

Supporting Information

New 3-Acyl Tetramic acid Derivatives from the Deep-Sea-Derived Fungus *Lecanicillium fuisporum*

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CAACCCTTATGTGACATACCAATCGTTGCTTCGGCGGACTCGCCCCGGTGTCCGGCAGGCCCTCGCGG
 CCGGCCGCGACCCGGATCCAGGCGGACGCCGGAGACCATCCAAAACTCTTTGTATTCTAGCAAGTCT
 TCTGAATGAGCCGCAAGGCAACACAAATGAATCAAACTTTCAACAACGGATCTCTTGGTTCTGGCATC
 GATGACTAGCGCAGCGAAAGGAGATAACTAATGTGAATTGCACACATCCGCGAACTTGAACACACTGG
 AGGCCCATAGCGGTCCCAACCAATCGCCCCGGTTGAGGAGGACGGGCAACGACTTGATTCTGTGGCTA
 CCATGCAGGCACCCTTTTCGGCCGTGAAGCTTCACCCACAACCTTGCAATCAAAGACTCGATGGTTCCCG
 GGATTCTGCAATTCGCTCATGATTGACATGACTTCAGCATTATTTAGCTTTGTACTGTAATGAAAAAGG
 AAAGGAACGTTTAATTTATTGGGCTTGAAAAATCCGATCAATACTCCACTCTTTTATTTCCCAAAAAAGAG
 AATACTACTTGTATATAGTTACA

Figure S1. The colonies and the ITS rRNA sequences data of deep-sea-derived fungus
Lecanicillium fusisporum GXIMD00542

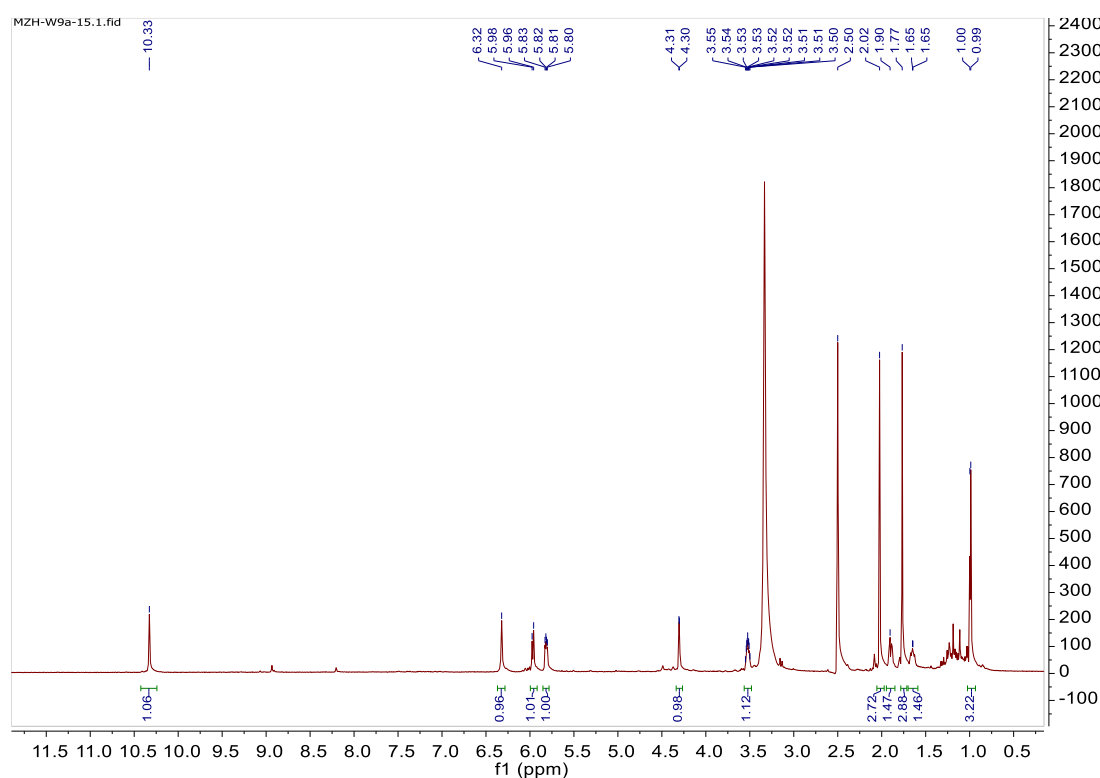


Figure S2. ^1H NMR spectrum of **1**

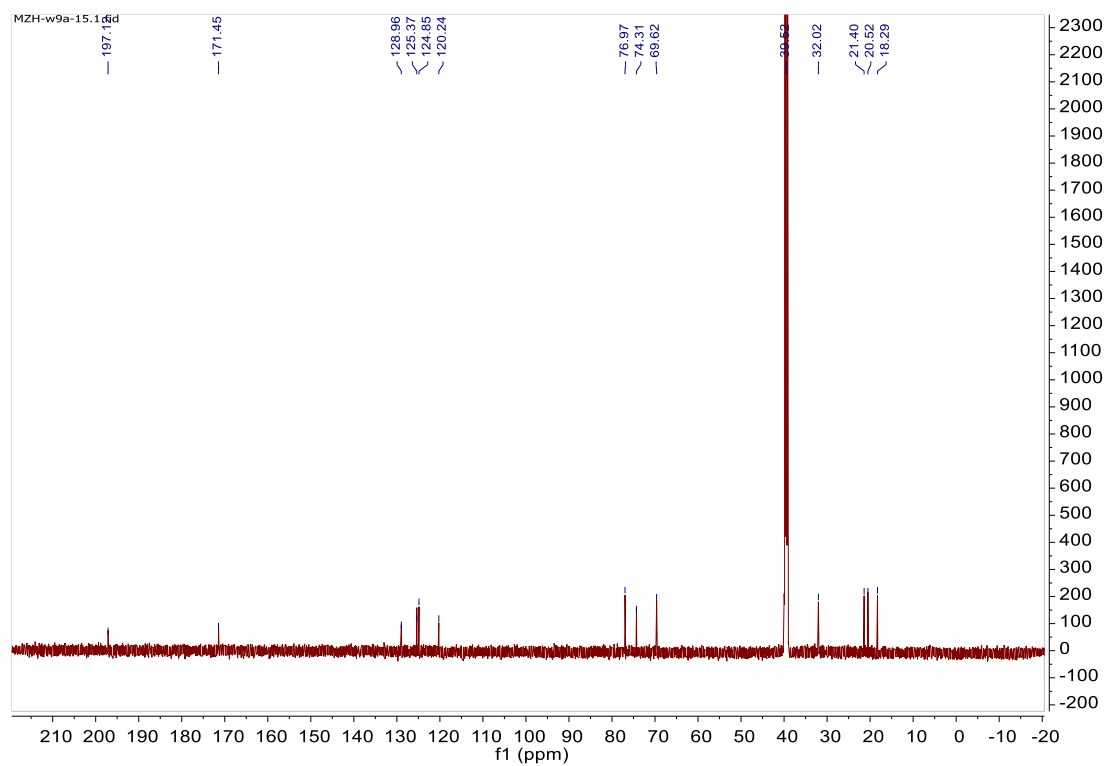


Figure S3. ^{13}C NMR spectrum of **1**

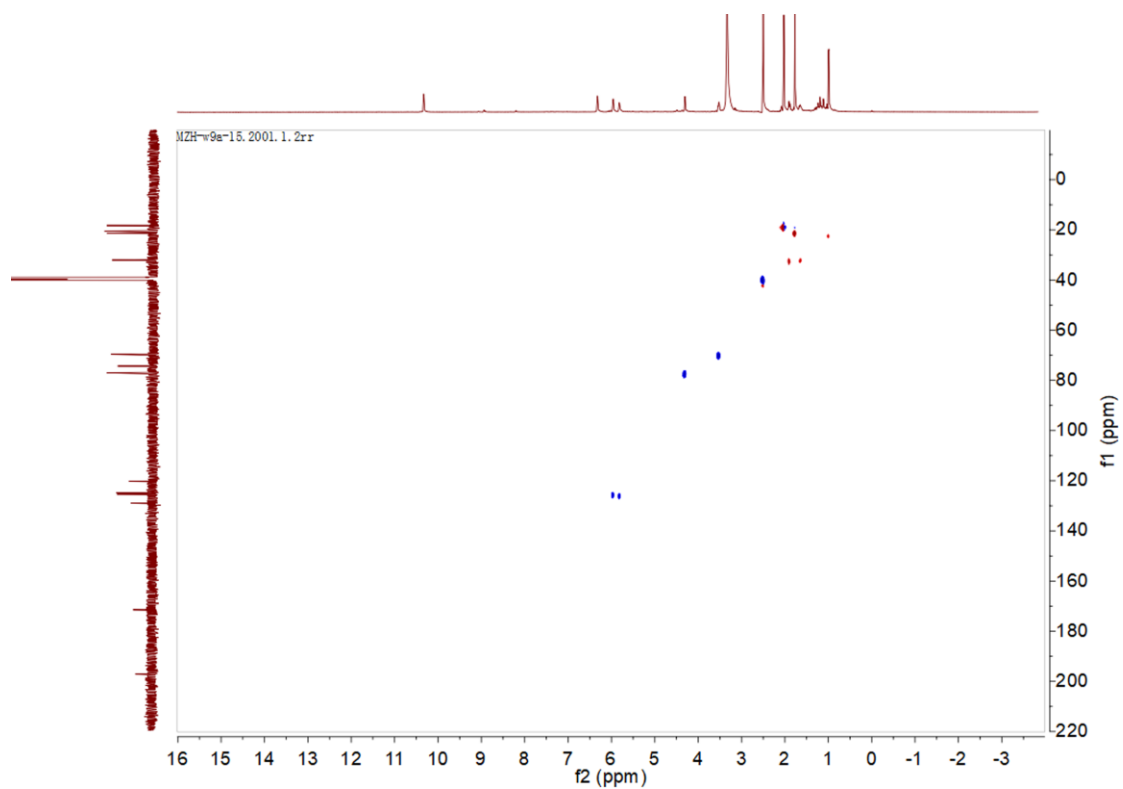


Figure S4. HSQC spectrum of **1**

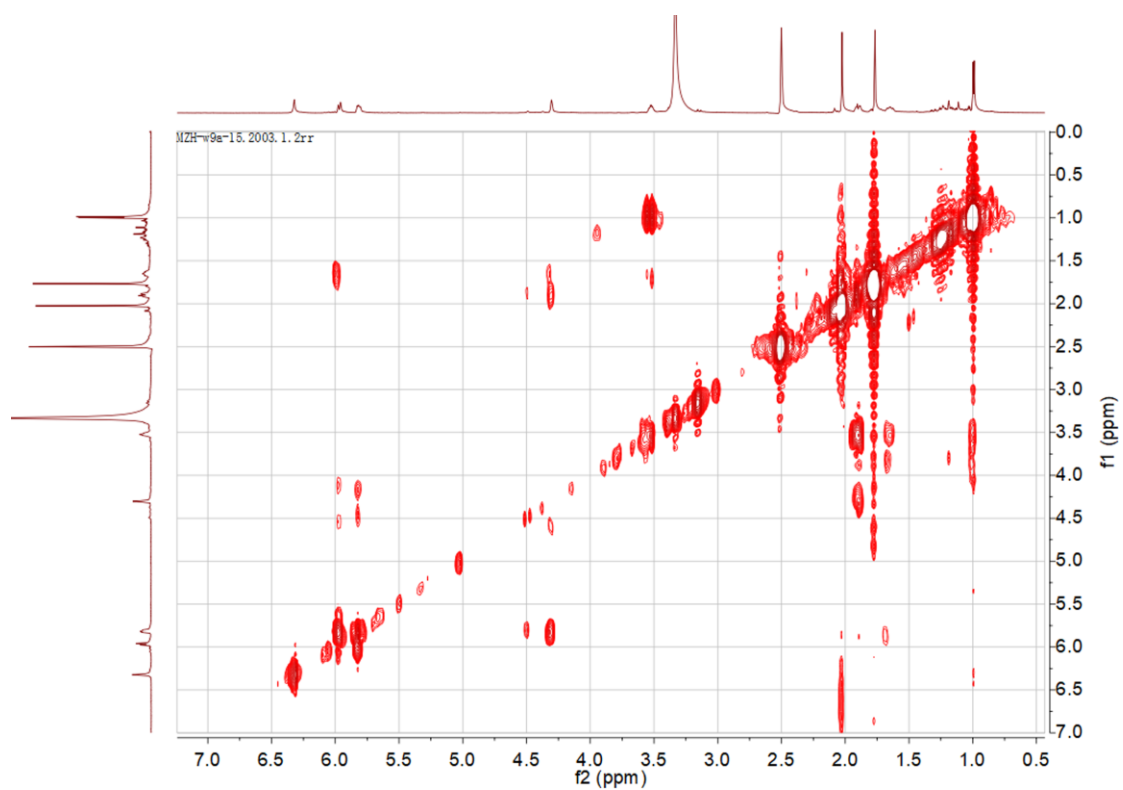


Figure S5. ^1H - ^1H COSY spectrum of **1**

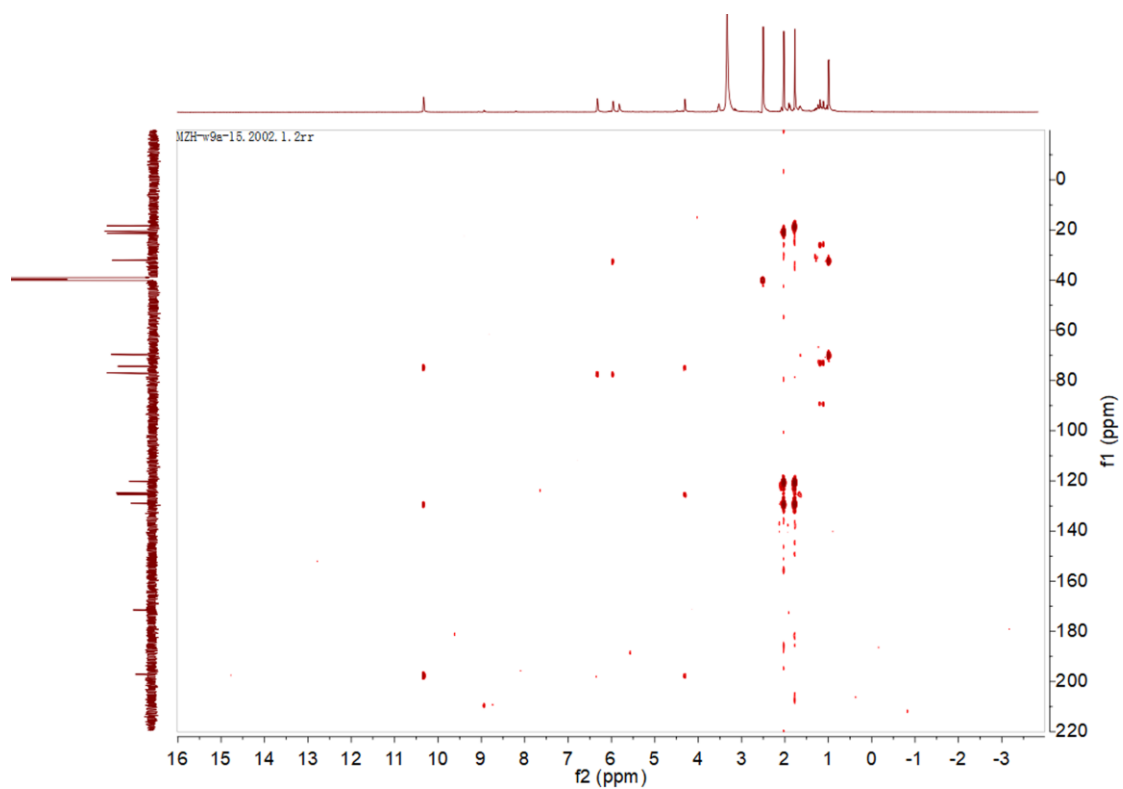


Figure S6. HMBC spectrum of **1**

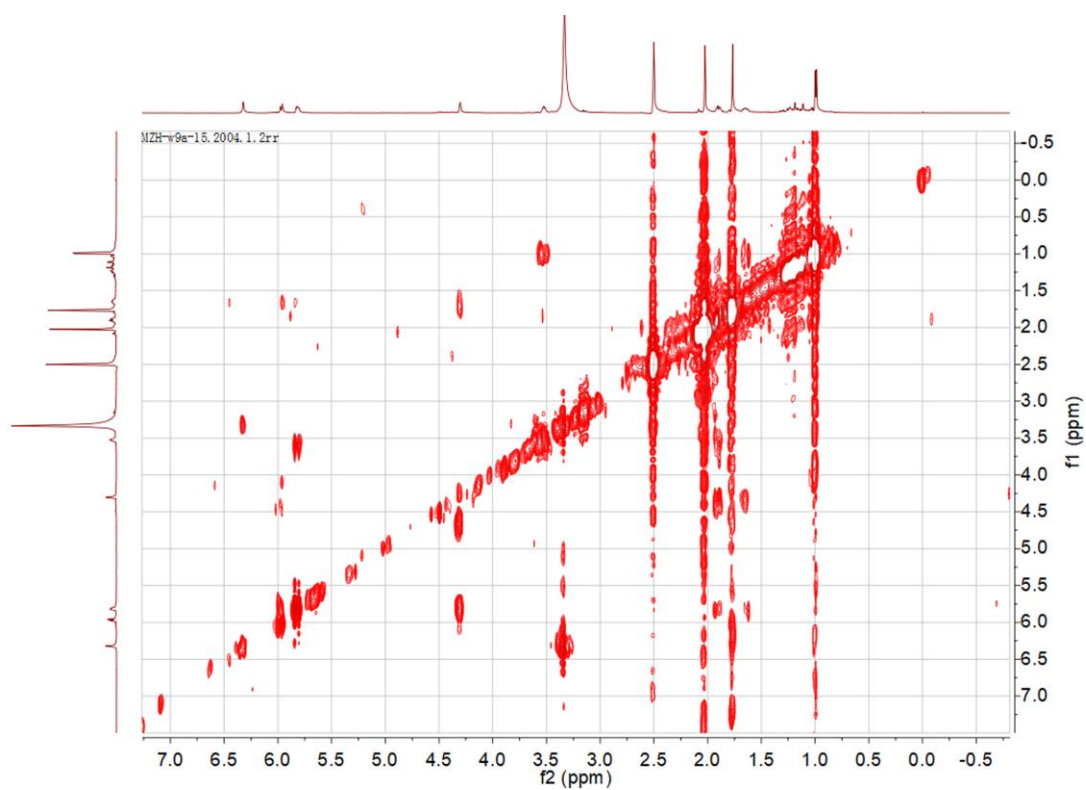


Figure S7. NOESY spectrum of **1**

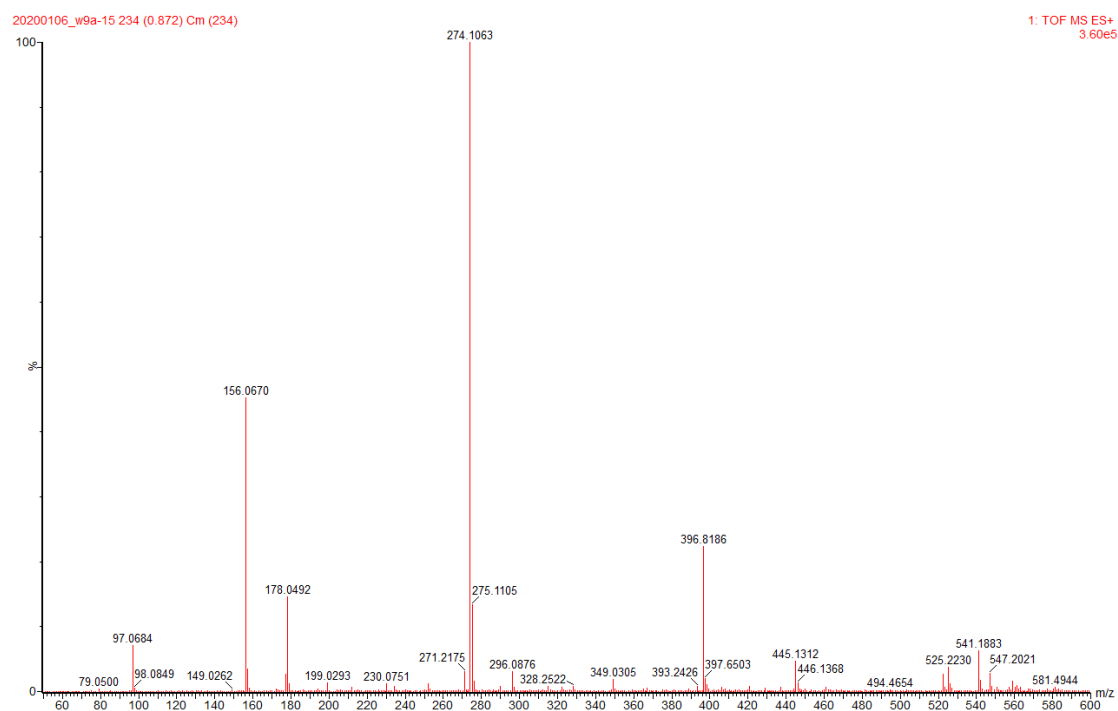


Figure S8. HR-ESI-MS spectrum of **1**

Table S1. Crystal data and structure refinement for compound **1**

Empirical formula	C ₂₆ H ₃₆ N ₂ O ₉
Formula weight	520.57
Temperature/K	100.00(10)
Crystal system	orthorhombic
Space group	P2 ₁ 2 ₁ 2 ₁
a/Å	11.32560(10)
b/Å	12.19970(10)
c/Å	18.5233(2)
α/°	90
β/°	90
γ/°	90
Volume/Å ³	2559.34(4)
Z	4
ρ _{calc} /cm ³	1.351
μ/mm ⁻¹	0.851
F(000)	1112.0
Crystal size/mm ³	0.16 × 0.08 × 0.08
Radiation	Cu Kα (λ = 1.54184)
2θ range for data collection/°	8.678 to 148.214
Index ranges	-13 ≤ h ≤ 12, -13 ≤ k ≤ 14, -22 ≤ l ≤ 22
Reflections collected	14388
Independent reflections	5046 [R _{int} = 0.0182, R _{sigma} = 0.0192]
Data/restraints/parameters	5046/0/345
Goodness-of-fit on F ²	1.045
Final R indexes [I > 2σ (I)]	R ₁ = 0.0266, wR ₂ = 0.0702
Final R indexes [all data]	R ₁ = 0.0270, wR ₂ = 0.0705
Largest diff. peak/hole / e Å ⁻³	0.18/-0.27
Flack parameter	0.03(4)

