

**Xing-Chen Guo** <sup>1,†</sup>, **Ya-Hui Zhang** <sup>1,†</sup>, **Wen-Bin Gao** <sup>2</sup>, **Li Pan** <sup>3</sup>, **Hua-Jie Zhu** <sup>1,\*</sup> and **Fei Cao** <sup>1,\*</sup>

<sup>1</sup> College of Pharmaceutical Sciences, Institute of Life Science and Green Development, Key Laboratory of Medicinal Chemistry and Molecular Diagnosis of Ministry of Education, Hebei University, Baoding 071002, China; guoxingchen92@163.com (X.-C.G.); 15689932652@163.com (Y.-H.Z.)

<sup>2</sup> College of Life Science, Cangzhou Normal University, Cangzhou 061001, China; wenbinxing@yeah.net

<sup>3</sup> State Key Laboratory of NBC Protection for Civilian, Beijing, 102205, China; bk6180b@163.com

\* Correspondence: hjzhu2019@163.com (H.-J.Z.); caofei542927001@163.com (F.C.)

† These authors contributed equally to this work.

## List of Supporting Information

**Figure S1.** <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of **1**.

**Figure S2.** <sup>13</sup>C NMR (150 MHz, DMSO-*d*<sub>6</sub>) spectrum of **1**.

**Figure S3.** HSQC (DMSO-*d*<sub>6</sub>) spectrum of **1**.

**Figure S4.** <sup>1</sup>H-<sup>1</sup>H COSY (DMSO-*d*<sub>6</sub>) spectrum of **1**.

**Figure S5.** HMBC (DMSO-*d*<sub>6</sub>) spectrum of **1**.

**Figure S6.** HRESIMS for compound **1**.

**Figure S7.** <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of **2**.

**Figure S8.** <sup>13</sup>C NMR (150 MHz, DMSO-*d*<sub>6</sub>) spectrum of **2**.

**Figure S9.** HSQC (DMSO-*d*<sub>6</sub>) spectrum of **2**.

**Figure S10.** <sup>1</sup>H-<sup>1</sup>H COSY (DMSO-*d*<sub>6</sub>) spectrum of **2**.

**Figure S11.** HMBC (DMSO-*d*<sub>6</sub>) spectrum of **2**.

**Figure S12.** 1D NOE (DMSO-*d*<sub>6</sub>) spectrum of **compounds 1 and 2**.

**Figure S13.** HRESIMS for compound **2**.

**Figure S14.** <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of **3**.

**Figure S15.** <sup>1</sup>H NMR (600 MHz, CDCl<sub>3</sub>) spectrum of **3**.

**Figure S16.** <sup>13</sup>C NMR (150 MHz, DMSO-*d*<sub>6</sub>) spectrum of **3**.

**Figure S17.** HSQC (DMSO-*d*<sub>6</sub>) spectrum of **3**.

**Figure S18.** <sup>1</sup>H-<sup>1</sup>H COSY (DMSO-*d*<sub>6</sub>) spectrum of **3**.

**Figure S19.** HMBC (DMSO-*d*<sub>6</sub>) spectrum of **3**.

**Figure S20.** NOESY (CDCl<sub>3</sub>) spectrum of compound **3**.

**Figure S21.** HRESIMS for compound **3**.

- Figure S22.**  $^1\text{H}$  NMR (600 MHz,  $\text{DMSO-}d_6$ ) spectrum of **4**.
- Figure S23.**  $^1\text{H}$  NMR (600 MHz,  $\text{DMSO-}d_6$ ) spectrum of **5**.
- Figure S24.**  $^1\text{H}$  NMR (600 MHz,  $\text{DMSO-}d_6$ ) spectrum of **6**.
- Figure S25.**  $^1\text{H}$  NMR (600 MHz,  $\text{DMSO-}d_6$ ) spectrum of **7**.
- Figure S26.** Experimental ECD spectra of **1–5**.
- Figure S27.** Experimental and calculated ECD spectra of **1**.
- Figure S28.** Experimental and calculated ECD spectra of **2**.
- Figure S29.** Experimental and calculated ECD spectra of **3**.
- Figure S30.** Experimental and calculated ECD spectra of **4**.
- Figure S31.** Experimental and calculated ECD spectra of **5**.
- Figure S32.** The data of DP4plus method of compound **3** (B3lyp/6-311+G(d,p))
- Figure S33.** The data of DP4plus method of compound **5** (B3lyp/6-311+G(d,p))
- Figure S34.** Compound **1** tightly bind to the entire active pocket of *OfChi-h*
- Figure S35.** Three hydrogen bonds formed by **1** with the guanidine group of ARG439
- Figure S36.**  $\pi$ -Sulfur interaction between **1** and the sulfur atom of methionine MET381
- Figure S37.**  $\pi$ - $\pi$  Stacking between **1** and the benzene rings of Trp268 and Phe309
- Figure S38.** Alkyl hydrophobic interactions between **1** and Ala355/Met381, and between **1** and Val469
- Figure S39.** Mixed  $\pi$ /alkyl hydrophobic interactions between **1** and Tyr156, and between **1** and Phe184
- Figure S40.** Compound **4** tightly bind to the *OfHex1* in a "U" conformation
- Figure S41.** Hydrogen bonds formed between **4** and the guanidine group of ARG220, and between **4** and the carboxyl of ASP367
- Figure S42.** Compound **4** had a  $\pi$ -anion with the carboxyhydroxyl oxygen anion in the residue of GLU368
- Figure S43.**  $\pi$ - $\pi$  Stacking interaction between **4** and the indole ring of Trp490
- Figure S44.** Alkyl hydrophobic interaction between **4** and the isopropyl group of Val484
- Figure S45.** Mixed  $\pi$ /alkyl hydrophobic interactions between **4** and Trp322/Trp483, and between **4** and the isopropyl group of Val327
- Figure S46.** N-H hydrogen bond interaction between **2** and Arg220

**Table S1.** Comparison of the data of DP4plus method of compound 3 and 5

**Table S2.** Cytotoxic activity data of compounds 1–7.

**Table S3.** The coordinate for the lowest-energy conformer of compound 1 for ECD calculation.

**Table S4.** The coordinate for the lowest-energy conformer of compound 2 for ECD calculation.

**Table S5.** The coordinate for the lowest-energy conformer of compound 3 for ECD calculation.

**Table S6.** The coordinate for the lowest-energy conformer of compound 4 for ECD calculation.

**Table S7.** The coordinate for the lowest-energy conformer of compound 5 for ECD calculation.

**Table S8-S16.** The coordinates for the conformers of compound 3 for NMR calculation.

**Table S17-S32.** The coordinates for the conformers of compound 5 for NMR calculation.

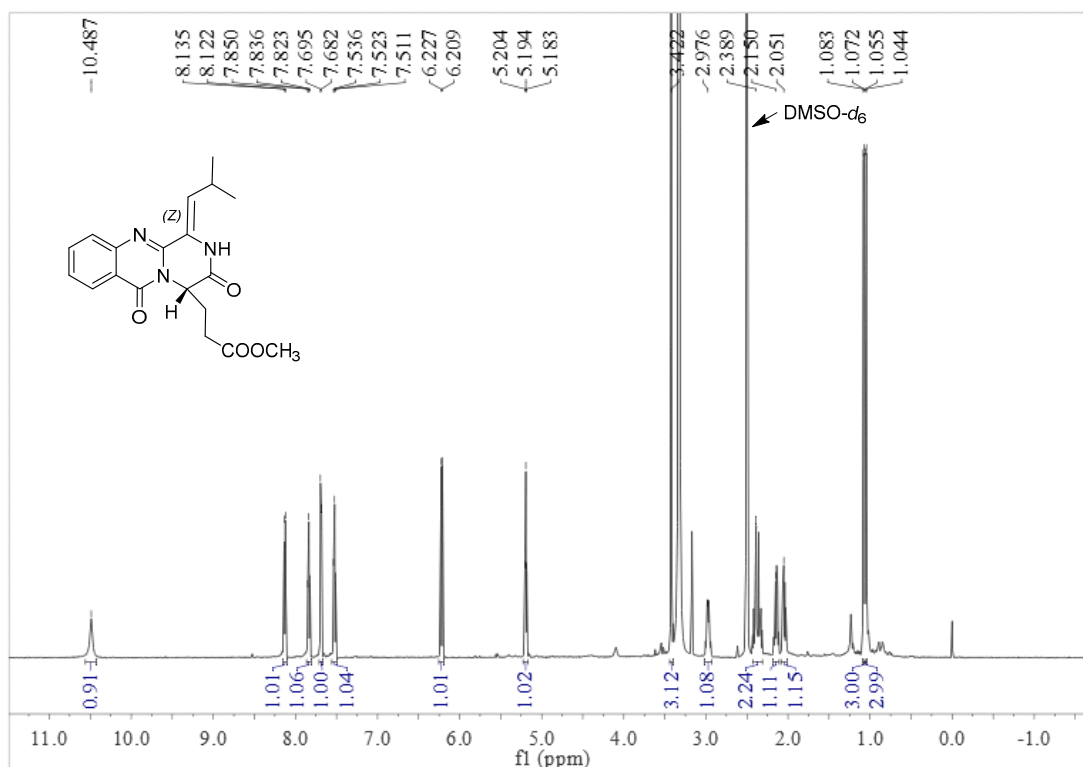


Figure S1. <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of 1.

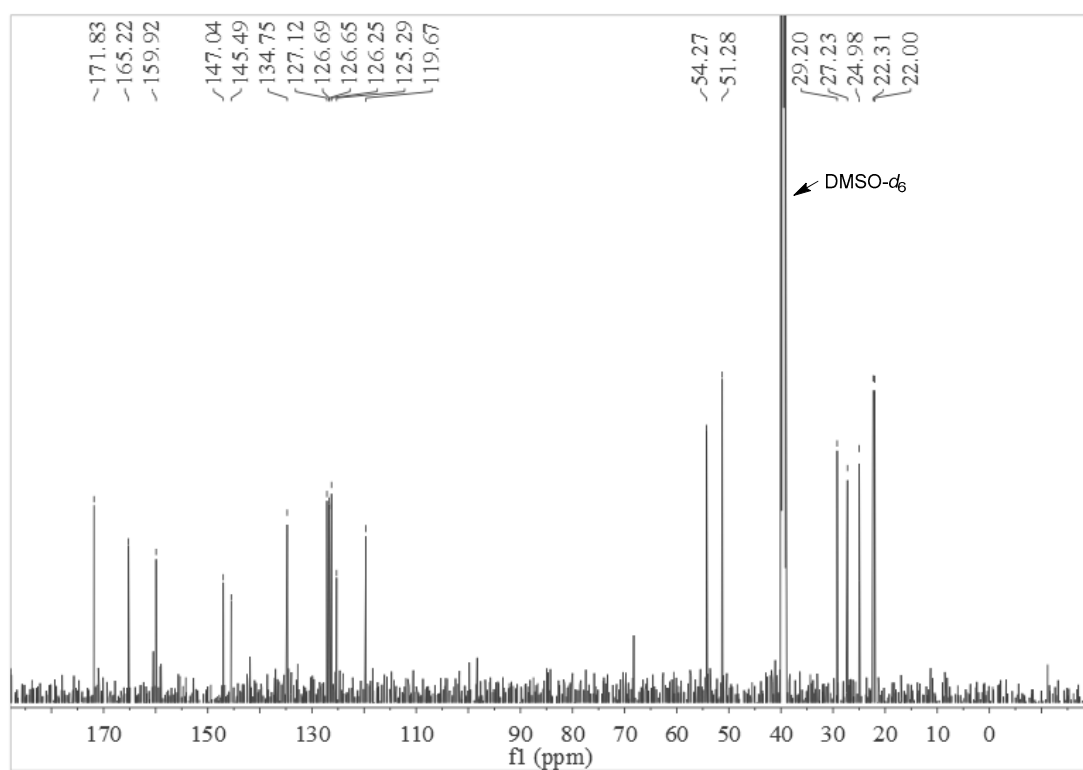
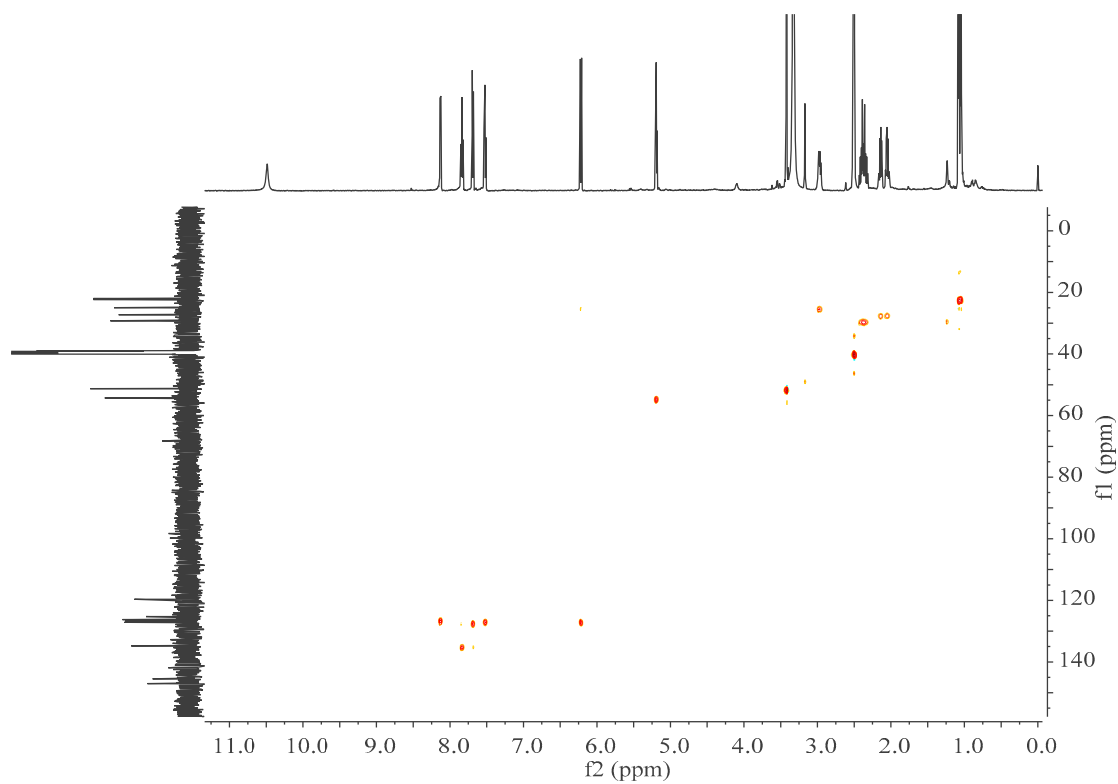
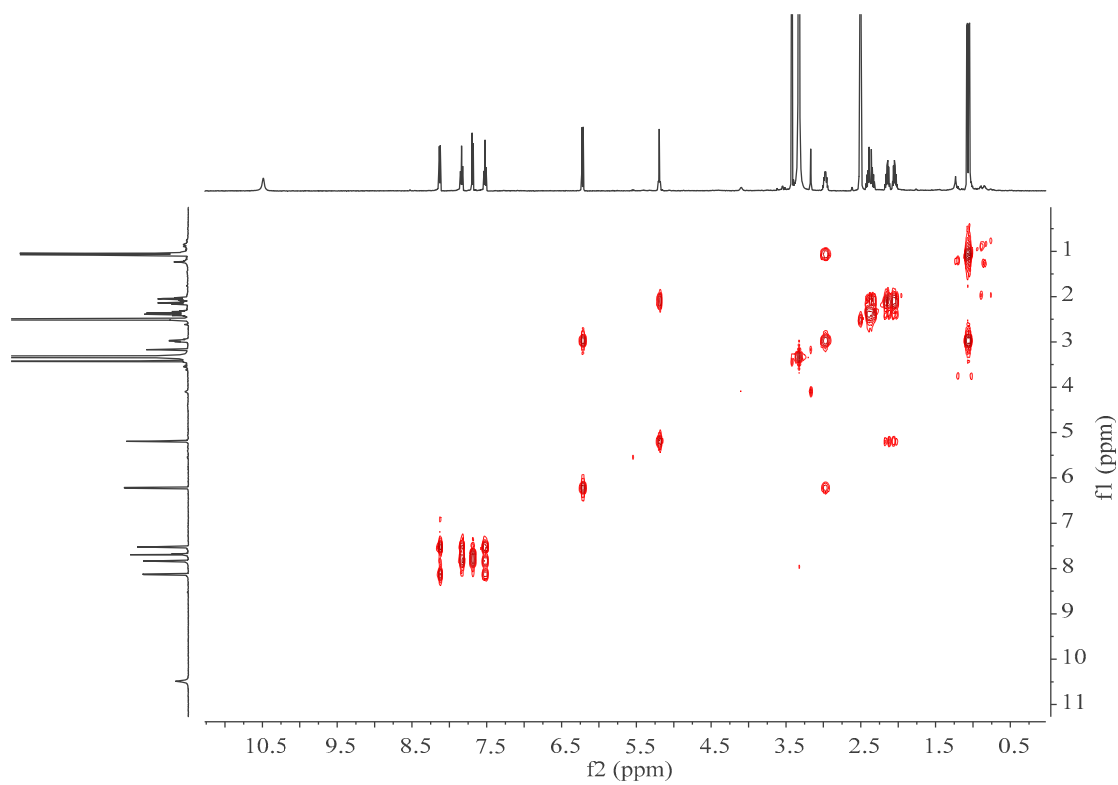


Figure S2. <sup>13</sup>C NMR (150 MHz, DMSO-*d*<sub>6</sub>) spectrum of compound 1.



**Figure S3.** HSQC (DMSO-*d*<sub>6</sub>) spectrum of **1**.



**Figure S4.** <sup>1</sup>H-<sup>1</sup>H COSY (DMSO-*d*<sub>6</sub>) spectrum of **1**.

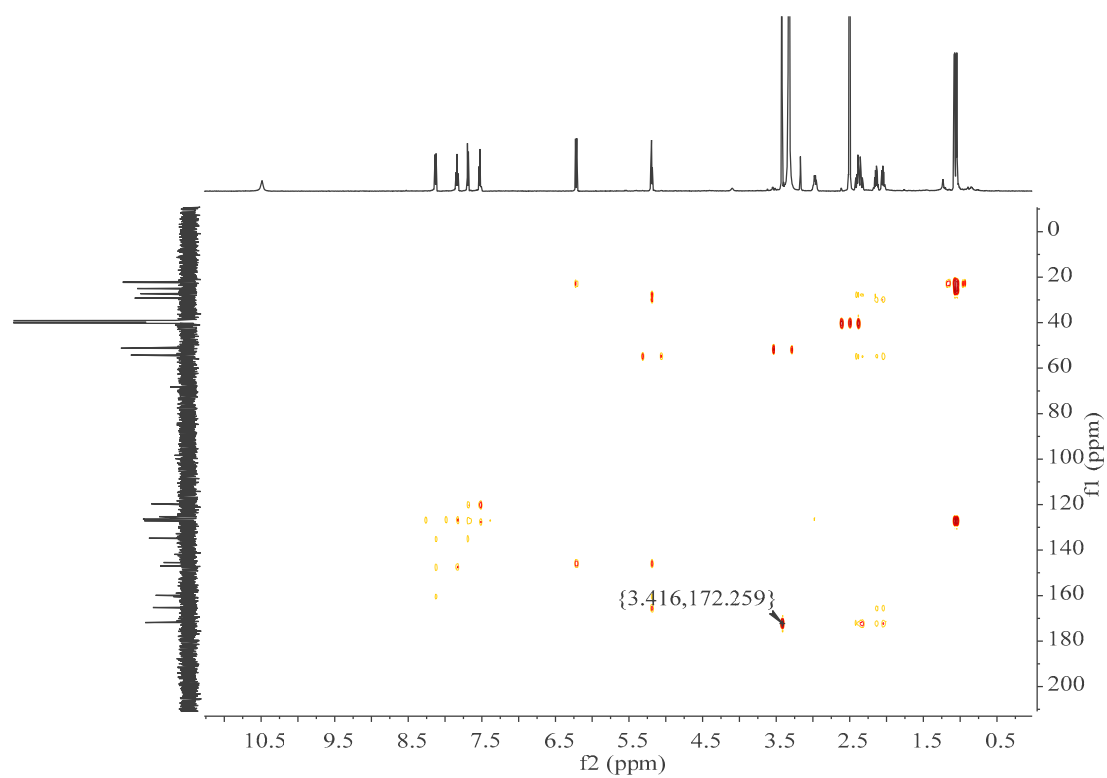


Figure S5. HMBC (DMSO- $d_6$ ) spectrum of **1**.

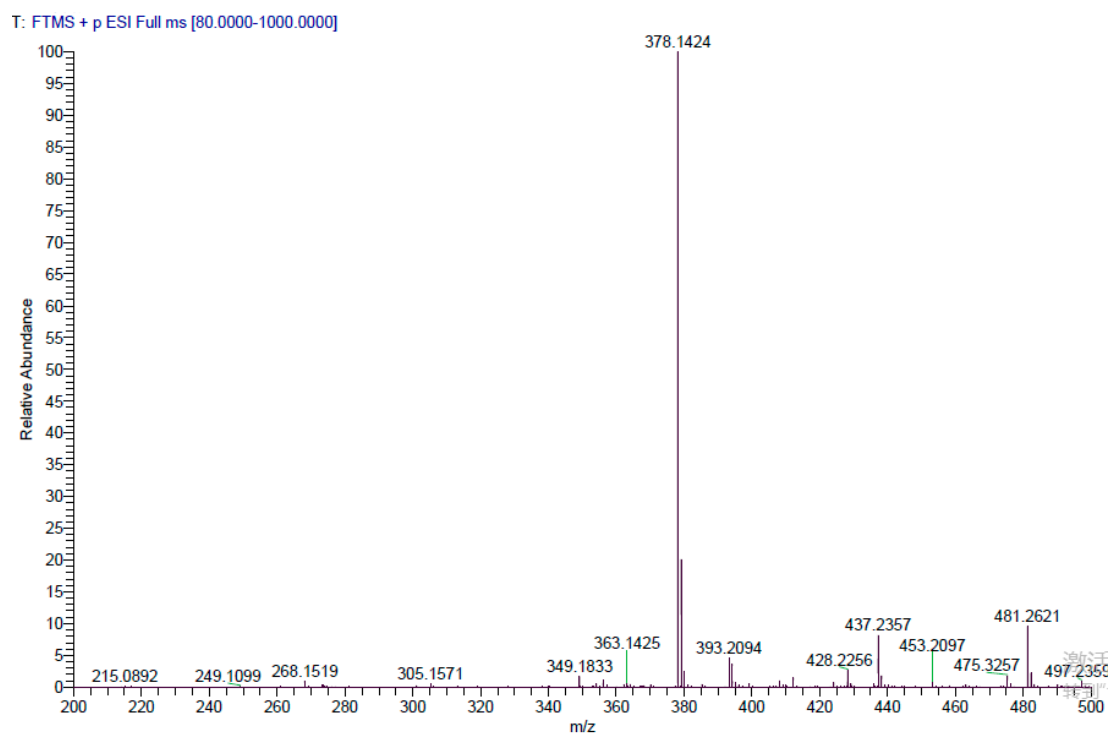


Figure S6. HRESIMS for compound **1**.

