New eremophilane derivatives produced by the marine-derived fungus *Emericellopsis maritima* BC17 in liquid culture media

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Table of contents

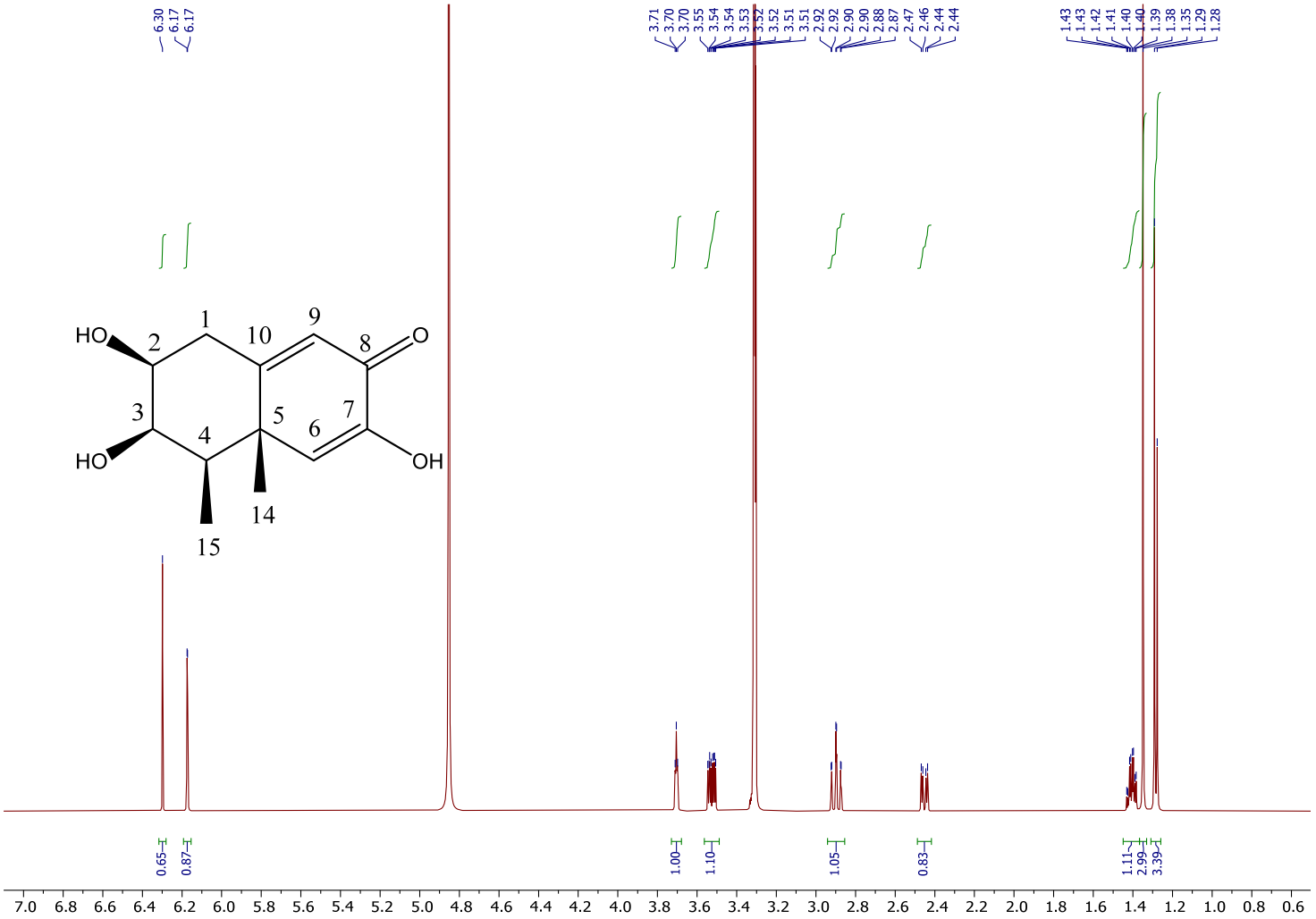


Figure S1. ^{1}H NMR spectrum (500 MHz, CD_3OD) of compound **1**.

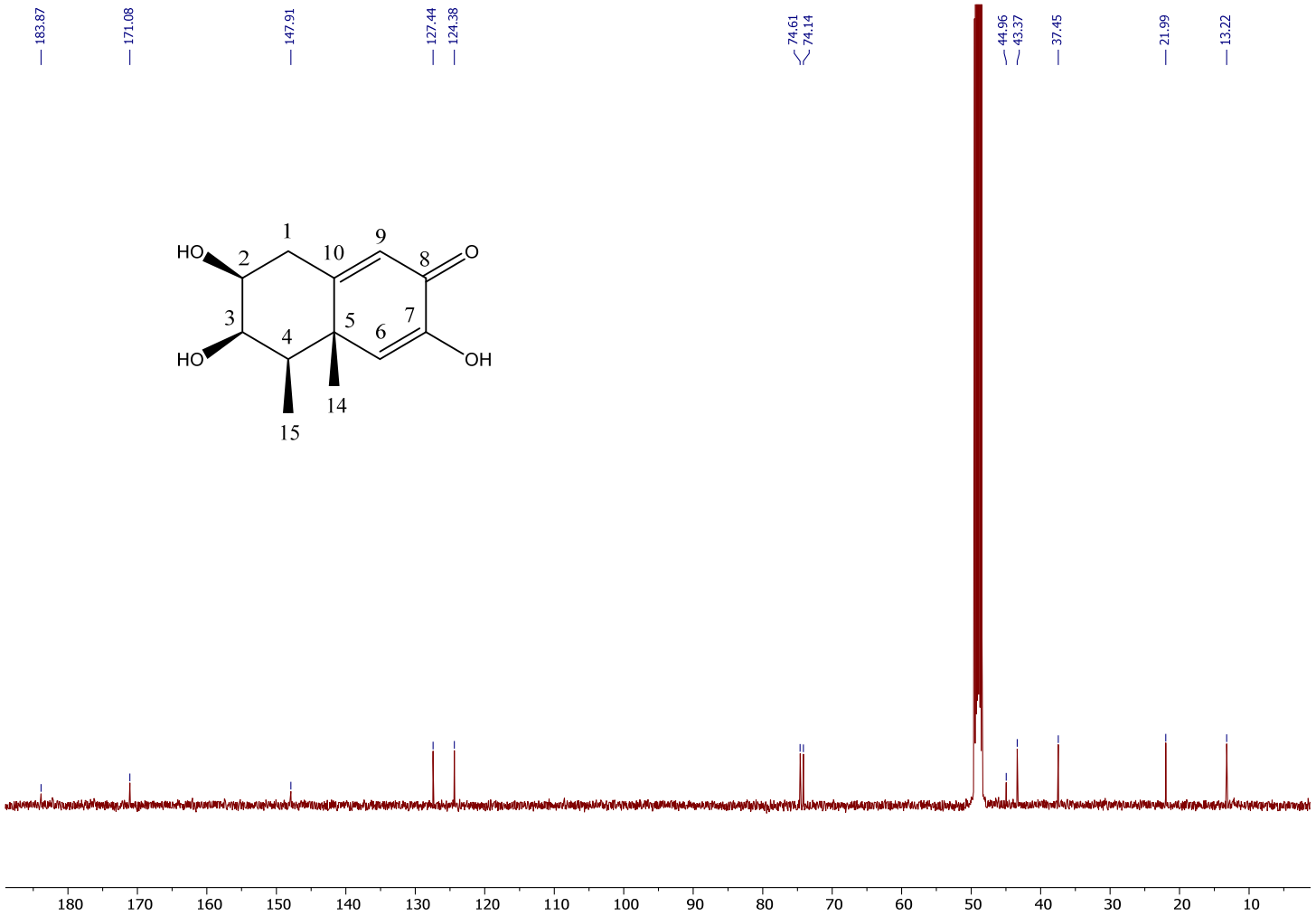


Figure S2. ^{13}C NMR spectrum (125 MHz, CD_3OD) of compound **1**.

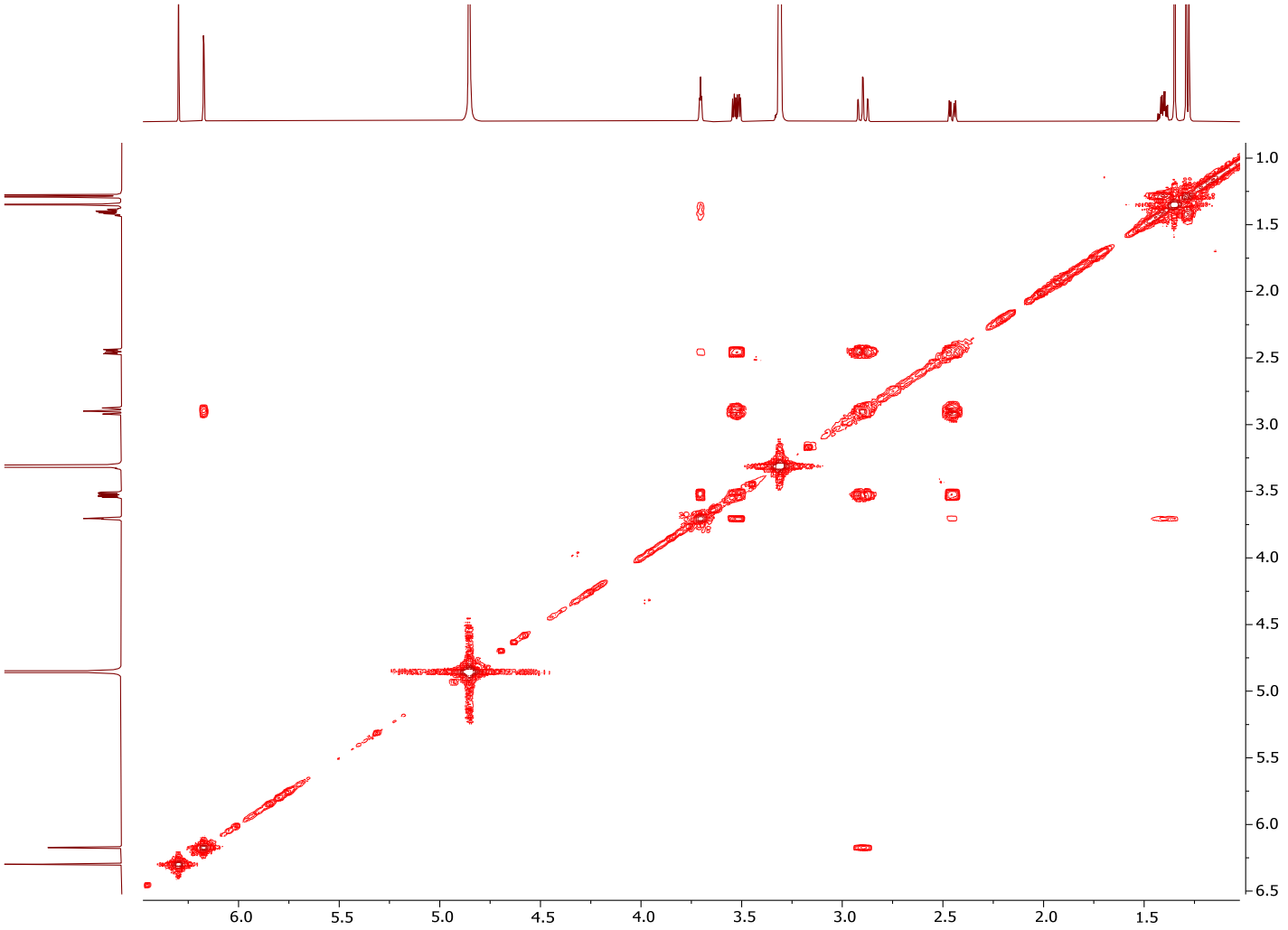


Figure S3. gCOSY spectrum of compound **1**.

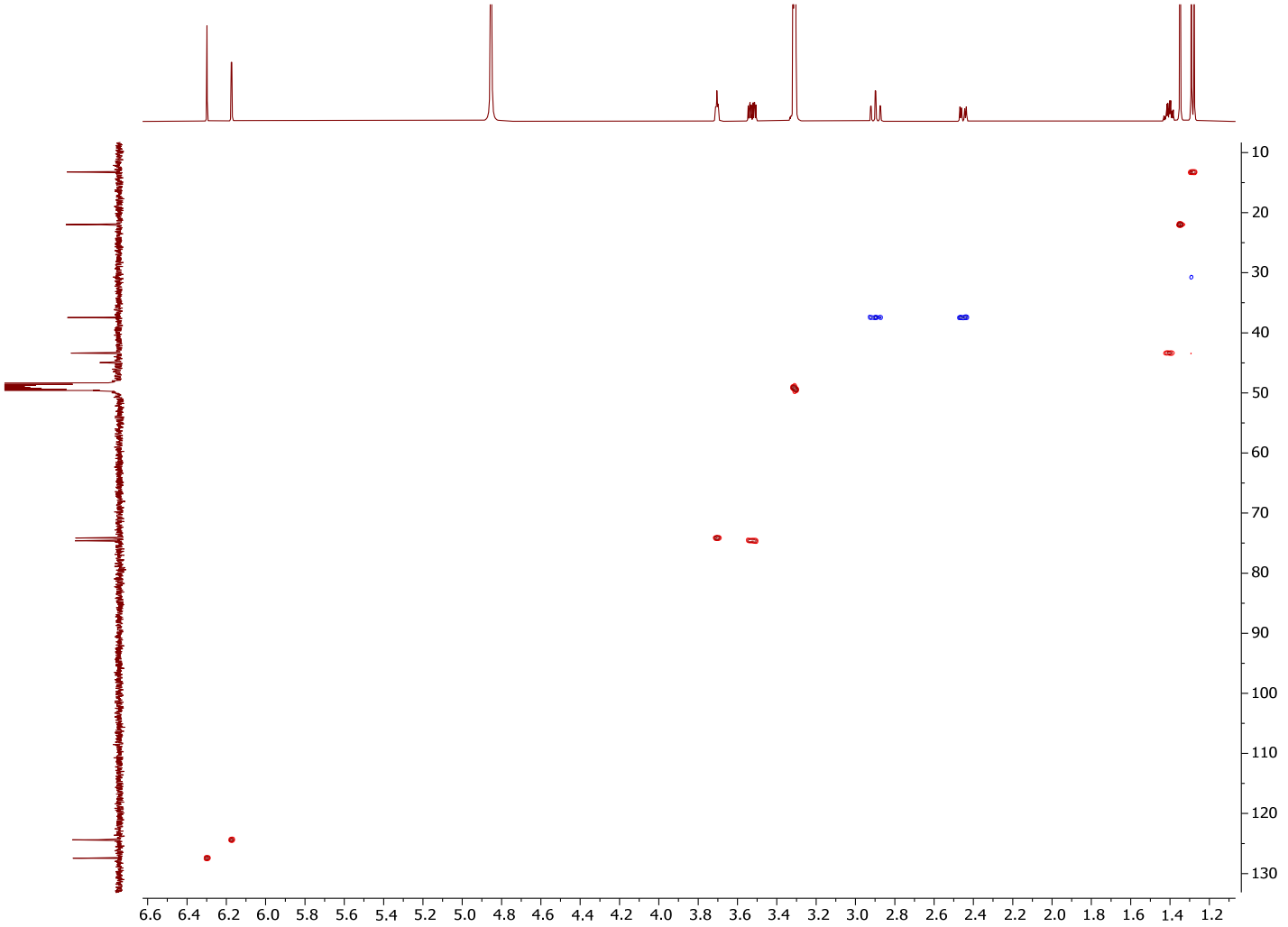


Figure S4. gHSQC spectrum of compound **1**.