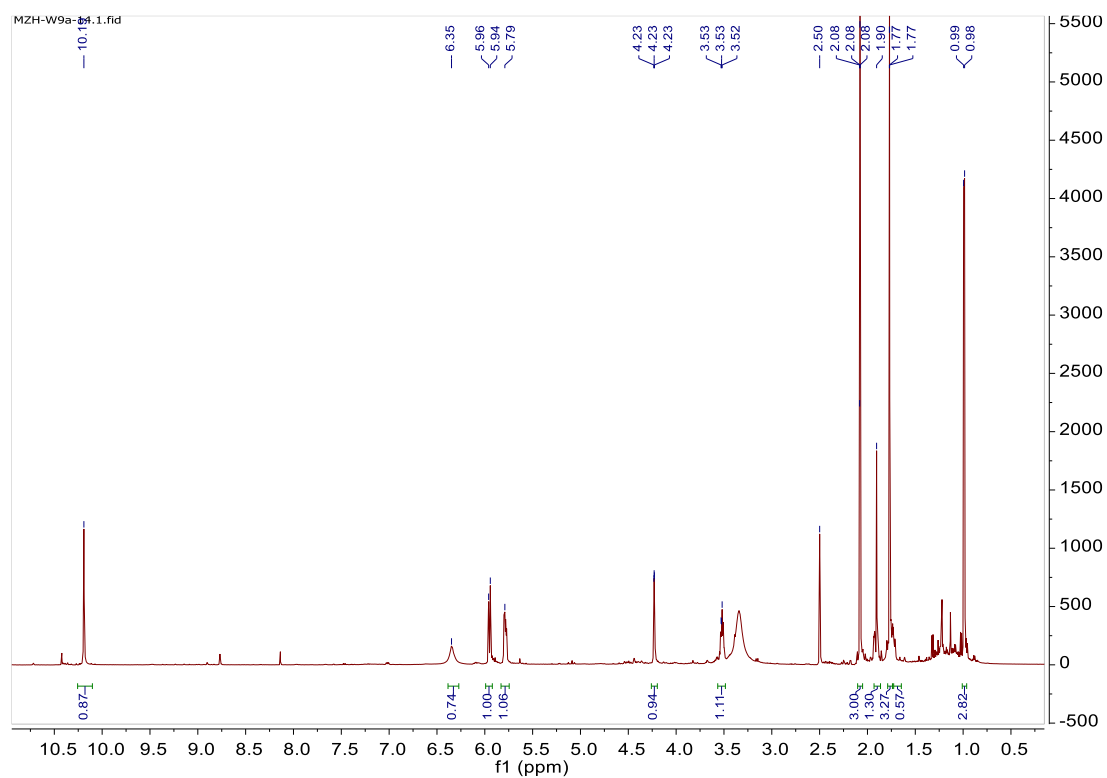


Figure S9. ^1H NMR spectrum of 2

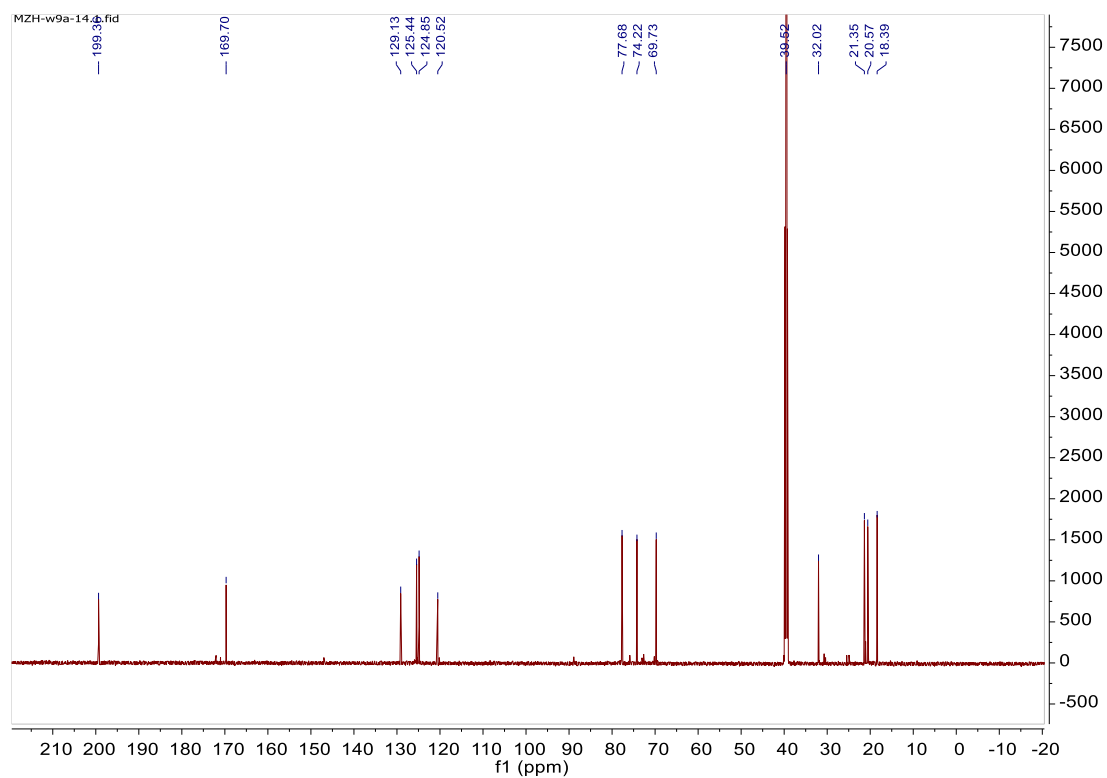


Figure S10. ^{13}C NMR spectrum of 2

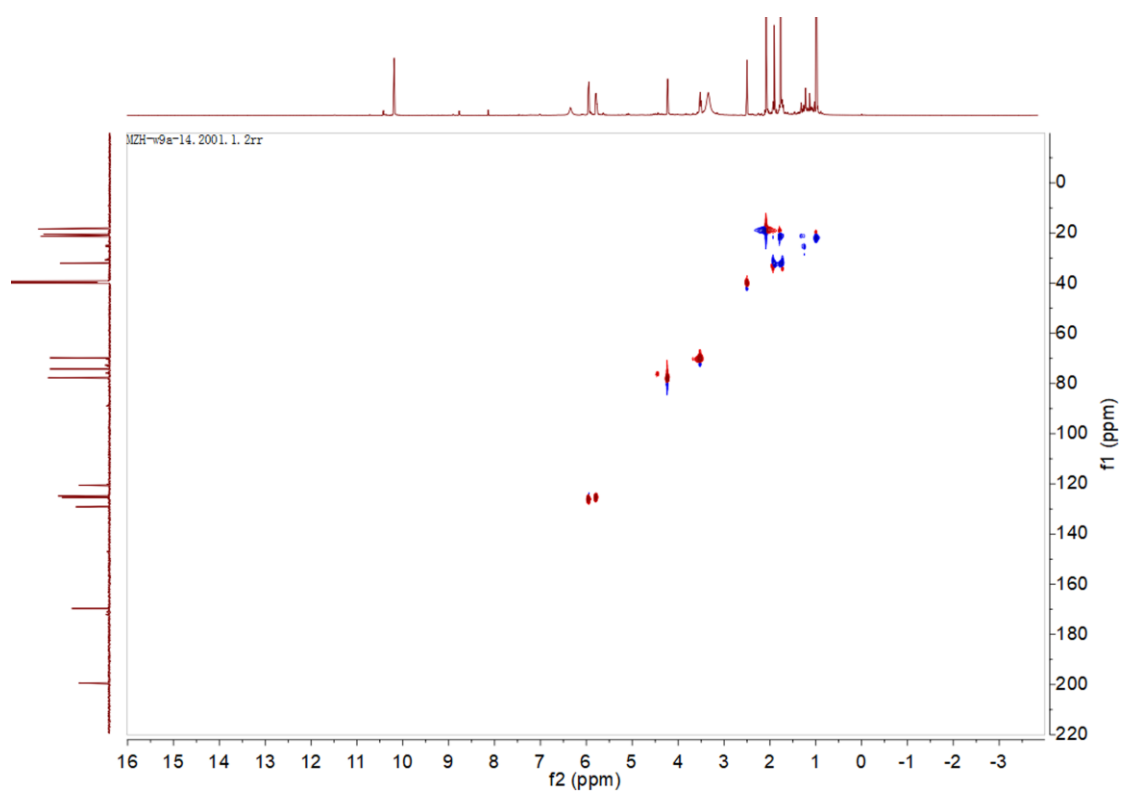


Figure S11. HSQC spectrum of 2

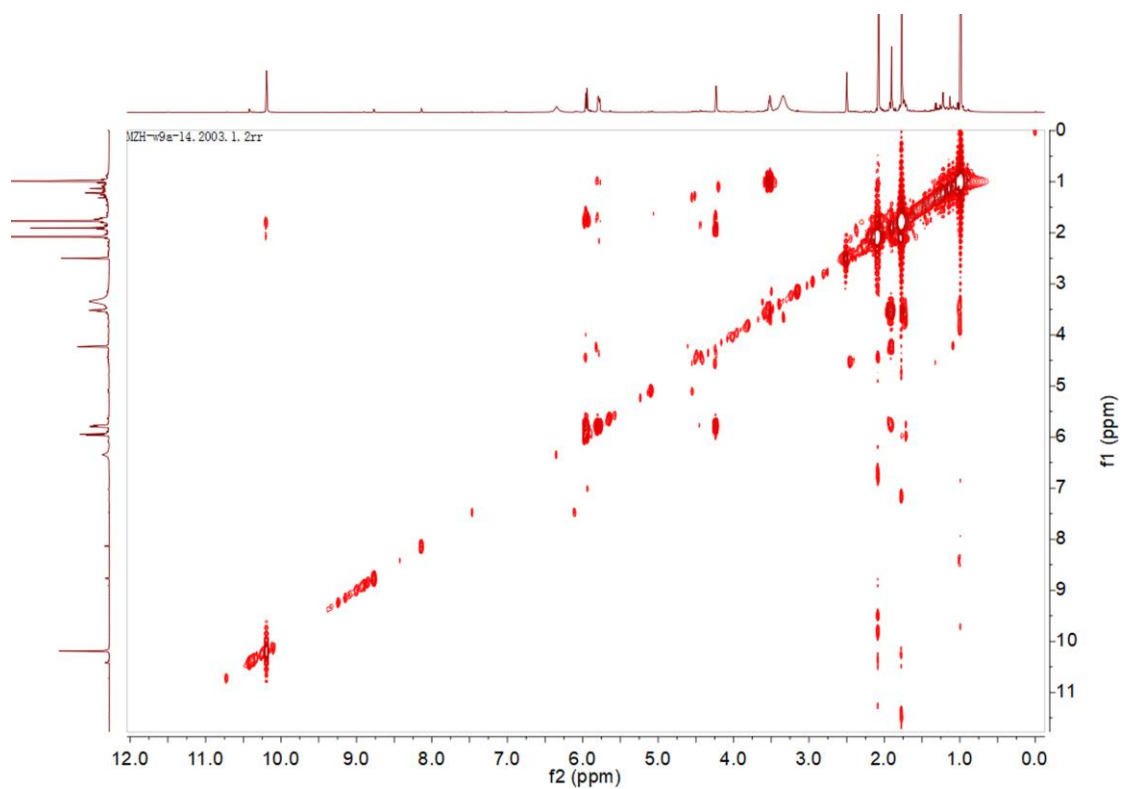


Figure S12. ^1H - ^1H COSY spectrum of 2

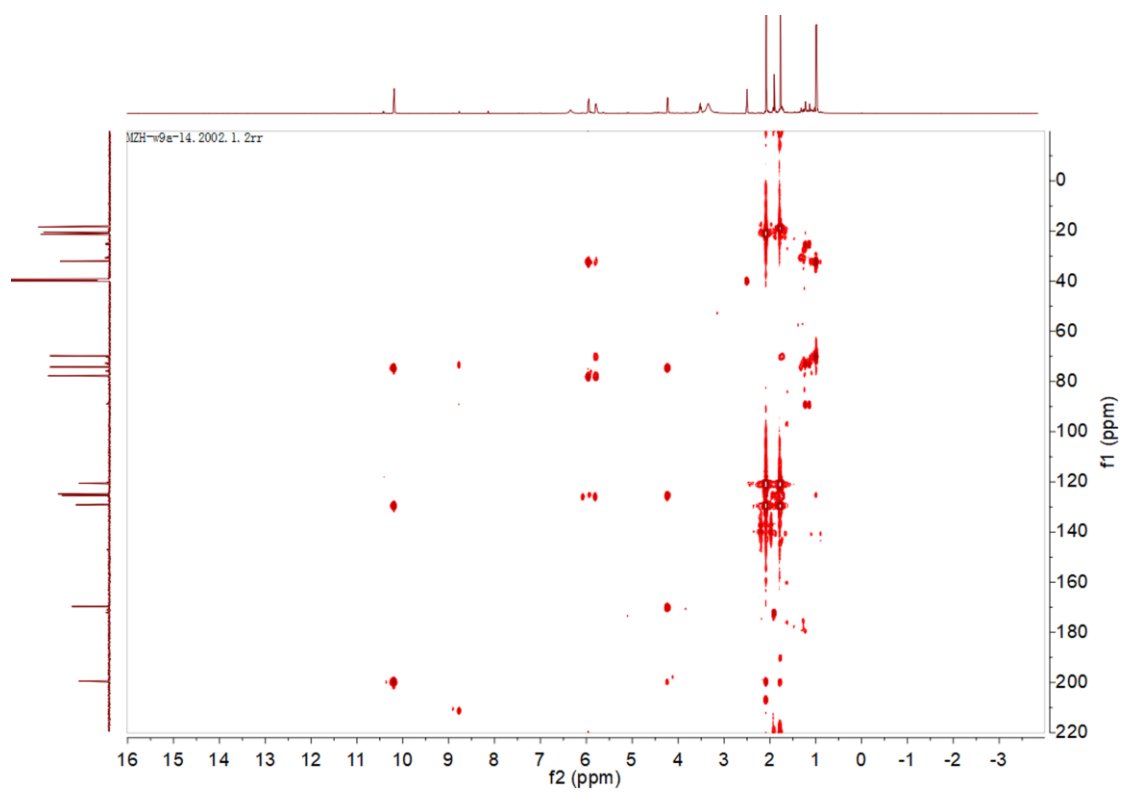


Figure S13. HMBC spectrum of 2

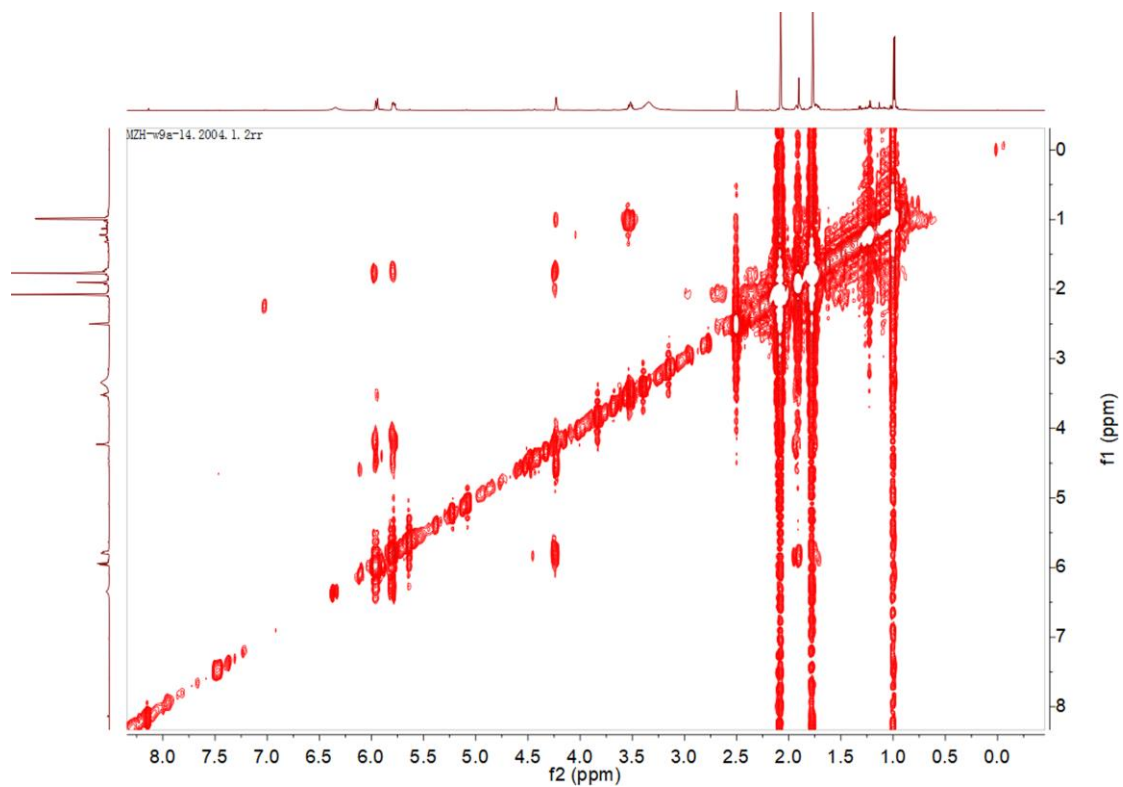


Figure S14. NOESY spectrum of 2

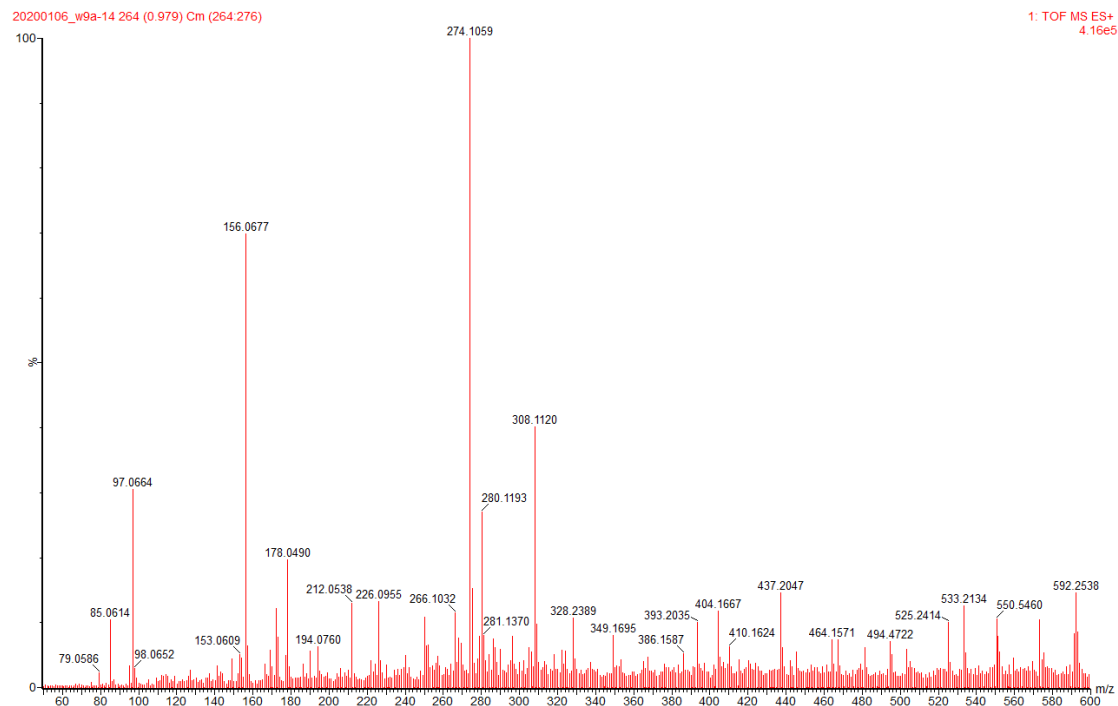


Figure S15. HR-ESI-MS spectrum of **2**

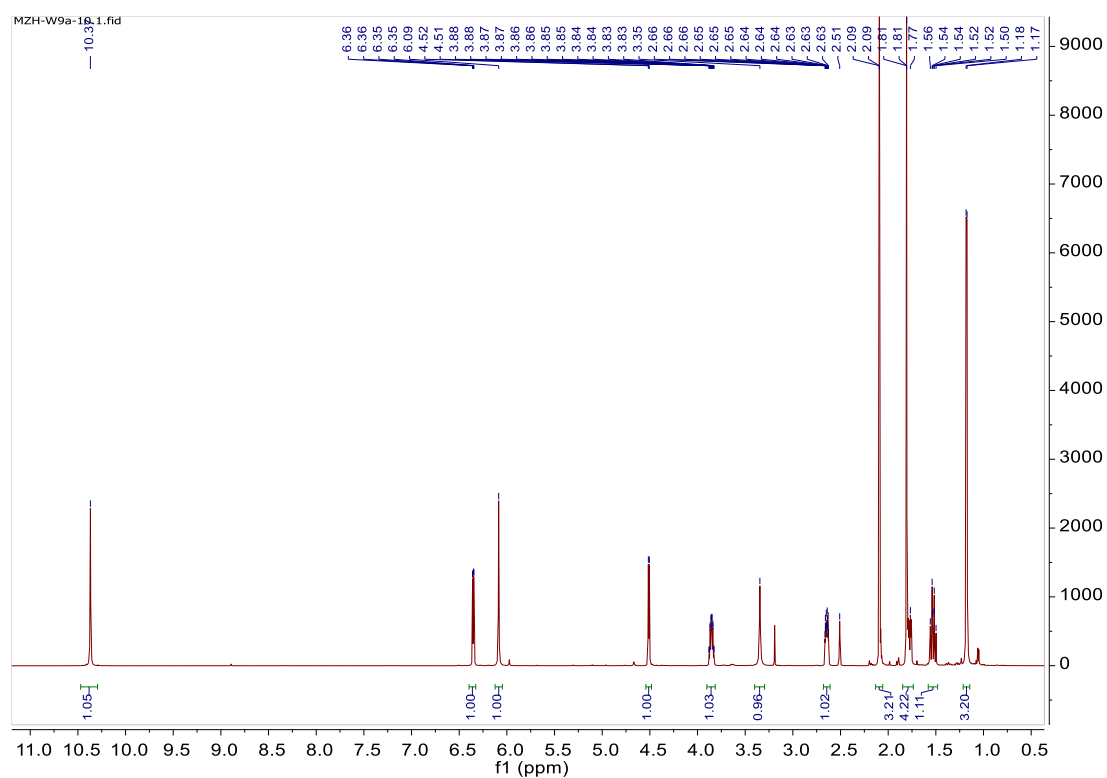


Figure S16. ^1H NMR spectrum of **3**

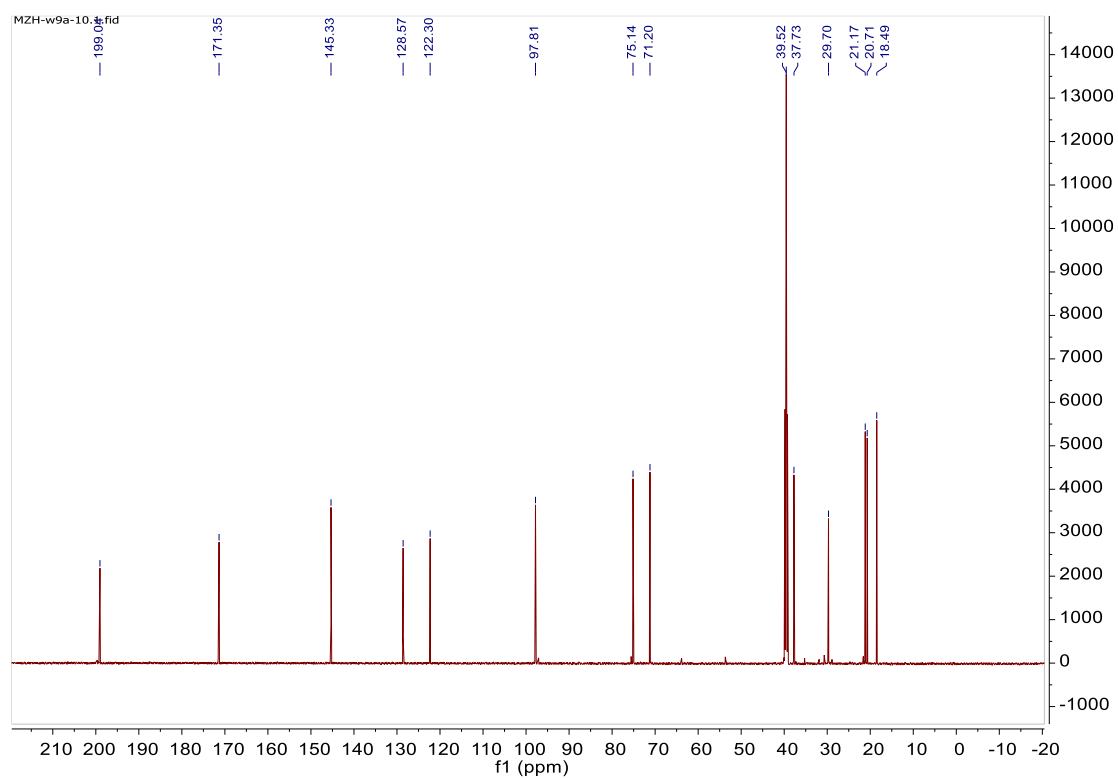


Figure S17. ^{13}C NMR spectrum of **3**

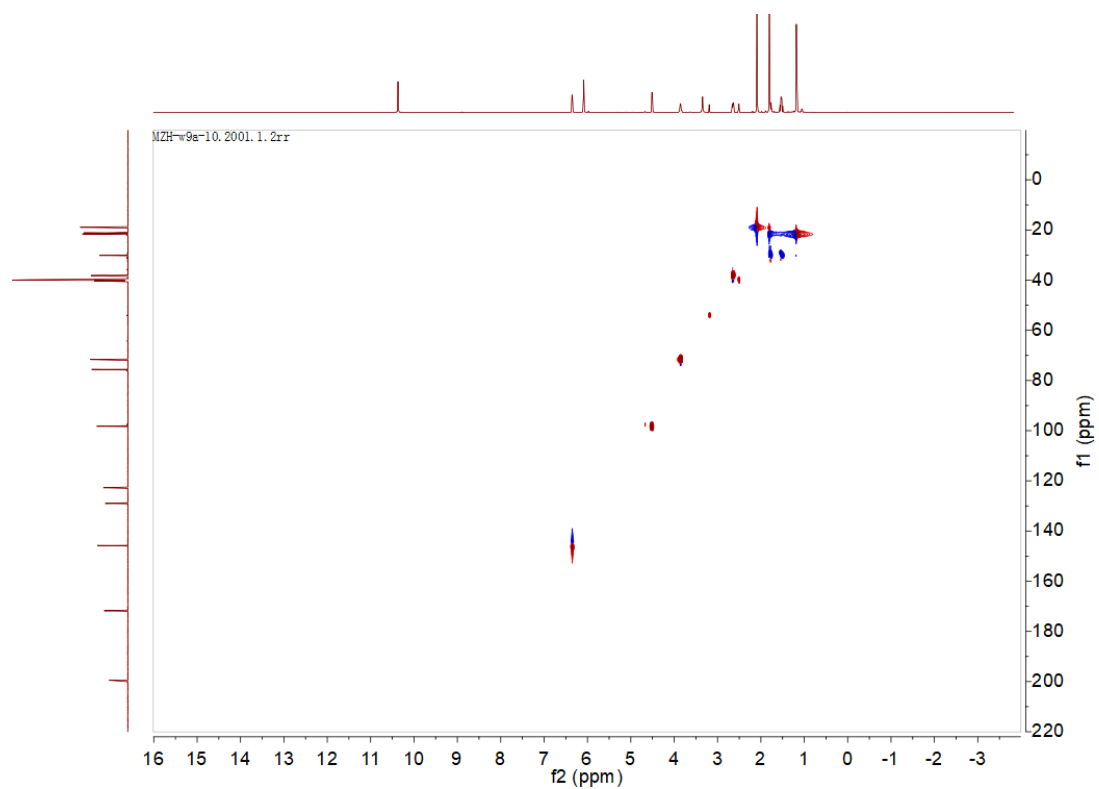


