

*Supporting information for*

## **Melodinines Y<sub>1</sub>–Y<sub>4</sub>, four monoterpene indole alkaloids from *Melodinus henryi***

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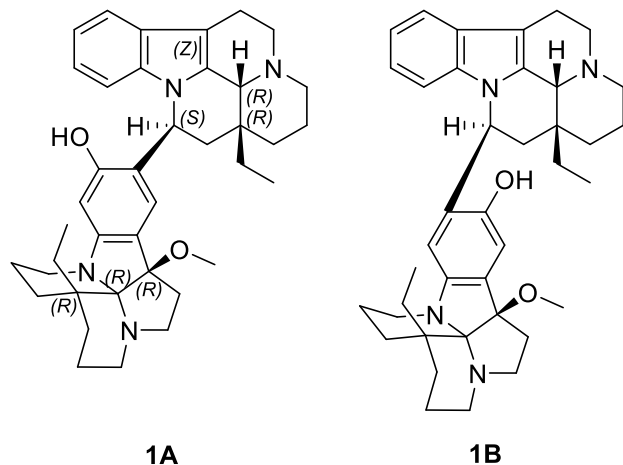
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## Section S1. Computational details for **1**

### S1.1. Computational details for compound **1** (NMR)



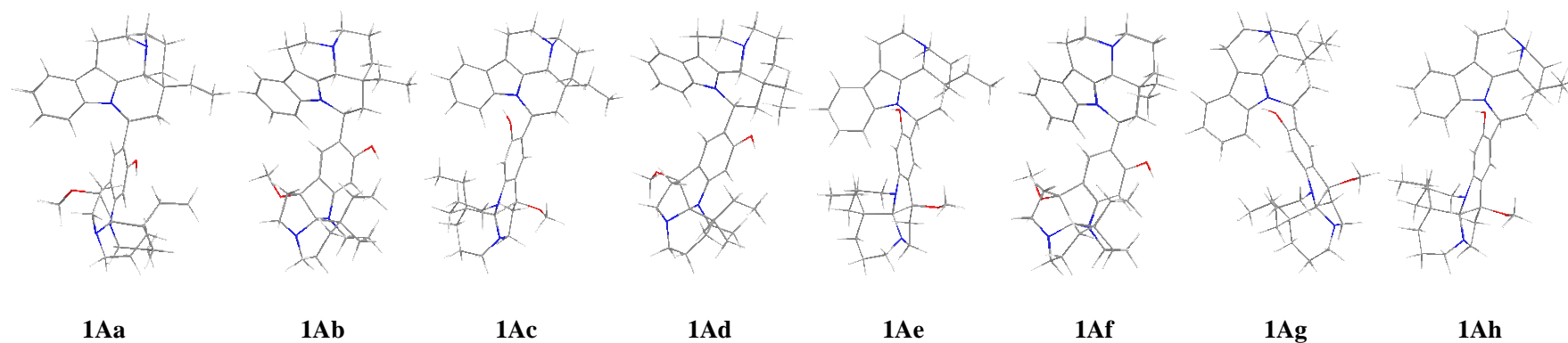
Conformation search based on molecular mechanics with MMFF force fields were performed for **1A** and **1B** gave 8 and 2 stable conformers with populations higher than 1%, respectively. All these conformers were further optimized by the density functional theory method at the B3LYP/6-31G(d) level by Gaussian 16 program package with g09 default keyword. Gauge Independent Atomic Orbital (GIAO) calculations of their  $^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts using density functional theory (DFT) at the mPW1PW91/6-311+G(d,p) level with the PCM model in methanol. The calculated NMR data of these conformers were averaged according to the Boltzmann distribution theory and their relative Gibbs free energy. The  $^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts for TMS were also calculated by the same procedures and used as the reference. After calculation, the experimental and calculated data were evaluated by linear correlation coefficients ( $R^2$ ) and the improved probability DP4+ method.

**Table S1.** Energy analysis for optimized geometries of dominant conformers **1Aa–1Ah** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>1Aa</b>	-1884.808178	-1884.770215	-1884.769271	-1884.875361	0.002622	1.645330	3.68%
<b>1Ab</b>	-1884.809349	-1884.771231	-1884.770287	-1884.876480	0.001503	0.943147	12.06%
<b>1Ac</b>	-1884.812073	-1884.774476	-1884.773532	-1884.877983	0	0	59.29%
<b>1Ad</b>	-1884.806825	-1884.768842	-1884.767897	-1884.873824	0.004159	2.609812	0.72%
<b>1Ae</b>	-1884.806714	-1884.768818	-1884.767874	-1884.873593	0.004390	2.754767	0.57%
<b>1Af</b>	-1884.808107	-1884.769971	-1884.769027	-1884.875017	0.002966	1.861193	2.56%
<b>1Ag</b>	-1884.810750	-1884.773088	-1884.772144	-1884.877005	0.000978	0.613704	21.03%
<b>1Ah</b>	-1884.805382	-1884.767470	-1884.766526	-1884.871942	0.006041	3.790785	0.10%

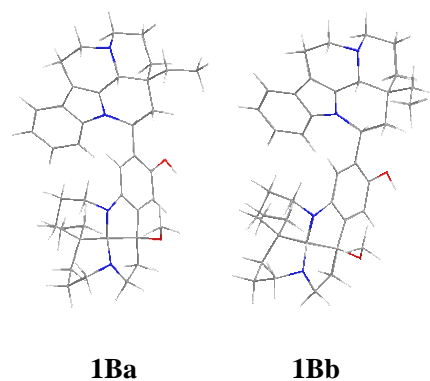
$E$ ,  $E'$ ,  $H$ ,  $G$ : total energy, total energy with zero point energy ( $ZPE$ ), enthalpy, and Gibbs free energy

**Figure S1.** Main conformers of **1A** in NMR and ECD calculations.



**Table S2.** Energy analysis for optimized geometries of dominant conformers **1Ba–1Bb** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>1Ba</b>	-1884.807065	-1884.769150	-1884.768206	-1884.873448	0	0	79.01%
<b>1Bb</b>	-1884.805710	-1884.767744	-1884.766799	-1884.872197	0.001251	0.785014	20.99%

**Figure S2.** Main conformers of **1B** in NMR calculation.**Table S3.** Calculated  $^{13}\text{C}$  NMR results for **1A**

No.	<b>1Aa</b>	<b>1Ab</b>	<b>1Ac</b>	<b>1Ad</b>	<b>1Ae</b>	<b>1Af</b>	<b>1Ag</b>	<b>1Ah</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	43.8155	43.6622	42.6273	43.3929	44.3148	43.1404	42.1378	43.8331	145.5	131.9	138.1	-6.2
3	141.1333	140.5529	138.7807	142.1730	139.9825	141.7676	139.1767	140.0338	48.9	45.5	47.7	-2.2
5	132.8267	133.1572	133.7171	133.1148	132.3610	133.3152	133.2178	132.4859	54.7	52.0	53.1	-1.1
6	167.4083	167.1311	167.1578	167.3096	167.0873	167.1782	167.1042	167.3670	21.0	17.6	21.6	-4.0
7	76.5263	76.8001	72.0424	76.4946	78.2087	76.7019	72.7938	78.0917	115.0	105.2	109.6	-4.4
8	53.0169	52.8209	51.6815	53.0291	53.6364	52.8697	51.8084	53.7005	136.2	129.3	129.5	-0.2
9	65.8718	65.8850	65.0040	65.8500	65.6984	65.9235	65.1764	65.6922	123.0	119.2	117.1	2.1
10	64.1157	63.9992	61.8716	63.9088	64.2758	63.8765	62.1227	64.5344	125.8	120.7	119.7	1.0

11	63.3999	63.1303	62.0009	63.4842	63.5846	63.1126	62.0563	63.7978	125.9	121.9	119.8	2.1
12	70.1313	70.0448	69.8822	69.5427	71.1770	69.5109	69.7349	71.2682	118.3	113.1	112.7	0.4
13	44.7158	44.6233	42.1339	45.6239	45.3714	45.3320	42.1187	45.2570	145.5	137.7	138.2	-0.5
14	163.3000	163.2943	162.8873	163.5044	162.8680	163.5494	163.3785	163.0179	25.1	20.6	25.4	-4.8
15	163.2791	162.9269	163.4354	156.0707	162.7263	156.2997	156.2767	155.9682	26.6	24.4	26.7	-2.3
16	134.5396	134.9375	126.2983	134.6211	128.0156	135.4756	126.3932	127.8352	60.2	49.9	58.3	-8.4
17	142.2479	142.5618	144.3825	143.3194	145.5394	143.6704	145.5056	146.7193	43.9	42.7	43.0	-0.3
18	178.5711	178.6659	178.6669	177.6490	178.9128	177.4698	177.2888	177.4463	9.8	7.7	11.1	-3.4
19	155.6013	155.1907	154.9892	156.9633	155.2830	157.0073	156.1506	156.1262	32.8	29.3	32.6	-3.3
20	148.5956	148.7386	148.0116	148.4842	147.7107	148.6739	148.5075	148.7831	39.9	36.4	39.3	-2.9
21	123.8228	123.5435	125.5633	130.0157	123.9896	130.0591	132.1584	130.8793	61.4	61.2	59.4	1.8
2'	143.4163	142.8184	143.6746	143.3703	143.6014	142.9225	143.6440	143.7273	44.6	58.2	43.7	14.5
3'	143.3455	143.7753	143.6143	142.6189	143.0523	143.6835	143.1440	142.6486	44.7	63.6	43.7	19.9
5'	124.2437	125.7541	124.4432	124.2493	124.3441	125.5940	124.2478	124.4377	63.6	64.5	61.4	3.1
6'	155.6124	162.6077	155.3228	155.5498	155.4600	162.7071	155.4009	155.6412	31.8	34.0	31.6	2.4
7'	83.6939	84.5654	83.9632	83.5959	83.9250	84.2065	83.8489	84.0144	104.2	94.3	99.5	-5.2
8'	62.9919	60.9399	62.8438	62.5864	63.3388	60.9208	62.8868	63.4819	125.6	117.4	119.5	-2.1
9'	60.0853	59.4859	58.8750	60.2396	56.6900	59.6629	59.0665	56.9409	129.1	124.9	122.8	2.1
10'	65.1296	67.7496	67.4285	64.9315	66.6981	67.5545	67.3657	66.7402	120.8	122.7	115.1	7.6
11'	26.6068	26.9020	23.0602	26.8190	24.5974	27.1919	22.7305	24.2852	164.5	158.5	155.9	2.6
12'	90.0227	89.7506	87.0925	89.8724	88.9909	89.5192	87.0933	88.9897	100.6	98.3	96.1	2.2
13'	24.9391	23.3038	23.5993	24.7622	23.9617	23.5321	23.4971	24.1932	164.6	150.8	156.0	-5.2
14'	167.3224	167.4382	167.3659	167.1072	166.6544	167.1307	166.3516	166.0066	21.0	26.3	21.6	4.7
15'	158.5885	158.0603	160.0394	159.1285	159.9295	159.2984	160.2610	160.2505	28.4	26.9	28.5	-1.6
16'	167.1504	167.3068	167.1129	167.0357	167.1533	166.9743	167.4742	167.0906	21.0	20.0	21.5	-1.5
17'	156.4030	156.6812	156.8002	156.1472	156.6707	156.6551	156.9604	156.5429	31.4	33.2	31.3	1.9
18'	179.0541	178.7881	178.5631	178.4839	178.1884	178.2497	178.5247	178.2841	9.6	7.1	10.8	-3.7
19'	158.3389	158.2044	158.1868	158.1922	158.6673	157.8750	158.2018	158.4950	30.0	34.5	30.0	4.5

20'	140.8938	140.7694	139.3168	140.8448	139.3834	140.3925	139.3655	139.3930	48.6	32.8	47.4	-14.6
21'	83.7195	79.0207	84.6484	83.8676	84.5397	79.8504	84.2893	84.4205	104.5	103.7	99.7	4.0
-OCH <sub>3</sub>	134.5929	138.8864	133.8743	134.1624	133.9411	138.7728	134.0304	134.0527	53.5	53.0	52.0	1.0
Population	3.68%	12.06%	59.29%	0.72%	0.57%	2.56%	21.03%	0.10%			RMSD	5.71

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

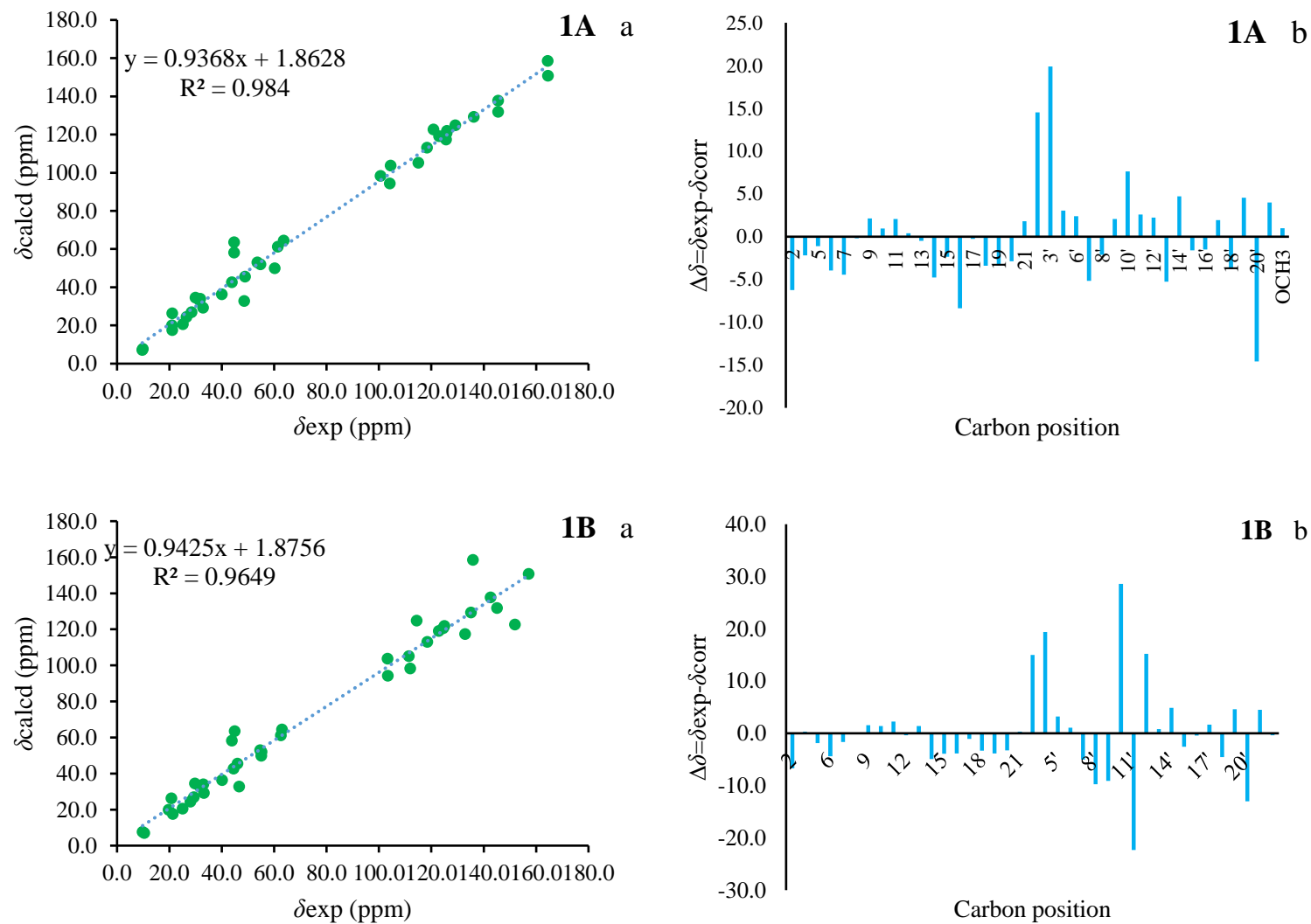
**Table S4.** Calculated <sup>13</sup>C NMR results for **1B**

No.	<b>1Ba</b>	<b>1Bb</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	43.1312	43.1532	145.0	131.9	138.6	-6.7
3	142.5085	141.2367	45.9	45.5	45.2	0.3
5	132.9010	133.5056	55.2	52.0	53.9	-1.9
6	167.0130	166.3282	21.3	17.6	22.0	-4.4
7	76.8020	76.6709	111.4	105.2	106.9	-1.7
8	53.1251	52.8565	135.1	129.3	129.2	0.1
9	65.3881	65.2540	122.8	119.2	117.6	1.6
10	63.5722	63.6879	124.6	120.7	119.3	1.4
11	63.2386	63.1275	125.0	121.9	119.7	2.2
12	69.8157	69.5210	118.4	113.1	113.5	-0.4
13	45.5483	45.6453	142.6	137.7	136.3	1.4
14	163.0851	163.2316	25.1	20.6	25.5	-4.9
15	161.4662	155.1986	28.0	24.4	28.3	-3.9
16	133.0036	133.6690	55.0	49.9	53.7	-3.8
17	143.2273	145.6493	44.4	42.7	43.8	-1.1
18	178.7125	177.5026	9.7	7.7	11.0	-3.3
19	154.7919	155.7944	33.2	29.3	33.1	-3.8
20	147.9192	148.7242	40.1	36.4	39.7	-3.3
21	124.5112	129.7295	62.6	61.2	60.9	0.3

2'	144.3497	144.3015	43.8	58.2	43.2	15.0
3'	143.2106	143.2866	45.0	63.6	44.2	19.4
5'	125.1541	125.2281	63.0	64.5	61.3	3.2
6'	155.1687	155.4165	33.0	34.0	32.9	1.1
7'	85.0082	83.8912	103.4	94.3	99.3	-5.0
8'	55.1539	55.7621	132.9	117.4	127.1	-9.7
9'	76.2547	76.0097	112.0	98.3	107.4	-9.1
10'	52.2437	52.5207	135.9	158.5	129.9	28.6
11'	36.3433	36.1454	151.9	122.7	145.0	-22.3
12'	73.7739	73.9181	114.4	124.9	109.7	15.2
13'	30.9768	31.3281	157.1	150.8	150.0	0.8
14'	167.4953	167.3047	20.7	26.3	21.4	4.9
15'	158.8980	158.9796	29.3	26.9	29.5	-2.6
16'	168.5091	168.4037	19.7	20.0	20.4	-0.4
17'	156.6775	156.9392	31.4	33.2	31.5	1.7
18'	177.8028	177.8630	10.4	7.1	11.6	-4.5
19'	158.4859	158.4520	29.7	34.5	29.9	4.6
20'	141.5538	141.6426	46.6	32.8	45.8	-13.0
21'	85.0326	84.3559	103.3	103.7	99.2	4.5
-OCH <sub>3</sub>	133.5116	133.6287	54.6	53.0	53.4	-0.4
Population	79.01%	20.99%			RMSD	8.47

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

**Figure S3.**  $^{13}\text{C}$  NMR calculation results of two possible isomers of **1**. (a) Linear correlation plots of predicted versus experimental  $^{13}\text{C}$  NMR chemical shifts. (b) Relative errors between the predicted  $^{13}\text{C}$  NMR chemical shifts of two potential structures and recorded  $^{13}\text{C}$  NMR data.





**Table S5.** DP4+ analysis results of **1A** (Isomer 1) and **1B** (Isomer 2)

	A	B	C	D	E	F	G	H
1	<b>Functional</b>		<b>Solvent?</b>		<b>Basis Set</b>		<b>Type of Data</b>	
2	mPW1PW91		PCM		6-311+G(d,p)		Shielding Tensors	
3								
12			<b>DP4+</b>	<b>100.00%</b>	<b>0.00%</b>	-	-	-
14	<b>Nuclei</b>	<b>sp2?</b>	<b>xperimental</b>	<b>Isomer 1</b>	<b>Isomer 2</b>	<b>Isomer 3</b>	<b>Isomer 4</b>	<b>Isomer 5</b>
15	C	x	131.9	42.72	43.14			
16	C		45.5	139.27	142.24			
17	C		52	133.49	133.03			
18	C		17.6	167.15	166.87			
19	C	x	105.2	73.13	76.77			
20	C	x	129.3	51.95	53.07			
21	C	x	119.2	65.21	65.36			
22	C	x	120.7	62.35	63.60			
23	C	x	121.9	62.25	63.22			
24	C	x	113.1	69.88	69.75			
25	C	x	137.7	42.65	45.57			
26	C		20.6	163.08	163.12			
27	C		24.4	161.62	160.15			
28	C		49.9	127.97	133.14			
29	C		42.7	144.30	143.74			
30	C		7.7	178.34	178.46			
31	C		29.3	155.35	155.00			
32	C		36.4	148.24	148.09			
33	C		61.2	126.79	125.61			
34	C		58.2	143.53	144.34			
35	C		63.6	143.52	143.23			
36	C		64.5	124.58	125.17			
37	C		34	156.42	155.22			
38	C		94.3	84.01	84.77			
39	C	x	117.4	62.58	55.28			
40	C	x	124.9	59.05	76.20			
41	C	x	122.7	67.35	52.30			
42	C	x	158.5	23.73	36.30			
43	C	x	98.3	87.62	73.80			
44	C	x	150.8	23.60	31.05			
45	C		26.3	167.15	167.46			
46	C		26.9	159.77	158.92			

47	C		20	167.21	168.49			
48	C		33.2	156.80	156.73			
49	C		7.1	178.59	177.82			
50	C		34.5	158.19	158.48			
51	C		32.8	139.60	141.57			
52	C		103.7	83.73	84.89			
53	C		53	134.67	133.54			
54								
55	H		2.58	29.09	29.28			
56	H		2.8	29.07	29.20			
57	H		3.46	28.51	28.57			
58	H		3.46	28.32	28.41			
59	H		2.82	29.30	29.12			
60	H		3.03	28.59	28.61			
61	H	x	7.41	24.04	23.87			
62	H	x	6.94	24.31	24.37			
63	H	x	6.81	24.48	24.52			
64	H	x	6.63	24.36	24.69			
65	H		1.46	30.27	30.33			
66	H		1.82	30.05	29.90			
67	H		1.15	30.60	30.55			
68	H		1.54	30.45	30.40			
69	H		5.55	26.77	26.15			
70	H		1.67	29.84	30.24			
71	H		2.37	29.74	29.42			
72	H		0.93	30.87	31.00			
73	H		0.93	30.88	30.87			
74	H		0.93	30.99	30.88			
75	H		1.54	30.30	30.41			
76	H		2.09	29.60	29.63			
77	H		4.38	27.65	27.47			
78	H		2.97	28.50	28.84			
79	H		3.26	28.33	28.82			
80	H		2.75	28.75	28.64			
81	H		3.45	28.42	28.60			
82	H		2.91	28.42	28.44			
83	H		3.33	27.75	27.84			

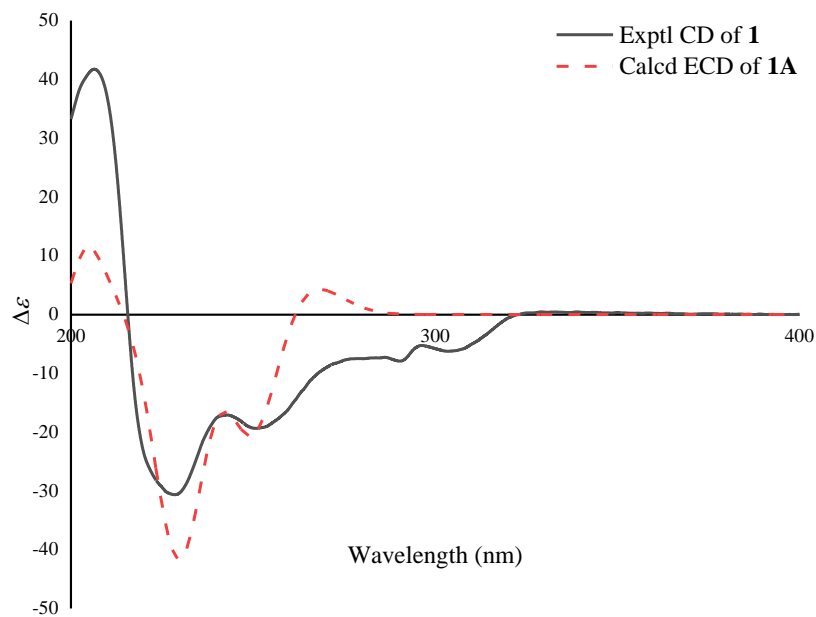
84	H		2.39	29.73	29.72			
85	H		2.48	29.63	29.55			
86	H	x	6.78	24.67	25.55			
87	H	x	6.48	25.71	25.03			
88	H		1.7	30.14	30.27			
89	H		1.7	29.73	29.68			
90	H		1.91	30.54	30.70			
91	H		2.41	30.28	30.21			
92	H		1.82	30.56	31.11			
93	H		2.3	29.80	30.61			
94	H		1.4	30.04	30.21			
95	H		1.61	29.85	30.03			
96	H		0.89	31.03	30.96			
97	H		0.89	31.07	31.10			
98	H		0.89	30.98	31.12			
99	H		1.34	30.82	30.94			
100	H		1.34	30.50	30.66			
101	H		2.68	28.82	29.20			
102	H		2.68	28.37	28.21			
103	H		2.68	29.15	28.81			

	A	B	C	D	E	F	G	H	
1	<b>Functional</b>		<b>Solvent?</b>		<b>Basis Set</b>		<b>Type of Data</b>		
2	mPW1PW91		PCM		6-311+G(d, p)		Shielding Tensors		
3									
4			Isomer 1	Isomer 2	Isomer 3	Isomer 4	Isomer 5	Isomer 6	
5	sDP4+ (H data)		2.27%		97.73%	-	-	-	-
6	sDP4+ (C data)		100.00%		0.00%	-	-	-	-
7	sDP4+ (all data)		100.00%		0.00%	-	-	-	-
8	uDP4+ (H data)		6.07%		93.93%	-	-	-	-
9	uDP4+ (C data)		100.00%		0.00%	-	-	-	-
10	uDP4+ (all data)		100.00%		0.00%	-	-	-	-
11	DP4+ (H data)		0.15%		99.85%	-	-	-	-
12	DP4+ (C data)		100.00%		0.00%	-	-	-	-
13	DP4+ (all data)		100.00%		0.00%	-	-	-	-

## S1.2. Computational details for compound 1 (ECD)

Conformation search based on molecular mechanics with MMFF force fields were performed for **1A** gave 8 stable conformers with populations higher than 1%. All these conformers were further optimized by the density functional theory method at the B3LYP/6-311G(2d,p) level by Gaussian 16 program package. The ECD were calculated using density functional theory (TDDFT) at B3LYP/6-311G(d) level in methanol with IEFPCM model. The calculated ECD curves were all generated using SpecDis 1.71 program package ( $\sigma = 0.15$  eV, and UV shift -31 nm) and the calculated ECD data of all conformers were Boltzmann averaged by Gibbs free energy.

