# **Supporting Information**

## Spiro Meroterpenoids from Ganoderma applanatum

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## 1. ECD and <sup>13</sup>C NMR chemical shift calculated methods

#### 1.1. ECD calculations of 1-9.

#### **Computational methods**

Molecular Merck force field (MMFF) and DFT/TDDFT calculations were performed with Spartan'14 software package (Wavefunction Inc., Irvine, CA, USA) and Gaussian09 program package,<sup>1</sup> respectively, using default grids and convergence criteria. MMFF conformational search generated low-energy conformers within a 10 kcal/mol energy window were subjected to geometry optimization using DFT method at the B3LYP/6-31G (d) level. Frequency calculations were run at the same level to estimate their relative thermal free energies ( $\Delta G$ ) at 298.15K. Energies of the low-energy conformers in MeOH were re-calculated at the B3LYP/def2-TZVP level. Solvent effects were taken into account by using polarizable continuum model (PCM). The TDDFT calculations were performed using the hybrid  $M06^2$  and BMK<sup>3</sup> and the long-range corrected hybrid CAM-B3LYP4 functionals, and the Ahlrichs' basis set TZVP (triple zeta valence plus polarization).<sup>5</sup> The number of excited states per each molecule was 40. The CD spectra were generated by the program SpecDis<sup>6</sup> using a Gaussian band shape with 0.24 or 0.28 eV exponential half-width from dipole-length dipolar and rotational strengths. The equilibrium population of each conformer at 298.15K was calculated from its relative free energies using Boltzmann statistics. The calculated spectra of compounds were generated from the low-energy conformers according to the Boltzmann weighting of each conformer in MeOH solution. (2'S,3'S)-1 and (2'R,3'S)-2 were subjected to MMFF, DFT, and TDDFT calculations. The calculated ECD spectra of (2'S, 3'R)-1 and (2'R, 3'R)-2 were the mirror images of calculated spectra of (2'R, 3'S)-2 and (2'S, 3'S)-1, respectively.

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**Table S1.** Relative and free energies<sup>*a*</sup> and equilibrium populations<sup>*b*</sup> of low-energy conformers of (2'S,3'S)-1 and (2'R,3'S)-2 in MeOH.

Conformer	$\Delta E$	$\Delta G$	P (%)
(2'S,3'S)-1			
SS-1a	0.0	0.0	60.6
SS-1b	1.01	0.87	13.9
SS-1c	1.34	1.21	7.9
SS-1d	1.41	1.34	6.3
SS-1e	0.87	1.45	5.2
SS-1f	2.11	1.56	4.3
$SS-1g^c$	2.33	2.11	1.7
(2'R,3'S)-2			
RS-2a	0.0	0.0	66.7
RS-2b	0.91	0.72	19.7
RS-2c	0.68	1.19	9.0
RS-2d	1.41	1.91	2.7
$RS-2e^{c}$	2.03	2.53	0.9
$RS-2f^{c}$	2.11	2.77	0.6
$RS-2g^{c}$	2.74	3.02	0.4

<sup>*a*</sup>At the B3LYP/def2-TZVP level, in kcal/mol. <sup>*b*</sup>From  $\Delta G$  values at 298.15 K.

<sup>*c*</sup>Conformer not applied to ECD/TDDFT calculations.

#### 1.2. ECD calculations of 10, 14, and 18.

#### **Computational Methods**

Molecular Merck force field (MMFF) and DFT/TDDFT calculations were performed with Spartan'14 software package (Wavefunction Inc., Irvine, CA, USA) and Gaussian09 program package, Conflex conformational search generated low-energy conformers within a 10 kcal/mol energy was finished by software CONFLEX 7. The predominant conformers of (2'R, 3'R, 7'S)-10 (90%). (90%), (2'S,3'R,6'R)-**14** (90%), (2'S, 3'R, 7'S)-10(2'S,3'S,6'S)-14 (90%), (2'R,3'S,6'R,7'S)-18 (90%) and (2'S,3'S,6'R,7'S)-18 (90%) were optimized by DFT calculation at B3LYP/6-311G (d, p) level with the PCM in MeOH. ECD calculations further were conducted at the B3LYP SCRF(PCM)/6-311G (d, p) level with the PCM in MeOH. For comparisons of the calculated curves and experimental CD spectra, the program SpecD is was used.