Figure S18. HSQC spectrum of **3**

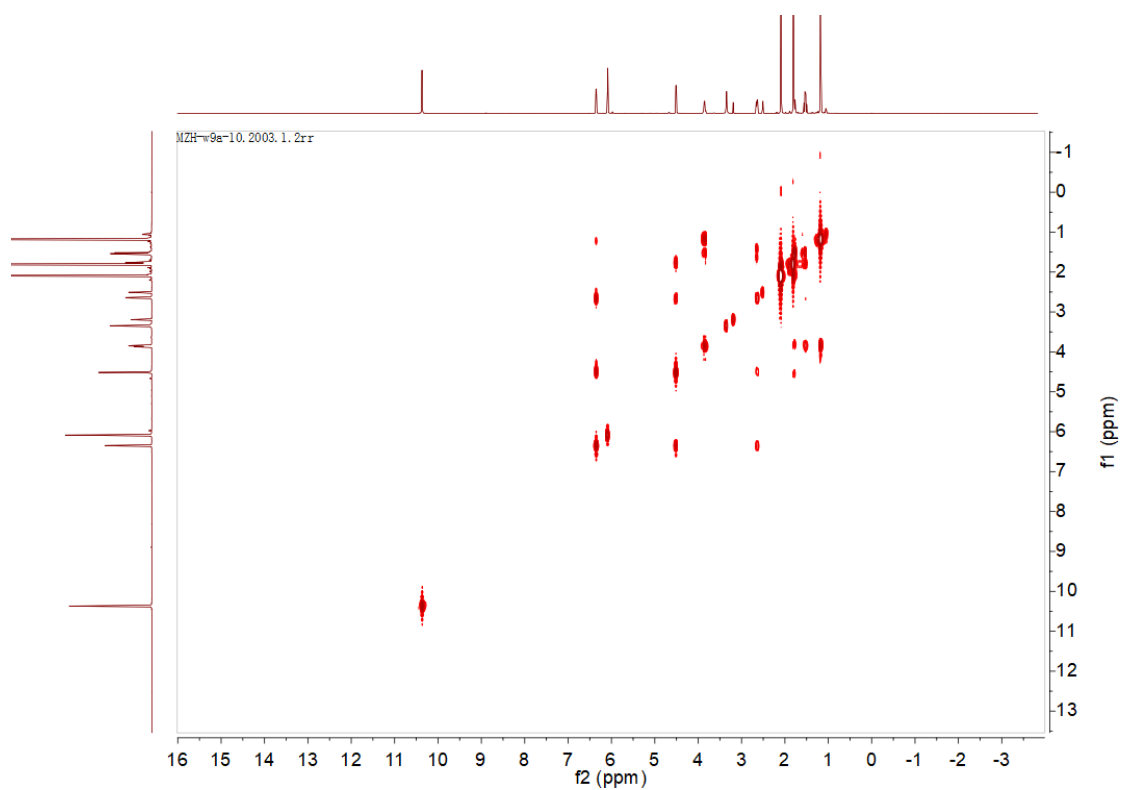


Figure S19. ^1H - ^1H COSY spectrum of **3**

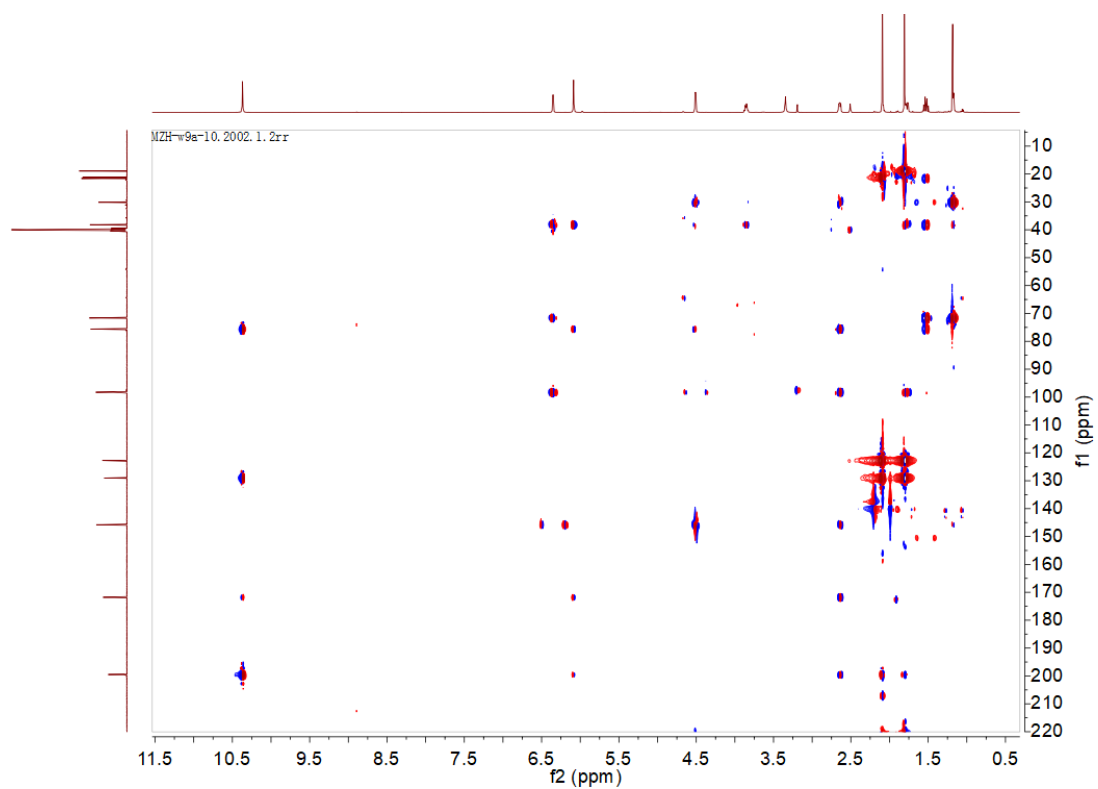


Figure S20. HMBC spectrum of **3**

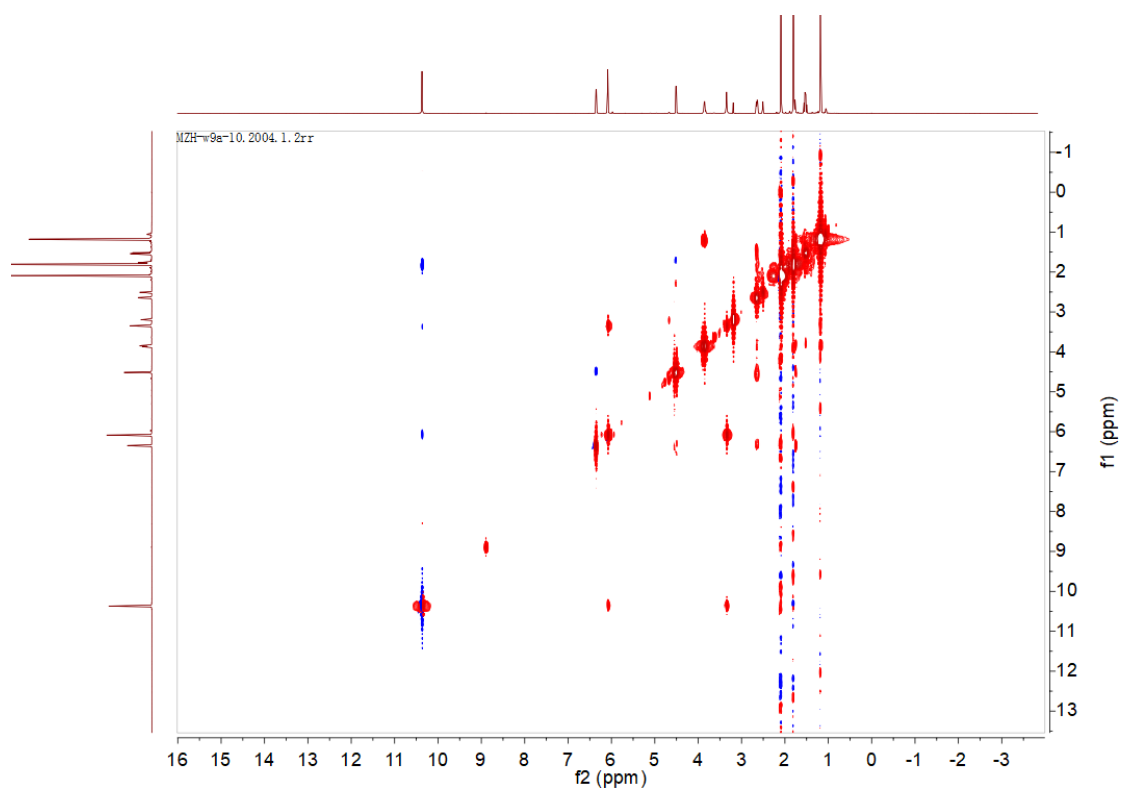


Figure S21. NOESY spectrum of **3**

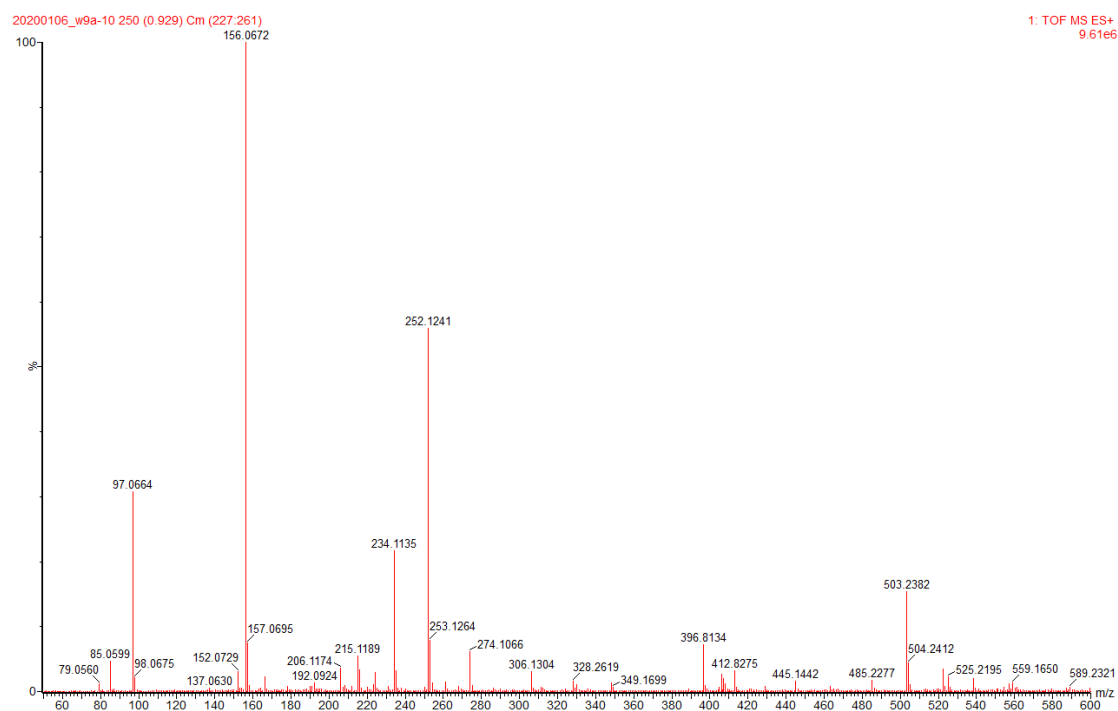


Figure S22. HR-ESI-MS spectrum of **3**

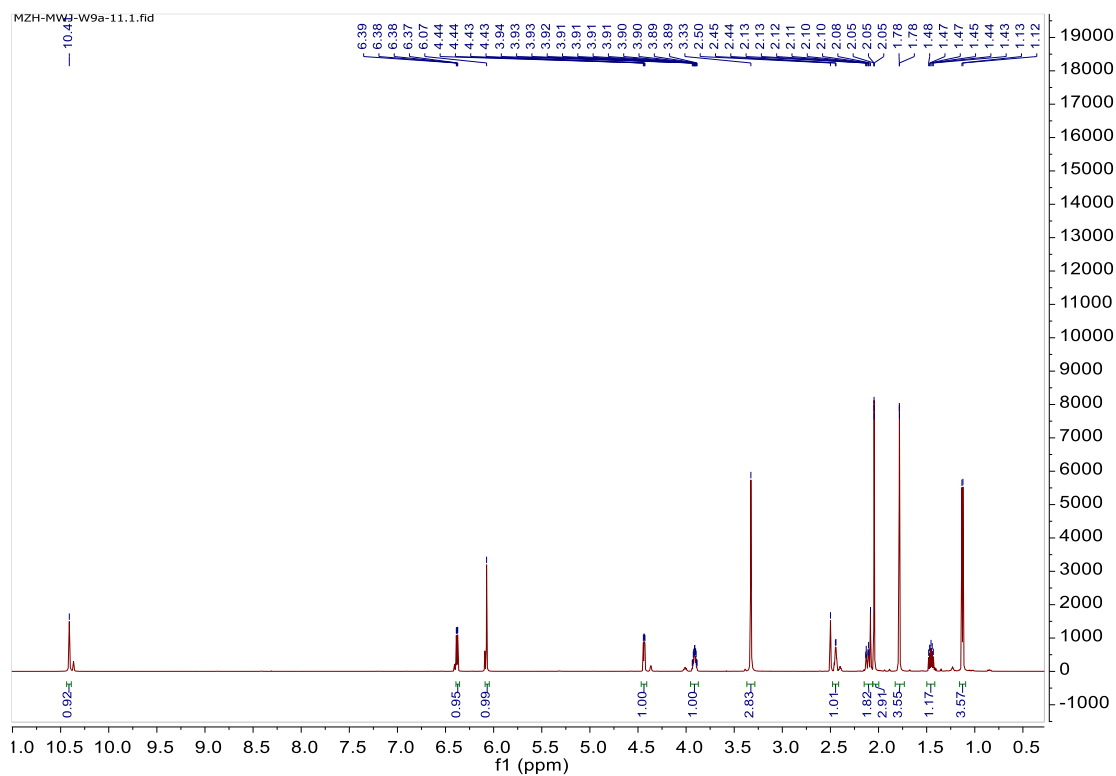


Figure S23. ^1H NMR spectrum of **4**

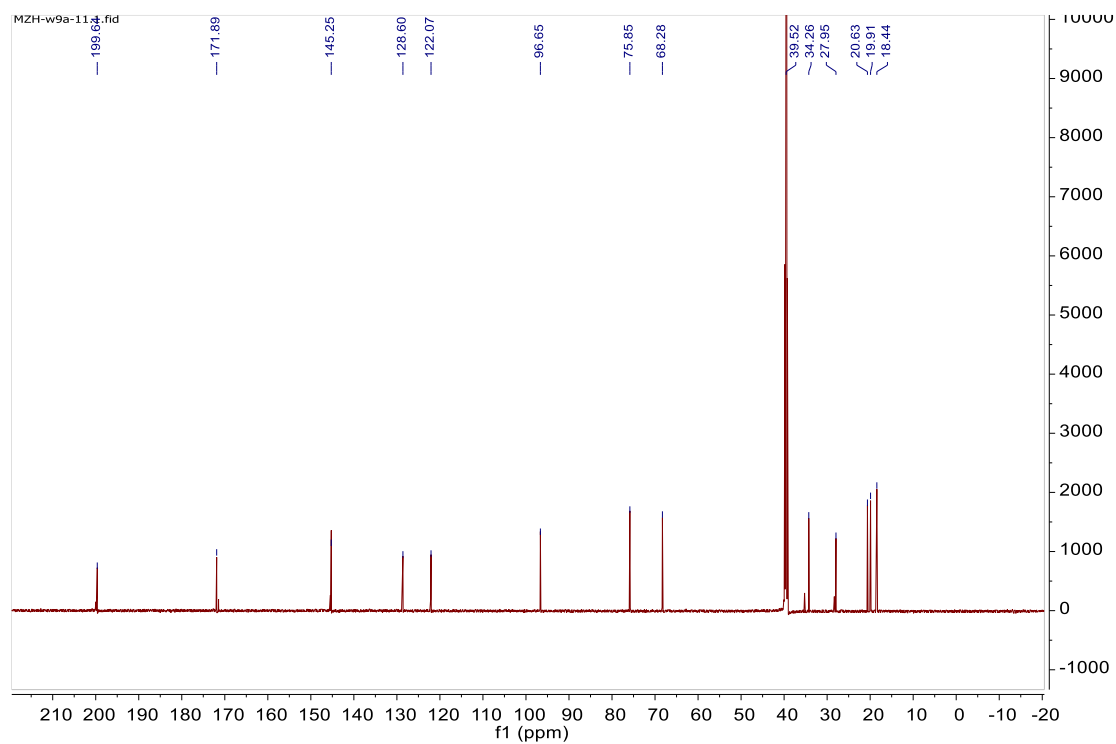


Figure S24. ^{13}C NMR spectrum of **4**

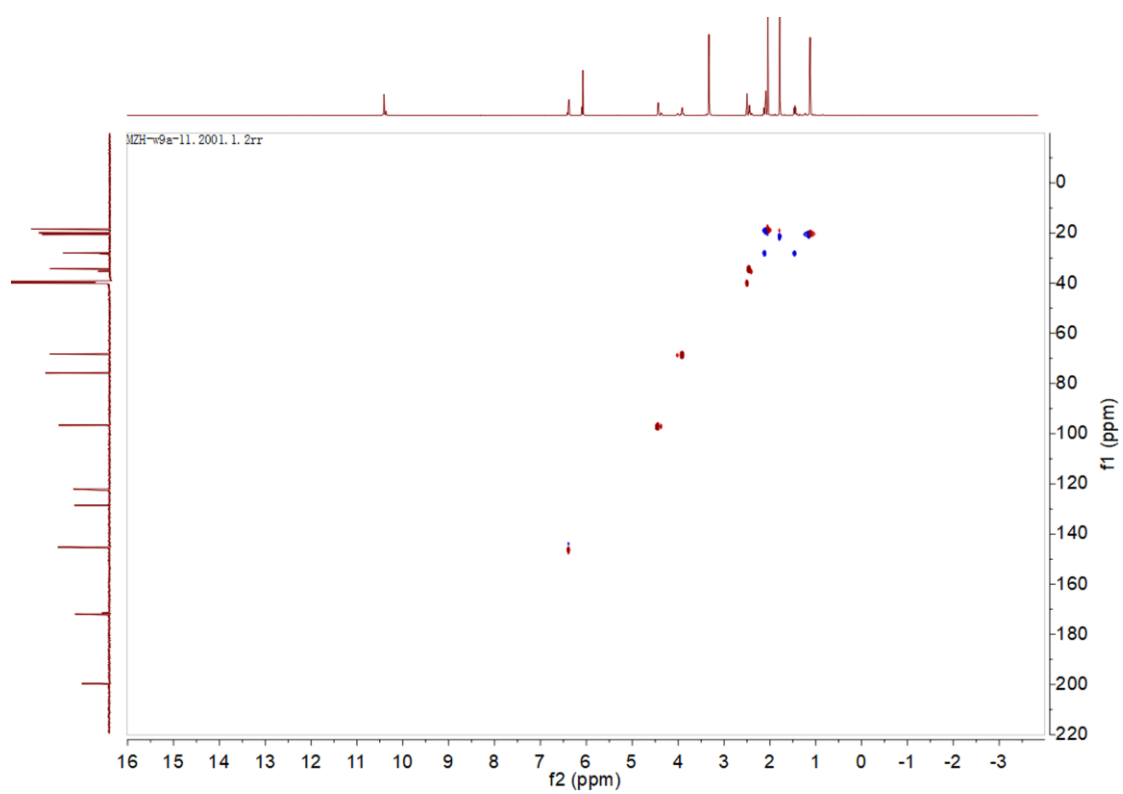


Figure S25. HSQC spectrum of 4

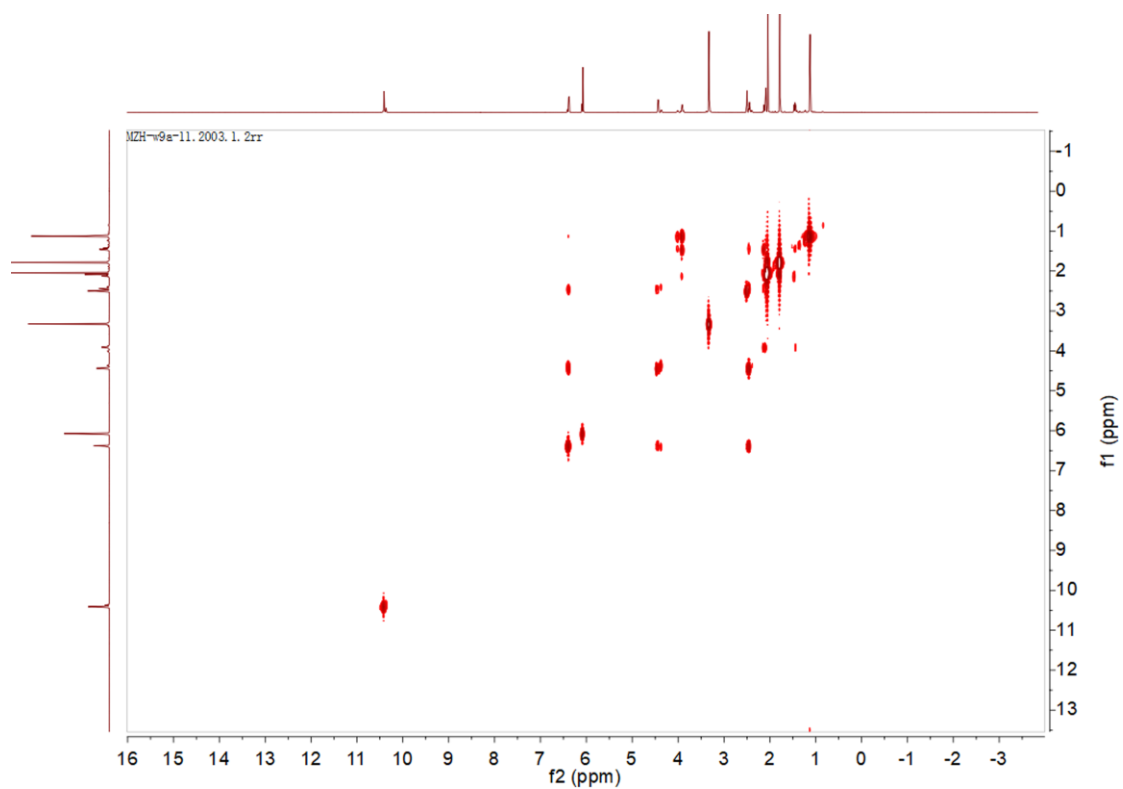


Figure S26. ^1H - ^1H COSY spectrum of 4

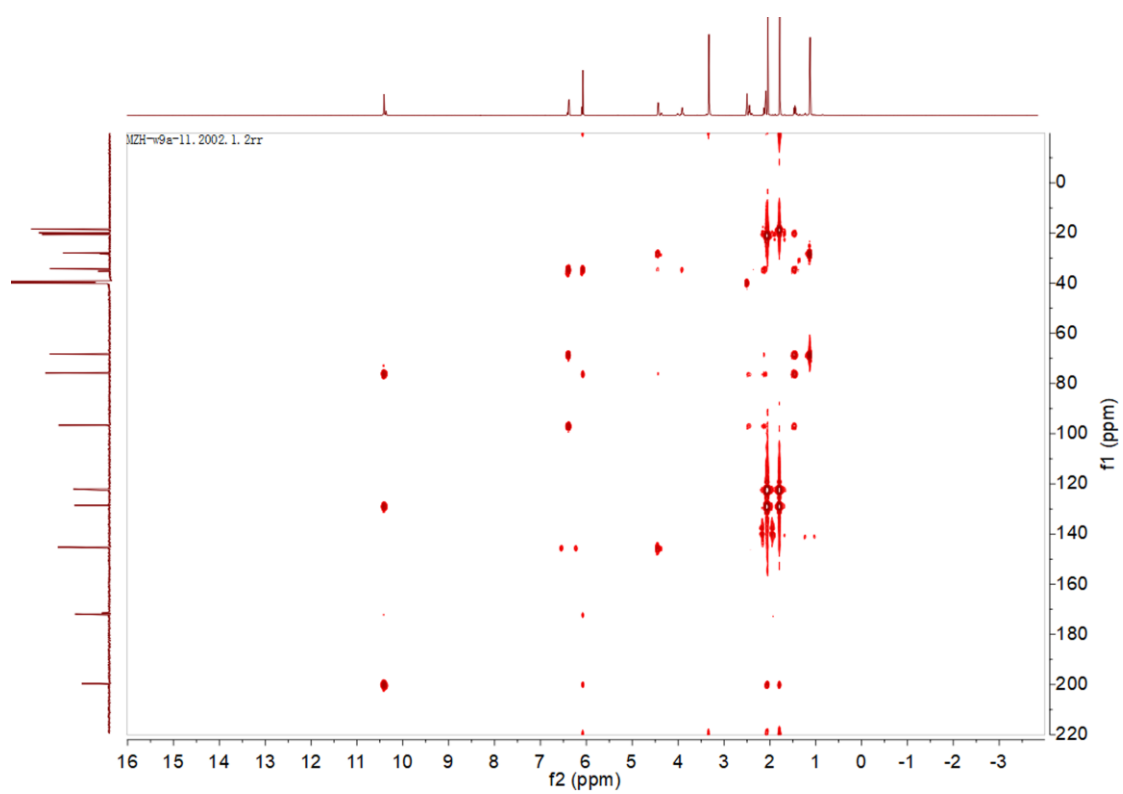


