

Enhancing structural diversity of lathyrane derivatives through biotransformation by the marine-derived actinomycete *Streptomyces puniceus* BC-5GB.11

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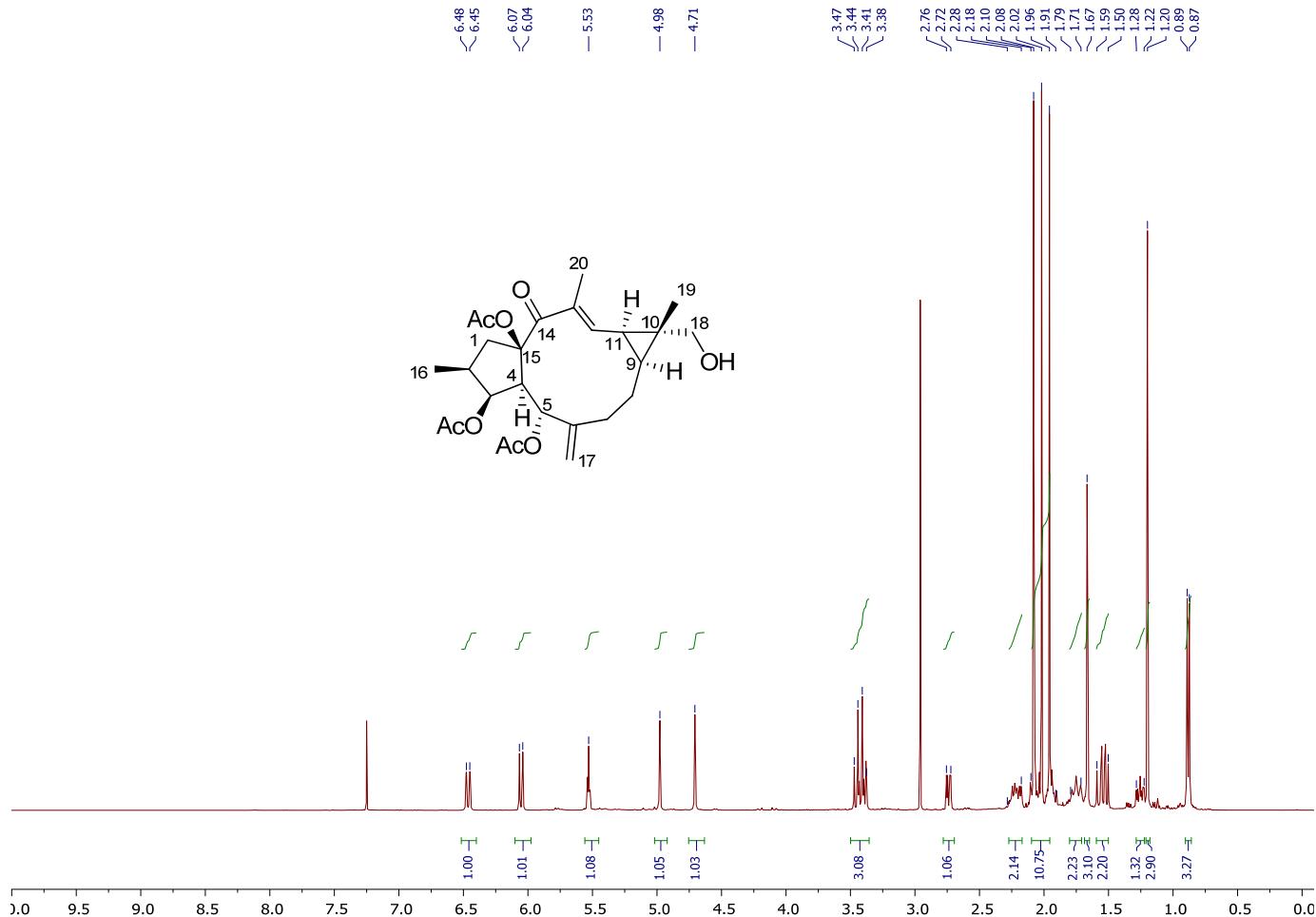


Figure S1. ^1H NMR spectrum (400 MHz) of compound 3 in CDCl_3 .

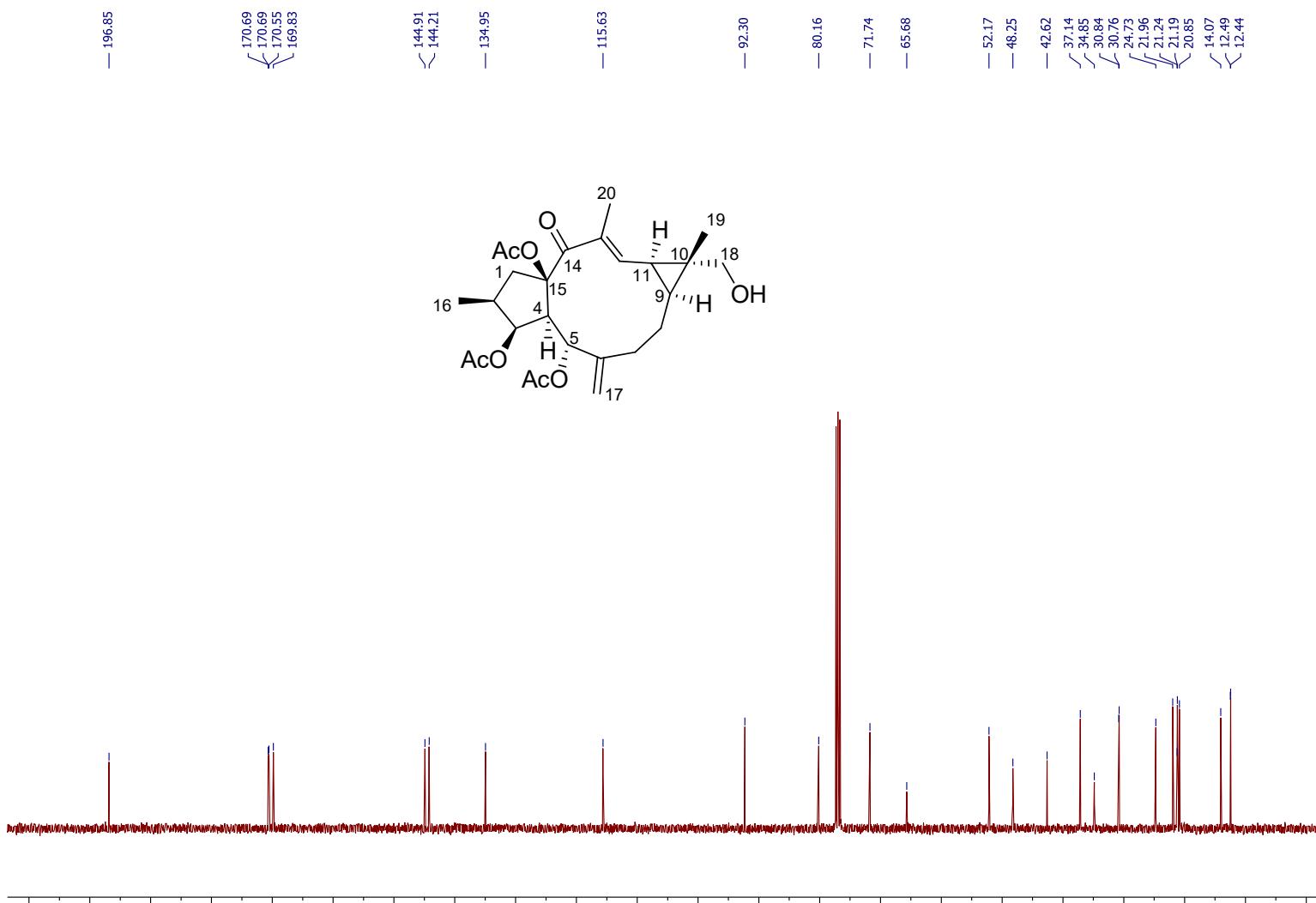


Figure S2. ^{13}C NMR spectrum (100 MHz) of compound 3 in CDCl_3 .

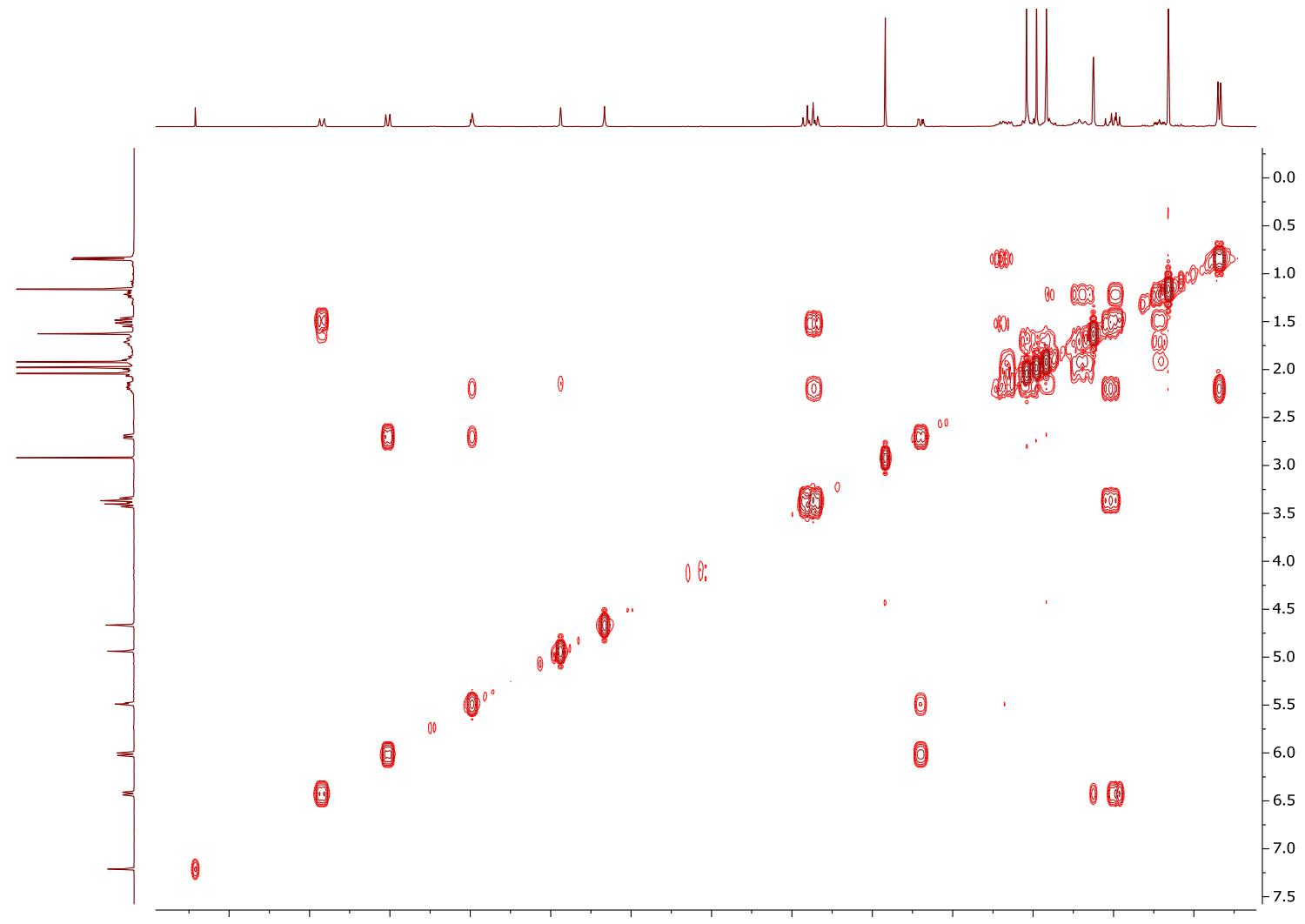


Figure S3. gCOSY spectrum of compound 3.

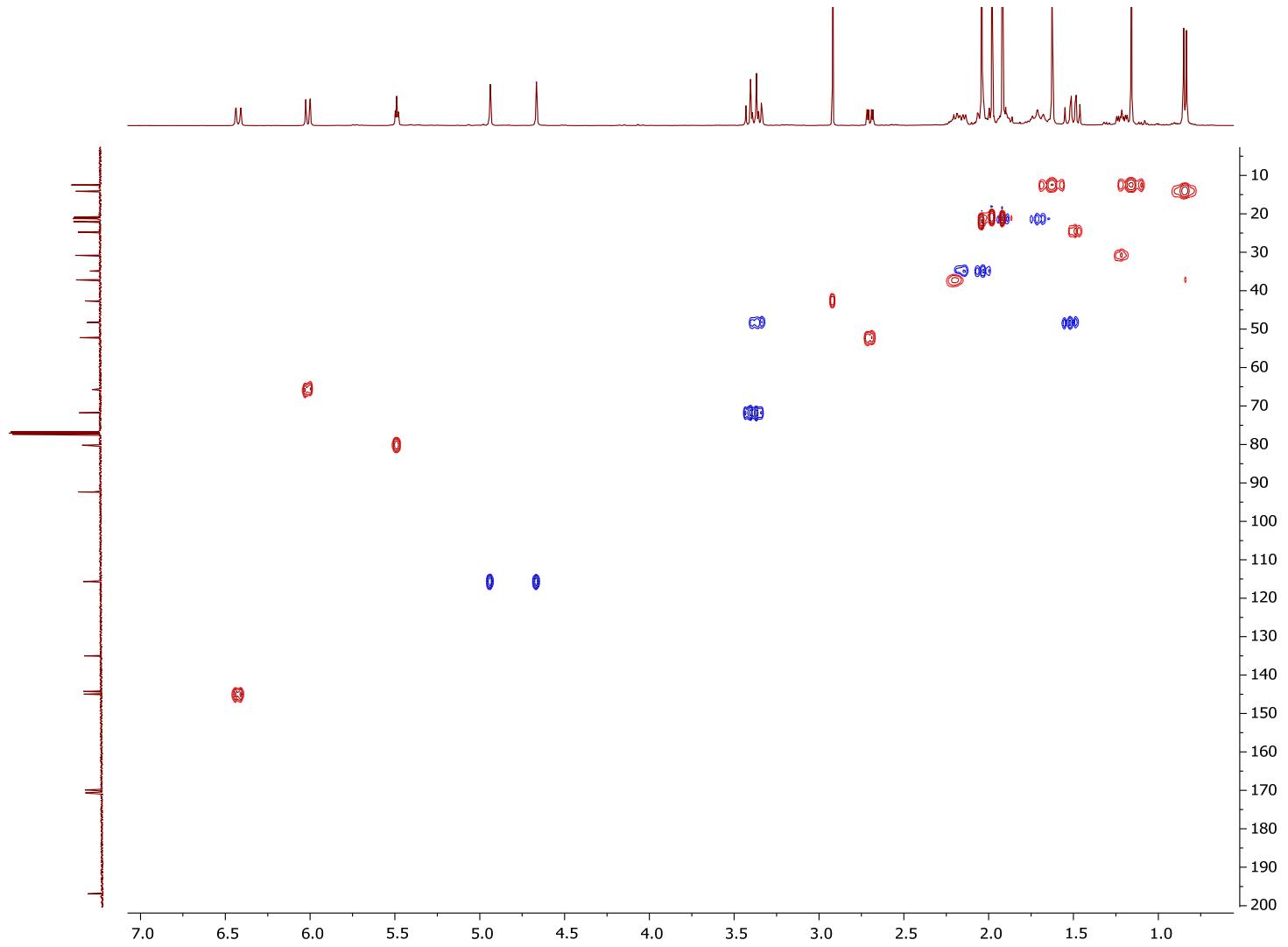


Figure S4. gHSQC spectrum of compound 3.

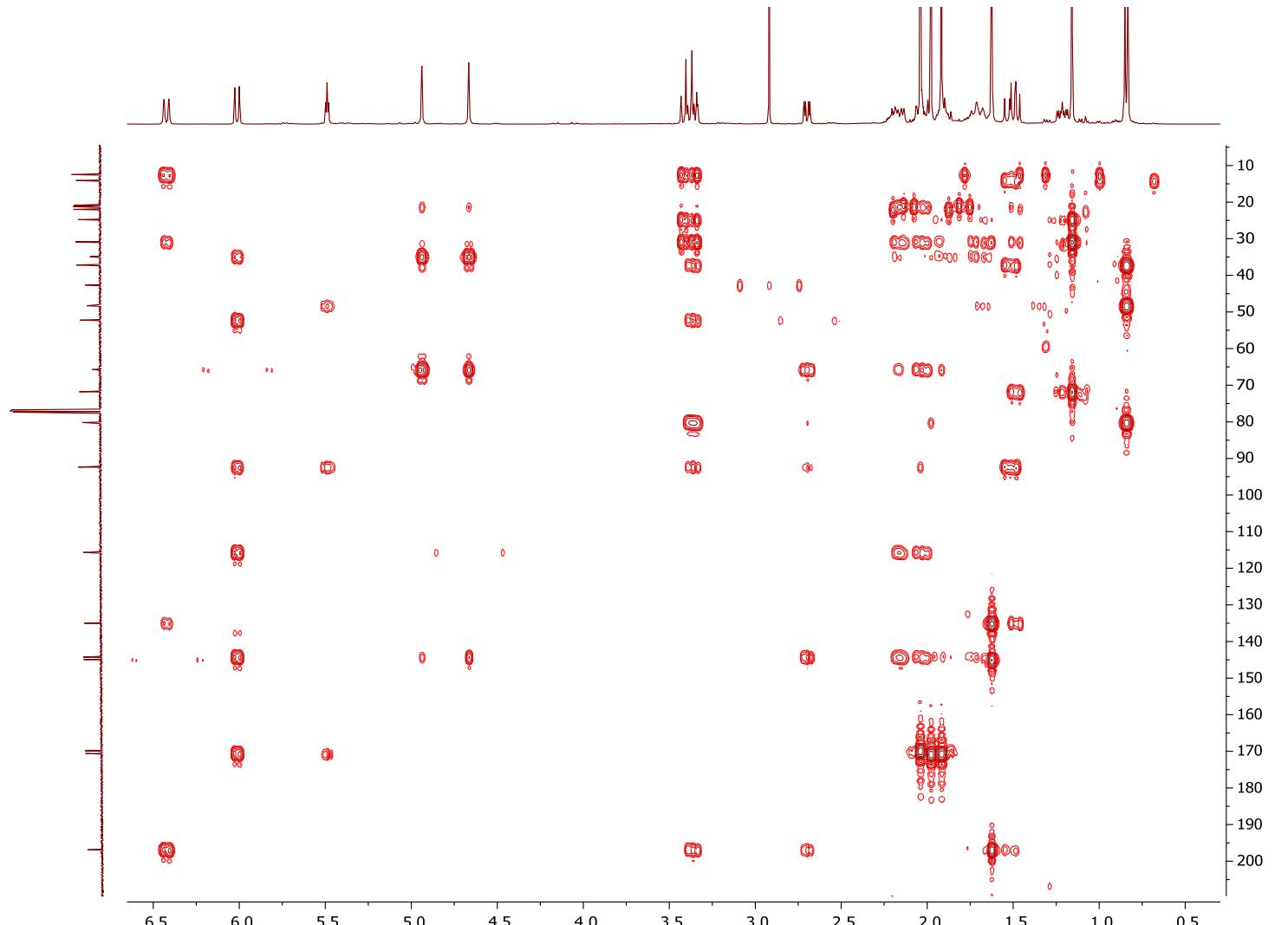


Figure S5. gHMBC spectrum of compound 3.

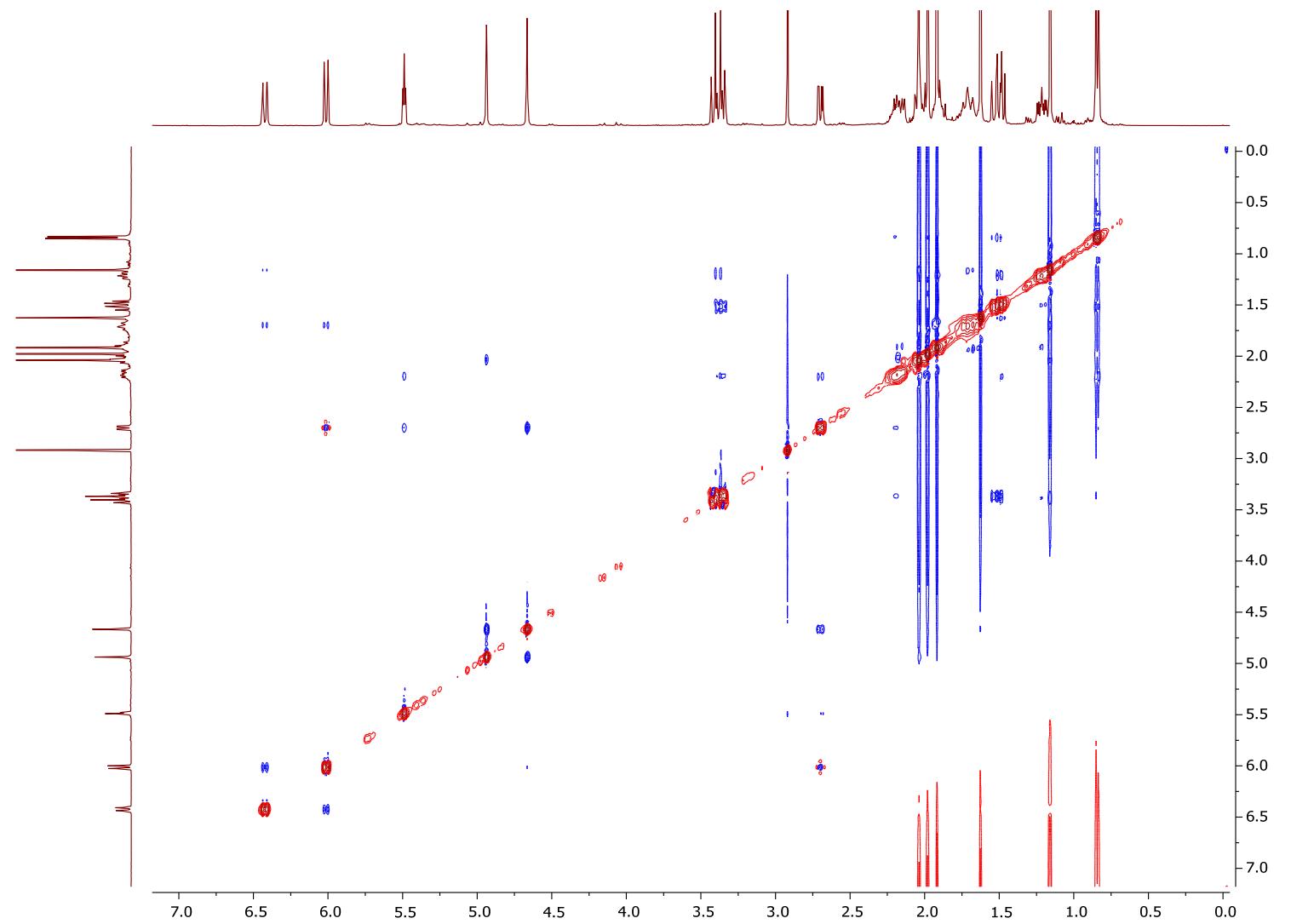


Figure S6. 2D NOESY spectrum of compound 3.

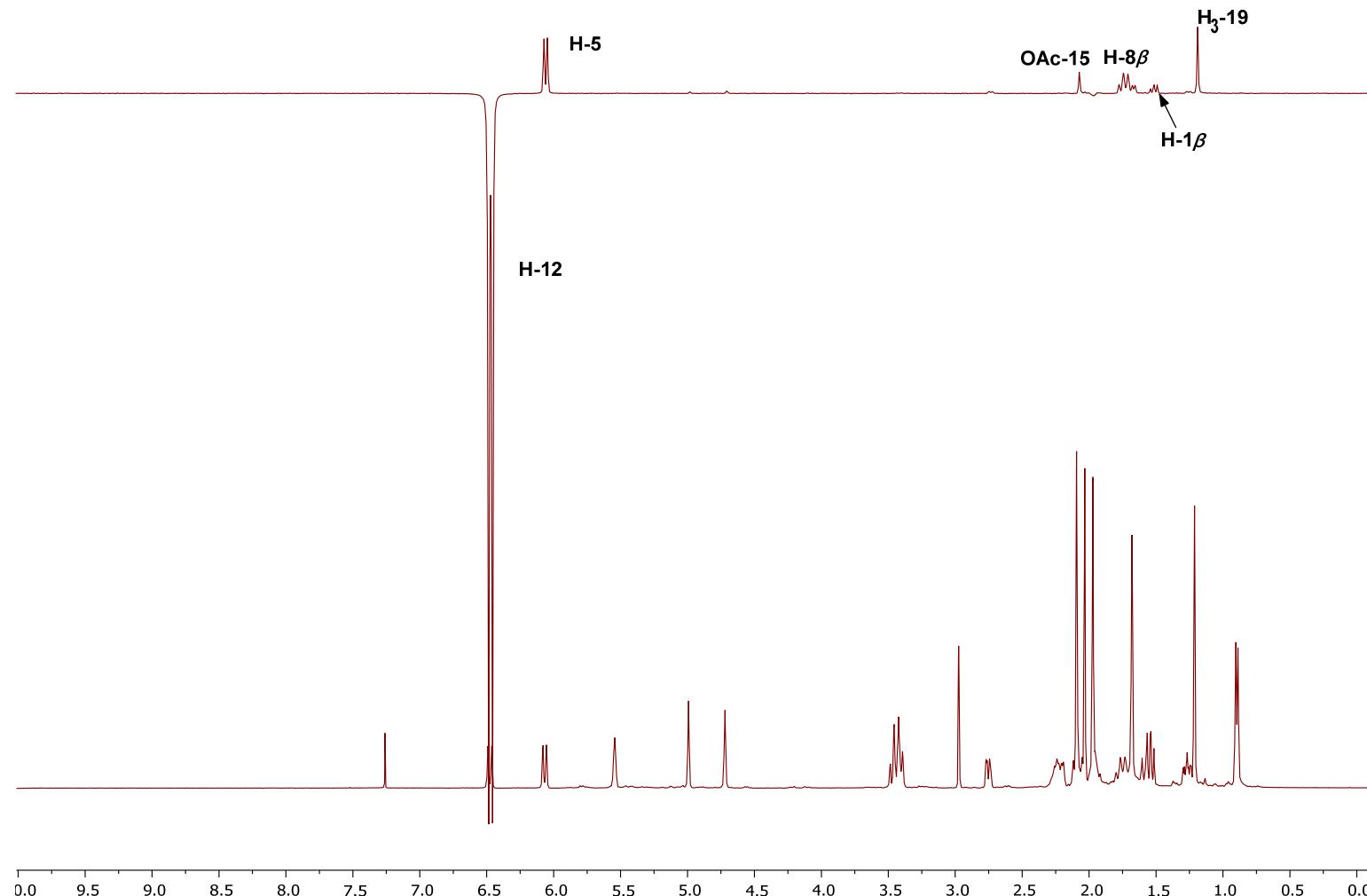


Figure S7a. 1D NOESY pectrum of compound 3.

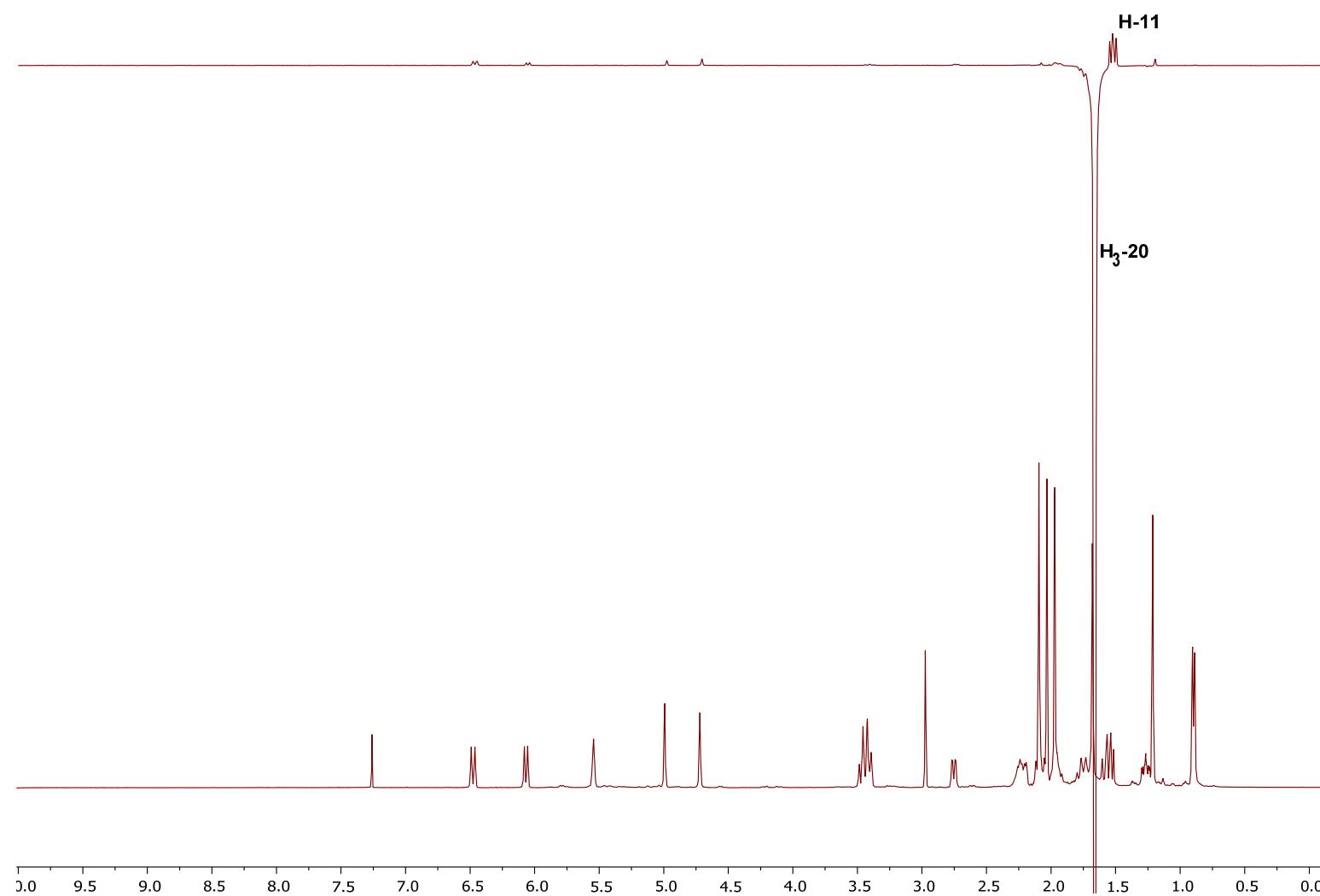


Figure S7b. 1D NOESY spectrum of compound 3.

