

Supplementary Information

Seco-tetracenomycins from the marine-derived actinomycete *Saccharothrix* sp. 10-10

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Supporting Information

Contents

Table S1. NMR Spectroscopic Data for 1 and 3–5 in DMSO- <i>d</i> ₆	S4
Table S2. NMR Spectroscopic Data for 13-de- <i>O</i> -methyltetracenomycin X (6) and tetracenomycin X (7) in CD ₃ OD	S5
Table S3. NMR Spectroscopic Data for Saccharothrixones I (5), B (8), and C (9).....	S6
Figure S1. The experimental ECD and UV spectra of saccharothrixones E–I, B, C (1–5 , 8 , 9), 13-de- <i>O</i> -methyltetracenomycin X (6) and Tcm X (7) recorded in MeOH.	S7
Scheme 1. Plausible biosynthesis pathway for saccharothrixones A, B (8), C (9) and E–I (1–5).....	S8
Figure S2. The (+)-HRESIMS spectrum of saccharothrixone E (1).....	S9
Figure S3. The IR spectrum of saccharothrixone E (1).	S10
Figure S4. The ¹ H NMR spectrum of saccharothrixone E (1) in acetone- <i>d</i> ₆ (500 MHz).	S11
Figure S5. The ¹ H NMR spectrum of saccharothrixone E (1) in DMSO- <i>d</i> ₆ (600 MHz).....	S12
Figure S6. The ¹³ C NMR spectrum of saccharothrixone E (1) in acetone- <i>d</i> ₆ (125 MHz).	S13
Figure S7. The HSQC spectrum of saccharothrixone E (1) in acetone- <i>d</i> ₆ (500 MHz).....	S14
Figure S8. The HMBC spectrum of saccharothrixone E (1) in acetone- <i>d</i> ₆ (500 MHz).....	S15
Figure S9. The ROESY spectrum of saccharothrixone E (1) in acetone- <i>d</i> ₆ (500 MHz).	S16
Figure S10. The ROESY spectrum of saccharothrixone E (1) in DMSO- <i>d</i> ₆ (600 MHz).	S17
Figure S11. The (+)-HRESIMS spectrum of saccharothrixone F (2).	S18
Figure S12. The IR spectrum of saccharothrixone F (2).....	S19
Figure S13. The ¹ H NMR spectrum of saccharothrixone F (2) in acetone- <i>d</i> ₆ (600 MHz).	S20
Figure S14. The ¹³ C NMR spectrum of saccharothrixone F (2) in acetone- <i>d</i> ₆ (150 MHz).	S21
Figure S15. The HSQC spectrum of saccharothrixone F (2) in acetone- <i>d</i> ₆ (600 MHz).	S22
Figure S16. The HMBC spectrum of saccharothrixone F (2) in acetone- <i>d</i> ₆ (600 MHz).....	S23
Figure S17. The (–)-HRESIMS spectrum of saccharothrixone G (3).	S24
Figure S18. The IR spectrum of saccharothrixone G (3).	S25
Figure S19. The ¹ H NMR spectrum of saccharothrixone G (3) in acetone- <i>d</i> ₆ (600 MHz).....	S26
Figure S20. The ¹ H NMR spectrum of saccharothrixone G (3) in DMSO- <i>d</i> ₆ (600 MHz).	S27
Figure S21. The ¹³ C NMR spectrum of saccharothrixone G (3) in acetone- <i>d</i> ₆ (150 MHz).....	S28
Figure S22. The HSQC spectrum of saccharothrixone G (3) in acetone- <i>d</i> ₆ (600 MHz).	S29
Figure S23. The HMBC spectrum of saccharothrixone G (3) in acetone- <i>d</i> ₆ (600 MHz).	S30
Figure S24. The ROESY spectrum of saccharothrixone G (3) in acetone- <i>d</i> ₆ (600 MHz).....	S31
Figure S25. The 1D NOE spectrum of saccharothrixone G (3) in DMSO- <i>d</i> ₆ (600 MHz).....	S32
Figure S26. The 1D NOE spectrum of saccharothrixone G (3) in DMSO- <i>d</i> ₆ (600 MHz).....	S33
Figure S27. The (–)-HRESIMS spectrum of saccharothrixone H (4).	S34
Figure S28. The IR spectrum of saccharothrixone H (4).	S35
Figure S29. The ¹ H NMR spectrum of saccharothrixone H (4) in acetone- <i>d</i> ₆ (600 MHz).....	S36
Figure S30. The ¹³ C NMR spectrum of saccharothrixone H (4) in acetone- <i>d</i> ₆ (150 MHz).....	S37
Figure S31. The HSQC spectrum of saccharothrixone H (4) in acetone- <i>d</i> ₆ (600 MHz).	S38
Figure S32. The HMBC spectrum of saccharothrixone H (4) in acetone- <i>d</i> ₆ (600 MHz).	S39
Figure S33. The ROESY spectrum of saccharothrixone H (4) in acetone- <i>d</i> ₆ (600 MHz).....	S40

Figure S34. The (–)-HRESIMS spectrum of saccharothrixone I (5).....	S41
Figure S35. The IR spectrum of saccharothrixone I (5).	S42
Figure S36. The ¹ H NMR spectrum of saccharothrixone I (5) in acetone- <i>d</i> ₆ (500 MHz).	S43
Figure S37. The ¹ H NMR spectrum of saccharothrixone I (5) in DMSO- <i>d</i> ₆ (600 MHz).....	S44
Figure S38. The ¹³ C NMR spectrum of saccharothrixone I (5) in acetone- <i>d</i> ₆ (125 MHz).	S45
Figure S39. The HSQC spectrum of saccharothrixone I (5) in acetone- <i>d</i> ₆ (500 MHz).	S46
Figure S40. The HMBC spectrum of saccharothrixone I (5) in acetone- <i>d</i> ₆ (500 MHz).	S47
Figure S41. The ROESY spectrum of saccharothrixone I (5) in acetone- <i>d</i> ₆ (600 MHz).	S48
Figure S42. The ROESY spectrum of saccharothrixone I (5) in DMSO- <i>d</i> ₆ (600 MHz).	S49
Figure S43. The (+)-HR ESIMS spectrum of 13-de- <i>O</i> -methyltetracenomycin X (6).	S50
Figure S44. The IR spectrum of 13-de- <i>O</i> -methyltetracenomycin X (6).....	S51
Figure S45. The ¹ H NMR spectrum of 13-de- <i>O</i> -methyltetracenomycin X (6) in CD ₃ OD.....	S52
Figure S46. The ¹³ C NMR spectrum of 13-de- <i>O</i> -methyltetracenomycin X (6) in CD ₃ OD.....	S53
Figure S47. The HSQC spectrum of 13-de- <i>O</i> -methyltetracenomycin X (6) in CD ₃ OD	S54
Figure S48. The HMBC spectrum of 13-de- <i>O</i> -methyltetracenomycin X (6) in CD ₃ OD.	S55
Figure S49. Optimized conformers (≥ 1%) of (4 <i>S</i> ,4 <i>aR</i> ,5 <i>S</i> ,12 <i>aR</i>)- 1a at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.....	S56
Figure S50. Optimized conformers (≥ 1%) of (4 <i>S</i> ,4 <i>aR</i> ,5 <i>S</i> ,12 <i>aR</i>)- 2a at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.....	S56
Figure S51. Optimized conformers (≥ 1%) of (4 <i>S</i> ,4 <i>aR</i> ,5 <i>S</i> ,12 <i>aS</i>)- 3a at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.....	S56
Figure S52. Optimized conformers (≥ 1%) of (4 <i>R</i> ,4 <i>aR</i> ,5 <i>S</i> ,12 <i>aS</i>)- 3c at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.....	S56
Figure S53. Optimized conformers (≥ 1%) of (4 <i>S</i> ,4 <i>aR</i> ,5 <i>R</i> ,12 <i>aR</i>)- 5a at the B3LYP/6-311+G (d,p) level with PCM model in MeOH.....	S57

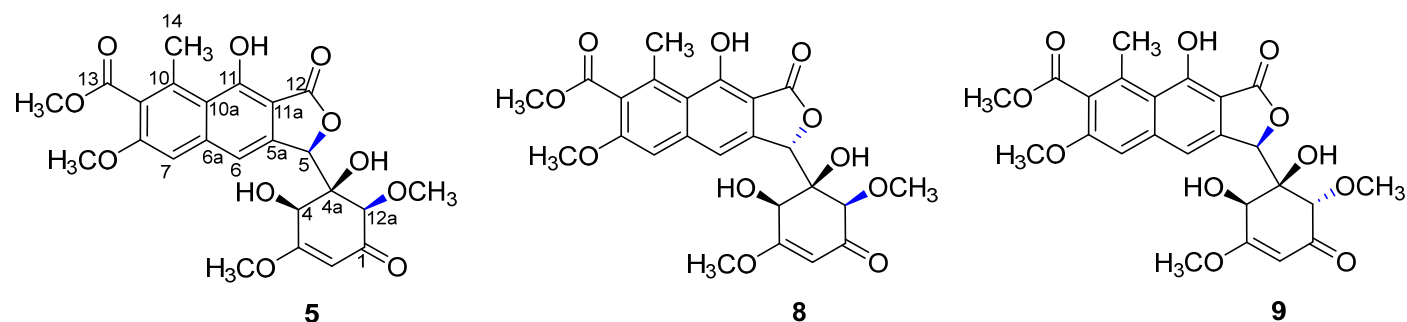
Table S1. NMR Spectroscopic Data for **1** and **3–5** in DMSO-*d*₆^a

no	1 (DMSO- <i>d</i> ₆)	3 (DMSO- <i>d</i> ₆)		4 (DMSO- <i>d</i> ₆)		5 (DMSO- <i>d</i> ₆)	
No.	δ_{H} , mult. (<i>J</i> in Hz)	δ_{C}	δ_{H} , mult. (<i>J</i> in Hz)	δ_{C}	δ_{H} , mult. (<i>J</i> in Hz)	δ_{C}	δ_{H} , mult. (<i>J</i> in Hz)
1		171.9		172.0		194.8	
2	5.62, s	88.6	5.38, d (1.2)	88.3	5.42, s	99.1	5.29, d (1.2)
3		183.4		181.3		173.2	
4	4.54, d (3.0)	77.0	5.16, d (1.2)	79.1	6.10, s	67.6	4.54, d (7.8)
4a		81.3		77.7		80.0	
5	4.58, d (3.0)	67.6	4.85, d (10.2)	66.5	5.18, d (9.6)	79.1	5.81, d (1.2)
5a		140.6		141.0		143.1	
6	7.41, s	116.7	7.42, s	116.6	7.42, s	113.8	7.52, d (1.2)
6a		140.6		141.0		141.2	
7	7.31, s	105.1	7.26, s	105.4	7.30, s	105.4	7.25, s
8		156.2		156.5		155.0	
9		126.0		125.9		125.9	
10		135.9		136.0		134.8	
10a		116.4		116.4		117.3	
11		163.9		164.0		156.9	
11a		108.6		107.8		105.6	
12		202.8		203.0		170.9	
12a		83.7	4.42, s	81.6	3.56, s	83.6	3.82, s
13		167.8		167.7		167.9	
14	2.73, s	20.5	2.72, s	20.6	2.72, s	20.4	2.73, s
3-OCH ₃	3.83, s	59.7	3.84, s	59.8	3.87, s	56.4	3.64, s
8-OCH ₃	3.89, s	56.1	3.90, s	56.2	3.91, s	55.9	3.88, s
12a-OCH ₃	3.60, s	60.6	3.63, s	58.4	3.25, s	59.7	3.36, s
13-OCH ₃	3.86, s	52.4	3.86, s	52.4	3.86, s	52.4	3.86, s
4-OH	5.47, d (3.0)						5.37, d (7.8)
4a-OH	5.23, s		5.26, s		5.31, s		5.74, s
5-OH	6.07, brs		5.81, d (10.2)		5.86, d (9.6)		
11-OH	14.70, brs		14.48, brs		n.o.		

^a ¹H NMR and ¹³C NMR data were recorded at 600 and 150 MHz, respectively. The assignments were based on HSQC and HMBC experiments. n.o.: not observed.

Table S2. NMR Spectroscopic Data for 13-de-*O*-methyltetracenomycin X (**6**) and tetracenomycin X (**7**) in CD₃OD ^a

No.	6 (CD ₃ OD)		7 (CD ₃ OD)	
	δ_C , type	δ_H , m	δ_C	δ_H , m
1	193.9, C		188.3	
2	101.9, CH	5.53, d (1.8)	101.4	5.32, brs
3	174.8, C		178.8	
4	70.9, CH	4.85, brs	70.2	4.88, brs
4a	86.1, C		84.8	
5	194.4, C		195.4	
5a	141.9, C		141.4	
6	121.8, CH	8.01, s	115.8	7.42, s
6a	128.7, C		127.2	
7	108.7, CH	7.46, s	108.0	7.20, s
8	159.3, C		157.8	
9	132.8, C		130.6	
10	138.0, C		140.2	
10a	121.9, C		127.2	
11	167.5, C		170.1	
11a	110.5, C		112.1	
12	197.6, C		195.6	
12a	89.0, C		88.9	
13	171.8, C		173.5	
14	21.1, CH ₃	2.88, s	20.5	2.63, s
3-OCH ₃	57.5, CH ₃	3.80, s	56.4	3.85, s
8-OCH ₃	56.7, CH ₃	4.01, s	55.7	3.91, s
12a-OCH ₃	56.7, CH ₃	3.56, s	55.6	3.48, s
13-OCH ₃			52.2	3.58, s

Table S3. NMR Spectroscopic Data for Saccharothrixones I (**5**), B (**8**), and C (**9**) ^a

No.	5 (acetone- <i>d</i> ₆)		8 (acetone- <i>d</i> ₆) ^b		9 (acetone- <i>d</i> ₆) ^b	
	δ_C , type	δ_H , m	δ_C , type	δ_H , m	δ_C , type	δ_H , m
1	194.9, C		196.0, C		193.6, C	
2	100.3, CH	5.32, d (1.0)	100.7, CH	5.26, s	100.6, CH	5.30, s
3	173.1, C		174.4, C		174.0, C	
4	68.3, CH	4.65, d (1.0)	72.2, CH	3.88, s	69.0, CH	4.62, s
4a	80.3, C		77.1, C		78.4, C	
5	81.0, CH	5.98, s	83.1, CH	5.81, s	84.7, CH	5.92, s
5a	143.0, C		142.1, C		142.4, C	
6	115.0, CH	7.61, s	114.3, CH	7.57, s	115.1, CH	7.61, s
6a	143.3, C		142.8, C		143.1, C	
7	106.2, CH	7.27, s	106.3, CH	7.37, s	106.3, CH	7.32, s
8	156.9, C		156.8, C		157.0, C	
9	127.7, C		127.7, C		127.9, C	
10	136.1, C		136.0, C		136.2, C	
10a	118.0, C		117.7, C		118.0, C	
11	158.5, C		158.2, C		158.6, C	
11a	106.2, C		107.7, C		105.9, C	
12	173.4, C		173.7, C		173.3, C	
12a	84.1, CH	3.93, s	81.6, CH	4.42, s	83.8, CH	3.84, s
13	168.6, C		168.7, C		168.5, C	
14	20.4, CH ₃	2.81, s	20.4, CH ₃	2.82, s	20.4, CH ₃	2.81, s
3-OCH ₃	56.8, CH ₃	3.72, s	56.7, CH ₃	3.65, s	56.9, CH ₃	3.75, s
8-OCH ₃	56.3, CH ₃	3.93, s	56.4, CH ₃	3.97, s	56.4, CH ₃	3.97, s
12a-OCH ₃	60.6, CH ₃	3.51, s	61.1, CH ₃	3.73, s	59.4, CH ₃	3.28, s
13-OCH ₃	52.5, CH ₃	3.90, s	52.5, CH ₃	3.91, s	52.5, CH ₃	3.91, s

^a ¹H NMR data were measured for **5**, **8** and **9** at 500, 800, and 600 MHz, respectively. ¹³C NMR data were measured for **5**, **8** and **9** at 125, 200, and 1500 MHz. The assignments were based on HSQC and HMBC experiments. ^b The NMR data were reported in our previous work: Gan, M. et al. *J. Nat. Prod.* 2015, 78, 2260-2265.

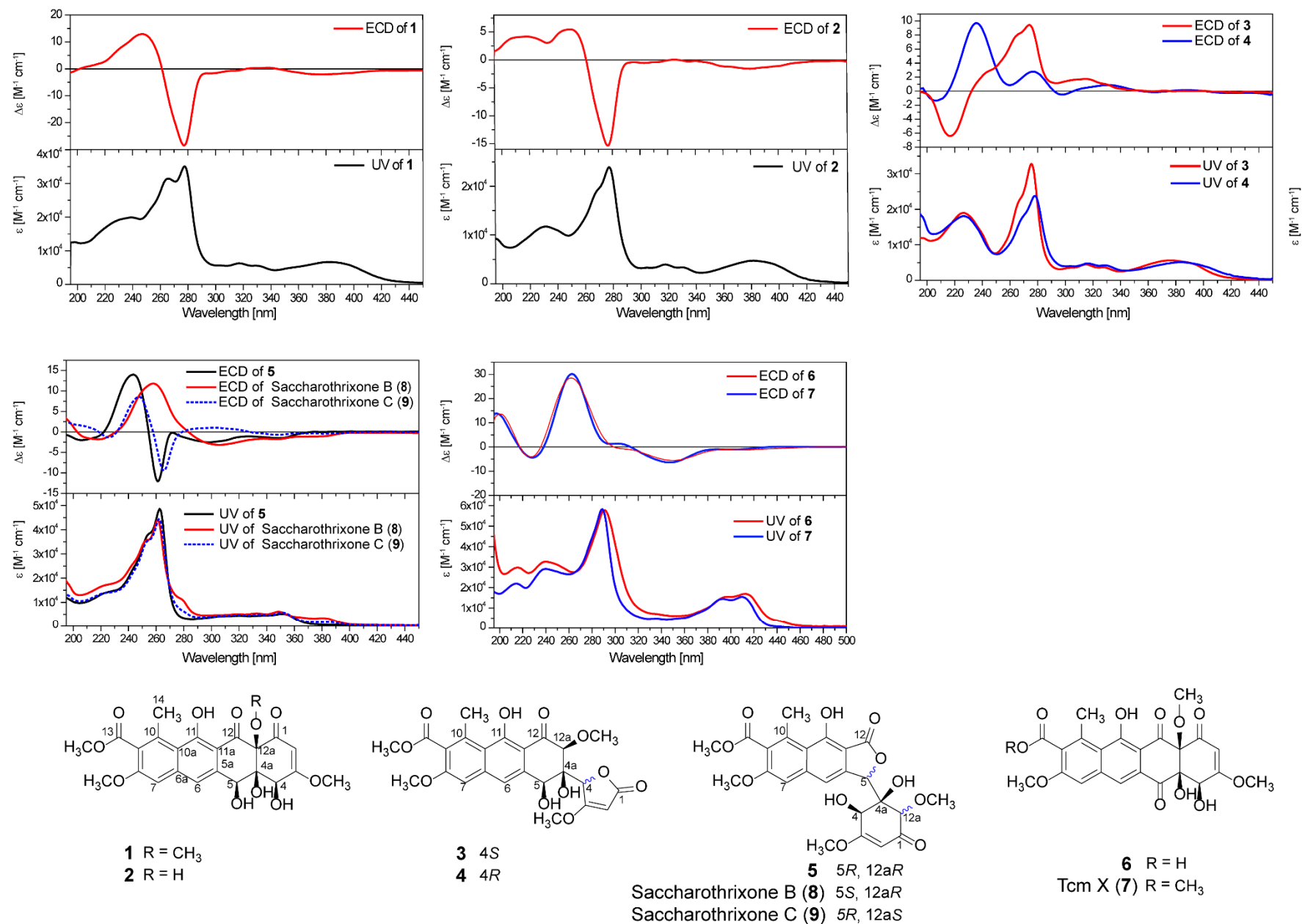
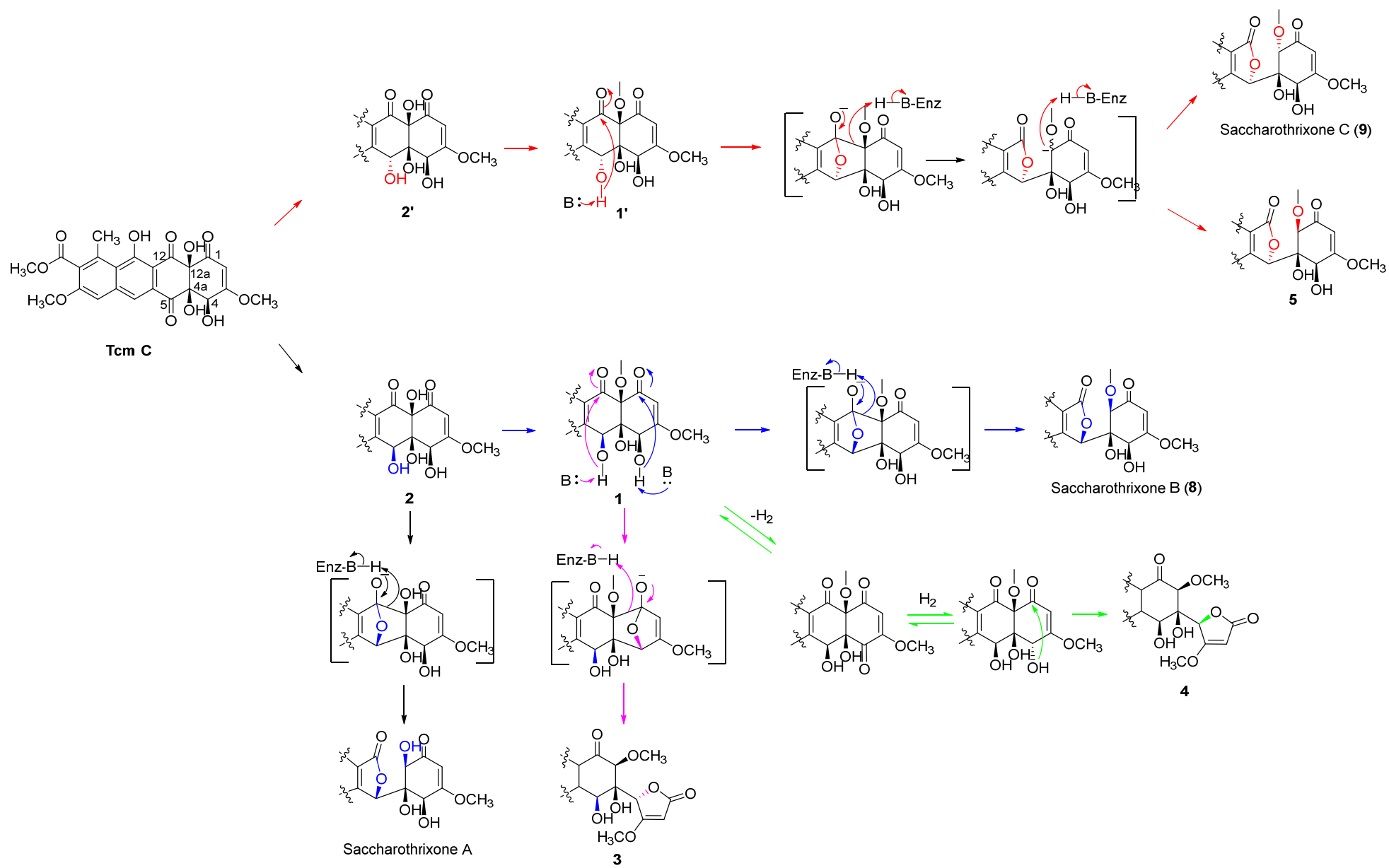


Figure S1. The experimental ECD and UV spectra of saccharothrixones E–I, B, C (**1–5**, **8**, **9**), 13-de-*O*-methyltetracenomycin X (**6**) and Tcm X (**7**) recorded in MeOH.



Scheme 1. Plausible biosynthesis pathway for saccharothrixones A, B (**8**), C (**9**) and E-I (**1–5**).

marine-20-6-HRESI #59 RT: 0.87 AV: 1 NL: 1.44E6
T: FTMS + c ESI Full ms [100.00-1000.00]

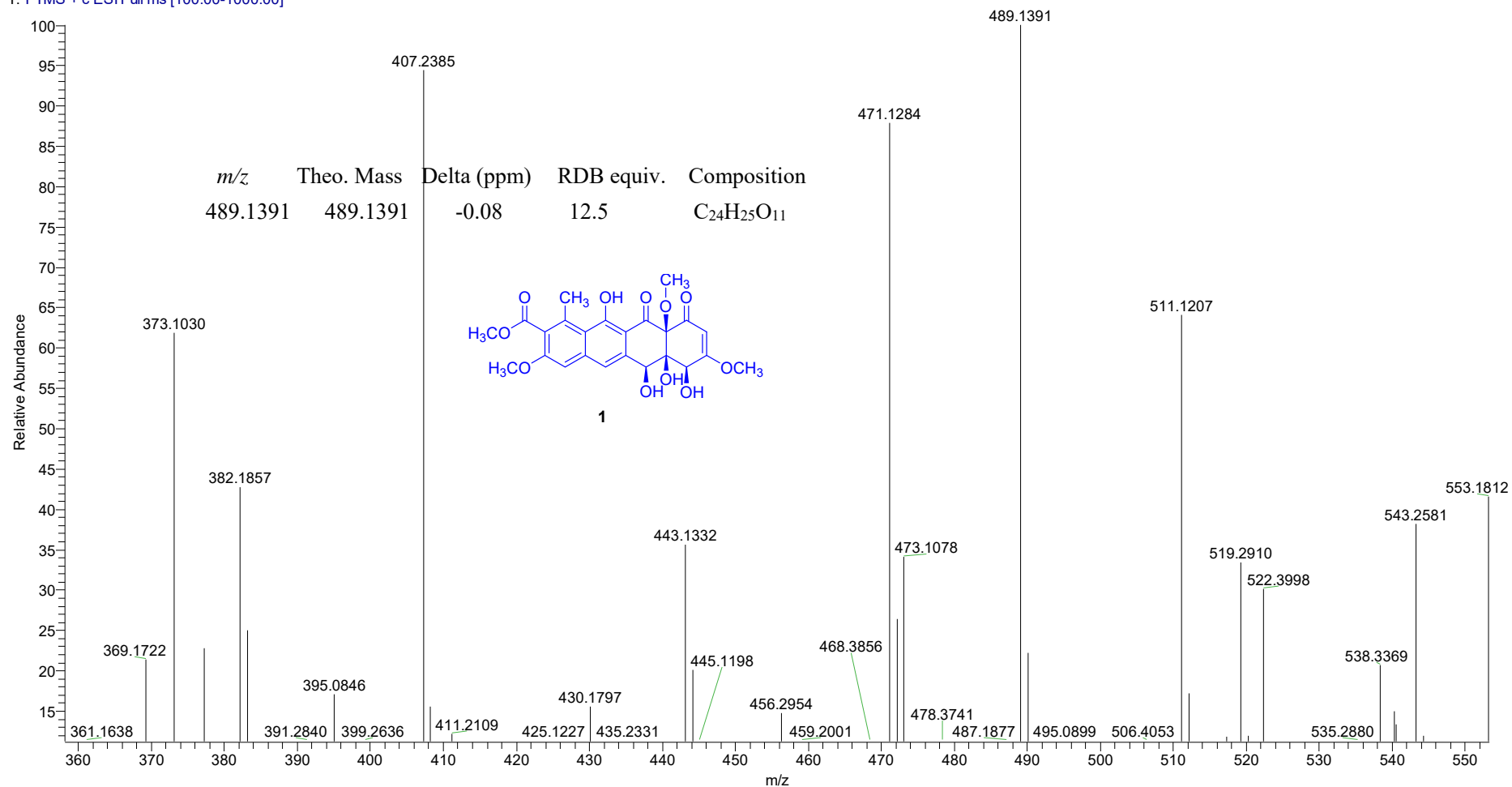


Figure S2. The (+)-HRESIMS spectrum of saccharothrixone E (**1**).