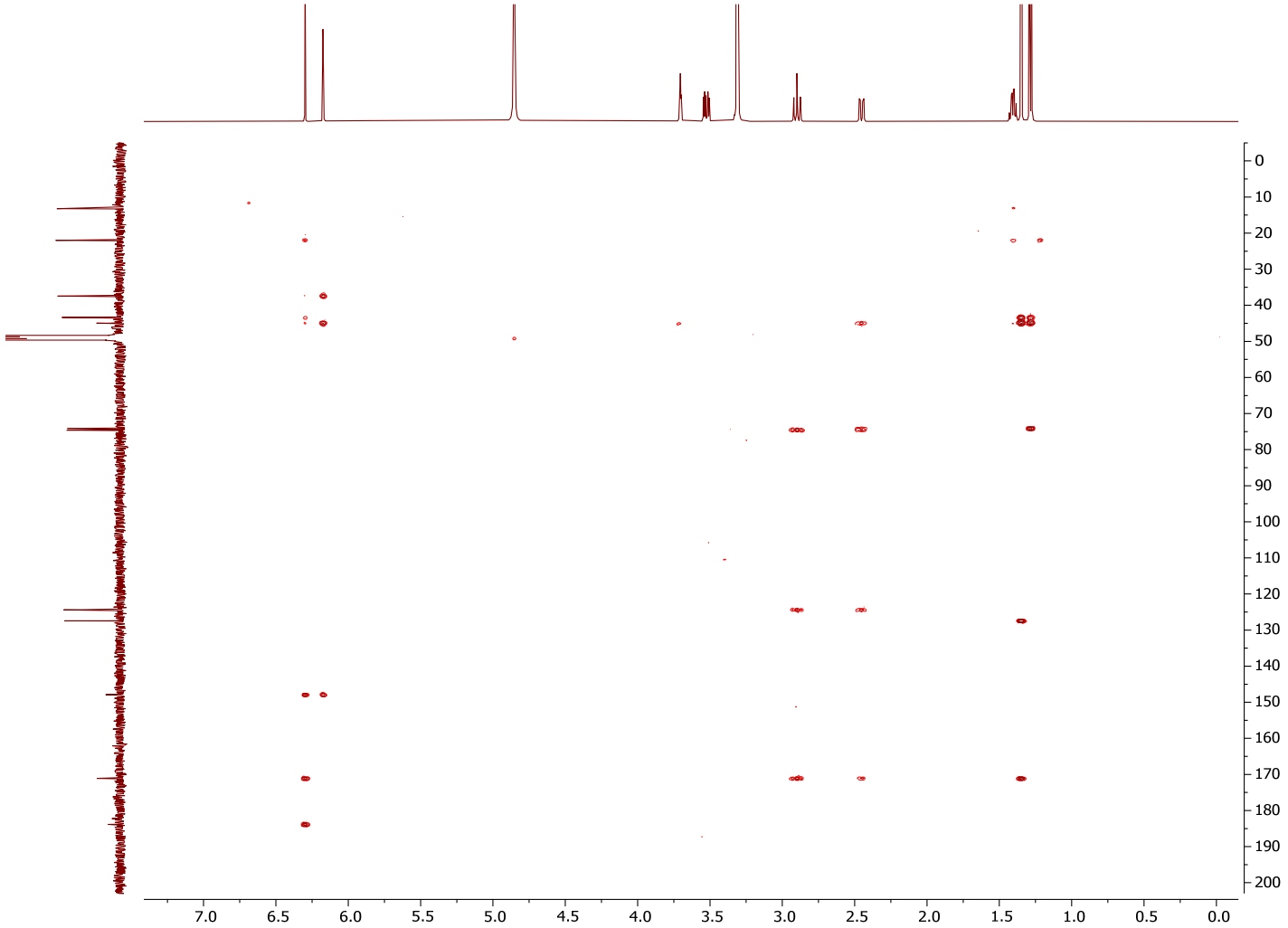


Figure S5. gHMBC spectrum of compound **1**.

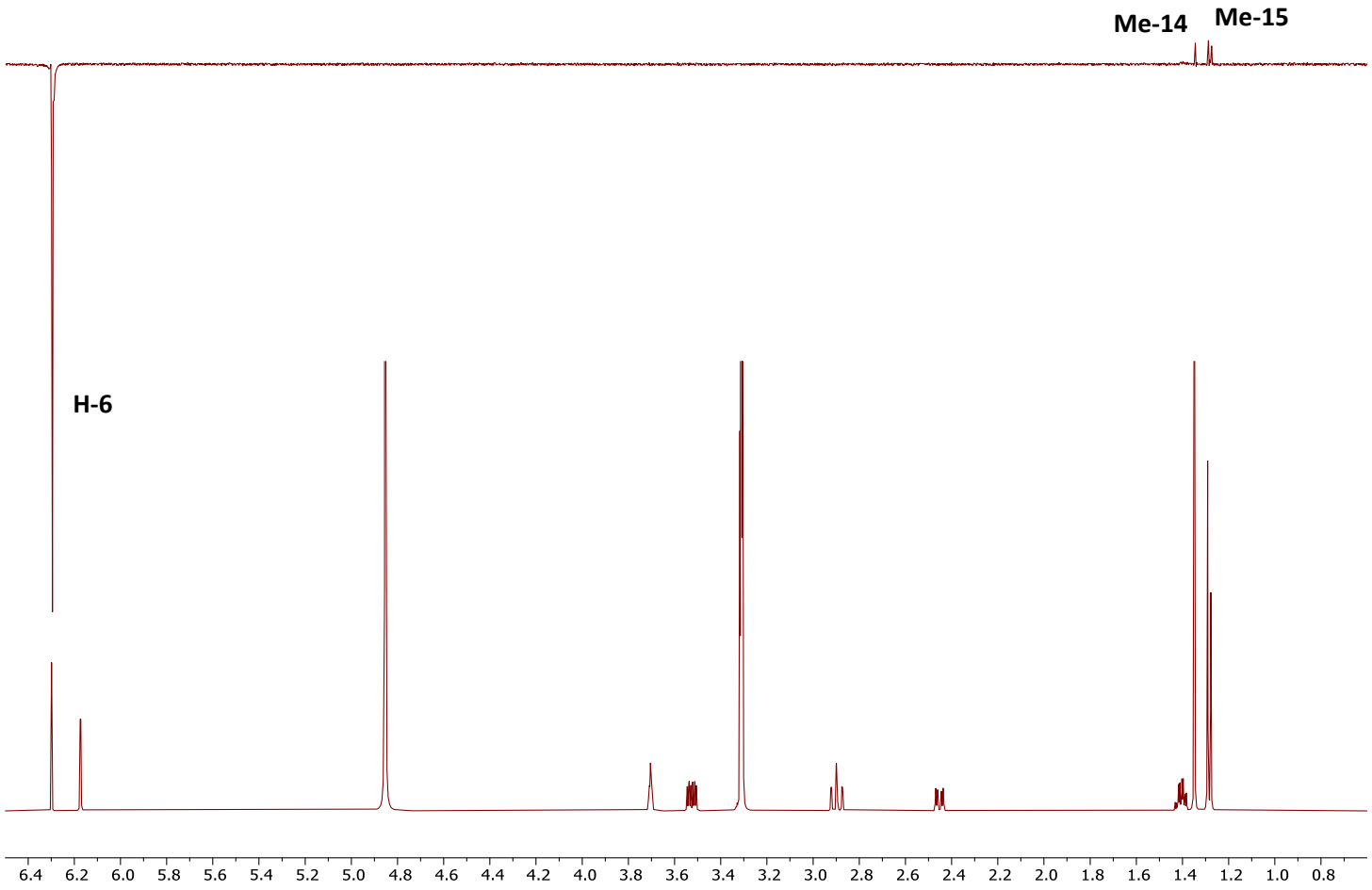


Figure S6a. 1D NOESY spectrum of compound **1**.

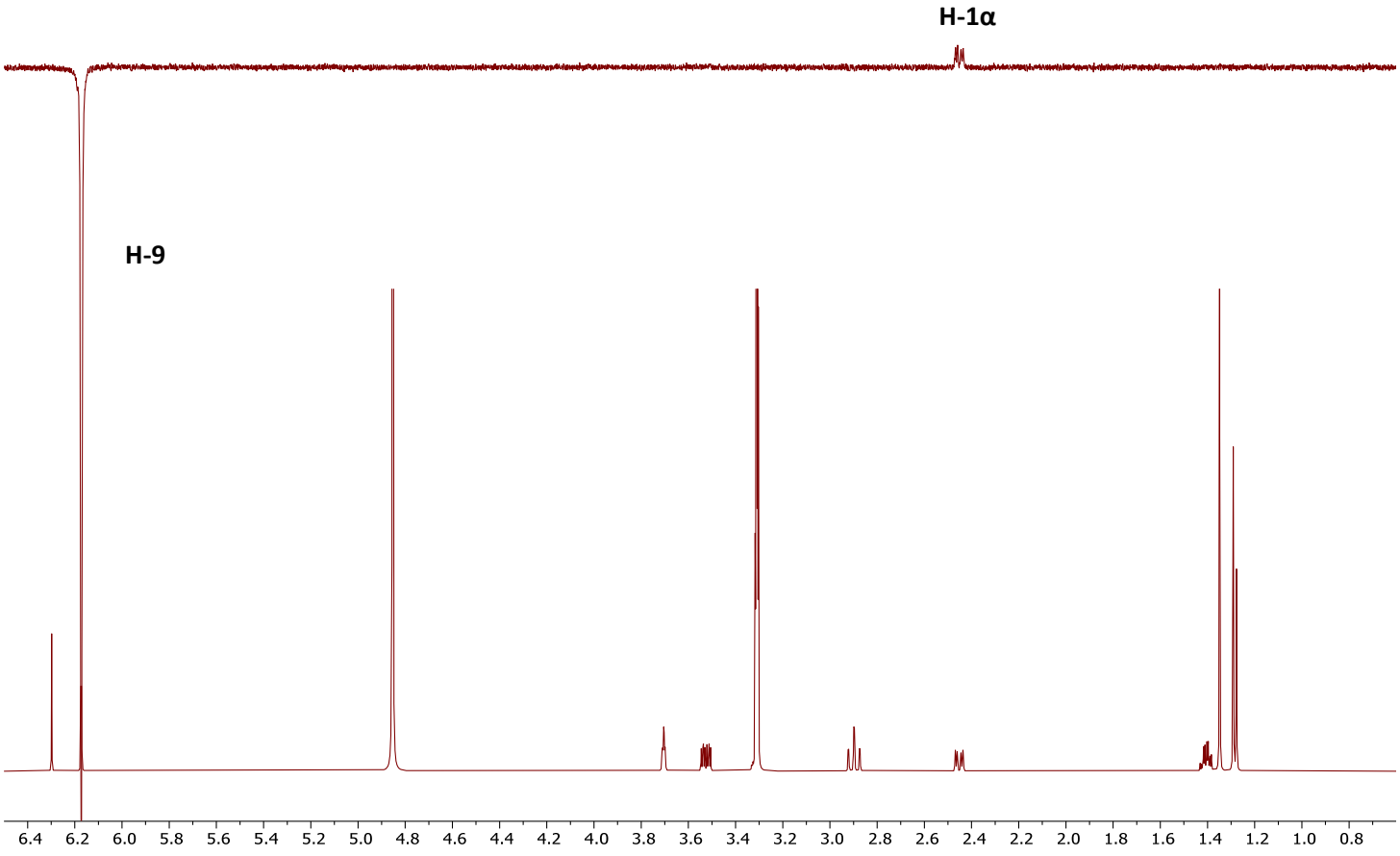


Figure S6b. 1D NOESY spectrum of compound **1**.

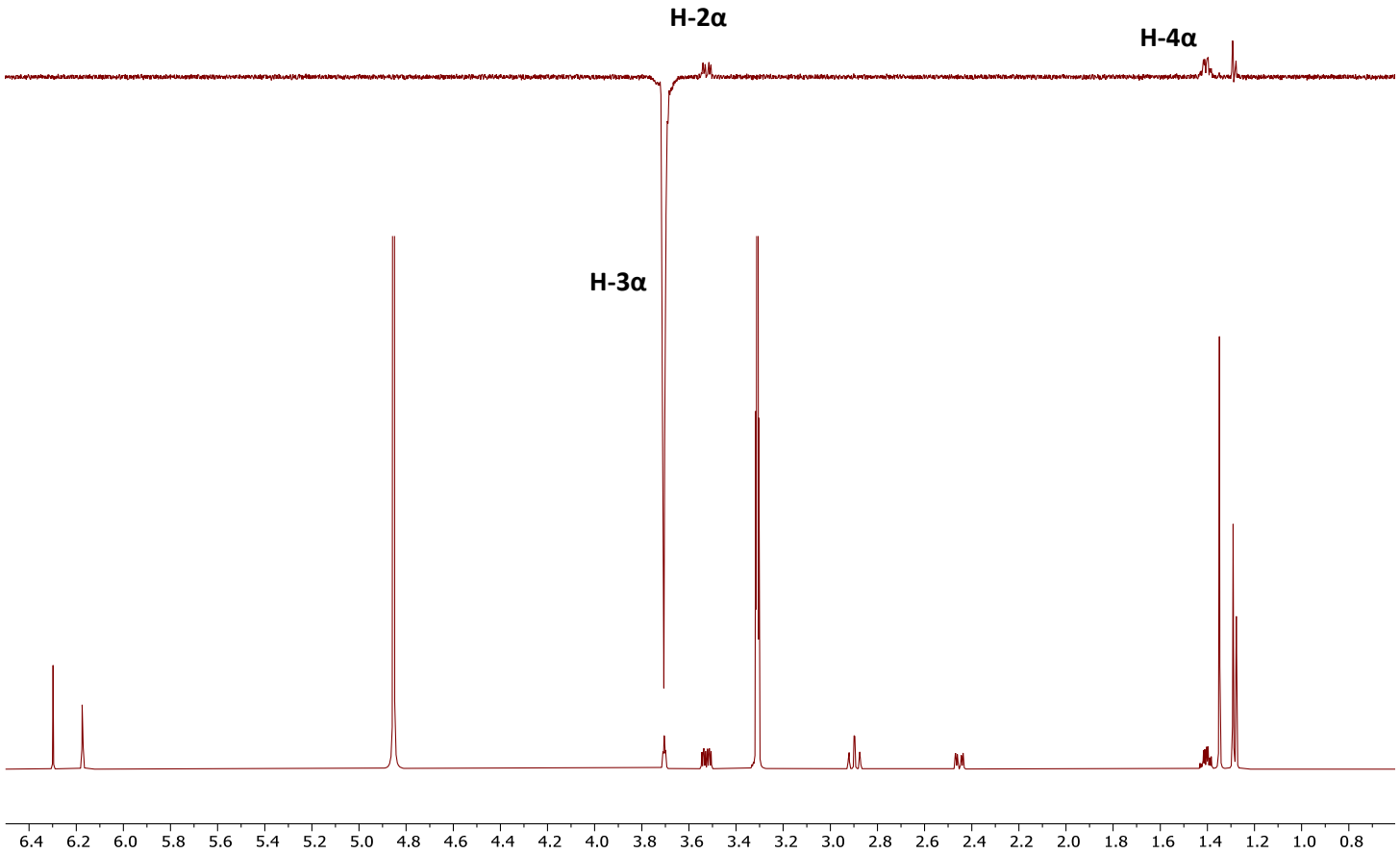


Figure S6c. 1D NOESY spectrum of compound **1**.

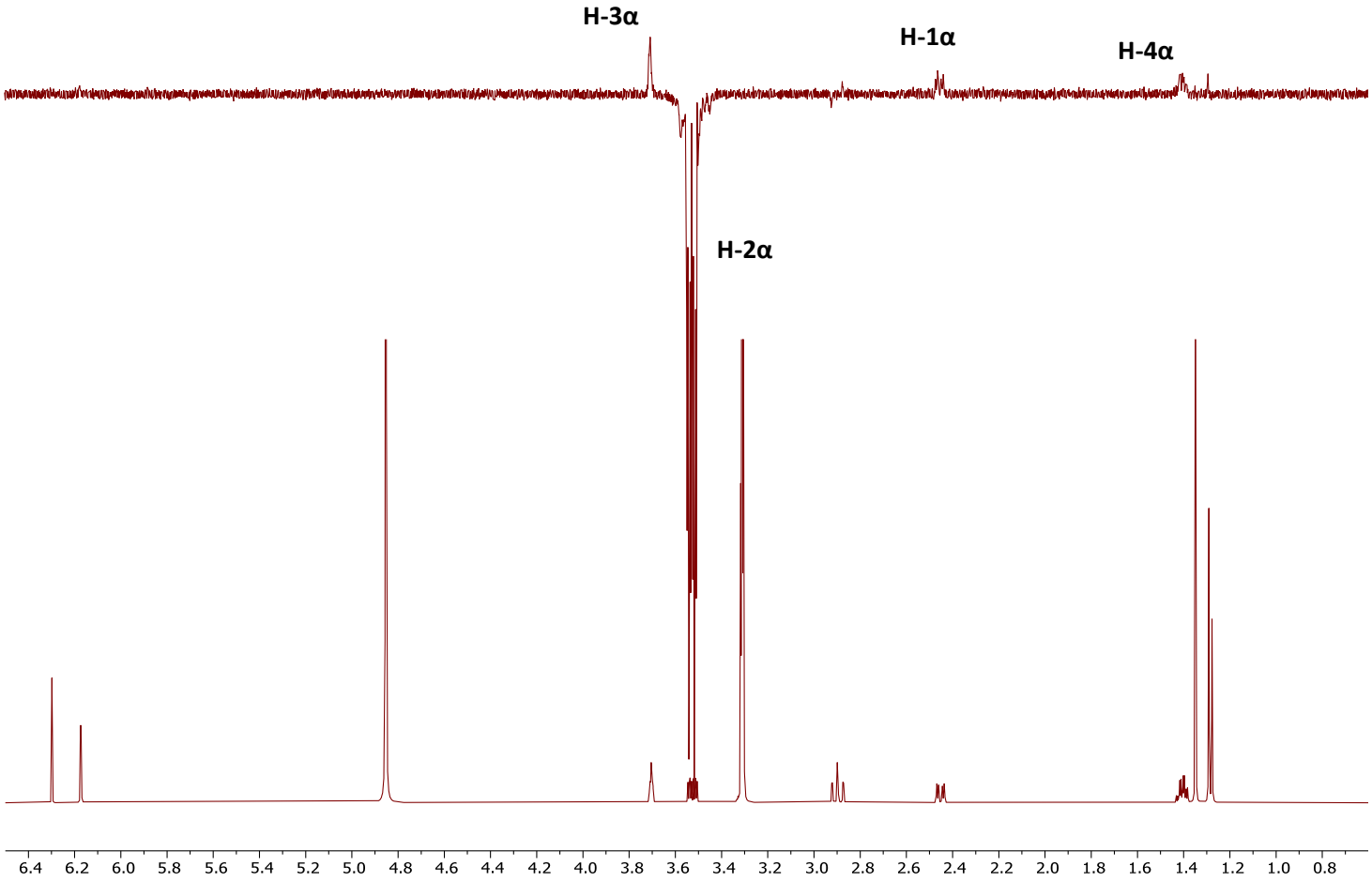


Figure S6d. 1D NOESY spectrum of compound **1**.