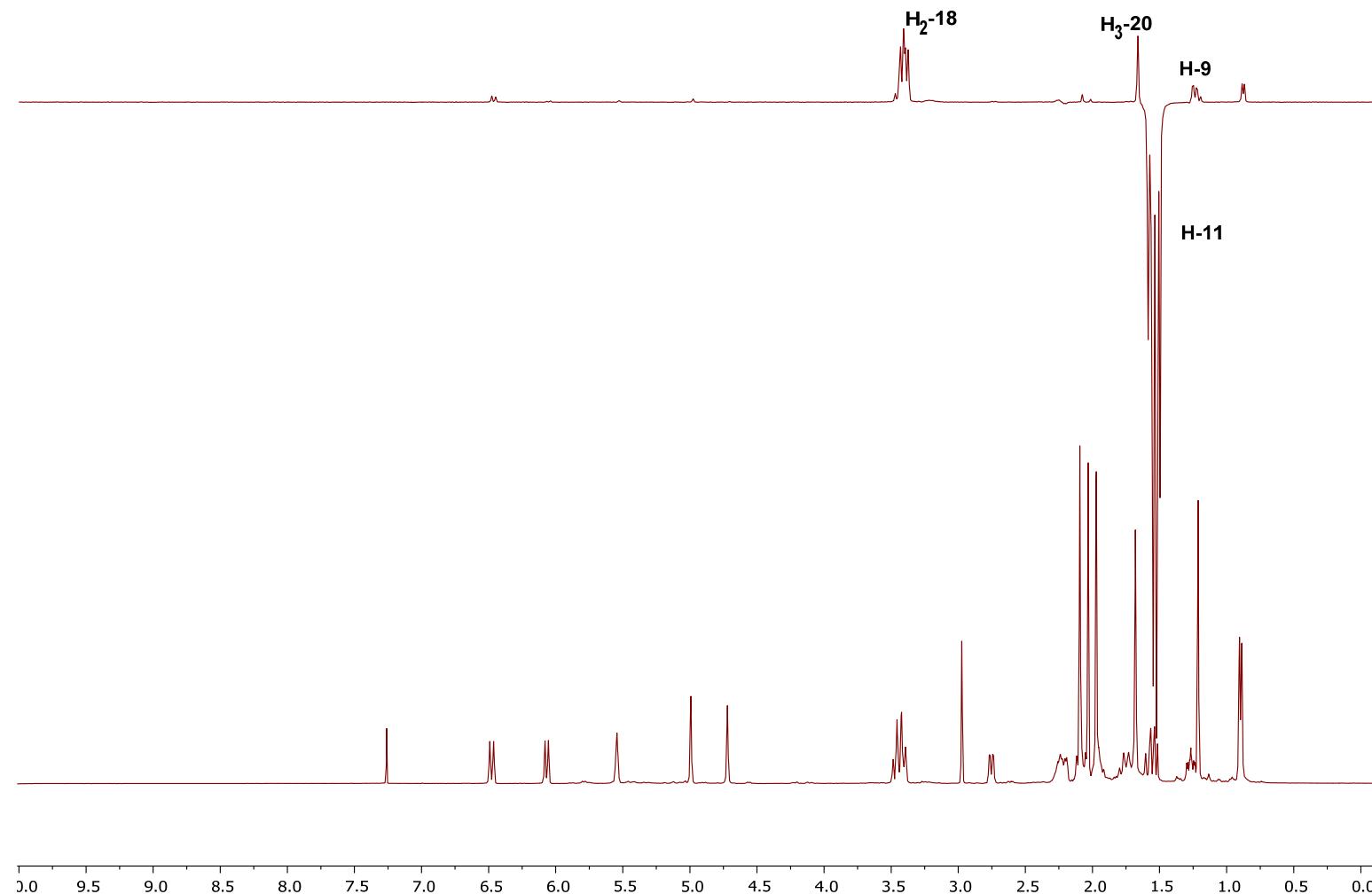


Figure S7c. 1D NOESY spectrum of compound 3.

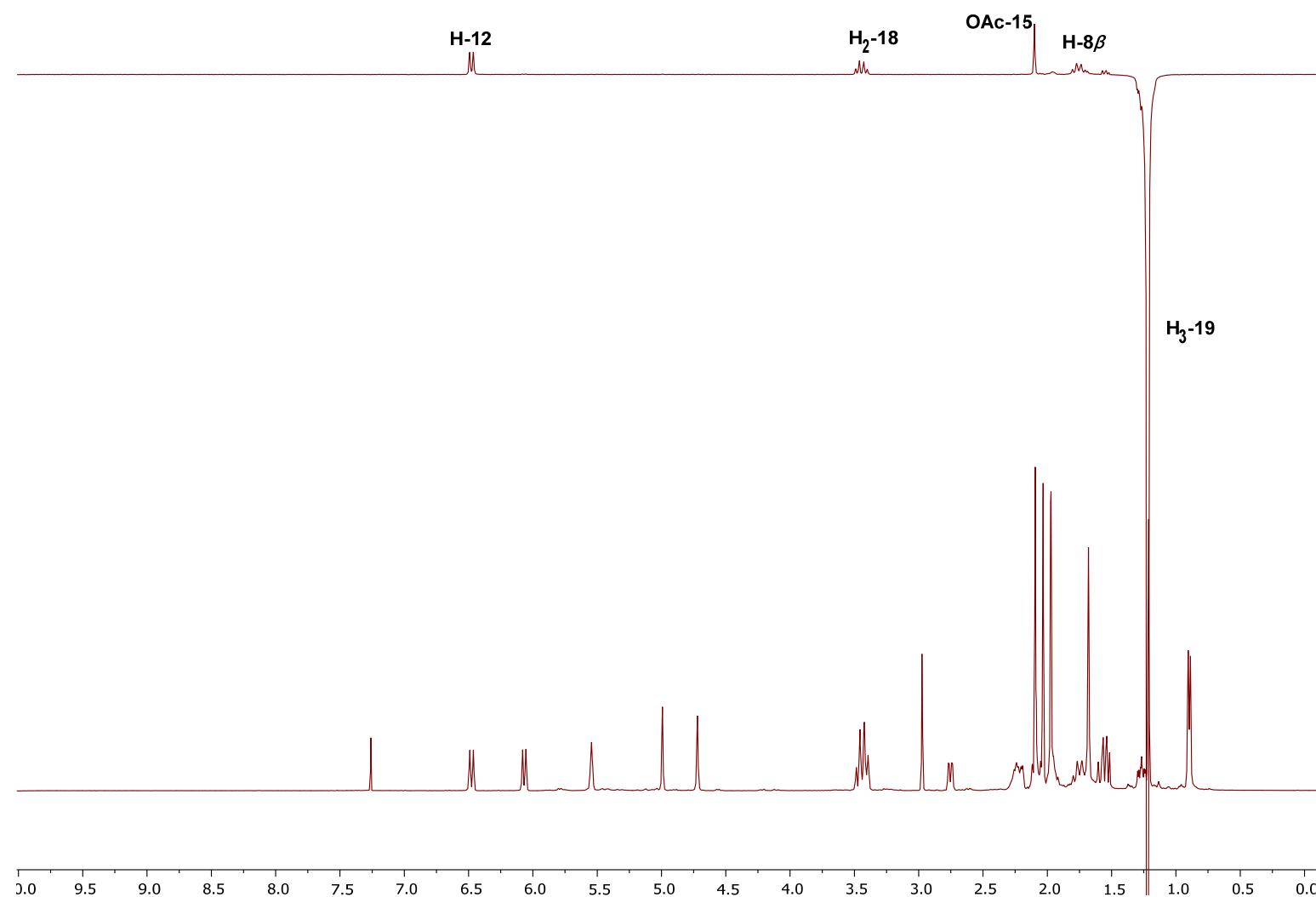


Figure S7d. 1D NOESY spectrum of compound 3.

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

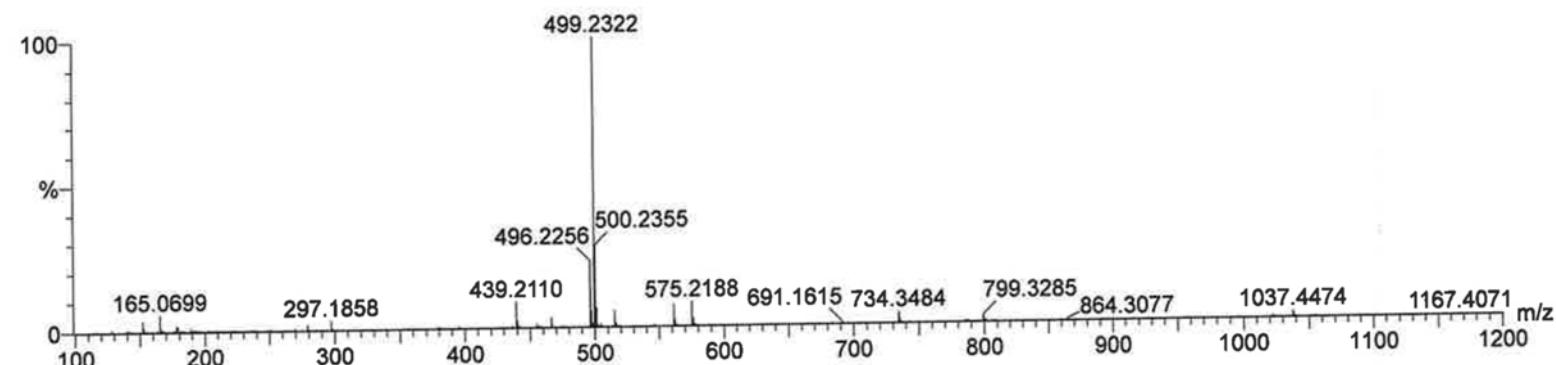
Monoisotopic Mass, Even Electron Ions

87 formula(e) evaluated with 3 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 0-30 H: 0-50 O: 0-15 23Na: 0-1

243_953_Strep-EB-12-MSe3pos 65 (1.215)

1: TOF MS ES+
3.53e+006

Minimum:	-1.5		
Maximum:	5.0	10.0	80.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
499.2322	499.2308	1.4	2.8	8.5	1690.9	0.198	82.03	C26 H36 O8 23Na
	499.2332	-1.0	-2.0	11.5	1692.4	1.719	17.92	C28 H35 O8
	499.2367	-4.5	-9.0	-0.5	1698.2	7.489	0.06	C19 H40 O13 23Na

Figure S8. HRMS of compound 3.

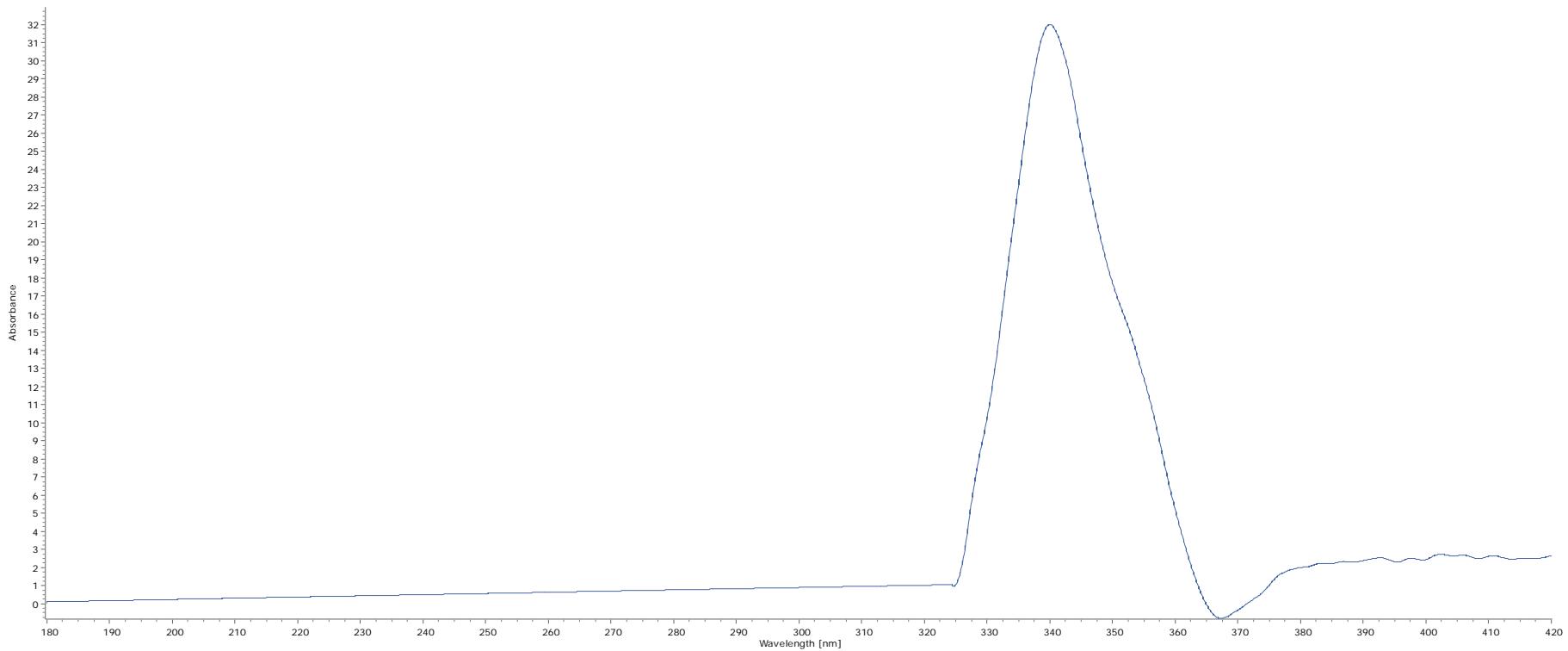


Figure S9. ECD of compound 3.

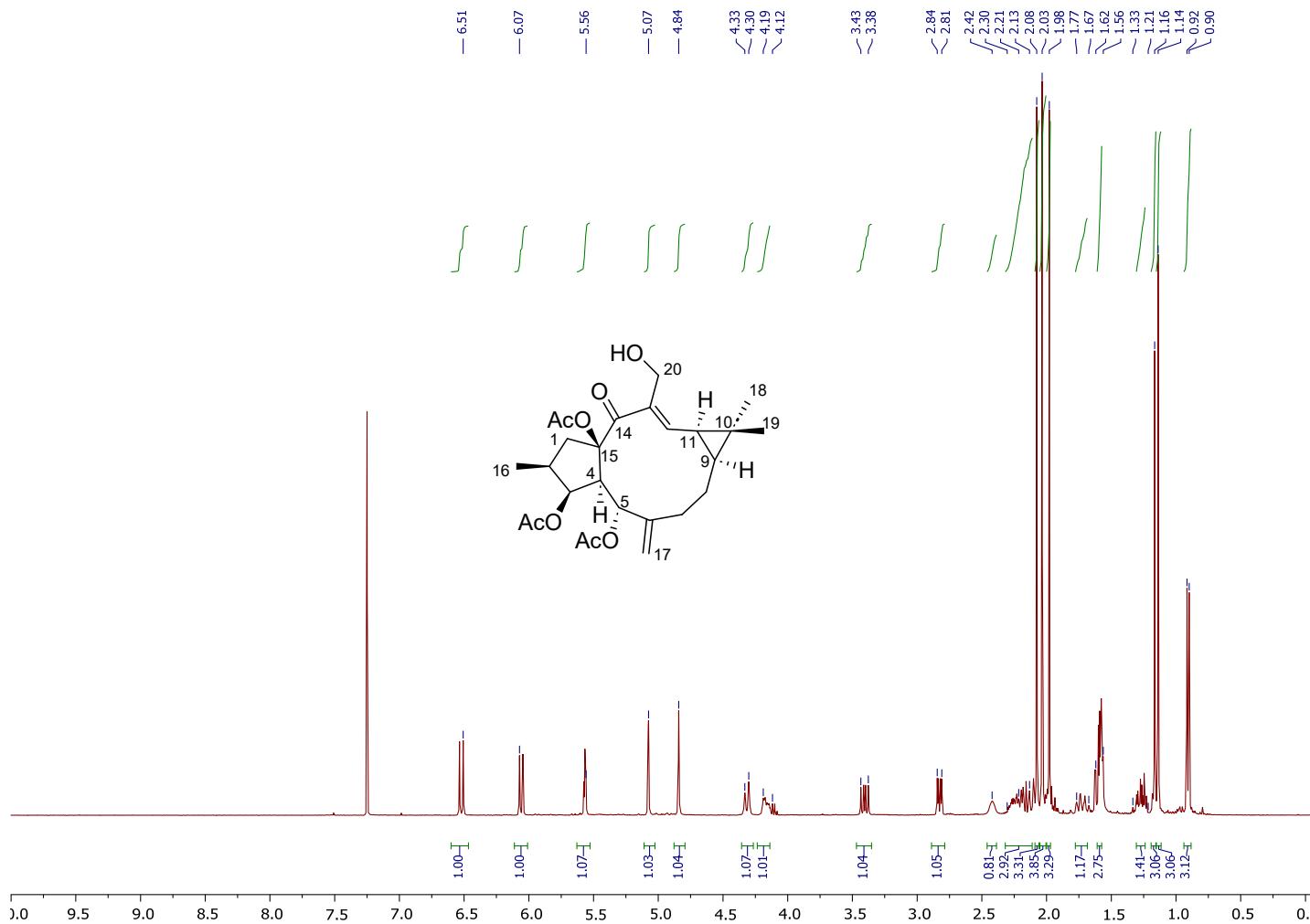


Figure S10. ^1H NMR spectrum (400 MHz) of compound 4 in CDCl_3 .

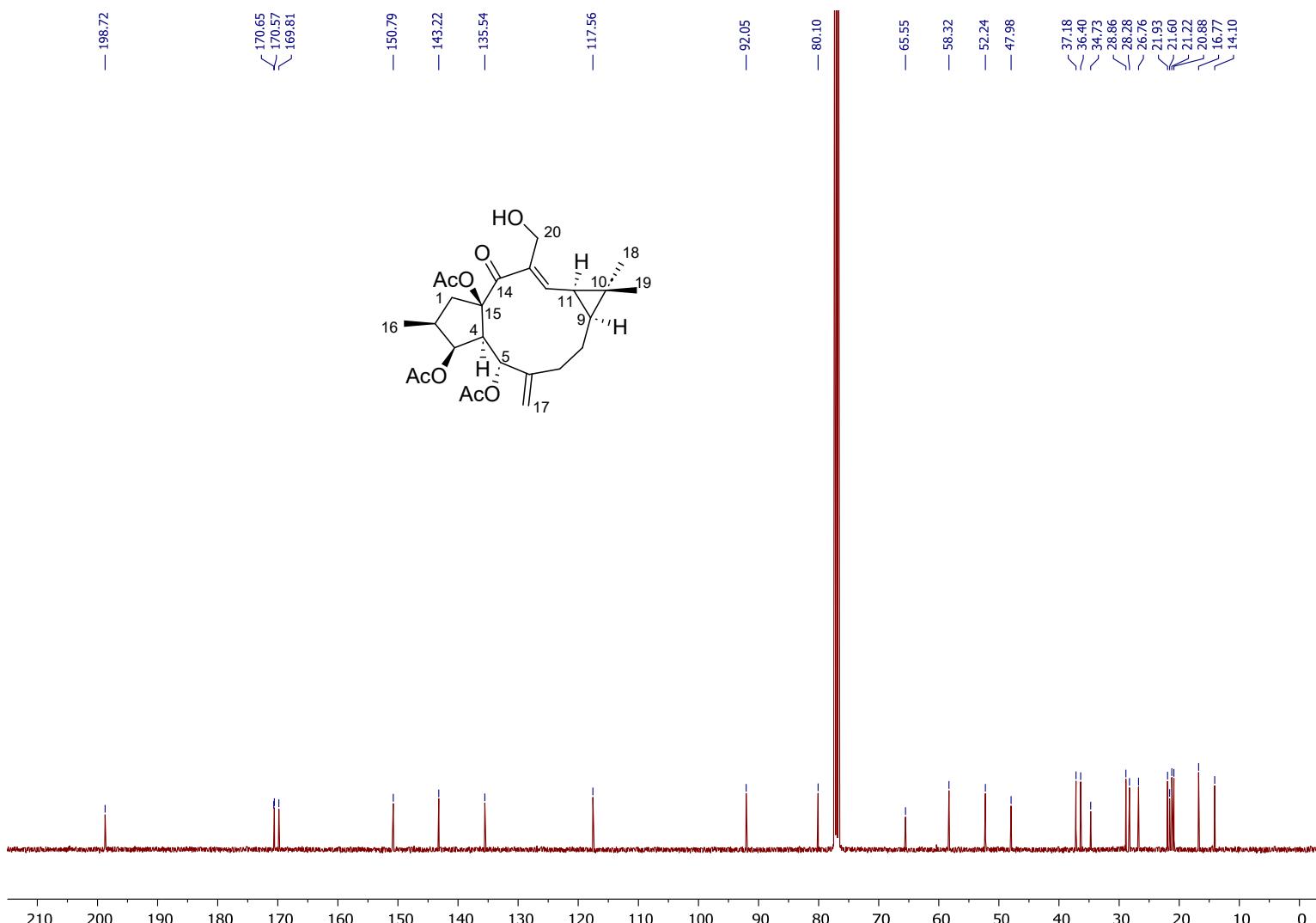


Figure S11. ^{13}C NMR spectrum (100 MHz) of compound 4 in CDCl_3 .

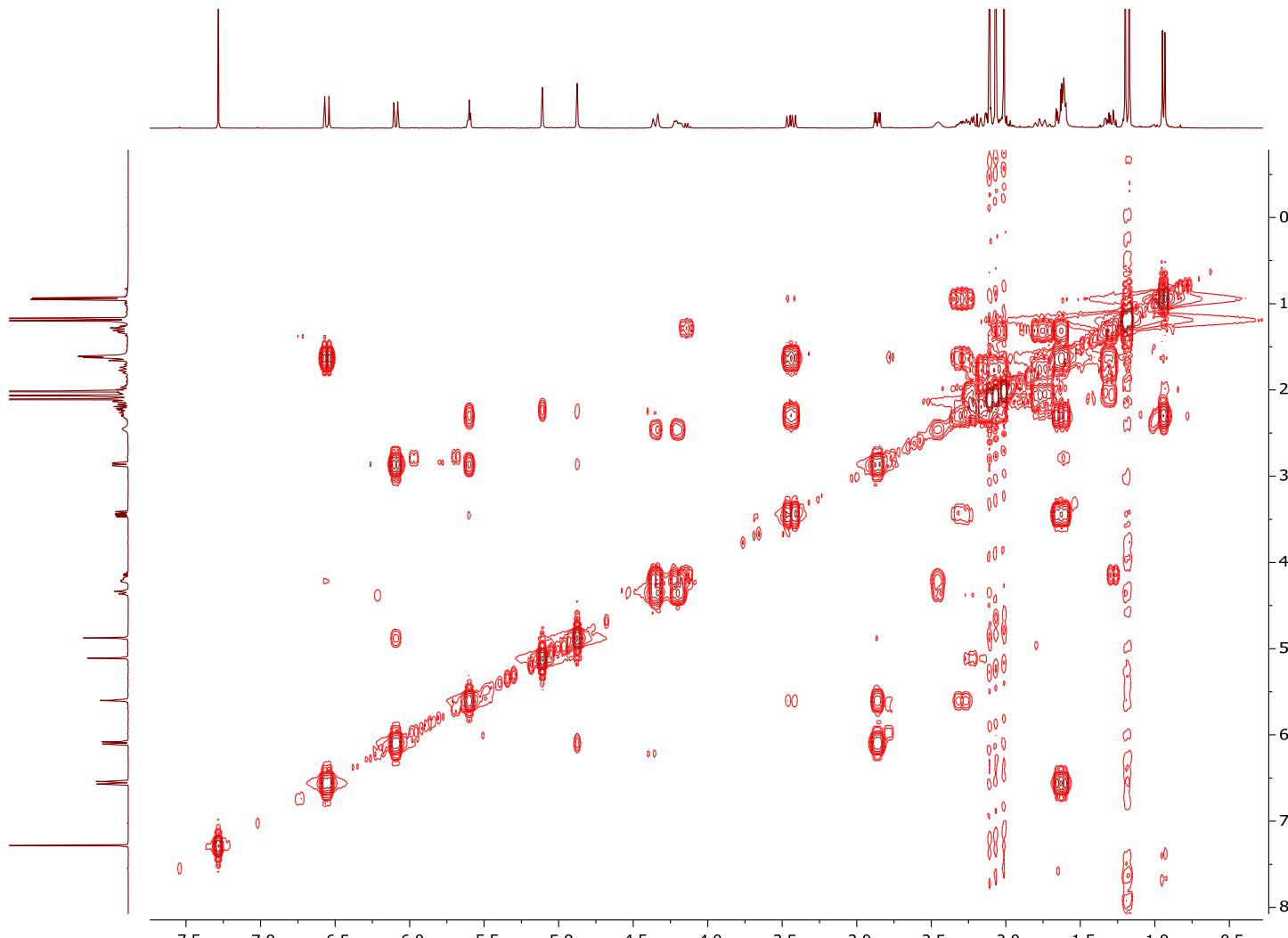


Figure S12. gCOSY spectrum of compound 4.

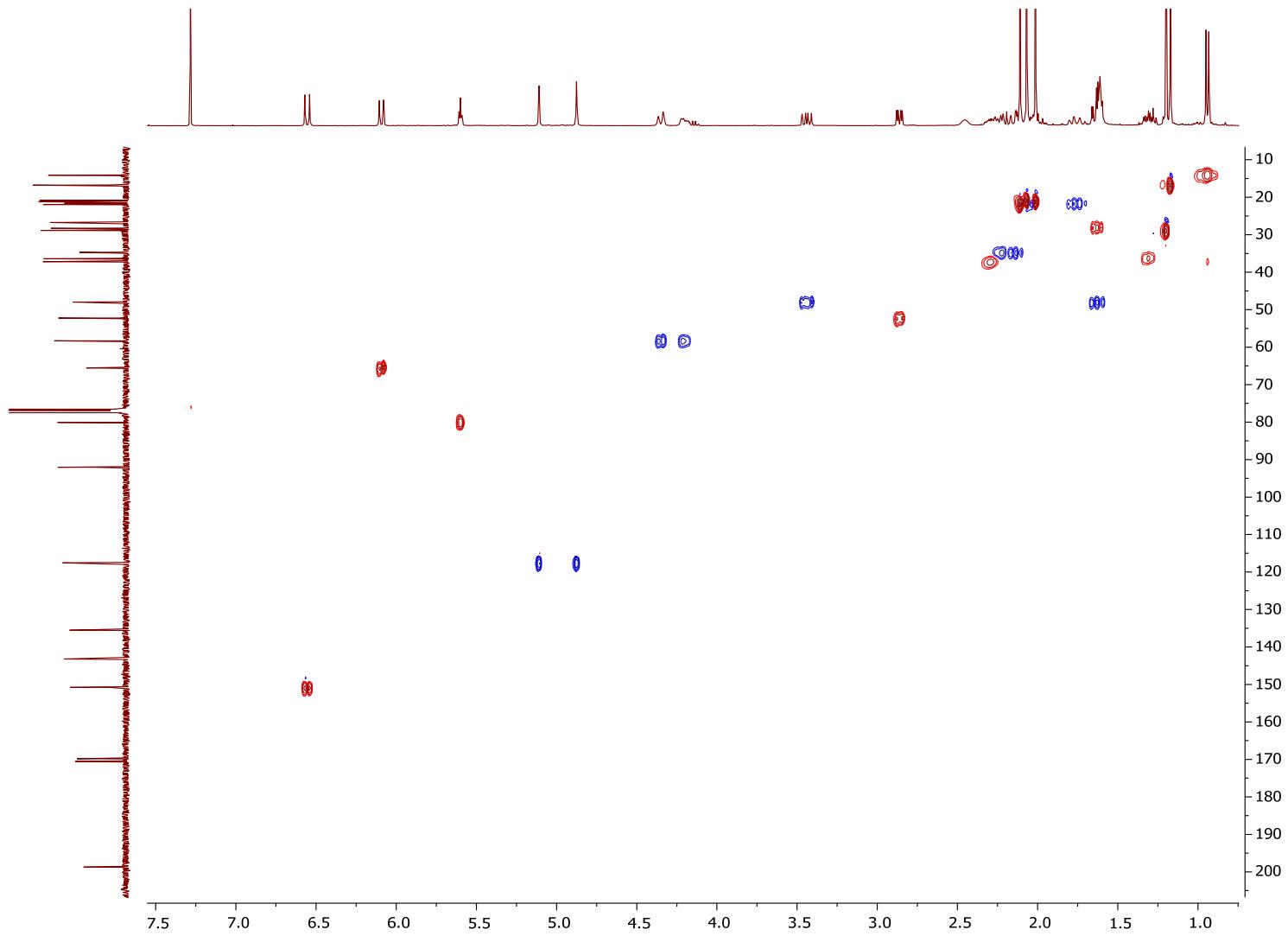


Figure S13. gHSQC spectrum of compound 4.

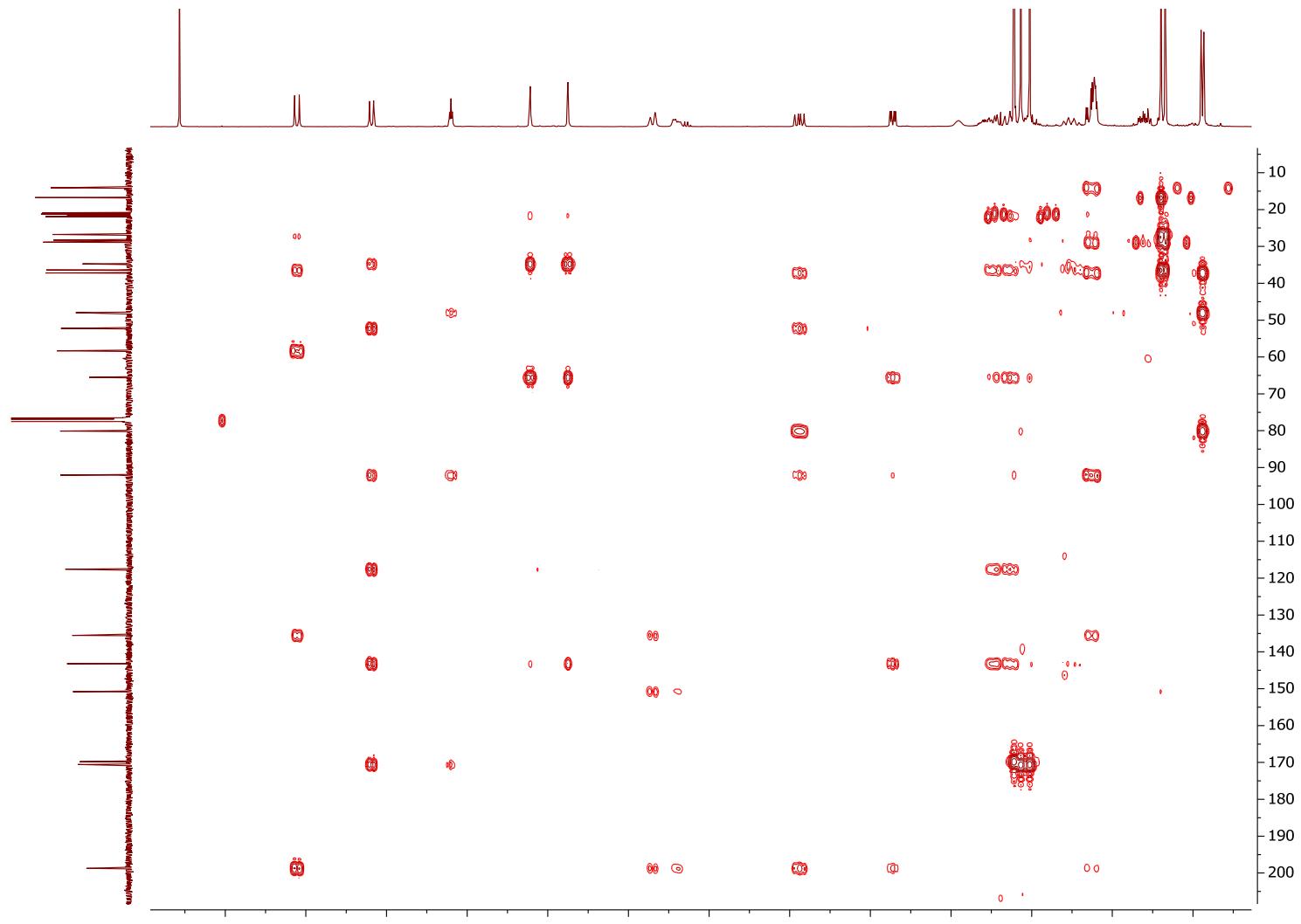


Figure S14. gHMBC spectrum of compound 4.

