

Supporting Information for

**Targeted Isolation of Two Disesquiterpenoids Macrocephadiolides A and B from
Ainsliaea macrocephala using Molecular Networking-based Dereplication
Strategy**

Yong-Mei Ren, ^{*a, b, c} Shuai-Zhen Zhou, ^a Tian Zhang, ^d Meijia Qian, ^e Rui Zhang, ^a Sheng Yao, ^a
Hong Zhu, ^e Chunping Tang ^a, Ligen Lin^{*d}, and Yang Ye ^{*a, b, c}

^a State Key Laboratory of Drug Research and Natural Products Chemistry Department, Shanghai
Institute of Materia Medica, Chinese Academy of Sciences, Shanghai 201203, China. E-mail:
yye@simm.ac.cn.

^b School of Life Science and Technology, ShanghaiTech University, Shanghai 201210, China.

^c University of Chinese Academy of Sciences, No.19A Yuquan Road, Beijing 100049, China.

^d State Key Laboratory of Quality Research in Chinese Medicine, Institute of Chinese Medical
Sciences, University of Macau, Taipa, Macau, China.

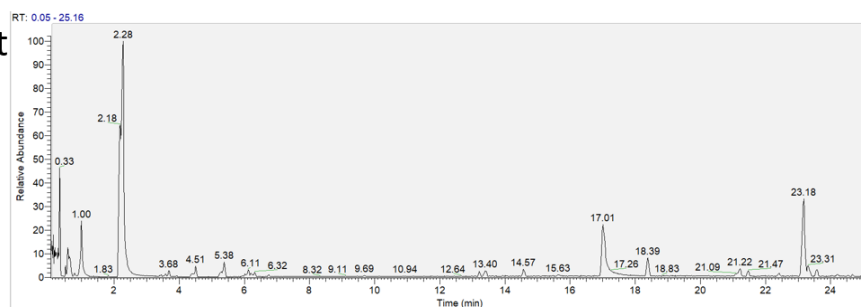
^e Institute of Pharmacology and Toxicology, College of Pharmaceutical Sciences, Zhejiang
University, Hangzhou, 310058, China.

List of contents

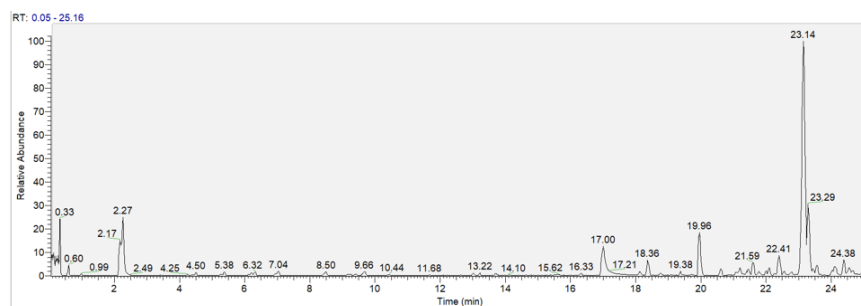
Scheme S1. UHPLC-MS chromatograms that were used for molecular networking.....	1
Table S1 Characterization of disesquiterpenoids from <i>A. macrocephala</i> by MN.	2
Table S2. X-ray crystallographic data for 1	2
Fig. S1. IR spectrum of compound 1	3
Fig. S2. HRESIMS spectrum of compound 1	4
Fig. S3. ¹ H NMR spectrum of compound 1 in CDCl ₃	5
Fig. S4. 1D TOCSY (4.095 PPM) spectrum of compound 1 in CDCl ₃	6
Fig. S5. 1D TOCSY (4.812 PPM) spectrum of compound 1 in CDCl ₃	7
Fig. S6. ¹³ C NMR spectrum of compound 1 in CDCl ₃	8
Fig. S7. HSQC spectrum of compound 1 in CDCl ₃	9
Fig. S8. HMBC spectrum of compound 1 in CDCl ₃	10
Fig. S9. ¹ H- ¹ H COSY spectrum of compound 1 in CDCl ₃	11
Fig. S10. ROESY spectrum of compound 1 in CDCl ₃	12
Fig. S11. IR spectrum of compound 2	13
Fig. S12. HRESIMS data of compound 2	14
Fig. S13. ¹ H NMR spectrum of compound 2 in CD ₃ OD	15
Fig. S14. 1D TOCSY (4.108 PPM) spectrum of compound 2 in CD ₃ OD	16
Fig. S15. 1D TOCSY (4.389 PPM) spectrum of compound 2 in CD ₃ OD	17
Fig. S16. ¹³ C NMR spectrum of compound 2 in CD ₃ OD	18
Fig. S17. HSQC spectrum of compound 2 in CD ₃ OD.....	19
Fig. S18. HMBC spectrum of compound 2 in CD ₃ OD.....	20
Fig. S19. ¹ H- ¹ H COSY spectrum of compound 2 in CD ₃ OD	21
Fig. S20. ROESY spectrum of compound 2 in CD ₃ OD.....	22
Fig. S21 HRESIMS data of compound 3	23
Fig. S22 ¹ H NMR spectrum of compound 3 in CDCl ₃	24
Fig. S23 ¹³ C NMR spectrum of compound 3 in CDCl ₃	25
Fig. S24. HRESIMS data of compound 4	26
Fig. S25 ¹ H NMR spectrum of compound 4 in CDCl ₃	27
Fig. S26 ¹³ C NMR spectrum of compound 4 in CDCl ₃	28
Fig. S27. HRESIMS data of compound 5	29
Fig. S28 ¹ H NMR spectrum of compound 5 in CDCl ₃	30
Fig. S29. ¹³ C NMR spectrum of compound 5 in CDCl ₃	31
Fig. S30 HRESIMS data of compound 6	32
Fig. S31 ¹ H NMR spectrum of compound 6 in CDCl ₃	33
Fig. S32 ¹³ C NMR spectrum of compound 6 in CDCl ₃	34
Fig. S33. HRESIMS data of compound 7	35
Fig. S34 ¹ H NMR spectrum of compound 7 in CDCl ₃	36
Fig. S35 ¹³ C NMR spectrum of compound 7 in CDCl ₃	37
Fig. S36. ¹ H NMR spectrum of compound 8 in CDCl ₃	38
Fig. S37 ¹³ C NMR spectrum of compound 8 in CDCl ₃	39
The complete western blot membranes and repeated results for Figure 7B.....	40
The complete western blot membranes and repeated results for Figure 7C.....	44

Scheme S1. UHPLC-MS chromatograms that were used for molecular networking

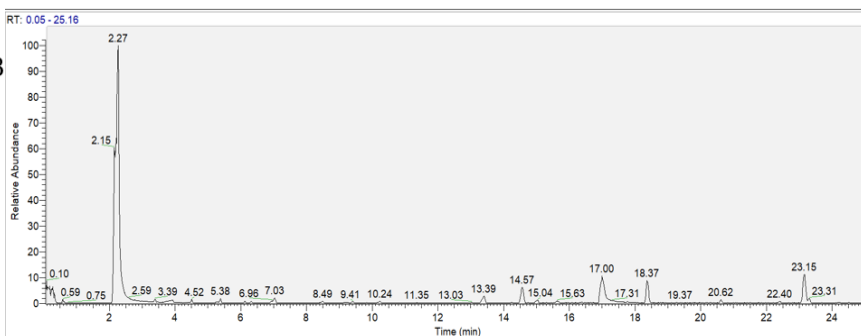
Extract



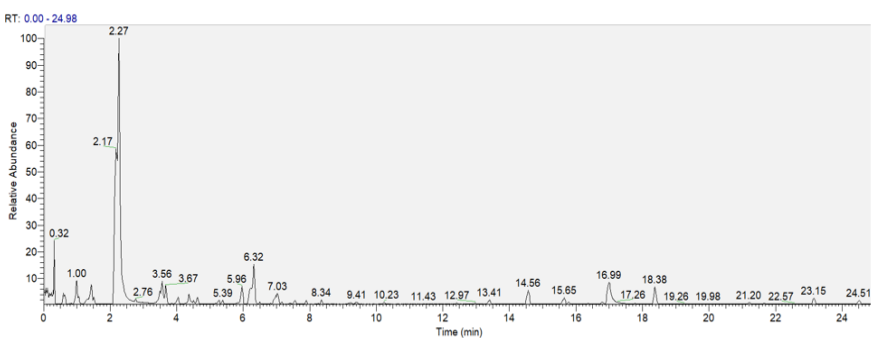
Fr. PE



Fr. CHCl₃



Fr. EA



Fr. H₂O

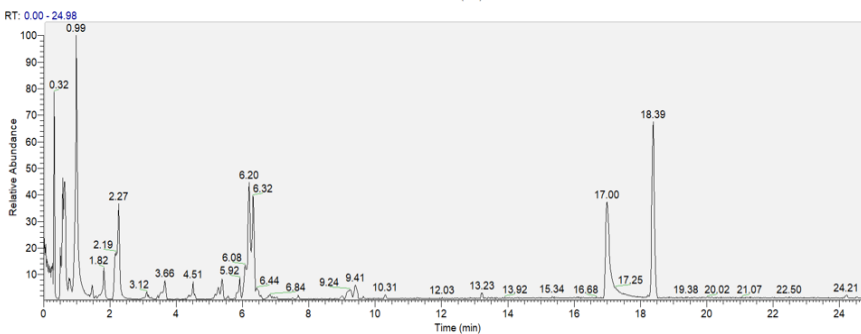


Table S1 Characterization of disesquiterpenoids from *A. macrocephala* by MN.

No.	m/z	δ_{ppm}	t_{R}	Molecular Formula	Ion description	Identification
1	521.21674	-1.540	14.13	C ₃₀ H ₃₂ O ₈	[M+H] ⁺	Macrocephadiolide A
2	523.23260	-1.133	14.97	C ₃₀ H ₃₄ O ₈	[M+H] ⁺	Macrocephadiolide B
3	520.23254	-0.844	13.79	C ₃₀ H ₃₀ O ₇	[M+NH ₄] ⁺	Gochnatolide A
4	485.19531	-1.144	13.34	C ₃₀ H ₃₀ O ₇	[M+H-H ₂ O] ⁺	Gochnatolide B
5	491.24268	-0.275	14.73	C ₃₀ H ₃₆ O ₇	[M+H-H ₂ O] ⁺	Ainsliadimer C
6	507.23740	-0.650	15.65	C ₃₀ H ₃₄ O ₇	[M+H] ⁺	Ainsliadimer A
7	524.26434	0.116	15.28	C ₃₀ H ₃₄ O ₇	[M+NH ₄] ⁺	Ainsliadimer J

These MS data were directly from the MN and they were not identical with the HRESIMS data in the following part, which were collected with pure compounds.

Table S2. X-ray crystallographic data for **1**

Identification code	Cu 20180655_0m
Empirical formula	C ₃₀ H ₃₂ O ₈
Formula weight	520.55
Temperature/K	173
Crystal system	monoclinic
Space group	P2 ₁
a/Å	8.8559(7)
b/Å	9.4486(9)
c/Å	15.3519(13)
$\alpha/^\circ$	90
$\beta/^\circ$	94.432(3)
$\gamma/^\circ$	90
Volume/Å ³	1280.74(19)
Z	2
$\rho_{\text{calc}}/\text{cm}^3$	1.350
μ/mm^{-1}	0.804
F(000)	552.0
Crystal size/mm ³	0.25 × 0.15 × 0.08
Radiation	Cu K α (λ = 1.54178)
2 θ range for data collection/ $^\circ$	5.774 to 127.998
Index ranges	-10 ≤ h ≤ 8, -10 ≤ k ≤ 10, -17 ≤ l ≤ 16
Reflections collected	14193
Independent reflections	3934 [R_{int} = 0.0281, R_{sigma} = 0.0245]
Data/restraints/parameters	3934/1/343
Goodness-of-fit on F ²	1.066
Final R indexes [$I \geq 2\sigma(I)$]	R_1 = 0.0292, wR_2 = 0.0767
Final R indexes [all data]	R_1 = 0.0295, wR_2 = 0.0769
Largest diff. peak/hole / e Å ⁻³	0.49/-0.17
Flack parameter	0.05(6)

Fig. S1. IR spectrum of compound **1**

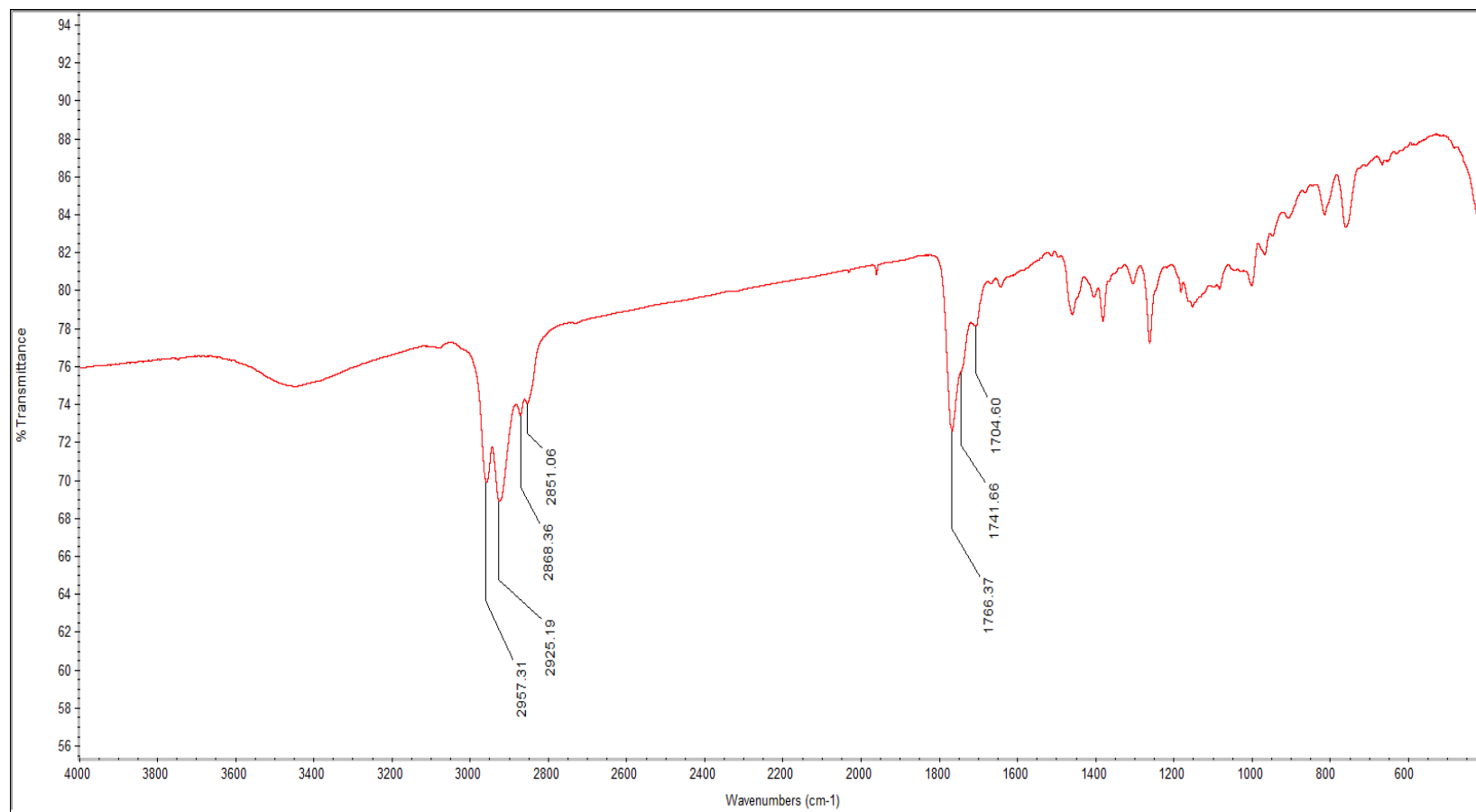


Fig. S2. HRESIMS spectrum of compound **1**

Elemental Composition Report

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

286 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-100 H: 0-100 O: 0-50 Na: 0-1

Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
543.1996	543.1995	0.1	0.2	14.5	221.7	0.156	85.59	C30 H32 O8 Na
	543.2019	-2.3	-4.2	17.5	223.6	1.985	13.74	C32 H31 O8
	543.1960	3.6	6.6	26.5	226.6	4.999	0.67	C39 H27 O3

AMB1F3D5C-pos
20180823-SA019 1647 (6.659)

2: TOF MS ES+
6.38e+004

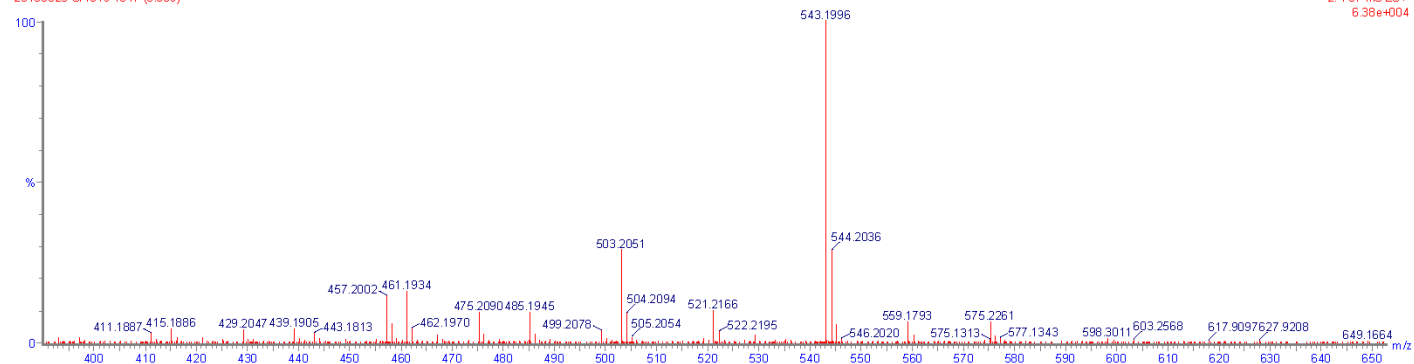


Fig. S3. ^1H NMR spectrum of compound **1** in CDCl_3

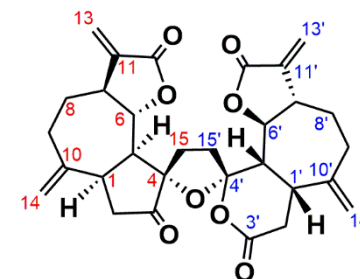
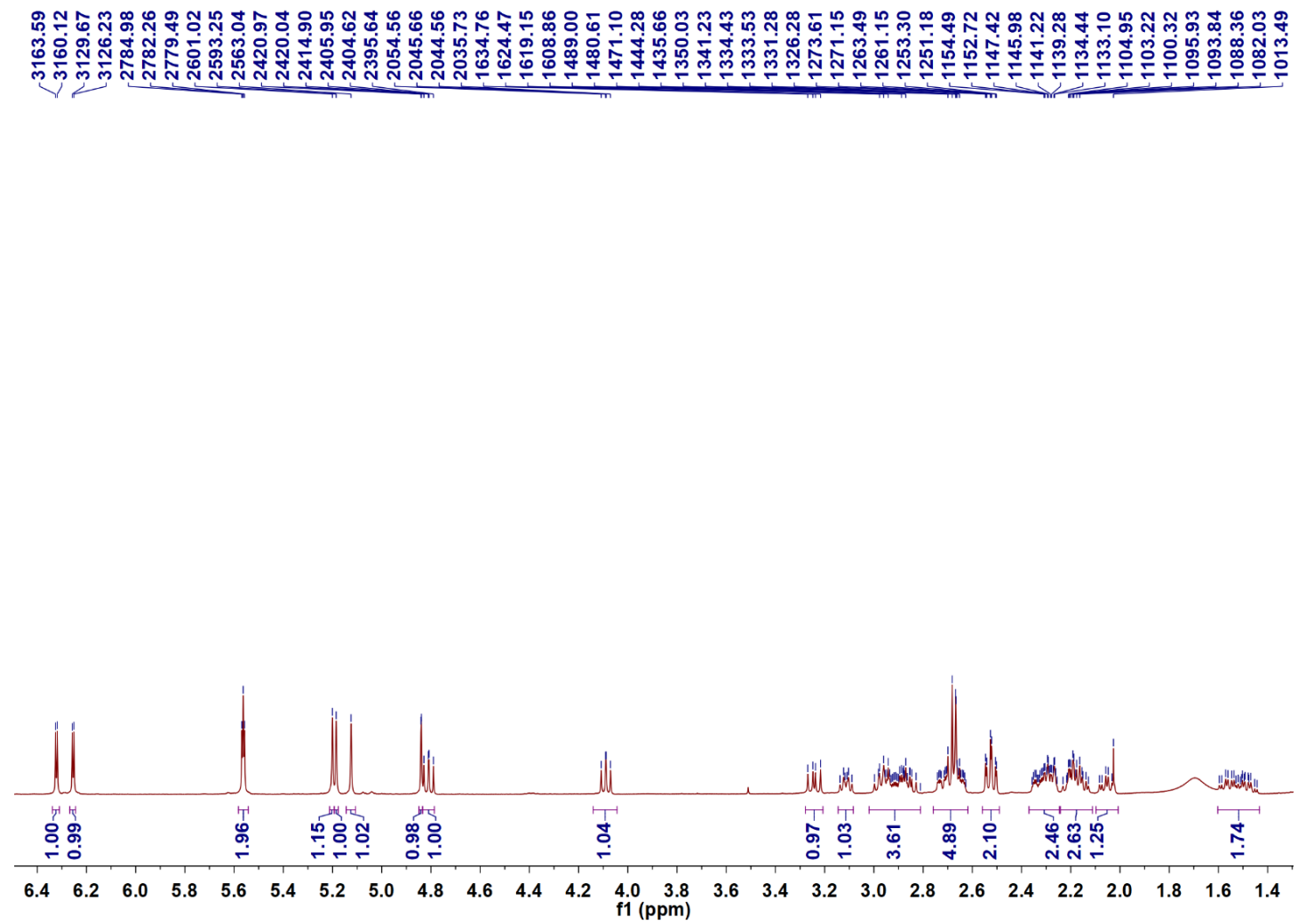