# 1.3. ECD and <sup>13</sup>C NMR chemical shift calculated methods of 12, 15, and 20

### **Computational Methods**

The CONFLEX 7 searches based on molecular mechanics with MMFF94S force fields were performed for model compounds of (2'R,3'R,6'S)-12, (2'S,3'R,6'S)-12, (2'R,3'S,6'R)-15, (2'R,3'R,6'S)-15, (2'R,3'S,6'R)-20 and (2'R,3'R,6'S)-20, respectively. All of the predominant conformers (90%) were optimized by DFT calculation at B3LYP/6-311G (d, p) level with the PCM in MeOH. All the above calculations were carried out with the Gaussian 09 package of programs. Under the circumstances, the calculations of their <sup>13</sup>C NMR chemical shifts at MPW1PW91-SCRF/6-311+G (2d, p) level with the PCM in MeOH were performed. Boltzmann averaging over all accessible conformers according to

$$\overline{\sigma}^{x} = \frac{\sum_{\text{confSi}} \sigma_{i}^{x} g_{i} \exp(-E_{i}/RT)}{\sum_{\text{confSi}} g_{i} \exp(-E_{i}/RT)}$$

Where  $\overline{\sigma}^x$  is the Boltzmann-averaged calculated shielding constant for portion *x*,  $\sigma_i^x$  is the shielding constant for portion *x* in conformer *i*, and *Ei* is the potential energy of conformer *i* (relative to the global minimum) kJ mol<sup>-1</sup>, obtained from a single-point solvent calculation on the pas-phase structures as discussed previously. *R* is the molar

gas constant (8.3145 J K<sup>-1</sup> mol<sup>-1</sup>),  $g_i$  is the degeneracy of conformer *i*, and the temperature *T* was taken as 298 K.

Chemical shifts were then calculated according to

$$\delta_{\rm calc}^{x} = \frac{\sigma_{\rm ref} - \overline{\sigma}^{x}}{1 - 10^{-6} \sigma_{\rm ref}}$$

Where  $\delta_{calc}^{x}$  is the calculated chemical shift for portion *x* (in ppm),  $\overline{\sigma}^{x}$  is the shielding constant for carbon *x* as calculated above (again in ppm) and  $\sigma_{ref}$  is the shielding constant for the carbon in tetramethylsilane (TMS). This last value was obtained by minimizing TMS in MeOH at the MPW1PW91-SCRF/6-311+G (2d, p) level and calculating the shielding constant for this structure again at the MPW1PW91-SCRF/6-311+G (2d, p) level; the value obtained was  $\sigma_{ref} = 188.0549$  ppm.

The parameters a and b of the linear regression  $\delta_{calcd} = a\delta_{exptl} + b$ ; the correlation coefficient,  $R^2$ ; the mean absolute error (MAE) defined as  $\Sigma n |\delta_{calcd} - \delta_{exptl}|/n$ ; the corrected mean absolute error (CMAE), defined as  $\Sigma n |\delta_{corr} - \delta_{exptl}|/n$ , where  $\delta_{corr} = (\delta_{calcd} - b)/a$  and therefore corrects for systematic errors were presented.

**Table S2**. The <sup>13</sup>C NMR experimental values and calculated chemical shifts of (2'R,3'R,6'S)-12 and (2'S,3'R,6'S)-12 are compared.

	12	(2' <i>R</i> ,3' <i>R</i> ,6' <i>S</i> )- <b>12</b>		(2'S,3'R,6'S)- <b>12</b>	
No.	$\delta_{ m exptl}$	$\delta_{ m corr}$	$\varDelta {\delta_{ m C}}^a$	$\delta_{ m corr}$	$\varDelta {\delta_{ m C}}^a$
1	167.11	165.77	1.34	166.43	0.68
2	123.32	121.81	1.51	121.81	1.51
3	107.88	107.12	0.76	107.72	0.16
4	153.68	152.71	0.97	150.97	2.71
5	128.16	127.54	0.62	127.90	0.26
6	114.62	113.22	1.40	113.30	1.32
1'	205.22	204.63	0.59	203.83	1.39
2'	98.83	98.87	0.04	102.08	3.25
3'	58.09	56.13	1.96	52.96	5.13
4′	28.37	31.05	2.68	29.75	1.38
5'	30.67	27.61	3.06	26.81	3.86
6'	53.87	53.19	0.68	57.58	3.71
7'	146.21	150.33	4.12	152.71	6.50
8'	113.14	117.64	4.50	116.64	3.50
9′	65.42	67.39	1.97	65.94	0.52
10'	174.3	173.87	0.43	172.46	1.84
CMAE			1.66		2.36

