

Supporting Information

Two new lactones from leaves of *Ardisia crenata* Sims with antibacterial and anti-inflammatory activities

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1 Computational methods

1.1 Conformational analysis

Conformational analysis for compound **1** (**Figure S1**) were performed using systematic algorithm by Confab^[1-3] at MMFF94 force field with RMSD threshold of 0.2 Å and energy window of 7 kcal/mol. The detailed experimental process is consistent with our previous work^[2-3]. The energies of all dominative conformers were provided in **Table S1**.

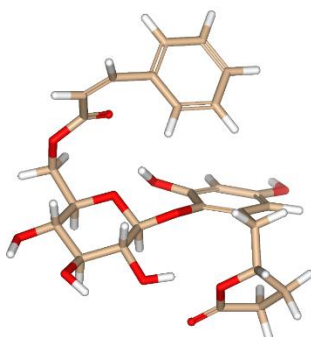
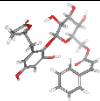

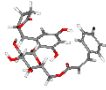
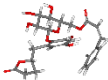
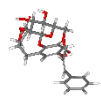
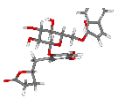
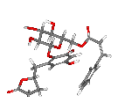
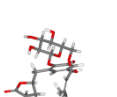
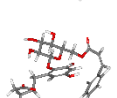
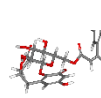


Figure S1 Chemical structure of compounds **1**.

Table S1 Energies of configurations **1**.

Configuration	Conformer	Structure	Energy (kcal/mol)	Population (%)
1	1a		110.4	38
1	1b		110.58	28
1	1c		111.05	12.7
1	1d		111.54	5.5

1	1e		111.94	2.8
1	1f		112.08	2.2
1	1g		112.18	1.9
1	1h		112.3	1.5
1	1i		112.51	1.1
1	1j		112.55	1

1.2 ECD calculation

The theoretical calculations were carried out using Gaussian 09 ^[4]. The detailed experimental process is consistent with our previous work^[2-3]. At first, all conformers were optimized at PM6. Room-temperature equilibrium populations were calculated according to Boltzmann distribution law (**eq. 1**), based on which dominative conformers of population over 5% were kept. The chosen conformers were further optimized at B3LYP/6-31G(d,p) in gas phase (**Tables S2 and S3**). Vibrational frequency analysis confirmed the stable structures. ECD calculations were conducted at B3LYP/6-311G(d,p) level in methanol with IEFPCM model using Time-dependent Density functional theory (TD-DFT). Rotatory strengths for 30 excited states were calculated. The ECD spectrum was simulated using the ECD/UV analysis tool by overlapping Gaussian functions for each transition according to (**eq. 2**).

$$\frac{N_i}{N} = \frac{g_i e^{-\frac{E_i}{k_B T}}}{\sum g_i e^{-\frac{E_i}{k_B T}}} \quad (1)$$

where N_i is the number of conformer i with energy E_i and degeneracy g_i at temperature T , and k_B is Boltzmann constant.

$$\Delta\varepsilon(E) = \frac{1}{2.297 \times 10^{-39}} \times \frac{1}{\sqrt{2\pi}\sigma} \sum_i^A \Delta E_i R_i e^{-\left(\frac{E-E_i}{2\sigma}\right)^2} \quad (2)$$

where σ represents the width of the band at $1/e$ height, while ΔE_i and R_i are the excitation energies and rotatory strengths for transition i , respectively.


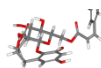
The σ and UV-shift values were set 0.27 eV and -13 nm, respectively. The spectrum of the enantiomers were produced directly by mirror inversion about the horizontal axis.

2 Energies and Coordinates

2.1 Energies at B3LYP theory level

Structures for ECD calculation were shown in **Table S2**.

Table S2 Energies of configurations **1** at B3LYP/6-311G(d,p) in methanol.

Configuration	Conformer	Structure	E (Hartree)	E (kcal/mol)	Population (%)
1	1e		-1835.589597	-1151849.85	78.16
1	1j		-1835.588394	-1151849.1	21.84

2.2 Coordinates at B3LYP theory level

Table S3 Standard orientations of configurations **1** for ECD calculation.

Conformer 1-1e					
Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-2.781397	-0.712892	-1.818198
2	8	0	-2.232896	0.053708	0.310402
3	6	0	-2.288897	-1.077892	-0.478098
4	8	0	-1.941497	-2.159092	-0.055798
5	8	0	0.925804	-0.209593	0.147102
6	6	0	2.026203	-1.020594	0.431302
7	6	0	2.859203	-1.435994	-0.628598
8	6	0	3.824403	-2.415694	-0.381798
9	6	0	3.964403	-2.944894	0.913802
10	6	0	3.169503	-2.528194	1.988002
11	6	0	2.195003	-1.554694	1.733602
12	8	0	1.445503	-1.175793	2.797102
13	8	0	4.937603	-3.890494	1.049802
14	6	0	2.695003	-0.863594	-2.005098
15	6	0	3.700104	0.233406	-2.358198
16	8	0	3.422204	1.368106	-1.449598
17	6	0	4.458104	1.544406	-0.535798
18	6	0	5.610104	0.646605	-0.877198
19	6	0	5.189304	-0.095995	-2.150598
20	8	0	4.277504	2.345506	0.349202
21	6	0	-4.069397	-0.651992	-2.171698
22	6	0	-5.219697	-0.893391	-1.283898
23	6	0	-5.316396	-0.265991	-0.031898
24	6	0	-6.429196	-0.497191	0.780902
25	6	0	-7.445797	-1.356591	0.352802
26	6	0	-7.352497	-1.982491	-0.895298
27	6	0	-6.247797	-1.748691	-1.716998
28	6	0	-1.043396	1.252907	2.029802
29	6	0	-1.645796	2.458307	1.282802
30	6	0	-0.704495	3.043507	0.213102
31	6	0	0.171804	1.960407	-0.454298
32	6	0	0.984604	1.162807	0.585002
33	8	0	0.397604	1.180807	1.892702
34	8	0	1.055304	2.728507	-1.274298
35	8	0	-1.589795	3.591607	-0.774598
36	8	0	-1.837795	3.432508	2.322202
37	6	0	-1.699196	-0.080793	1.663102
38	1	0	-1.973796	-0.517992	-2.536098
39	1	0	4.471303	-2.780494	-1.179298
40	1	0	3.287603	-2.936494	2.988602
41	1	0	0.817204	-0.391893	2.564702
42	1	0	4.996802	-4.255595	1.980202
43	1	0	2.763403	-1.673194	-2.769898
44	1	0	1.661104	-0.455594	-2.118598
45	1	0	3.488804	0.657506	-3.365198
46	1	0	6.544604	1.231005	-1.018598
47	1	0	5.829504	-0.050995	-0.039798
48	1	0	5.788404	0.232605	-3.022598
49	1	0	5.368503	-1.185695	-2.069498

50	1	0	-4.337796	-0.417292	-3.213998
51	1	0	-4.523896	0.406708	0.304302
52	1	0	-6.503296	-0.006191	1.751902
53	1	0	-8.311997	-1.537891	0.990002
54	1	0	-8.145797	-2.652291	-1.229298
55	1	0	-6.181797	-2.234691	-2.691298
56	1	0	-1.114496	1.427807	3.142402
57	1	0	-2.639096	2.212908	0.827202
58	1	0	-0.072595	3.864407	0.627002
59	1	0	-0.435296	1.276607	-1.096198
60	1	0	2.024804	1.517806	0.755202
61	1	0	1.721204	2.138306	-1.728998
62	1	0	-1.058295	4.024107	-1.490898
63	1	0	-2.423795	4.160508	2.000502
64	1	0	-0.972597	-0.919293	1.694702
65	1	0	-2.567796	-0.309592	2.310402

Conformer 1-lj

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-4.134402	-1.791591	-1.865899
2	8	0	-2.331196	-0.317299	-1.849599
3	6	0	-3.708996	-0.384993	-1.971799
4	8	0	-4.366892	0.613710	-2.158999
5	8	0	1.017607	0.367987	-1.002399
6	6	0	1.586301	-0.881916	-0.746599
7	6	0	2.956101	-1.069021	-1.024899
8	6	0	3.497395	-2.350124	-0.896199
9	6	0	2.677291	-3.410220	-0.468899
10	6	0	1.317791	-3.245914	-0.177599
11	6	0	0.773697	-1.963812	-0.328399
12	8	0	-0.544503	-1.839507	-0.039399
13	8	0	3.302986	-4.617823	-0.359099
14	6	0	3.796605	0.083275	-1.488899
15	6	0	4.796208	0.608071	-0.459699
16	8	0	4.021409	1.011074	0.735401
17	6	0	4.374406	0.251973	1.847401
18	6	0	5.482302	-0.698532	1.502101
19	6	0	5.850803	-0.393234	0.047001
20	8	0	3.775007	0.463375	2.873201
21	6	0	-4.539605	-2.370490	-0.731099
22	6	0	-4.582102	-1.716689	0.589001
23	6	0	-5.760702	-1.786784	1.350801
24	6	0	-5.815900	-1.175784	2.605601
25	6	0	-4.696897	-0.505989	3.112801
26	6	0	-3.519497	-0.444694	2.360701
27	6	0	-3.460799	-1.045194	1.100701
28	6	0	-1.684188	1.624298	-0.558899
29	6	0	-1.419582	3.142997	-0.616999
30	6	0	-0.136880	3.566192	0.125601
31	6	0	1.030016	2.590987	-0.134999
32	6	0	0.635310	1.130288	0.164601
33	8	0	-0.770591	0.927294	0.318401
34	8	0	2.019118	2.994183	0.816201
35	8	0	0.188226	4.842390	-0.442999
36	8	0	-2.559779	3.692302	0.063801
37	6	0	-1.691191	0.983498	-1.950099