Figure S27. HMBC spectrum of **4**

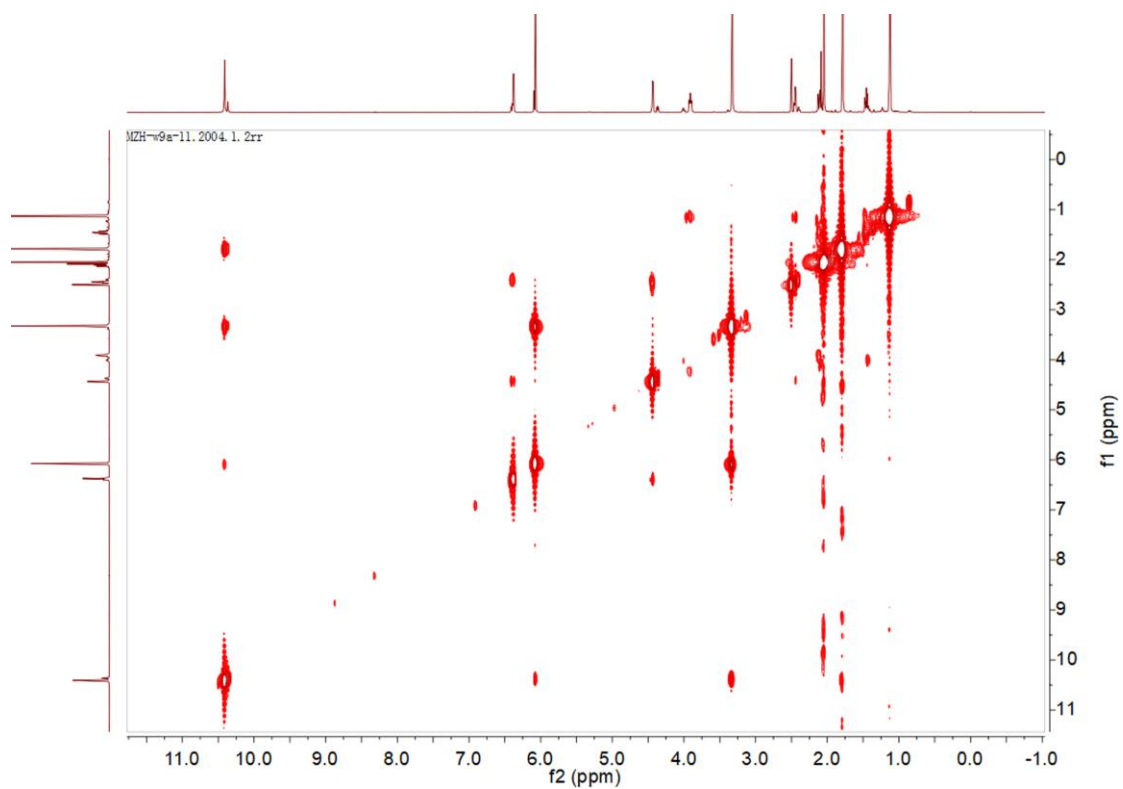


Figure S28. NOESY spectrum of **4**

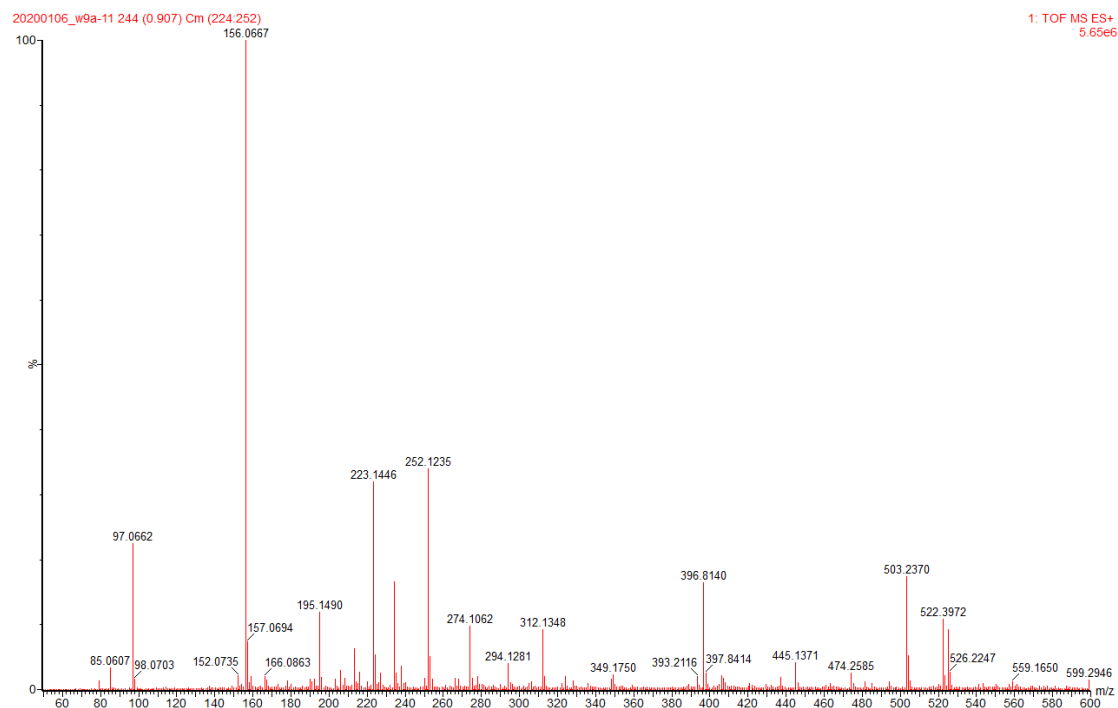


Figure S29. HR-ESI-MS spectrum of 4

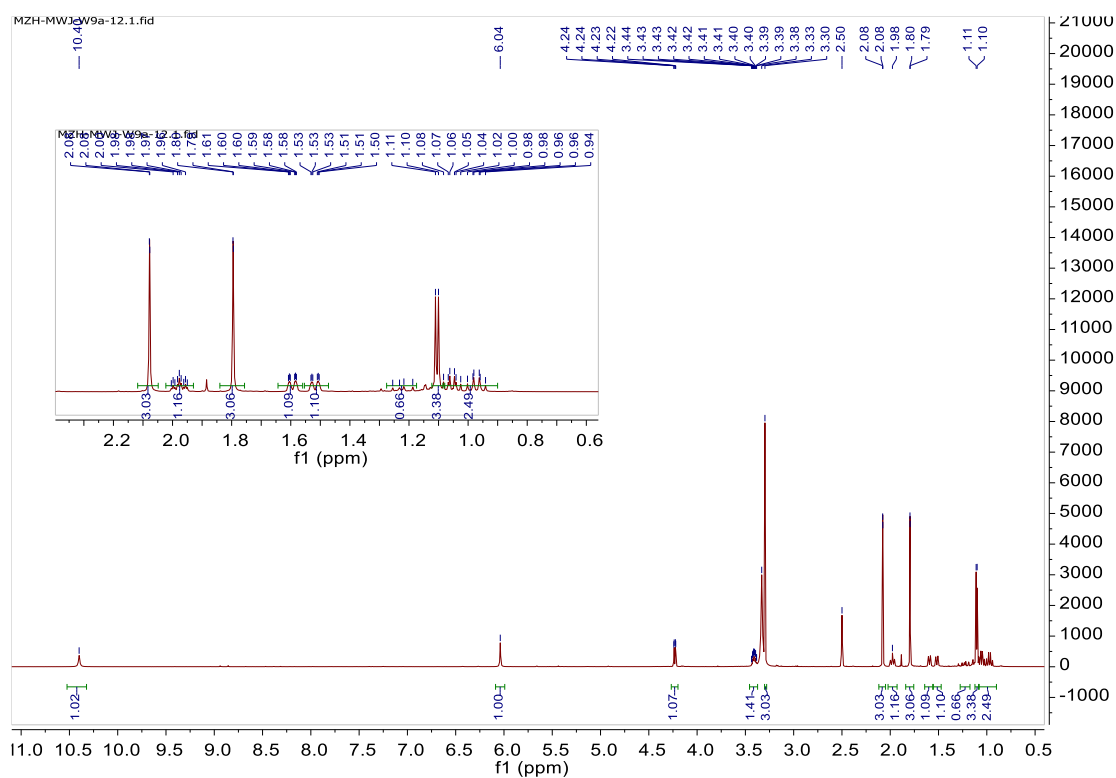


Figure S30. ¹H NMR spectrum of 5

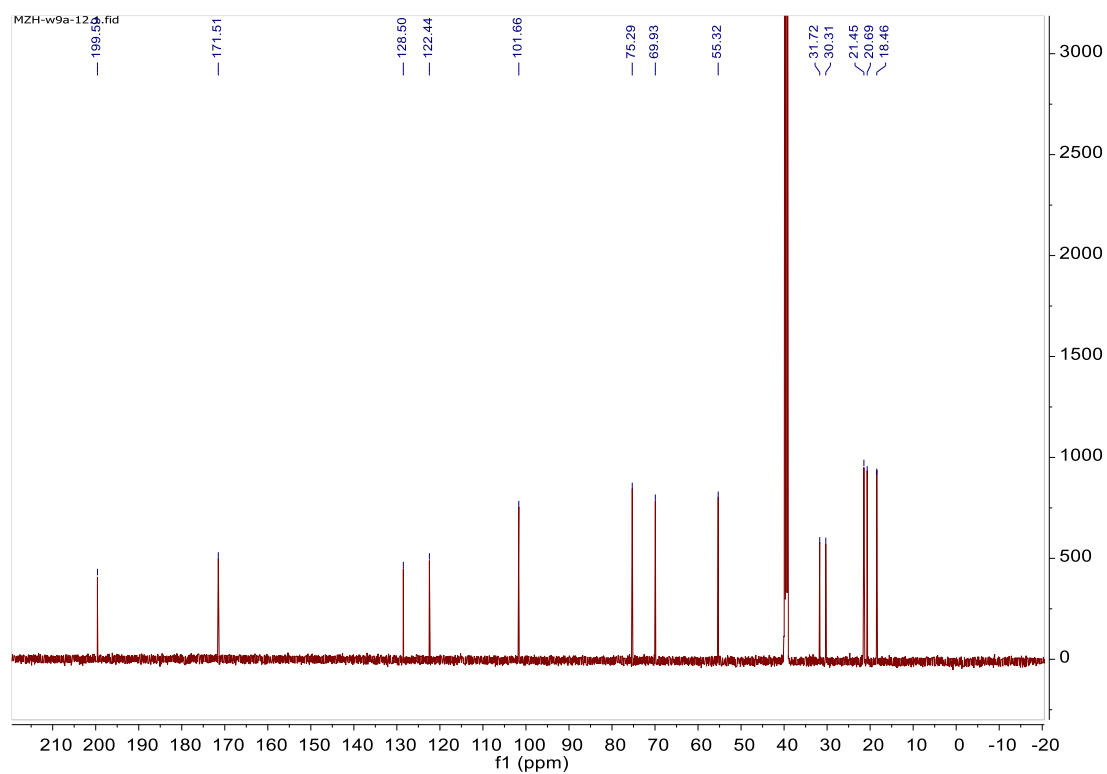


Figure S31. ^{13}C NMR spectrum of 5

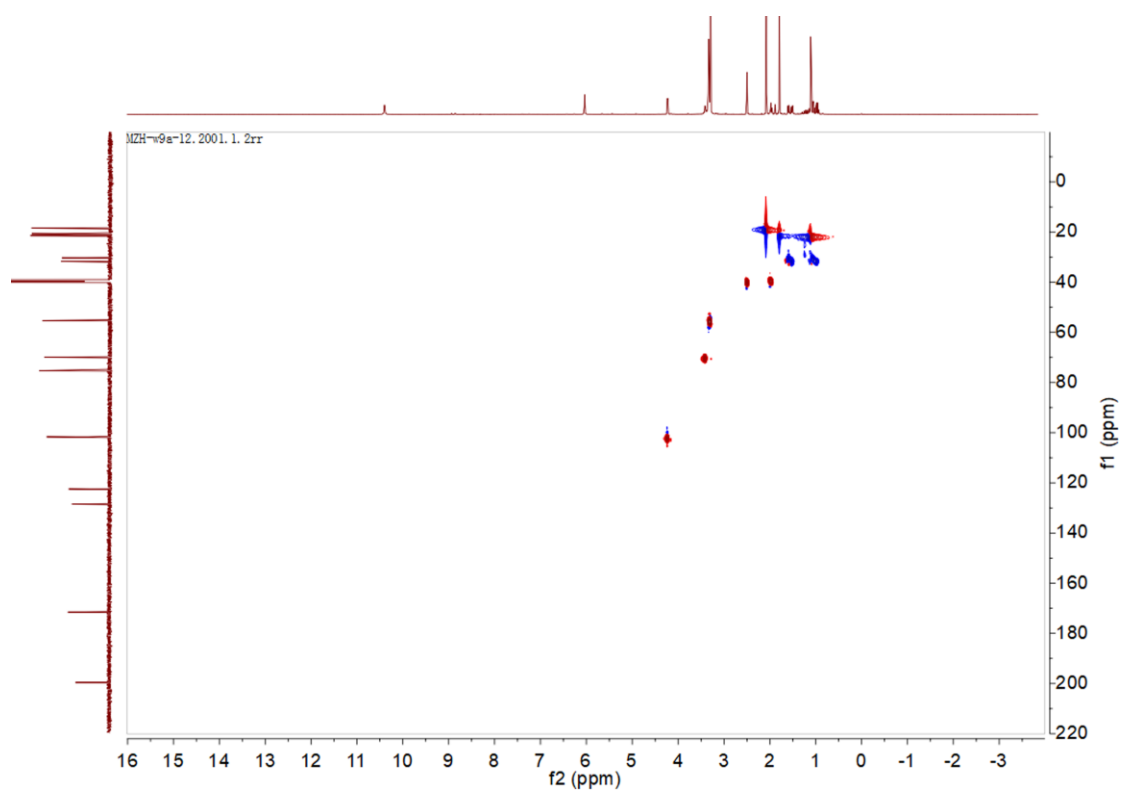


Figure S32. HSQC spectrum of 5

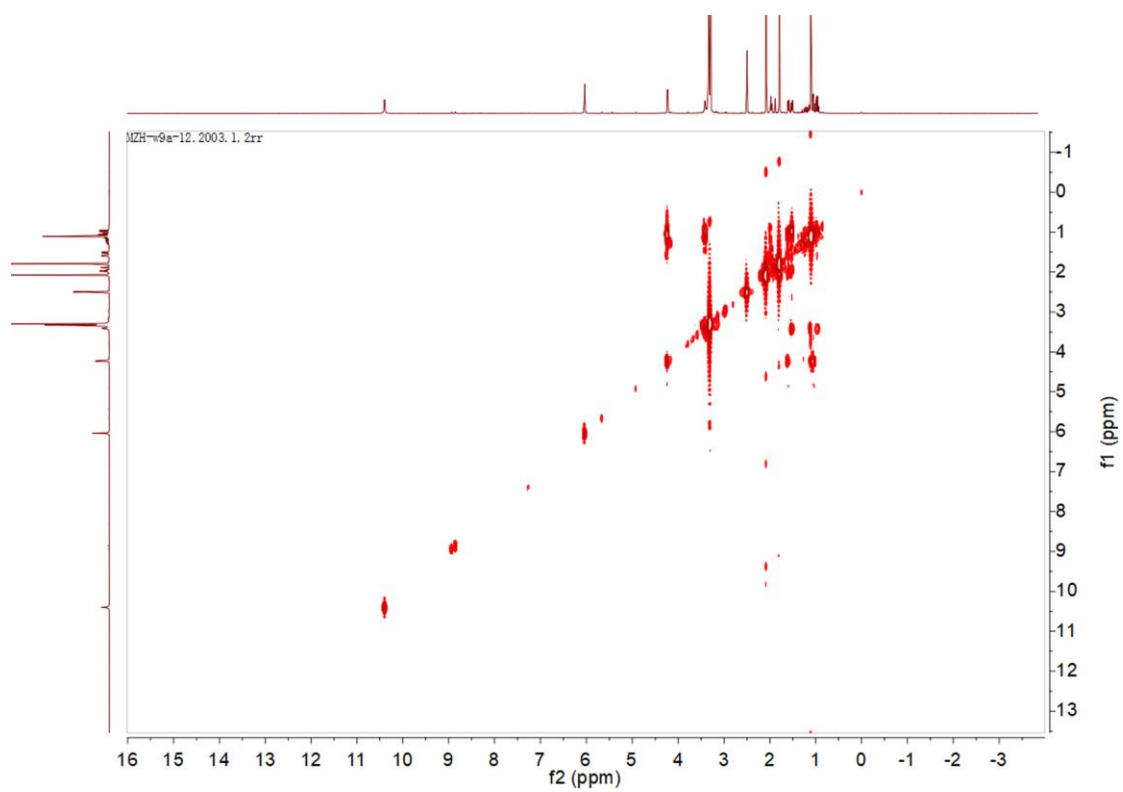


Figure S33. ^1H - ^1H COSY spectrum of **5**

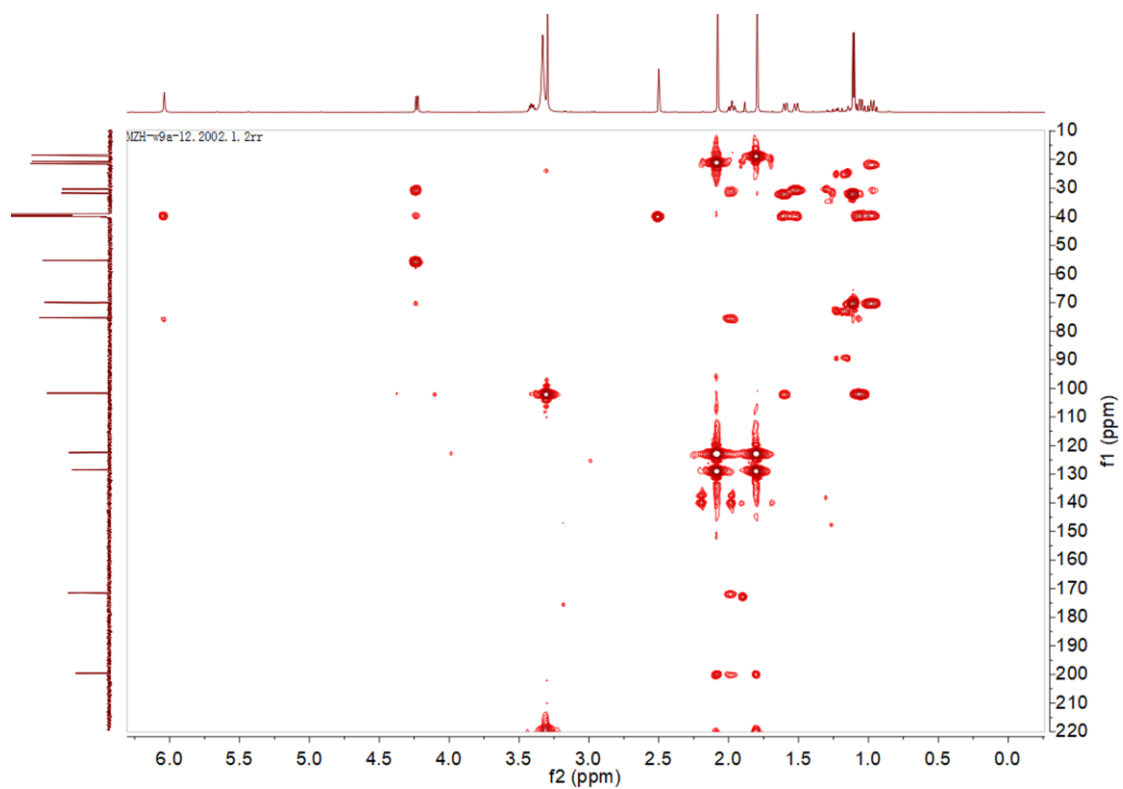


Figure S34. HMBC spectrum of **5**

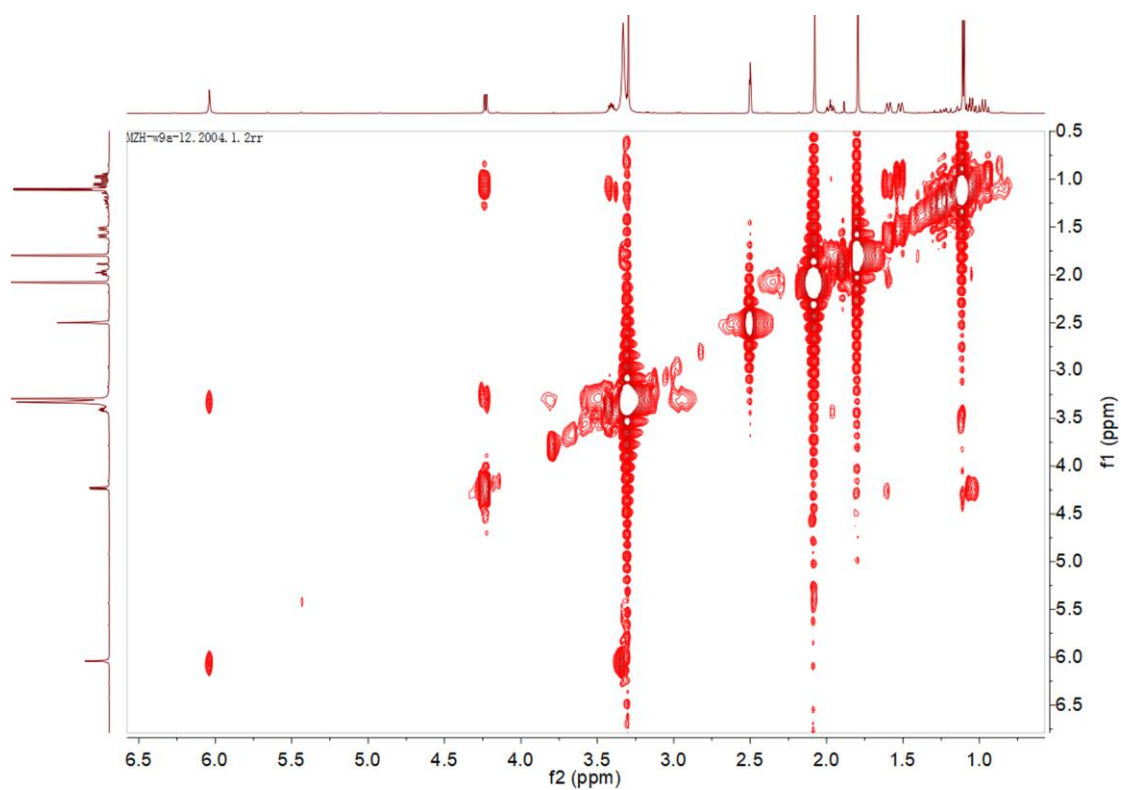


Figure S35. NOESY spectrum of 5

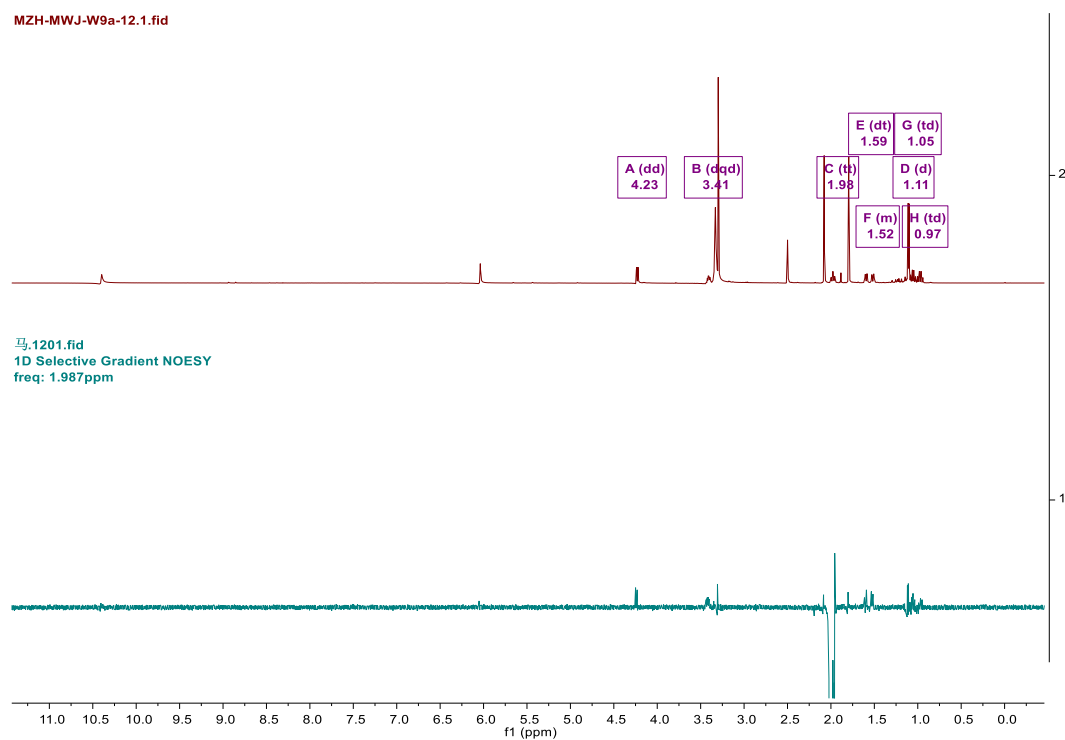


Figure S36. 1D selective NOESY spectrum of 5 (1.98ppm)