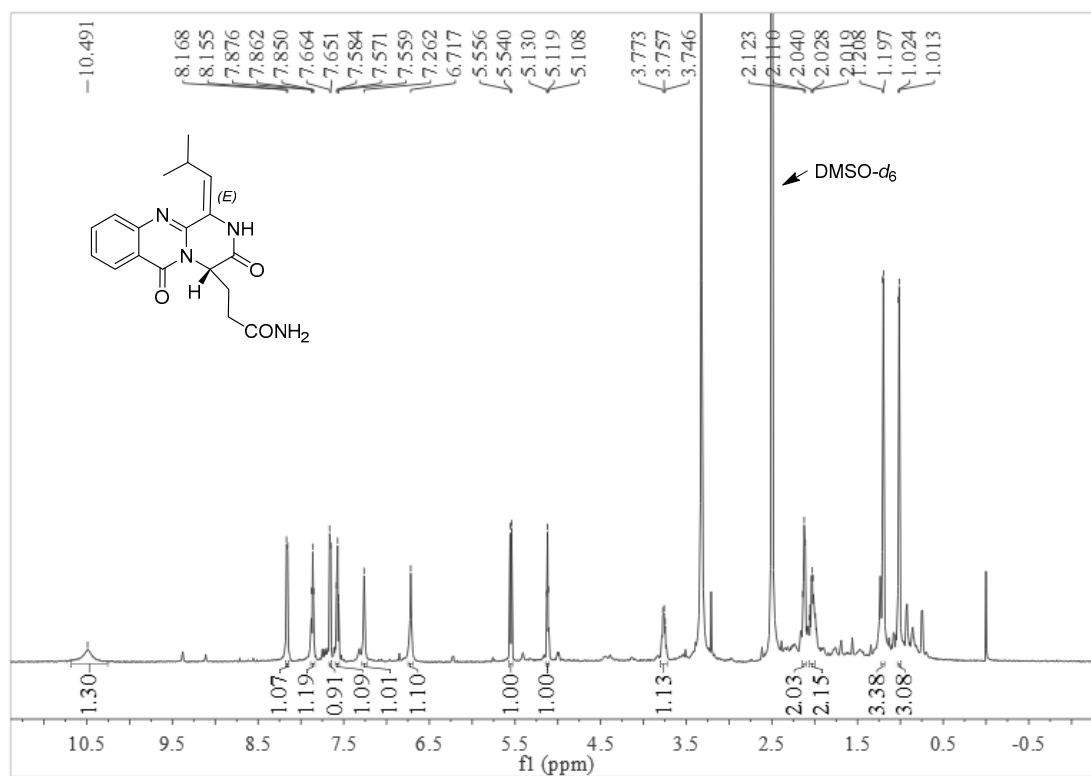


Figure S7. <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of 2

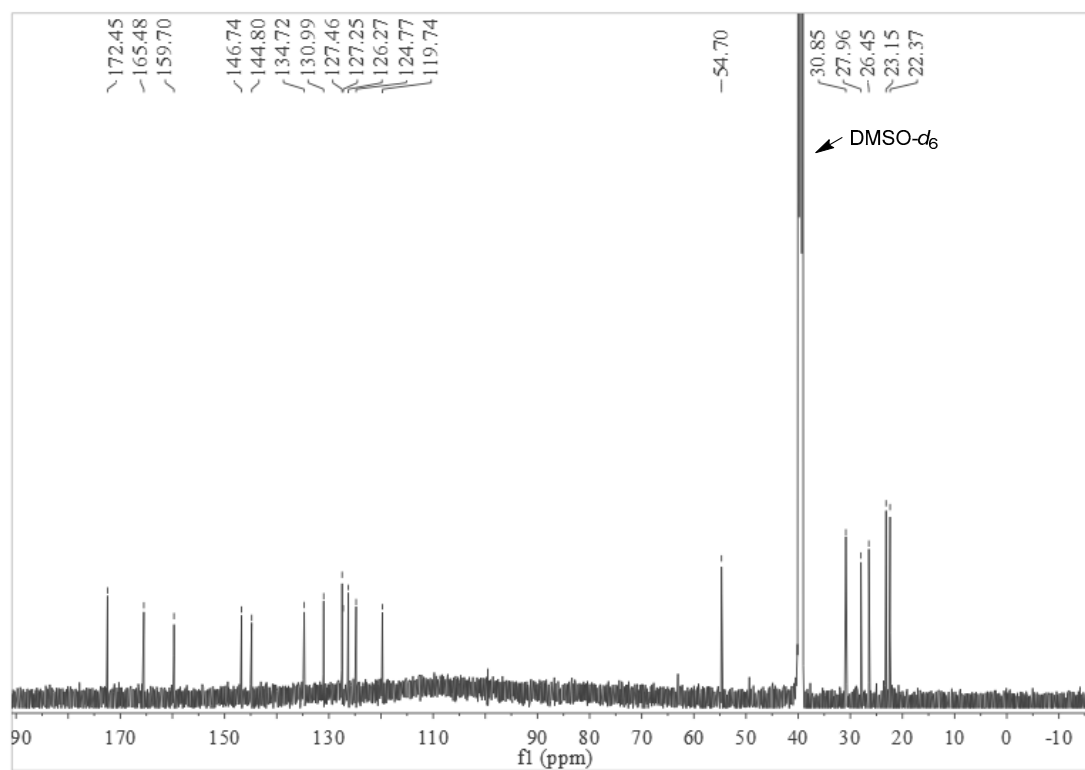
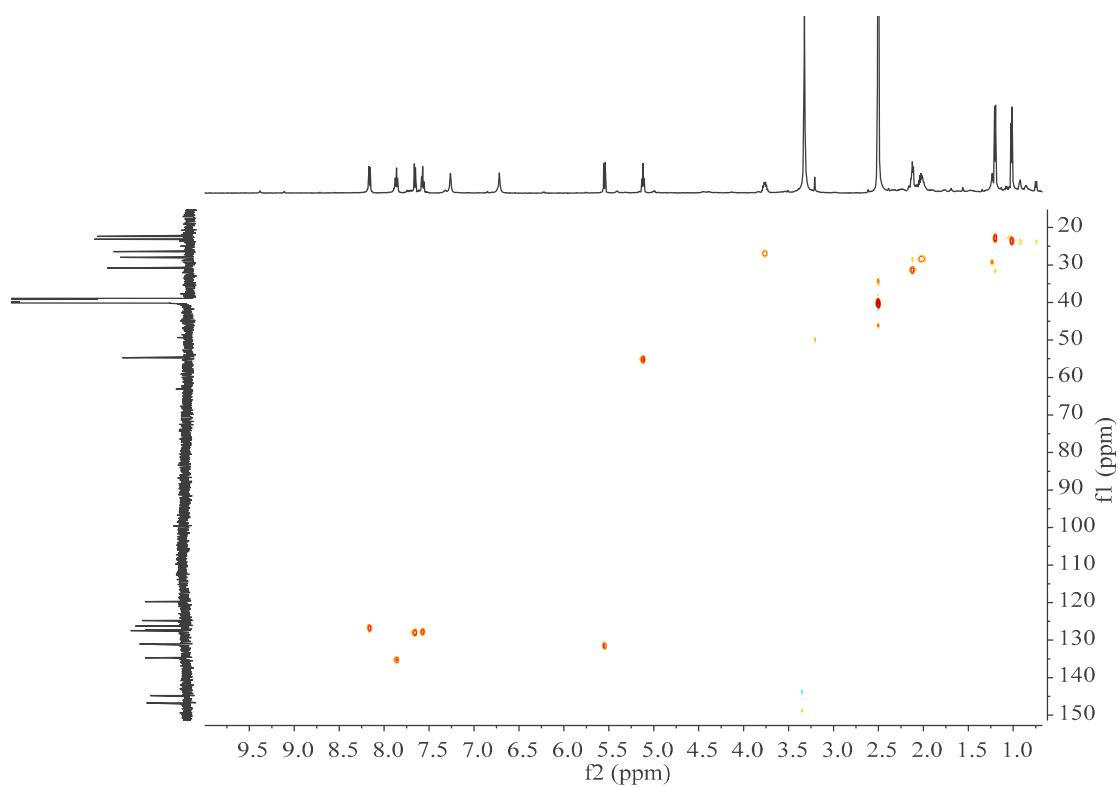
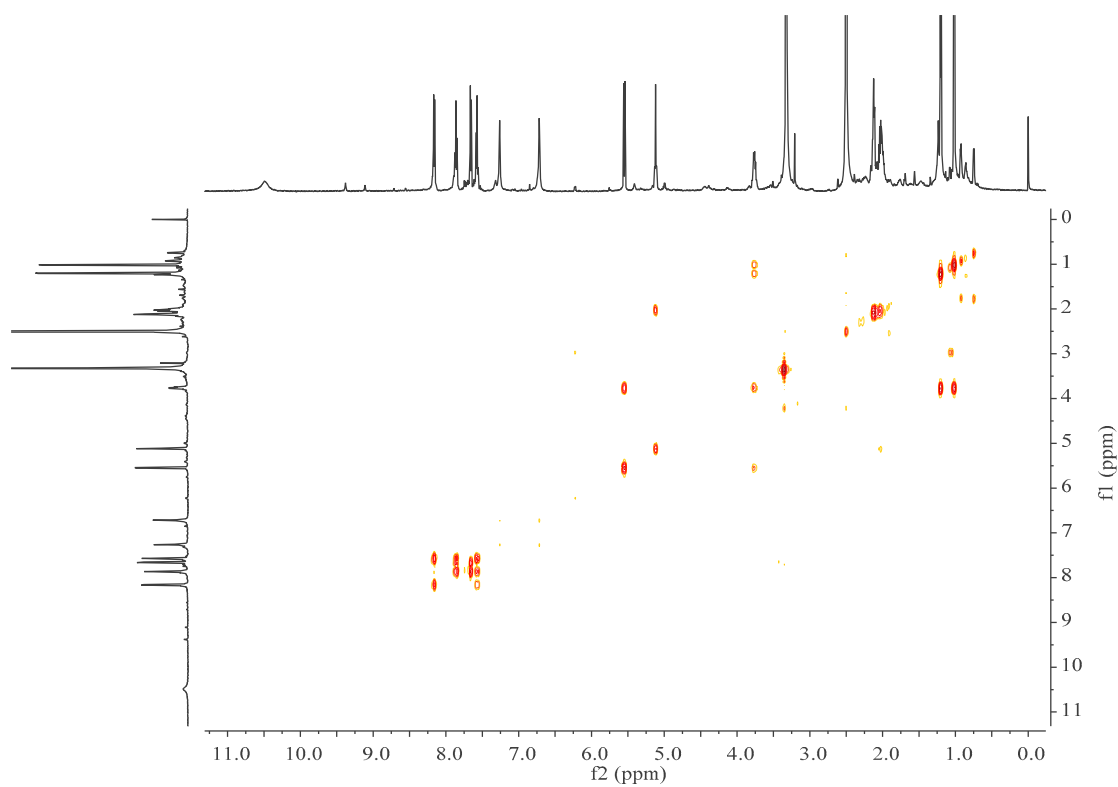


Figure S8. <sup>13</sup>C NMR (150 MHz, DMSO-*d*<sub>6</sub>) spectrum of 2.

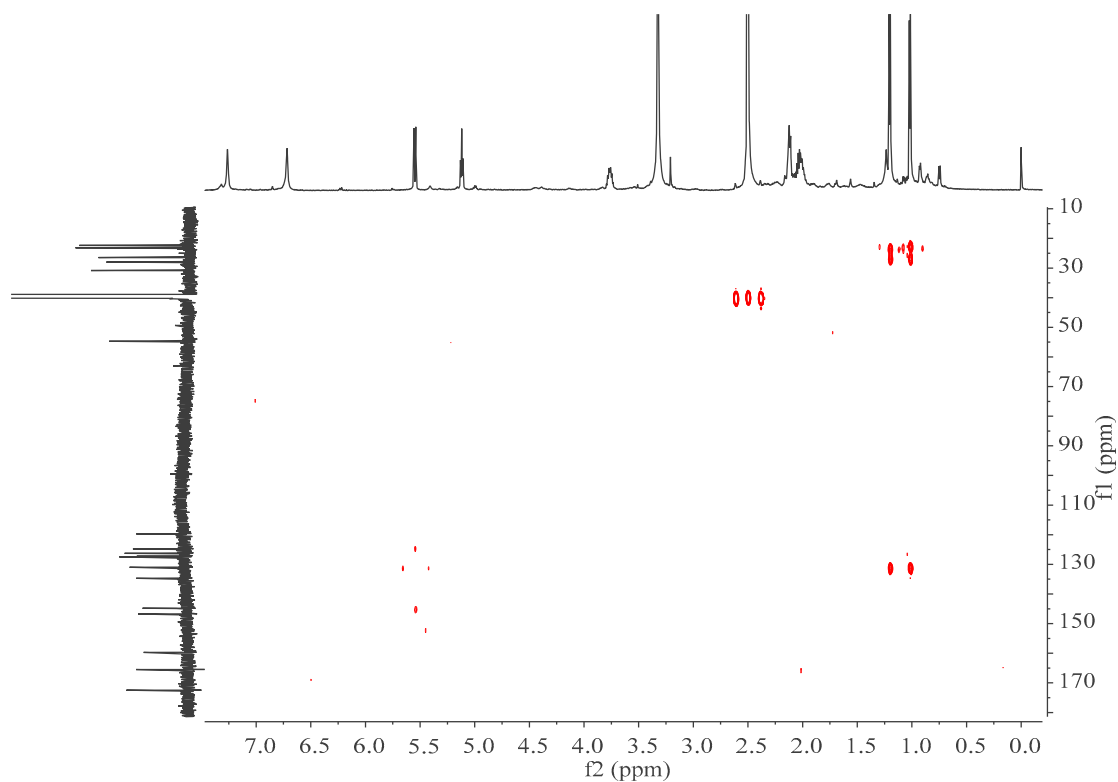


**Figure S9.** HSQC (DMSO-*d*<sub>6</sub>) spectrum of **2**.

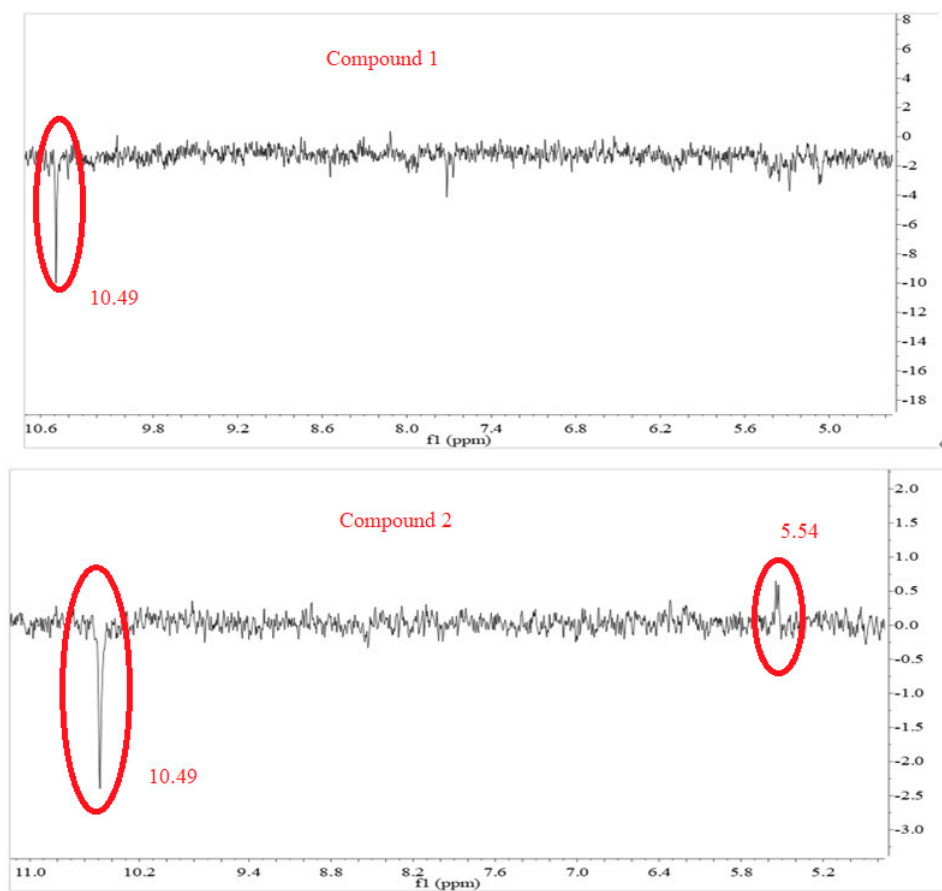


**Figure S10.** <sup>1</sup>H-<sup>1</sup>H COSY (DMSO-*d*<sub>6</sub>) spectrum of **2**.





**Figure S11.** HMBC (DMSO-*d*<sub>6</sub>) spectrum of **2**.



**Figure S12.** 1D NOE (DMSO-*d*<sub>6</sub>) spectrum of compounds **1** and **2**.

T: FTMS + p ESI Full ms [133.4000-2000.0000]

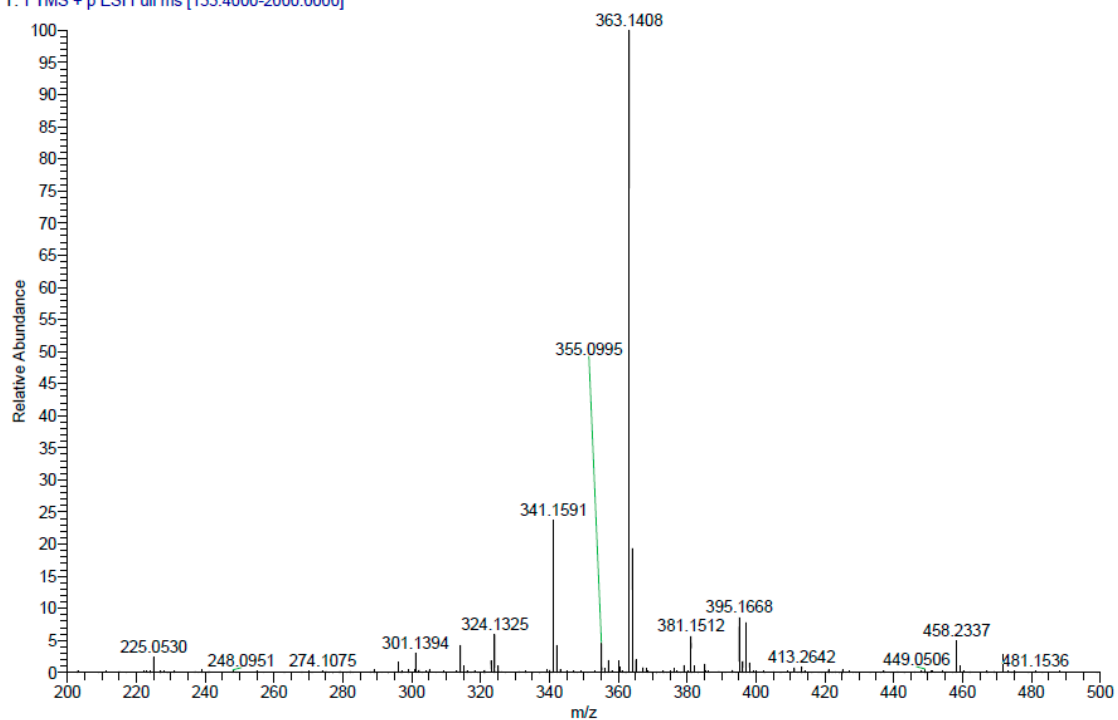
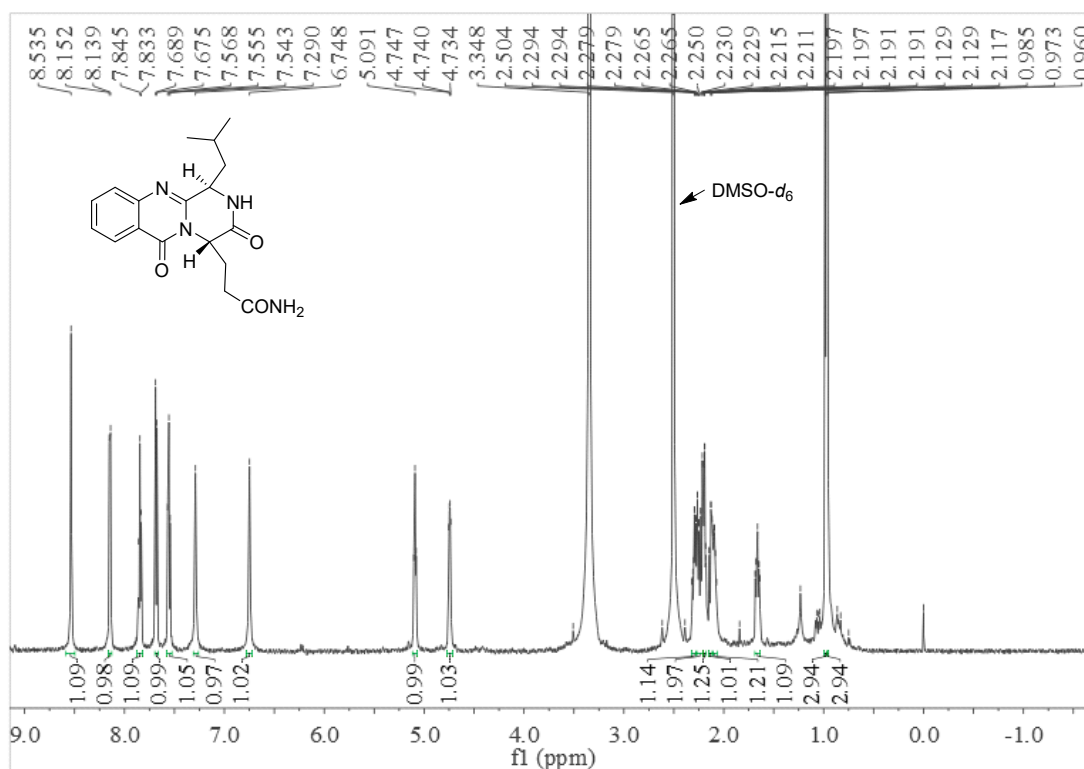
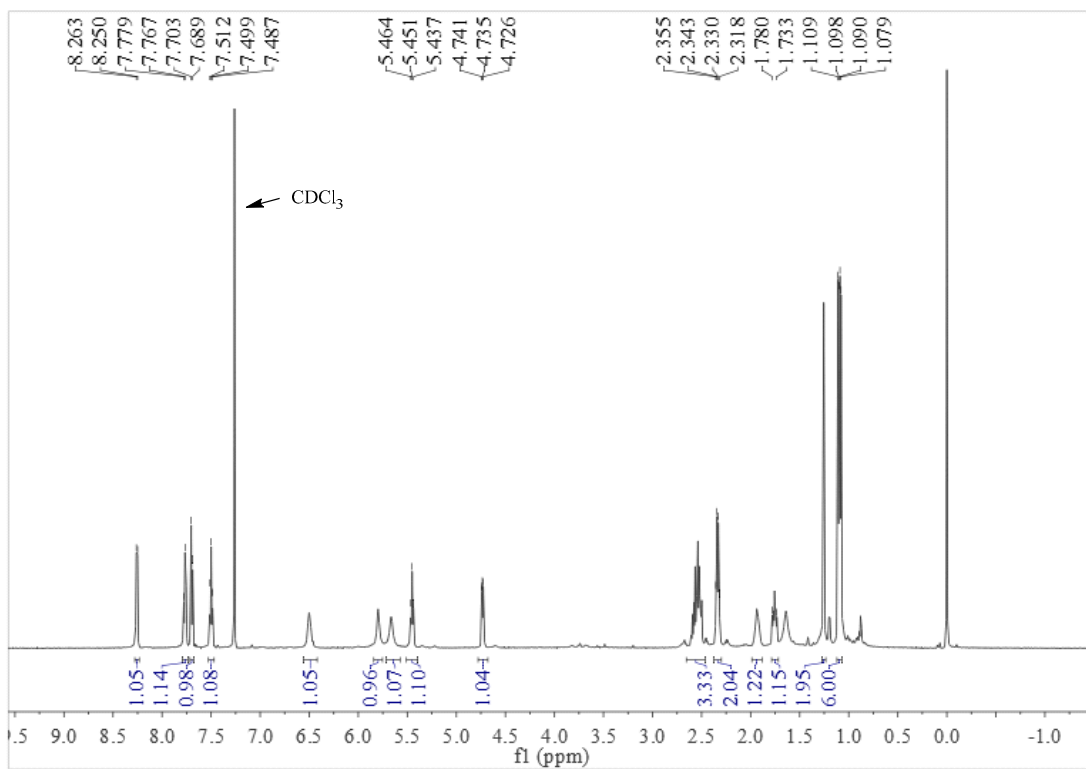
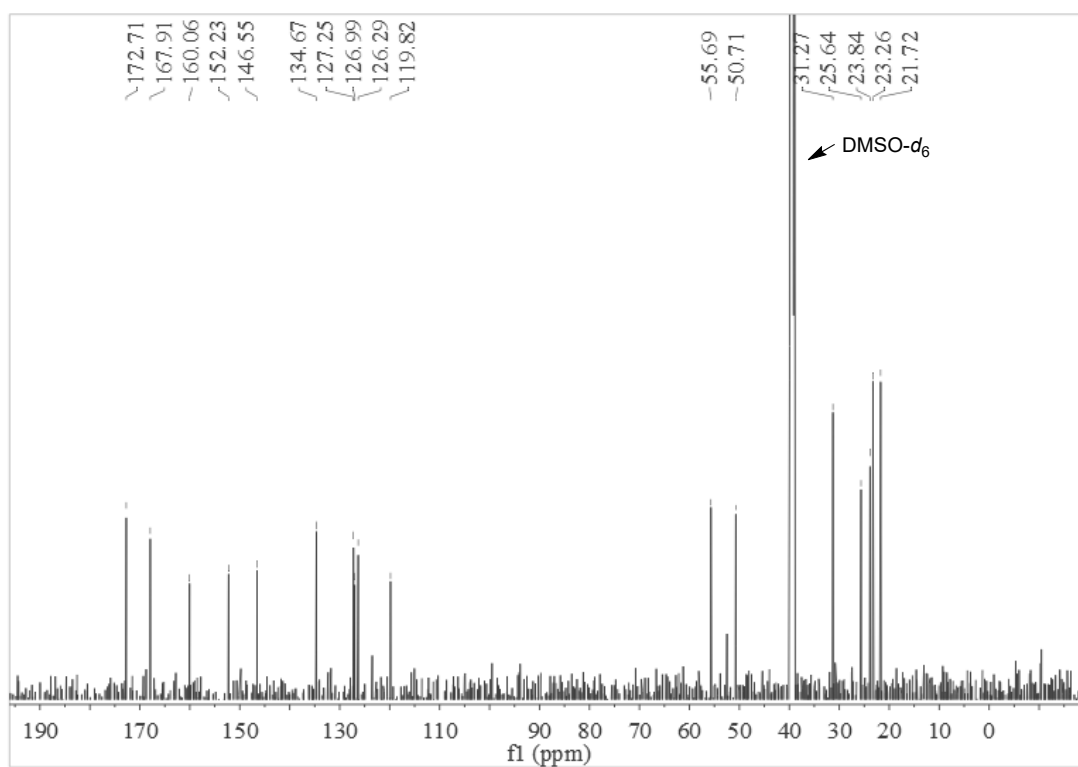


Figure S13. HRMSIMS for compound 2.