Fig. S4. 1D TOCSY (4.095 PPM) spectrum of compound **1** in CDCl₃

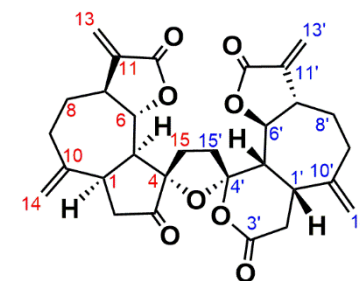
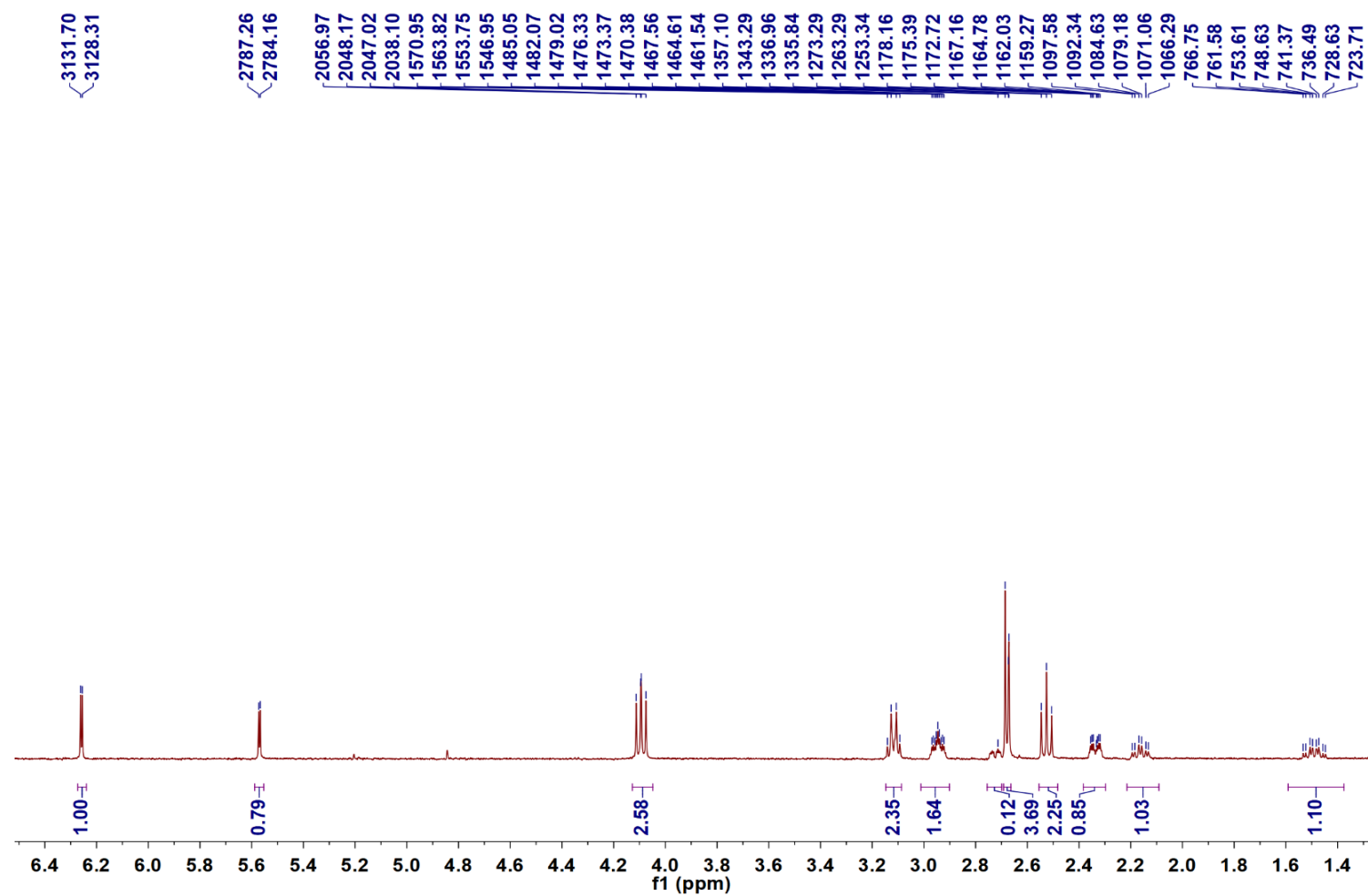


Fig. S5. 1D TOCSY (4.812 PPM) spectrum of compound **1** in CDCl₃

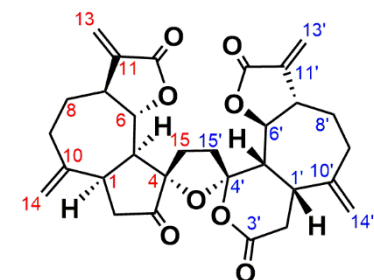
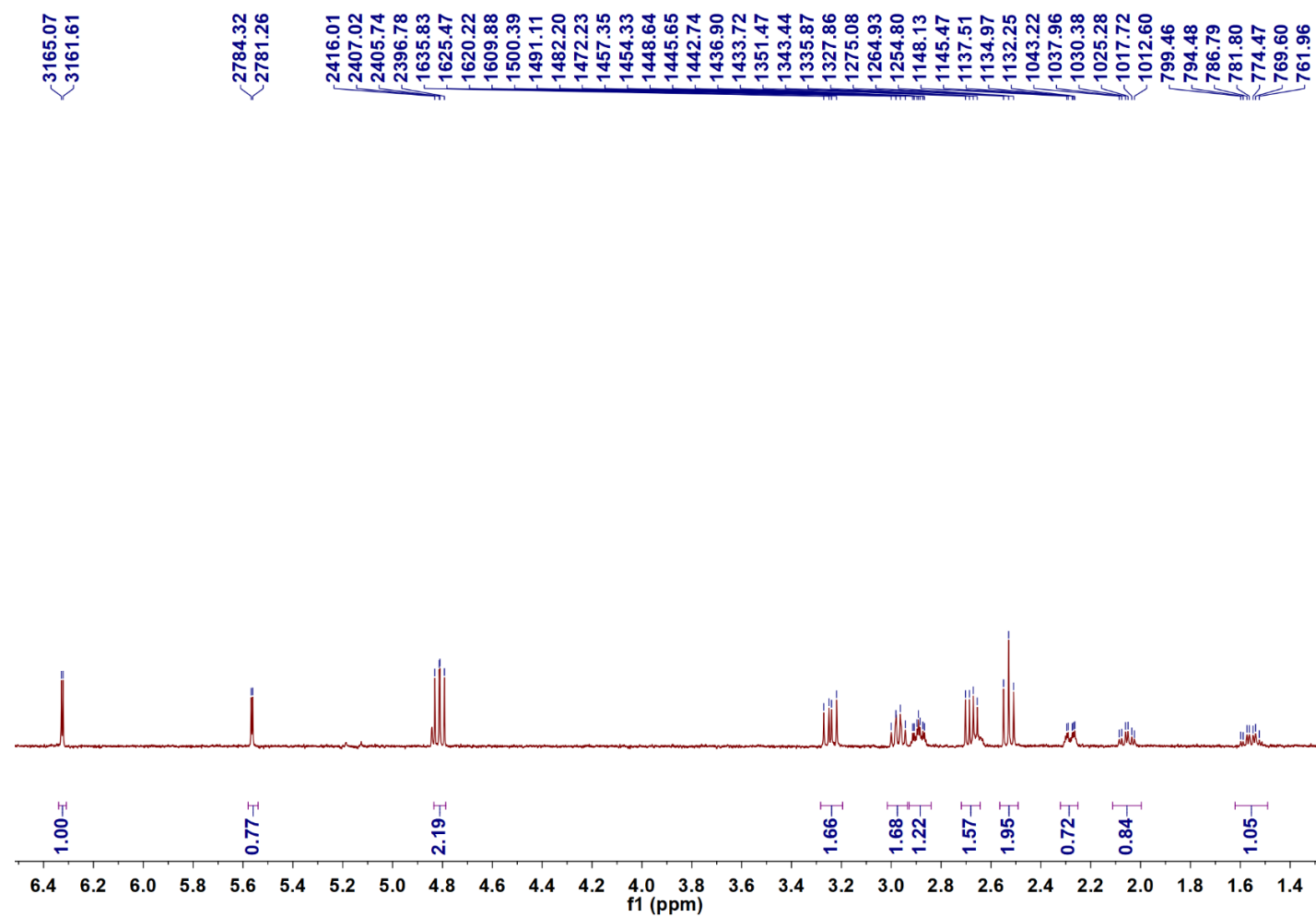


Fig. S6. ^{13}C NMR spectrum of compound **1** in CDCl_3

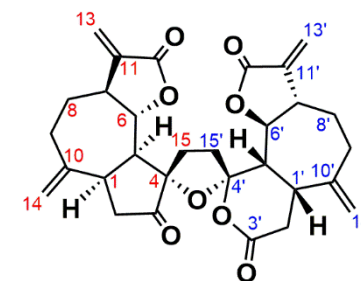
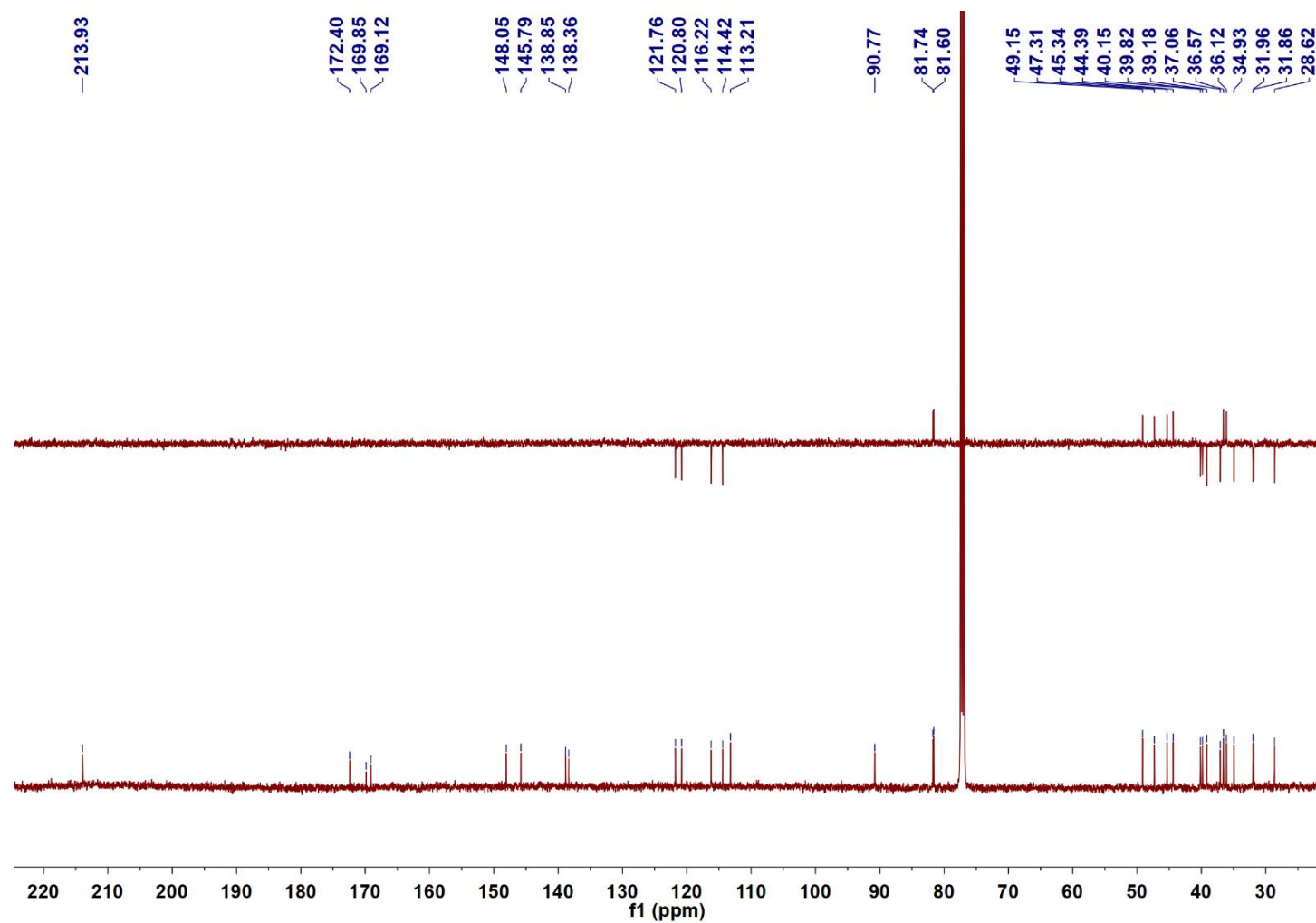


Fig. S7. HSQC spectrum of compound **1** in CDCl₃

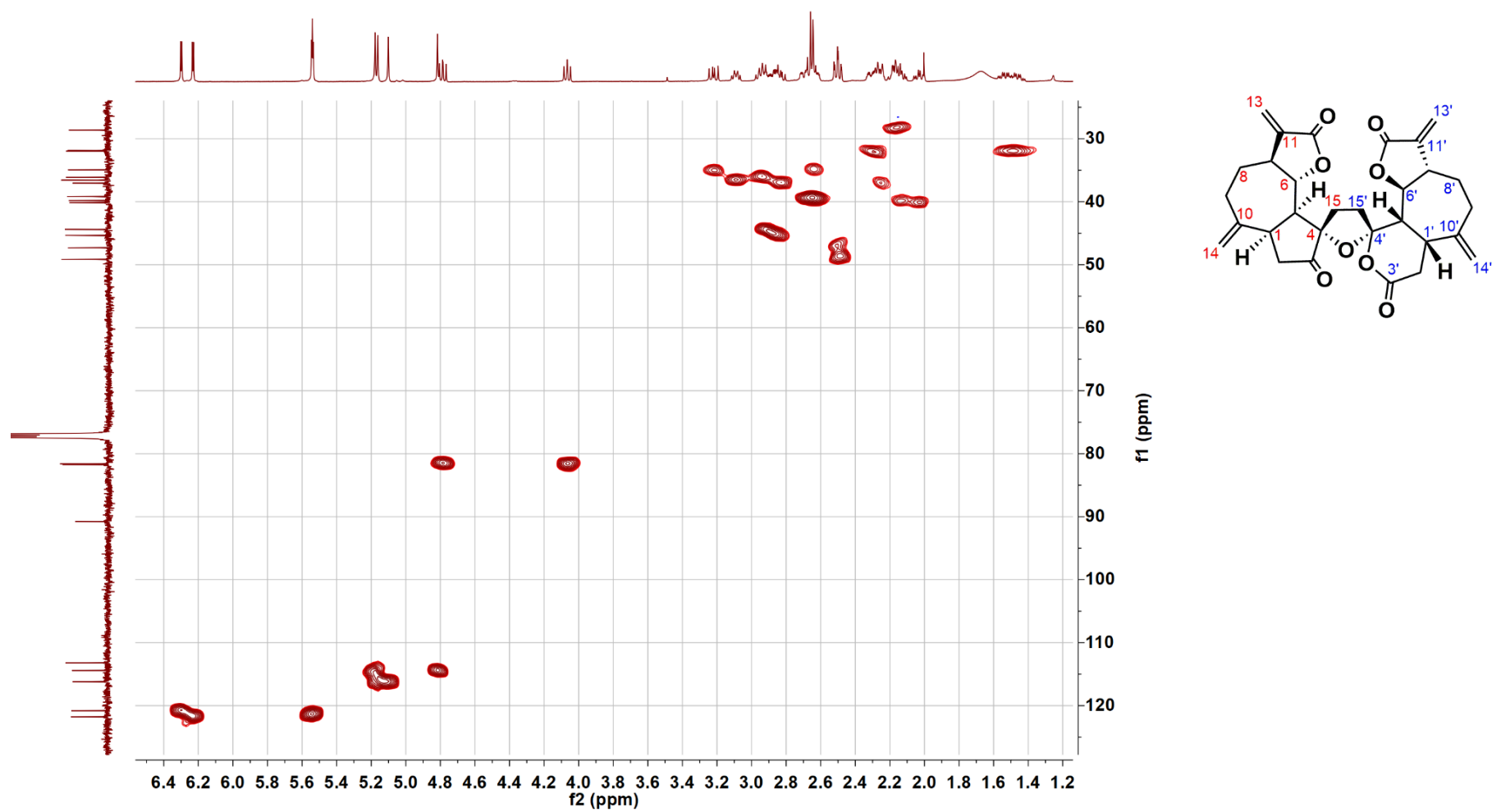


Fig. S8. HMBC spectrum of compound **1** in CDCl₃

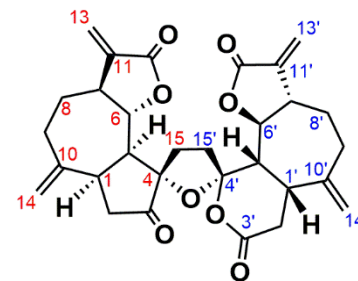
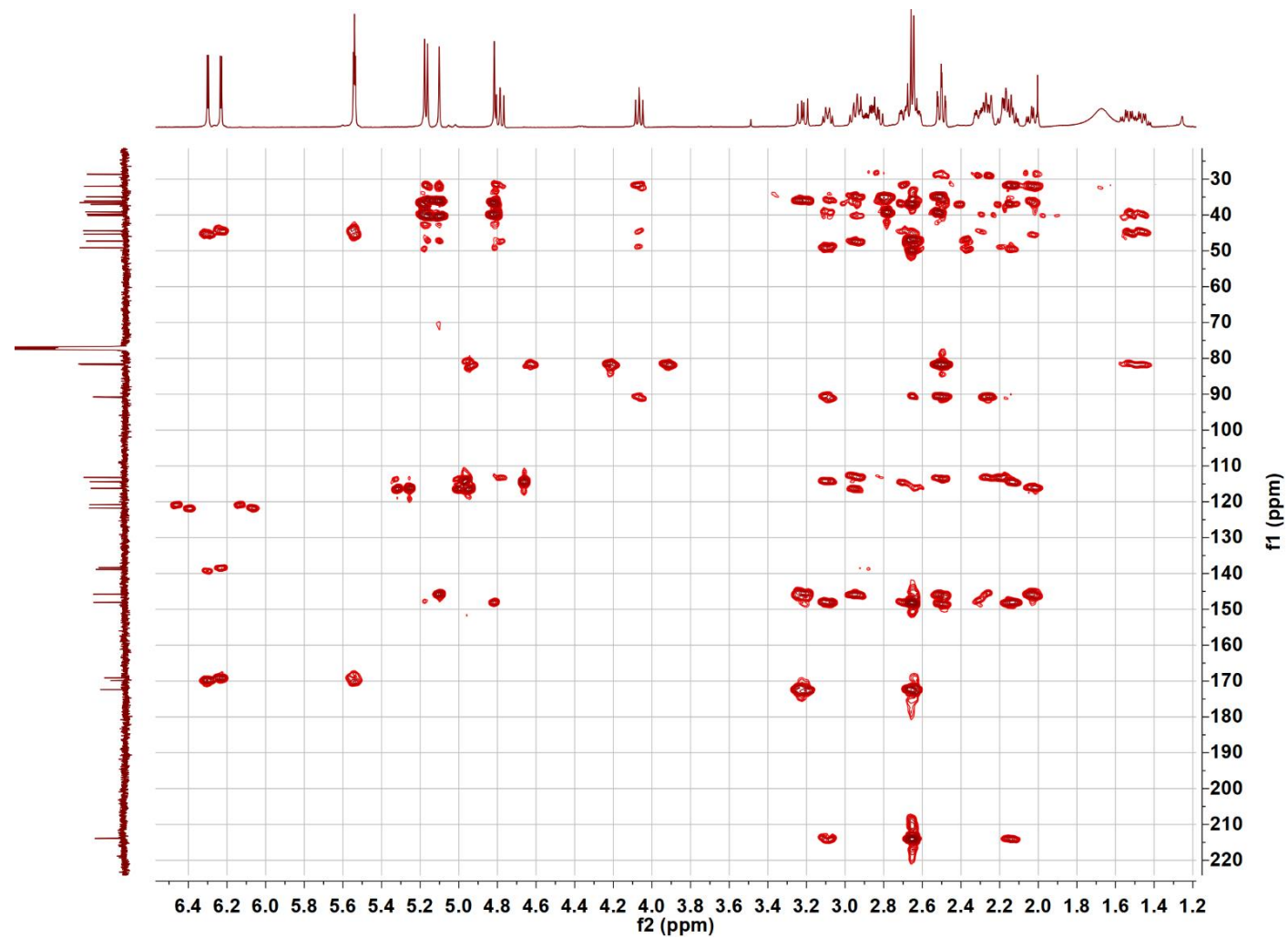