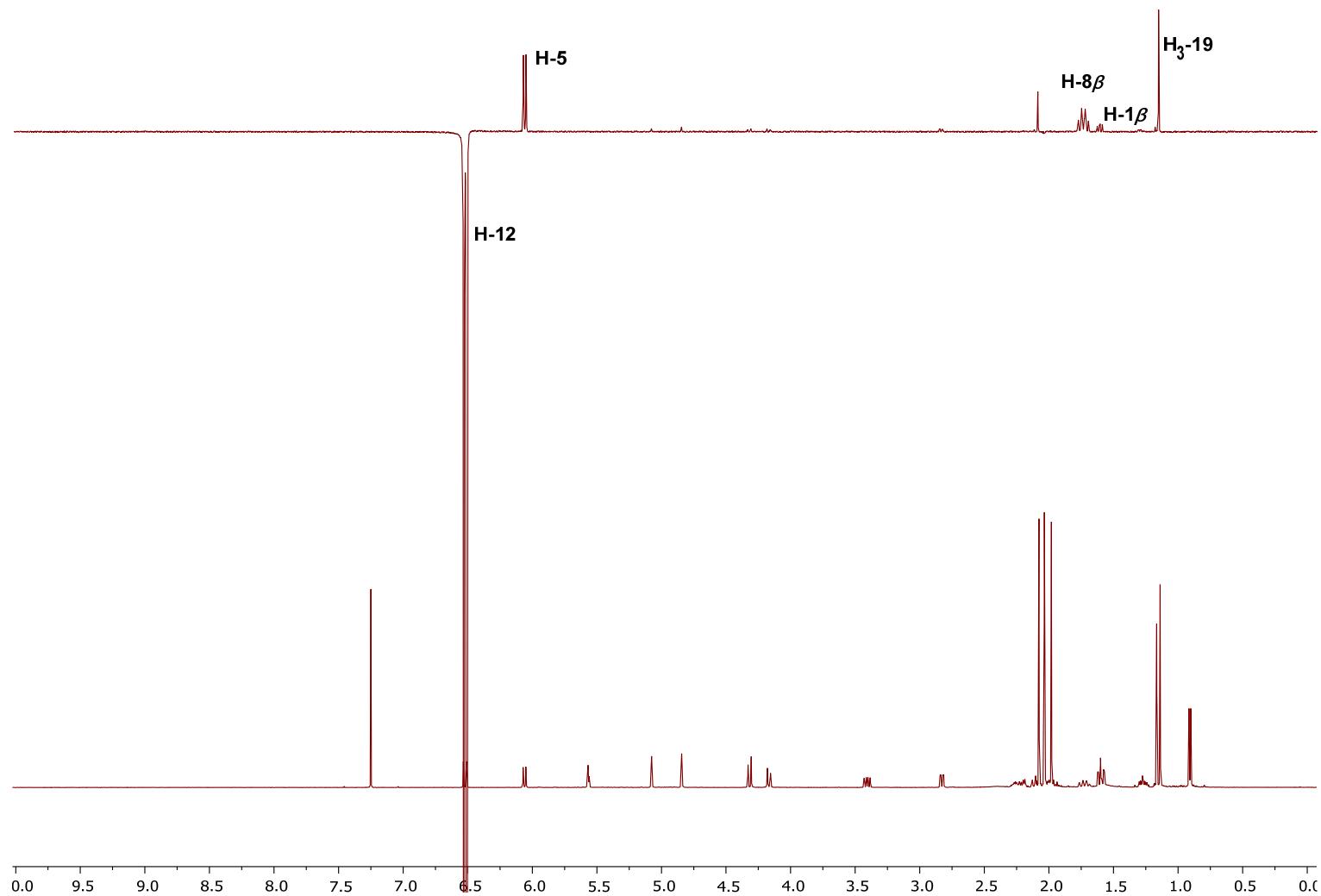


Figure S15a. 1D NOESY spectrum of compound 4.

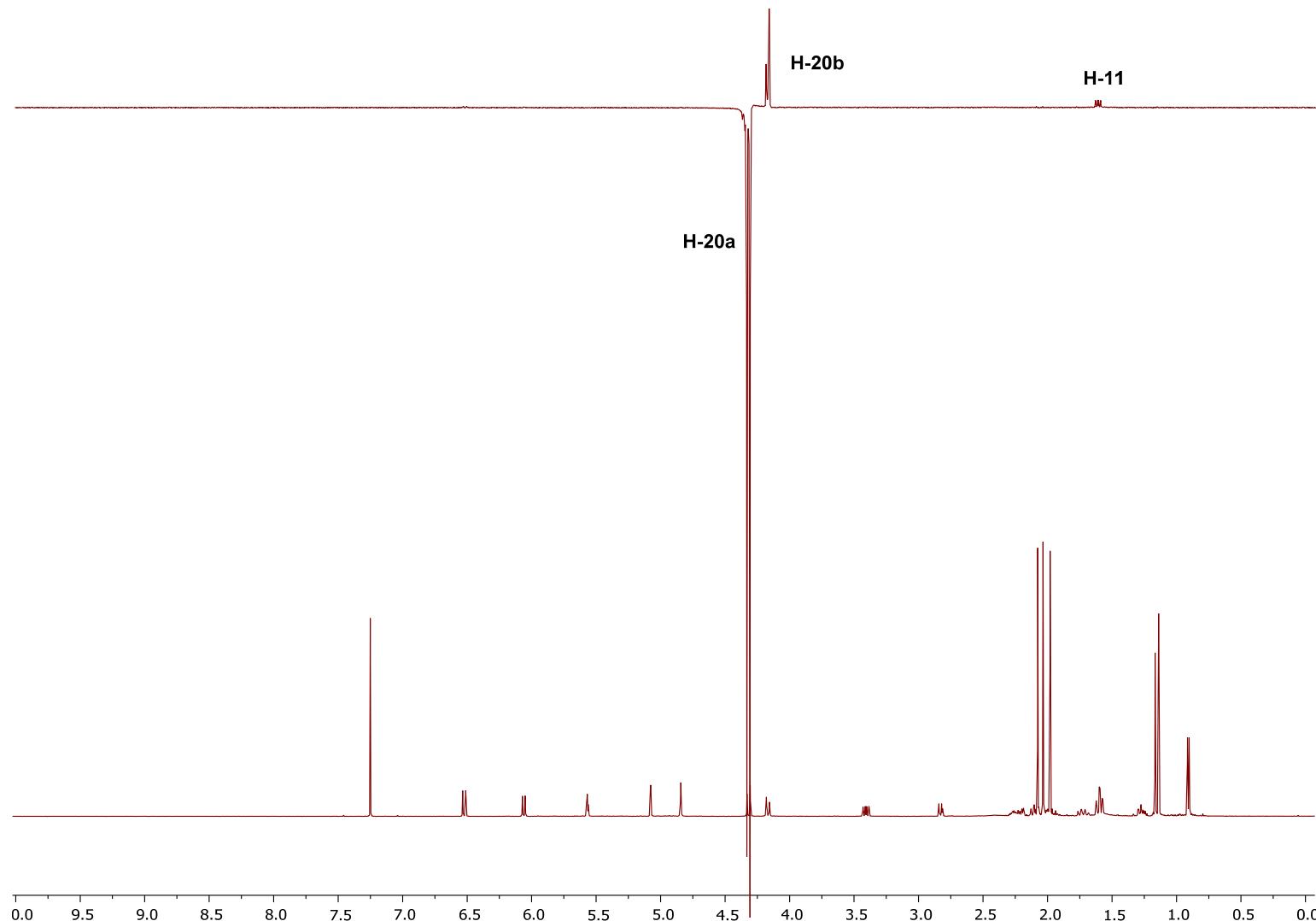


Figure S15b. 1D NOESY spectrum of compound 4.

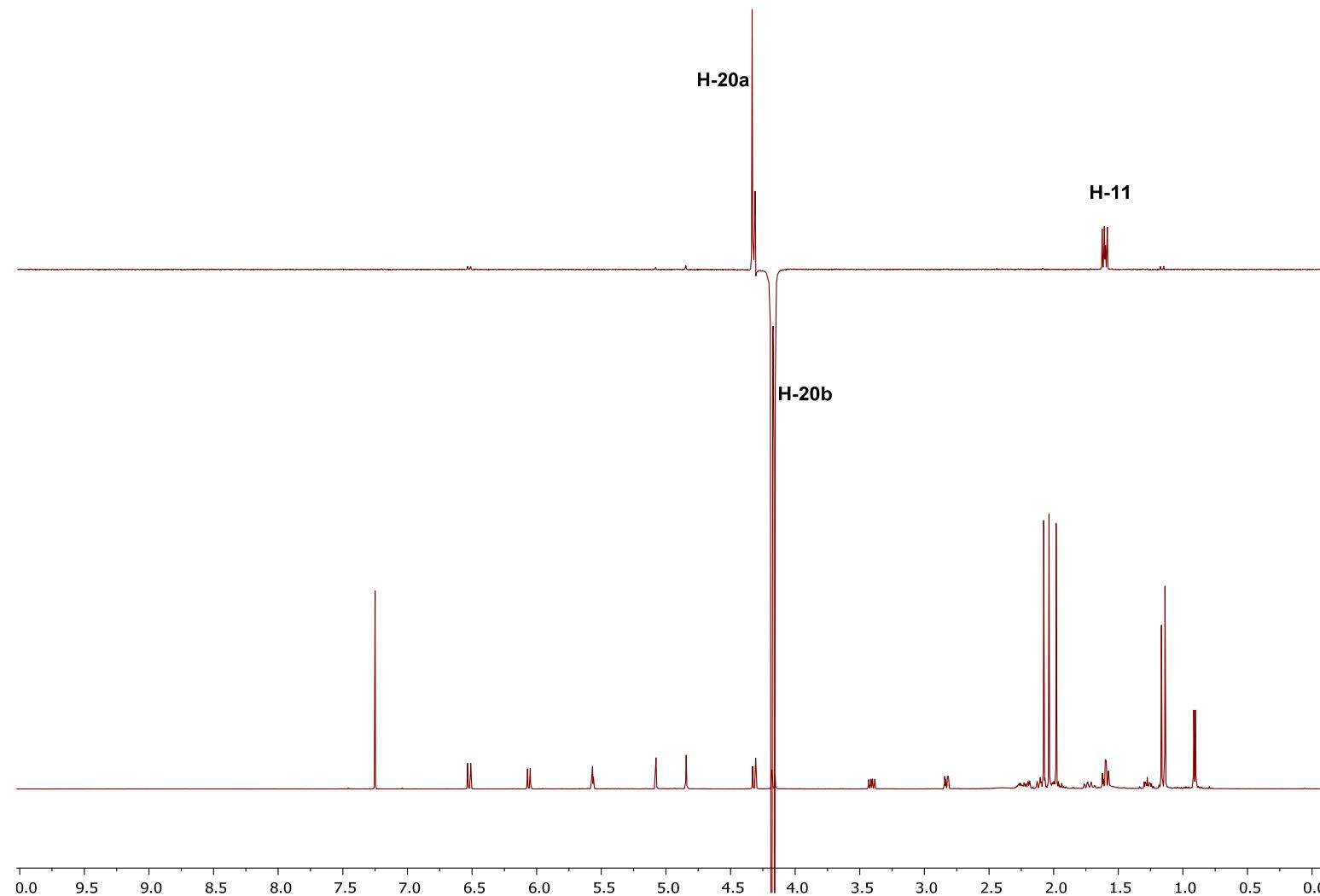


Figure S15c. 1D NOESY spectrum of compound 4.

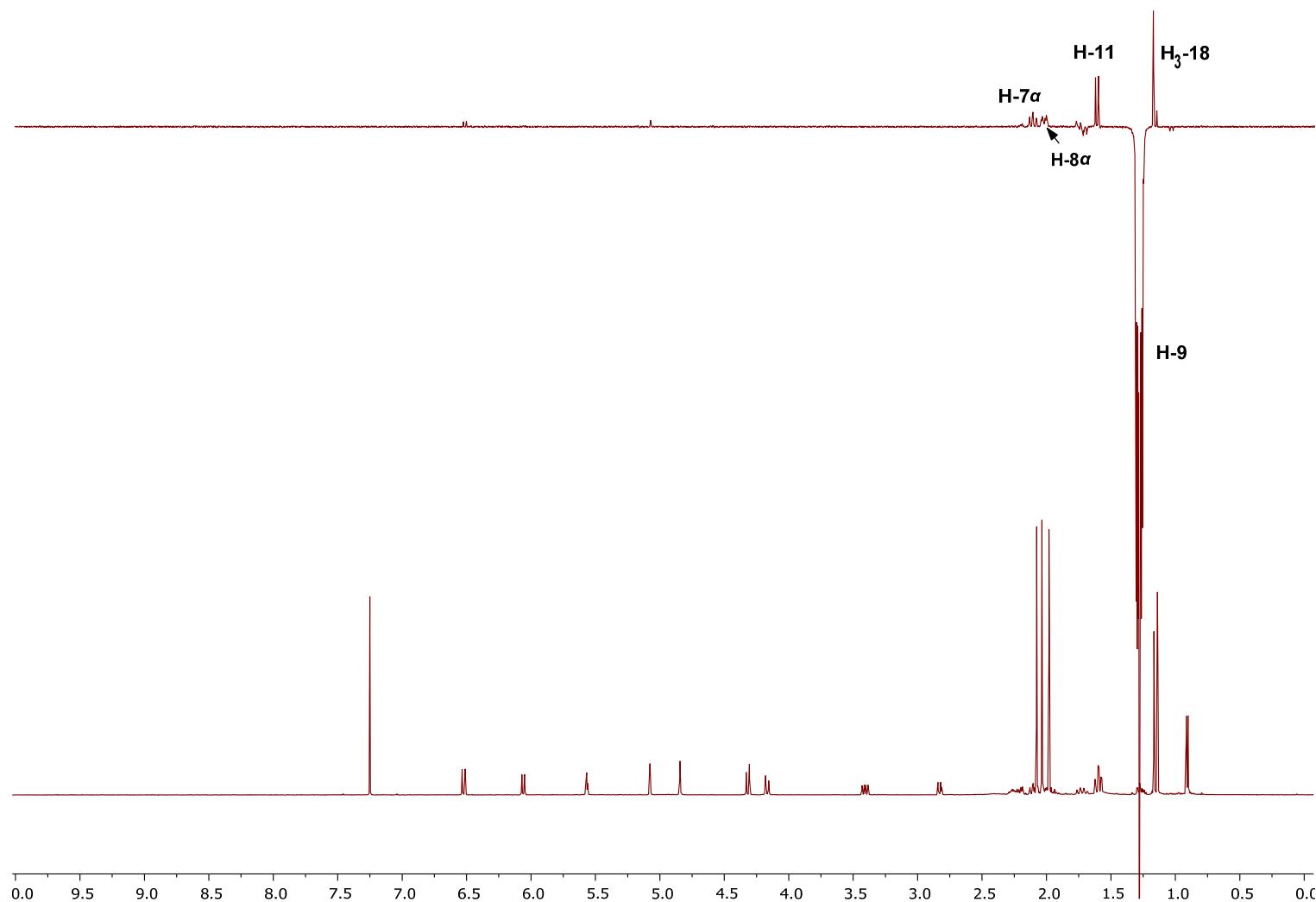


Figure S15d. 1D NOESY spectrum of compound 4.

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

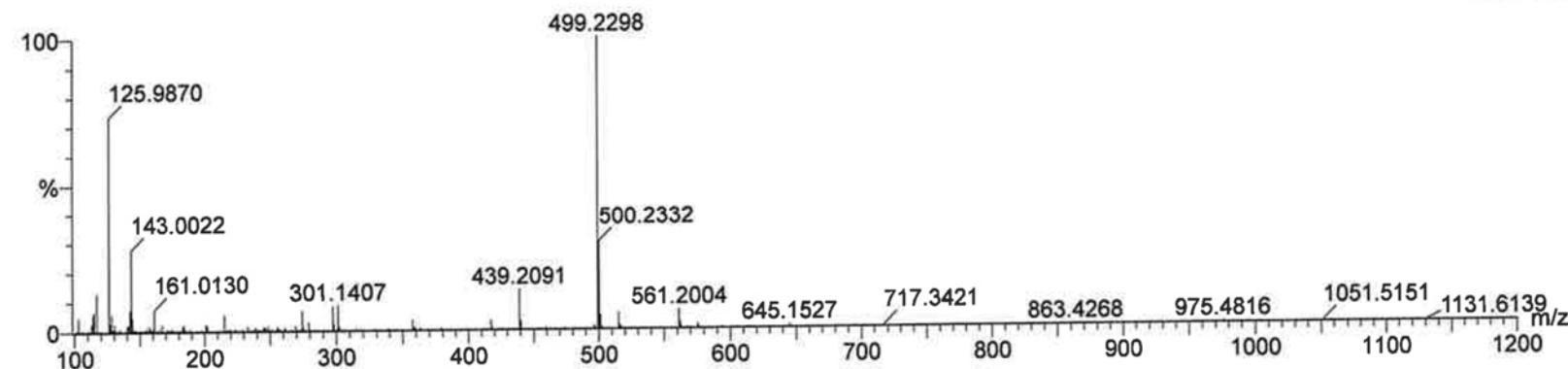
Monoisotopic Mass, Even Electron Ions

87 formula(e) evaluated with 2 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 0-30 H: 0-50 O: 0-15 23Na: 0-1

246_1009_Strep-EB12- MSe3pos 148 (2.741)

1: TOF MS ES+
3.42e+005

Minimum:				-1.5
Maximum:	5.0	10.0	80.0	

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
499.2298	499.2308	-1.0	-2.0	8.5	1285.1	0.063	93.85	C26 H36 O8 23Na
	499.2332	-3.4	-6.8	11.5	1287.8	2.789	6.15	C28 H35 O8

Figure S16. HRMS of compound 4.

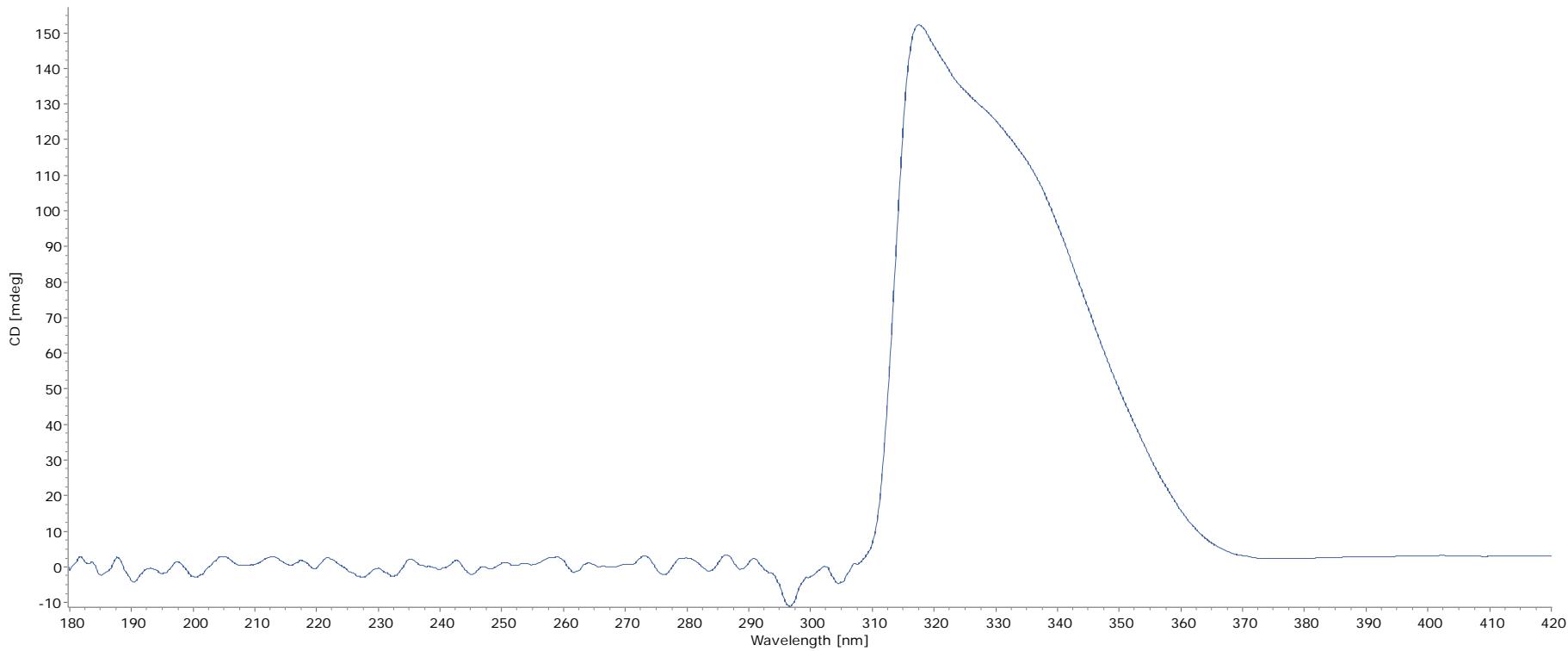


Figure S17. ECD of compound 4.

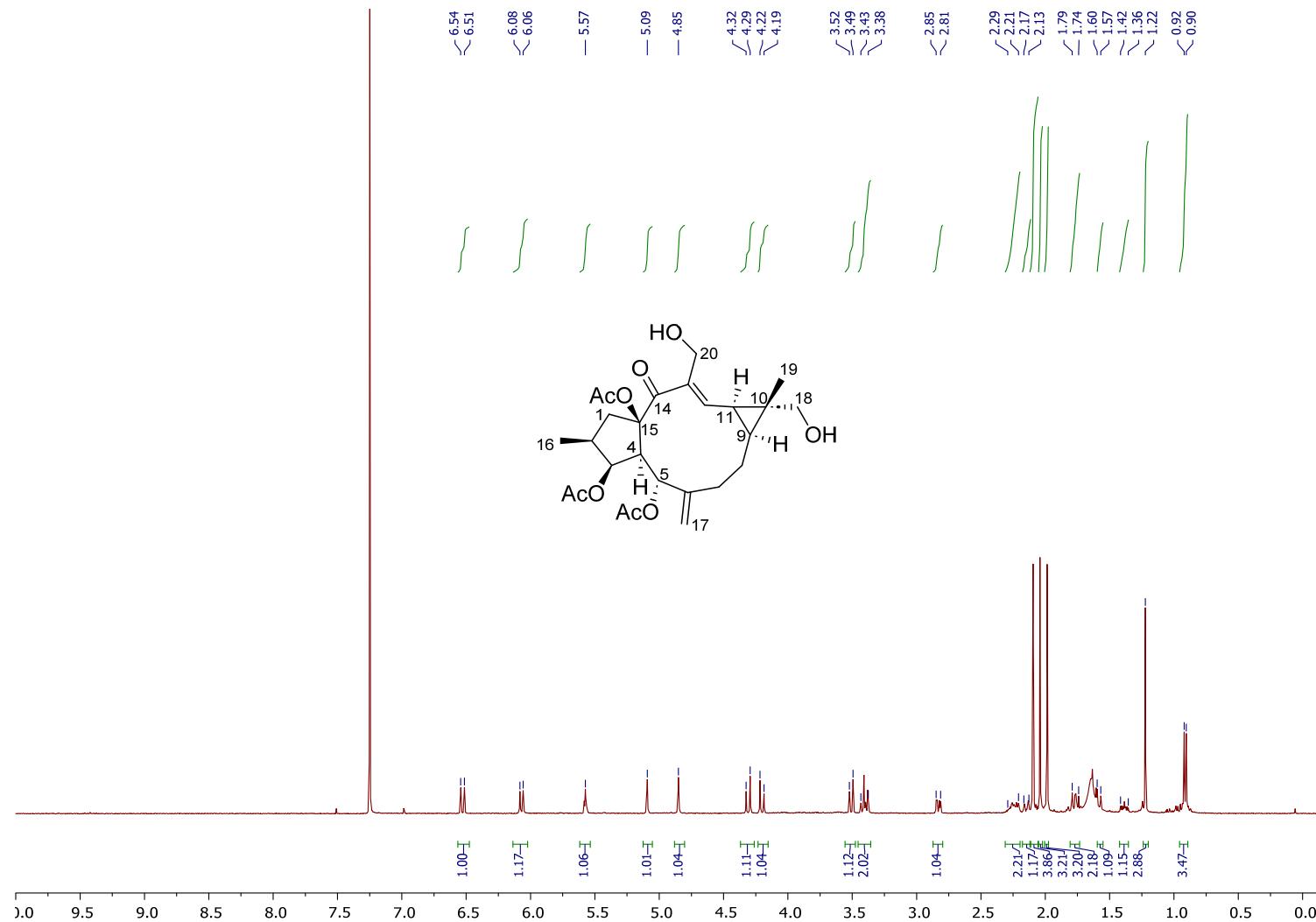


Figure S18. ¹H NMR spectrum (400 MHz) of compound 5 in CDCl₃.

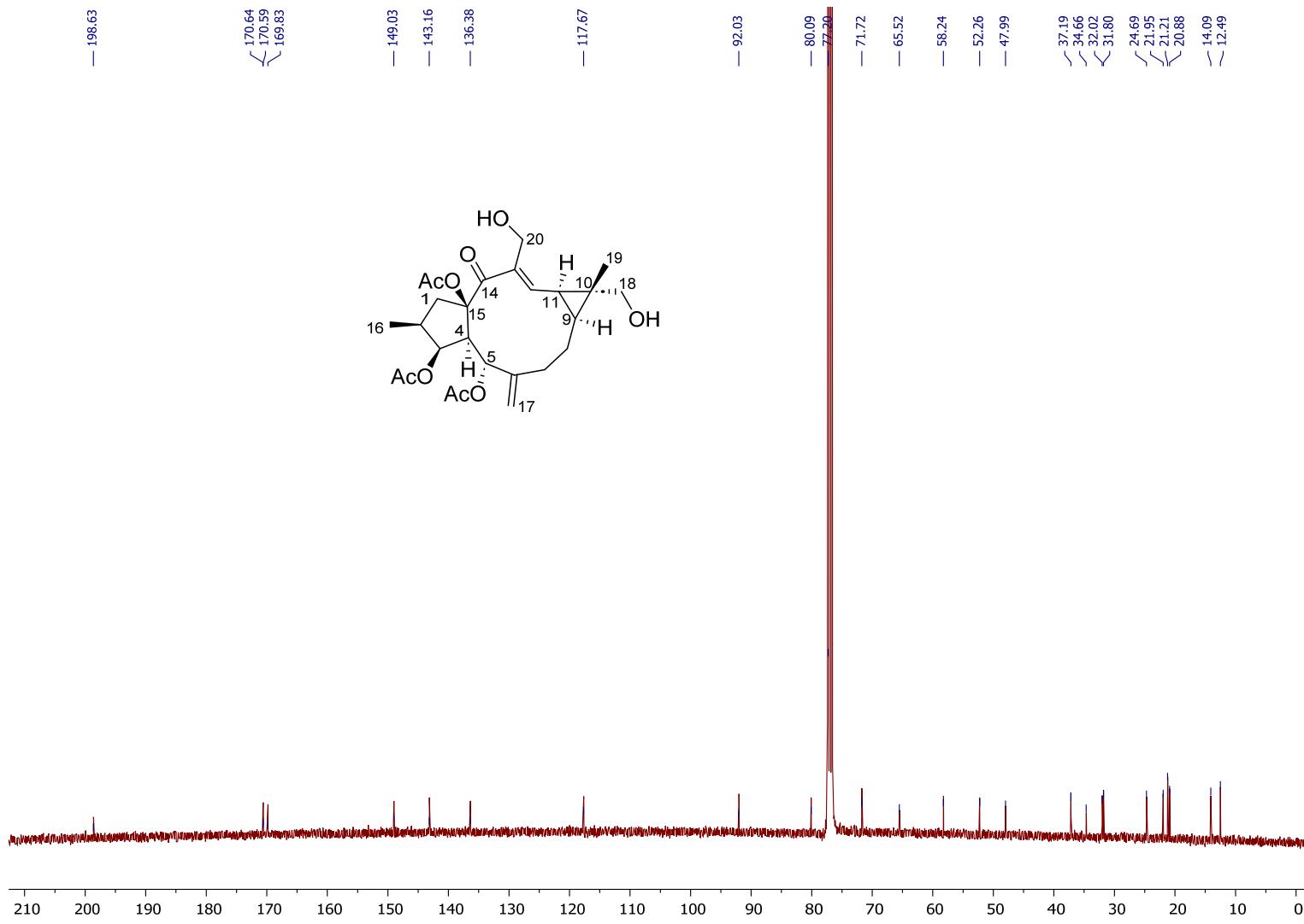


Figure S19. ^{13}C NMR spectrum (100 MHz) of compound 5 in CDCl_3 .