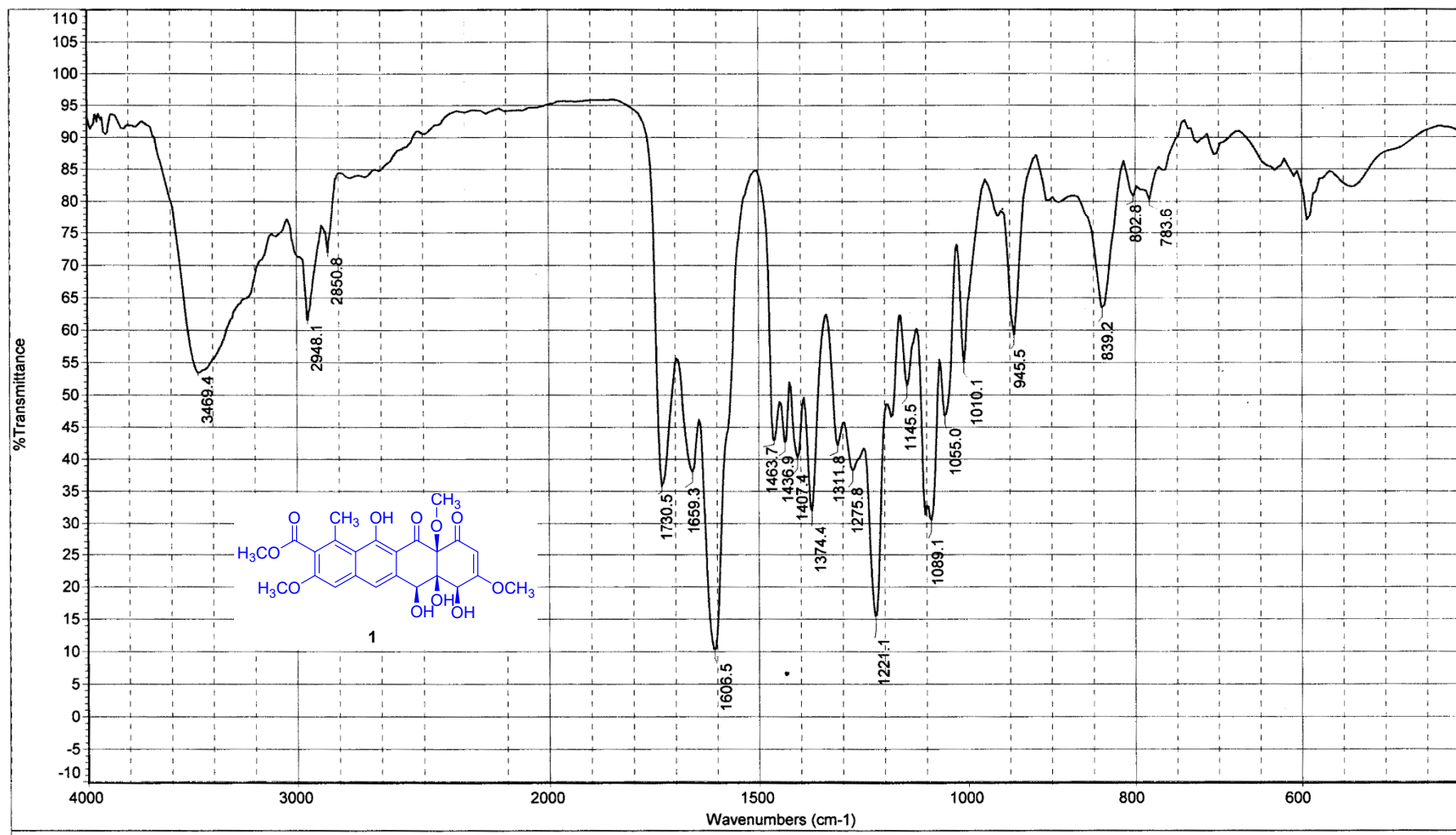


Figure S3. The IR spectrum of saccharothrixone E (**1**).

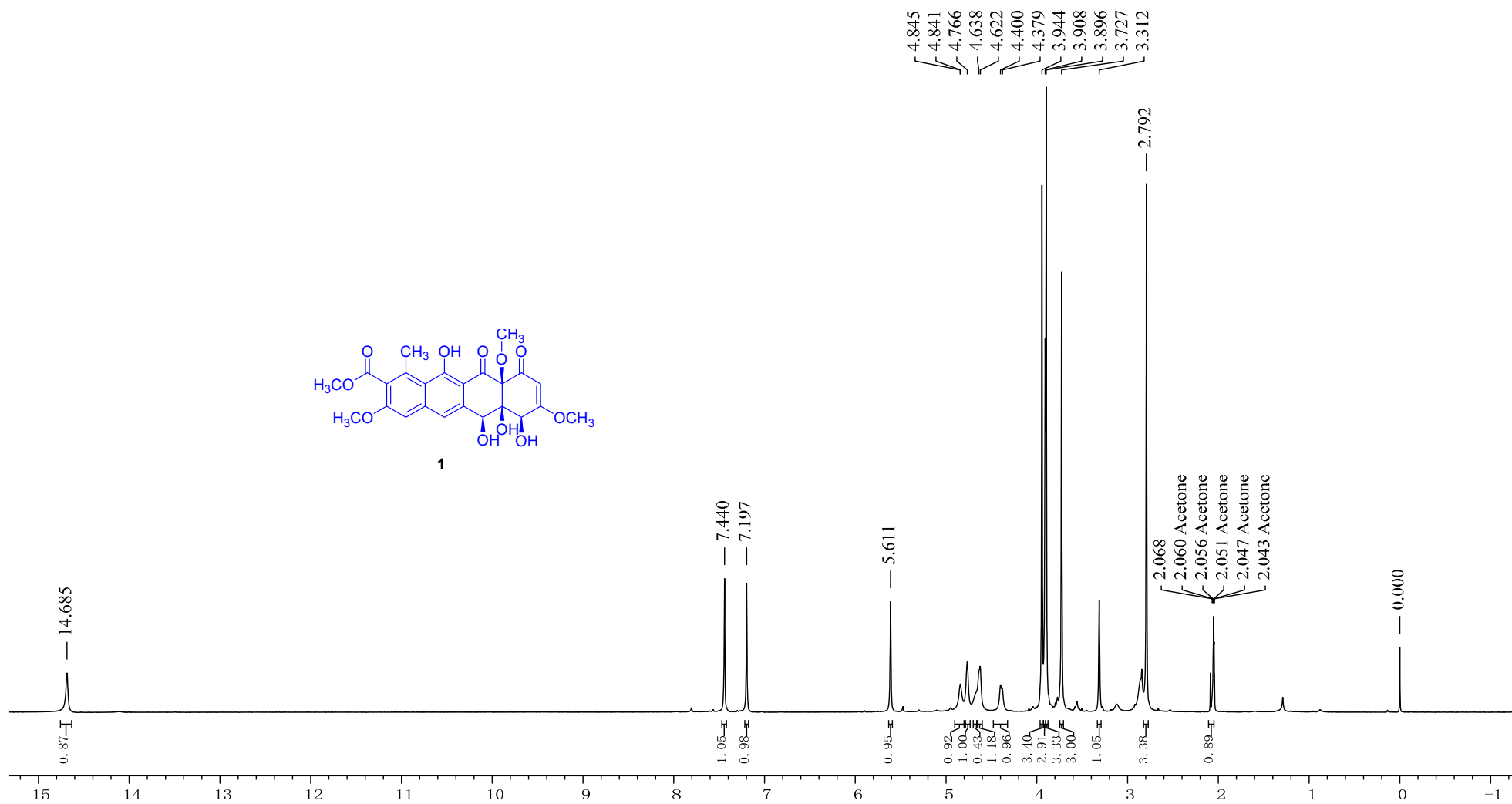


Figure S4. The ¹H NMR spectrum of saccharothrixone E (**1**) in acetone-*d*₆ (500 MHz).

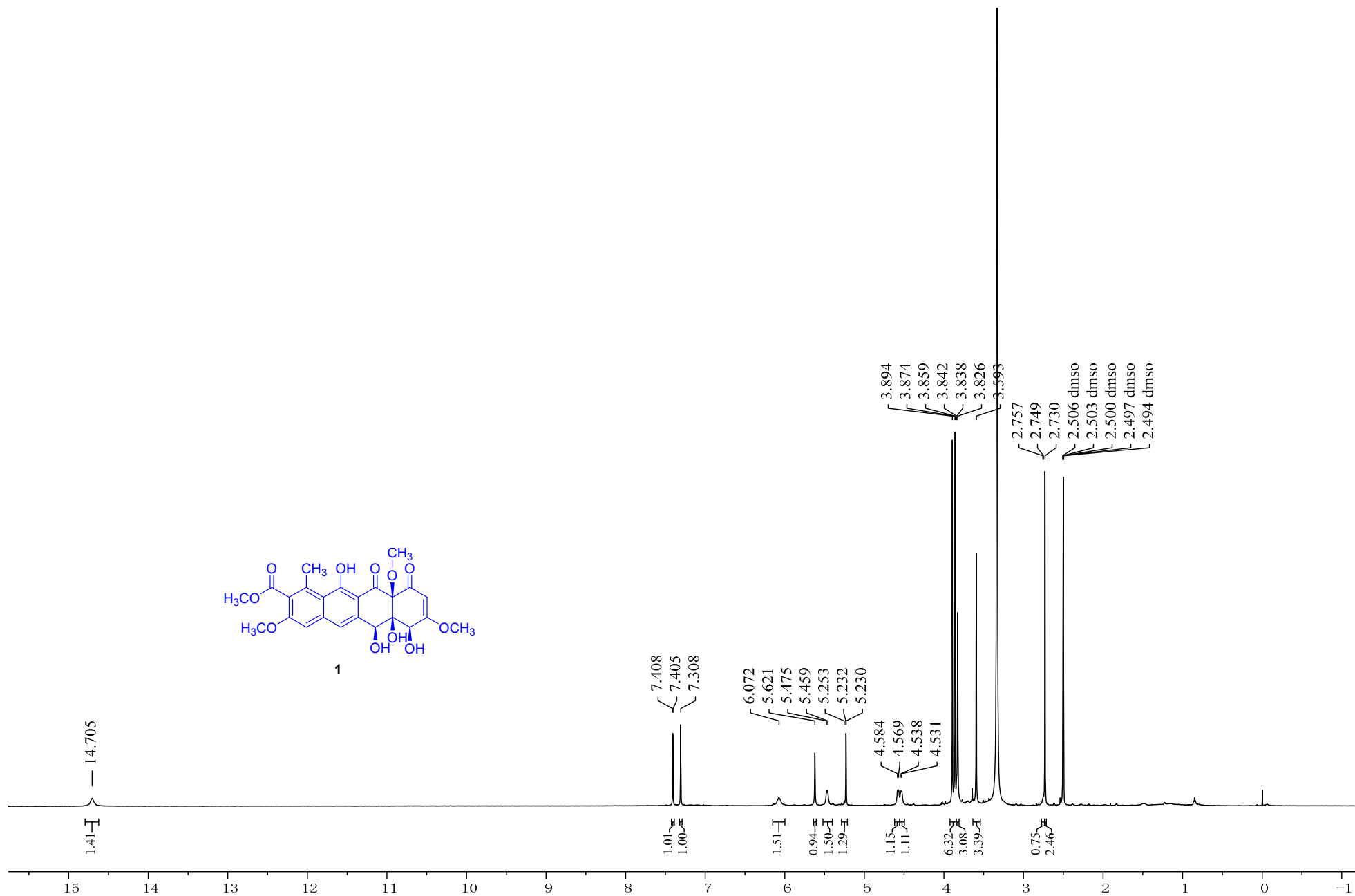


Figure S5. The ¹H NMR spectrum of saccharothrixone E (**1**) in DMSO-*d*₆ (600 MHz).

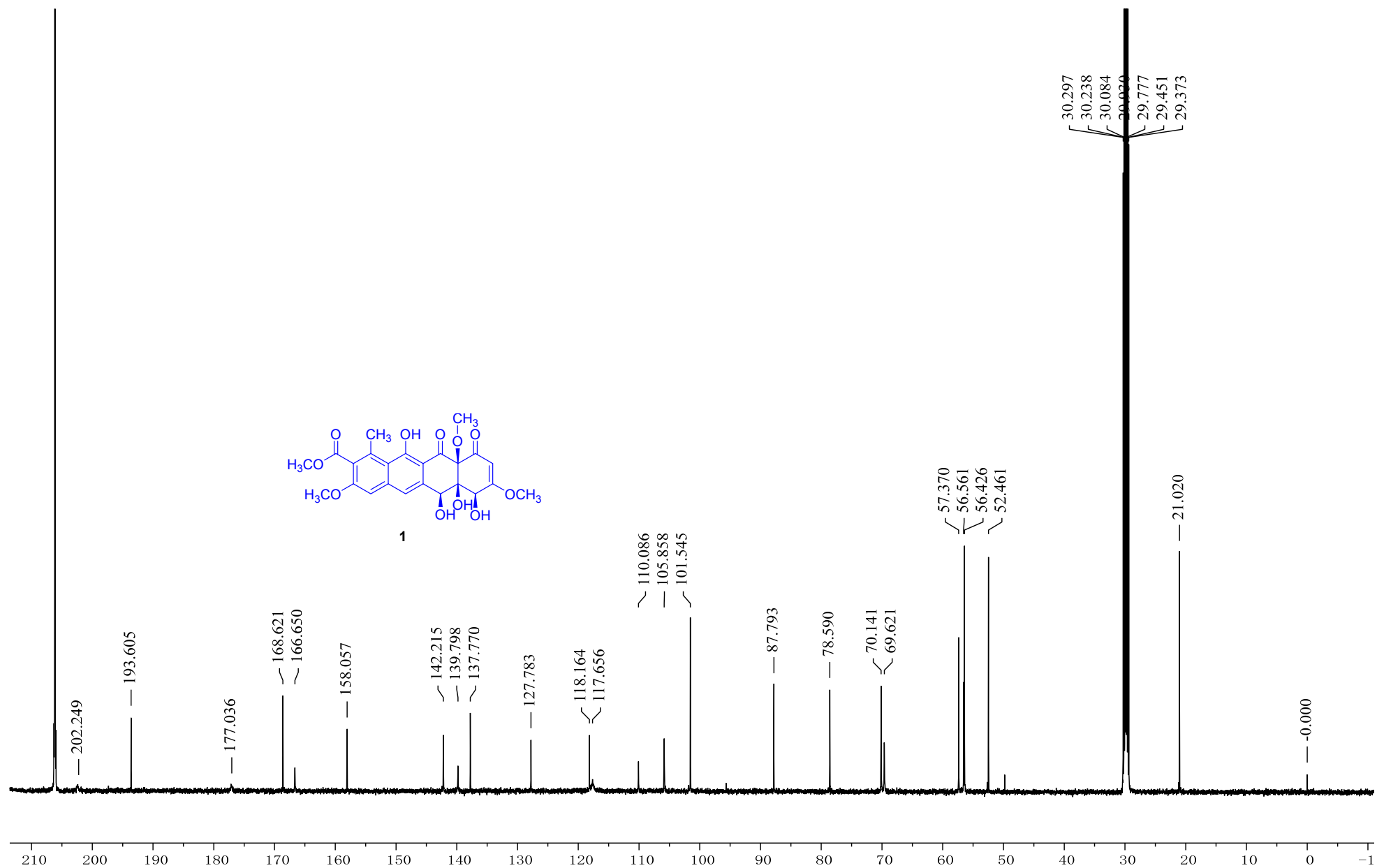


Figure S6. The ^{13}C NMR spectrum of saccharothrixone E (**1**) in acetone- d_6 (125 MHz).

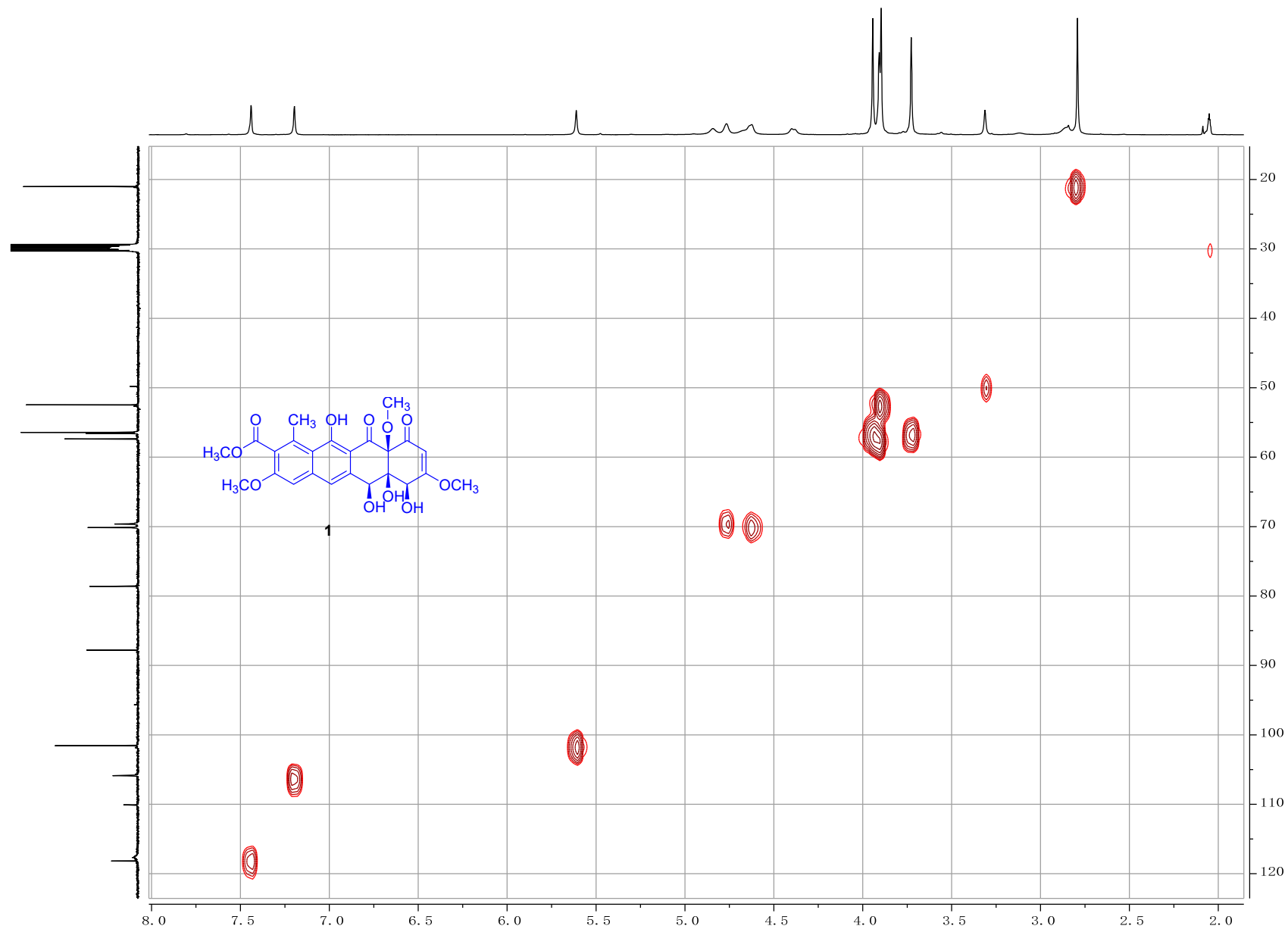


Figure S7. The HSQC spectrum of saccharothrixone E (**1**) in acetone-*d*₆ (500 MHz).

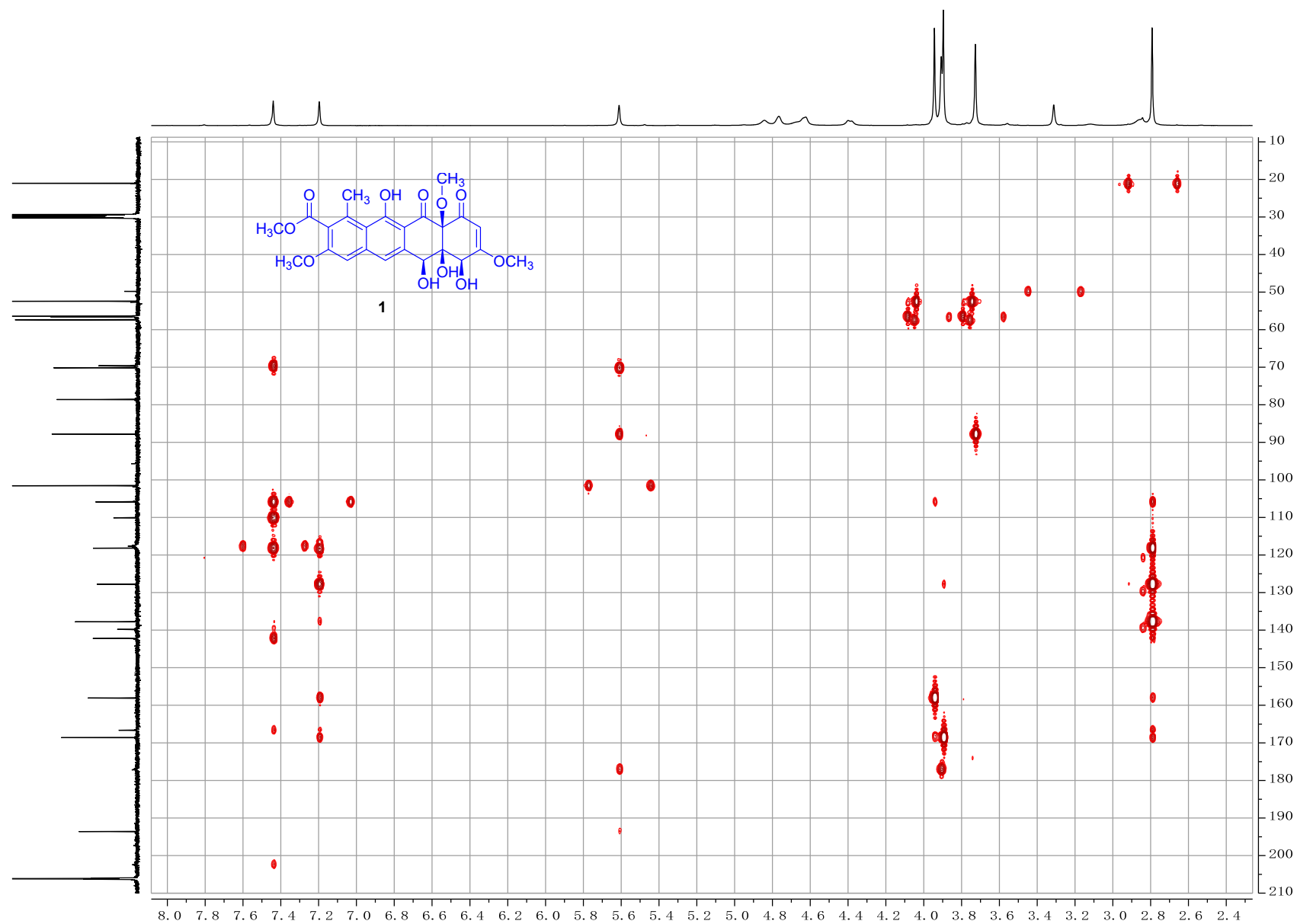


Figure S8. The HMBC spectrum of saccharothrixone E (**1**) in acetone- d_6 (500 MHz).