**Figure S4.** Comparison of the calculated ECD spectra for with the experimental spectrum of **1** in methanol with PCM model.



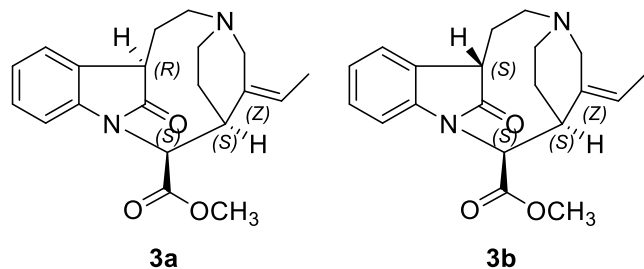
**Table S6.** Energy analysis for optimized geometries of dominant conformers **1Aa–1Ah** at B3LYP/6-311G(2d,p) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>1Aa</b>	-1885.319891	-1885.281611	-1885.280667	-1885.387417	0.002103	1.319652	4.78%
<b>1Ab</b>	-1885.321033	-1885.282544	-1885.2816	-1885.388853	0.000667	0.418549	21.87%
<b>1Ac</b>	-1885.323068	-1885.285164	-1885.28422	-1885.38952	0	0	44.35%
<b>1Ad</b>	-1885.318496	-1885.280166	-1885.279222	-1885.386149	0.003371	2.115335	1.25%
<b>1Ae</b>	-1885.318432	-1885.280223	-1885.279279	-1885.385334	0.004186	2.626755	0.53%
<b>1Af</b>	-1885.319663	-1885.281107	-1885.280162	-1885.387746	0.001774	1.113202	6.77%
<b>1Ag</b>	-1885.321953	-1885.283946	-1885.283002	-1885.388782	0.000738	0.463102	20.29%
<b>1Ah</b>	-1885.317177	-1885.278892	-1885.277948	-1885.384282	0.005238	3.286895	0.17%

$E, E', H, G$ : total energy, total energy with zero point energy ( $ZPE$ ), enthalpy, and Gibbs free energy

## Section S2. Computational details for **3**

### S2.1. Computational details for compound **3** (NMR)

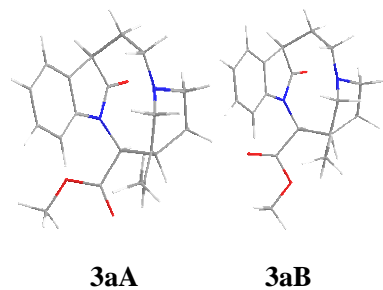


Conformation search based on molecular mechanics with MMFF force fields were performed for **3a**, and **3b** gave 2, and 1 stable conformers with populations higher than 1%, respectively. All these conformers were further optimized by the density functional theory method at the B3LYP/6-31G(d) level by Gaussian 16 program package with g09 default keyword. Gauge Independent Atomic Orbital (GIAO) calculations of their  $^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts using density functional theory (DFT) at the mPW1PW91/6-311+G(d,p) level with the PCM model in methanol. The calculated NMR data of these conformers were averaged according to the Boltzmann distribution theory and their relative Gibbs free energy. The  $^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts for TMS were also calculated by the same procedures and used as the reference. After calculation, the experimental and calculated data were evaluated by linear correlation coefficients ( $R^2$ ) and the improved probability DP4+ method.

**Table S7.** Energy analysis for optimized geometries of dominant conformers **3aA–3aB** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>3aA</b>	-1111.332647	-1111.310645	-1111.309701	-1111.382716	0	0	91.65%
<b>3aB</b>	-1111.330373	-1111.308376	-1111.307431	-1111.380455	0.002261	1.418799	8.35%

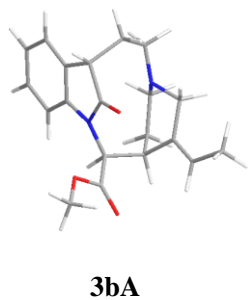
**Figure S5.** Main conformers of **3a** in NMR and ECD calculations.



**Table S8.** Energy analysis for optimized geometries of dominant conformers **3bA** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>3bA</b>	-1111.331138	-1111.309237	-1111.308293	-1111.380773	0	0	100.00%

**Figure S6.** Main conformers of **3b** in NMR calculation.



**Table S9.** Calculated  $^{13}\text{C}$  NMR results for **3a**

No.	<b>3aA</b>	<b>3aB</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	-1.4116	-2.2914	189.67	184.1	181.4	2.7
3	141.9232	142.2114	46.23	46.1	43.0	3.1
5	134.2861	134.4603	53.88	51.7	50.4	1.3
6	151.2742	151.6327	36.88	31.9	34.0	-2.1
7	138.1072	137.8675	50.09	45.2	46.7	-1.5
8	51.3223	51.1203	136.88	130.1	130.4	-0.3
9	56.7021	57.4232	131.42	125.0	125.2	-0.2
10	60.9598	61.0471	127.21	123.2	121.1	2.1
11	55.4534	55.5887	132.72	127.3	126.4	0.9
12	70.1696	67.2080	118.26	114.3	112.5	1.8
13	35.6491	35.4396	152.55	145.8	145.6	0.2
14	154.5225	153.8186	33.72	31.0	30.9	0.1
15	143.5664	140.4704	44.87	35.2	41.7	-6.5
16	121.2378	118.1563	67.20	63.2	63.2	0.0
17	5.2266	6.4705	182.85	172.7	174.8	-2.1
18	172.2769	172.6176	15.88	13.9	13.7	0.2
19	56.2750	57.7685	131.78	121.7	125.5	-3.8
20	42.6596	41.4894	145.62	136.5	138.9	-2.4
21	129.8492	130.3397	58.29	59.6	54.6	5.0
-OCH <sub>3</sub>	132.9766	132.7825	55.22	53.0	51.7	1.3
Population	91.65%	8.35%			RMSD	2.5

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

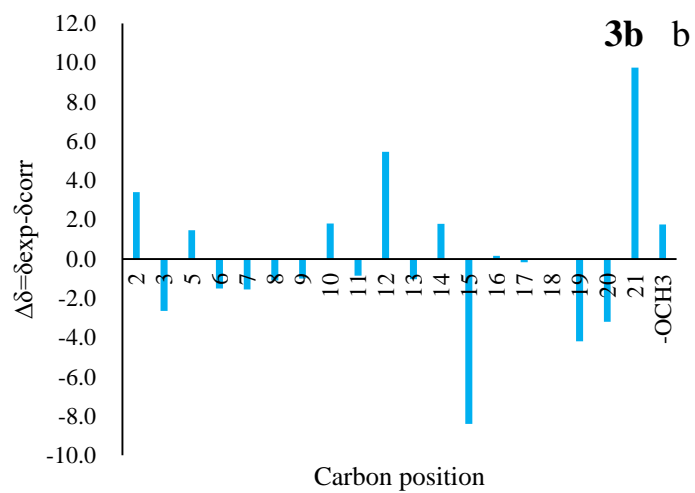
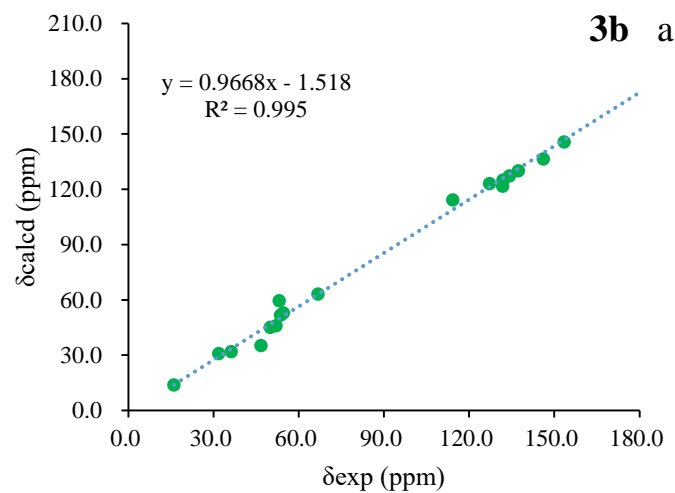
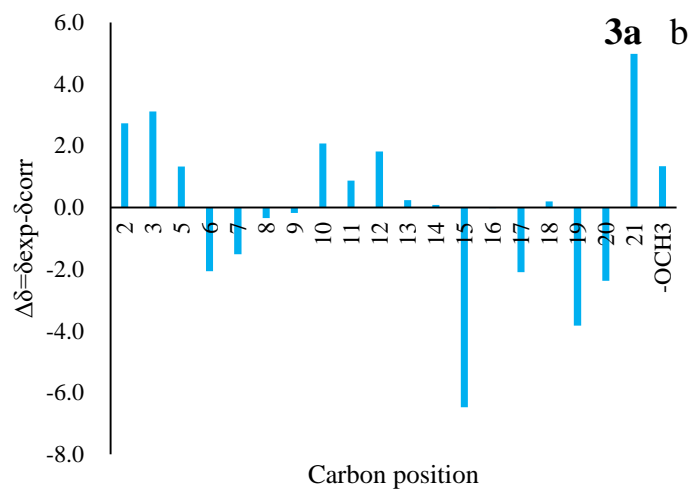
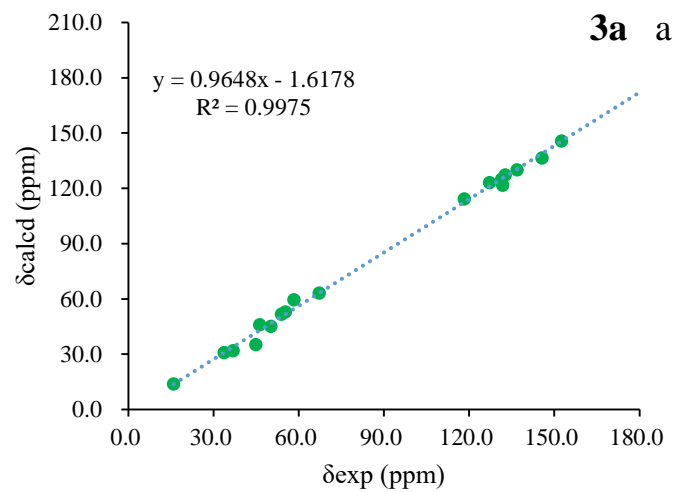


**Table S10.** Calculated  $^{13}\text{C}$  NMR results for **3b**

No.	<b>3bA</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	-0.2863	188.47	184.1	180.7	3.4
3	136.1902	51.99	46.1	48.7	-2.6
5	134.6476	53.53	51.7	50.2	1.5
6	152.0603	36.12	31.9	33.4	-1.5
7	138.2472	49.93	45.2	46.8	-1.6
8	50.9131	137.27	130.1	131.2	-1.1
9	56.2807	131.90	125.0	126.0	-1.0
10	61.0550	127.13	123.2	121.4	1.8
11	54.0685	134.11	127.3	128.1	-0.8
12	74.0292	114.15	114.3	108.8	5.5
13	34.7420	153.44	145.8	146.8	-1.0
14	156.3956	31.79	31.0	29.2	1.8
15	141.5133	46.67	35.2	43.6	-8.4
16	121.4071	66.77	63.2	63.0	0.2
17	7.8113	180.37	172.7	172.9	-0.2
18	172.1963	15.98	13.9	13.9	0.0
19	56.3919	131.79	121.7	125.9	-4.2
20	42.1198	146.06	136.5	139.7	-3.2
21	135.0487	53.13	59.6	49.9	9.7
-OCH <sub>3</sub>	133.6053	54.58	53.0	51.2	1.8
Population	100%			RMSD	3.6

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

**Figure S7.**  $^{13}\text{C}$  NMR calculation results of two possible isomers of **3**. (a) Linear correlation plots of predicted versus experimental  $^{13}\text{C}$  NMR chemical shifts. (b) Relative errors between the predicted  $^{13}\text{C}$  NMR chemical shifts of two potential structures and recorded  $^{13}\text{C}$  NMR data.



**Table S11.** DP4+ analysis results of **3a** (Isomer 1) and **3b** (Isomer 2)

	A	B	C	D	E	F	G	H
1	<b>Functional</b>		<b>Solvent?</b>		<b>Basis Set</b>		<b>Type of Data</b>	
2	mPW1PW91		PCM		6-311+G(d,p)		Shielding Tensors	
3								
12			DP4+	100.00%	0.00%	-	-	-
14	<b>Nuclei</b>	<b>sp2?</b>	<b>xperimenta</b>	<b>Isomer 1</b>	<b>Isomer 2</b>	<b>Isomer 3</b>	<b>Isomer 4</b>	<b>Isomer 5</b>
15	C	x	184.1	-1.49	-0.29			
16	C		46.1	141.95	136.19			
17	C		51.7	134.30	134.65			
18	C		31.9	151.30	152.06			
19	C		45.2	138.09	138.25			
20	C	x	130.1	51.31	50.91			
21	C	x	125	56.76	56.28			
22	C	x	123.2	60.97	61.06			
23	C	x	127.3	55.46	54.07			
24	C	x	114.3	69.92	74.03			
25	C	x	145.8	35.63	34.74			
26	C		31	154.46	156.40			
27	C		35.2	143.31	141.51			
28	C		63.2	120.98	121.41			
29	C	x	172.7	5.33	7.81			
30	C		13.9	172.31	172.20			
31	C	x	121.7	56.40	56.39			
32	C	x	136.5	42.56	42.12			
33	C		59.6	129.89	135.05			
34	C		53	132.96	133.61			
35								
36	H		1.96	29.77	29.54			
37	H		2.72	29.12	30.39			
38	H		1.94	29.74	29.76			
39	H		2.21	29.35	29.44			
40	H		2.19	29.52	29.56			
41	H		2.55	29.12	29.15			
42	H		3.39	28.31	28.51			
43	H	x	7.27	24.11	24.09			
44	H	x	7.03	24.37	24.31			
45	H	x	7.05	24.32	24.09			
46	H	x	6.68	25.03	24.47			

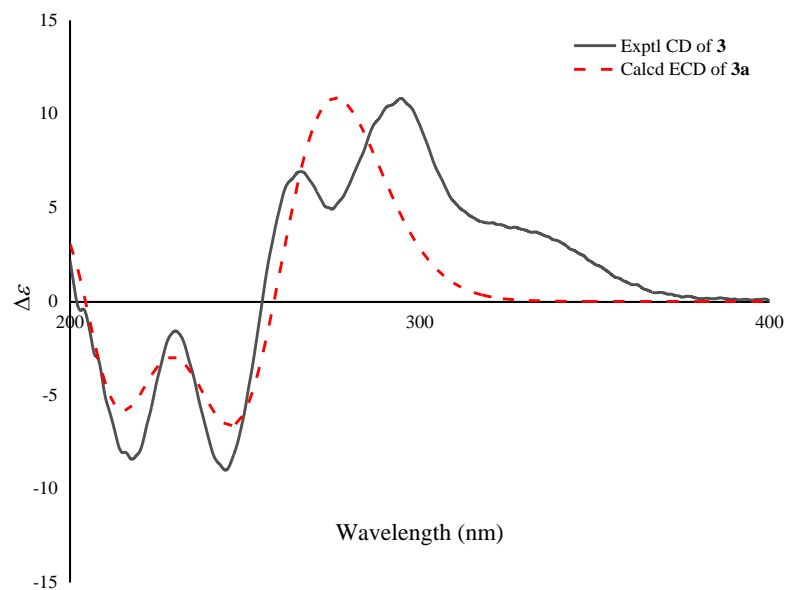
47	H		1.67	30.07	29.84			
48	H		2.25	29.68	30.11			
49	H		3.78	28.24	28.61			
50	H		5.43	26.37	27.17			
51	H		1.73	30.35	30.03			
52	H		1.73	30.47	30.13			
53	H		1.73	30.44	30.18			
54	H	x	4.86	25.95	25.80			
55	H		1.6	29.93	28.46			
56	H		2.68	28.85	28.96			
57	H		3.84	28.08	28.20			
58	H		3.84	27.89	27.88			
59	H		3.84	27.91	28.19			

	A	B	C	D	E	F	G	H
1	<b>Functional</b>		<b>Solvent?</b>		<b>Basis Set</b>		<b>Type of Data</b>	
2	mPW1PW91		PCM		6-311+G(d, p)		Shielding Tensors	
3								
4			Isomer 1	Isomer 2	Isomer 3	Isomer 4	Isomer 5	Isomer 6
5	sDP4+ (H data)		100.00%		0.00%	-	-	-
6	sDP4+ (C data)		100.00%		0.00%	-	-	-
7	sDP4+ (all data)		100.00%		0.00%	-	-	-
8	uDP4+ (H data)		100.00%		0.00%	-	-	-
9	uDP4+ (C data)		99.98%		0.02%	-	-	-
10	uDP4+ (all data)		100.00%		0.00%	-	-	-
11	DP4+ (H data)		100.00%		0.00%	-	-	-
12	DP4+ (C data)		100.00%		0.00%	-	-	-
13	DP4+ (all data)		100.00%		0.00%	-	-	-

## S2.2. Computational details for compound 3 (ECD)

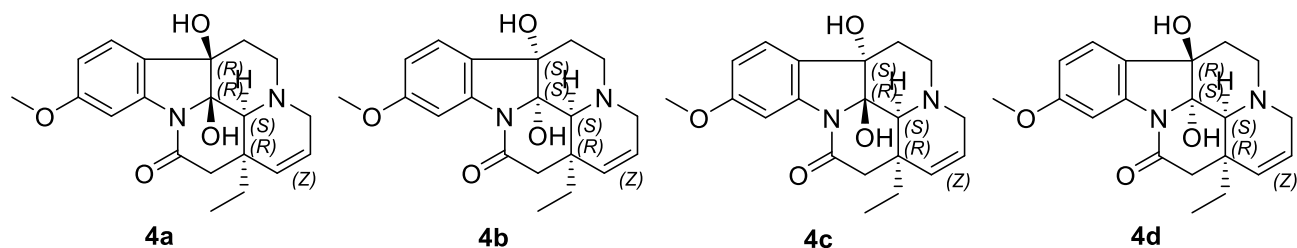
Conformation search based on molecular mechanics with MMFF force fields were performed for **3a** gave 2 stable conformers with populations higher than 1%. All these conformers were further optimized by the density functional theory method at the B3LYP/6-31G(d) level by Gaussian 16 program package with g09 default keyword. The ECD were calculated using density functional theory (TDDFT) at B3LYP/6-31+G(d,p) level in methanol with IEFPCM model. The calculated ECD curves were all generated using SpecDis 1.71 program package ( $\sigma = 0.30$  eV, and UV shift -22 nm) and the calculated ECD data of all conformers were Boltzmann averaged by Gibbs free energy.

**Figure S8.** Comparison of the calculated ECD spectra with the experimental spectrum of **3** in methanol with PCM model.



## Section S3. Computational details for 4

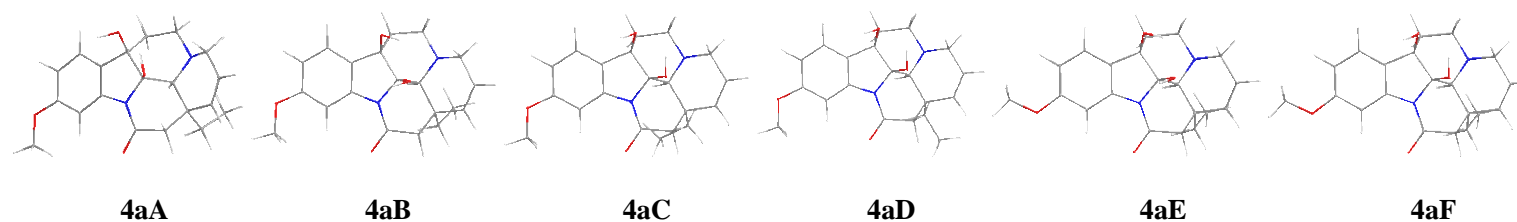
### S3.1. Computational details for compound 4 (NMR)



Conformation search based on molecular mechanics with MMFF force fields were performed for **4a**, **4b**, **4c**, **4d** gave 6, 4, 5, and 4 stable conformers with populations higher than 1%, respectively. All these conformers were further optimized by the density functional theory method at the B3LYP/6-31G(d) level by Gaussian 16 program package with g09 default keyword. Gauge Independent Atomic Orbital (GIAO) calculations of their  $^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts using density functional theory (DFT) at the mPW1PW91/6-311+G(d,p) level with the PCM model in methanol. The calculated NMR data of these conformers were averaged according to the Boltzmann distribution theory and their relative Gibbs free energy. The  $^1\text{H}$  and  $^{13}\text{C}$  NMR chemical shifts for TMS were also calculated by the same procedures and used as the reference. After calculation, the experimental and calculated data were evaluated by linear correlation coefficients ( $R^2$ ) and the improved probability DP4+ method.