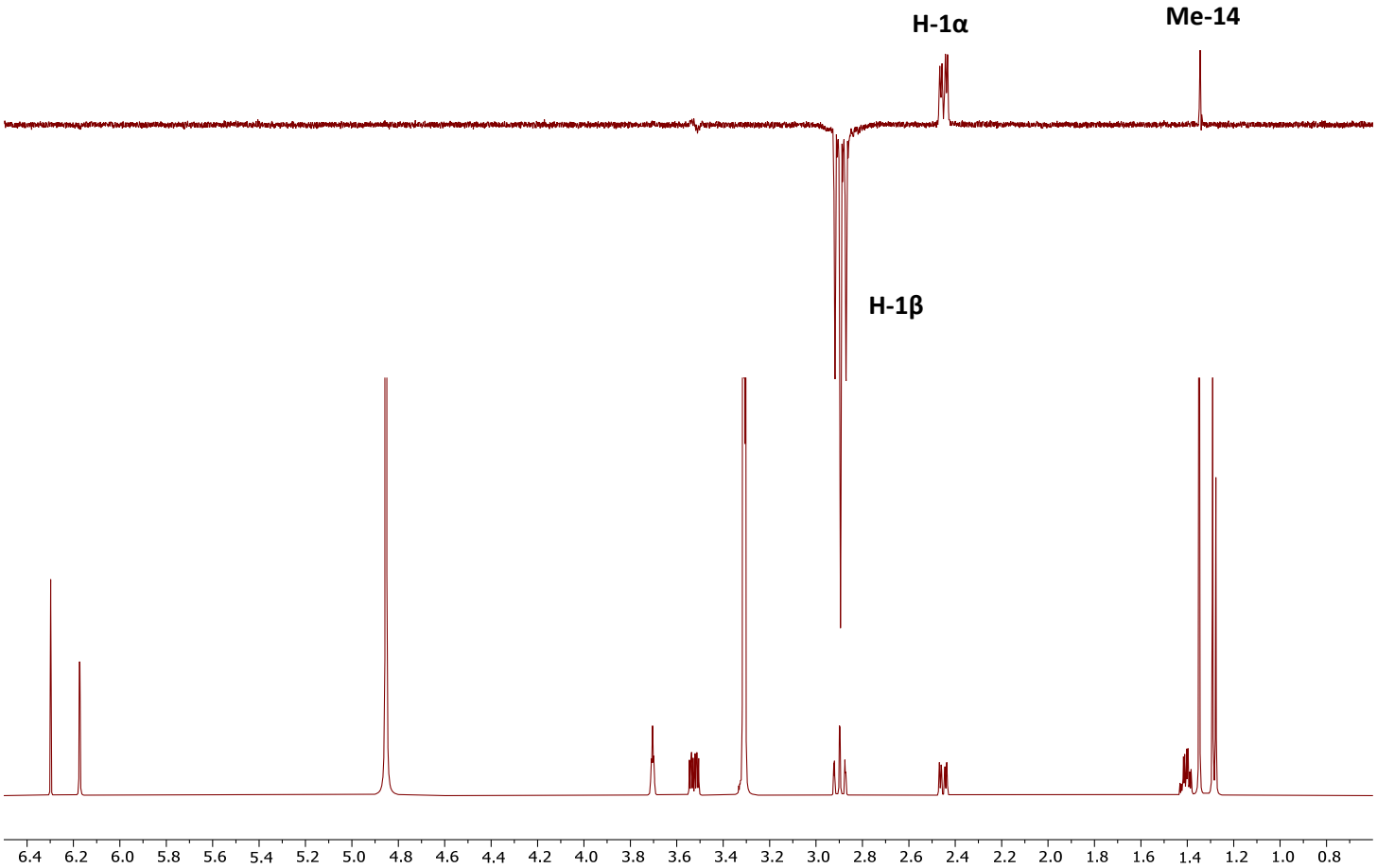


Figure S6e. 1D NOESY spectrum of compound **1**.

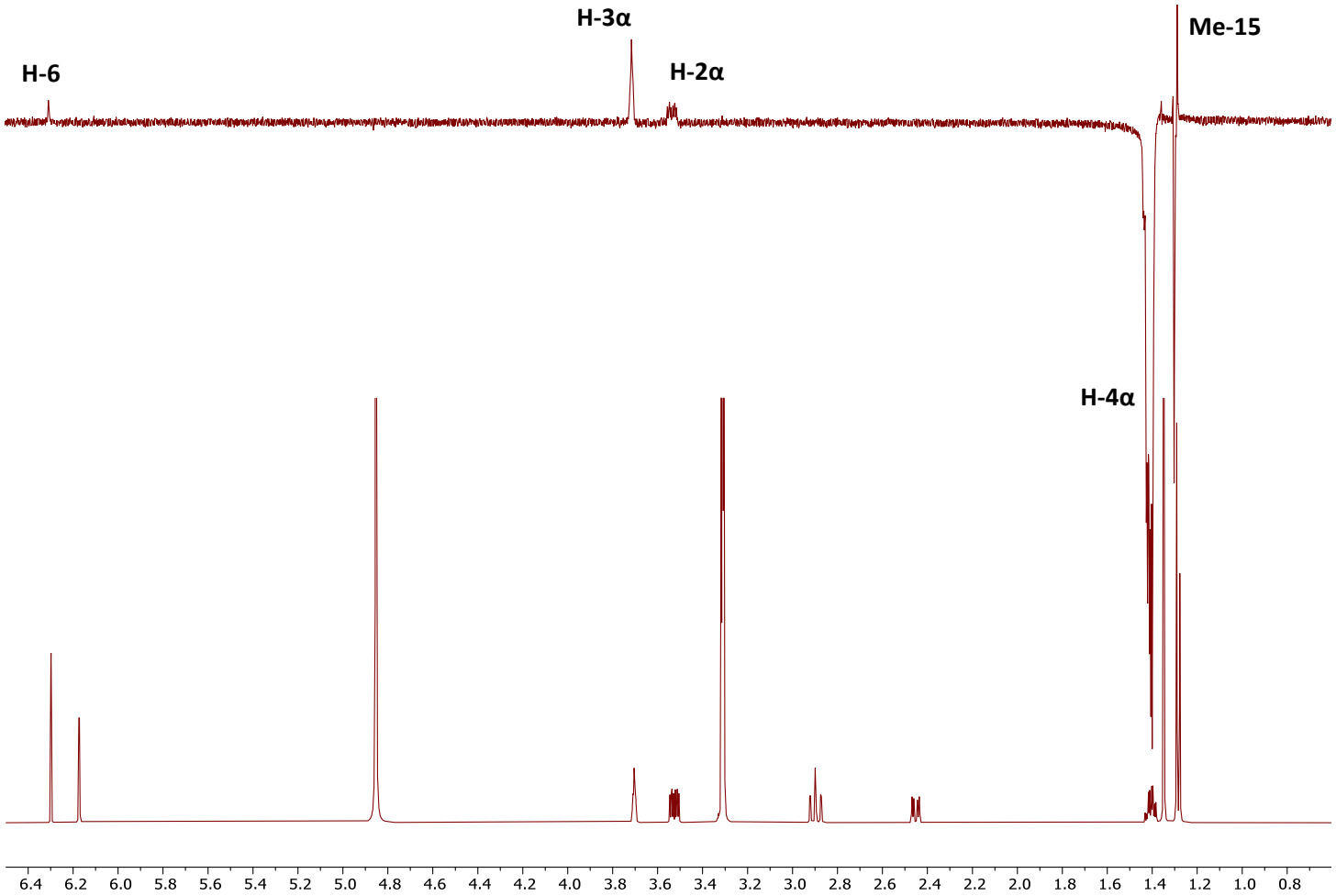


Figure S6f. 1D NOESY spectrum of compound **1**.

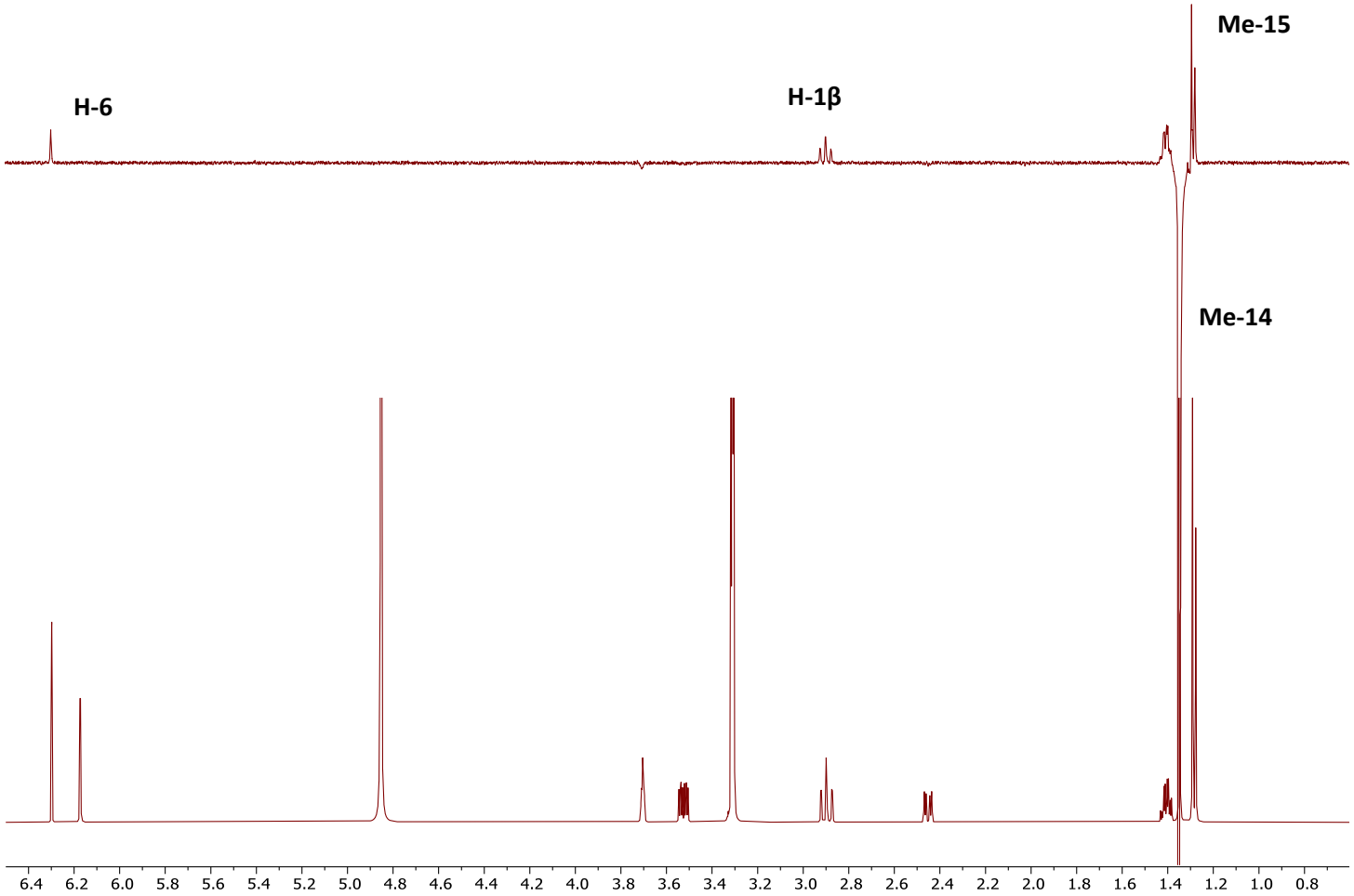
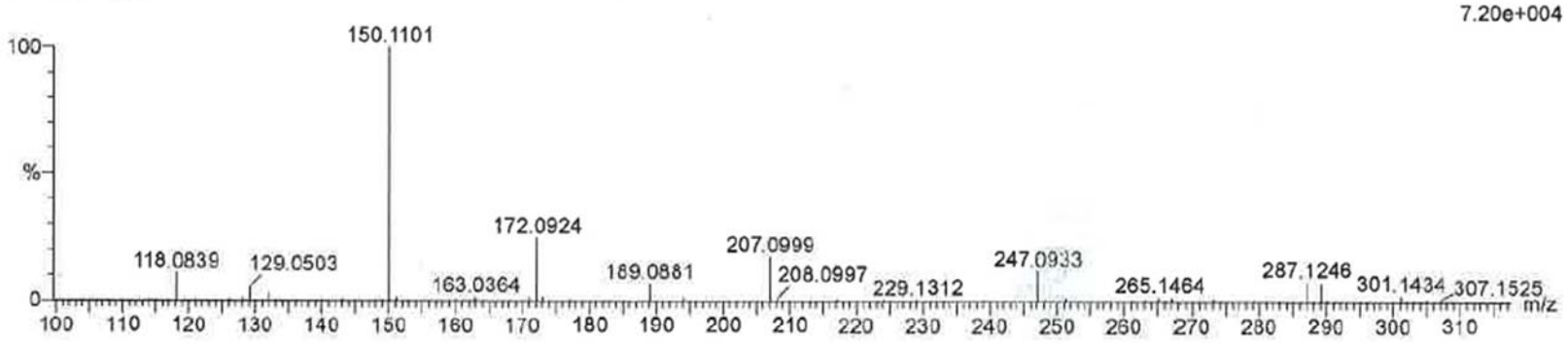


Figure S6g. 1D NOESY difference spectrum of compound **1**.

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
76 formula(e) evaluated with 2 results within limits (up to 50 best isotopic matches for each mass)
Elements Used:
C: 0-500 H: 0-1000 O: 0-200 Na: 0-1
CDI-80-P8 26 (0.489)
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
247.0933	247.0946	-1.3	-5.3	4.5	323.1	0.316	72.93	C12 H16 O4 Na
	247.0970	-3.7	-15.0	7.5	324.1	1.307	27.07	C14 H15 O4

Figure S7. HRESIMS spectrum of compound **1**.

