

## *Supplementary Material*

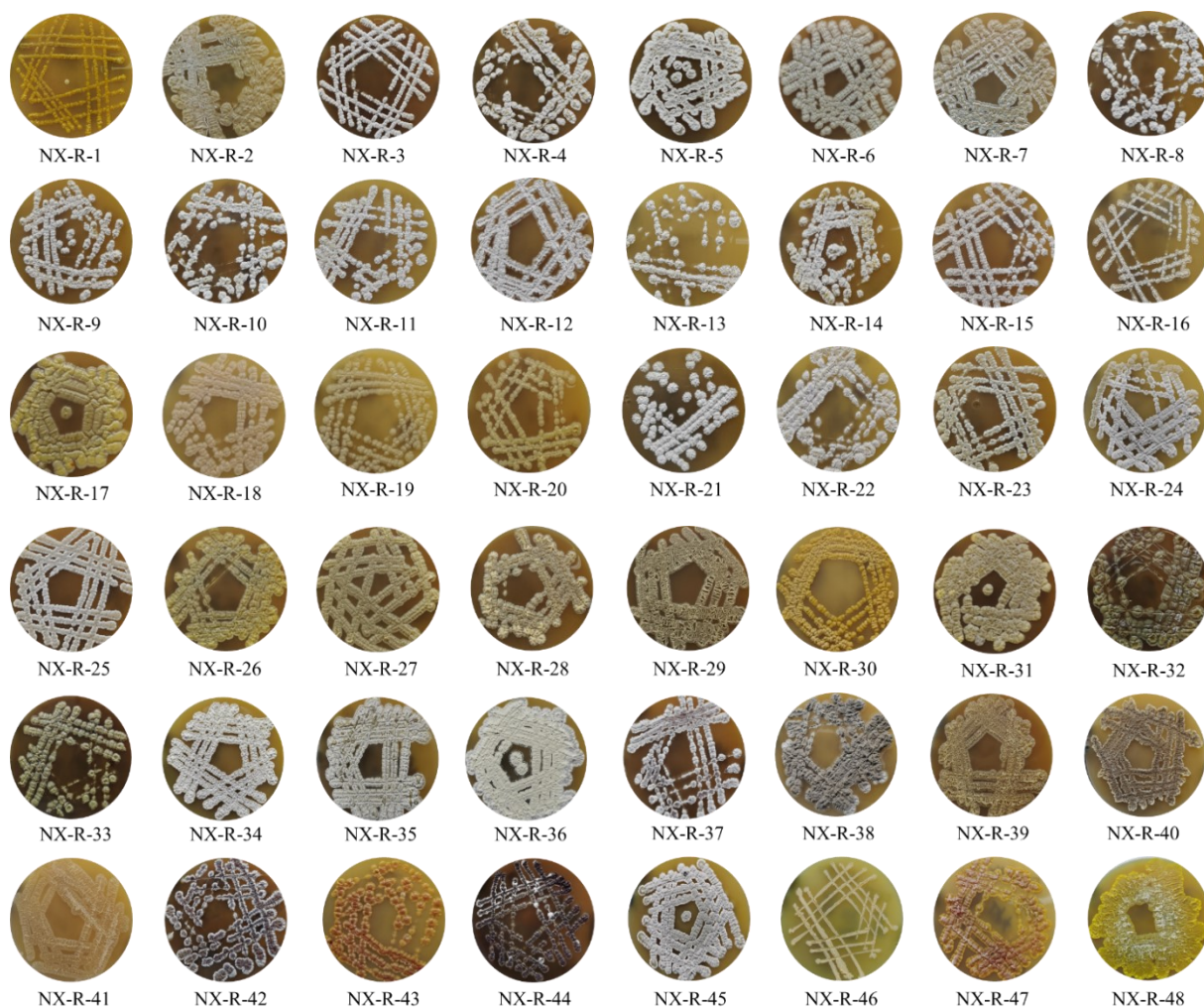
### **Endophytic Microbes from Medicinal Plants in Fenghuang Mountain as a Source of Antibiotics**

Aiping Yang, Yu Hong, Fengjuan Zhou, Ling Zhang, Youjuan Zhu, Chang Wang, Yang Hu, Li Yu, Lihong Chen\* and Xiachang Wang\*

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Lihong Chen: [clh\\_helen@njucm.edu.cn](mailto:clh_helen@njucm.edu.cn)

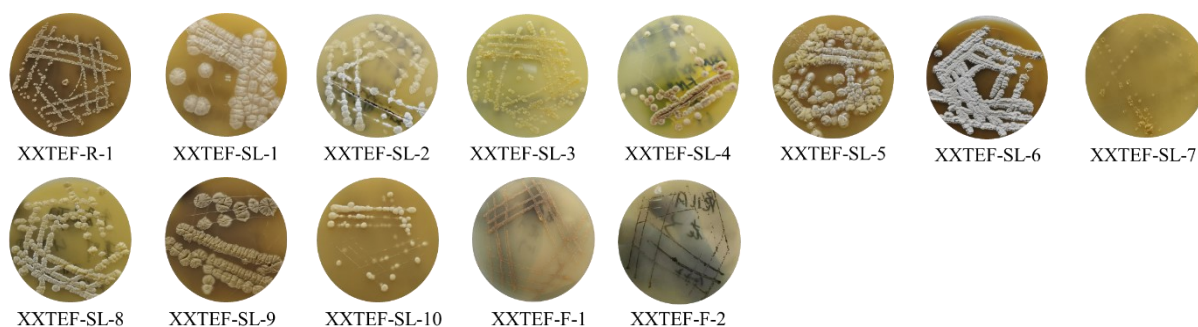
#### **1 Supplementary Figures and Tables**

##### **1.1 Supplementary Figures**

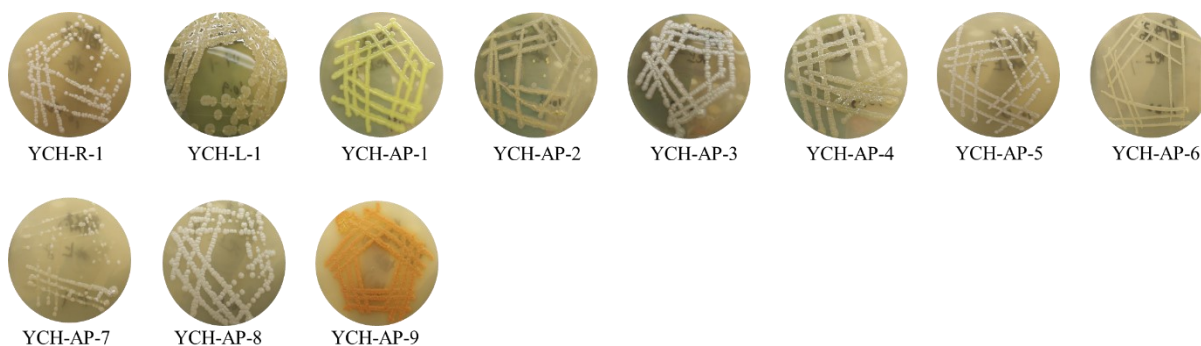




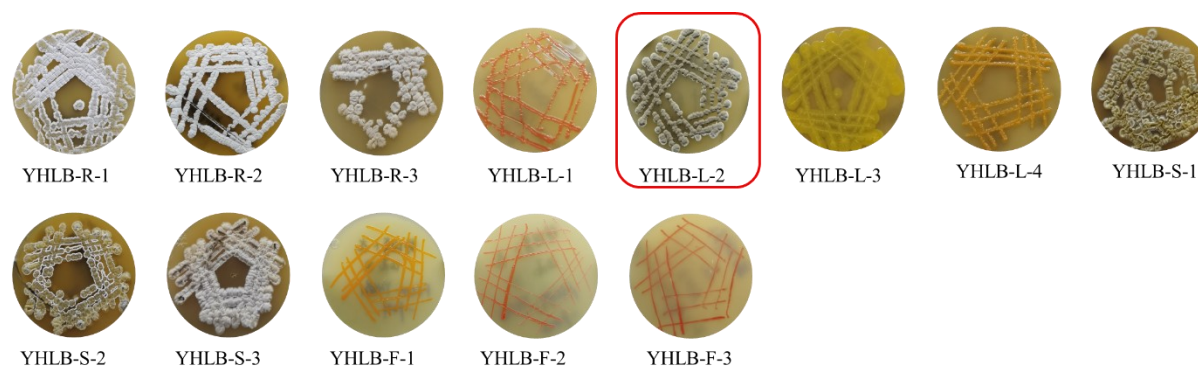
**Supplementary Figure S1.** Morphological features of 124 endophytic strains from *Achyranthes bidentata* (Red box: selected *Penicillium* sp. NX-S-6 for further study).



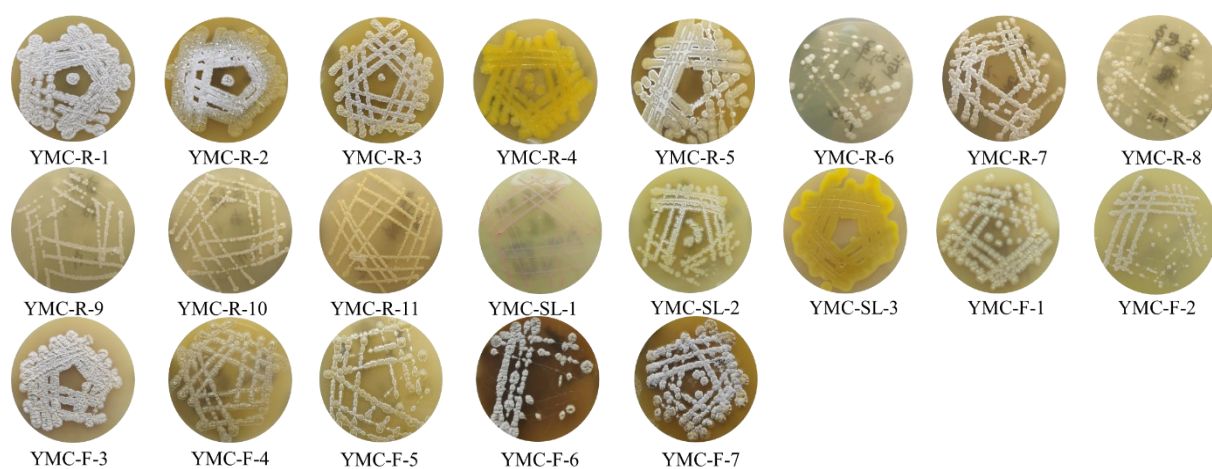
**Supplementary Figure S2.** Morphological features of 13 endophytic strains from *Ainsliaea fragrans*.



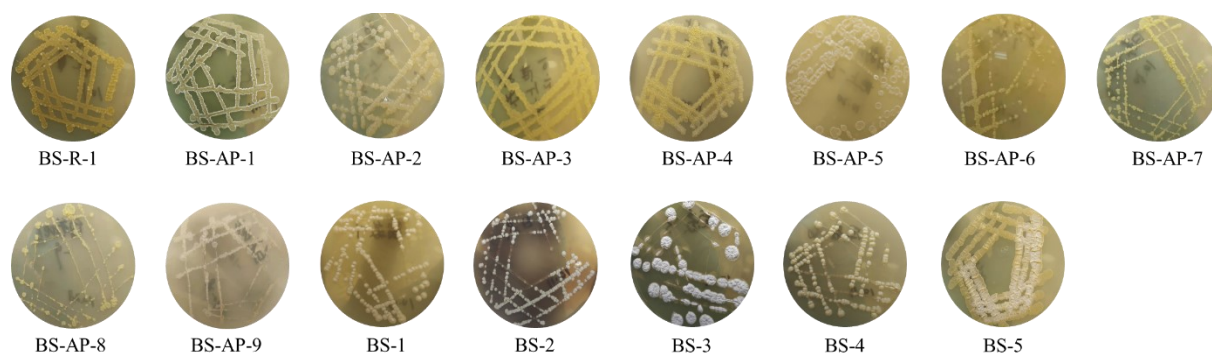
**Supplementary Figure S3.** Morphological features of 11 endophytic strains from *Artemisia capillaris*.



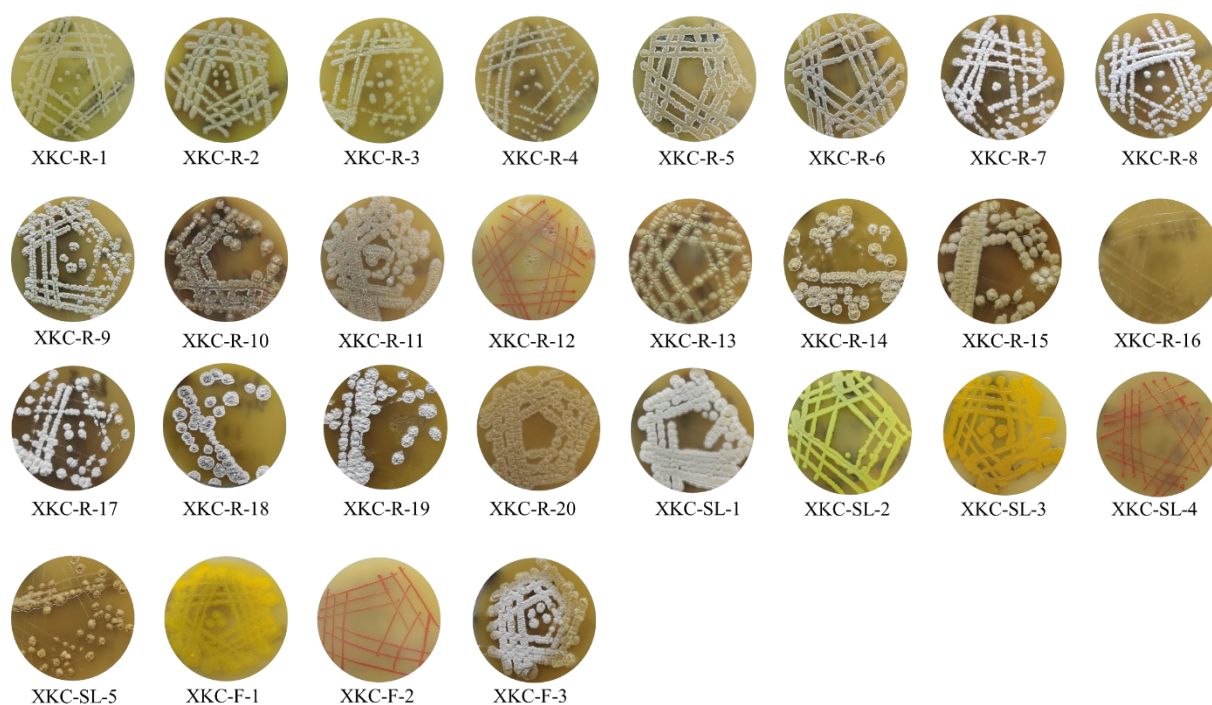
**Supplementary Figure S4.** Morphological features of 13 endophytic strains from *Daucus carota* (Red box: selected *Streptomyces* sp. YHLB-L-2 for further study).



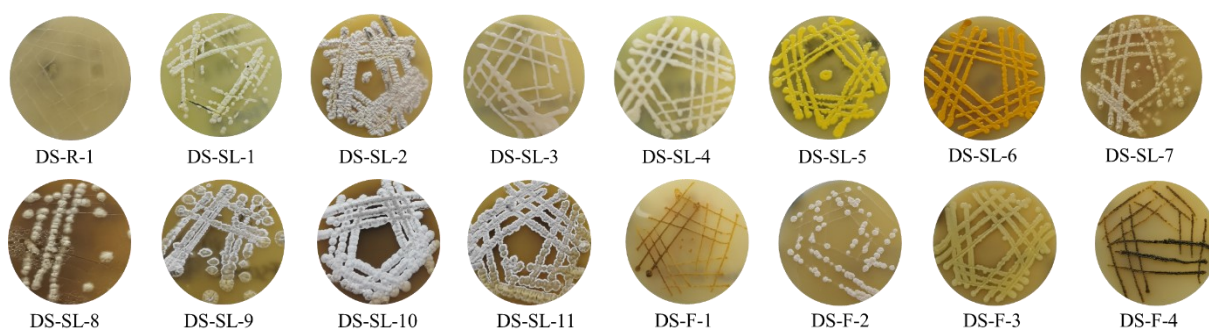
**Supplementary Figure S5.** Morphological features of 21 endophytic strains from *Leonurus japonicus*.



**Supplementary Figure S6.** Morphological features of 15 endophytic strains from *Perilla frutescens*.



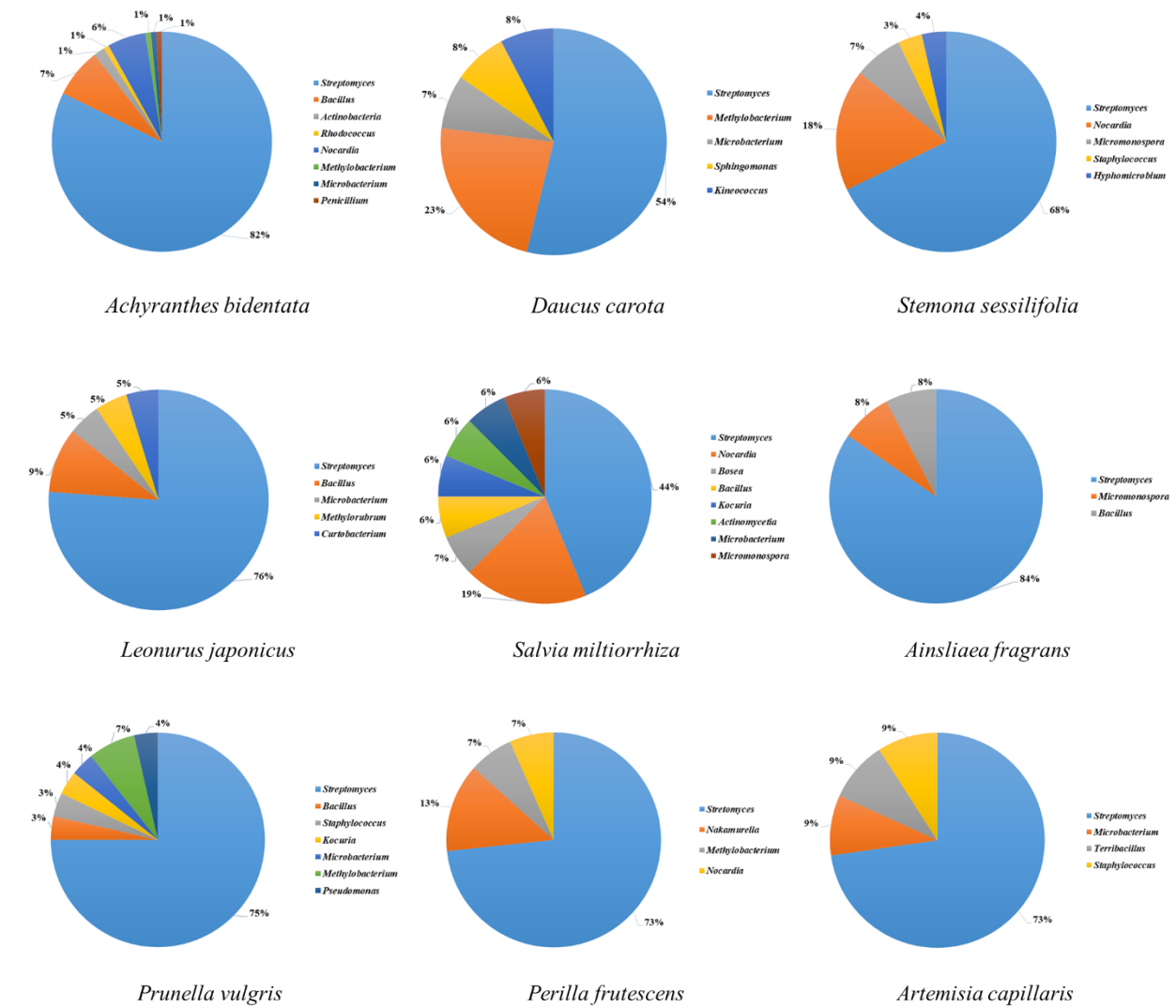
**Supplementary Figure S7.** Morphological features of 28 endophytic strains from *Prunella vulgaris*.



**Supplementary Figure S8.** Morphological features of 16 endophytic strains from *Salvia miltiorrhiza*.



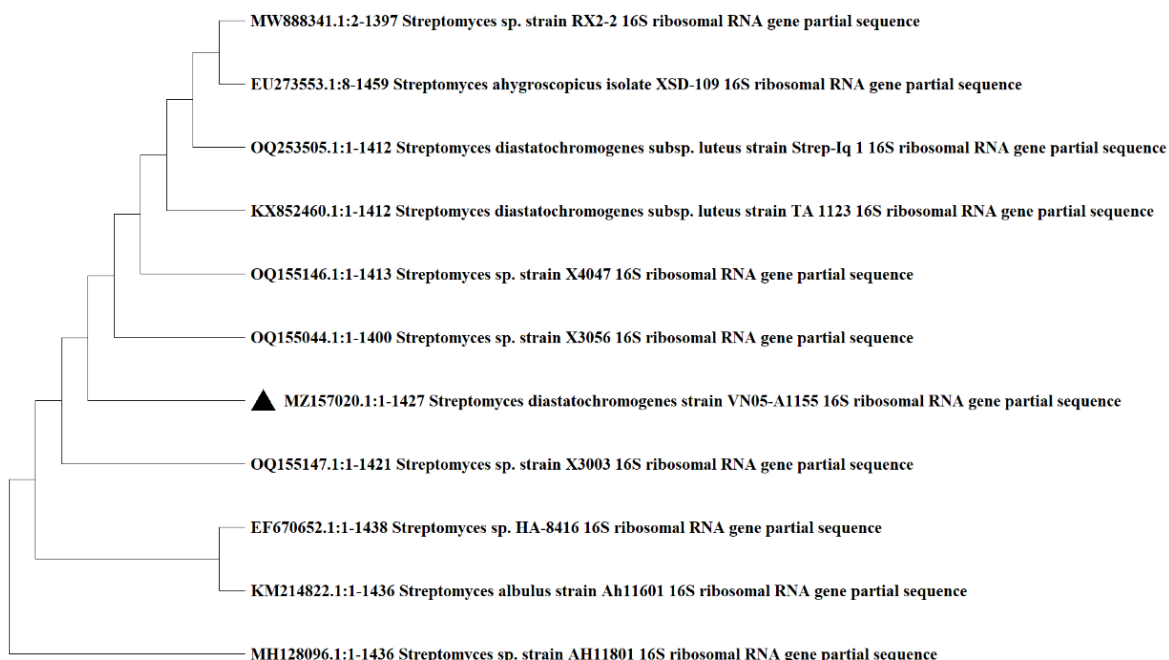
**Supplementary Figure S9.** Morphological features of 28 endophytic strains from *Stemona sessilifolia* (Red box: selected *Streptomyces* sp. ZLBB-S-6 for further study).



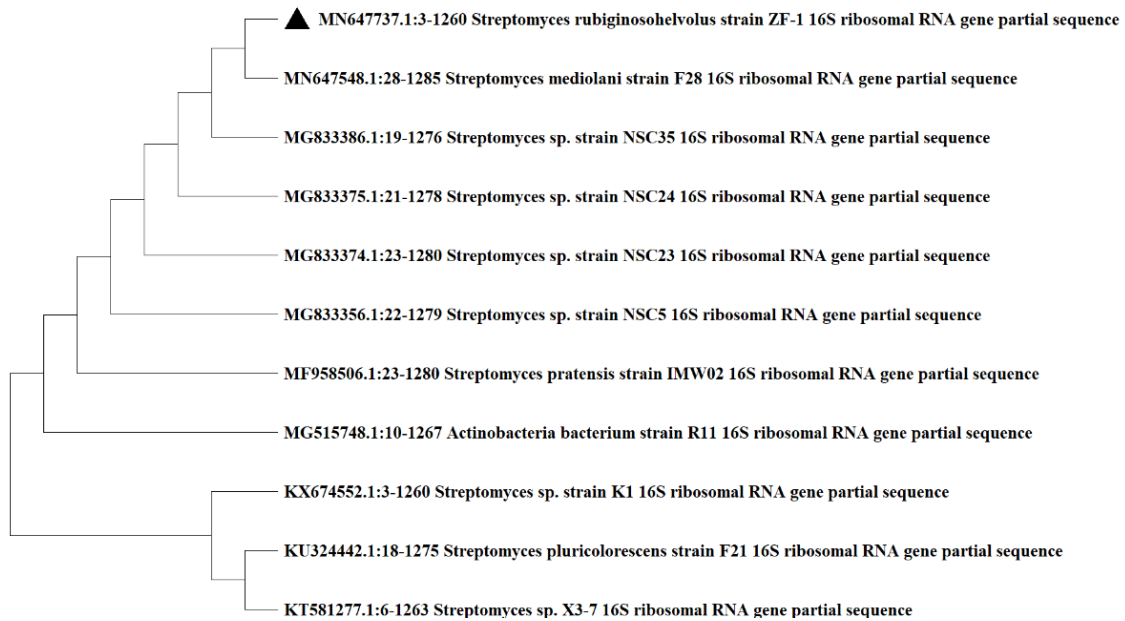
**Supplementary Figure S10.** Diversity analyses of endophytes from nine medicinal plants.



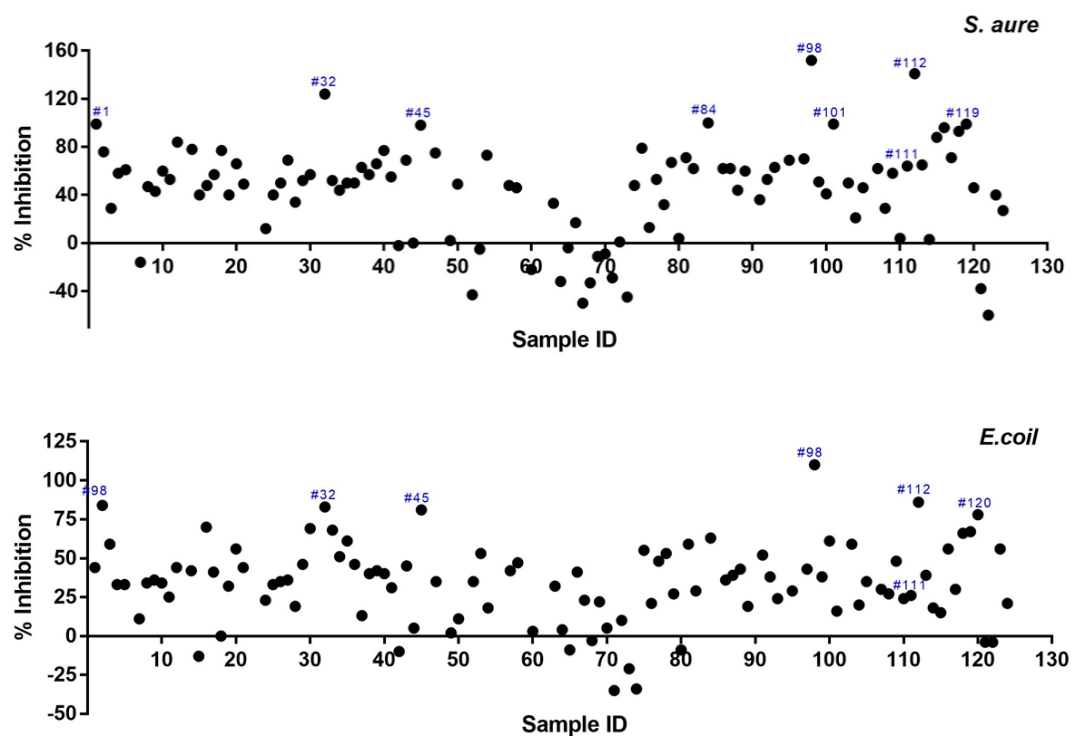
**Supplementary Figure S11.** Evolutionary relationship analysis of *Penicillium* sp. NX-S-6 (▲).