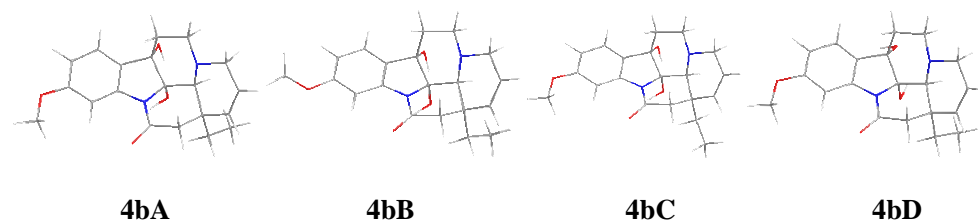
**Table S12.** Energy analysis for optimized geometries of dominant conformers **4aA–4aF** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>4aA</b>	-1186.528244	-1186.506097	-1186.505153	-1186.577142	0	0	59.12%
<b>4aB</b>	-1186.525993	-1186.503870	-1186.502926	-1186.574886	0.002256	1.415661	5.41%
<b>4aC</b>	-1186.526787	-1186.504603	-1186.503659	-1186.575685	0.001457	0.914281	12.62%
<b>4aD</b>	-1186.526534	-1186.504343	-1186.503399	-1186.575567	0.001575	0.988327	11.14%
<b>4aE</b>	-1186.524416	-1186.502260	-1186.501316	-1186.573463	0.003679	2.308607	1.20%
<b>4aF</b>	-1186.526495	-1186.504312	-1186.503368	-1186.575513	0.001629	1.022213	10.52%

**Figure S9.** Main conformers of **4a** in NMR and ECD calculations.**Table S13.** Energy analysis for optimized geometries of dominant conformers **4bA–4bD** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>4bA</b>	-1186.534528	-1186.512406	-1186.511462	-1186.583146	0	0	66.88%
<b>4bB</b>	-1186.533465	-1186.511333	-1186.510389	-1186.582148	0.000998	0.626254	23.23%
<b>4bC</b>	-1186.531817	-1186.509571	-1186.508627	-1186.581101	0.002045	1.283257	7.66%
<b>4bD</b>	-1186.531036	-1186.508745	-1186.507801	-1186.579939	0.003207	2.012423	2.23%

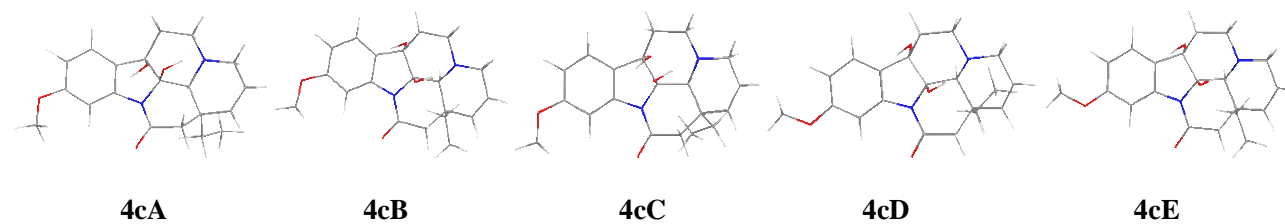
**Figure S10.** Main conformers of **4b** in NMR and ECD calculations.



**Table S14.** Energy analysis for optimized geometries of dominant conformers **4cA–4cE** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>4cA</b>	-1186.519276	-1186.497403	-1186.496459	-1186.567421	0	0	35.86%
<b>4cB</b>	-1186.518530	-1186.496628	-1186.495684	-1186.566970	0.000451	0.283007	22.23%
<b>4cC</b>	-1186.518832	-1186.496958	-1186.496013	-1186.566969	0.000452	0.283634	22.21%
<b>4cD</b>	-1186.518146	-1186.496216	-1186.495271	-1186.566439	0.000982	0.616214	12.66%
<b>4cE</b>	-1186.517382	-1186.495455	-1186.494511	-1186.565884	0.001537	0.964482	7.03%

**Figure S11.** Main conformers of **4c** in NMR and ECD calculations.

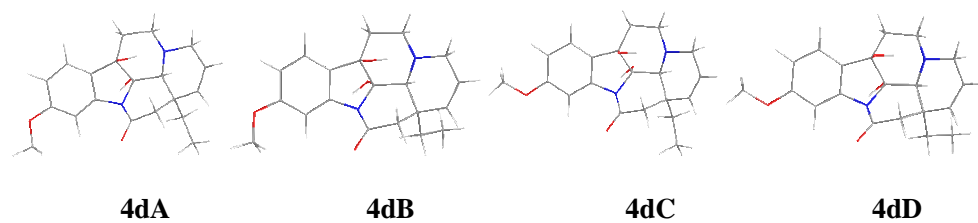




**Table S15.** Energy analysis for optimized geometries of dominant conformers **4dA–4dD** at B3LYP/6-31G(d) level in the gas phase

Species	$E'=E+ZPE$	$E$	$H$	$G$	$\Delta G$	$\Delta E(\text{kcal/mol})$	$PE\%$
<b>4dA</b>	-1186.510799	-1186.488883	-1186.487939	-1186.559356	0.000782	0.490712	22.61%
<b>4dB</b>	-1186.511231	-1186.489191	-1186.488247	-1186.560138	0	0	51.78%
<b>4dC</b>	-1186.509823	-1186.487913	-1186.486969	-1186.558397	0.001741	1.092494	8.18%
<b>4dD</b>	-1186.510254	-1186.488225	-1186.487281	-1186.559111	0.001027	0.644452	17.44%

**Figure S12.** Main conformers of **4d** in NMR and ECD calculations.



**Table S16.** Calculated  $^{13}\text{C}$  NMR results for **4a**

No.	<b>4aA</b>	<b>4aB</b>	<b>4aC</b>	<b>4aD</b>	<b>4aE</b>	<b>4aF</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	89.0177	89.5332	89.9805	89.3992	89.5428	88.7698	98.99	93.1	95.1	-2.0
3	133.6251	133.5503	133.4514	132.9541	133.5633	133.6545	54.65	55.7	52.7	3.0
5	137.6755	138.2344	137.2596	137.4291	137.9413	137.3635	50.58	50.3	48.8	1.5
6	152.1772	152.2745	152.2488	152.0981	152.5340	152.2037	35.99	37.5	34.9	2.6
7	104.3283	104.3076	104.3050	103.9541	104.5018	104.0856	83.92	79.3	80.7	-1.4
8	58.4256	59.4498	57.8696	58.0508	57.8628	56.1965	130.05	130.9	124.8	6.1
9	56.9562	55.9223	57.2173	57.4222	55.8074	56.6506	131.24	124.0	126.0	-2.0
10	69.2695	69.7522	69.3952	68.8567	78.2474	78.1369	117.88	111.3	113.2	-1.9
11	18.0590	18.7775	18.2972	18.1358	18.3554	17.5278	170.10	161.6	163.2	-1.6
12	86.1268	86.2579	85.6547	85.8326	79.0918	78.6414	103.01	106.1	99.0	7.1
13	37.2620	37.8712	37.8784	37.6444	38.9139	38.2124	150.65	141.2	144.5	-3.3
14	55.2473	55.5883	58.6018	56.9179	55.6711	55.3142	132.29	125.2	127.0	-1.8
15	49.9002	50.3988	47.2530	50.8275	50.6877	50.2417	138.44	135.4	132.9	2.5
16	9.5194	10.1536	9.3113	9.8039	10.3477	9.8263	178.58	170.2	171.3	-1.1
17	136.6527	136.7124	140.2162	143.0717	136.6333	136.5770	50.37	45.1	48.6	-3.5
18	176.0864	176.1143	177.0339	177.4802	176.4510	176.3818	11.78	9.5	11.7	-2.2
19	147.6877	147.4271	148.4931	148.0706	147.0265	147.3145	40.41	34.0	39.1	-5.1
20	144.9104	145.3587	145.8976	145.9838	145.3559	144.9426	42.99	40.4	41.6	-1.2
21	126.3261	126.5133	129.8561	122.4294	126.1545	125.8239	61.89	61.8	59.6	2.2
-OCH <sub>3</sub>	132.3480	132.3539	132.1058	132.5203	131.7331	131.7411	55.92	56.0	53.9	2.1
Population	59.12%	5.41%	12.62%	11.14%	1.20%	10.52%			RMSD	3.1

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

**Table S17.** Calculated  $^{13}\text{C}$  NMR results for **4b**

No.	<b>4bA</b>	<b>4bB</b>	<b>4bC</b>	<b>4bD</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	89.2092	89.1430	89.3242	90.9718	98.94	93.1	94.8	-1.7
3	130.8292	130.8784	130.6541	130.9717	57.35	55.7	54.8	0.9
5	137.2352	136.7294	136.9658	136.7054	51.10	50.3	48.8	1.5
6	149.5743	149.6063	149.8007	148.1514	38.61	37.5	36.9	0.6
7	104.6621	104.7230	104.3950	102.3634	83.58	79.3	80.0	-0.7
8	55.1282	53.4416	54.9969	54.8533	133.46	130.9	127.9	3.0
9	57.1346	56.9793	57.0098	59.8330	131.03	124.0	125.6	-1.6
10	68.4434	77.4543	68.4361	69.3366	117.63	111.3	112.7	-1.4
11	19.8935	20.1358	19.8556	19.5813	168.24	161.6	161.3	0.3
12	82.5999	75.9516	82.4166	82.7730	107.14	106.1	102.6	3.5
13	41.8657	42.3290	42.0452	38.6640	146.27	141.2	140.2	1.0
14	52.8505	52.7511	53.7829	53.2440	135.27	125.2	129.7	-4.5
15	45.3875	45.3915	49.2412	45.3368	142.50	135.4	136.6	-1.2
16	11.7760	11.7412	11.6758	11.8231	176.42	170.2	169.2	1.0
17	139.0748	139.2158	144.5157	139.2289	48.65	45.1	46.5	-1.4
18	176.5640	176.3911	179.3350	176.5019	11.45	9.5	10.8	-1.3
19	151.7688	151.6772	155.5089	152.2258	36.14	34.0	34.5	-0.5
20	142.6134	142.9094	143.2746	143.0104	45.44	40.4	43.4	-3.0
21	127.9784	128.2626	122.0192	123.4066	60.70	61.8	58.1	3.7
-OCH <sub>3</sub>	131.5552	131.6037	131.2107	131.6207	56.64	56.0	54.2	1.8
Population	66.88%	23.23%	7.66%	2.23%			RMSD	2.1

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

**Table S18.** Calculated  $^{13}\text{C}$  NMR results for **4c**

No.	<b>4cA</b>	<b>4cB</b>	<b>4cC</b>	<b>4cD</b>	<b>4cE</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	92.4096	92.5280	91.8986	92.0786	92.4535	95.90	93.1	91.9	1.2
3	136.6348	136.4538	137.3523	136.4798	136.5256	51.45	55.7	50.1	5.6
5	135.1209	134.6970	134.9328	135.1732	134.7981	53.21	50.3	51.8	-1.5
6	154.1896	154.3355	154.5519	154.1612	154.3024	33.87	37.5	33.6	3.9
7	105.0537	105.3532	104.8041	105.0590	105.1196	83.11	79.3	79.9	-0.6
8	55.5867	55.6083	55.7197	54.0527	54.1316	132.86	130.9	126.6	4.3
9	55.9566	55.9357	56.1577	55.6819	55.6334	132.24	124.0	126.1	-2.1
10	69.9172	69.8629	69.9018	78.6156	78.5655	116.57	111.3	111.3	0.0
11	18.6166	18.7137	18.6904	18.6499	18.9610	169.50	161.6	161.1	0.5
12	84.9383	85.0306	84.7737	77.2812	77.8188	104.73	106.1	100.2	5.9
13	35.0535	35.1127	35.3381	35.7557	35.8236	152.91	141.2	145.5	-4.3
14	52.0986	53.9120	55.0934	52.2241	53.9534	134.87	125.2	128.5	-3.3
15	48.3921	47.8674	44.3066	48.7933	47.8510	140.80	135.4	134.1	1.3
16	7.8659	8.4225	7.5004	8.1377	8.6616	180.18	170.2	171.1	-0.9
17	136.5053	142.8966	139.7183	136.1307	142.6987	49.15	45.1	48.0	-2.9
18	178.0841	177.9557	177.2136	178.0253	177.7266	10.35	9.5	11.5	-2.0
19	149.3819	147.5538	145.2269	149.1272	147.0171	40.33	34.0	39.7	-5.7
20	143.9726	145.0842	144.4794	143.9453	144.5618	43.81	40.4	43.0	-2.6
21	129.0168	122.6914	127.0535	128.8618	122.2869	61.50	61.8	59.6	2.2
-OCH <sub>3</sub>	131.5043	131.8302	131.5289	131.6014	131.7760	56.57	56.0	54.9	1.1
Population	35.86%	22.23%	22.21%	12.66%	7.03%			RMSD	3.1

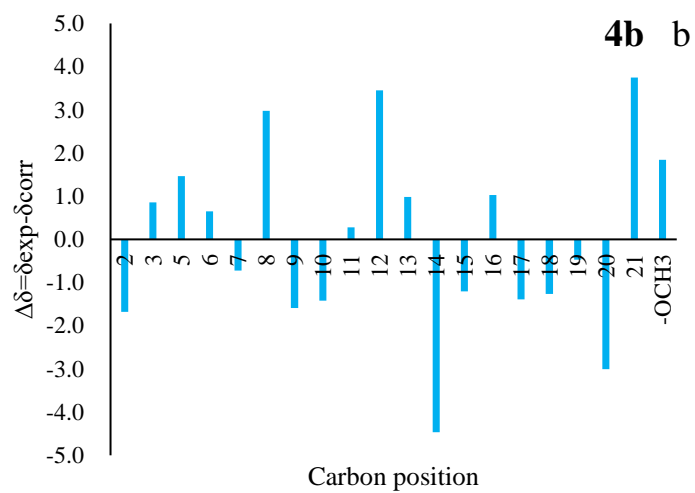
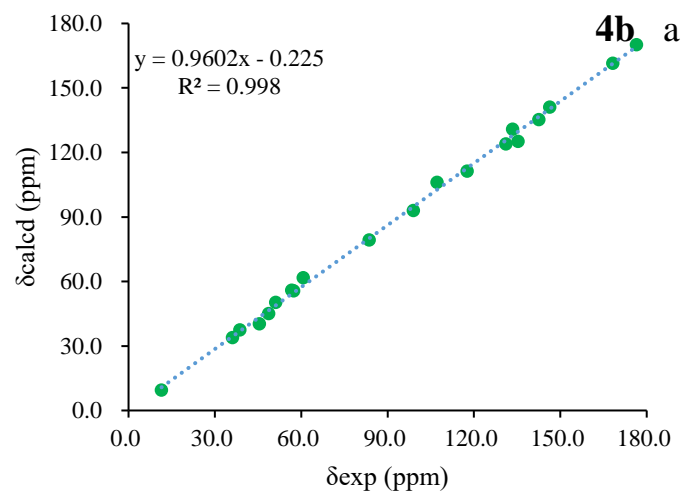
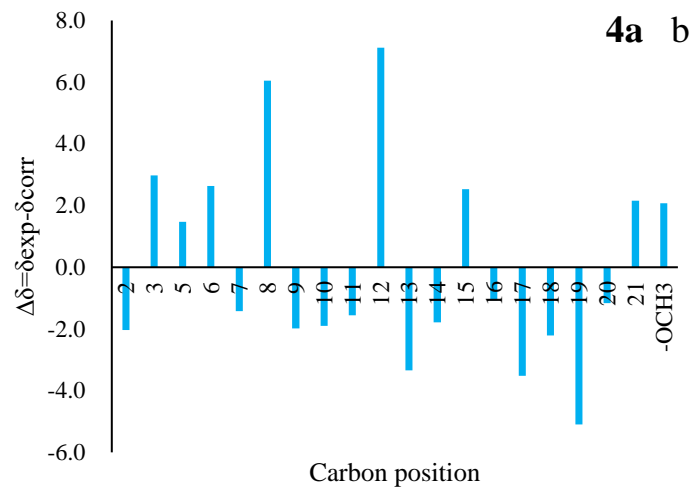
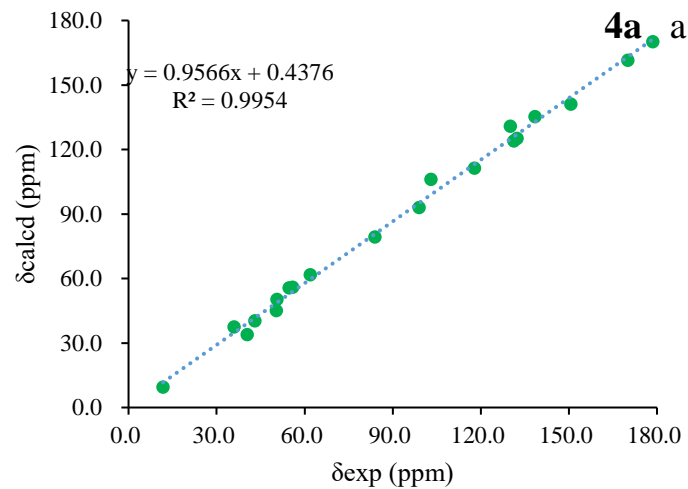
<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

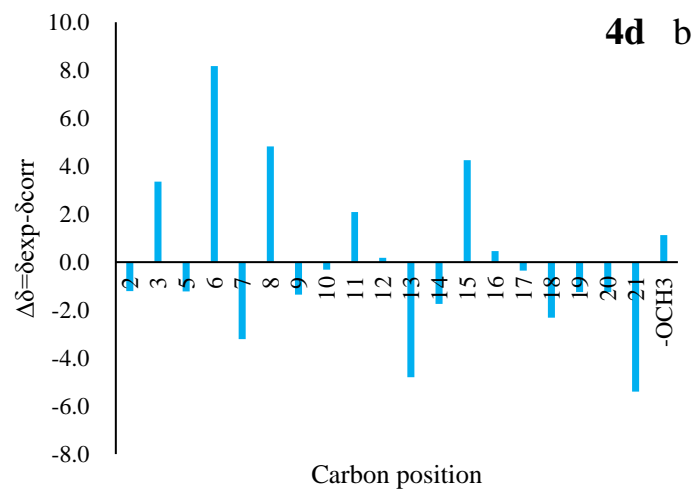
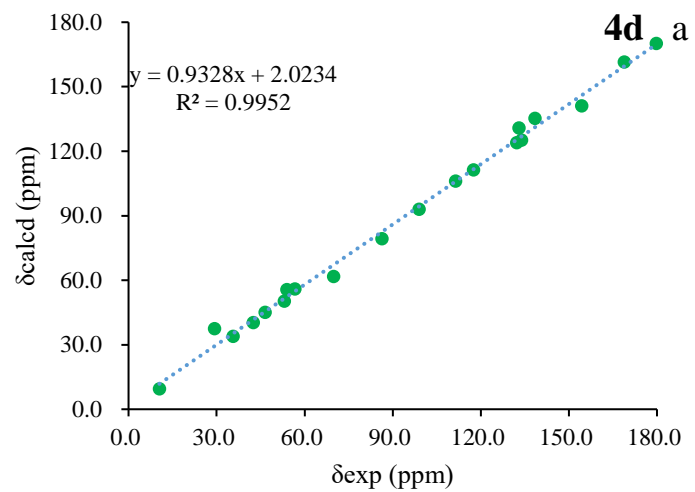
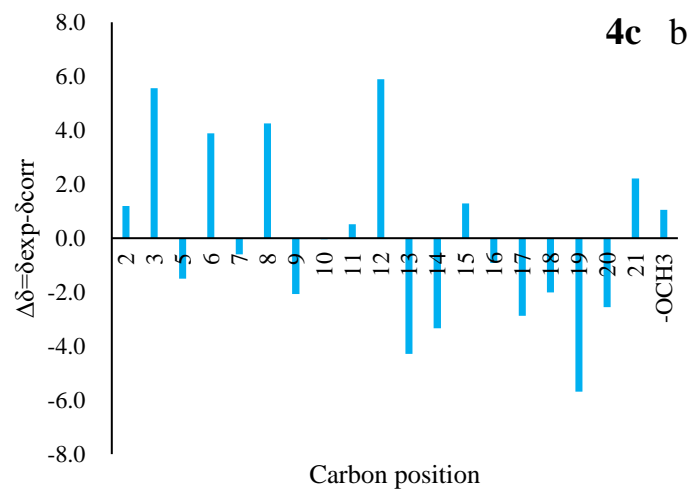
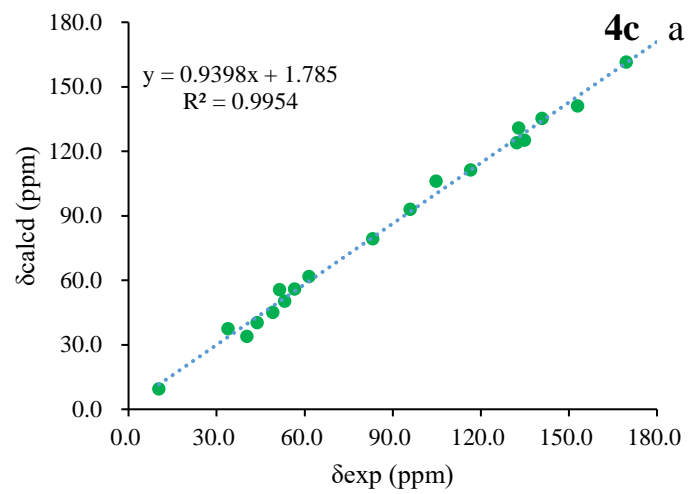
**Table S19.** Calculated  $^{13}\text{C}$  NMR results for **4d**

No.	<b>4dA</b>	<b>4dB</b>	<b>4dC</b>	<b>4dD</b>	$\delta_{\text{Calcd}}^a$	$\delta_{\text{Exp}}$	$\delta_{\text{Corr}}^b$	Relative errors <sup>c</sup>
2	89.6777	89.0505	89.7498	89.0167	98.94	93.1	94.3	-1.2
3	134.1612	134.3055	133.9316	134.2523	53.95	55.7	52.3	3.4
5	134.9985	135.1200	135.1446	135.2425	53.07	50.3	51.5	-1.2
6	158.9561	158.9986	158.6318	158.6779	29.28	37.5	29.3	8.2
7	101.8580	101.9320	101.8417	101.8671	86.28	79.3	82.5	-3.2
8	55.6014	55.4457	54.4665	54.2527	132.99	130.9	126.1	4.8
9	55.9095	55.9498	56.0335	56.0471	132.22	124.0	125.4	-1.4
10	68.4665	68.4436	77.1029	77.2751	117.48	111.3	111.6	-0.3
11	19.3492	19.4181	19.2517	19.1955	168.83	161.6	159.5	2.1
12	79.0999	78.7711	71.0959	70.6026	111.39	106.1	105.9	0.2
13	33.6100	33.7169	34.3396	34.2427	154.35	141.2	146.0	-4.8
14	55.5220	53.6686	55.6507	53.7456	133.92	125.2	126.9	-1.7
15	50.9389	49.2935	50.7199	49.0984	138.43	135.4	131.2	4.2
16	8.0095	8.4351	8.3766	8.7156	179.80	170.2	169.7	0.5
17	145.3910	139.9229	145.3395	140.0015	46.57	45.1	45.5	-0.4
18	178.5562	177.3207	178.1597	177.3726	10.50	9.5	11.8	-2.3
19	152.0918	152.8630	151.3424	152.8356	35.62	34.0	35.3	-1.3
20	146.1676	145.5353	145.8422	145.4948	42.48	40.4	41.7	-1.3
21	113.8386	120.2023	114.0531	120.5280	69.86	61.8	67.2	-5.4
-OCH <sub>3</sub>	131.6204	131.4389	131.6028	131.5975	56.66	56.0	54.9	1.1
Population	22.61%	51.78%	8.18%	17.44%			RMSD	3.2

<sup>a</sup>Weighted average from the calculated shifts; <sup>b</sup>Obtained by linear fit  $\delta_{\text{exp}}$  versus  $\delta_{\text{calcd}}$ ; <sup>c</sup> $\Delta\delta = \delta_{\text{exp}} - \delta_{\text{corr}}$ .