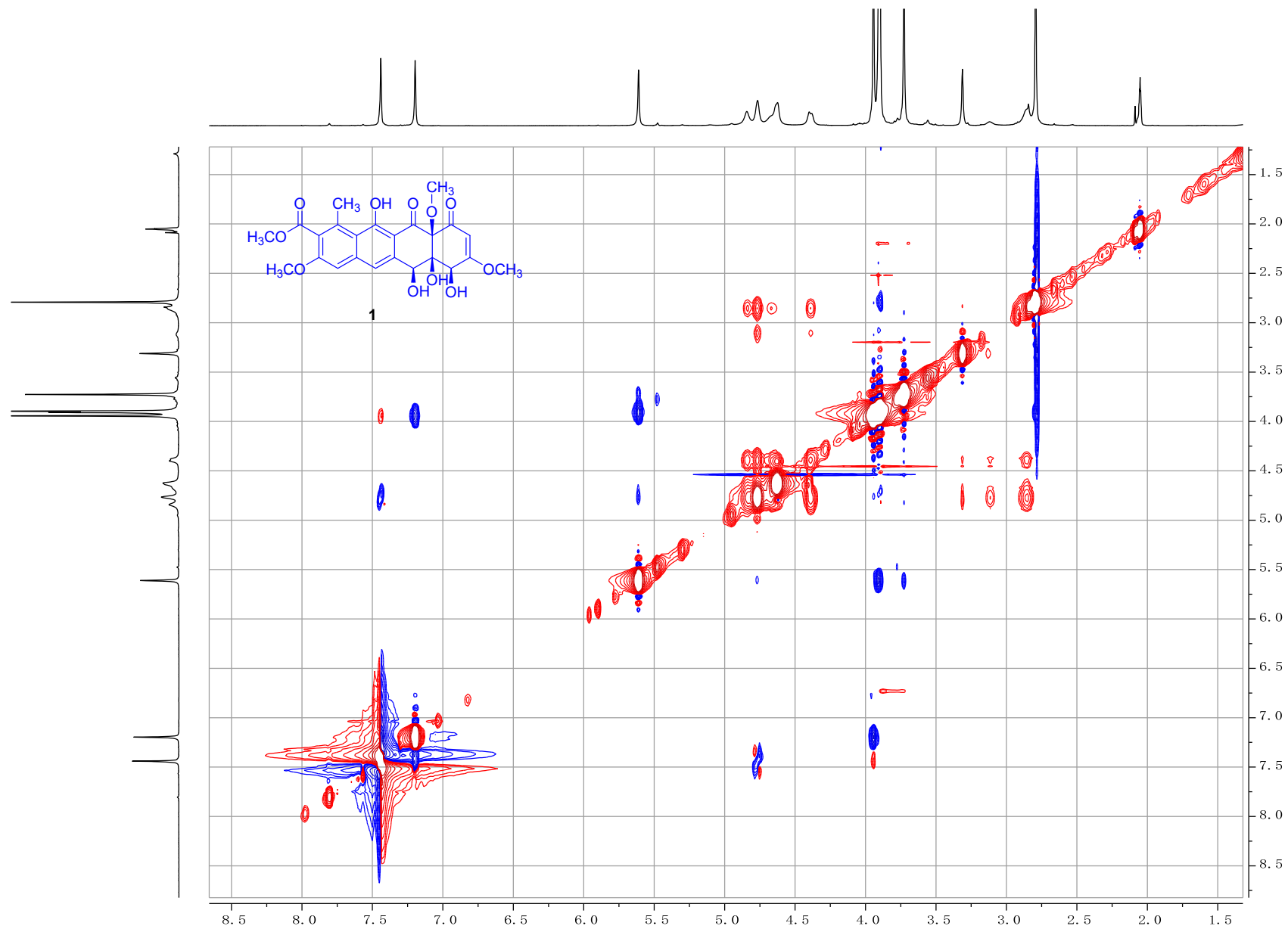


Figure S9. The ROESY spectrum of saccharothrixone E (**1**) in acetone- d_6 (500 MHz).

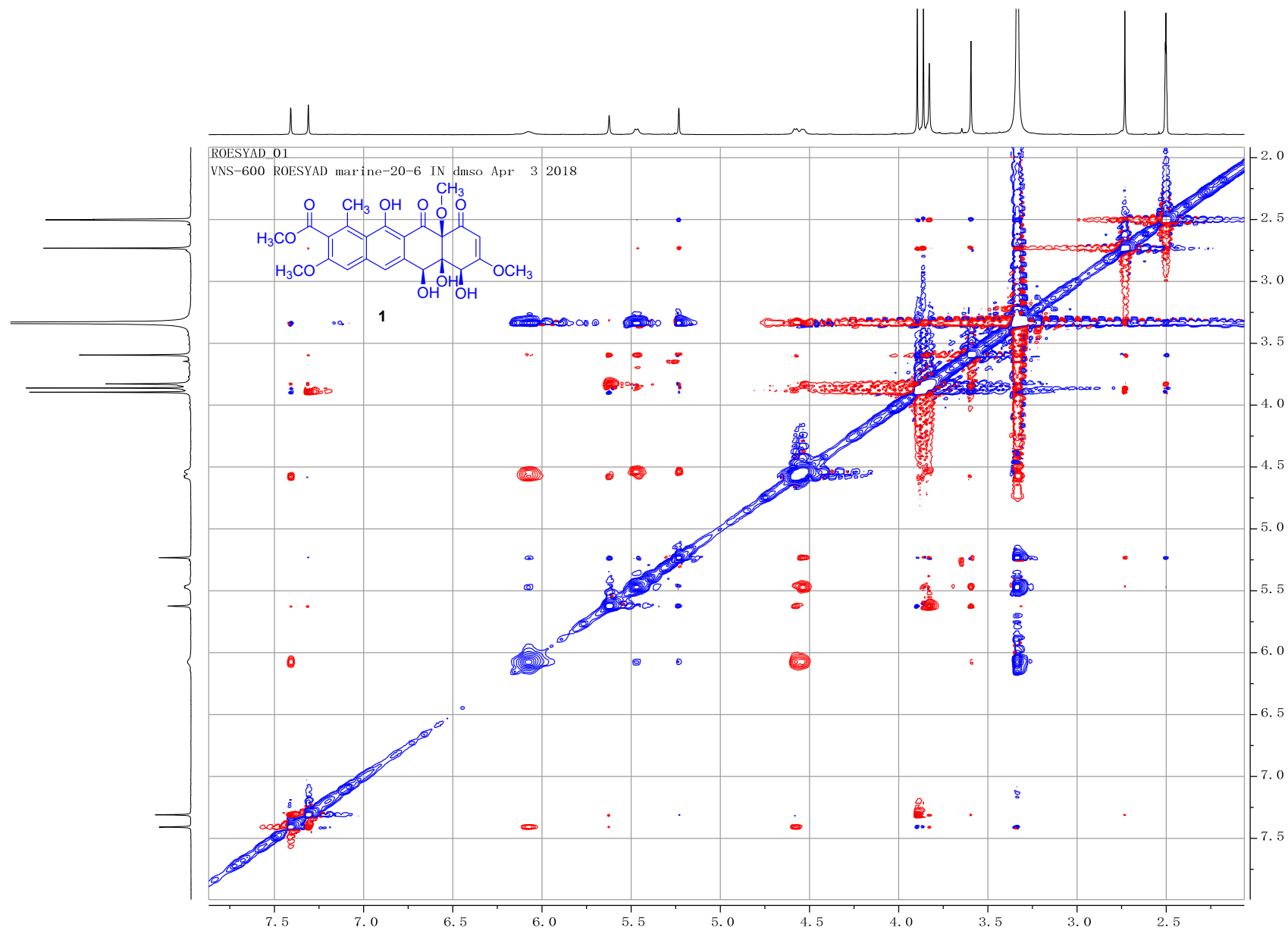


Figure S10. The ROESY spectrum of saccharothrixone E (**1**) in DMSO- d_6 (600 MHz).

marine-20-7 HRESI #13 RT: 0.20 AV: 1 NL: 2.80E6
T: FTMS + c ESI Full ms [100.00-1000.00]

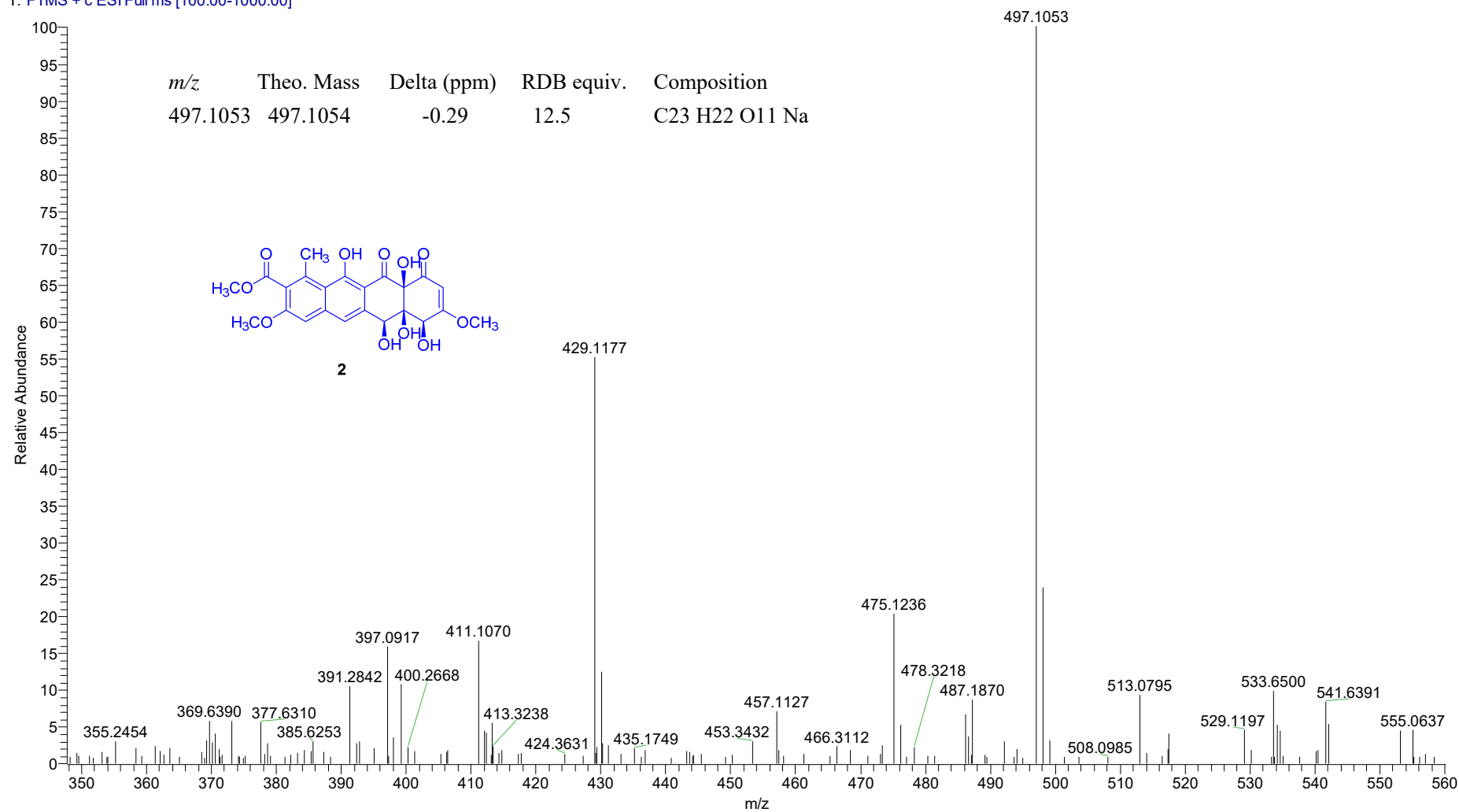


Figure S11. The (+)-HRESIMS spectrum of saccharothrixone F (**2**).

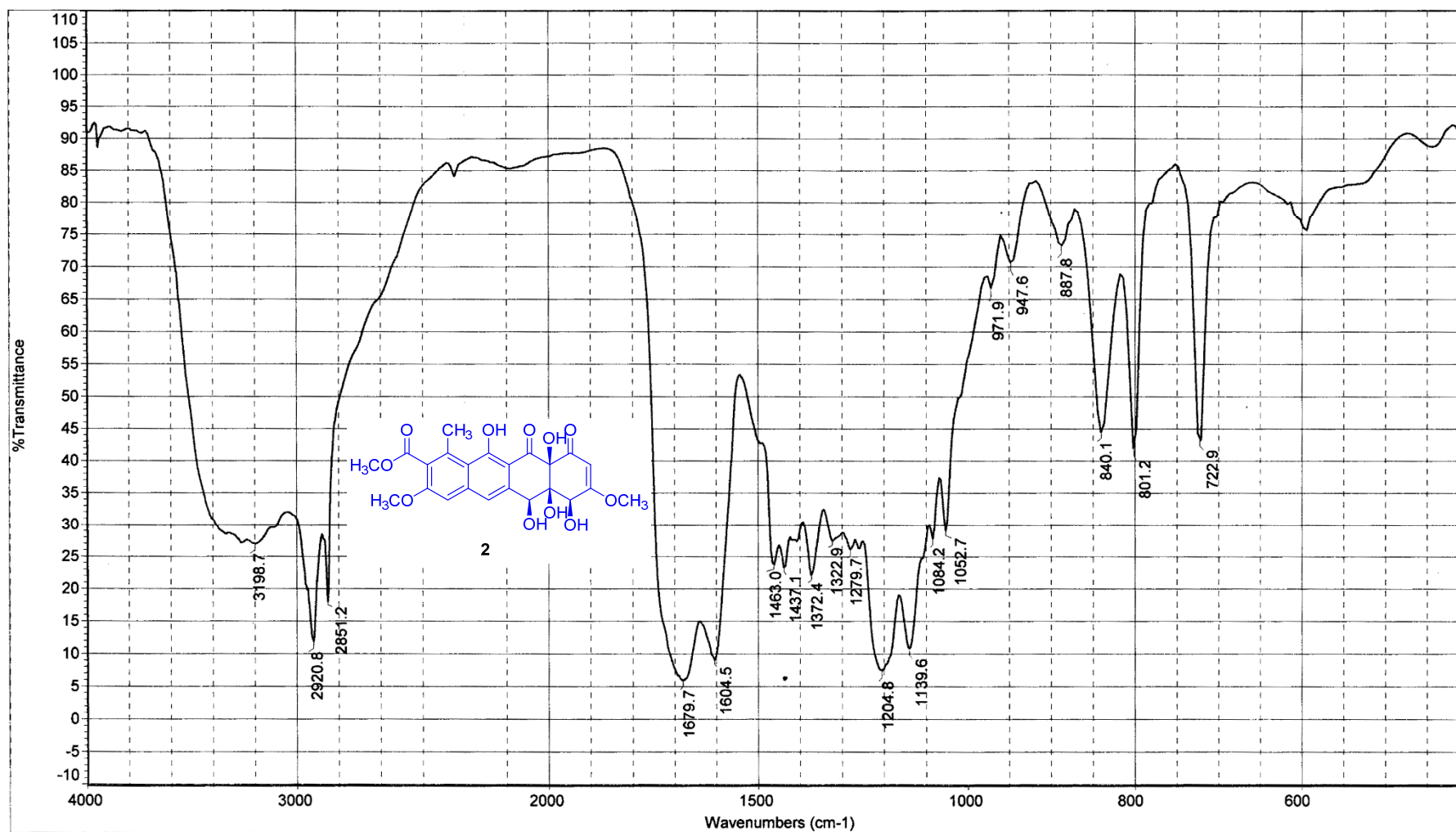


Figure S12. The IR spectrum of saccharothrixone F (2).

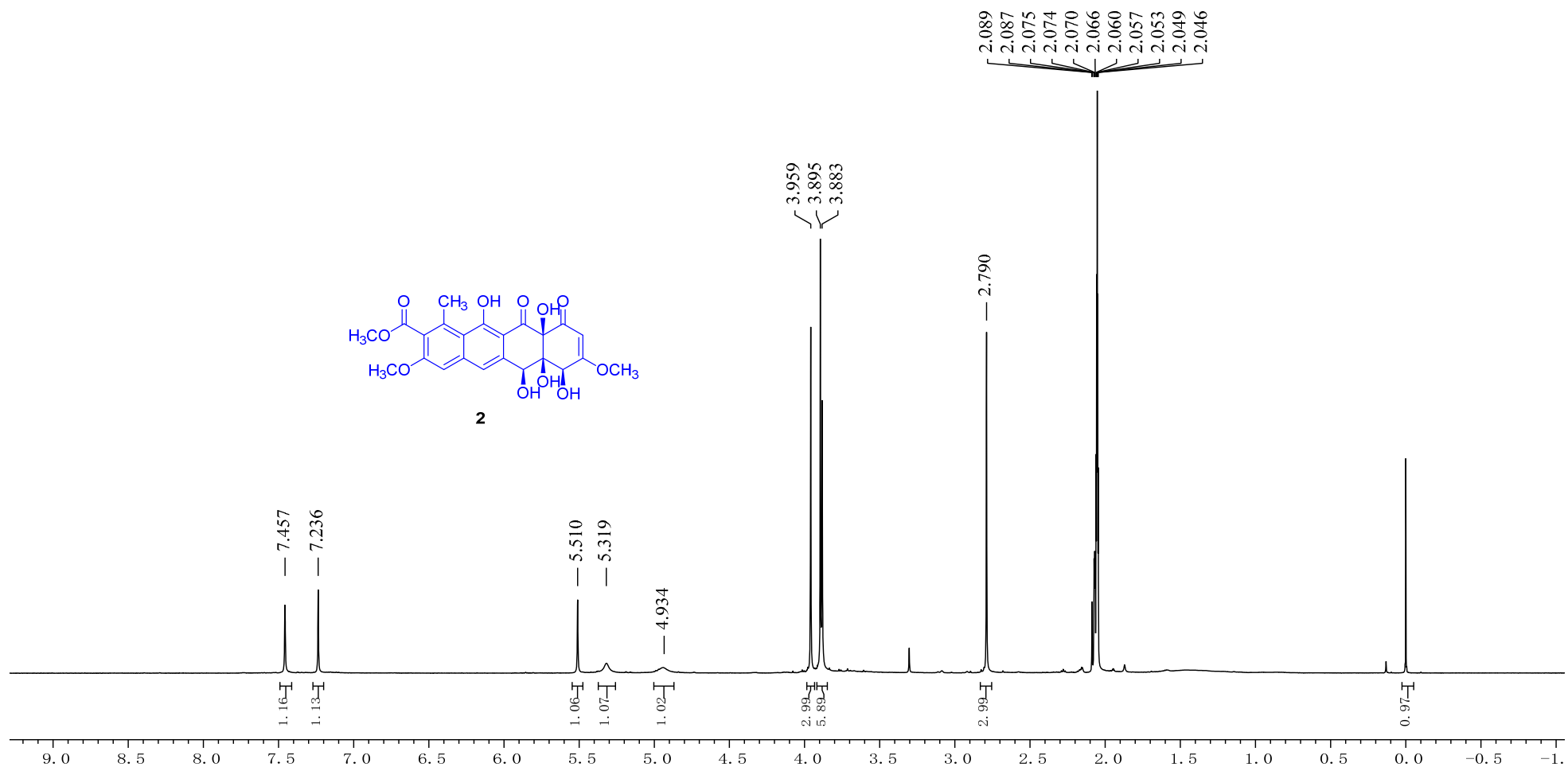


Figure S13. The ^1H NMR spectrum of saccharothrixone F (**2**) in acetone- d_6 (600 MHz).

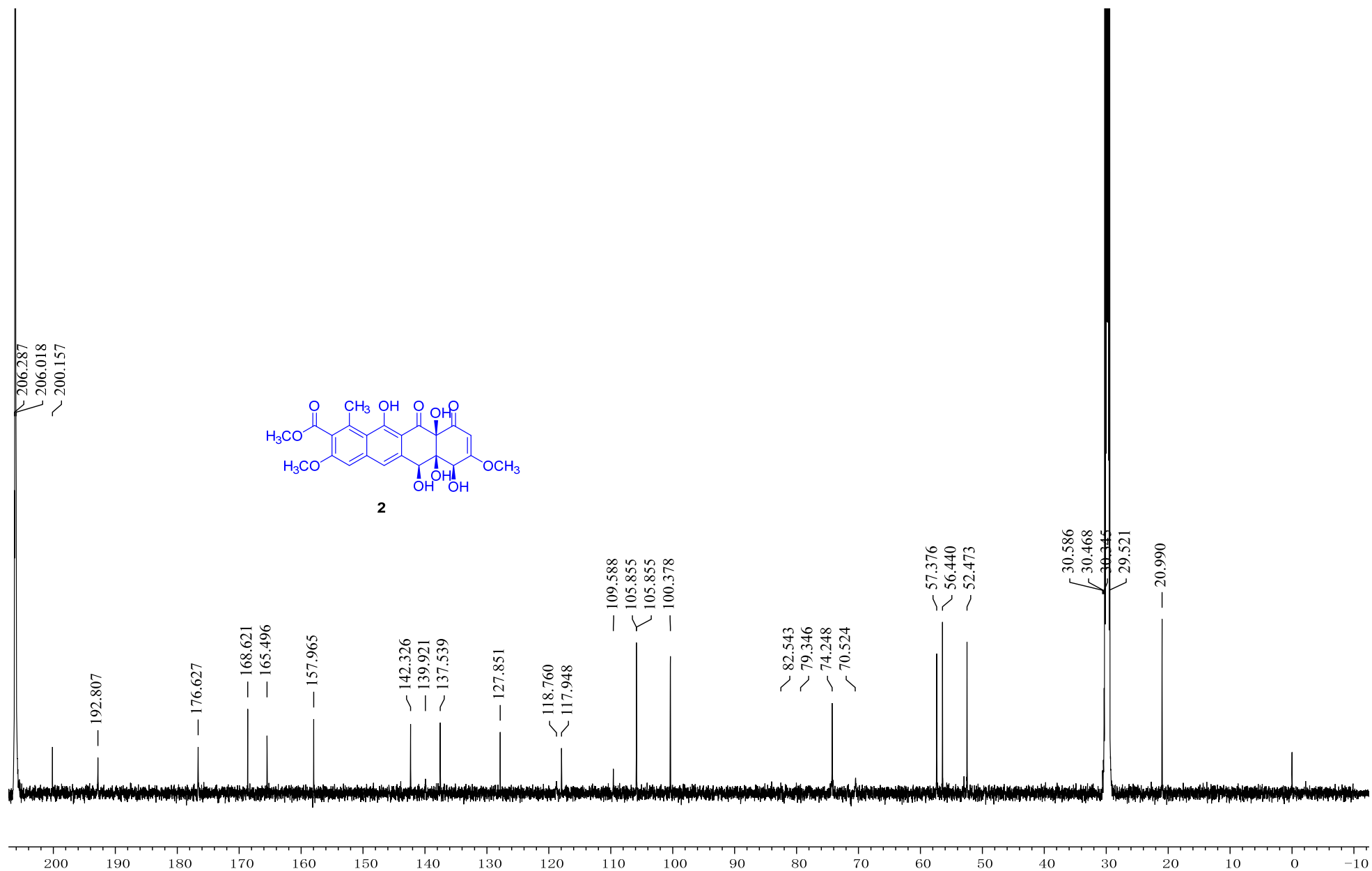


Figure S14. The ^{13}C NMR spectrum of saccharothrixone F (**2**) in acetone- d_6 (150 MHz).

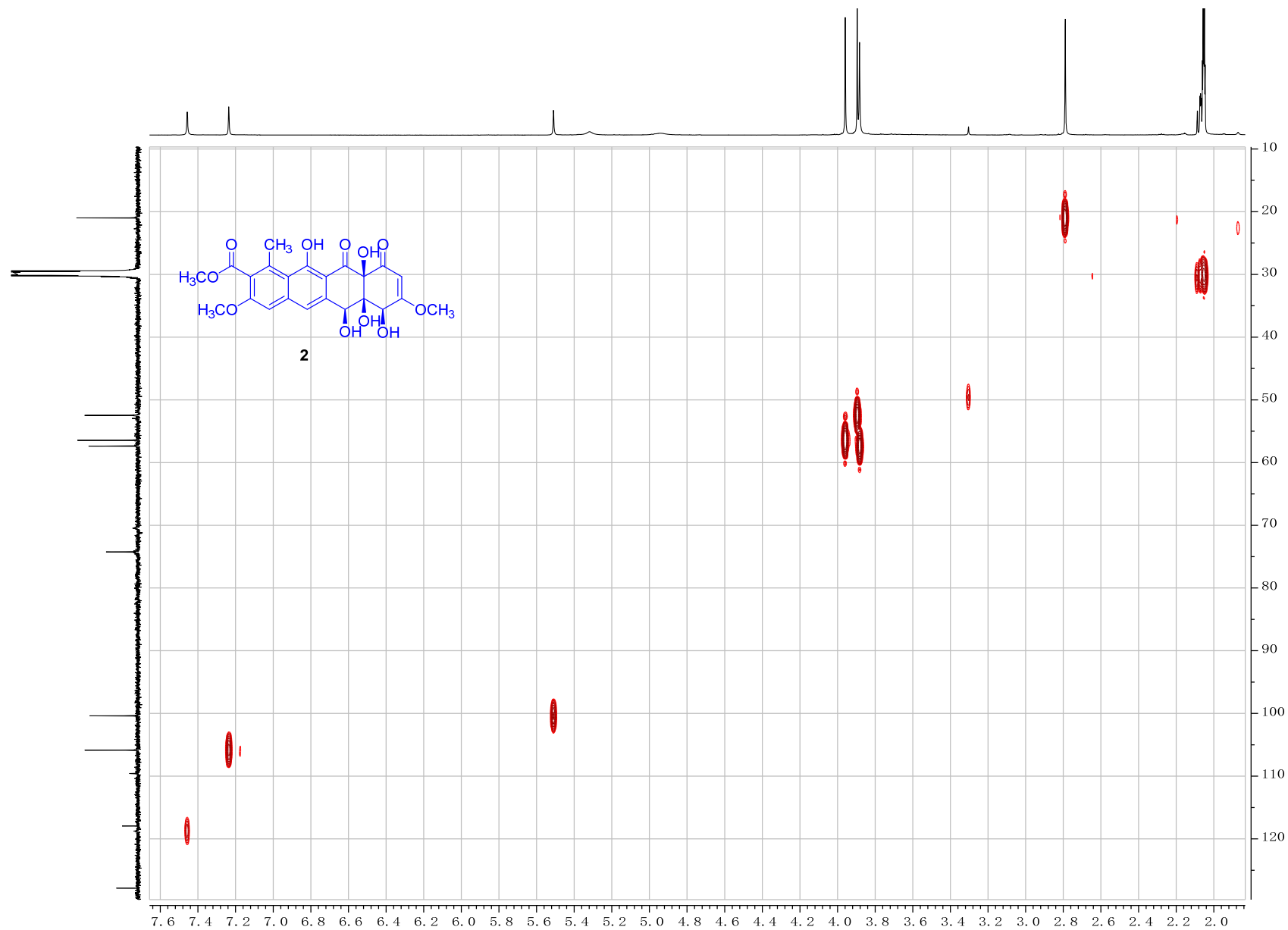


Figure S15. The HSQC spectrum of saccharothrixone F (**2**) in acetone- d_6 (600 MHz).