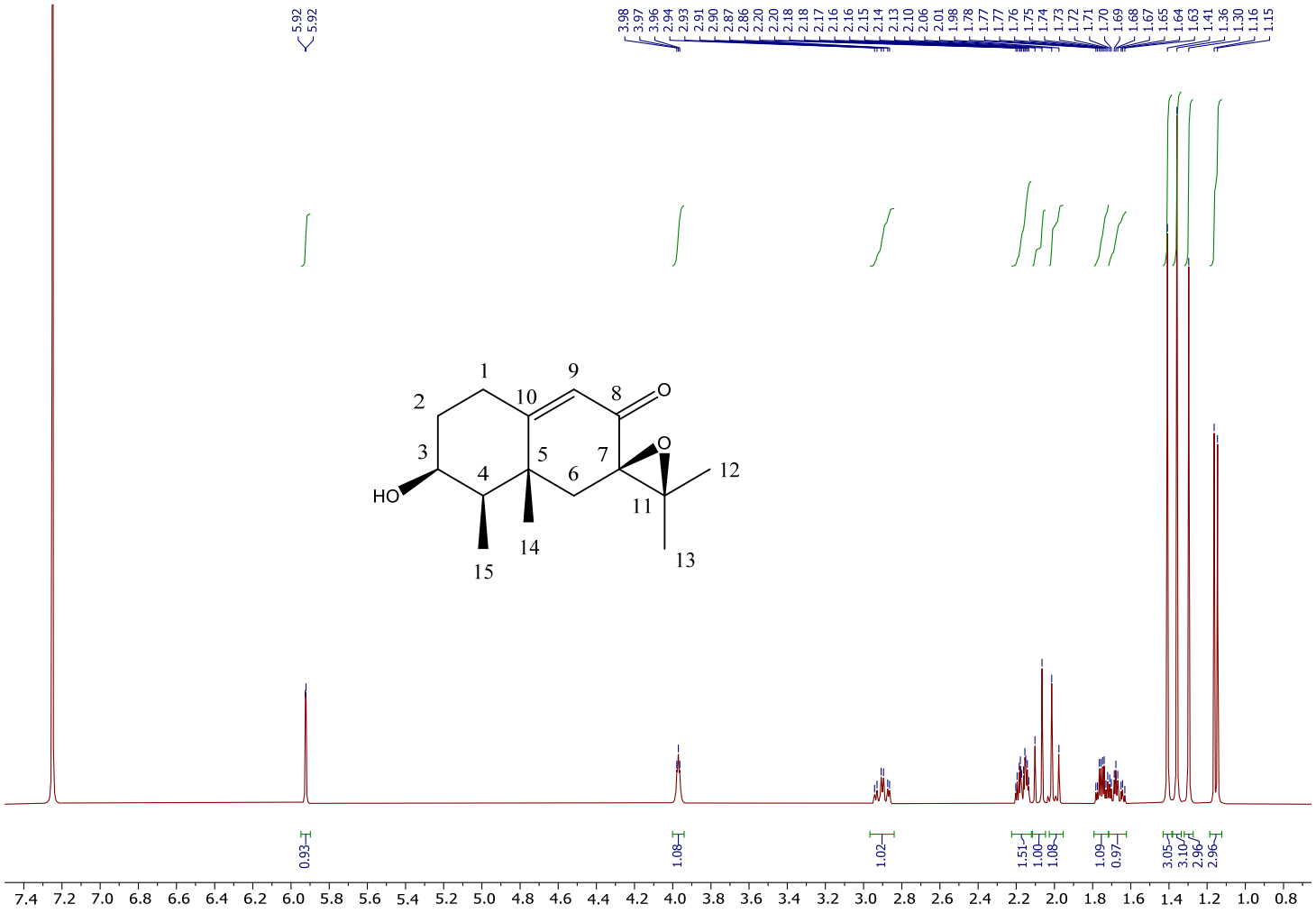


Figure S8. ^{1}H NMR spectrum (400 MHz, $CDCl_3$) of compound **2**.

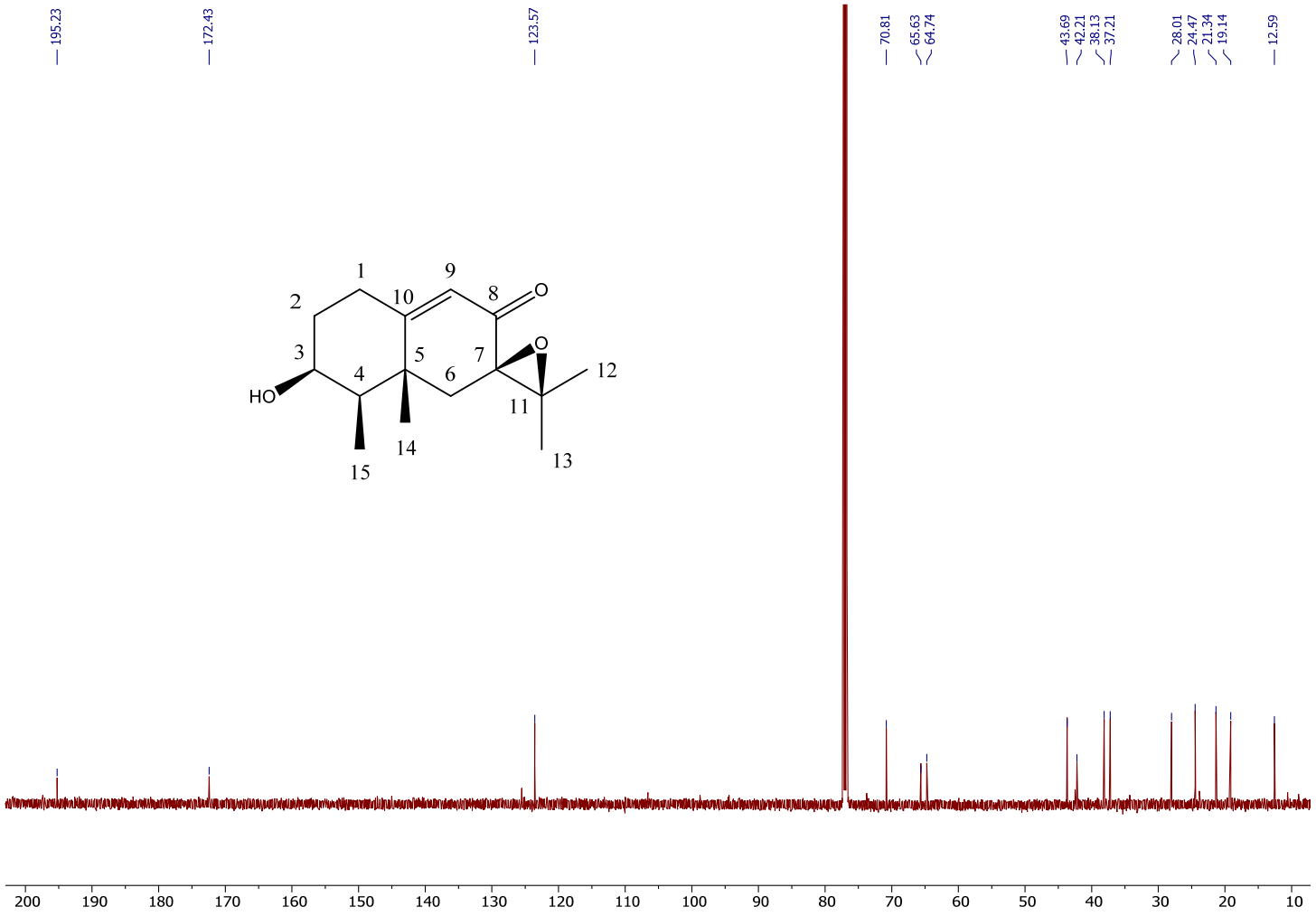


Figure S9. ^{13}C NMR spectrum (125 MHz, $CDCl_3$) of compound **2**.

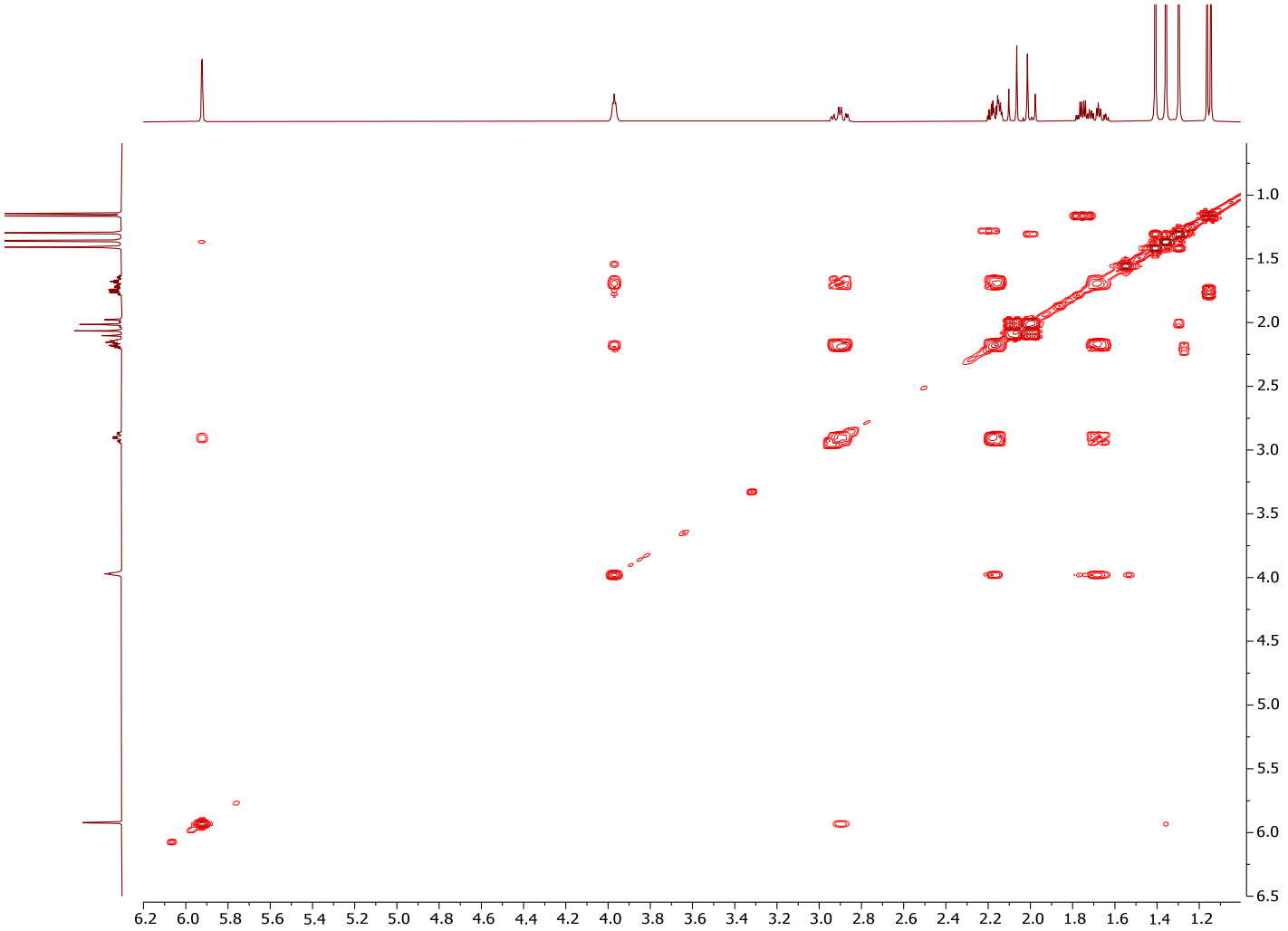


Figure S10. gCOSY spectrum of compound **2**.

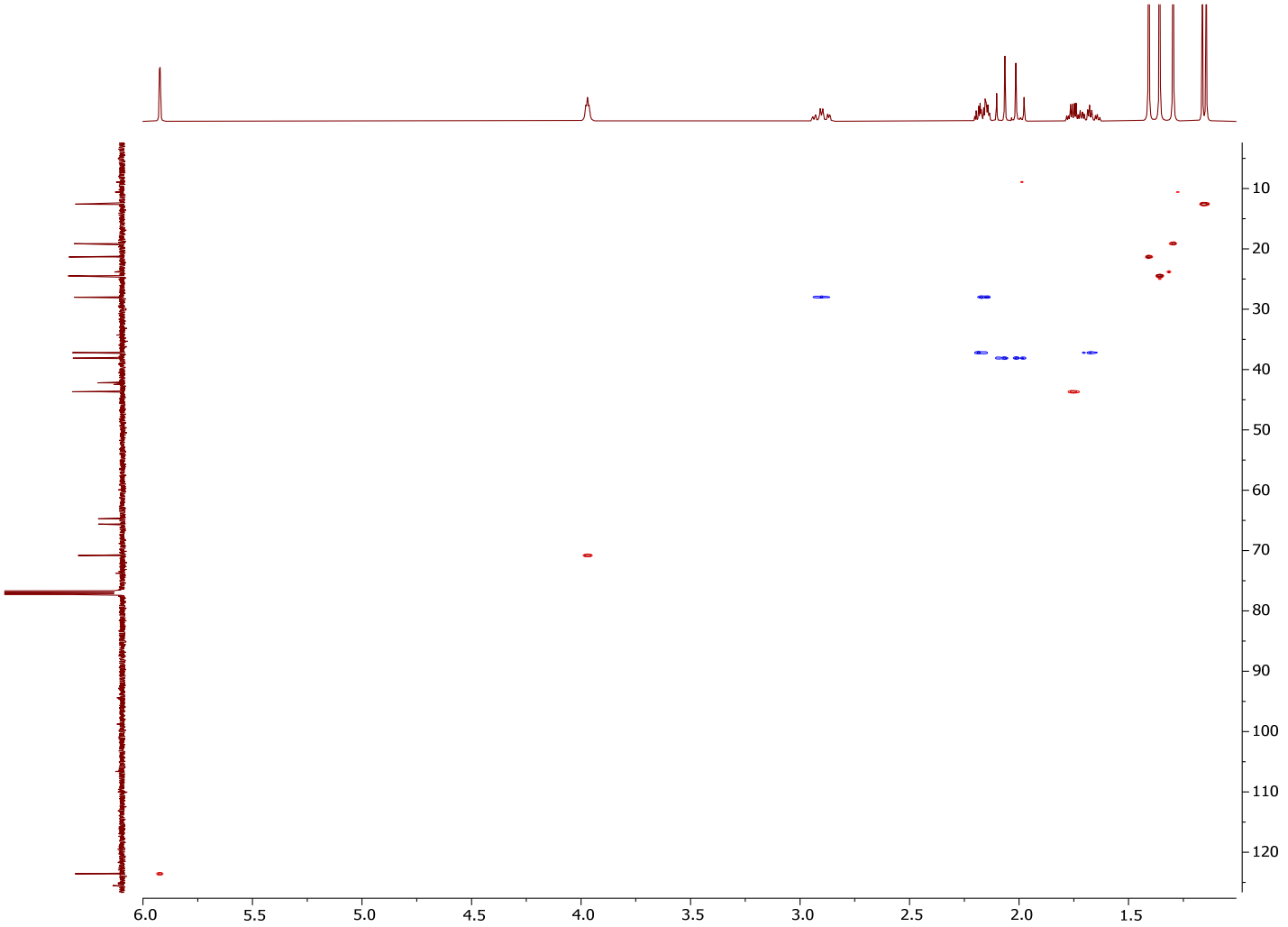


Figure S11. gHSQC spectrum of compound **2**.

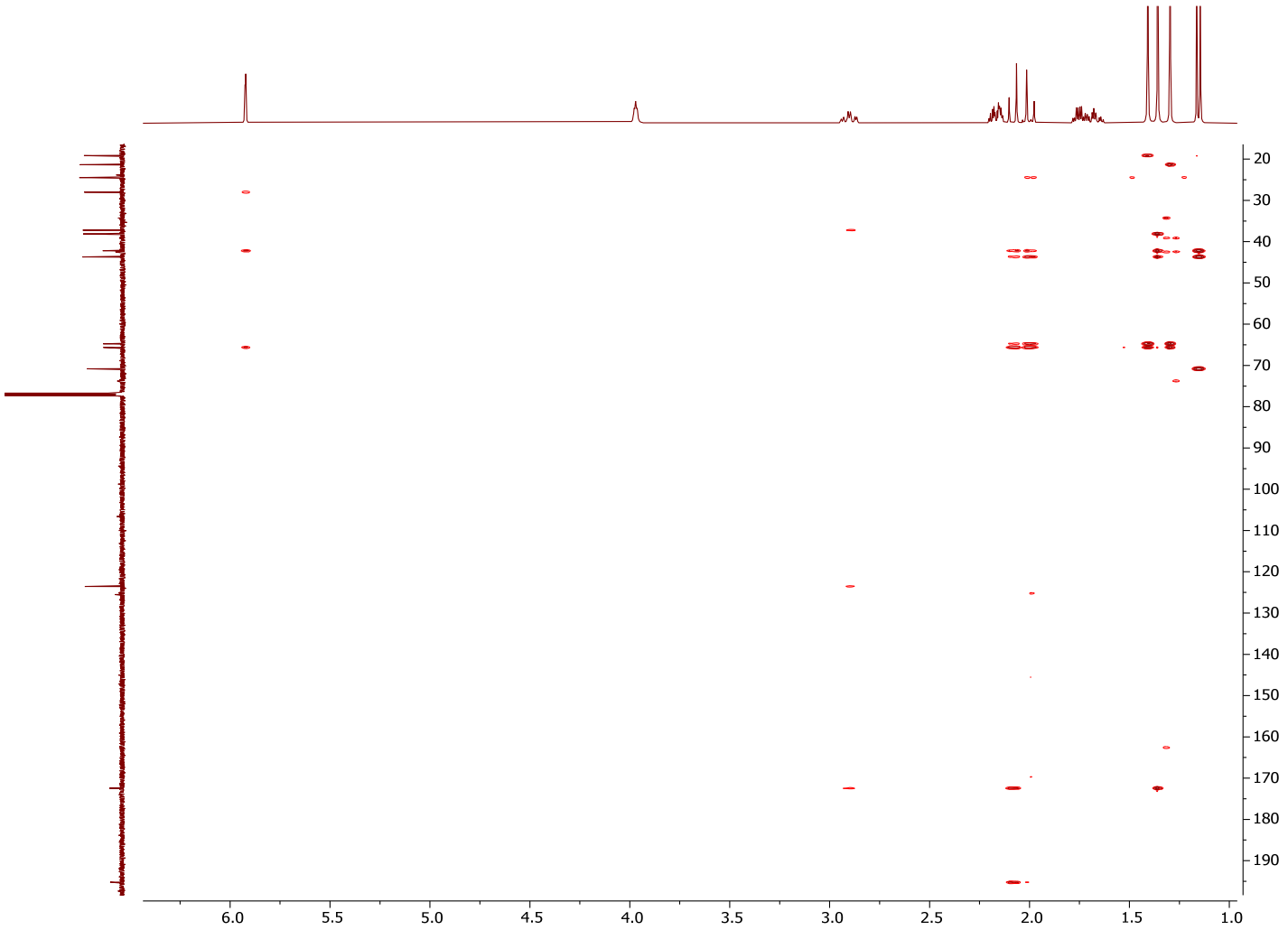


Figure S12. gHMBC spectrum of compound **2**.