**Figure S13.**  $^{13}\text{C}$  NMR calculation results of four possible isomers of **4**. (a) Linear correlation plots of predicted versus experimental  $^{13}\text{C}$  NMR chemical shifts. (b) Relative errors between the predicted  $^{13}\text{C}$  NMR chemical shifts of two potential structures and recorded  $^{13}\text{C}$  NMR data.





**Table S20.** DP4+ analysis results of **4a** (Isomer 1), **4b** (Isomer 2), **4c** (Isomer 3), and **4d** (Isomer 4)

	A	B	C	D	E	F	G	H
1	Functional		Solvent?		Basis Set		Type of Data	
2	mPW1PW91		PCM		6-311+G(d,p)		Shielding Tensors	
3								
12			DP4+	0.00%	100.00%	0.00%	0.00%	-
14	Nuclei	sp2?	xperimental	Isomer 1	Isomer 2	Isomer 3	Isomer 4	Isomer 5
15	C		93.1	89.19	89.24	92.28	89.24	
16	C		55.7	133.53	130.83	136.73	134.23	
17	C		50.3	137.60	137.09	134.97	135.12	
18	C		37.5	152.19	149.57	154.31	158.90	
19	C		79.3	104.26	104.60	105.07	101.90	
20	C	x	130.9	58.13	54.72	55.32	55.19	
21	C	x	124	56.94	57.15	55.94	55.96	
22	C	x	111.3	70.31	70.56	71.61	70.70	
23	C	x	161.6	18.08	19.94	18.68	19.35	
24	C	x	106.1	85.17	81.05	83.45	76.79	
25	C	x	141.2	37.54	41.92	35.27	33.84	
26	C	x	125.2	55.89	52.91	53.31	54.26	
27	C	x	135.4	49.74	45.68	47.38	49.75	
28	C	x	170.2	9.60	11.76	8.00	8.38	
29	C		45.1	137.81	139.53	139.03	141.62	
30	C		9.5	176.40	176.73	177.83	177.68	
31	C		34	147.77	152.04	147.85	152.56	
32	C		40.4	145.19	142.74	144.37	145.70	
33	C		61.8	126.29	127.49	126.68	118.32	
34	C		56	132.27	131.54	131.61	131.52	
35								
36	H		2.88	28.42	28.60	28.58	28.73	
37	H		3.29	28.41	28.40	28.57	28.37	
38	H		2.59	28.87	29.12	28.95	28.66	
39	H		2.7	28.22	28.96	28.78	29.03	
40	H		1.6	29.39	30.29	29.39	29.49	
41	H		2.01	29.39	29.74	29.61	29.36	
42	H	x	7.26	24.10	24.11	24.06	24.04	
43	H	x	6.71	24.74	24.80	24.77	24.68	
44	H	x	7.65	23.80	23.68	23.86	24.26	
45	H	x	5.77	25.52	25.57	25.32	25.49	
46	H	x	5.47	25.96	25.92	25.85	25.87	



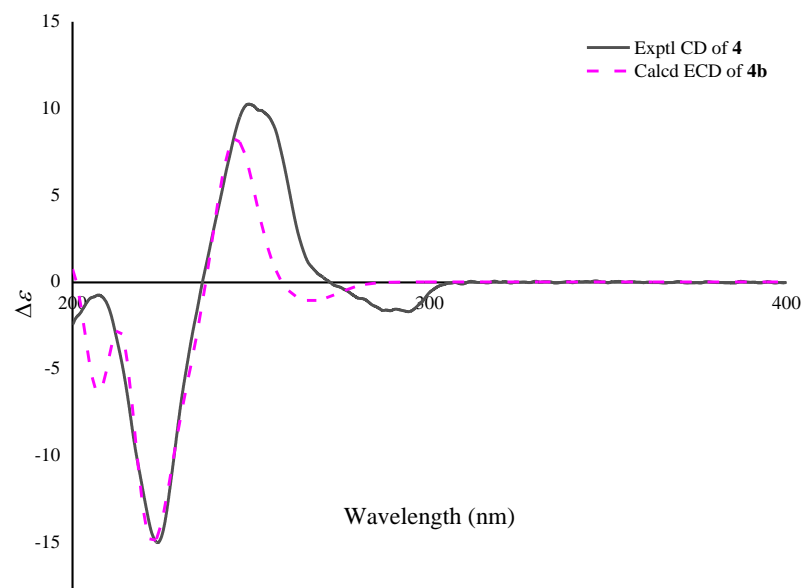
47	H		2.35	29.49	29.51	29.45	29.49	
48	H		2.91	28.64	28.73	28.65	29.07	
49	H		0.96	30.94	30.90	30.85	30.84	
50	H		0.96	30.86	30.84	30.81	30.81	
51	H		0.96	30.89	30.55	30.83	30.71	
52	H		1.42	30.34	30.26	30.31	30.11	
53	H		2.28	30.32	29.62	30.20	30.08	
54	H		2.79	28.71	28.94	28.17	28.81	
55	H		3.79	27.67	27.68	27.66	27.67	
56	H		3.79	28.05	28.02	28.01	27.99	
57	H		3.79	28.03	28.03	28.03	27.99	

	A	B	C	D	E	F	G	H			
1	<b>Functional</b>		<b>Solvent?</b>		<b>Basis Set</b>		<b>Type of Data</b>				
2	mPW1PW91		PCM		6-311+G(d, p)		Shielding Tensors				
3											
4			Isomer 1	Isomer 2	Isomer 3	Isomer 4	Isomer 5	Isomer 6			
5	sDP4+ (H data)		0.00%		100.00%		0.00%		0.00%	-	-
6	sDP4+ (C data)		0.00%		100.00%		0.00%		0.00%	-	-
7	sDP4+ (all data)		0.00%		100.00%		0.00%		0.00%	-	-
8	uDP4+ (H data)		0.00%		100.00%		0.00%		0.00%	-	-
9	uDP4+ (C data)		0.00%		100.00%		0.00%		0.00%	-	-
10	uDP4+ (all data)		0.00%		100.00%		0.00%		0.00%	-	-
11	DP4+ (H data)		0.00%		100.00%		0.00%		0.00%	-	-
12	DP4+ (C data)		0.00%		100.00%		0.00%		0.00%	-	-
13	DP4+ (all data)		0.00%		100.00%		0.00%		0.00%	-	-

### S3.2. Computational details for compound **4** (ECD)

Conformation search based on molecular mechanics with MMFF force fields were performed for **4a**, **4b**, **4c**, **4d** gave 6, 4, 5, and 4 stable conformers with populations higher than 1%, respectively. All these conformers were further optimized by the density functional theory method at the B3LYP/6-31G(d) level by Gaussian 16 program package with g09 default keyword. The ECD were calculated using density functional theory (TDDFT) at B3LYP/6-31+G(d,p) level in methanol with IEFPCM model. The calculated ECD curves were all generated using SpecDis 1.71 program package and the calculated ECD data of all conformers were Boltzmann averaged by Gibbs free energy.

**Figure S14.** Comparison of the calculated ECD spectra with the experimental spectrum of **4** in methanol with PCM model.



## Section S4. NMR, MS, IR, and CD spectra for 1

Figure S15.  $^1\text{H}$  NMR of compound 1 in  $\text{CD}_3\text{OD}$

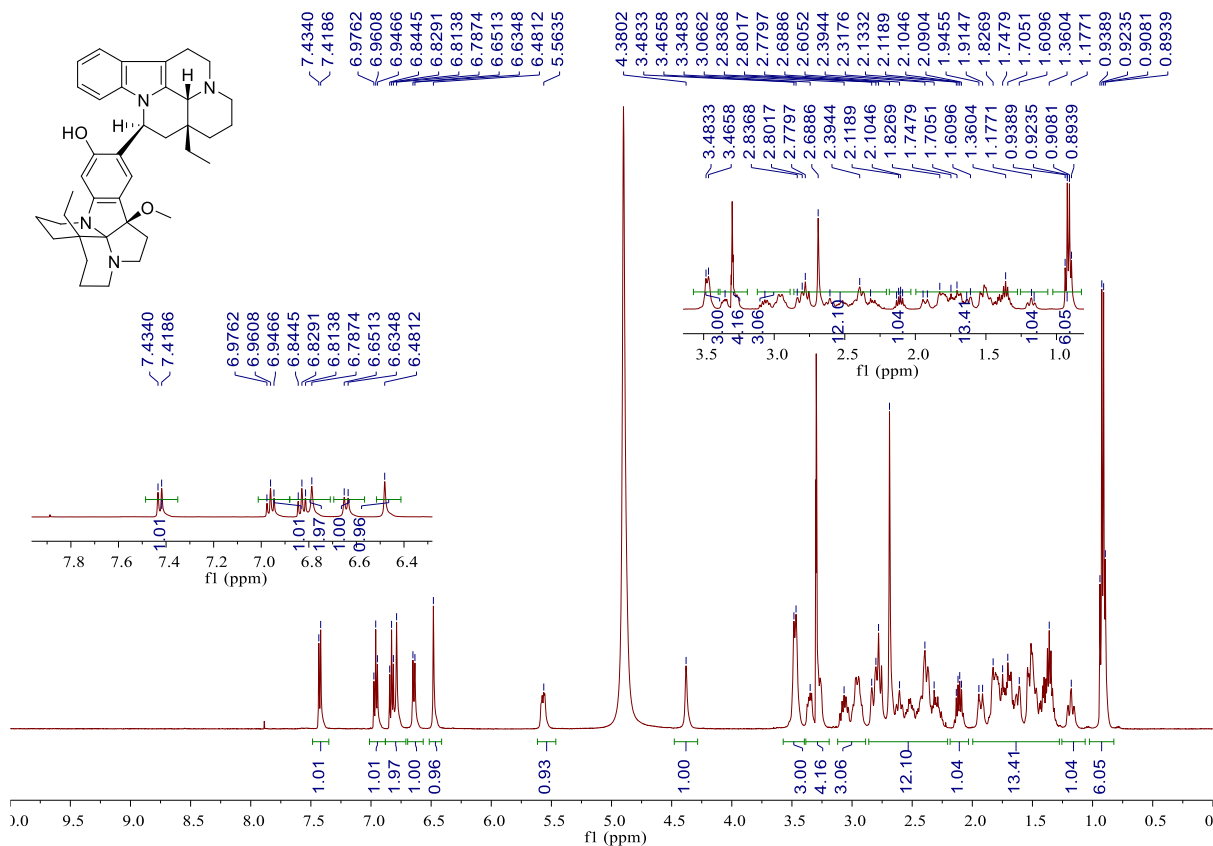


Figure S16.  $^{13}\text{C}$  NMR and DEPT of compound 1 in  $\text{CD}_3\text{OD}$

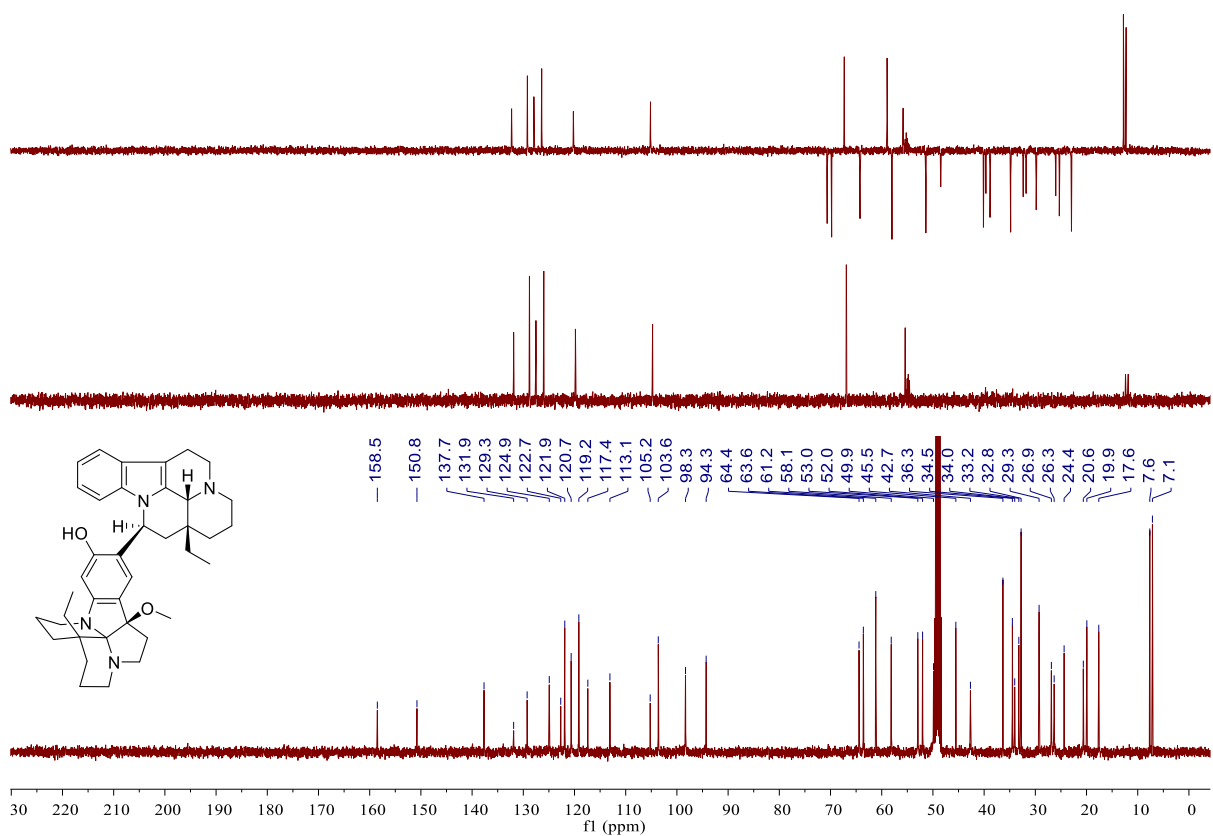


Figure S17. HSQC of compound 1 in CD<sub>3</sub>OD

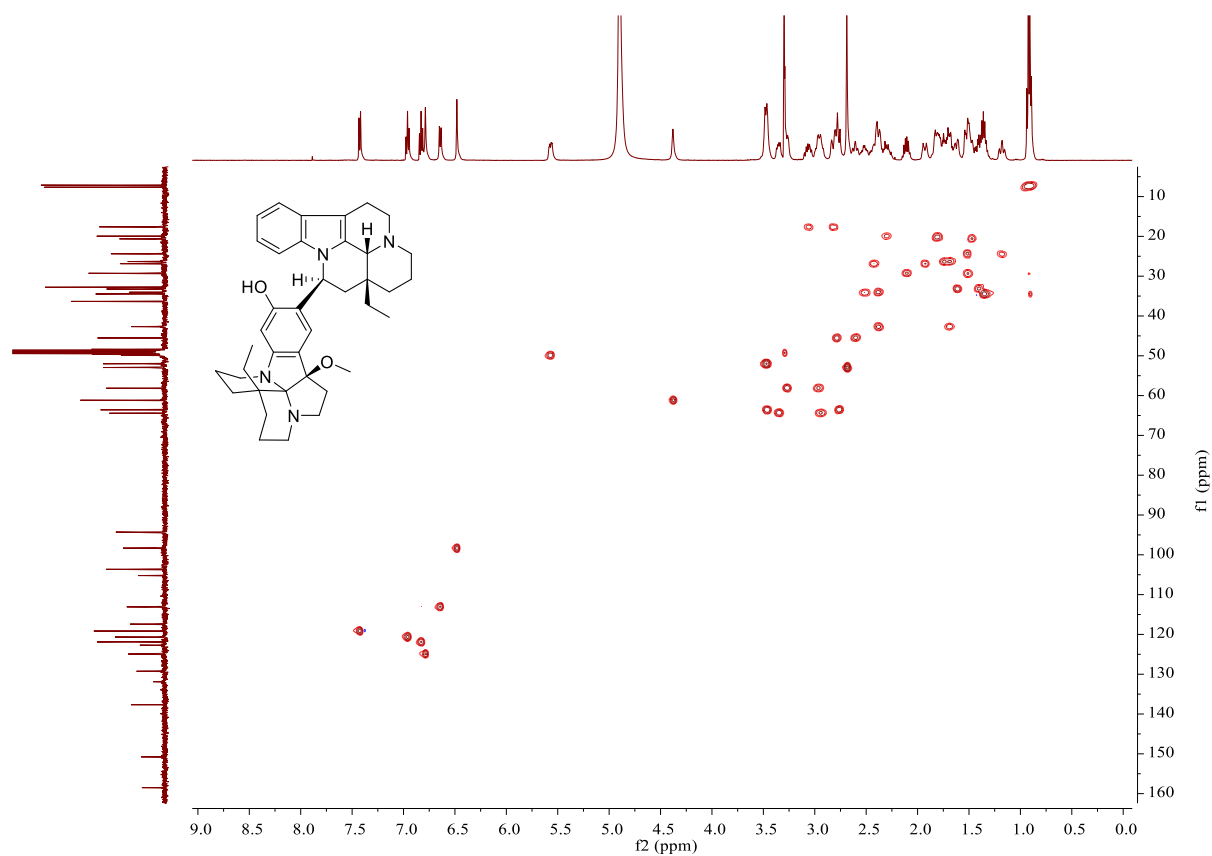
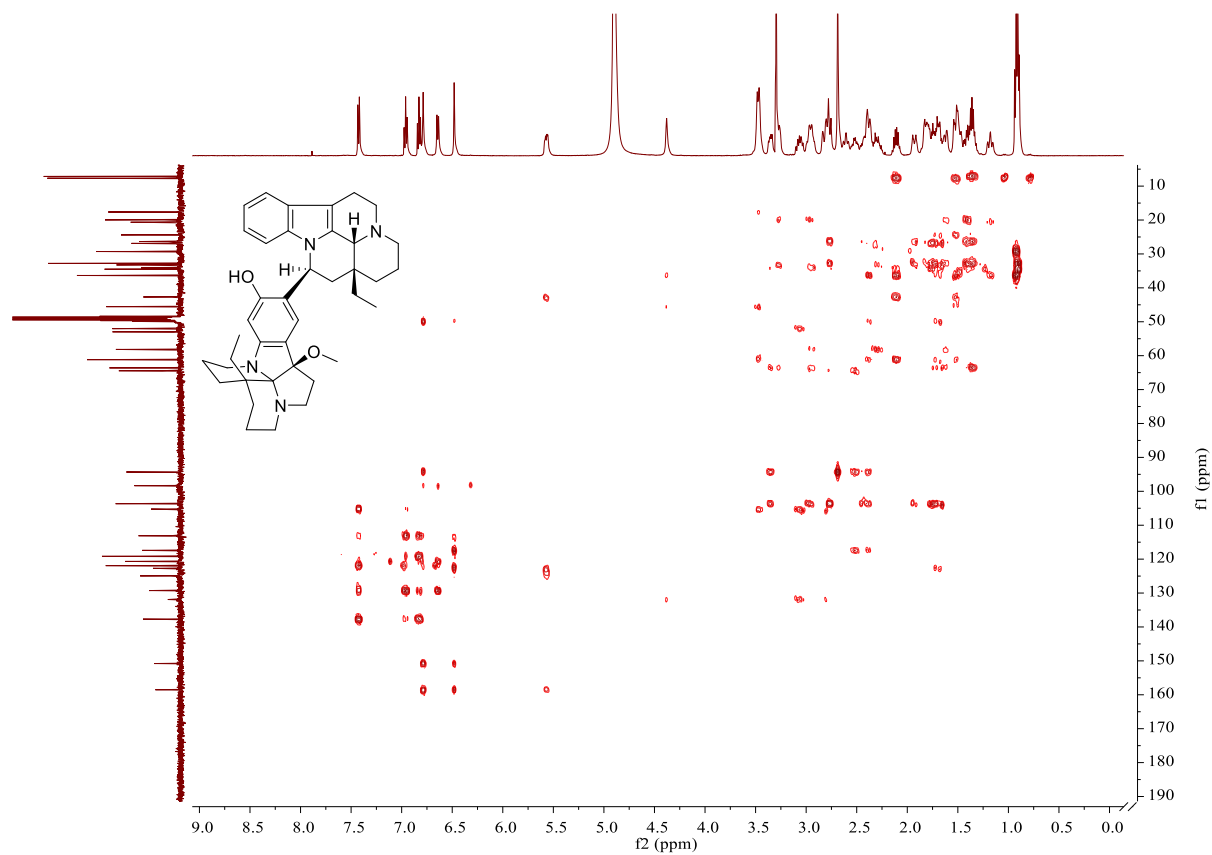
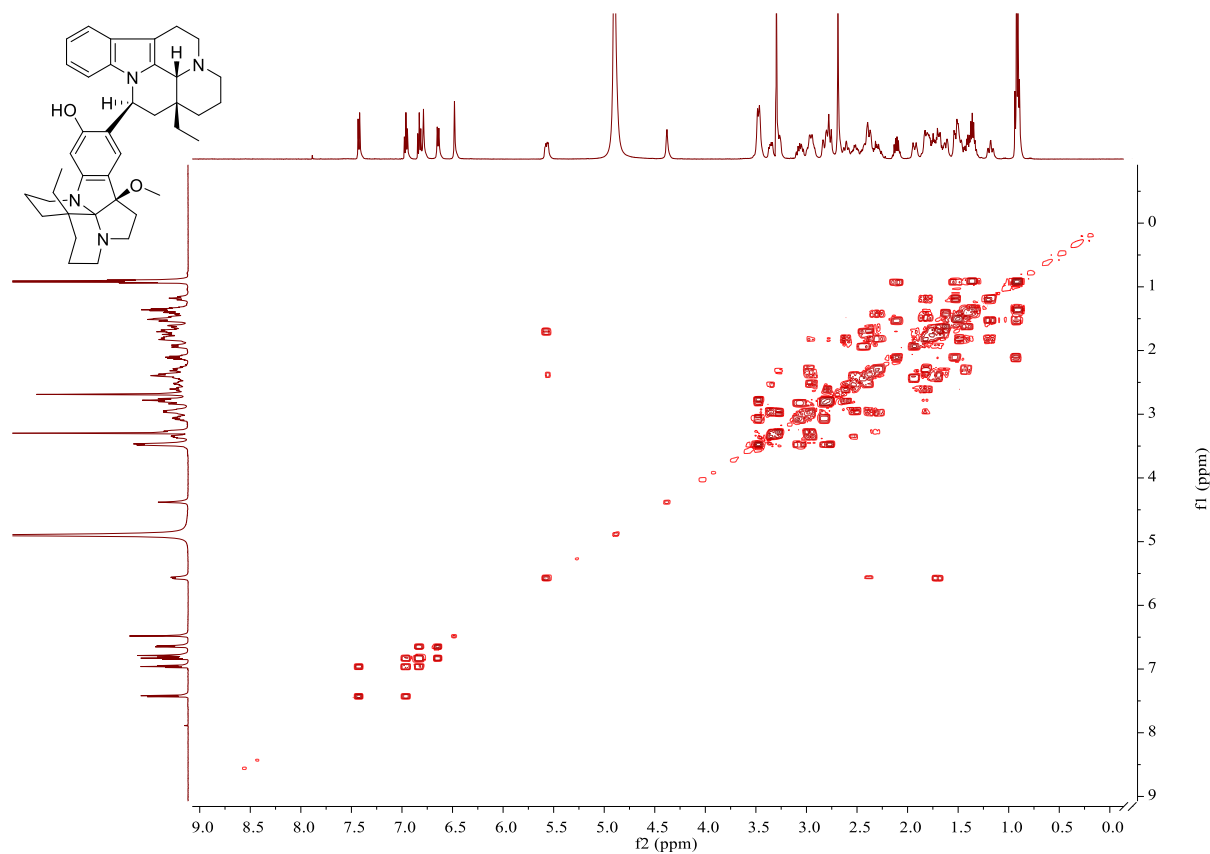