38	1	0	-4.111505	-2.329691	-2.822499
39	1	0	4.547594	-2.540728	-1.118799
40	1	0	0.693288	-4.071312	0.154401
41	1	0	-0.900399	-0.884905	-0.216799
42	1	0	2.689083	-5.341820	-0.041199
43	1	0	4.350104	-0.200627	-2.415799
44	1	0	3.132209	0.923778	-1.806599
45	1	0	5.248512	1.566569	-0.800899
46	1	0	6.345303	-0.575636	2.190501
47	1	0	5.152098	-1.752031	1.637201
48	1	0	6.868605	0.036962	-0.028799
49	1	0	5.877100	-1.315234	-0.568199
50	1	0	-4.887109	-3.415088	-0.739599
51	1	0	-6.634305	-2.311481	0.961601
52	1	0	-6.734200	-1.223980	3.192701
53	1	0	-4.742595	-0.034589	4.095101
54	1	0	-2.644495	0.071402	2.757901
55	1	0	-2.534799	-1.000198	0.518301
56	1	0	-2.655489	1.434602	-0.020399
57	1	0	-1.402480	3.543197	-1.660699
58	1	0	-0.313679	3.689393	1.220301
59	1	0	1.433717	2.698285	-1.169999
60	1	0	1.057708	0.720787	1.107301
61	1	0	2.883616	2.516179	0.643201
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3 Figures

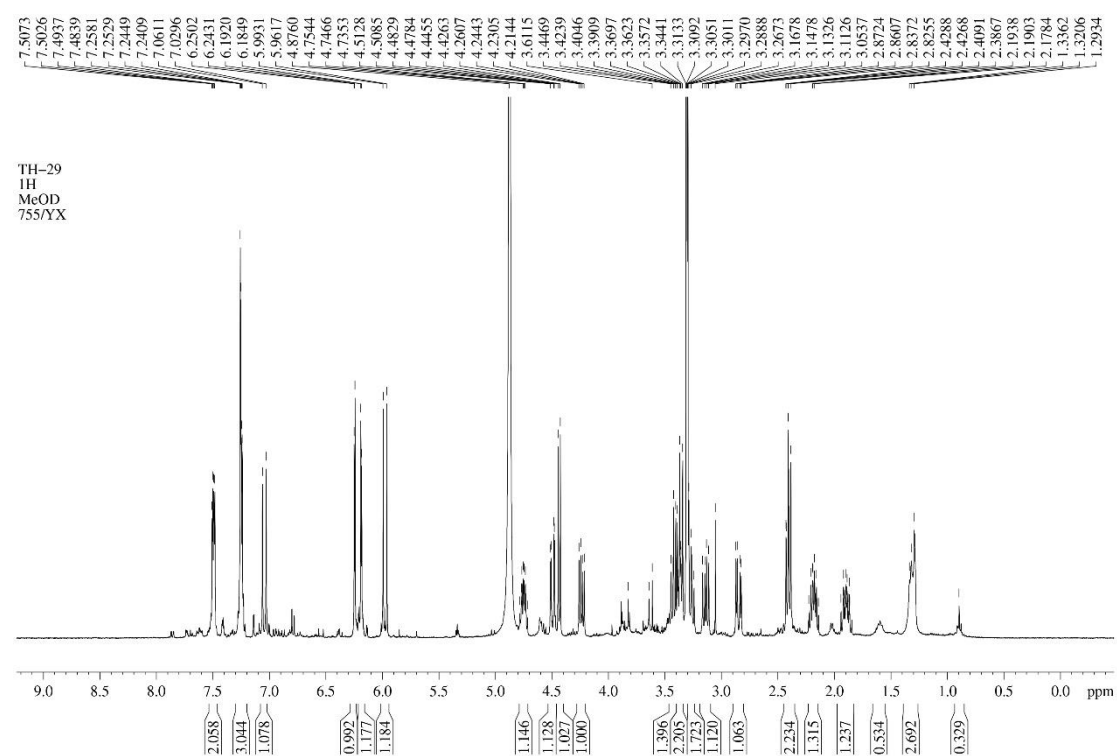


Figure S2. ^1H NMR spectrum of **1** in CD_3OD (400 MHz)

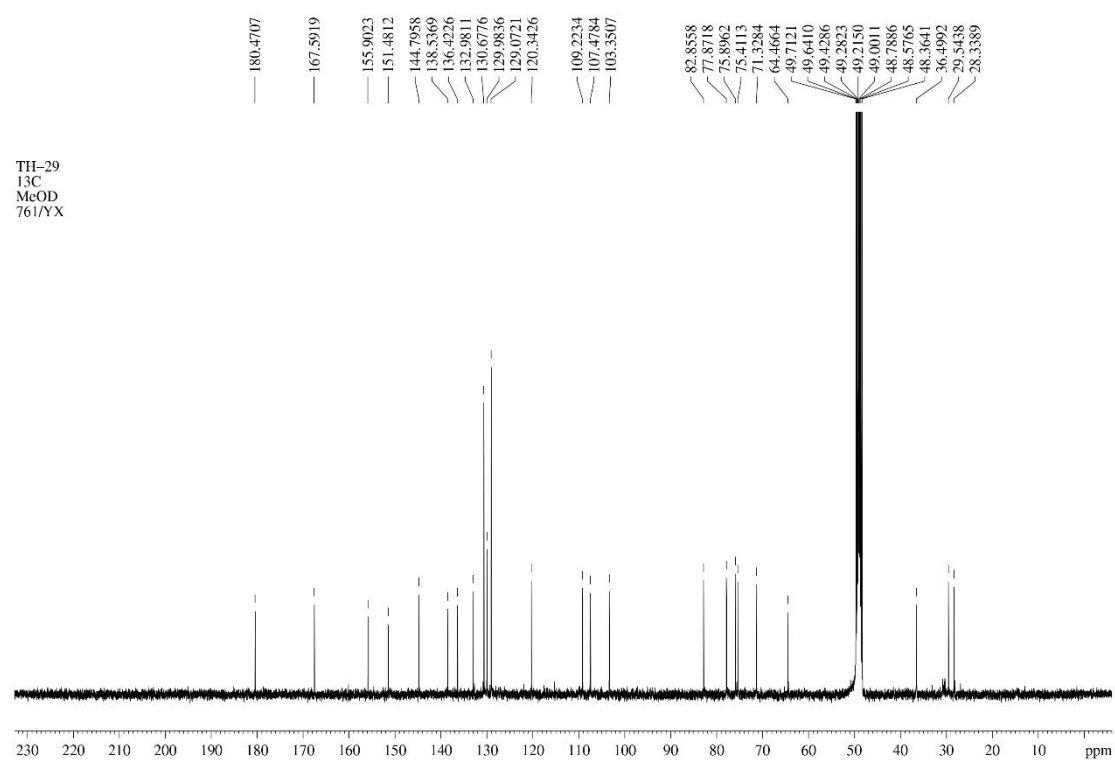


Figure S3. ¹³C NMR spectrum of 1 in CD₃OD (100 MHz)

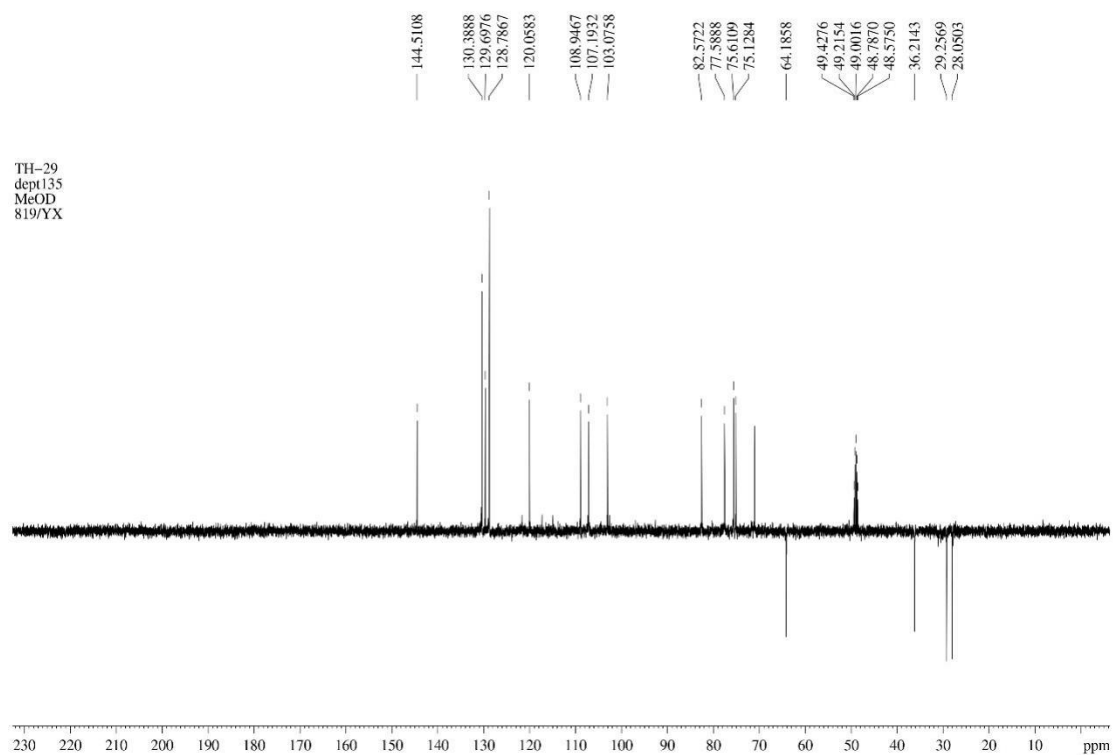


Figure S4. dept135° spectrum of 1 in CD₃OD (100 MHz)