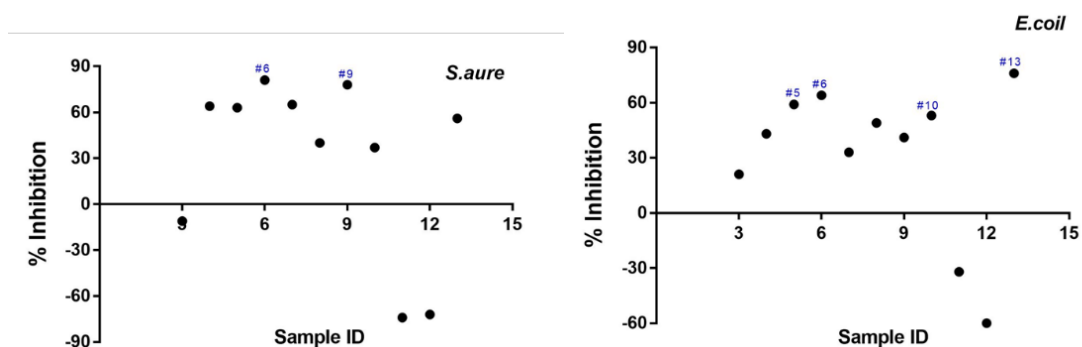
**Supplementary Figure S12.** Evolutionary relationship analysis of *Streptomyces* sp. YHLB-L-2 (▲).



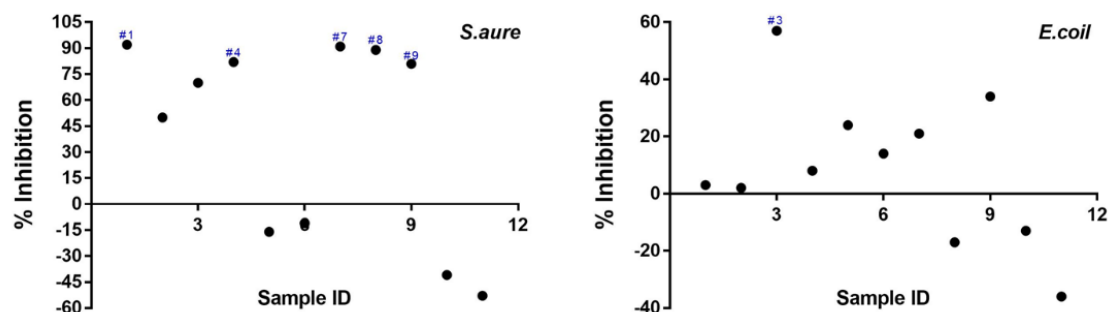
**Supplementary Figure S13.** Evolutionary relationship analysis of *Streptomyces* sp. ZLBB-S-6 (▲).



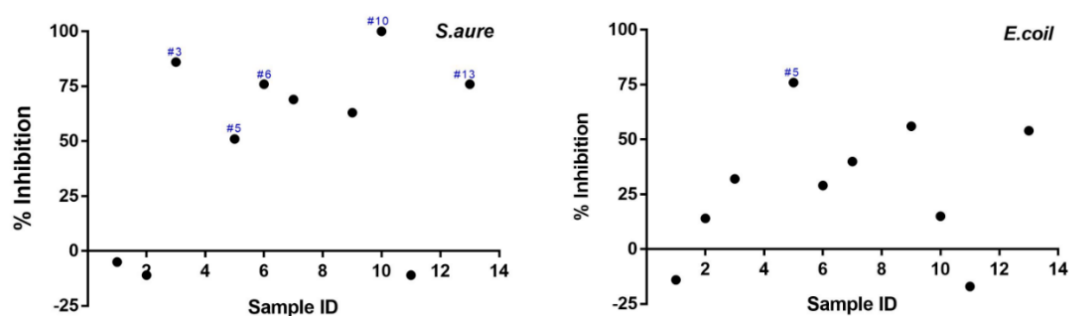
**Supplementary Figure S14.** Scatterplots of 124 strain extracts (1.0 mg/mL) from *Achyranthes bidentata* screened in *S. aureus* and *E. coli* inhibitory assay.



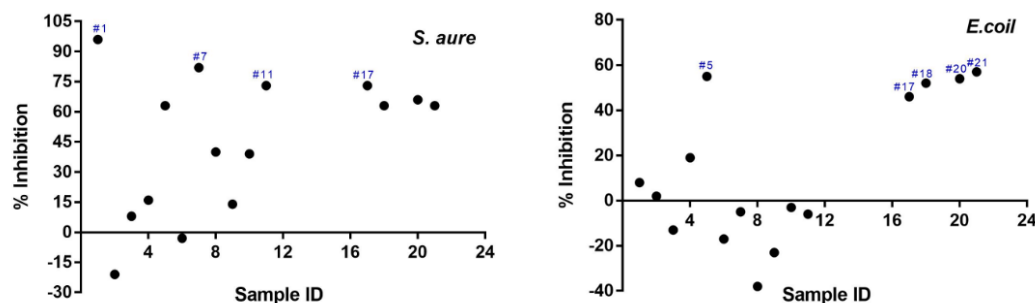
**Supplementary Figure S15.** Scatterplots of 13 strain extracts (1.0 mg/mL) from *Ainsliaea fragrans* screened in *S. aureus* and *E. coli* inhibitory assay.



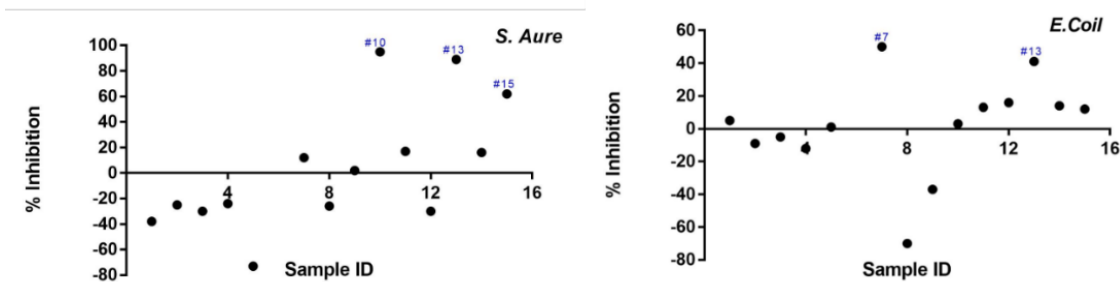
**Supplementary Figure S16.** Scatterplots of 11 strain extracts (1.0 mg/mL) from *Artemisia capillaris* screened in *S. aureus* and *E. coli* inhibitory assay.



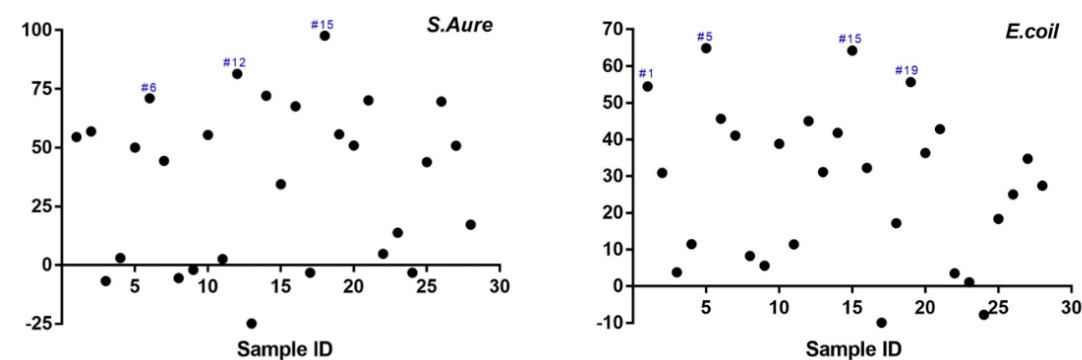
**Supplementary Figure S17.** Scatterplots of 13 strain extracts (1.0 mg/mL) from *Daucus carota* screened in *S. aureus* and *E. coli* inhibitory assay.



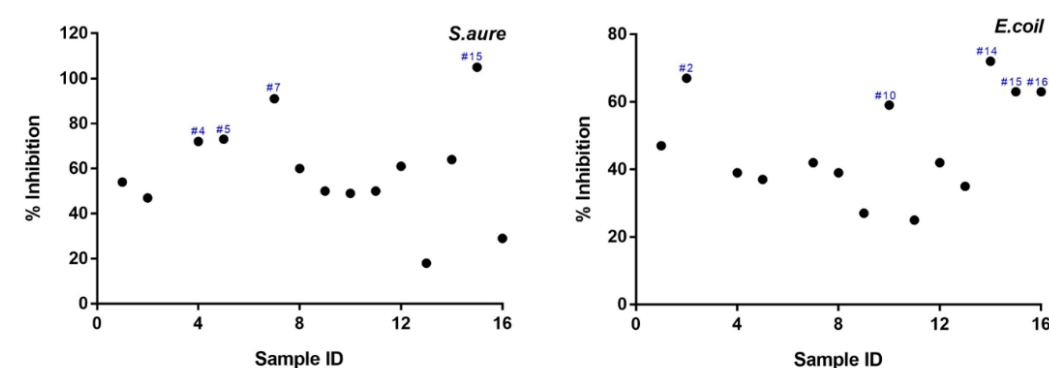
**Supplementary Figure S18.** Scatterplots of 21 strain extracts (1.0 mg/mL) from *Leonurus japonicus* screened in *S. aureus* and *E. coli* inhibitory analyses.



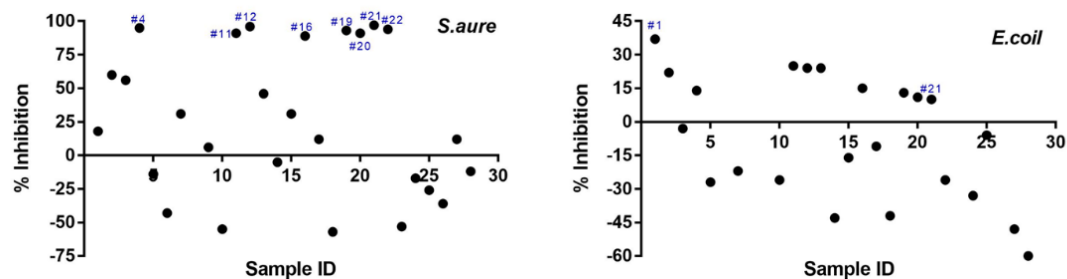
**Supplementary Figure S19.** Scatterplots of 15 strain extracts (1.0 mg/mL) from *Perilla frutescens* screened in *S. aureus* and *E. coli* inhibitory assay.



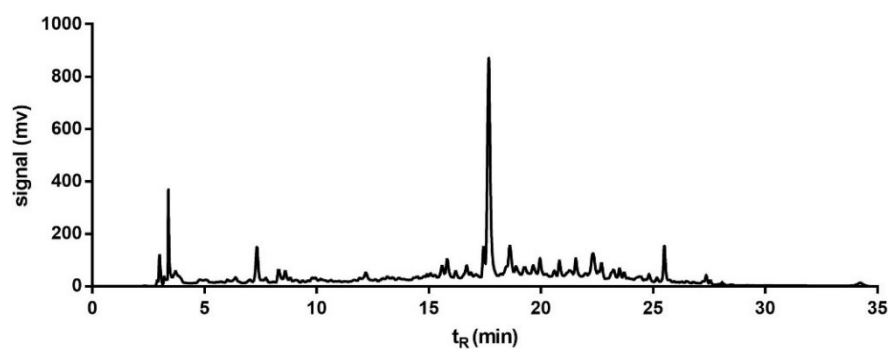
**Supplementary Figure S20.** Scatterplots of 28 strain extracts (1.0 mg/mL) from *Prunella vulgaris* screened in *S. aureus* and *E. coli* inhibitory assay.



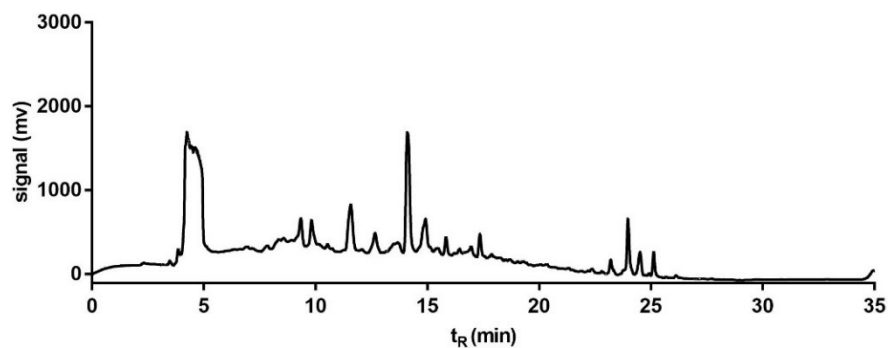
**Supplementary Figure S21.** Scatterplots of 16 strain extracts (1.0 mg/mL) from *Salvia miltiorrhiza* screened in *S. aureus* and *E. coli* inhibitory assay.



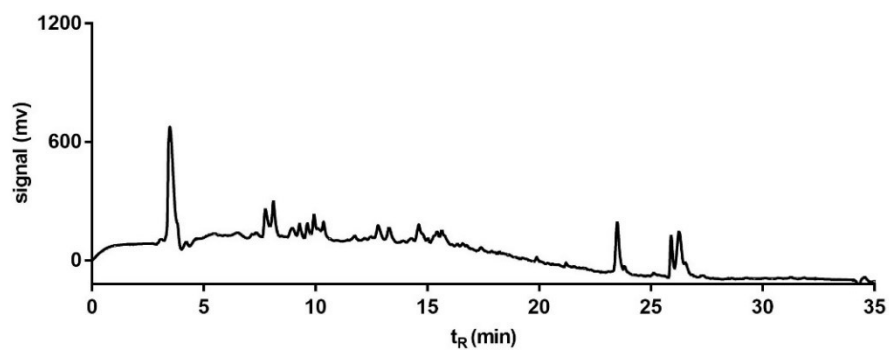
**Supplementary Figure S22.** Scatterplots of 28 strain extracts (1.0 mg/mL) from *Stemona sessilifolia* screened in *S. aureus* and *E. coli* inhibitory assay.



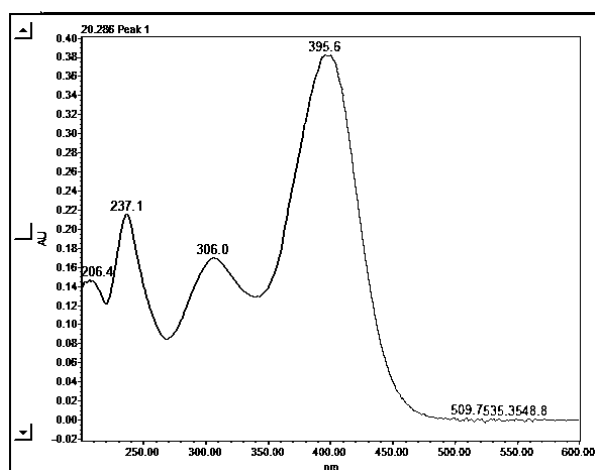
**Supplementary Figure S23.** HPLC spectrum of *Penicillium sp.* NX-S-6



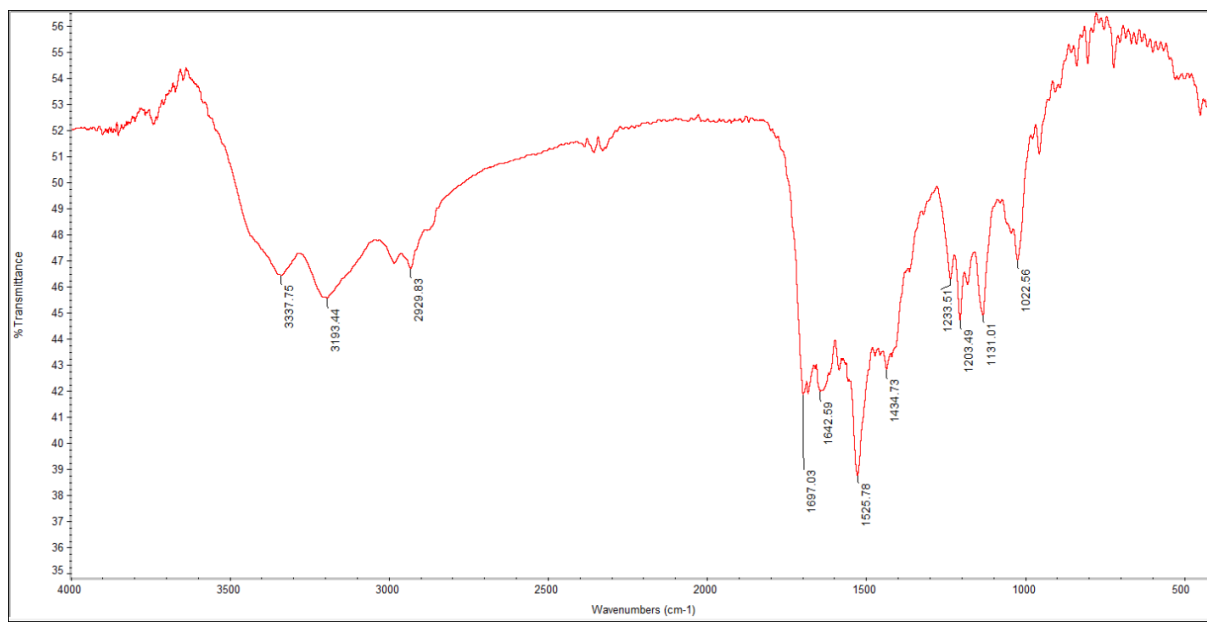
**Supplementary Figure S24.** HPLC spectrum of *Streptomyces sp.* YHLB-L-2



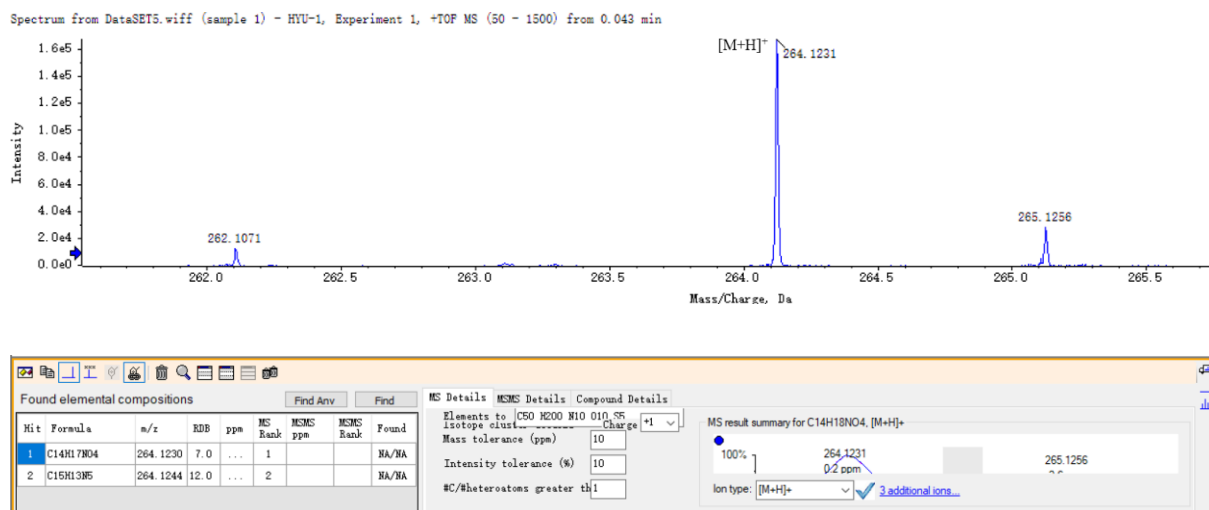
**Supplementary Figure S25.** HPLC spectrum of *Streptomyces* sp. ZLBB-S-6



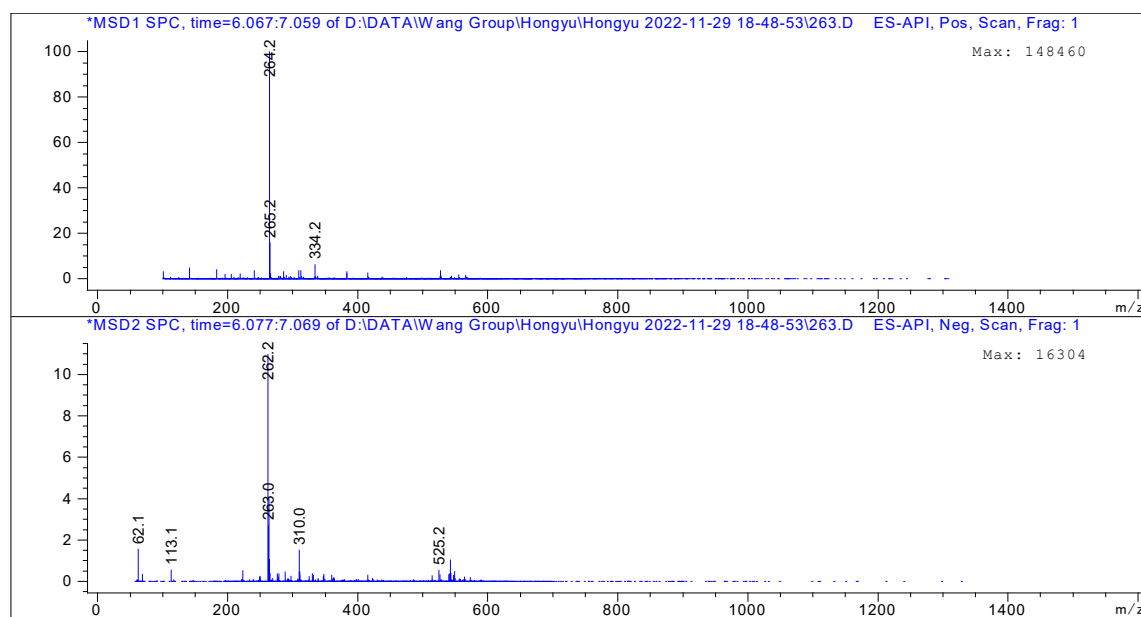
**Supplementary Figure S26.** UV spectrum of compound 1.



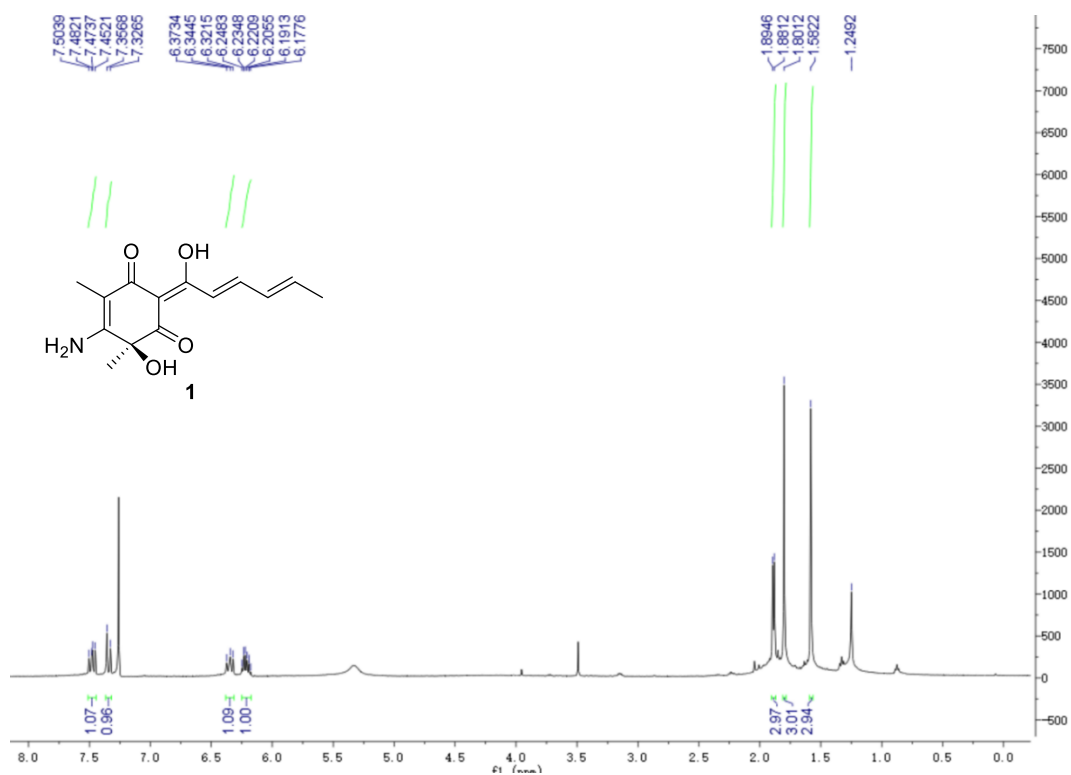
**Supplementary Figure S27.** IR spectrum of compound 1.



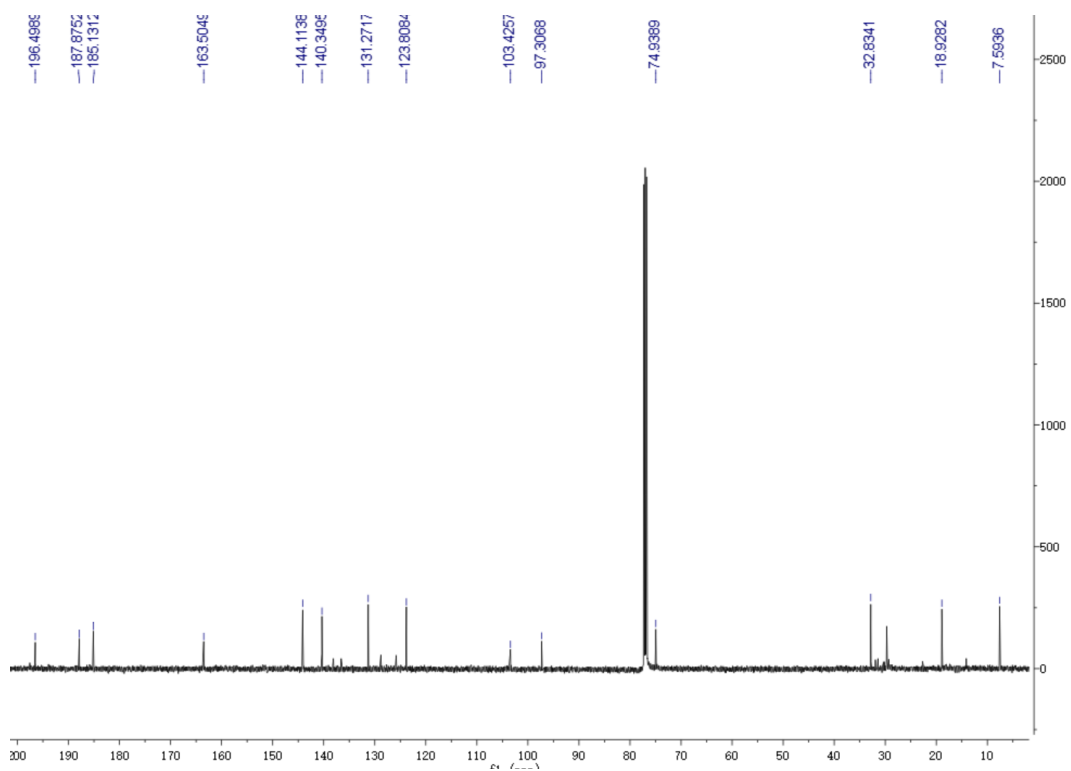
Supplementary Figure S28. (+)-HRESI-MS spectrum of compound 1.