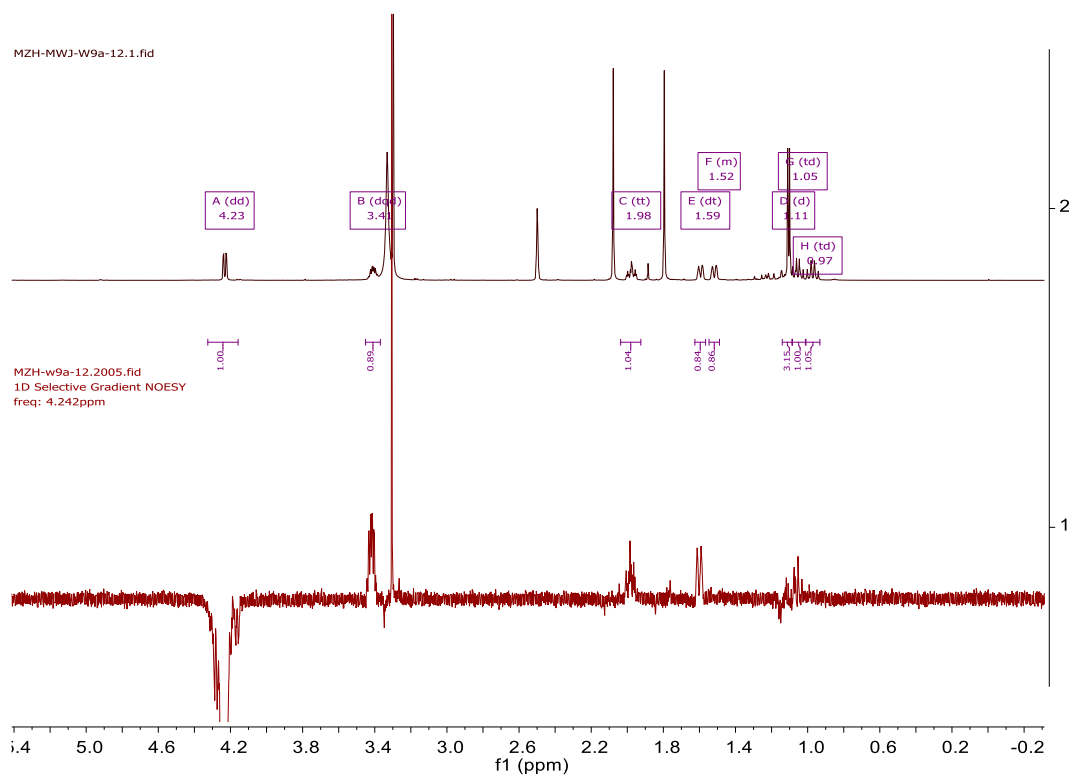


Figure S37. 1D selective NOESY spectrum of **5** (4.24ppm)