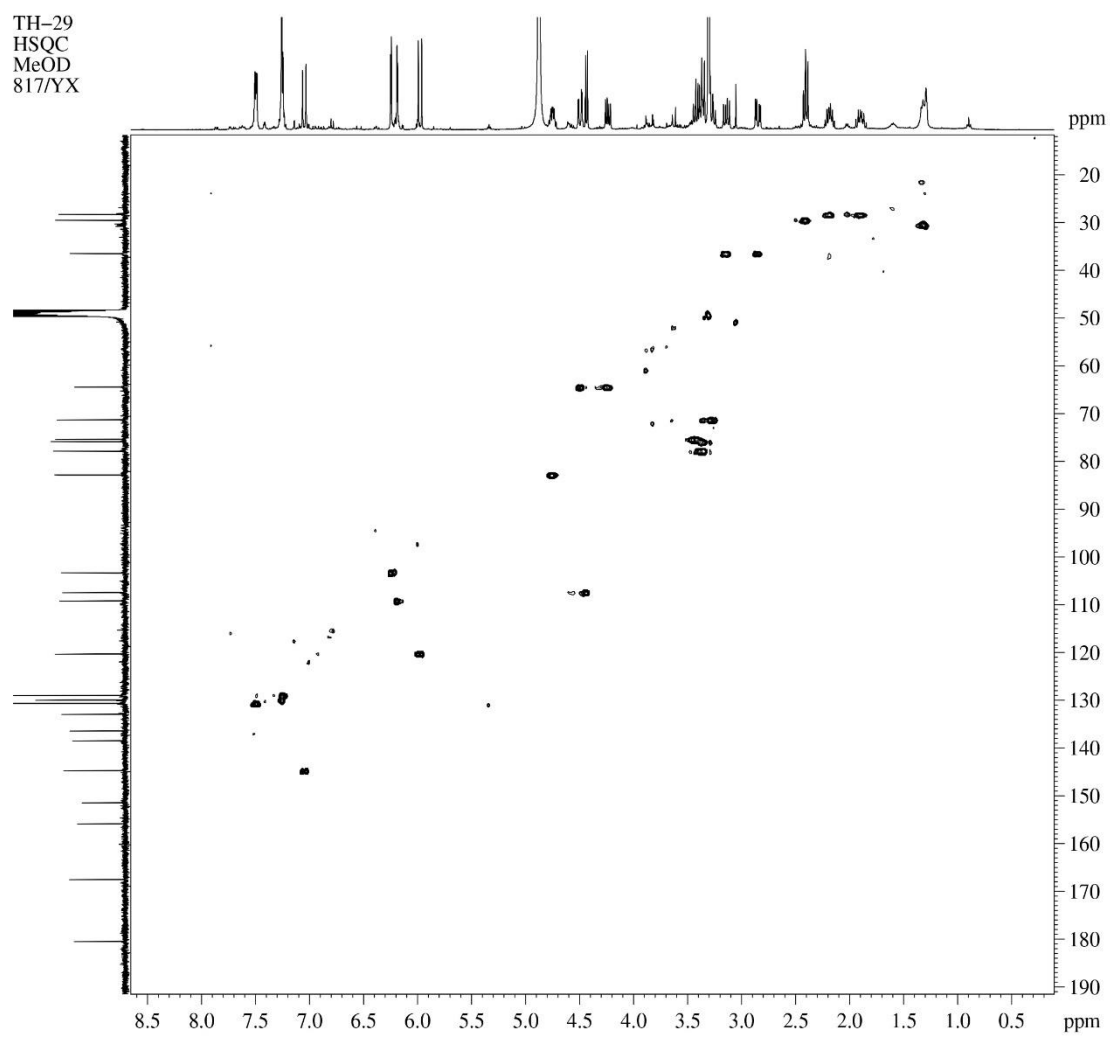


Figure S5. HSQC spectrum of **1** in CD₃OD (100 MHz)

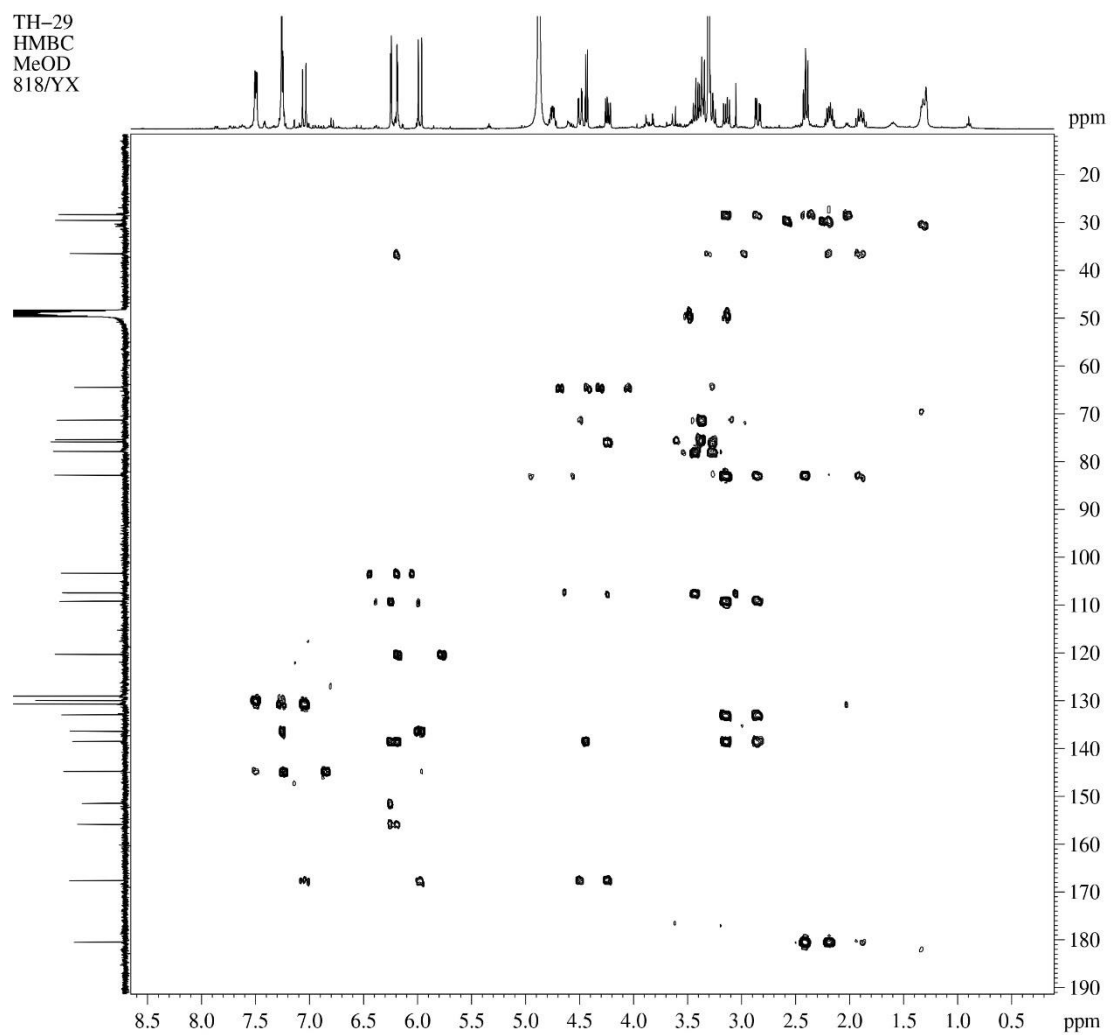


Figure S6. HMBC spectrum of **1** in CD₃OD (100 MHz)

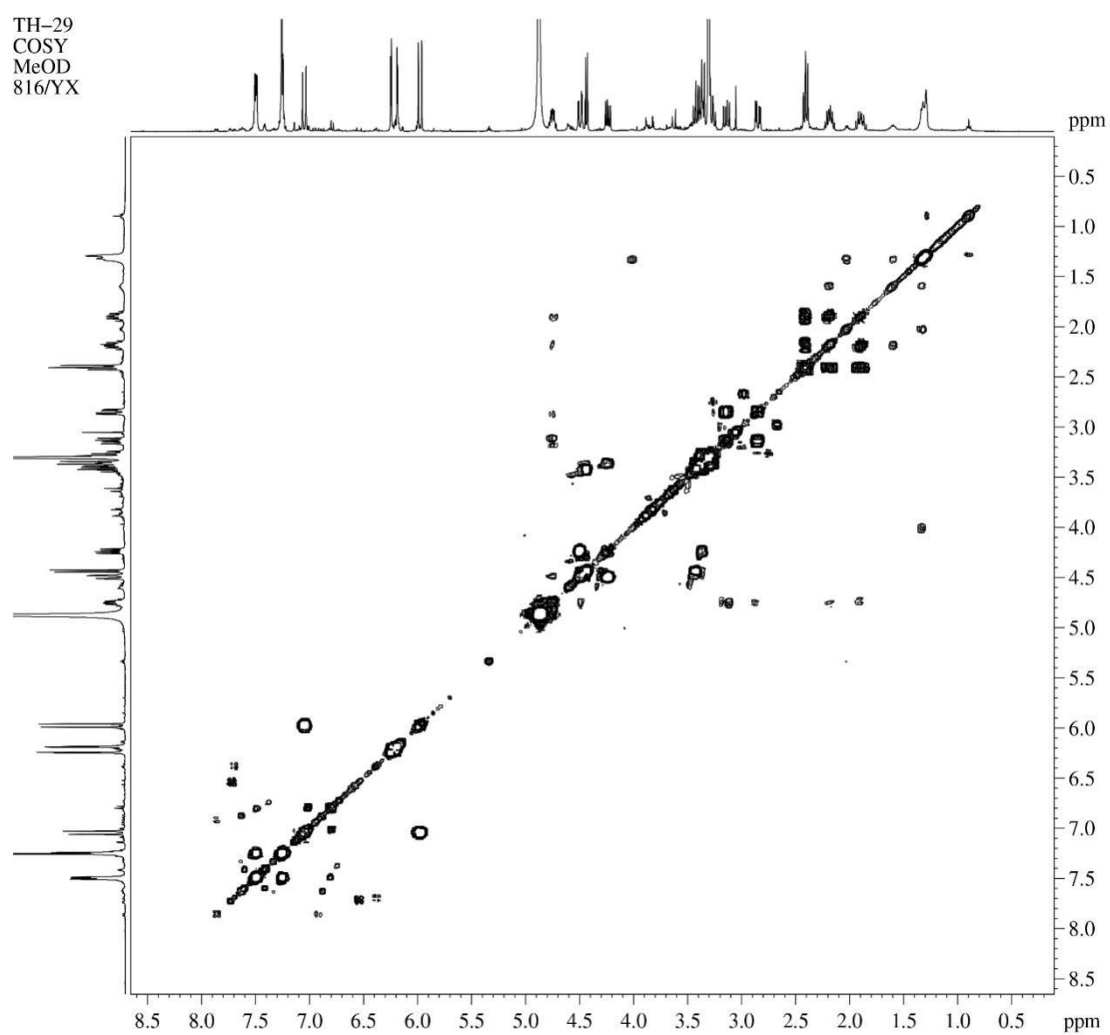


Figure S7. ^1H - ^1H COSY spectrum of **1** in CD_3OD (400 MHz).