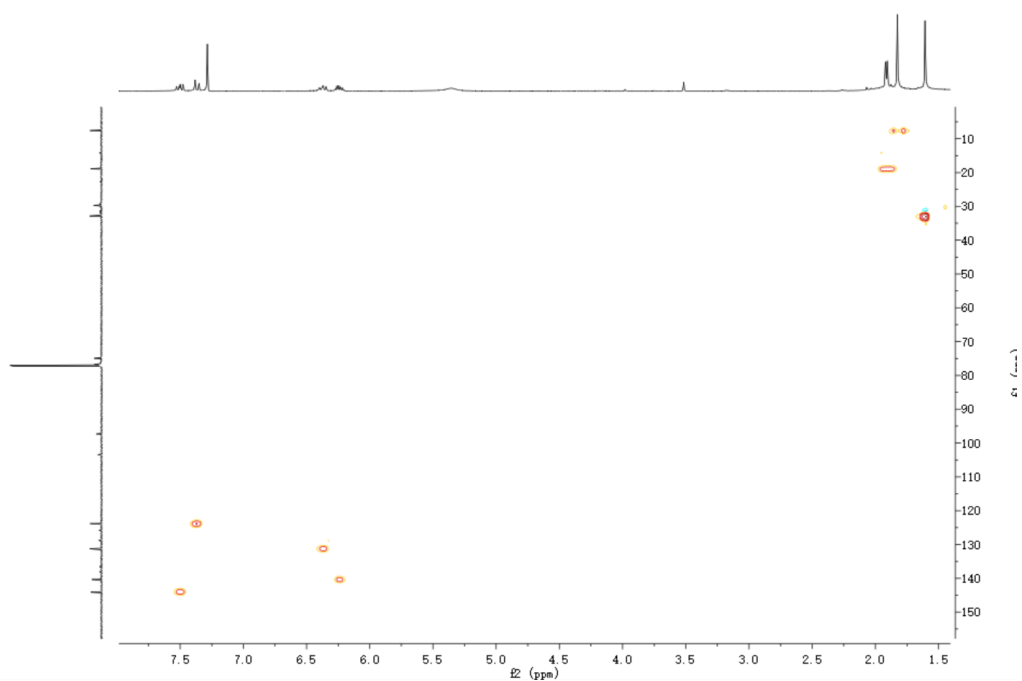
Supplementary Figure S29. (+) and (-)-ESI-MS of compound 1.



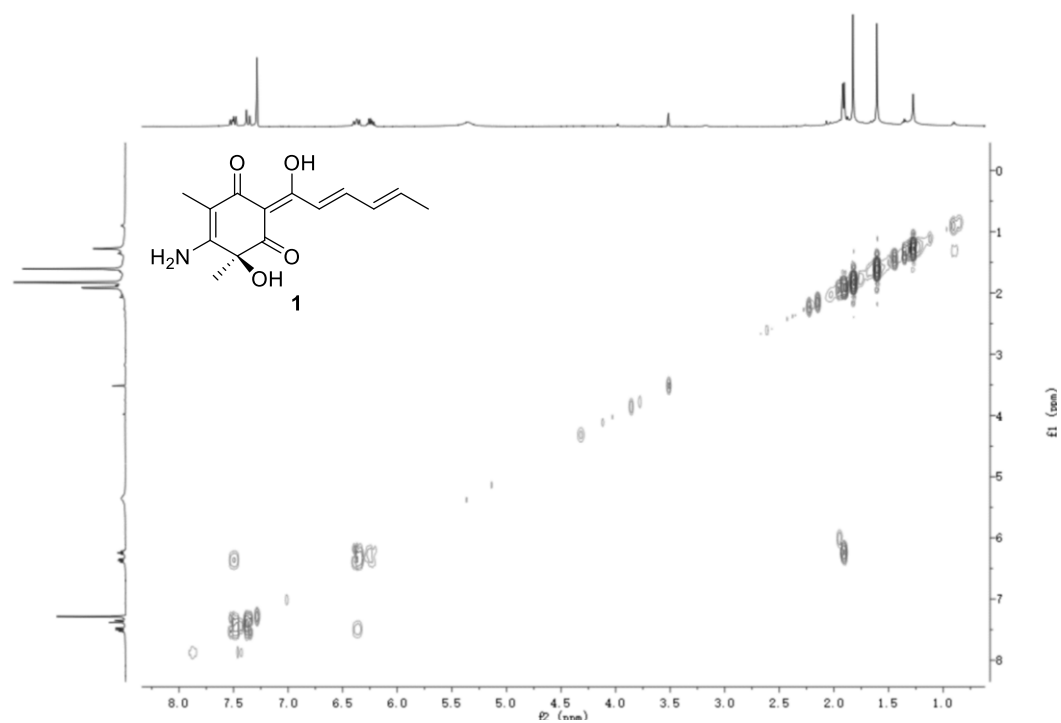
**Supplementary Figure S30.**  $^1\text{H}$  NMR (500 MHz,  $\text{CDCl}_3$ ) spectrum of compound **1**.



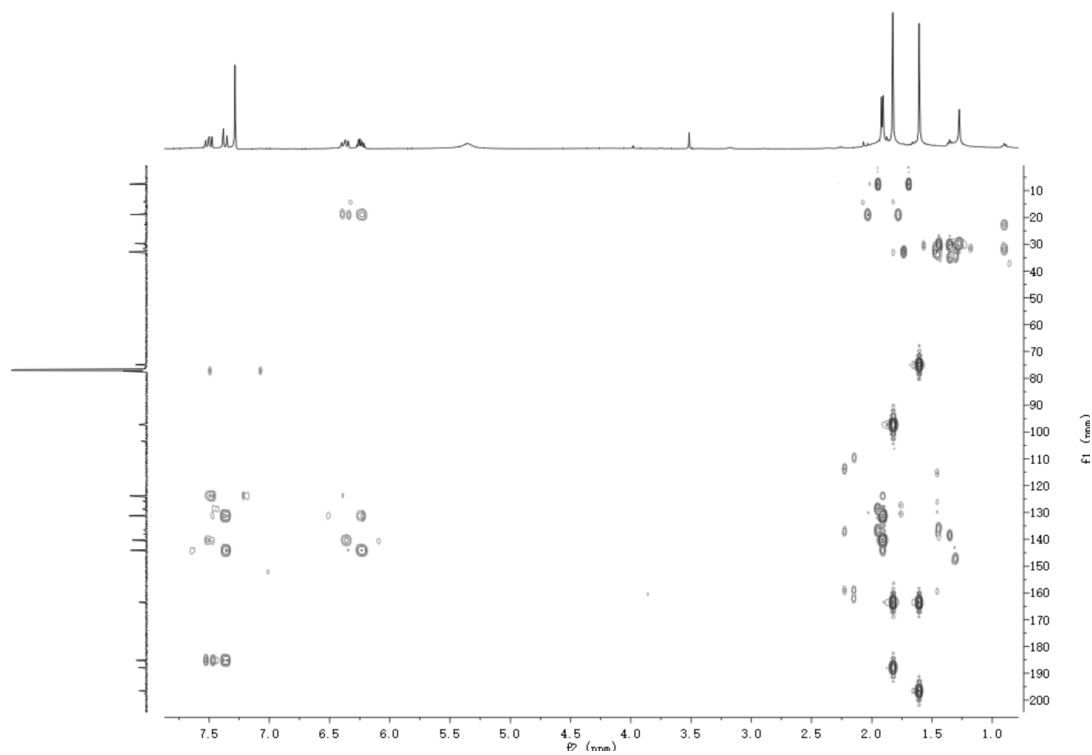
**Supplementary Figure S31.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{CDCl}_3$ ) spectrum of compound **1**.



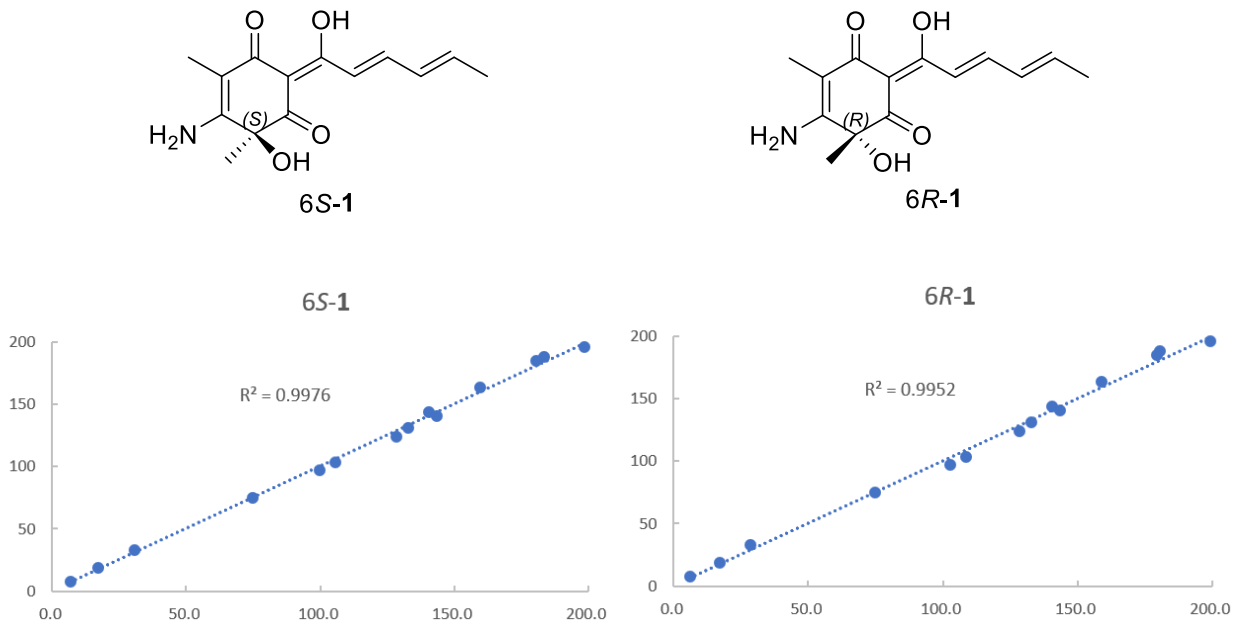
**Supplementary Figure S32.** HSQC (500 MHz, CDCl<sub>3</sub>) spectrum of compound **1**.



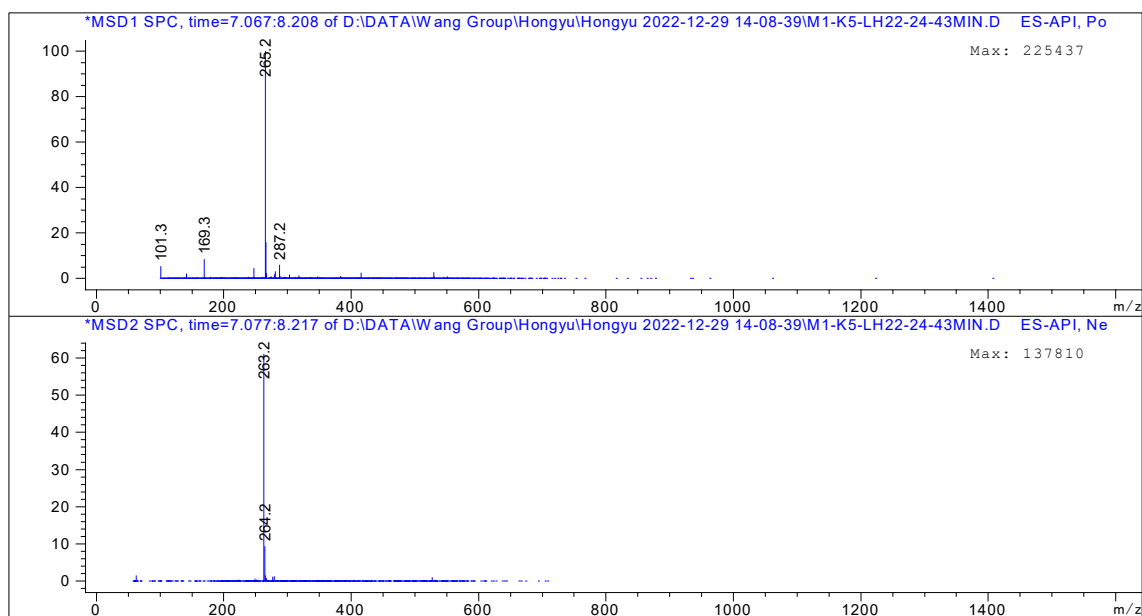
**Supplementary Figure S33.** <sup>1</sup>H-<sup>1</sup>H COSY (500 MHz, CDCl<sub>3</sub>) spectrum of compound **1**.



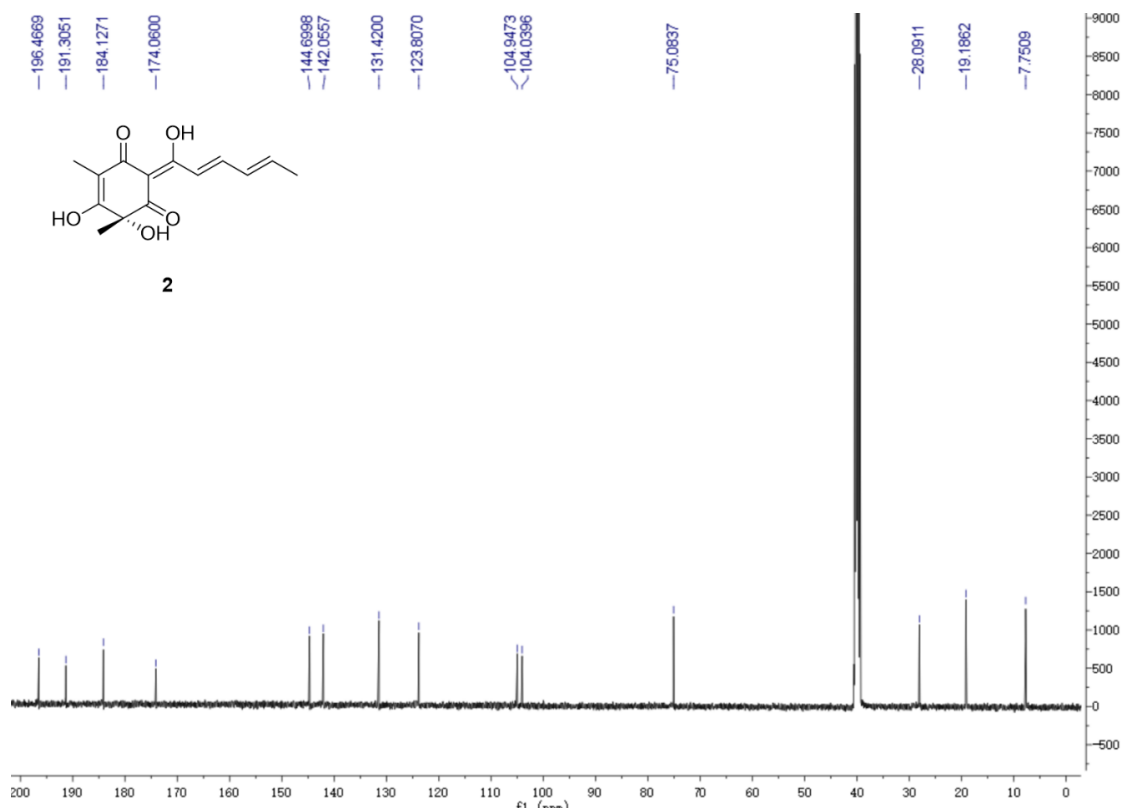
**Supplementary Figure S34.** HMBC (500 MHz, CDCl<sub>3</sub>) spectrum of compound **1**.



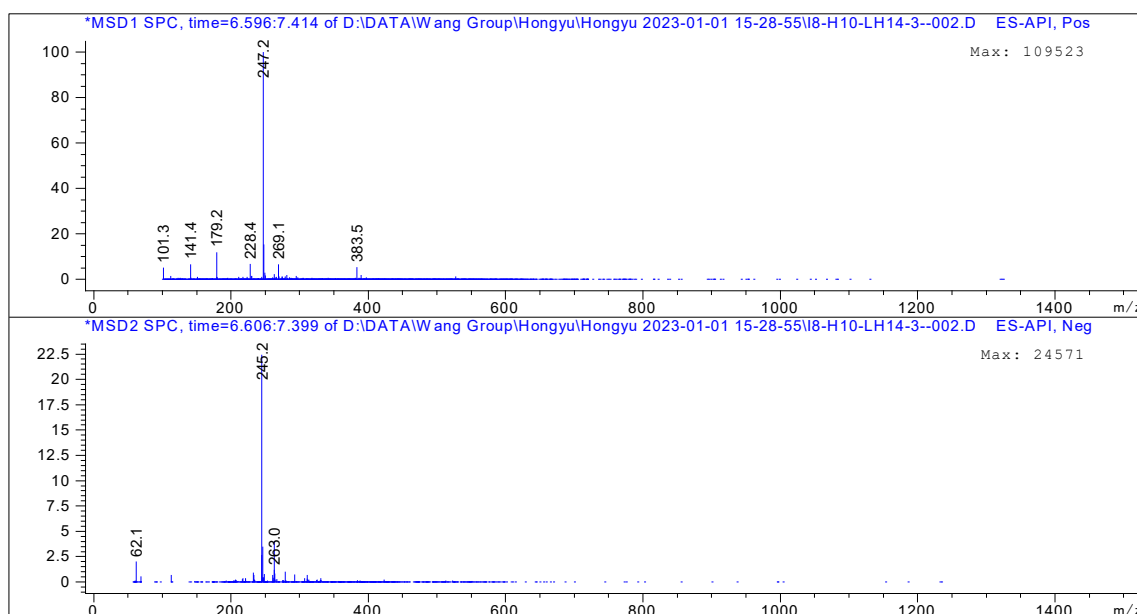
**Supplementary Figure S35.** Regression analysis of the experimental vs calculated <sup>13</sup>C NMR chemical shifts of **1** at the B3LYP/6-311+G(d,p) level in CDCl<sub>3</sub>.



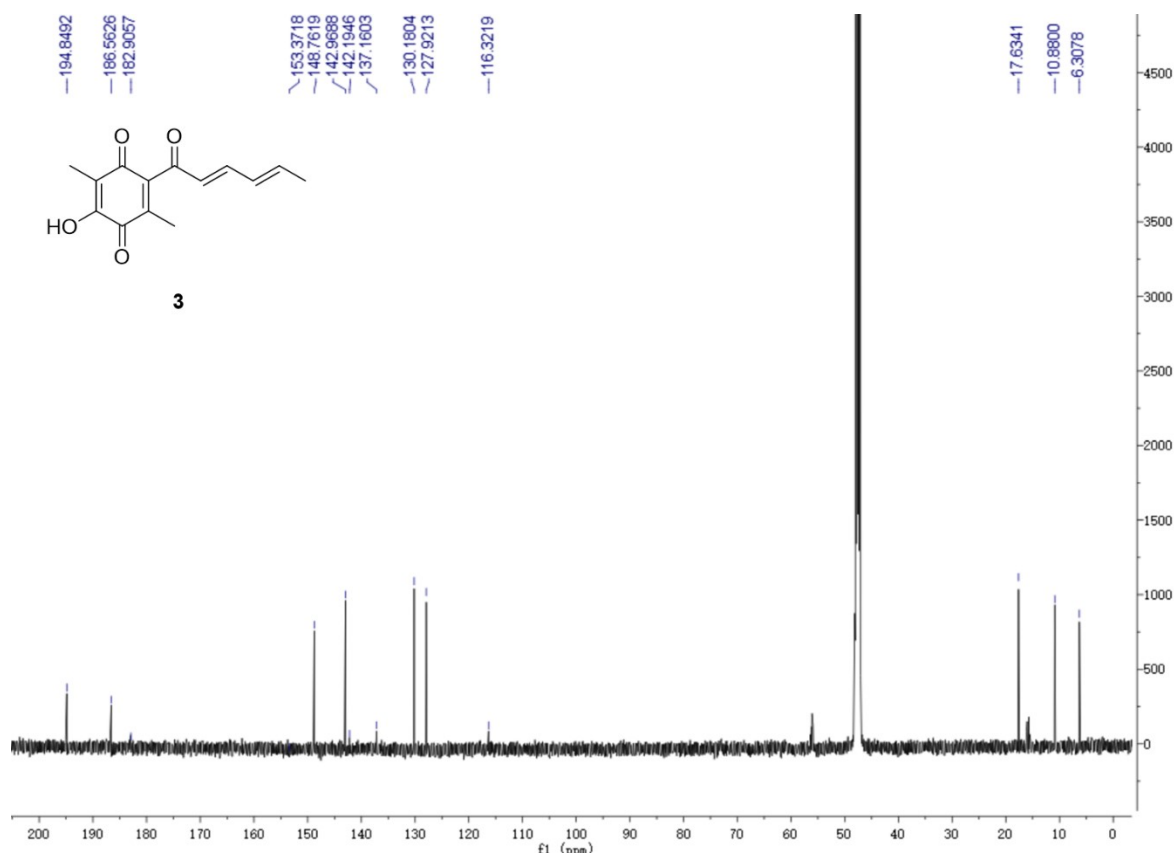
**Supplementary Figure S36.** (+) and (–)-ESI-MS of compound **2**.



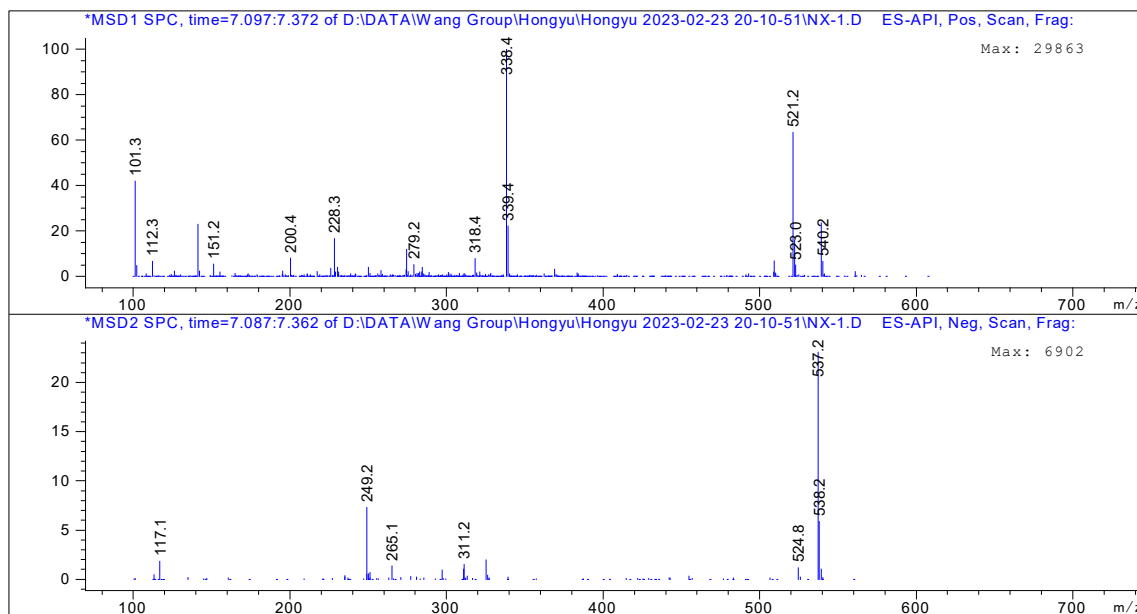
**Supplementary Figure S37.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **2**.



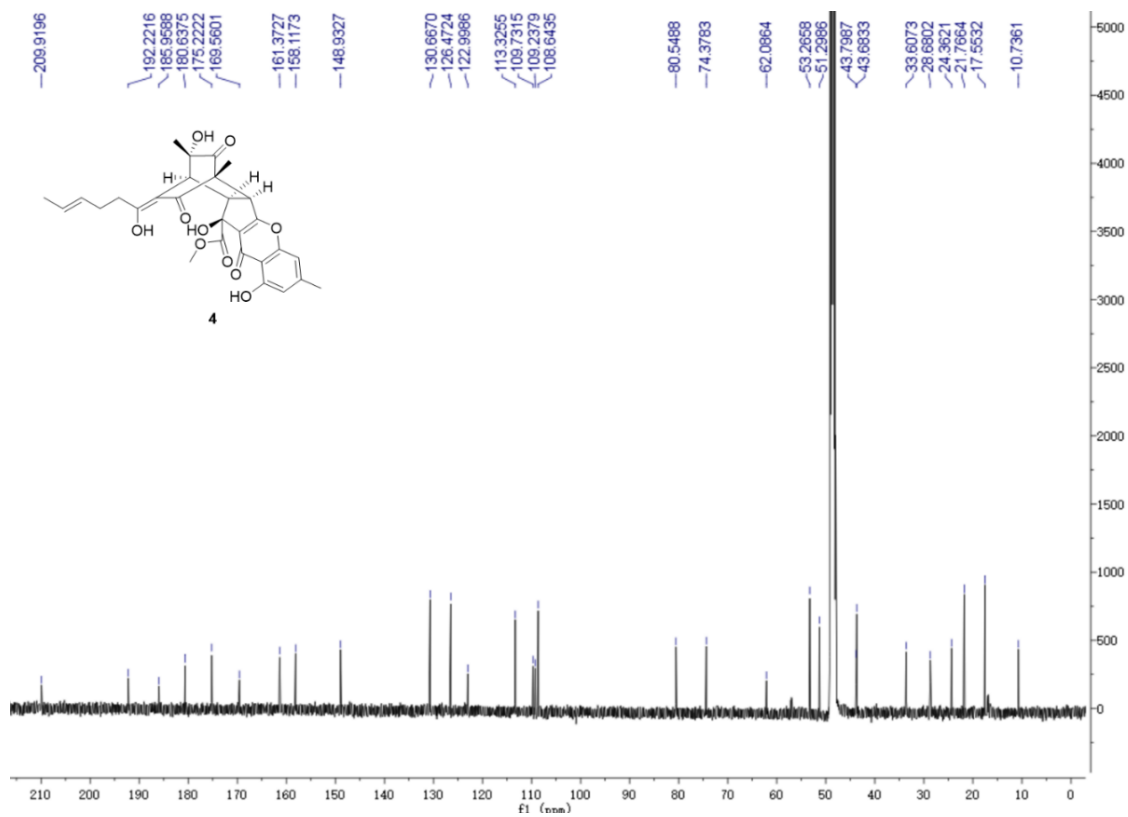
**Supplementary Figure S38.** (+) and (–)-ESI-MS of compound **3**.



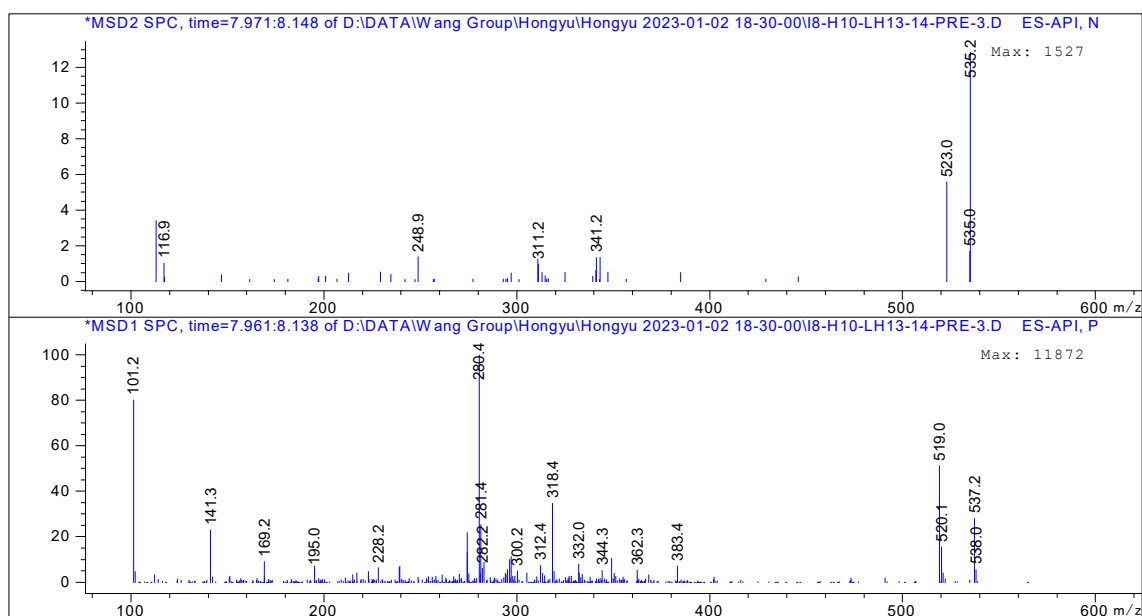
**Supplementary Figure S39.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{CD}_3\text{OD}$ ) spectrum of compound **3**.