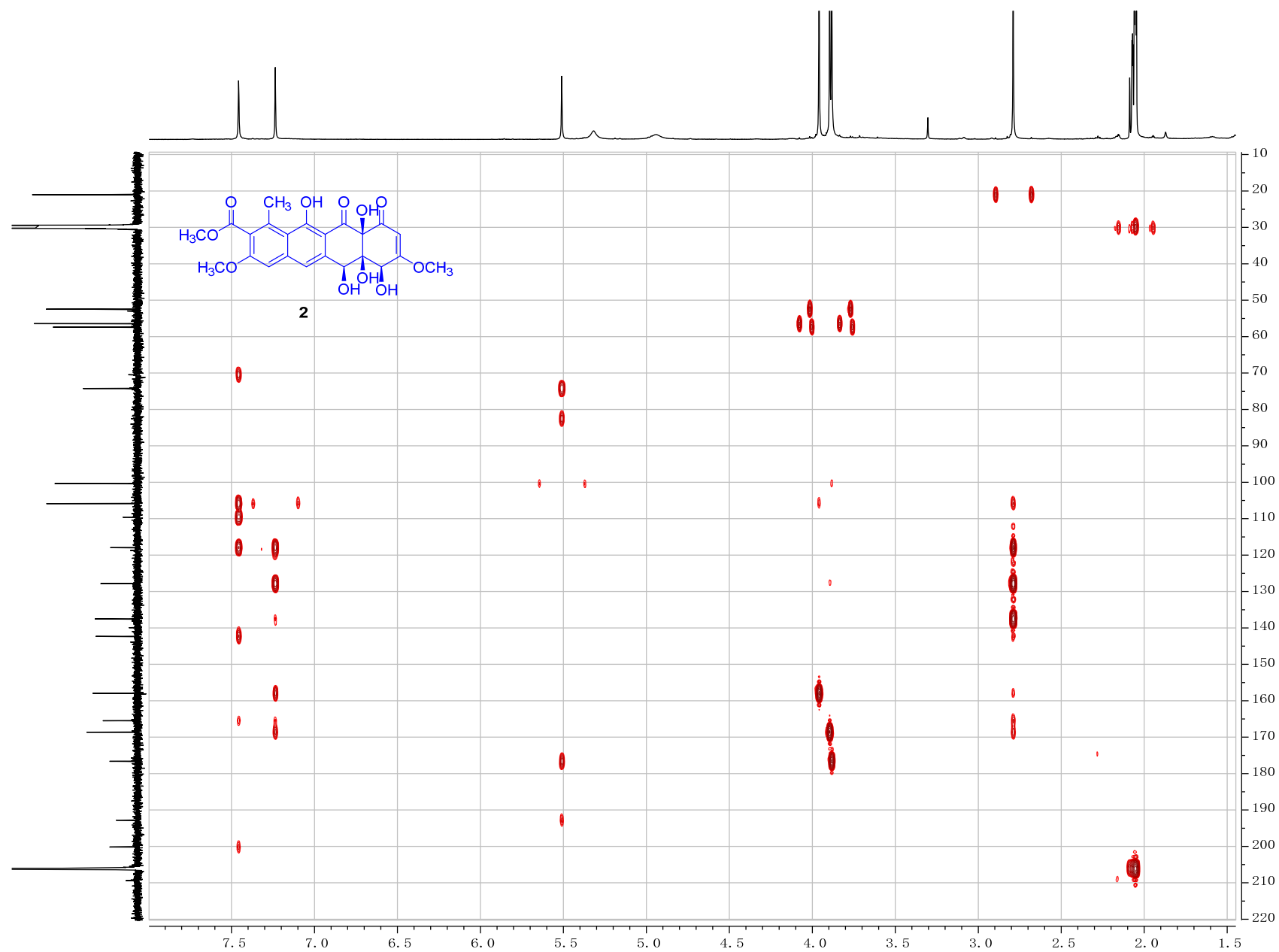


Figure S16. The HMBC spectrum of saccharothrixone F (**2**) in acetone- d_6 (600 MHz).

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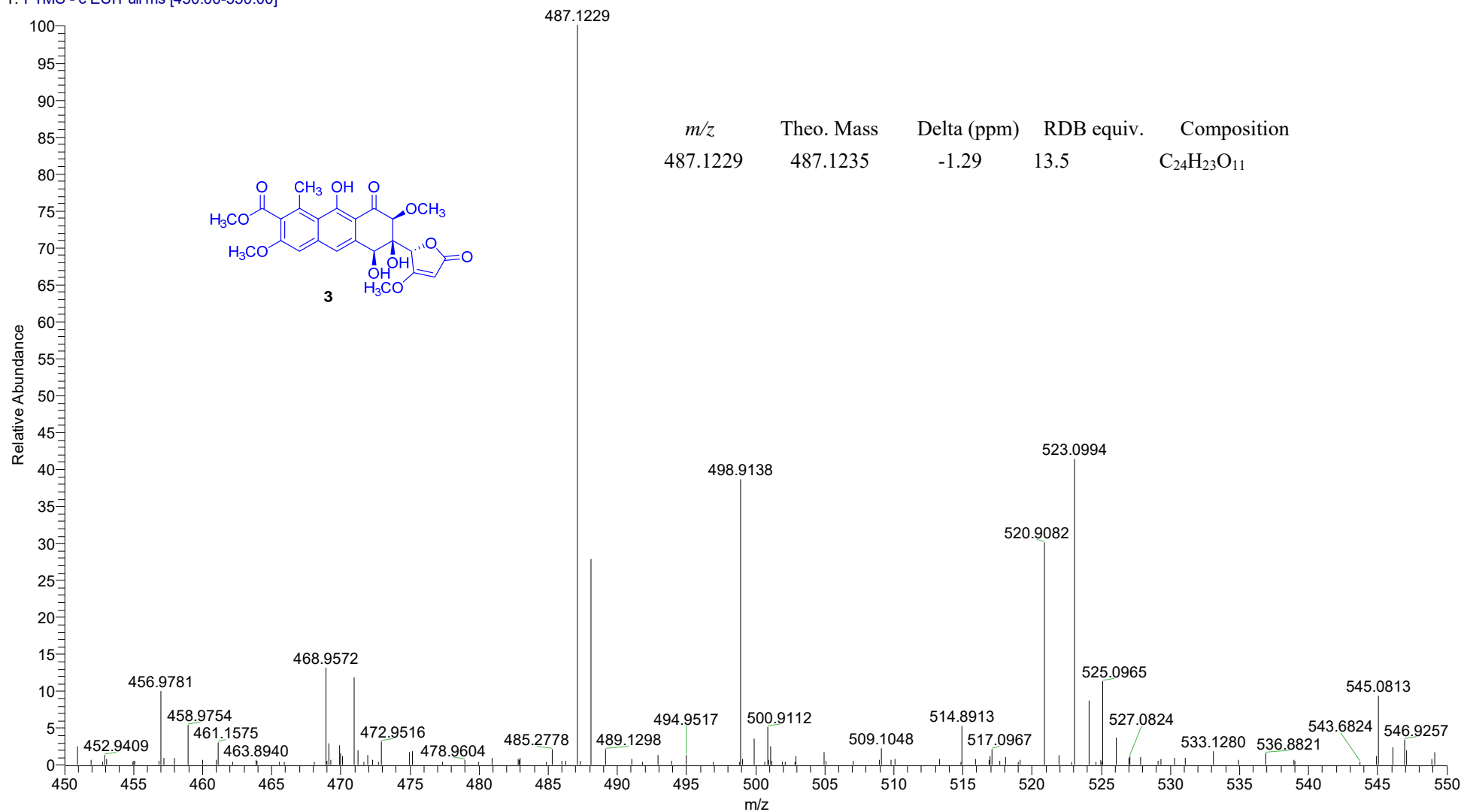


Figure S17. The (–)-HRESIMS spectrum of saccharothrixone G (3).

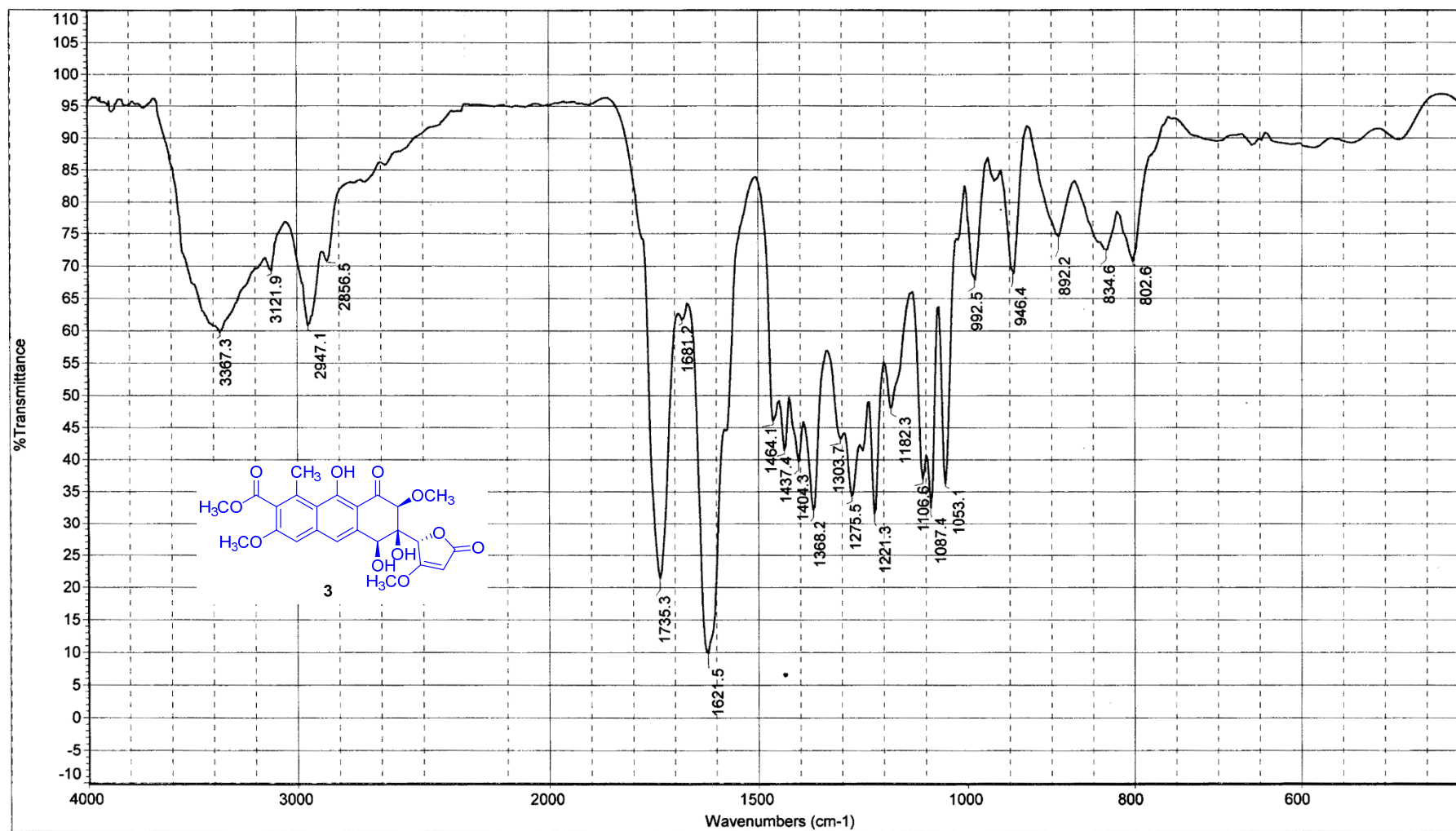


Figure S18. The IR spectrum of saccharothrixone G (**3**).

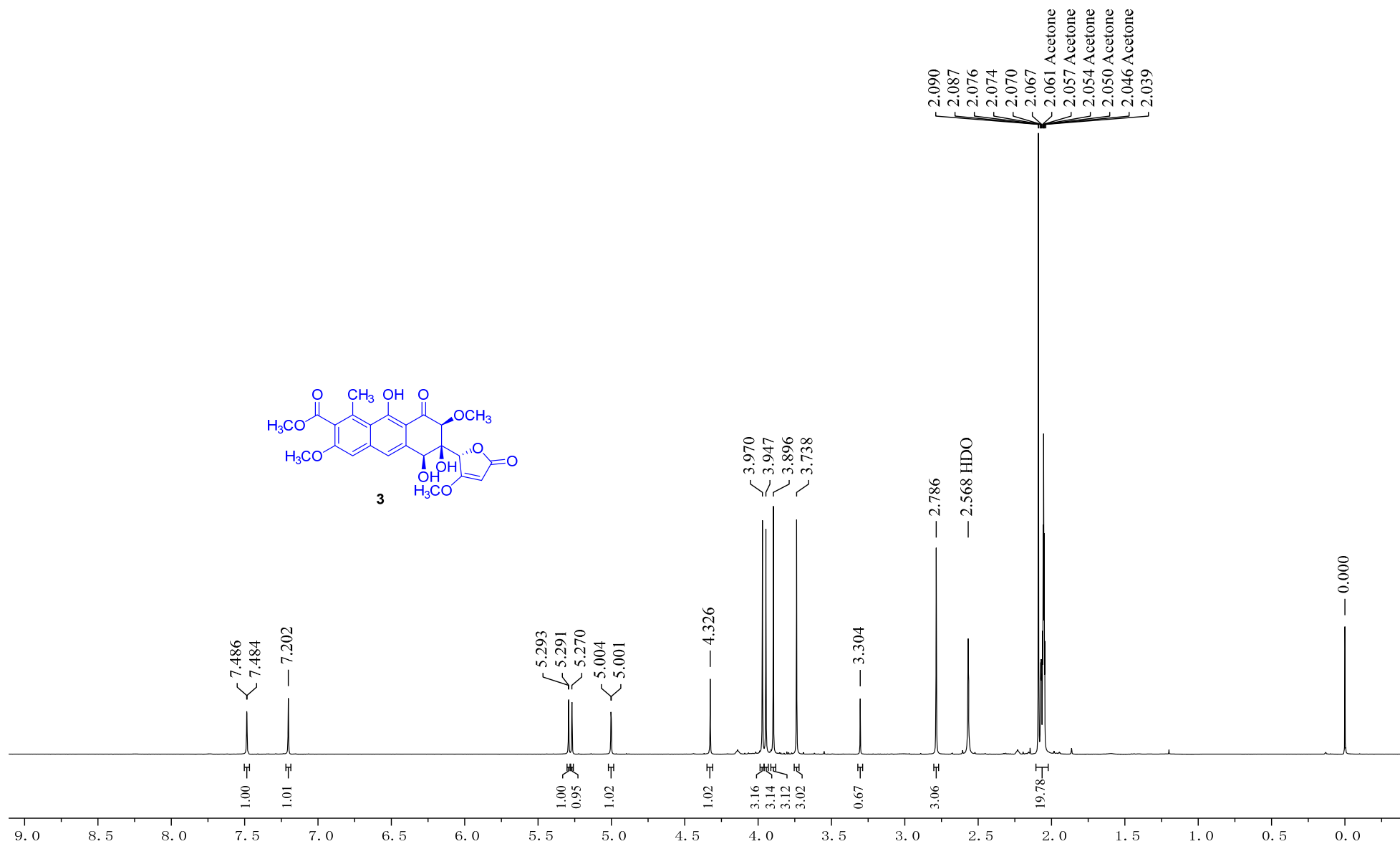


Figure S19. The ^1H NMR spectrum of saccharothrixone G (**3**) in acetone- d_6 (600 MHz).

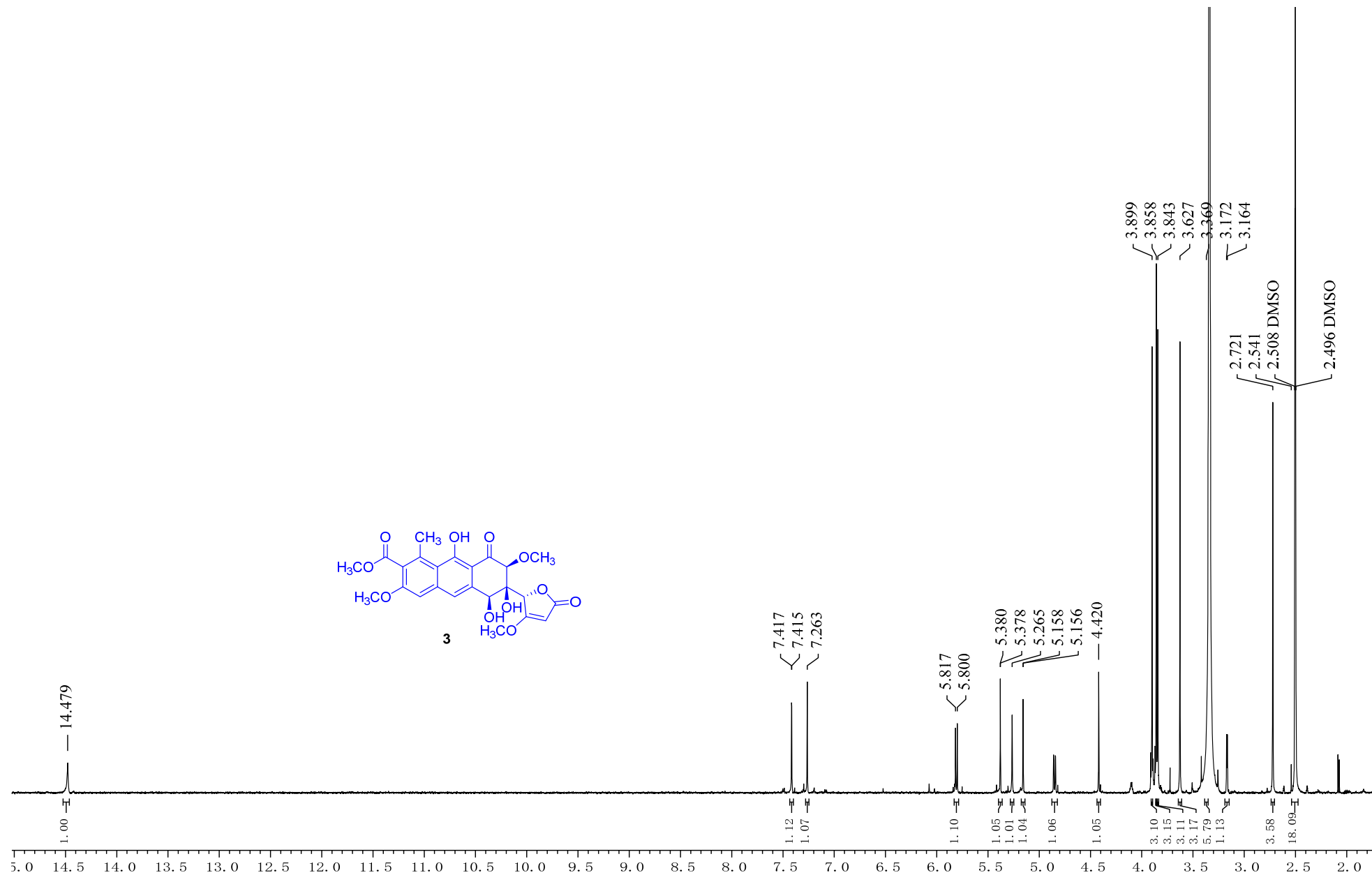


Figure S20. The ¹H NMR spectrum of saccharothrixone G (3) in DMSO-*d*₆ (600 MHz).

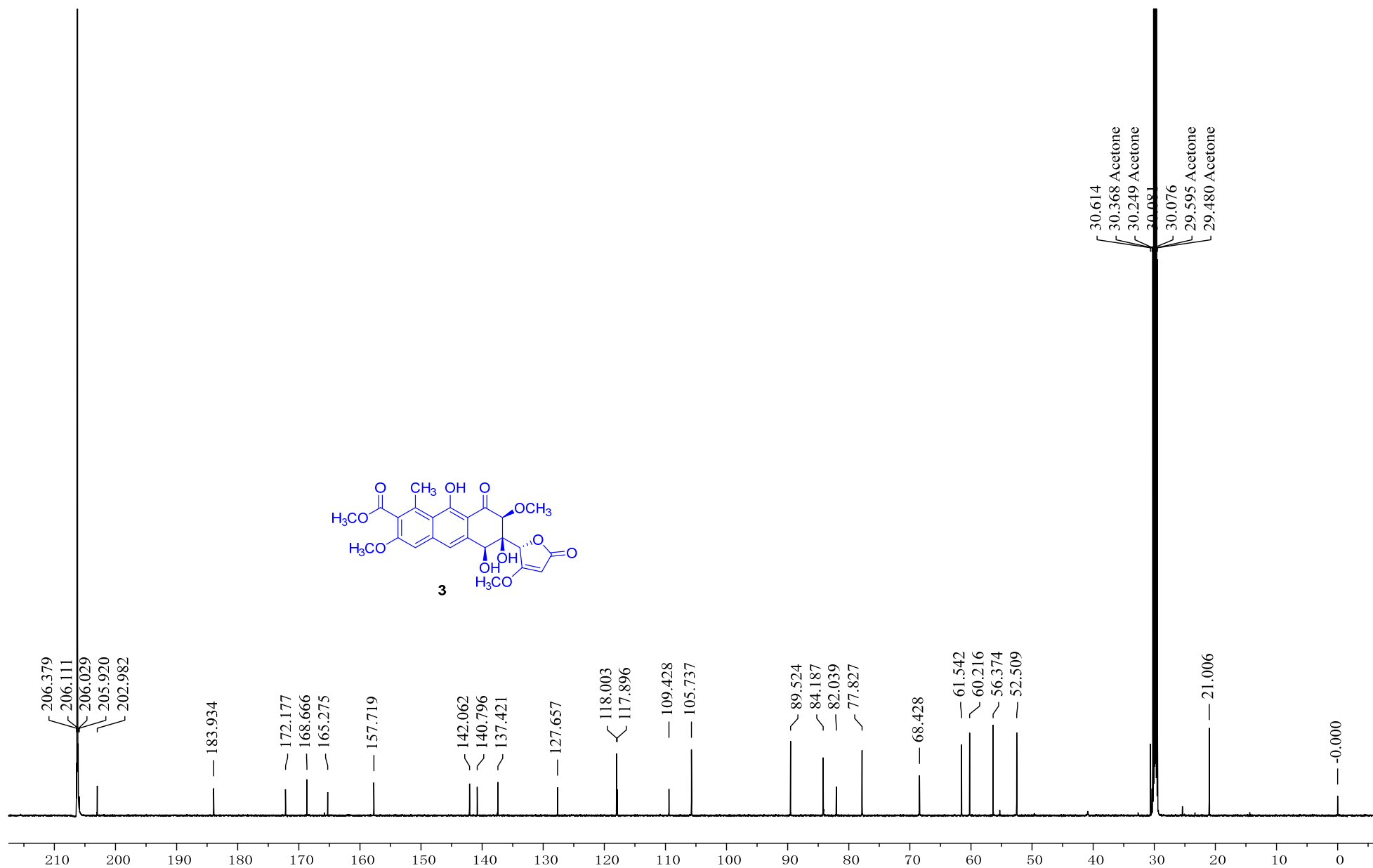


Figure S21. The ^{13}C NMR spectrum of saccharothrixone G (**3**) in acetone- d_6 (150 MHz).

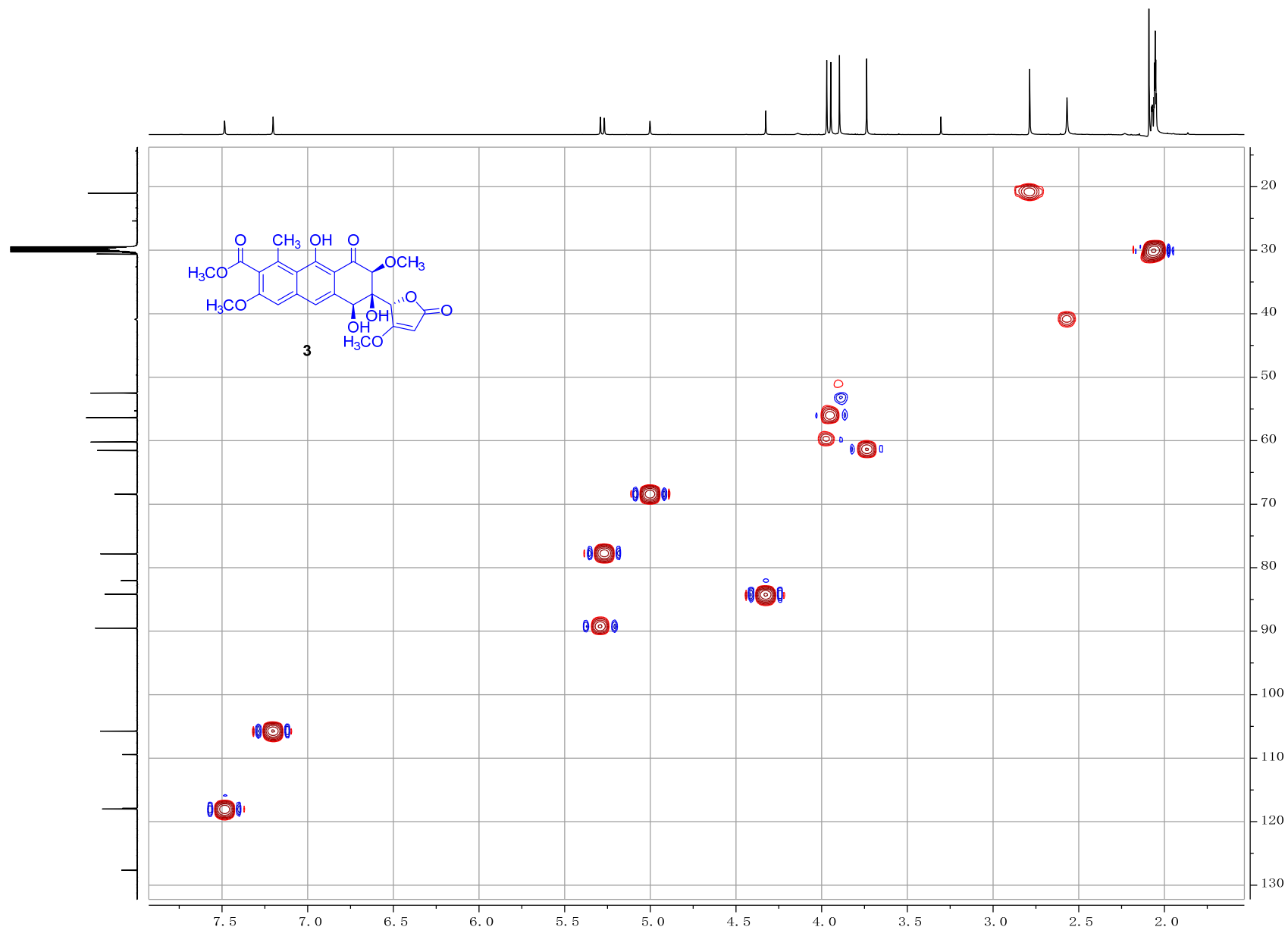


Figure S22. The HSQC spectrum of saccharothrixone G (**3**) in acetone- d_6 (600 MHz).