Fig. S9. ^1H - ^1H COSY spectrum of compound **1** in CDCl_3

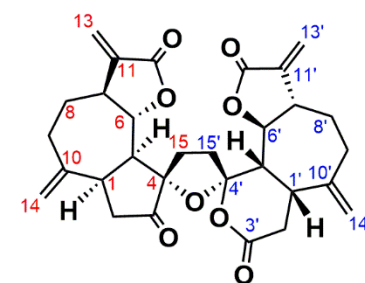
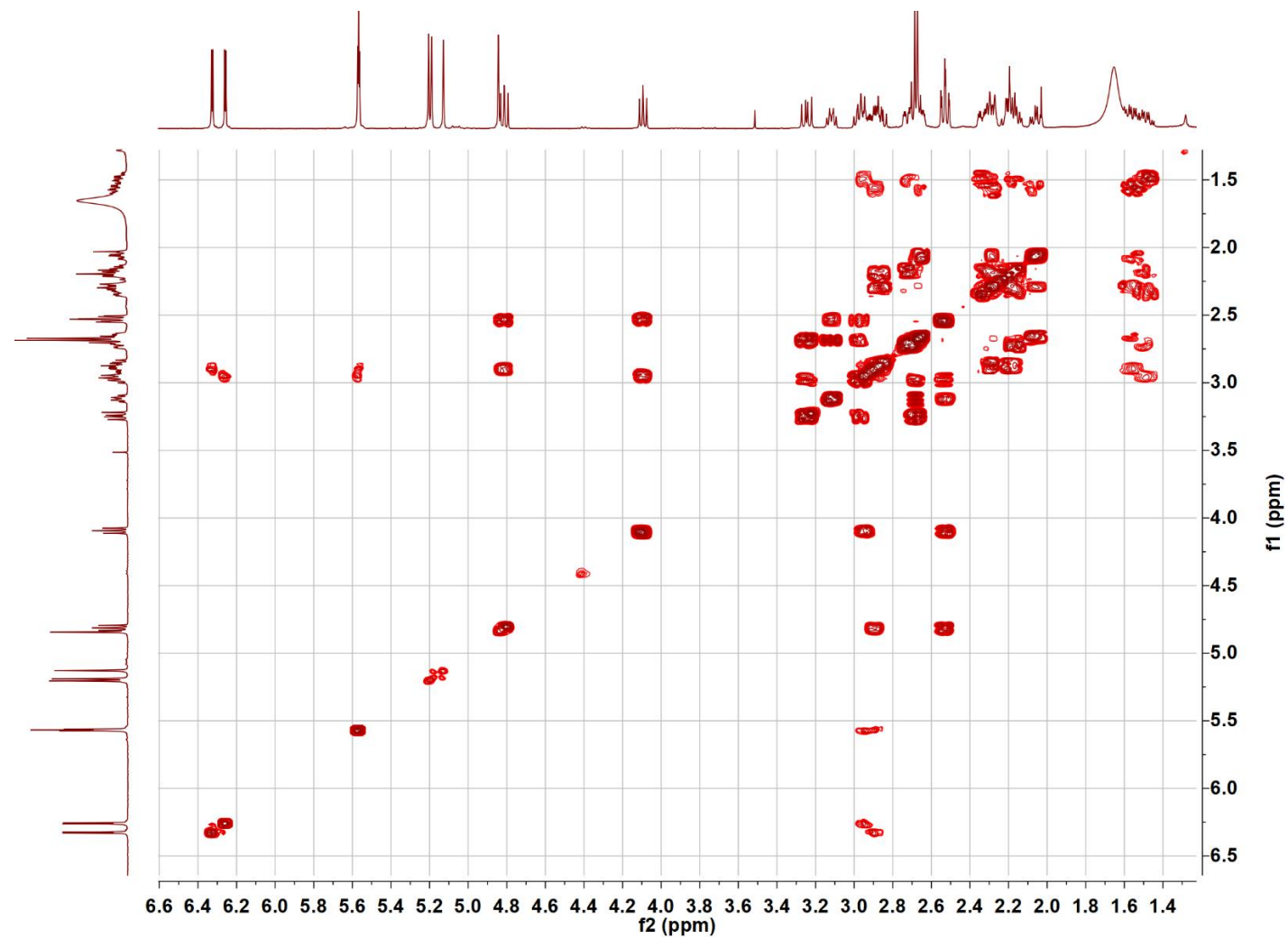


Fig. S10. ROESY spectrum of compound **1** in CDCl₃

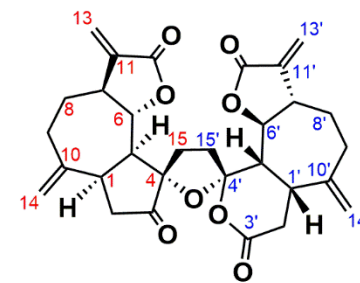
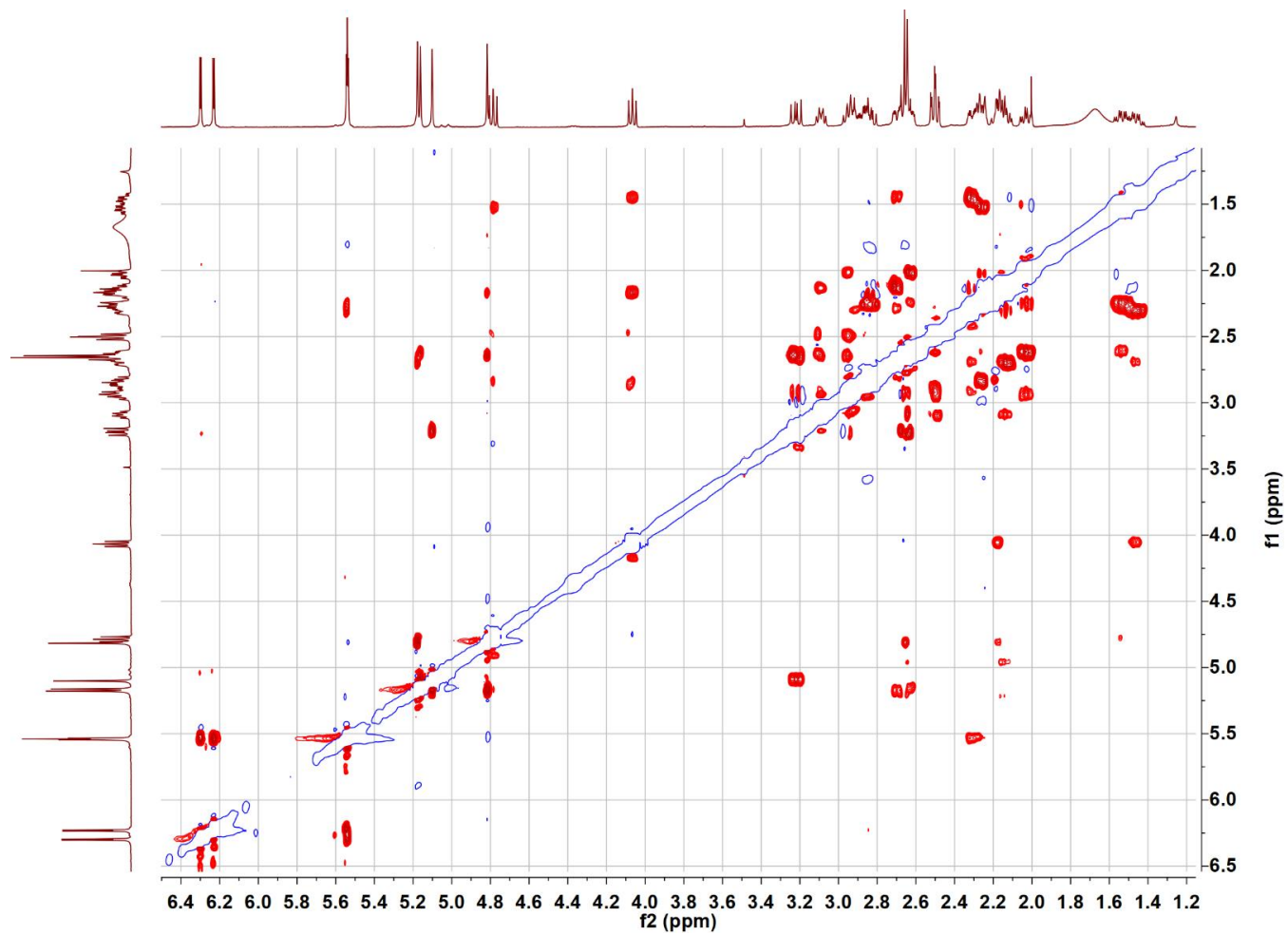


Fig. S11. IR spectrum of compound **2**

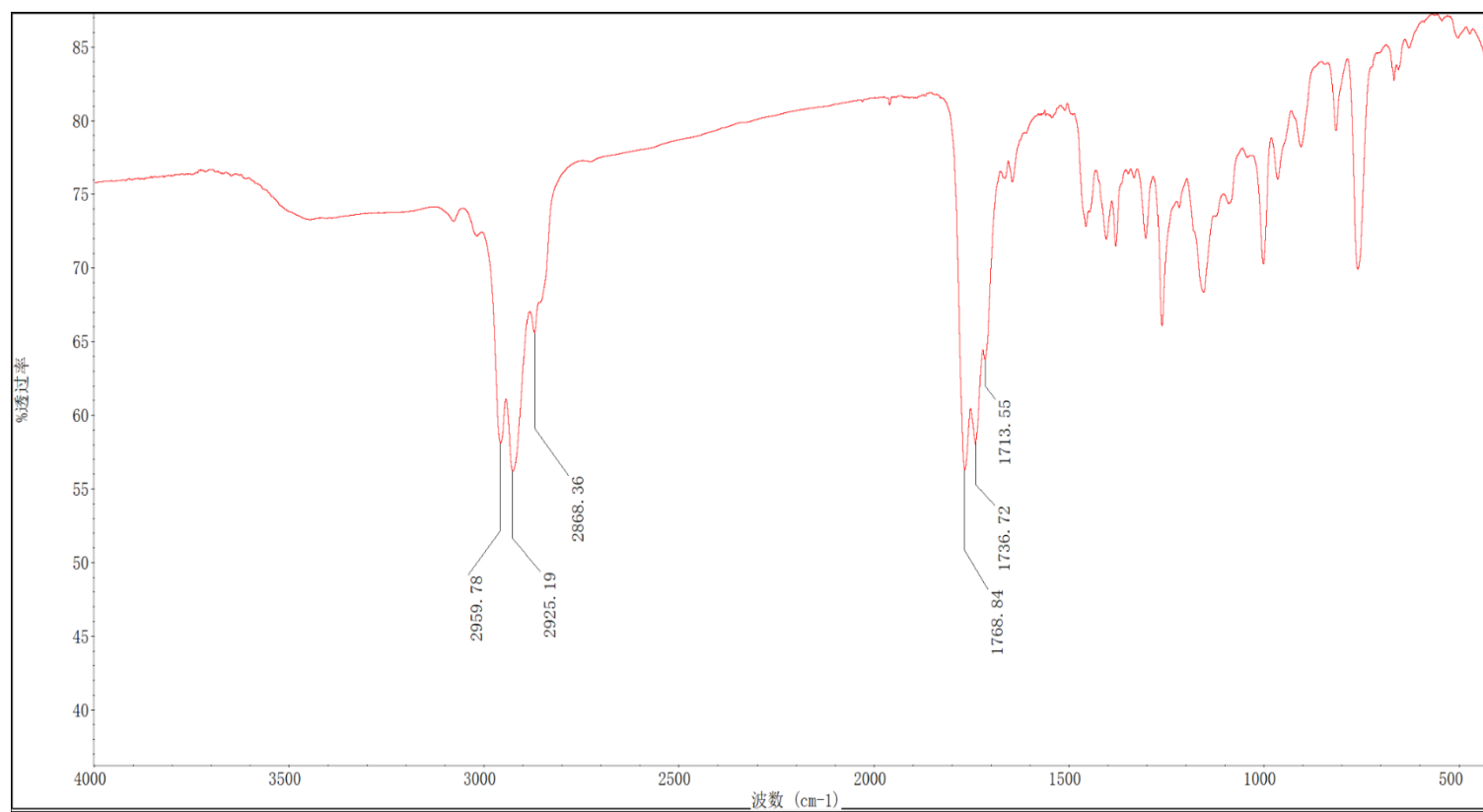


Fig. S12. HRESIMS data of compound **2**

Elemental Composition Report

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

277 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-100 H: 0-100 O: 0-50 Na: 0-1

Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
545.2147	545.2151	-0.4	-0.7	13.5	129.1	0.126	88.14	C30 H34 O8 Na
	545.2175	-2.8	-5.1	16.5	131.1	2.169	11.43	C32 H33 O8
	545.2117	3.0	5.5	25.5	134.4	5.437	0.44	C39 H29 O3

AM-BIF3C2C1-pos
20180823-SA021 1604 (6.494)

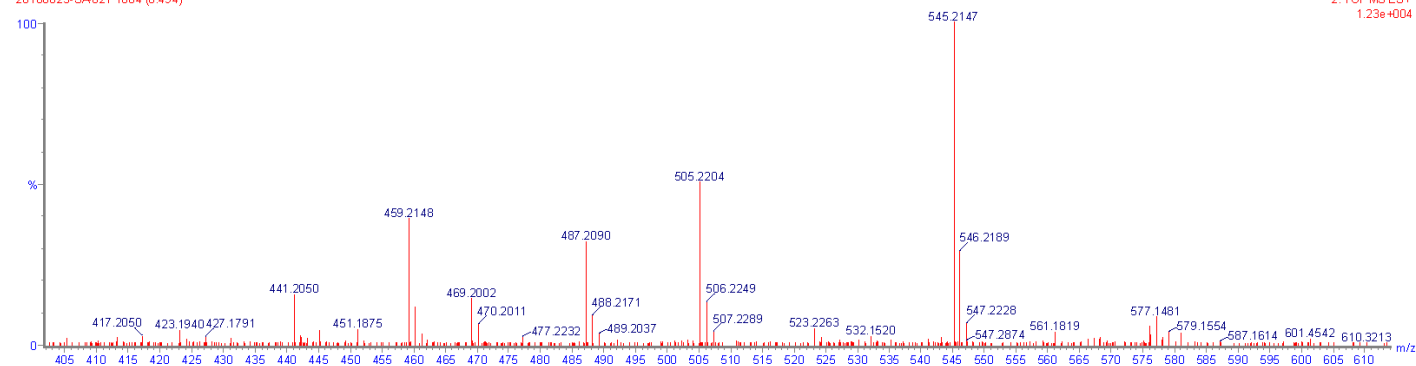


Fig. S13. ^1H NMR spectrum of compound **2** in CD_3OD

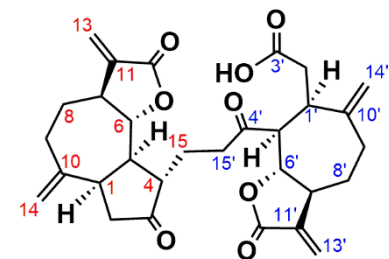
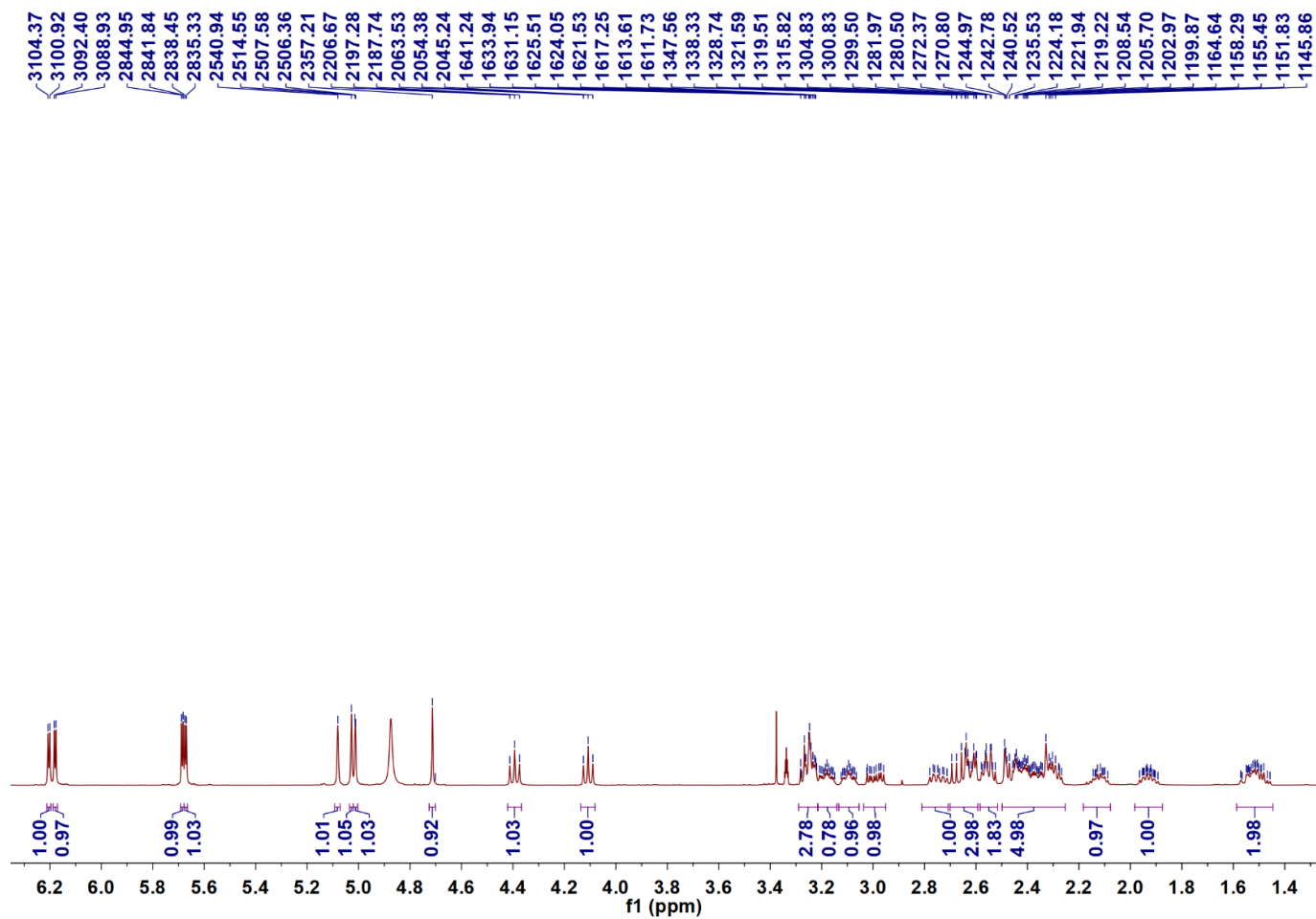


Fig. S14. 1D TOCSY (4.108 PPM) spectrum of compound **2** in CD₃OD

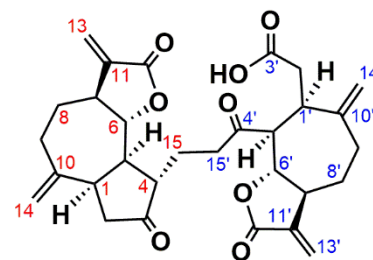
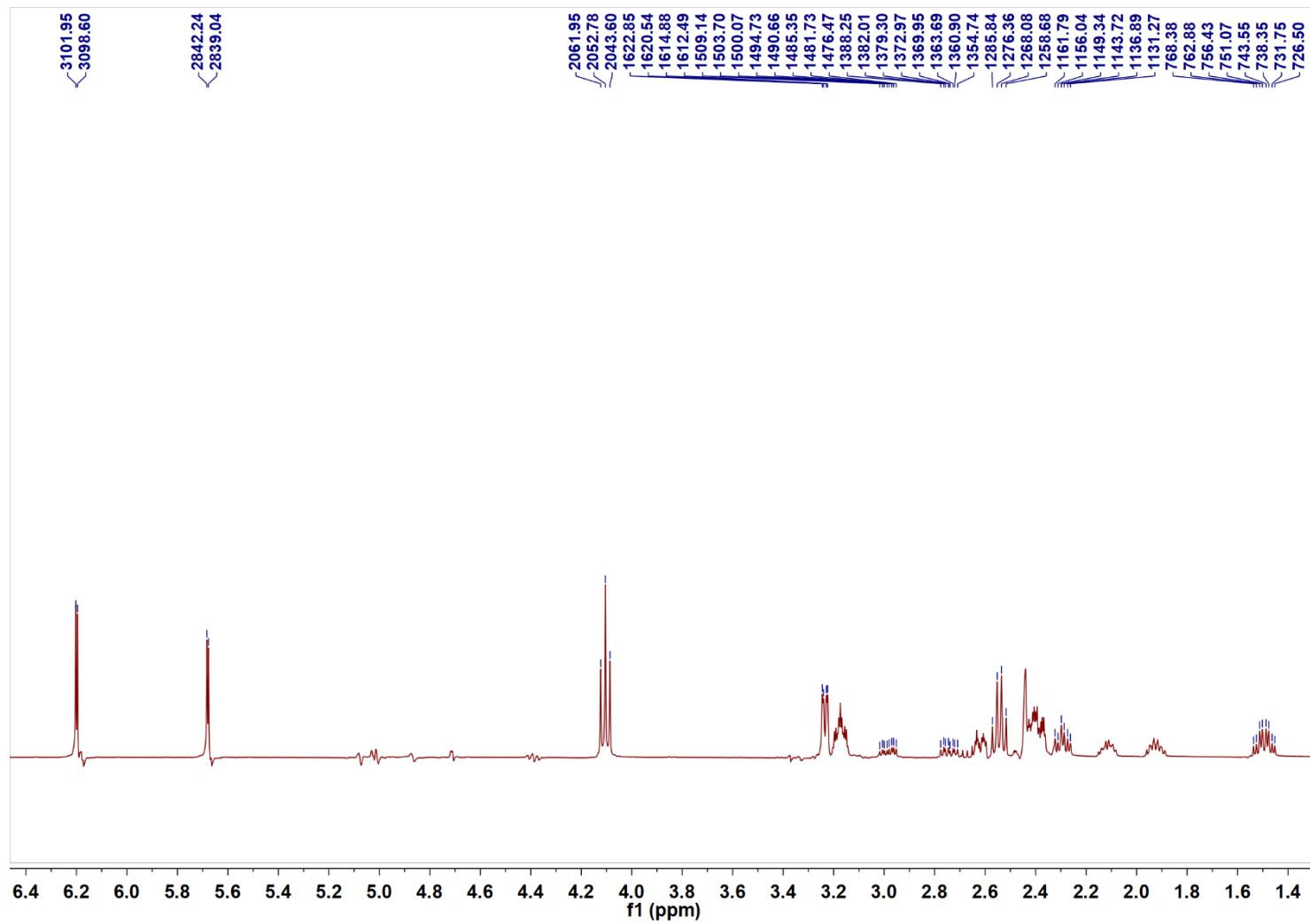