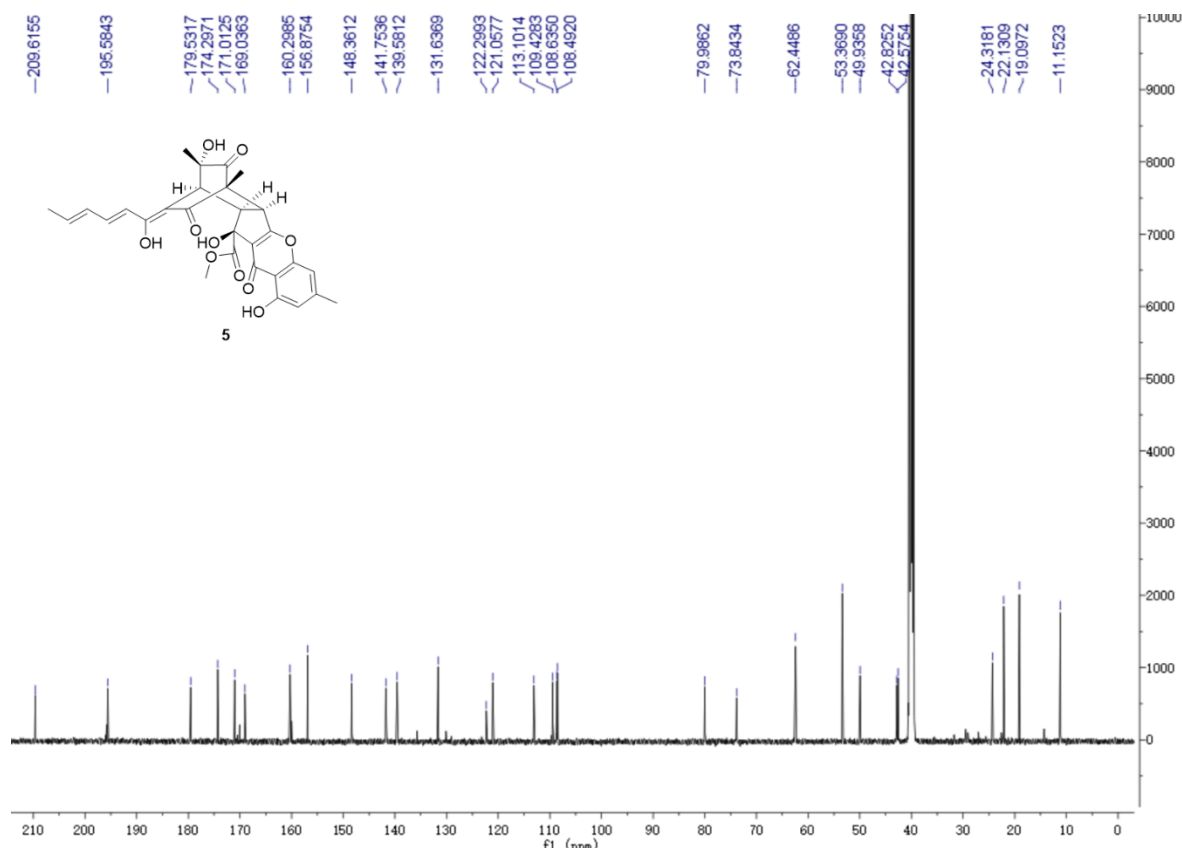
Supplementary Figure S40. (+) and (-)-ESI-MS of compound **4**.



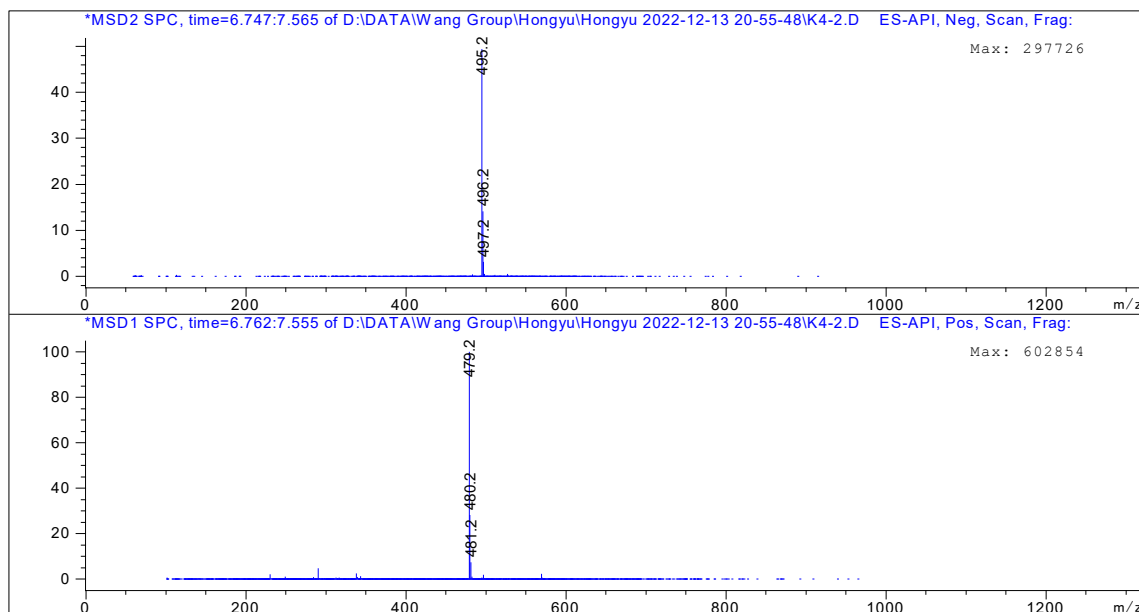
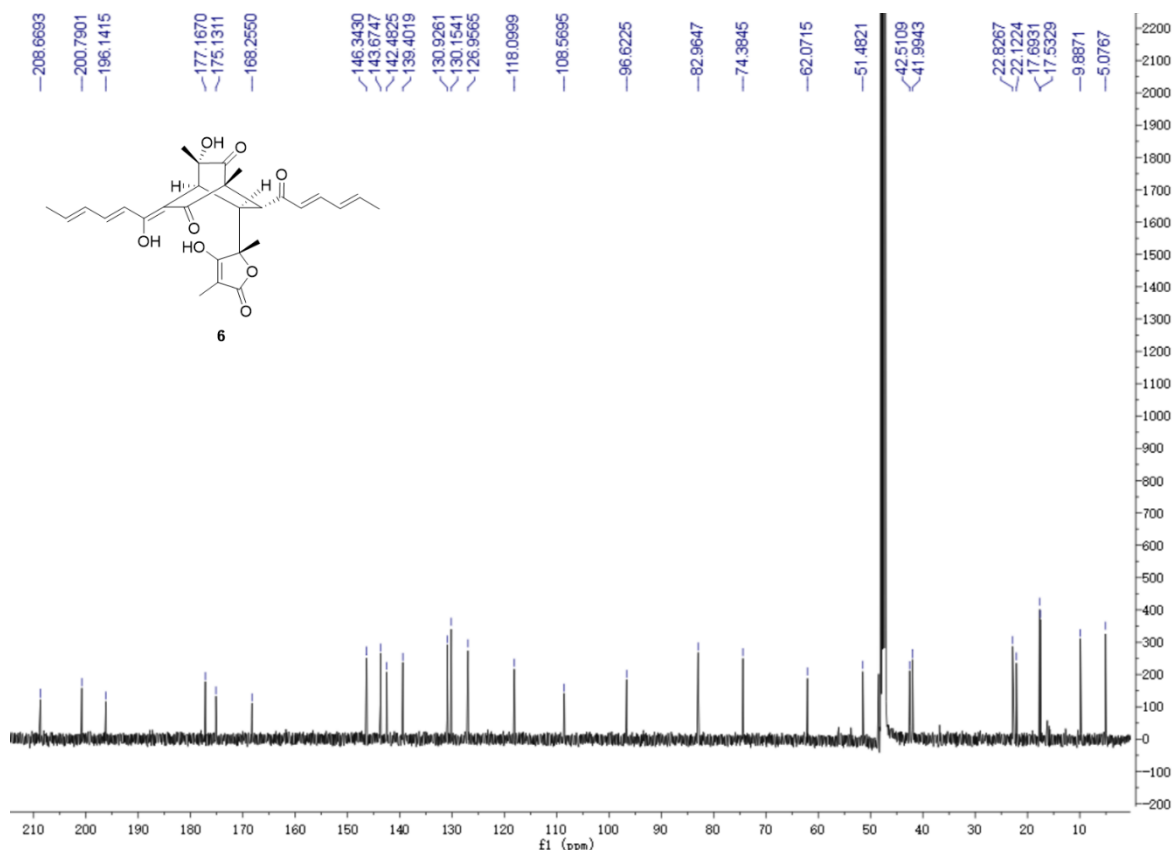
Supplementary Figure S41.  $^{13}\text{C}$  NMR (125 MHz,  $\text{CD}_3\text{OD}$ ) spectrum of compound **4**.

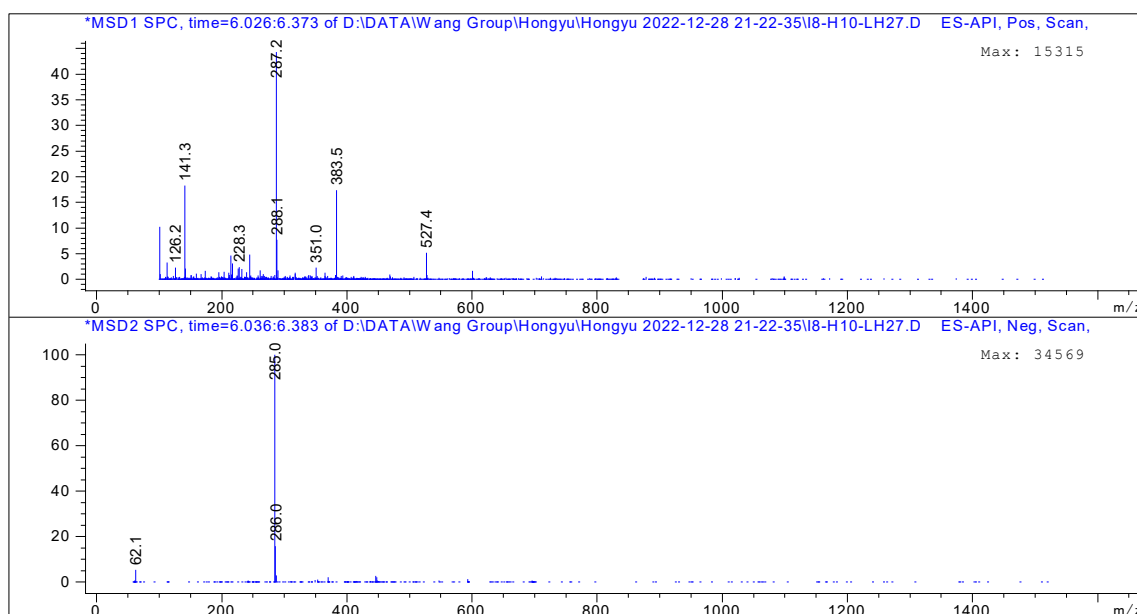


**Supplementary Figure S42.** (+) and (-)-ESI-MS of compound **5**.

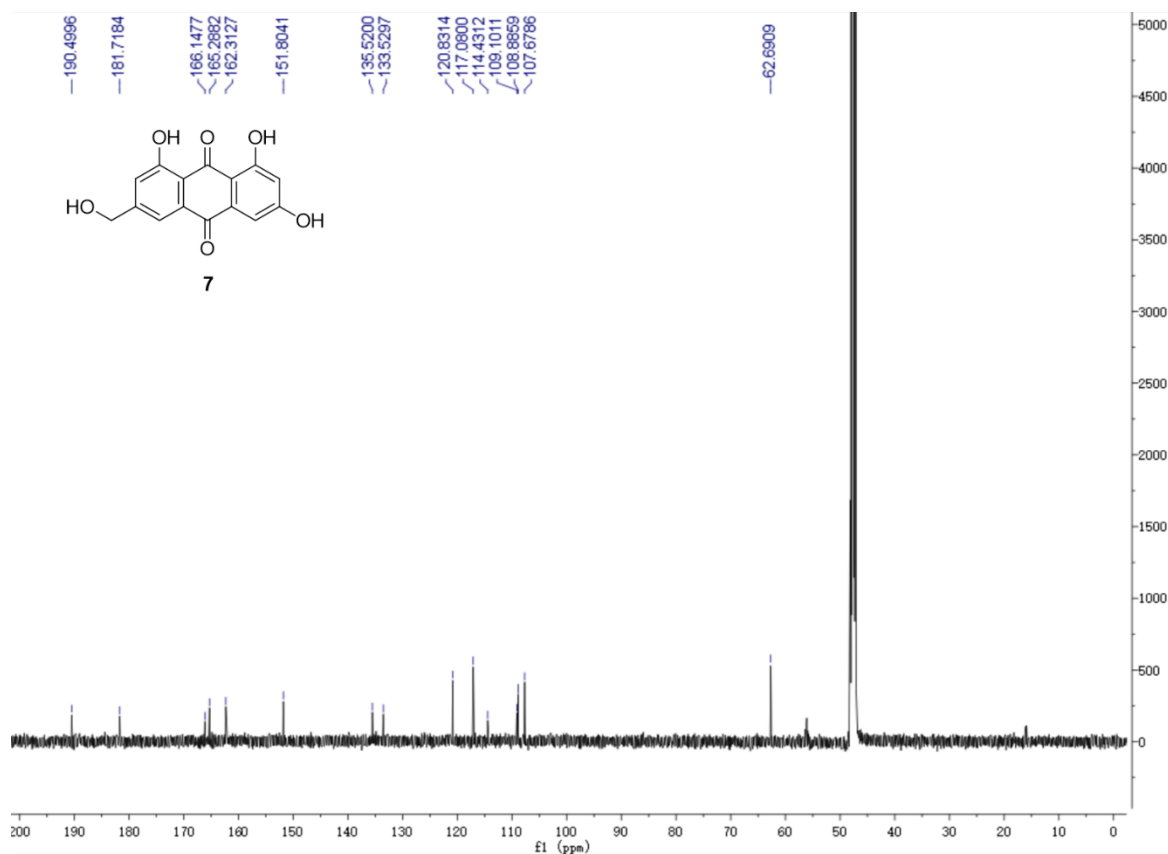


**Supplementary Figure S43.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **5**.

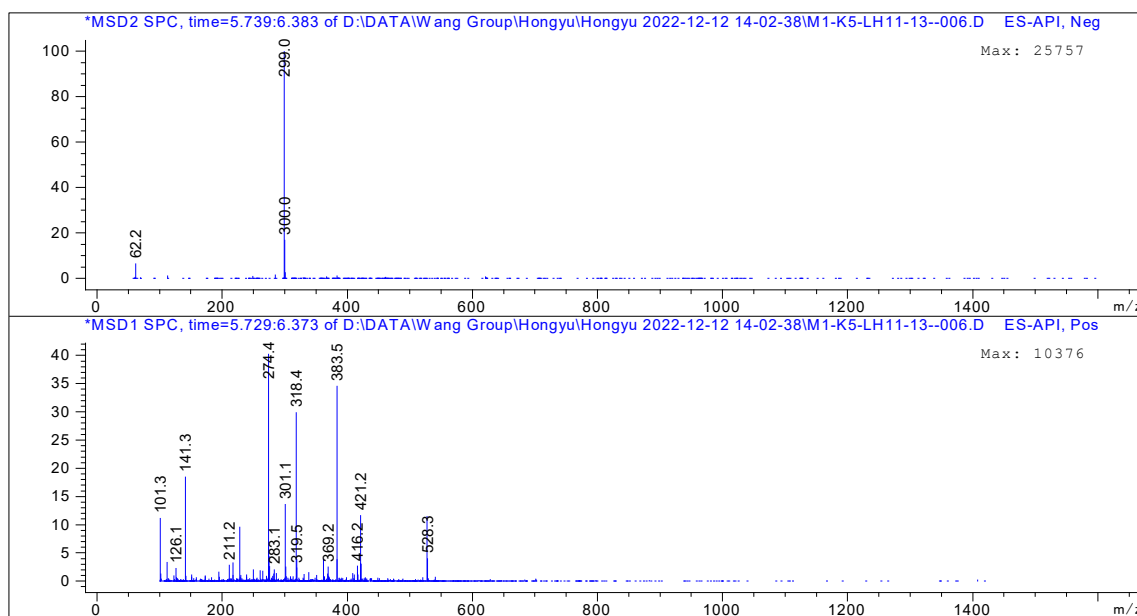
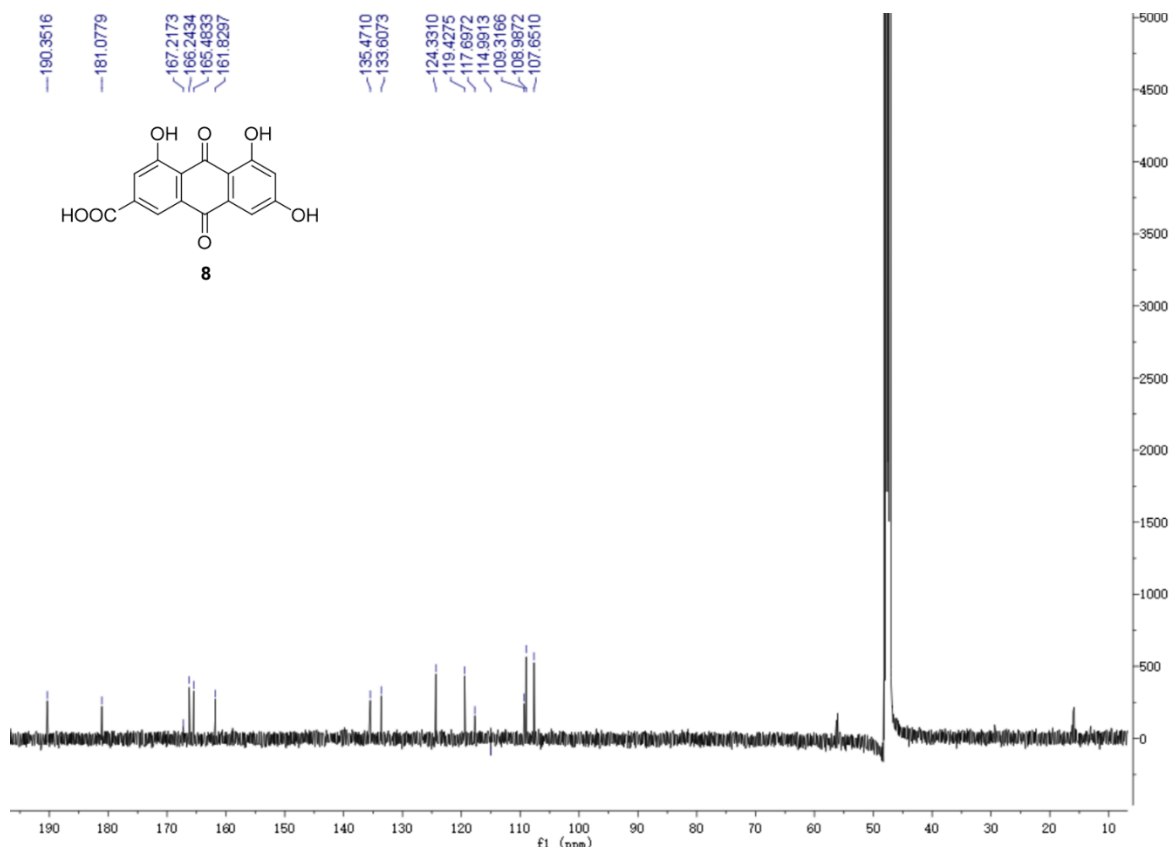
Supplementary Figure S44. (+) and (-)-ESI-MS of compound **6**.Supplementary Figure S45.  $^{13}\text{C}$  NMR (125 MHz,  $\text{CD}_3\text{OD}$ ) spectrum of compound **6**.

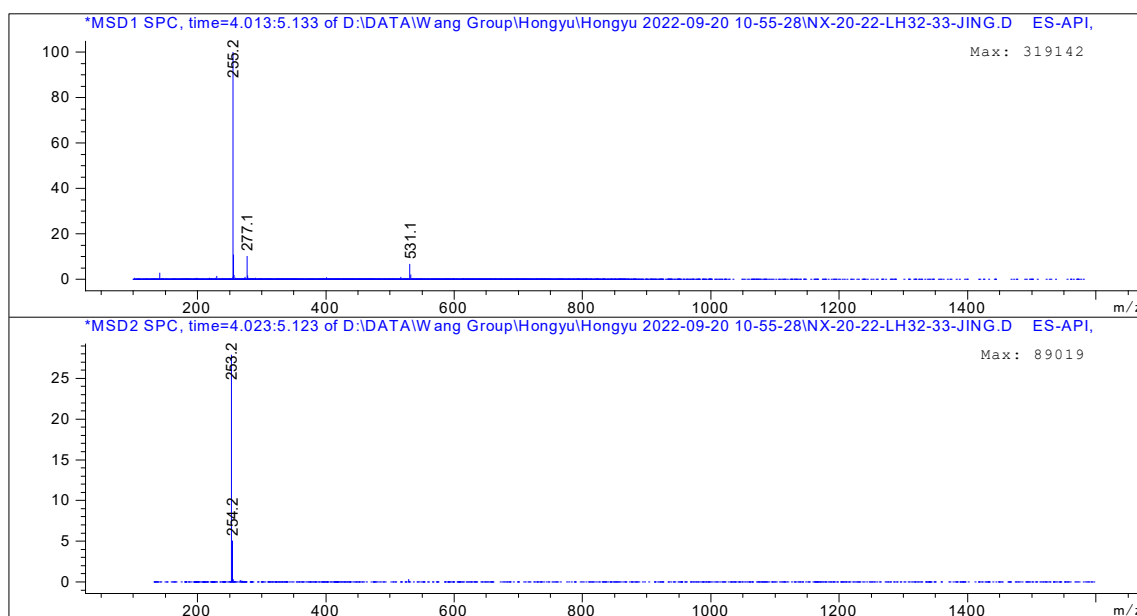


**Supplementary Figure S46.** (+) and (-)-ESI-MS of compound **7**.

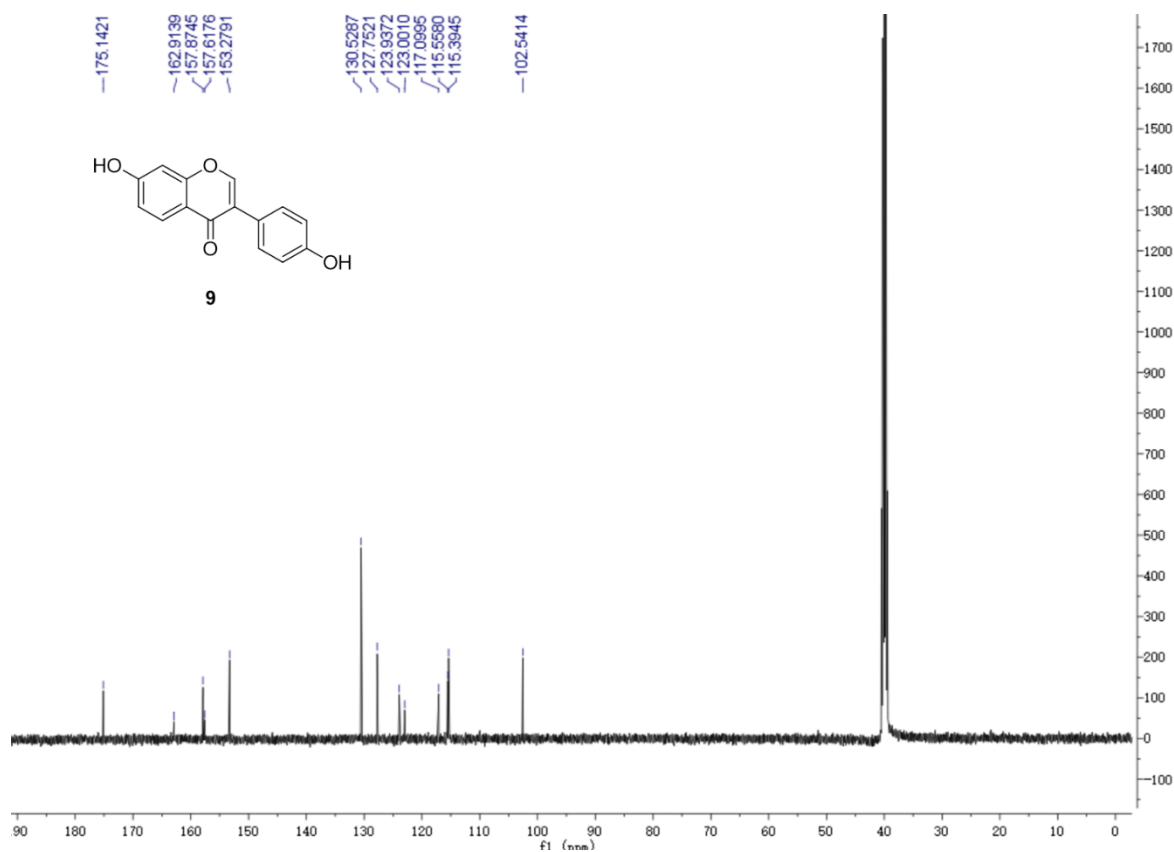


**Supplementary Figure S47.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{CD}_3\text{OD}$ ) spectrum of compound **7**.

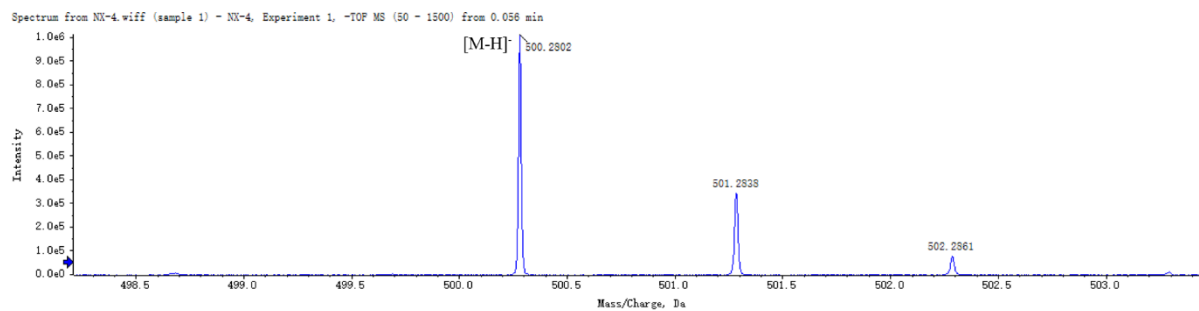
Supplementary Figure S48. (+) and (-)-ESI-MS of compound **8**.Supplementary Figure S49. <sup>13</sup>C NMR (125 MHz, CD<sub>3</sub>OD) spectrum of compound **8**.



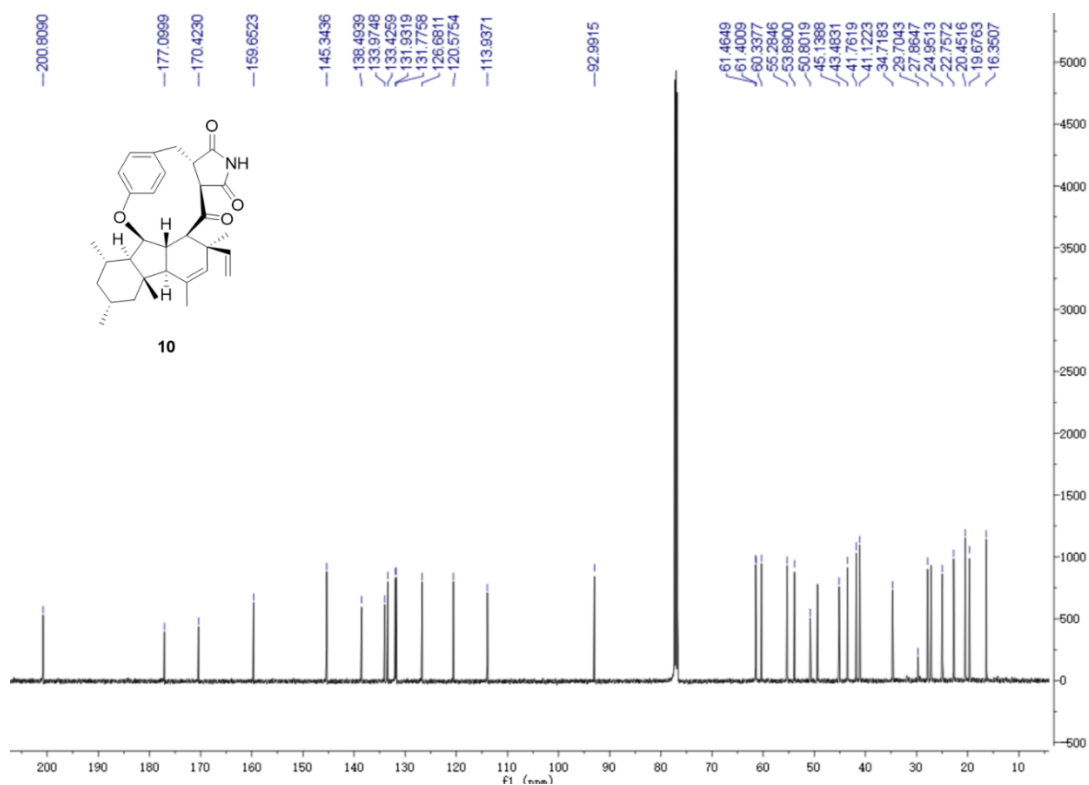
**Supplementary Figure S50.** (+) and (–)-ESI-MS of compound **9**.