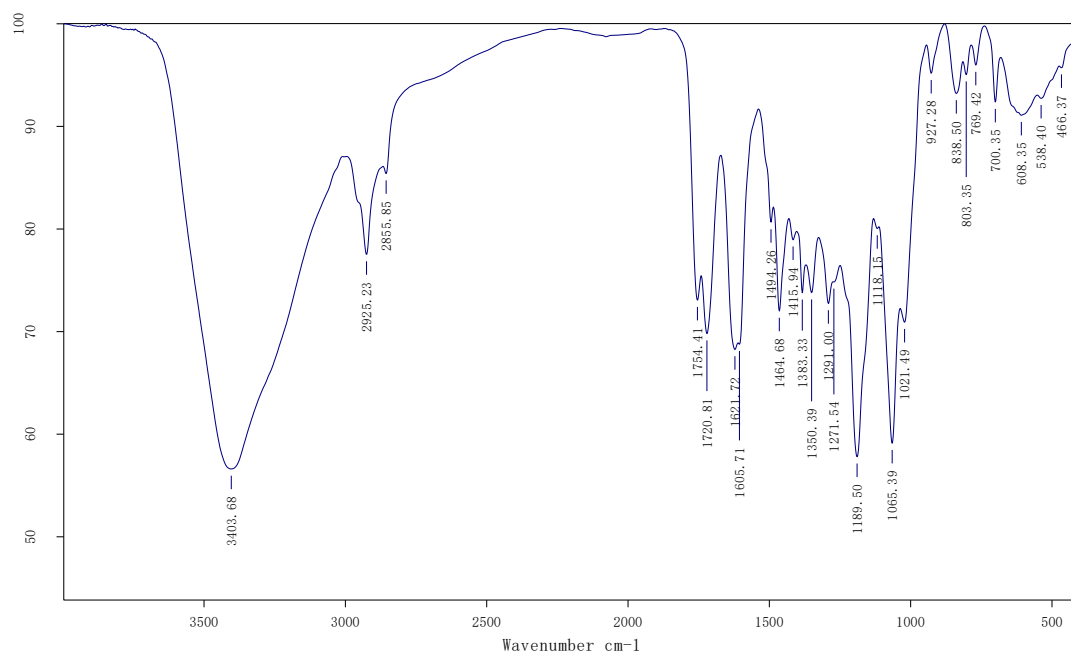


Figure S8. IR spectrum of 1.

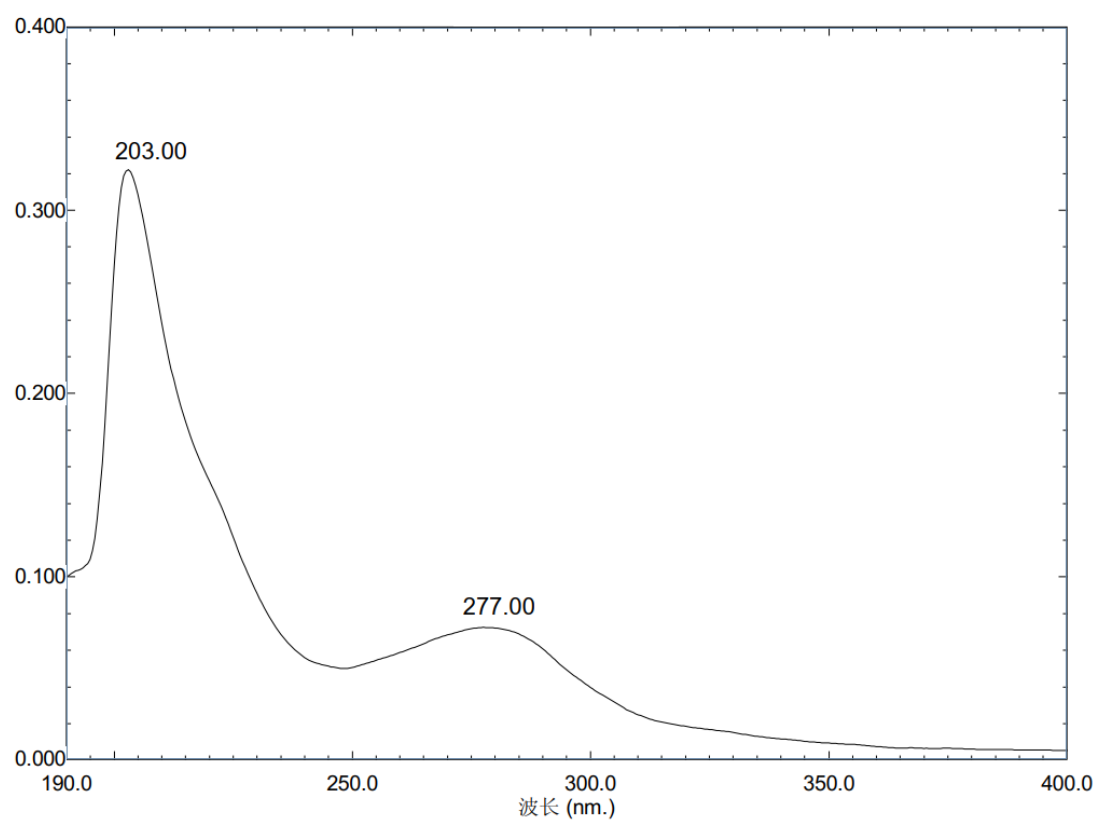


Figure S9. UV spectrum of 1.

Rudolph Research Analytical

This sample was measured on an Autopol VI, Serial #91058
Manufactured by Rudolph Research Analytical, Hackettstown, NJ, USA.

Measurement Date : Wednesday, 15-DEC-2021

Set Temperature : OFF

Time Delay : Disabled

Delay between Measurement : Disabled

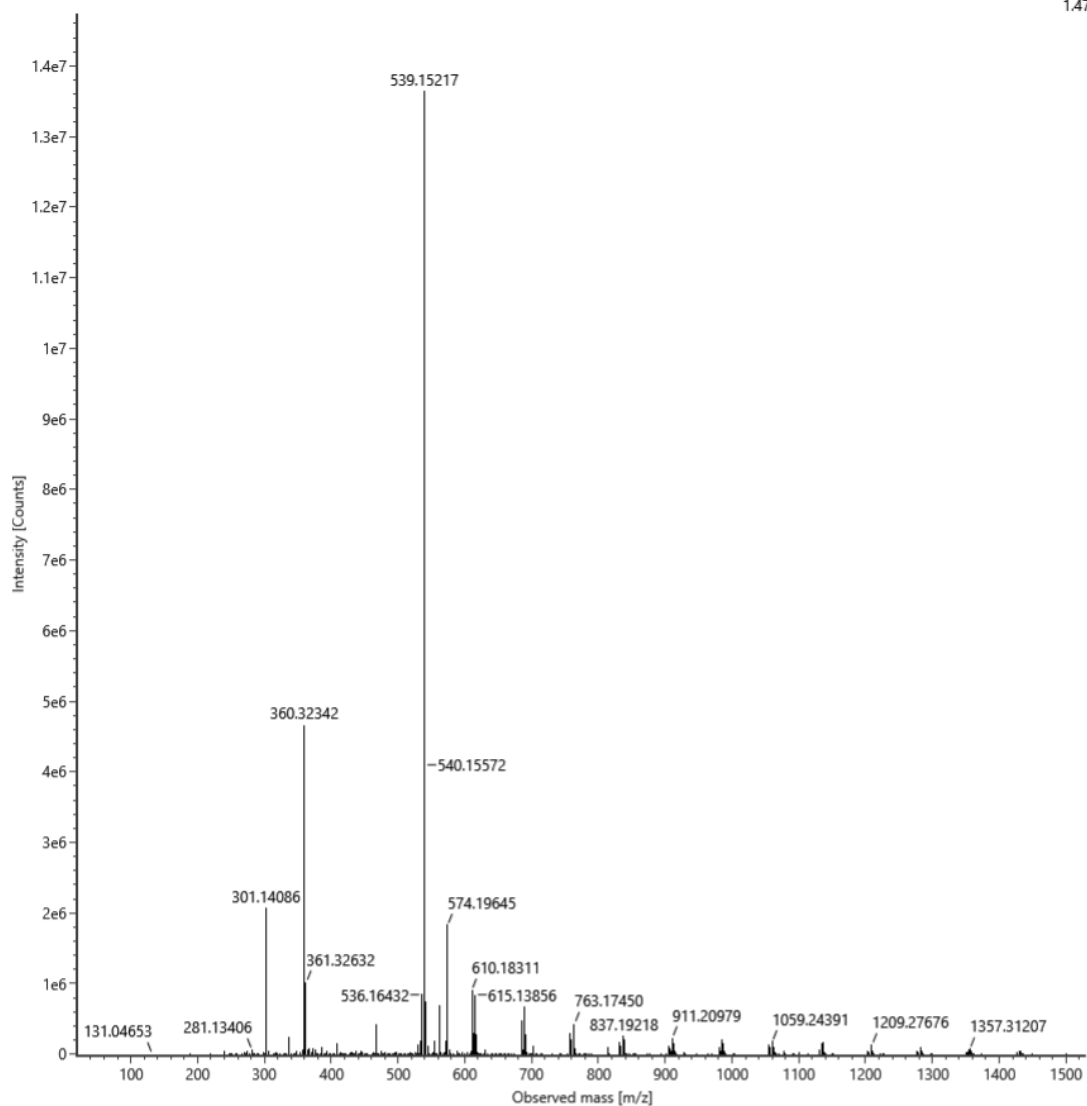
<u>n</u>	<u>Average</u>	<u>Std.Dev.</u>	<u>% RSD</u>	<u>Maximum</u>	<u>Minimum</u>					
5	14.23	0.45	3.16	14.62	13.65					
<u>S.No</u>	<u>Sample ID</u>	<u>Time</u>	<u>Result</u>	<u>Scale</u>	<u>OR °Arc</u>	<u>WLG.nm</u>	<u>Lg.mm</u>	<u>Conc.g/100ml</u>	<u>Temp.</u>	
1	TH-29	03:21:27 PM	13.65	SR	0.0071	589	100.00	0.052	19.6	
2	TH-29	03:21:35 PM	14.62	SR	0.0076	589	100.00	0.052	19.6	
3	TH-29	03:21:43 PM	14.42	SR	0.0075	589	100.00	0.052	19.6	
4	TH-29	03:21:51 PM	14.62	SR	0.0076	589	100.00	0.052	19.6	
5	TH-29	03:22:00 PM	13.85	SR	0.0072	589	100.00	0.052	19.6	