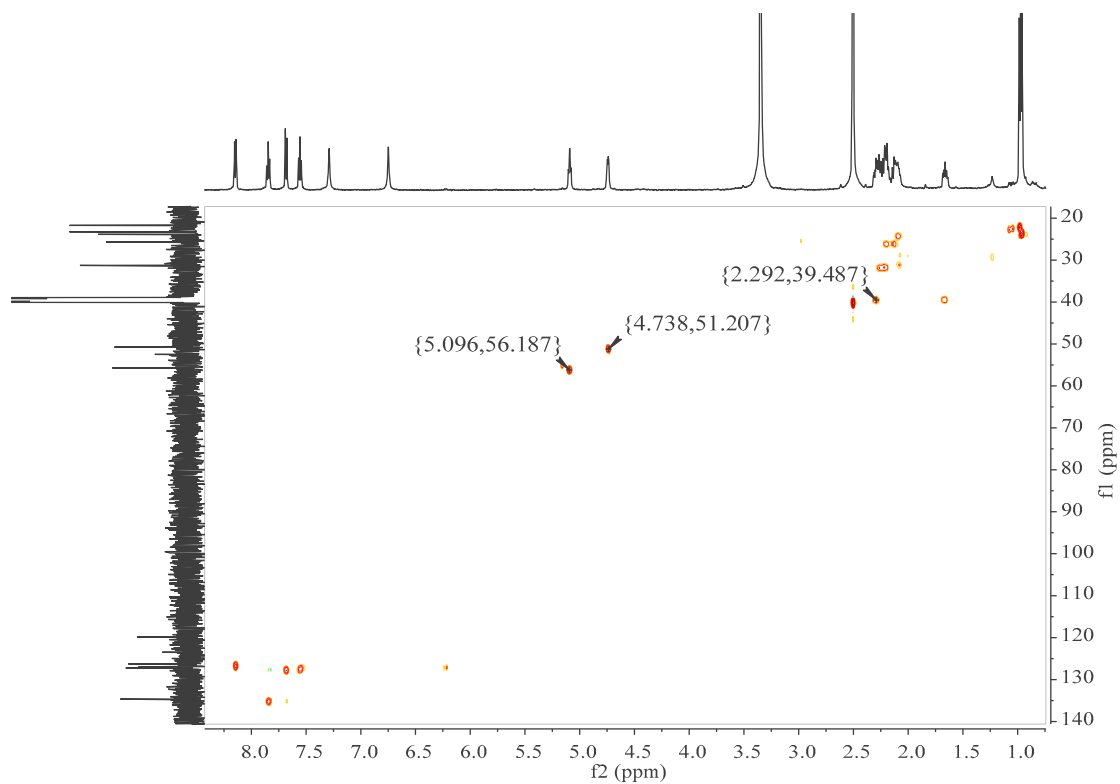
Figure S14. <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of 3.



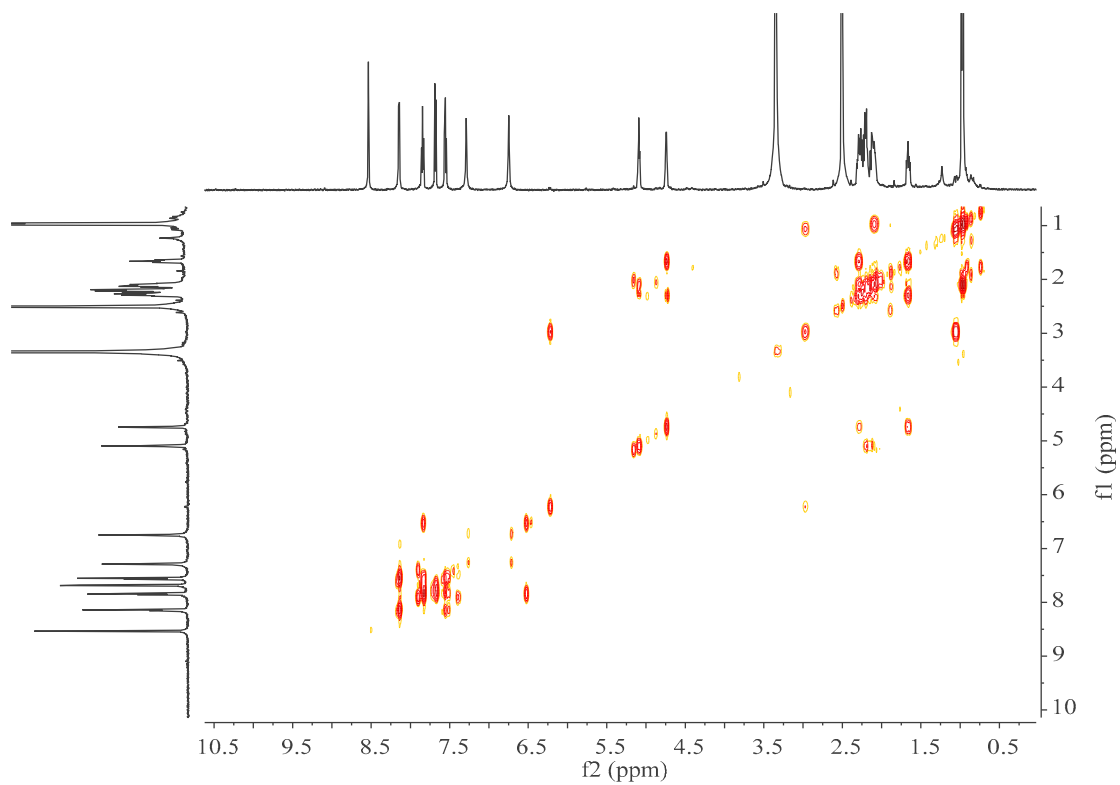
**Figure S15.** <sup>1</sup>H NMR (600 MHz, CDCl<sub>3</sub>) spectrum of **3**.



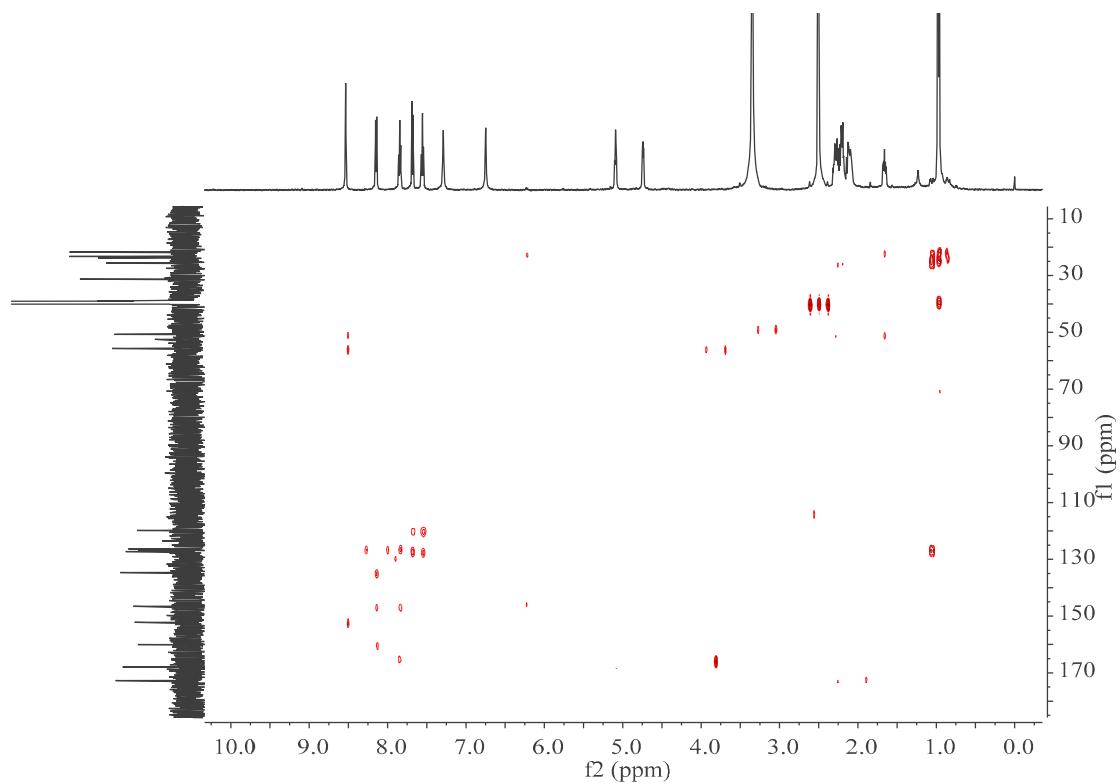
**Figure S16.** <sup>13</sup>C NMR (150 MHz, DMSO-*d*<sub>6</sub>) spectrum of **3**.



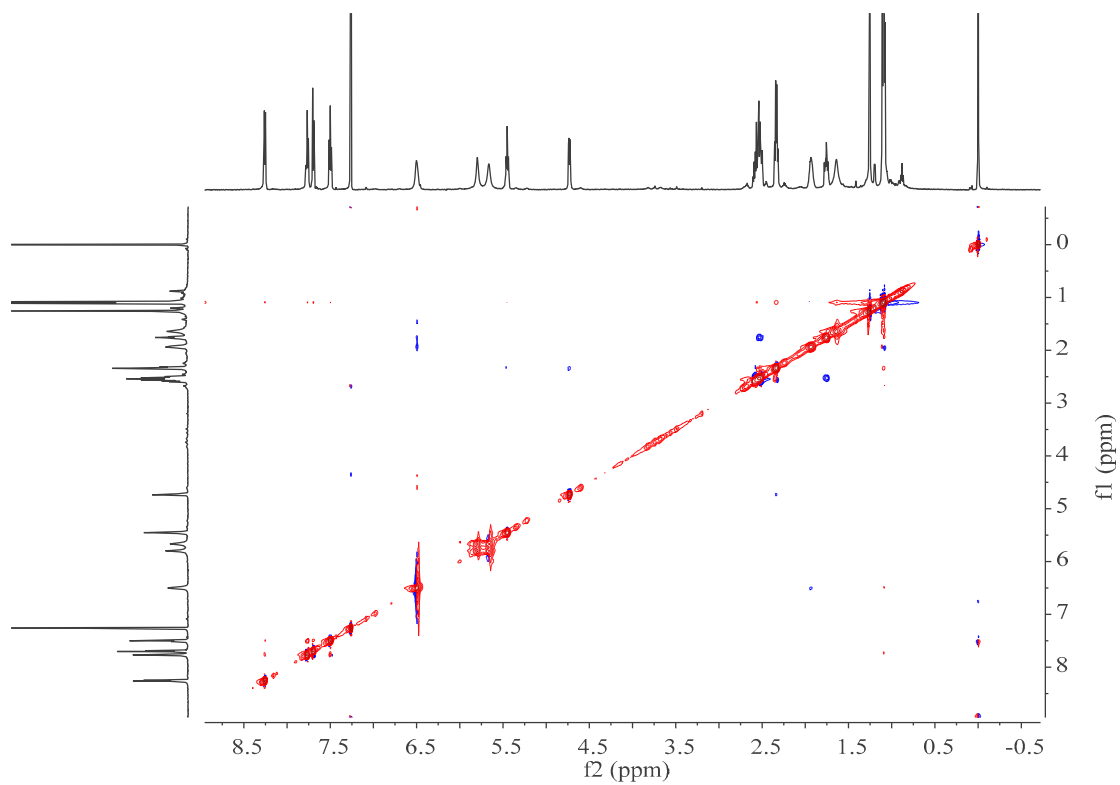
**Figure S17.** HSQC (DMSO-*d*<sub>6</sub>) spectrum of **3**.



**Figure S18.** <sup>1</sup>H-<sup>1</sup>H COSY (DMSO-*d*<sub>6</sub>) spectrum of **3**.



**Figure S19.** HMBC (DMSO-*d*<sub>6</sub>) spectrum of **3**.



**Figure S20.** NOESY (CDCl<sub>3</sub>) spectrum of compound **3**.

T: FTMS + p ESI Full ms [133.4000-2000.0000]

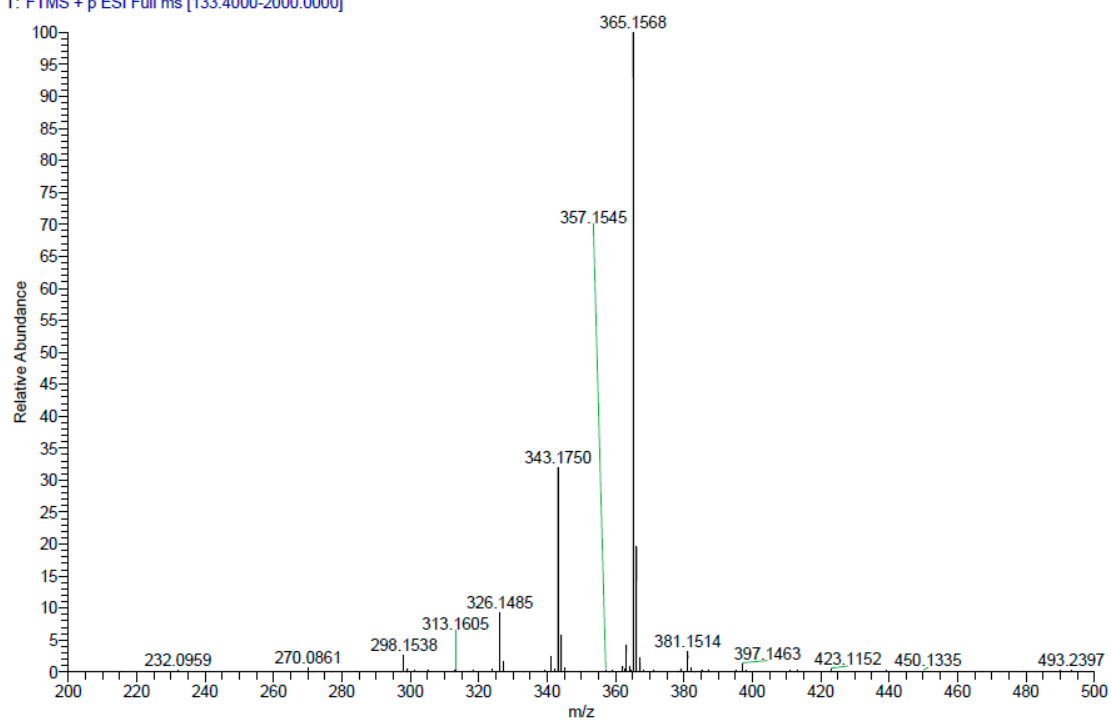
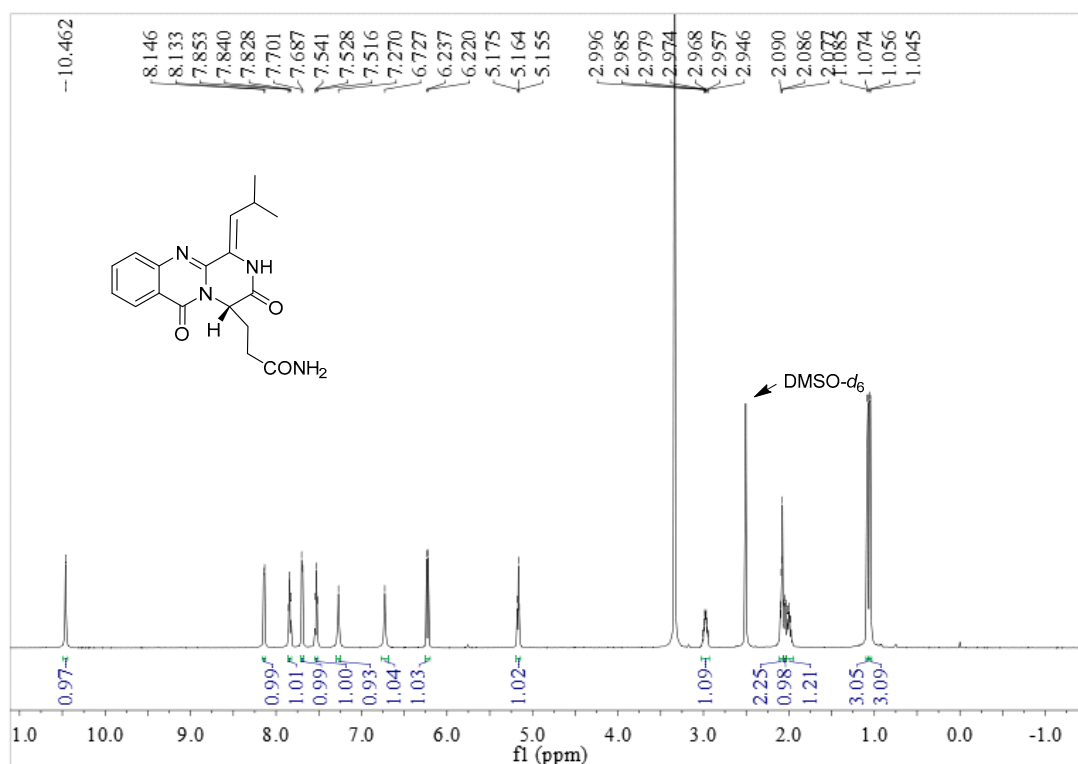
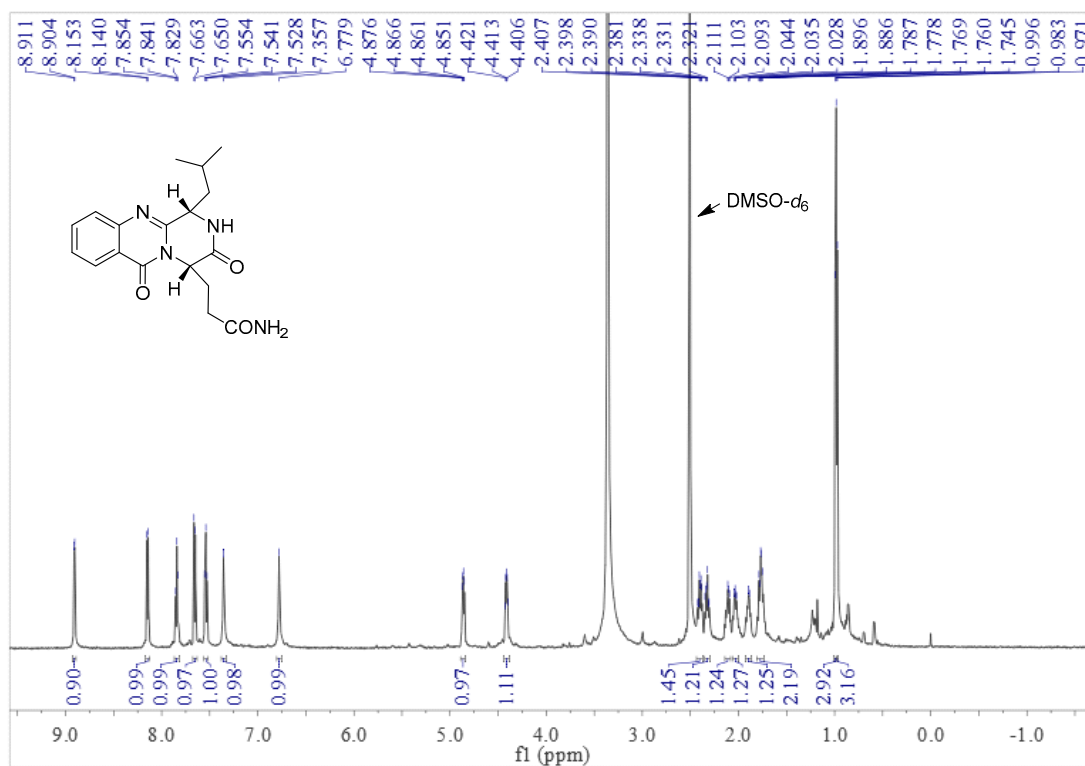
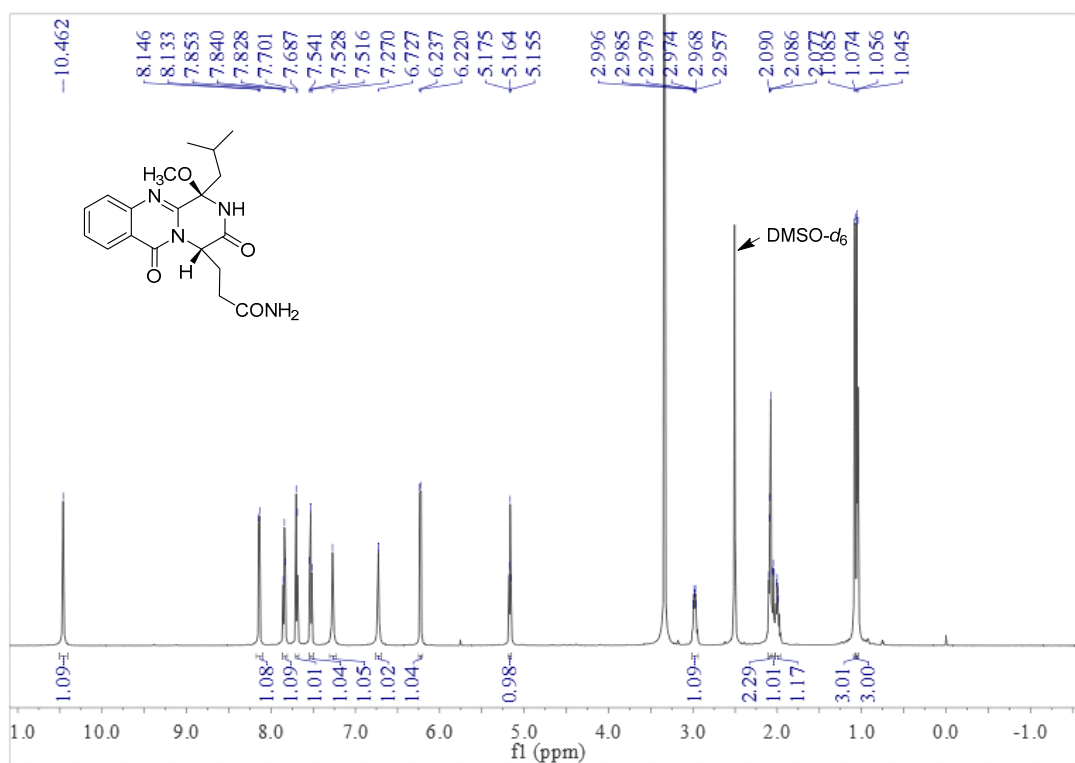


Figure S21. HRESIMS for compound 3.

Figure S22. <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of 4.



**Figure S23.**  $^1\text{H}$  NMR (600 MHz,  $\text{DMSO-}d_6$ ) spectrum of **5**.



**Figure S24.**  $^1\text{H}$  NMR (600 MHz,  $\text{DMSO-}d_6$ ) spectrum of **6**.

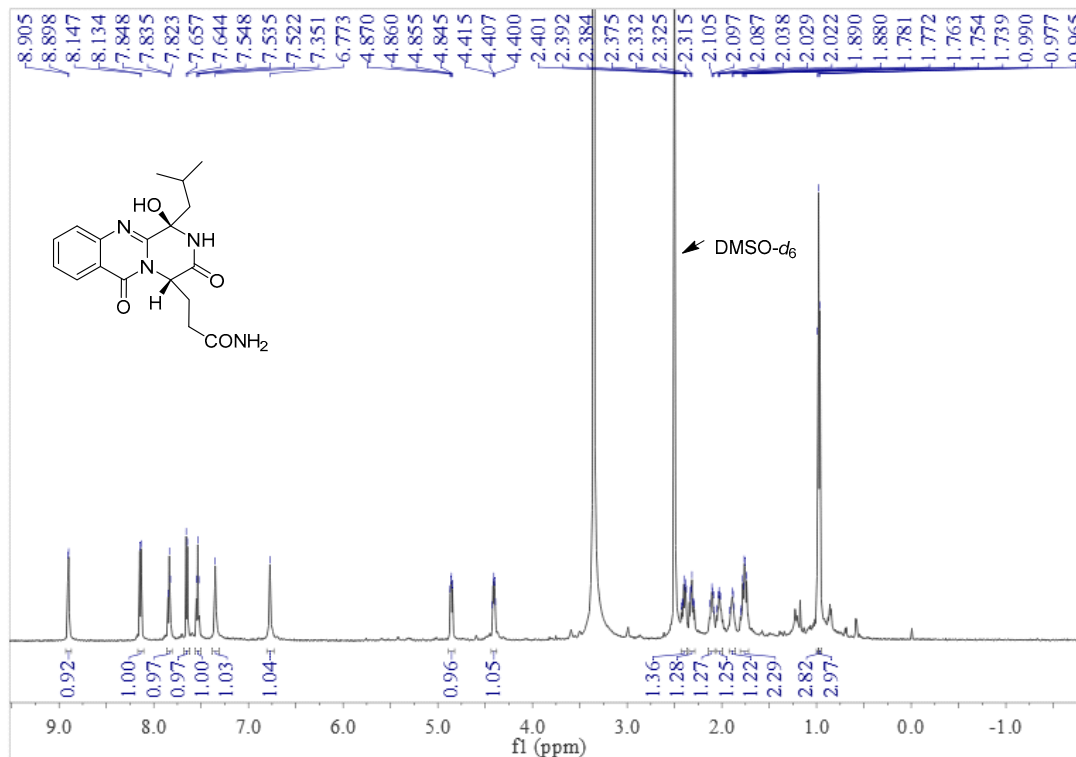


Figure S25. <sup>1</sup>H NMR (600 MHz, DMSO-*d*<sub>6</sub>) spectrum of 7.