Fig. S15. 1D TOCSY (4.389 PPM) spectrum of compound **2** in CD₃OD

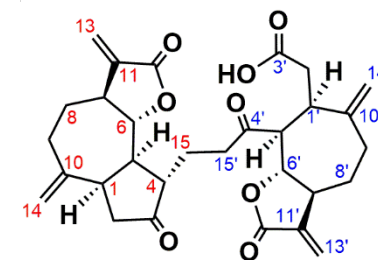
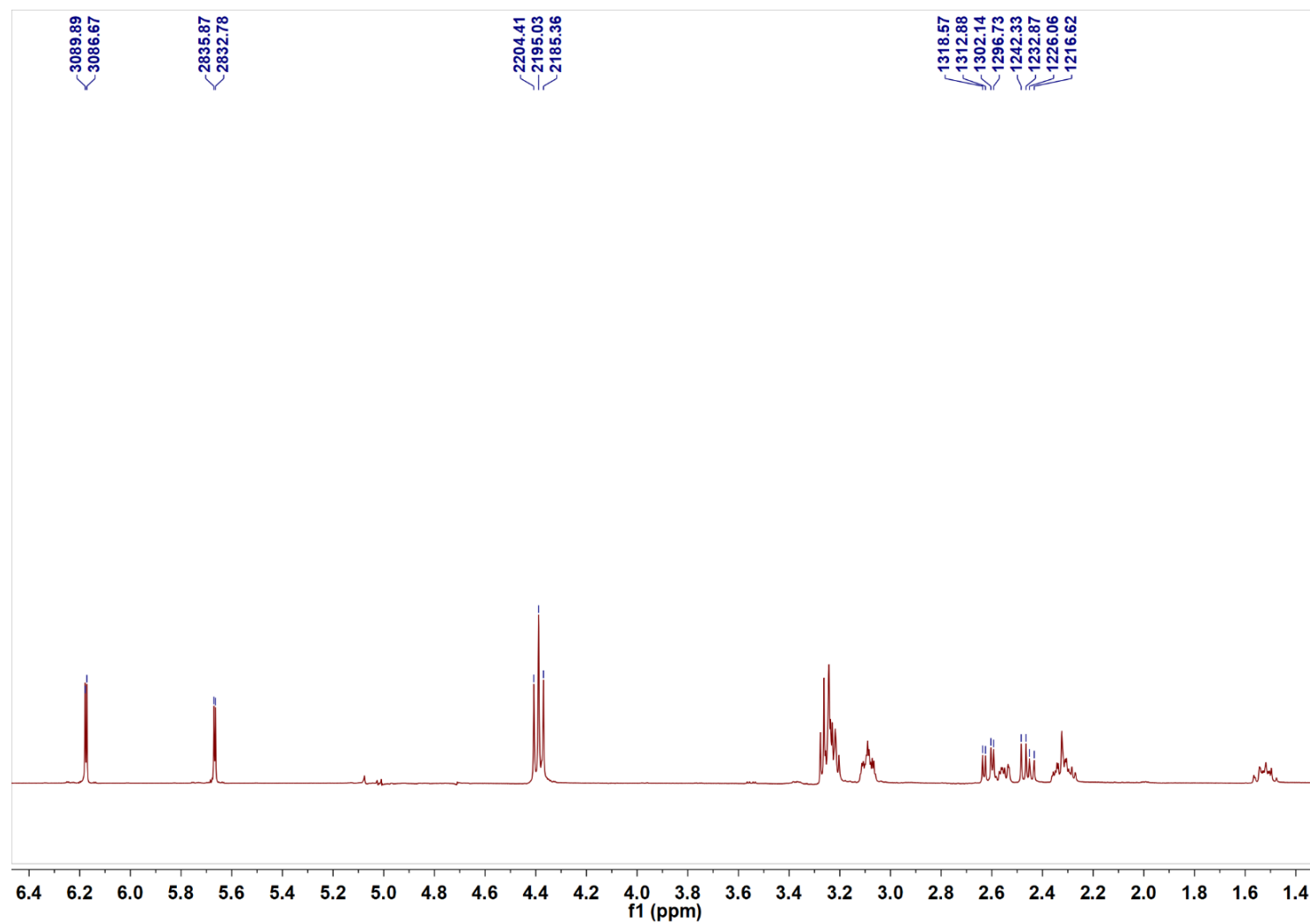


Fig. S16. ^{13}C NMR spectrum of compound **2** in CD_3OD

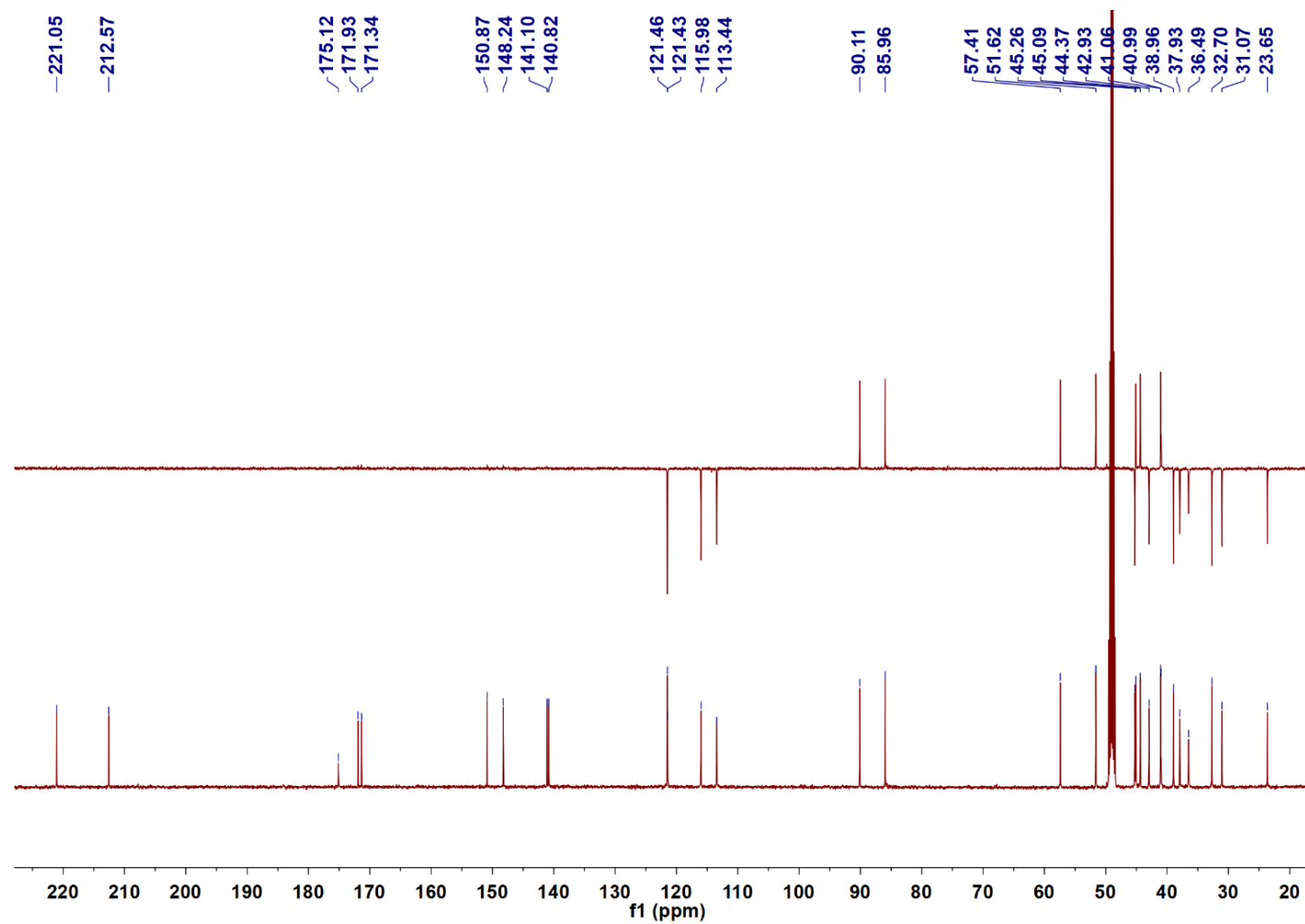


Fig. S17. HSQC spectrum of compound **2** in CD₃OD

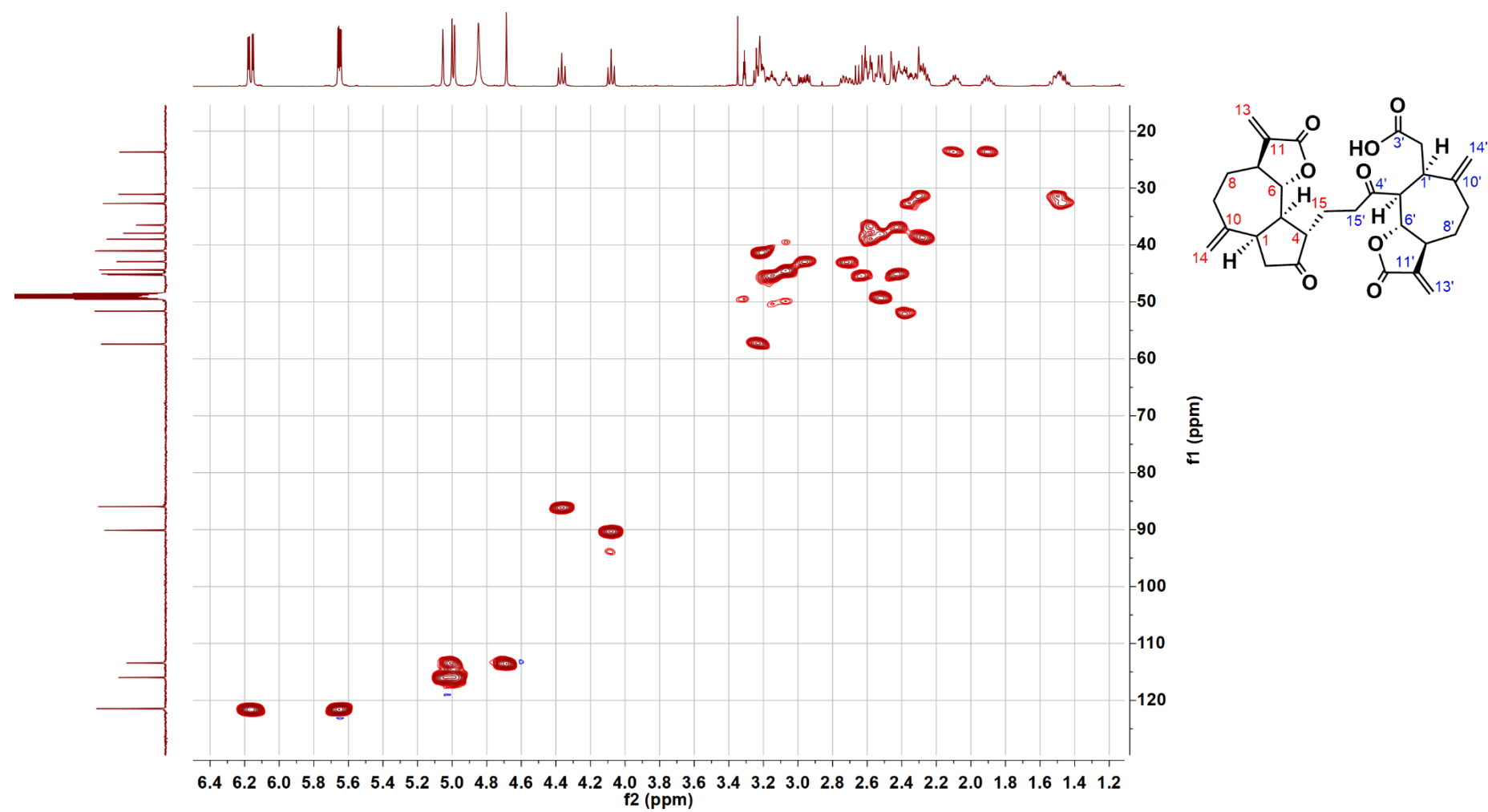


Fig. S18. HMBC spectrum of compound **2** in CD₃OD

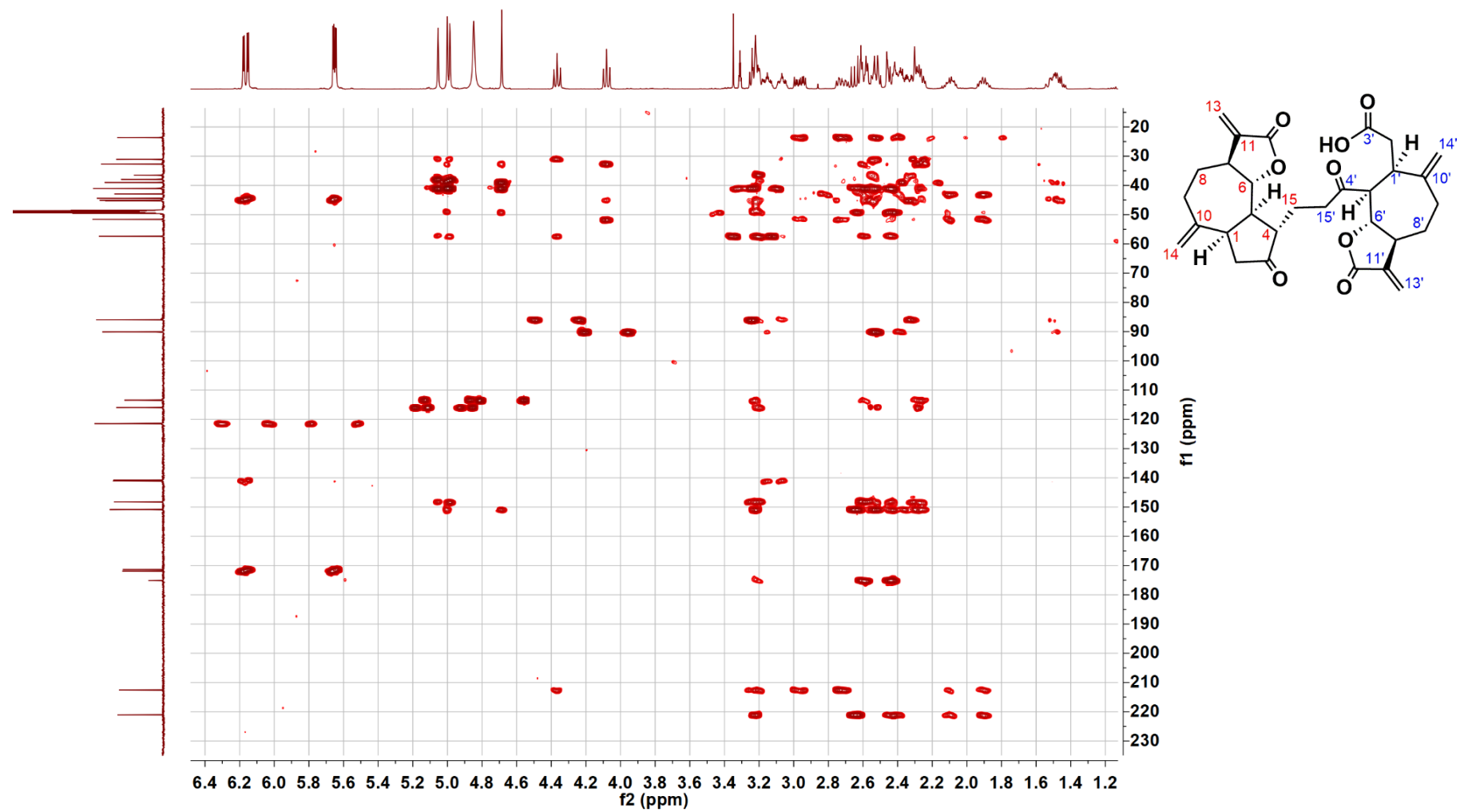


Fig. S19. ^1H - ^1H COSY spectrum of compound **2** in CD_3OD

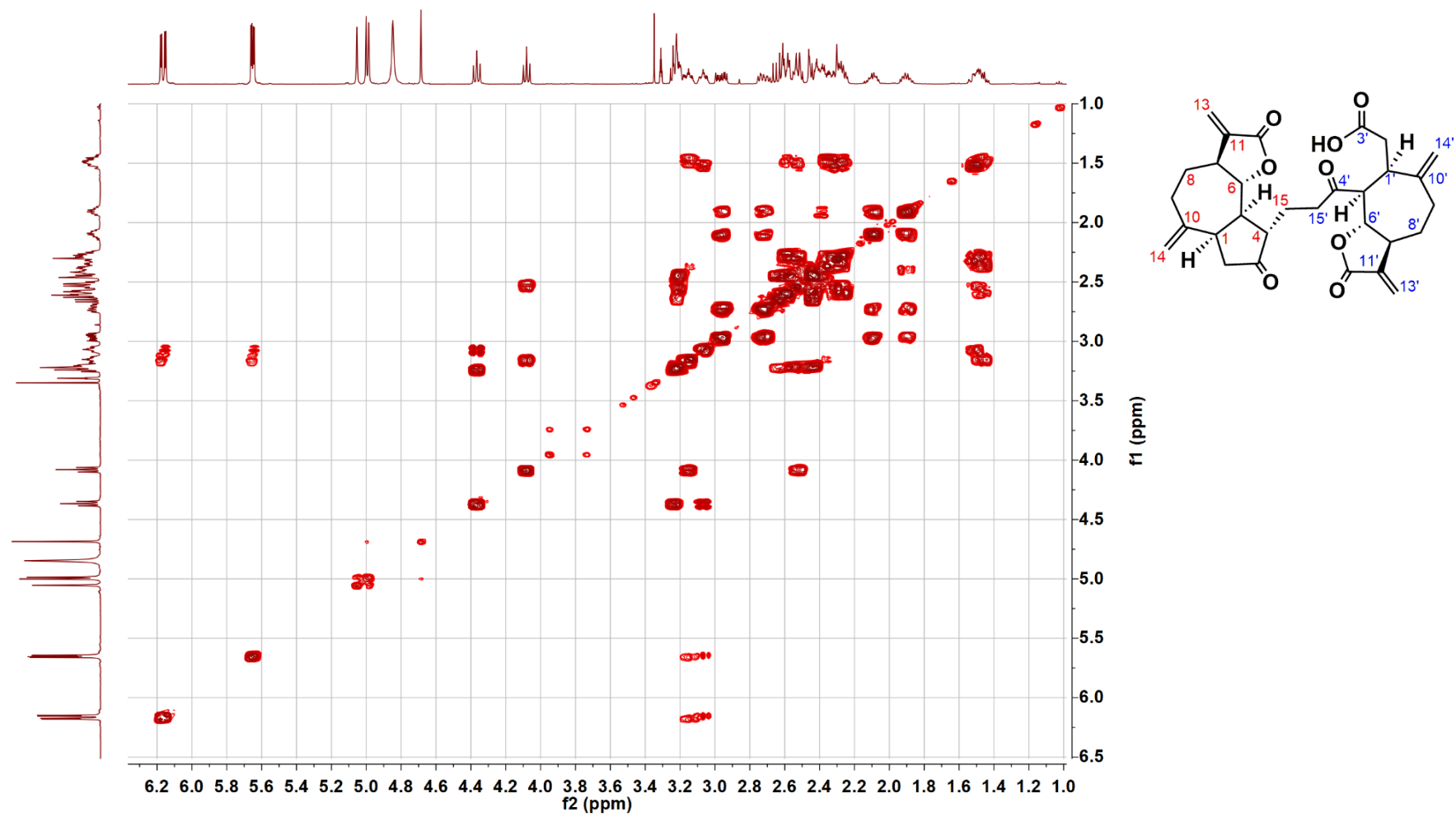


Fig. S20. ROESY spectrum of compound **2** in CD₃OD

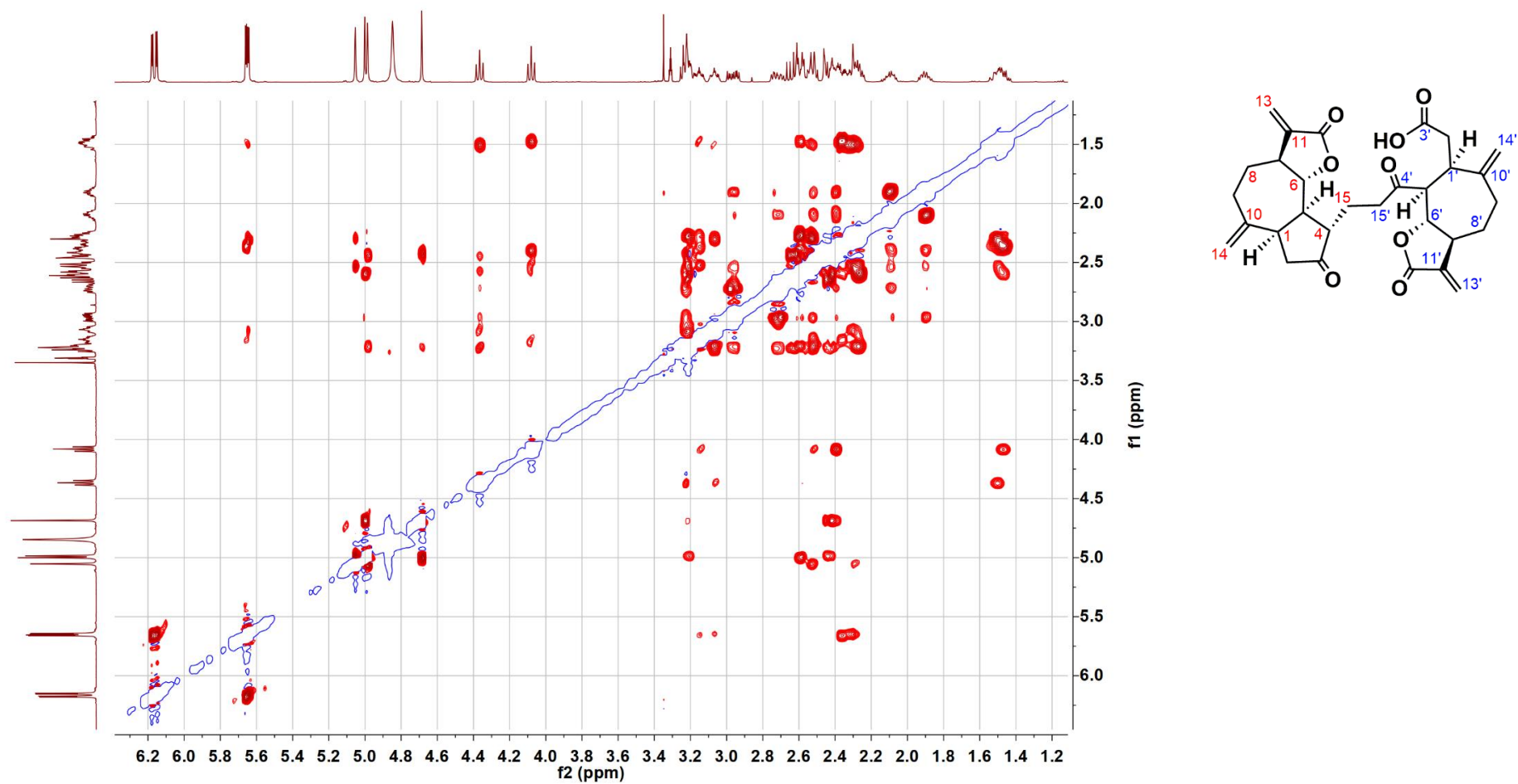


Fig. S21 HRESIMS data of compound **3**

Elemental Composition Report

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

258 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

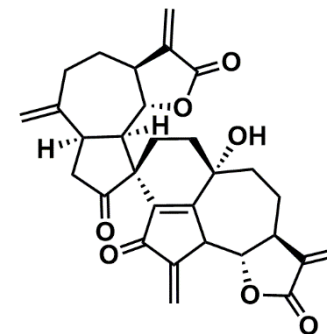
Elements Used:

C: 0-100 H: 0-100 O: 0-50 Na: 0-1

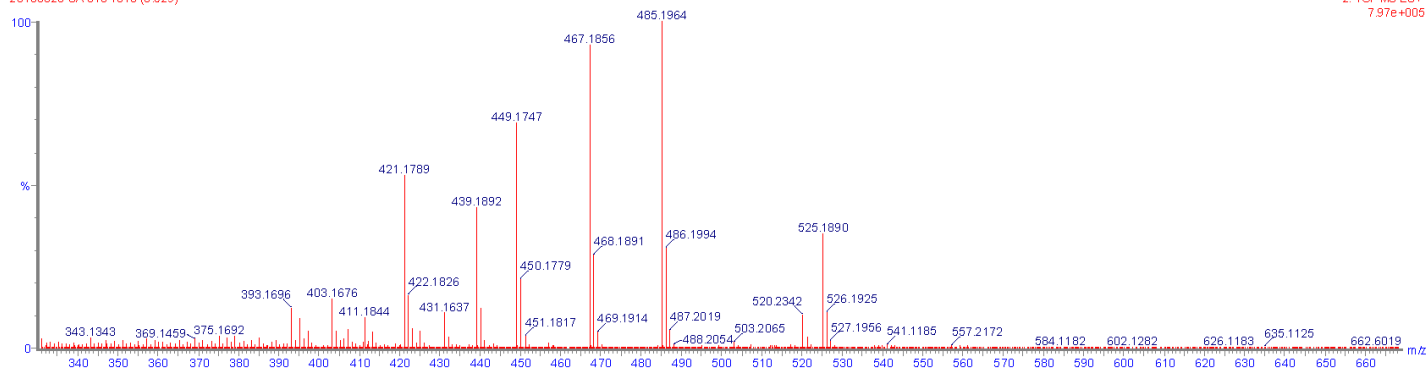
Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
525.1890	525.1889	0.1	0.2	15.5	425.6	0.089	91.47	C ₃₀ H ₃₀ O ₇ Na
	525.1913	-2.3	-4.4	18.5	428.0	2.487	8.31	C ₃₂ H ₂₉ O ₇
	525.1855	3.5	6.7	27.5	431.6	6.128	0.22	C ₃₉ H ₂₅ O ₂



AM-B1F3D2-pos
20180823-SA-010 1613 (6 529)



2: TOF MS ES+
7.97e+005

Fig. S22 ^1H NMR spectrum of compound **3** in CDCl_3

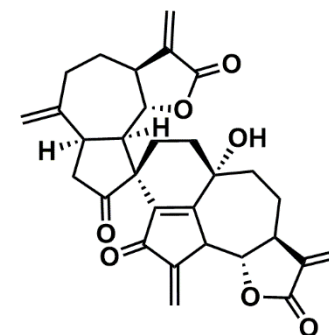
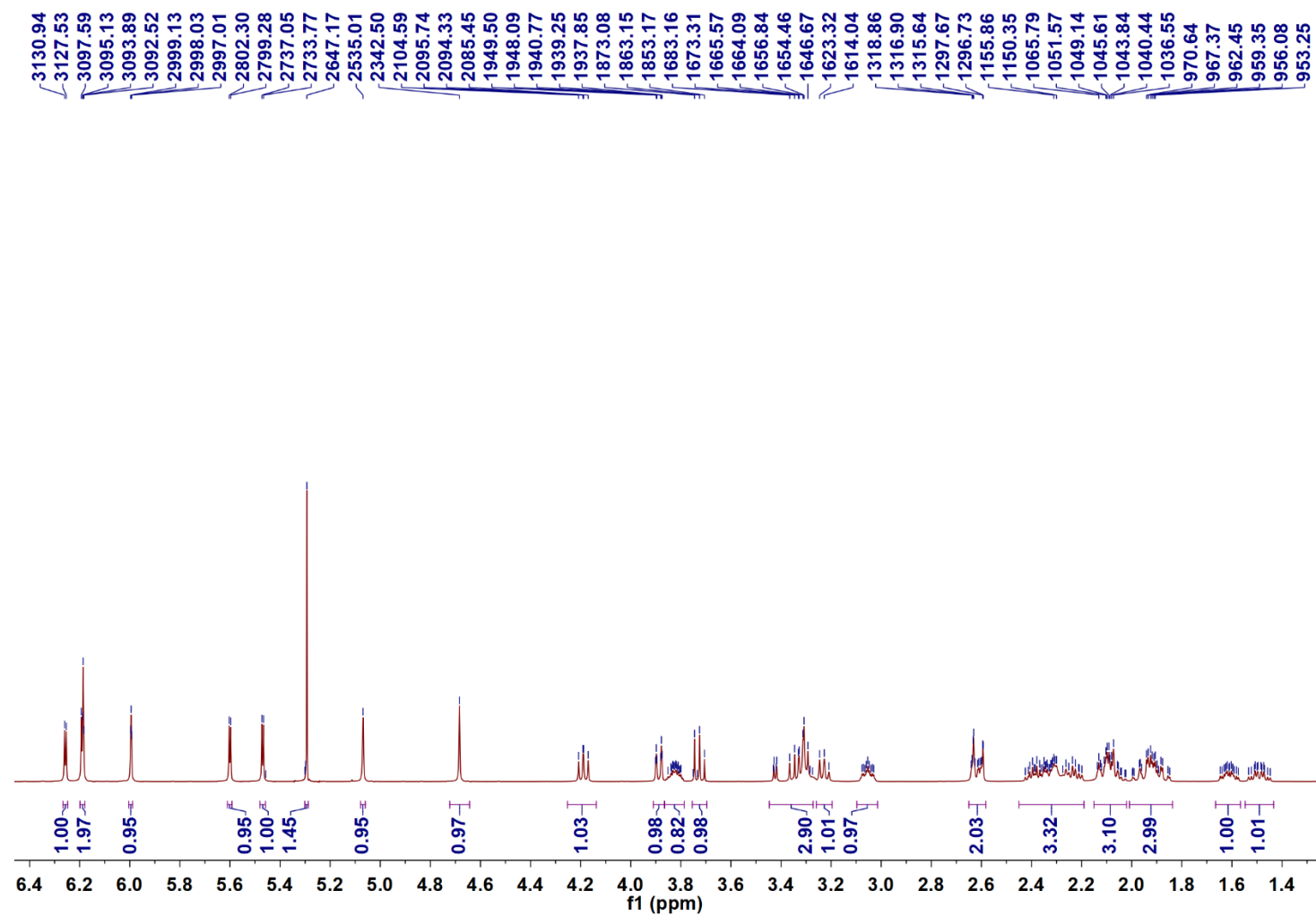


Fig. S23 ^{13}C NMR spectrum of compound **3** in CDCl_3

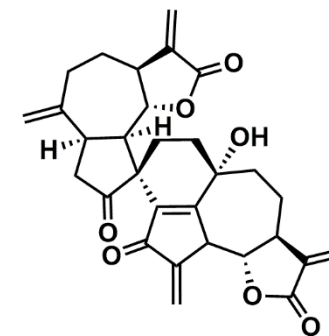
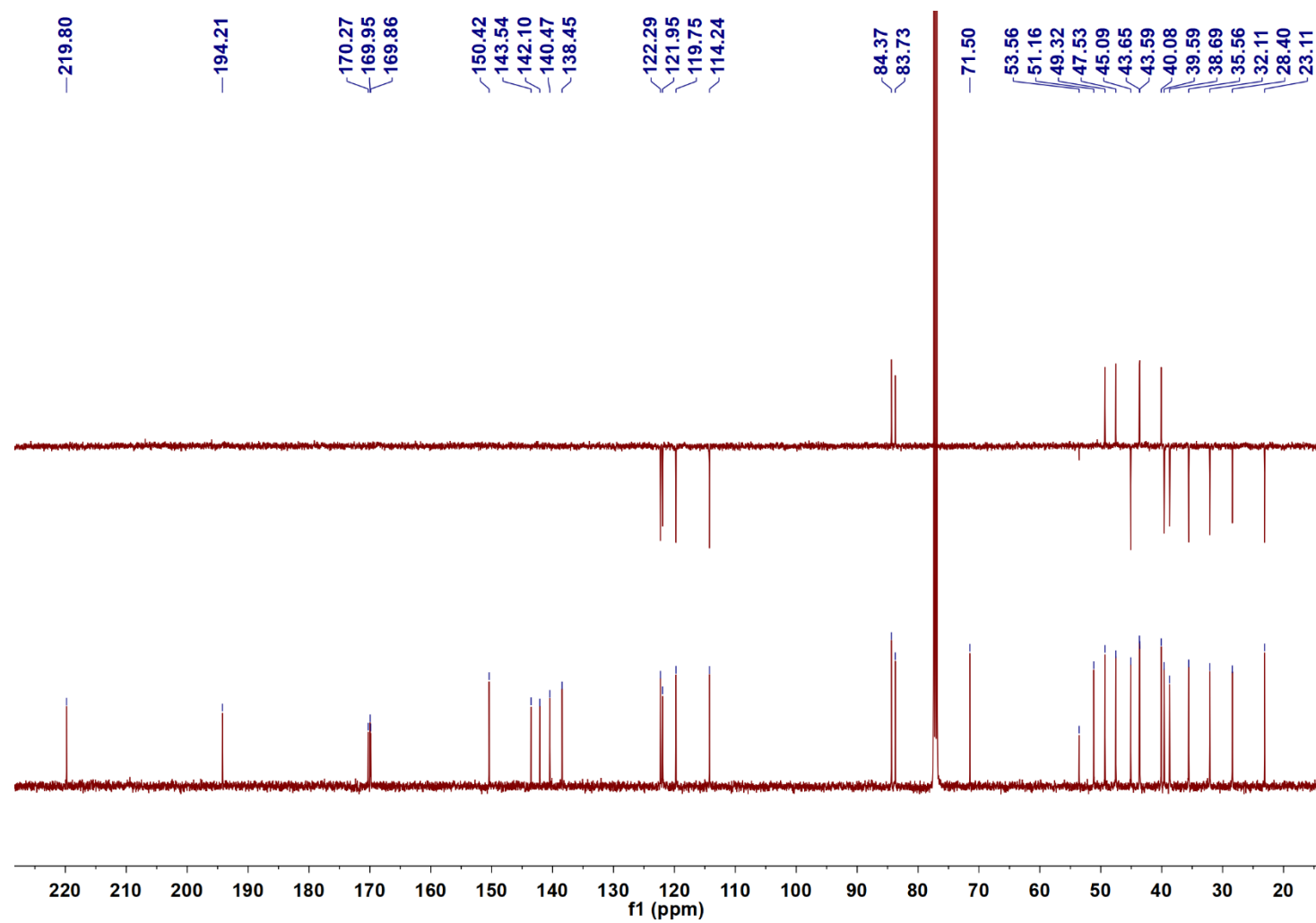


Fig. S24. HRESIMS data of compound **4**

Elemental Composition Report

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

258 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Elements Used:

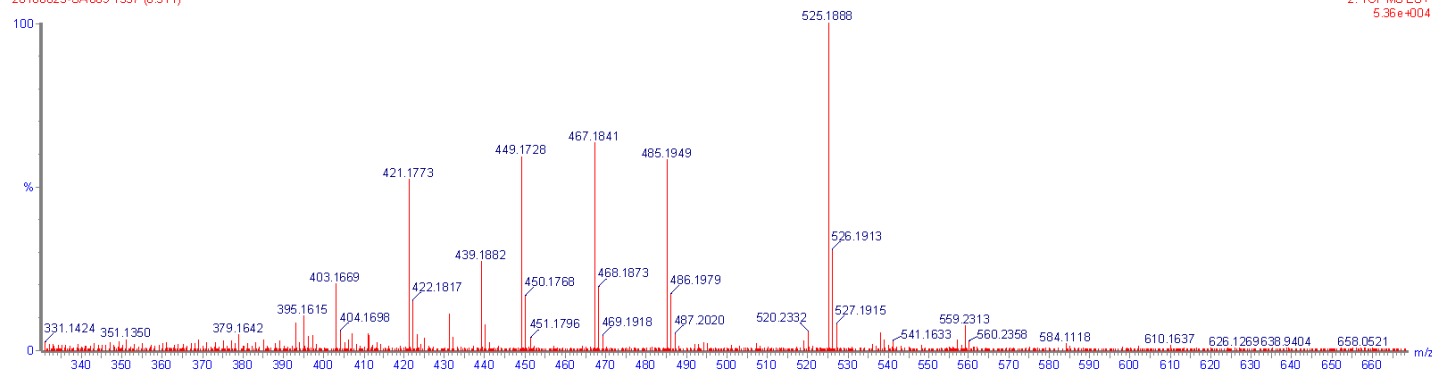
C: 0-100 H: 0-100 O: 0-50 Na: 0-1

Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
525.1888	525.1889	-0.1	-0.2	15.5	218.0	0.166	84.67	C30 H30 O7 Na
	525.1913	-2.5	-4.8	18.5	219.8	1.903	14.91	C32 H29 O7
	525.1855	3.3	6.3	27.5	223.3	5.472	0.42	C39 H25 O2

AMB1F3D1-pos
20180823-SA009 1557 (6.311)



2: TOF MS ES+
5.36e+004

Fig. S25 ^1H NMR spectrum of compound **4** in CDCl_3

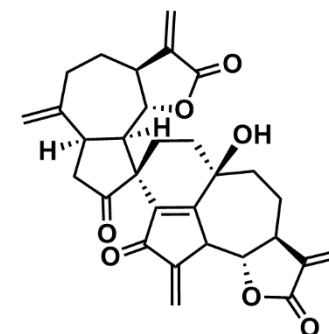
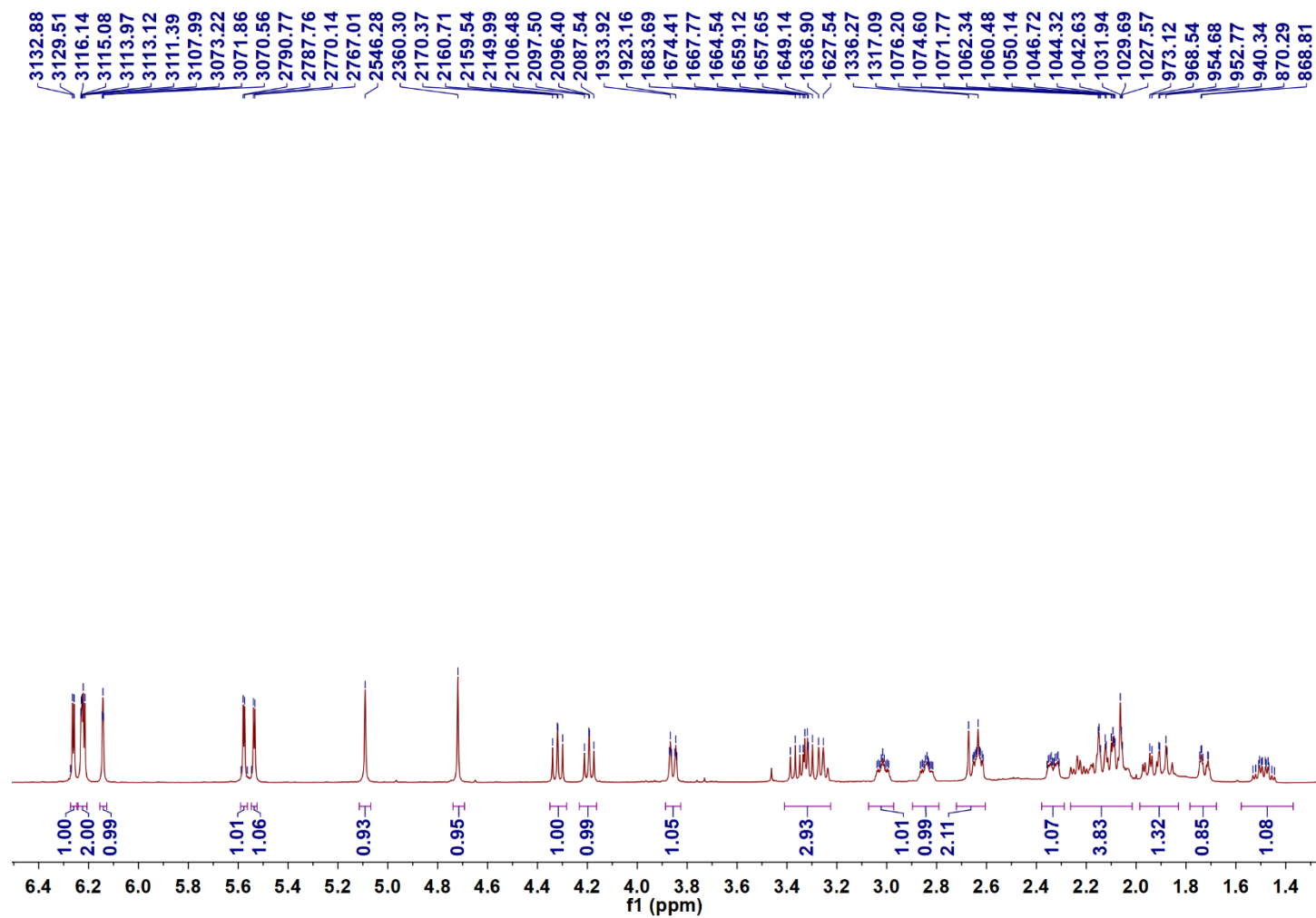


Fig. S26 ^{13}C NMR spectrum of compound **4** in CDCl_3

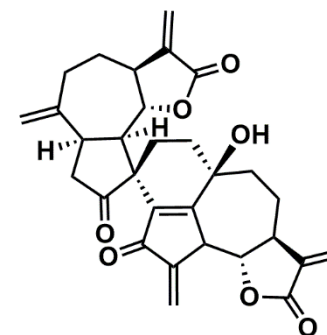
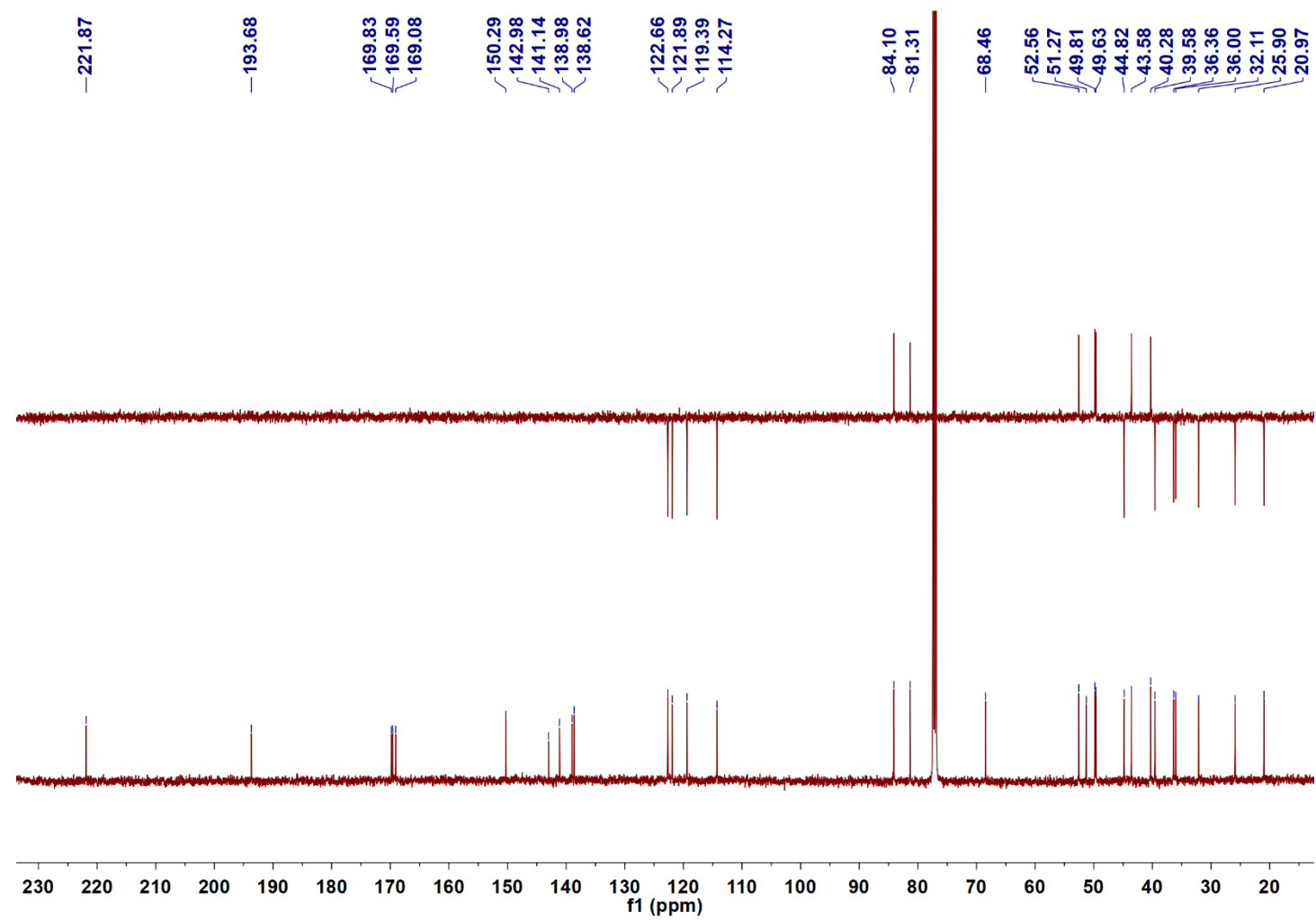