Figure S10. ORD spectrum of 1.

Item name: TH-29
Item description:

Channel name: 1: Average Time 0.1089 min : TOF MS (50-1500) ESI+ : Centroided : Combined

1.47e7



Add:Na⁺

Composition	i-FIT Confidence (%)	Predicted m/z	m/z error (PPM)
C ₂₆ H ₂₈ O ₁₁	100.000000	539.152383	-0.395467

Figure S11. HRESIMS spectrum of 1.

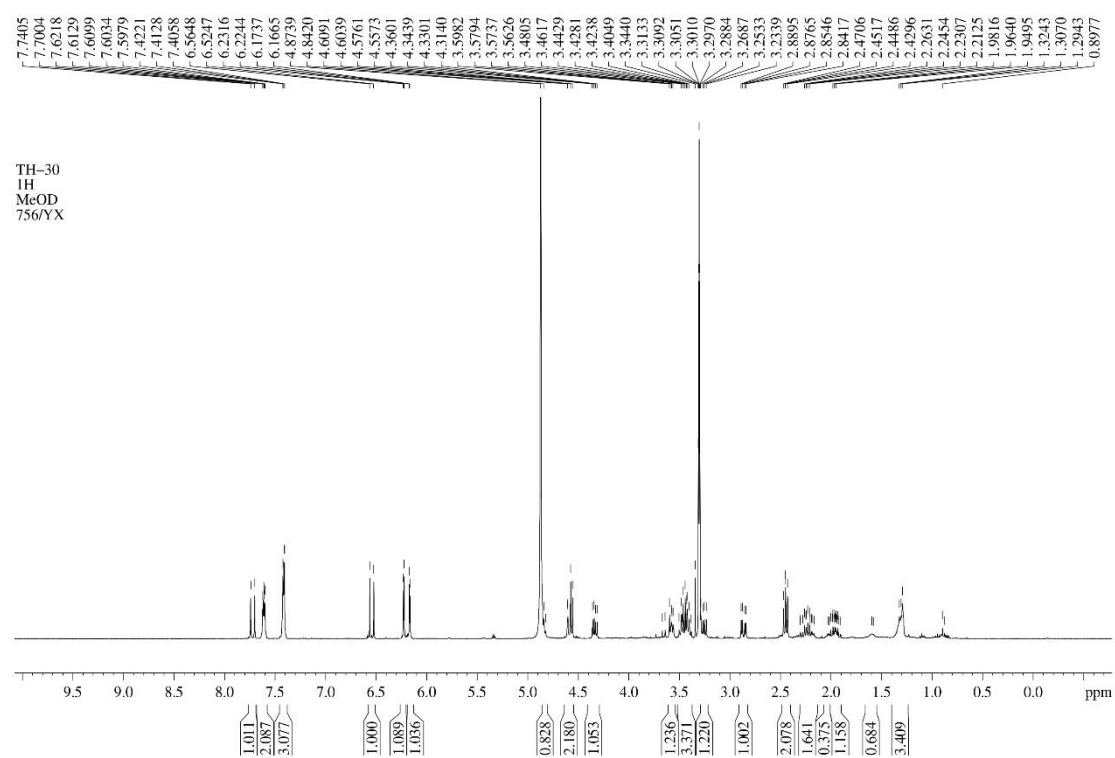


Figure S12. ^1H NMR spectrum of **2** in CD_3OD (400 MHz)

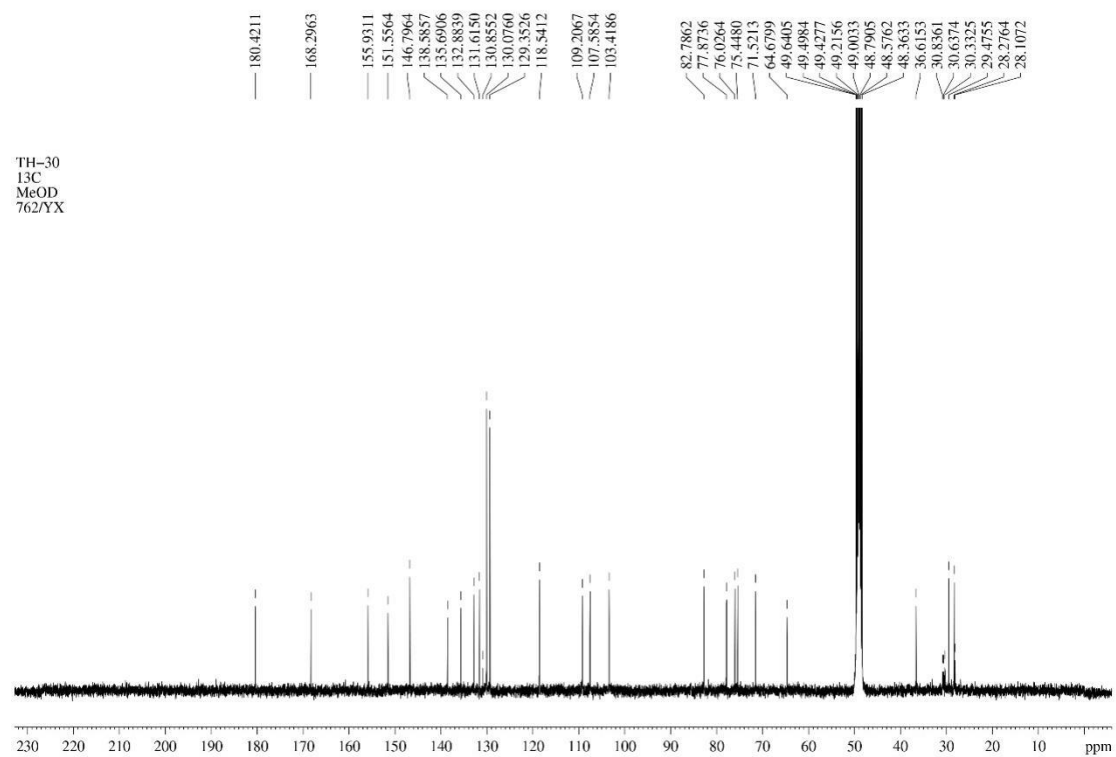


Figure S13. ^{13}C NMR spectrum of 2 in CD_3OD (100 MHz)