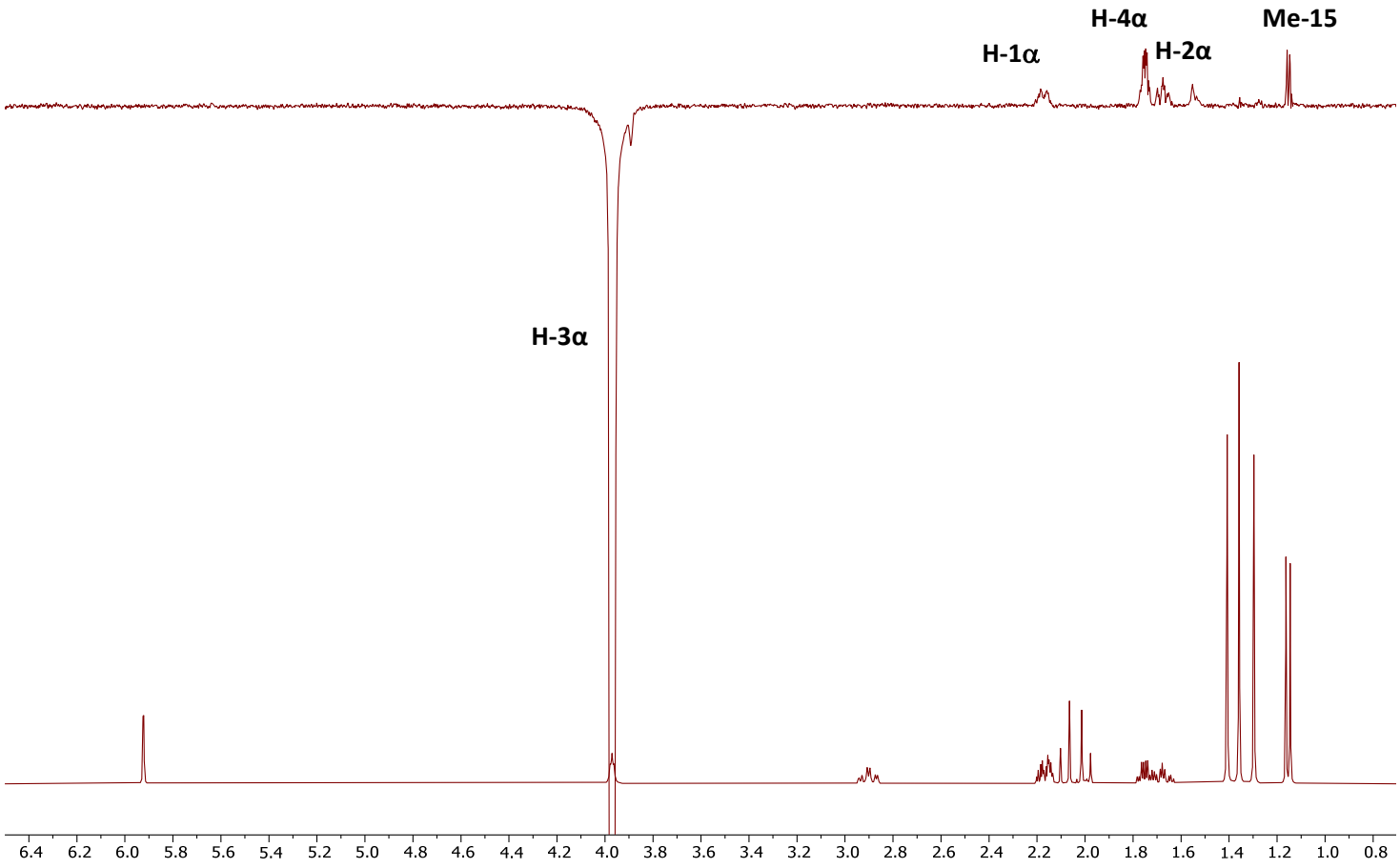


Figure S13a. 1D NOESY spectrum of compound **2**.

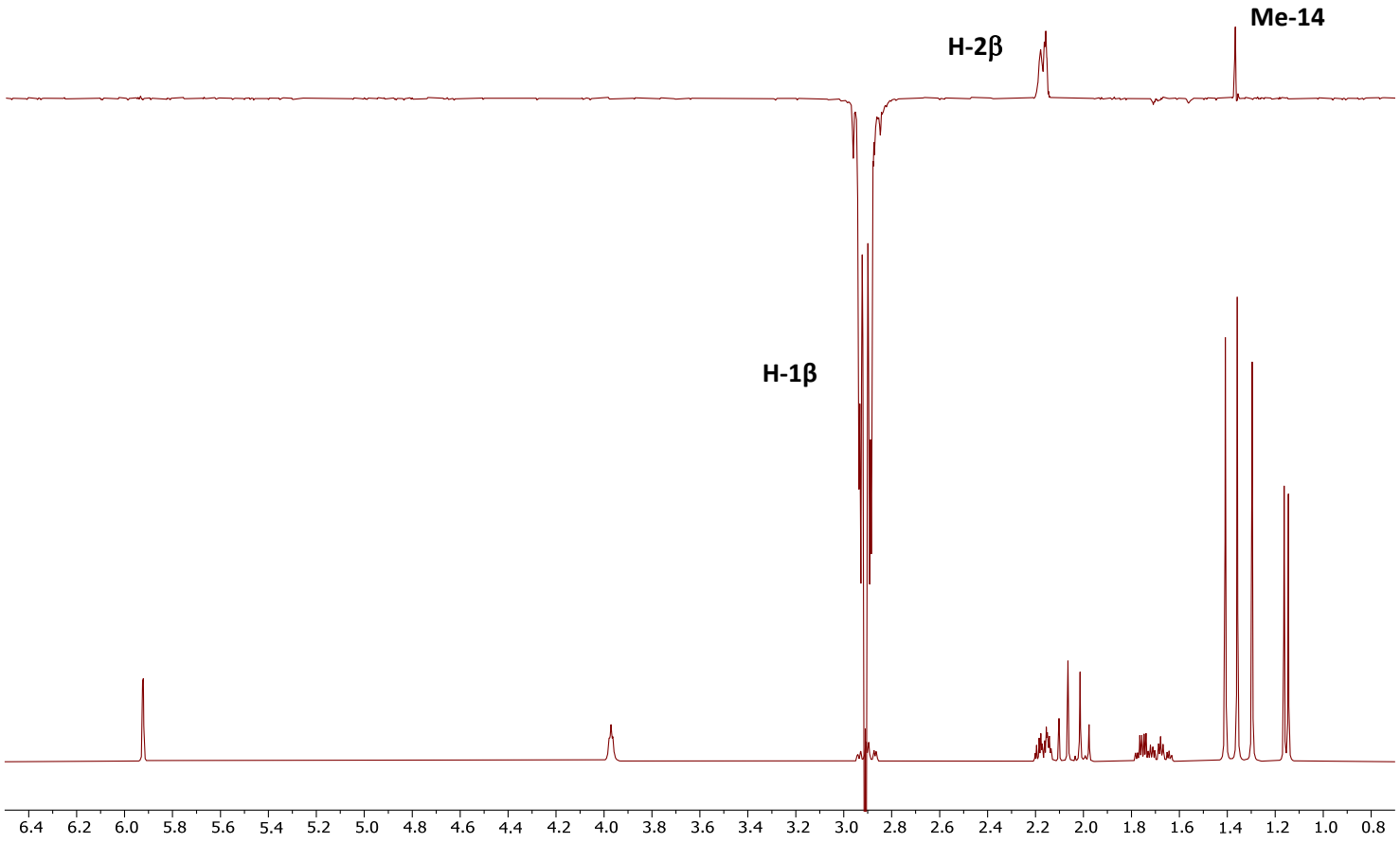


Figure S13b. 1D NOESY spectrum of compound **2**.

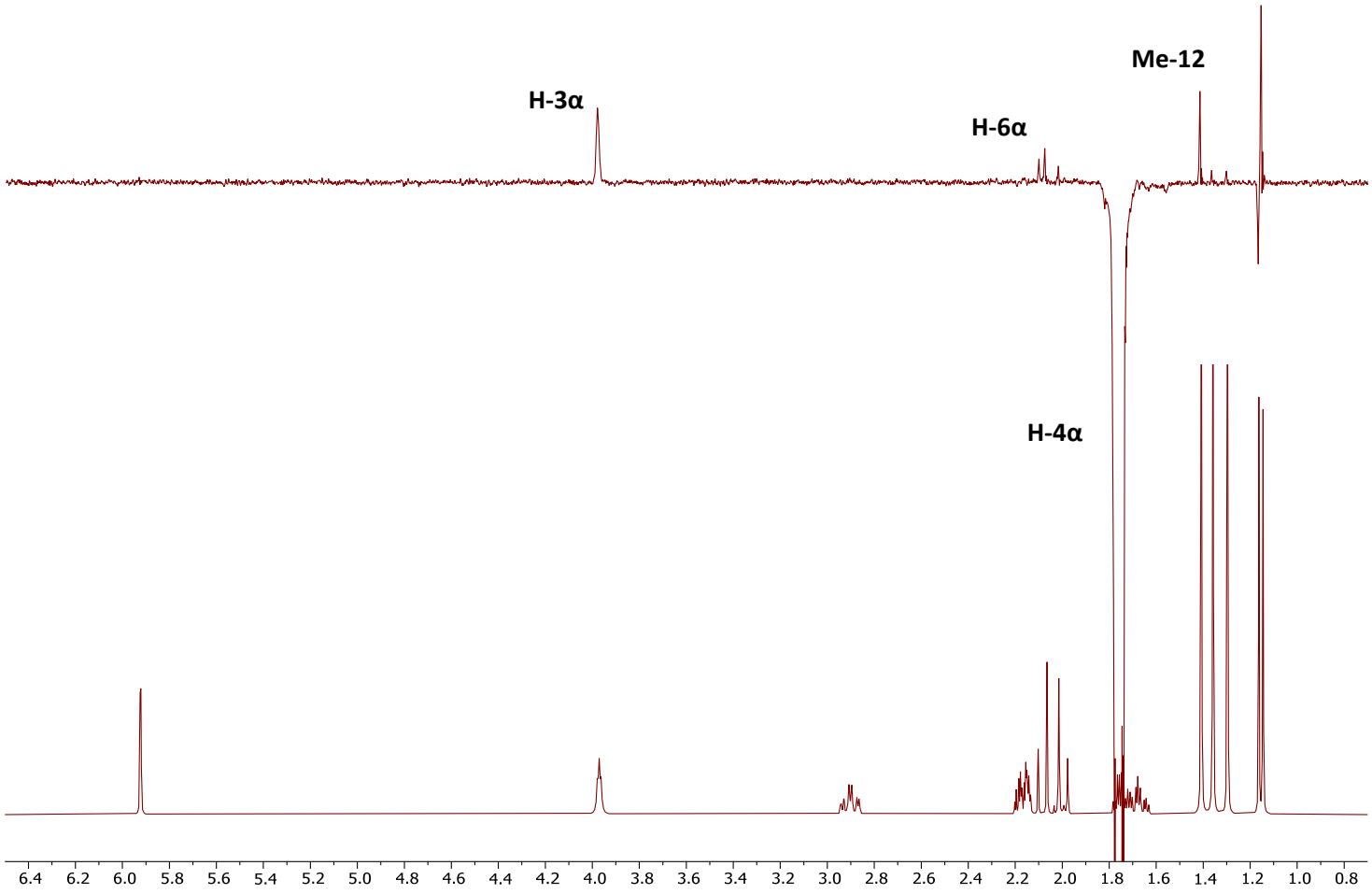


Figure S13c. 1D NOESY spectrum of compound **2**.

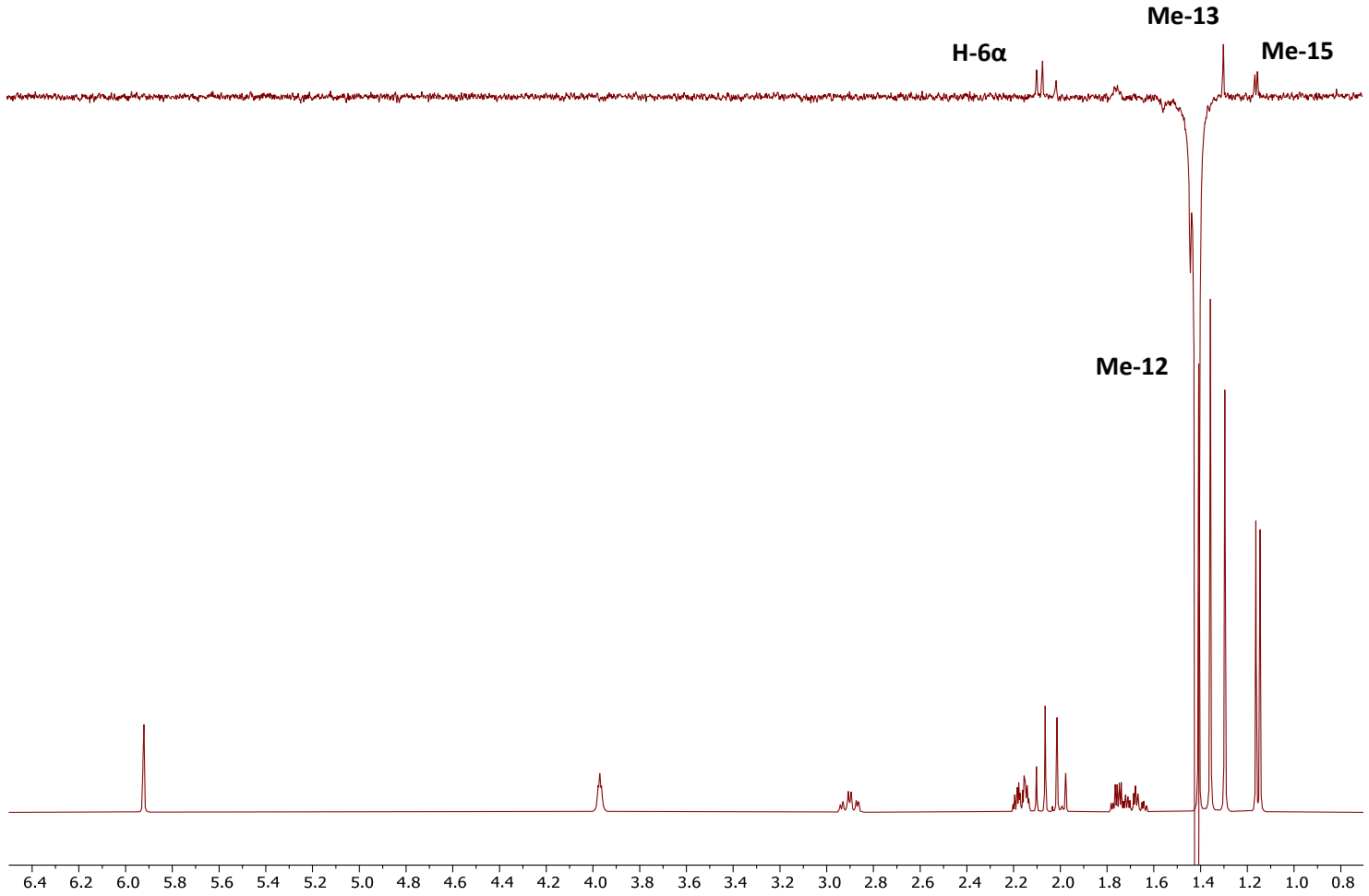


Figure S13d. 1D NOESY spectrum of compound **2**.