 ${}^{a}\Delta\delta_{\rm C} = |\delta_{\rm corr} - \delta_{\rm exptl}|$ 

	15	(2' <i>R</i> ,3' <i>S</i> ,6' <i>R</i> )-15		(2' <i>R</i> ,3' <i>R</i> ,6' <i>S</i> )- <b>15</b>	
No.	$\delta_{ m exptl}$	$\delta_{ m corr}$	$\varDelta {\delta_{ m C}}^a$	$\delta_{ m corr}$	$\varDelta \delta_{ m C}{}^a$
1	166.6	167.35	0.75	166.34	0.26
2	123.17	122.13	1.04	121.89	1.28
3	108.11	108.14	0.03	108.04	0.07
4	153.71	152.44	1.27	152.22	1.49
5	127.86	127.78	0.08	127.70	0.16
6	114.3	113.69	0.61	113.79	0.51
1'	204.62	202.69	1.93	204.54	0.08
2'	96.16	100.96	4.80	97.74	1.58
3'	56.07	53.95	2.12	55.35	0.72
4′	27.98	26.68	1.30	28.12	0.14
5'	29.97	31.00	1.03	31.22	1.25
6'	47.27	47.47	0.20	46.81	0.46
7'	147.91	147.91	0	147.91	0
8'	138.09	138.09	0	138.09	0
9'	195.17	197.77	2.60	197.19	2.02
10'	174.32	173.28	1.04	174.37	0.05
CMAE			1.18		0.63

**Table S3.** The <sup>13</sup>C NMR experimental values and calculated chemical shifts of (2'R,3'S,6'R)-15 and (2'R,3'R,6'S)-15 are compared.

 ${}^{a}\Delta\delta_{\rm C} = |\delta_{\rm corr} - \delta_{\rm exptl}|.$ 

**Table S4**. The <sup>13</sup>C NMR experimental data ( $\delta$  in ppm) and calculated chemical shifts of (2'*R*,3'*S*,6'*R*)-**20** and (2'*R*,3'*R*,6'*S*)-**20** are compared.

	20	(2' <i>R</i> ,3' <i>S</i> ,6' <i>R</i> )- <b>20</b>		(2' <i>R</i> ,3' <i>R</i> ,6' <i>S</i> )- <b>20</b>	
No.	$\delta_{ m exptl}$	$\delta_{ m corr}$	$\varDelta {\delta_{ m C}}^a$	$\delta_{ m corr}$	$\varDelta {\delta_{ m C}}^a$
1	167.99	167.54	0.45	167.46	0.53
2	123.81	122.79	1.02	121.91	1.90
3	107.75	107.59	0.16	107.63	0.12
4	153.8	151.76	2.04	152.46	1.34
5	128.21	127.61	0.60	127.82	0.39
6	114.67	113.65	1.02	113.53	1.14
1′	203.47	202.95	0.52	203.29	0.18
2'	97.48	99.70	2.22	98.74	1.26
3'	49.96	50.82	0.86	50.24	0.28
4′	22.24	22.62	0.38	23.28	1.04
5'	32.1	29.74	2.36	32.16	0.06
6'	79.19	80.17	0.98	78.77	0.42
7'	171.77	174.85	3.08	175.58	3.81
OCH <sub>3</sub>	52	52.65	0.65	51.58	0.42
CMAE			1.17		0.92

 ${}^{a}\Delta\delta_{\rm C} = |\delta_{\rm corr} - \delta_{\rm exptl}|$ 

## 2. Supplementary Figures



Figure S1. <sup>1</sup>H NMR spectrum of **1** in methanol- $d_4$ 











Figure S4. HMBC spectrum of 1 in methanol- $d_4$ 



Figure S5. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **1** in methanol- $d_4$ 



Figure S6. ROESY spectrum of 1 in methanol- $d_4$ 



Figure S7. HRESIMS spectrum of 1



Figure S8. CD spectrum of 1 in methanol













Figure S12. HMBC spectrum of 2 in methanol- $d_4$ 







Figure S14. ROESY spectrum of 2 in methanol- $d_4$ 







Figure S16. CD spectrum of 2 in methanol















Figure S20. HMBC spectrum of  $\mathbf{3}$  in methanol- $d_4$ 



Figure S21. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **3** in methanol- $d_4$ 



Figure S22. ROESY spectrum of **3** in methanol- $d_4$ 



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Figure S23. HRESIMS spectrum of 3