Fig. S27. HRESIMS data of compound **5**

Elemental Composition Report

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

275 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-100 H: 0-100 O: 0-50 Na: 0-1

Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
531.2360	531.2359	0.1	0.2	12.5	282.2	0.046	95.47	C30 H36 O7 Na
	531.2383	-2.3	-4.3	15.5	285.2	3.122	4.41	C32 H35 O7
	531.2324	3.6	6.8	24.5	288.8	6.700	0.12	C39 H31 O2

AMB1F3C2B1-pos
20180823-SA-013 1597 (6.467)

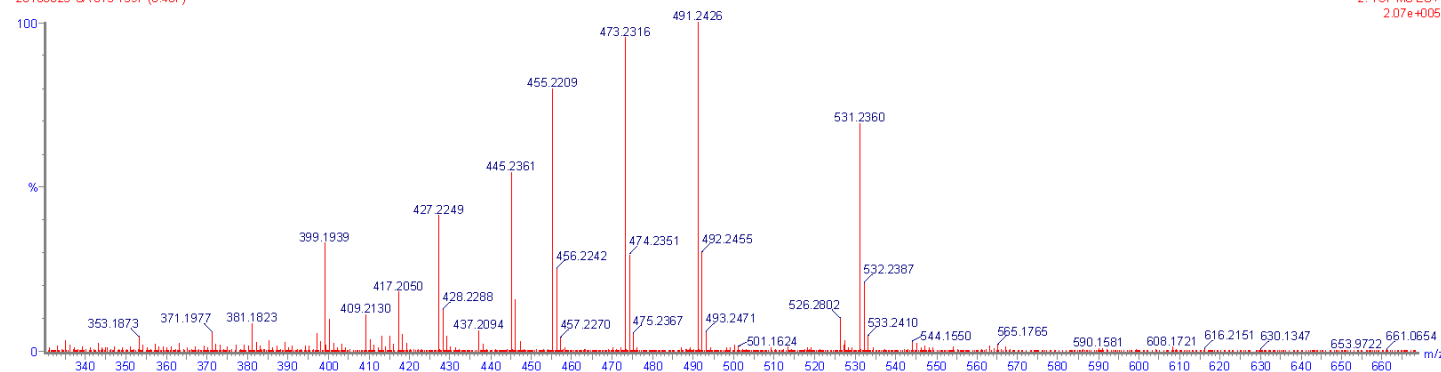


Fig. S28 ^1H NMR spectrum of compound **5** in CDCl_3

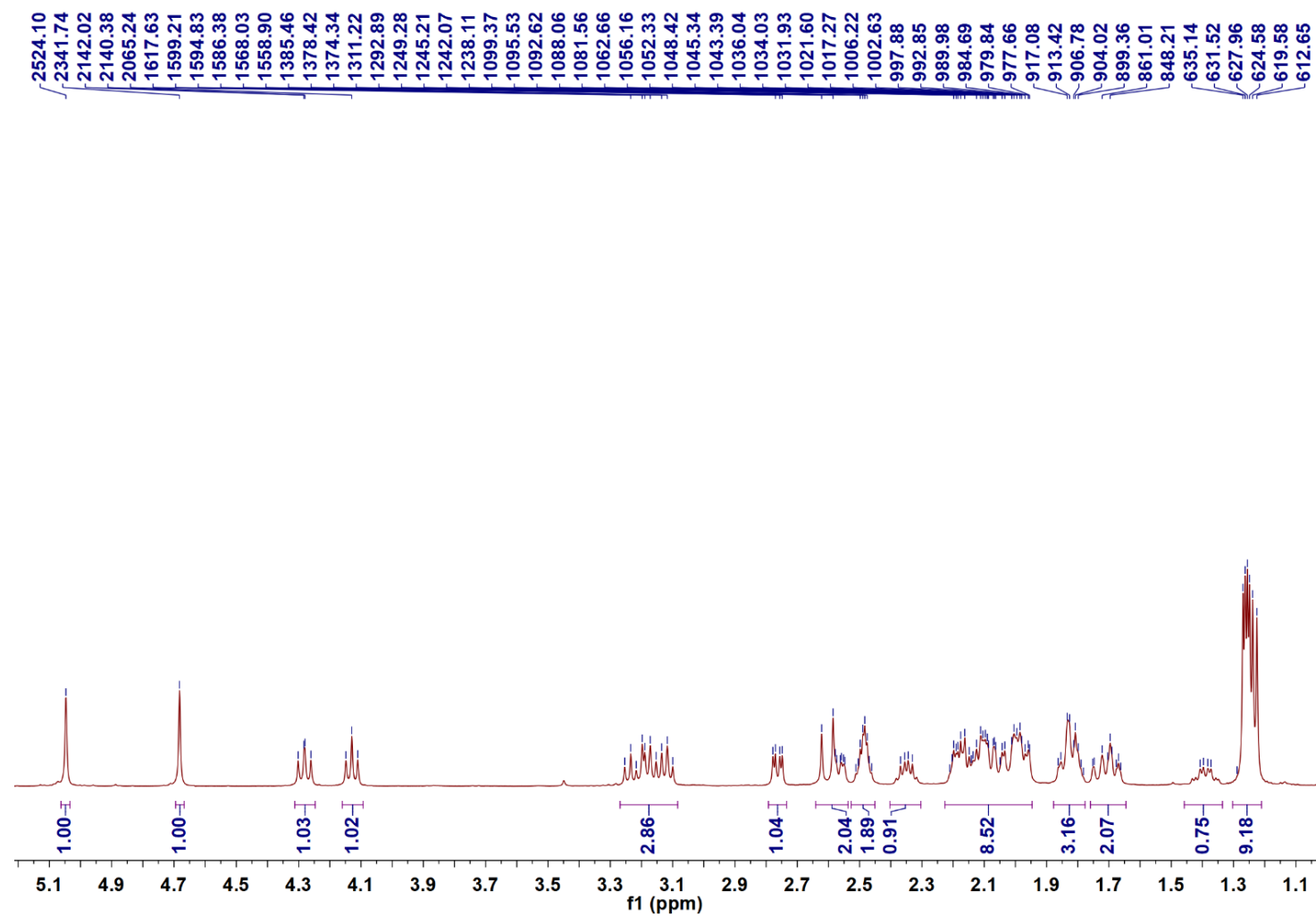


Fig. S29. ^{13}C NMR spectrum of compound **5** in CDCl_3

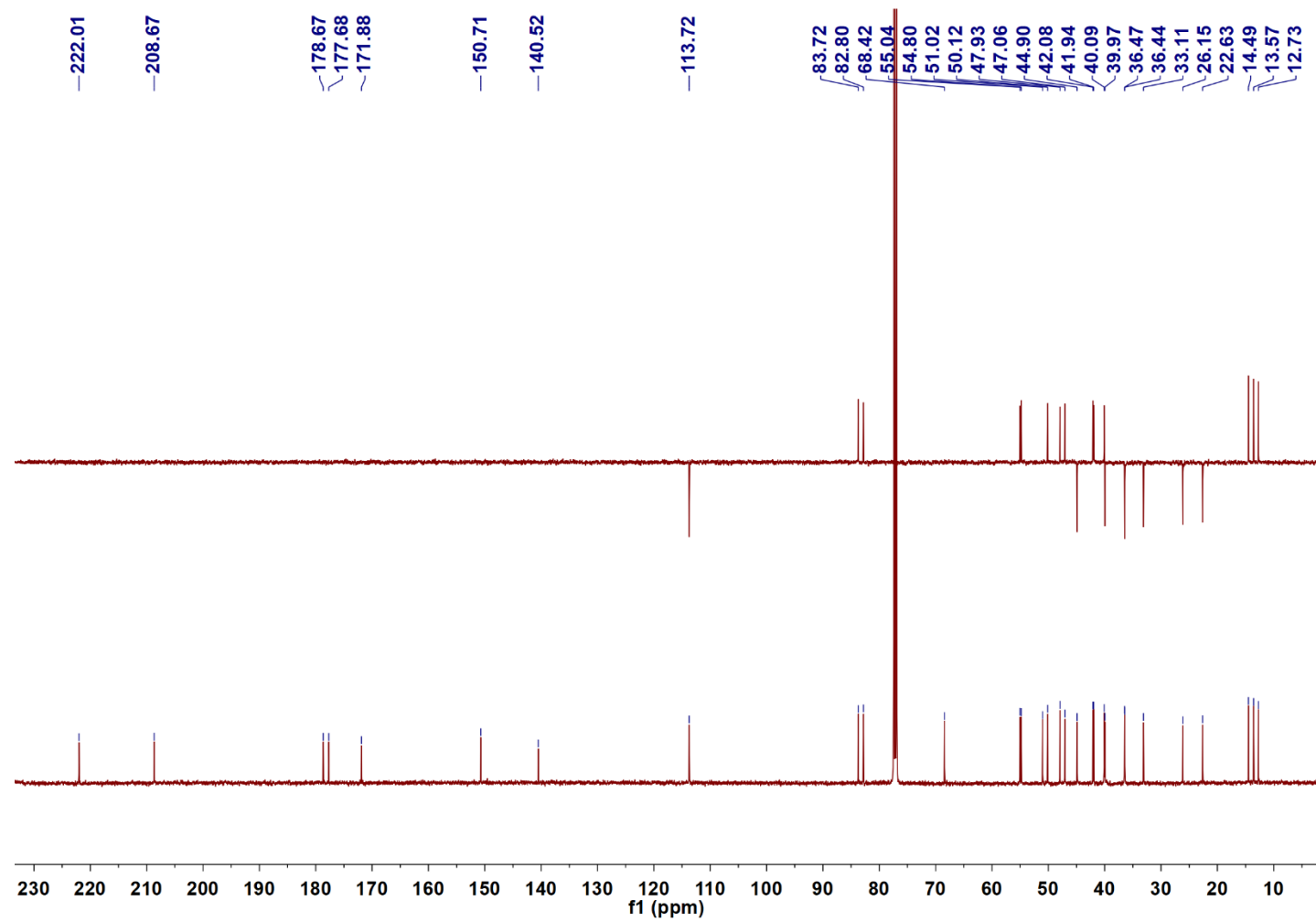


Fig. S30 HRESIMS data of compound 6

Elemental Composition Report

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

263 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-100 H: 0-100 O: 0-50 Na: 0-1

Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
529.2200	529.2202	-0.2	-0.4	13.5	334.8	0.211	80.96	C30 H34 O7 Na
	529.2226	-2.6	-4.9	16.5	336.3	1.674	18.76	C32 H33 O7
	529.2168	3.2	6.0	25.5	340.5	5.850	0.29	C39 H29 O2

AM-B1F3C2E5-pos
20180823-SA-012 1703 (6.875)

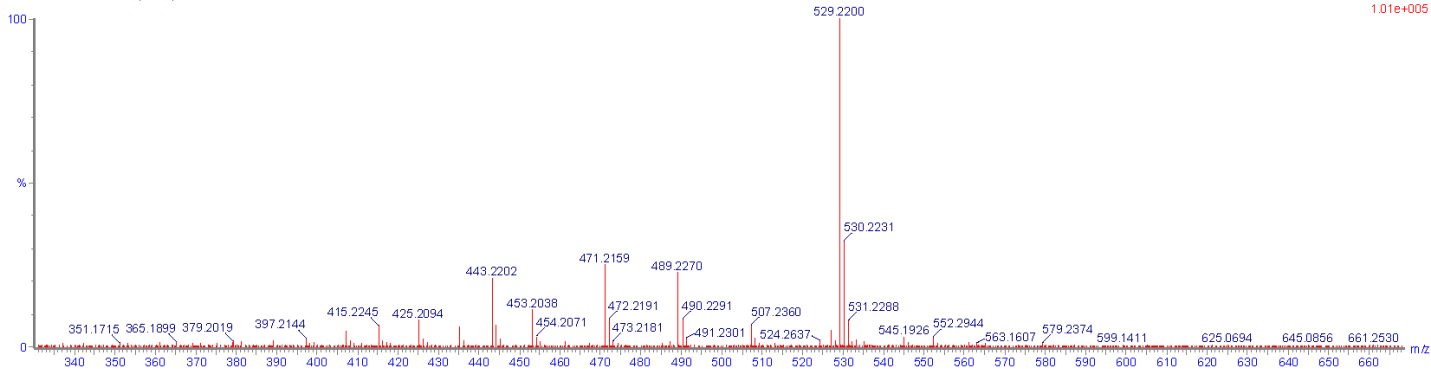


Fig. S31 ^1H NMR spectrum of compound **6** in CDCl_3

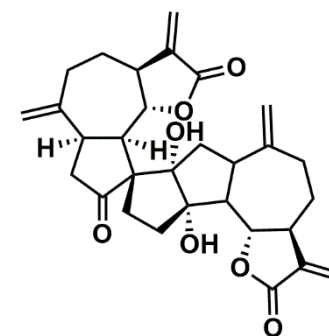
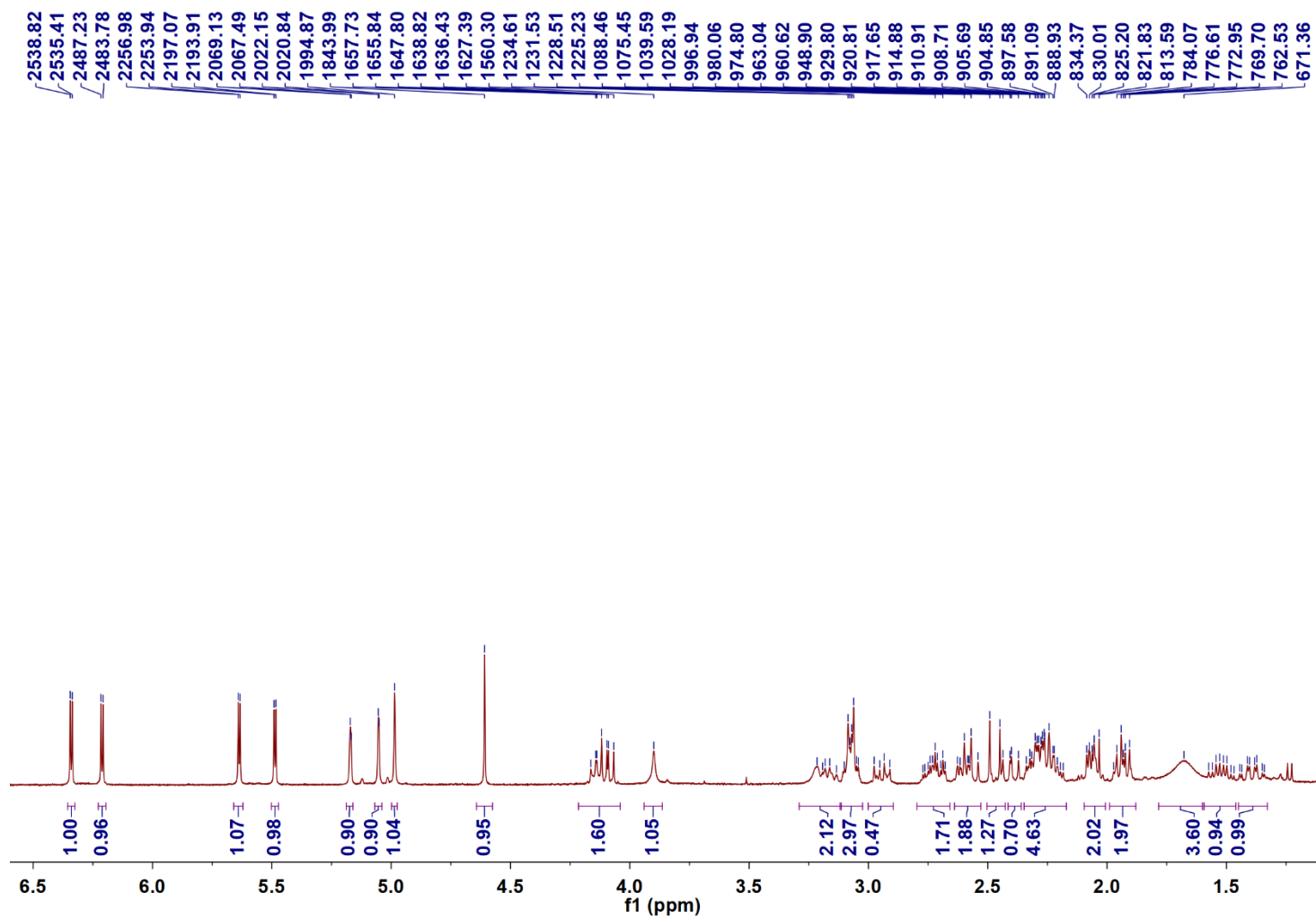


Fig. S32 ^{13}C NMR spectrum of compound **6** in CDCl_3

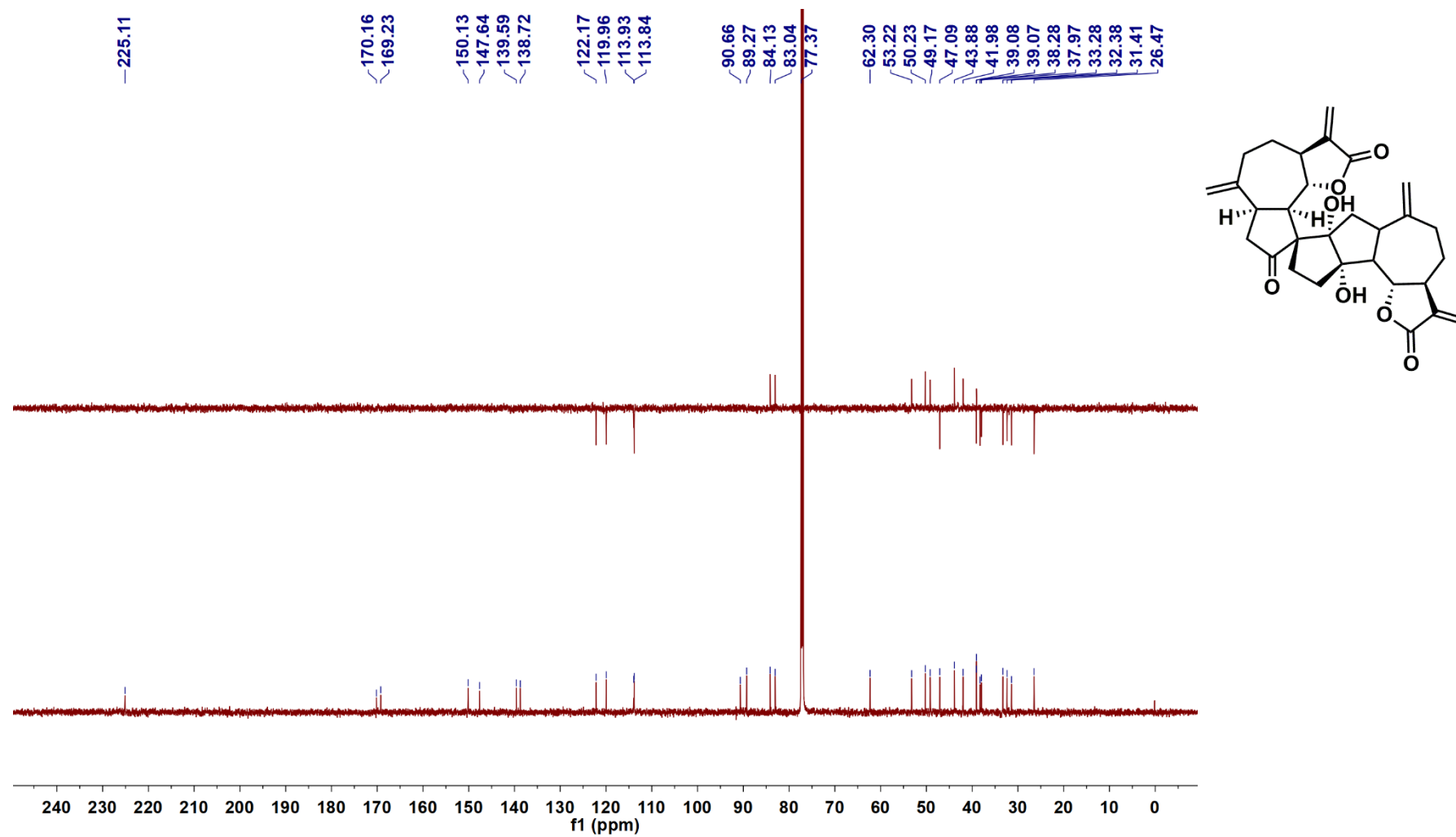


Fig. S33. HRESIMS data of compound **7**

Elemental Composition Report

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

263 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-100 H: 0-100 O: 0-50 Na: 0-1