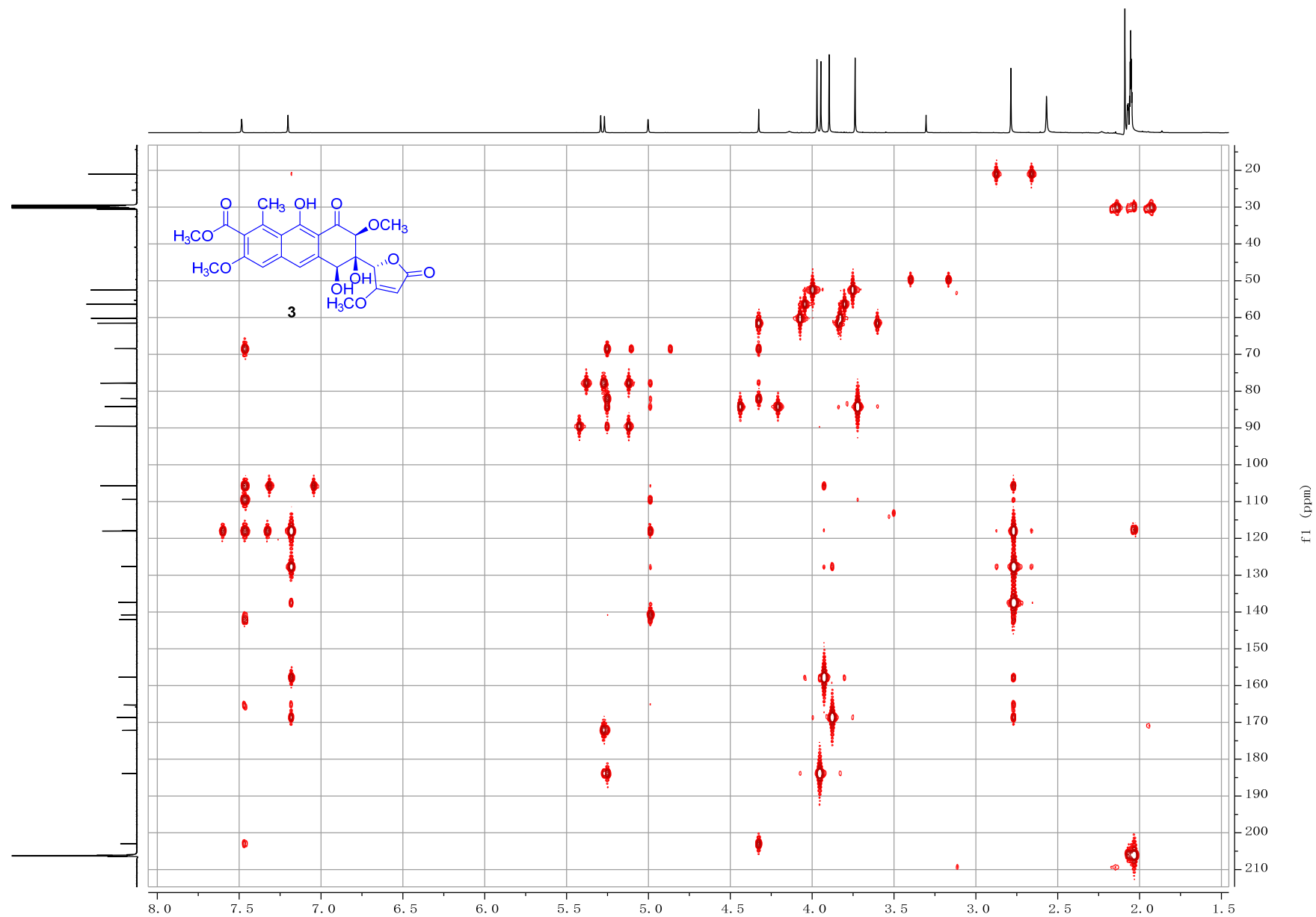


Figure S23. The HMBC spectrum of saccharothrixone G (**3**) in acetone- d_6 (600 MHz).

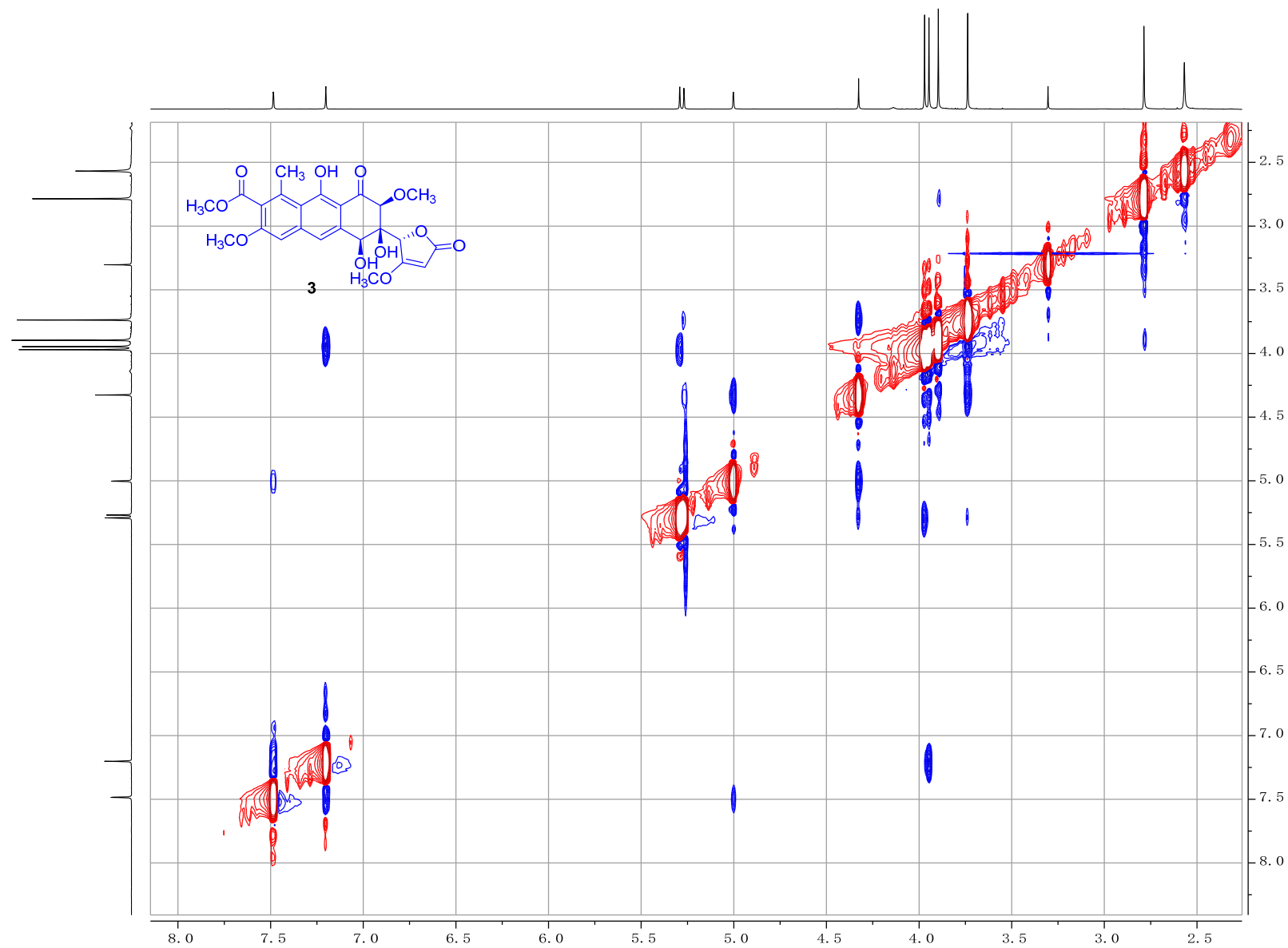


Figure S24. The ROESY spectrum of saccharothrixone G (**3**) in acetone-*d*₆ (600 MHz).

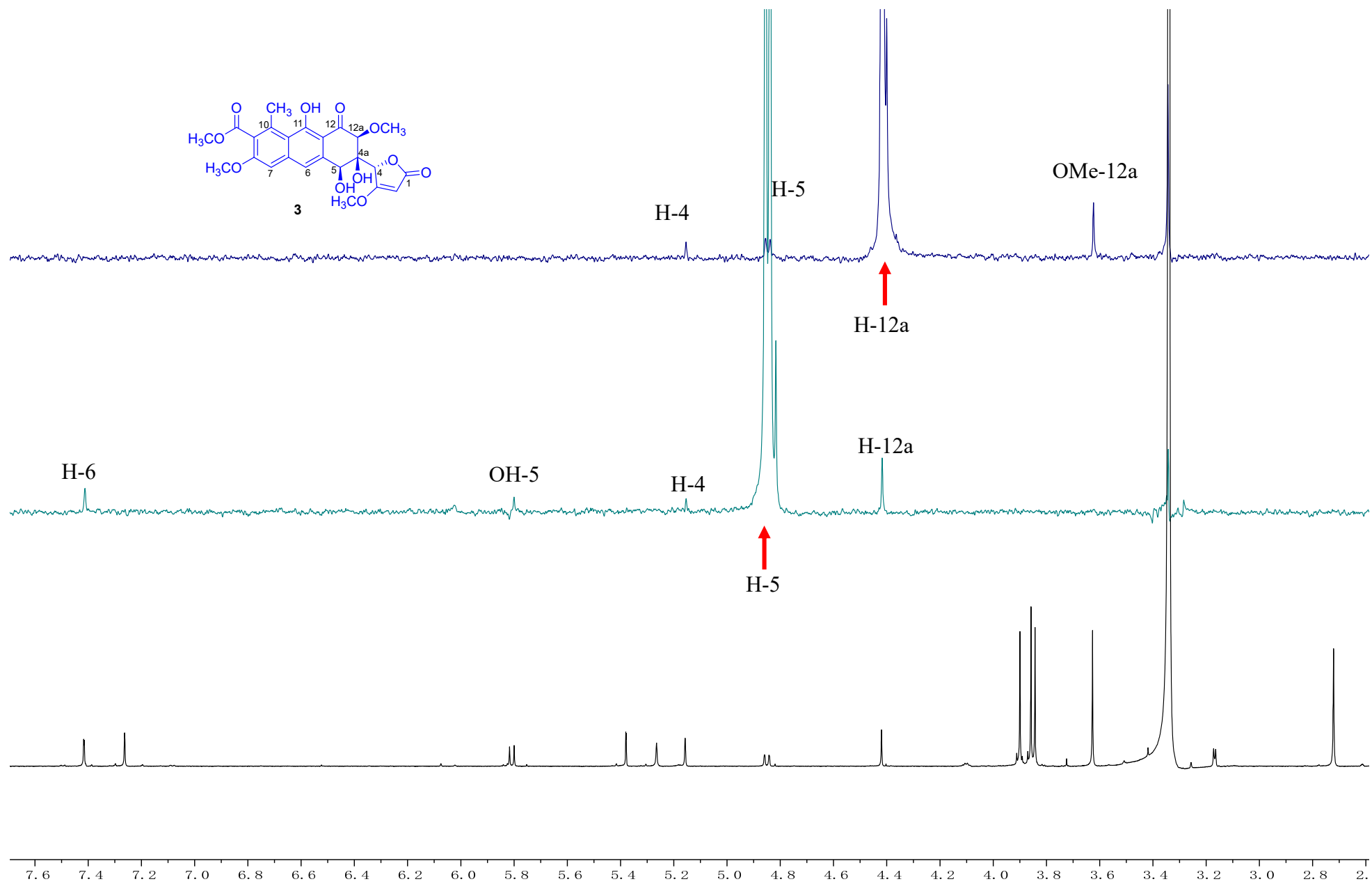


Figure S25. The 1D NOE spectrum of saccharothrixone G (**3**) in DMSO- d_6 (600 MHz).

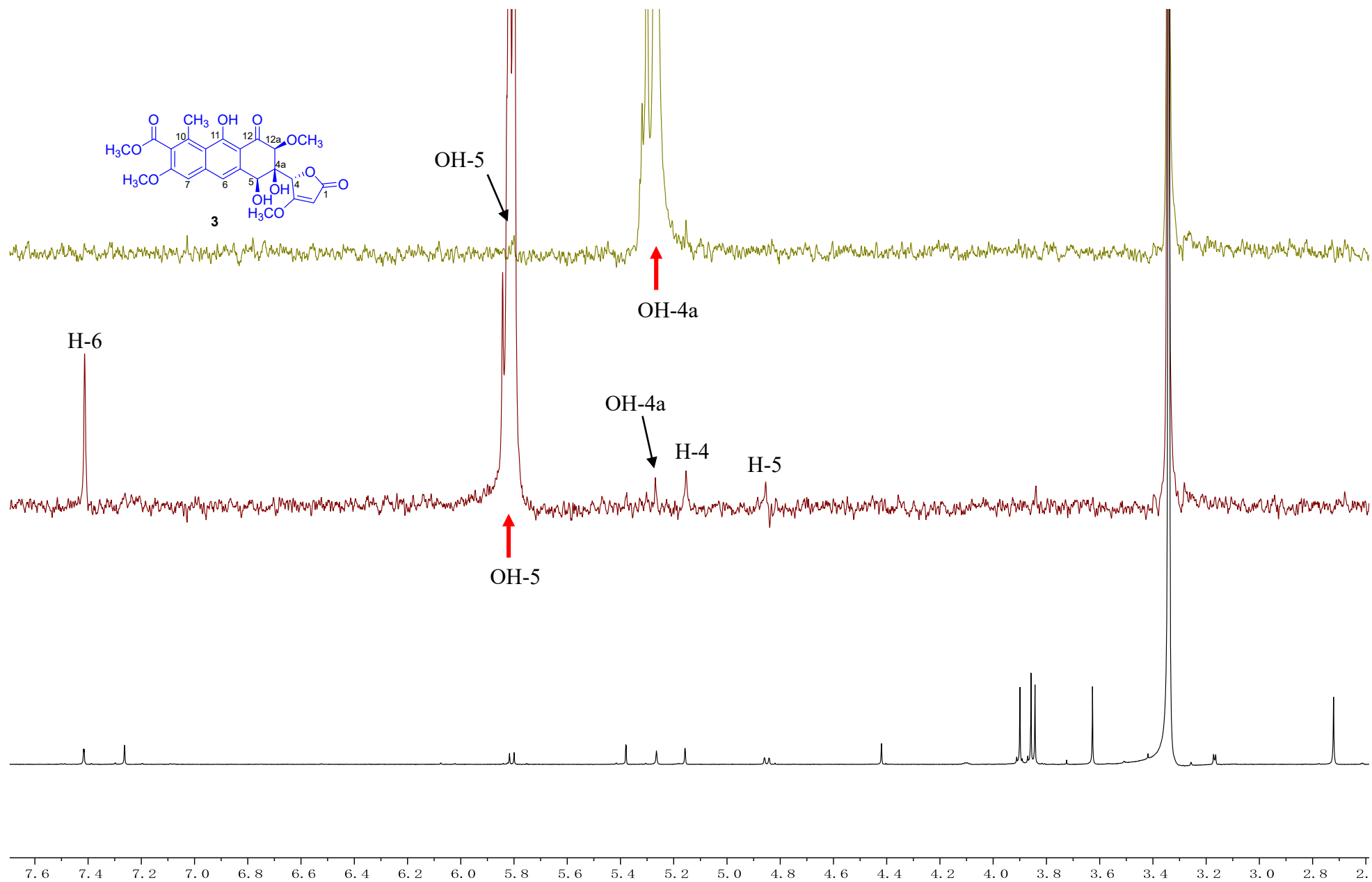


Figure S26. The 1D NOE spectrum of saccharothrixone G (**3**) in DMSO-d₆ (600 MHz).

marine-20-12_-#52 RT: 1.16 AV: 1 NL: 1.73E5
T: FTMS - c ESI Full ms [300.00-1000.00]

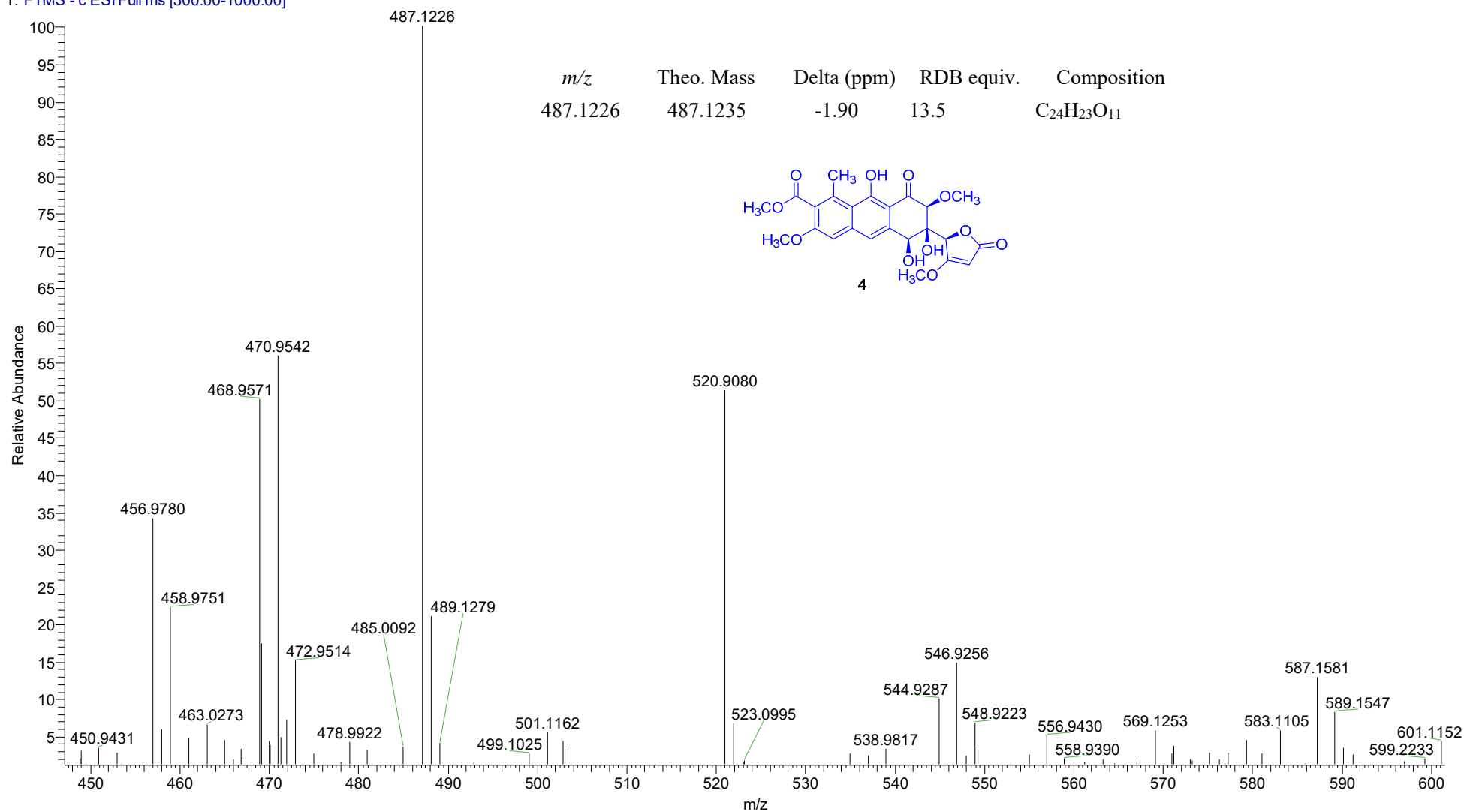


Figure S27. The (–)-HRESIMS spectrum of saccharothrixone H (4).

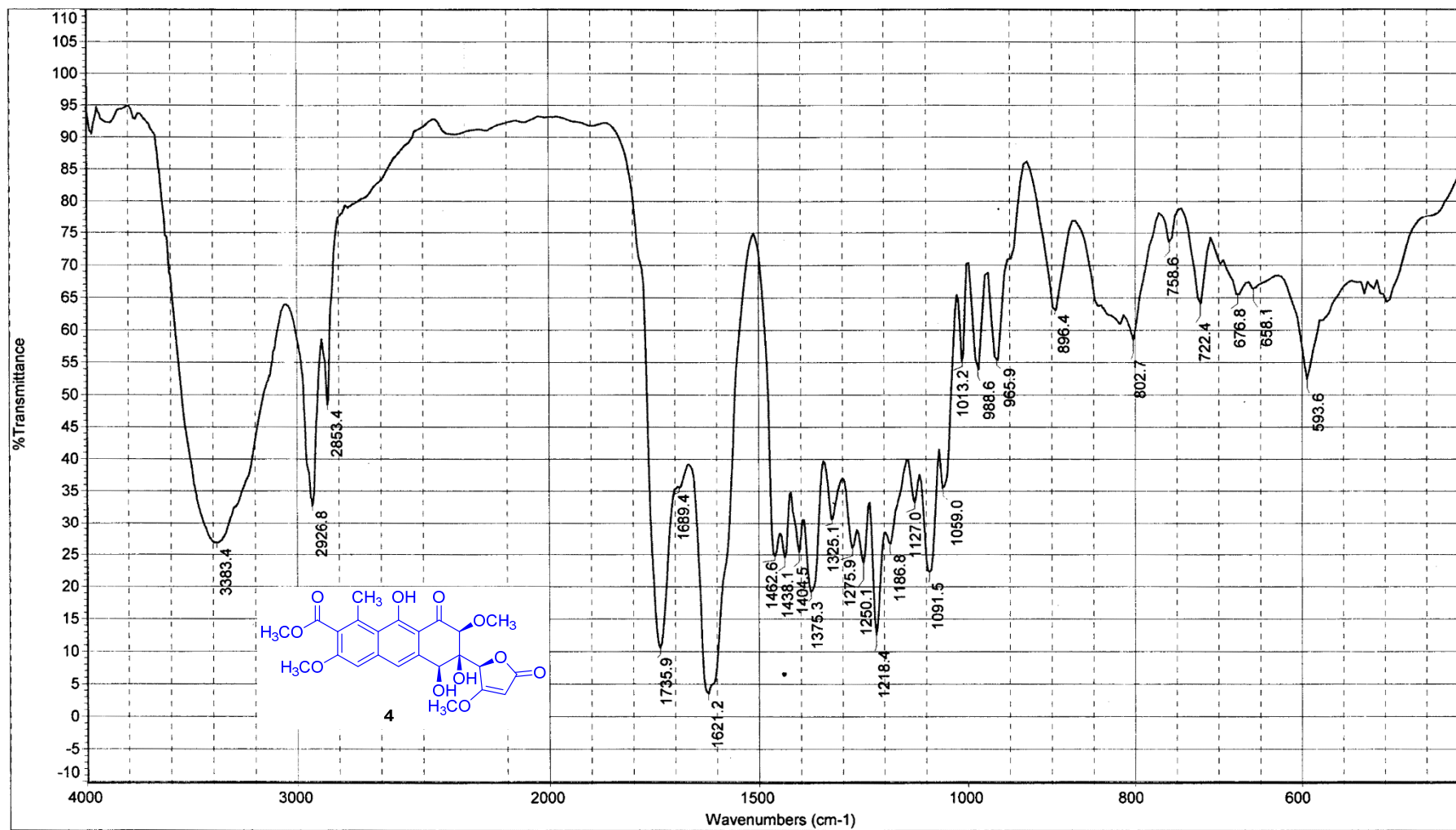


Figure S28. The IR spectrum of saccharothrixone H (4).

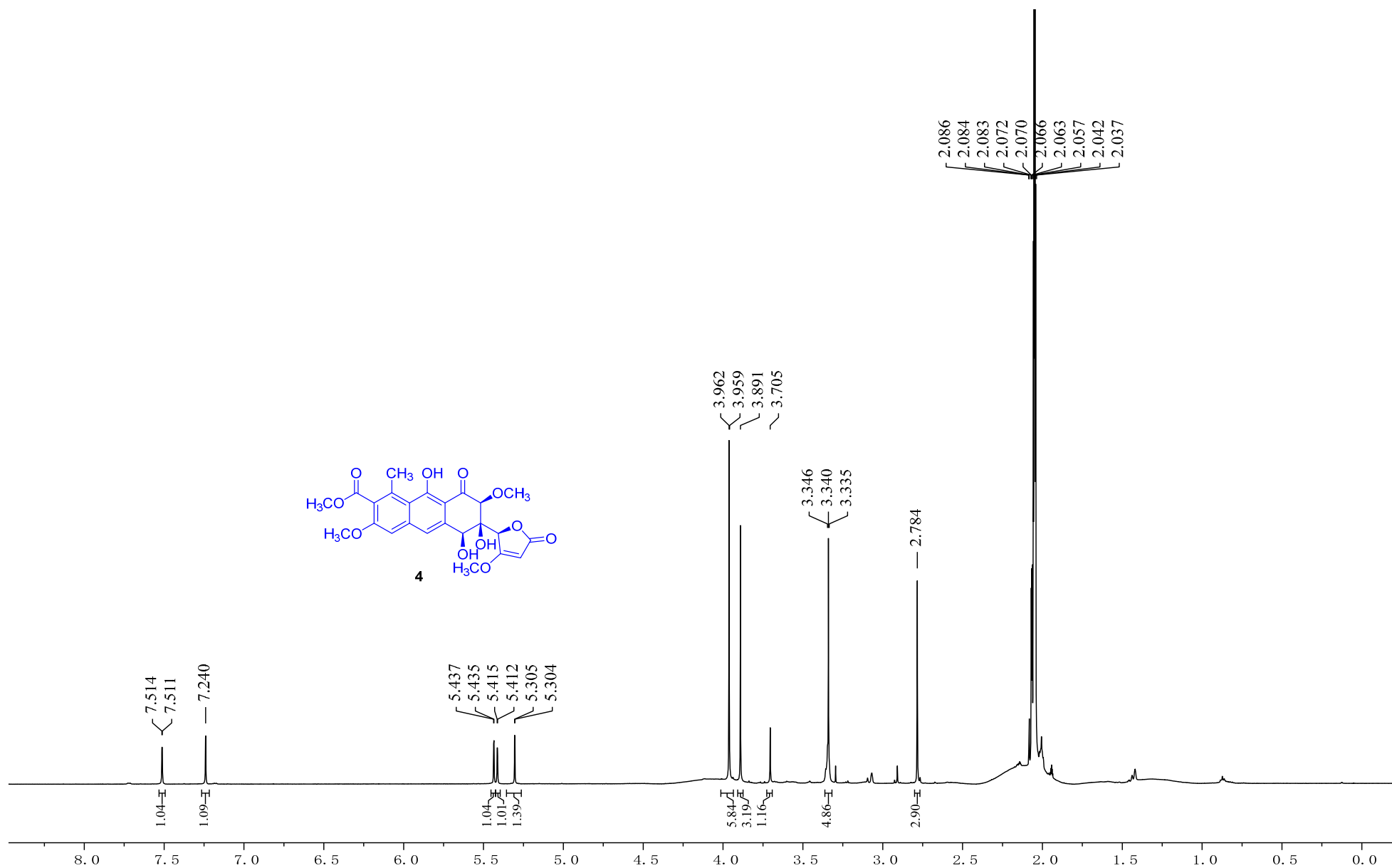


Figure S29. The ^1H NMR spectrum of saccharothrixone H (**4**) in acetone- d_6 (600 MHz).