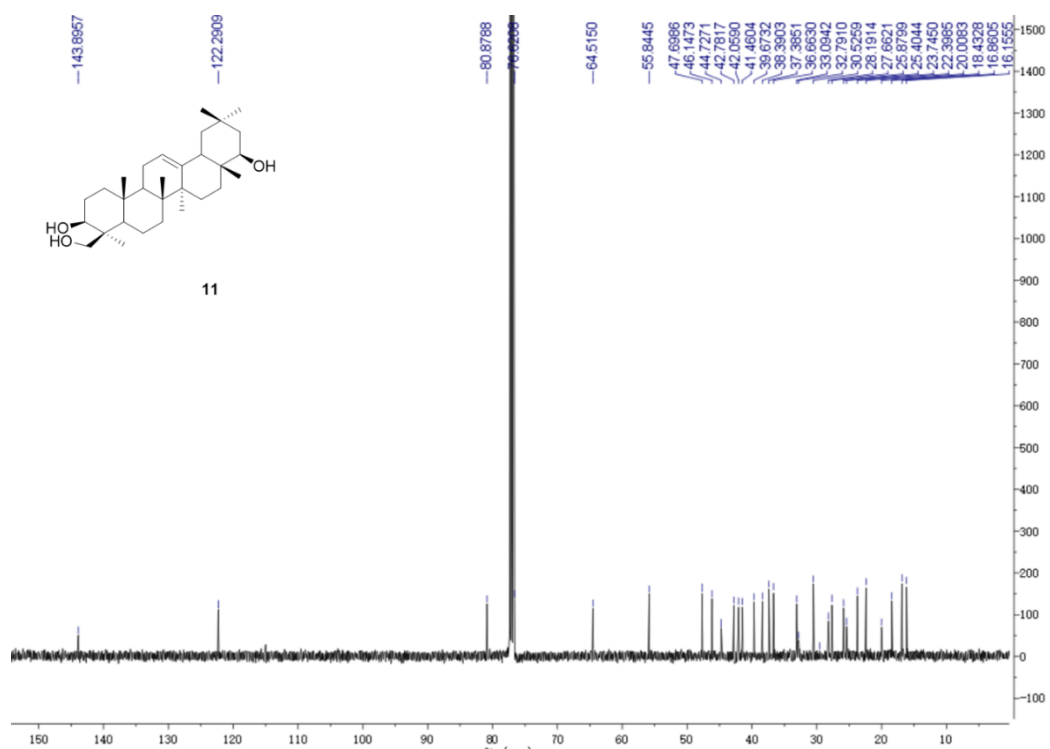
**Supplementary Figure S51.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **9**.



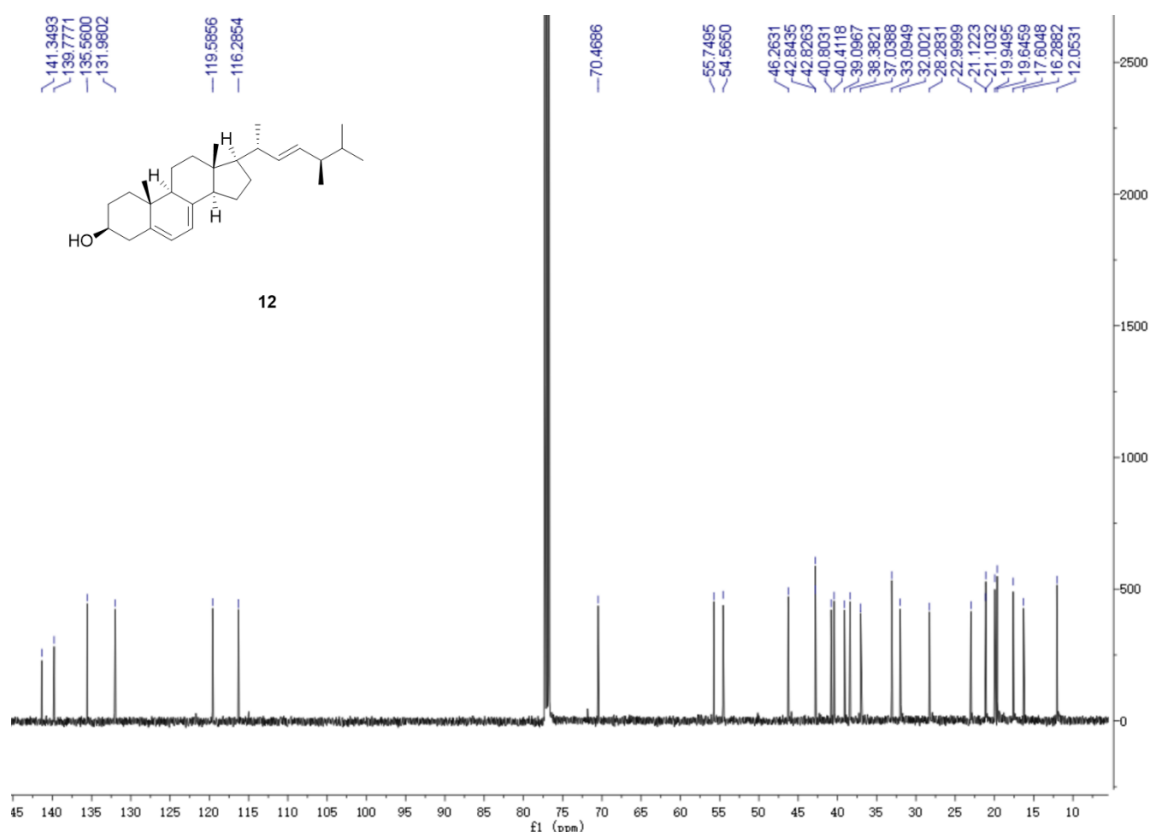
**Supplementary Figure S52.** (-)-HRESI-MS of compound **10**.



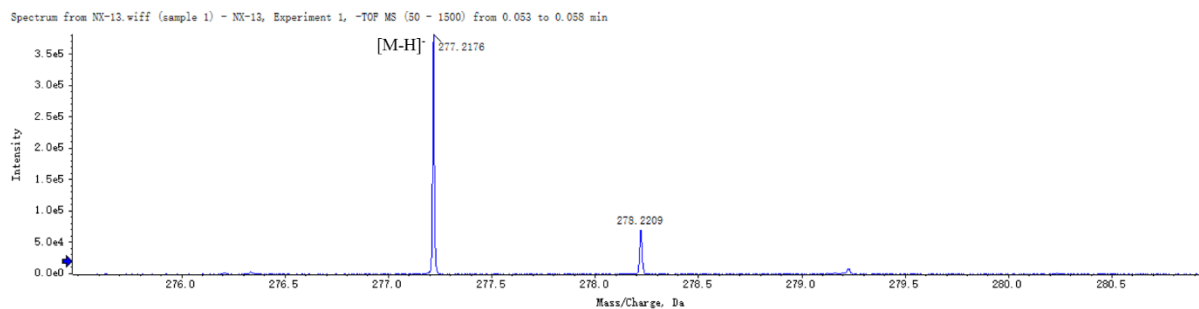
**Supplementary Figure S53.** <sup>13</sup>C NMR (125 MHz, CD<sub>3</sub>OD) spectrum of compound **10**.



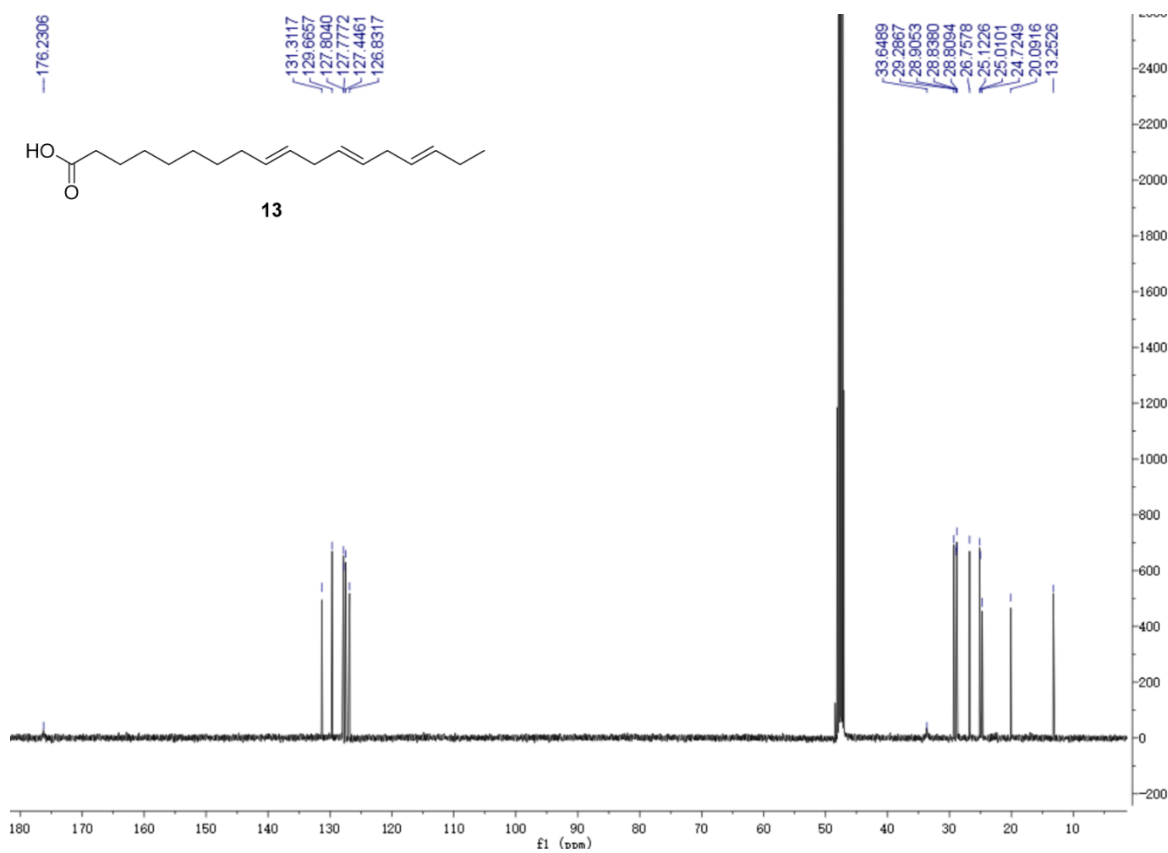
Supplementary Figure S54.  $^{13}\text{C}$  NMR (125 MHz,  $\text{CD}_3\text{Cl}_3$ ) spectrum of compound 11.



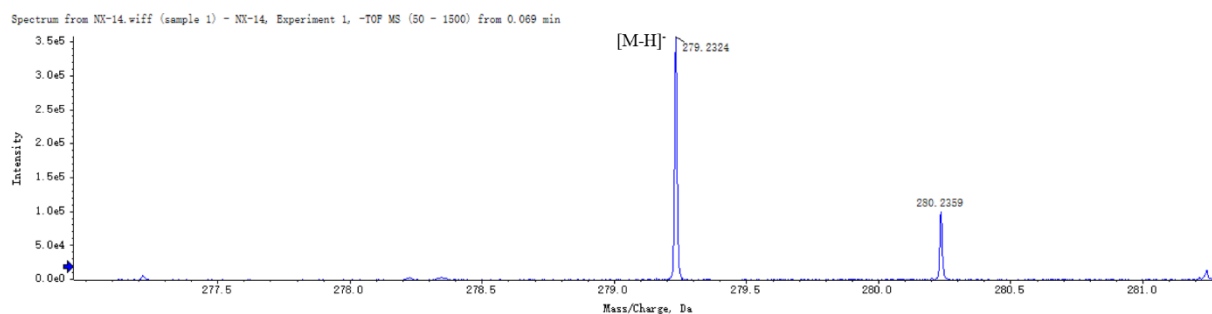
Supplementary Figure S55.  $^{13}\text{C}$  NMR (125 MHz,  $\text{CD}_3\text{Cl}_3$ ) spectrum of compound 12.



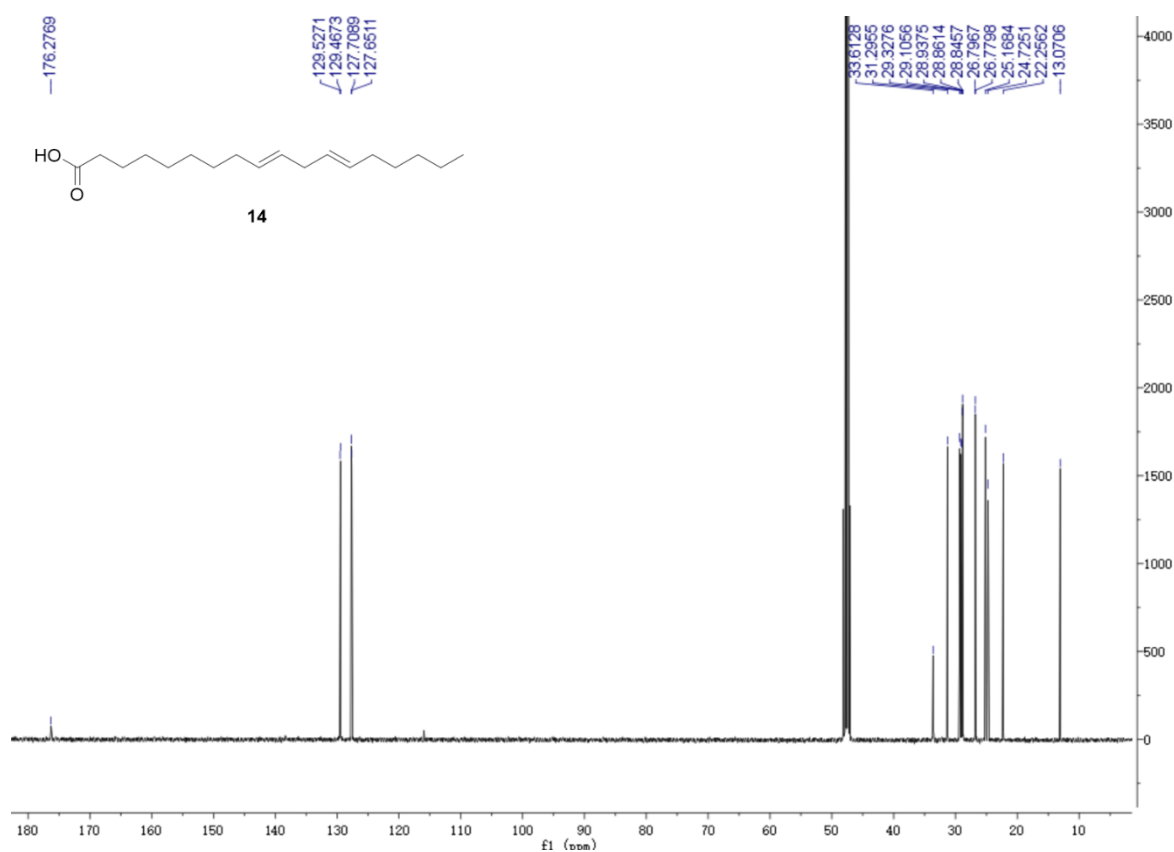
**Supplementary Figure S56.** (-)-HRESI-MS of compound **13**.



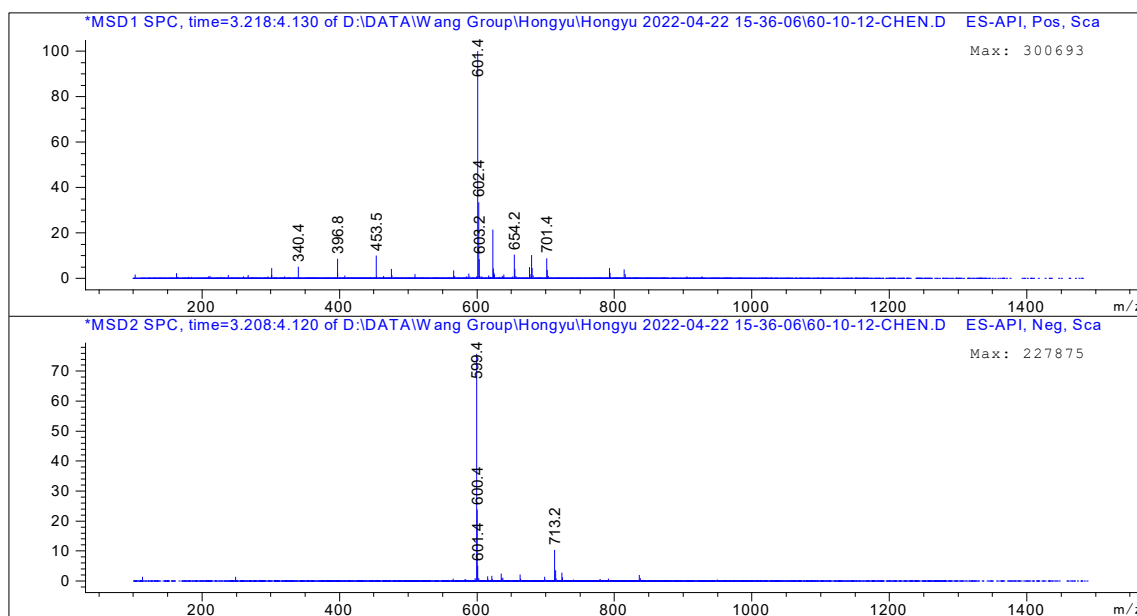
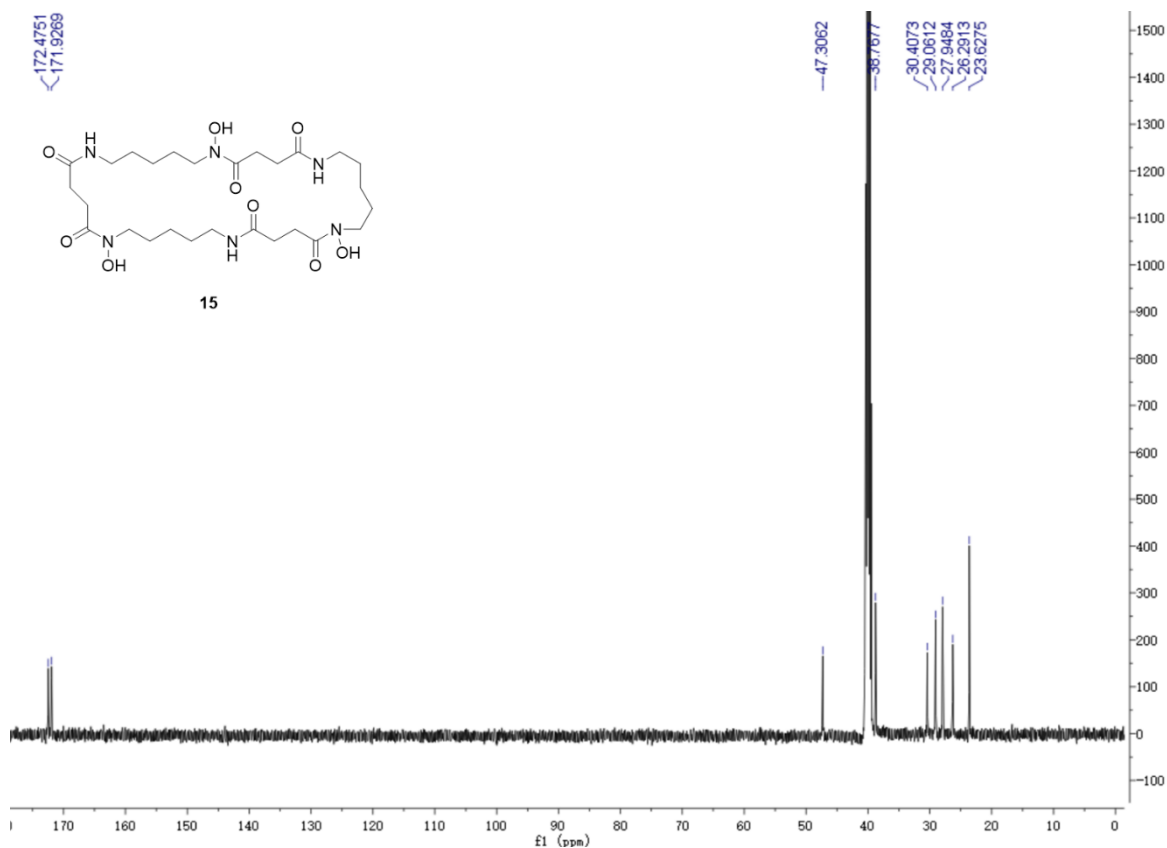
**Supplementary Figure S57.** <sup>13</sup>C NMR (125 MHz, CD<sub>3</sub>OD) spectrum of compound **13**.

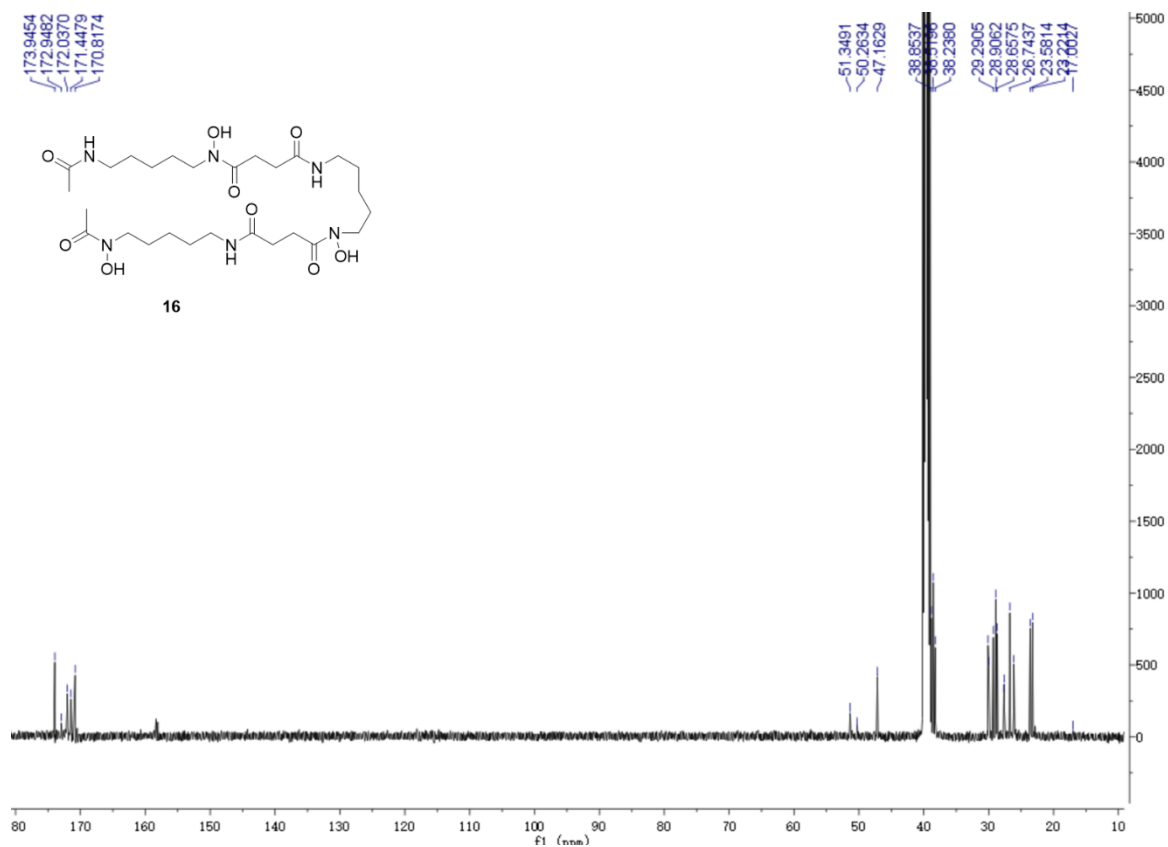
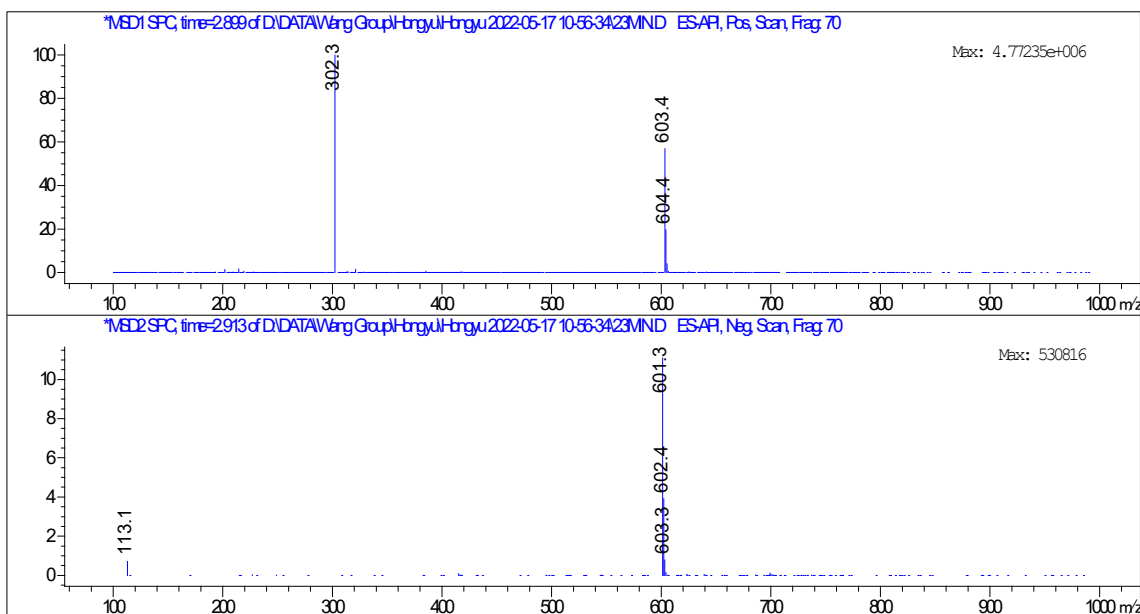


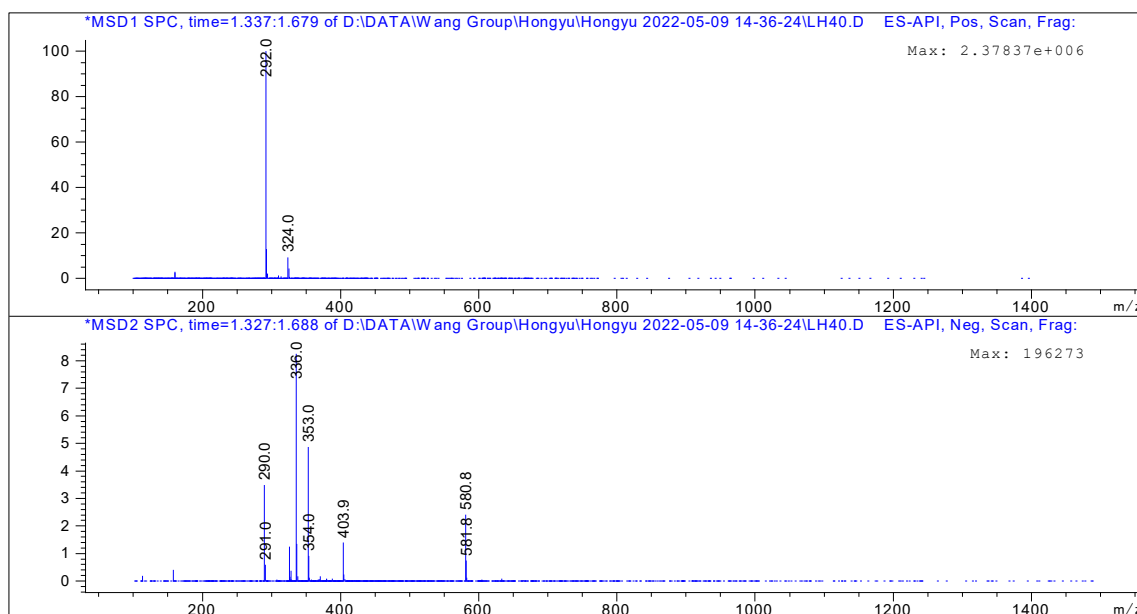
**Supplementary Figure S58.** (–)-HRESI-MS of compound **14**.