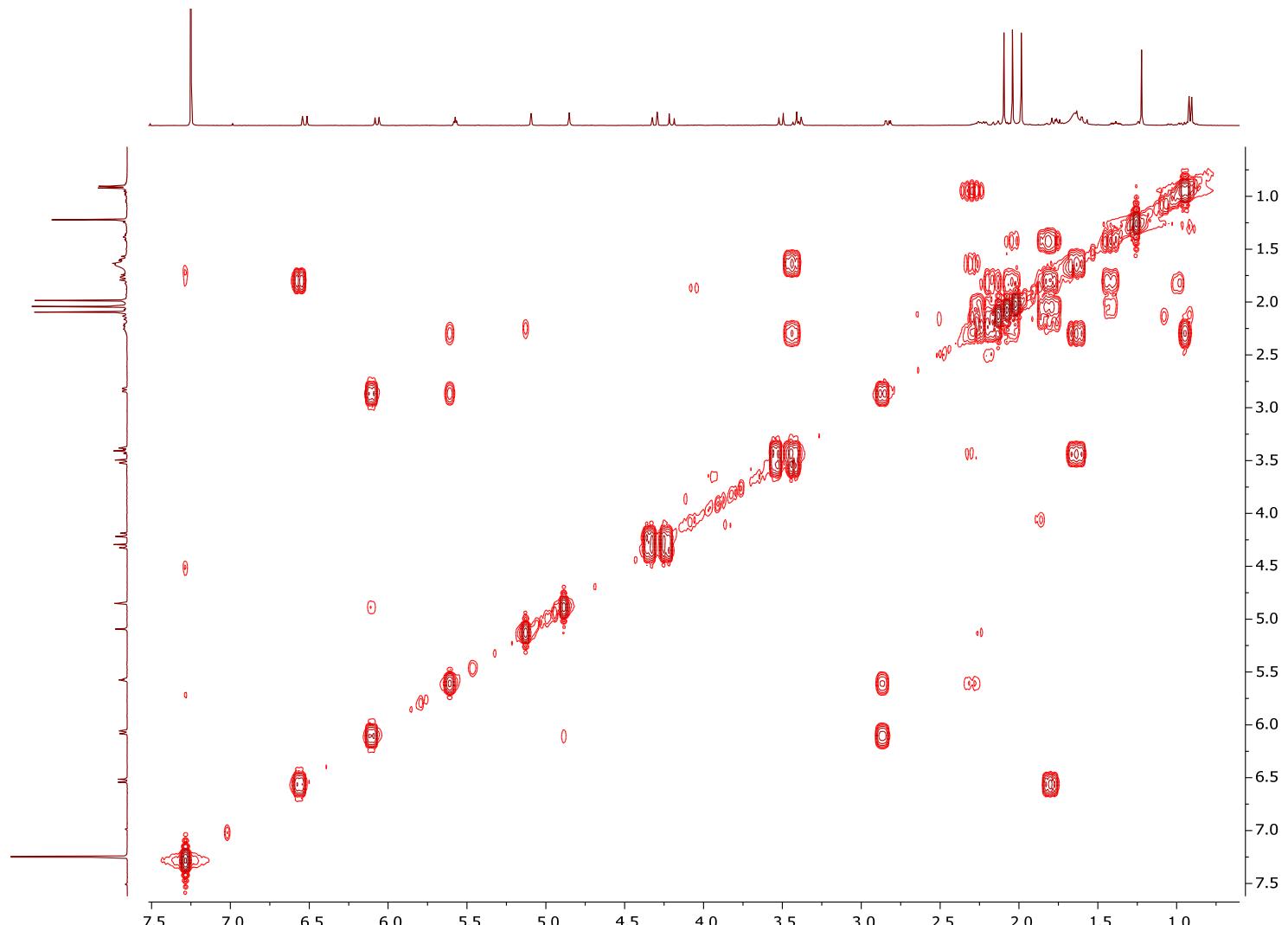


Figure S20. gCOSY spectrum of compound 5.

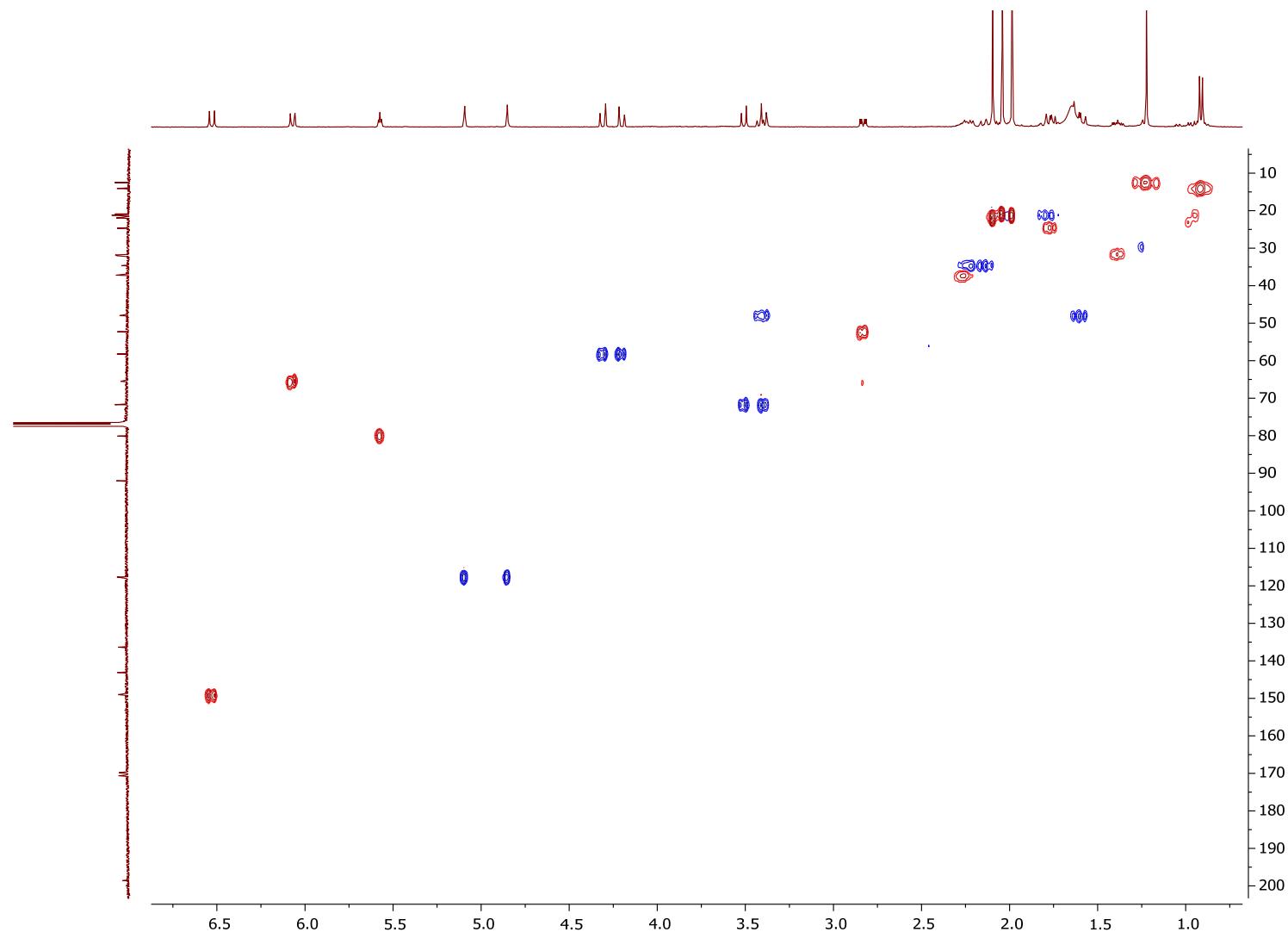


Figure S21. gHSQC spectrum of compound 5.

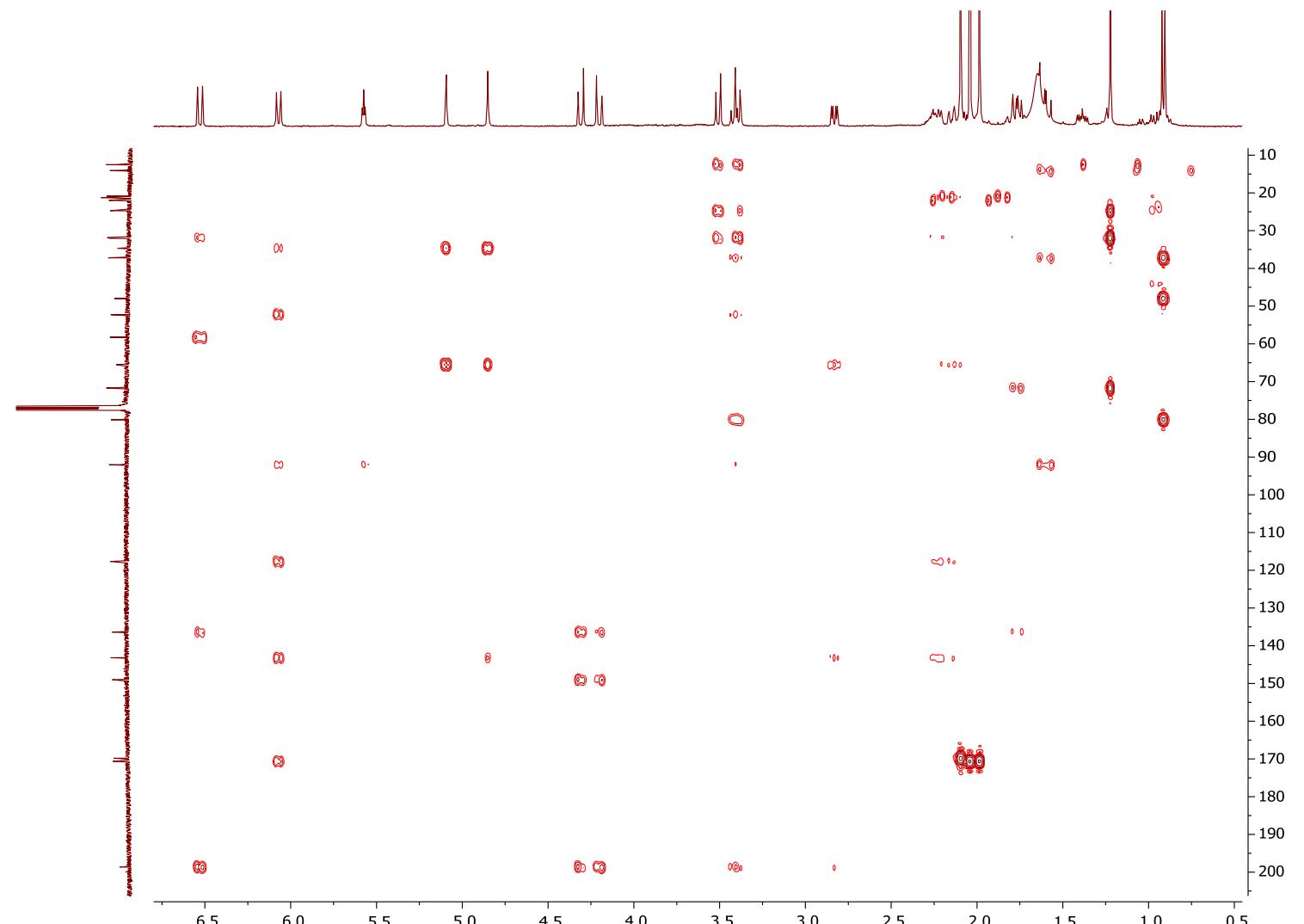


Figure S22. gHMBC spectrum of compound 5.

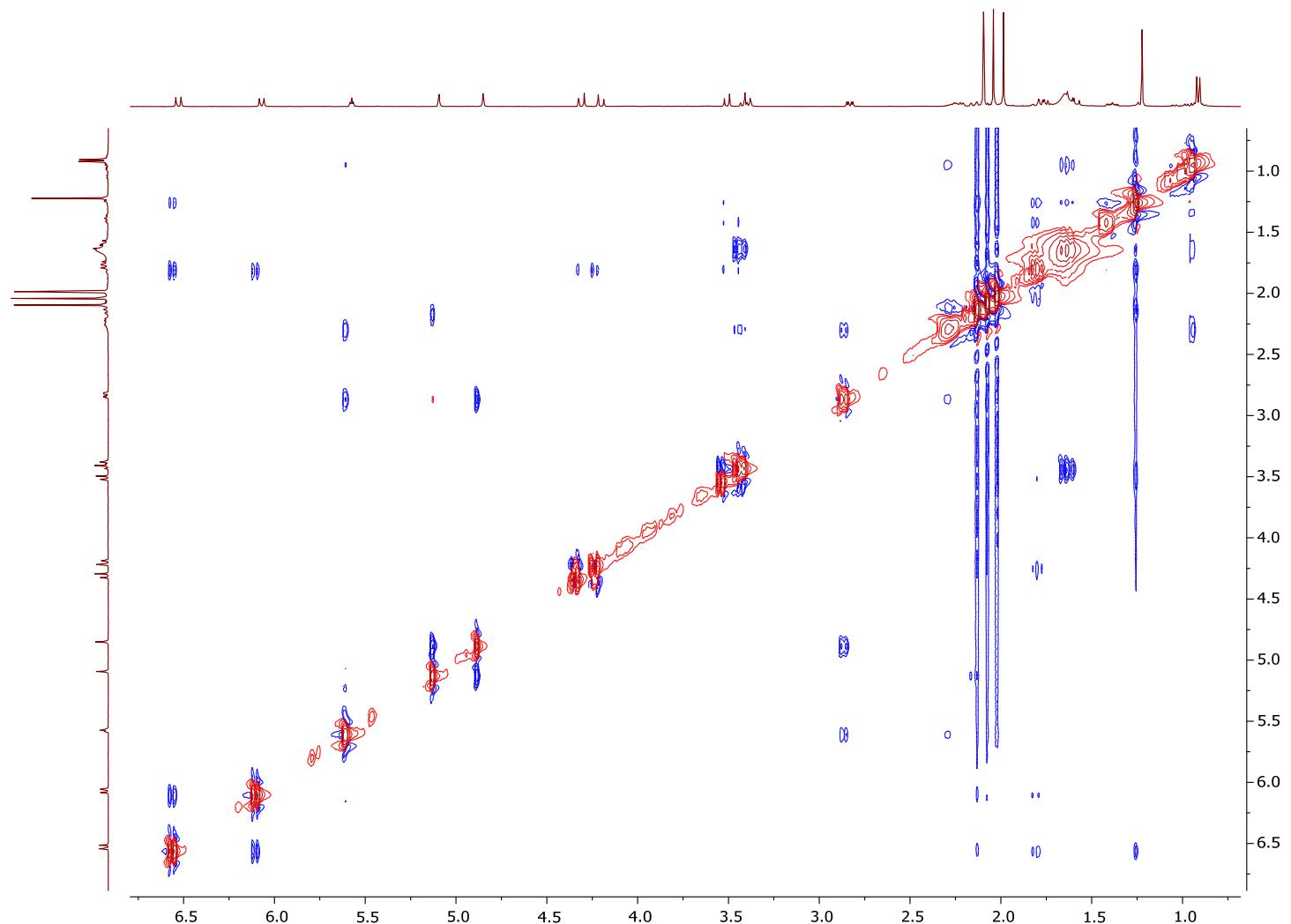


Figure S23. 2D NOESY spectrum of compound 5.

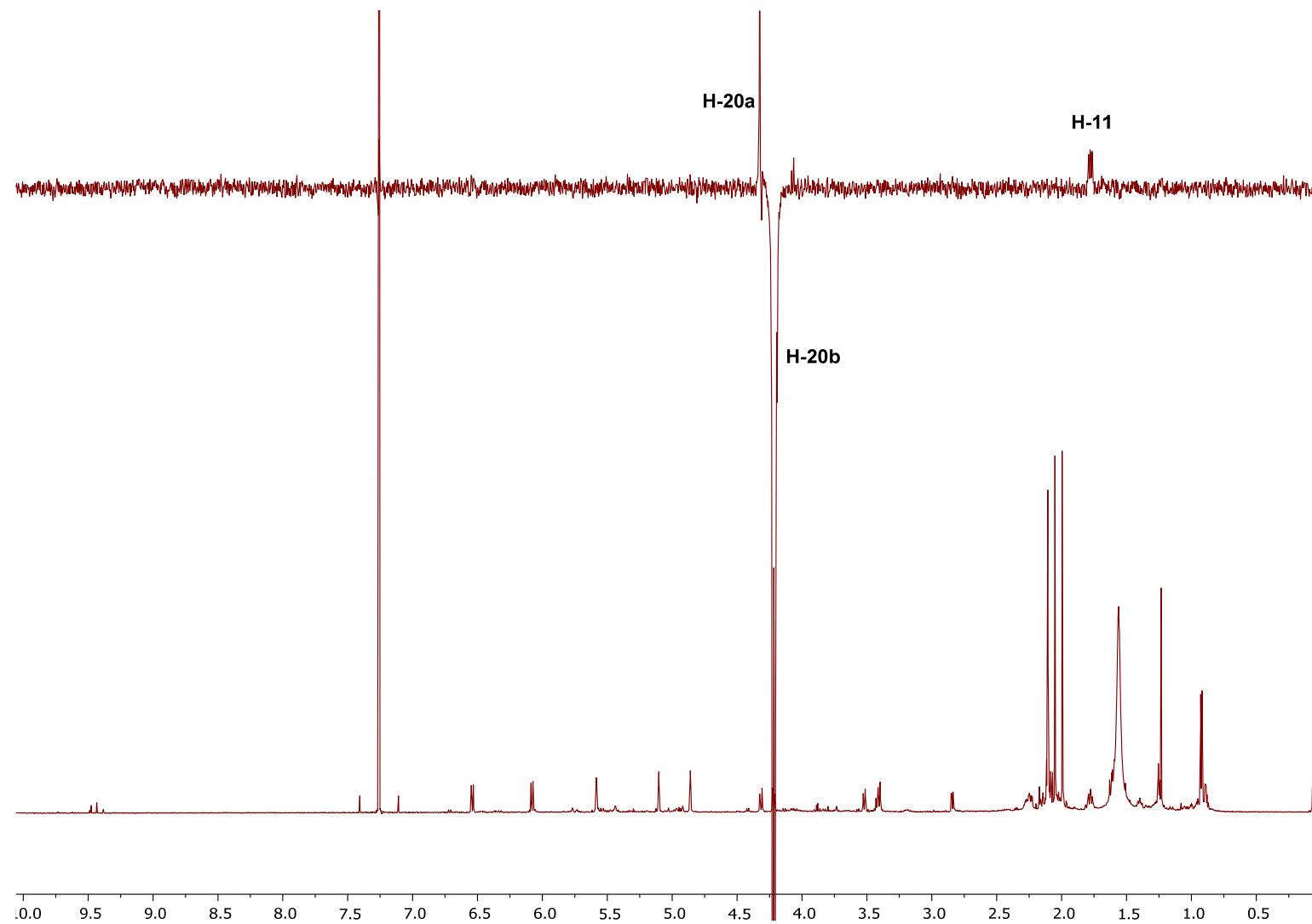


Figure S24a. 1D NOESY spectrum of compound 5.

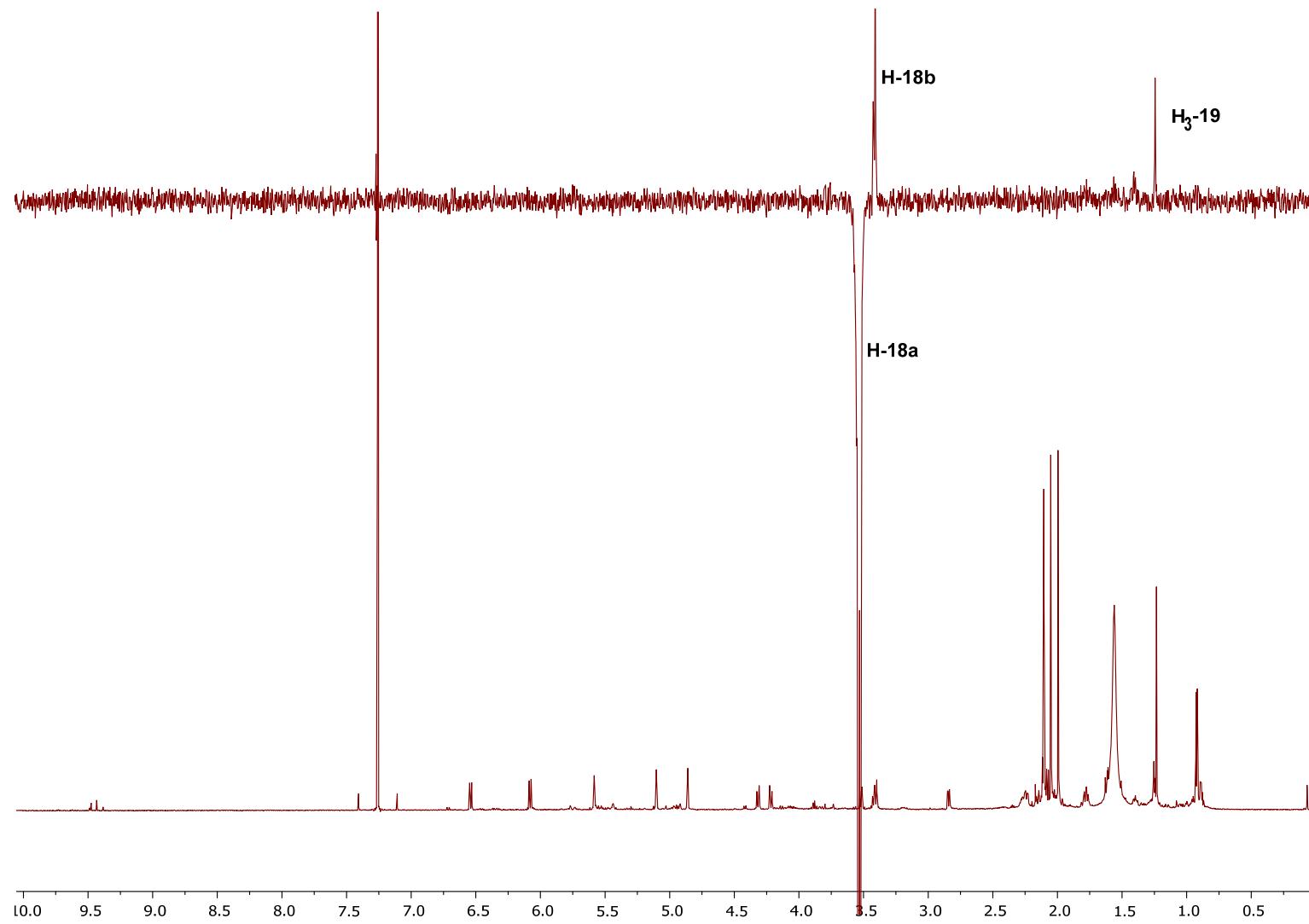


Figure S24b. 1D NOESY spectrum of compound 5.

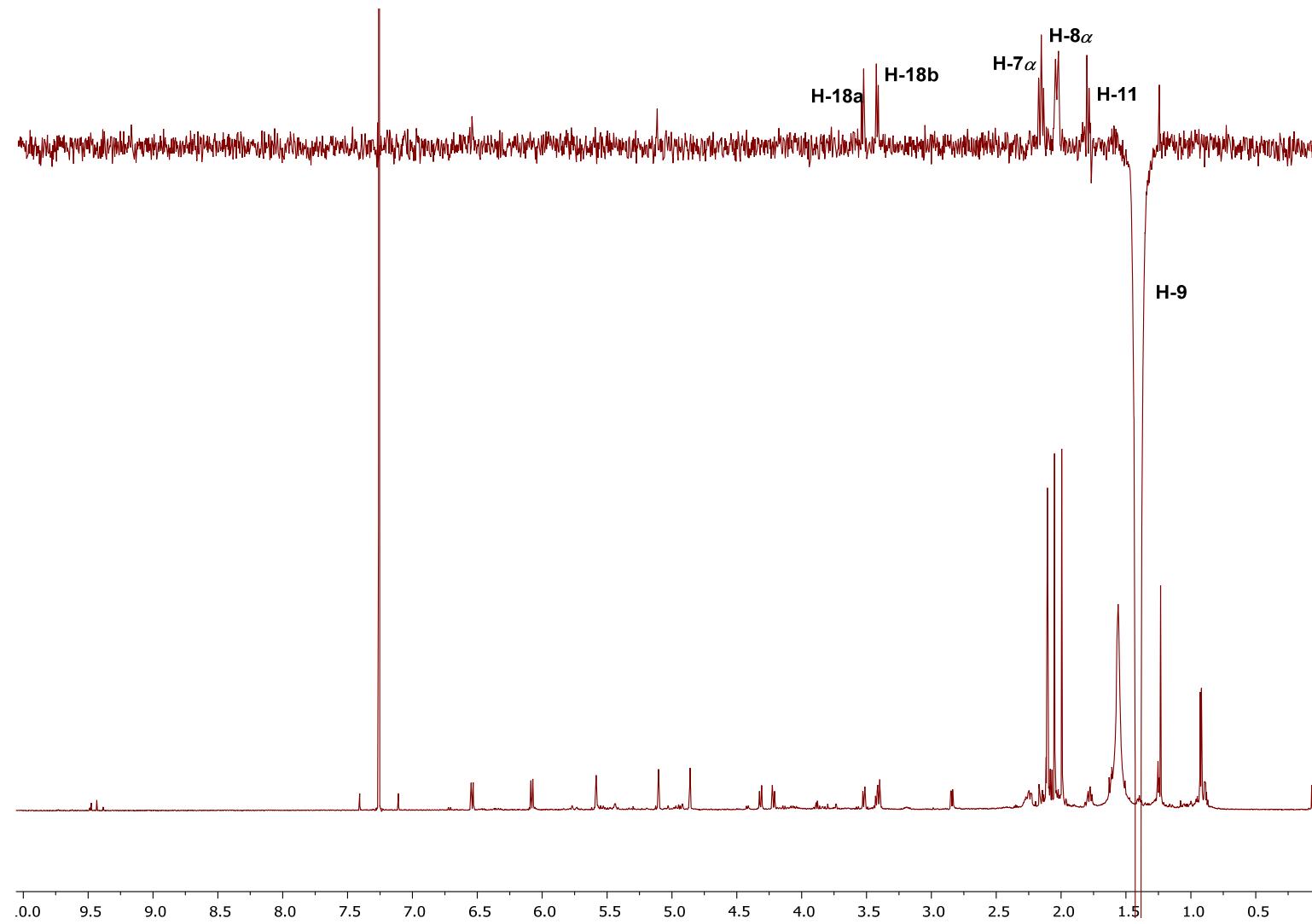


Figure S24c. 1D NOESY spectrum of compound 5.

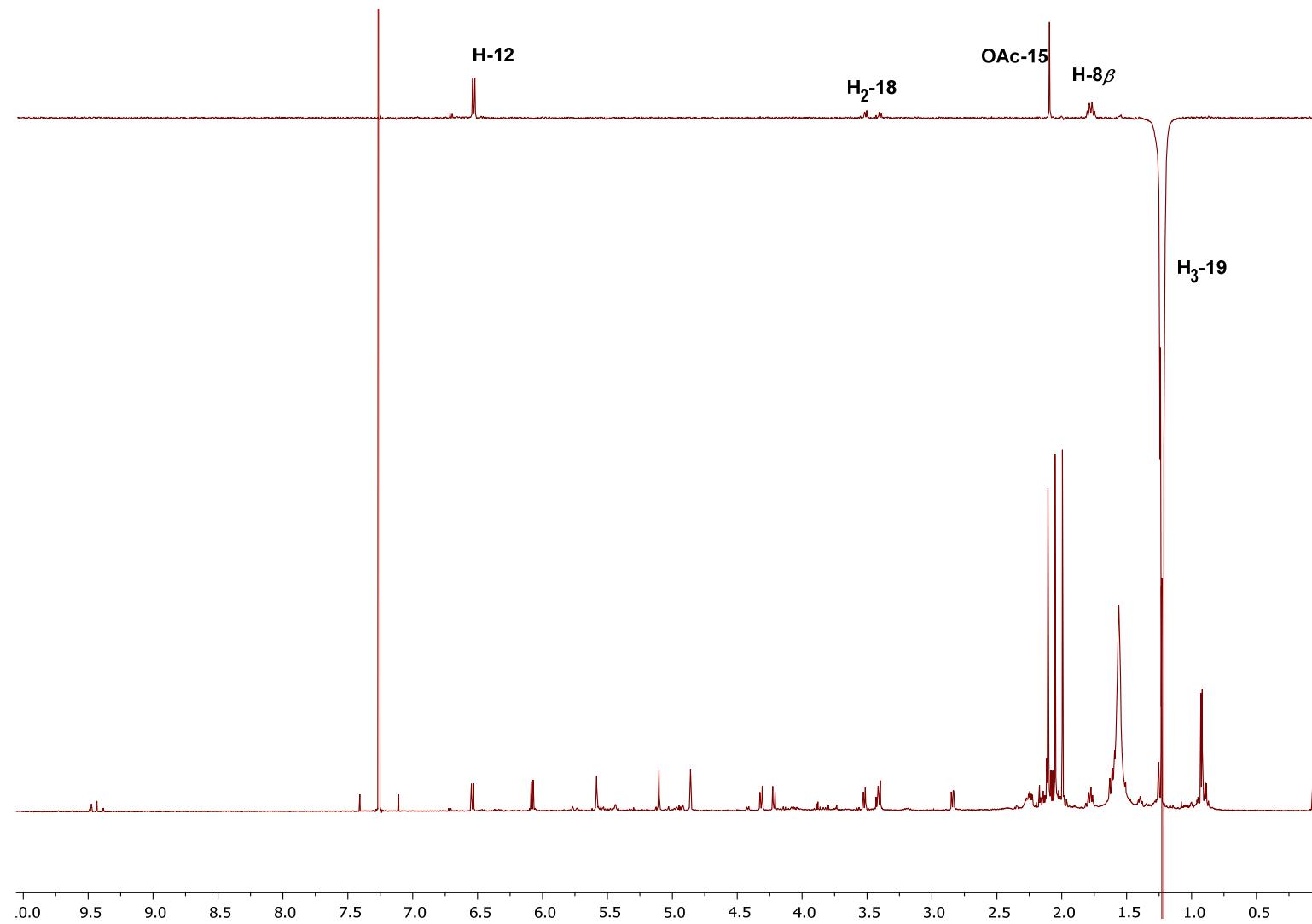


Figure S24d. 1D NOESY spectrum of compound 5.

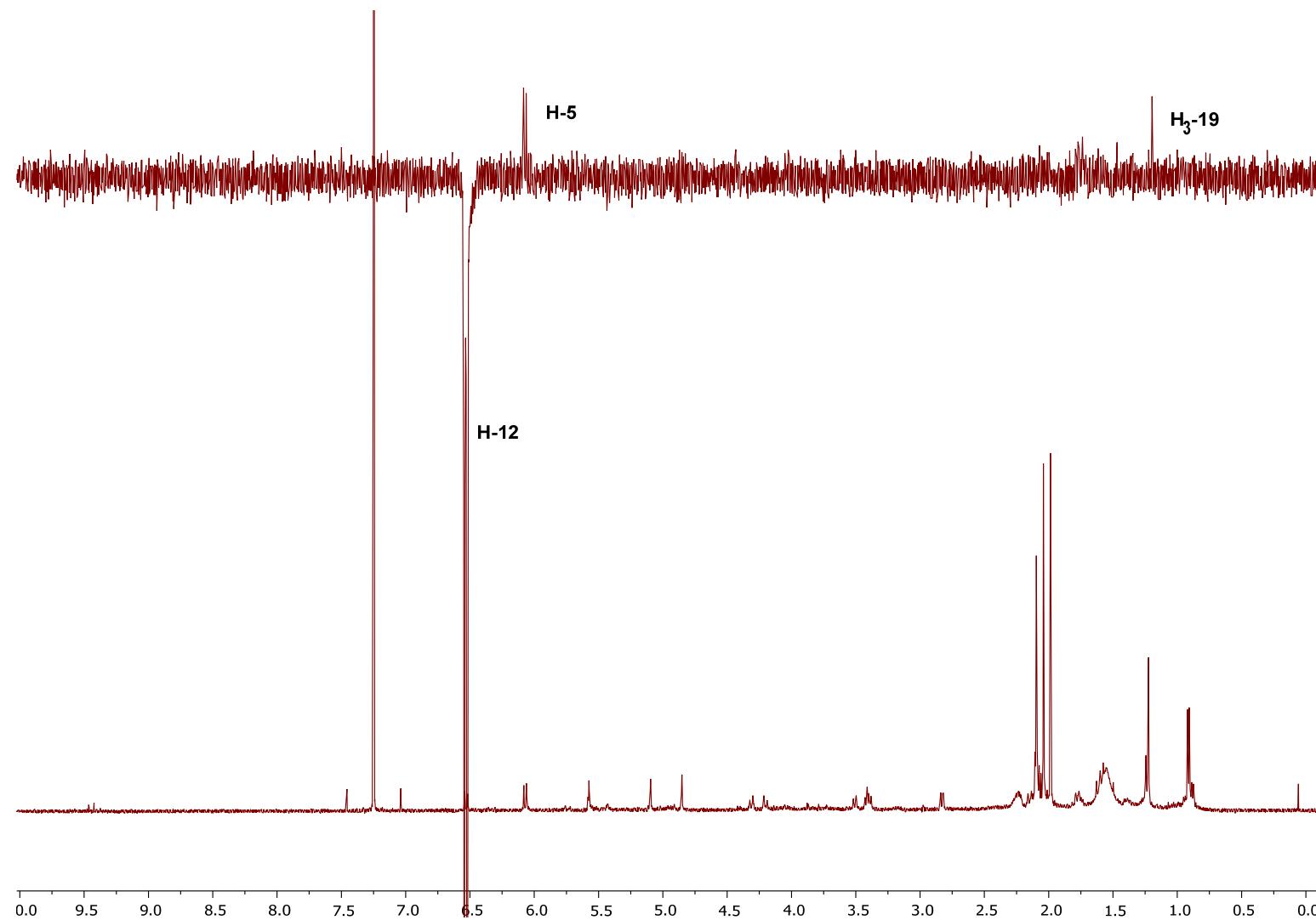


Figure S24e. 1D NOESY spectrum of compound 5.