Minimum: -1.5

Maximum: 5.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
529.2199	529.2202	-0.3	-0.6	13.5	580.3	0.038	96.26	C30 H34 O7 Na
	529.2226	-2.7	-5.1	16.5	583.6	3.330	3.58	C32 H33 O7
	529.2168	3.1	5.9	25.5	586.7	6.421	0.16	C39 H29 O2

AM-BI F3C2C2-pos
20180823-SA011 1657 (6.701)

2: TOF MS ES+
8.90e+005

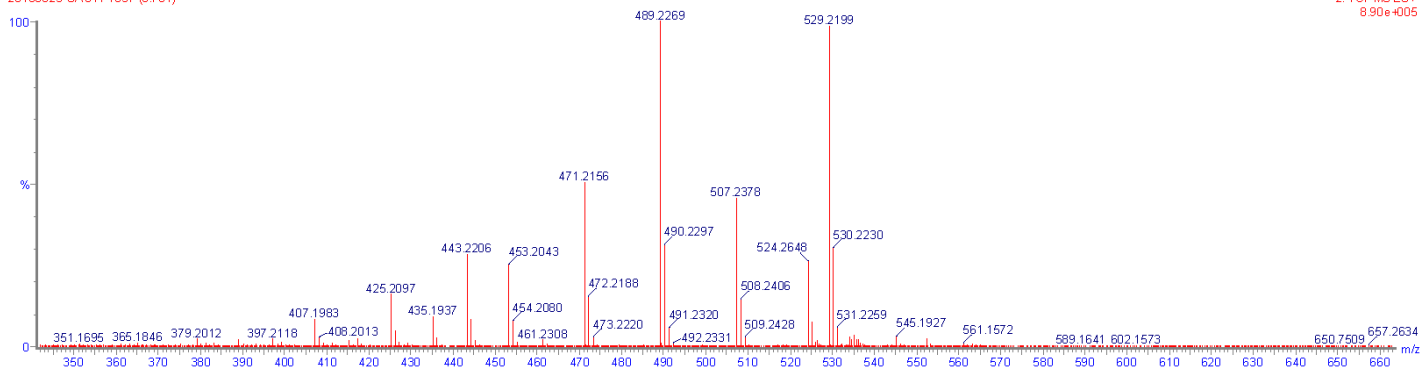


Fig. S34 ^1H NMR spectrum of compound **7** in CDCl_3

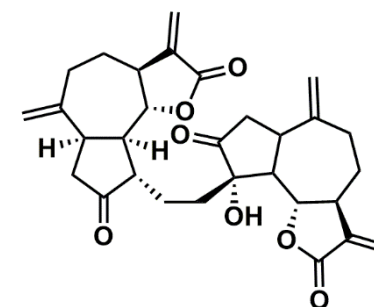
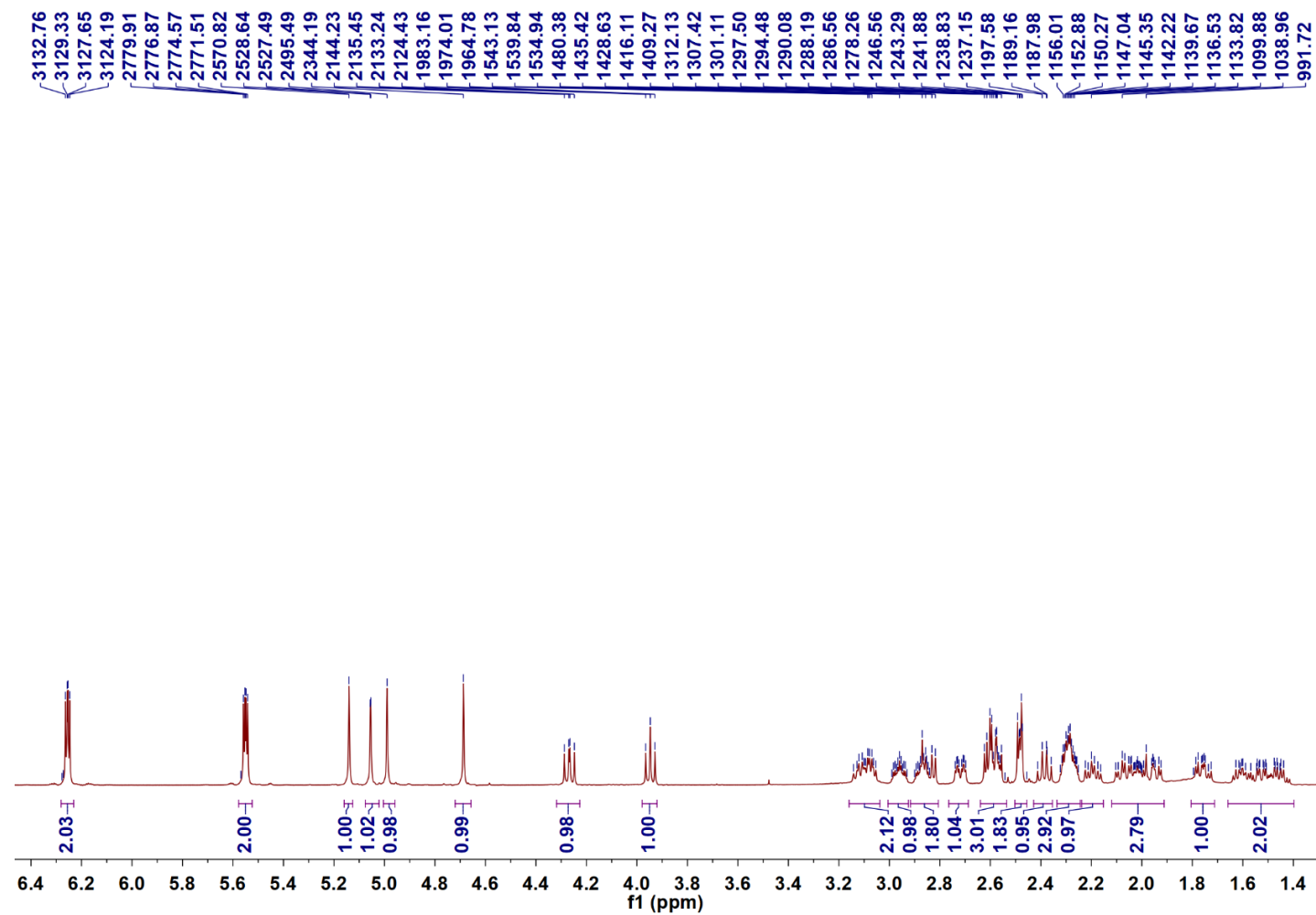


Fig. S35 ^{13}C NMR spectrum of compound **7** in CDCl_3

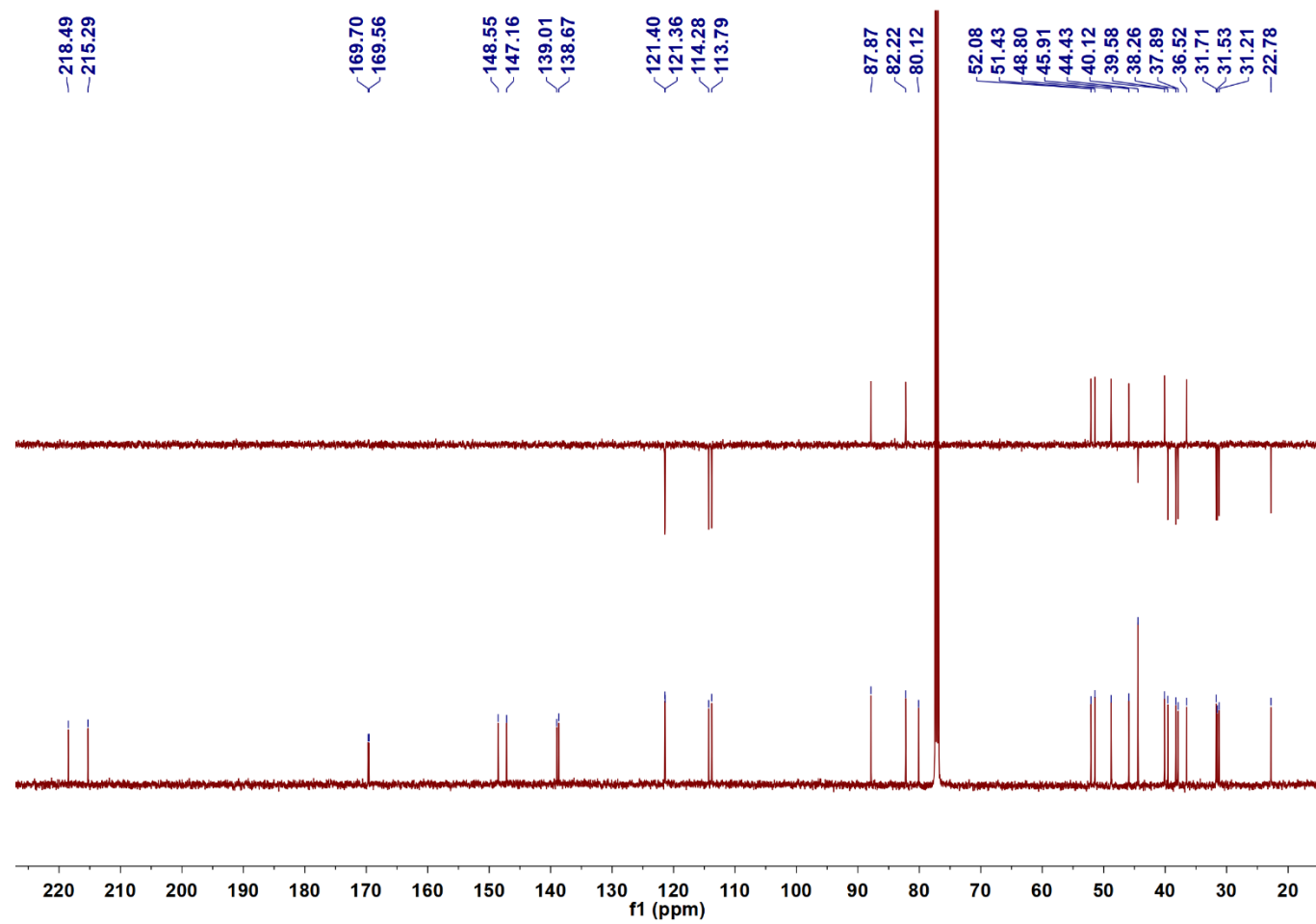
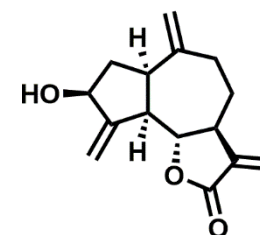
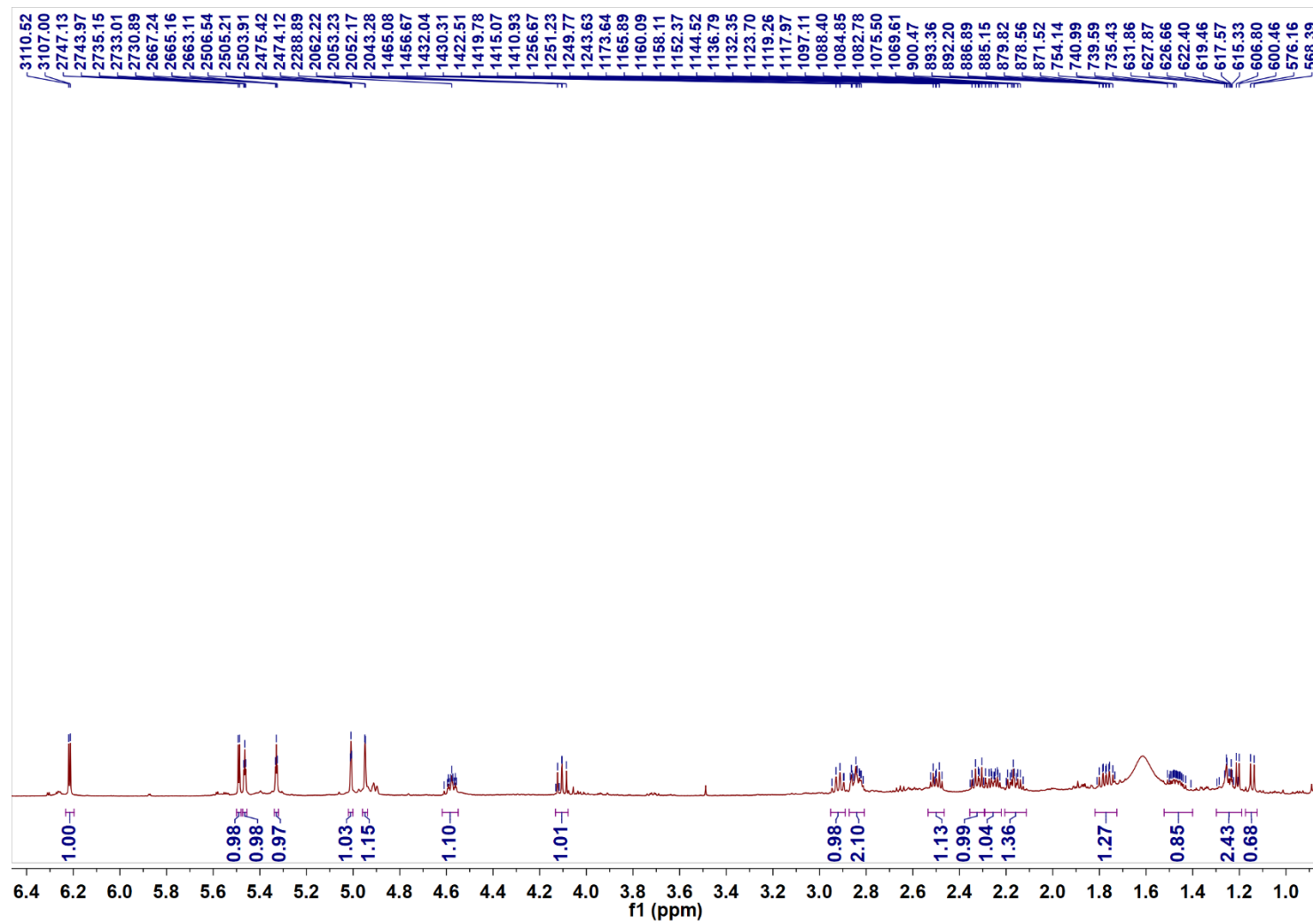
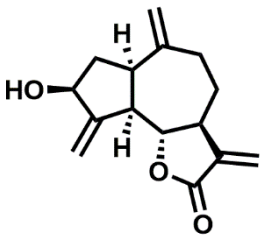


Fig. S36. ^1H NMR spectrum of compound **8** in CDCl_3

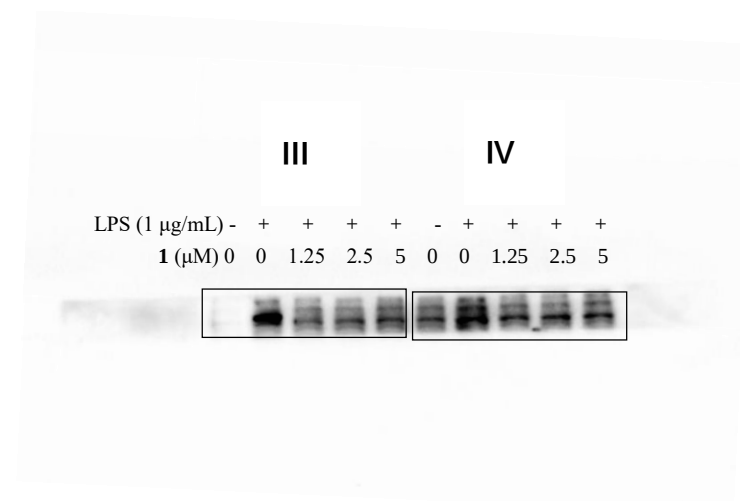
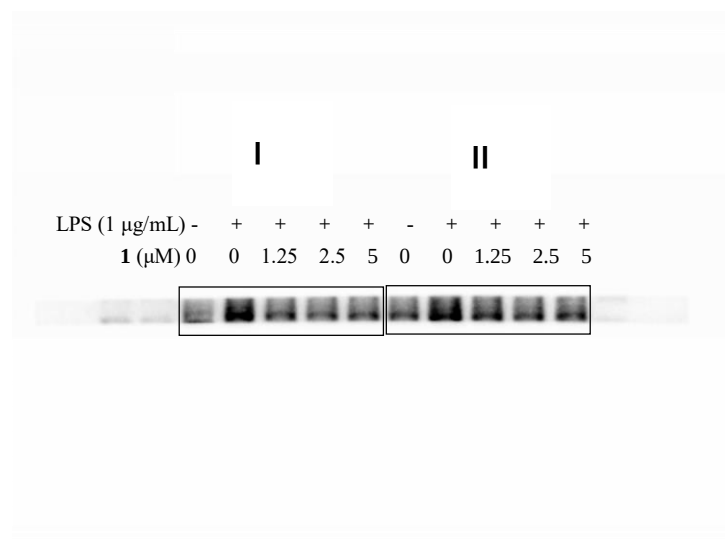


Chemical Shift (ppm)
170.17
153.26
148.13
139.85
120.36
114.60
111.51
84.06
73.79
50.16
45.78
44.42
39.22
34.39
30.75

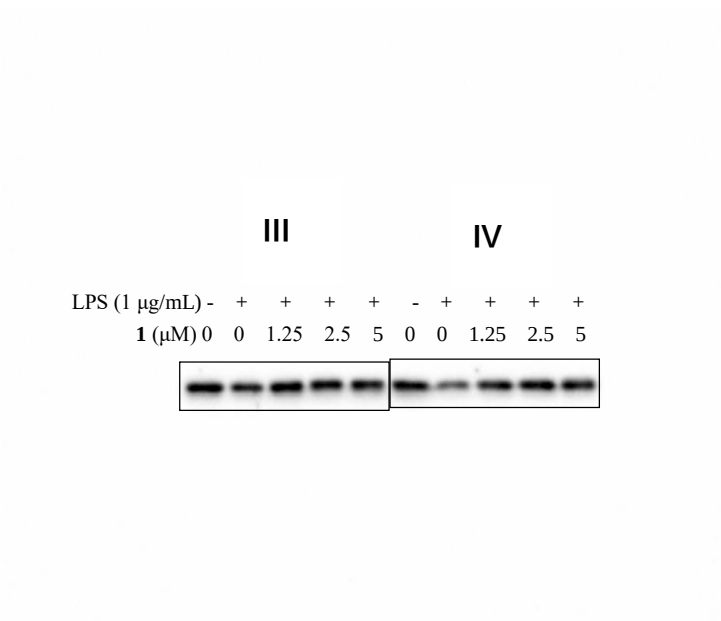
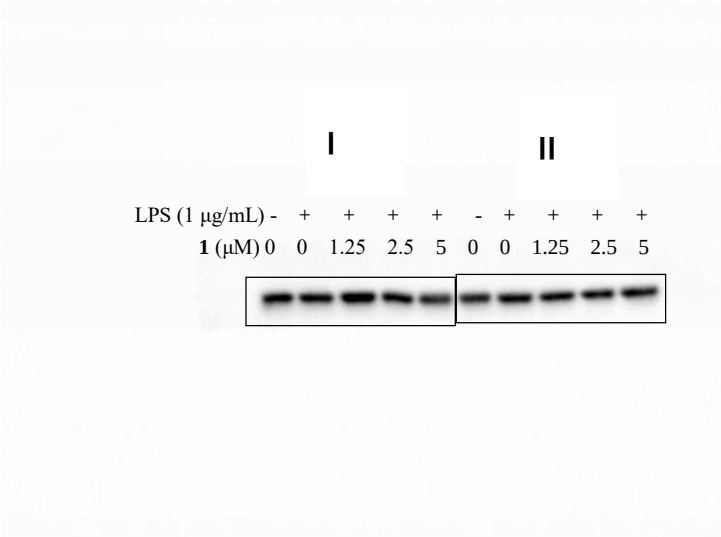