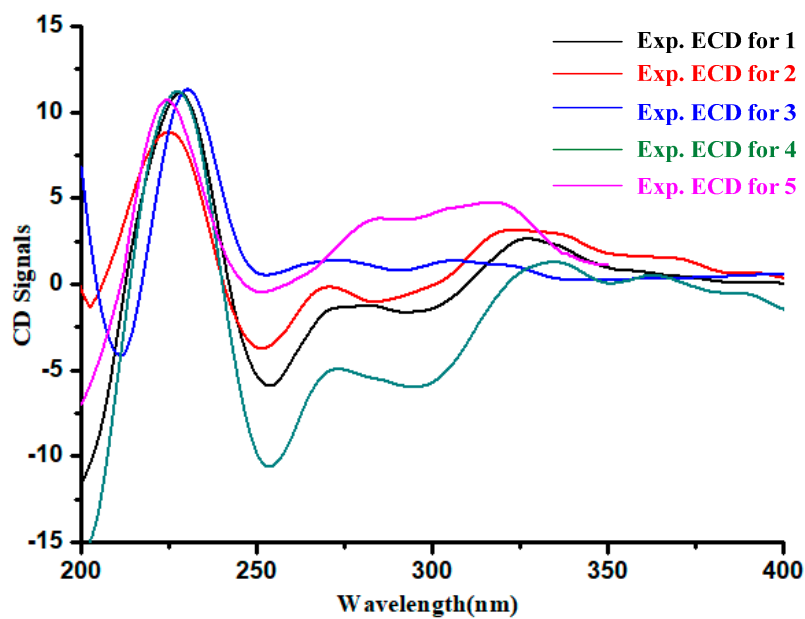


Figure S26. Experimental ECD spectra of 1–5.



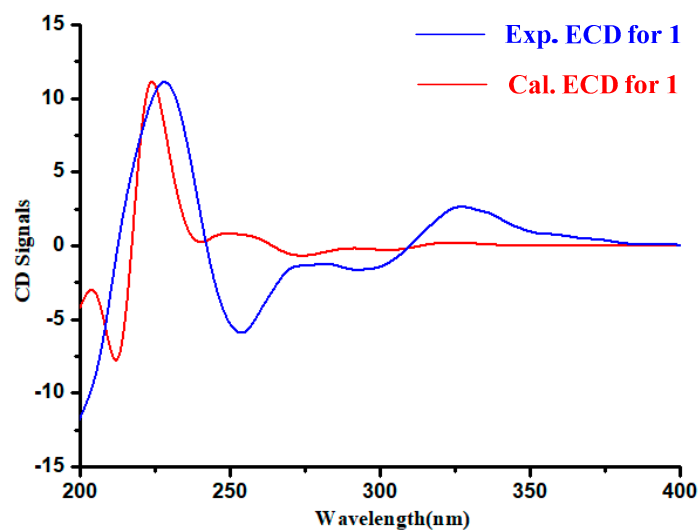


Figure S27. Experimental and calculated ECD spectra of 1.

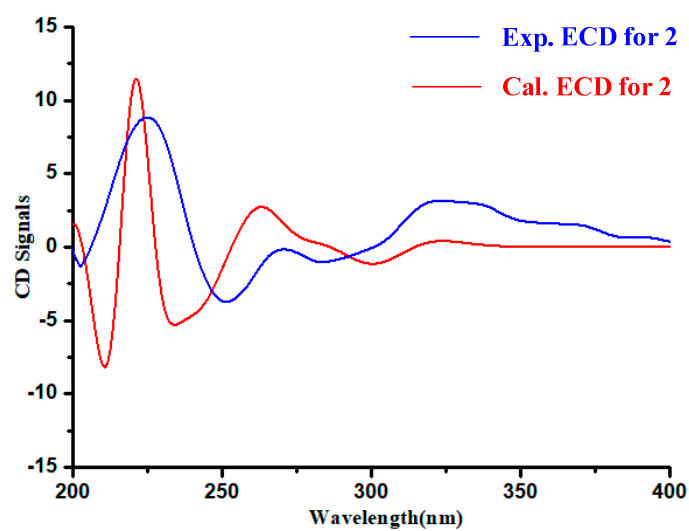


Figure S28. Experimental and calculated ECD spectra of 2.

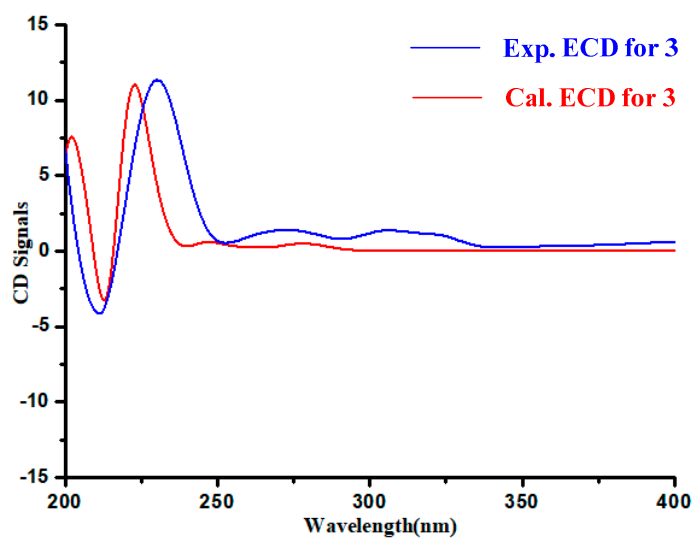
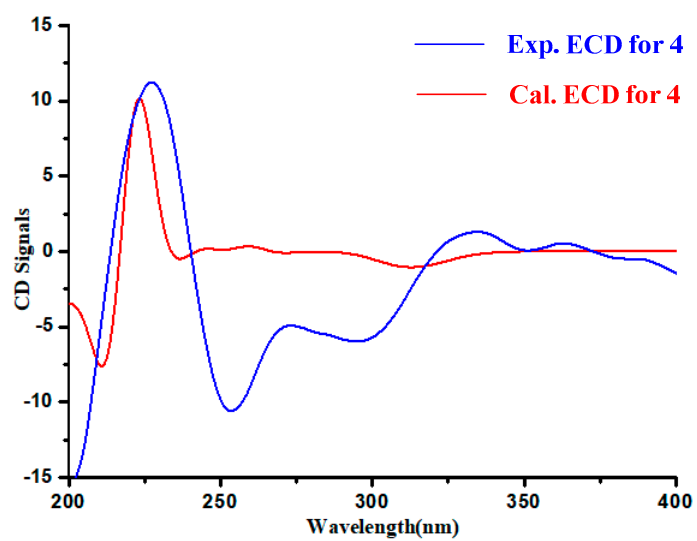
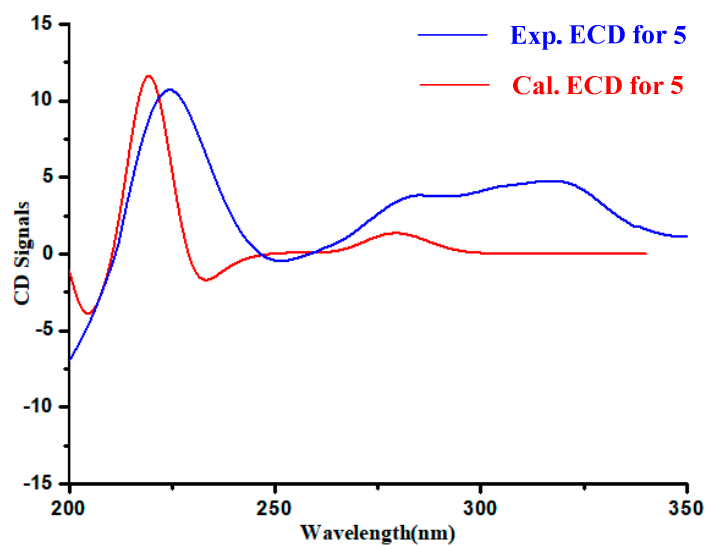


Figure S29. Experimental and calculated ECD spectra of 3.



**Figure S30.** Experimental and calculated ECD spectra of 4.



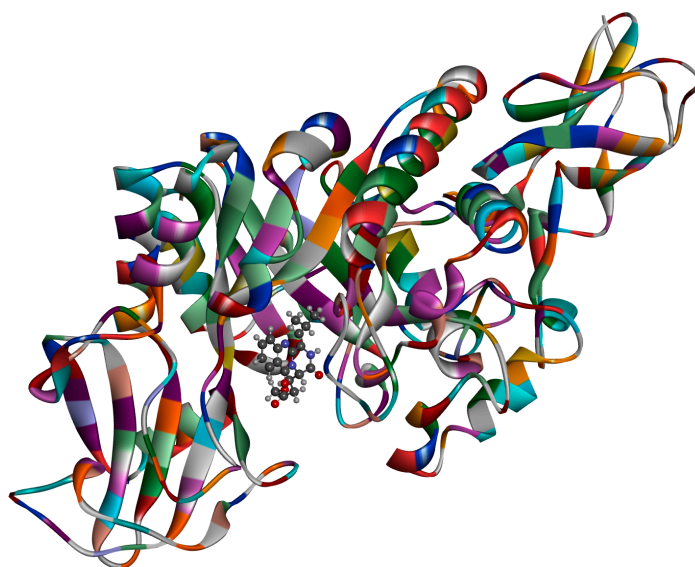
**Figure S31.** Experimental and calculated ECD spectra of 5.

Functional B3LYP		Solvent? Gas Phase		Basis Set 6-311+G(d,p)	
		DP4+	100.00%	0.00%	-
Nuclei	sp2?	Experimental	Isomer 1	Isomer 2	Isomer 3
C	x	134.7	134.2420294	134.3632431	
C	x	127	126.4808853	126.171998	
C	x	126.3	128.4799392	128.3022902	
C	x	119.8	121.2847373	120.6404755	
C	x	146.6	148.4599039	148.7865657	
C	x	127.3	128.5309245	128.0625951	
C	x	160.1	159.654048	159.6477137	
C	x	152.2	152.921151	152.6898206	
C		55.7	56.4478451	55.7540951	
C	x	167.9	167.820049	166.9695422	
C		50.7	51.55953235	57.17189902	
C		39	38.68519118	46.50892157	
C		23.8	27.0402049	26.99855392	
C		21.7	17.36431176	20.85926961	
C		23.3	22.18061765	18.03609902	
C		25.6	26.0581402	29.29893529	
C		31.3	31.09409118	31.59072451	
C	x	172.7	171.0076245	171.0901745	
H		5.09	4.479535981	3.94216221	
H	x	7.68	7.385941048	7.283663257	
H	x	7.85	7.386634648	7.247590695	
H	x	7.56	7.133378952	6.986158467	
H	x	8.15	8.125107962	7.984918019	
H		4.74	5.040505524	4.7519782	
H		2.2	2.011418514	2.0901142	
H		2.24	2.109524038	2.035519295	
H		2.29	2.237236648	1.772153333	
H		1.66	1.432108762	1.504773133	
H		0.98	0.998012781	0.736730771	
H		0.97	0.967706933	0.833866333	

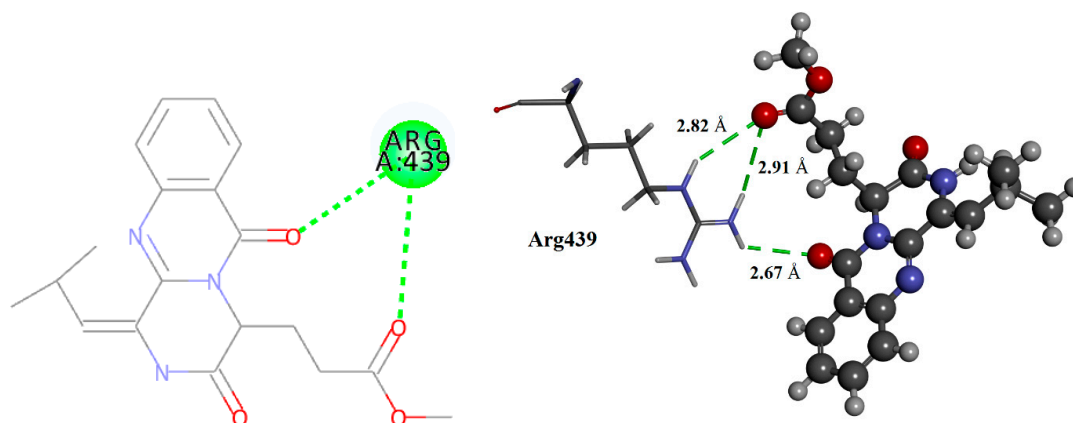
**Figure S32.** The data of DP4plus method of compound **3** (B3lyp/6-311+G(d,p))

Functional		Solvent?		Basis Set	
B3LYP		Gas Phase		6-311+G(d,p)	
		DP4+	0.00%	100.00%	-
Nuclei	sp2?	Experimental	Isomer 1	Isomer 2	Isomer 3
C	x	134.7	134.2420294	134.3632431	
C	x	126.7	126.4808853	126.171998	
C	x	126.2	128.4799392	128.3022902	
C	x	119.7	121.2847373	120.6404755	
C	x	147	148.4599039	148.7865657	
C	x	126.7	128.5309245	128.0625951	
C	x	160.1	159.654048	159.6477137	
C	x	152	152.921151	152.6898206	
C		54.9	56.4478451	55.7540951	
C	x	166.6	167.820049	166.9695422	
C		53.8	51.55953235	57.17189902	
C		47.2	38.68519118	46.50892157	
C		24	27.0402049	26.99855392	
C		21.4	17.36431176	20.85926961	
C		23	22.18061765	18.03609902	
C		29.4	26.0581402	29.29893529	
C		32.2	31.09409118	31.59072451	
C	x	172.8	171.0076245	171.0901745	
H		4.41	4.479535981	3.94216221	
H	x	7.66	7.385941048	7.283663257	
H	x	7.84	7.386634648	7.247590695	
H	x	7.54	7.133378952	6.986158467	
H	x	8.15	8.125107962	7.984918019	
H		4.86	5.040505524	4.7519782	
H		2.06	2.011418514	2.0901142	
H		2.35	2.109524038	2.035519295	
H		1.76	1.50710861	1.772153333	
H		1.9	1.432108762	1.504773133	
H		0.97	0.998012781	0.736730771	
H		0.99	0.967706933	0.833866333	

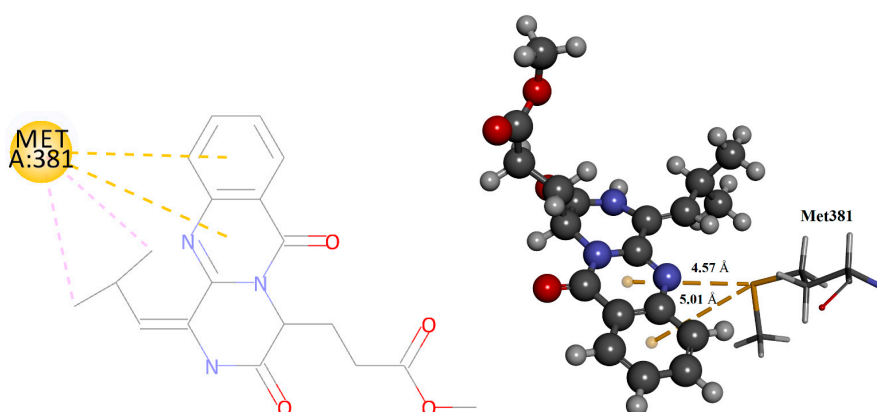
**Figure S33.** The data of DP4plus method of compound **5** (B3lyp/6-311+G(d,p))



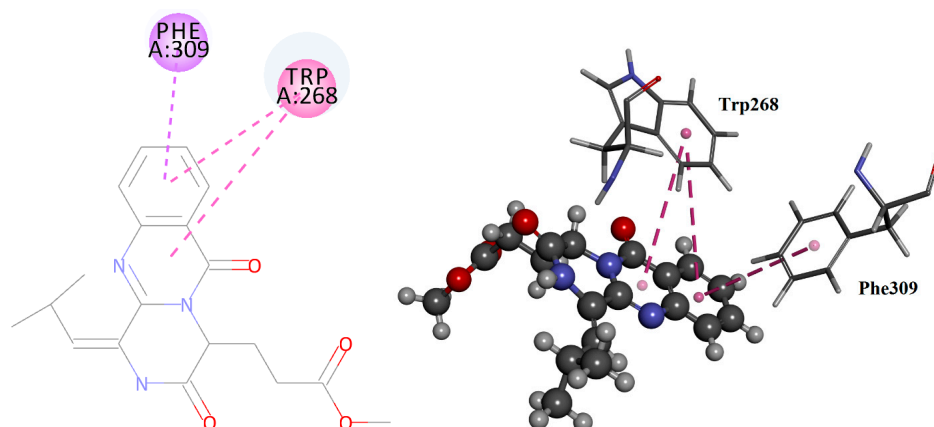
**Figure S34.** Compound 1 tightly bind to the entire active pocket of *OfChi-h*



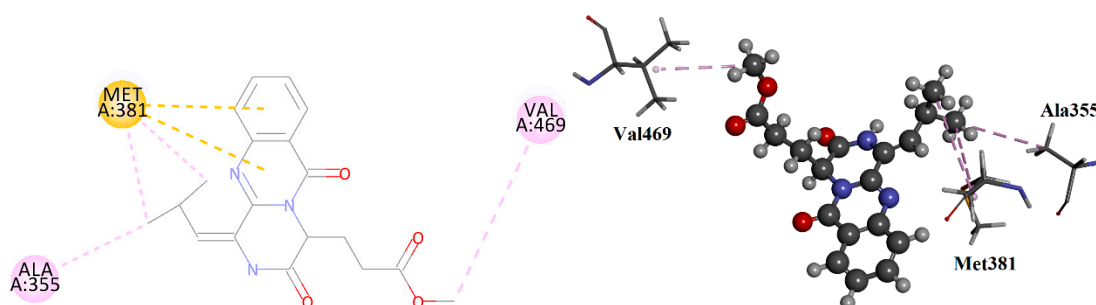
**Figure S35.** Three hydrogen bonds formed by 1 with the guanidine group of ARG439



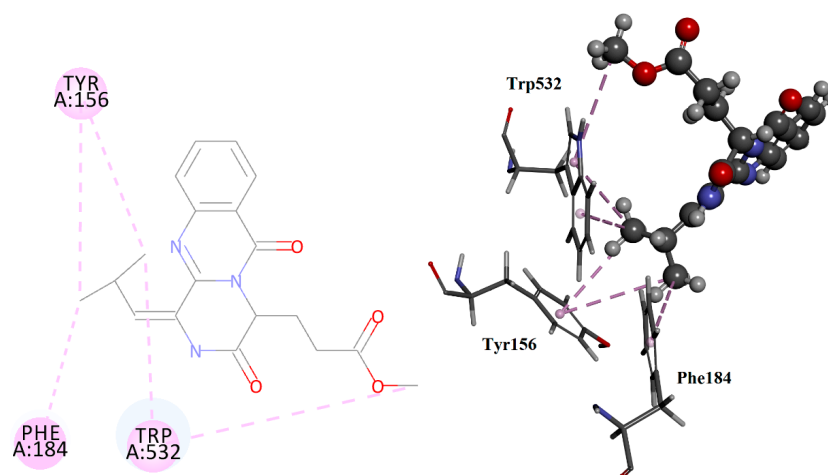
**Figure S36.**  $\pi$ -Sulfur interaction between 1 and the sulfur atom of methionine MET381



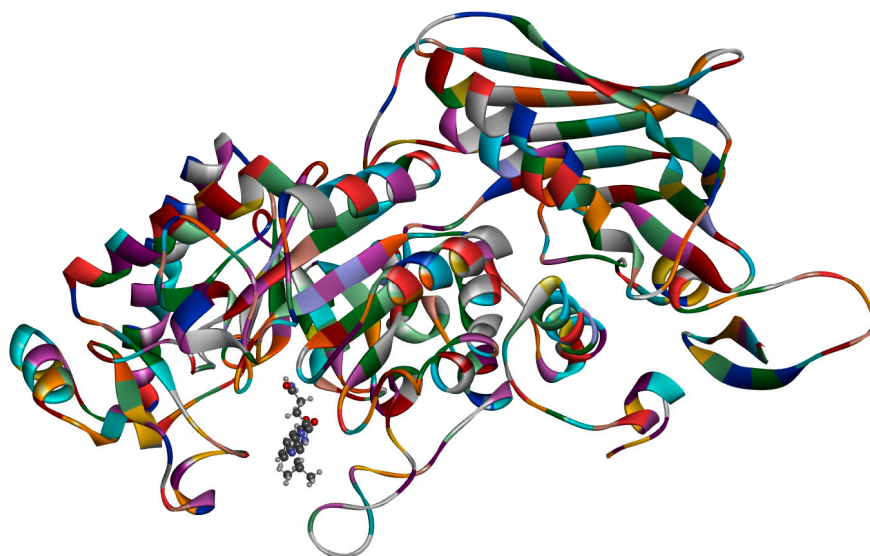
**Figure S37.**  $\pi$ - $\pi$  Stacking between **1** and the benzene rings of Trp268 and Phe309



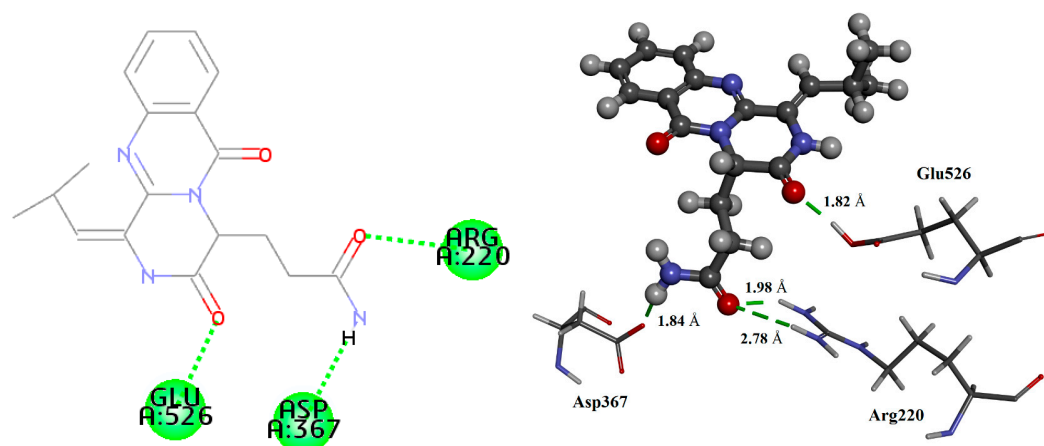
**Figure S38.** Alkyl hydrophobic interactions between **1** and Ala355/Met381, and between **1** and Val469



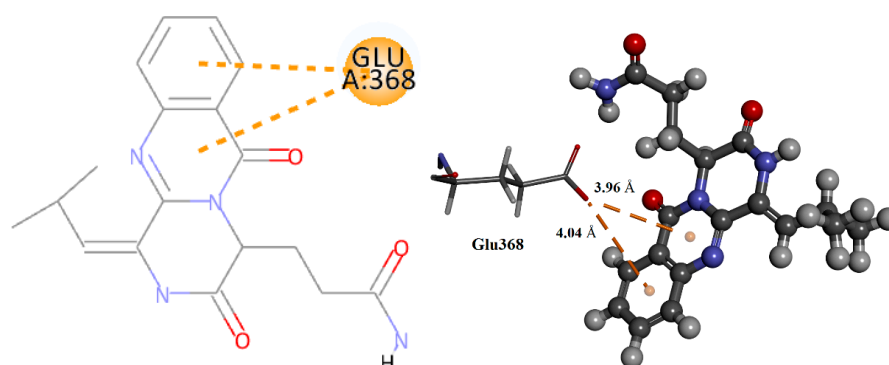
**Figure S39.** Mixed  $\pi$ /alkyl hydrophobic interactions between **1** and Tyr156, and between **1** and Phe184



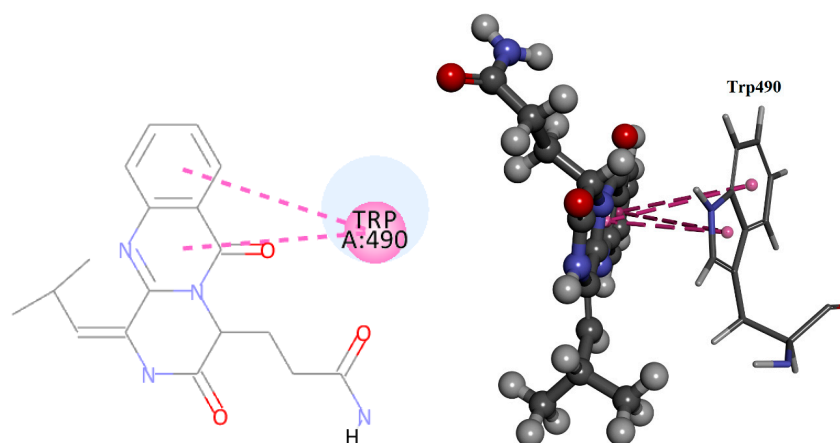
**Figure S40.** Compound 4 tightly bind to the *OfHex1* in a "U" conformation