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

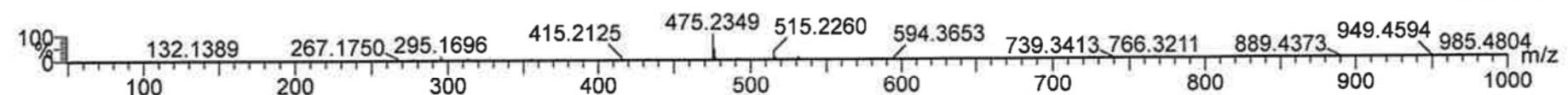
215 formula(e) evaluated with 3 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 0-50 H: 0-200 O: 0-20 Na: 0-1

C26H36O9 FELIPE

267 191 (0.749)

1: TOF MS ES+
1.64e+007

Minimum: -1.5
 Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
515.2260	515.2257	0.3	0.6	8.5	717.7	0.042	95.89	C26 H36 O9 Na
	515.2281	-2.1	-4.1	11.5	720.9	3.197	4.09	C28 H35 O9
	515.2222	3.8	7.4	20.5	725.9	8.264	0.03	C35 H31 O4

Figure S25. HRMS of compound 5.

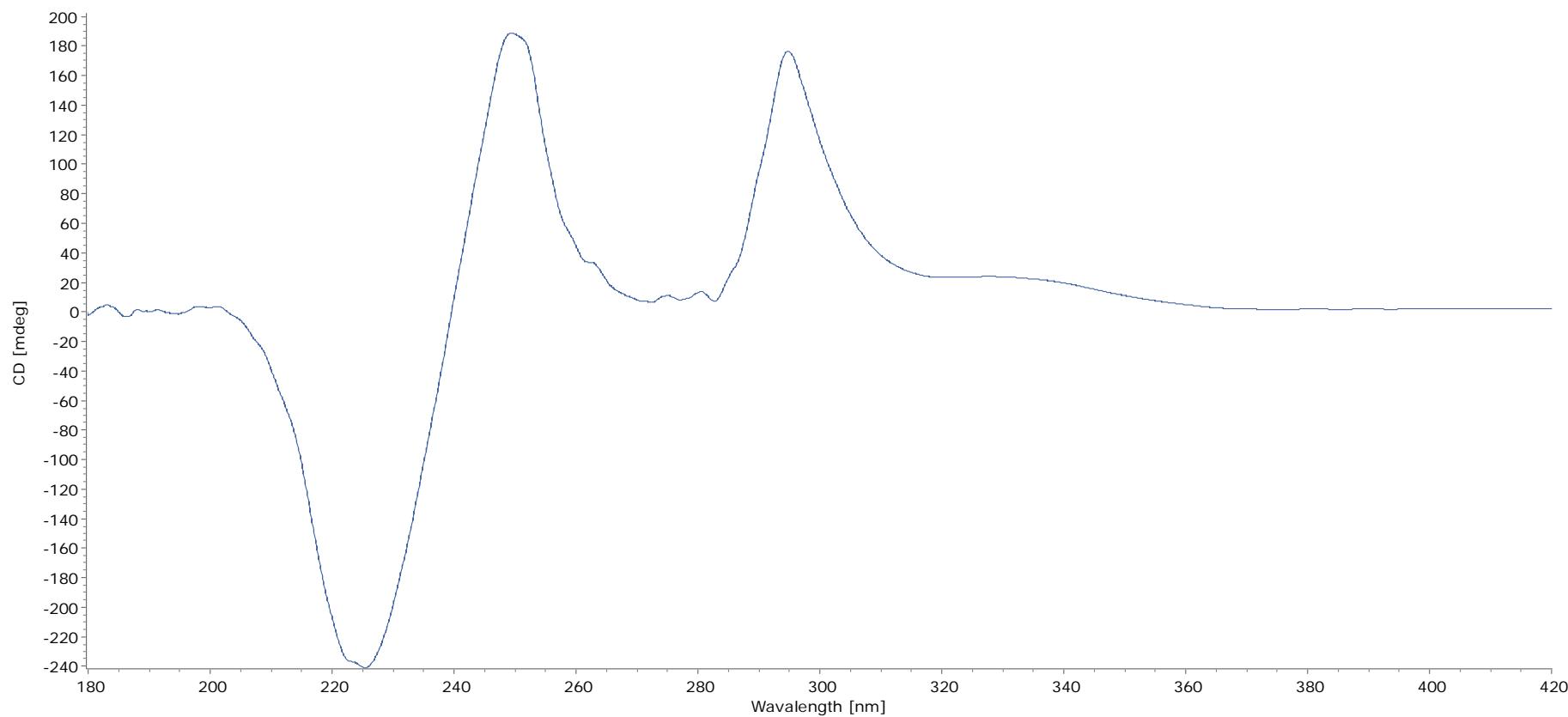


Figure S26. ECD of compound **5**.

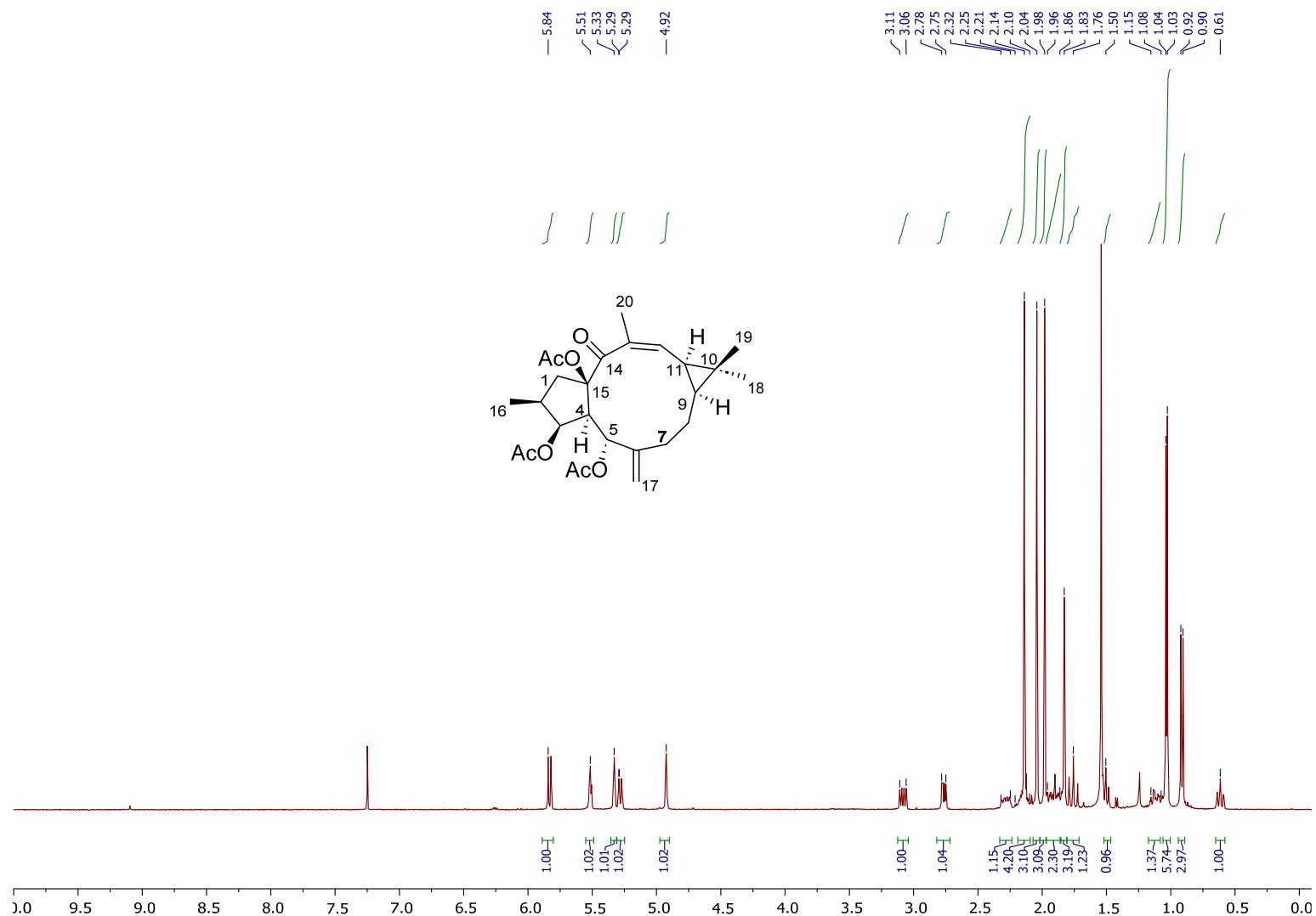


Figure S27. ^1H NMR spectrum (400 MHz) of compound **6** in CDCl_3 .

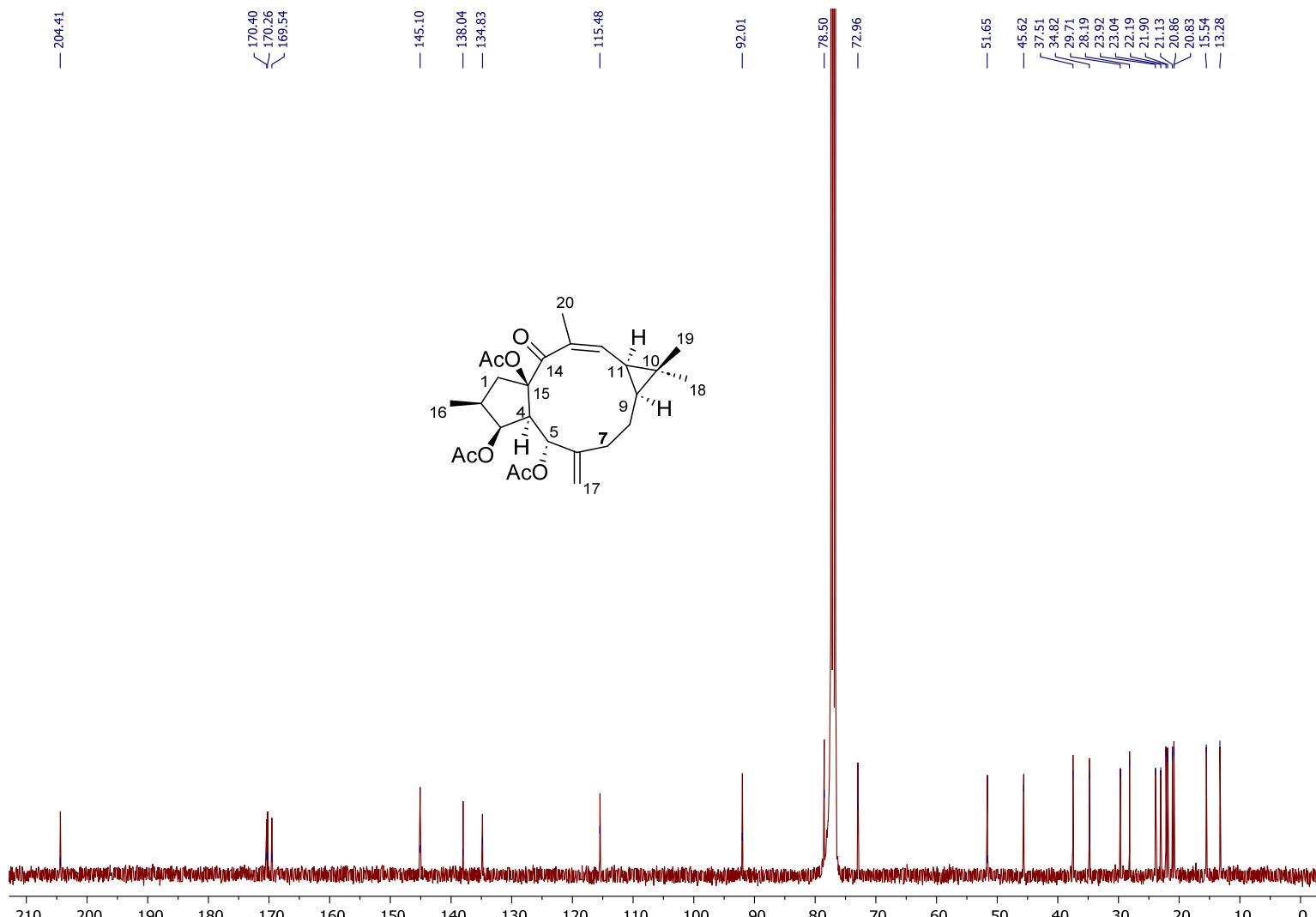


Figure S28. ^{13}C NMR spectrum (100 Mz) of compound 6 in CDCl_3 .

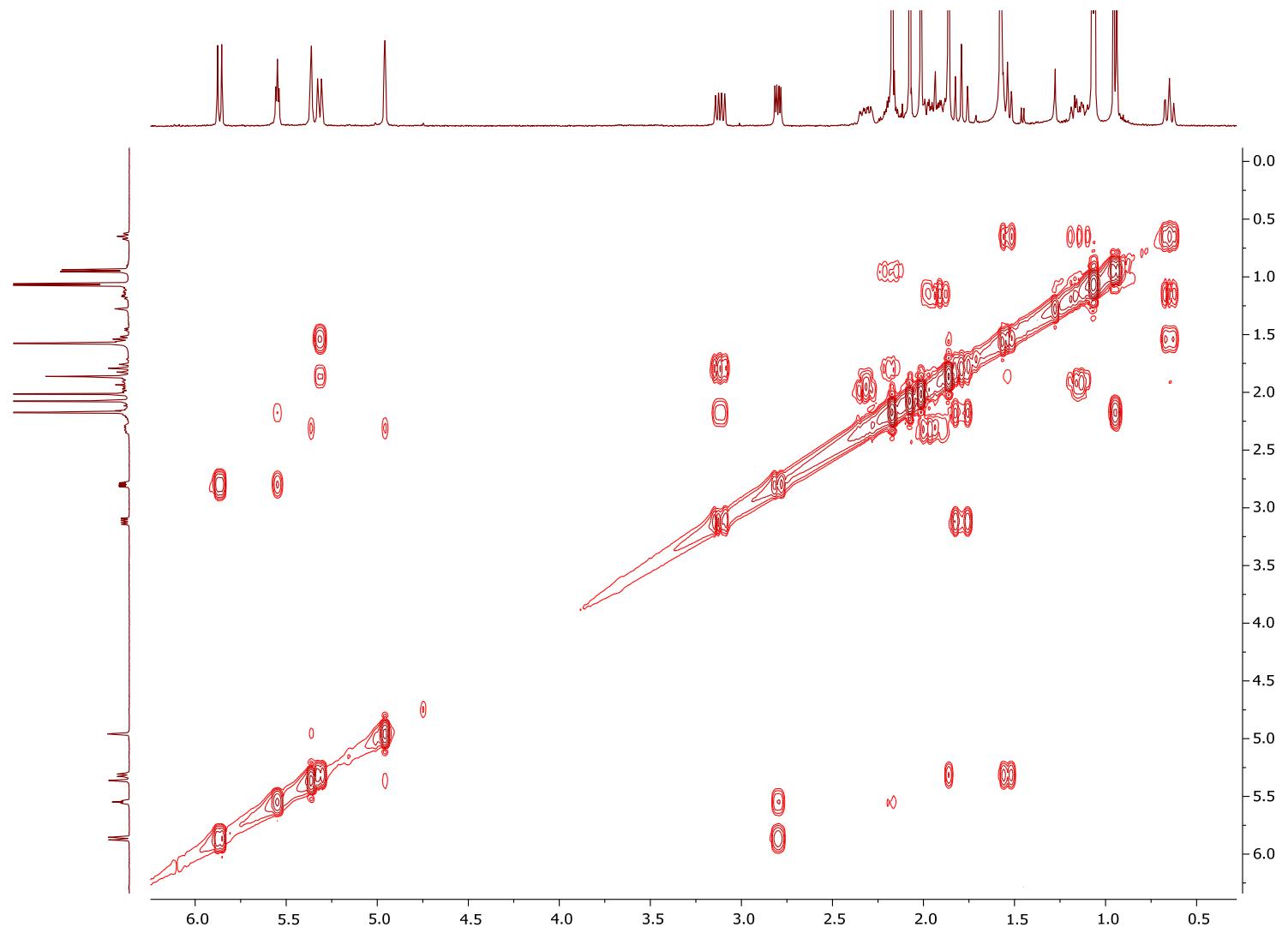


Figure S29. gCOSY spectrum of compound 6.

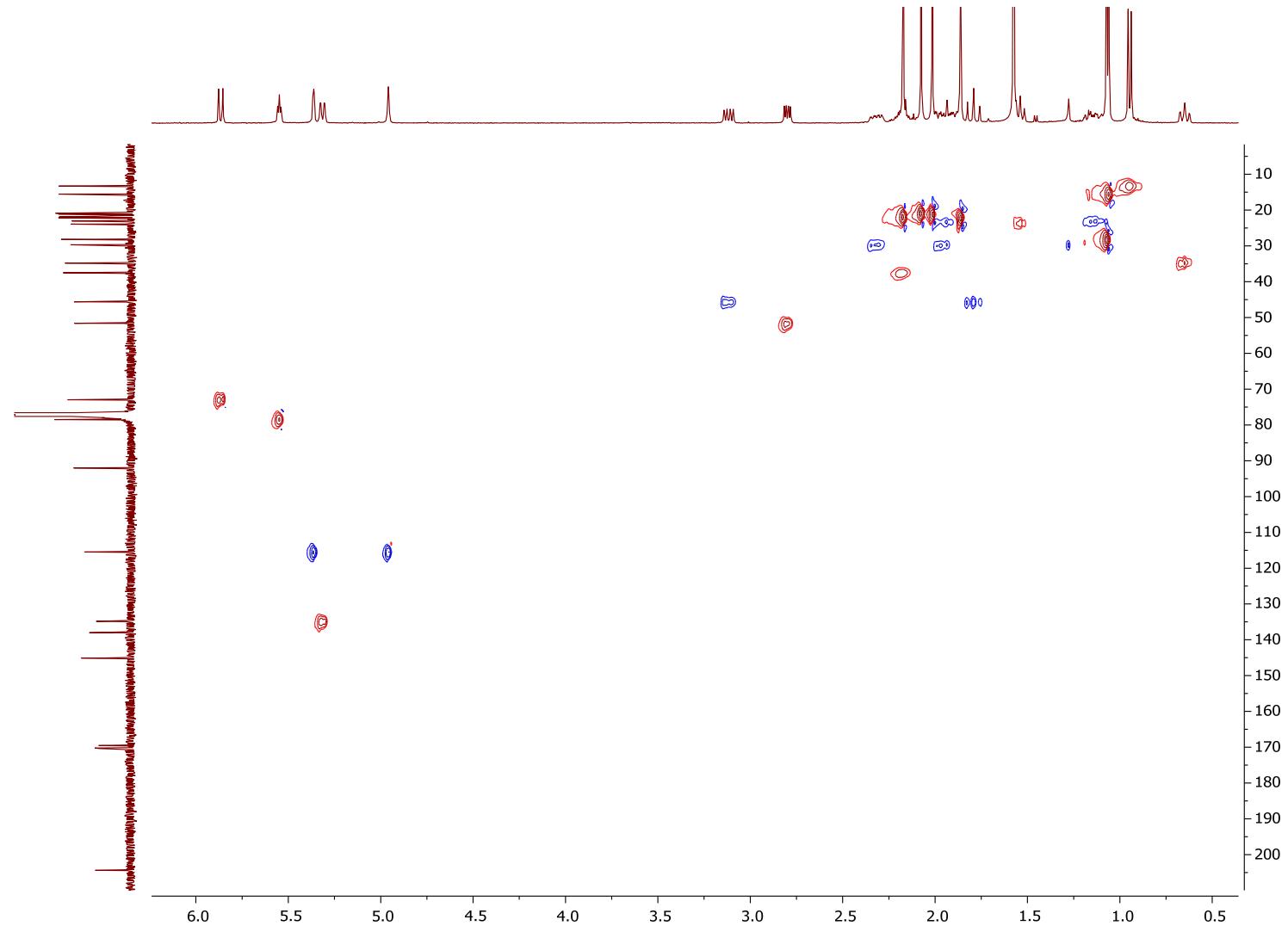


Figure S30. gHSQC spectrum of compound 6.

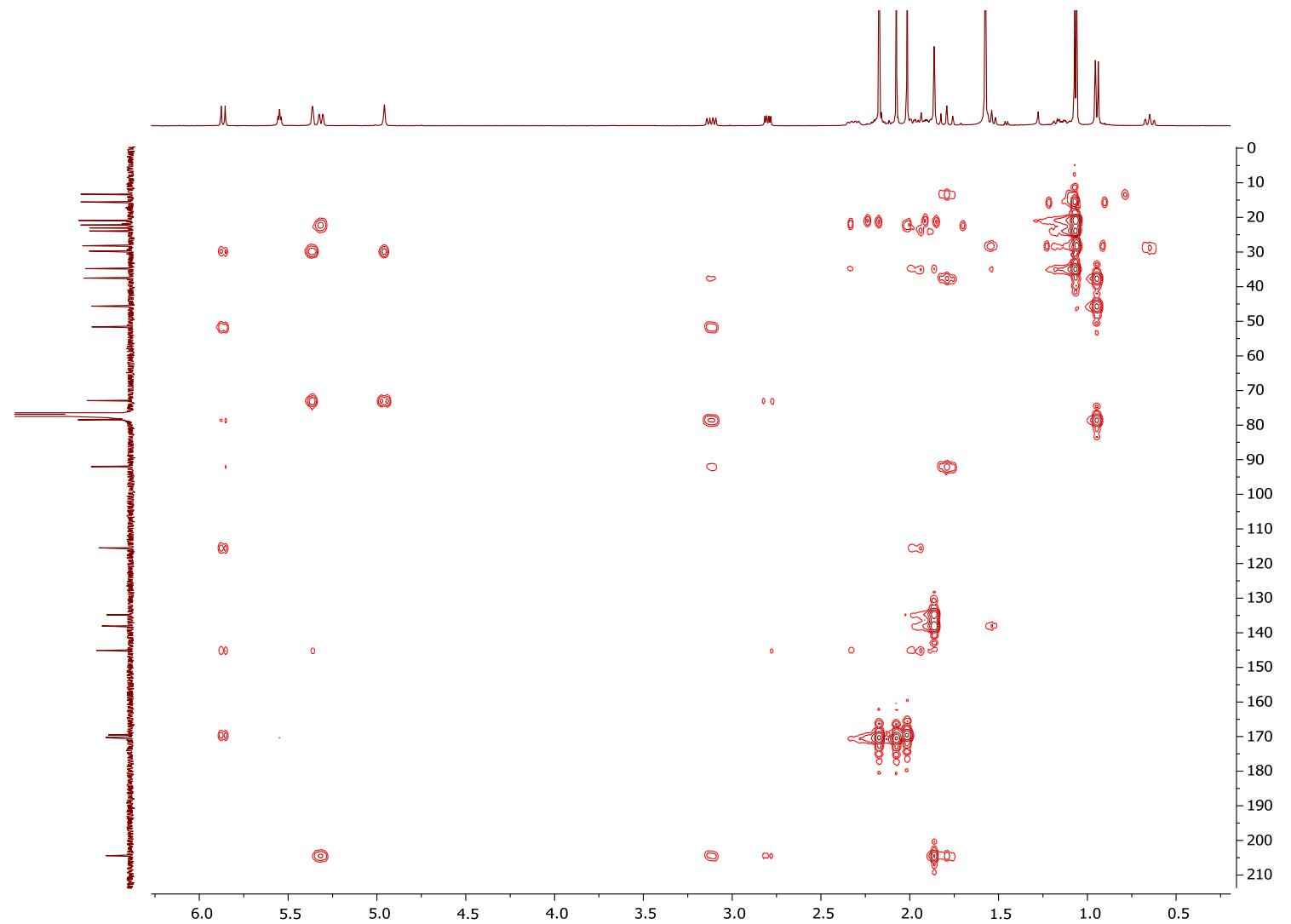


Figure S31. gHMBC spectrum of compound 6.

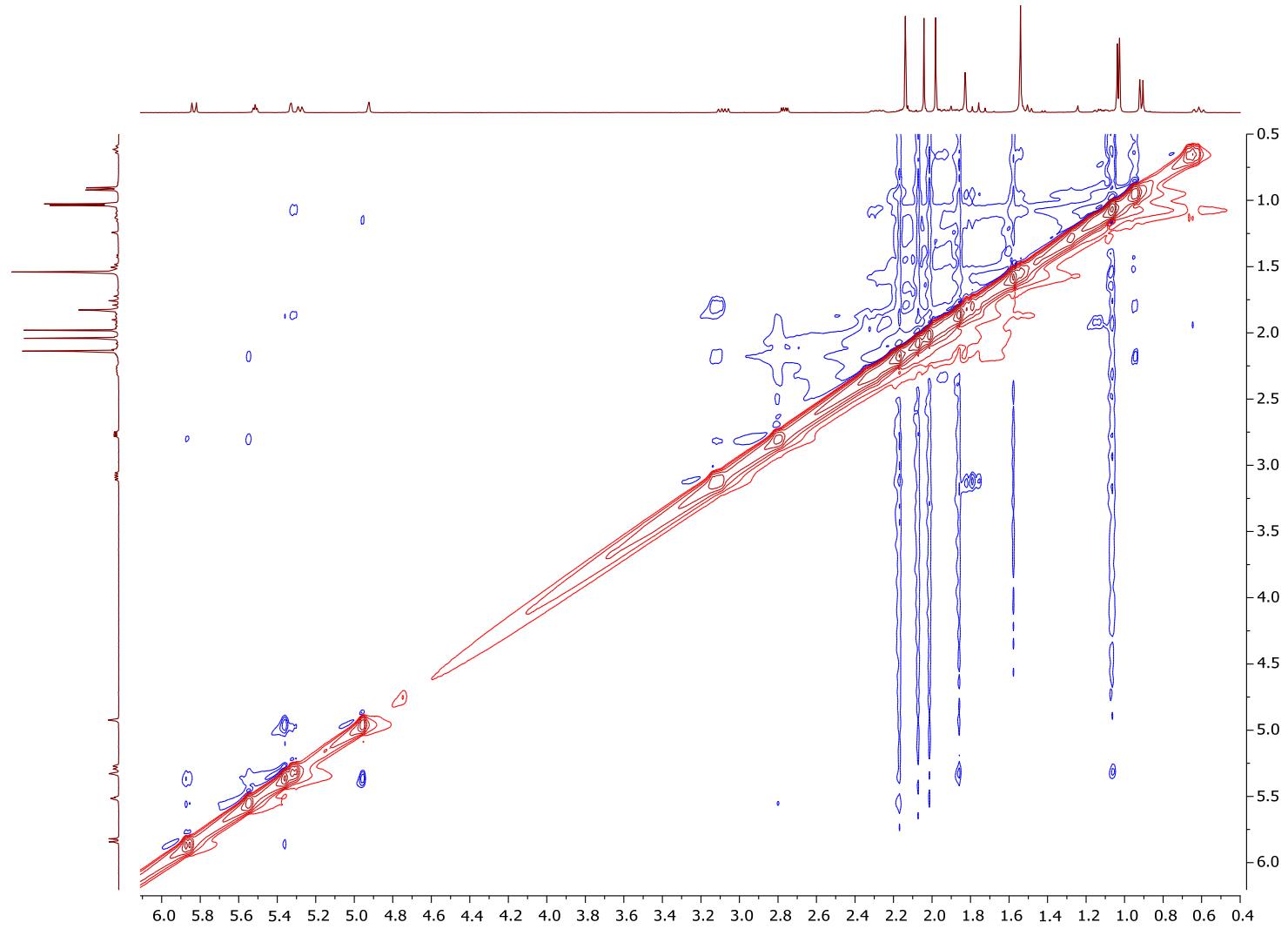


Figure S32. 2D NOESY spectrum of compound 6.

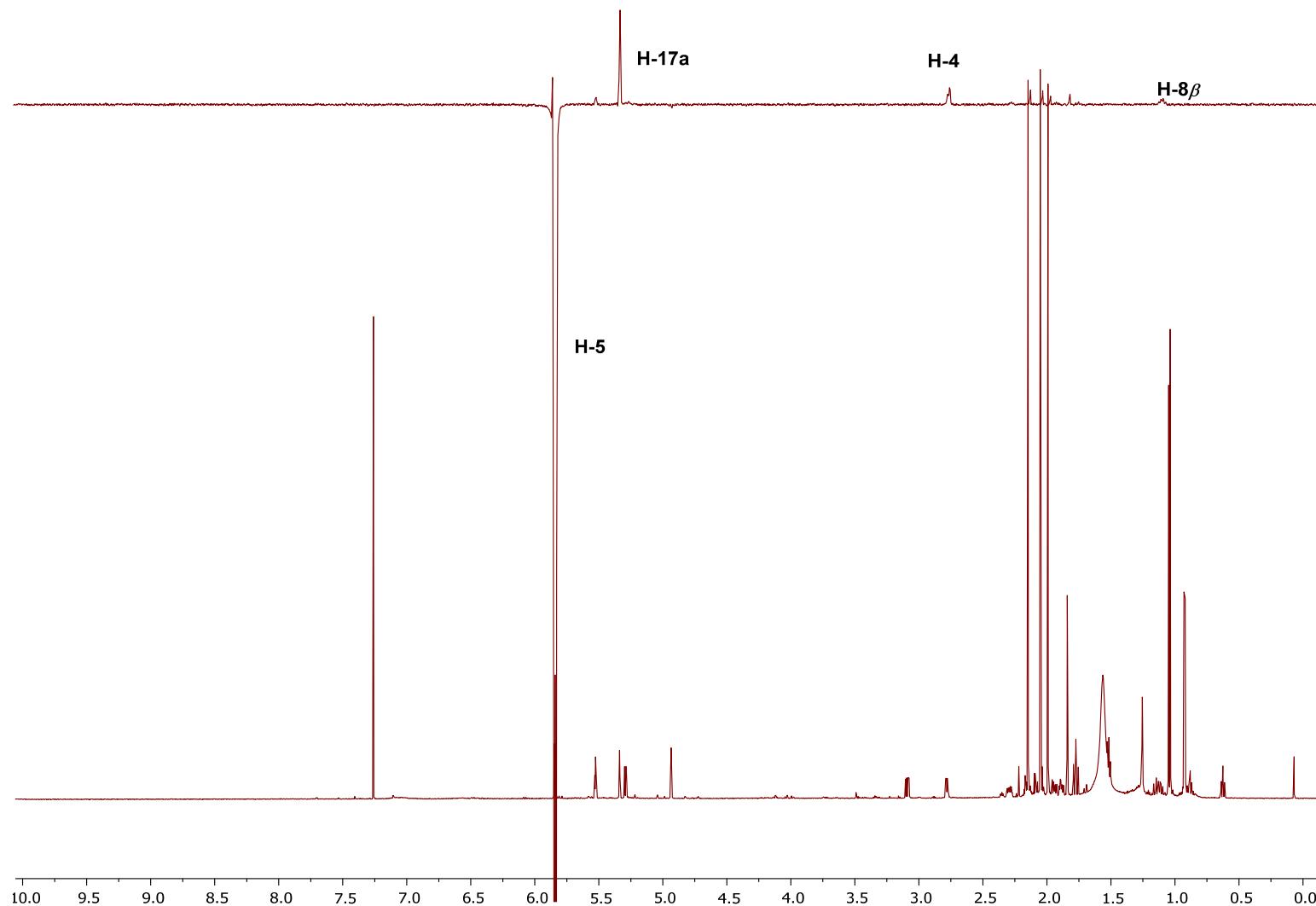


Figure S33a. 1D NOESY spectrum of compound 6.