The complete western blot membranes and repeated results for Figure 7B

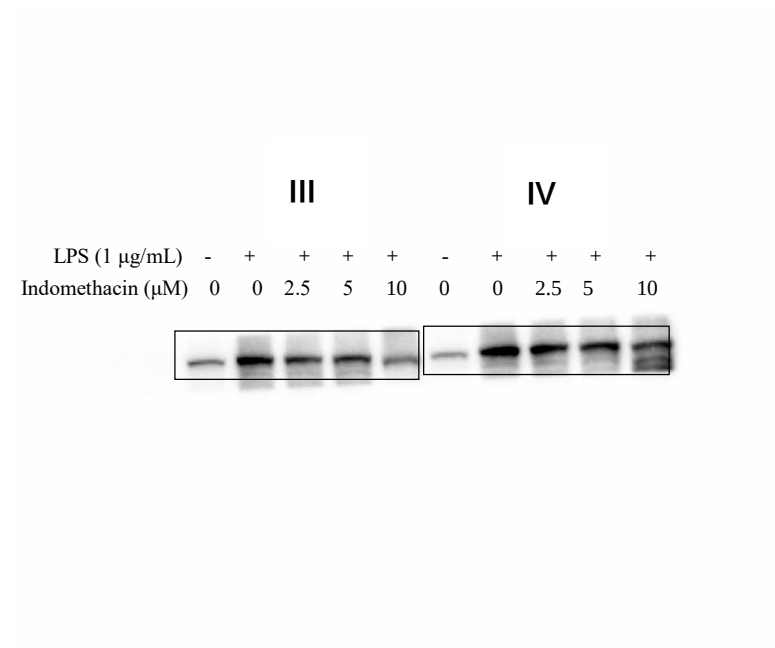
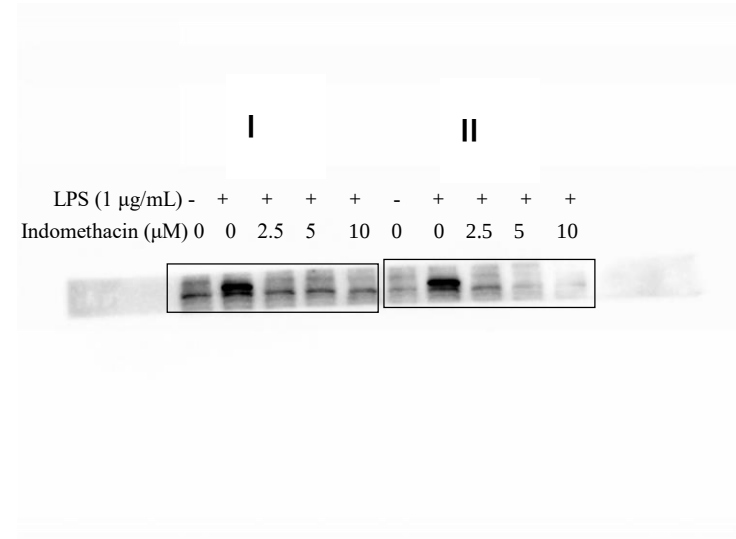
iNOS



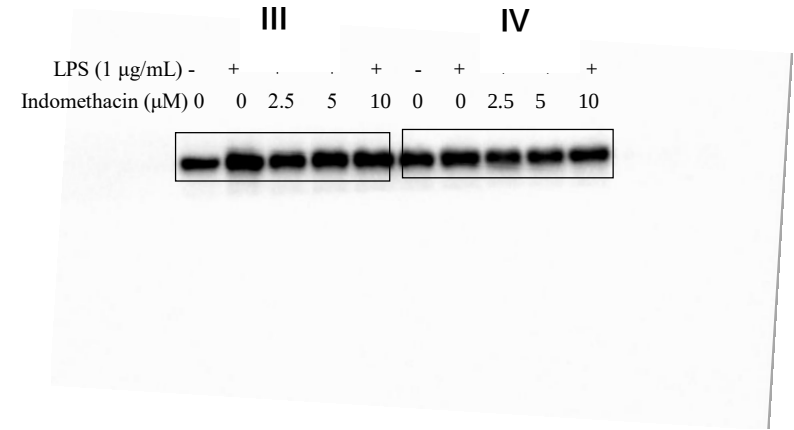
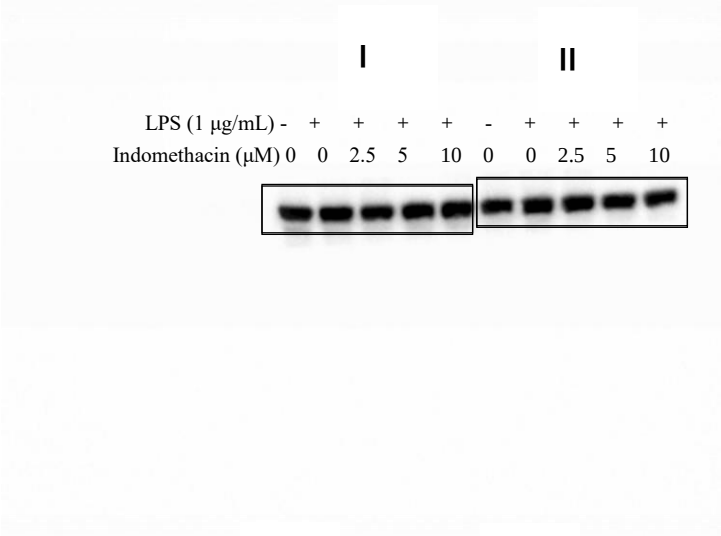
GAPDH



iNOS

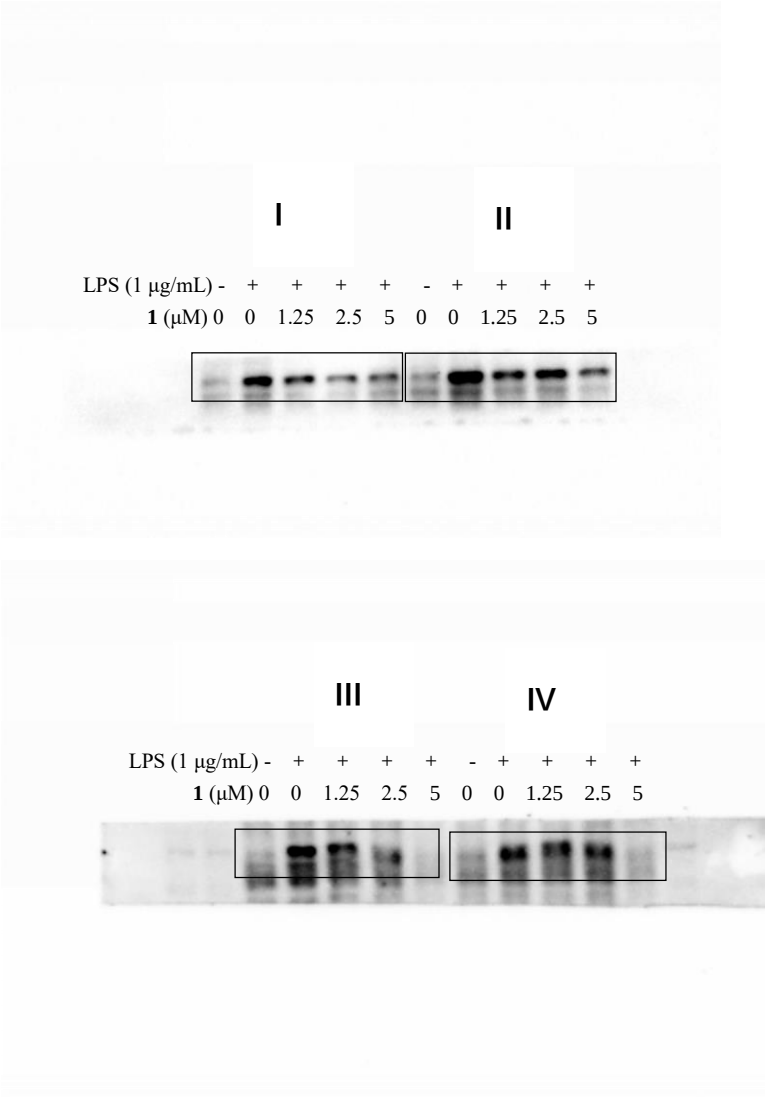


GAPDH

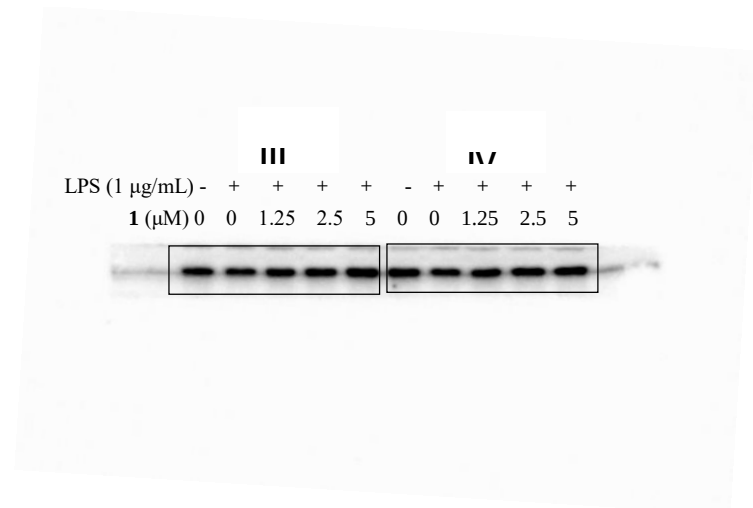
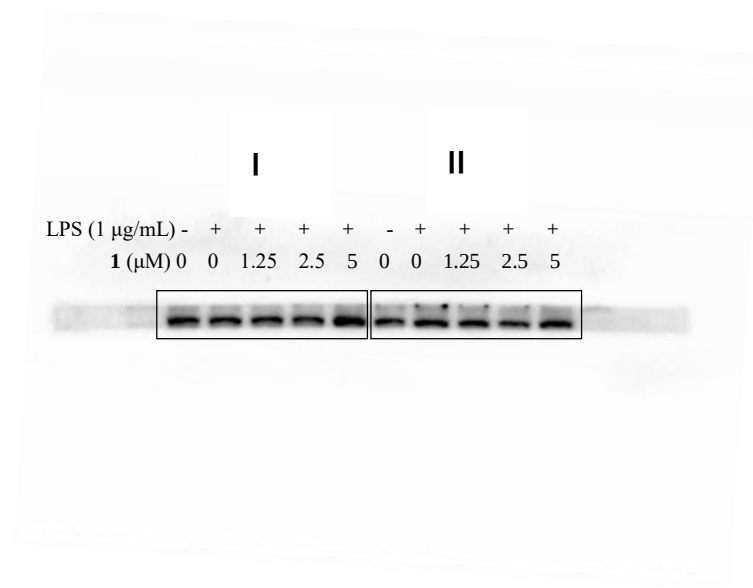


The complete western blot membranes and repeated results for Figure7C

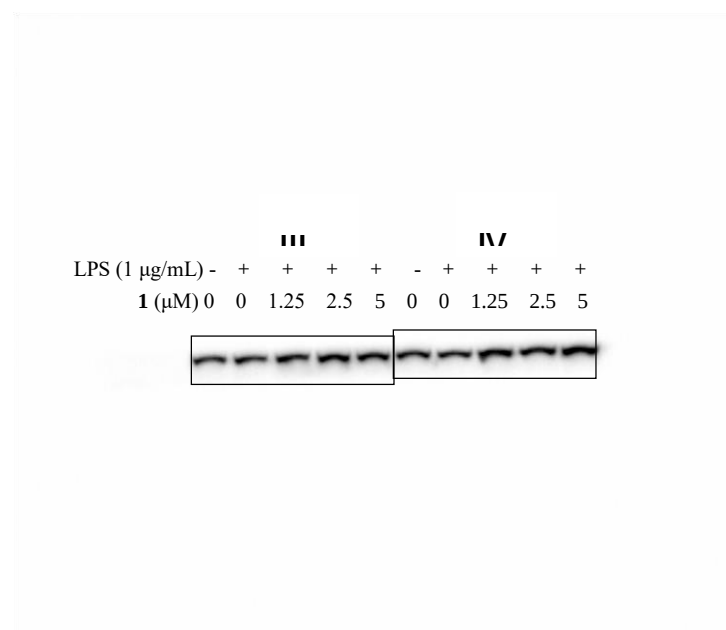
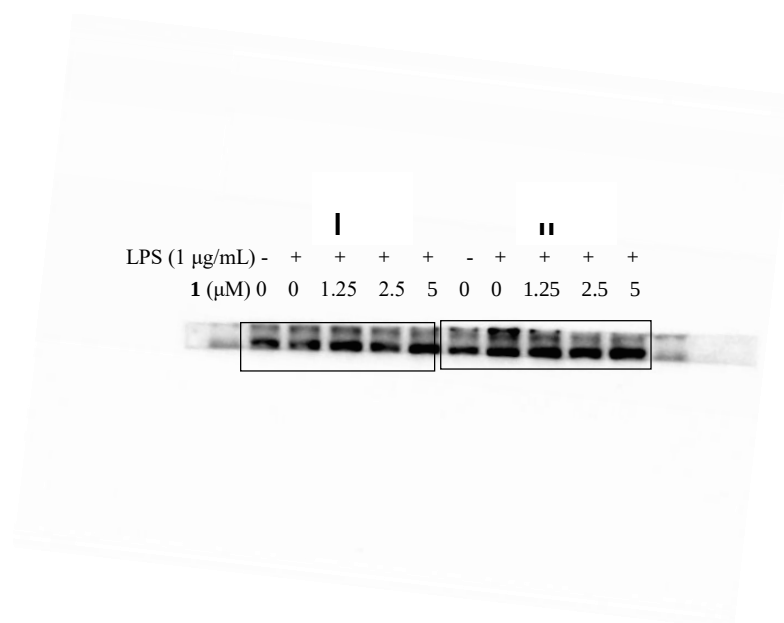
p-IKKα/β



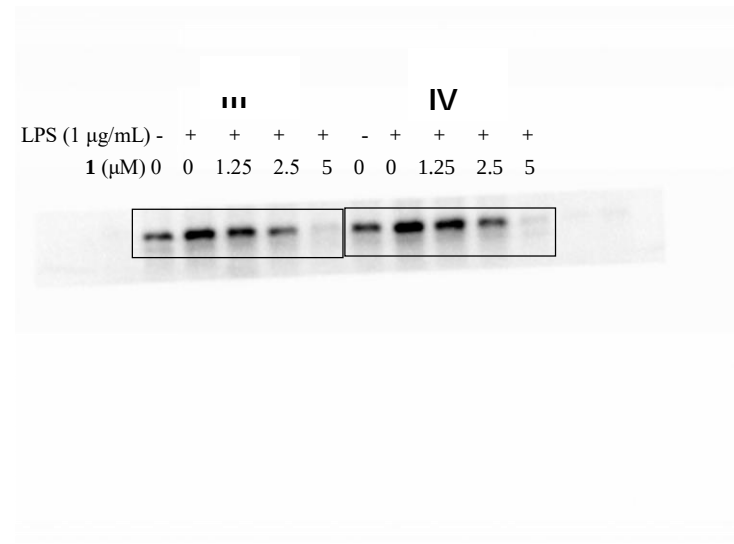
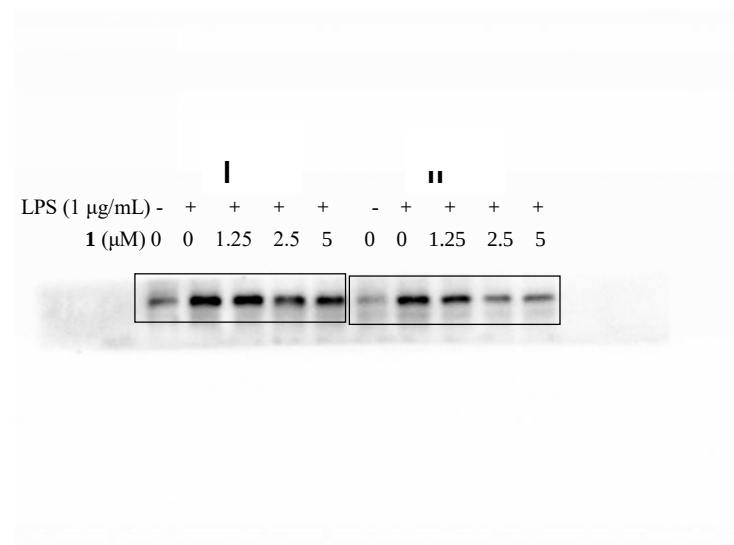
IKKα



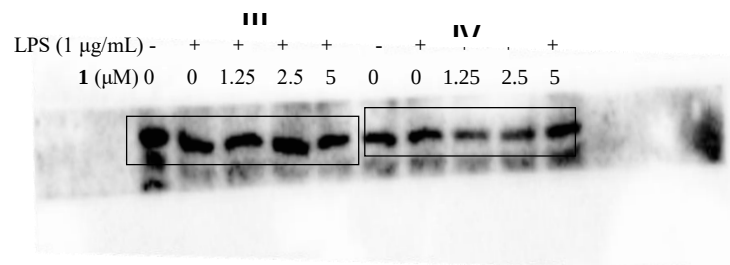
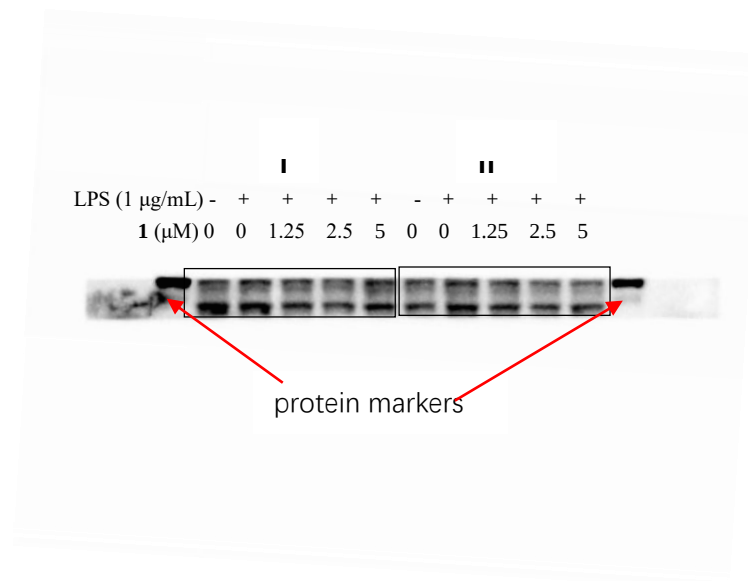
IKK β



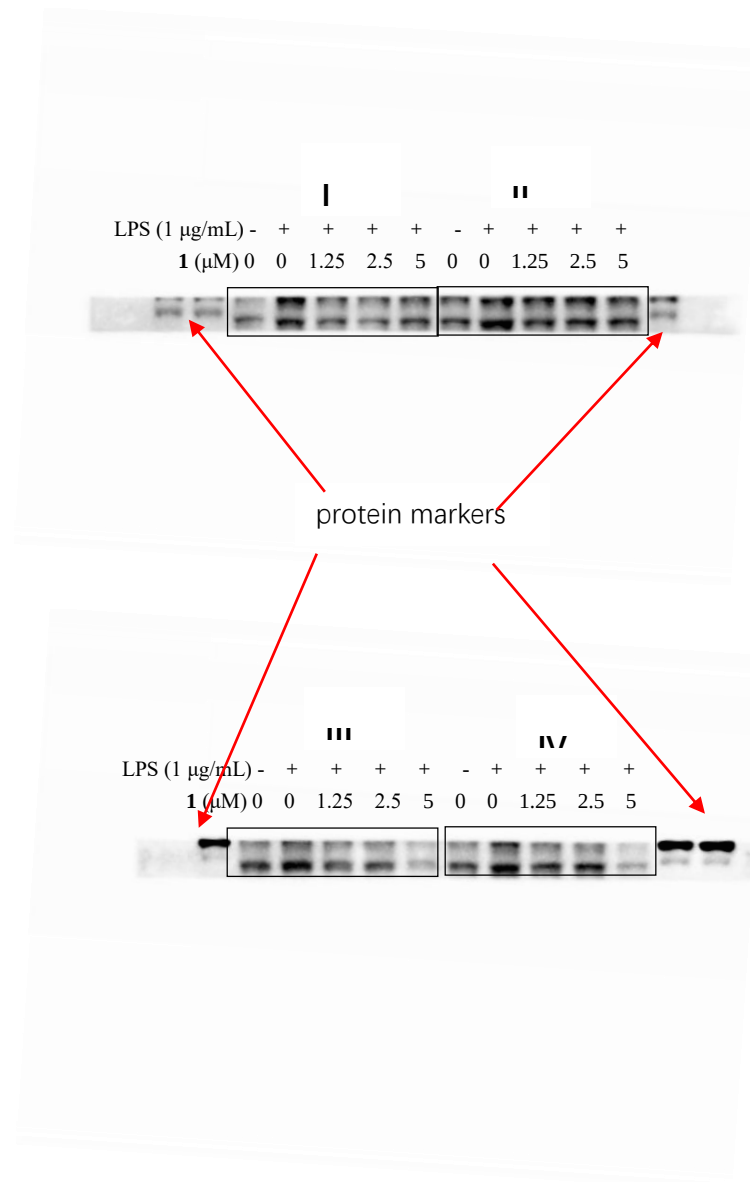
p-IκBα



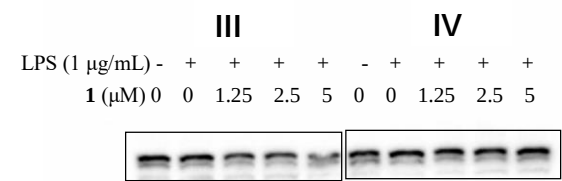
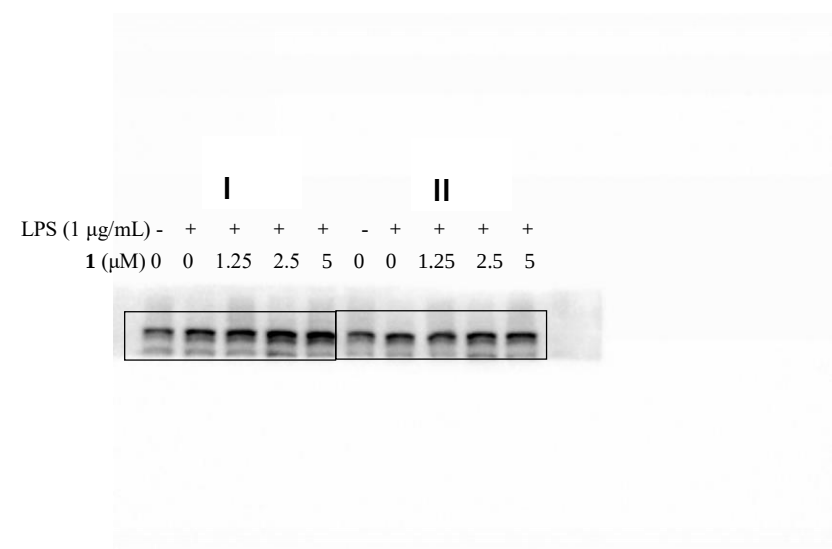
I κ B α



p-p65



p65



GAPDH