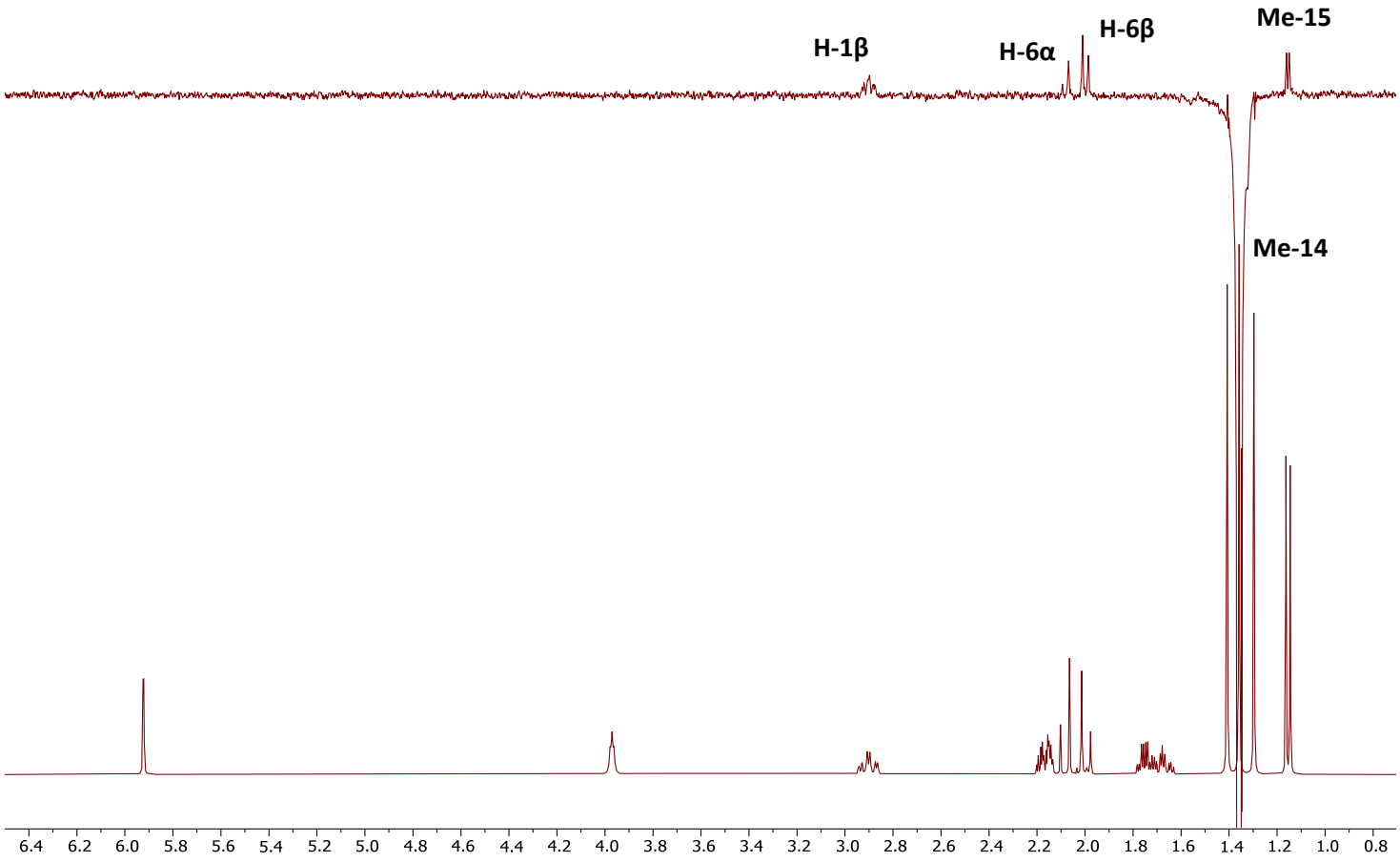


Figure S13e. 1D NOESY spectrum of compound **2**.

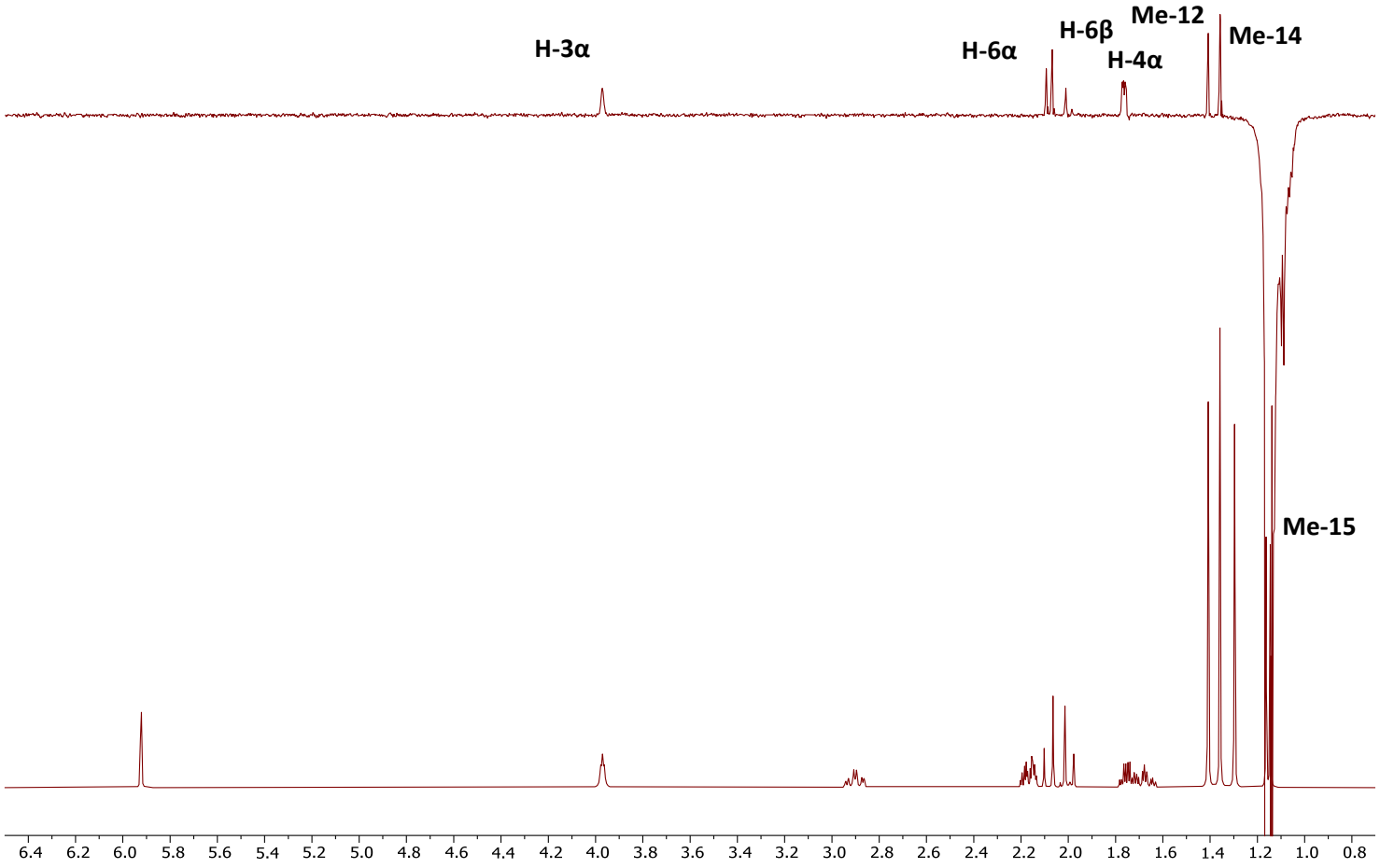
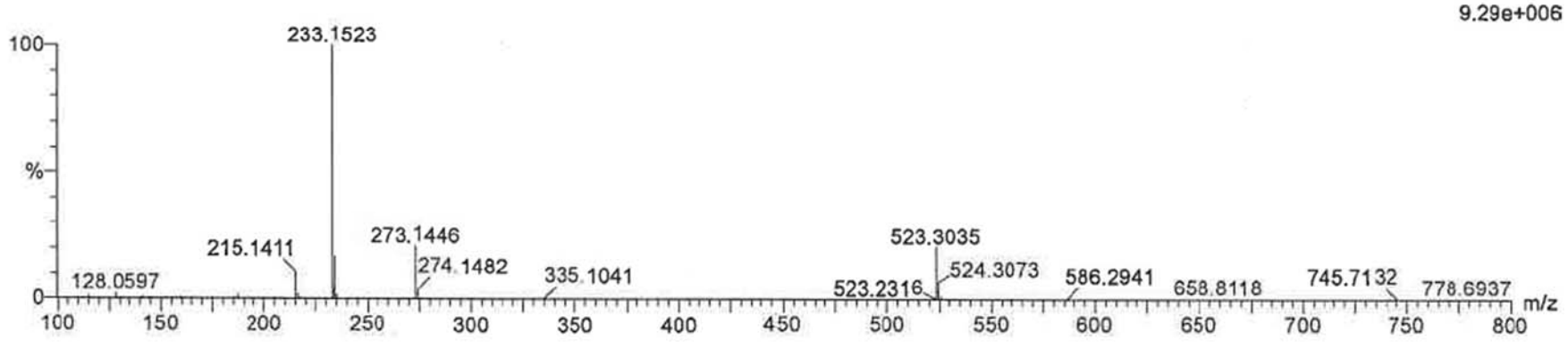


Figure S13f. 1D NOESY spectrum of compound **2**.

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
83 formula(e) evaluated with 2 results within limits (up to 50 best isotopic matches for each mass)
Elements Used:
C: 0-500 H: 0-1000 O: 0-200 Na: 0-1
CD2-80-P6 43 (0.803)
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
273.1446	273.1467	-2.1	-7.7	4.5	685.4	0.150	86.07	C15 H22 O3 Na
	273.1491	-4.5	-16.5	7.5	687.2	1.971	13.93	C17 H21 O3

Figure S14. HRESIMS spectrum of compound **2**.

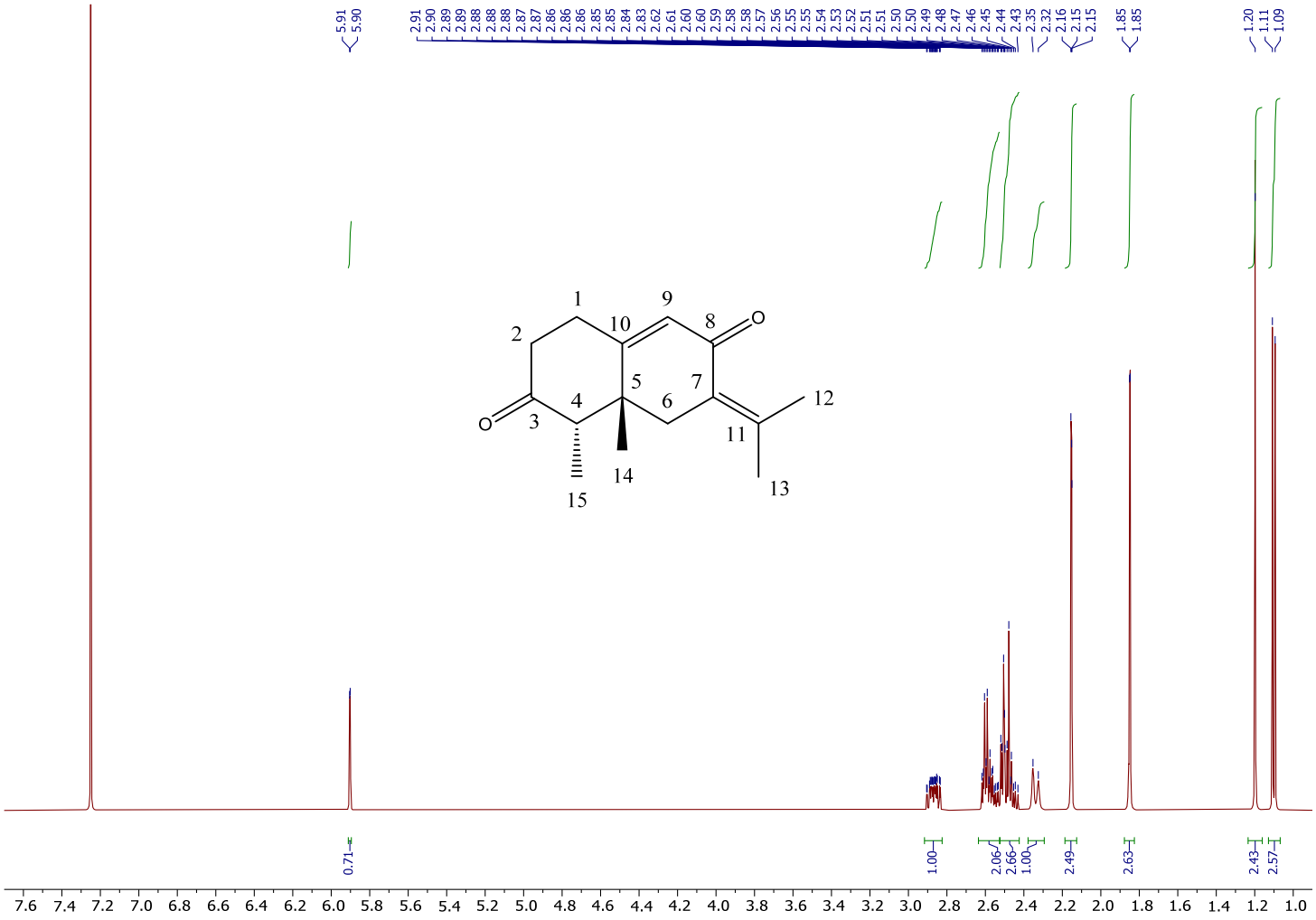


Figure S15. ^{1}H NMR spectrum (500 MHz, $CDCl_3$) of compound **3**.

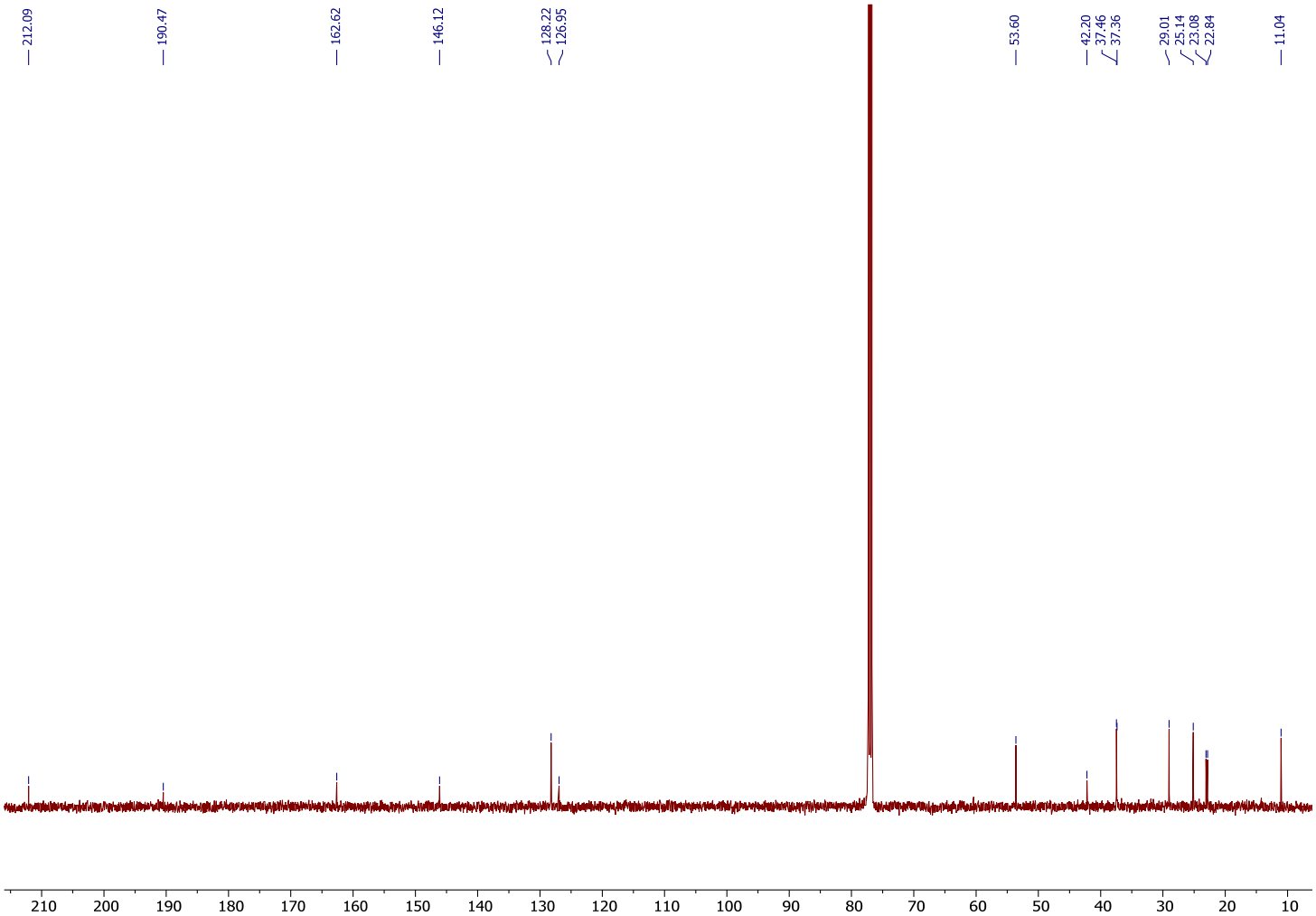


Figure S16. ^{13}C NMR spectrum (125 MHz, $CDCl_3$) of compound **3**.