Figure S24. CD spectrum of 3 in methanol











Figure S27. HSQC spectrum of 4 in methanol- $d_4$ 



Figure S28. HMBC spectrum of **4** in methanol- $d_4$ 



Figure S29. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **4** in methanol- $d_4$ 



Figure S30. HRESIMS spectrum of 4



Figure S31. CD spectrum of 4 in methanol



Figure S32. <sup>1</sup>H NMR spectrum of **5** in methanol- $d_4$ 



Figure S33. <sup>13</sup>C NMR and DEPT spectra of **5** in methanol- $d_4$ 



Figure S34. HSQC spectrum of **5** in methanol- $d_4$ 







Figure S36. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **5** in methanol- $d_4$ 







Figure S38. CD spectrum of 5 in methanol















Figure S42. HMBC spectrum of **6** in methanol- $d_4$ 



Figure S43. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **6** in methanol- $d_4$ 



Figure S44. HRESIMS spectrum of 6



Figure S45. CD spectrum of 6 in methanol



Figure S46. <sup>1</sup>H NMR spectrum of **7** in methanol-*d*<sub>4</sub>











Figure S50. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **7** in methanol- $d_4$ 



Figure S51. HRESIMS spectrum of 7



Figure S52. CD spectrum of 7 in methanol



Figure S53. <sup>1</sup>H NMR spectrum of **8** in methanol- $d_4$ 



Figure S54. <sup>13</sup>C NMR spectrum of **8** in methanol- $d_4$






Figure S56. HMBC spectrum of **8** in methanol- $d_4$ 



Figure S57. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **8** in methanol- $d_4$ 



Figure S58. HRESIMS spectrum of 8



















Figure S64. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **9** in methanol- $d_4$ 







Figure S66. CD spectrum of 9 in methanol

















Figure S72. ROESY spectrum of 10 in methanol- $d_4$ 



Enlarged ROESY spectrum of 10 (up-field region) in methanol- $d_4$ 



Figure S73. HRESIMS spectrum of 10







Figure S75. <sup>1</sup>H NMR spectrum of **11** in methanol- $d_4$ 



Figure S77. HSQC spectrum of 11 in methanol- $d_4$ 



Figure S78. HMBC spectrum of 11 in methanol- $d_4$ 



Figure S79. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **11** in methanol- $d_4$ 



Figure S80. ROESY spectrum of 11 in methanol- $d_4$ 



Figure S81. HRESIMS spectrum of 11



Figure S82. CD spectrum of 11 in methanol



Figure S83. <sup>1</sup>H NMR spectrum of **12** in methanol- $d_4$ 











Figure S87. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **12** in methanol- $d_4$ 



Figure S88. ROESY spectrum of 12 in methanol- $d_4$ 



Figure S89. HREIMS spectrum of 12



Figure S90. CD spectrum of 12 in methanol



Figure S91. <sup>1</sup>H NMR spectrum of **14** in methanol- $d_4$ 



Figure S92. <sup>13</sup>C NMR and DEPT spectra of **14** in methanol- $d_4$ 



Figure S93. HSQC spectrum of 14 in methanol- $d_4$ 



Figure S95. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **14** in methanol- $d_4$ 







Figure S97. HRESIMS spectrum of 14



Figure S98. CD spectrum of 14 in methanol



Figure S99. <sup>1</sup>H NMR spectrum of **15** in methanol- $d_4$ 



Figure S101. HSQC spectrum of 15 in methanol- $d_4$ 



Figure S103. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **15** in methanol- $d_4$ 







Figure S105. HRESIMS spectrum of 15



Figure S106. CD spectrum of 15a in methanol



Figure S107. CD spectrum of **15b** in methanol











Figure S110. HSQC spectrum of 16 in methanol- $d_4$ 



Figure S111. HMBC spectrum of 16 in methanol- $d_4$ 







Figure S113. HRESIMS spectrum of 16















Figure S118. HMBC spectrum of 18 in methanol- $d_4$ 



Figure S119. <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **18** in methanol- $d_4$ 



Figure S120. ROESY spectrum of 18 in methanol- $d_4$ 



Enlarged ROESY spectrum of 18 (up-field region) in methanol- $d_4$ 



Figure S122. CD spectrum of 18 in methanol














Figure S126. HMBC spectrum of 20 in methanol- $d_4$ 







Figure S128. ROESY spectrum of **20** in methanol- $d_4$ 







Figure S130. CD spectrum of 20 in methanol