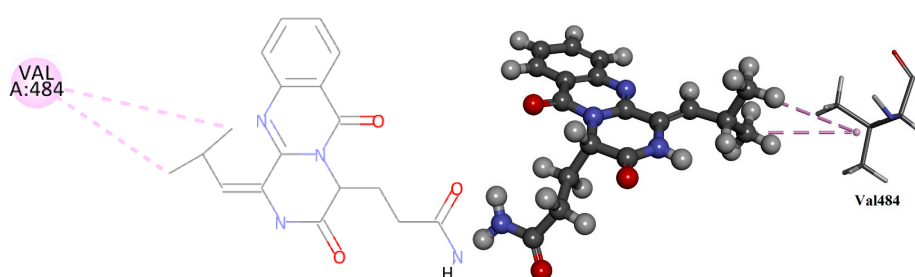
**Figure S41.** Hydrogen bonds formed between 4 and the guanidine group of ARG220, and between 4 and the carboxyl of ASP367



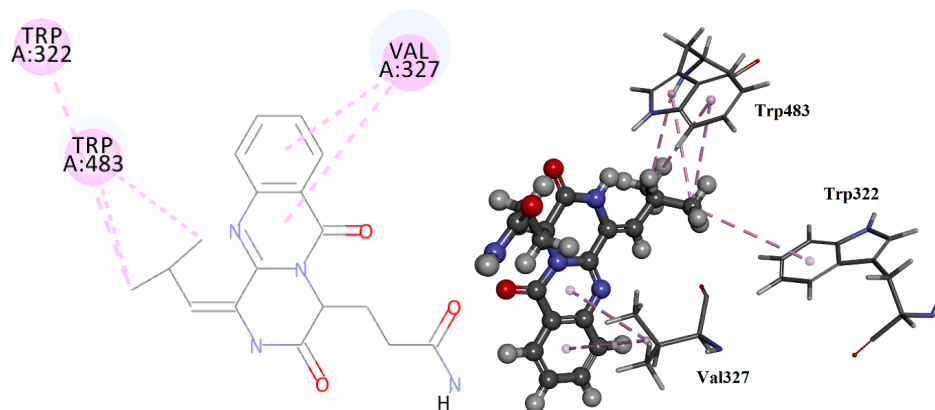
**Figure S42.** Compound 4 had a  $\pi$ -anion with the carboxyhydroxyl oxygen anion in the residue of GLU368



**Figure S43.**  $\pi$ - $\pi$  Stacking interaction between **4** and the indole ring of Trp490

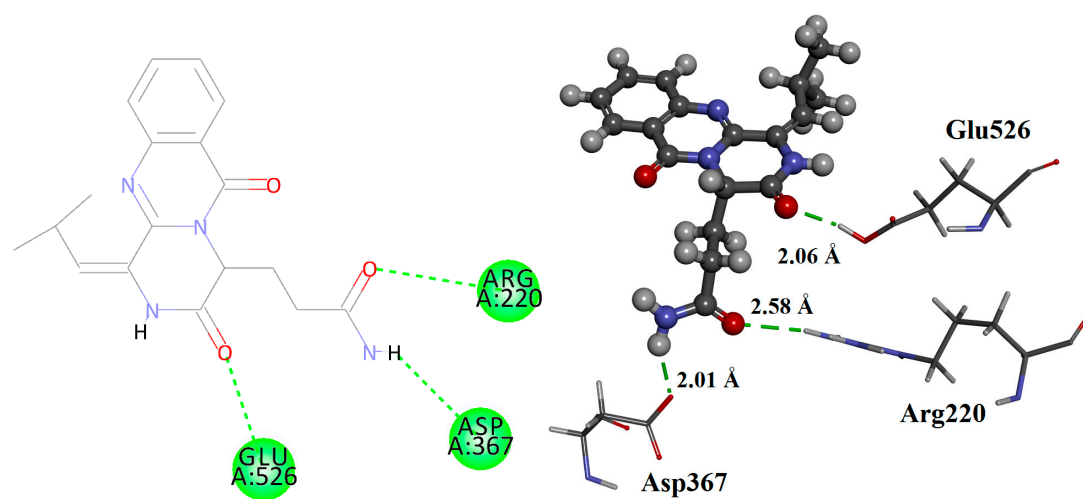


**Figure S44.** Alkyl hydrophobic interaction between **4** and the isopropyl group of Val484



**Figure S45.** Mixed  $\pi$ /alkyl hydrophobic interactions between **4** and Trp322/Trp483, and between **4** and the isopropyl group of Val327





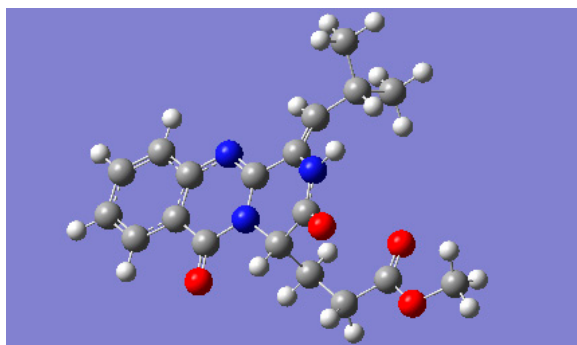
**Figure S46.** N-H hydrogen bond interaction between **2** and Arg220

**Table S1.** The data of DP4plus method of compound **3** and **5**

Nuclei	3-Exp.	3-Cal.3R,14S	5-Exp.	5-Cal.3S,14S
C-8	134.7	134.242	134.7	134.3632
C-9	127	126.4809	126.7	126.172
C-10	126.3	128.4799	126.2	128.3023
C-11	119.8	121.2847	119.7	120.6405
C-6	146.6	148.4599	147	148.7866
C-7	127.3	128.5309	126.7	128.0626
C-12	160.1	159.654	160.1	159.6477
C-4	152.2	152.9212	152	152.6898
C-3	55.7	56.44785	54.9	55.7541
C-1	167.9	167.82	166.6	166.9695
C-14	50.7	51.55953	53.8	57.1719
C-18	39	38.68519	47.2	46.50892
C-19	23.8	27.0402	24	26.99855
C-21	21.7	17.36431	21.4	20.85927
C-20	23.3	22.18062	23	18.0361
C-15	25.6	26.05814	29.4	29.29894
C-16	31.3	31.09409	32.2	31.59072
C-17	172.7	171.0076	172.8	171.0902
H-3	4.74	4.479536	4.41	3.942162
H-7	7.68	7.385941	7.66	7.283663
H-8	7.85	7.386635	7.84	7.247591
H-9	7.56	7.133379	7.54	6.986158
H-10	8.15	8.125108	8.15	7.984918
H-14	5.09	5.040506	4.86	4.751978
H-15	2.18	2.011419	2.06	2.090114
H-16	2.24	2.109524	2.35	2.035519
H-18	2.29	2.237237	1.76	1.772153
H-19	2.1	1.432109	1.9	1.504773
H-21	0.98	0.998013	0.97	0.736731
H-20	0.97	0.967707	0.99	0.833866

**Table S2.** Cytotoxic activity data of compounds 1–7

Compd.	IC <sub>50</sub> (μM)		
	A549	HGC-27	UMUC-3
<b>1</b>	>10	>10	>10
<b>2</b>	>10	>10	>10
<b>3</b>	>10	>10	>10
<b>4</b>	>10	>10	>10
<b>5</b>	6.0	6.2	7.2
<b>6</b>	>10	>10	>10
<b>7</b>	>10	>10	>10
<b>DDP</b>	3.1	0.8	1.8

**Table S3.** The coordinate for the lowest-energy conformer of compound **1** for ECD calculation.

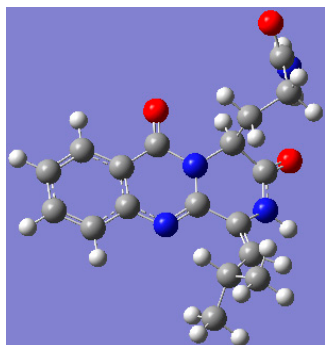
E=-1202.2385103 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.712213	0.122134	-0.606088
2	6	0	-5.645965	-1.266240	-0.390490
3	6	0	-4.436573	-1.857697	-0.073716
4	6	0	-3.279302	-1.070387	0.034140
5	6	0	-3.339313	0.323463	-0.177906
6	6	0	-4.580173	0.908676	-0.501551
7	6	0	-1.992437	-1.690273	0.337346
8	7	0	-0.913627	-0.776280	0.412868
9	6	0	-1.092404	0.584867	0.225586
10	7	0	-2.230805	1.133145	-0.058420
11	6	0	0.430214	-1.369961	0.579110
12	6	0	1.376786	-0.445246	1.325816
13	7	0	1.143582	0.887338	1.165696
14	6	0	0.103603	1.447800	0.397243
15	8	0	-1.803560	-2.885775	0.498808
16	6	0	0.140339	2.693743	-0.101104
17	8	0	2.304341	-0.881035	1.985534
18	1	0	0.299749	-2.256852	1.196618
19	6	0	0.992491	-1.801797	-0.798481
20	6	0	2.357860	-2.492331	-0.743409
21	6	0	1.264237	3.687003	-0.010664
22	6	0	1.688219	4.134896	-1.421630
23	6	0	0.847324	4.896934	0.848280
24	6	0	3.534107	-1.535705	-0.722251
25	8	0	3.489333	-0.363455	-1.014620
26	8	0	4.666246	-2.175304	-0.380948
27	6	0	5.858514	-1.373333	-0.327721

28	1	0	-6.663612	0.581403	-0.855196
29	1	0	-6.542868	-1.870545	-0.473671
30	1	0	-4.354487	-2.925381	0.093623
31	1	0	-4.617457	1.980625	-0.659409
32	1	0	1.847612	1.498563	1.555254
33	1	0	-0.758103	3.021264	-0.614381
34	1	0	0.261805	-2.493688	-1.221831
35	1	0	1.047143	-0.932914	-1.459399
36	1	0	2.441364	-3.173181	0.106446
37	1	0	2.483664	-3.114469	-1.637068
38	1	0	2.141522	3.224928	0.457642
39	1	0	2.502744	4.863072	-1.366713
40	1	0	2.030246	3.288062	-2.020747
41	1	0	0.853907	4.607786	-1.949472
42	1	0	1.663579	5.622213	0.916108
43	1	0	0.574508	4.597077	1.863421
44	1	0	-0.016215	5.407070	0.410556
45	1	0	6.061919	-0.923232	-1.299979
46	1	0	5.750380	-0.587837	0.420529
47	1	0	6.655781	-2.058121	-0.048581

---

**Table S4.** The coordinate for the lowest-energy conformer of compound **2** for ECD calculation

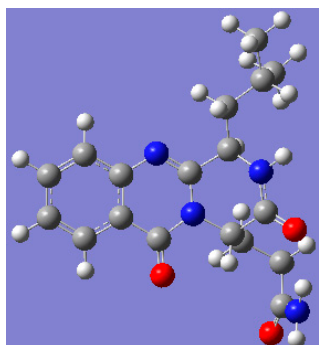
E=-1142.4900148 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	4.101073	-3.000198	0.171249
2	6	0	3.113465	-3.999989	0.137728
3	6	0	1.781447	-3.651130	0.001691
4	6	0	1.420462	-2.299242	-0.103004
5	6	0	2.406535	-1.291832	-0.071903
6	6	0	3.757627	-1.664517	0.067172
7	6	0	0.014292	-1.921530	-0.232966
8	7	0	-0.201420	-0.522696	-0.319282
9	6	0	0.848822	0.383222	-0.309801
10	7	0	2.094322	0.047957	-0.196846
11	6	0	-1.610643	-0.078121	-0.345740
12	6	0	-1.781794	1.156143	-1.215851
13	7	0	-0.721111	2.003217	-1.227797
14	6	0	0.488675	1.808635	-0.506974
15	8	0	-0.925740	-2.696024	-0.256802
16	6	0	1.210128	2.876729	-0.124906
17	8	0	-2.821239	1.391913	-1.818538
18	6	0	-2.135347	0.155366	1.094560
19	6	0	-3.597052	0.628500	1.196301
20	6	0	-4.610105	-0.421129	0.732658
21	8	0	-5.057393	-1.259271	1.499021
22	7	0	-4.966861	-0.344592	-0.579084
23	6	0	2.528293	2.949947	0.593127
24	6	0	3.662122	3.253946	-0.405647
25	6	0	2.463865	4.019048	1.696726
26	1	0	5.144368	-3.279542	0.277840
27	1	0	3.396873	-5.043740	0.219044

28	1	0	0.999088	-4.400622	-0.024472
29	1	0	4.509169	-0.883338	0.085208
30	1	0	-2.174321	-0.886788	-0.806001
31	1	0	-0.851486	2.872025	-1.729945
32	1	0	0.781728	3.847820	-0.380945
33	1	0	-1.491475	0.890619	1.588911
34	1	0	-2.020741	-0.787744	1.634579
35	1	0	-3.736671	1.567367	0.654263
36	1	0	-3.815296	0.819439	2.248377
37	1	0	-5.574283	-1.060959	-0.946441
38	1	0	-4.494585	0.278946	-1.222141
39	1	0	2.739747	1.981335	1.046620
40	1	0	4.620592	3.329074	0.116979
41	1	0	3.743687	2.464176	-1.154574
42	1	0	3.493802	4.203550	-0.924723
43	1	0	3.414148	4.072519	2.235136
44	1	0	1.679341	3.799352	2.426020
45	1	0	2.265589	5.013287	1.281344

---

**Table S5.** The coordinate for the lowest-energy conformer of compound **3** for ECD calculation.

E=-1144.293861 a.u.

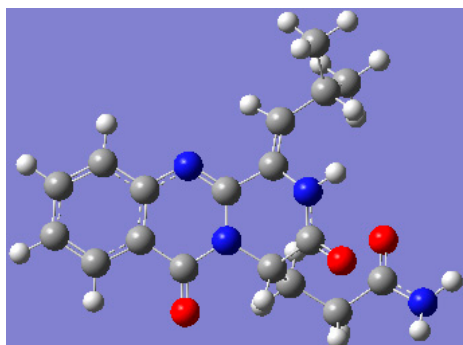
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	4.423436	-2.950361	0.212409
2	6	0	3.506887	-3.992575	-0.008359
3	6	0	2.162691	-3.707731	-0.170130
4	6	0	1.718811	-2.377411	-0.113785
5	6	0	2.634125	-1.328691	0.105156
6	6	0	3.997563	-1.635072	0.268209
7	6	0	0.297529	-2.070236	-0.269743
8	7	0	-0.006347	-0.687688	-0.173477
9	6	0	0.979554	0.261294	0.020095
10	7	0	2.233416	-0.003430	0.155389
11	6	0	-1.438129	-0.322385	-0.262526
12	6	0	-1.628242	1.014425	-0.968238
13	7	0	-0.626988	1.906060	-0.797913
14	6	0	0.509050	1.709409	0.105639
15	8	0	-0.586106	-2.887654	-0.456798
16	1	0	0.176698	1.869185	1.141198
17	6	0	1.626342	2.711025	-0.207205
18	6	0	1.273730	4.197223	0.003518
19	6	0	0.888948	4.517866	1.454095
20	6	0	2.455745	5.066431	-0.447980
21	8	0	-2.634283	1.253942	-1.627363
22	1	0	-1.901431	-1.083299	-0.887216
23	6	0	-2.102556	-0.355296	1.136991
24	6	0	-3.601318	-0.001139	1.163138
25	6	0	-4.484072	-1.028814	0.449874
26	8	0	-4.934468	-1.999807	1.036897
27	7	0	-4.724495	-0.773756	-0.865354



28	1	0	5.476597	-3.179548	0.339816
29	1	0	3.854370	-5.019169	-0.050698
30	1	0	1.433434	-4.491468	-0.338915
31	1	0	4.693247	-0.820541	0.435434
32	1	0	-0.758277	2.812143	-1.226253
33	1	0	2.481634	2.450221	0.418901
34	1	0	1.949975	2.549602	-1.241024
35	1	0	0.420147	4.466061	-0.636323
36	1	0	0.697453	5.587282	1.577566
37	1	0	-0.014821	3.993375	1.776947
38	1	0	1.695543	4.246917	2.143729
39	1	0	2.223279	6.130402	-0.350513
40	1	0	2.719109	4.877474	-1.492628
41	1	0	3.344800	4.865368	0.158897
42	1	0	-1.961341	-1.362484	1.536680
43	1	0	-1.570271	0.330011	1.805568
44	1	0	-3.925723	0.006993	2.205116
45	1	0	-3.768455	0.999815	0.756748
46	1	0	-4.239313	-0.033527	-1.358526
47	1	0	-5.237462	-1.462734	-1.393865

---

**Table S6.** The coordinate for the lowest-energy conformer of compound **4** for ECD calculation.

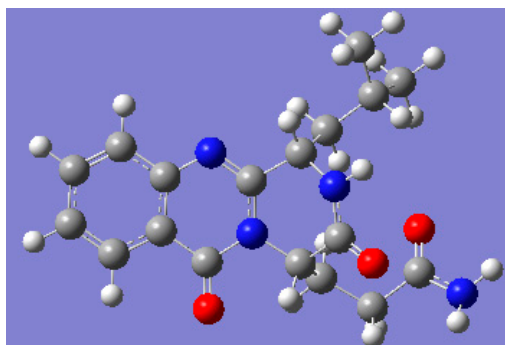
E=-1142.4931498 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.407963	0.744444	-0.541343
2	6	0	-5.516841	-0.645553	-0.356043
3	6	0	-4.389227	-1.393137	-0.068823
4	6	0	-3.139834	-0.762258	0.040315
5	6	0	-3.024382	0.632639	-0.140044
6	6	0	-4.183858	1.377942	-0.435029
7	6	0	-1.938728	-1.547877	0.310584
8	7	0	-0.753275	-0.779090	0.395288
9	6	0	-0.759827	0.598043	0.243878
10	7	0	-1.821479	1.293179	-0.014296
11	6	0	0.509044	-1.537621	0.520838
12	6	0	1.562531	-0.765715	1.296450
13	7	0	1.495032	0.588828	1.185432
14	6	0	0.537649	1.295594	0.427960
15	8	0	-1.900772	-2.761691	0.439151
16	6	0	0.739371	2.532612	-0.051440
17	6	0	1.986193	3.365632	0.042918
18	6	0	2.567608	3.601054	-1.364873
19	6	0	1.689787	4.700360	0.751202
20	8	0	2.431747	-1.337648	1.934071
21	1	0	0.275465	-2.428612	1.100751
22	6	0	1.011343	-1.967290	-0.879954
23	6	0	2.317315	-2.766803	-0.871297
24	6	0	3.567966	-1.887405	-0.827541
25	8	0	3.585532	-0.756865	-1.287722
26	7	0	4.687673	-2.491658	-0.330150
27	1	0	-6.295637	1.326618	-0.767784

28	1	0	-6.484289	-1.128892	-0.439590
29	1	0	-4.442391	-2.466266	0.073671
30	1	0	-4.085665	2.449338	-0.568782
31	1	0	2.269864	1.096251	1.589410
32	1	0	-0.105165	2.980526	-0.565418
33	1	0	2.753691	2.840511	0.623971
34	1	0	3.468526	4.219397	-1.310576
35	1	0	2.831442	2.657046	-1.846400
36	1	0	1.845769	4.119030	-2.004372
37	1	0	2.596064	5.308710	0.824763
38	1	0	1.303645	4.543647	1.761795
39	1	0	0.944599	5.279932	0.197458
40	1	0	0.216962	-2.574573	-1.318498
41	1	0	1.142327	-1.084007	-1.509811
42	1	0	2.328118	-3.501766	-0.061420
43	1	0	2.383902	-3.338055	-1.804349
44	1	0	4.599147	-3.273201	0.299749
45	1	0	5.497026	-1.902767	-0.199034

---

**Table S7.** The coordinate for the lowest-energy conformer of compound **3** for ECD calculation.

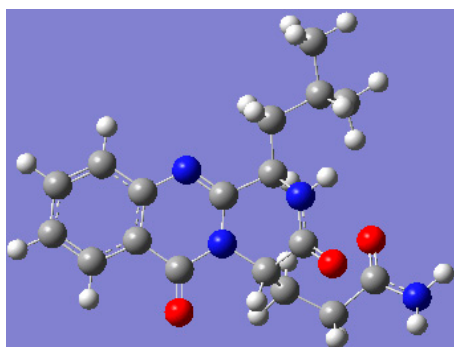
E=-1144.2721485 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.474130	0.926211	-0.468557
2	6	0	-5.571434	-0.474214	-0.551403
3	6	0	-4.448316	-1.256609	-0.354057
4	6	0	-3.213557	-0.649939	-0.071979
5	6	0	-3.111582	0.753625	0.016309
6	6	0	-4.264819	1.535319	-0.187674
7	6	0	-2.018612	-1.467768	0.121540
8	7	0	-0.840531	-0.721862	0.375487
9	6	0	-0.864384	0.653706	0.476390
10	7	0	-1.917987	1.383645	0.315177
11	6	0	0.409044	-1.505134	0.512071
12	6	0	1.432789	-0.831052	1.419442
13	7	0	1.308047	0.505519	1.588250
14	6	0	0.427803	1.374769	0.809983
15	8	0	-1.979663	-2.687014	0.069895
16	1	0	0.121818	2.192537	1.463431
17	6	0	1.102693	1.967584	-0.449444
18	6	0	2.319179	2.870272	-0.169929
19	6	0	3.010471	3.221456	-1.494157
20	6	0	1.950122	4.141726	0.606534
21	8	0	2.298926	-1.499542	1.967716
22	1	0	0.128915	-2.427025	1.021292
23	6	0	0.959203	-1.883487	-0.885549
24	6	0	2.287064	-2.646021	-0.872492
25	6	0	3.507080	-1.729516	-0.790642
26	8	0	3.509485	-0.608703	-1.278314
27	7	0	4.621233	-2.284393	-0.229587

28	1	0	-6.358583	1.535633	-0.624925
29	1	0	-6.527047	-0.938504	-0.770087
30	1	0	-4.493404	-2.337881	-0.413077
31	1	0	-4.175120	2.613399	-0.116106
32	1	0	2.021239	0.934333	2.162235
33	1	0	1.421959	1.149010	-1.099524
34	1	0	0.340730	2.539753	-0.990182
35	1	0	3.041938	2.295282	0.423732
36	1	0	3.892686	3.845632	-1.323043
37	1	0	3.334507	2.319993	-2.019543
38	1	0	2.337400	3.778356	-2.155678
39	1	0	2.831990	4.768796	0.765618
40	1	0	1.525752	3.932718	1.593333
41	1	0	1.216341	4.740070	0.055794
42	1	0	0.191809	-2.499319	-1.357084
43	1	0	1.078159	-0.985442	-1.494216
44	1	0	2.308301	-3.399221	-0.081272
45	1	0	2.385267	-3.188589	-1.819894
46	1	0	4.516221	-3.030992	0.439438
47	1	0	5.402139	-1.662385	-0.078228

---

**Table S8.** The coordinate for the conformer C1 of compound 3 for NMR calculation.

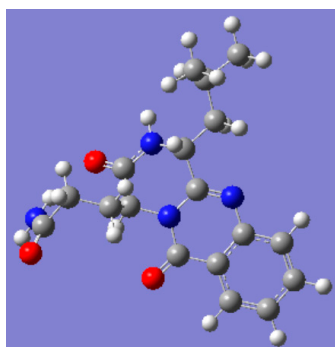
E=-1144.2460465 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.399386	0.822946	-0.495915
2	6	0	-5.548607	-0.554476	-0.258338
3	6	0	-4.436085	-1.334417	0.001479
4	6	0	-3.161092	-0.747469	0.030152
5	6	0	-3.006304	0.633991	-0.204133
6	6	0	-4.148255	1.412569	-0.470091
7	6	0	-1.977856	-1.567756	0.286793
8	7	0	-0.766518	-0.839652	0.293722
9	6	0	-0.729830	0.524832	0.066132
10	7	0	-1.767354	1.251618	-0.173110
11	6	0	0.475748	-1.617535	0.497881
12	6	0	1.515737	-0.826125	1.287249
13	7	0	1.436590	0.519076	1.154787
14	6	0	0.655998	1.156157	0.089914
15	8	0	-1.981649	-2.774506	0.473379
16	1	0	1.127225	0.932757	-0.877106
17	6	0	0.615053	2.675038	0.277722
18	6	0	1.978923	3.390331	0.196567
19	6	0	2.698276	3.154377	-1.138421
20	6	0	1.778848	4.890721	0.451962
21	8	0	2.340972	-1.399258	1.983340
22	1	0	0.201382	-2.465809	1.122570
23	6	0	0.998637	-2.155631	-0.857535
24	6	0	2.364197	-2.842856	-0.792405
25	6	0	3.530594	-1.854412	-0.799917
26	8	0	3.431499	-0.732648	-1.276107
27	7	0	4.711898	-2.341739	-0.322954
28	1	0	-6.275154	1.430430	-0.701083

29	1	0	-6.535627	-1.003866	-0.280076
30	1	0	-4.521087	-2.399234	0.184911
31	1	0	-4.017217	2.473830	-0.649105
32	1	0	2.191565	1.045380	1.572702
33	1	0	-0.049295	3.081339	-0.488085
34	1	0	0.136809	2.893656	1.238450
35	1	0	2.631920	3.017994	0.999866
36	1	0	3.622307	3.737895	-1.185786
37	1	0	2.970941	2.107592	-1.293104
38	1	0	2.070998	3.467687	-1.980521
39	1	0	2.735077	5.421507	0.450275
40	1	0	1.296178	5.076833	1.416040
41	1	0	1.150891	5.340202	-0.324727
42	1	0	0.249448	-2.863025	-1.219162
43	1	0	1.055195	-1.341409	-1.582656
44	1	0	2.433003	-3.515530	0.066918
45	1	0	2.485783	-3.467493	-1.684928
46	1	0	4.714049	-3.124506	0.311600
47	1	0	5.469137	-1.681755	-0.223559

---

**Table S9.** The coordinate for the conformer C2 of compound 3 for NMR calculation.

E=-1144.2455425 a.u.