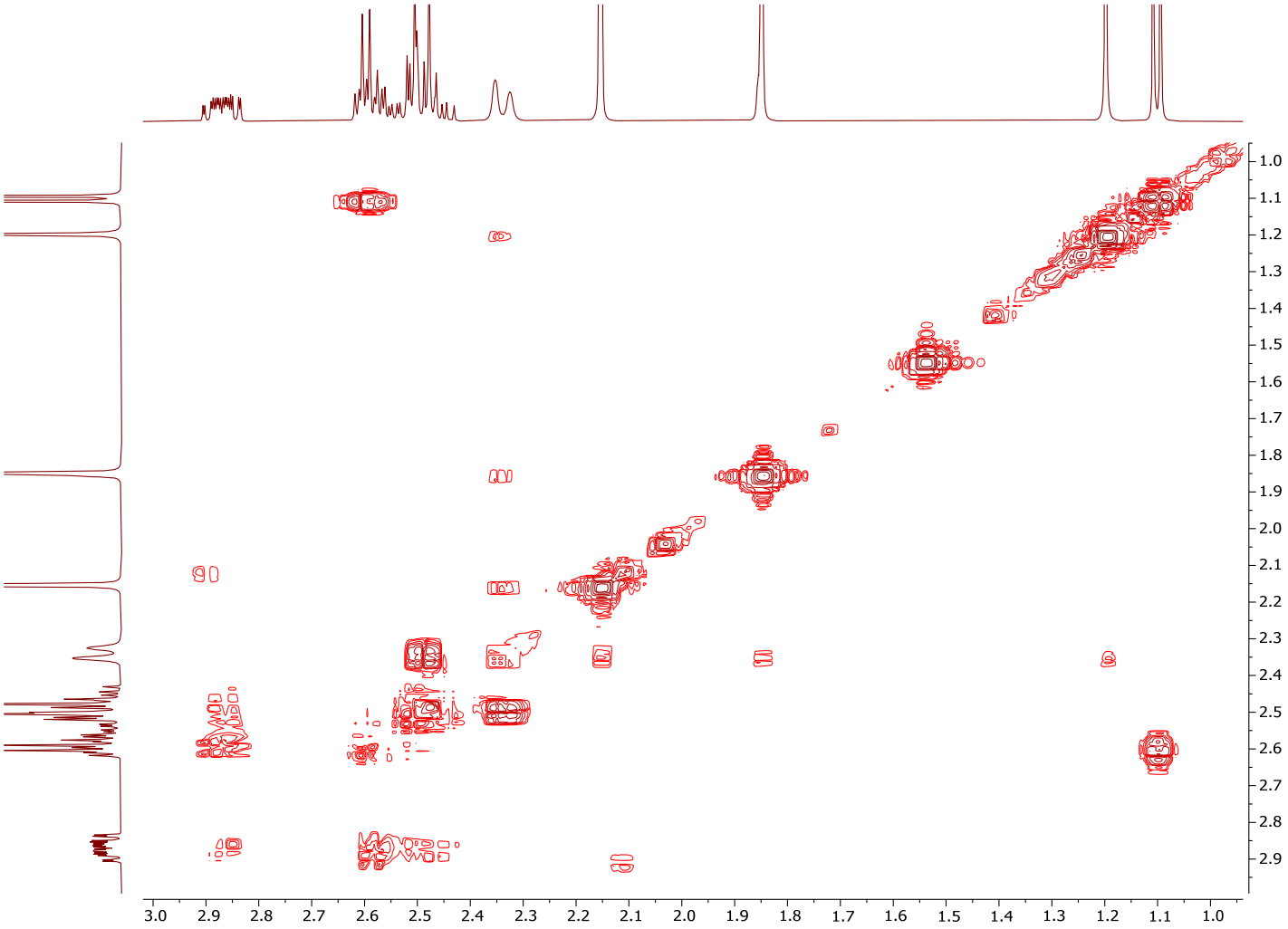


Figure S17. gCOSY spectrum of compound **3**.

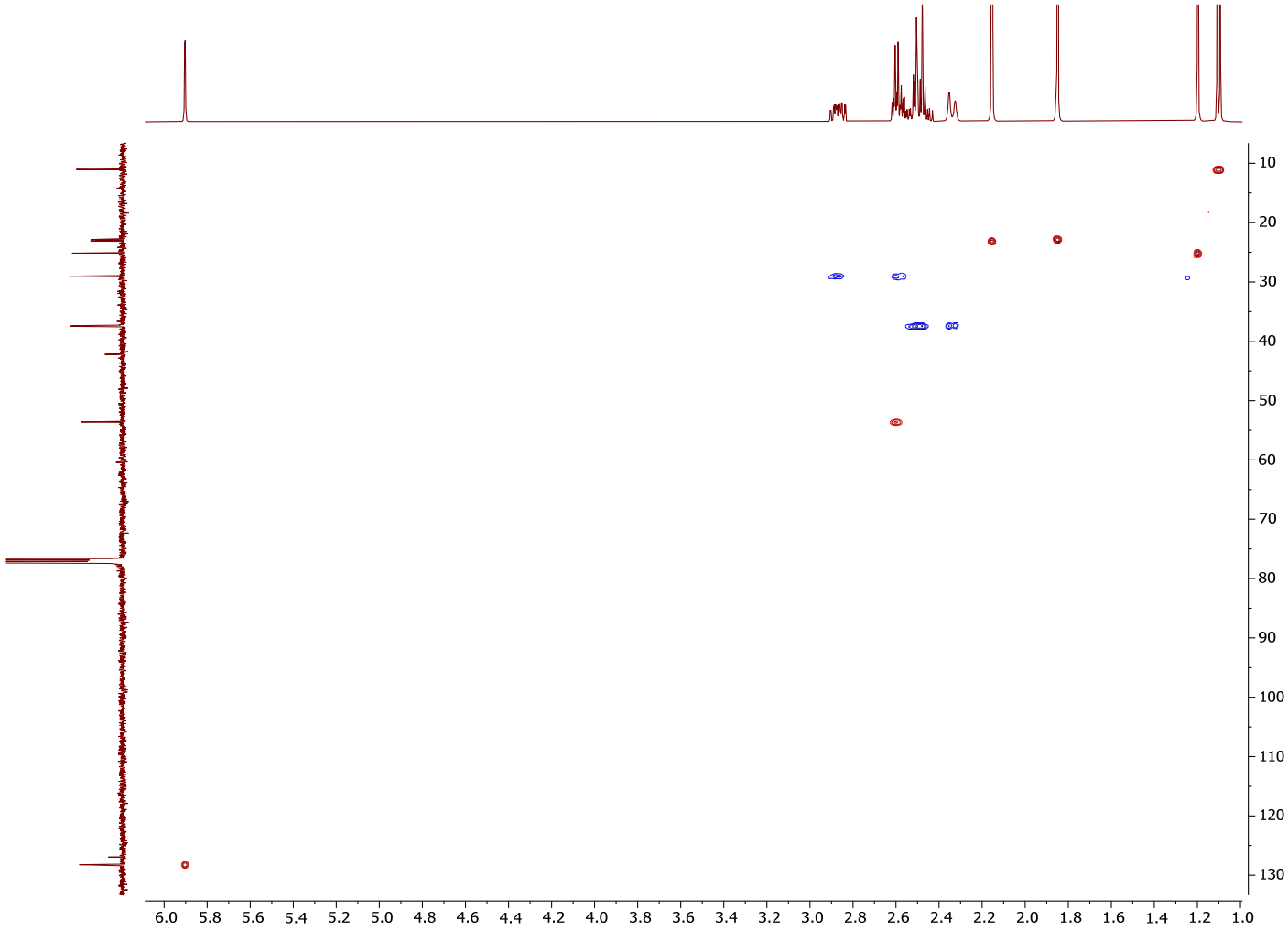


Figure S18. gHSQC spectrum of compound **3**.

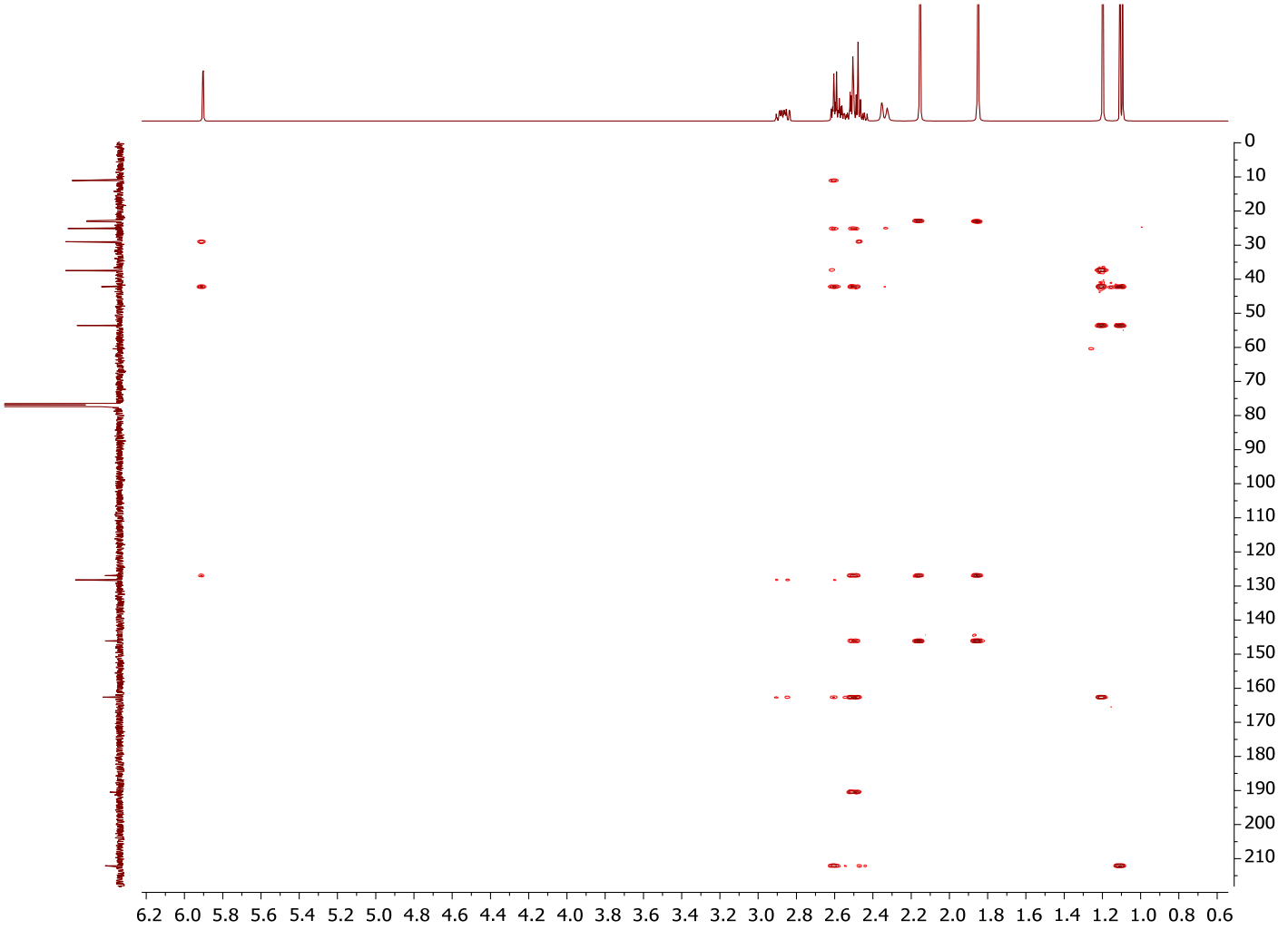


Figure S19. gHMBC spectrum of compound **3**.

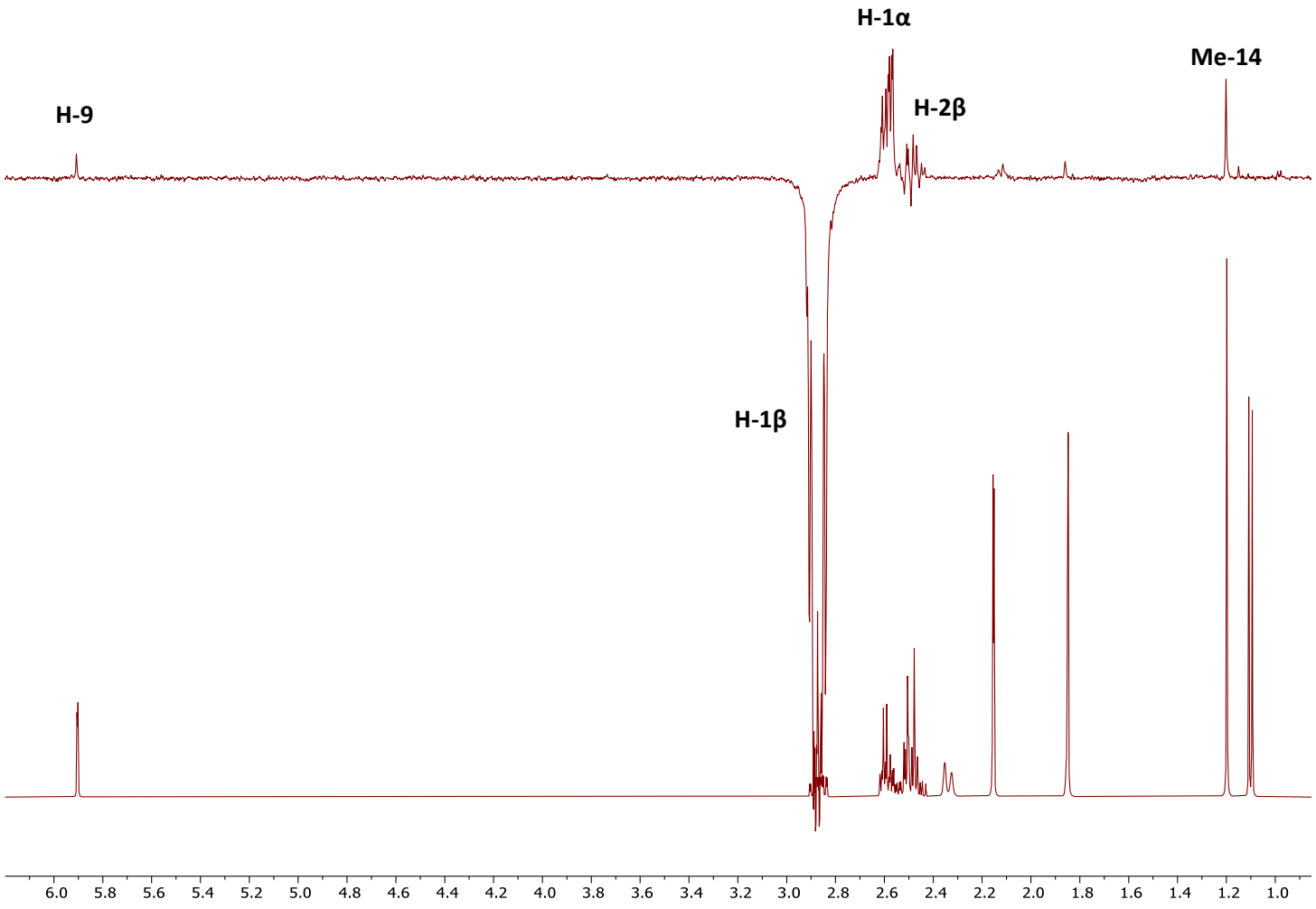


Figure S20a. 1D NOESY spectrum of compound **3**.