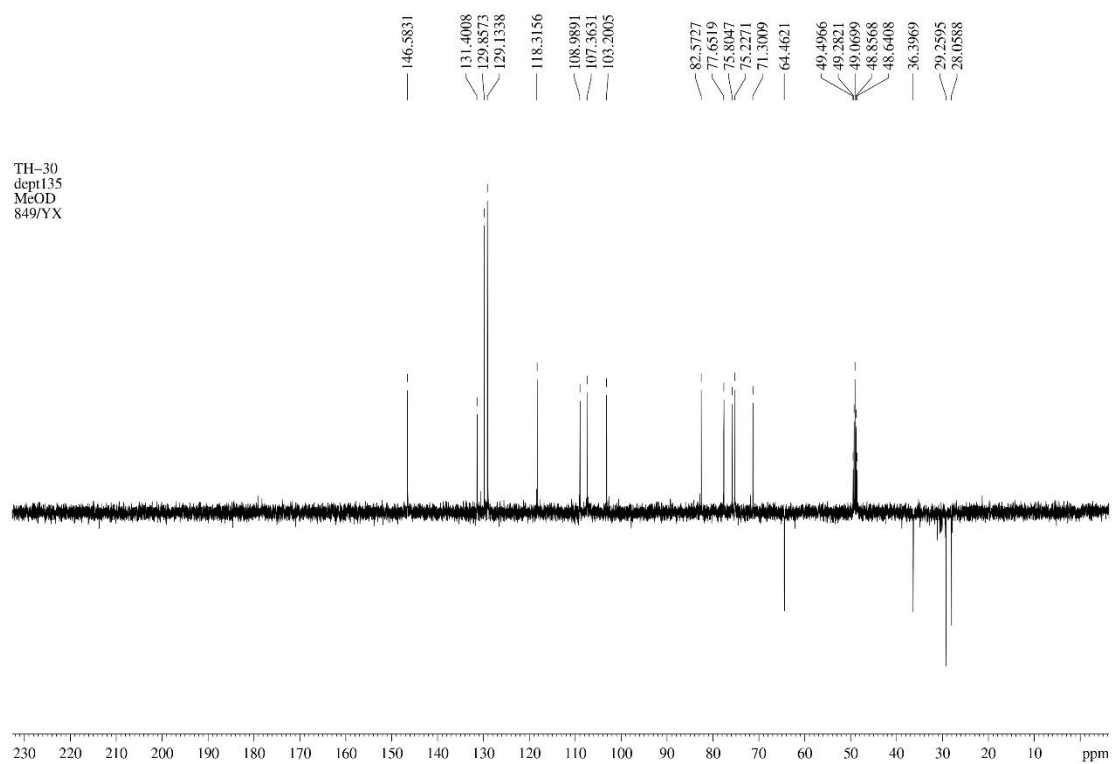


Figure S14. dept135° spectrum of 2 in CD₃OD (100 MHz)

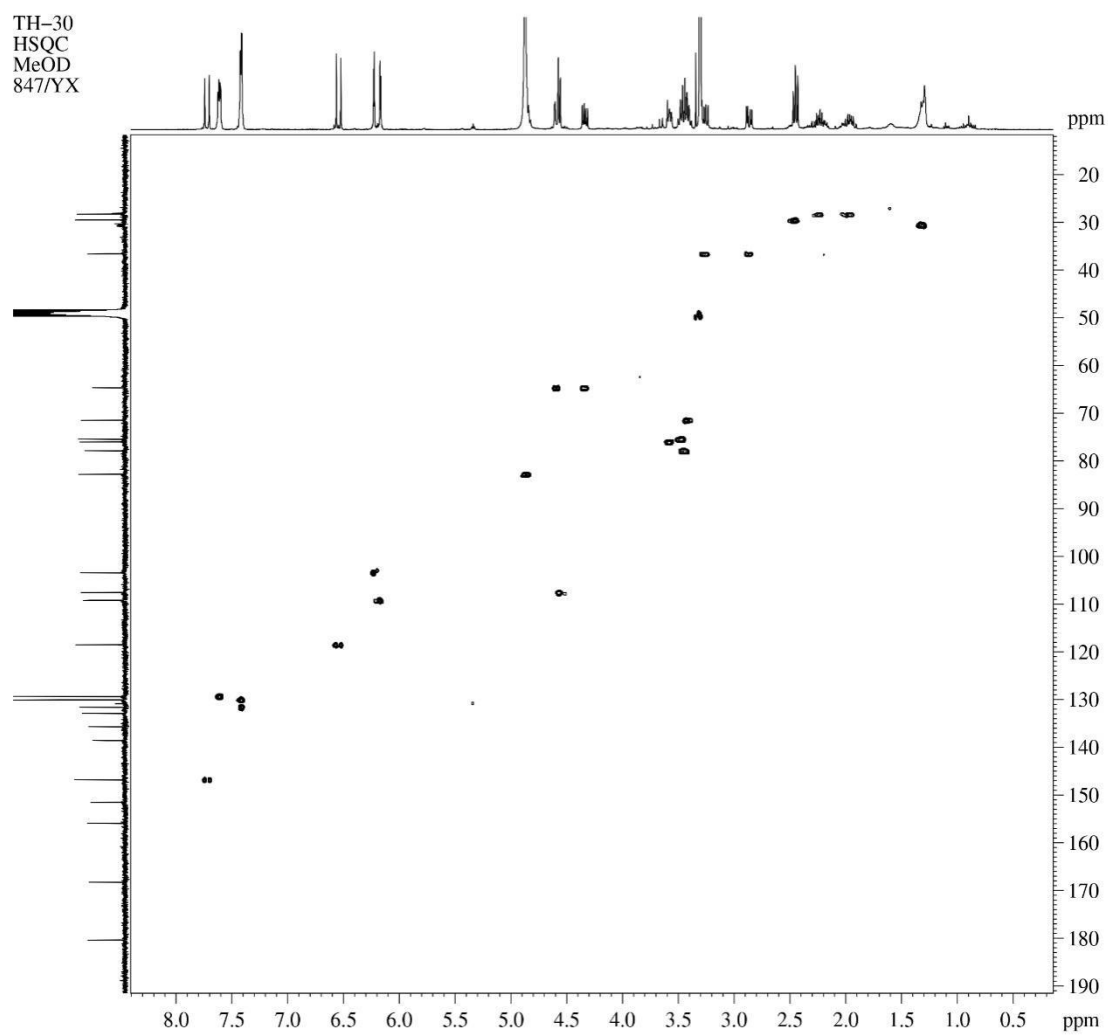


Figure S15. HSQC spectrum of **2** in CD₃OD (100 MHz)

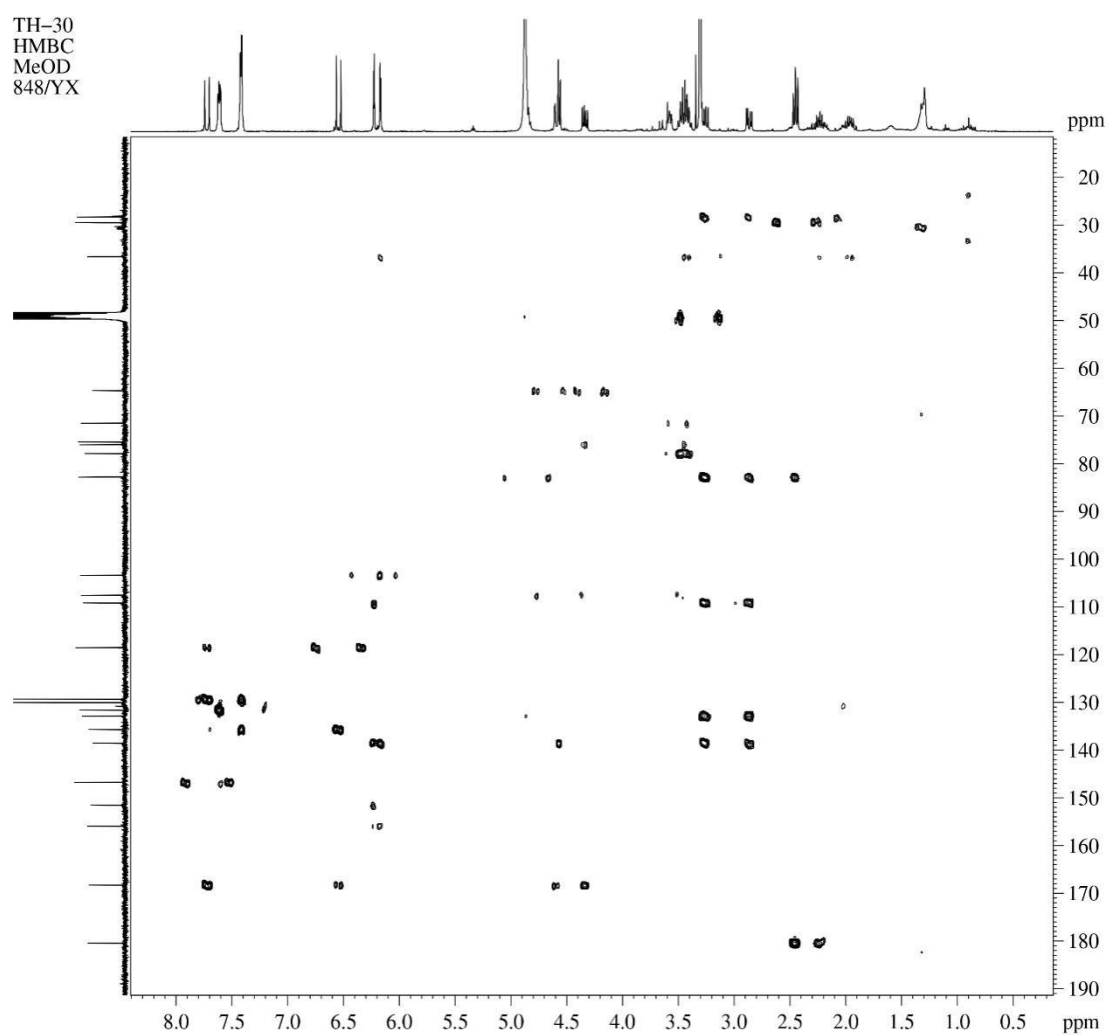


Figure S16. HMBC spectrum of **2** in CD₃OD (100 MHz)