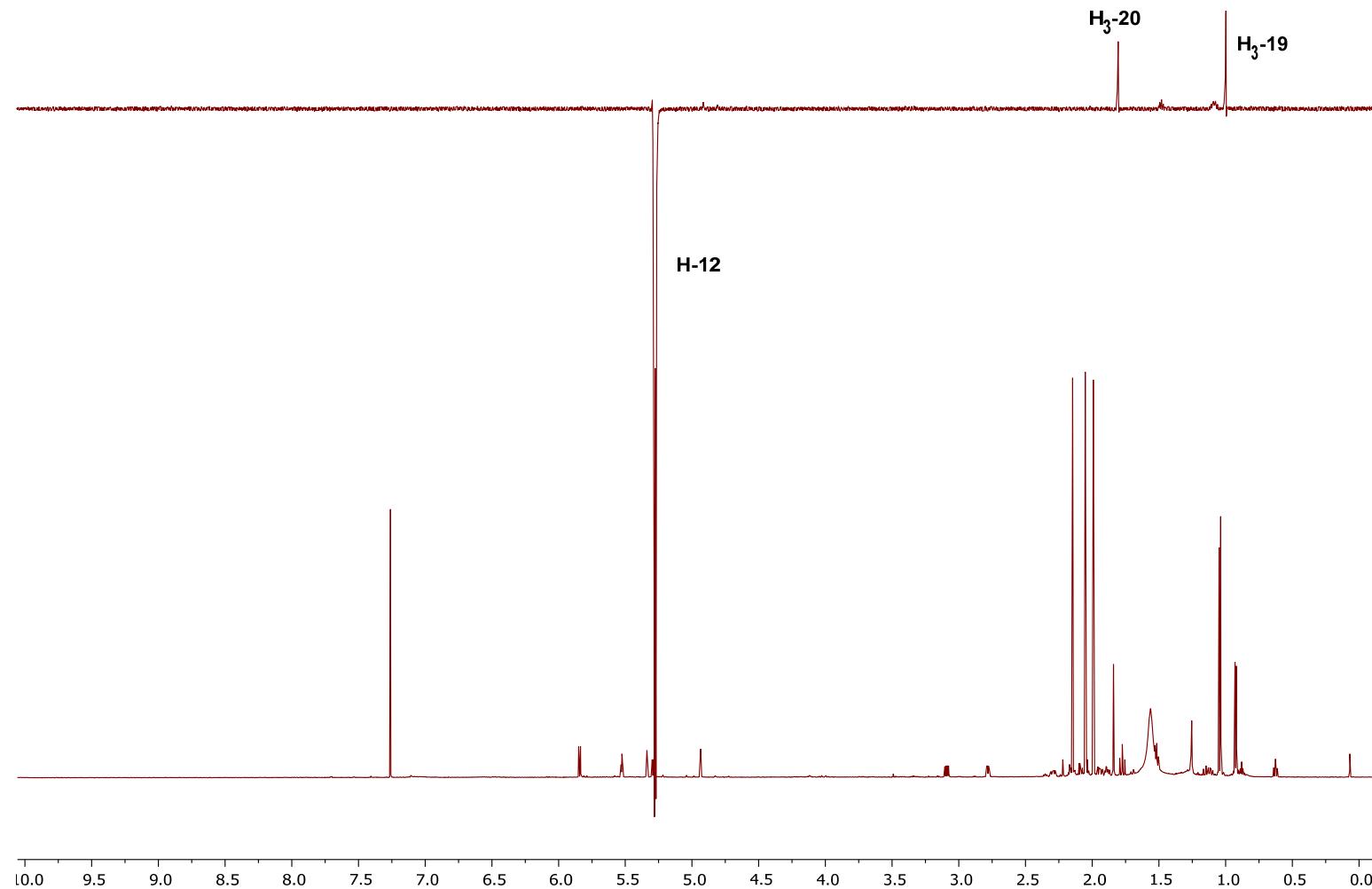


Figure S33b. 1D NOESY spectrum of compound 6.

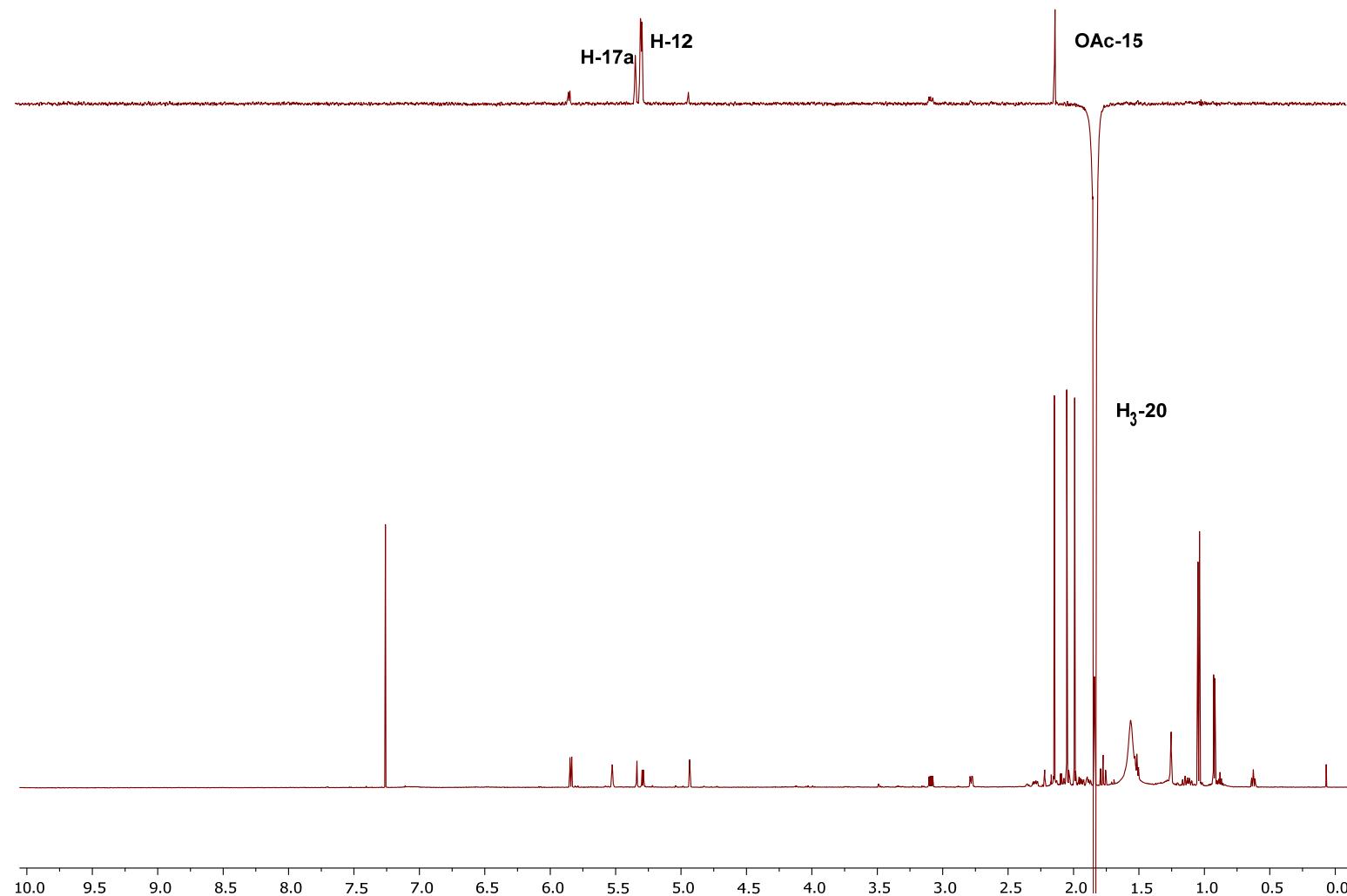


Figure S33c. 1D NOESY spectrum of compound 6.

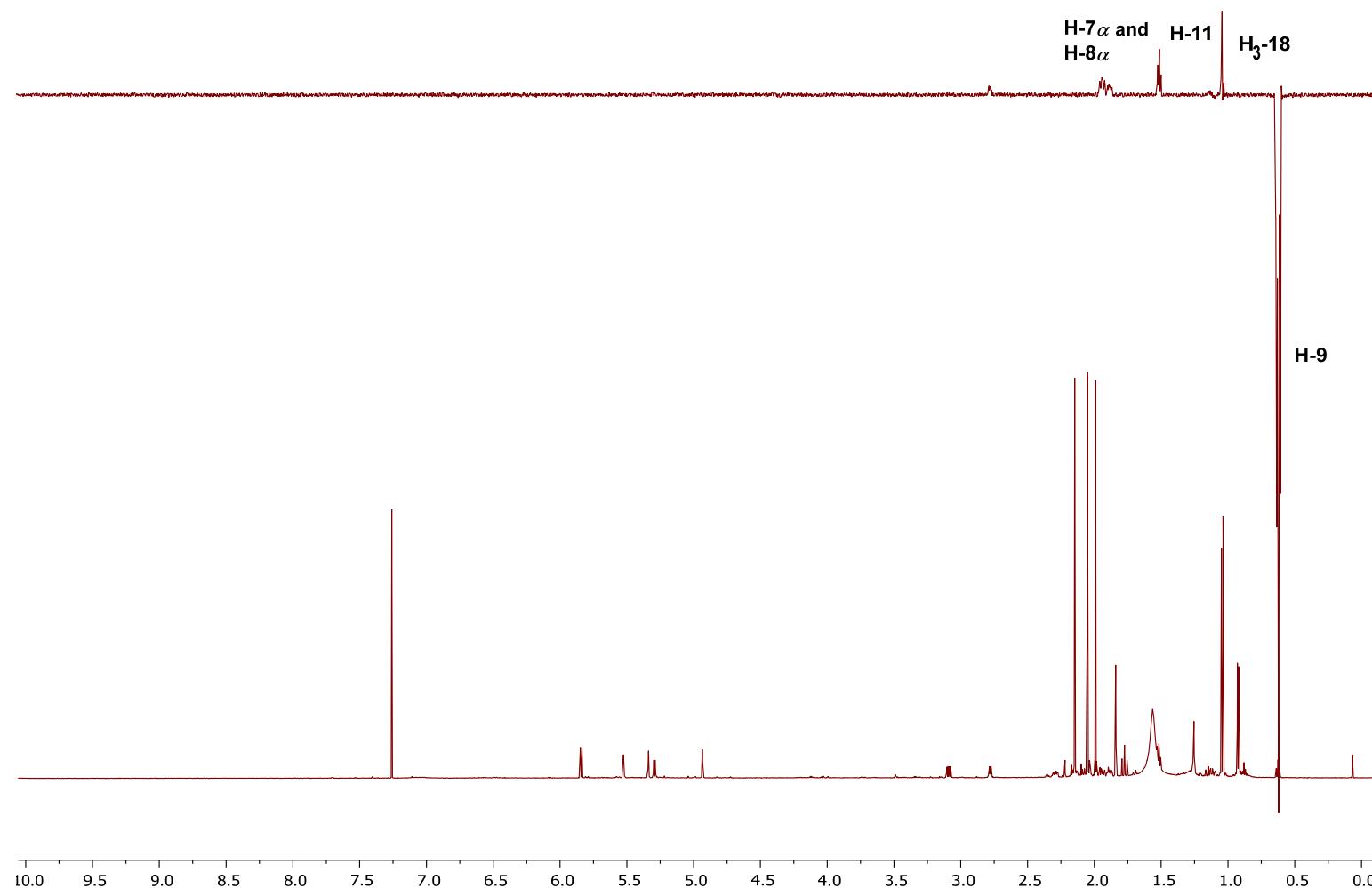


Figure S33d. 1D NOESY spectrum of compound 6.

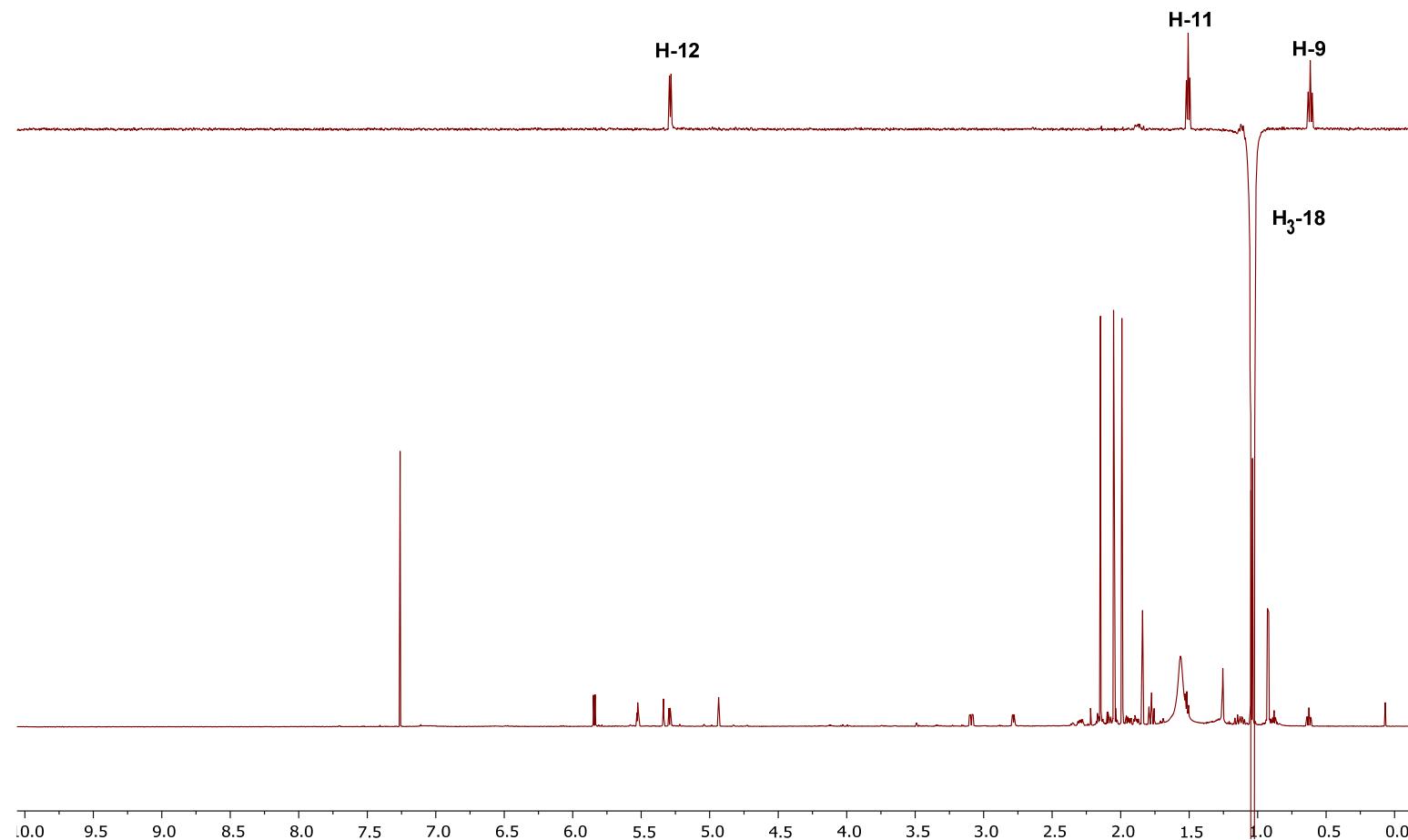


Figure S33e. 1D NOESY spectrum of compound 6.

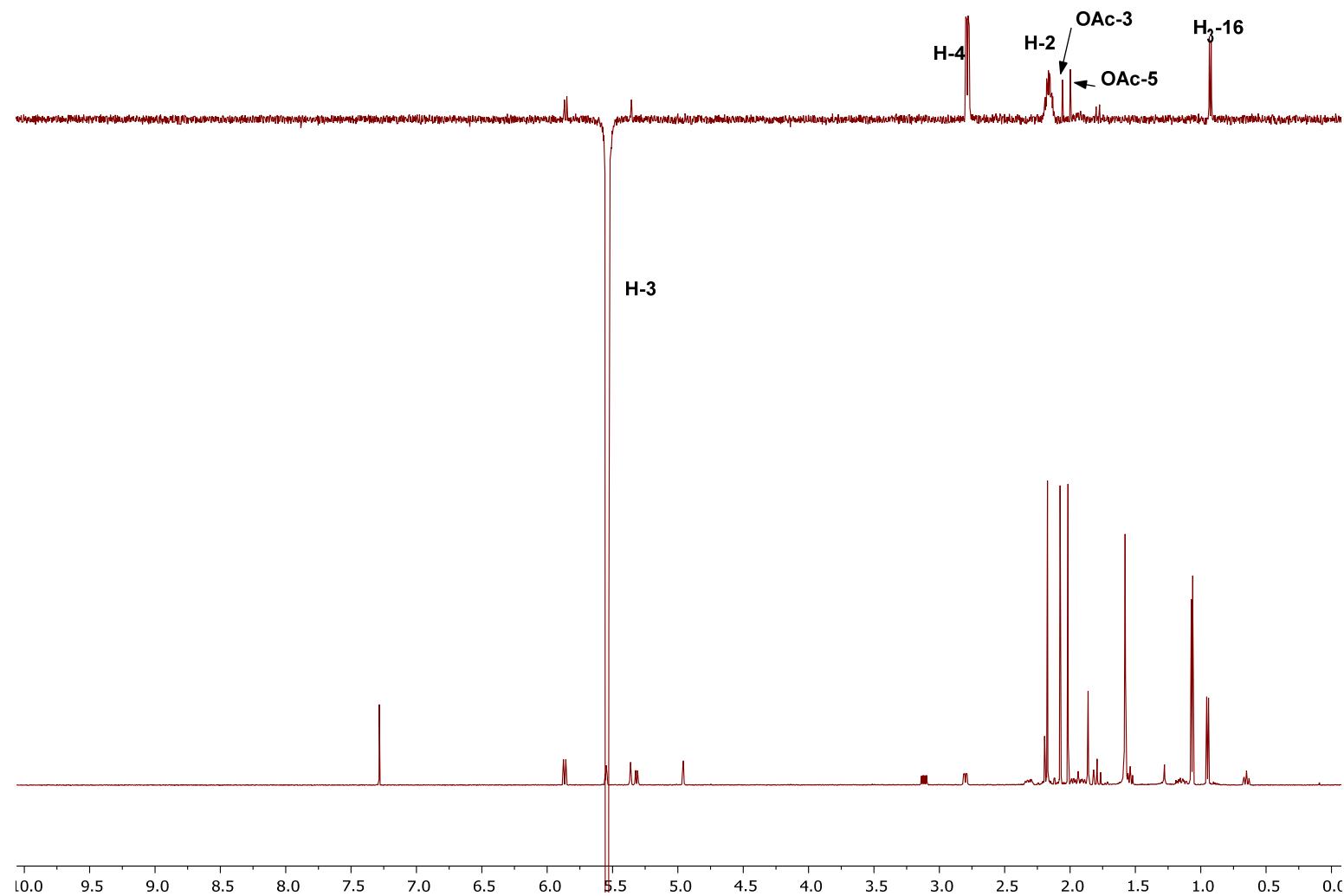


Figure S33f. 1D NOESY spectrum of compound 6.

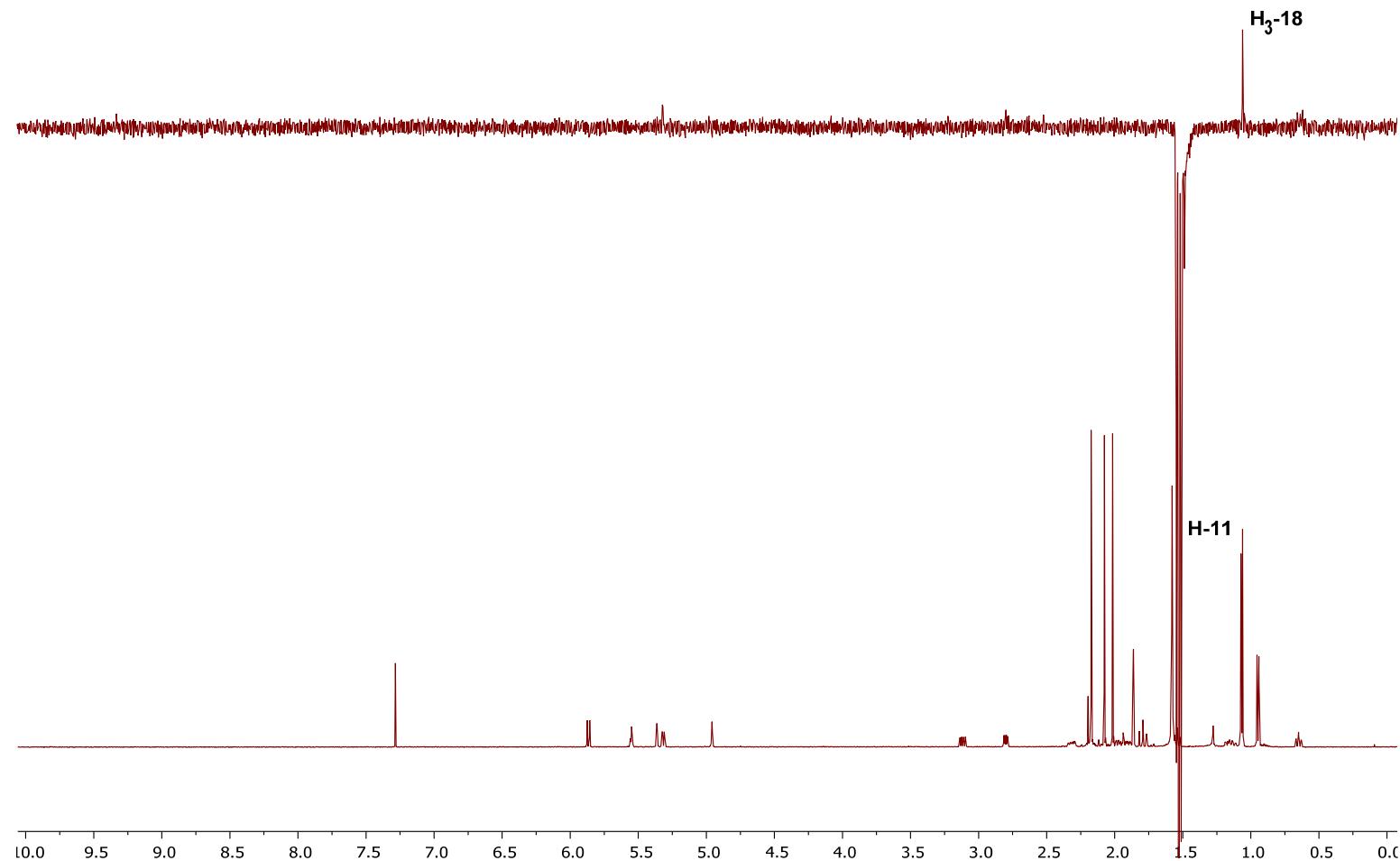


Figure S33g. 1D NOESY spectrum of compound 6.

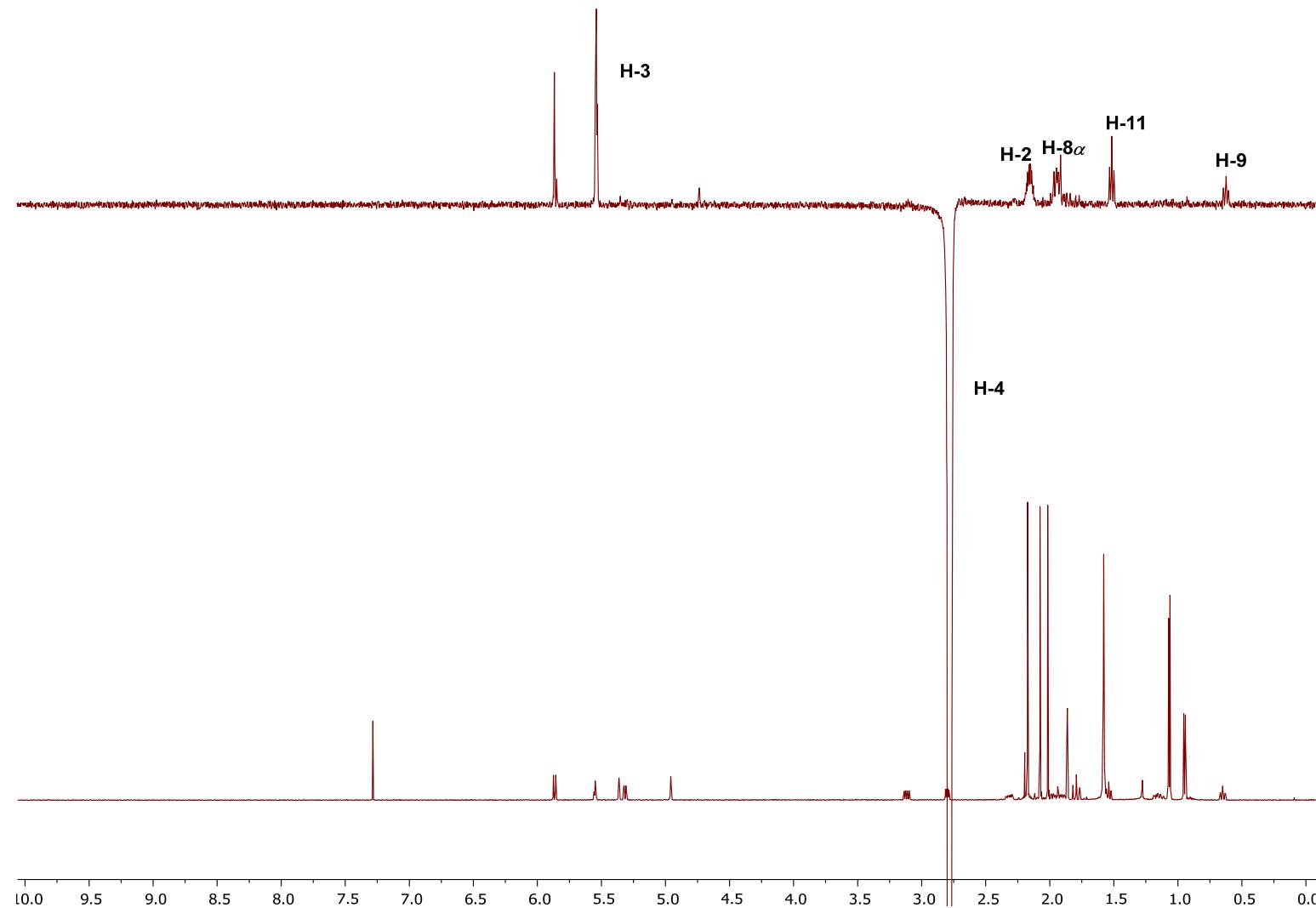


Figure S33h. 1D NOESY spectrum of compound 6.

Elemental Composition Report

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

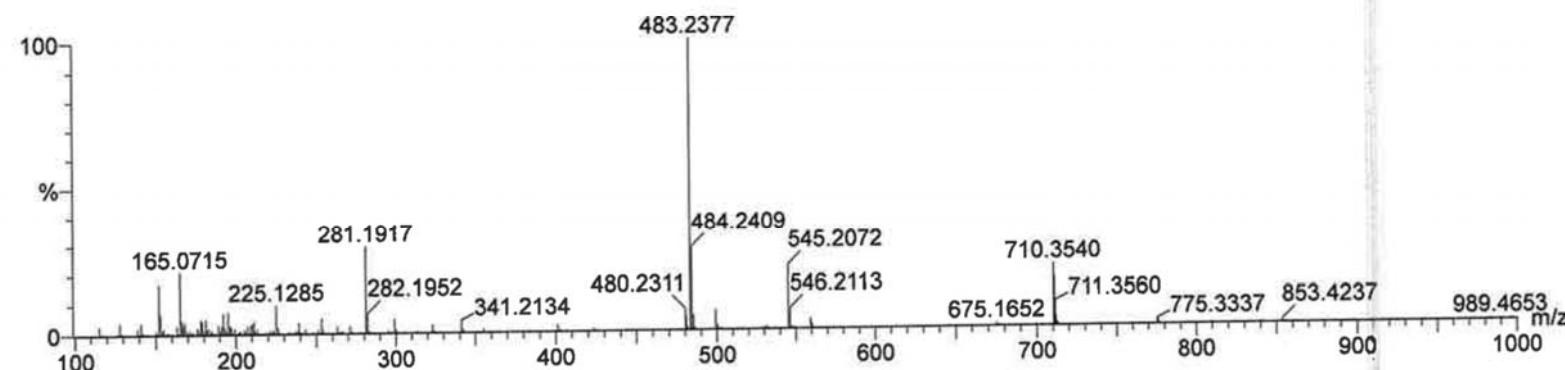
Monoisotopic Mass, Even Electron Ions

93 formula(e) evaluated with 3 results within limits (all results (up to 1000) for each mass)

Elements Used:

C: 0-30 H: 0-50 O: 0-15 23Na: 0-1

FEM-253 520 (4.808)

1: TOF MS ES+
6.86e+005

Minimum:			-1.5
Maximum:	5.0	10.0	80.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
483.2377	483.2359	1.8	3.7	8.5	1058.8	0.001	99.86	C26 H36 O7 23Na
	483.2383	-0.6	-1.2	11.5	1065.4	6.598	0.14	C28 H35 O7
	483.2417	-4.0	-8.3	-0.5	1068.7	9.859	0.01	C19 H40 O12 23Na

Figure S34. HRMS of compound 6.