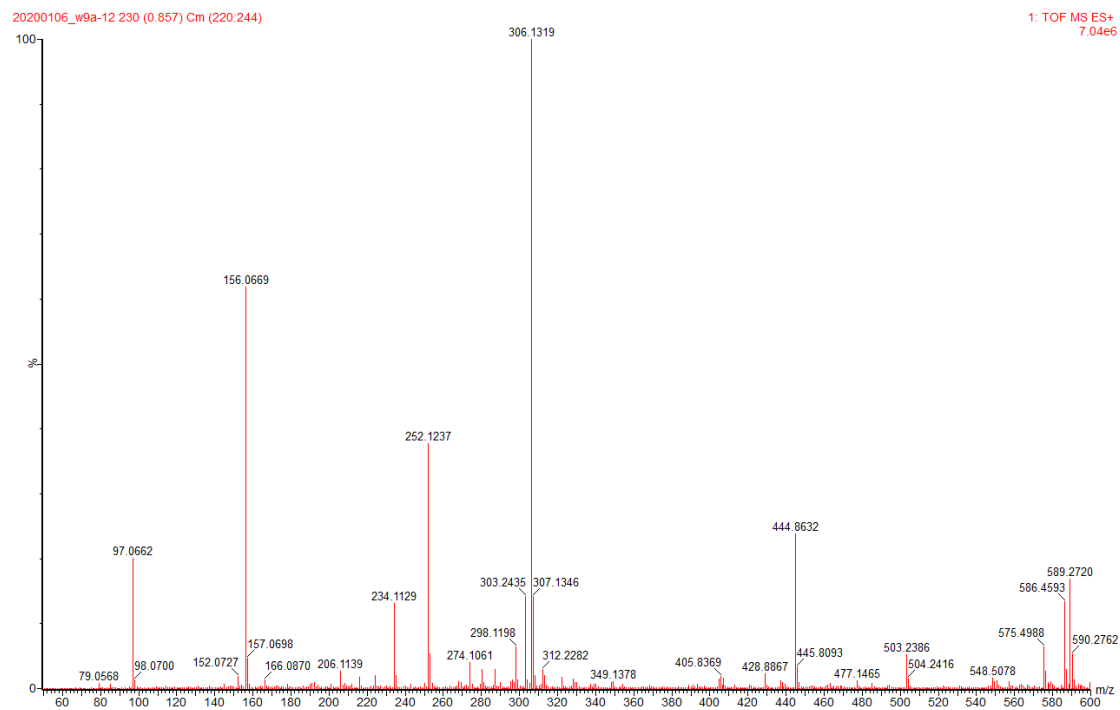


Figure S38. HR-ESI-MS spectrum of **5**

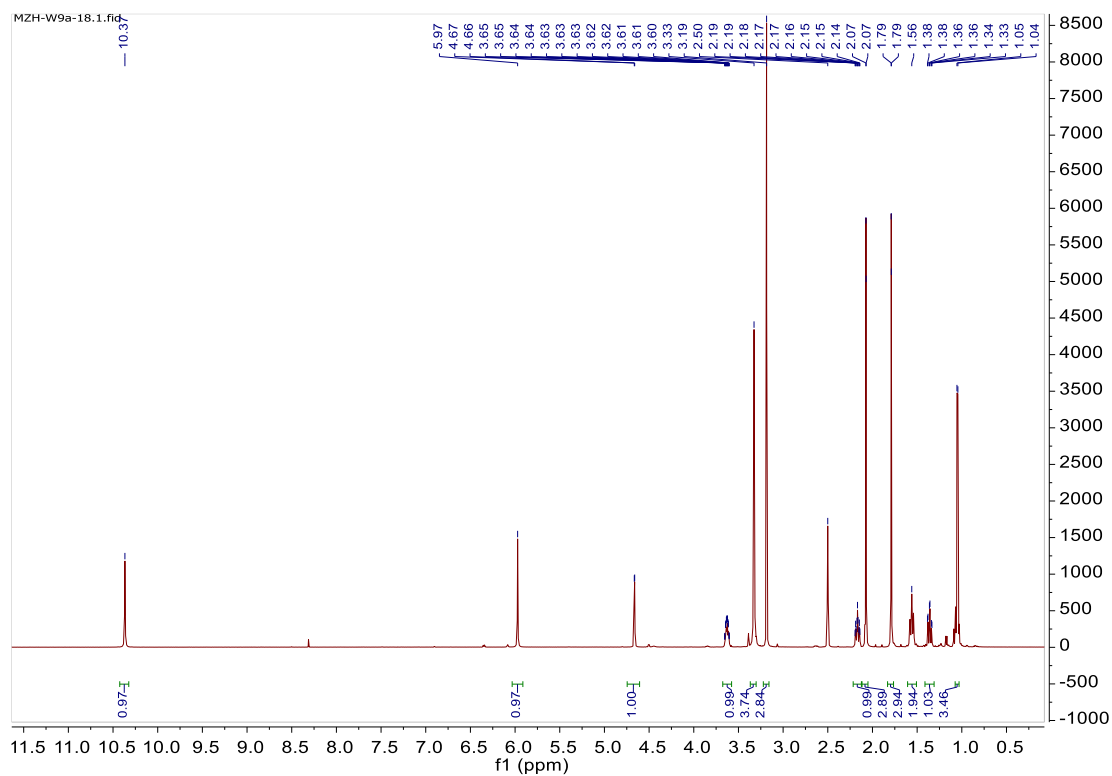


Figure S39. ¹H NMR spectrum of **6**

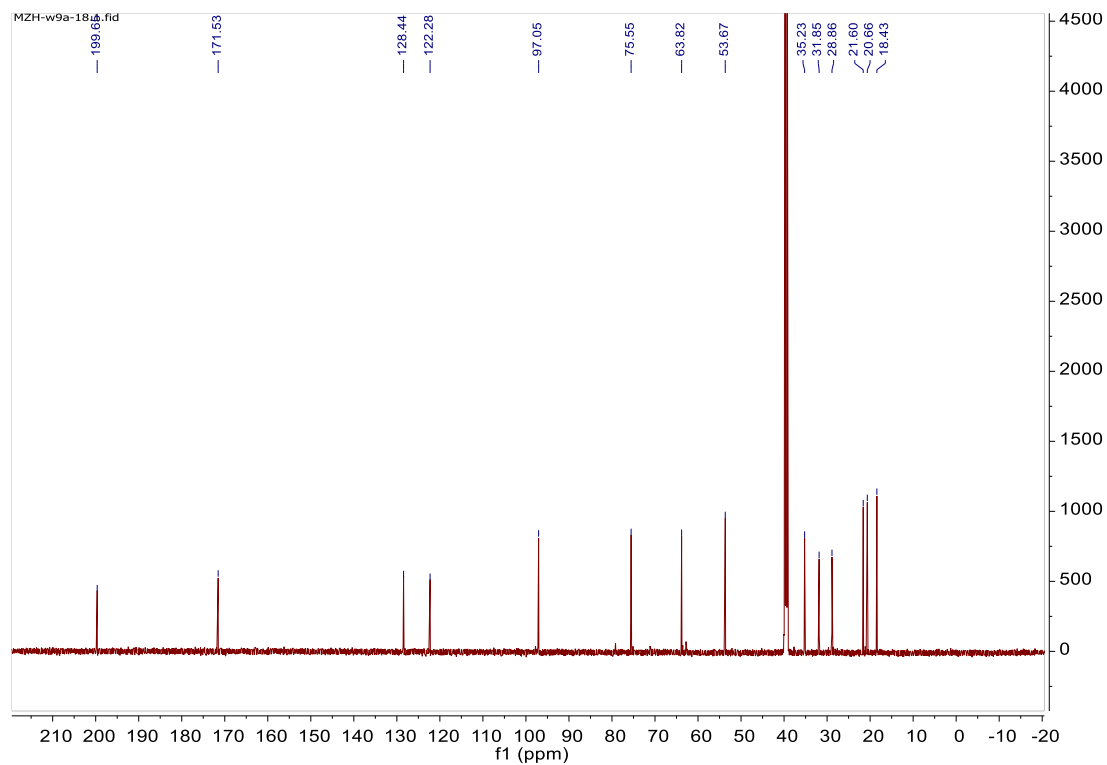


Figure S40. ¹³C NMR spectrum of **6**

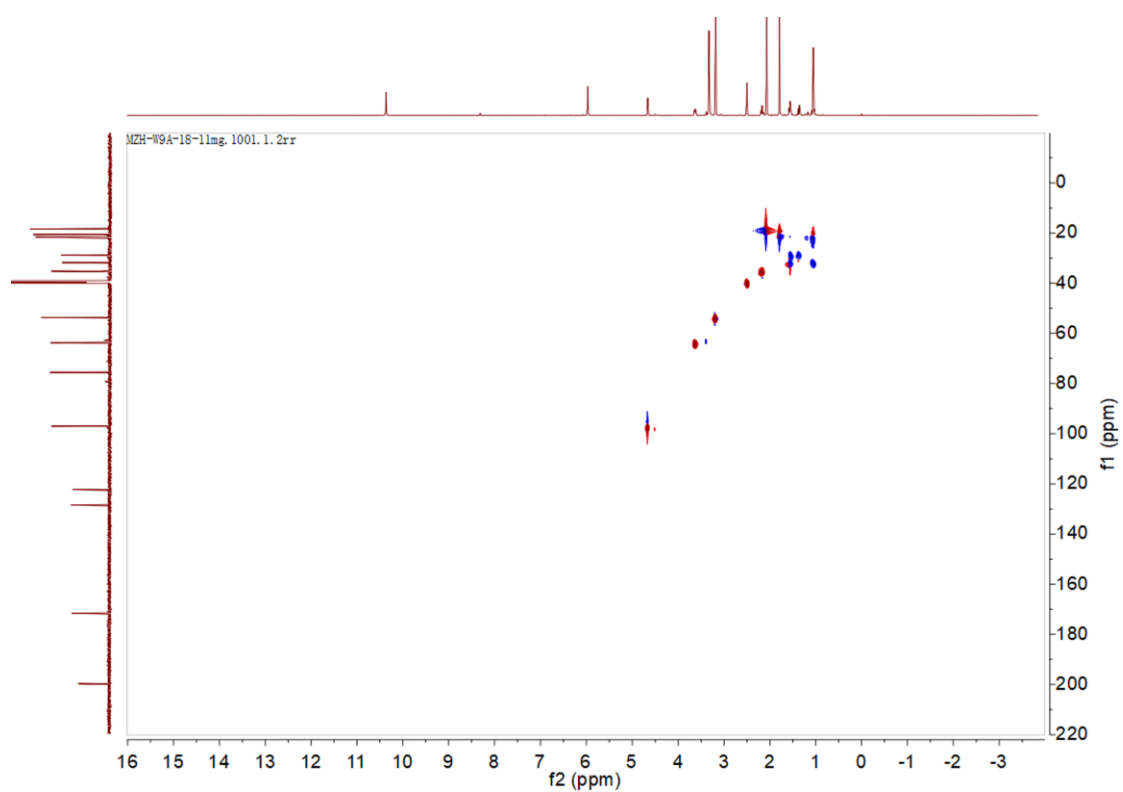


Figure S41. HSQC spectrum of 6

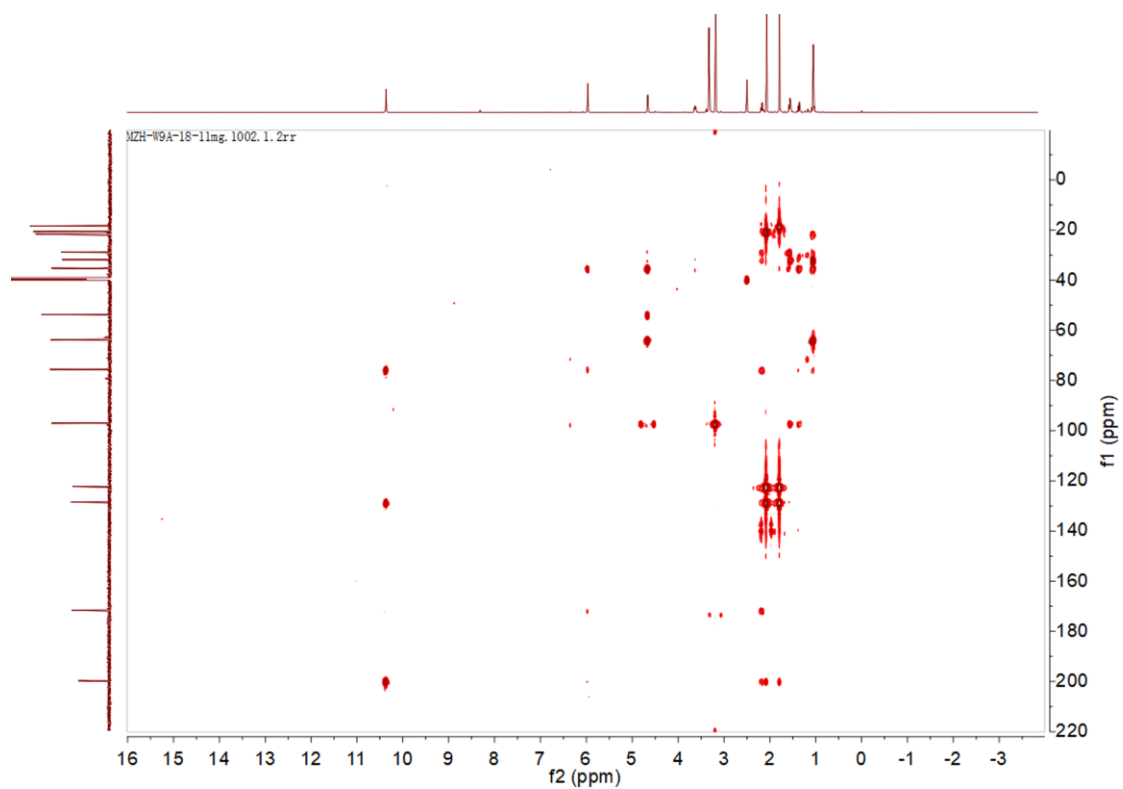


Figure S42. ^1H - ^1H COSY spectrum of 6

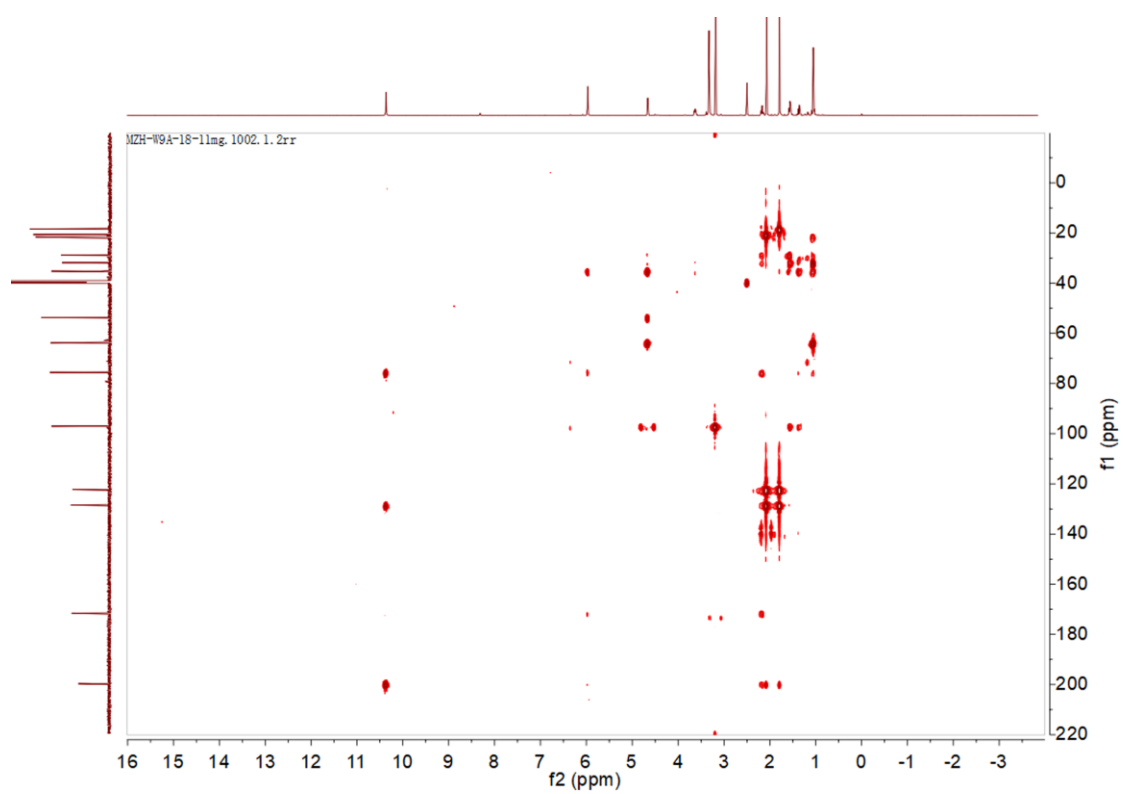


Figure S43. HMBC spectrum of **6**

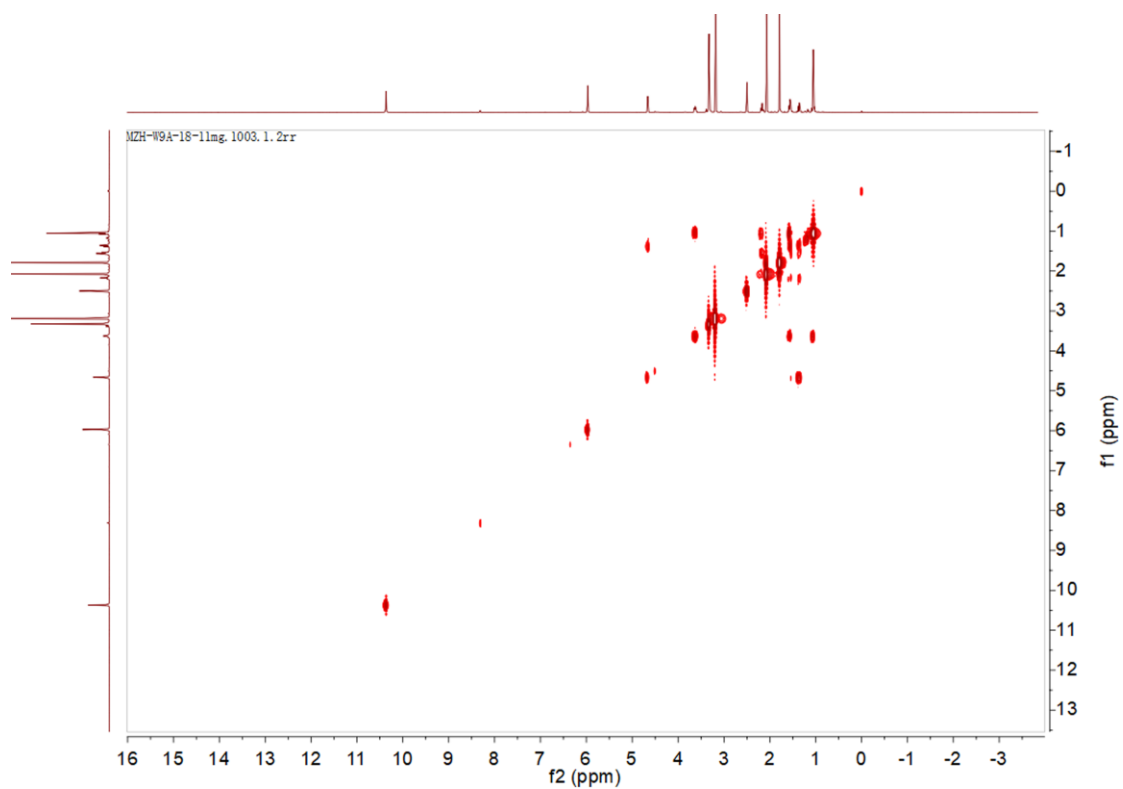


Figure S44. NOESY spectrum of **6**

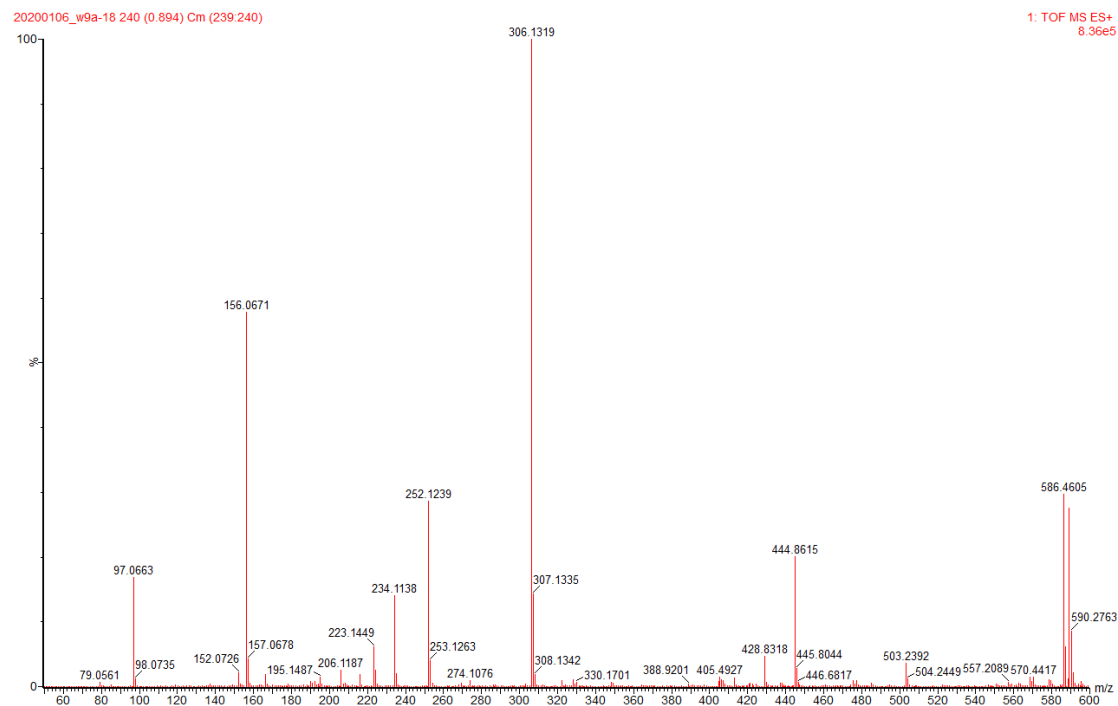


Figure S45. HR-ESI-MS spectrum of **6**

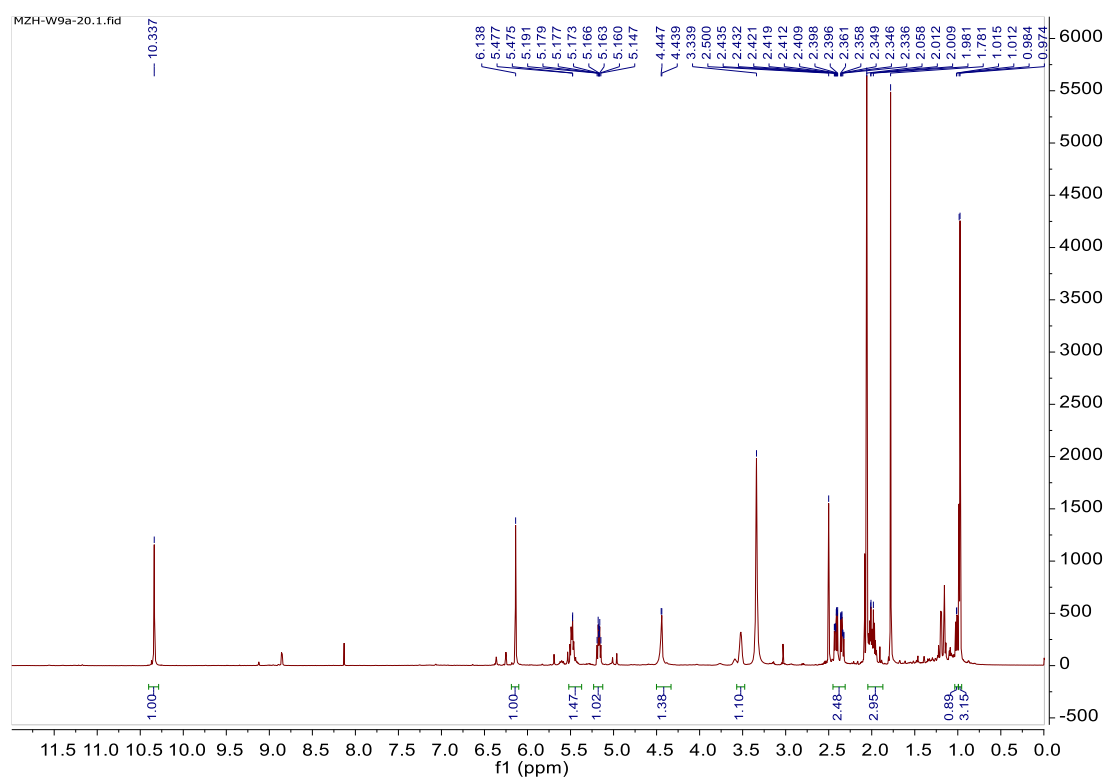


Figure S46. ¹H NMR spectrum of **7**

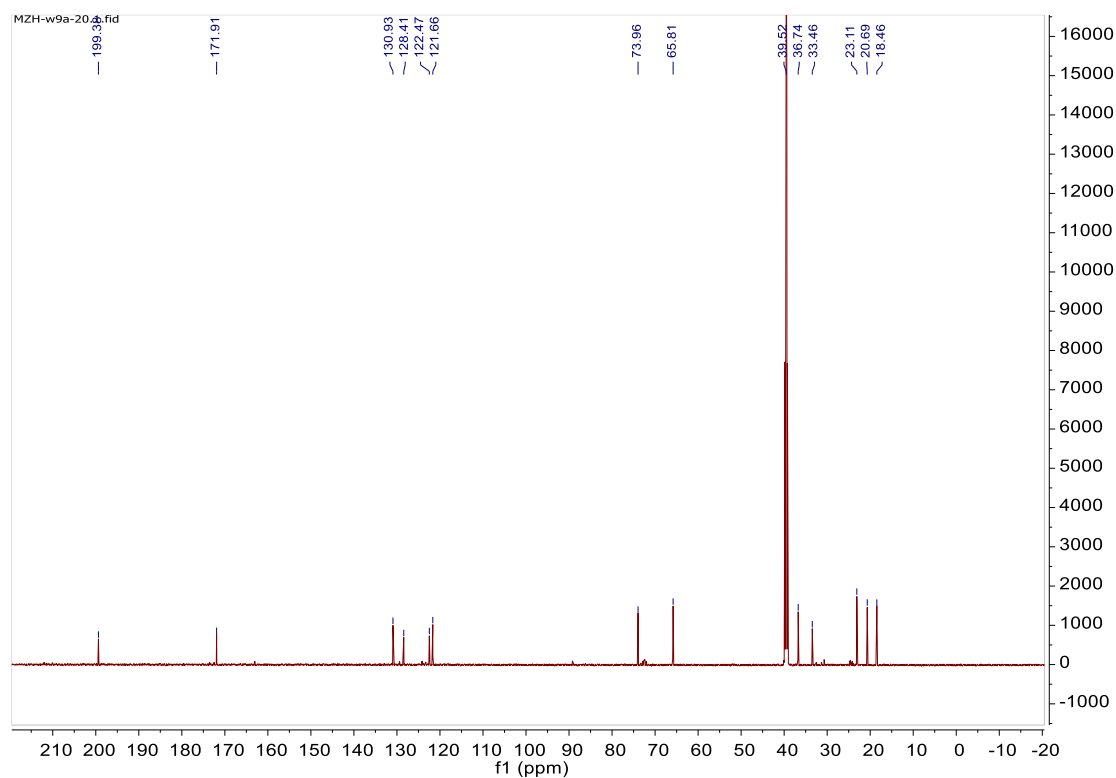


Figure S47. ^{13}C NMR spectrum of 7

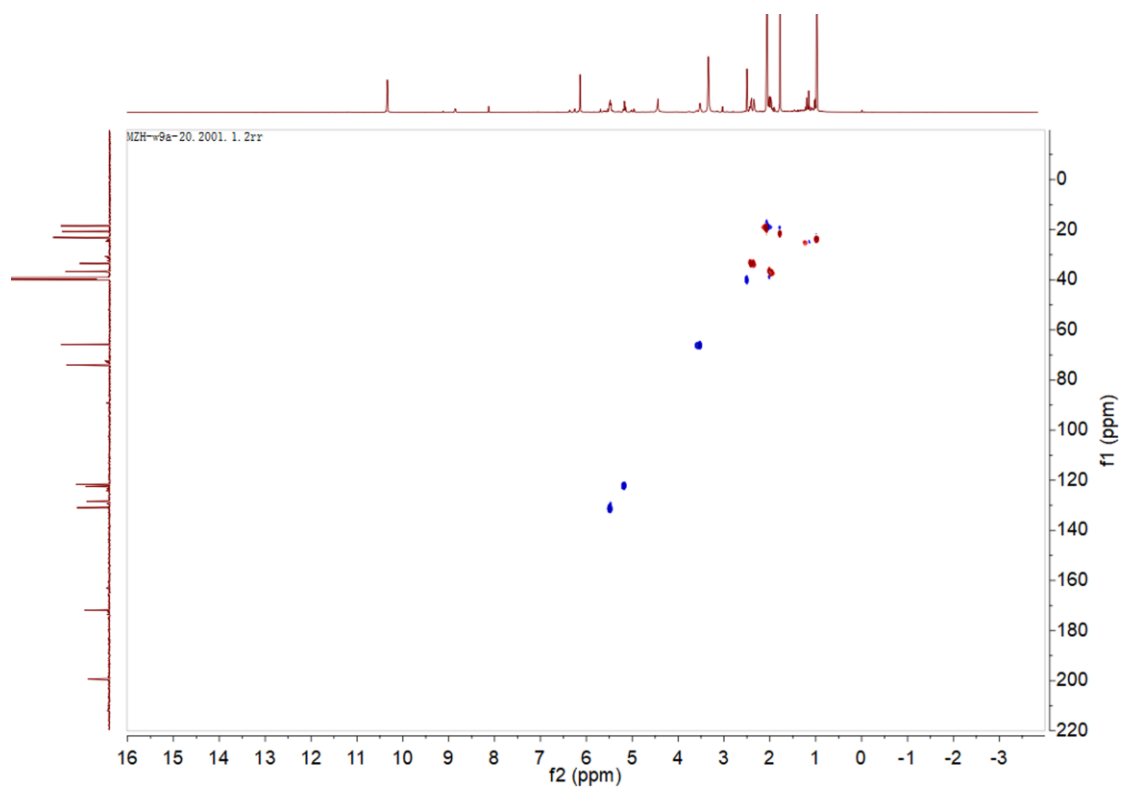


Figure S48. HSQC spectrum of 7

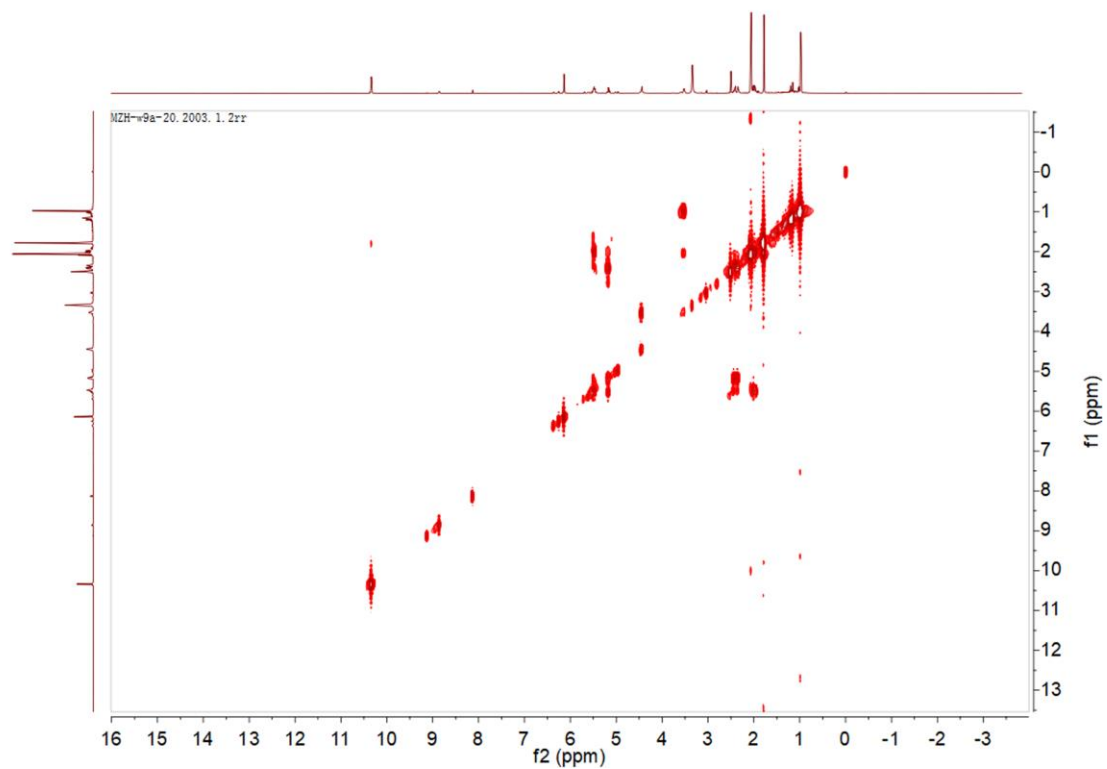


Figure S49. ^1H - ^1H COSY spectrum of **7**

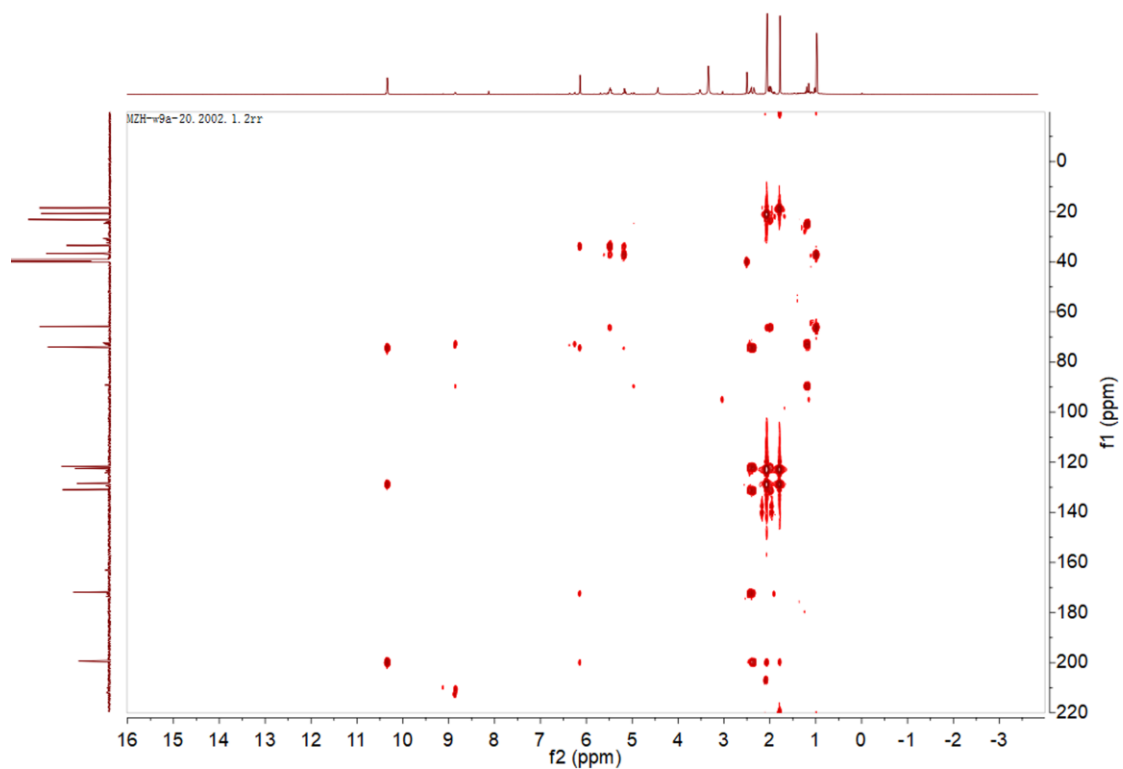


Figure S50. HMBC spectrum of **7**

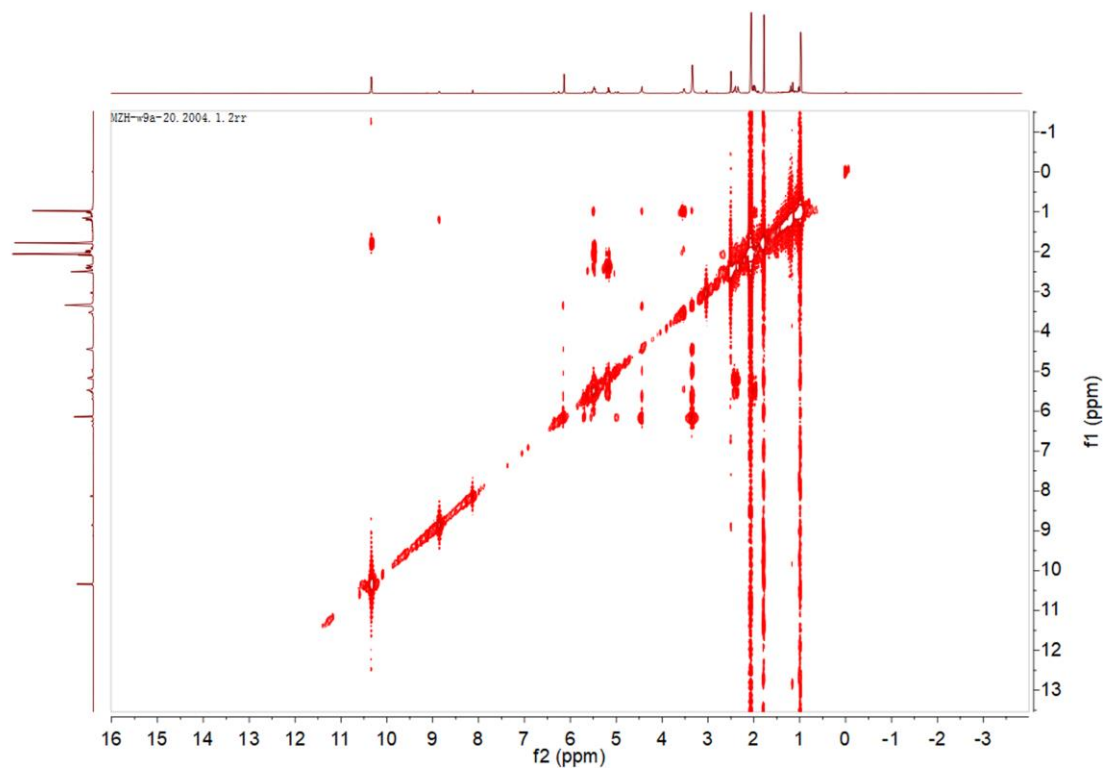


Figure S51. NOESY spectrum of 7

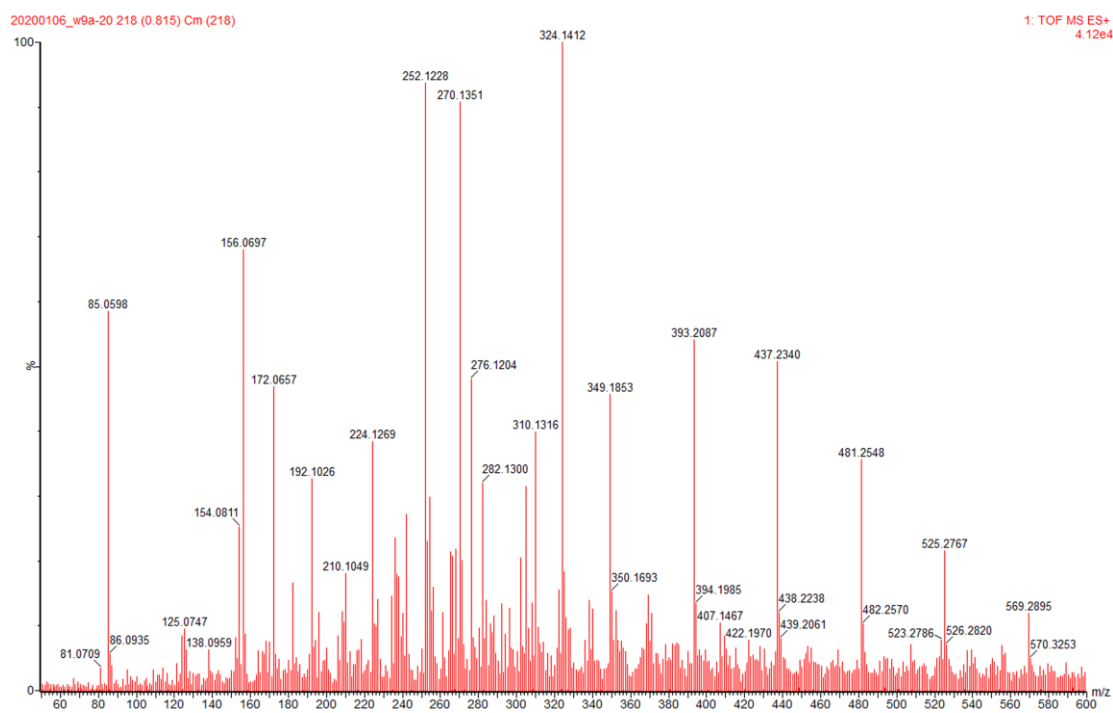


Figure S52. HR-ESI-MS spectrum of 7

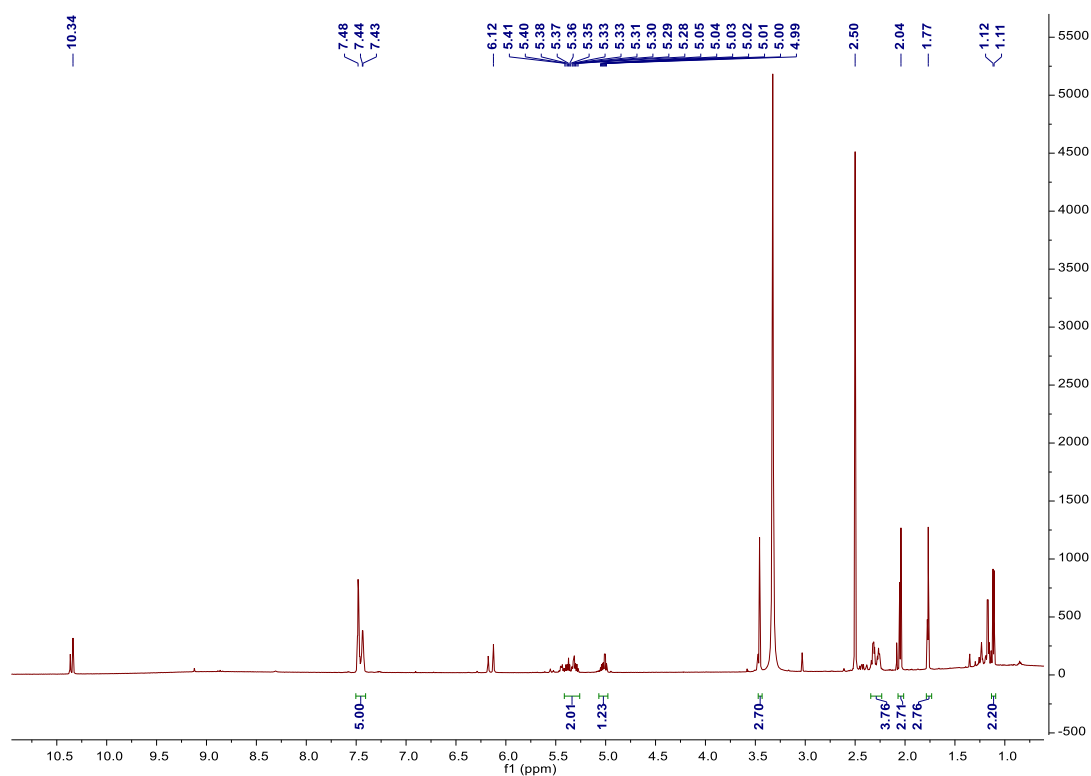


Figure S53. ^1H NMR spectrum of (R)-MTPA ester of 7

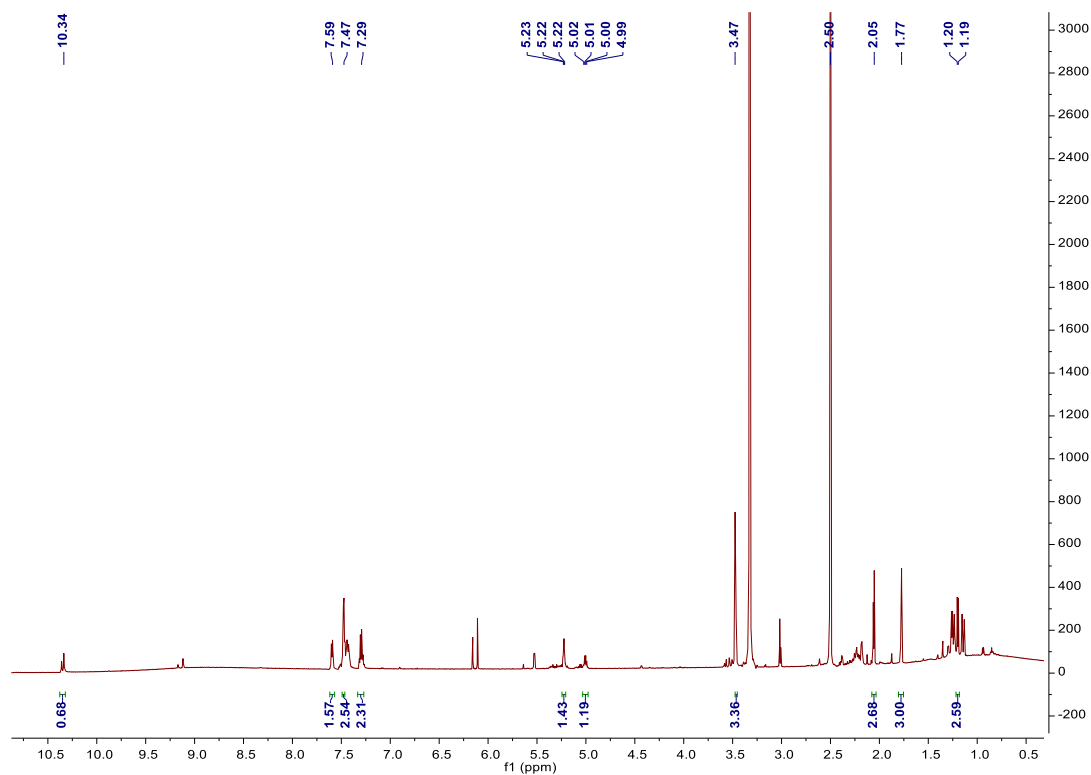
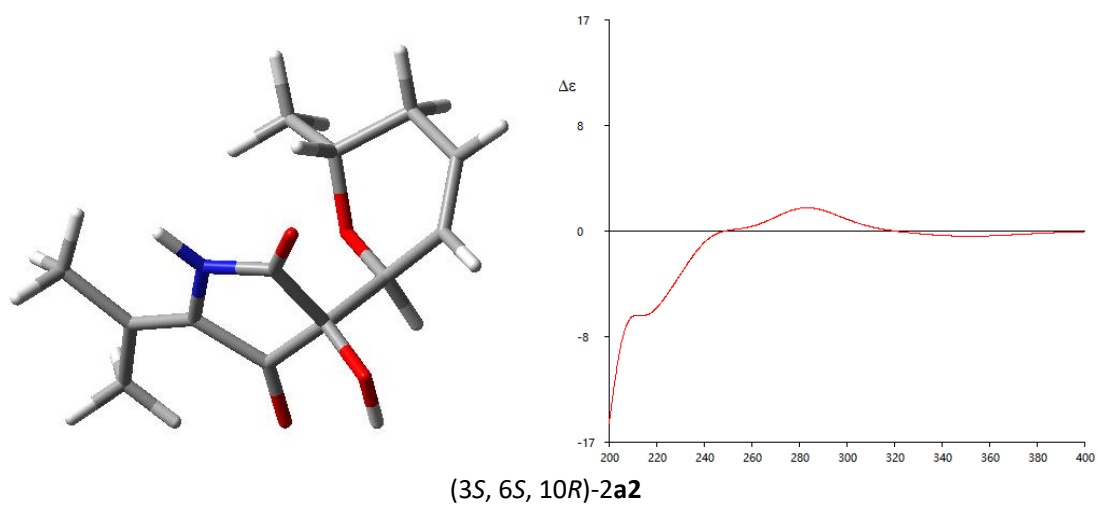
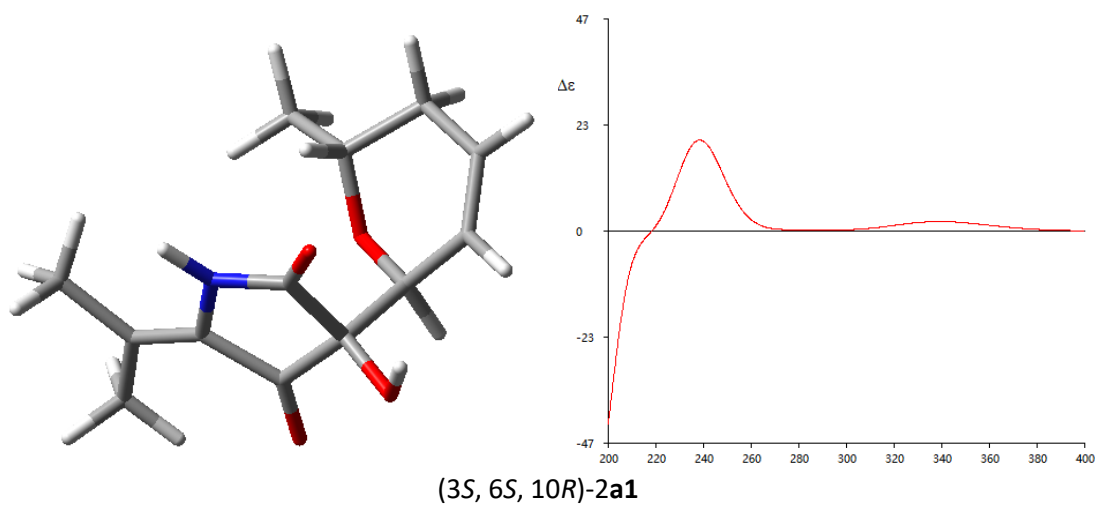


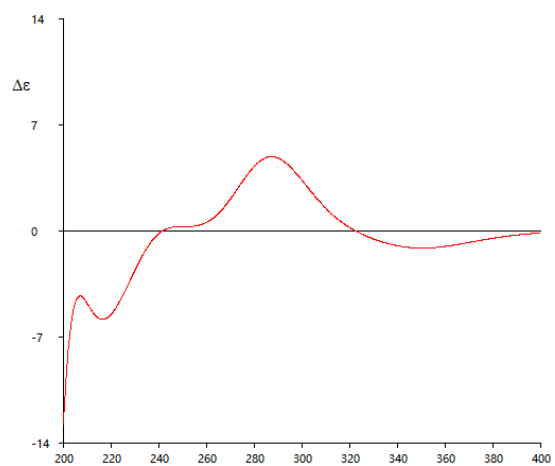
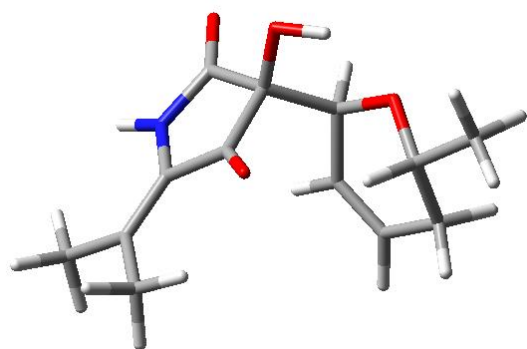
Figure S54. ^1H NMR spectrum of (S)-MTPA ester of 7

Table S2. Relative free energies^a and equilibrium populations^b of conformers for (3S, 6S, 10R)-2a^c

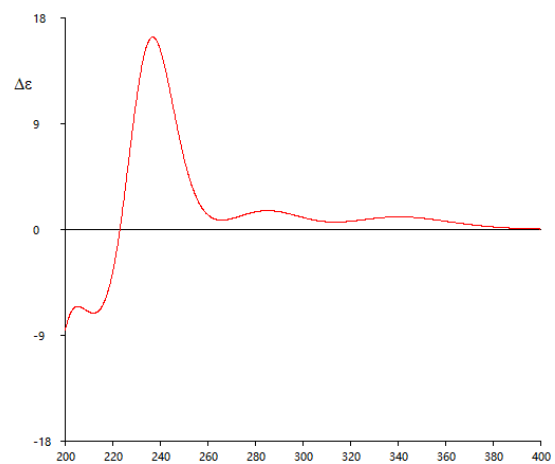
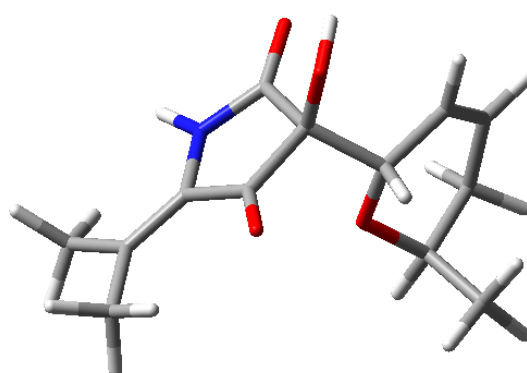
Conformer	ΔG	P (%)
2a1	0.00	42.97