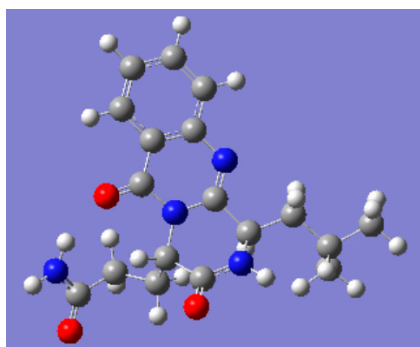
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	4.423436	-2.950361	0.212409
2	6	0	3.506887	-3.992575	-0.008359
3	6	0	2.162691	-3.707731	-0.170130
4	6	0	1.718811	-2.377411	-0.113785
5	6	0	2.634125	-1.328691	0.105156
6	6	0	3.997563	-1.635072	0.268209
7	6	0	0.297529	-2.070236	-0.269743
8	7	0	-0.006347	-0.687688	-0.173477
9	6	0	0.979554	0.261294	0.020095
10	7	0	2.233416	-0.003430	0.155389
11	6	0	-1.438129	-0.322385	-0.262526
12	6	0	-1.628242	1.014425	-0.968238
13	7	0	-0.626988	1.906060	-0.797913
14	6	0	0.509050	1.709409	0.105639
15	8	0	-0.586106	-2.887654	-0.456798
16	1	0	0.176698	1.869185	1.141198
17	6	0	1.626342	2.711025	-0.207205
18	6	0	1.273730	4.197223	0.003518
19	6	0	0.888948	4.517866	1.454095
20	6	0	2.455745	5.066431	-0.447980
21	8	0	-2.634283	1.253942	-1.627363
22	1	0	-1.901431	-1.083299	-0.887216
23	6	0	-2.102556	-0.355296	1.136991
24	6	0	-3.601318	-0.001139	1.163138
25	6	0	-4.484072	-1.028814	0.449874
26	8	0	-4.934468	-1.999807	1.036897
27	7	0	-4.724495	-0.773756	-0.865354
28	1	0	5.476597	-3.179548	0.339816



29	1	0	3.854370	-5.019169	-0.050698
30	1	0	1.433434	-4.491468	-0.338915
31	1	0	4.693247	-0.820541	0.435434
32	1	0	-0.758277	2.812143	-1.226253
33	1	0	2.481634	2.450221	0.418901
34	1	0	1.949975	2.549602	-1.241024
35	1	0	0.420147	4.466061	-0.636323
36	1	0	0.697453	5.587282	1.577566
37	1	0	-0.014821	3.993375	1.776947
38	1	0	1.695543	4.246917	2.143729
39	1	0	2.223279	6.130402	-0.350513
40	1	0	2.719109	4.877474	-1.492628
41	1	0	3.344800	4.865368	0.158897
42	1	0	-1.961341	-1.362484	1.536680
43	1	0	-1.570271	0.330011	1.805568
44	1	0	-3.925723	0.006993	2.205116
45	1	0	-3.768455	0.999815	0.756748
46	1	0	-4.239313	-0.033527	-1.358526
47	1	0	-5.237462	-1.462734	-1.393865

---

**Table S10.** The coordinate for the conformer C3 of compound 3 for NMR calculation.

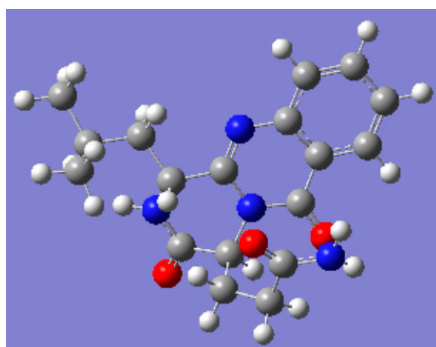
E=-1144.2447972 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-0.527385	5.022303	-0.359329
2	6	0	-1.841234	4.698370	0.022851
3	6	0	-2.175901	3.386426	0.303653
4	6	0	-1.200214	2.380608	0.206185
5	6	0	0.118881	2.701823	-0.170515
6	6	0	0.442970	4.041366	-0.454338
7	6	0	-1.546545	0.988846	0.476361
8	7	0	-0.483112	0.079272	0.317494
9	6	0	0.795499	0.515015	-0.001574
10	7	0	1.109468	1.739836	-0.247313
11	6	0	-0.758928	-1.364411	0.552453
12	6	0	0.393430	-2.024642	1.312868
13	7	0	1.623622	-1.536023	1.003860
14	6	0	1.872207	-0.563861	-0.060159
15	8	0	-2.665250	0.606318	0.802802
16	1	0	1.775812	-1.055880	-1.038403
17	6	0	3.288115	0.012512	0.050083
18	6	0	4.436647	-1.002110	-0.122433
19	6	0	4.419989	-1.692615	-1.492819
20	6	0	5.777537	-0.291860	0.109882
21	8	0	0.199390	-2.924587	2.108018
22	1	0	-1.622949	-1.406917	1.211003
23	6	0	-1.046556	-2.145937	-0.751097
24	6	0	-2.388271	-1.825148	-1.438494
25	6	0	-3.567161	-2.531609	-0.765088
26	8	0	-3.827288	-3.698513	-1.016423
27	7	0	-4.282785	-1.781746	0.114124
28	1	0	-0.271624	6.053805	-0.579174

29	1	0	-2.591056	5.478547	0.096271
30	1	0	-3.181225	3.109541	0.598158
31	1	0	1.461810	4.276134	-0.740856
32	1	0	2.409745	-1.992681	1.444411
33	1	0	3.384102	0.795241	-0.705039
34	1	0	3.379930	0.515310	1.018471
35	1	0	4.353237	-1.783129	0.647079
36	1	0	5.281792	-2.356293	-1.603523
37	1	0	3.526093	-2.303566	-1.646403
38	1	0	4.465410	-0.957500	-2.303671
39	1	0	6.613347	-0.993107	0.037123
40	1	0	5.820568	0.175406	1.097939
41	1	0	5.939667	0.494089	-0.635251
42	1	0	-0.233583	-1.968760	-1.461865
43	1	0	-1.020954	-3.212221	-0.509576
44	1	0	-2.549323	-0.746752	-1.505789
45	1	0	-2.350160	-2.215260	-2.457400
46	1	0	-3.962101	-0.870464	0.415313
47	1	0	-5.012181	-2.241113	0.637915

---

**Table S11.** The coordinate for the conformer C4 of compound **3** for NMR calculation.

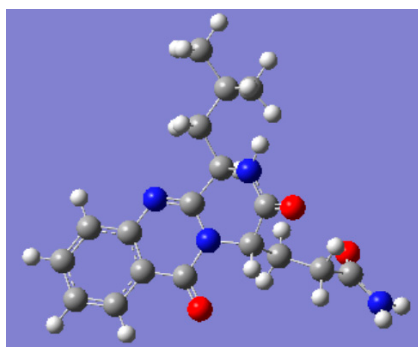
E=-1144.2444319 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	3.340116	3.445290	0.092548
2	6	0	4.210436	2.460025	-0.406419
3	6	0	3.720976	1.208184	-0.732988
4	6	0	2.355374	0.925208	-0.567307
5	6	0	1.478124	1.912256	-0.075900
6	6	0	1.992592	3.179889	0.255475
7	6	0	1.833273	-0.402013	-0.886859
8	7	0	0.443482	-0.540884	-0.685413
9	6	0	-0.331600	0.511973	-0.227242
10	7	0	0.123648	1.678055	0.075615
11	6	0	-0.169687	-1.844302	-1.046388
12	6	0	-1.531889	-1.645986	-1.712537
13	7	0	-2.244231	-0.599407	-1.222253
14	6	0	-1.817217	0.213211	-0.078992
15	8	0	2.510084	-1.342736	-1.281110
16	1	0	-1.923979	-0.366724	0.847622
17	6	0	-2.667867	1.481193	0.036176
18	6	0	-4.167091	1.254376	0.316116
19	6	0	-4.420738	0.510087	1.633802
20	6	0	-4.894166	2.606434	0.308441
21	8	0	-1.932874	-2.398398	-2.583291
22	1	0	0.490161	-2.282989	-1.791444
23	6	0	-0.331179	-2.837440	0.128355
24	6	0	0.957983	-3.177844	0.881858
25	6	0	1.338302	-2.121250	1.917546
26	8	0	0.500106	-1.545322	2.594739
27	7	0	2.676998	-1.919789	2.082831
28	1	0	3.729570	4.425339	0.349109

29	1	0	5.263948	2.683797	-0.535850
30	1	0	4.368950	0.431365	-1.122196
31	1	0	1.307752	3.930218	0.634080
32	1	0	-3.174248	-0.473503	-1.596095
33	1	0	-2.240725	2.089158	0.836345
34	1	0	-2.546114	2.064746	-0.882470
35	1	0	-4.603435	0.654812	-0.496282
36	1	0	-5.493344	0.415091	1.824807
37	1	0	-4.004264	-0.500728	1.633337
38	1	0	-3.981557	1.048369	2.480371
39	1	0	-5.969347	2.478430	0.462095
40	1	0	-4.754402	3.134277	-0.639565
41	1	0	-4.521477	3.256776	1.106889
42	1	0	-1.065529	-2.469771	0.849084
43	1	0	-0.746656	-3.747833	-0.311654
44	1	0	0.791964	-4.099936	1.450728
45	1	0	1.778459	-3.375709	0.189387
46	1	0	3.334704	-2.209569	1.376700
47	1	0	2.967299	-1.195531	2.722722

---

**Table S12.** The coordinate for the conformer C5 of compound 3 for NMR calculation.

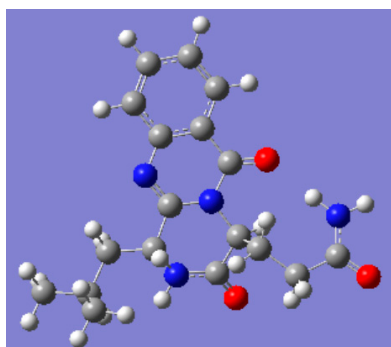
E=-1144.243994 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.143445	-1.389299	-0.808823
2	6	0	-4.711388	-2.655489	-0.377446
3	6	0	-3.411470	-2.828438	0.062678
4	6	0	-2.528103	-1.737474	0.080219
5	6	0	-2.957252	-0.465325	-0.348452
6	6	0	-4.281997	-0.307012	-0.796256
7	6	0	-1.148902	-1.913492	0.532279
8	7	0	-0.378880	-0.725994	0.507415
9	6	0	-0.906655	0.475977	0.075184
10	7	0	-2.117742	0.635602	-0.336639
11	6	0	1.033718	-0.853269	0.933919
12	6	0	1.523848	0.405539	1.642280
13	7	0	0.950879	1.560373	1.218489
14	6	0	0.052624	1.660988	0.067212
15	8	0	-0.655544	-2.967326	0.900848
16	1	0	0.637477	1.579275	-0.860463
17	6	0	-0.676521	3.008749	0.065405
18	6	0	0.219045	4.254339	-0.090568
19	6	0	1.013262	4.256277	-1.403578
20	6	0	-0.647322	5.516806	0.020770
21	8	0	2.382080	0.355057	2.508910
22	1	0	1.052434	-1.658146	1.666118
23	6	0	1.939546	-1.233115	-0.259451
24	6	0	3.341148	-1.670006	0.170195
25	6	0	4.273928	-1.827277	-1.026756
26	8	0	4.178885	-1.142838	-2.032001
27	7	0	5.245990	-2.774982	-0.884394
28	1	0	-6.163963	-1.259423	-1.154787

29	1	0	-5.397748	-3.495325	-0.391249
30	1	0	-3.050410	-3.793767	0.397937
31	1	0	-4.601105	0.675791	-1.124416
32	1	0	1.308884	2.410042	1.632014
33	1	0	-1.404794	2.984856	-0.747576
34	1	0	-1.258668	3.084302	0.990061
35	1	0	0.941397	4.291450	0.738054
36	1	0	1.591128	5.178981	-1.505984
37	1	0	1.722836	3.427062	-1.469365
38	1	0	0.343832	4.190480	-2.268040
39	1	0	-0.036805	6.421787	-0.043391
40	1	0	-1.192719	5.549598	0.968466
41	1	0	-1.385313	5.558082	-0.787285
42	1	0	1.450805	-2.050584	-0.795465
43	1	0	2.023238	-0.400994	-0.962434
44	1	0	3.789217	-0.923255	0.834517
45	1	0	3.287570	-2.597883	0.748687
46	1	0	5.352788	-3.319402	-0.045058
47	1	0	5.940426	-2.867933	-1.609341

---

**Table S13.** The coordinate for the conformer C6 of compound 3 for NMR calculation.

E=-1144.2433829 a.u.

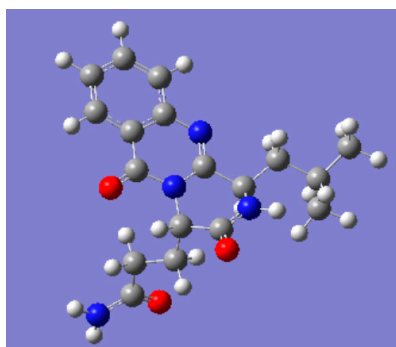
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-1.721192	4.917395	0.225409
2	6	0	-0.353539	5.168125	0.014945
3	6	0	0.526090	4.113261	-0.143631
4	6	0	0.047535	2.793418	-0.098514
5	6	0	-1.322619	2.536771	0.111325
6	6	0	-2.203429	3.622100	0.274669
7	6	0	0.961998	1.669370	-0.254298
8	7	0	0.346665	0.403460	-0.224727
9	6	0	-1.012513	0.263684	-0.001382
10	7	0	-1.826264	1.249412	0.162485
11	6	0	1.249204	-0.761451	-0.396316
12	6	0	0.541954	-1.950626	-1.039945
13	7	0	-0.796764	-2.022024	-0.821143
14	6	0	-1.534924	-1.163654	0.103305
15	8	0	2.176879	1.767339	-0.395938
16	1	0	-1.341404	-1.485447	1.137804
17	6	0	-3.043088	-1.254820	-0.159388
18	6	0	-3.676932	-2.645045	0.045839
19	6	0	-3.506311	-3.174242	1.476075
20	6	0	-5.162885	-2.584256	-0.334803
21	8	0	1.162301	-2.771173	-1.692714
22	1	0	2.010966	-0.449815	-1.107707
23	6	0	1.919631	-1.150899	0.947501
24	6	0	3.168417	-2.033085	0.811887
25	6	0	4.504390	-1.395288	0.414925
26	8	0	5.518276	-2.076598	0.426463
27	7	0	4.520493	-0.076982	0.079812
28	1	0	-2.405701	5.750126	0.350887



29	1	0	0.009555	6.189447	-0.021643
30	1	0	1.584884	4.278196	-0.304775
31	1	0	-3.254876	3.413678	0.436442
32	1	0	-1.258181	-2.839796	-1.195035
33	1	0	-3.534032	-0.534684	0.498014
34	1	0	-3.232865	-0.910290	-1.181282
35	1	0	-3.205332	-3.366271	-0.637468
36	1	0	-4.017141	-4.133111	1.598461
37	1	0	-2.458416	-3.335720	1.744415
38	1	0	-3.934216	-2.478872	2.206230
39	1	0	-5.633946	-3.567065	-0.246393
40	1	0	-5.301872	-2.241114	-1.364022
41	1	0	-5.707972	-1.896301	0.320040
42	1	0	2.162417	-0.230831	1.485249
43	1	0	1.185224	-1.674854	1.569016
44	1	0	3.364838	-2.520333	1.770889
45	1	0	2.993639	-2.845269	0.101306
46	1	0	3.705256	0.515019	-0.019847
47	1	0	5.409881	0.305168	-0.203673

---

**Table S14.** The coordinate for the conformer C7 of compound 3 for NMR calculation.

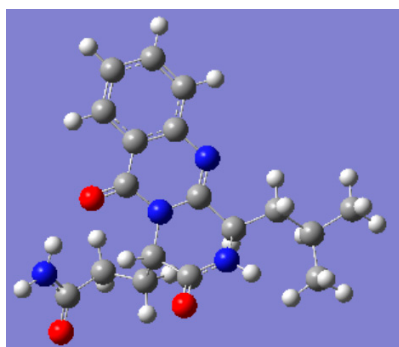
E=-1144.2428681 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-2.266329	4.440534	-0.929245
2	6	0	-3.356688	3.856669	-0.261230
3	6	0	-3.215472	2.619153	0.340606
4	6	0	-1.983032	1.949103	0.282199
5	6	0	-0.885593	2.533128	-0.381360
6	6	0	-1.045843	3.792197	-0.989331
7	6	0	-1.831203	0.632116	0.900009
8	7	0	-0.545111	0.064099	0.735616
9	6	0	0.477960	0.752819	0.106682
10	7	0	0.351432	1.914416	-0.436901
11	6	0	-0.319923	-1.281904	1.316857
12	6	0	1.061737	-1.382794	1.963614
13	7	0	2.026037	-0.655539	1.340907
14	6	0	1.826320	0.042065	0.068959
15	8	0	-2.710447	0.032050	1.497857
16	1	0	1.772281	-0.691906	-0.747451
17	6	0	2.989684	0.996721	-0.217429
18	6	0	4.368524	0.331712	-0.403250
19	6	0	4.394470	-0.672849	-1.563290
20	6	0	5.432051	1.420121	-0.604282
21	8	0	1.260070	-2.082534	2.940561
22	1	0	-1.052650	-1.389640	2.113369
23	6	0	-0.493242	-2.429166	0.298615
24	6	0	-1.898753	-2.536734	-0.296304
25	6	0	-1.969080	-3.628872	-1.359741
26	8	0	-1.046419	-3.863819	-2.122243
27	7	0	-3.150374	-4.309781	-1.416747
28	1	0	-2.382606	5.411270	-1.400563

29	1	0	-4.307242	4.377452	-0.218675
30	1	0	-4.039326	2.145899	0.862049
31	1	0	-0.193931	4.231774	-1.495727
32	1	0	2.964573	-0.757182	1.700917
33	1	0	2.736951	1.558712	-1.118877
34	1	0	3.039074	1.732854	0.591838
35	1	0	4.638748	-0.210180	0.514893
36	1	0	5.400338	-1.078603	-1.702520
37	1	0	3.727313	-1.523554	-1.400140
38	1	0	4.100011	-0.195136	-2.504024
39	1	0	6.431224	0.984556	-0.692212
40	1	0	5.448748	2.126142	0.231081
41	1	0	5.239310	1.992694	-1.517670
42	1	0	0.227909	-2.338808	-0.517005
43	1	0	-0.245948	-3.355268	0.826035
44	1	0	-2.640392	-2.694619	0.490850
45	1	0	-2.177249	-1.600985	-0.793171
46	1	0	-3.922034	-4.108697	-0.803342
47	1	0	-3.278158	-4.998827	-2.141656

---

**Table S15.** The coordinate for the conformer C8 of compound **3** for NMR calculation.

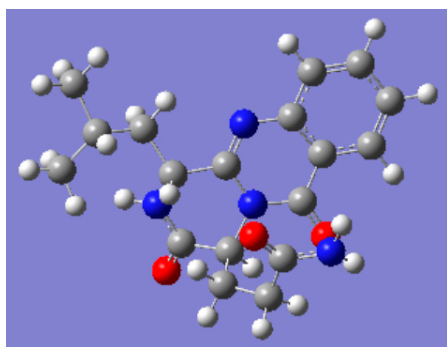
E=-1144.2428074 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-0.724982	5.014538	-0.286268
2	6	0	-2.025115	4.633477	0.089971
3	6	0	-2.307926	3.305326	0.351083
4	6	0	-1.293092	2.340553	0.239988
5	6	0	0.012202	2.719008	-0.130024
6	6	0	0.283439	4.074183	-0.394402
7	6	0	-1.583564	0.932030	0.486914
8	7	0	-0.484500	0.067267	0.315777
9	6	0	0.777117	0.557243	0.008711
10	7	0	1.039301	1.797525	-0.219216
11	6	0	-0.709825	-1.389991	0.520086
12	6	0	0.462757	-2.022189	1.271778
13	7	0	1.674482	-1.482287	0.978335
14	6	0	1.896132	-0.483083	-0.068748
15	8	0	-2.686795	0.500102	0.803618
16	1	0	1.812087	-0.960599	-1.057427
17	6	0	3.302163	0.128387	0.037747
18	6	0	4.418329	-0.638704	-0.698639
19	6	0	5.728186	0.155736	-0.609519
20	6	0	4.628674	-2.076713	-0.204532
21	8	0	0.298905	-2.942430	2.050780
22	1	0	-1.573141	-1.476283	1.175244
23	6	0	-0.967526	-2.154280	-0.799658
24	6	0	-2.319242	-1.868955	-1.483181
25	6	0	-3.472449	-2.632560	-0.827826
26	8	0	-3.686453	-3.803720	-1.101766
27	7	0	-4.220335	-1.927255	0.061435
28	1	0	-0.510021	6.058377	-0.490995

29	1	0	-2.805159	5.382316	0.173965
30	1	0	-3.301776	2.984608	0.639965
31	1	0	1.292102	4.353672	-0.676720
32	1	0	2.474795	-1.910998	1.420694
33	1	0	3.246583	1.138434	-0.367379
34	1	0	3.554240	0.243807	1.099481
35	1	0	4.132280	-0.689513	-1.758763
36	1	0	6.524804	-0.330848	-1.179169
37	1	0	5.611538	1.170397	-1.000779
38	1	0	6.070165	0.238864	0.427842
39	1	0	5.453524	-2.553460	-0.741266
40	1	0	3.748096	-2.707409	-0.354410
41	1	0	4.887444	-2.099018	0.861007
42	1	0	-0.160853	-1.933132	-1.505161
43	1	0	-0.903315	-3.223656	-0.579872
44	1	0	-2.519976	-0.796133	-1.528834
45	1	0	-2.265067	-2.236183	-2.509861
46	1	0	-3.934891	-1.011193	0.382607
47	1	0	-4.933023	-2.424456	0.573581

---

**Table S16.** The coordinate for the conformer C9 of compound 3 for NMR calculation.

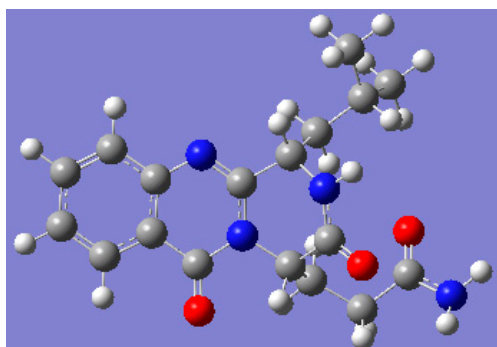
E=-1144.2425858 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	3.450967	3.362192	-0.050231
2	6	0	4.290131	2.333382	-0.513292
3	6	0	3.763188	1.084452	-0.788462
4	6	0	2.390871	0.848090	-0.606236
5	6	0	1.544983	1.878199	-0.150622
6	6	0	2.097135	3.142557	0.128202
7	6	0	1.828797	-0.474890	-0.869371
8	7	0	0.436133	-0.565668	-0.658148
9	6	0	-0.307087	0.526081	-0.239411
10	7	0	0.185505	1.687977	0.016834
11	6	0	-0.212942	-1.865201	-0.968651
12	6	0	-1.568660	-1.652333	-1.642114
13	7	0	-2.249596	-0.569596	-1.188809
14	6	0	-1.801837	0.267599	-0.068987
15	8	0	2.476892	-1.449723	-1.226986
16	1	0	-1.908144	-0.288793	0.873879
17	6	0	-2.638530	1.550351	0.035246
18	6	0	-3.939384	1.430309	0.854060
19	6	0	-4.595743	2.812204	0.974050
20	6	0	-4.940480	0.407081	0.300194
21	8	0	-1.989961	-2.421074	-2.489031
22	1	0	0.433379	-2.348215	-1.697961
23	6	0	-0.397744	-2.811805	0.240364
24	6	0	0.883017	-3.156199	1.006461
25	6	0	1.287228	-2.072614	2.004471
26	8	0	0.462538	-1.459647	2.665679
27	7	0	2.629888	-1.888144	2.155744
28	1	0	3.869612	4.339966	0.165773

29	1	0	5.348945	2.521349	-0.655673
30	1	0	4.386501	0.274452	-1.149260
31	1	0	1.436493	3.927035	0.479559
32	1	0	-3.171318	-0.421670	-1.573594
33	1	0	-2.006198	2.310575	0.493548
34	1	0	-2.856624	1.913640	-0.977281
35	1	0	-3.657070	1.106630	1.865227
36	1	0	-5.490506	2.772555	1.601726
37	1	0	-3.912905	3.544540	1.414215
38	1	0	-4.899575	3.191923	-0.007746
39	1	0	-5.851073	0.389141	0.905672
40	1	0	-4.542160	-0.611151	0.297116
41	1	0	-5.243470	0.658531	-0.723705
42	1	0	-1.123386	-2.403430	0.947923
43	1	0	-0.834392	-3.726911	-0.168341
44	1	0	0.694944	-4.053313	1.607443
45	1	0	1.698426	-3.397574	0.321994
46	1	0	3.279172	-2.215299	1.458238
47	1	0	2.936634	-1.149125	2.770565

---

**Table S17.** The coordinate for the conformer C1 of compound 5 for NMR calculation.

E=-1144.2458631 a.u.