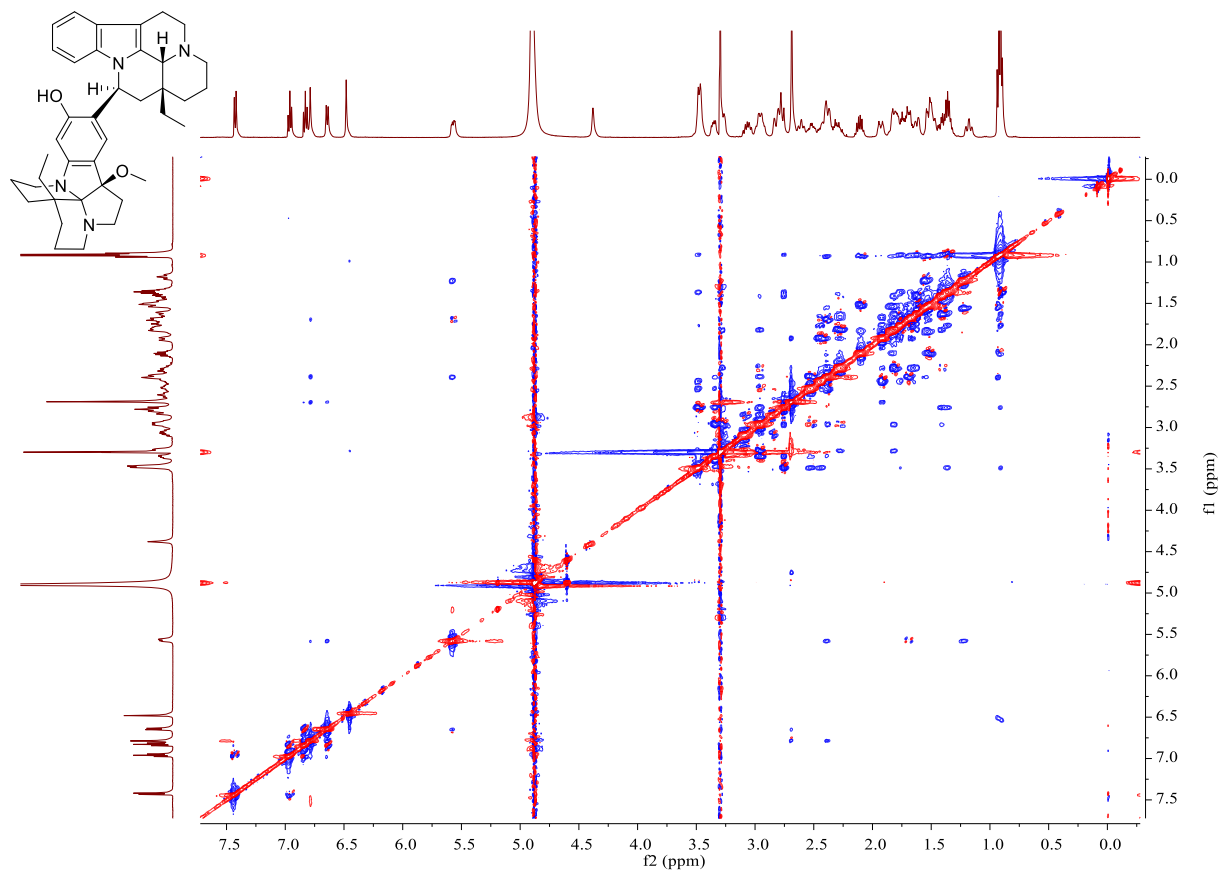
Figure S18. HMBC of compound 1 in CD<sub>3</sub>OD



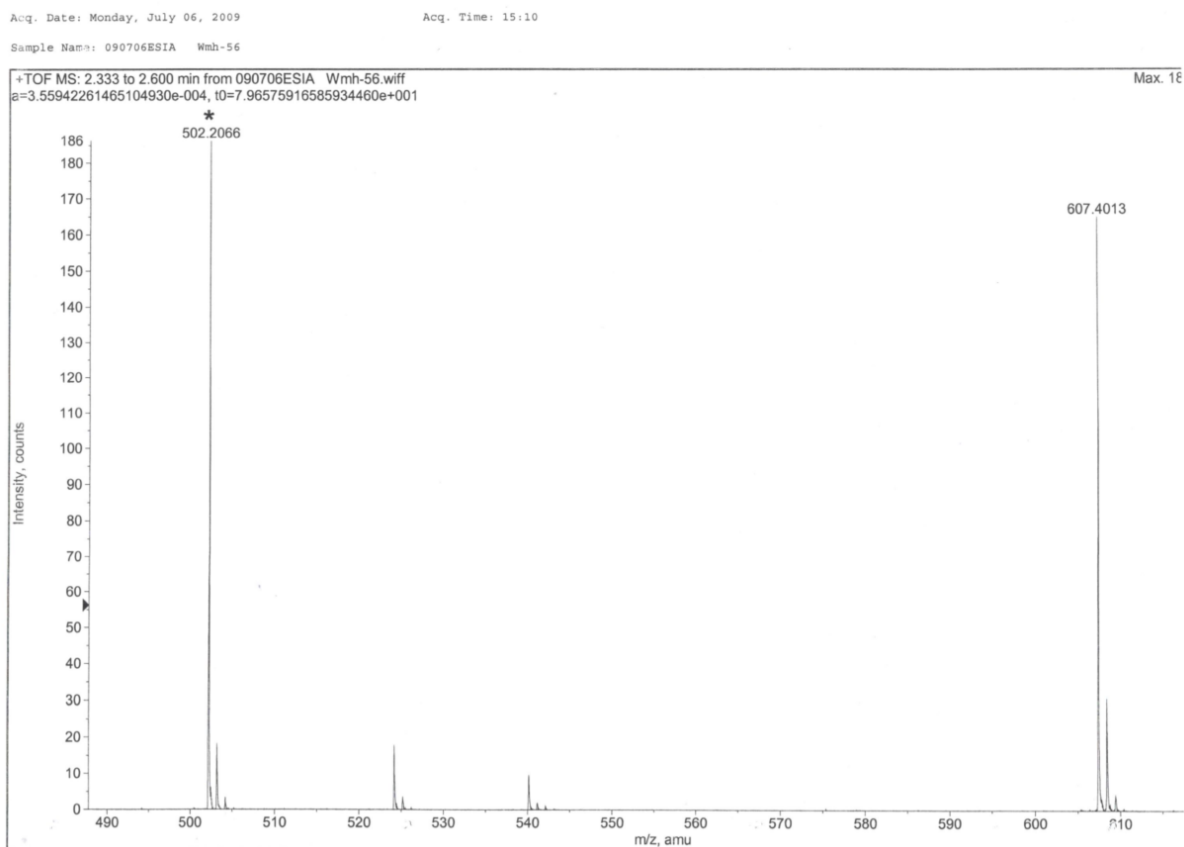
**Figure S19.**  $^1\text{H}$ - $^1\text{H}$  COSY of compound **1** in  $\text{CD}_3\text{OD}$



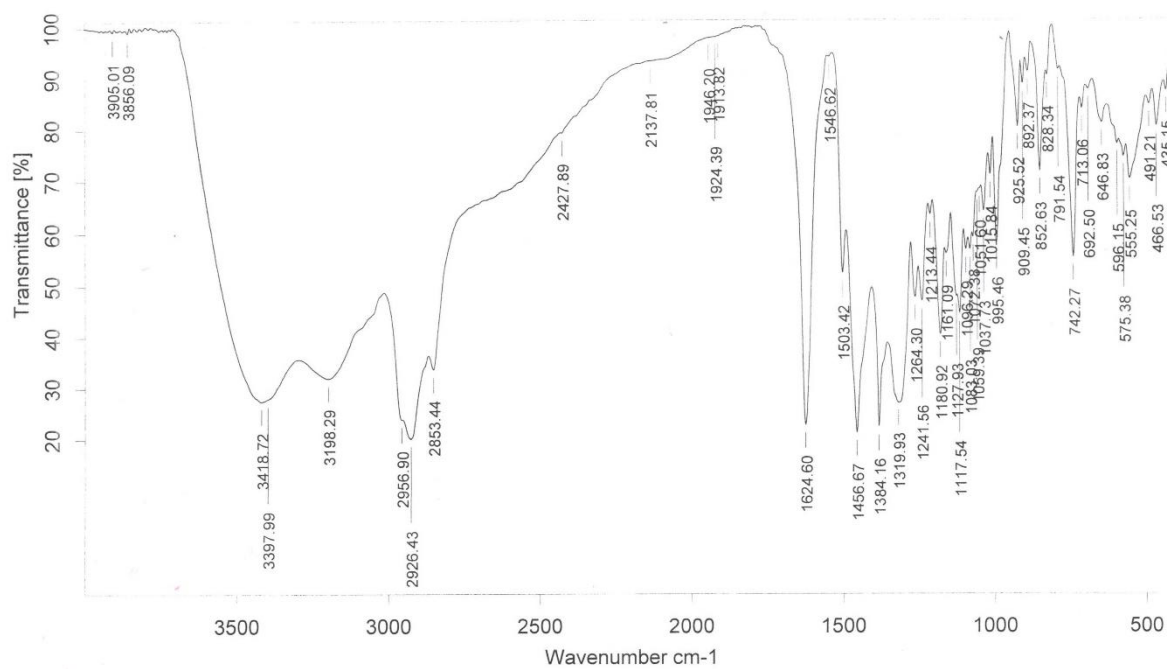
**Figure S20.** ROESY of compound **1** in  $\text{CD}_3\text{OD}$



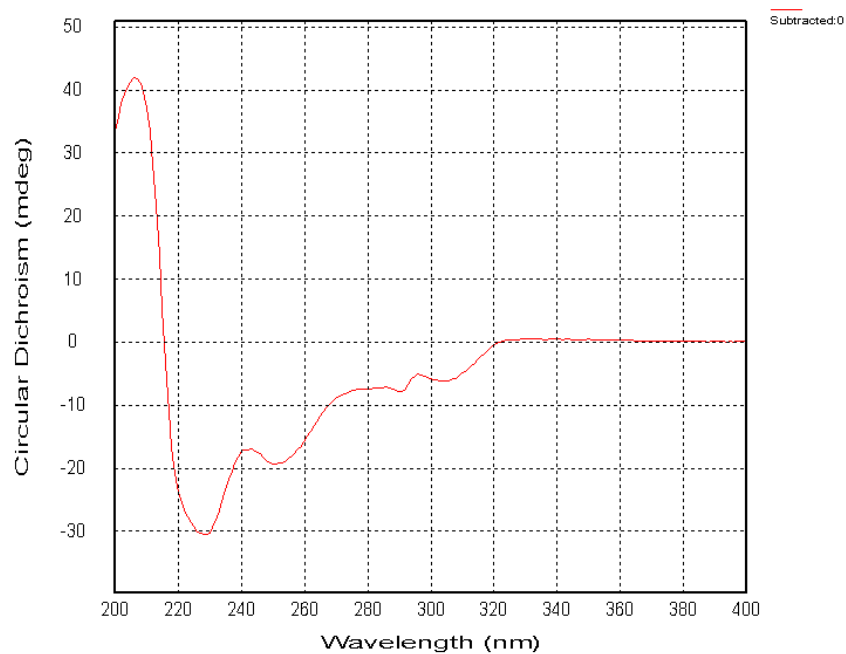
**Figure S21.** HR-ESIMS of compound **1**



**Figure S22.** IR spectra of compound **1**



**Figure S23.** CD spectra of compound **1** in CH<sub>3</sub>OH



## Section S5. NMR, MS, IR, and CD spectra for 2

Figure S24.  $^1\text{H}$  NMR of compound 2 in  $\text{CD}_3\text{OD}$

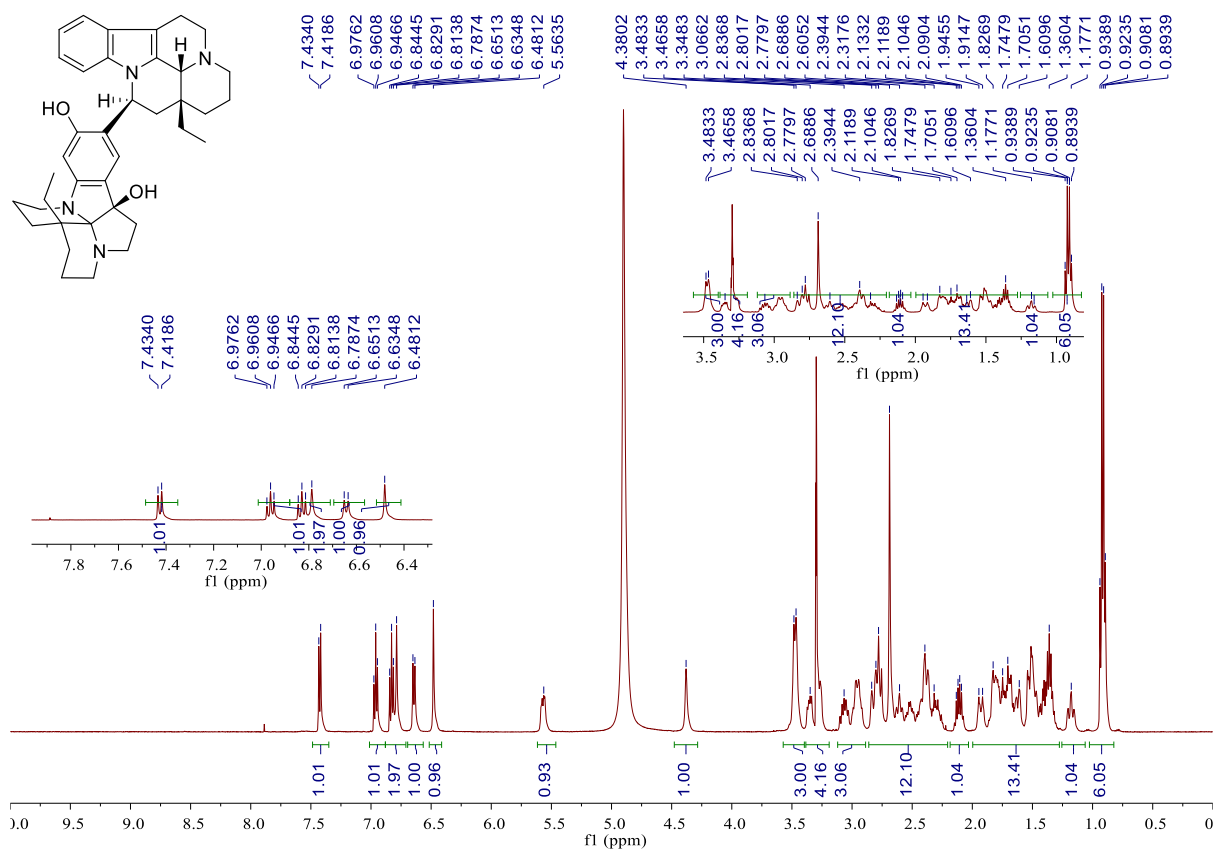


Figure S25.  $^{13}\text{C}$  NMR and DEPT of compound 2 in  $\text{CD}_3\text{OD}$