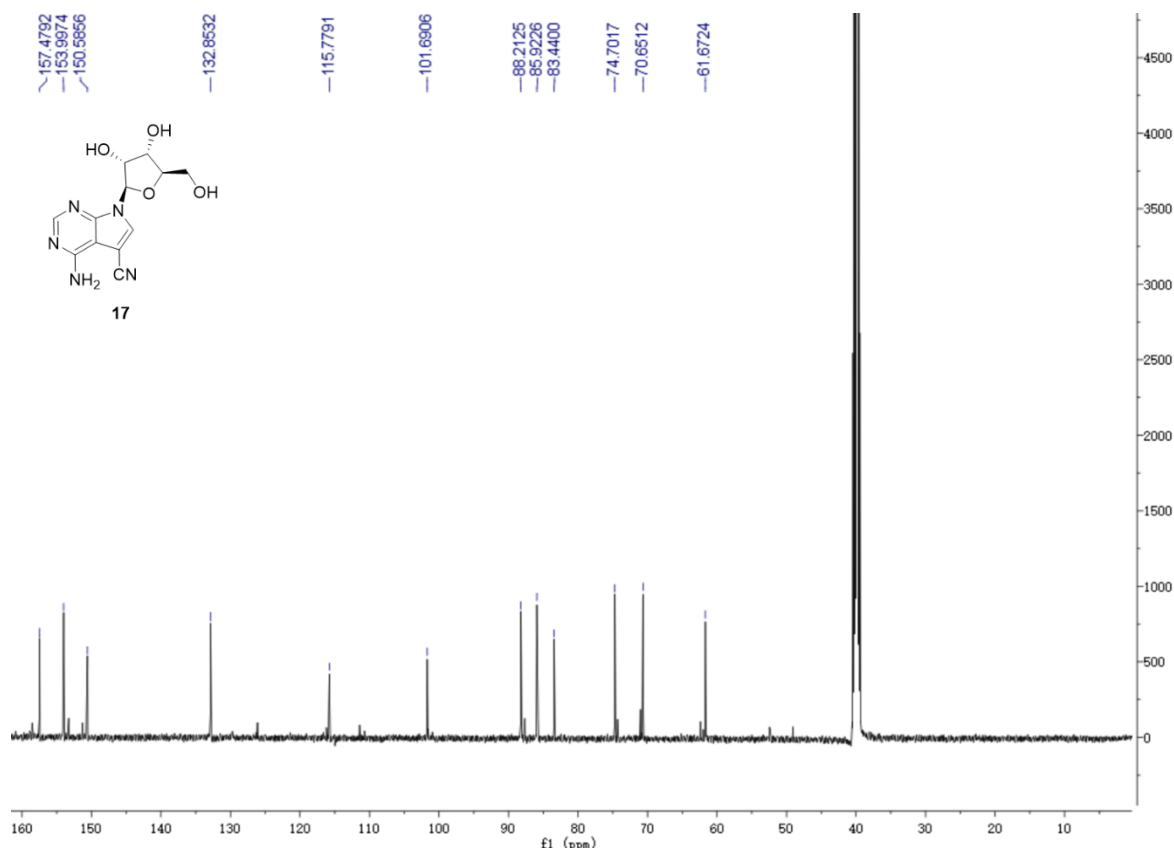
**Supplementary Figure S59.** <sup>13</sup>C NMR (125 MHz, CD<sub>3</sub>OD) spectrum of compound **14**.

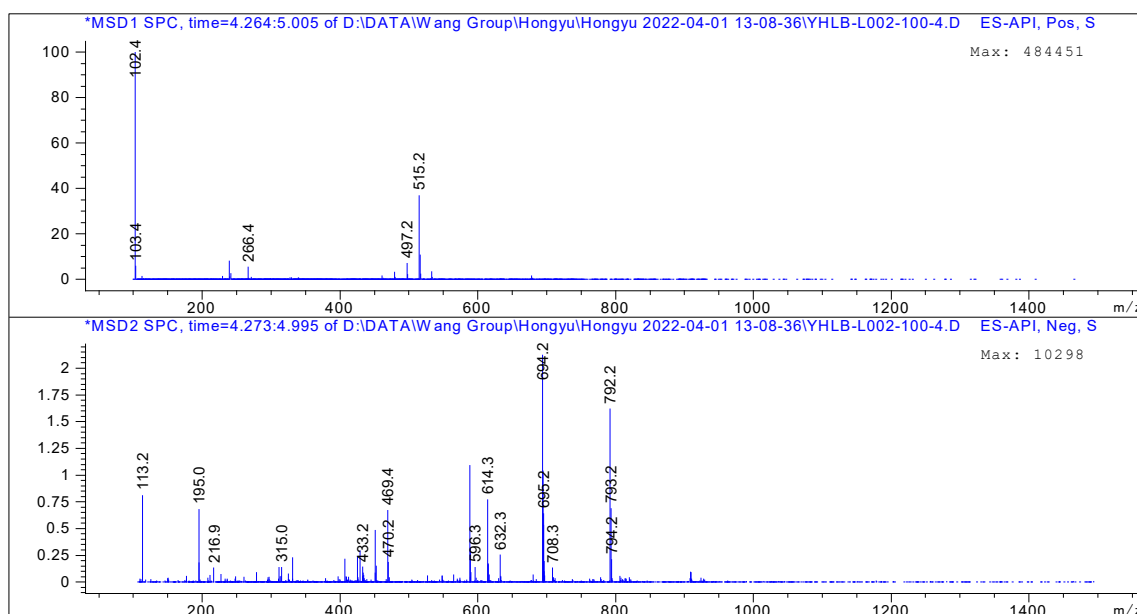
Supplementary Figure S60. (+) and (-)-ESI-MS of compound **15**.Supplementary Figure S61.  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **15**.



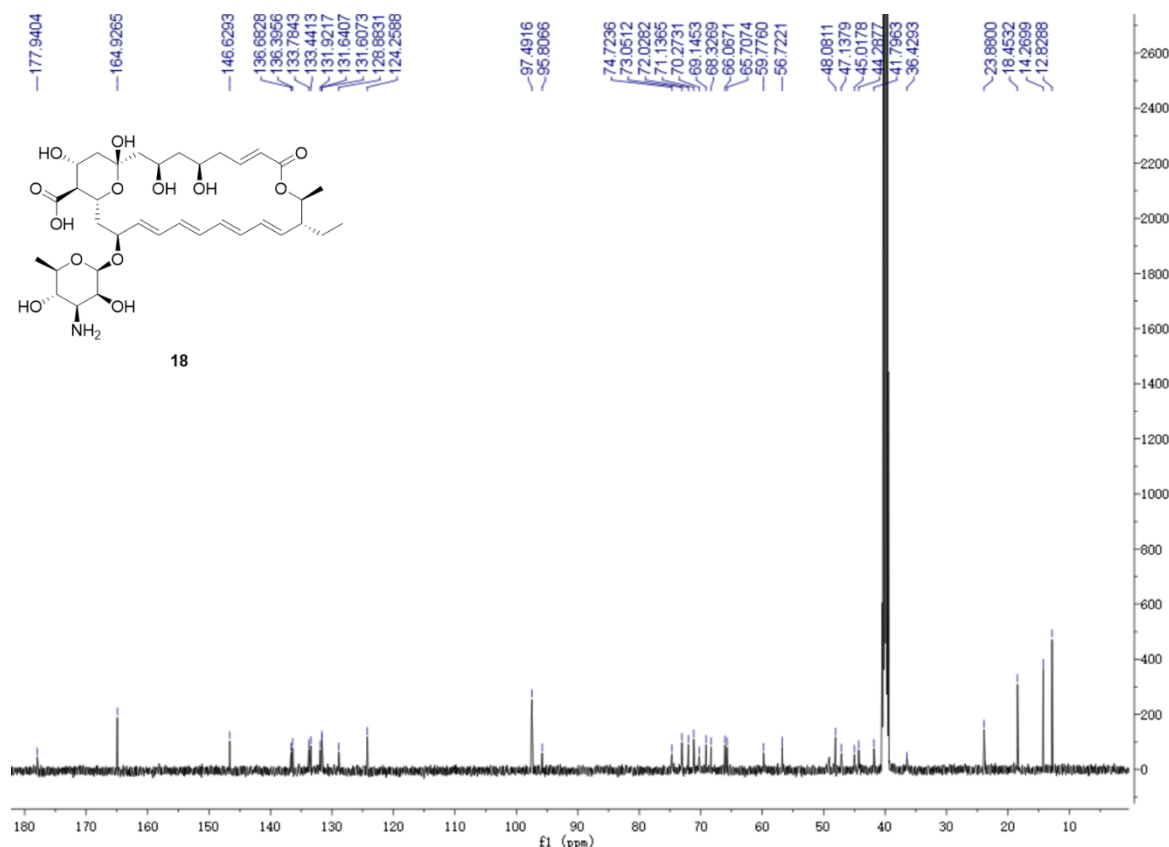


Supplementary Figure S64. (+) and (-)-ESI-MS of compound 17.

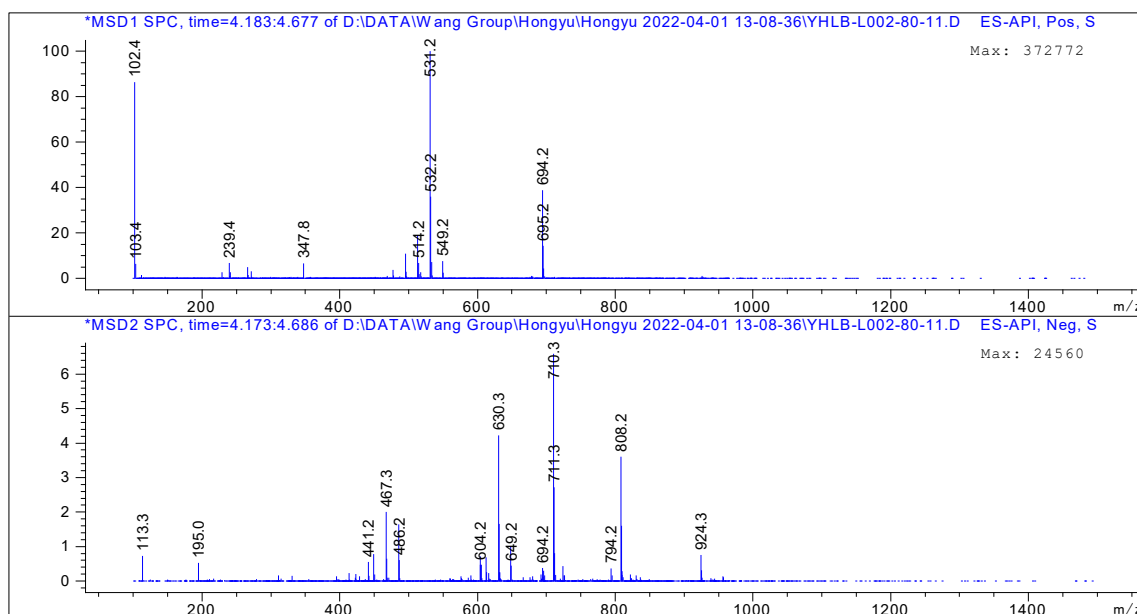
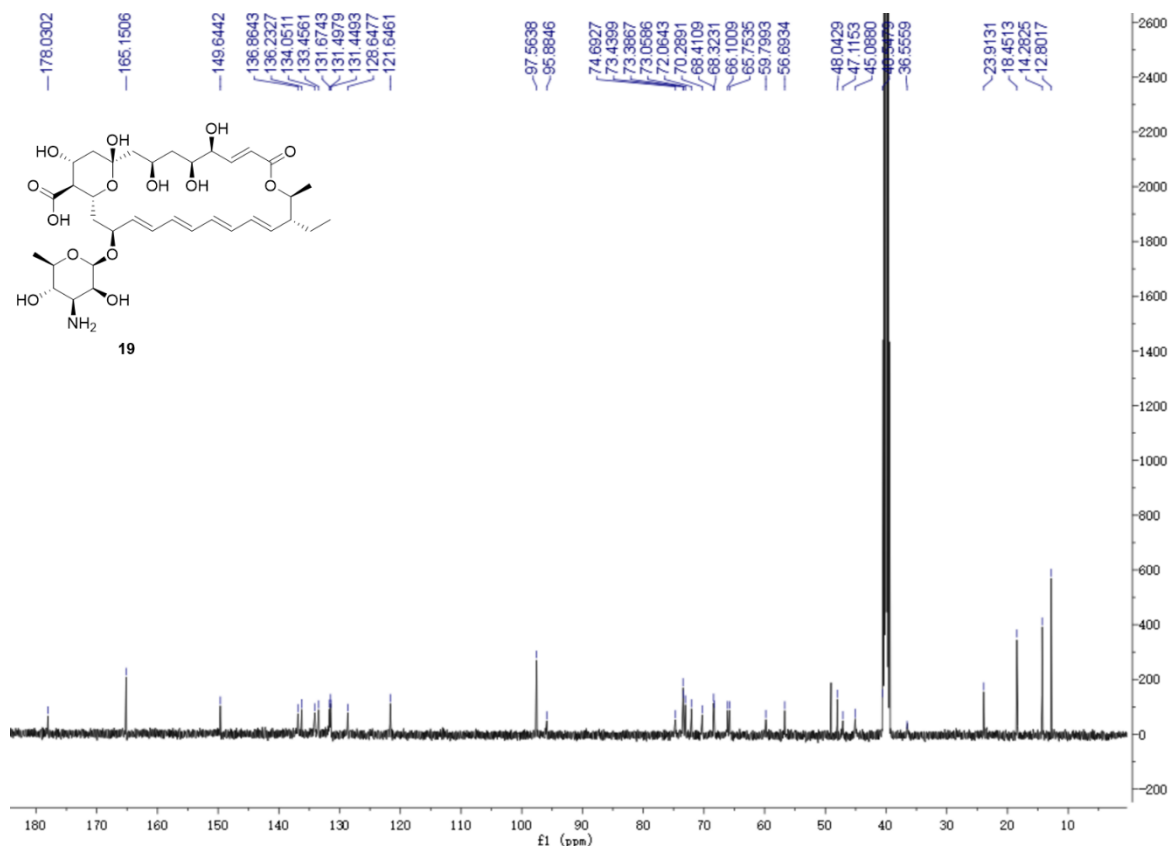
Supplementary Figure S65.  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound 17.

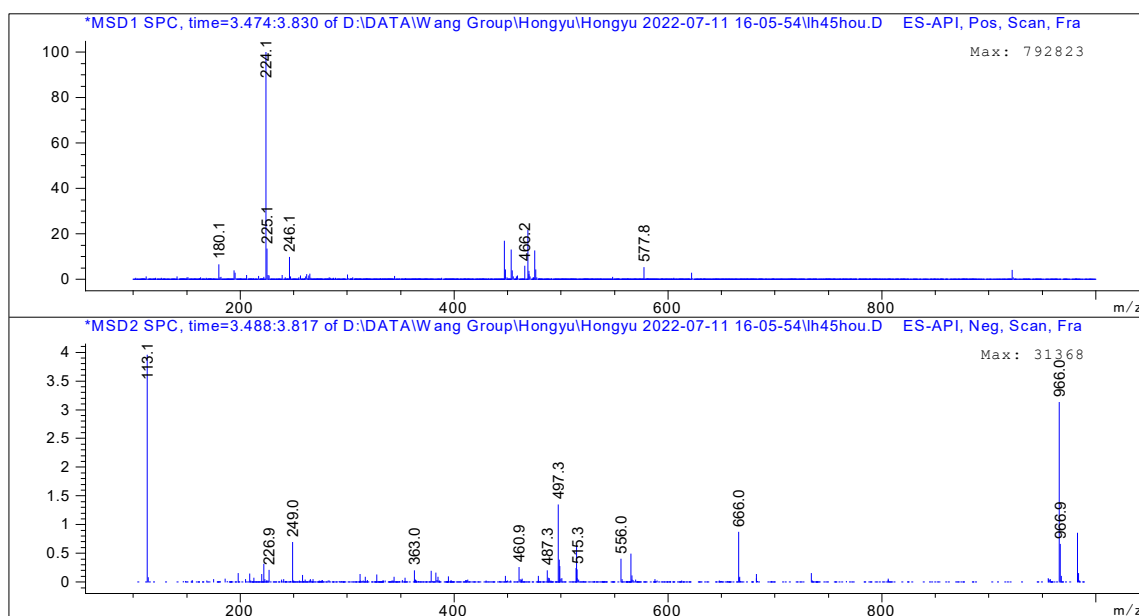


Supplementary Figure S66. (+) and (-)-ESI-MS of compound **18**.

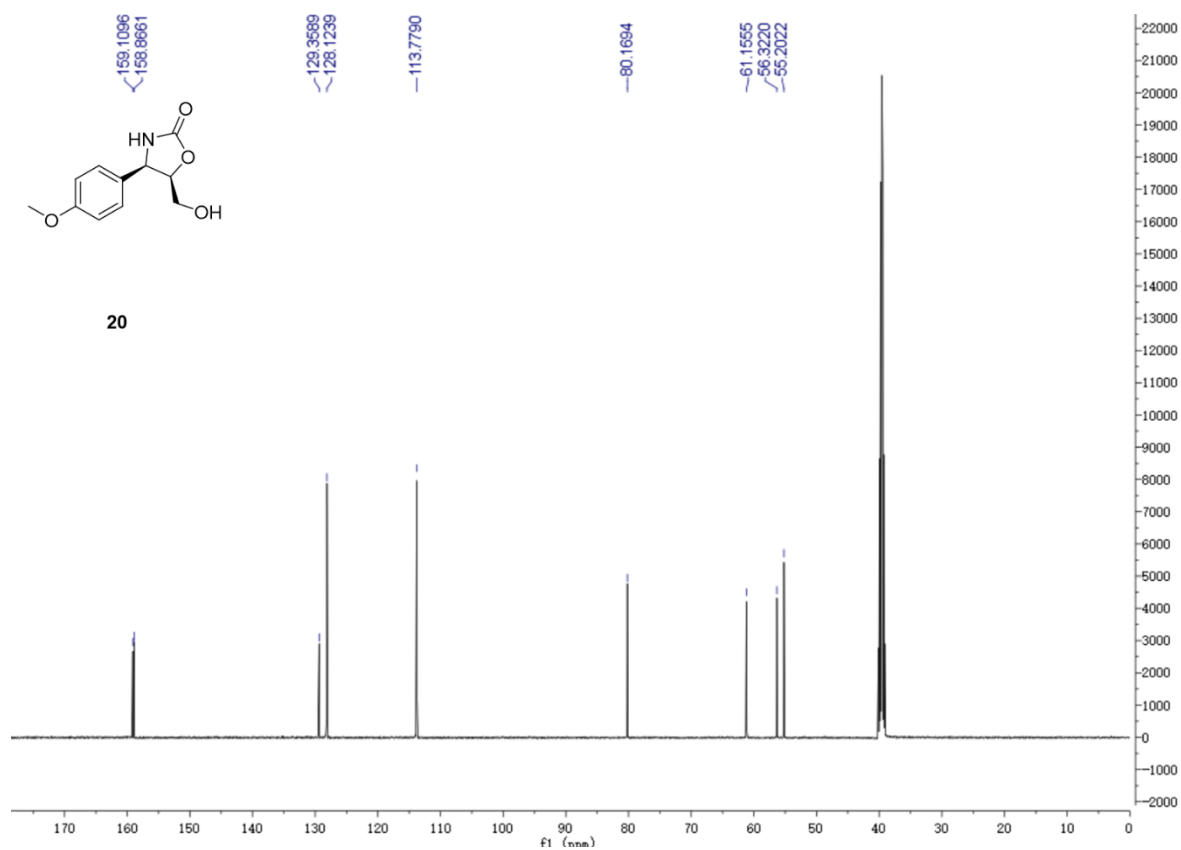


Supplementary Figure S67.  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **18**.

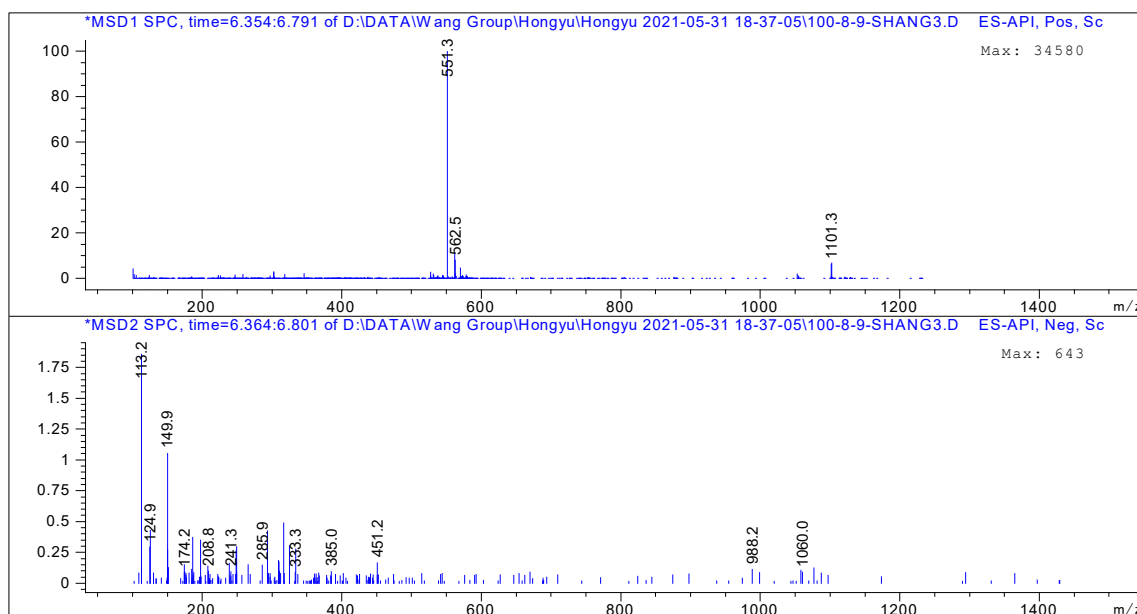
Supplementary Figure S68. (+) and (-)-ESI-MS of compound **19**.Supplementary Figure S69.  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **19**.



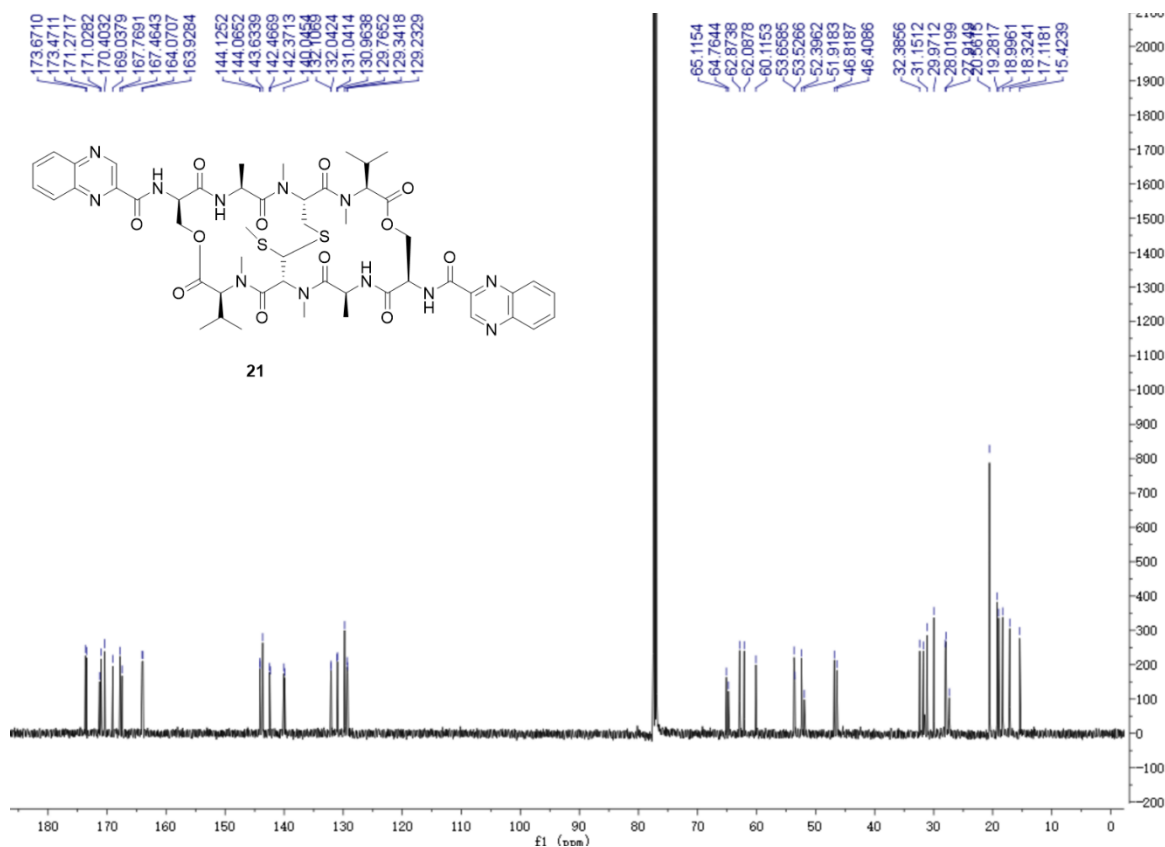
**Supplementary Figure S70.** (+) and (-)-ESI-MS of compound **20**.



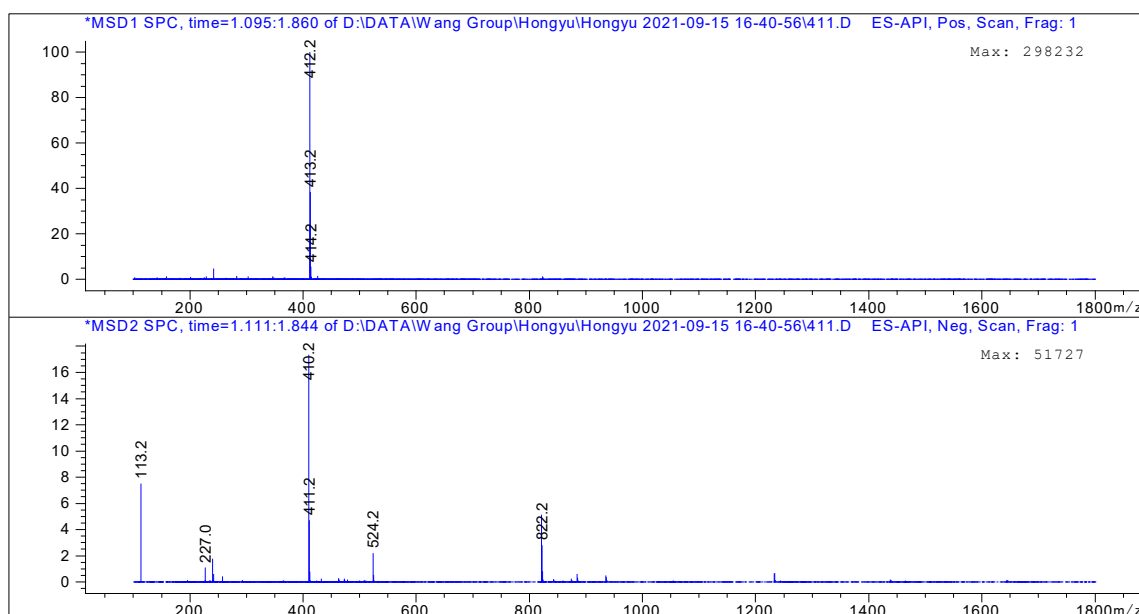
**Supplementary Figure S71.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **20**.