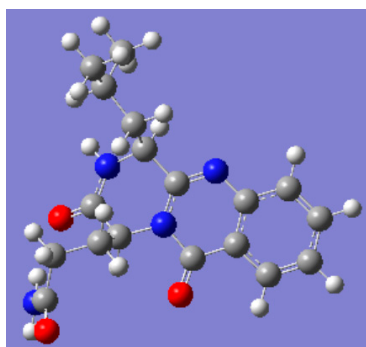
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.474130	0.926211	-0.468557
2	6	0	-5.571434	-0.474214	-0.551403
3	6	0	-4.448316	-1.256609	-0.354057
4	6	0	-3.213557	-0.649939	-0.071979
5	6	0	-3.111582	0.753625	0.016309
6	6	0	-4.264819	1.535319	-0.187674
7	6	0	-2.018612	-1.467768	0.121540
8	7	0	-0.840531	-0.721862	0.375487
9	6	0	-0.864384	0.653706	0.476390
10	7	0	-1.917987	1.383645	0.315177
11	6	0	0.409044	-1.505134	0.512071
12	6	0	1.432789	-0.831052	1.419442
13	7	0	1.308047	0.505519	1.588250
14	6	0	0.427803	1.374769	0.809983
15	8	0	-1.979663	-2.687014	0.069895
16	1	0	0.121818	2.192537	1.463431
17	6	0	1.102693	1.967584	-0.449444
18	6	0	2.319179	2.870272	-0.169929
19	6	0	3.010471	3.221456	-1.494157
20	6	0	1.950122	4.141726	0.606534
21	8	0	2.298926	-1.499542	1.967716
22	1	0	0.128915	-2.427025	1.021292
23	6	0	0.959203	-1.883487	-0.885549
24	6	0	2.287064	-2.646021	-0.872492
25	6	0	3.507080	-1.729516	-0.790642
26	8	0	3.509485	-0.608703	-1.278314
27	7	0	4.621233	-2.284393	-0.229587
28	1	0	-6.358583	1.535633	-0.624925



29	1	0	-6.527047	-0.938504	-0.770087
30	1	0	-4.493404	-2.337881	-0.413077
31	1	0	-4.175120	2.613399	-0.116106
32	1	0	2.021239	0.934333	2.162235
33	1	0	1.421959	1.149010	-1.099524
34	1	0	0.340730	2.539753	-0.990182
35	1	0	3.041938	2.295282	0.423732
36	1	0	3.892686	3.845632	-1.323043
37	1	0	3.334507	2.319993	-2.019543
38	1	0	2.337400	3.778356	-2.155678
39	1	0	2.831990	4.768796	0.765618
40	1	0	1.525752	3.932718	1.593333
41	1	0	1.216341	4.740070	0.055794
42	1	0	0.191809	-2.499319	-1.357084
43	1	0	1.078159	-0.985442	-1.494216
44	1	0	2.308301	-3.399221	-0.081272
45	1	0	2.385267	-3.188589	-1.819894
46	1	0	4.516221	-3.030992	0.439438
47	1	0	5.402139	-1.662385	-0.078228

---

**Table S18.** The coordinate for the conformer C2 of compound 5 for NMR calculation.

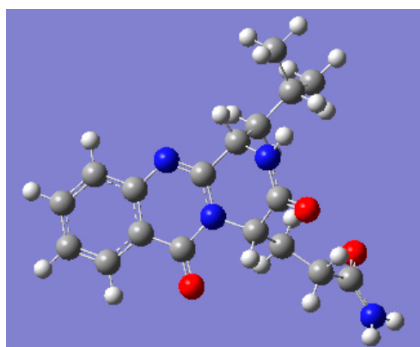
E=-1144.2455392 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	5.400768	-0.329043	0.256239
2	6	0	5.083554	-1.696208	0.337851
3	6	0	3.770185	-2.109149	0.205115
4	6	0	2.757616	-1.160511	-0.009489
5	6	0	3.071035	0.211362	-0.094470
6	6	0	4.412395	0.614732	0.041808
7	6	0	1.366650	-1.588945	-0.141023
8	7	0	0.445746	-0.521773	-0.322995
9	6	0	0.871971	0.785320	-0.424397
10	7	0	2.100409	1.170100	-0.325148
11	6	0	-0.983181	-0.905191	-0.397966
12	6	0	-1.792843	0.054944	-1.259050
13	7	0	-1.335457	1.322950	-1.352760
14	6	0	-0.160880	1.868451	-0.676124
15	8	0	0.972492	-2.740744	-0.098430
16	1	0	0.322408	2.554620	-1.372784
17	6	0	-0.497842	2.644964	0.617910
18	6	0	-1.448685	3.842869	0.437352
19	6	0	-1.796061	4.436579	1.809483
20	6	0	-0.871638	4.923512	-0.487433
21	8	0	-2.817781	-0.317686	-1.822214
22	1	0	-1.006625	-1.870122	-0.902123
23	6	0	-1.574576	-1.086284	1.024976
24	6	0	-3.056208	-1.500966	1.091166
25	6	0	-3.323015	-2.914415	0.566498
26	8	0	-3.271053	-3.889176	1.299672
27	7	0	-3.620341	-2.992149	-0.759164
28	1	0	6.433008	-0.010518	0.361086

29	1	0	5.869064	-2.425176	0.505077
30	1	0	3.496130	-3.156021	0.264676
31	1	0	4.642767	1.671857	-0.027694
32	1	0	-1.893449	1.940629	-1.926703
33	1	0	-0.931961	1.950751	1.344989
34	1	0	0.450829	2.990855	1.041705
35	1	0	-2.384988	3.474875	-0.003748
36	1	0	-2.506035	5.262701	1.713559
37	1	0	-2.245643	3.688906	2.469345
38	1	0	-0.902889	4.825646	2.309660
39	1	0	-1.556628	5.772092	-0.566541
40	1	0	-0.692517	4.563984	-1.505298
41	1	0	0.080688	5.303956	-0.103215
42	1	0	-0.967013	-1.842695	1.526665
43	1	0	-1.448223	-0.153474	1.582426
44	1	0	-3.352159	-1.507773	2.141666
45	1	0	-3.684992	-0.772526	0.573594
46	1	0	-3.556340	-2.181115	-1.362748
47	1	0	-3.742678	-3.906631	-1.166019

---

**Table S19.** The coordinate for the conformer C3 of compound 5 for NMR calculation.

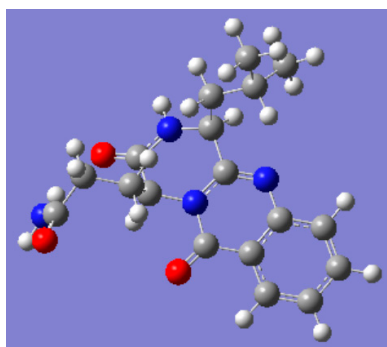
E=-1144.2443127 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.426716	-0.080108	-0.875774
2	6	0	-5.226167	-1.462291	-0.712048
3	6	0	-3.995291	-1.936405	-0.296592
4	6	0	-2.949236	-1.035624	-0.038608
5	6	0	-3.146064	0.351369	-0.198923
6	6	0	-4.405046	0.817116	-0.623477
7	6	0	-1.643759	-1.528041	0.393831
8	7	0	-0.681006	-0.509441	0.608629
9	6	0	-0.991794	0.823135	0.439800
10	7	0	-2.144280	1.267910	0.061869
11	6	0	0.663496	-0.973128	1.025567
12	6	0	1.404630	0.058027	1.868784
13	7	0	1.048048	1.354125	1.686258
14	6	0	0.088892	1.853215	0.704754
15	8	0	-1.350473	-2.700452	0.565869
16	1	0	-0.434494	2.694026	1.161516
17	6	0	0.736147	2.336749	-0.615508
18	6	0	1.804413	3.434654	-0.461428
19	6	0	2.444927	3.723289	-1.826134
20	6	0	1.244170	4.723493	0.156336
21	8	0	2.280624	-0.287357	2.647428
22	1	0	0.491958	-1.835056	1.669041
23	6	0	1.499194	-1.424593	-0.195531
24	6	0	2.722567	-2.258659	0.188119
25	6	0	3.651531	-2.471749	-1.002812
26	8	0	3.771752	-1.656118	-1.902225
27	7	0	4.365401	-3.635221	-0.980713
28	1	0	-6.394668	0.286549	-1.202447

29	1	0	-6.037016	-2.154390	-0.912234
30	1	0	-3.812016	-2.996285	-0.163486
31	1	0	-4.545520	1.885612	-0.741867
32	1	0	1.570628	2.023488	2.234620
33	1	0	1.179721	1.479094	-1.130756
34	1	0	-0.074211	2.702171	-1.255491
35	1	0	2.596546	3.051927	0.195584
36	1	0	3.237845	4.471598	-1.738956
37	1	0	2.885621	2.822365	-2.261838
38	1	0	1.704539	4.108337	-2.535695
39	1	0	2.019483	5.491870	0.221560
40	1	0	0.854694	4.577307	1.168558
41	1	0	0.428997	5.129344	-0.452220
42	1	0	0.845627	-2.015194	-0.841223
43	1	0	1.825525	-0.560983	-0.777549
44	1	0	3.301146	-1.750442	0.966517
45	1	0	2.409765	-3.217224	0.614837
46	1	0	4.301638	-4.294860	-0.223685
47	1	0	5.056450	-3.792143	-1.697811

---

**Table S20.** The coordinate for the conformer C4 of compound **5** for NMR calculation.

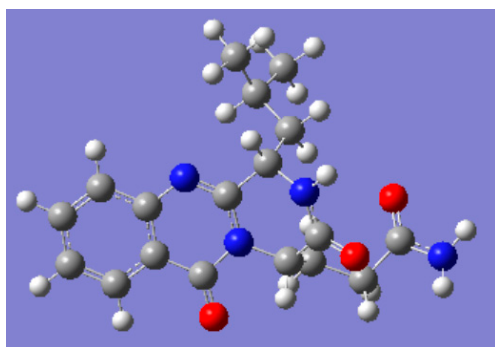
E=-1144.2442476 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	4.479926	-2.585860	0.085208
2	6	0	3.583488	-3.636443	0.347174
3	6	0	2.219875	-3.409472	0.297376
4	6	0	1.735746	-2.128414	-0.011806
5	6	0	2.630453	-1.072355	-0.276805
6	6	0	4.014653	-1.320396	-0.224566
7	6	0	0.296395	-1.879850	-0.064051
8	7	0	-0.050402	-0.532875	-0.354933
9	6	0	0.914475	0.416861	-0.622499
10	7	0	2.187314	0.197963	-0.602673
11	6	0	-1.501476	-0.235763	-0.399485
12	6	0	-1.830212	0.886799	-1.376015
13	7	0	-0.845540	1.774962	-1.636319
14	6	0	0.462672	1.819785	-0.983005
15	8	0	-0.572581	-2.713425	0.119560
16	1	0	1.179533	2.164734	-1.728299
17	6	0	0.467250	2.815976	0.205366
18	6	0	1.831503	3.131133	0.845635
19	6	0	2.804559	3.792667	-0.138813
20	6	0	1.621587	4.017508	2.081294
21	8	0	-2.941984	0.966578	-1.889577
22	1	0	-1.973672	-1.134558	-0.794184
23	6	0	-2.056890	0.011687	1.027471
24	6	0	-3.565849	0.303856	1.120650
25	6	0	-4.445765	-0.891942	0.747010
26	8	0	-4.788434	-1.718319	1.577693
27	7	0	-4.810643	-0.953977	-0.562631
28	1	0	5.548788	-2.770053	0.124472

29	1	0	3.962210	-4.623952	0.587542
30	1	0	1.505855	-4.200656	0.494297
31	1	0	4.694327	-0.502040	-0.433770
32	1	0	-1.094920	2.526314	-2.266242
33	1	0	0.025772	3.749426	-0.167684
34	1	0	-0.219565	2.447756	0.973773
35	1	0	2.280255	2.188468	1.176355
36	1	0	3.738036	4.061002	0.363771
37	1	0	3.067067	3.130382	-0.966655
38	1	0	2.383297	4.714911	-0.555985
39	1	0	2.571865	4.224997	2.580964
40	1	0	0.961142	3.542770	2.813320
41	1	0	1.176466	4.981326	1.810144
42	1	0	-1.827087	-0.879987	1.614872
43	1	0	-1.512145	0.843935	1.482246
44	1	0	-3.795096	0.533778	2.162715
45	1	0	-3.828236	1.179080	0.521724
46	1	0	-4.423979	-0.315922	-1.247748
47	1	0	-5.335367	-1.757567	-0.871975

---

**Table S21.** The coordinate for the conformer C5 of compound 5 for NMR calculation.

E=-1144.2439025 a.u.

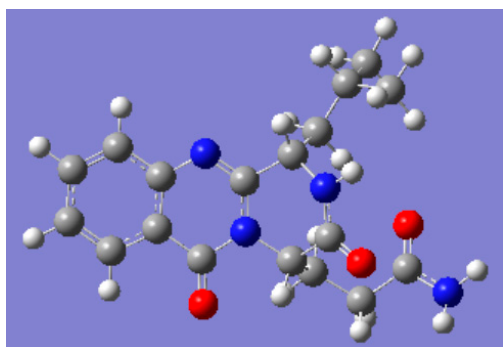
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.335907	-0.172053	-0.234602
2	6	0	-5.126845	-1.511593	-0.607412
3	6	0	-3.853429	-2.049345	-0.563234
4	6	0	-2.772700	-1.254887	-0.147541
5	6	0	-2.977112	0.087951	0.228853
6	6	0	-4.279927	0.619507	0.178975
7	6	0	-1.423886	-1.813925	-0.097739
8	7	0	-0.429500	-0.895427	0.321119
9	6	0	-0.746763	0.397763	0.686568
10	7	0	-1.939925	0.893826	0.661824
11	6	0	0.955206	-1.419665	0.374310
12	6	0	1.808361	-0.727494	1.433025
13	7	0	1.396704	0.496147	1.840118
14	6	0	0.371866	1.299395	1.172308
15	8	0	-1.124649	-2.962814	-0.382757
16	1	0	-0.088155	1.928533	1.934594
17	6	0	0.992071	2.208346	0.080032
18	6	0	0.054963	3.227134	-0.593427
19	6	0	-0.517406	4.247433	0.399176
20	6	0	0.809288	3.936814	-1.726630
21	8	0	2.794898	-1.290543	1.887519
22	1	0	0.866136	-2.454956	0.703818
23	6	0	1.585768	-1.430276	-1.040077
24	6	0	3.030881	-1.934102	-1.096774
25	6	0	4.060326	-0.846188	-0.794154
26	8	0	3.869175	0.323860	-1.088709
27	7	0	5.243949	-1.285832	-0.272297
28	1	0	-6.337453	0.244732	-0.270224



29	1	0	-5.964609	-2.121292	-0.928338
30	1	0	-3.662876	-3.078727	-0.843752
31	1	0	-4.426657	1.652545	0.473453
32	1	0	2.013205	0.961069	2.493740
33	1	0	1.815950	2.750099	0.562633
34	1	0	1.470955	1.579774	-0.675120
35	1	0	-0.786923	2.682887	-1.036114
36	1	0	-1.130608	4.990577	-0.118914
37	1	0	-1.153386	3.777616	1.152764
38	1	0	0.284070	4.789514	0.914976
39	1	0	0.156636	4.639647	-2.252626
40	1	0	1.196670	3.225919	-2.461866
41	1	0	1.662144	4.505115	-1.339561
42	1	0	0.952335	-2.077764	-1.648228
43	1	0	1.547712	-0.430423	-1.475884
44	1	0	3.176319	-2.805050	-0.453133
45	1	0	3.241939	-2.266552	-2.119760
46	1	0	5.255051	-2.138072	0.266101
47	1	0	5.889041	-0.569102	0.027729

---

**Table S22.** The coordinate for the conformer C6 of compound **5** for NMR calculation.

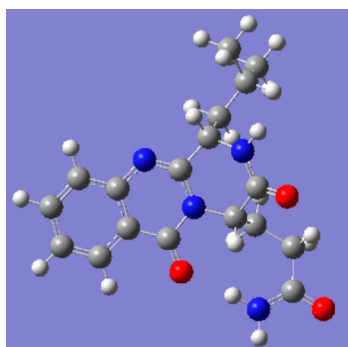
E=-1144.2438034 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.505320	0.950561	-0.322466
2	6	0	-5.600535	-0.444346	-0.473306
3	6	0	-4.470169	-1.230626	-0.345221
4	6	0	-3.230440	-0.633604	-0.064422
5	6	0	-3.130266	0.764313	0.091993
6	6	0	-4.290910	1.550089	-0.042921
7	6	0	-2.028943	-1.455261	0.059419
8	7	0	-0.849749	-0.718290	0.331619
9	6	0	-0.872931	0.650697	0.494303
10	7	0	-1.931077	1.384426	0.390042
11	6	0	0.403336	-1.500680	0.424036
12	6	0	1.419043	-0.868003	1.368882
13	7	0	1.301797	0.462937	1.581133
14	6	0	0.430251	1.358292	0.820227
15	8	0	-1.986150	-2.669689	-0.059496
16	1	0	0.137606	2.170714	1.488600
17	6	0	1.092824	1.975075	-0.442386
18	6	0	1.894091	3.268918	-0.196319
19	6	0	3.051803	3.098381	0.797767
20	6	0	2.416624	3.804961	-1.536294
21	8	0	2.275495	-1.562062	1.900405
22	1	0	0.128243	-2.452249	0.878057
23	6	0	0.963855	-1.791385	-0.989899
24	6	0	2.275882	-2.580448	-1.008820
25	6	0	3.513069	-1.707198	-0.805606
26	8	0	3.545726	-0.535700	-1.153184
27	7	0	4.608984	-2.350700	-0.307428
28	1	0	-6.395426	1.563214	-0.424863

29	1	0	-6.560160	-0.901266	-0.689956
30	1	0	-4.513361	-2.307767	-0.457414
31	1	0	-4.202701	2.623624	0.080254
32	1	0	2.030136	0.871311	2.150021
33	1	0	1.742548	1.226998	-0.906139
34	1	0	0.295641	2.206281	-1.155534
35	1	0	1.199636	4.014563	0.216272
36	1	0	3.650860	4.012078	0.852222
37	1	0	2.702488	2.901098	1.816943
38	1	0	3.715849	2.281968	0.497806
39	1	0	2.938186	4.757350	-1.403040
40	1	0	1.602429	3.969468	-2.248348
41	1	0	3.120408	3.101385	-1.992242
42	1	0	0.193125	-2.361330	-1.511710
43	1	0	1.106729	-0.856681	-1.535343
44	1	0	2.255478	-3.402197	-0.288301
45	1	0	2.391387	-3.039647	-1.997543
46	1	0	4.484428	-3.177611	0.255300
47	1	0	5.398621	-1.768718	-0.068219

---

**Table S23.** The coordinate for the conformer C7 of compound 5 for NMR calculation.

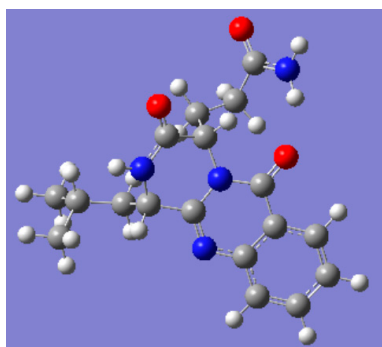
E=-1144.2437512 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.304605	-0.387507	-0.291140
2	6	0	-4.934745	-1.742444	-0.369942
3	6	0	-3.609397	-2.106213	-0.222085
4	6	0	-2.636043	-1.119236	0.007846
5	6	0	-3.002039	0.240194	0.088145
6	6	0	-4.356253	0.592913	-0.065267
7	6	0	-1.234731	-1.486126	0.161770
8	7	0	-0.361459	-0.402706	0.391490
9	6	0	-0.831804	0.894422	0.456391
10	7	0	-2.070270	1.233336	0.320289
11	6	0	1.072353	-0.751560	0.563058
12	6	0	1.841740	0.280843	1.382465
13	7	0	1.329117	1.538142	1.413636
14	6	0	0.164787	2.014352	0.675860
15	8	0	-0.798865	-2.631342	0.100591
16	1	0	-0.355314	2.730505	1.313249
17	6	0	0.514181	2.712544	-0.660817
18	6	0	1.454407	3.925896	-0.540815
19	6	0	1.813234	4.441873	-1.941056
20	6	0	0.858780	5.053619	0.313270
21	8	0	2.868673	-0.031802	1.960186
22	1	0	1.089366	-1.665717	1.154470
23	6	0	1.753556	-1.013290	-0.807728
24	6	0	3.081886	-1.779890	-0.740098
25	6	0	3.075147	-3.297295	-0.528973
26	8	0	4.131067	-3.910822	-0.564603
27	7	0	1.882045	-3.917653	-0.326602
28	1	0	-6.346893	-0.108853	-0.408469

29	1	0	-5.690465	-2.499969	-0.546805
30	1	0	-3.296943	-3.142273	-0.278427
31	1	0	-4.626696	1.640686	-0.000336
32	1	0	1.868723	2.205494	1.947863
33	1	0	0.963026	1.977464	-1.337598
34	1	0	-0.430502	3.024735	-1.118821
35	1	0	2.387155	3.590718	-0.067967
36	1	0	2.517577	5.276404	-1.884864
37	1	0	2.274813	3.660869	-2.552208
38	1	0	0.923451	4.796317	-2.472193
39	1	0	1.534153	5.912811	0.346255
40	1	0	0.676327	4.753534	1.349634
41	1	0	-0.094752	5.400076	-0.099344
42	1	0	1.041931	-1.544034	-1.444587
43	1	0	1.932909	-0.050188	-1.295555
44	1	0	3.621469	-1.630175	-1.679719
45	1	0	3.727452	-1.366017	0.038122
46	1	0	0.987600	-3.452719	-0.234522
47	1	0	1.909871	-4.914334	-0.174561

---

**Table S24.** The coordinate for the conformer C8 of compound 5 for NMR calculation.

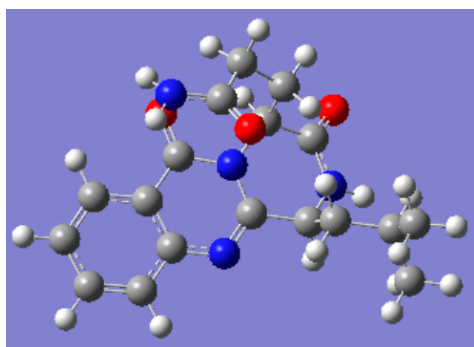
E=-1144.2437043 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	3.034693	-4.158045	-0.461293
2	6	0	4.012723	-3.161583	-0.290364
3	6	0	3.634198	-1.866514	0.010366
4	6	0	2.271516	-1.549249	0.140927
5	6	0	1.288874	-2.546411	-0.022096
6	6	0	1.691139	-3.860602	-0.326960
7	6	0	1.858047	-0.180893	0.428862
8	7	0	0.460723	0.017012	0.470240
9	6	0	-0.409654	-1.055678	0.384189
10	7	0	-0.054214	-2.273758	0.145125
11	6	0	-0.034400	1.406536	0.670434
12	6	0	-1.294783	1.447579	1.534758
13	7	0	-2.097768	0.353280	1.469533
14	6	0	-1.890170	-0.818995	0.626064
15	8	0	2.632864	0.751960	0.609700
16	1	0	-2.204067	-1.689673	1.203710
17	6	0	-2.703144	-0.803949	-0.690006
18	6	0	-4.229930	-0.710781	-0.511037
19	6	0	-4.902573	-0.572610	-1.883797
20	6	0	-4.811507	-1.907124	0.255090
21	8	0	-1.551911	2.425178	2.213098
22	1	0	0.733080	1.923406	1.242468
23	6	0	-0.290289	2.159401	-0.658796
24	6	0	0.958730	2.606221	-1.442827
25	6	0	1.604122	3.856762	-0.841580
26	8	0	1.162343	4.970418	-1.079855
27	7	0	2.679332	3.638089	-0.039052
28	1	0	3.338902	-5.172785	-0.696999

29	1	0	5.063015	-3.411038	-0.394877
30	1	0	4.367184	-1.080060	0.145768
31	1	0	0.926377	-4.619750	-0.445507
32	1	0	-2.935269	0.400592	2.033388
33	1	0	-2.367295	0.033038	-1.309806
34	1	0	-2.453601	-1.720766	-1.234966
35	1	0	-4.452404	0.203642	0.054800
36	1	0	-5.985807	-0.462470	-1.782751
37	1	0	-4.530678	0.300143	-2.428191
38	1	0	-4.717192	-1.455443	-2.505014
39	1	0	-5.898924	-1.824714	0.334343
40	1	0	-4.424161	-1.991063	1.274959
41	1	0	-4.589366	-2.849013	-0.257725
42	1	0	-0.905607	1.528299	-1.305244
43	1	0	-0.887539	3.044535	-0.423516
44	1	0	1.677233	1.790702	-1.542897
45	1	0	0.639951	2.890201	-2.447845
46	1	0	2.937675	2.704337	0.251001
47	1	0	3.080068	4.430691	0.438830

---

**Table S25.** The coordinate for the conformer C9 of compound 5 for NMR calculation.

E=-1144.2435726 a.u.