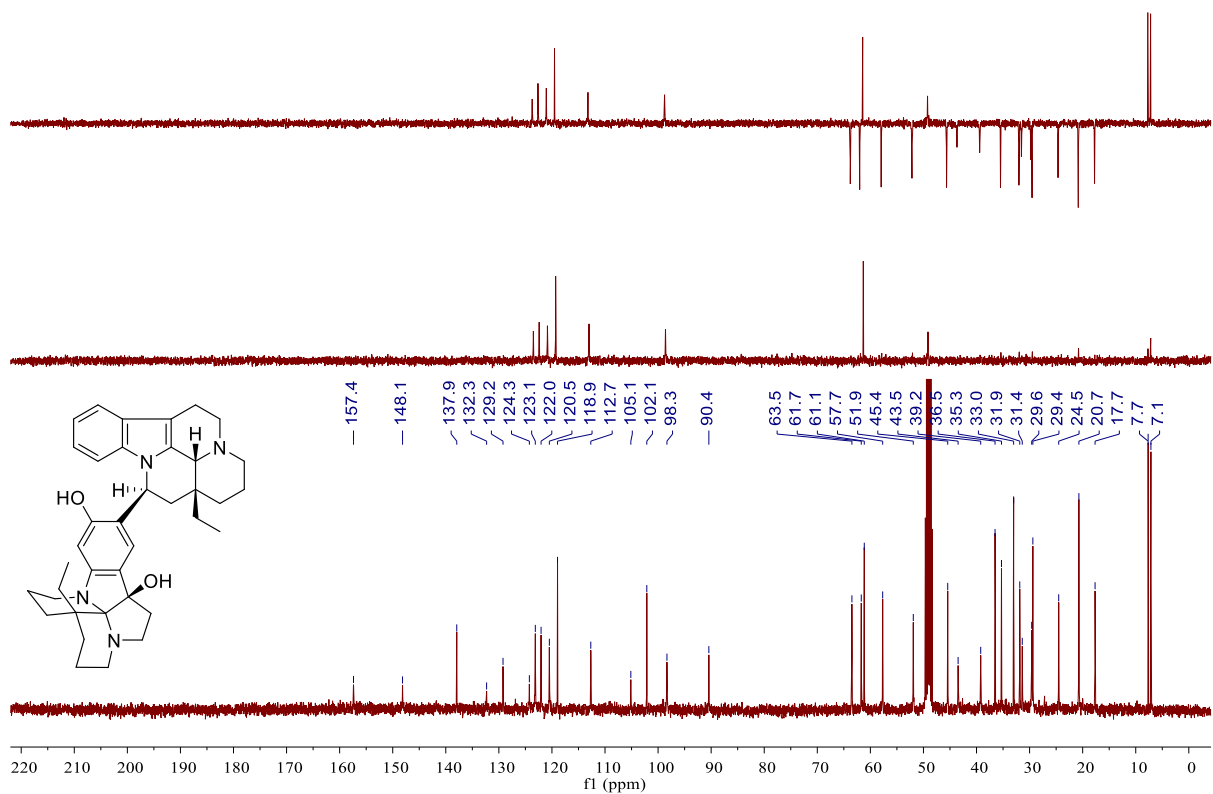




Figure S26. HSQC of compound 2 in CD<sub>3</sub>OD

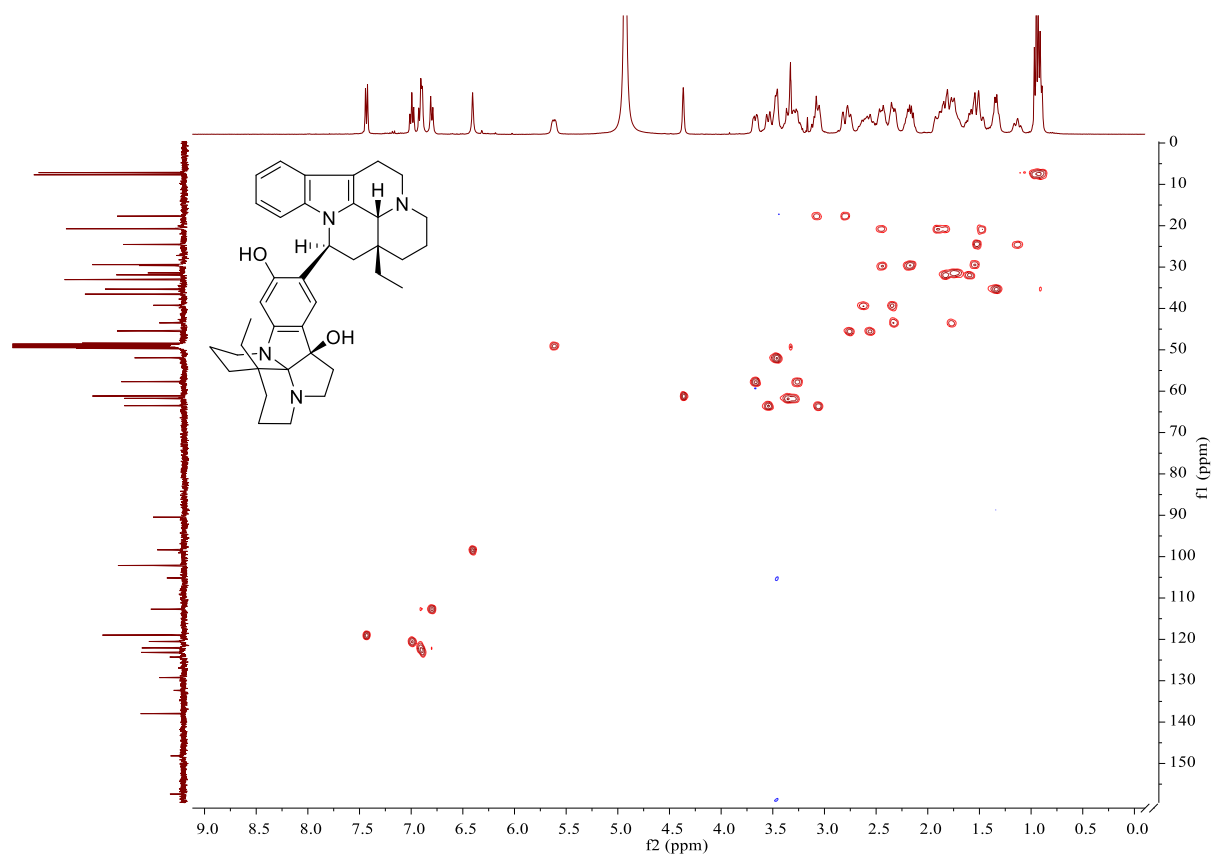
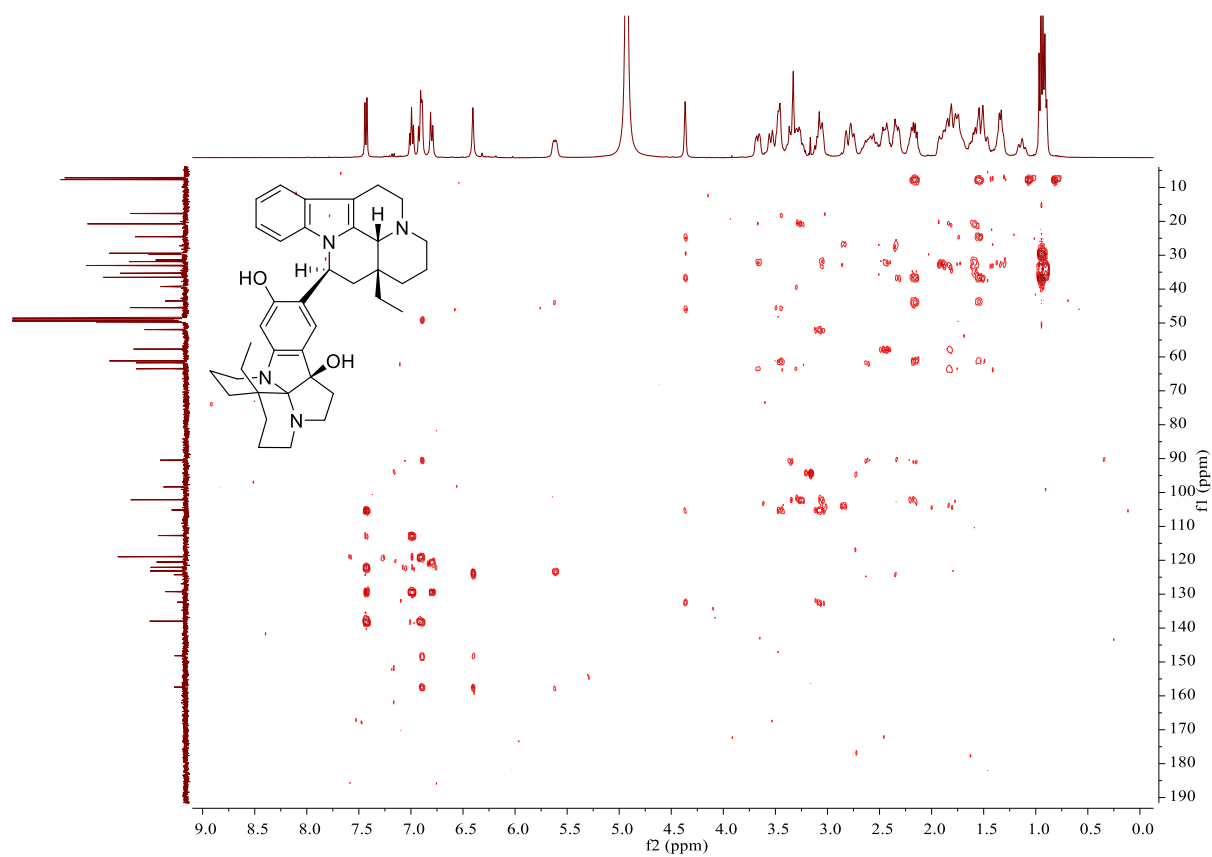
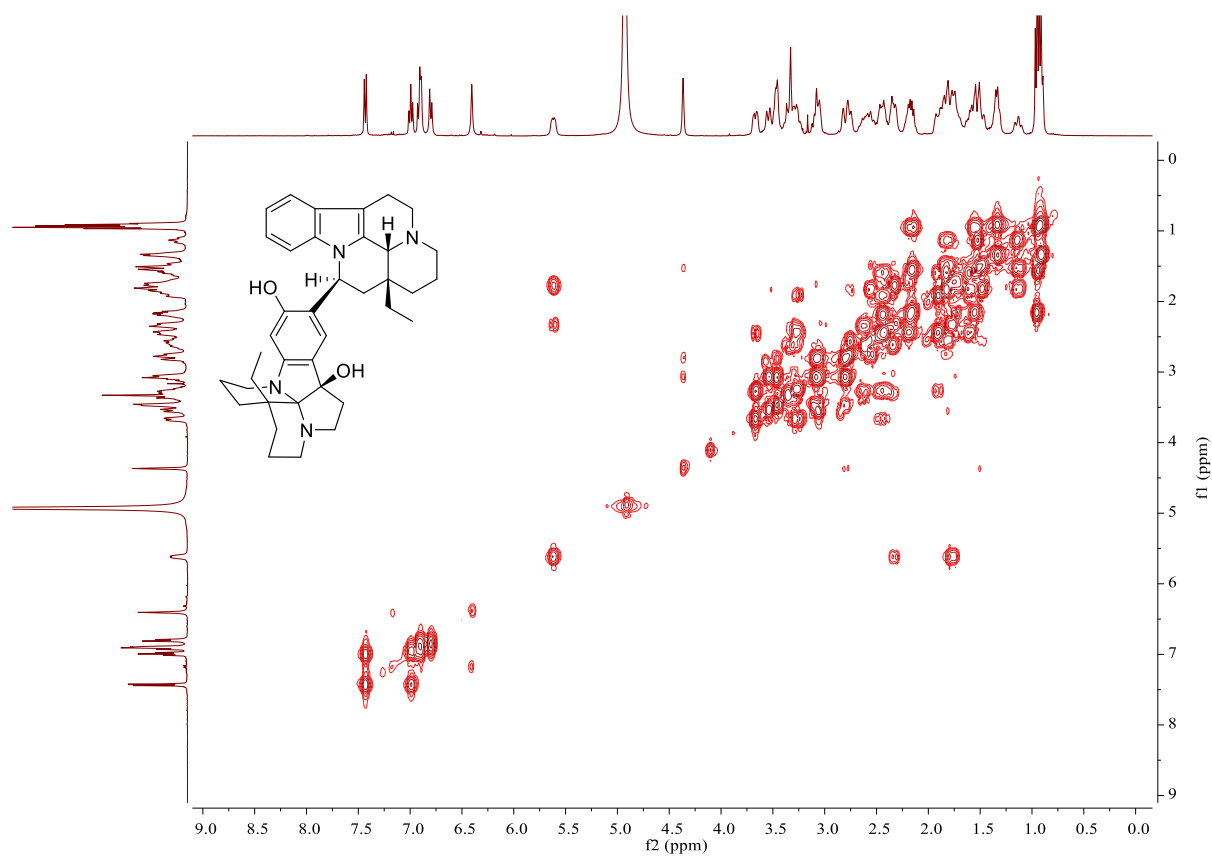


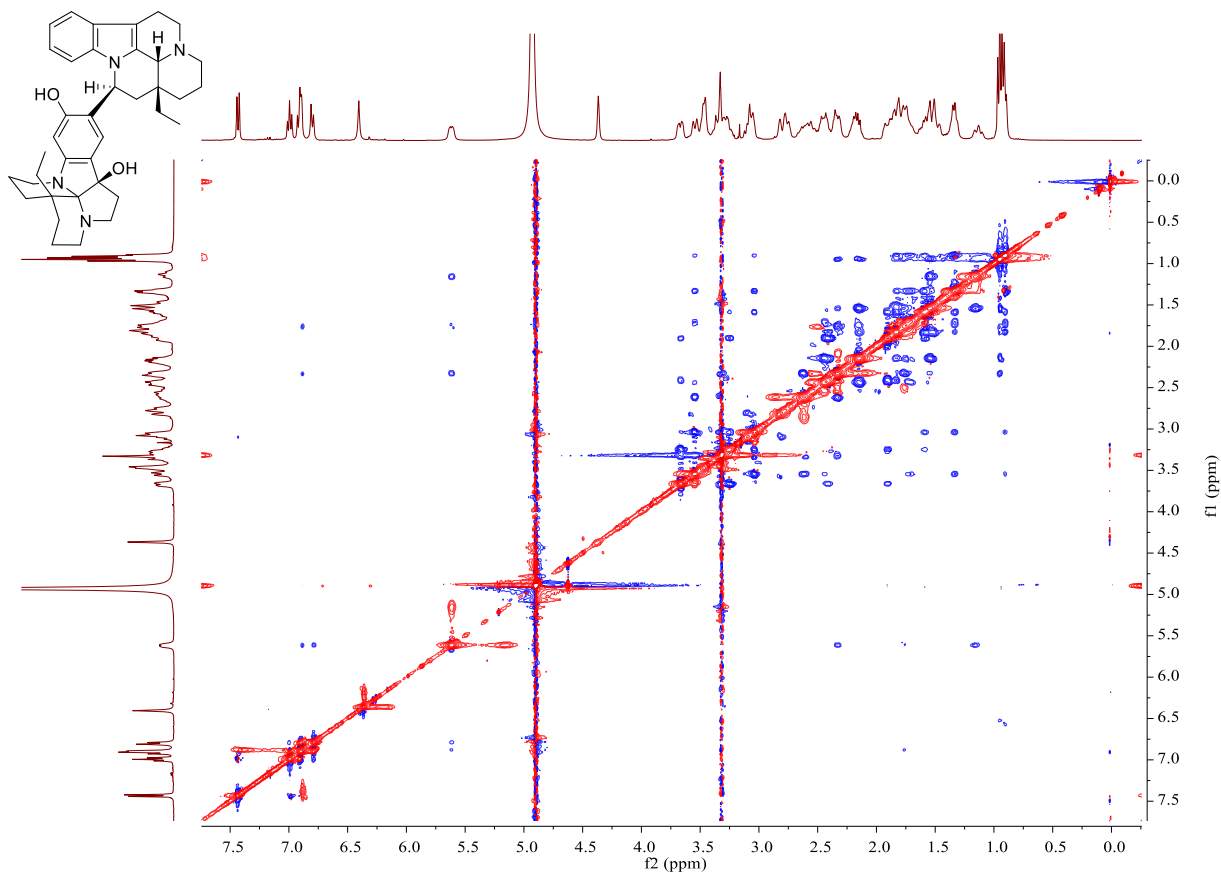
Figure S27. HMBC of compound 2 in CD<sub>3</sub>OD



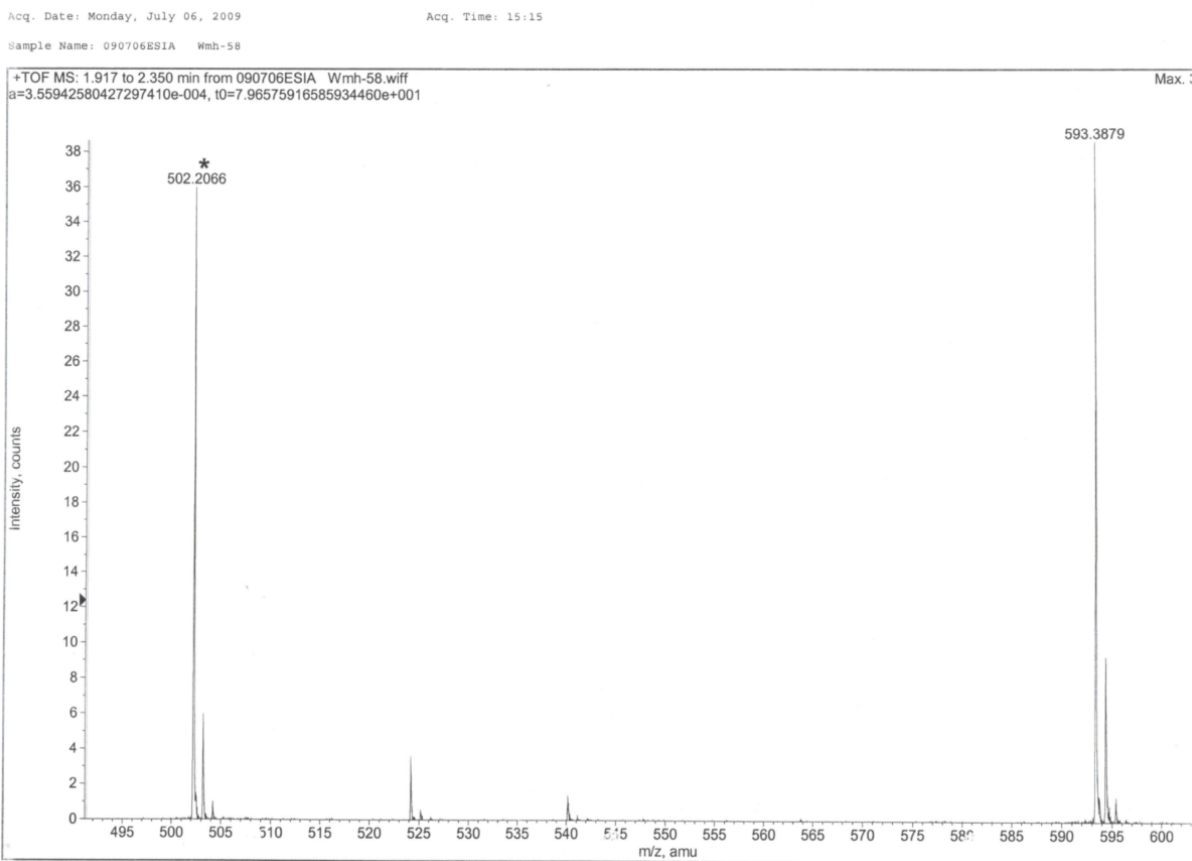
**Figure S28.**  $^1\text{H}$ - $^1\text{H}$  COSY of compound **2** in  $\text{CD}_3\text{OD}$



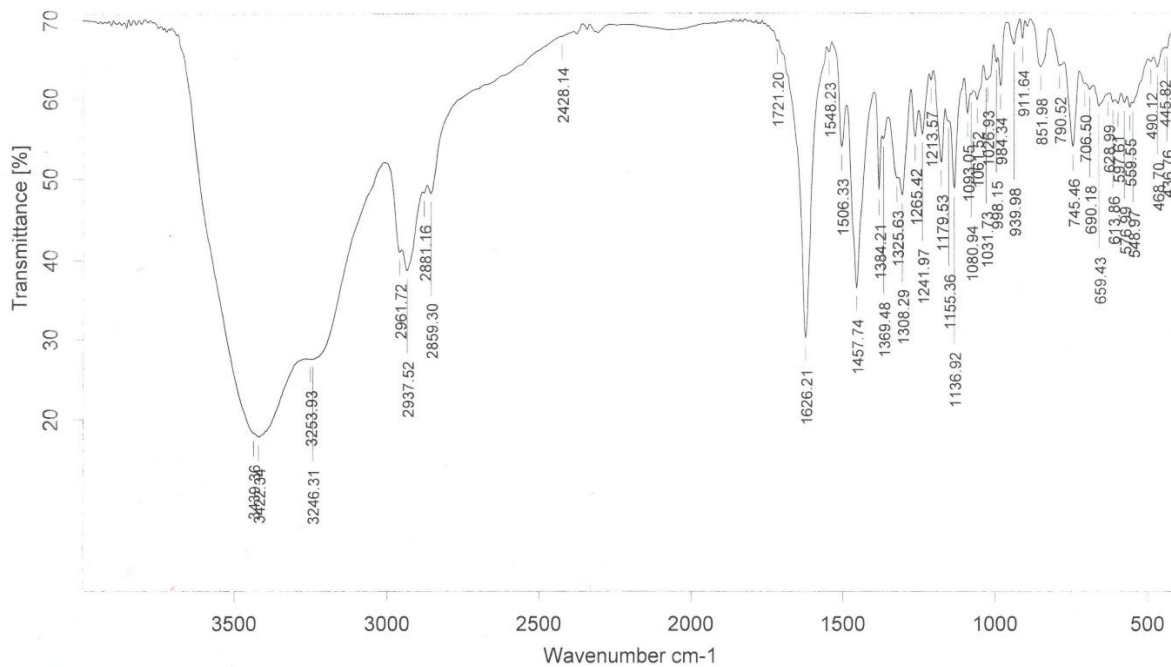
**Figure S29.** ROESY of compound **2** in  $\text{CD}_3\text{OD}$