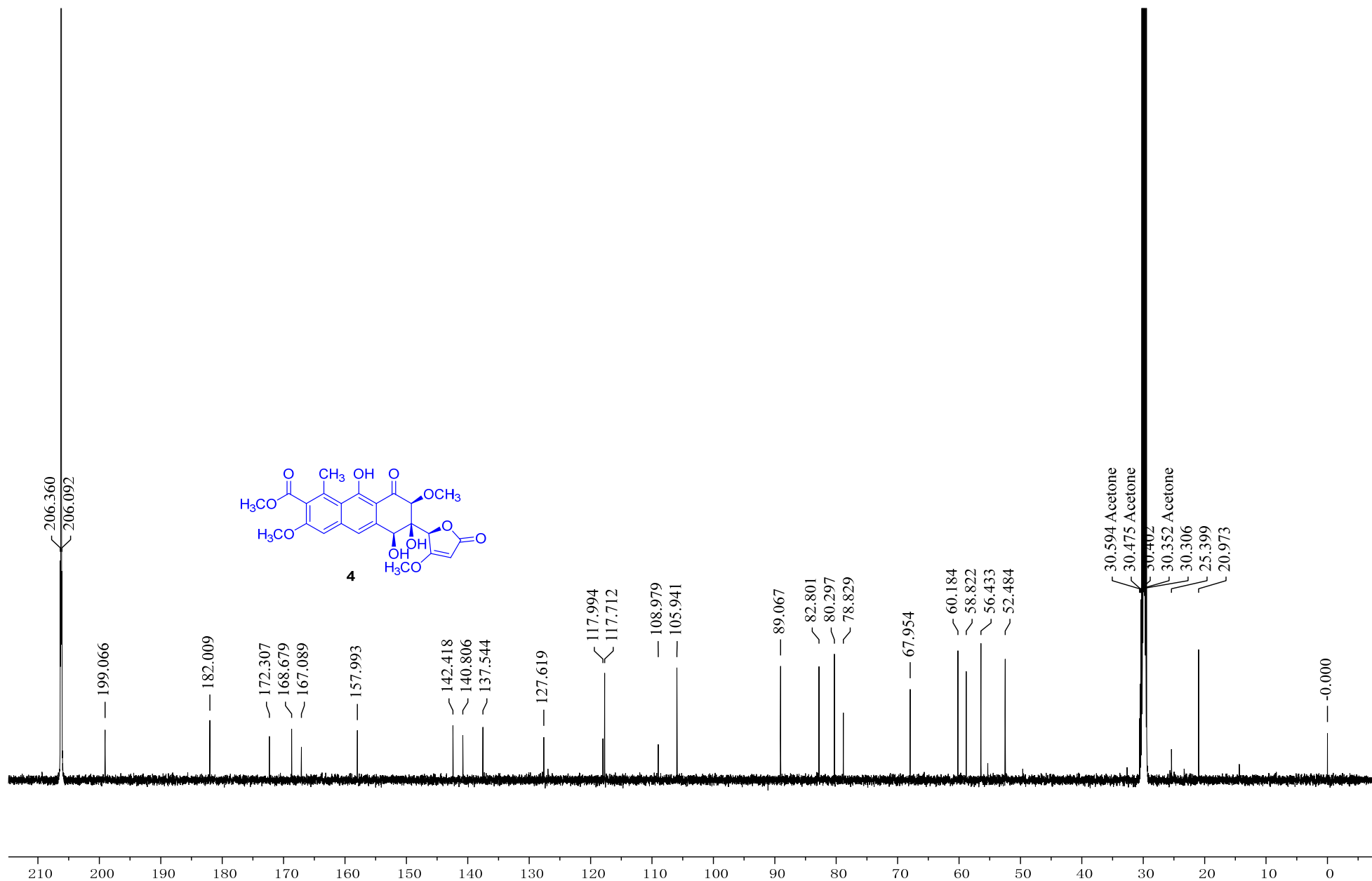


Figure S30. The ^{13}C NMR spectrum of saccharothrixone H (**4**) in acetone- d_6 (150 MHz).

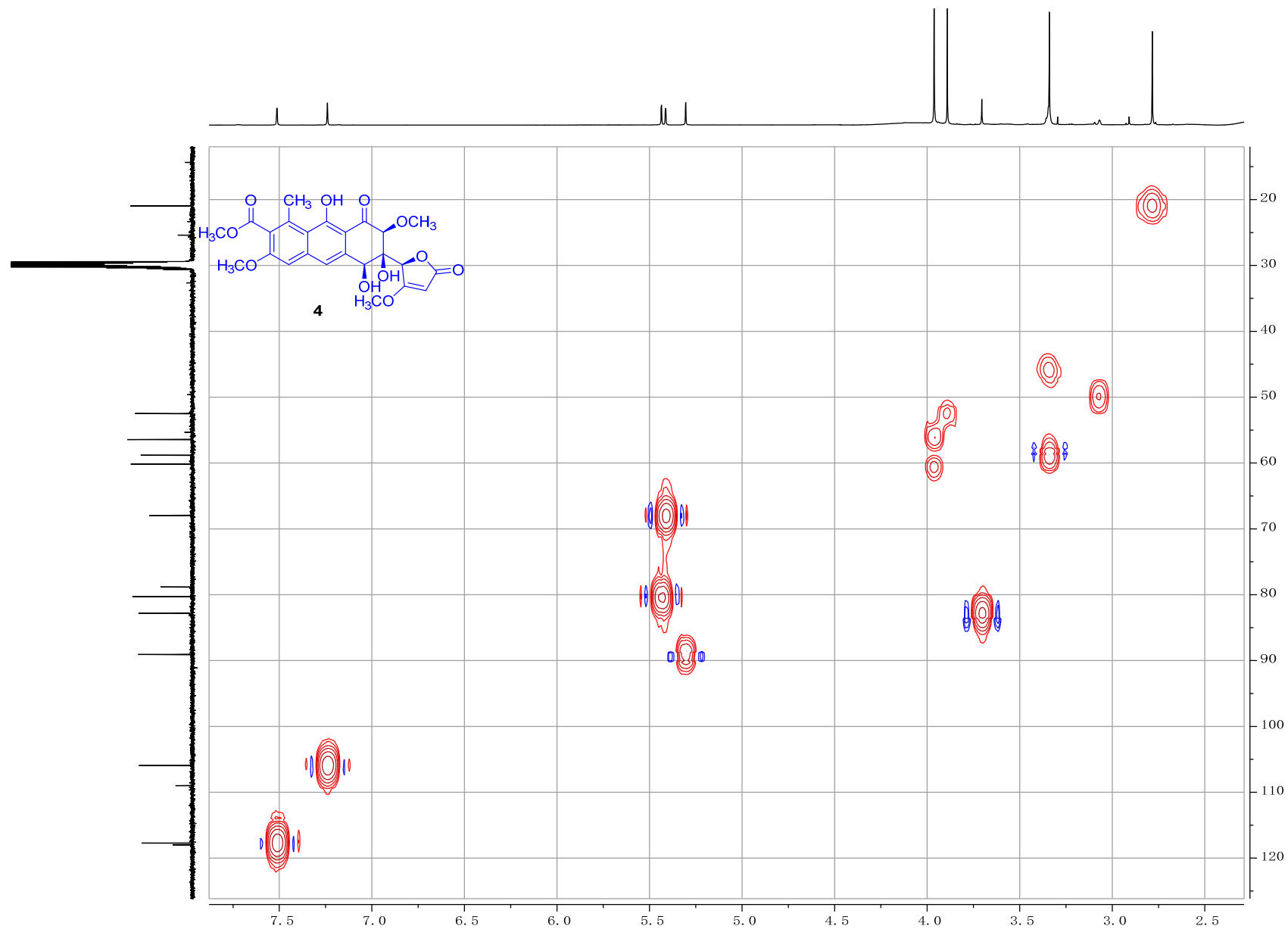


Figure S31. The HSQC spectrum of saccharothrixone H (**4**) in acetone- d_6 (600 MHz).

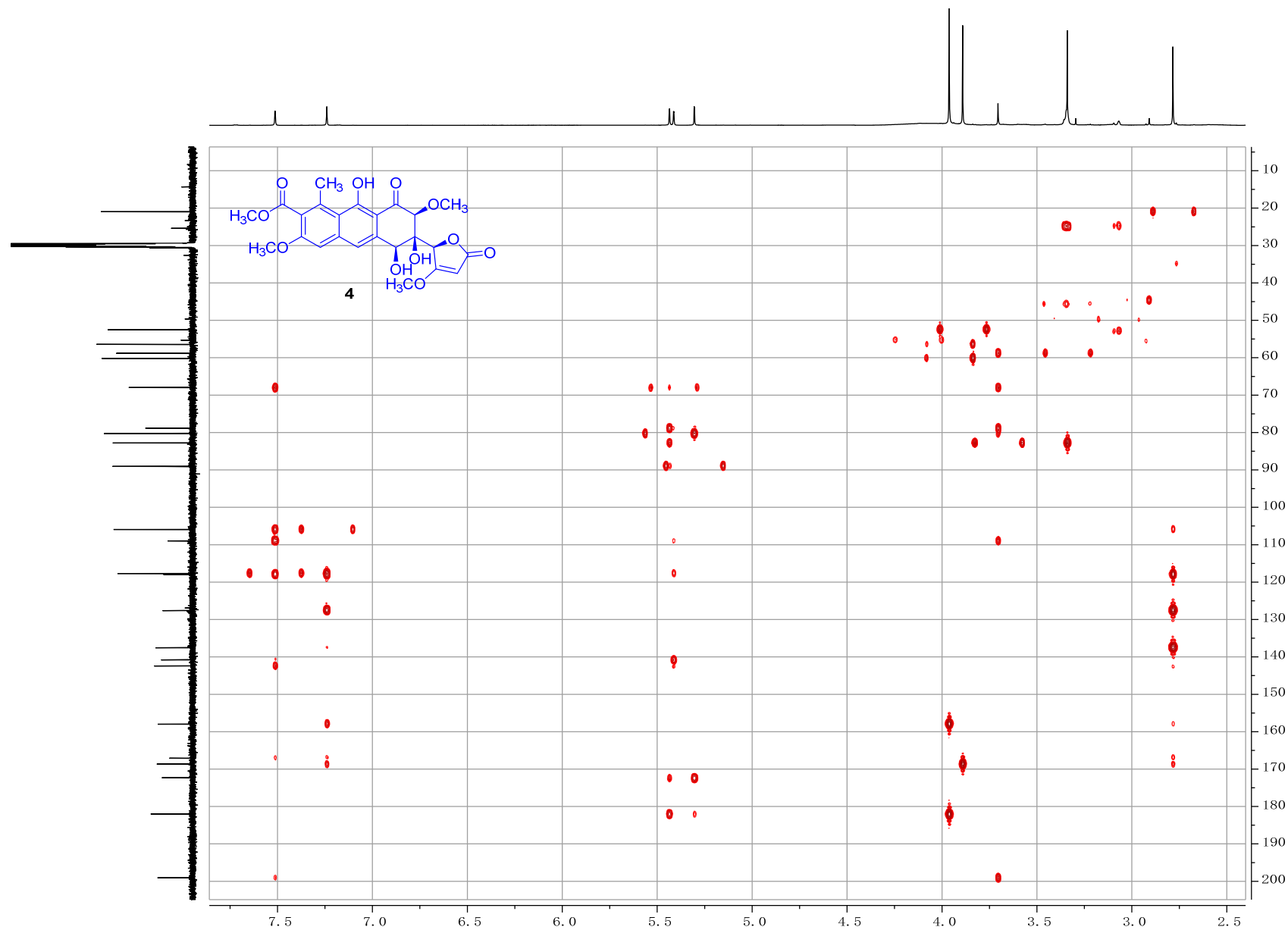


Figure S32. The HMBC spectrum of saccharothrixone H (4) in acetone- d_6 (600 MHz).

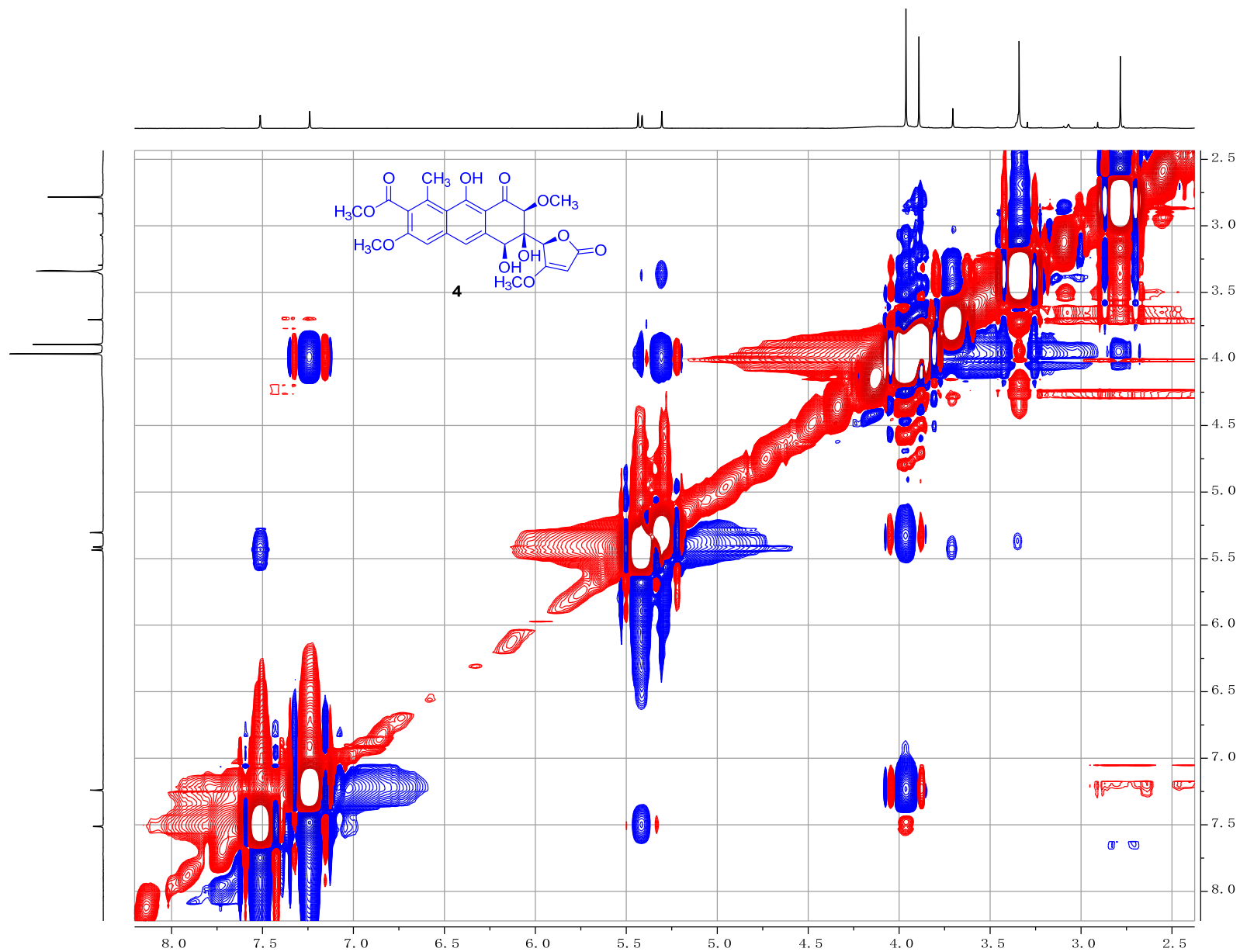


Figure S33. The ROESY spectrum of saccharothrixone H (**4**) in acetone-*d*₆ (600 MHz).

marine-20-11_HRESI#27 RT: 1.01 AV: 1 NL: 1.12E5
T: FTMS - c ESI Full ms [450.00-550.00]

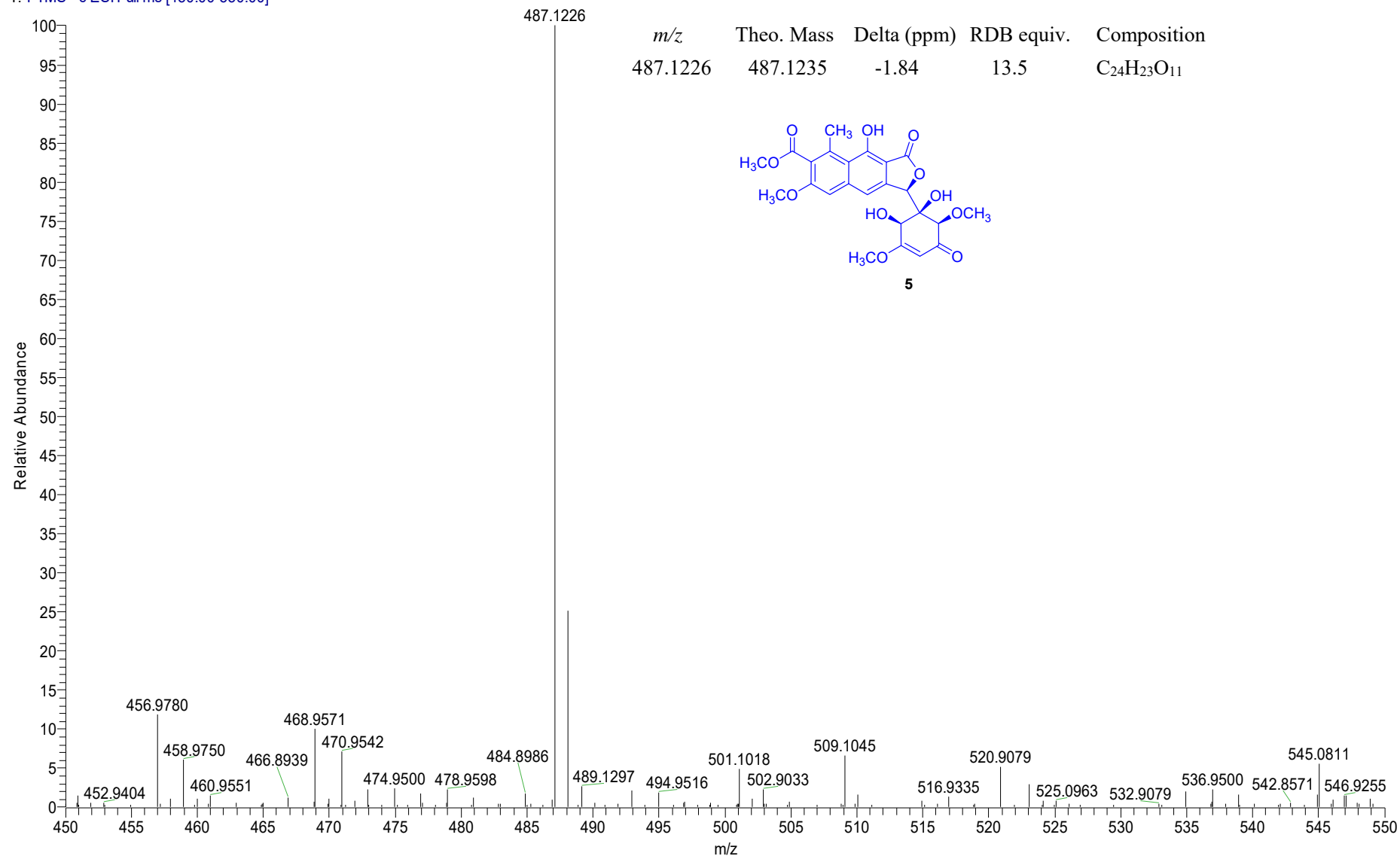


Figure S34. The (–)-HRESIMS spectrum of saccharothrixone I (**5**).

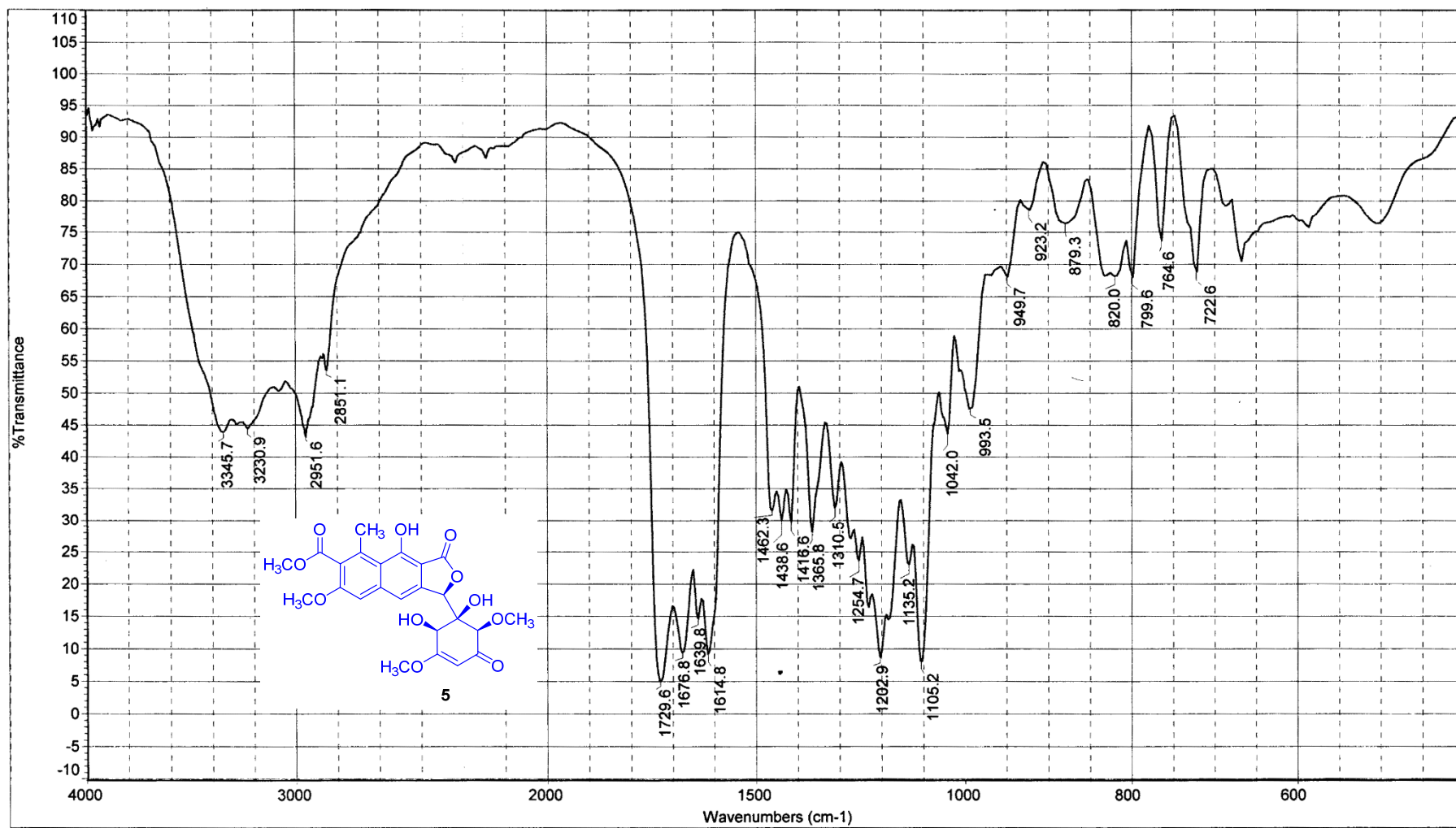


Figure S35. The IR spectrum of saccharothrixone I (5).

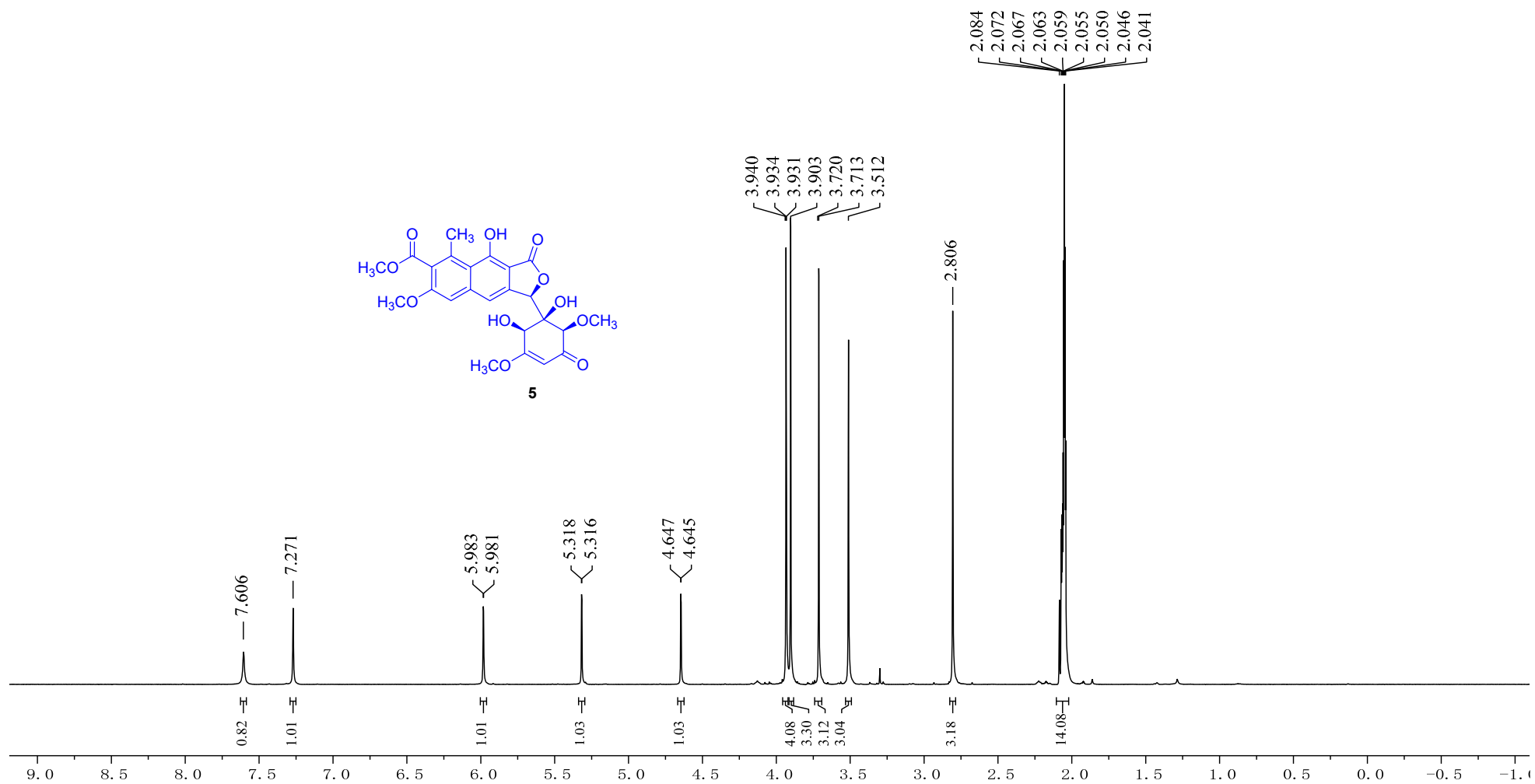


Figure S36. The ^1H NMR spectrum of saccharothrixone I (**5**) in acetone- d_6 (500 MHz).