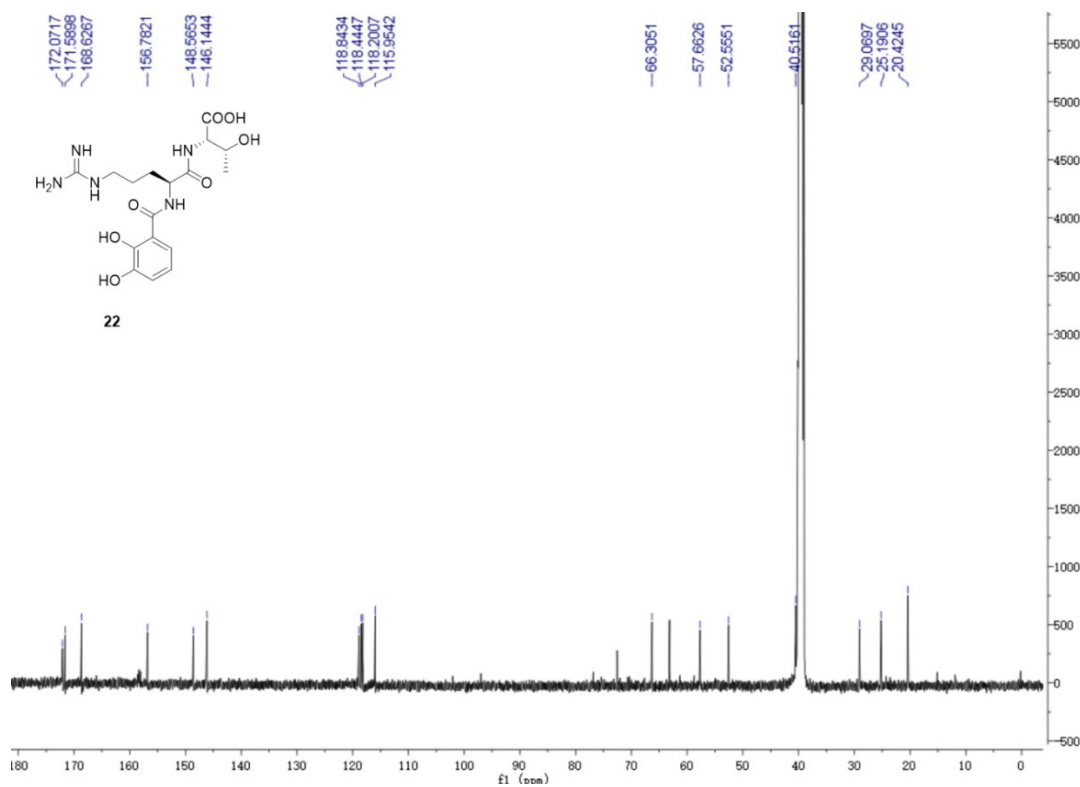
**Supplementary Figure S72.** (+) and (-)-ESI-MS of compound **21**.



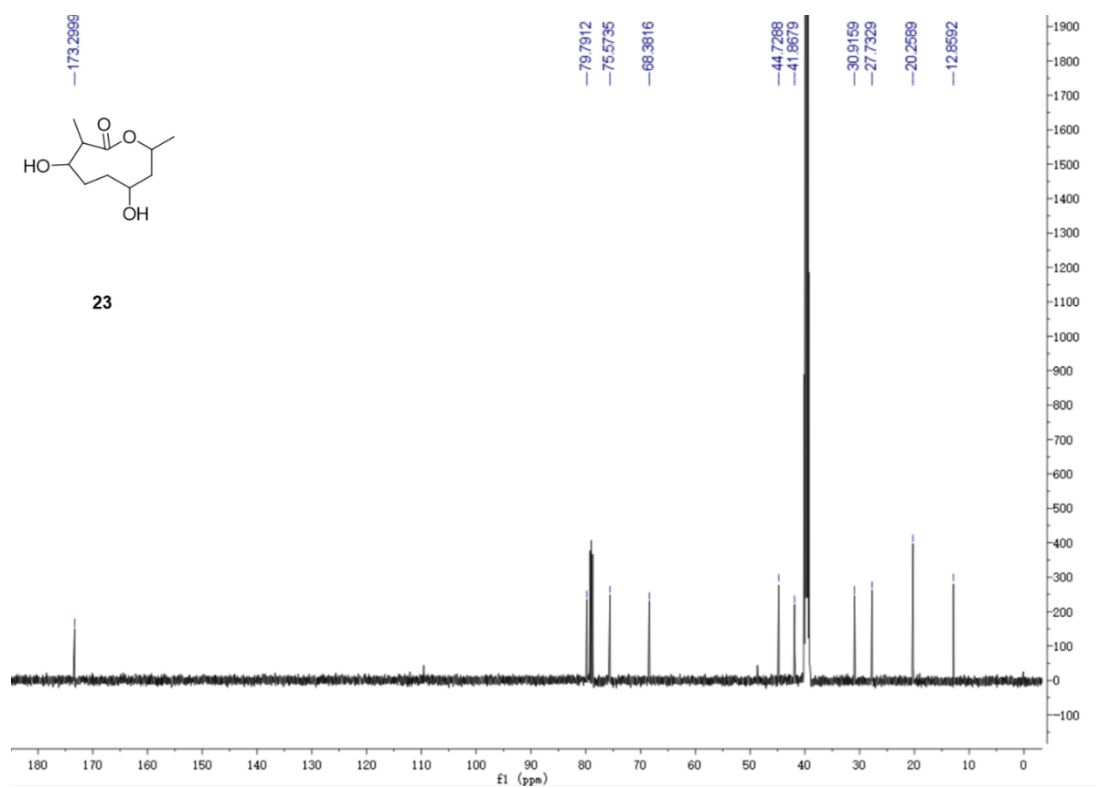
**Supplementary Figure S73.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{CDCl}_3$ ) spectrum of compound **21**.



**Supplementary Figure S74.** (+) and (-)-ESI-MS of compound **22**.



**Supplementary Figure S75.**  $^{13}\text{C}$  NMR (125 MHz,  $\text{DMSO}-d_6$ ) spectrum of compound **22**.



Supplementary Figure S76.  $^{13}\text{C}$  NMR (125 MHz, DMSO- $d_6$ ) spectrum of compound **23**.

## 1.2 Supplementary Tables

**Supplementary Table S1.** Phylogenetic analyses of 124 endophytic strains from *Achyranthes bidentata*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	NX-R-1	Root	<i>Streptomyces triostinicus</i>	100.00%	OQ438164
2	NX-R-2	Root	<i>Streptomyces manipurensis</i>	99.93%	OQ438165
3	NX-R-3	Root	<i>Streptomyces</i> sp. CPE338	100.00%	OQ438166
4	NX-R-4	Root	<i>Streptomyces paradoxus</i>	96.61%	OQ438167
5	NX-R-5	Root	Same as 45	-	-
6	NX-R-6	Root	<i>Streptomyces flavogriseus</i>	99.93%	OQ438168
7	NX-R-7	Root	<i>Bacillus thuringiensis</i>	99.93%	OQ438169
8	NX-R-8	Root	<i>Streptomyces</i> sp.	98.04%	OQ438170
9	NX-R-9	Root	<i>Streptomyces phaeovorticillatus</i> subsp. <i>takatsukiensis</i>	98.24%	OQ438171
10	NX-R-10	Root	<i>Streptomyces collinus</i>	99.93%	OQ438172
11	NX-R-11	Root	<i>Streptomyces</i> sp.	99.86%	OQ438173
12	NX-R-12	Root	<i>Streptomyces</i> sp.	99.93%	OQ438174
13	NX-R-13	Root	<i>Streptomyces</i> sp. X2-21	99.93%	OQ438175
14	NX-R-14	Root	Same as 16	-	-
15	NX-R-15	Root	<i>Streptomyces</i> sp. MG62	99.72%	OQ438176
16	NX-R-16	Root	<i>Streptomyces sporovirgulis</i>	99.72%	OQ438177
17	NX-R-17	Root	<i>Streptomyces</i> sp.	95.07%	OQ438178
18	NX-R-18	Root	<i>Streptomyces</i> sp.	99.81%	OQ438179
19	NX-R-19	Root	<i>Streptomyces chartreusis</i>	96.33%	OQ438180
20	NX-R-20	Root	<i>Streptomyces cadmiisoli</i>	99.93%	OQ438181
21	NX-R-21	Root	<i>Bacillus anthracis</i>	100.00%	OQ438182
22	NX-R-22	Root	<i>Streptomyces</i> sp. RM395	100.00%	OQ438183
23	NX-R-23	Root	<i>Streptomyces phaeochromogenes</i>	99.93%	OQ438184
24	NX-R-24	Root	<i>Streptomyces halstedii</i>	99.79%	OQ438185
25	NX-R-25	Root	<i>Streptomyces</i> sp.	98.72%	OQ438186
26	NX-R-26	Root	<i>Streptomyces canus</i>	99.72%	OQ438187
27	NX-R-27	Root	<i>Streptomyces</i> sp. SOK6/25-05	99.93%	OQ438188
28	NX-R-28	Root	<i>Streptomyces canus</i>	99.93%	OQ438189
29	NX-R-29	Root	<i>Streptomyces</i> sp. SOK6/25-05	99.93%	OQ438190
30	NX-R-30	Root	<i>Actinobacteria</i> bacterium	100.00%	OQ438191
31	NX-R-31	Root	<i>Streptomyces canus</i>	99.86%	OQ438192
32	NX-R-32	Root	<i>Streptomyces</i> sp. GLY-P2	99.39%	OQ438193
33	NX-R-33	Root	<i>Streptomyces</i> sp.	97.43%	OQ438194
34	NX-R-34	Root	<i>Streptomyces parvus</i>	99.86%	OQ438195
35	NX-R-35	Root	<i>Streptomyces violascens</i>	100.00%	OQ438196
36	NX-R-36	Root	<i>Streptomyces flavogriseus</i>	99.93%	OQ438197
37	NX-R-37	Root	<i>Streptomyces</i> sp. Sn-13	99.86%	OQ438198
38	NX-R-38	Root	<i>Streptomyces flavogriseus</i>	99.93%	OQ438199
39	NX-R-39	Root	<i>Streptomyces canus</i>	100.00%	OQ438200
40	NX-R-40	Root	<i>Streptomyces beijiangensis</i>	99.81%	OQ438201
41	NX-R-41	Root	<i>Streptomyces phaeochromogenes</i>	99.71%	OQ438202
42	NX-R-42	Root	<i>Streptomyces</i> sp.	100.00%	OQ438203
43	NX-R-43	Root	<i>Streptomyces tauricus</i>	100.00%	OQ438204
44	NX-R-44	Root	<i>Streptomyces</i> sp. X1-2	99.66%	OQ438205
45	NX-R-45	Root	<i>Streptomyces</i> sp.	99.59%	OQ438206
46	NX-R-46	Root	<i>Rhodococcus</i> sp. 7-181	99.93%	OQ438207

47	NX-R-47	Root	<i>Streptomyces tauricus</i>	100.00%	OQ438208
48	NX-R-48	Root	<i>Bacillus thuringiensis</i>	99.93%	OQ438209
49	NX-R-49	Root	<i>Bacillus cereus</i>	99.93%	OQ438210
50	NX-R-50	Root	<i>Streptomyces parvus</i>	99.79%	OQ438211
51	NX-R-51	Root	<i>Streptomyces</i> sp.	99.37%	OQ438212
52	NX-R-52	Root	<i>Streptomyces</i> sp.	99.93%	OQ438213
53	NX-R-53	Root	<i>Streptomyces</i> sp.	100.00%	OQ438214
54	NX-R-54	Root	<i>Streptomyces</i> sp. SOK6/25-05	100.00%	OQ438215
55	NX-R-55	Root	<i>Streptomyces</i> sp.	99.93%	OQ438216
56	NX-R-56	Root	<i>Streptomyces manipurensis</i>	99.86%	OQ438217
57	NX-R-57	Root	<i>Streptomyces canus</i>	99.79%	OQ438218
58	NX-R-58	Root	<i>Streptomyces collinus</i>	100.00%	OQ438219
59	NX-R-59	Root	<i>Streptomyces microflavus</i>	99.86%	OQ438220
60	NX-R-60	Root	<i>Streptomyces anulatus</i>	99.93%	OQ438221
61	NX-R-61	Root	<i>Streptomyces manipurensis</i>	99.93%	OQ438222
62	NX-R-62	Root	<i>Streptomyces</i> sp.	100.00%	OQ438223
63	NX-R-63	Root	<i>Streptomyces</i> sp.	99.93%	OQ438224
64	NX-R-64	Root	<i>Streptomyces collinus</i>	100.00%	OQ438225
65	NX-R-65	Root	<i>Streptomyces</i> sp. CPE338	100.00%	OQ438226
66	NX-R-66	Root	<i>Streptomyces</i> sp. GLY-P2	99.86%	OQ438227
67	NX-R-67	Root	<i>Nocardia</i> sp. 30903	99.93%	OQ438228
68	NX-R-68	Root	<i>Streptomyces</i> sp. YRA166	98.94%	OQ438229
69	NX-R-69	Root	<i>Streptomyces</i> sp. A333 Ydz-QZ	100.00%	OQ438230
70	NX-R-70	Root	<i>Nocardia asteroides</i>	100.00%	OQ438231
71	NX-R-71	Root	<i>Bacillus cereus</i>	100.00%	OQ438232
72	NX-R-72	Root	<i>Streptomyces</i> sp. 0616206	99.86%	OQ438233
73	NX-R-73	Root	<i>Streptomyces neyagawaensis</i>	99.79%	OQ438234
74	NX-R-74	Root	<i>Streptomyces anulatus</i>	99.86%	OQ438235
75	NX-L-1	Leaf	<i>Streptomyces</i> sp. NBRC 3111	99.93%	OQ438236
76	NX-L-2	Leaf	<i>Streptomyces</i> sp. X4-1	99.64%	OQ438237
77	NX-L-3	Leaf	<i>Actinobacteria</i> bacterium	99.26%	OQ438238
78	NX-L-4	Leaf	<i>Streptomyces</i> sp. CPE314	97.31%	OQ438239
79	NX-L-5	Leaf	<i>Streptomyces</i> sp. BF-9	99.93%	OQ438240
80	NX-L-6	Leaf	<i>Streptomyces</i> sp. MM117	99.86%	OQ438241
81	NX-L-7	Leaf	<i>Streptomyces griseolus</i>	99.51%	OQ438242
82	NX-L-8	Leaf	<i>Streptomyces</i> sp.	99.93%	OQ438243
83	NX-L-9	Leaf	<i>Streptomyces</i> sp.	99.42%	OQ438244
84	NX-L-10	Leaf	<i>Streptomyces rubiginosohelvolus</i>	99.86%	OQ438245
85	NX-L-11	Leaf	<i>Streptomyces tricolor</i>	99.79%	OQ438246
86	NX-L-12	Leaf	<i>Streptomyces zaomyceticus</i>	99.93%	OQ438247
87	NX-L-13	Leaf	<i>Streptomyces parvus</i>	99.72%	OQ438248
88	NX-L-14	Leaf	<i>Streptomyces</i> sp.	98.65%	OQ438249
89	NX-L-15	Leaf	<i>Streptomyces</i> sp. QZGY-A35	99.79%	OQ438250
90	NX-L-16	Leaf	Same as 123	-	-
91	NX-L-17	Leaf	<i>Streptomyces variabilis</i>	100.00%	OQ438251
92	NX-L-18	Leaf	<i>Streptomyces zaomyceticus</i>	99.93%	OQ438252
93	NX-L-19	Leaf	<i>Nocardia asteroides</i>	100.00%	OQ438253
94	NX-L-20	Leaf	<i>Methylobacterium brachiatum</i>	99.90%	OQ438254
95	NX-L-21	Leaf	<i>Bacillus cereus</i>	99.16%	OQ438255
96	NX-L-22	Leaf	<i>Microbacterium</i> sp. PTS2402	94.75%	OQ438256
97	NX-L-23	Leaf	<i>Streptomyces</i> sp. STR43	98.75%	OQ438257
98	NX-L-24	Leaf	<i>Streptomyces</i> sp.	99.93%	OQ438258
99	NX-L-25	Leaf	<i>Streptomyces violascens</i>	100.00%	OQ438259

100	NX-L-26	Leaf	<i>Streptomyces</i> sp. A160 Ydz-ZZ	100.00%	OQ438260
101	NX-L-27	Leaf	<i>Streptomyces microflavus</i>	99.93%	OQ438261
102	NX-L-28	Leaf	<i>Streptomyces parvus</i>	99.93%	OQ438262
103	NX-L-29	Leaf	<i>Streptomyces halstedii</i>	99.79%	OQ438263
104	NX-L-30	Leaf	<i>Streptomyces akiyoshiensis</i>	99.93%	OQ438264
105	NX-L-31	Leaf	<i>Streptomyces</i> sp. MM14	99.86%	OQ438265
106	NX-S-1	Stem	<i>Streptomyces</i> sp. E5N187	99.86%	OQ438266
107	NX-S-2	Stem	<i>Streptomyces</i> sp. wa22	98.28%	OQ438267
108	NX-S-3	Stem	<i>Streptomyces venezuelae</i>	99.86%	OQ438268
109	NX-S-4	Stem	<i>Streptomyces</i> sp. X54	99.79%	OQ438269
110	NX-S-5	Stem	<i>Streptomyces halstedii</i>	99.23%	OQ438270
111	NX-S-6	Stem	<i>Penicillium</i> sp.	99.63%	OQ438028
112	NX-S-7	Stem	<i>Nocardia asteroides</i>	100.00%	OQ438271
113	NX-S-8	Stem	<i>Streptomyces anulatus</i>	99.86%	OQ438272
114	NX-S-9	Stem	<i>Streptomyces</i> sp. SM18	96.96%	OQ438273
115	NX-S-10	Stem	<i>Nocardia asteroides</i>	100.00%	OQ438274
116	NX-S-11	Stem	<i>Streptomyces</i> sp.	98.64%	OQ438275
117	NX-S-12	Stem	<i>Streptomyces</i> sp.	99.86%	OQ438276
118	NX-S-13	Stem	<i>Streptomyces flavogriseus</i>	100.00%	OQ438277
119	NX-S-14	Stem	<i>Nocardia asteroides</i>	100.00%	OQ438278
120	NX-S-15	Stem	<i>Streptomyces</i> sp. A2-16	99.86%	OQ438279
121	NX-S-16	Stem	<i>Nocardia asteroides</i>	100.00%	OQ438280
122	NX-S-17	Stem	<i>Streptomyces griseolus</i>	99.51%	OQ438281
123	NX-S-18	Stem	<i>Bacillus</i> sp. CSB5	83.70%	OQ438282
124	NX-AP-1	Aerial Part	<i>Bacillus cereus</i>	99.86%	OQ438283

(Red color: selected for further study).

**Supplementary Table S2.** Phylogenetic analyses of 13 endophytic strains from *Ainsliaea fragrans*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	XXTEF-R-1	Root	<i>Streptomyces anulatus</i>	99.93%	OQ438816
2	XXTEF-SL-1	Stem-Leaf	<i>Streptomyces</i> sp. INBio_4506AN	99.93%	OQ438817
3	XXTEF-SL-2	Stem-Leaf	<i>Streptomyces</i> sp. SM18	99.93%	OQ438818
4	XXTEF-SL-3	Stem-Leaf	<i>Streptomyces mucoflavus</i>	98.56%	OQ438819
5	XXTEF-SL-4	Stem-Leaf	<i>Streptomyces mucoflavus</i>	98.56%	OQ438820
6	XXTEF-SL-5	Stem-Leaf	<i>Streptomyces</i> sp. SM18	99.93%	OQ438821
7	XXTEF-SL-6	Stem-Leaf	<i>Streptomyces halstedii</i>	99.93%	OQ438822
8	XXTEF-SL-7	Stem-Leaf	<i>Streptomyces variabilis</i>	100.00%	OQ438823
9	XXTEF-SL-8	Stem-Leaf	<i>Streptomyces</i> sp. SM18	99.93%	OQ438824
10	XXTEF-SL-9	Stem-Leaf	<i>Streptomyces</i> sp. MJM4235	99.86%	OQ438825
11	XXTEF-SL-10	Stem-Leaf	<i>Streptomyces mucoflavus</i>	98.56%	OQ438826
12	XXTEF-F-1	Flower	<i>Micromonospora chokoriensis</i>	99.79%	OQ438827
13	XXTEF-F-2	Flower	<i>Bacillus thuringiensis</i>	99.93%	OQ438828

**Supplementary Table S3.** Phylogenetic analyses of 11 endophytic strains from *Artemisia capillaris*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	YCH-R-1	Root	<i>Streptomyces chartreusis</i> NRRL 3882	100.00%	OQ430664
2	YCH-L-1	Leaf	<i>Streptomyces zaomyceticus</i>	100.00%	OQ430665
3	YCH-AP-1	Aerial Part	<i>Microbacterium hydrocarbonoxydans</i>	99.93%	OQ430666
4	YCH-AP-2	Aerial Part	<i>Terribacillus goriensis</i>	100.00%	OQ430667
5	YCH-AP-3	Aerial Part	<i>Streptomyces</i> sp.	100.00%	OQ430668
6	YCH-AP-4	Aerial Part	<i>Streptomyces</i> sp. A311Ydz-AH	99.86%	OQ430669
7	YCH-AP-5	Aerial Part	<i>Staphylococcus xylosus</i>	99.93%	OQ430670
8	YCH-AP-6	Aerial Part	<i>Streptomyces</i> sp.	99.17%	OQ430671
9	YCH-AP-7	Aerial Part	<i>Streptomyces</i> sp. DUT_AHX	99.93%	OQ430672
10	YCH-AP-8	Aerial Part	<i>Streptomyces neopeptini</i>	99.79%	OQ430673
11	YCH-AP-9	Aerial Part	<i>Streptomyces pseudovenezuelae</i>	99.93%	OQ430674

**Supplementary Table S4.** Phylogenetic analyses of 13 endophytic strains from *Daucus carota*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	YHLB-R-1	Root	<i>Streptomyces griseolus</i>	98.38%	OQ438148
2	YHLB-R-2	Root	<i>Streptomyces halstedii</i>	99.86%	OQ438149
3	YHLB-R-3	Root	<i>Streptomyces flavogriseus</i>	100.00%	OQ438150
4	YHLB-L-1	Leaf	<i>Methylobacterium aquaticum</i>	97.75%	OQ438151
5	YHLB-L-2	Leaf	<i>Streptomyces diastatochromogenes</i>	100.00%	OQ438152
6	YHLB-L-3	Leaf	<i>Microbacterium oxydans</i>	94.18%	OQ438153
7	YHLB-L-4	Leaf	<i>Sphingomonas</i> sp.	99.93%	OQ438154
8	YHLB-S-1	Stem	<i>Streptomyces flaveus</i>	99.93%	OQ438155
9	YHLB-S-2	Stem	<i>Streptomyces griseolus</i>	99.79%	OQ438156
10	YHLB-S-3	Stem	<i>Streptomyces flavogriseus</i>	100.00%	OQ438157
11	YHLB-F-1	Flower	<i>Kineococcus tegulae</i>	99.54%	OQ438158
12	YHLB-F-2	Flower	Same as 13	-	-
13	YHLB-F-3	Flower	<i>Methylobacterium</i> sp. N7	99.93%	OQ438159

(Red color: selected for further study).

**Supplementary Table S5.** Phylogenetic analyses of 21 endophytic strains from *Leonurus japonicus*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	YMC-R-1	Root	<i>Streptomyces globisporus</i>	99.93%	OQ438707
2	YMC-R-2	Root	<i>Bacillus albus</i>	99.93%	OQ438708
3	YMC-R-3	Root	<i>Streptomyces venezuelae</i>	99.72%	OQ438709
4	YMC-R-4	Root	<i>Microbacterium</i> sp. SuP-E064	99.64%	OQ438710
5	YMC-R-5	Root	<i>Streptomyces zaomyceticus</i>	99.79%	OQ438711
6	YMC-R-6	Root	<i>Streptomyces</i> sp. NJYH2618	99.86%	OQ438712
7	YMC-R-7	Root	<i>Streptomyces</i> sp. X1-2	98.86%	OQ438713
8	YMC-R-8	Root	<i>Streptomyces</i> sp.	99.38%	OQ438714
9	YMC-R-9	Root	<i>Streptomyces albogriseolus</i>	99.93%	OQ438715
10	YMC-R-10	Root	<i>Streptomyces albogriseolus</i>	100.00%	OQ438716
11	YMC-R-11	Root	<i>Bacillus cereus</i>	99.93%	OQ438717
12	YMC-SL-1	Stem-Leaf	<i>Methylobacterium populi</i>	97.96%	OQ438718
13	YMC-SL-2	Stem-Leaf	<i>Streptomyces lateritius</i>	99.86%	OQ438719
14	YMC-SL-3	Stem-Leaf	<i>Curtobacterium</i> sp. b163	100.00%	OQ438720
15	YMC-F-1	Flower	<i>Streptomyces</i> sp. HF-0	100.00%	OQ438721
16	YMC-F-2	Flower	<i>Streptomyces</i> sp. G49	99.03%	OQ438722
17	YMC-F-3	Flower	<i>Streptomyces</i> sp. 3076	99.79%	OQ438742
18	YMC-F-4	Flower	<i>Streptomyces</i> sp. FSRh8	99.86%	OQ438723
19	YMC-F-5	Flower	<i>Streptomyces griseolus</i>	99.93%	OQ438724
20	YMC-F-6	Flower	<i>Streptomyces</i> sp. SM18	99.86%	OQ438725
21	YMC-F-7	Flower	<i>Streptomyces</i> sp. SM18	99.79%	OQ438726

**Supplementary Table S6.** Phylogenetic analyses of 15 endophytic strains from *Perilla frutescens*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	BS-R-1	Root	<i>Streptomyces fragilis</i>	99.65%	OQ438801
2	BS-AP-1	Aerial-Part	<i>Streptomyces</i> sp.	99.79%	OQ438802
3	BS-AP-2	Aerial-Part	<i>Streptomyces</i> sp. JSM 147702	99.93%	OQ438803
4	BS-AP-3	Aerial-Part	<i>Streptomyces tanashiensis</i>	99.36%	OQ438804
5	BS-AP-4	Aerial-Part	<i>Streptomyces kunmingensis</i>	99.93%	OQ438805
6	BS-AP-5	Aerial-Part	<i>Streptomyces</i> sp.	99.57%	OQ438806
7	BS-AP-6	Aerial-Part	<i>Methylobacterium</i> sp. MG-2011-78-DC	99.86%	OQ438807
8	BS-AP-7	Aerial-Part	<i>Nakamurella</i> sp.	99.63%	OQ438808
9	BS-AP-8	Aerial-Part	<i>Nakamurella</i> sp.	99.92%	OQ438809
10	BS-AP-9	Aerial-Part	<i>Streptomyces kunmingensis</i>	99.79%	OQ438810
11	BS-1	Whole Plant	<i>Nocardia</i> sp. HBUM 79084	100.00%	OQ438811
12	BS-2	Whole Plant	<i>Streptomyces</i> sp. ACTYS11	99.93%	OQ438812
13	BS-3	Whole Plant	<i>Streptomyces</i> sp. FSRh6	99.86%	OQ438813
14	BS-4	Whole Plant	<i>Streptomyces akiyoshiensis</i>	99.52%	OQ438814
15	BS-5	Whole Plant	<i>Streptomyces</i> sp. JSM 147702	99.93%	OQ438815