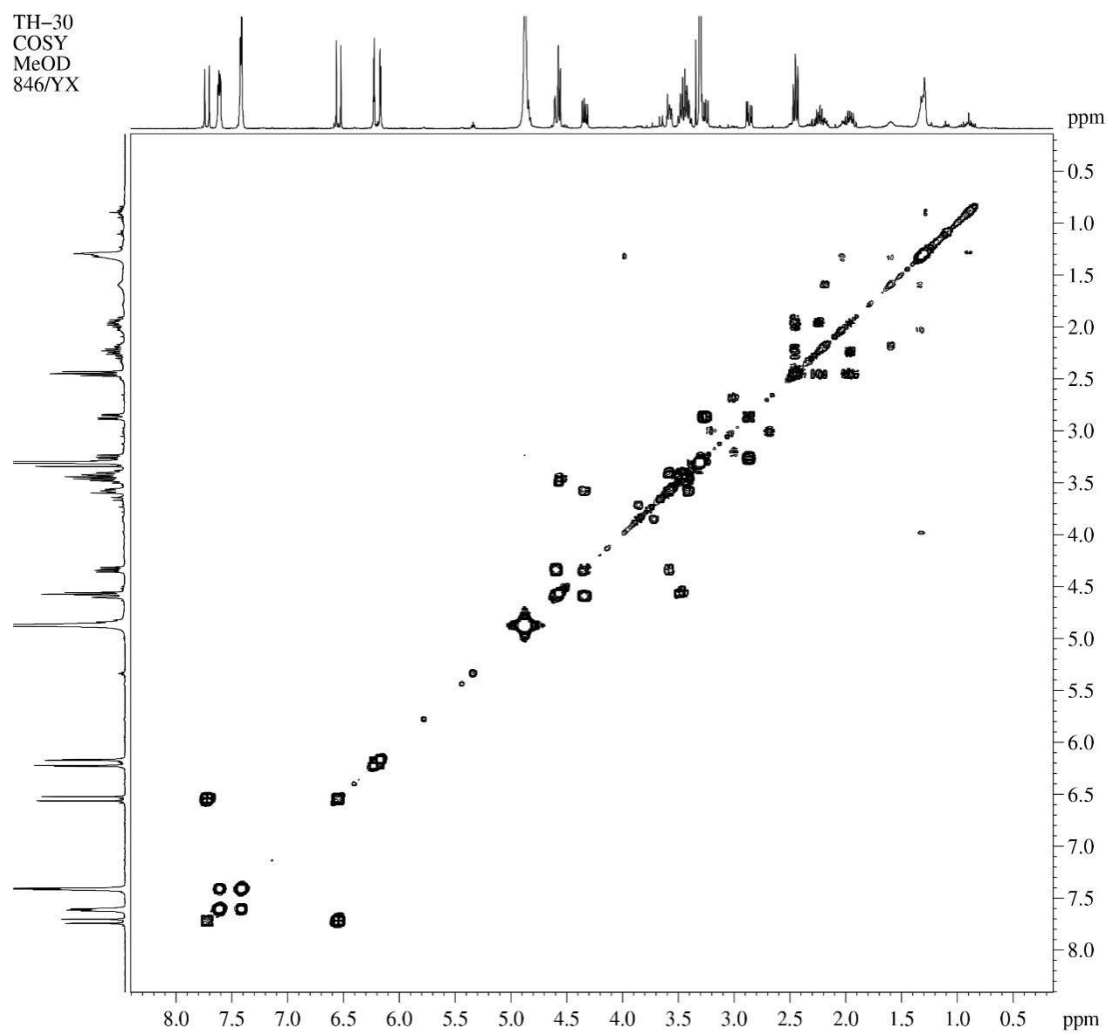


Figure S17. ^1H - ^1H COSY spectrum of **2** in CD_3OD (400 MHz).

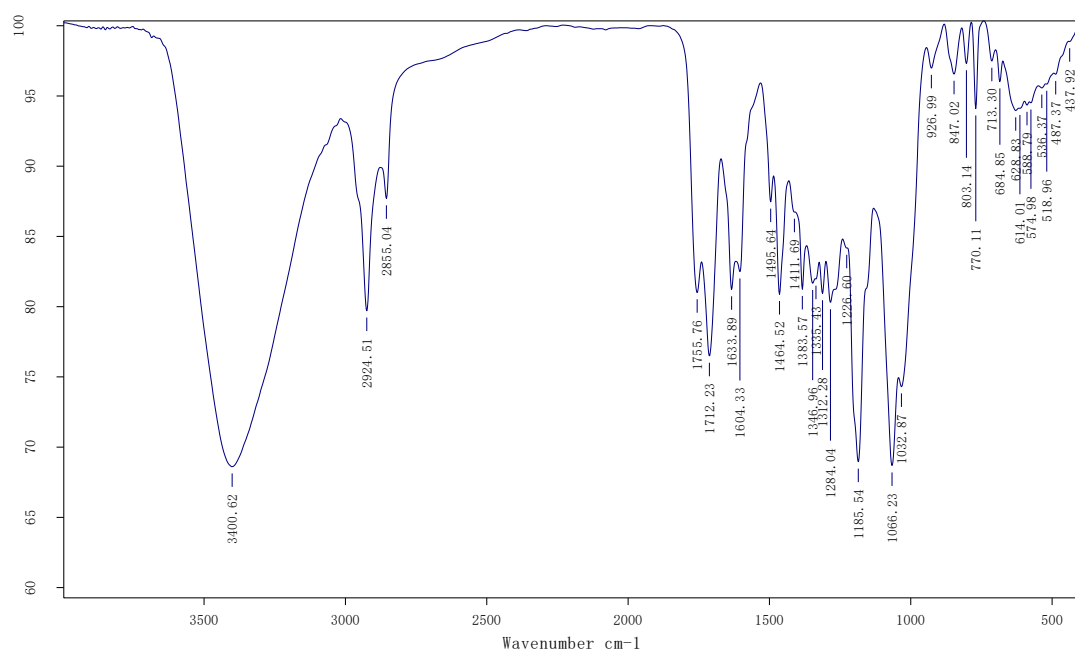


Figure S18. IR spectrum of 2.

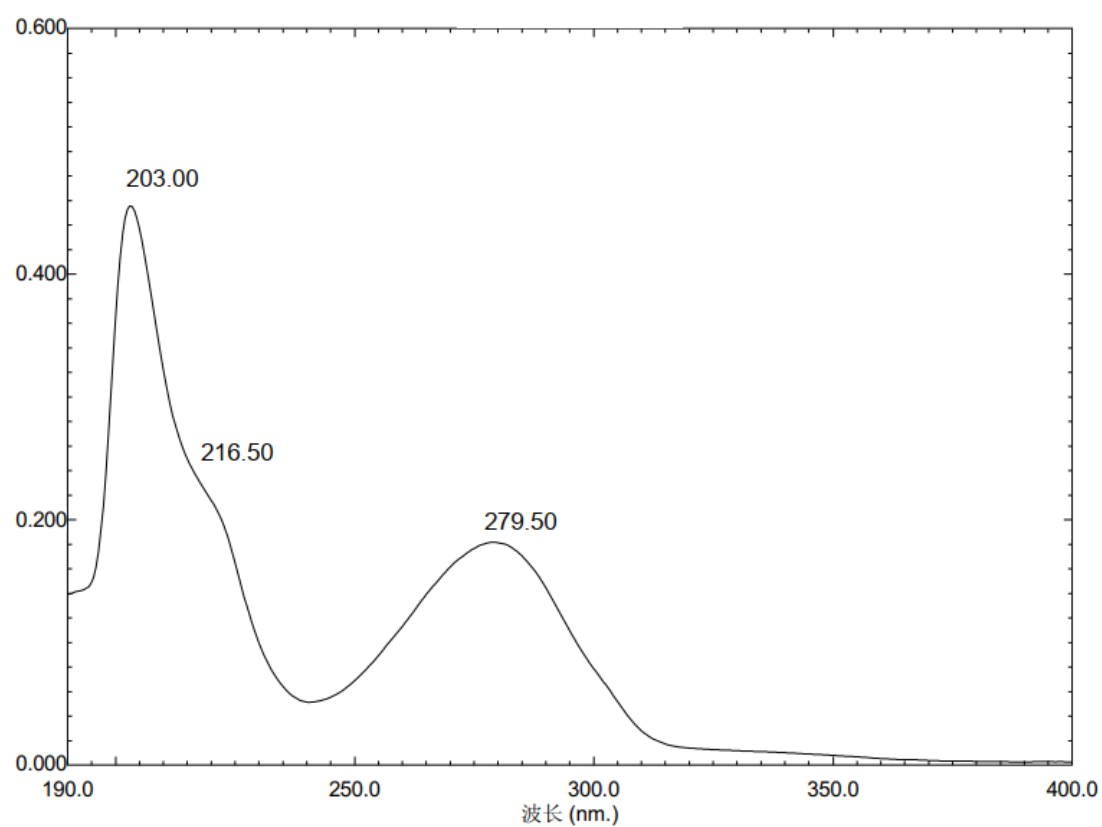


Figure S19. UV spectrum of 2.

Rudolph Research Analytical

This sample was measured on an Autopol VI, Serial #91058
Manufactured by Rudolph Research Analytical, Hackettstown, NJ, USA.