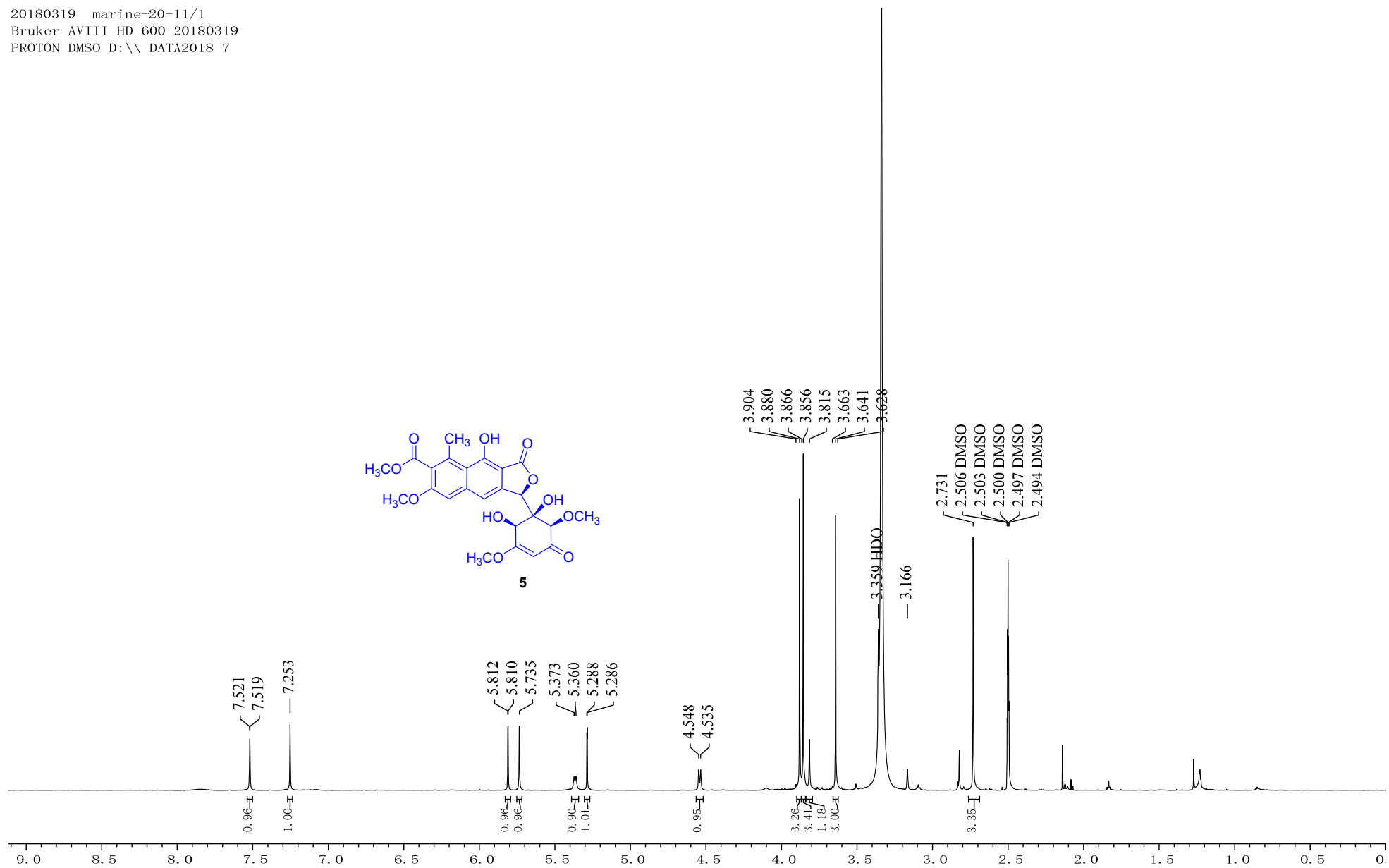


Figure S37. The ¹H NMR spectrum of saccharothrixone I (**5**) in DMSO-d₆ (600 MHz).

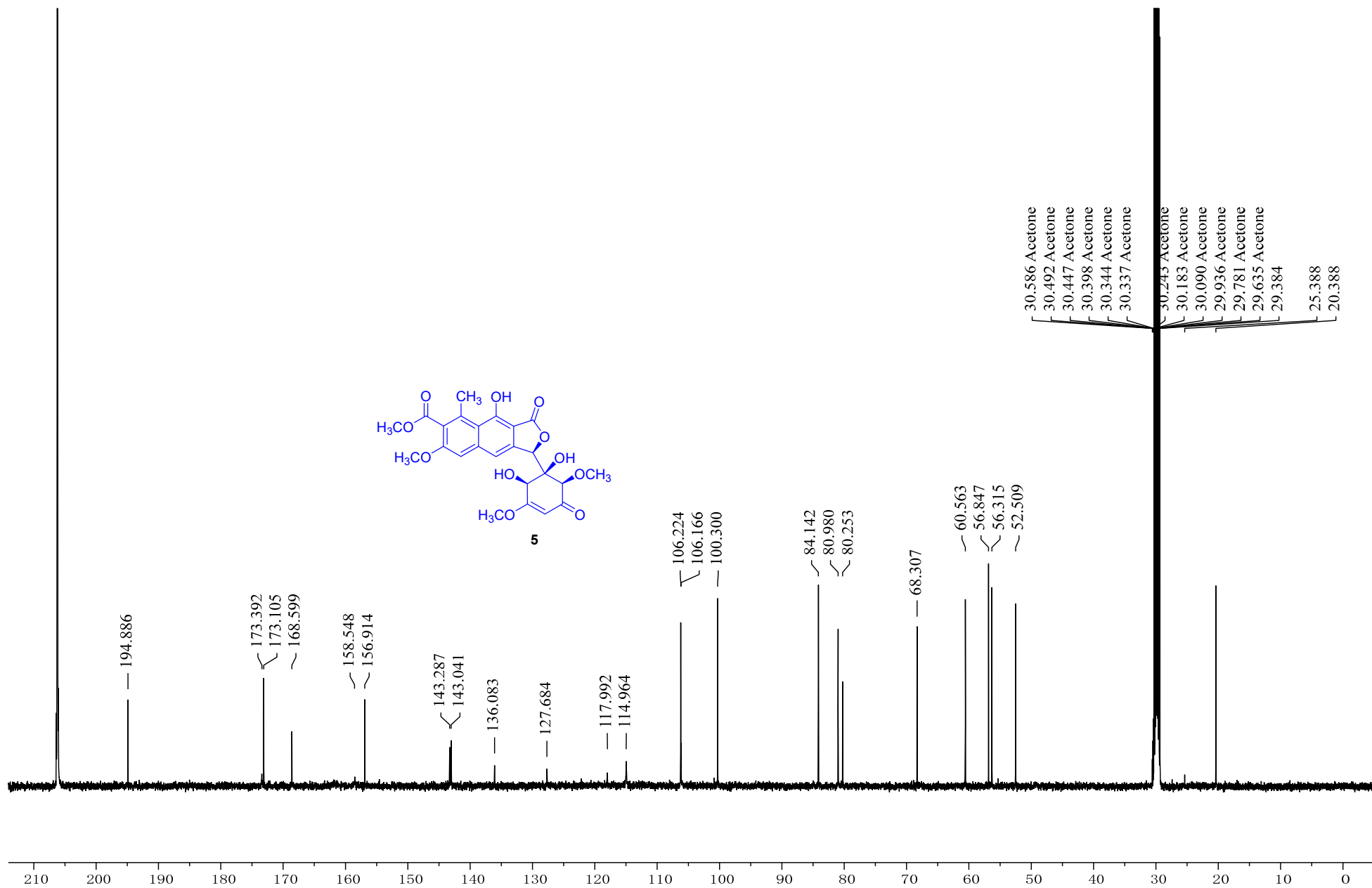


Figure S38. The ^{13}C NMR spectrum of saccharothrixone I (**5**) in acetone- d_6 (125 MHz).

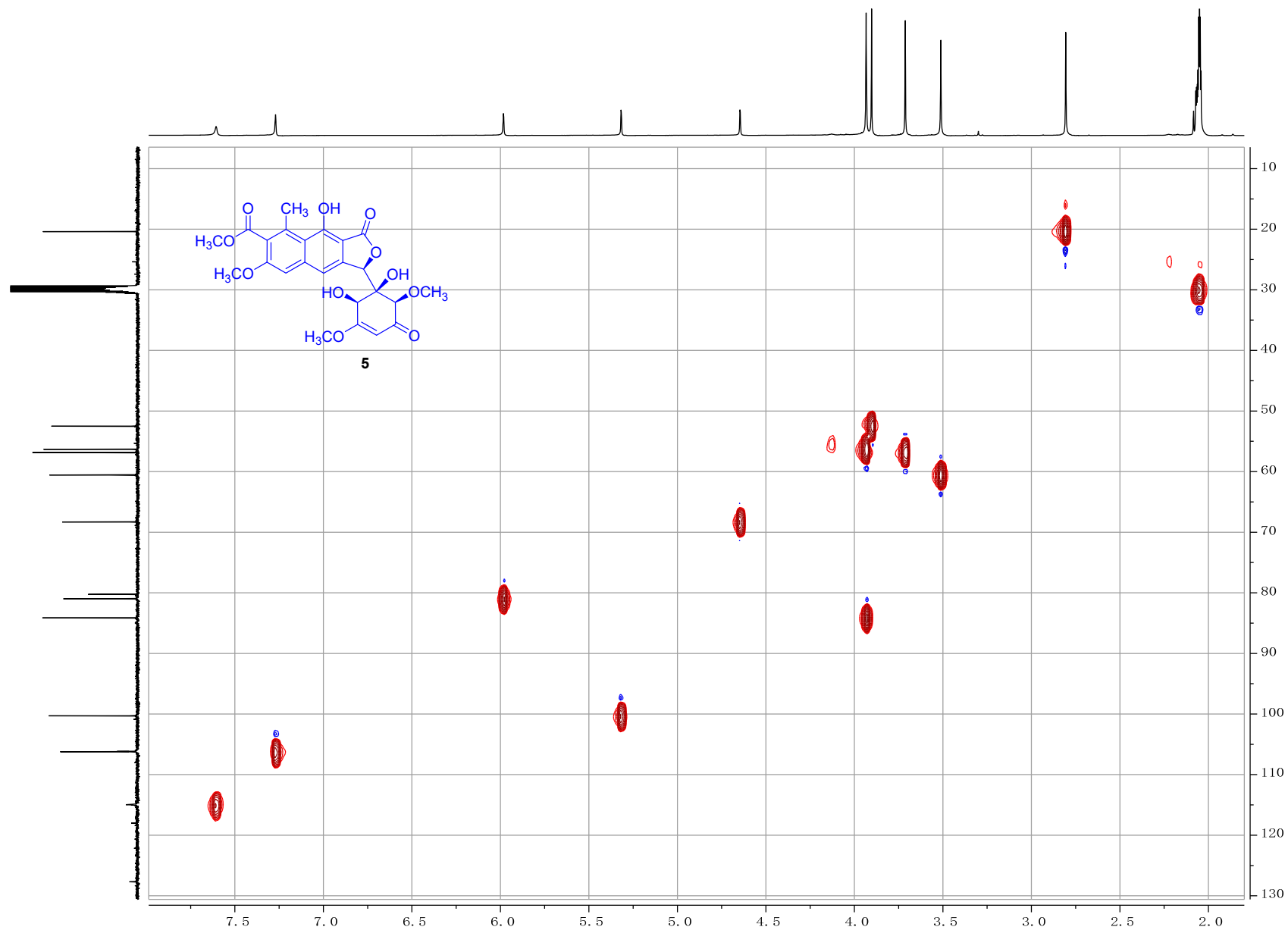


Figure S39. The HSQC spectrum of saccharothrixone I (**5**) in acetone- d_6 (500 MHz).

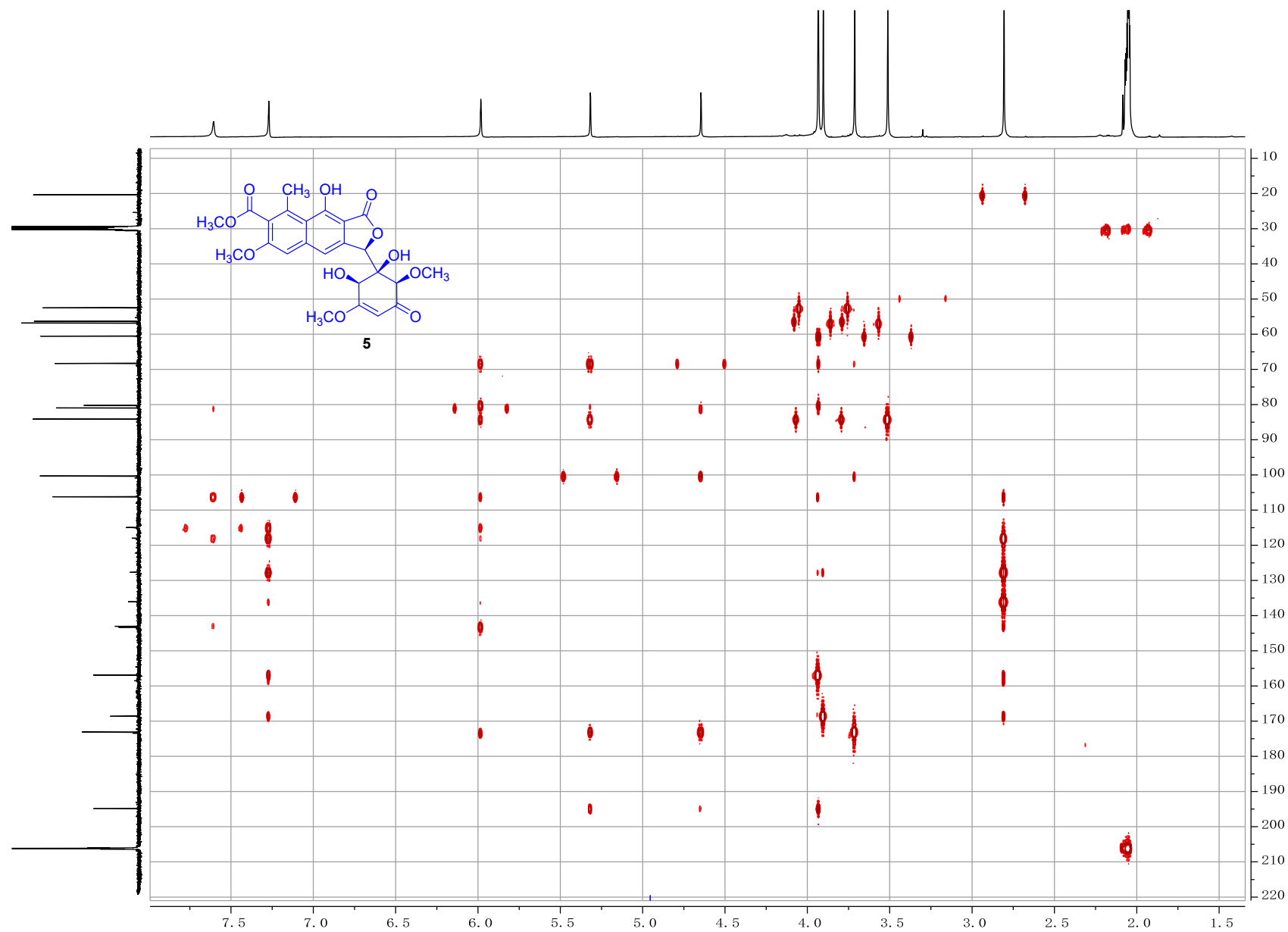


Figure S40. The HMBC spectrum of saccharothrixone I (**5**) in acetone- d_6 (500 MHz).

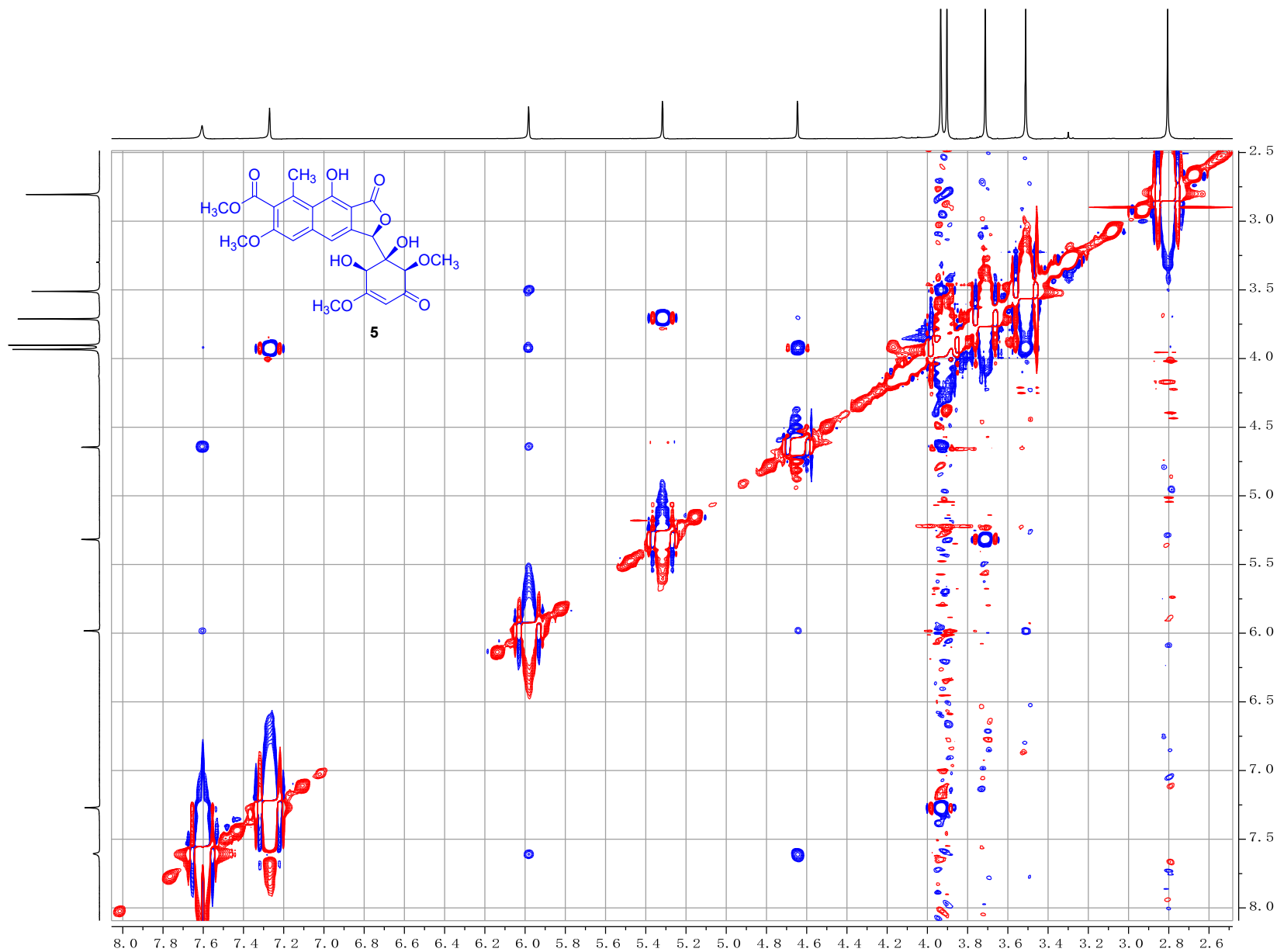


Figure S41. The ROESY spectrum of saccharothrixone I (**5**) in acetone- d_6 (600 MHz).

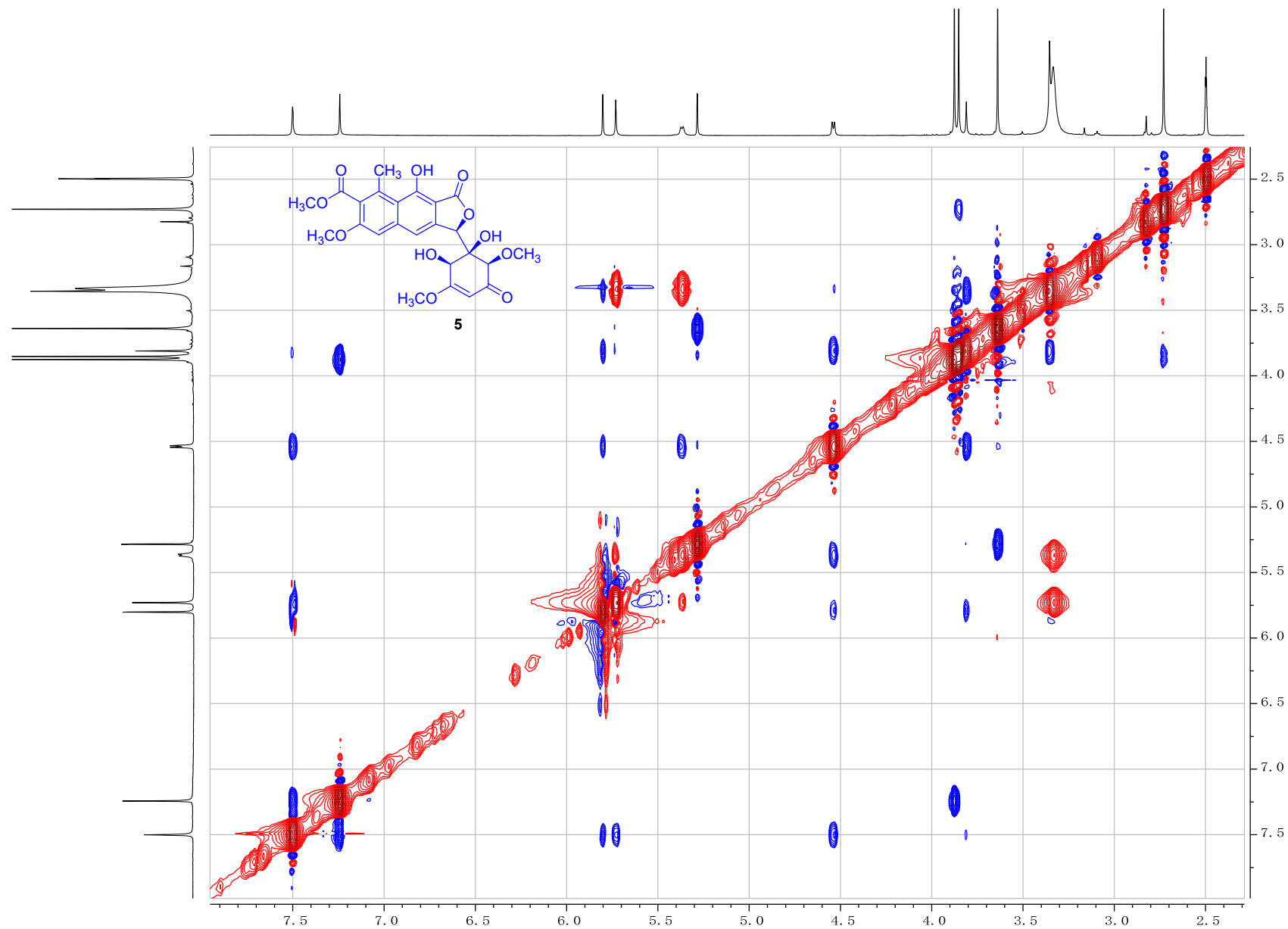


Figure S42. The ROESY spectrum of saccharothrixone I (**5**) in DMSO-*d*₆ (600 MHz).