2a2	0.52	17.90
2a3	0.78	11.50
2a4	0.98	8.26
2a5	1.11	6.62
2a6	1.11	6.56

^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.

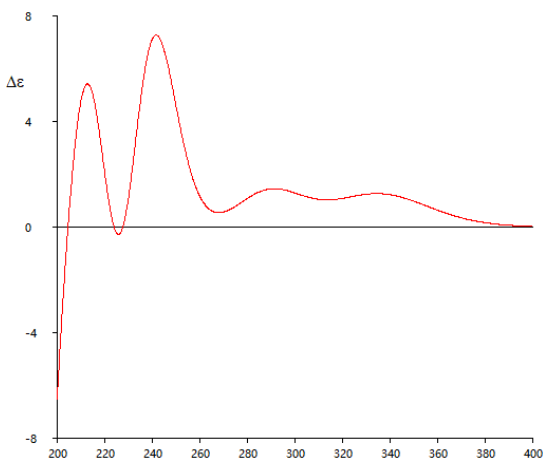
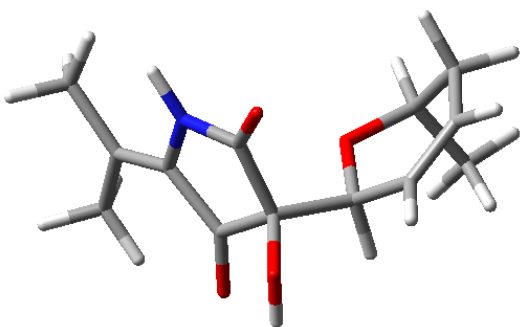




(3*S*, 6*S*, 10*R*)-2a3



(3*S*, 6*S*, 10*R*)-2a4



(3*S*, 6*S*, 10*R*)-2a5

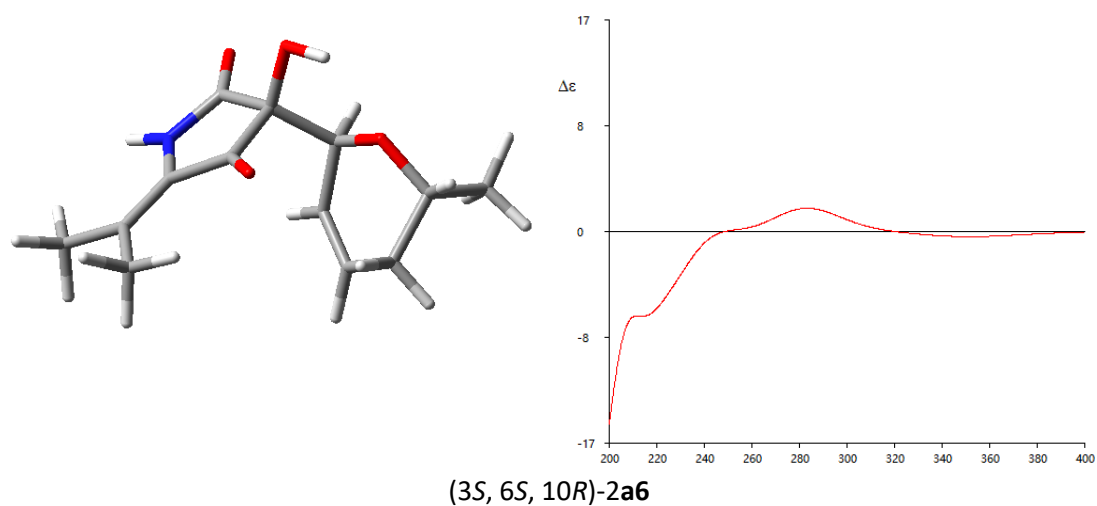
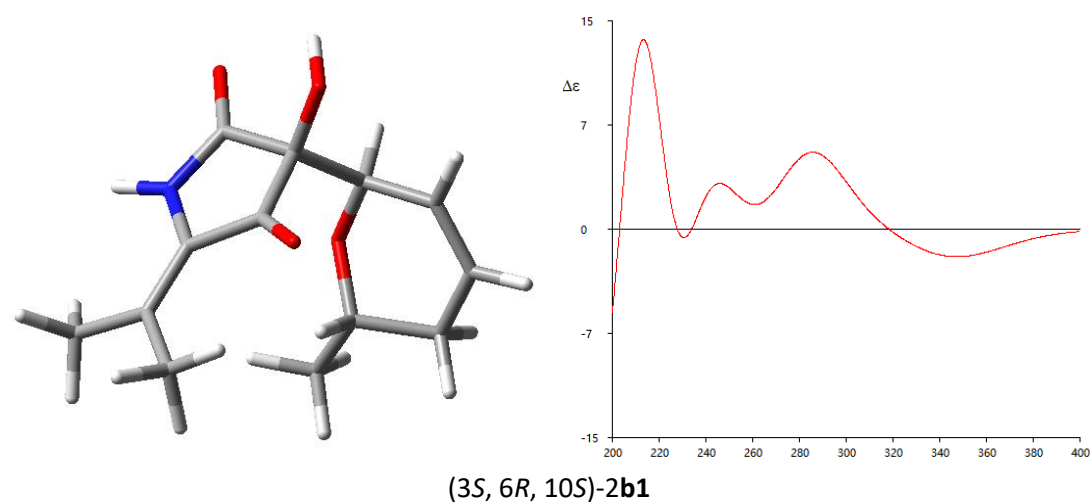


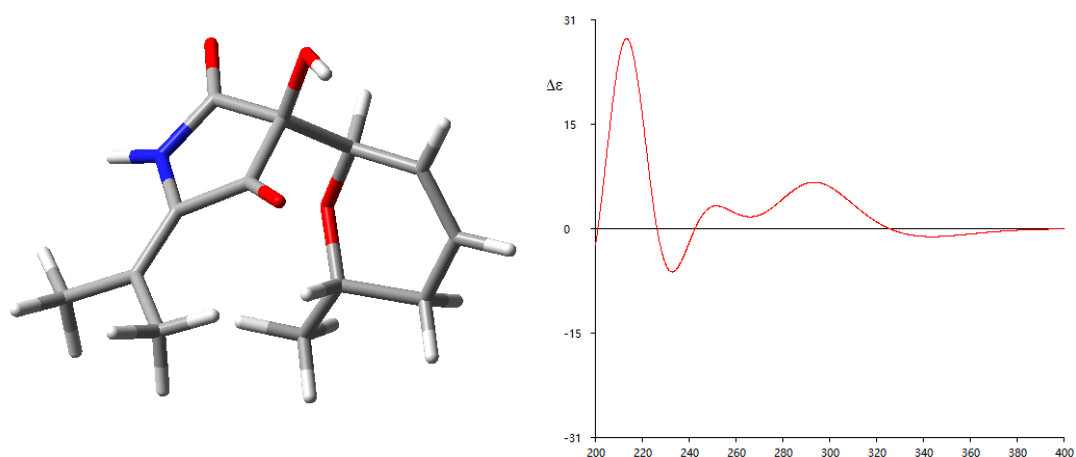
Figure S55. The optimized structures (left) and the calculated CD spectra of conformers (3S, 6S, 10R)-2a in MeOH at M06-2X/def2TZVP level (right). $\sigma=0.30$ eV.

Table S3. Relative free energies^a and equilibrium populations^b of conformers for (3S, 6R, 10S)-2b^c

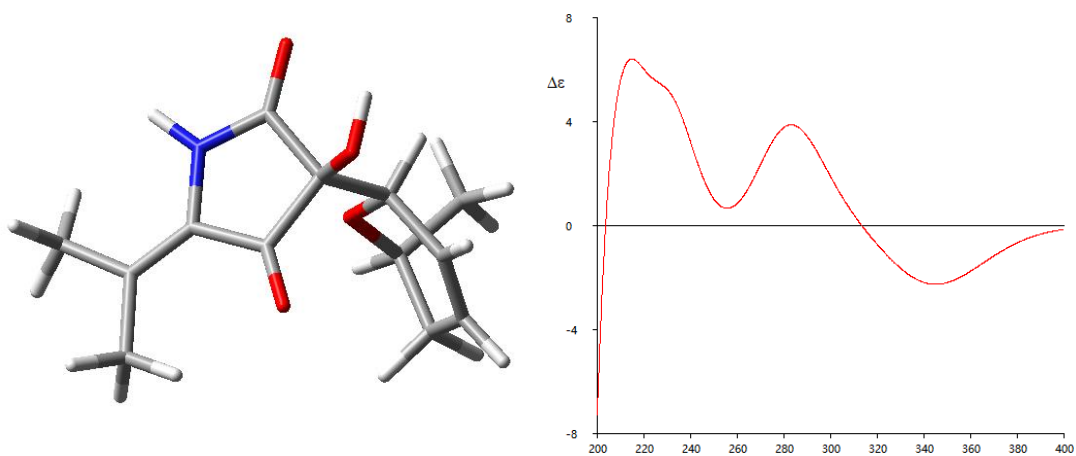
Conformer	ΔG	P (%)
2b1	0.00	51.42
2b2	0.61	18.24
2b3	0.65	17.19
2b4	0.81	13.15

^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.

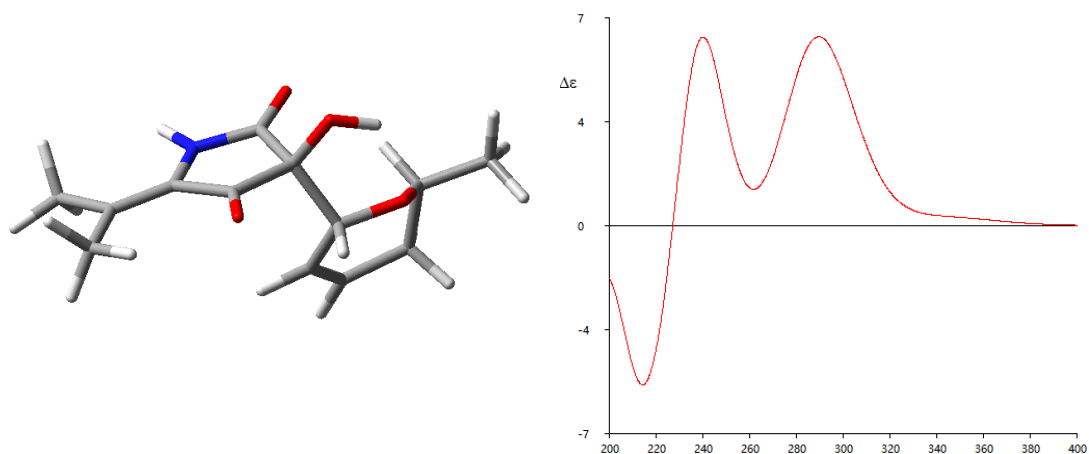




(3S, 6R, 10S)-2b2



(3S, 6R, 10S)-2b3



(3S, 6R, 10S)-2b4

Figure S56. The optimized structures (left) and the calculated CD spectra of conformers (3S, 6R, 10S)-2b in MeOH at M06-2X/def2TZVP level (right). $\sigma=0.30$ eV.

Table S4. Relative free energies^a and equilibrium populations^b of conformers for (3S, 6S, 9R)-3a^c

Conformer	ΔG	P (%)
3a1	0.00	45.78
3a2	0.47	20.56
3a3	0.48	20.22
3a4	0.73	13.45

^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.

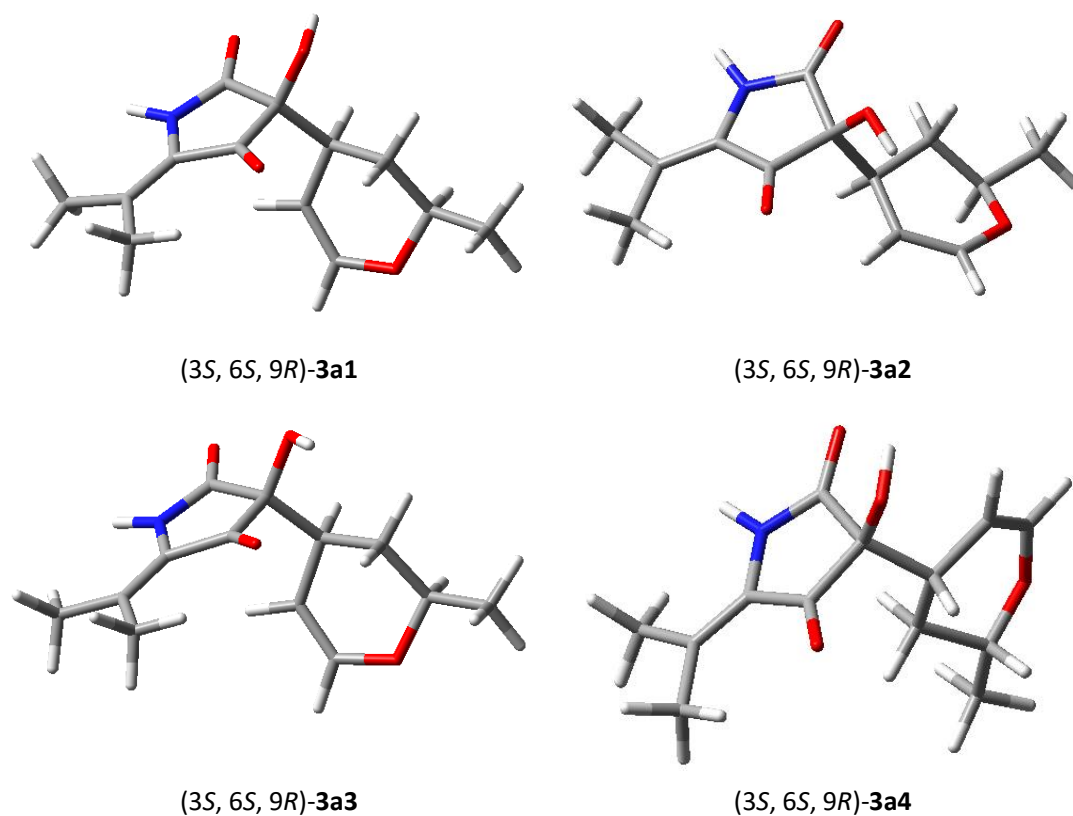


Figure S57. The optimized structure of conformers for (3*S*, 6*S*, 9*R*)-**3a** in MeOH at B3LYP/6-31G (d) level

Table S5. Relative free energies^a and equilibrium populations^b of conformers for (3*S*, 6*R*, 9*S*)-**3b**^c

Conformer	ΔG	P (%)
3b1	0.00	46.24
3b2	0.36	25.13
3b3	0.52	19.29
3b4	0.95	9.34

^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.