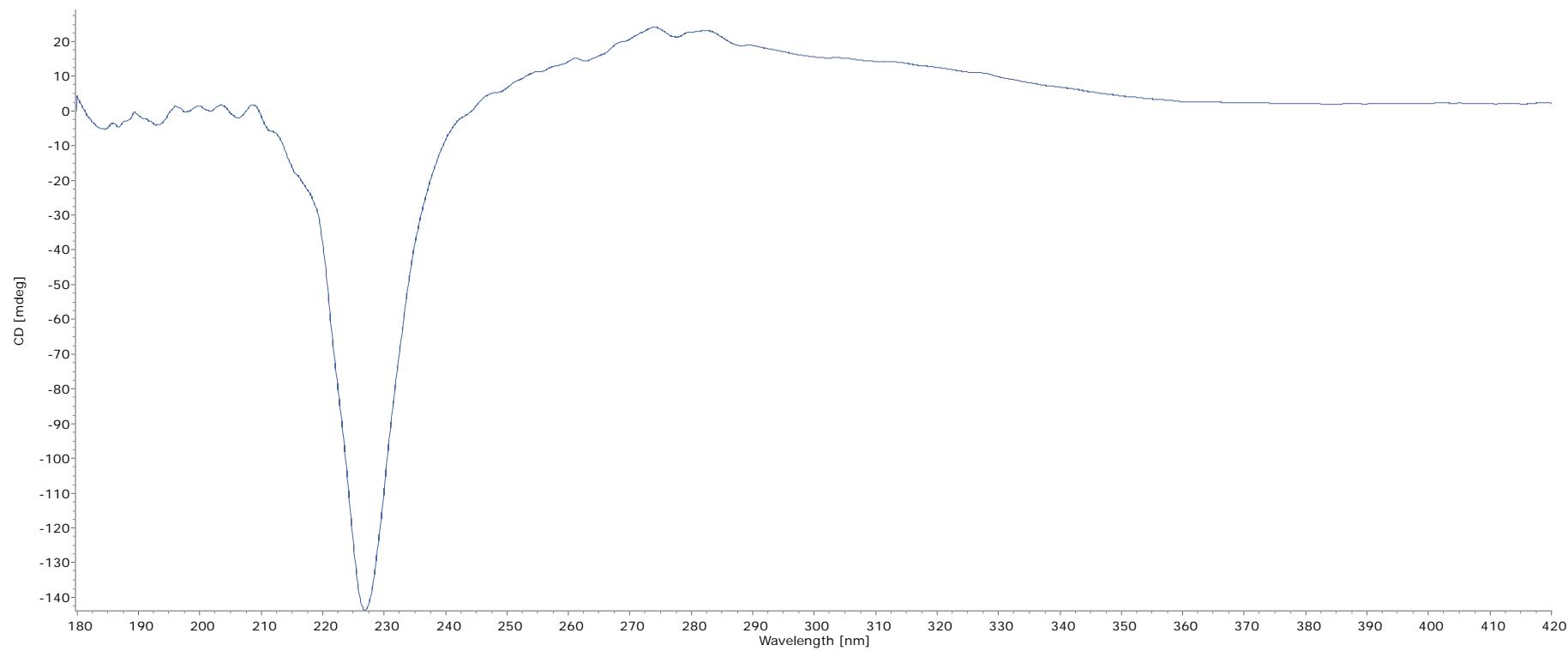


Figure S35. ECD of compound **6**.

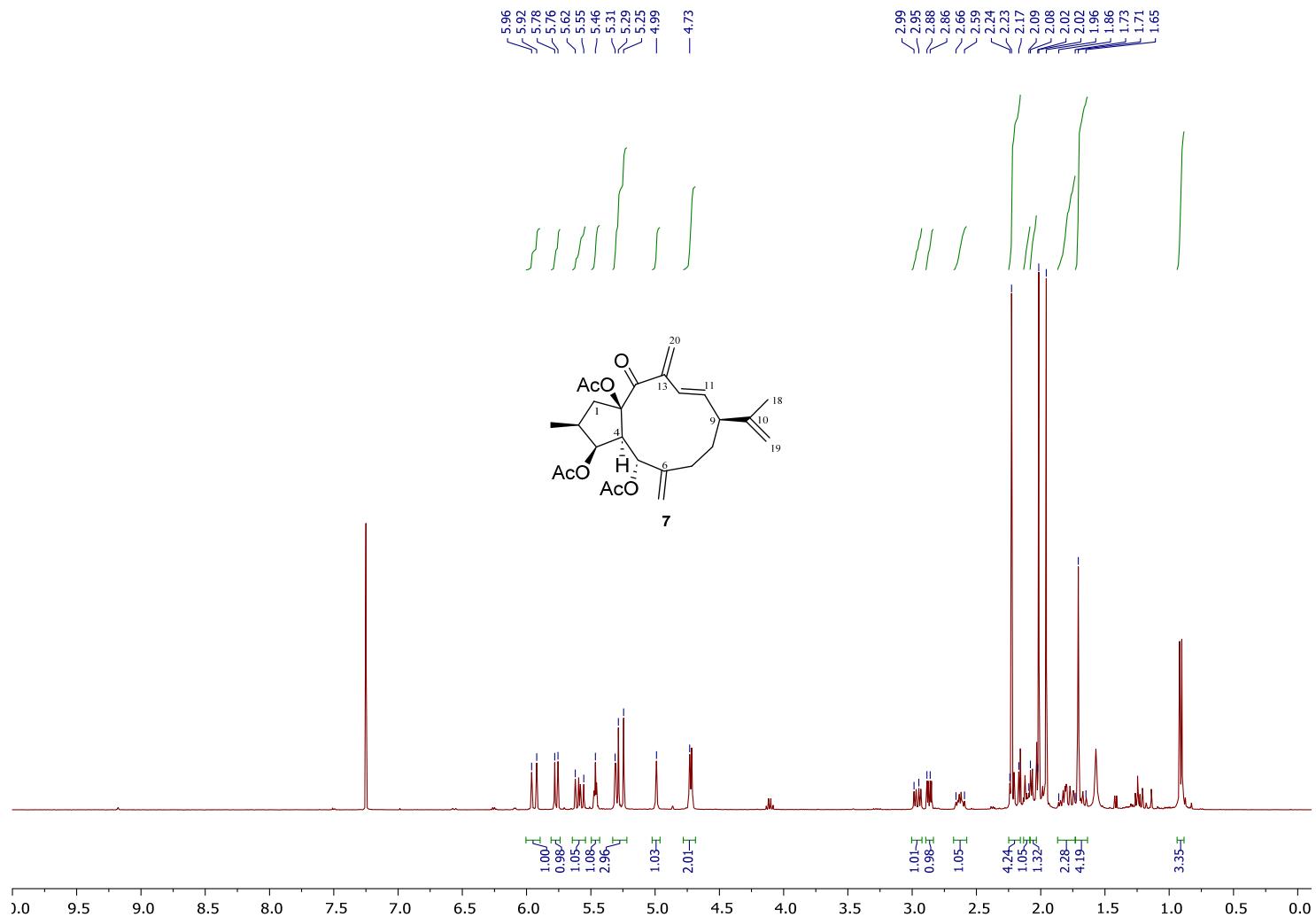


Figure S36. ^1H NMR spectrum (400 MHz) of compound 7 in CDCl_3 .

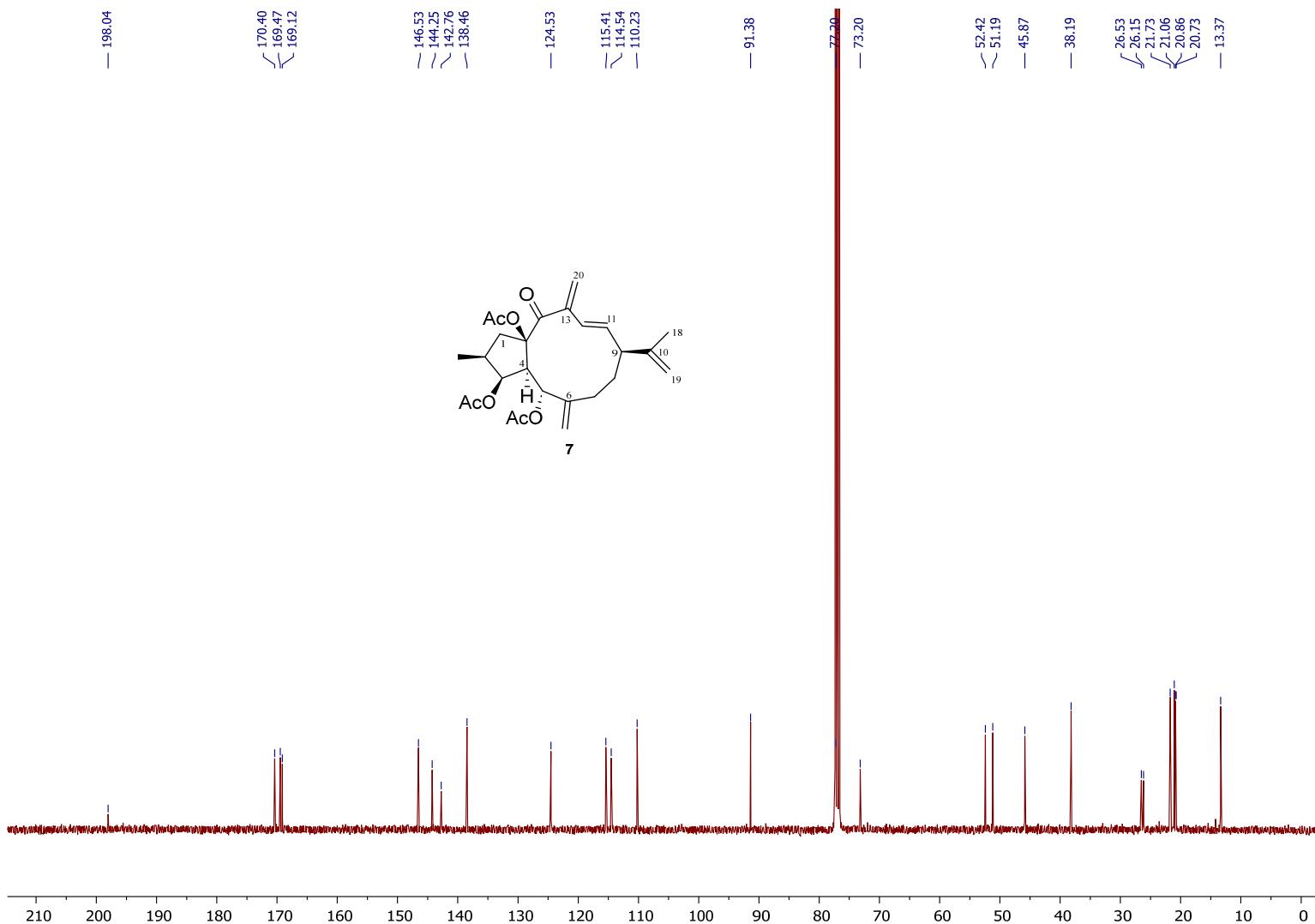


Figure S37. ^{13}C NMR spectrum (100 MHz) of compound **7** in CDCl_3 .

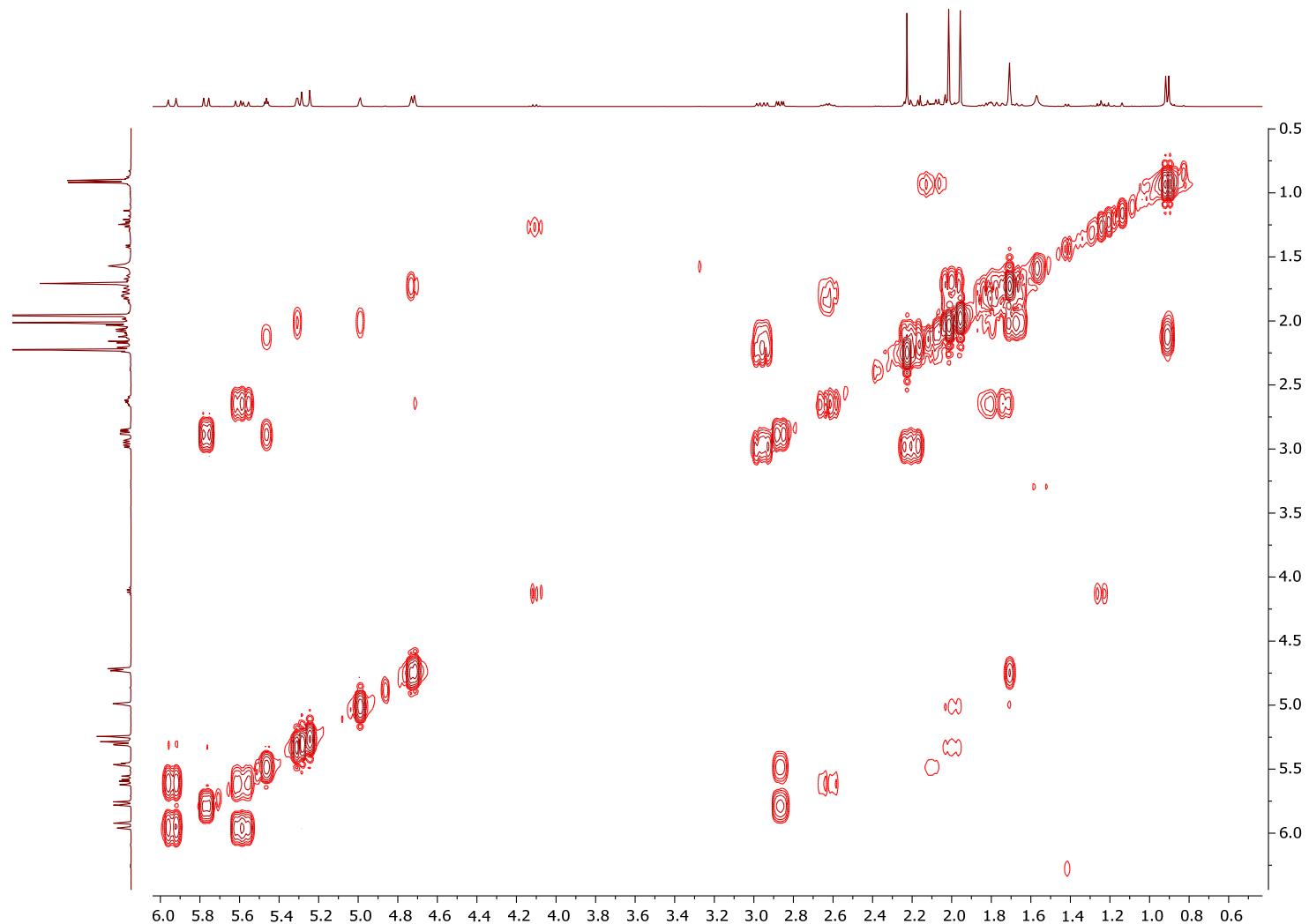


Figure S38. gCOSY spectrum of compound 7.

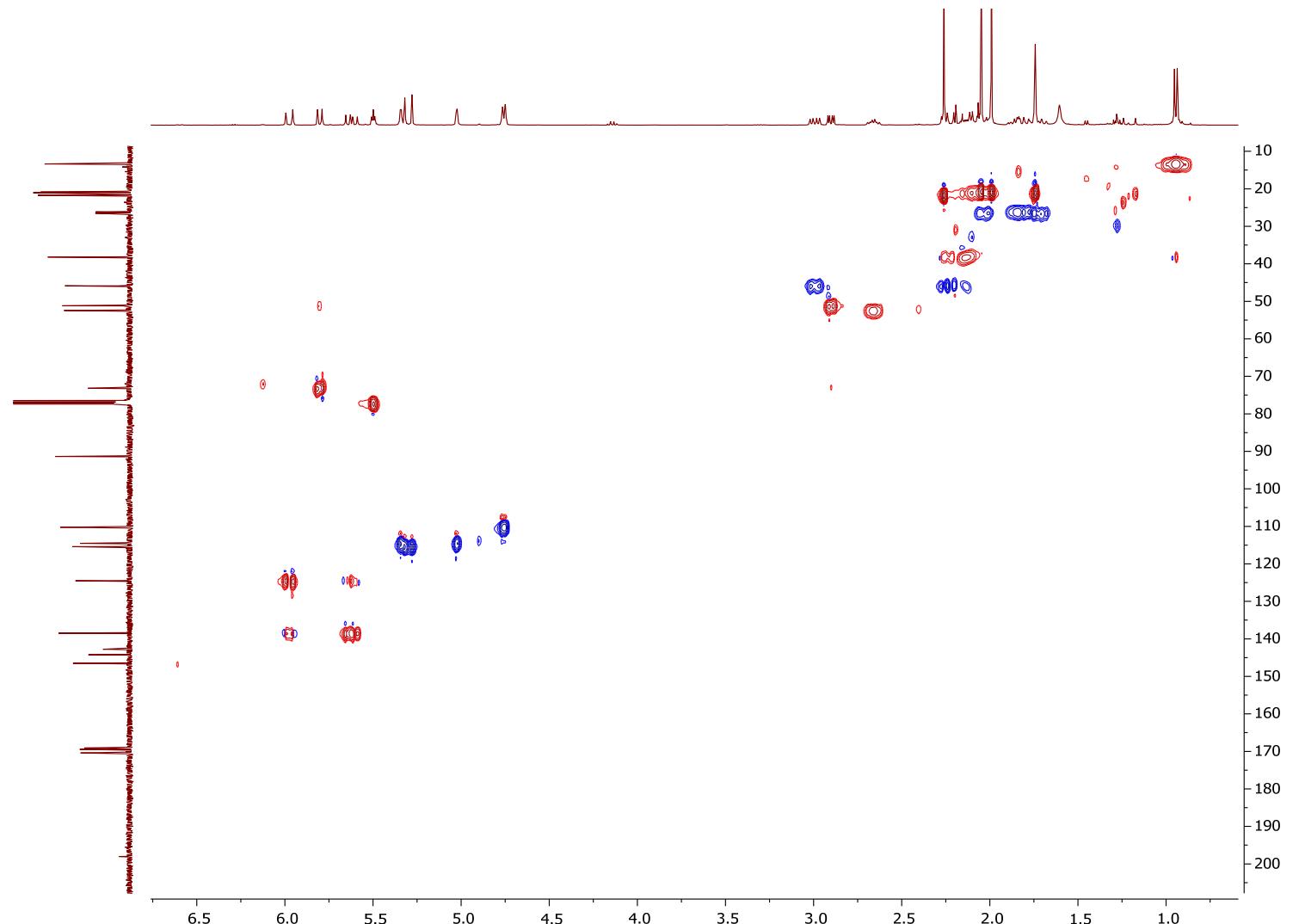


Figure S39. gHSQC spectrum of compound 7.

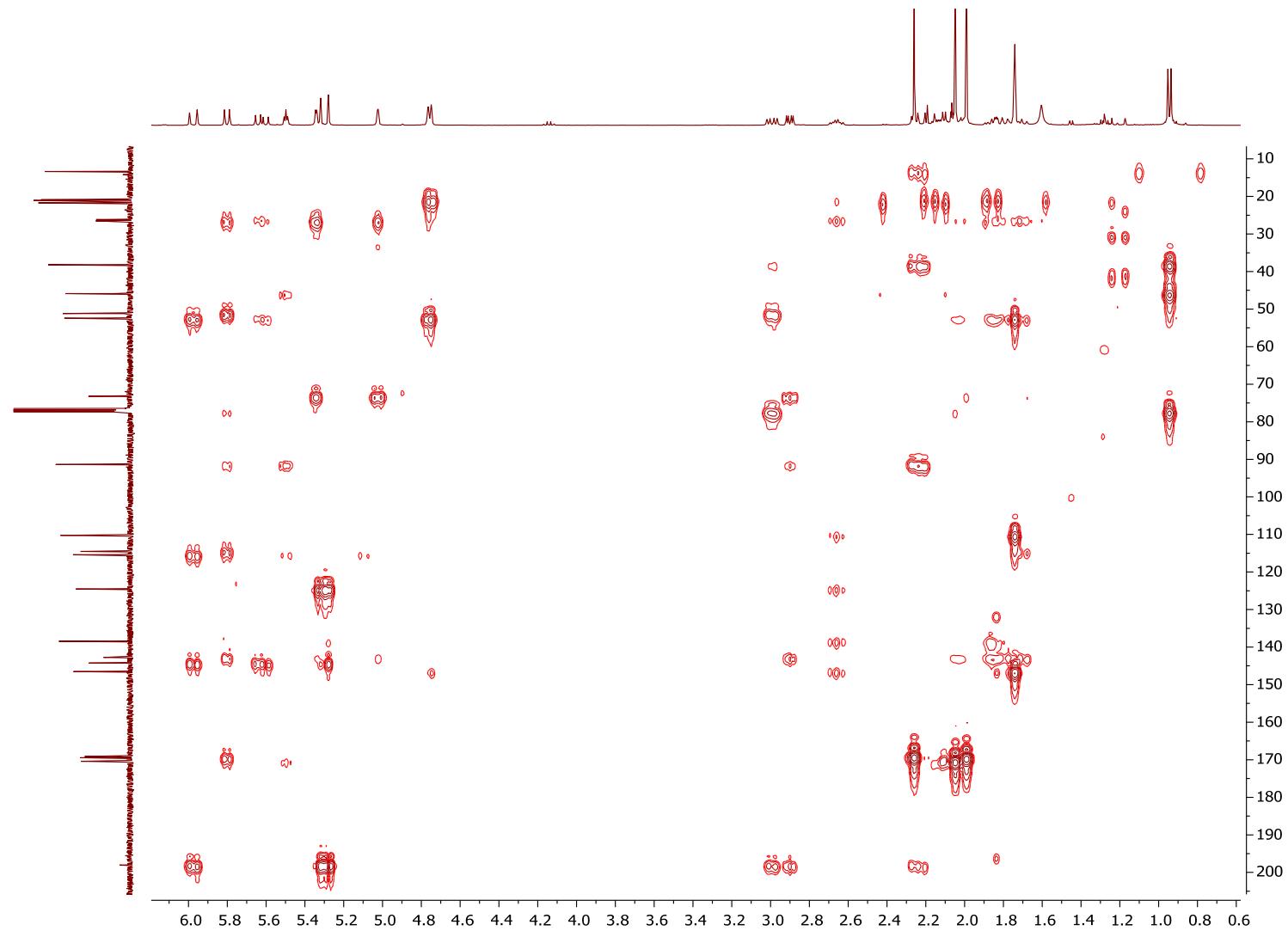


Figure S40. gHMBC spectrum of compound 7.

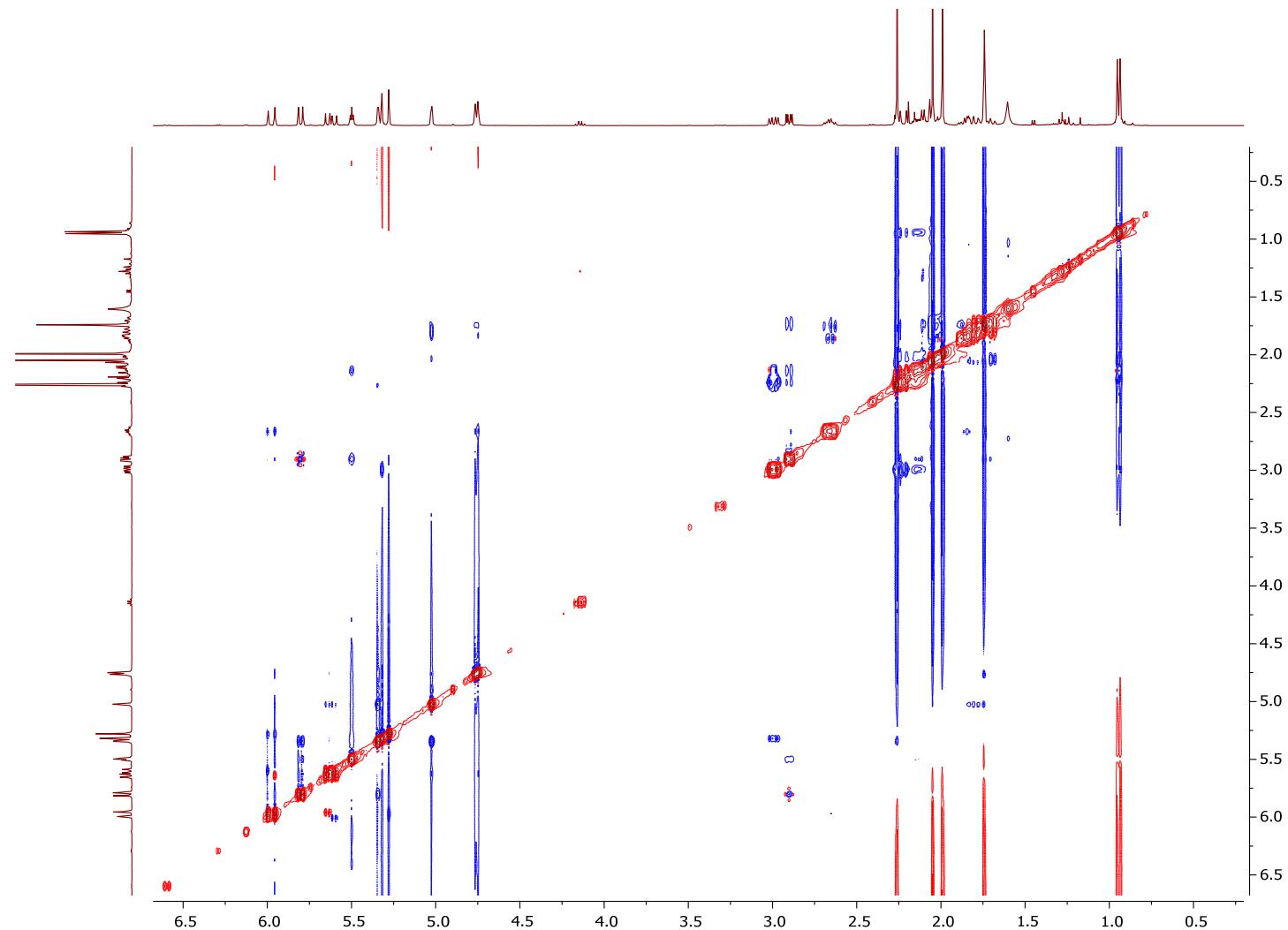


Figure S41. 2D NOESY spectrum of compound 7.

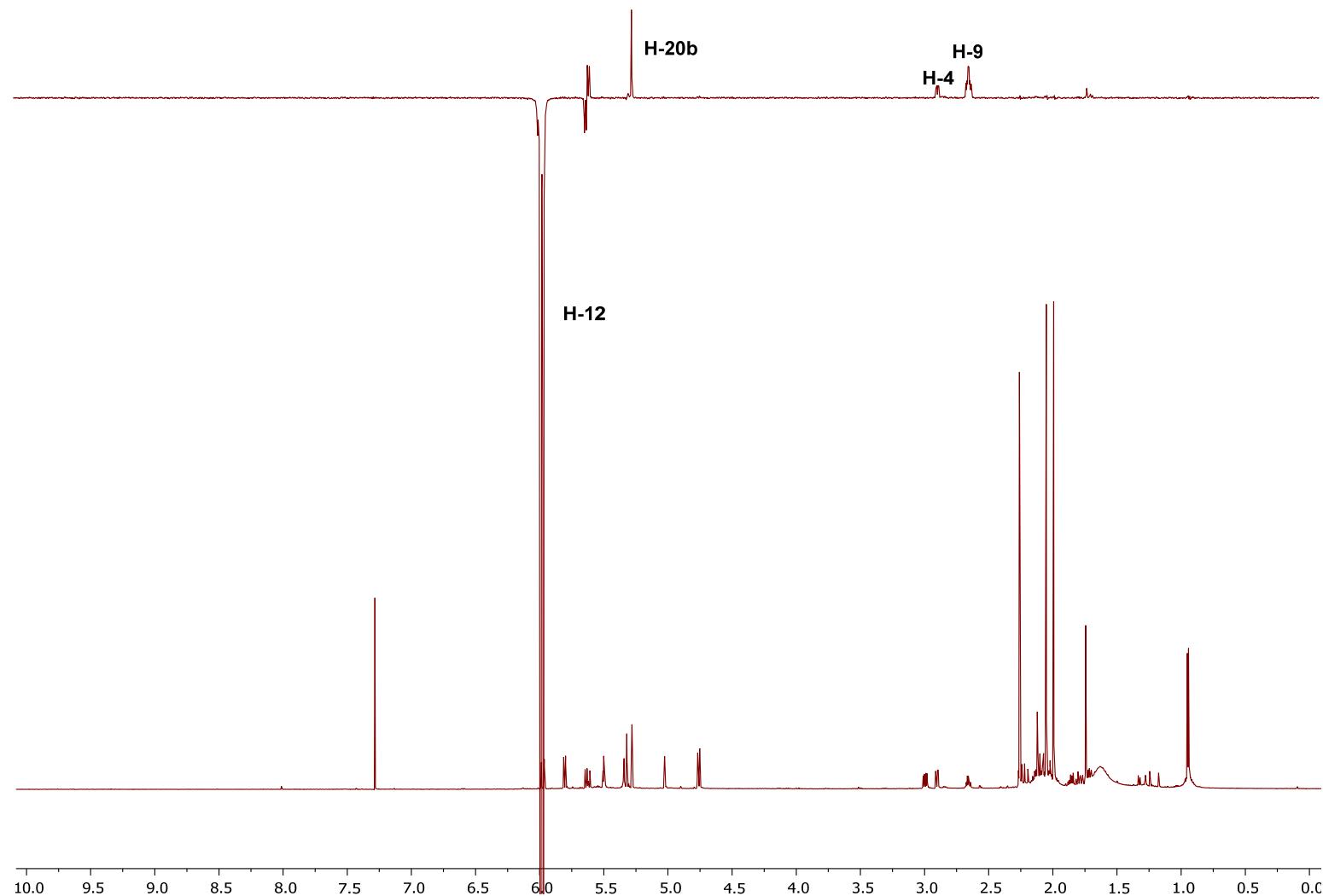


Figure S42a. 1D NOESY spectrum of compound 7.

