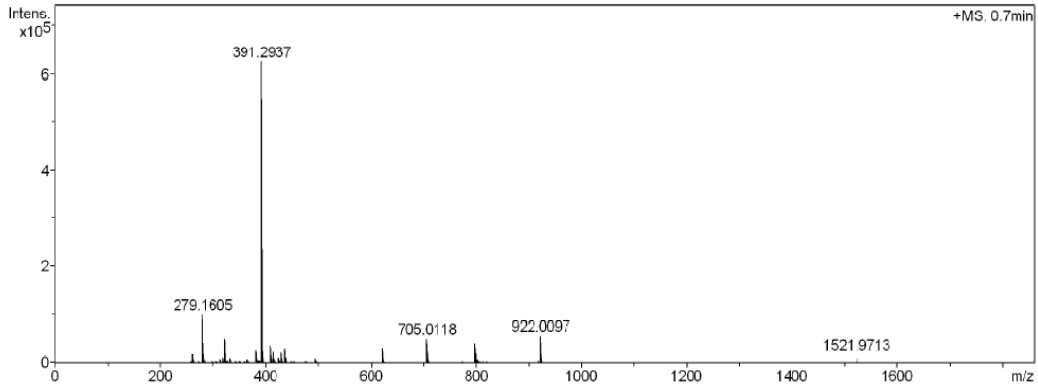
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Operator BDAL@DE
Instrument / Ser# micrOTOF-Q 10139

Acquisition Parameter

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Scan End	1800 m/z	Set Collision Cell RF	500.0 Vpp	Set Divert Valve	Source



Meas. m/z	#	Formula	Score	m/z	err [mDa]	err [ppm]	mSigma	rdb	e ⁻ Cont	N-Rule
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