**Supplementary Table S7.** Phylogenetic analyses of 28 endophytic strains from *Prunella vulgaris*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	XKC-R-1	Root	<i>Streptomyces</i> sp. SM18	99.66%	OQ449506
2	XKC-R-2	Root	<i>Streptomyces</i> sp. W	99.93%	OQ449507
3	XKC-R-3	Root	<i>Streptomyces zaomyceticus</i>	99.93%	OQ449508
4	XKC-R-4	Root	<i>Streptomyces</i> sp. 18(2008)	99.93%	OQ449509
5	XKC-R-5	Root	<i>Streptomyces</i> sp. SM18	99.86%	OQ449510
6	XKC-R-6	Root	<i>Bacillus cereus</i>	99.86%	OQ449511
7	XKC-R-7	Root	<i>Streptomyces</i> sp. ZG634	99.72%	OQ449512
8	XKC-R-8	Root	<i>Streptomyces</i> sp. TY68-3	100.00%	OQ449519
9	XKC-R-9	Root	<i>Streptomyces</i> sp.	98.00%	OQ449513
10	XKC-R-10	Root	<i>Streptomyces canus</i>	96.09%	OQ449514
11	XKC-R-11	Root	<i>Streptomyces lateritius</i>	99.93%	OQ449515
12	XKC-R-12	Root	<i>Staphylococcus</i> sp. MRSA154B15_13_3E	100.00%	OQ449516
13	XKC-R-13	Root	<i>Streptomyces</i> sp. H	99.86%	OQ449517
14	XKC-R-14	Root	<i>Streptomyces parvus</i>	97.92%	OQ449518
15	XKC-R-15	Root	<i>Streptomyces</i> sp. H	99.93%	OQ449536
16	XKC-R-16	Root	<i>Streptomyces luteogriseus</i>	99.93%	OQ449552
17	XKC-R-17	Root	<i>Streptomyces</i> sp. TY68-3	100.00%	OQ449519
18	XKC-R-18	Root	<i>Streptomyces hygroscopicus</i>	99.86%	OQ449520
19	XKC-R-19	Root	<i>Streptomyces</i> sp.	99.15%	OQ449521
20	XKC-R-20	Root	<i>Streptomyces flavovariabilis</i>	99.86%	OQ449522
21	XKC-SL-1	Stem-Leaf	<i>Streptomyces</i> sp.	97.88%	OQ449523
22	XKC-SL-2	Stem--Leaf	<i>Kocuria</i> sp.	100.00%	OQ449524
23	XKC-SL-3	Stem-Leaf	<i>Microbacterium</i> sp. HP4S	100.00%	OQ449525
24	XKC-SL-4	Stem-Leaf	<i>Methylobacterium</i> sp. N7	99.93%	OQ449526
25	XKC-SL-5	Stem-Leaf	<i>Streptomyces sannanensis</i>	99.65%	OQ449527
26	XKC-F-1	Flower	<i>Pseudomonas parafulva</i>	99.86%	OQ449503
27	XKC-F-2	Flower	<i>Methylobacterium</i> sp. PB280	99.93%	OQ449504
28	XKC-F-3	Flower	<i>Streptomyces</i> sp. SM18	99.86%	OQ449505

**Supplementary Table S8.** Phylogenetic analyses of 16 endophytic strains from *Salvia miltiorrhiza*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	DS-R-1	Root	<i>Nocardia tengchongensis</i>	99.93%	OQ438792
2	DS-SL-1	Stem-Leaf	<i>Streptomyces</i> sp. E5N142	98.27%	OQ438769
3	DS-SL-2	Stem-Leaf	<i>Streptomyces caviscabies</i>	99.79%	OQ438770
4	DS-SL-3	Stem-Leaf	<i>Bosea</i> sp.	99.93%	OQ438771
5	DS-SL-4	Stem-Leaf	<i>Bacillus subtilis</i>	99.86%	OQ438772
6	DS-SL-5	Stem-Leaf	<i>Kocuria carniphila</i>	99.93%	OQ438773
7	DS-SL-6	Stem-Leaf	<i>Streptomyces</i> sp.	99.86%	OQ438774
8	DS-SL-7	Stem-Leaf	<i>Streptomyces glauciniger</i>	99.93%	OQ438775
9	DS-SL-8	Stem-Leaf	<i>Nocardia</i> sp. HBUM 79084	100.00%	OQ438776
10	DS-SL-9	Stem-Leaf	<i>Streptomyces verne</i>	99.86%	OQ438777
11	DS-SL-10	Stem-Leaf	<i>Streptomyces sannanensis</i>	99.79%	OQ438778
12	DS-SL-11	Stem-Leaf	<i>Streptomyces</i> sp. NBRC 3111	99.79%	OQ438779
13	DS-F-1	Flower	<i>Actinomycetia</i> bacterium	100.00%	OQ438780
14	DS-F-2	Flower	<i>Nocardia</i> sp. HBUM 79084	100.00%	OQ438781
15	DS-F-3	Flower	<i>Microbacterium saperdae</i>	99.44%	OQ438782
16	DS-F-4	Flower	<i>Micromonospora aurantiaca</i>	100.00%	OQ438783

**Supplementary Table S9.** Phylogenetic analyses of 28 endophytic strains from *Stemona sessilifolia*.

NO.	Strains	Parts	Closest sequences by BlastN	Percent Identity	Genbank accession
1	ZLBB-RT-1	Root-Tuber	<i>Streptomyces</i> sp. KGS-6-31	99.85%	OQ438618
2	ZLBB-RT-2	Root-Tuber	<i>Streptomyces pseudovenezuelae</i>	99.64%	OQ438619
3	ZLBB-RT-3	Root-Tuber	<i>Streptomyces diastatochromogenes</i>	99.57%	OQ438620
4	ZLBB-RT-4	Root-Tuber	<i>Streptomyces microsporus</i>	99.79%	OQ438621
5	ZLBB-RT-5	Root-Tuber	<i>Streptomyces</i> sp. STR30	99.86%	OQ438622
6	ZLBB-RT-6	Root-Tuber	Same as 9	-	-
7	ZLBB-RT-7	Root-Tuber	<i>Staphylococcus xylosus</i>	99.93%	OQ438623
8	ZLBB-RT-8	Root-Tuber	<i>Nocardia</i> sp. FXJ1.491	100.00%	OQ438624
9	ZLBB-RT-9	Root-Tuber	<i>Nocardia niigatensis</i>	99.85%	OQ438625
10	ZLBB-L-1	Leaf	<i>Streptomyces</i> sp. A46Ydz-XG	99.93%	OQ438626
11	ZLBB-L-2	Leaf	<i>Streptomyces</i> sp. INBio_4515Q	98.78%	OQ438627
12	ZLBB-L-3	Leaf	<i>Streptomyces griseoruber</i>	97.10%	OQ438628
13	ZLBB-L-4	Leaf	<i>Micromonospora</i> sp. A6-9	100.00%	OQ438629
14	ZLBB-L-5	Leaf	<i>Streptomyces</i> sp.	99.79%	OQ438630
15	ZLBB-L-6	Leaf	<i>Streptomyces</i> sp. BF-9	99.72%	OQ438631
16	ZLBB-S-1	Stem	<i>Streptomyces tauricus</i>	100.00%	OQ438632
17	ZLBB-S-2	Stem	<i>Streptomyces flavofungini</i>	99.86%	OQ438633
18	ZLBB-S-3	Stem	<i>Streptomyces</i> sp. A240 Ydz-QZ	99.93%	OQ438634
19	ZLBB-S-4	Stem	<i>Streptomyces thinghirensis</i>	100.00%	OQ438635
20	ZLBB-S-5	Stem	<i>Streptomyces</i> sp. CLI2509	99.93%	OQ438636
21	ZLBB-S-6	Stem	<i>Streptomyces rubiginosohelvolus</i>	100.00%	OQ438637
22	ZLBB-S-7	Stem	<i>Streptomyces</i> sp.	99.93%	OQ438638
23	ZLBB-S-8	Stem	<i>Hyphomicrobium</i> sp.	98.98%	OQ438639
24	ZLBB-S-9	Stem	<i>Streptomyces</i> sp. RI104-LiC104	99.49%	OQ438640
25	ZLBB-S-10	Stem	<i>Streptomyces</i> sp. RI104-LiC104	99.31%	OQ438641
26	ZLBB-S-11	Stem	<i>Nocardia niigatensis</i>	98.00%	OQ438642
27	ZLBB-S-12	Stem	<i>Micromonospora</i> sp. A6-9	100.00%	OQ438643
28	ZLBB-S-13	Stem	<i>Nocardia niigatensis</i>	99.57%	OQ438644

(Red color: selected for further study).

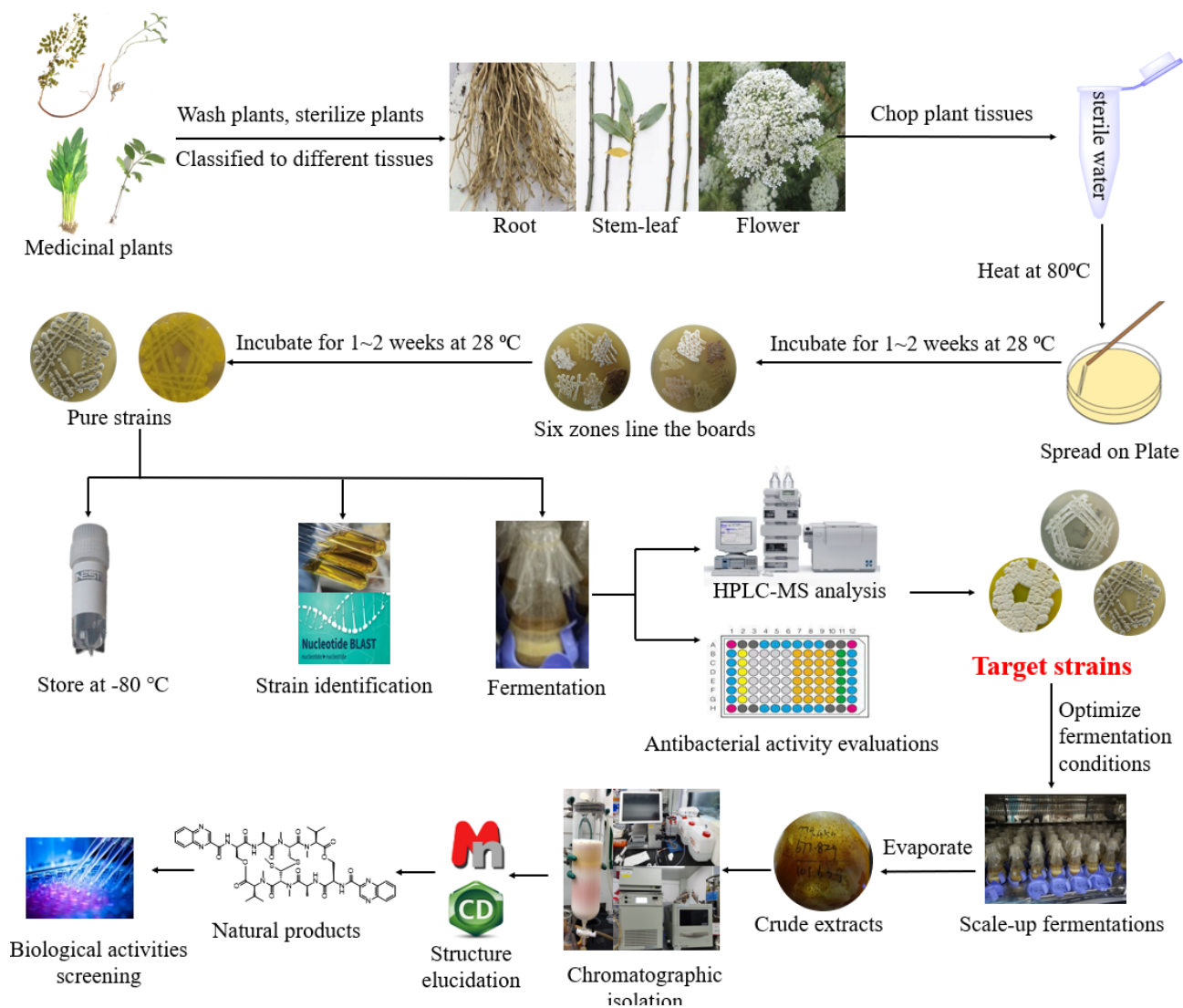
**Supplementary Table S10.** Experimental and calculated  $^{13}\text{C}$  NMR for **1** in  $\text{CDCl}_3$ .

	<b>1</b>	<b>6S-1</b>		<b>6R-1</b>	
No.	$\delta_{\text{exp}}$	$\delta_{\text{cal}}$	$\Delta\delta$	$\delta_{\text{cal}}$	$\Delta\delta$
1	196.5	198.5	-2.0	199.5	-3.0
2	103.4	105.5	-2.1	108.5	-5.1
3	187.9	183.5	4.4	180.5	7.4
4	97.3	99.8	-2.5	102.8	-5.5
5	163.5	159.8	3.7	158.8	4.7
6	75.0	74.9	0.1	74.9	0.1
7	7.6	6.8	0.8	6.1	1.5
8	32.8	30.7	2.1	28.7	4.1
1'	185.1	180.7	4.4	179.7	5.4
2'	123.8	128.5	-4.7	128.6	-4.8
3'	144.1	140.6	3.5	140.6	3.5
4'	131.2	132.9	-1.7	132.9	-1.7
5'	140.4	143.6	-3.2	143.6	-3.2
6'	18.9	17.2	1.7	17.2	1.7

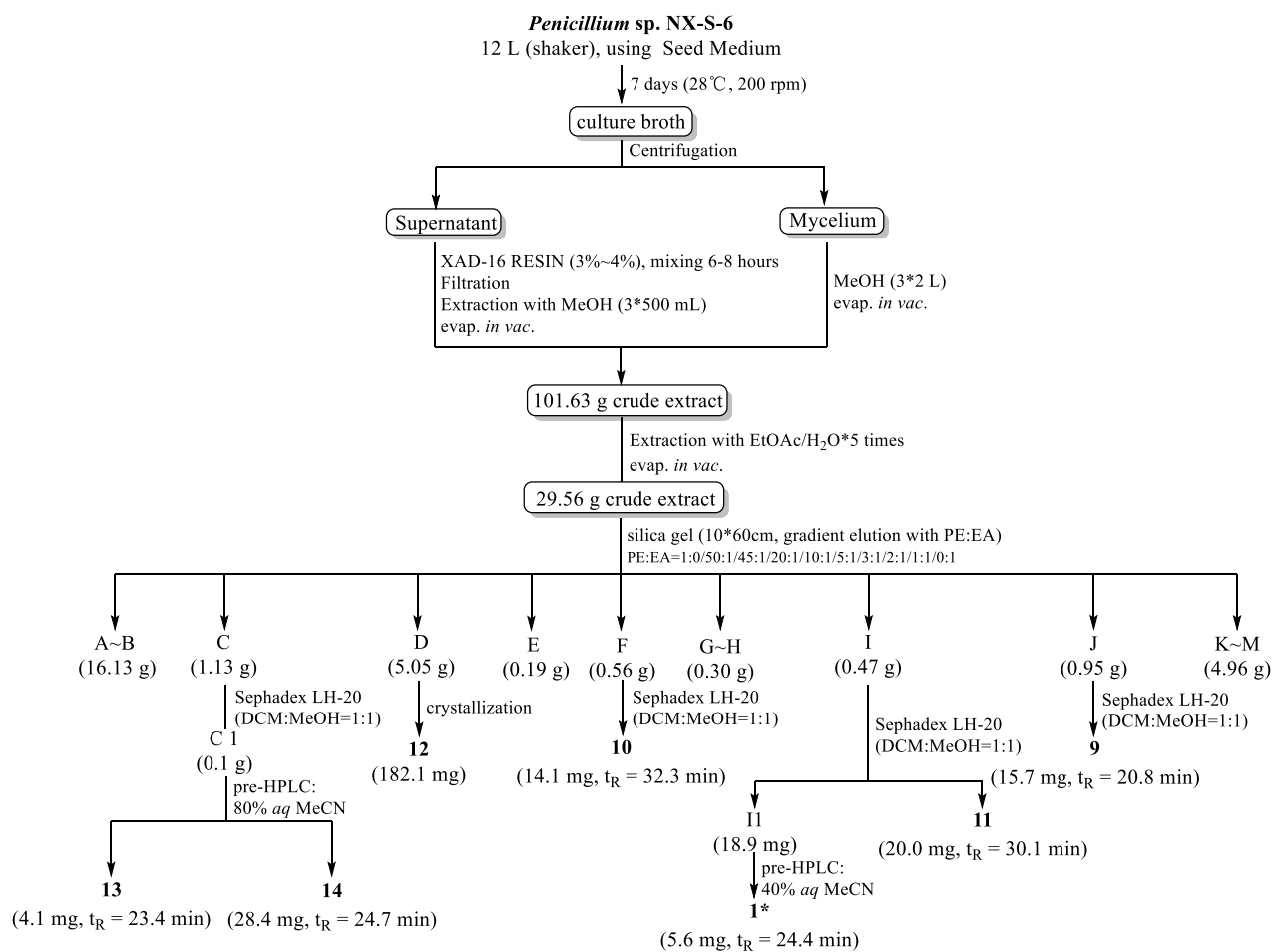
**Supplementary Table S11.** Antibacterial screening results of three selected endophytes (inhibition, %).

Strains	Organisms	
	<i>S. aureus</i>	<i>E. coli</i>
NX-S-6	64	26
YHLB-L-2	51	72
ZLBB-S-6	97	10

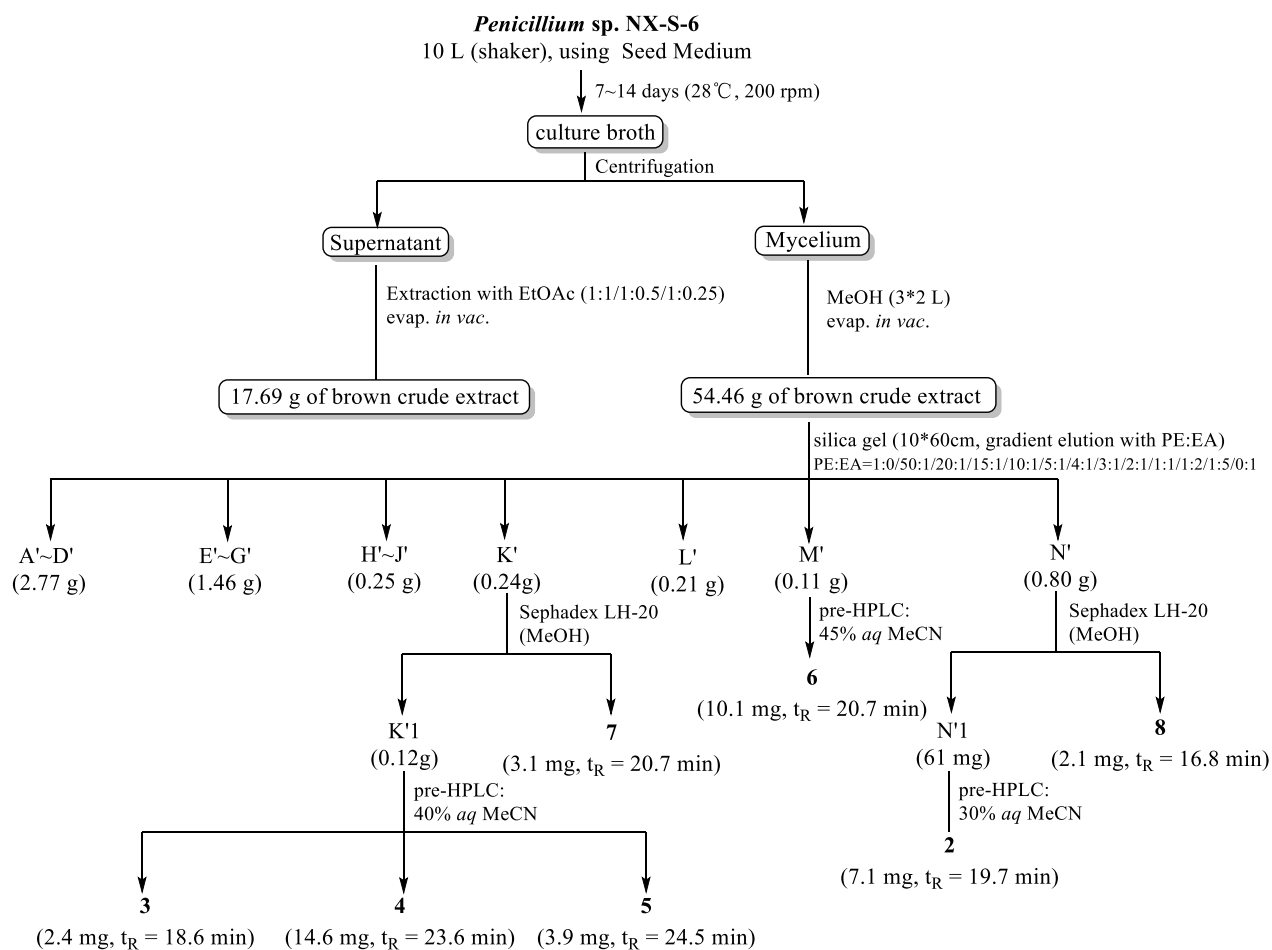
### 1.3 Supplementary Schemes



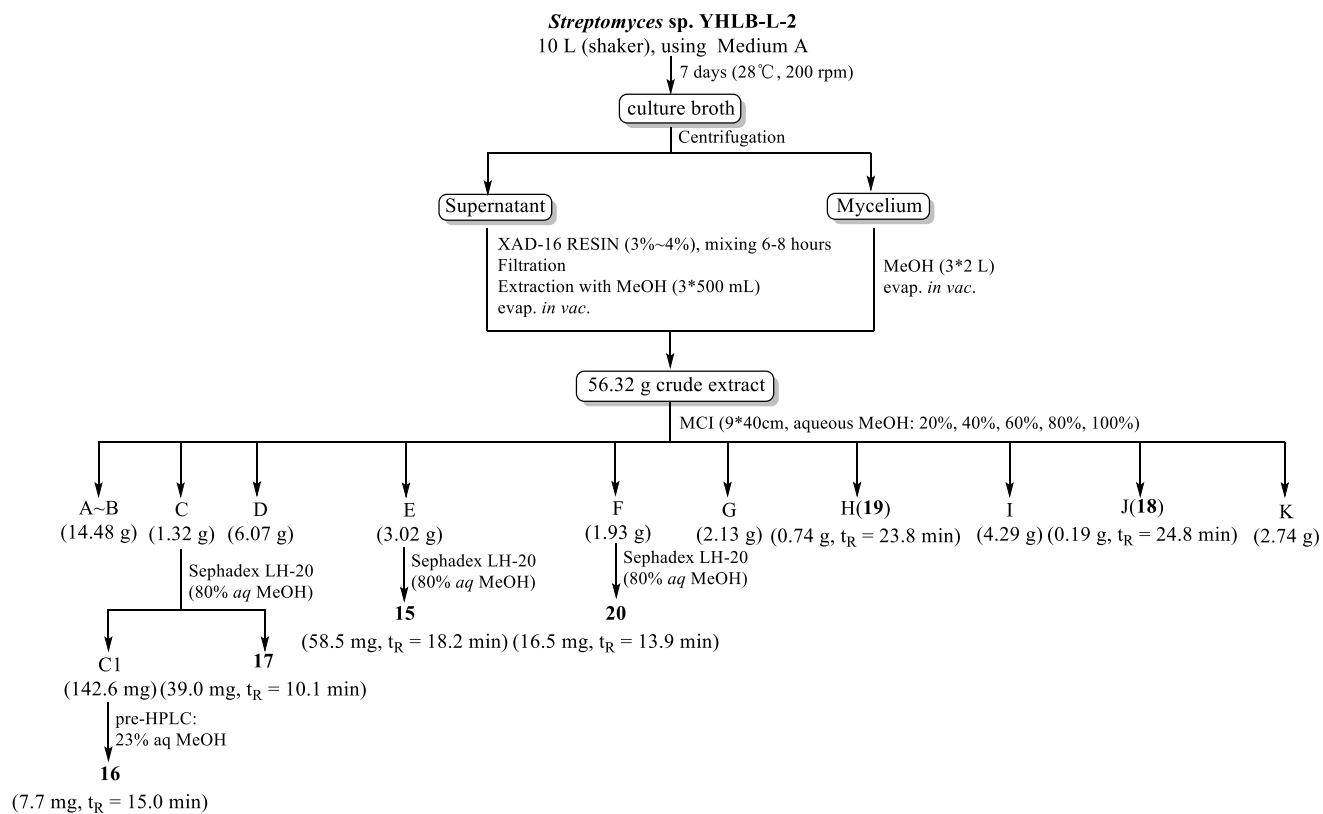
**Supplementary Scheme S1.** Working scheme for this project.



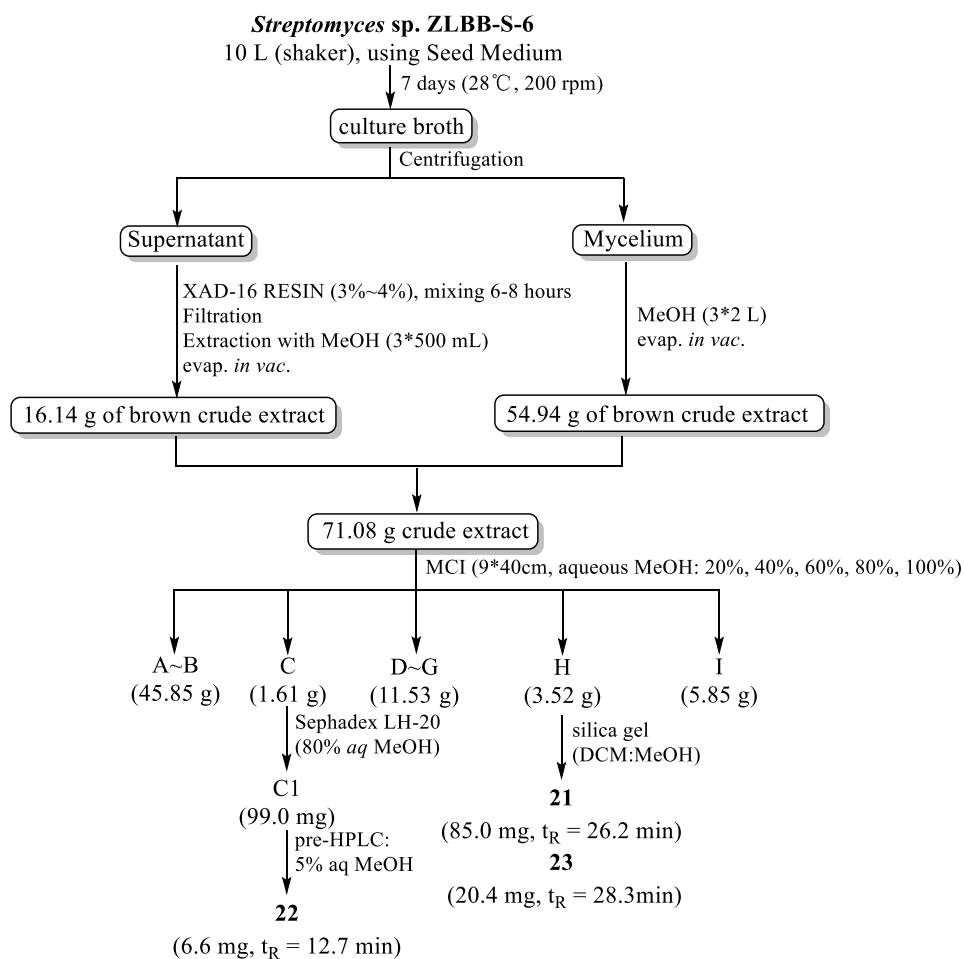
**Supplementary Scheme S2.** Work-up scheme for the *Penicillium* sp. NX-S-6 culture (1<sup>st</sup>).



**Supplementary Scheme S3.** Work-up scheme for the *Penicillium sp.* NX-S-6 culture (2<sup>nd</sup>).



**Supplementary Scheme S4.** Work-up scheme for the *Streptomyces* sp. YHLB-L-2 culture.



**Supplementary Scheme S5.** Work-up scheme for the *Streptomyces* sp. ZLBB-S-6 culture.