Measurement Date : Wednesday, 15-DEC-2021

Set Temperature : OFF

Time Delay : Disabled

Delay between Measurement : Disabled

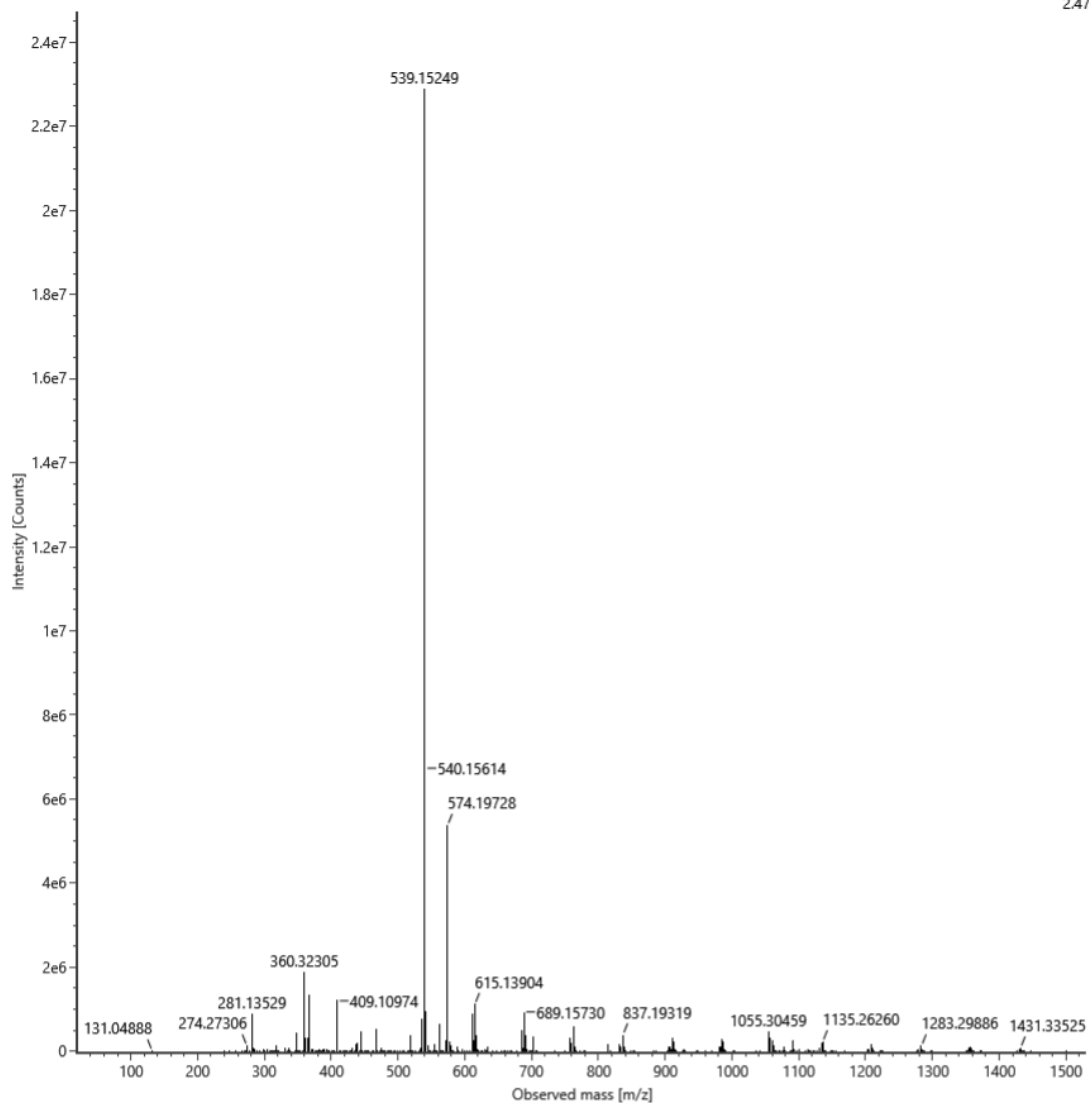
<u>n</u>	<u>Average</u>	<u>Std.Dev.</u>	<u>% RSD</u>	<u>Maximum</u>	<u>Minimum</u>					
5	-22.65	1.03	-4.54	-21.57	-23.98					
<u>S.No</u>	<u>Sample ID</u>	<u>Time</u>	<u>Result</u>	<u>Scale</u>	<u>OR °Arc</u>	<u>WLG.nm</u>	<u>Lq.mm</u>	<u>Conc.g/100ml</u>	<u>Temp.</u>	
1	TH-30	03:39:26 PM	-23.98	SR	-0.0199	589	100.00	0.083	19.7	
2	TH-30	03:39:34 PM	-23.25	SR	-0.0193	589	100.00	0.083	19.7	
3	TH-30	03:39:43 PM	-22.77	SR	-0.0189	589	100.00	0.083	19.7	
4	TH-30	03:39:51 PM	-21.69	SR	-0.0180	589	100.00	0.083	19.7	
5	TH-30	03:39:59 PM	-21.57	SR	-0.0179	589	100.00	0.083	19.7	

Figure S20. ORD spectrum of 2.

Item name: TH-30
Item description:

Channel name: 1: Average Time 0.1132 min : TOF MS (50-1500) ESI+ : Centroided : Combined

2.47e7



Add:Na+

Composition i-FIT Confidence (%) Predicted m/z m/z error (PPM)

C ₂₆ H ₂₈ O ₁₁	100.000000	539.152383	0.199168
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Figure S21. HRESIMS spectrum of 2.

Table S4 ^{13}C -NMR (100 MHz) and ^1H -NMR (400 MHz) Data of 3-4 in CD_3OD

Myricetrin (3)			Quercitrin (4)	
No	C	H	C	H
2	159.4	-	158.4	-
3	136.3	-	136.2	-
4	179.7	-	179.6	-
5	163.2	-	159.2	-
6	99.8	6.20 (1H, d, $J = 2.1$ Hz)	99.8	6.18 (1H, s),
7	165.9	-	165.8	-
8	94.7	6.36 (1H, d, $J = 2.1$ Hz)	94.7	6.34 (1H, s)
9	158.5	-	163.1	-
10	105.9	-	105.8	-
1'	121.9	-	122.9	-
2'	109.6	6.95 (2H, s)	116.3	7.22 – 7.37 (2H, m)
3'	146.9	-	146.3	-
4'	109.6	-	149.7	-
5'	146.9	-	116.9	6.90 (1H, d, $J = 8.3$ Hz)
6'	109.6	6.95 (s)	122.9	7.22 – 7.37 (2H, m).
1''	103.6	5.31 (1H, d, $J = 1.7$ Hz)	103.5	5.34 (1H, d, $J = 1.5$ Hz)
2''	71.9	4.22 (1H, dd, $J = 1.7, 3.4$ Hz)	72.0	4.23-3.37 (sugar H)
3''	72.1	3.79 (1H, dd, $J = 3.4, 9.5$ Hz)	72.1	-
4''	73.3	3.51 (1H, dd, $J = 2.6, 7.0$ Hz)	73.2	-
5''	72.0	3.51 – 3.69 (1H, m)	71.9	-
6''	17.7	0.96 (3H, d, $J = 6.2$ Hz)	17.6	0.94 (3H, d, $J = 6.0$ Hz)

Table S5 ^{13}C -NMR (100 MHz) and ^1H -NMR (400 MHz) Data of 5-6 in CD_3OD

No	Quercetin (5)		Tamarixetin-3-O-rhamnoside (6)	
	C	H	C	H
2	148.0	-	158.9	-
3	137.2	-	136.4	-
4	177.3	-	179.6	-
5	162.5	-	158.6	-
6	99.2	6.17 (1H, s)	94.7	6.38 (1H, d, $J = 2.1$ Hz)
7	165.6	-	166.0	-
8	94.4	6.38 (1H, s)	99.9	6.20 (1H, d, $J = 2.1$ Hz)
9	158.2	-	163.2	-
10	104.5	-	105.9	-
1'	124.2	-	124.2	-
2'	116.0	7.73 (1H, d, $J = 2.1$ Hz)	116.6	7.34 (1H, d, $J = 2.1$ Hz)
3'	146.2	-	147.7	-
4'	148.8	-	151.6	-
5'	116.2	6.88 (1H, d, $J = 8.4$ Hz)	112.3	7.07 (1H, d, $J = 8.5$ Hz)
6'	121.7	7.62 (1H, dd, $J = 8.5, 1.7$ Hz)	122.7	7.41 (1H, dd, $J = 8.5, 2.2$ Hz)
Rha-1	-	-	103.5	5.37 (1H, d, $J = 1.7$ Hz)
2	-	-	71.9	4.21 (1H, dd, $J = 1.7, 3.4$ Hz)
3	-	-	72.0	3.73 (1H, s)
4	-	-	72.1	3.35 (2H, m)
5	-	-	73.2	
6	-	-	17.7	0.92 (3H, d, $J = 5.6$ Hz)
C ₄ -Me	-	-	56.4	3.94 (3H, s)

Table S6. Effects of ardisicreolide A and B on NO release of raw cells

Compound	C(μ M/mL)	Mean \pm SD(μ M/mL)
Blank	-	12.80 \pm 1.97
LPS	-	55.25 \pm 6.59
DXM ^b	-	25.88 \pm 2.55
Ardisicreolide A	5	37.17 \pm 3.11
	20	40.35 \pm 3.95
	80	37.88 \pm 3.92
Ardisicreolide B	5	35.04 \pm 3.47
	20	36.75 \pm 2.69
	80	38.99 \pm 2.98

^b Dexamethasone (DXM) with 40 μ g/ml was used as the positive control.

Table S7. The effects of ardisicreolide A and B on production release of TNF- α , IL-1 β , IL-4 and IL-10

Compound	C(μ M/mL)	Mean \pm SD(pg/mL)			
		TNF- α	IL-1 β	IL-4	IL-10
Blank	-	43.82 \pm 6.22	53.06 \pm 4.54	32.5 \pm 4.88	89.94 \pm 10.00
LPS	-	162.34 \pm 8.64	192.02 \pm 18.67	118.81 \pm 6.34	365.33 \pm 28.36
DXM ^b	-	53.95 \pm 6.06	71.72 \pm 6.81	44.60 \pm 5.41	128.02 \pm 10.72
1	5	83.51 \pm 6.86	83.16 \pm 6.66	73.89 \pm 5.37	139.25 \pm 12.18
	20	81.64 \pm 7.32	75.21 \pm 7.40	67.75 \pm 5.58	131.26 \pm 11.93
	80	78.18 \pm 7.05	69.48 \pm 6.35	65.80 \pm 5.55	124.07 \pm 11.75
2	5	93.51 \pm 7.65	97.80 \pm 8.15	61.20 \pm 5.32	152.83 \pm 10.94
	20	90.00 \pm 7.39	95.55 \pm 8.80	56.18 \pm 5.11	146.79 \pm 11.35
	80	83.11 \pm 6.74	91.06 \pm 8.11	53.82 \pm 5.37	134.48 \pm 11.22

^bDexamethasone (DXM) with 40 μ g/ml was used as the positive control.

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