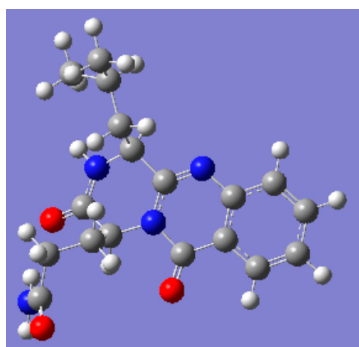
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-4.099350	-2.722146	0.254596
2	6	0	-4.732378	-1.503174	-0.048861
3	6	0	-3.974844	-0.402304	-0.404120
4	6	0	-2.574697	-0.503964	-0.460587
5	6	0	-1.937500	-1.726379	-0.168827
6	6	0	-2.723016	-2.837597	0.193546
7	6	0	-1.765132	0.659675	-0.809686
8	7	0	-0.370170	0.424162	-0.810795
9	6	0	0.140078	-0.837448	-0.568458
10	7	0	-0.567680	-1.871861	-0.257543
11	6	0	0.501484	1.565871	-1.193479
12	6	0	1.737330	1.113183	-1.969544
13	7	0	2.209680	-0.120317	-1.660209
14	6	0	1.636937	-1.044550	-0.682816
15	8	0	-2.213170	1.768795	-1.070304
16	1	0	1.741552	-2.048400	-1.096355
17	6	0	2.314430	-1.020912	0.708183
18	6	0	3.839266	-1.229016	0.694493
19	6	0	4.388730	-1.095865	2.121407
20	6	0	4.251153	-2.575355	0.082003
21	8	0	2.272503	1.855336	-2.776003
22	1	0	-0.094118	2.170011	-1.875193
23	6	0	0.979248	2.465232	-0.027346
24	6	0	-0.113311	3.087202	0.846811
25	6	0	-0.633838	2.135749	1.922083
26	8	0	0.104739	1.369579	2.522921
27	7	0	-1.960232	2.254969	2.216843
28	1	0	-4.698687	-3.582753	0.534200



29	1	0	-5.813786	-1.429736	-0.006073
30	1	0	-4.437545	0.547158	-0.647520
31	1	0	-2.220675	-3.772287	0.415555
32	1	0	3.047168	-0.398215	-2.153001
33	1	0	2.074678	-0.084157	1.218121
34	1	0	1.841942	-1.812500	1.299887
35	1	0	4.288875	-0.426658	0.094235
36	1	0	5.478956	-1.185025	2.134676
37	1	0	4.126503	-0.130902	2.563695
38	1	0	3.984910	-1.877269	2.773864
39	1	0	5.336476	-2.704866	0.120398
40	1	0	3.956322	-2.676384	-0.967385
41	1	0	3.800773	-3.410946	0.628614
42	1	0	1.670306	1.918700	0.617130
43	1	0	1.556375	3.262435	-0.502568
44	1	0	0.327307	3.929367	1.393096
45	1	0	-0.924306	3.491570	0.239717
46	1	0	-2.593787	2.686468	1.562782
47	1	0	-2.350538	1.606835	2.884748

---

**Table S26.** The coordinate for the conformer C10 of compound 5 for NMR calculation.

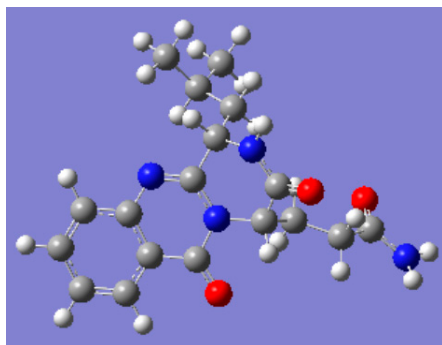
E=-1144.2431876 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	5.417479	-0.153039	0.190116
2	6	0	5.149297	-1.530259	0.278378
3	6	0	3.849858	-1.989429	0.161691
4	6	0	2.802205	-1.077601	-0.043927
5	6	0	3.066445	0.304386	-0.135792
6	6	0	4.394281	0.754898	-0.015434
7	6	0	1.425409	-1.554608	-0.155377
8	7	0	0.467401	-0.520165	-0.333507
9	6	0	0.845410	0.799889	-0.441947
10	7	0	2.060364	1.228312	-0.357326
11	6	0	-0.949491	-0.945722	-0.381998
12	6	0	-1.792548	-0.027087	-1.256130
13	7	0	-1.369904	1.250303	-1.380789
14	6	0	-0.230044	1.845643	-0.684933
15	8	0	1.070754	-2.718496	-0.098352
16	1	0	0.234206	2.555856	-1.372216
17	6	0	-0.608876	2.609374	0.613910
18	6	0	-0.997306	4.088128	0.419935
19	6	0	-2.186064	4.300948	-0.527540
20	6	0	-1.277433	4.729142	1.785793
21	8	0	-2.811852	-0.439348	-1.802067
22	1	0	-0.951395	-1.923269	-0.861479
23	6	0	-1.517539	-1.102349	1.053181
24	6	0	-2.988492	-1.547414	1.149840
25	6	0	-3.229085	-2.980014	0.666050
26	8	0	-3.147088	-3.933196	1.424493
27	7	0	-3.538276	-3.099972	-0.653606

28	1	0	6.439025	0.201697	0.282718
29	1	0	5.961766	-2.230751	0.438344
30	1	0	3.613496	-3.045082	0.227397
31	1	0	4.586488	1.819290	-0.089919
32	1	0	-1.965832	1.848933	-1.935615
33	1	0	-1.423977	2.075465	1.114924
34	1	0	0.253667	2.574444	1.285002
35	1	0	-0.125689	4.596247	-0.015050
36	1	0	-2.468624	5.356735	-0.560301
37	1	0	-1.958055	4.012725	-1.558760
38	1	0	-3.066575	3.736576	-0.202948
39	1	0	-1.478426	5.799323	1.685343
40	1	0	-0.429120	4.613848	2.466466
41	1	0	-2.151273	4.274452	2.264873
42	1	0	-0.886776	-1.832116	1.565906
43	1	0	-1.402615	-0.152562	1.584339
44	1	0	-3.271866	-1.532531	2.203708
45	1	0	-3.639729	-0.847559	0.620593
46	1	0	-3.497822	-2.304424	-1.279614
47	1	0	-3.643515	-4.027641	-1.034602

---

**Table S27.** The coordinate for the conformer C11 of compound 5 for NMR calculation.

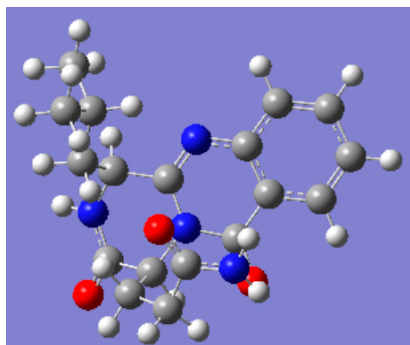
E=-1144.2429465 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-4.965639	-1.549227	-0.721214
2	6	0	-4.325516	-2.797265	-0.819455
3	6	0	-2.984105	-2.911566	-0.502866
4	6	0	-2.265066	-1.779902	-0.085595
5	6	0	-2.902514	-0.526906	0.016200
6	6	0	-4.268738	-0.428010	-0.308911
7	6	0	-0.848086	-1.891018	0.251656
8	7	0	-0.245451	-0.668566	0.642389
9	6	0	-0.971793	0.501894	0.718710
10	7	0	-2.229187	0.605335	0.437997
11	6	0	1.190753	-0.749024	1.000870
12	6	0	1.604713	0.307878	2.019616
13	7	0	0.825969	1.417006	2.090164
14	6	0	-0.255101	1.760175	1.167228
15	8	0	-0.191748	-2.919630	0.217455
16	1	0	-0.996911	2.320458	1.736788
17	6	0	0.252286	2.647678	0.000330
18	6	0	-0.812030	3.254432	-0.931988
19	6	0	-1.779527	4.188205	-0.193202
20	6	0	-0.116057	3.993964	-2.083276
21	8	0	2.591928	0.141614	2.719888
22	1	0	1.315394	-1.713833	1.491689
23	6	0	2.084336	-0.723212	-0.261465
24	6	0	3.510033	-1.208174	0.002530
25	6	0	4.413230	-0.989062	-1.206807
26	8	0	4.217416	-0.112633	-2.032412
27	7	0	5.481136	-1.836353	-1.290185

28	1	0	-6.018606	-1.465138	-0.970708
29	1	0	-4.884804	-3.668444	-1.143147
30	1	0	-2.465290	-3.860823	-0.568633
31	1	0	-4.749935	0.539907	-0.225014
32	1	0	1.148265	2.127510	2.734019
33	1	0	0.828536	3.462567	0.458101
34	1	0	0.969470	2.072650	-0.592898
35	1	0	-1.399469	2.435490	-1.361146
36	1	0	-2.481955	4.648946	-0.893755
37	1	0	-2.375036	3.658486	0.553657
38	1	0	-1.241623	5.001021	0.308895
39	1	0	-0.848037	4.400287	-2.787020
40	1	0	0.552162	3.333947	-2.643820
41	1	0	0.483040	4.832791	-1.711759
42	1	0	1.616242	-1.368326	-1.007735
43	1	0	2.120210	0.280195	-0.688261
44	1	0	3.948433	-0.664478	0.845786
45	1	0	3.503030	-2.264643	0.290816
46	1	0	5.677993	-2.527785	-0.586106
47	1	0	6.163773	-1.682728	-2.015923

---

**Table S28.** The coordinate for the conformer C12 of compound 5 for NMR calculation.

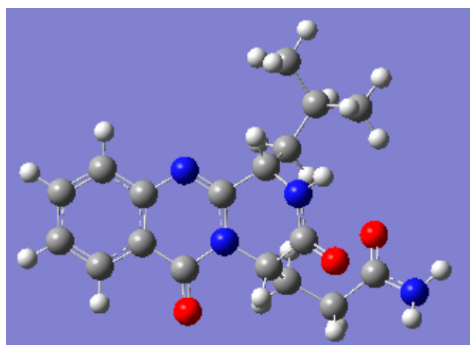
E=-1144.2429299 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	4.003019	2.430755	-0.327753
2	6	0	4.587186	1.152651	-0.267477
3	6	0	3.792671	0.025340	-0.368957
4	6	0	2.403575	0.159586	-0.530040
5	6	0	1.815770	1.437786	-0.600755
6	6	0	2.638233	2.575892	-0.495012
7	6	0	1.553827	-1.024300	-0.620798
8	7	0	0.173871	-0.746739	-0.760524
9	6	0	-0.286304	0.553349	-0.870697
10	7	0	0.459400	1.605869	-0.798910
11	6	0	-0.727972	-1.917662	-0.919197
12	6	0	-1.892862	-1.625168	-1.864502
13	7	0	-2.301792	-0.332848	-1.914804
14	6	0	-1.765916	0.769862	-1.115271
15	8	0	1.958610	-2.178483	-0.571522
16	1	0	-1.817289	1.662573	-1.738625
17	6	0	-2.578171	1.021932	0.182468
18	6	0	-2.338680	2.377204	0.873967
19	6	0	-2.808870	3.564846	0.023700
20	6	0	-3.033141	2.378476	2.242904
21	8	0	-2.426345	-2.525831	-2.490491
22	1	0	-0.121718	-2.685007	-1.397197
23	6	0	-1.312689	-2.502935	0.390288
24	6	0	-0.304207	-2.905239	1.468946
25	6	0	0.193126	-1.722847	2.297424
26	8	0	-0.533438	-0.786186	2.596935
27	7	0	1.478247	-1.820094	2.742279

28	1	0	4.631187	3.312189	-0.247289
29	1	0	5.660417	1.053565	-0.144849
30	1	0	4.218058	-0.970878	-0.332325
31	1	0	2.173700	3.553782	-0.552195
32	1	0	-3.110599	-0.155482	-2.495258
33	1	0	-3.639406	0.938861	-0.085990
34	1	0	-2.370379	0.222015	0.896411
35	1	0	-1.262629	2.486690	1.044348
36	1	0	-2.658111	4.507318	0.558004
37	1	0	-2.264619	3.646130	-0.920986
38	1	0	-3.878027	3.488989	-0.207143
39	1	0	-2.860393	3.321796	2.769601
40	1	0	-2.661817	1.568988	2.876575
41	1	0	-4.117567	2.255487	2.139463
42	1	0	-2.032930	-1.810791	0.829108
43	1	0	-1.876681	-3.385445	0.078337
44	1	0	-0.813798	-3.566017	2.180274
45	1	0	0.524185	-3.474672	1.045094
46	1	0	2.129450	-2.449504	2.300983
47	1	0	1.858695	-1.037059	3.252580

---

**Table S29.** The coordinate for the conformer C13 of compound 5 for NMR calculation.

E=-1144.2425119 a.u.

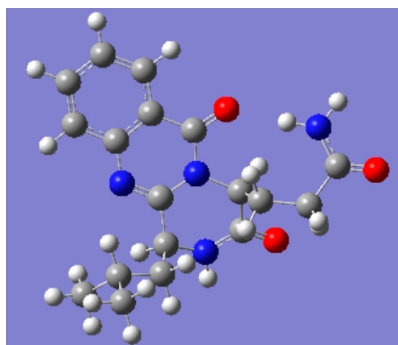
Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.487627	0.679675	-0.366516
2	6	0	-5.532100	-0.725868	-0.373604
3	6	0	-4.370753	-1.453646	-0.189221
4	6	0	-3.150052	-0.786701	0.004209
5	6	0	-3.100544	0.622494	0.015310
6	6	0	-4.292420	1.348035	-0.174066
7	6	0	-1.915998	-1.547153	0.185568
8	7	0	-0.760259	-0.744148	0.352264
9	6	0	-0.833046	0.632718	0.375307
10	7	0	-1.920743	1.312575	0.221791
11	6	0	0.523694	-1.468526	0.488339
12	6	0	1.539824	-0.705842	1.331050
13	7	0	1.376497	0.634312	1.411620
14	6	0	0.447085	1.418204	0.597645
15	8	0	-1.829384	-2.765043	0.192191
16	1	0	0.136614	2.273312	1.199015
17	6	0	1.051879	1.932540	-0.737639
18	6	0	1.887950	3.229368	-0.654744
19	6	0	1.040563	4.457670	-0.296028
20	6	0	3.121259	3.119851	0.254007
21	8	0	2.437597	-1.310140	1.902948
22	1	0	0.296985	-2.376071	1.047115
23	6	0	1.055994	-1.886336	-0.904522
24	6	0	2.401713	-2.616489	-0.885609
25	6	0	3.603166	-1.674556	-0.825892
26	8	0	3.572419	-0.550924	-1.306501
27	7	0	4.741067	-2.209476	-0.294491



28	1	0	-6.401996	1.246117	-0.512142
29	1	0	-6.477050	-1.236862	-0.523806
30	1	0	-4.374943	-2.537461	-0.190923
31	1	0	-4.242910	2.431017	-0.161930
32	1	0	2.101035	1.124544	1.916939
33	1	0	1.668625	1.139956	-1.168699
34	1	0	0.221010	2.114901	-1.426335
35	1	0	2.261326	3.381458	-1.675687
36	1	0	1.636599	5.372702	-0.362236
37	1	0	0.186010	4.567644	-0.969707
38	1	0	0.647796	4.407260	0.725077
39	1	0	3.777170	3.984417	0.116118
40	1	0	3.703850	2.220599	0.038119
41	1	0	2.848689	3.110819	1.316582
42	1	0	0.297198	-2.540667	-1.337062
43	1	0	1.139152	-1.009828	-1.549637
44	1	0	2.443170	-3.353926	-0.079921
45	1	0	2.506310	-3.176446	-1.822257
46	1	0	4.669883	-2.968295	0.365044
47	1	0	5.510957	-1.571820	-0.152605

---

**Table S30.** The coordinate for the conformer C14 of compound 5 for NMR calculation.

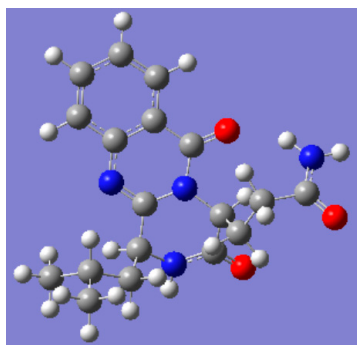
E=-1144.242472 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-3.907904	3.257637	0.085235
2	6	0	-2.818571	4.116496	0.317909
3	6	0	-1.526877	3.628770	0.248745
4	6	0	-1.307815	2.274312	-0.053586
5	6	0	-2.396309	1.409704	-0.287270
6	6	0	-3.705025	1.922959	-0.213845
7	6	0	0.047171	1.745381	-0.133865
8	7	0	0.129732	0.373376	-0.447970
9	6	0	-1.008525	-0.384789	-0.653951
10	7	0	-2.214451	0.073922	-0.592057
11	6	0	1.503758	-0.176972	-0.584449
12	6	0	1.566971	-1.396294	-1.501513
13	7	0	0.408407	-2.081564	-1.683214
14	6	0	-0.840453	-1.858441	-0.960466
15	8	0	1.066850	2.402786	0.048147
16	1	0	-1.655400	-2.102997	-1.641661
17	6	0	-0.948041	-2.767131	0.292752
18	6	0	-2.318766	-2.835461	0.991224
19	6	0	-3.411475	-3.414827	0.083268
20	6	0	-2.192269	-3.660019	2.280101
21	8	0	2.618093	-1.715043	-2.029961
22	1	0	2.084088	0.594927	-1.088048
23	6	0	2.137652	-0.466422	0.803236
24	6	0	3.665459	-0.619725	0.804632
25	6	0	4.557203	0.623657	0.729901
26	8	0	5.769229	0.496542	0.816367
27	7	0	3.963532	1.838964	0.588421

28	1	0	-4.919140	3.647720	0.140053
29	1	0	-2.993863	5.161171	0.550624
30	1	0	-0.670174	4.269233	0.422074
31	1	0	-4.534371	1.249584	-0.398242
32	1	0	0.487060	-2.908909	-2.259943
33	1	0	-0.662929	-3.777169	-0.028794
34	1	0	-0.185670	-2.457960	1.015197
35	1	0	-2.616431	-1.818068	1.267543
36	1	0	-4.357889	-3.499862	0.624525
37	1	0	-3.599695	-2.787847	-0.791154
38	1	0	-3.145496	-4.419028	-0.266740
39	1	0	-3.143721	-3.696219	2.817937
40	1	0	-1.445178	-3.238586	2.959457
41	1	0	-1.898120	-4.693099	2.064472
42	1	0	1.835734	0.331044	1.486150
43	1	0	1.699088	-1.385915	1.202377
44	1	0	3.965318	-1.124415	1.727630
45	1	0	3.987384	-1.276572	-0.006761
46	1	0	2.972047	1.995289	0.457771
47	1	0	4.576342	2.637495	0.524492

---

**Table S31.** The coordinate for the conformer C15 of compound **5** for NMR calculation.

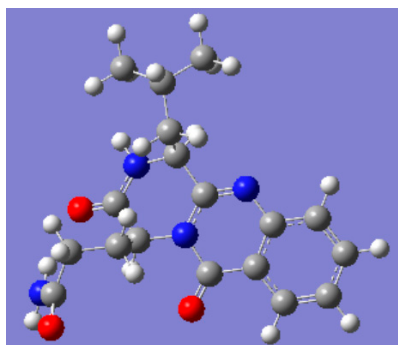
E=-1144.2423185 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-3.004467	3.915670	0.240173
2	6	0	-1.749108	4.549880	0.256559
3	6	0	-0.599277	3.812539	0.043120
4	6	0	-0.689009	2.428852	-0.184332
5	6	0	-1.944880	1.791032	-0.207711
6	6	0	-3.106450	2.556363	0.008215
7	6	0	0.517131	1.635175	-0.388761
8	7	0	0.295396	0.248719	-0.536704
9	6	0	-0.991346	-0.258393	-0.623125
10	7	0	-2.066867	0.440698	-0.470702
11	6	0	1.491227	-0.625689	-0.691161
12	6	0	1.239173	-1.780173	-1.663445
13	7	0	-0.051007	-2.188336	-1.784282
14	6	0	-1.173386	-1.724572	-0.973250
15	8	0	1.650912	2.100303	-0.422622
16	1	0	-2.055488	-1.740477	-1.614063
17	6	0	-1.438937	-2.658005	0.236881
18	6	0	-2.736749	-2.422188	1.031090
19	6	0	-3.994563	-2.619816	0.174937
20	6	0	-2.761456	-3.352144	2.252293
21	8	0	2.162577	-2.290570	-2.270406
22	1	0	2.256065	-0.010119	-1.160345
23	6	0	2.025715	-1.178441	0.653739
24	6	0	2.771945	-0.177788	1.557174
25	6	0	4.197406	0.092765	1.070362
26	8	0	5.107035	-0.682932	1.320978
27	7	0	4.369034	1.234796	0.352989

28	1	0	-3.903024	4.500838	0.407290
29	1	0	-1.685014	5.617513	0.436344
30	1	0	0.379543	4.277231	0.050401
31	1	0	-4.067276	2.054985	-0.017695
32	1	0	-0.197152	-2.981279	-2.394903
33	1	0	-1.448840	-3.681320	-0.160666
34	1	0	-0.581923	-2.613395	0.915167
35	1	0	-2.736776	-1.387957	1.392233
36	1	0	-4.896823	-2.504843	0.782322
37	1	0	-4.061474	-1.891746	-0.636533
38	1	0	-4.023308	-3.624682	-0.262454
39	1	0	-3.656405	-3.181374	2.856982
40	1	0	-1.891889	-3.196958	2.898037
41	1	0	-2.765289	-4.405263	1.950404
42	1	0	1.190095	-1.603084	1.215650
43	1	0	2.701163	-2.006070	0.421333
44	1	0	2.205053	0.747028	1.677912
45	1	0	2.876449	-0.632453	2.544516
46	1	0	3.578327	1.783580	0.043026
47	1	0	5.276089	1.403894	-0.054208

---

**Table S32.** The coordinate for the conformer C16 of compound **5** for NMR calculation.

E=-1144.2418739 a.u.

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	5.246711	-1.142858	0.155390
2	6	0	4.734640	-2.452028	0.135352
3	6	0	3.372945	-2.659008	0.008511
4	6	0	2.506007	-1.559826	-0.099277
5	6	0	3.015031	-0.245194	-0.082931
6	6	0	4.403100	-0.052019	0.046592
7	6	0	1.065003	-1.770431	-0.220356
8	7	0	0.308310	-0.569274	-0.292687
9	6	0	0.918788	0.665702	-0.304408
10	7	0	2.191665	0.859452	-0.209135
11	6	0	-1.162379	-0.729223	-0.346803
12	6	0	-1.831694	0.394923	-1.125823
13	7	0	-1.187637	1.582290	-1.155786
14	6	0	0.050840	1.904828	-0.447908
15	8	0	0.506365	-2.852338	-0.252468
16	1	0	0.629556	2.563665	-1.096527
17	6	0	-0.170255	2.619878	0.913369
18	6	0	-0.387621	4.149090	0.859635
19	6	0	0.878275	4.903521	0.430264
20	6	0	-1.598511	4.582614	0.021527
21	8	0	-2.911307	0.217392	-1.682760
22	1	0	-1.341259	-1.646345	-0.906046
23	6	0	-1.742983	-0.903509	1.081159
24	6	0	-3.268589	-1.091493	1.167673
25	6	0	-3.755870	-2.415258	0.573366
26	8	0	-3.826264	-3.431083	1.246884
27	7	0	-4.100198	-2.365868	-0.742220

28	1	0	6.316100	-0.986493	0.255070
29	1	0	5.408076	-3.298041	0.219928
30	1	0	2.950339	-3.656880	-0.007681
31	1	0	4.783698	0.963130	0.054922
32	1	0	-1.665755	2.319253	-1.655324
33	1	0	-1.021778	2.149731	1.415363
34	1	0	0.709650	2.428386	1.533935
35	1	0	-0.597695	4.438306	1.897629
36	1	0	0.729628	5.984218	0.508961
37	1	0	1.735477	4.638880	1.055162
38	1	0	1.153489	4.694663	-0.608599
39	1	0	-1.808686	5.645060	0.172172
40	1	0	-2.503509	4.028059	0.285730
41	1	0	-1.421257	4.455492	-1.052810
42	1	0	-1.248678	-1.772508	1.521490
43	1	0	-1.461472	-0.035896	1.685653
44	1	0	-3.541395	-1.115422	2.224227
45	1	0	-3.789443	-0.247836	0.708557
46	1	0	-3.928282	-1.538808	-1.301487
47	1	0	-4.362746	-3.226840	-1.196783

---



© 2020 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (<http://creativecommons.org/licenses/by/4.0/>).