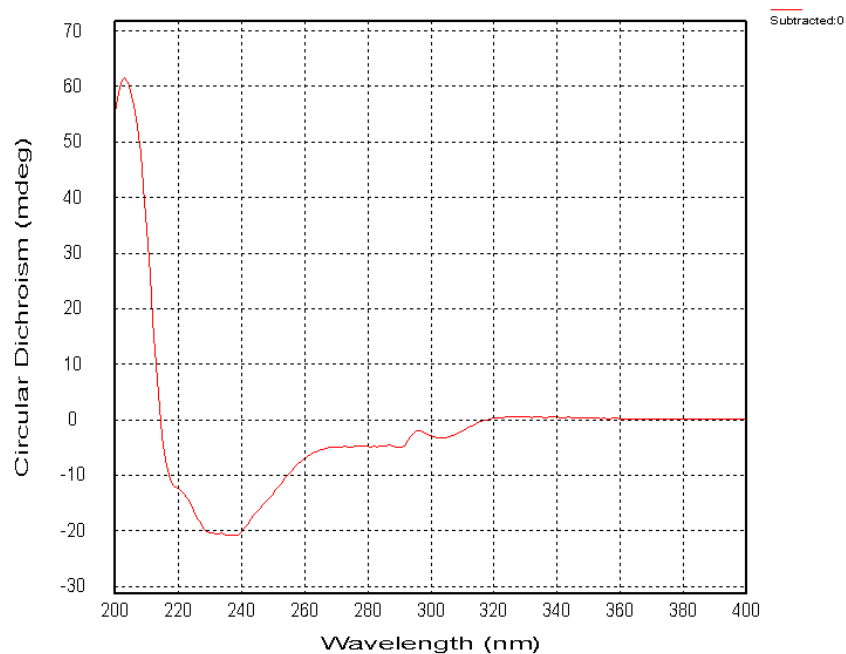
**Figure S30. HR-ESIMS of compound 2**



**Figure S31. IR spectra of compound 2**

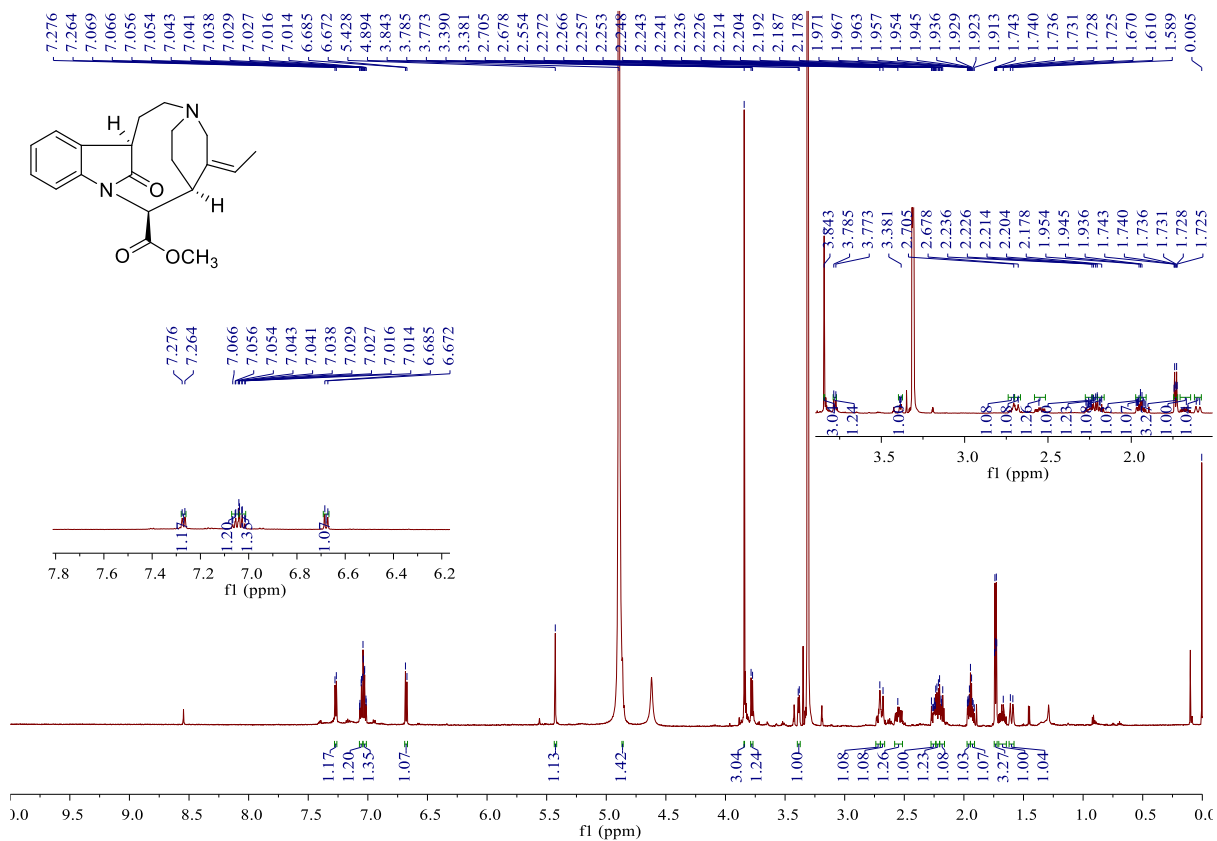


**Figure S32.** CD spectra of compound **2** in CH<sub>3</sub>OH



## Section S6. NMR, MS, IR, and CD spectra for 3

**Figure S33.**  $^1\text{H}$  NMR of compound 3 in  $\text{CD}_3\text{OD}$



**Figure S34.**  $^{13}\text{C}$  NMR and DEPT of compound 3 in  $\text{CD}_3\text{OD}$

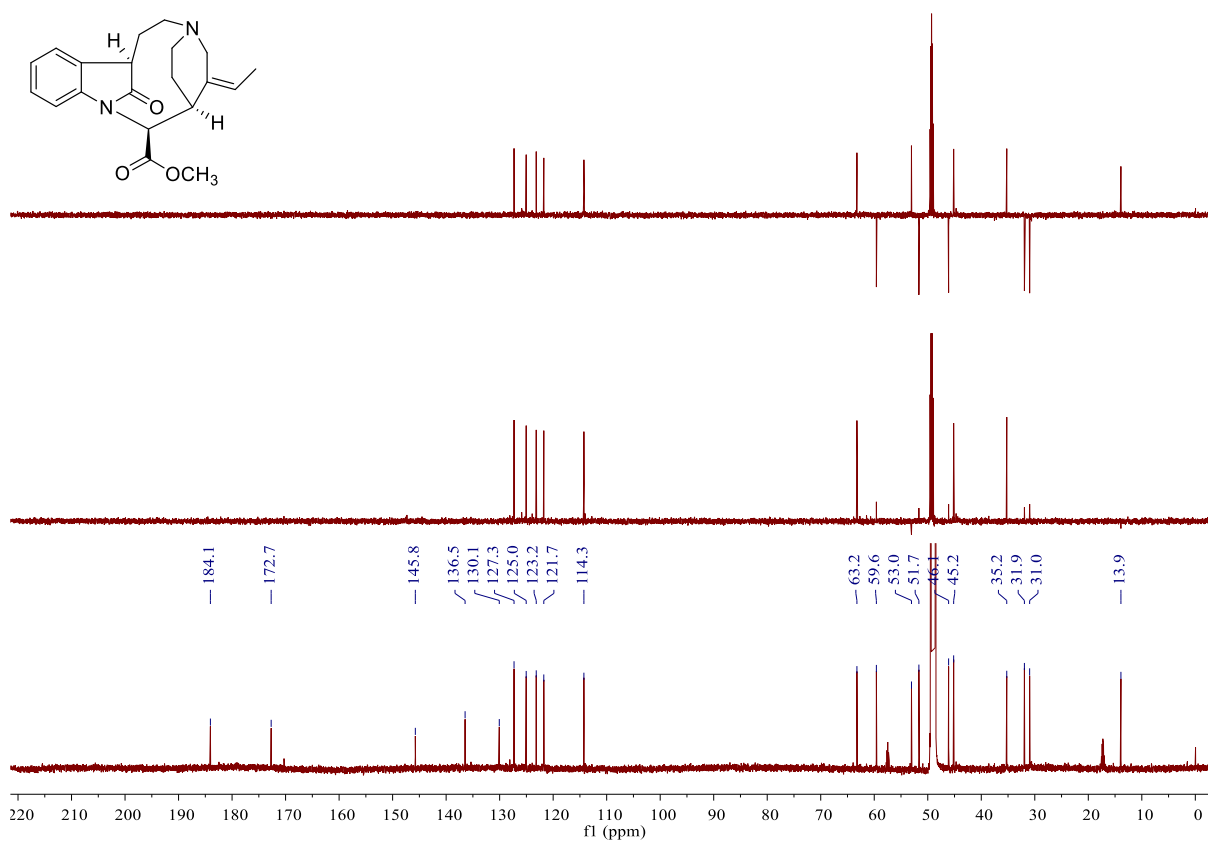


Figure S35. HSQC of compound 3 in CD<sub>3</sub>OD

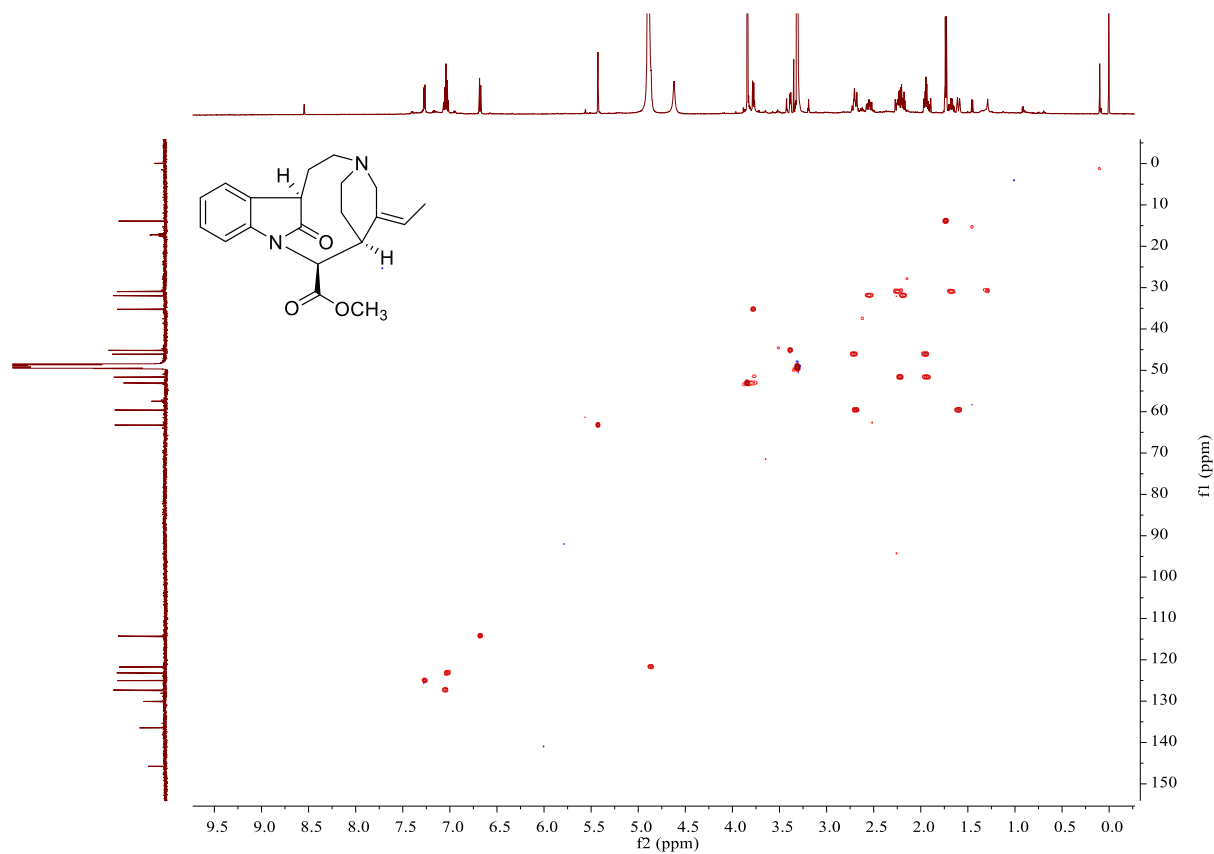


Figure S36. HMBC of compound 3 in CD<sub>3</sub>OD

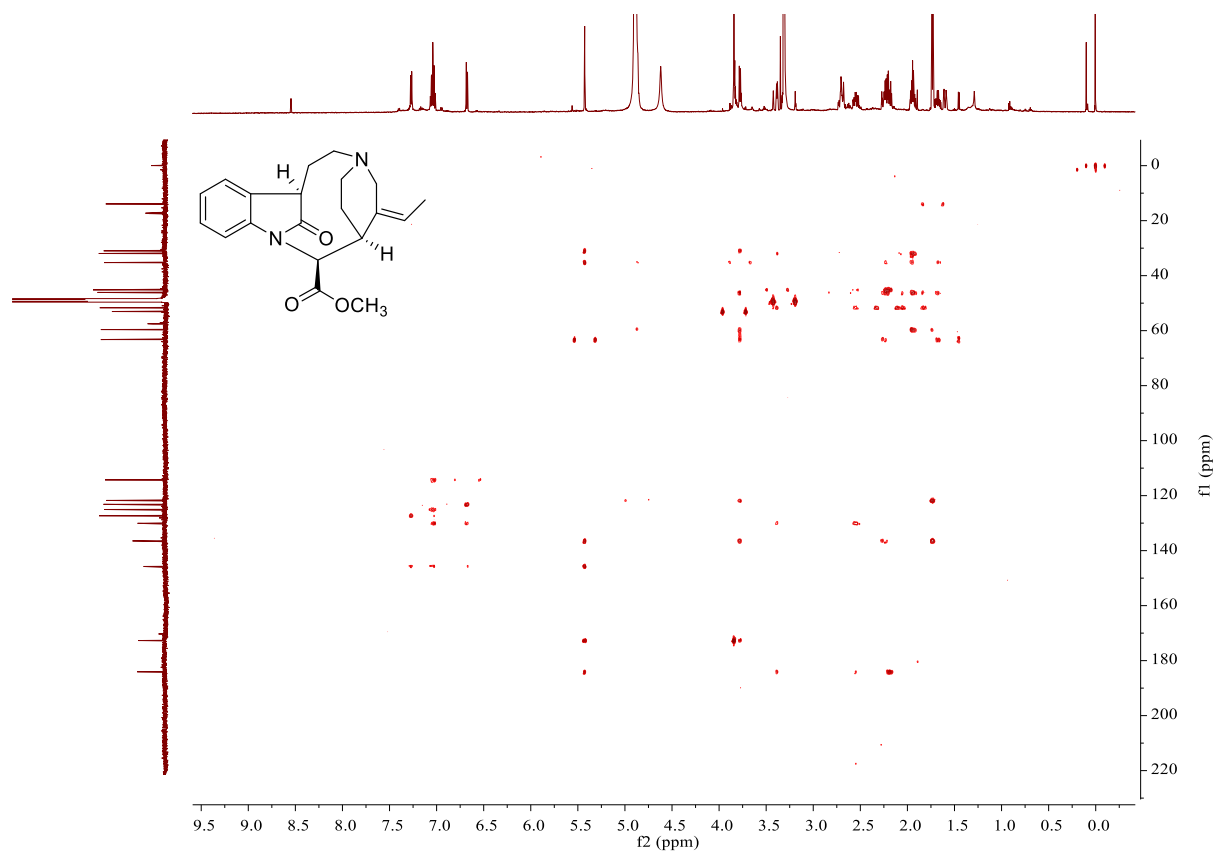


Figure S37.  $^1\text{H}$ - $^1\text{H}$  COSY of compound **3** in  $\text{CD}_3\text{OD}$

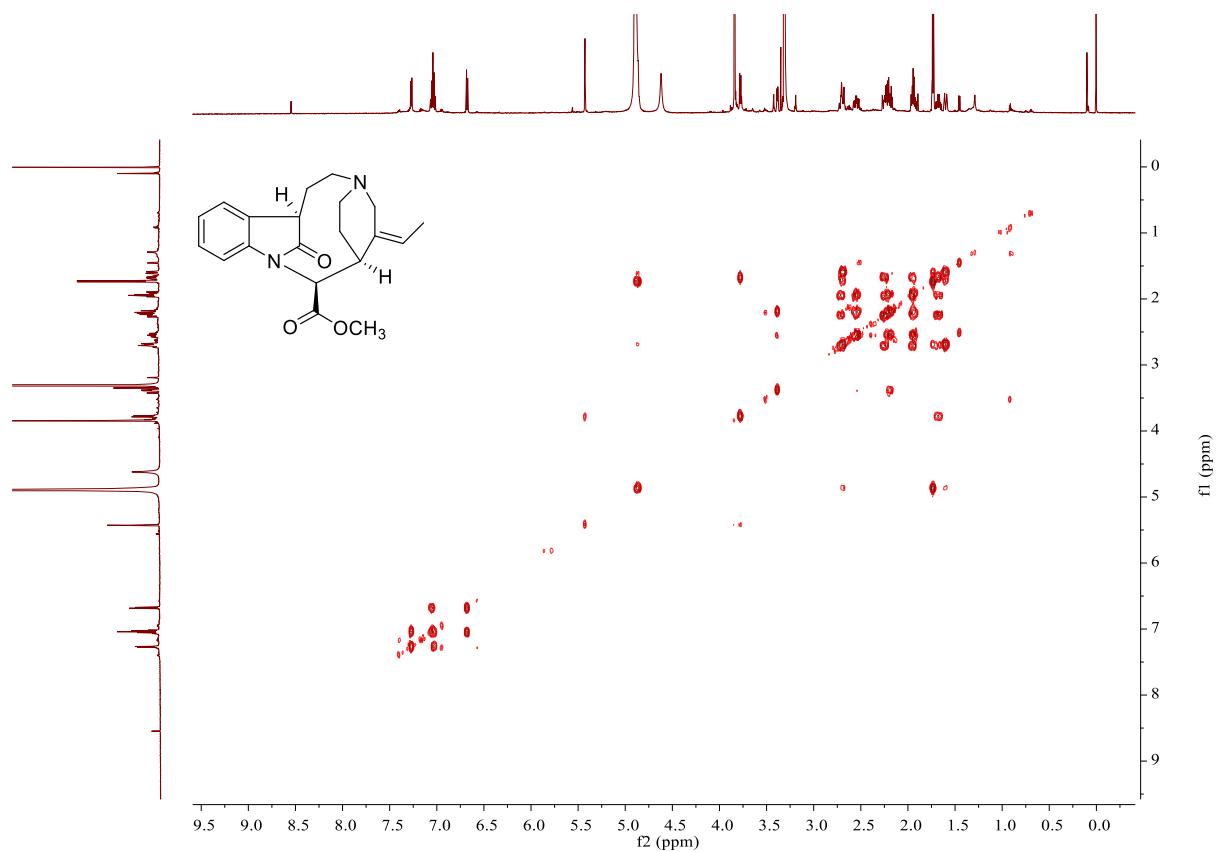
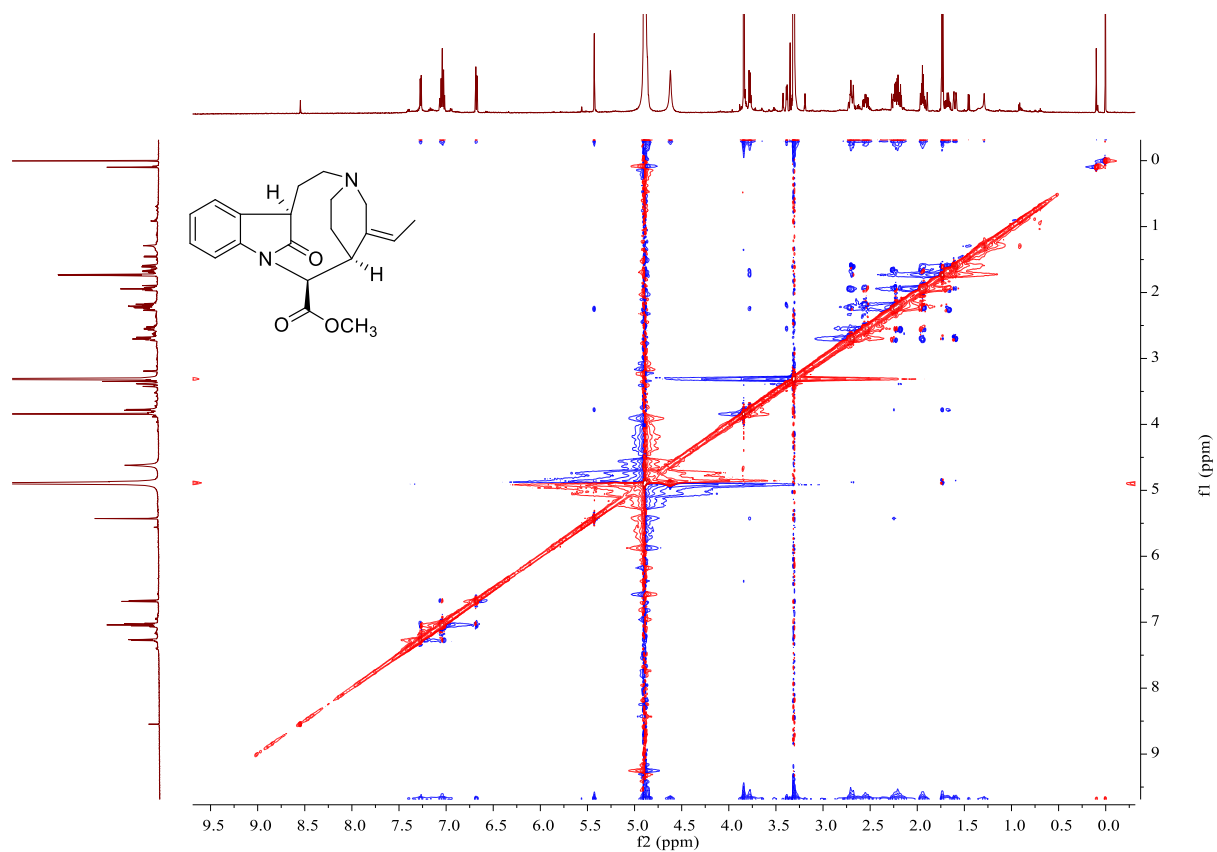
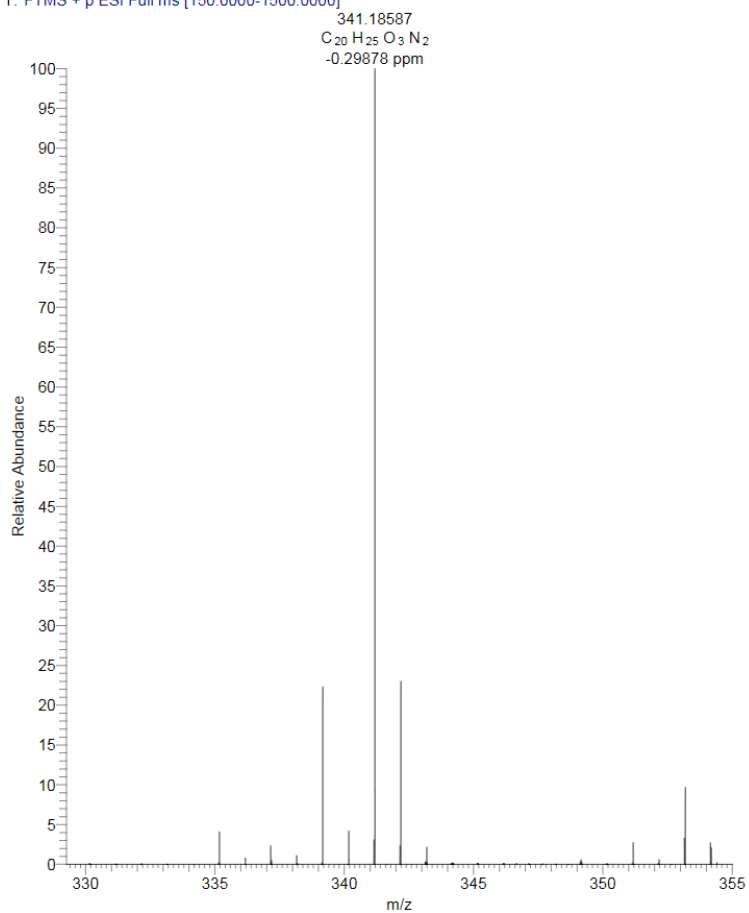


Figure S38. ROESY of compound **3** in  $\text{CD}_3\text{OD}$

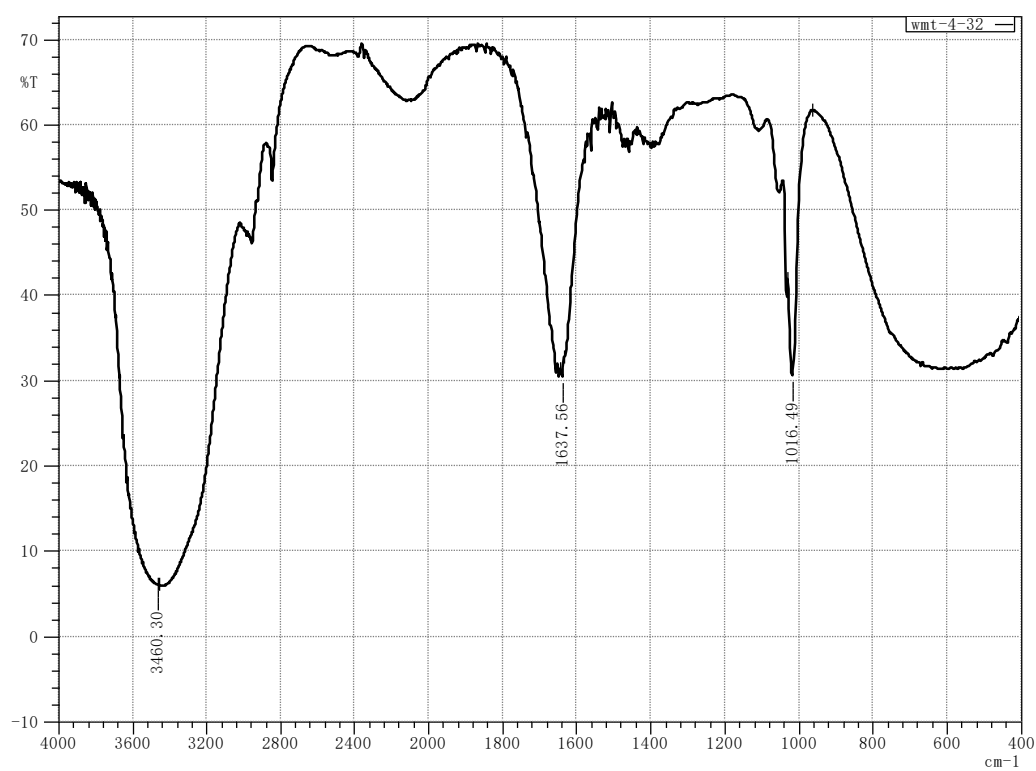


**Figure S39.** HR-ESIMS of compound **3**

T: FTMS + p ESI Full ms [150.0000-1500.0000]