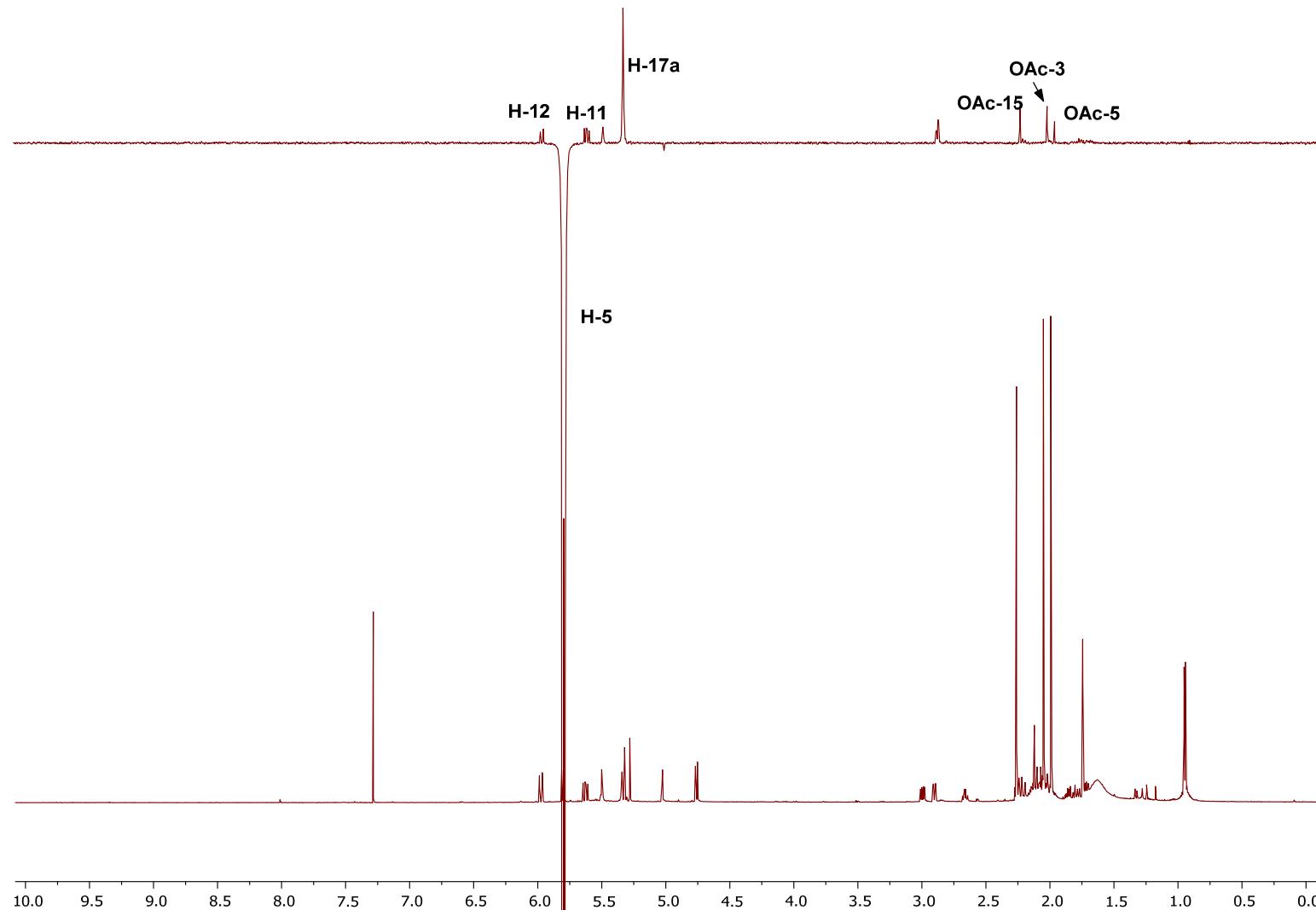


Figure S42b. 1D NOESY spectrum of compound 7.

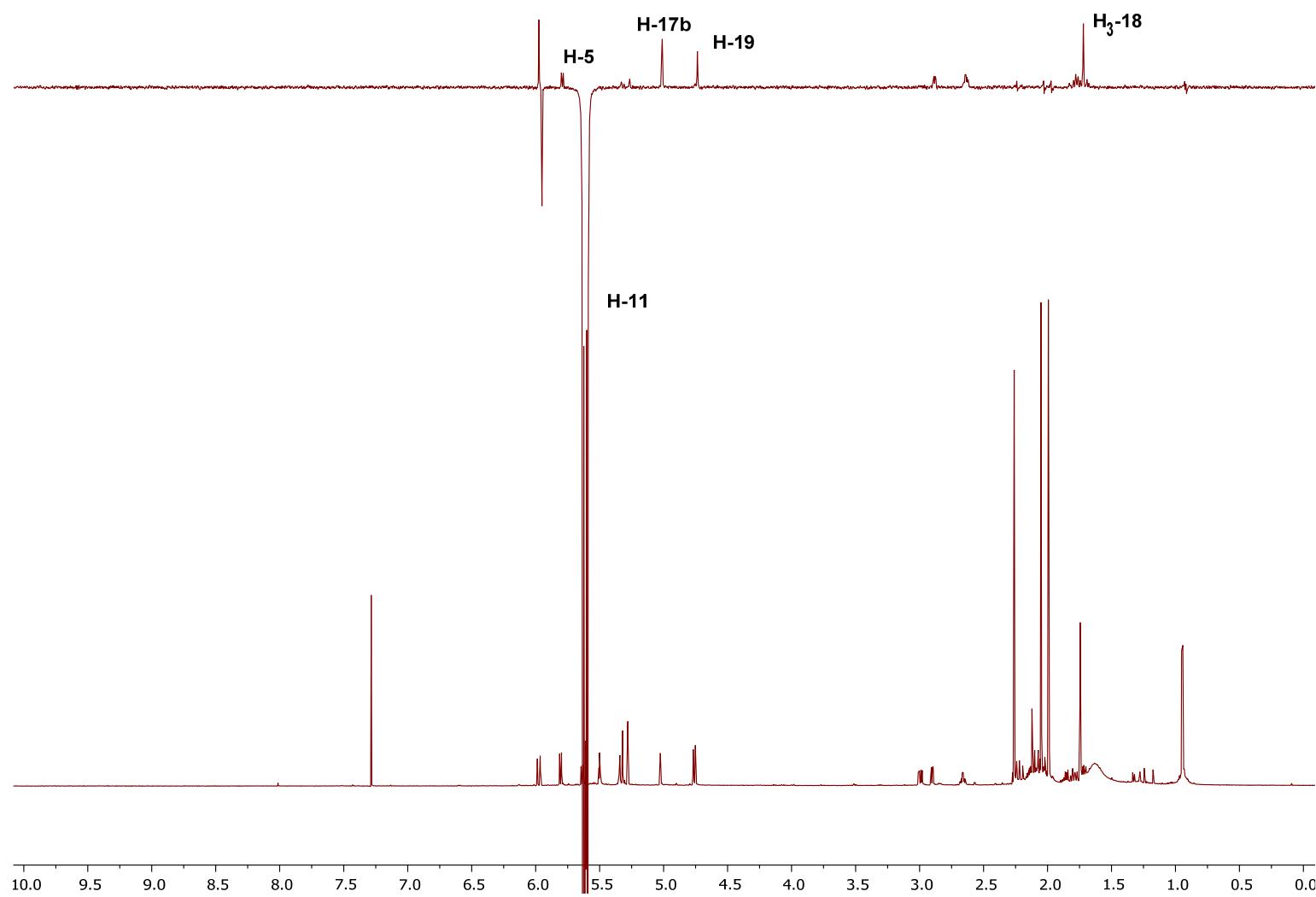


Figure S42c. 1D NOESY spectrum of compound 7.

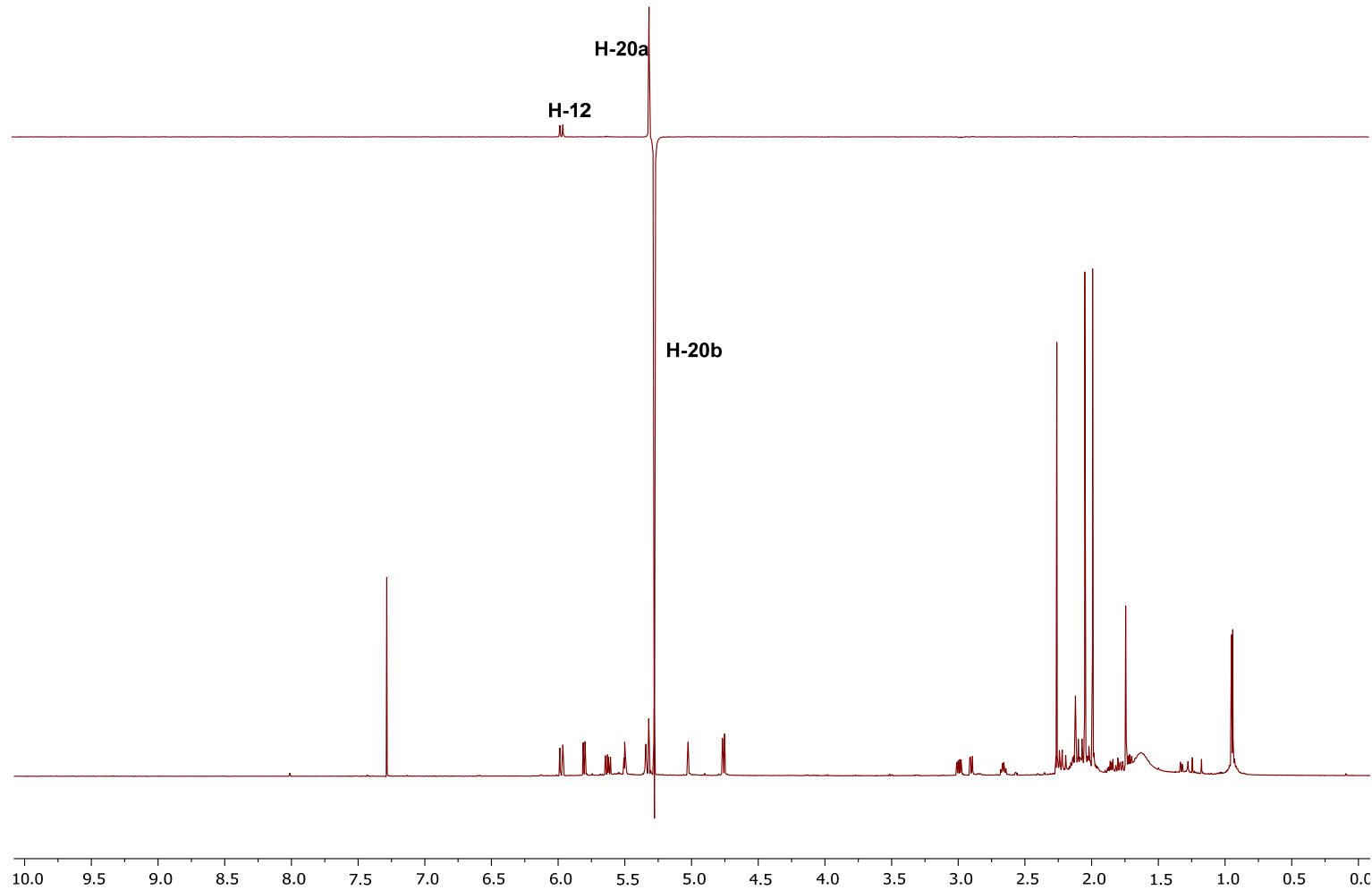


Figure S42d. 1D NOESY spectrum of compound 7.

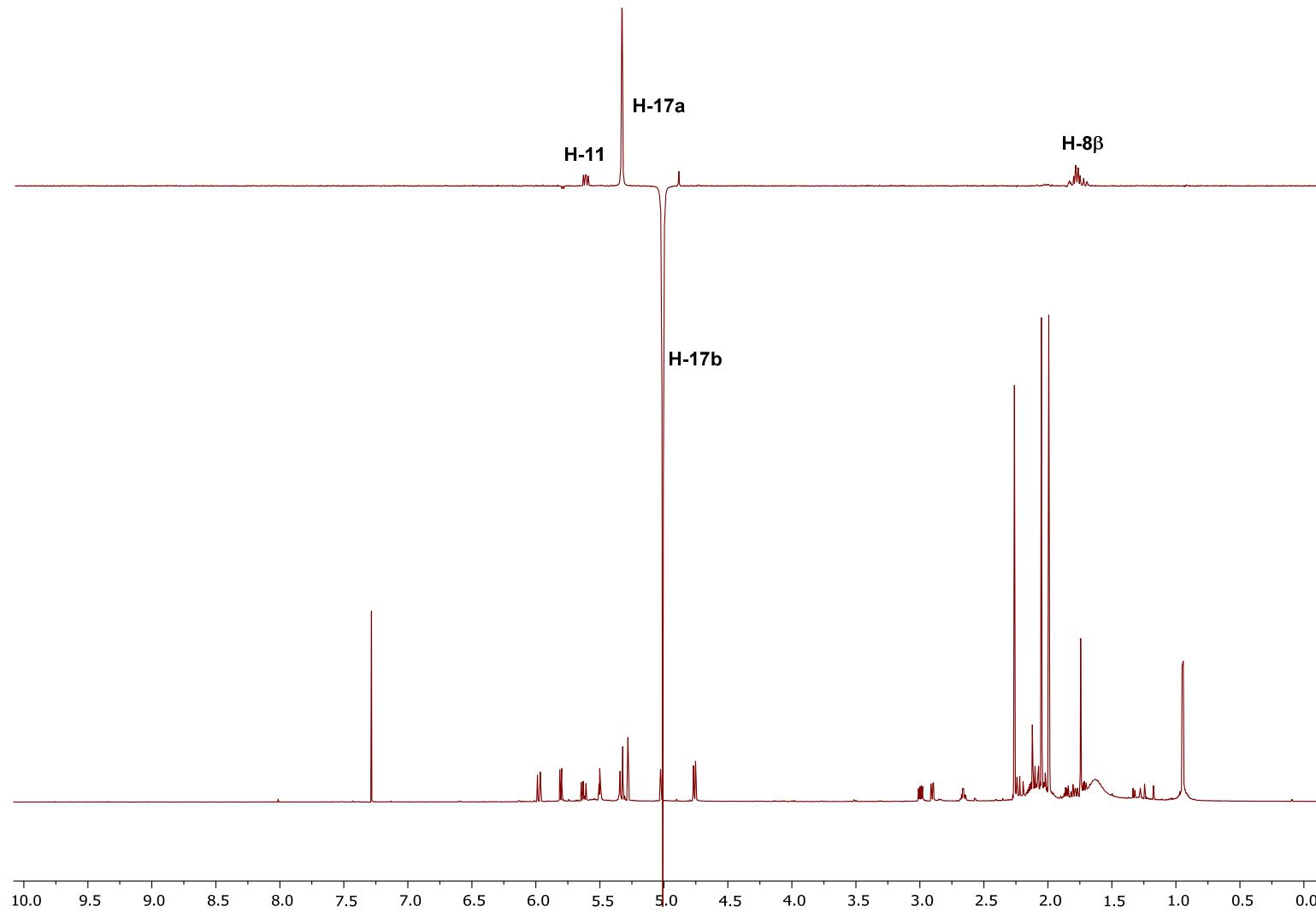


Figure S42e. 1D NOESY spectrum of compound 7.

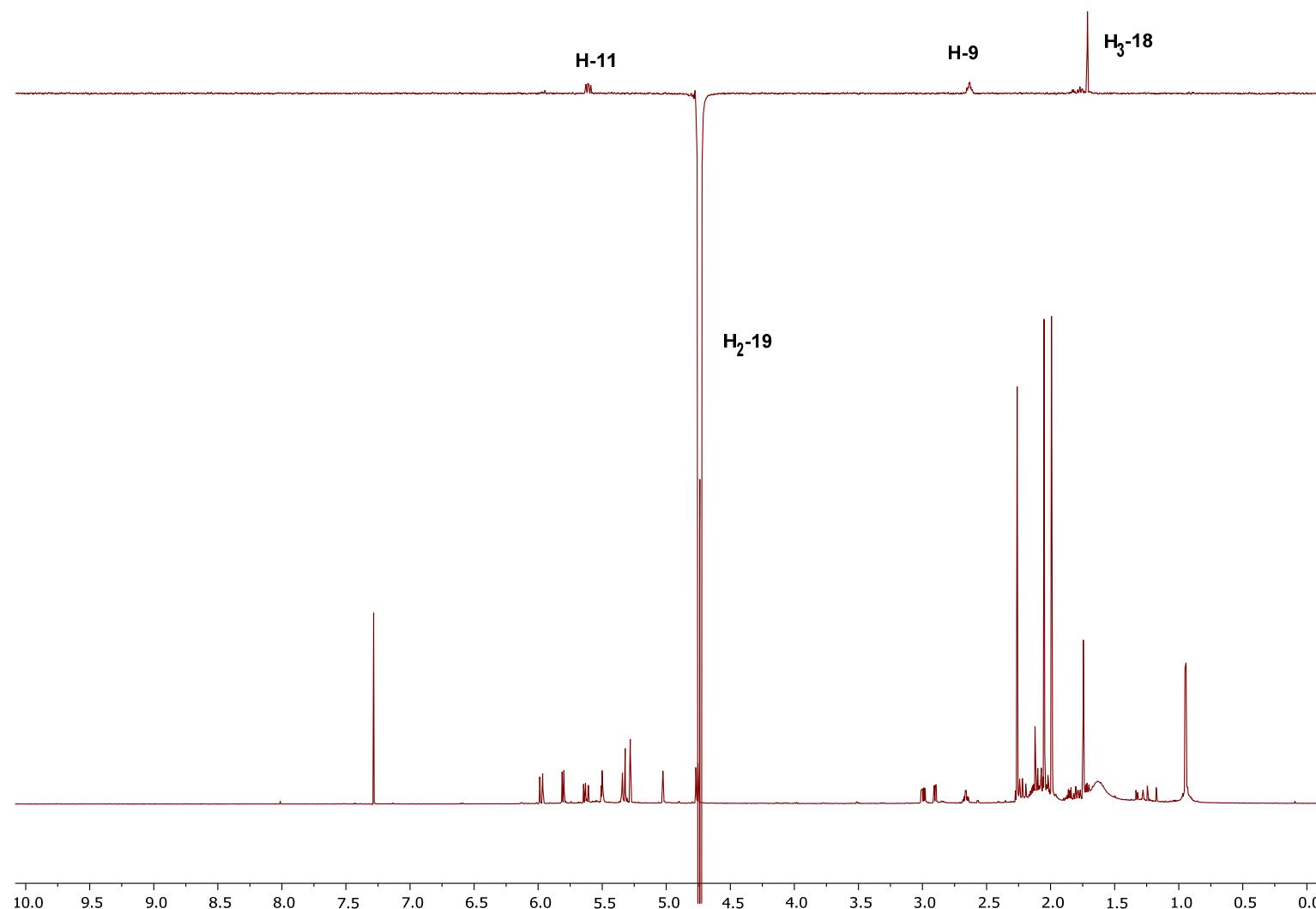


Figure S42f. 1D NOESY spectrum of compound 7.

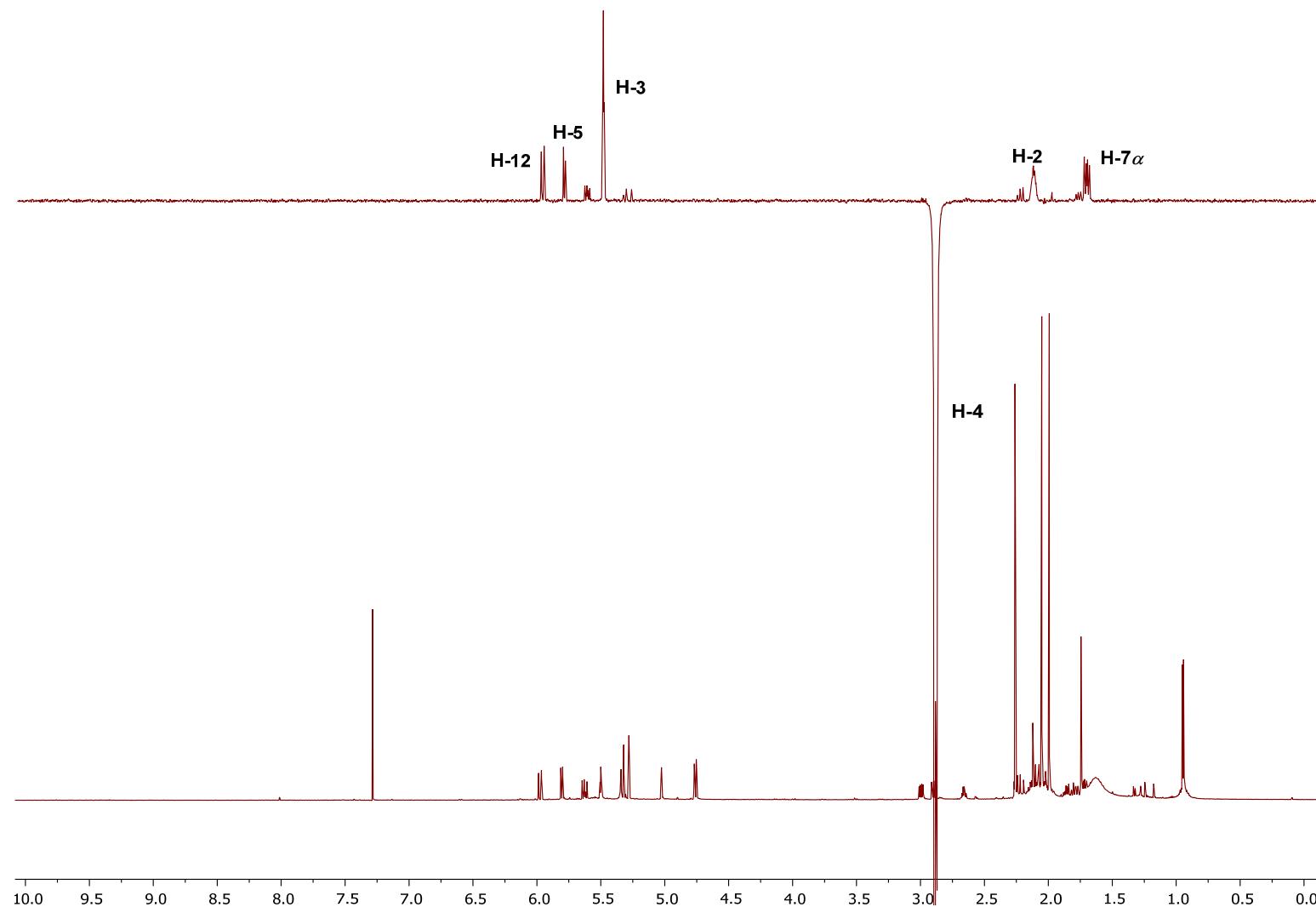


Figure S42g. 1D NOESY spectrum of compound 7.

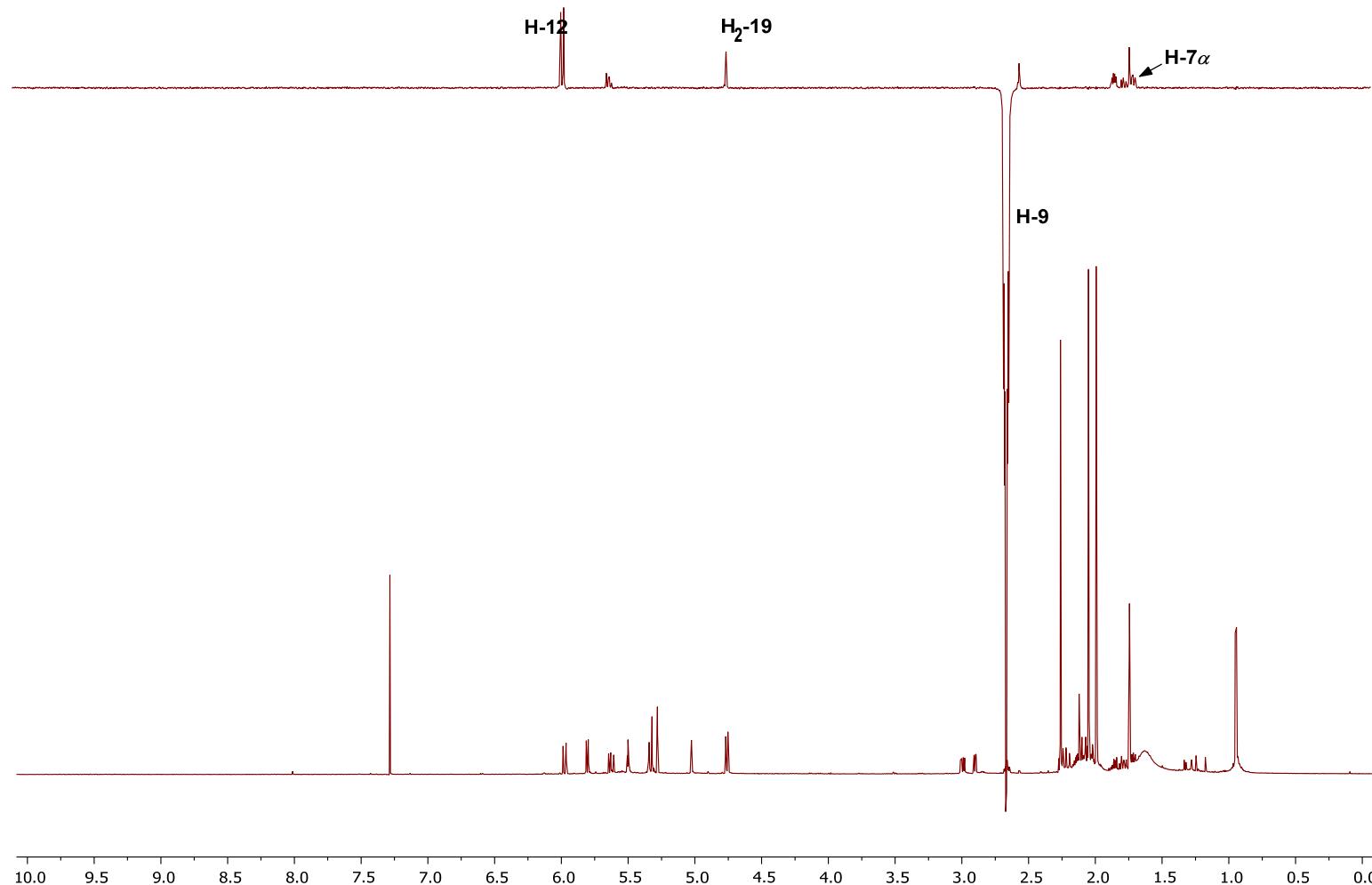


Figure S42h. 1D NOESY spectrum of compound 7.

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Even Electron Ions

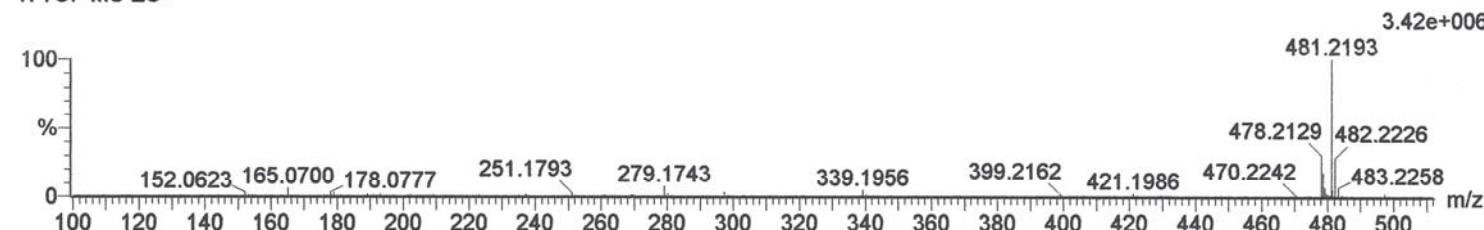
411 formula(e) evaluated with 7 results within limits (up to 10 closest results for each mass)

Elements Used:

C: 1-500 H: 0-1000 O: 0-200 Na: 0-1 K: 0-1

242_946-980_sTREP-EB12-MSe2pos 109 (2.026)

1: TOF MS ES+



Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
481.2193	481.2202	-0.9	-1.9	9.5	1638.8	0.147	86.29	C26 H34 O7 Na
	481.2204	-1.1	-2.3	4.5	1654.8	16.067	0.00	C23 H38 O8 K
	481.2180	1.3	2.7	1.5	1654.5	15.837	0.00	C21 H39 O8 Na K
	481.2168	2.5	5.2	21.5	1646.3	7.586	0.05	C35 H29 O2
	481.2226	-3.3	-6.9	12.5	1640.7	2.003	13.50	C28 H33 O7
	481.2145	4.8	10.0	13.5	1655.5	16.811	0.00	C30 H34 O3 K
	481.2143	5.0	10.4	18.5	1645.2	6.464	0.16	C33 H30 O2 Na

Figure S43. HRMS of compound 7.

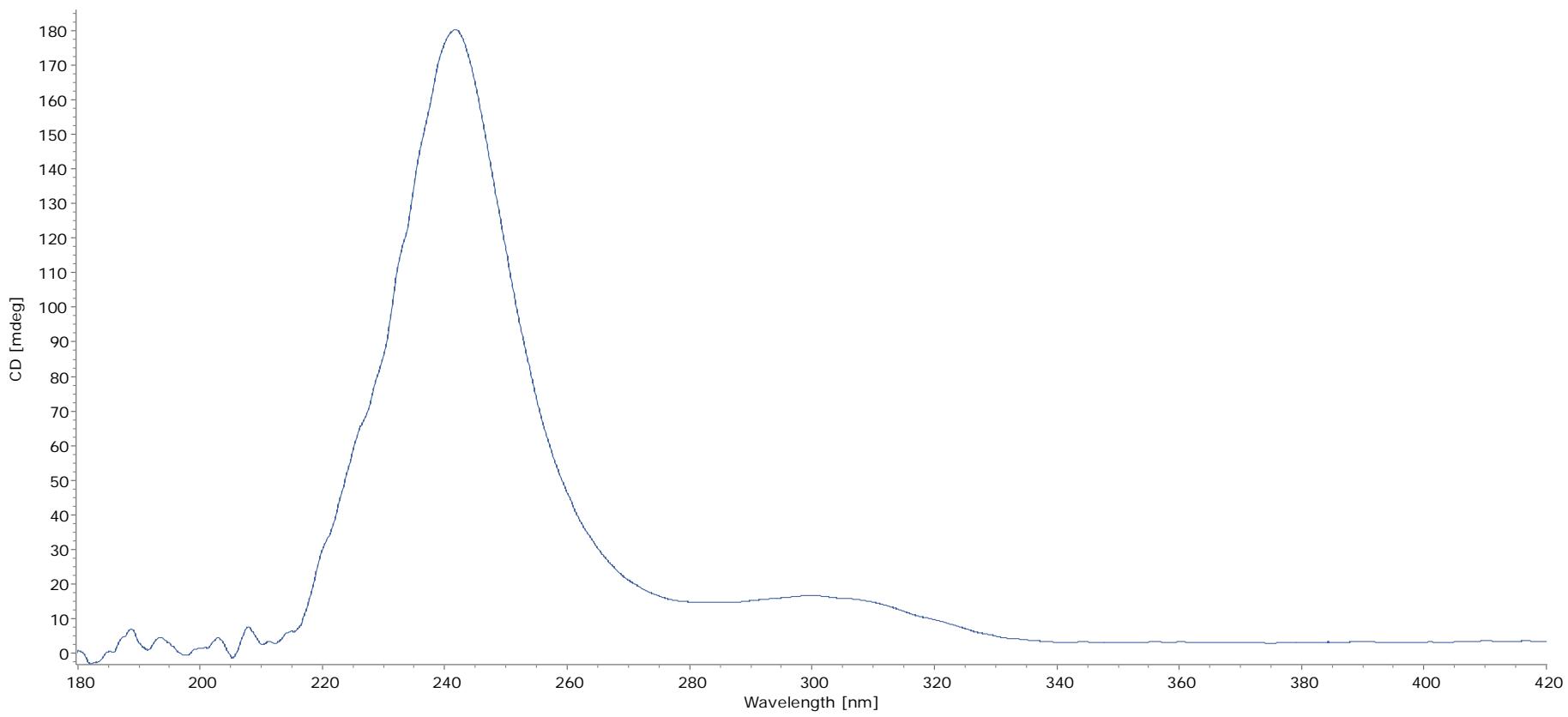


Figure S44. ECD of compound 7.