MARINE-20-16 #35 RT: 1.17 AV: 1 NL: 7.73E4

T: FTMS + c ESI Full ms [450.00-550.00]

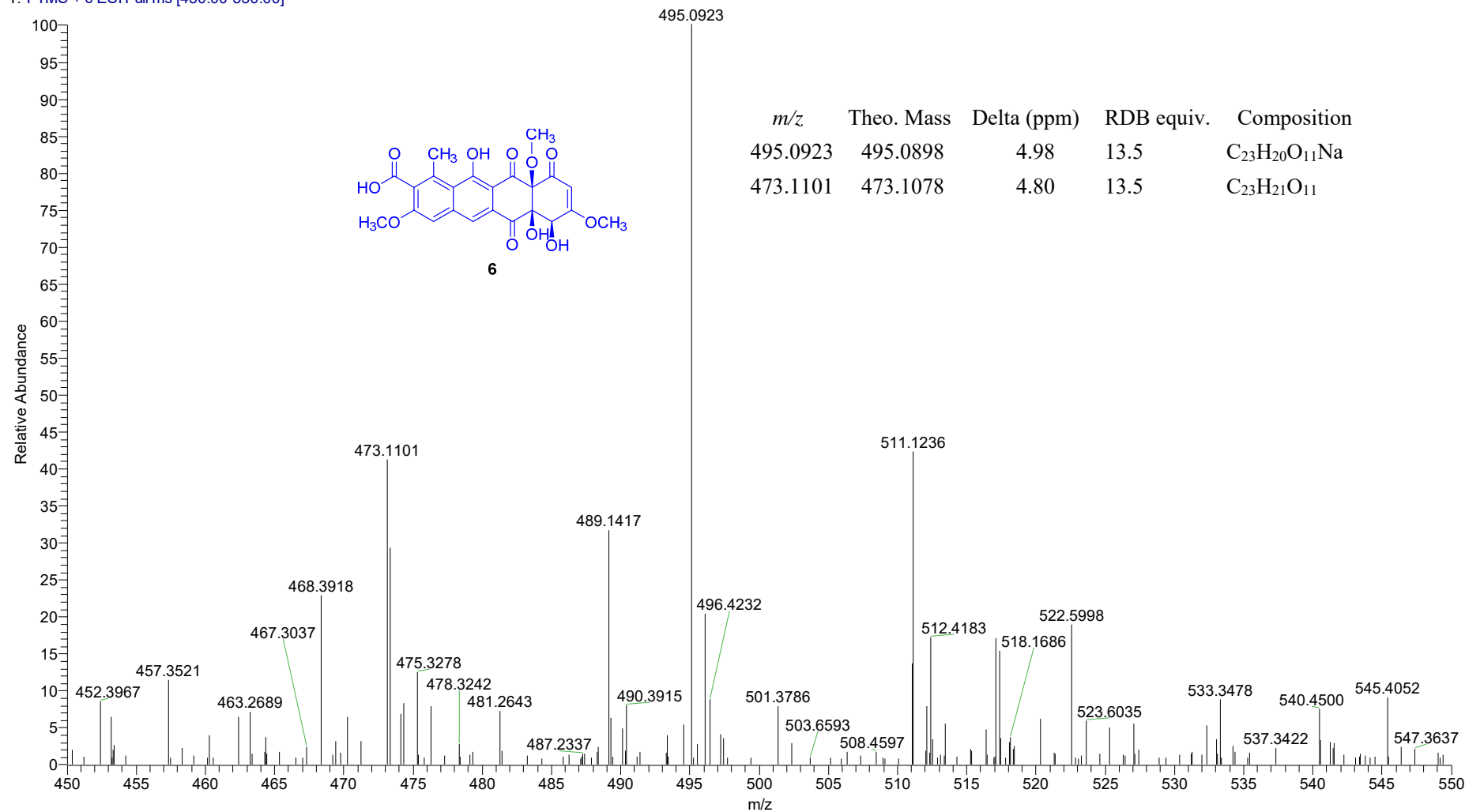


Figure S43. The (+)-HR ESIMS spectrum of 13-de-*O*-methyltetracenomycin X (**6**).

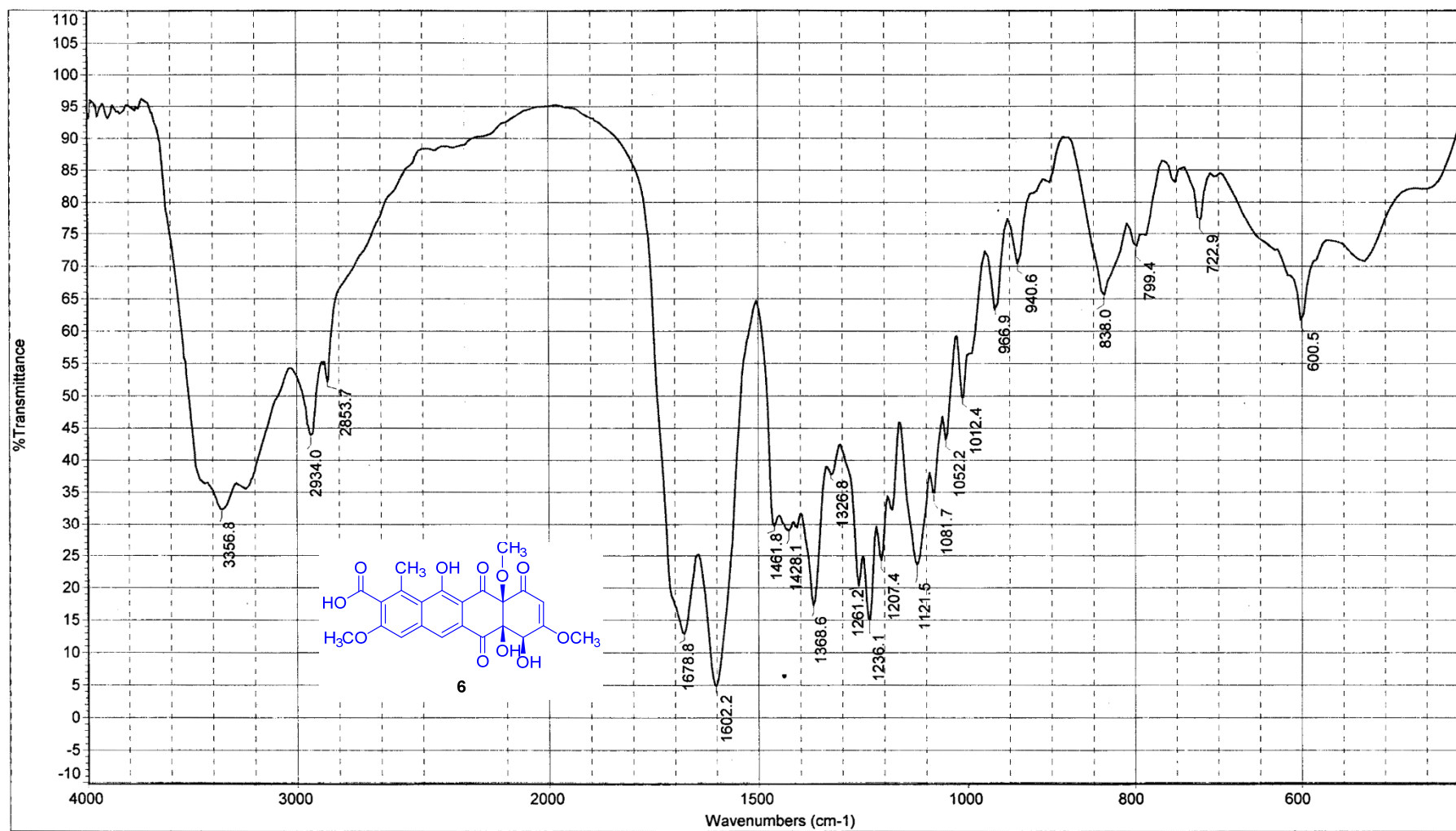


Figure S44. The IR spectrum of 13-de-O-methyltetracenomycin X (6).

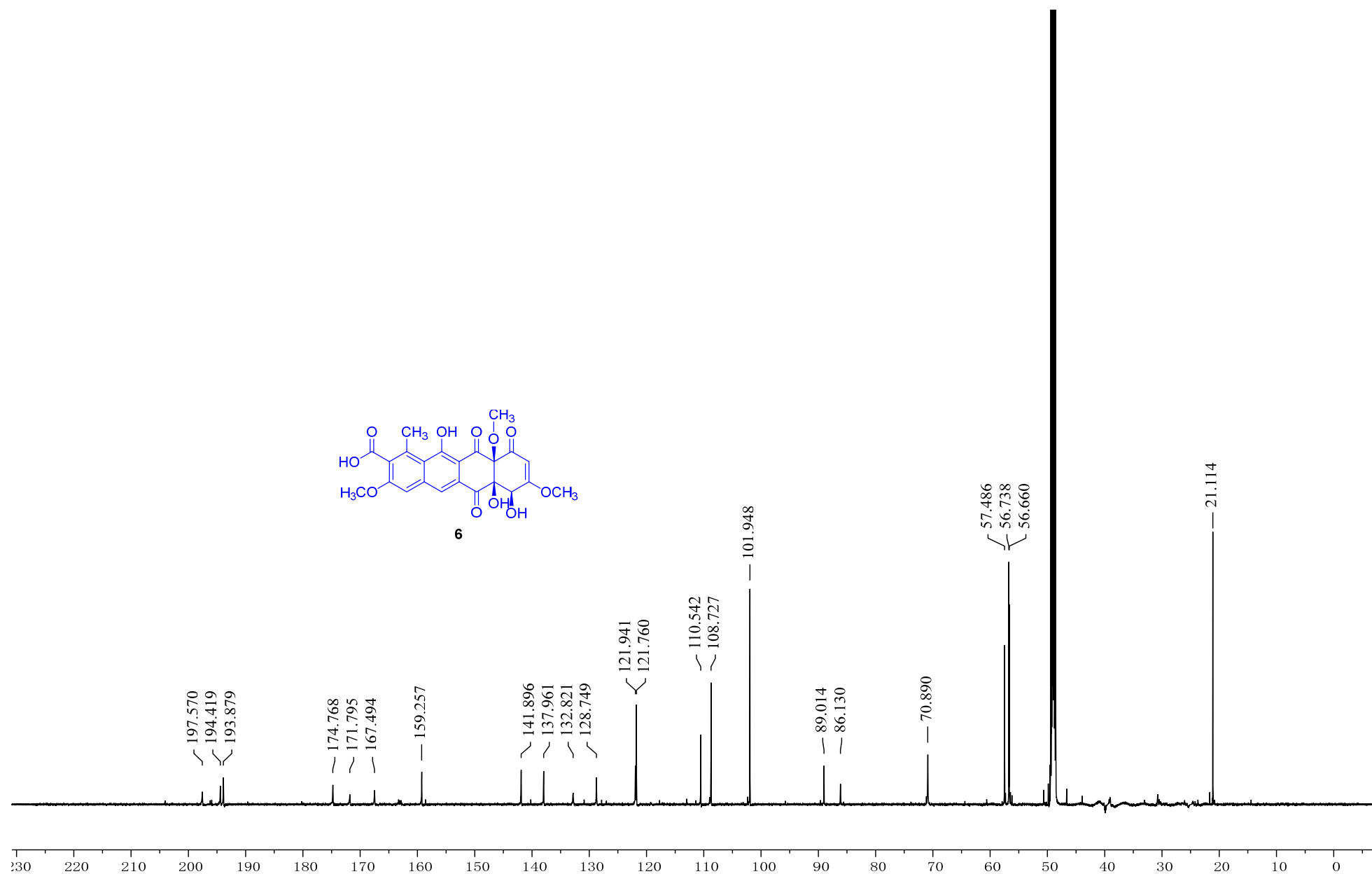


Figure S46. The ^{13}C NMR spectrum of 13-de-O-methyltetracenomycin X (**6**) in CD_3OD (150 MHz).

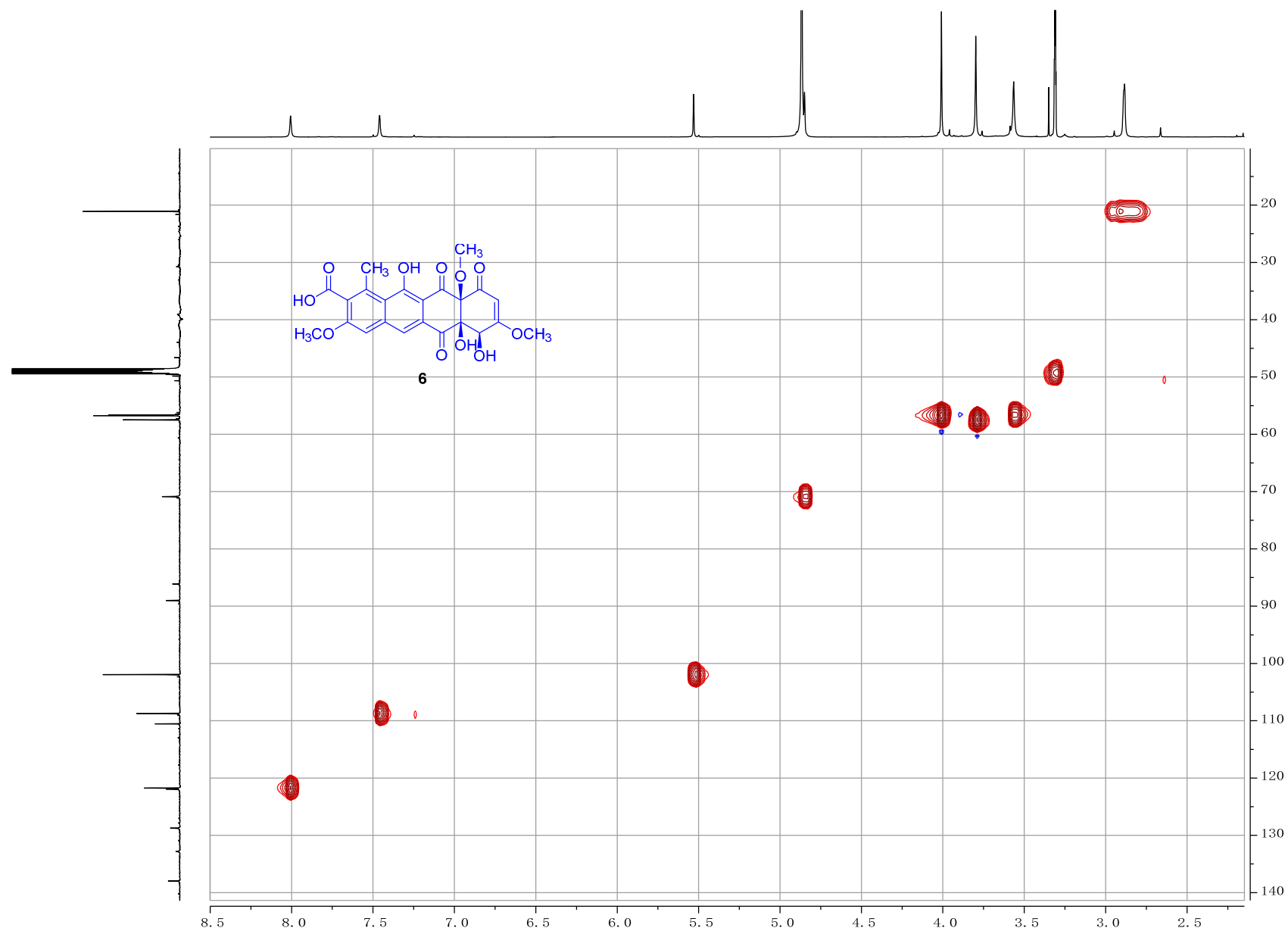


Figure S47. The HSQC spectrum of 13-de-*O*-methyltetracenomycin X (**6**) in CD₃OD (600 MHz).

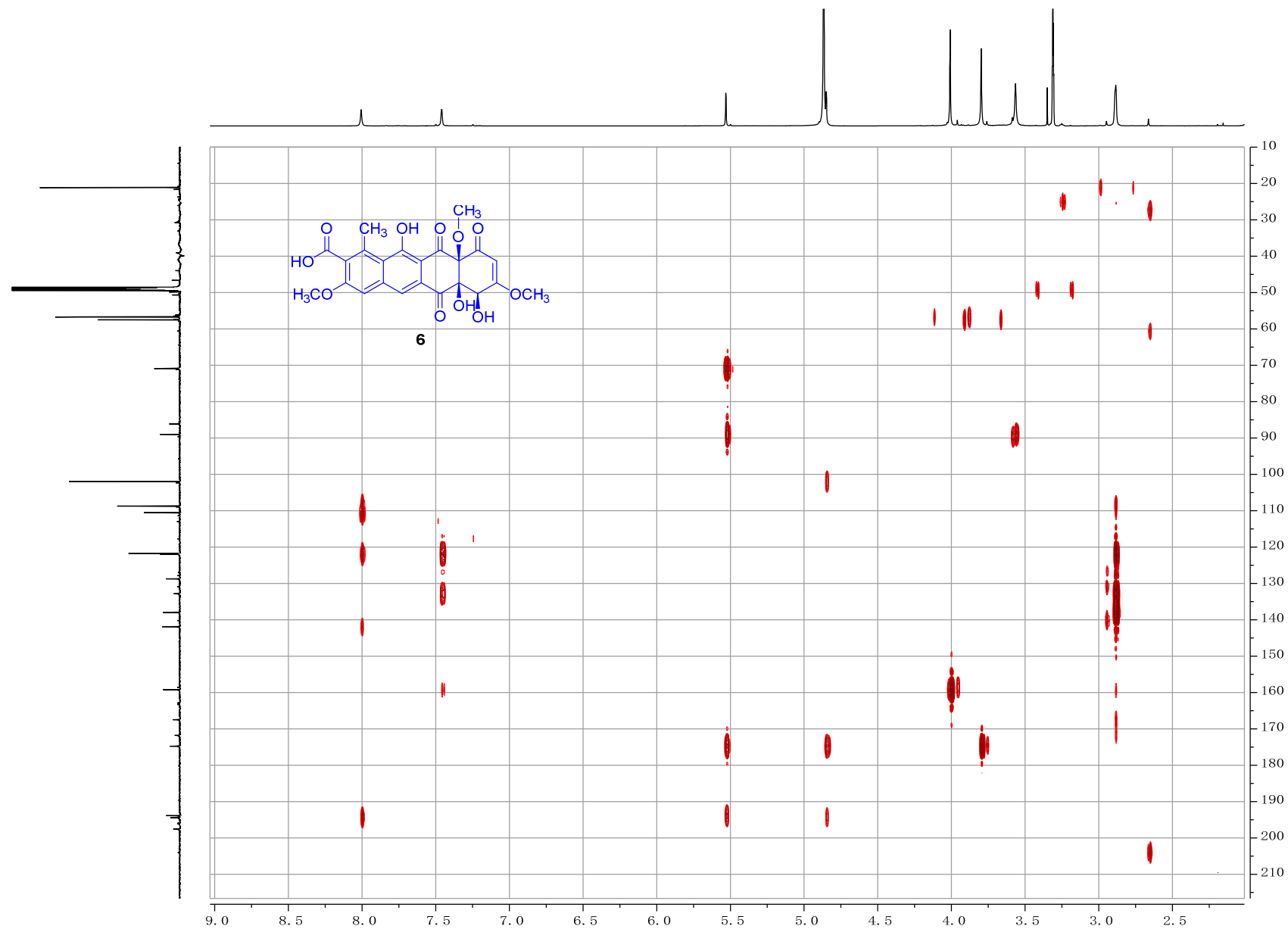


Figure S48. The HMBC spectrum of 13-de-*O*-methyltetracenomycin X (**6**) in CD₃OD (600 MHz).

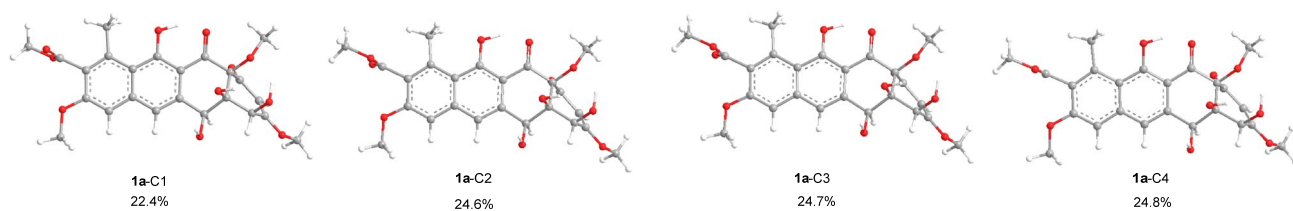


Figure S49. Optimized conformers ($\geq 1\%$) of (4*S*,4*aR*,5*S*,12*aR*)-**1a** at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.

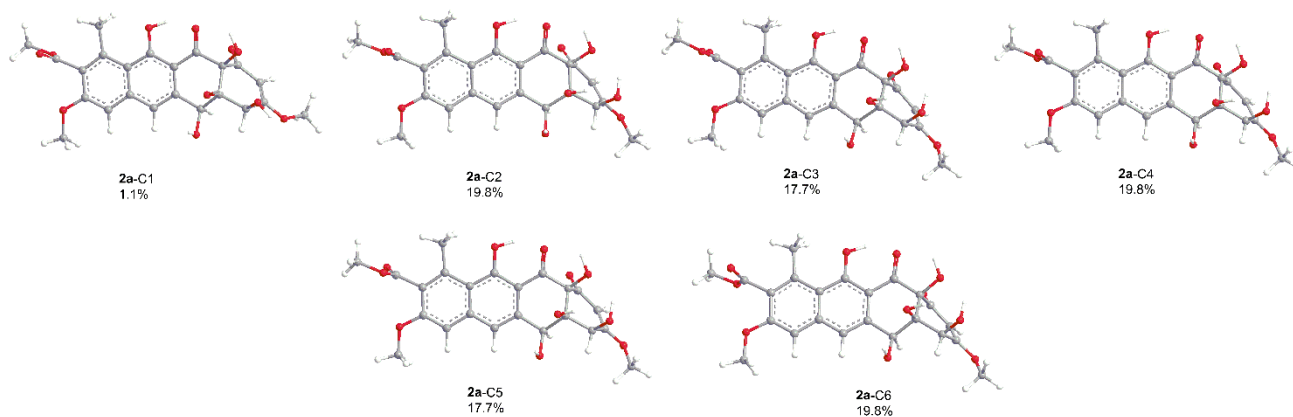


Figure S50. Optimized conformers ($\geq 1\%$) of (4*S*,4*aR*,5*S*,12*aR*)-**2a** at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.

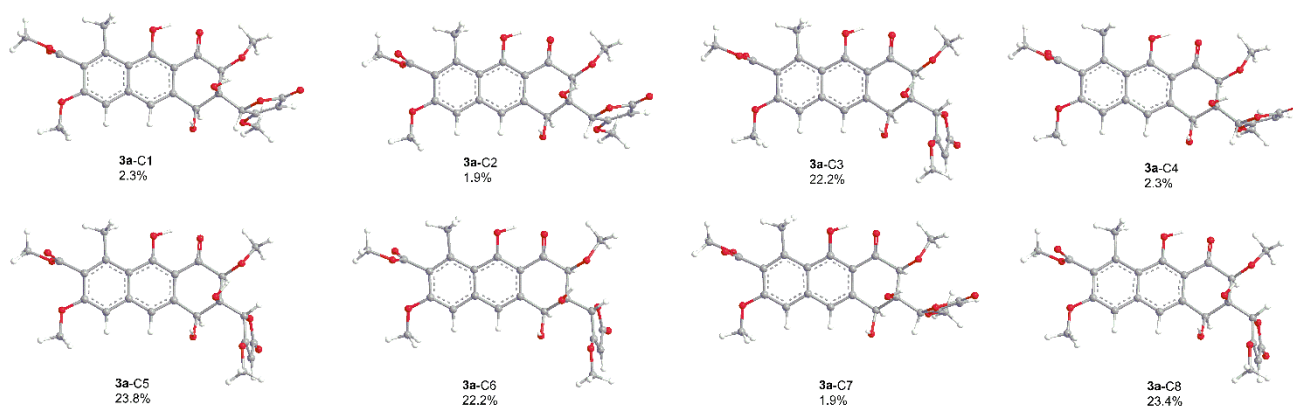


Figure S51. Optimized conformers ($\geq 1\%$) of (4*S*,4*aR*,5*S*,12*aS*)-**3a** at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.

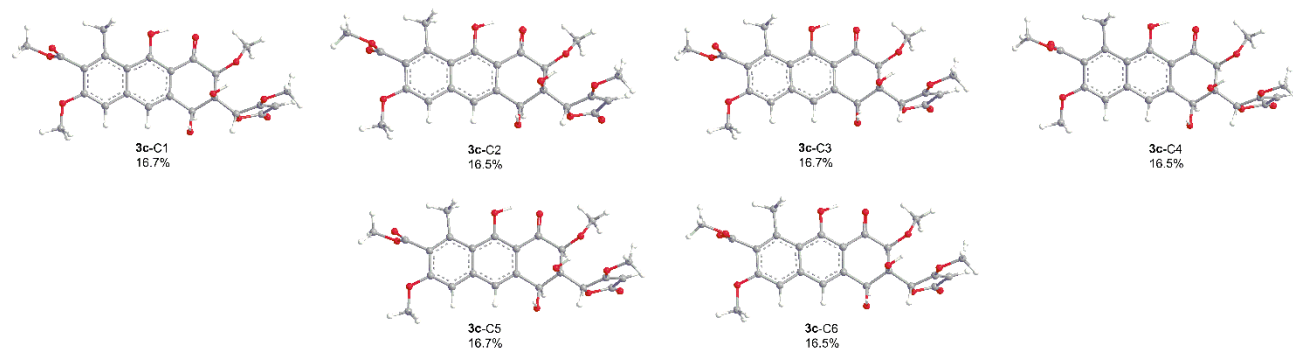


Figure S52. Optimized conformers ($\geq 1\%$) of (4*R*,4*aR*,5*S*,12*aS*)-**3c** at the B3LYP/6-311+G(d,p) level with PCM model in MeOH.

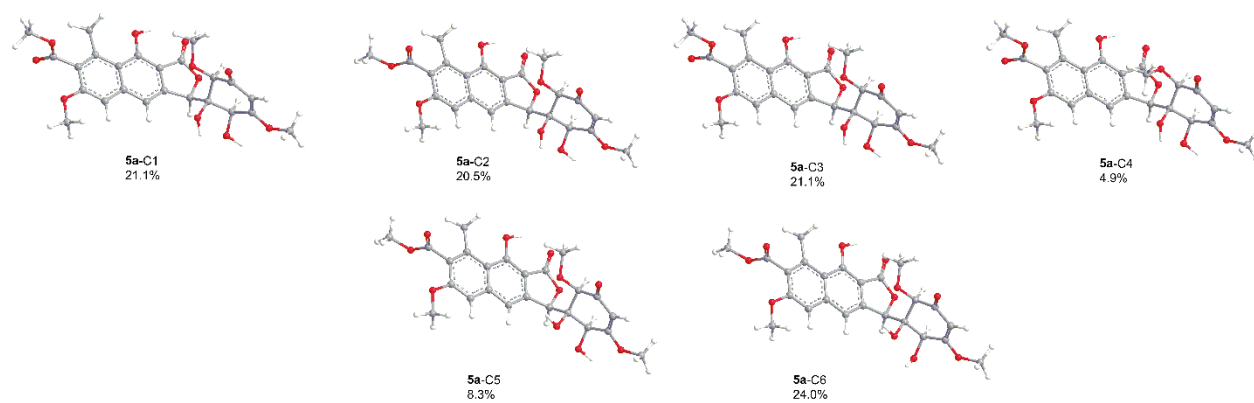


Figure S53. Optimized conformers ($\geq 1\%$) of (4*S*,4*aR*,5*R*,12*aR*)-**5a** at the B3LYP/6-311+G (d,p) level with PCM model in MeOH.