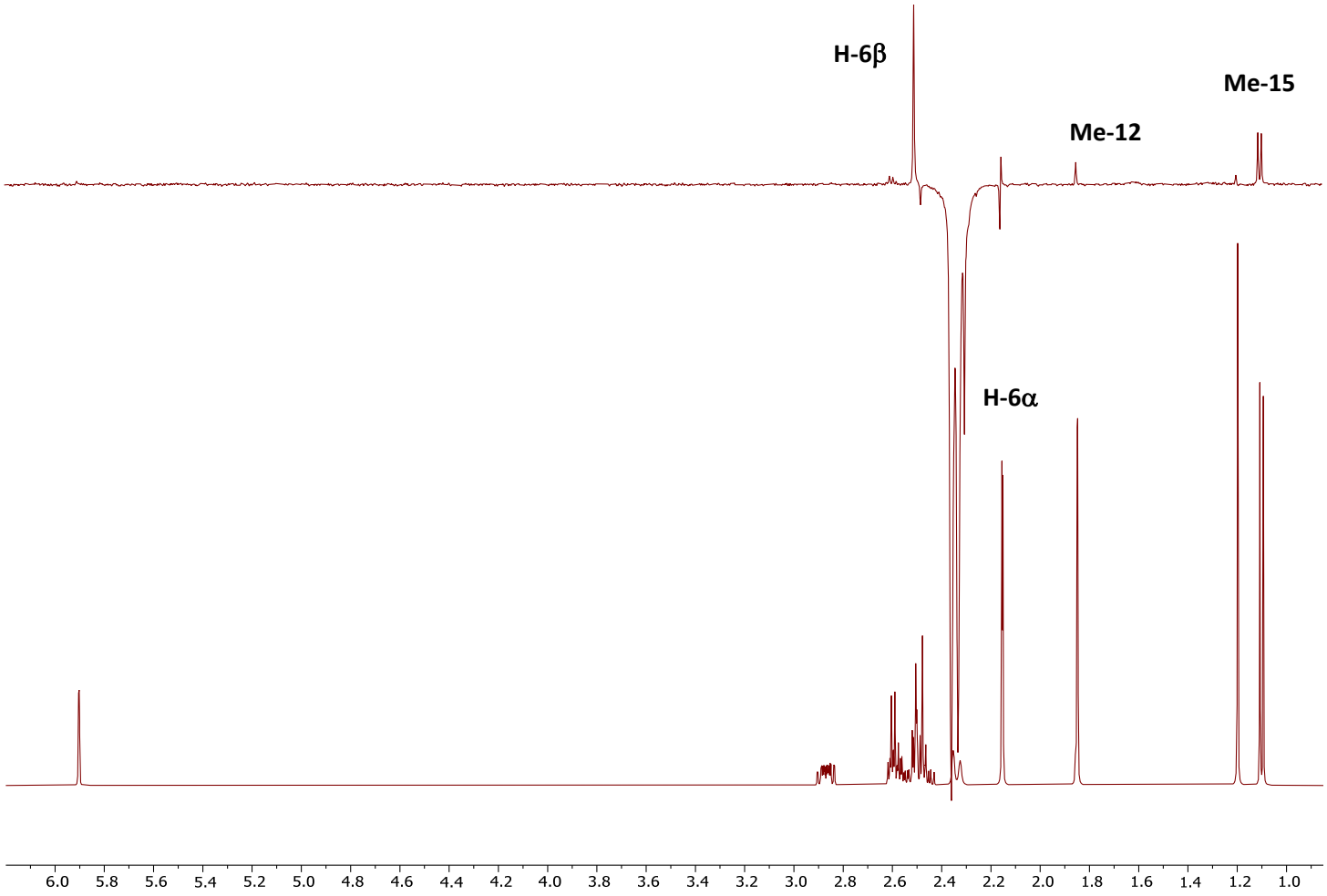


Figure S20b. 1D NOESY spectrum of compound **3**.

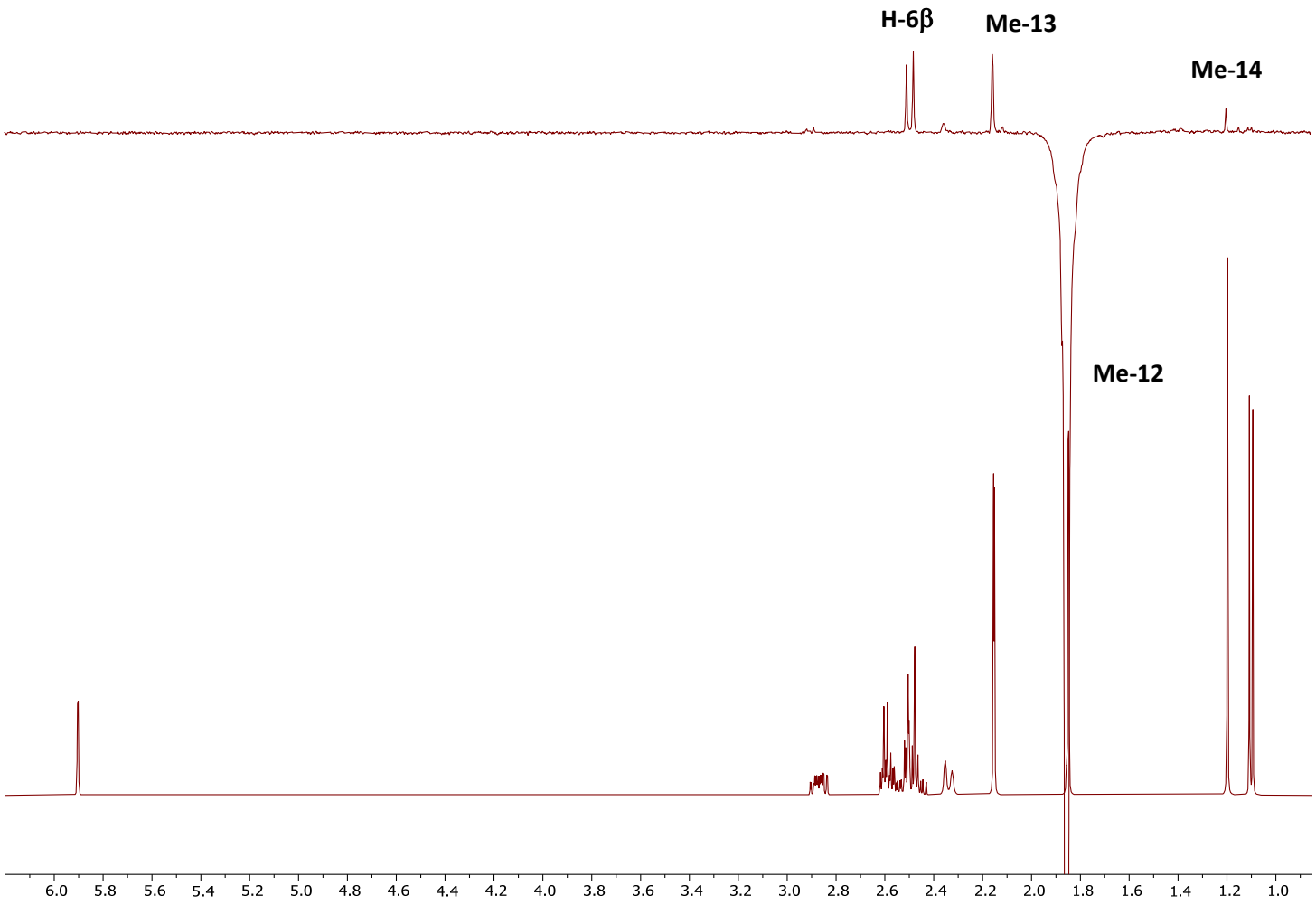


Figure S20c. 1D NOESY spectrum of compound **3**.

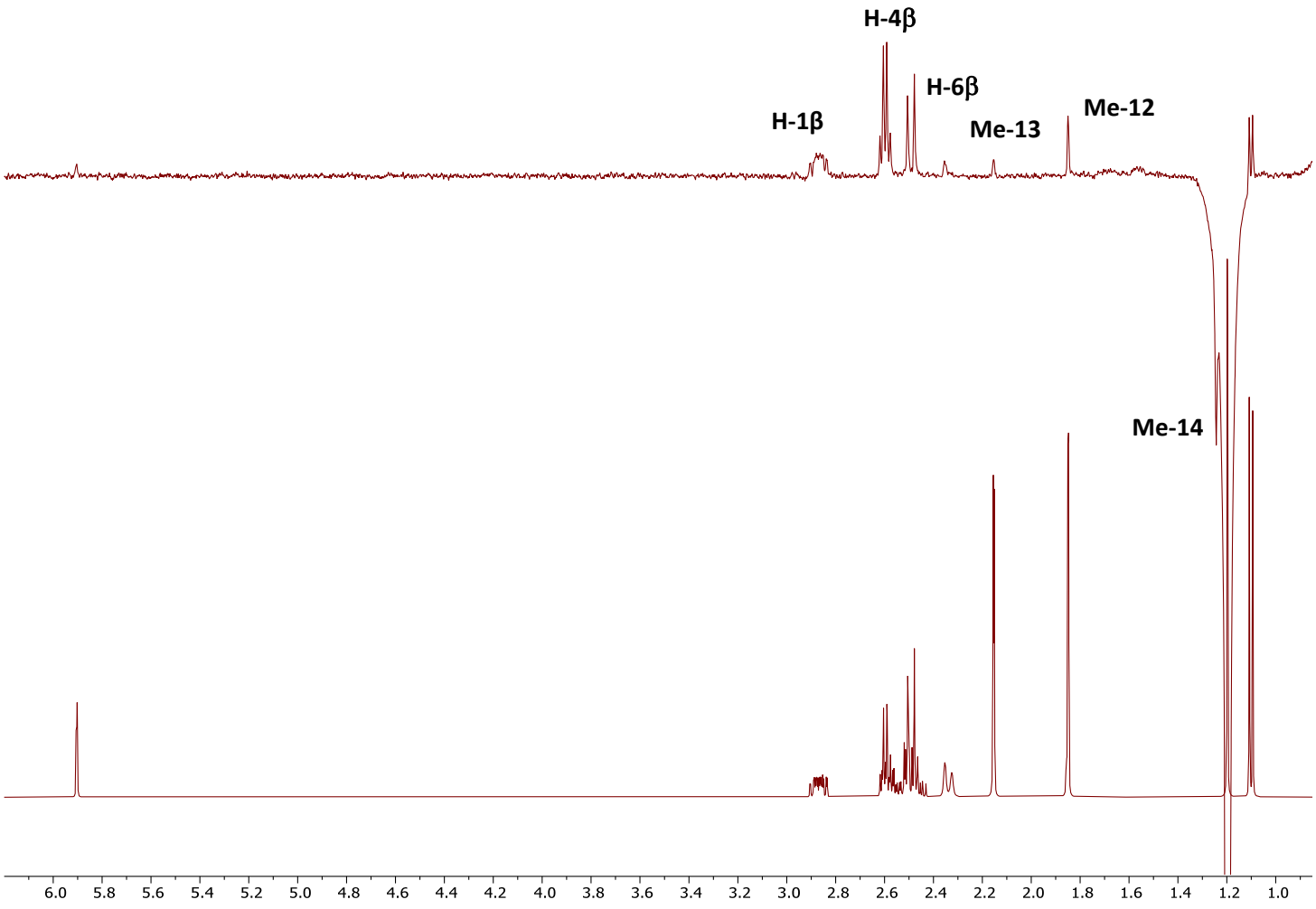


Figure S20d. 1D NOESY spectrum of compound **3**.

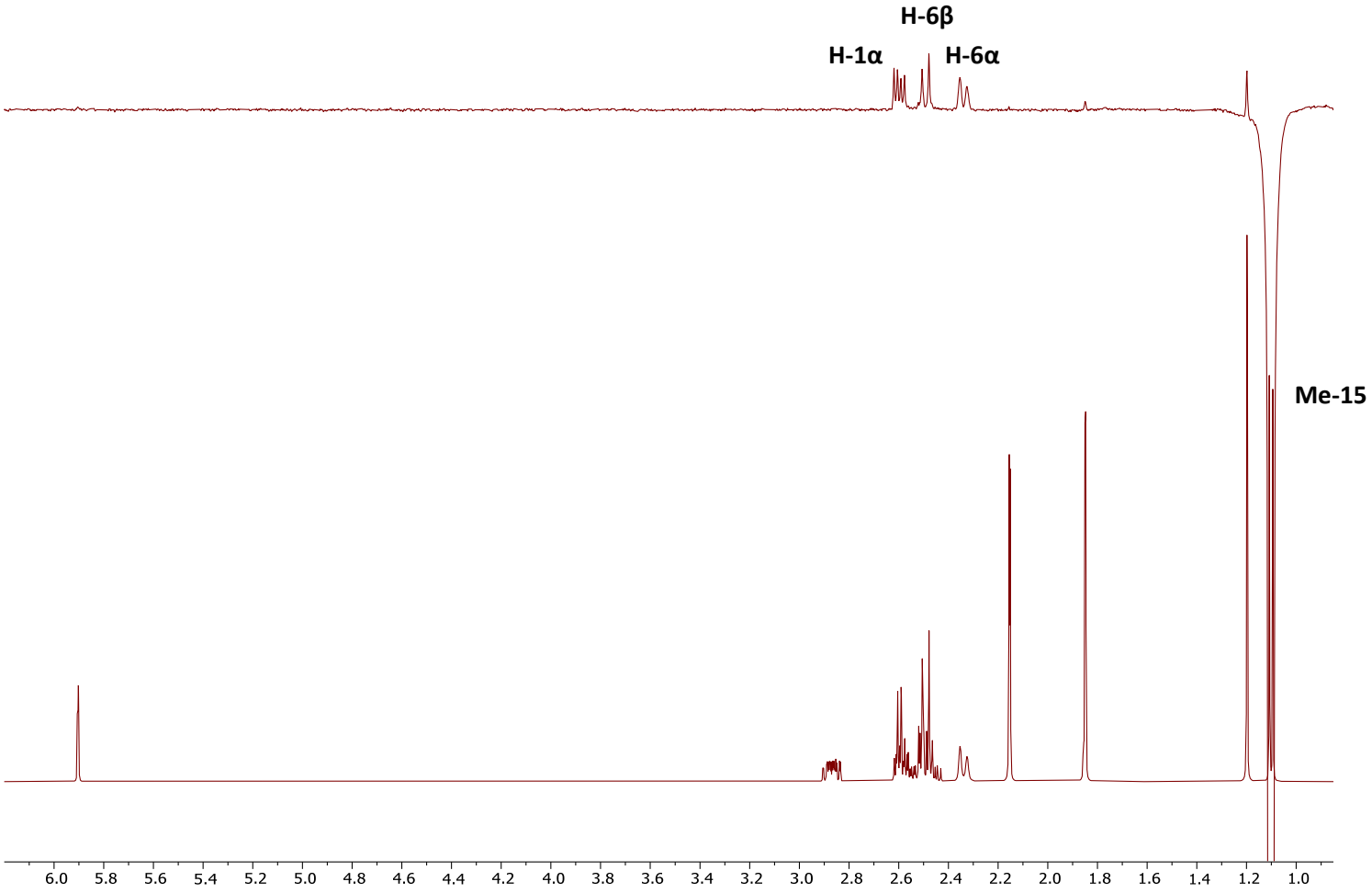
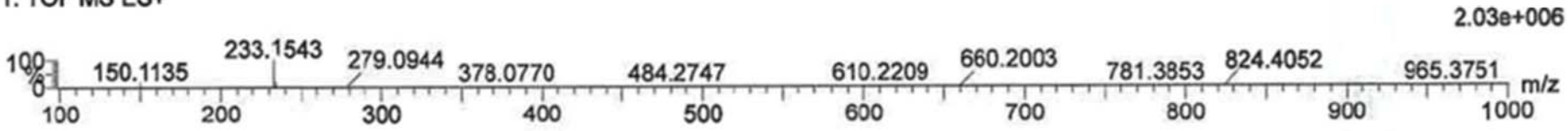


Figure S20e. 1D NOESY spectrum of compound **3**.

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 = 3

Monoisotopic Mass, Even Electron Ions
36 formula(e) evaluated with 1 results within limits (up to 5 best isotopic matches for each mass)
Elements Used:
C: 0-100 H: 0-200 O: 0-50
CD1-40-P8 288 (2.670)
1: TOF MS ES+



Minimum:				-1.5				
Maximum:		5.0	10.0	80.0				
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
233.1543	233.1542	0.1	0.4	5.5	1387.8	n/a	n/a	C15 H21 O2

Figure S21. HRESIMS spectrum of compound **3**.