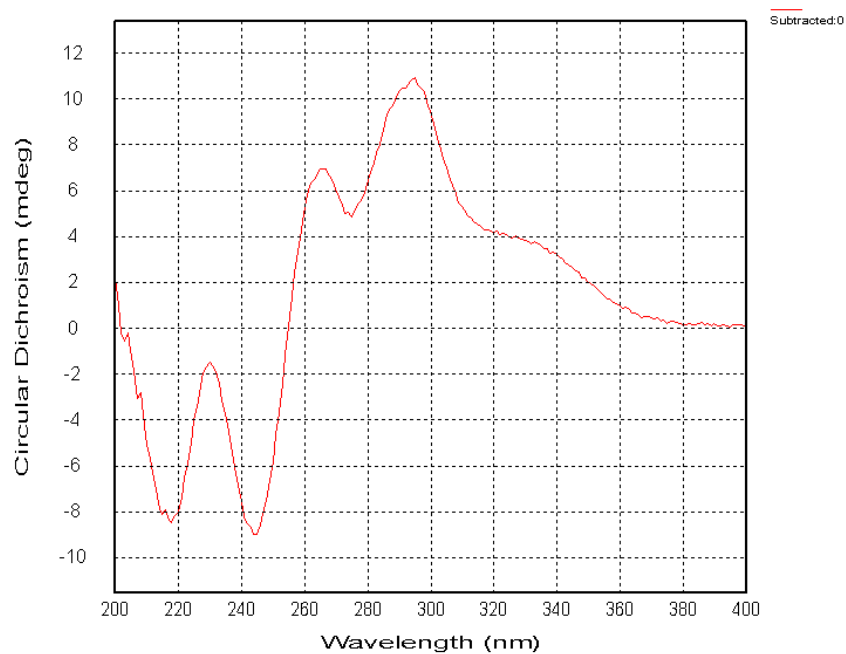


**Figure S40.** IR spectra of compound **3**





**Figure S41.** CD spectra of compound **3** in CH<sub>3</sub>OH



## Section S7. NMR, MS, IR, and CD spectra for 4

Figure S42. <sup>1</sup>H NMR of compound 4 in CD<sub>3</sub>OD

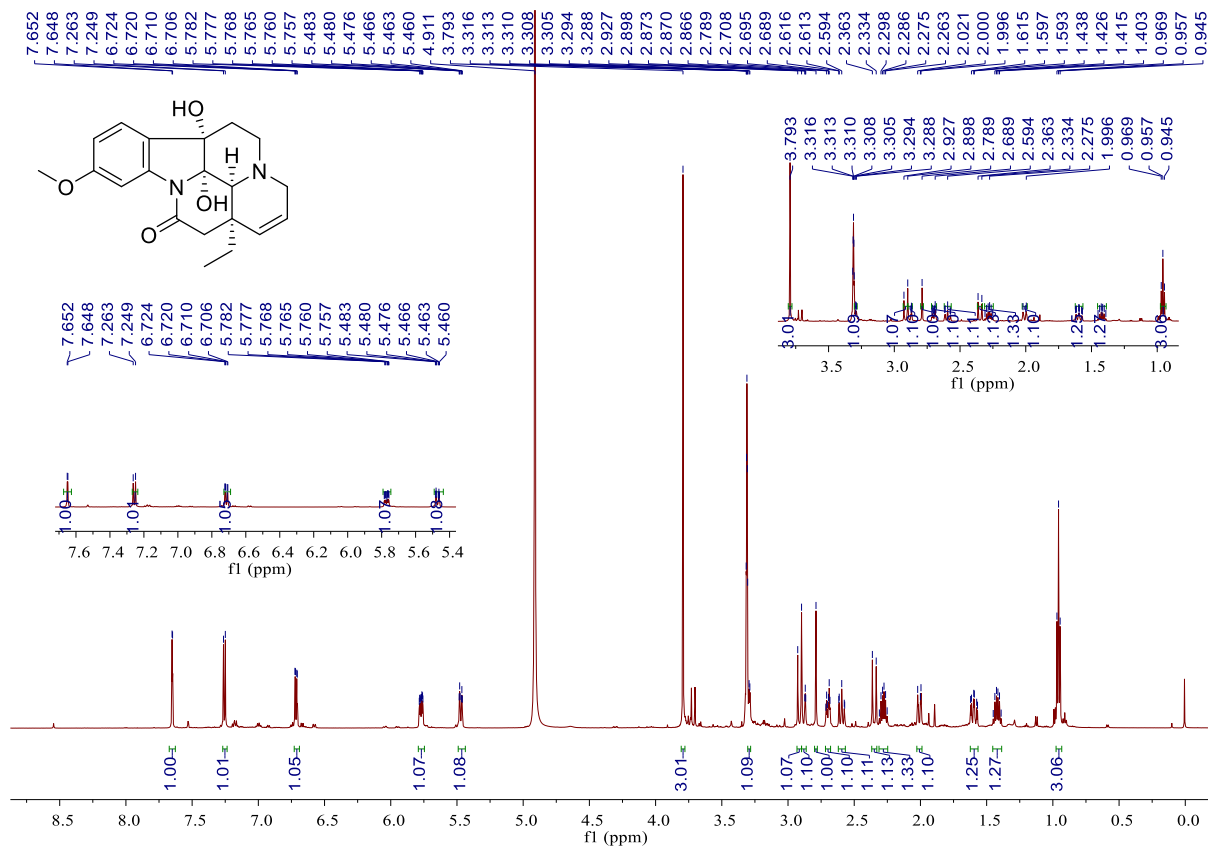


Figure S43. <sup>13</sup>C NMR and DEPT of compound 4 in CD<sub>3</sub>OD

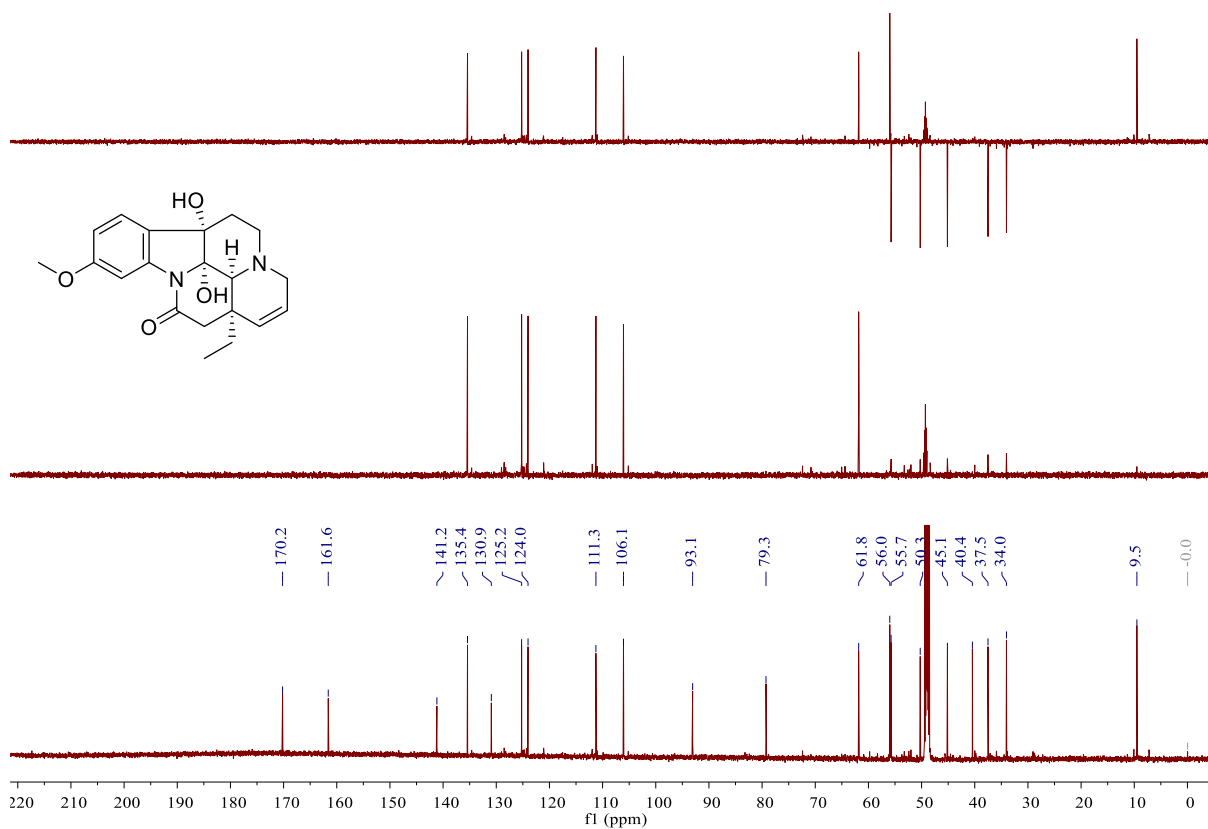


Figure S44. HSQC of compound **4** in CD<sub>3</sub>OD

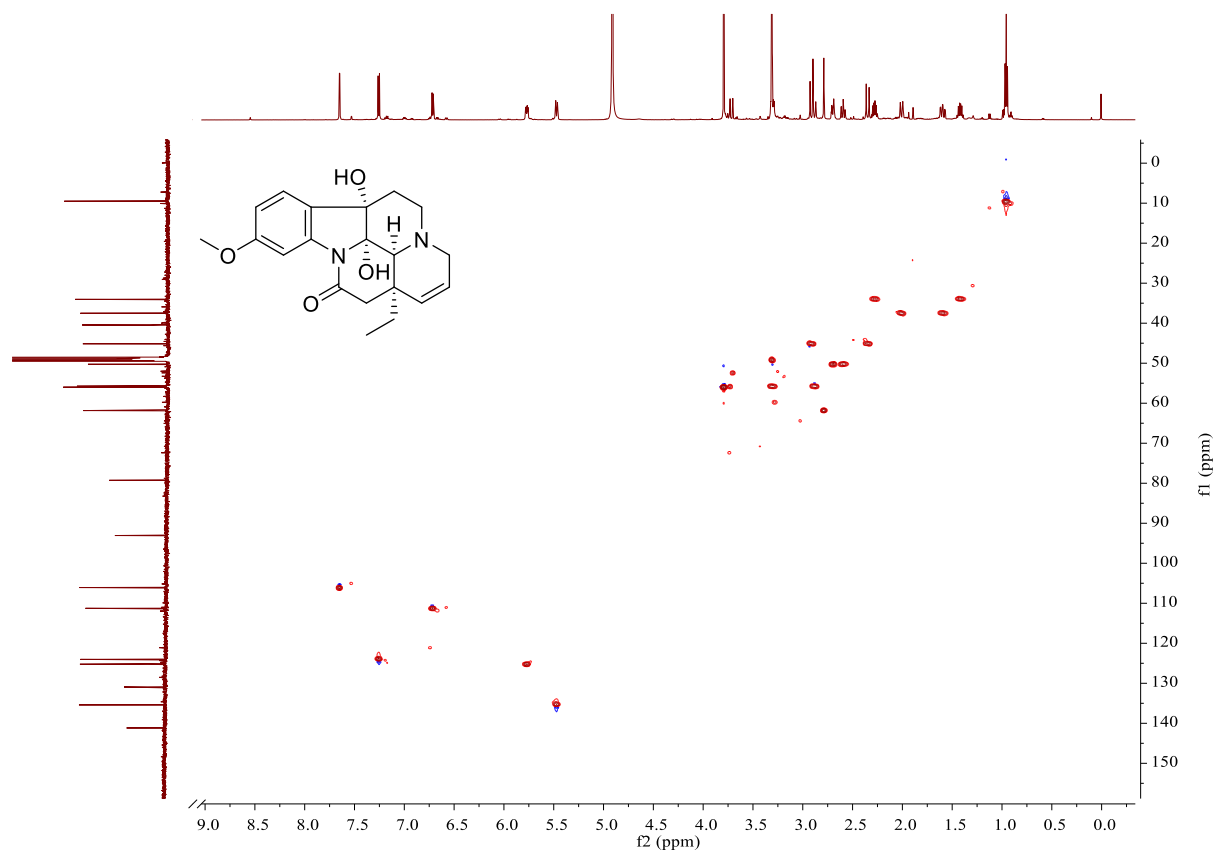


Figure S45. HMBC of compound **4** in CD<sub>3</sub>OD

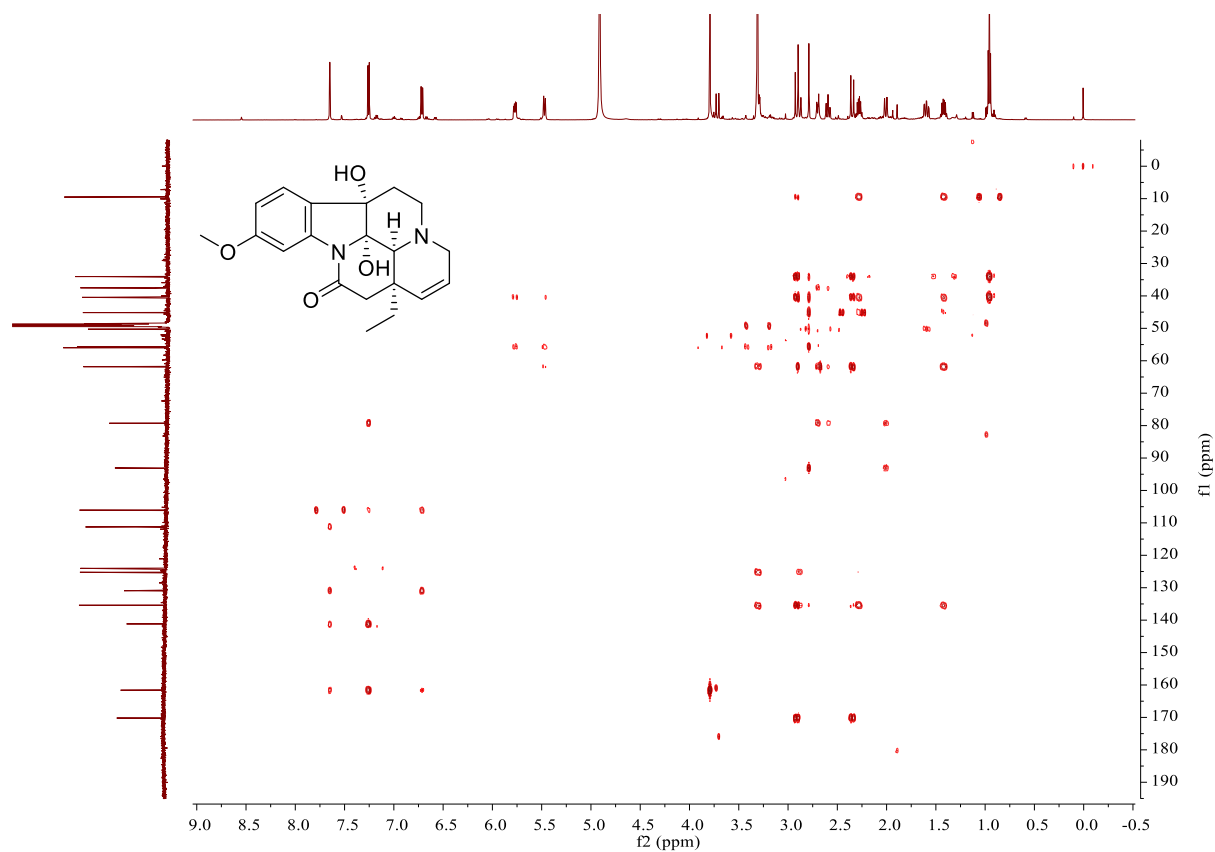


Figure S46.  $^1\text{H}$ - $^1\text{H}$  COSY of compound **4** in  $\text{CD}_3\text{OD}$

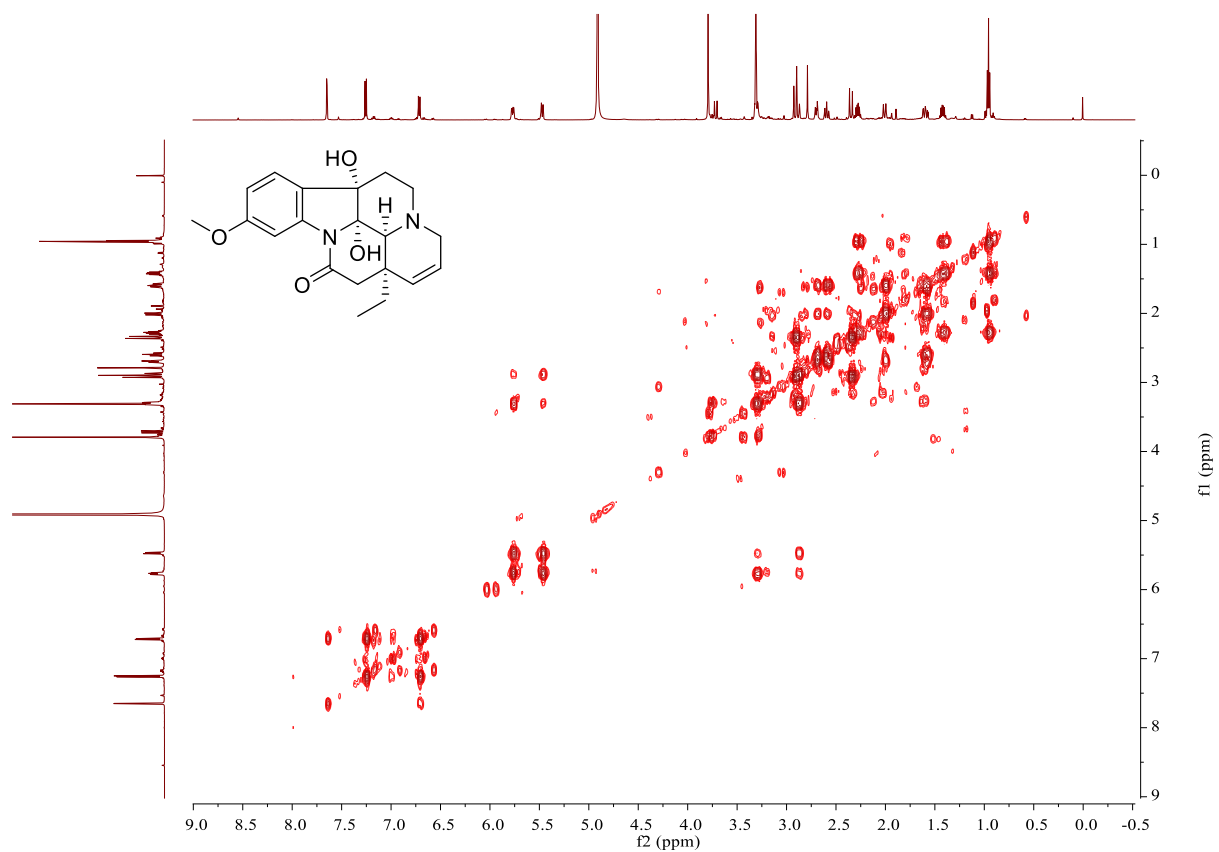
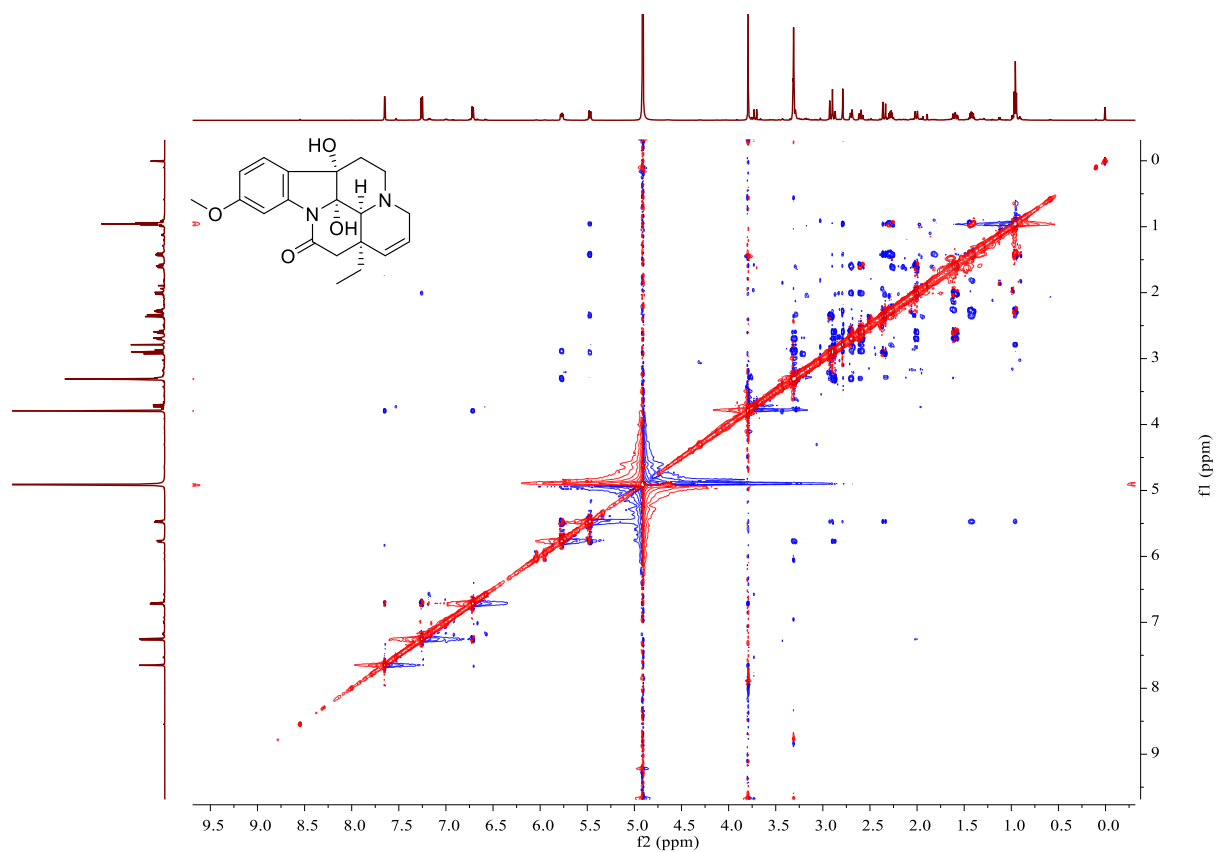
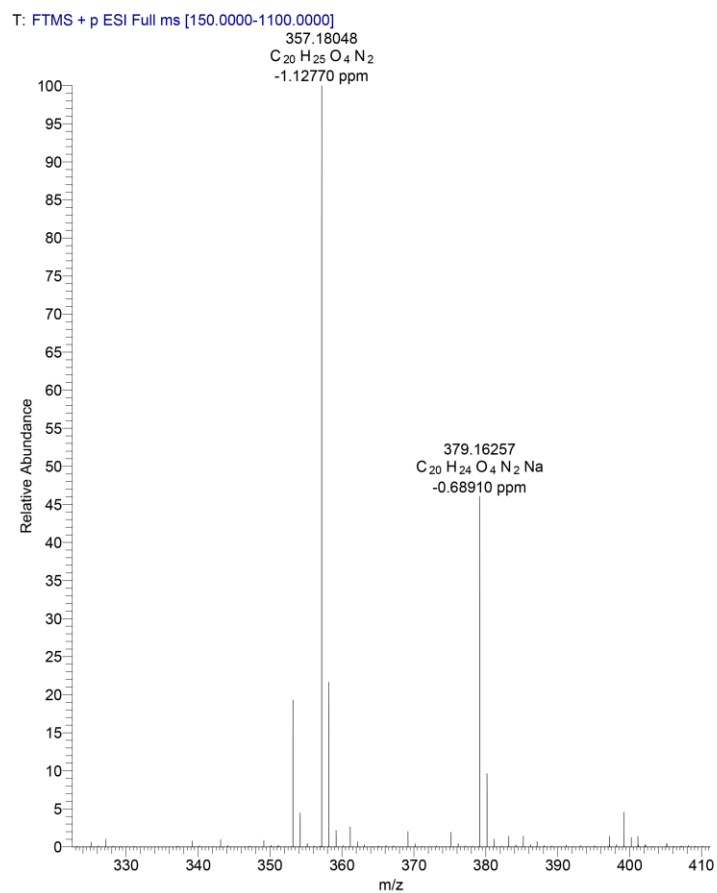


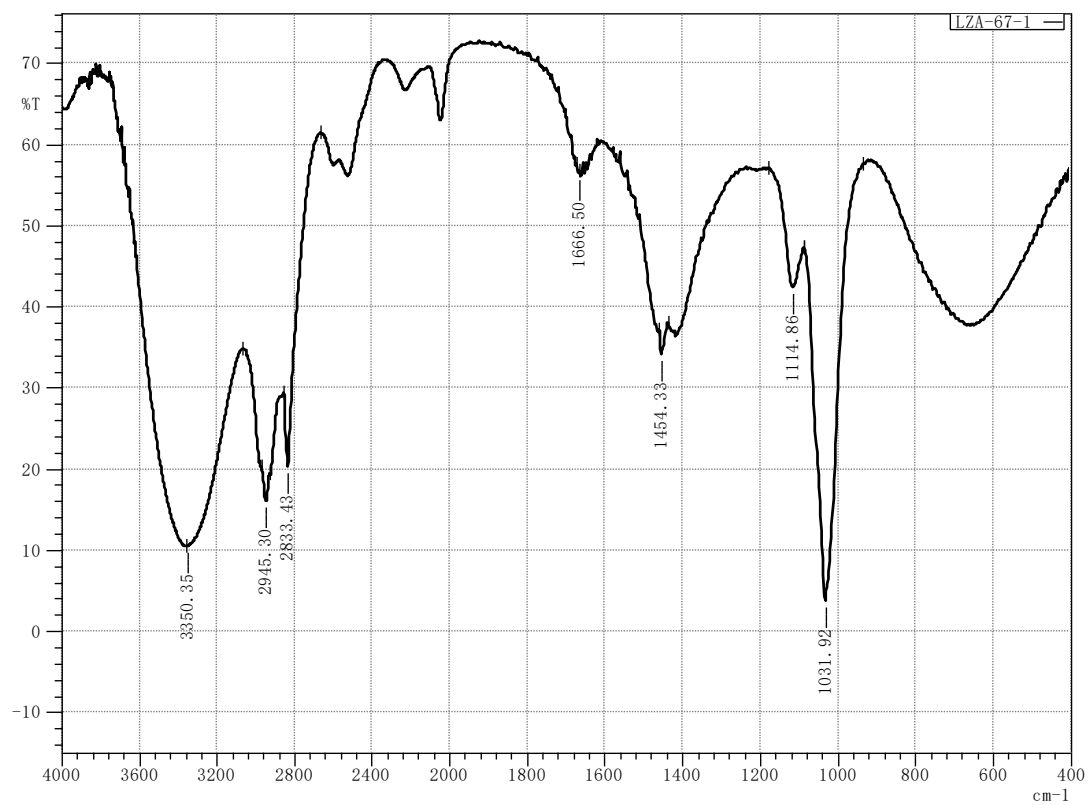
Figure S47. ROESY of compound **4** in  $\text{CD}_3\text{OD}$



**Figure S48.** HR-ESIMS of compound 4



**Figure S49.** IR spectra of compound 4



**Figure S50.** CD spectra of compound **4** in CH<sub>3</sub>OH