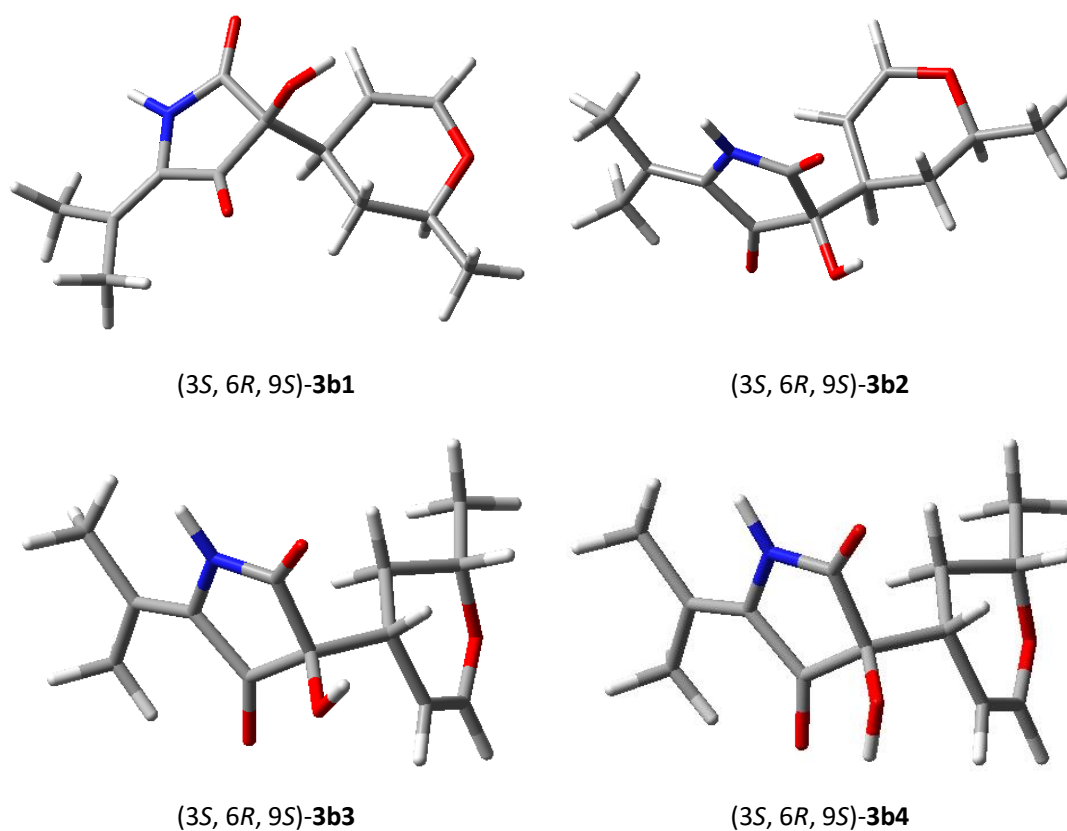
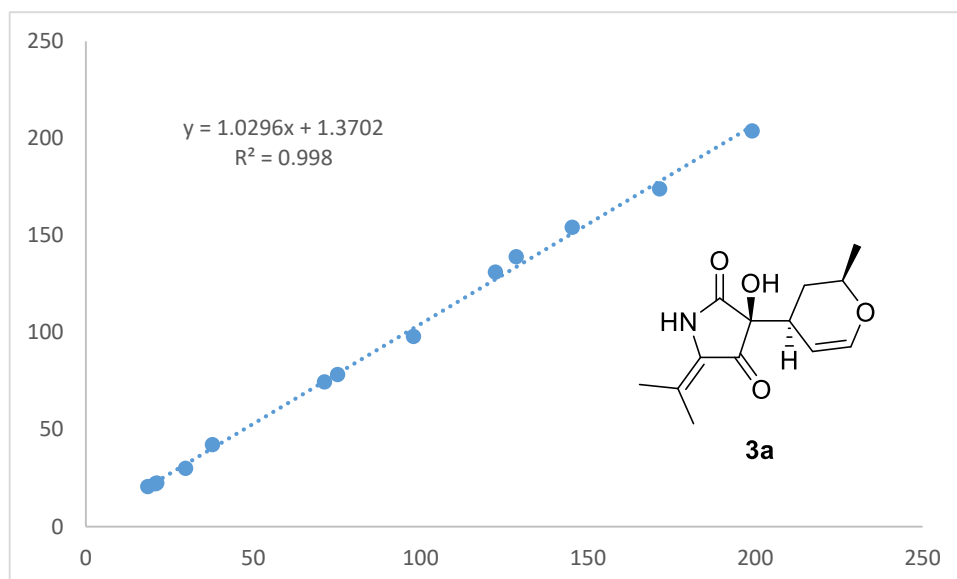


Figure S58. The optimized structure of conformers for (3*S*, 6*R*, 9*S*)-**3b** in MeOH at B3LYP/6-31G (d) level



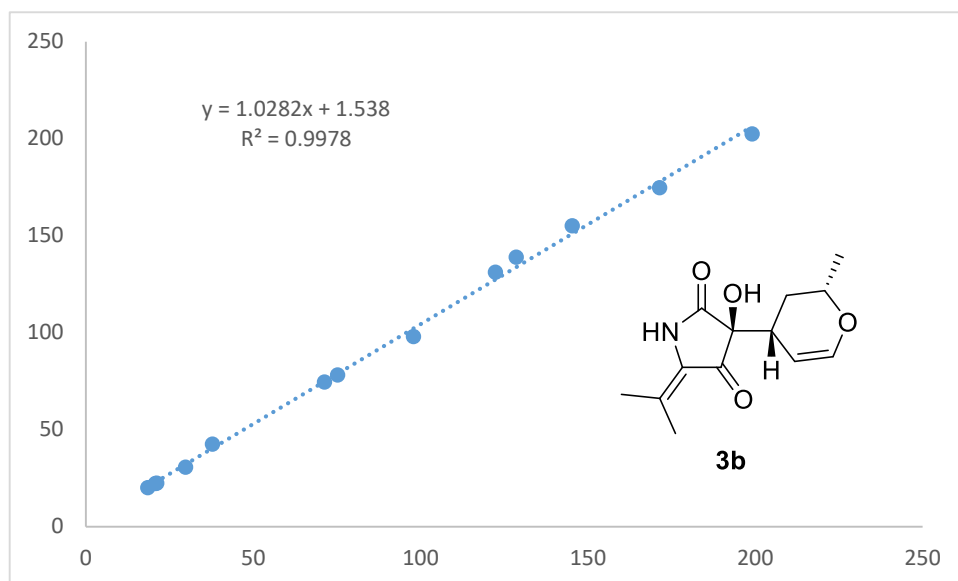


Figure S59. Correlation plots of experimental ^{13}C NMR chemical shifts versus the corresponding calculated data for (3S, 6S, 9R)-3a and (3S, 6R, 9S)-3b

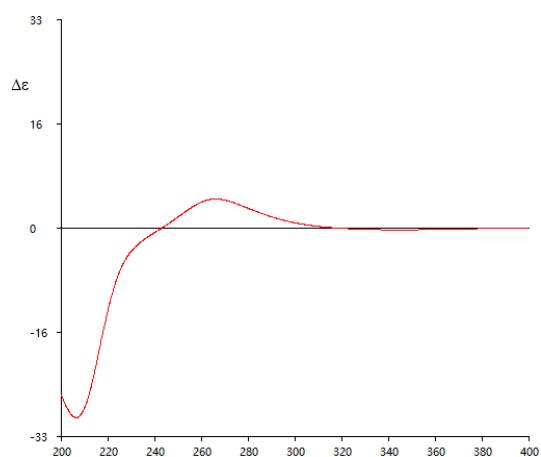
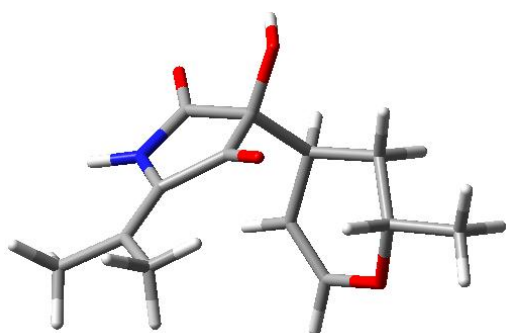
Nuclei	sp2?	DP4+	89.66%	10.34%
		xperimental	Isomer 1	Isomer 2
C	x	122.3	131.118	131.161
C	x	199.04	203.806	202.455
C		75.14	78.319	78.102
C	x	171.35	174.002	174.743
C	x	128.57	139.058	138.82
C		18.49	20.49	20.138
C		20.71	22.077	22.151
C		37.73	42.155	42.442
C	x	97.81	97.961	98.007
C	x	145.33	154.144	155.104
C		71.2	74.538	74.579
C		29.7	30.025	30.613
C		21.17	22.413	22.364

Figure S60. DP4+ probabilities (%) for conformers (3S, 6S, 9R)-3a (isomer 1) and (3S, 6R, 9S)-3b (isomer 2)

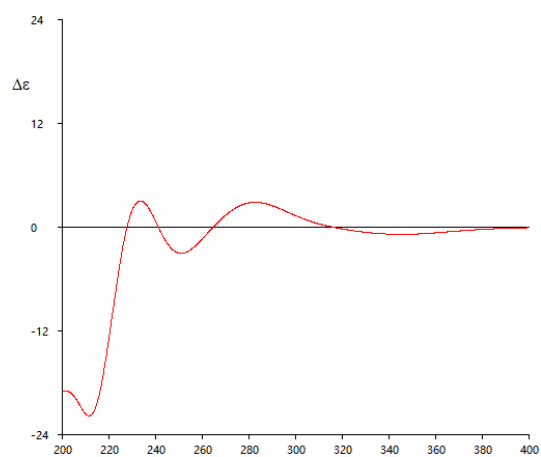
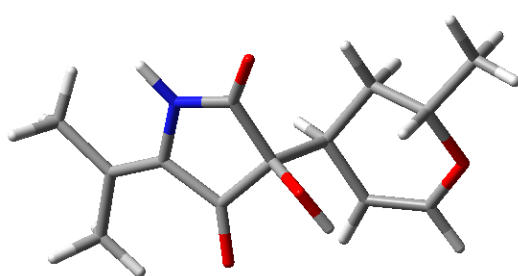
Table S6. Relative free energies^a and equilibrium populations^b of conformers for (3S,6S,9S)-4a^c

Conformer	ΔG	P (%)
4a1	0.00	56.61
4a2	0.35	31.47
4a3	1.30	6.28
4a4	1.37	5.64

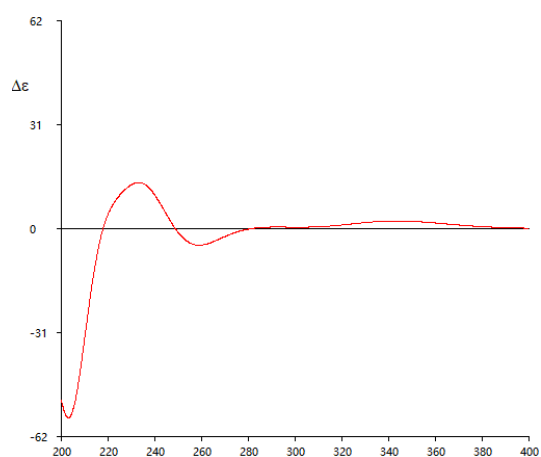
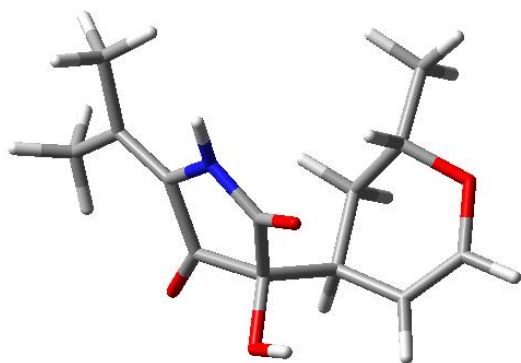
^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



(3S,6S,9S)-4a1



(3S,6S,9S)-4a2



(3S,6S,9S)-4a3

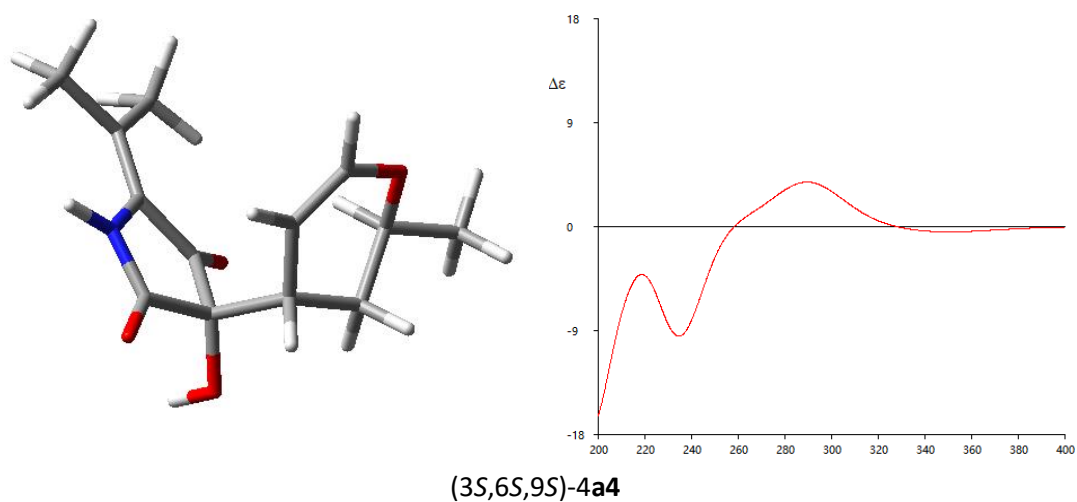
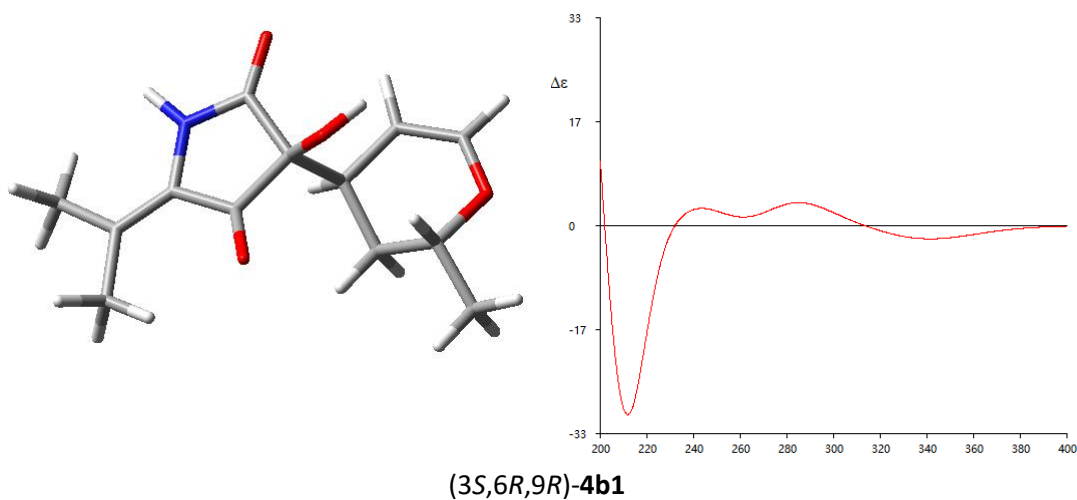


Figure S61. The optimized structures (left) and the calculated CD spectra for conformers (3S,6S,9S)-4a in MeOH at M06-2X/def2TZVP level (right). $\sigma=0.30$ eV.

Table S7. Relative free energies^a and equilibrium populations^b of conformers for (3S,6R,9R)-4b^c

Conformer	ΔG	P (%)
4b1	0.00	58.58
4b2	0.34	32.79
4b3	1.13	8.63

^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



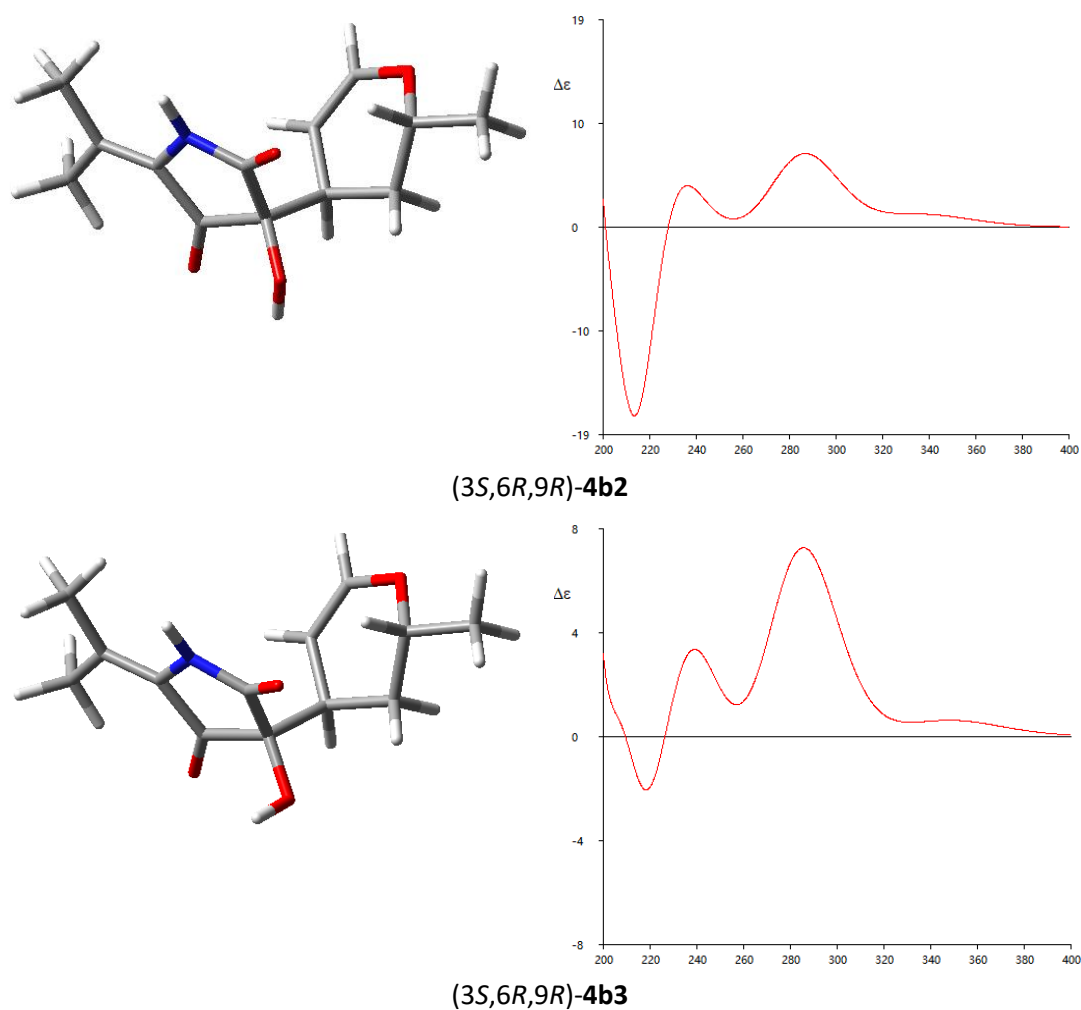
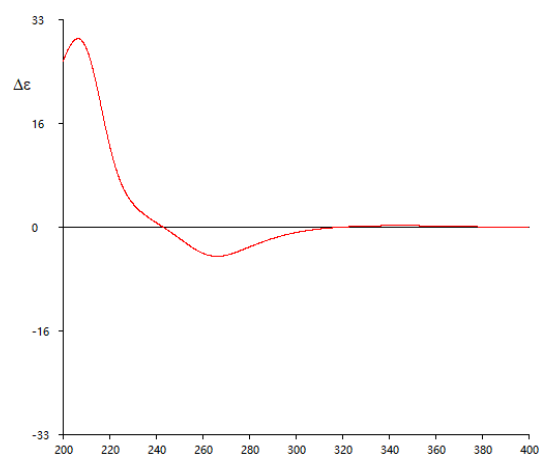
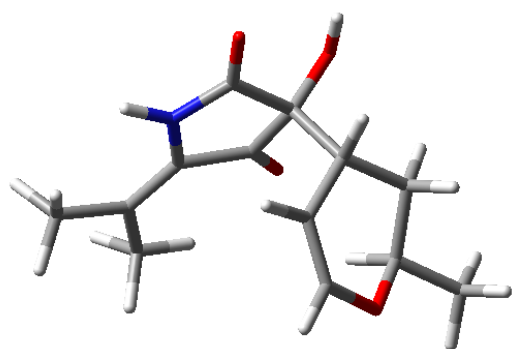


Figure S62. The optimized structures (left) and the calculated CD spectra for conformers (3S,6R,9R)-**4b** in MeOH at M06-2X/def2TZVP level (right). $\sigma=0.30$ eV.

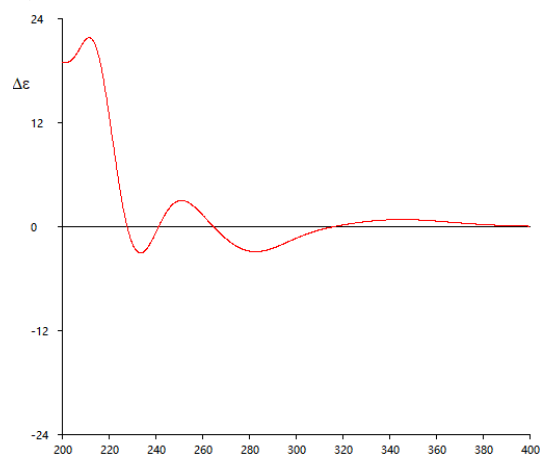
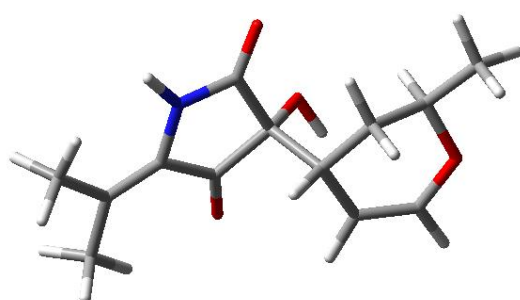
Table S8. Relative free energies^a and equilibrium populations^b of conformers for (3R,6R,9R)-**4c**^c

Conformer	ΔG	P (%)
4c1	0.00	57.30
4c2	0.36	31.09
4c3	1.34	5.95
4c4	1.37	5.65

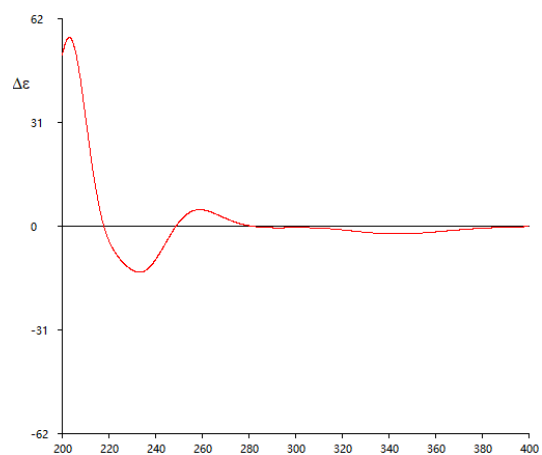
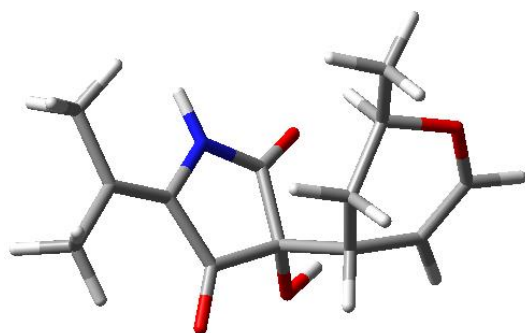
^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



(3*R*,6*R*,9*R*)-4c1



(3*R*,6*R*,9*R*)-4c2



(3*R*,6*R*,9*R*)-4c3

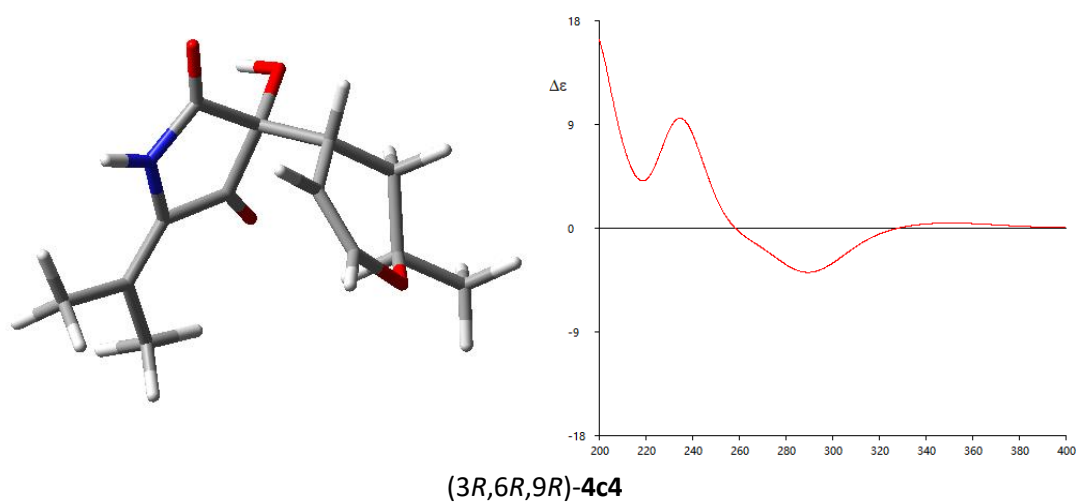
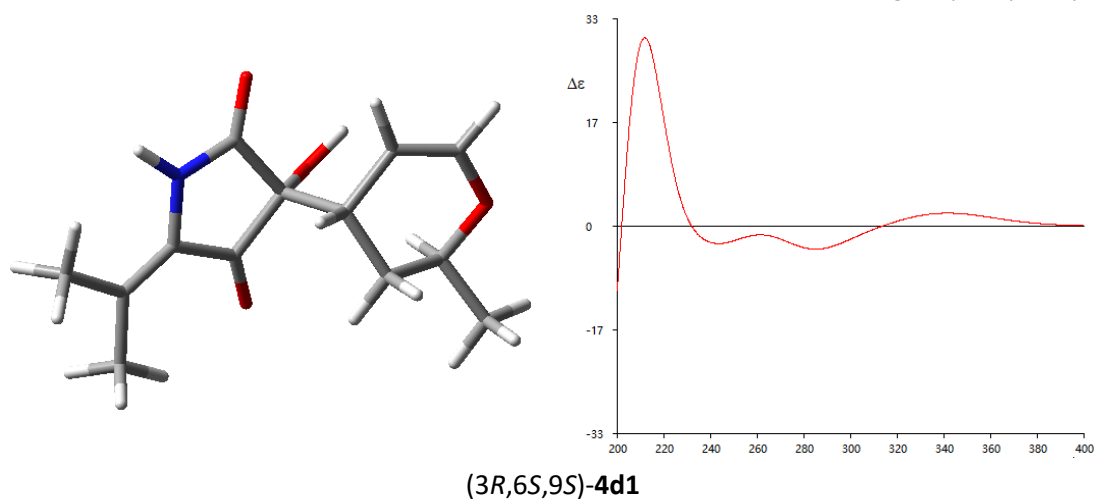


Figure S63. The optimized structures (left) and the calculated CD spectra for conformers (3R,6R,9R)-4c in MeOH at M06-2X/def2TZVP level (right). $\sigma=0.30$ eV.

Table S9. Relative free energies^a and equilibrium populations^b of conformers for (3R,6S,9S)-4d^c

Conformer	ΔG	P (%)
4d1	0.00	54.38
4d2	0.34	30.43
4d4	1.13	8.00
4d4	1.20	7.19

^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



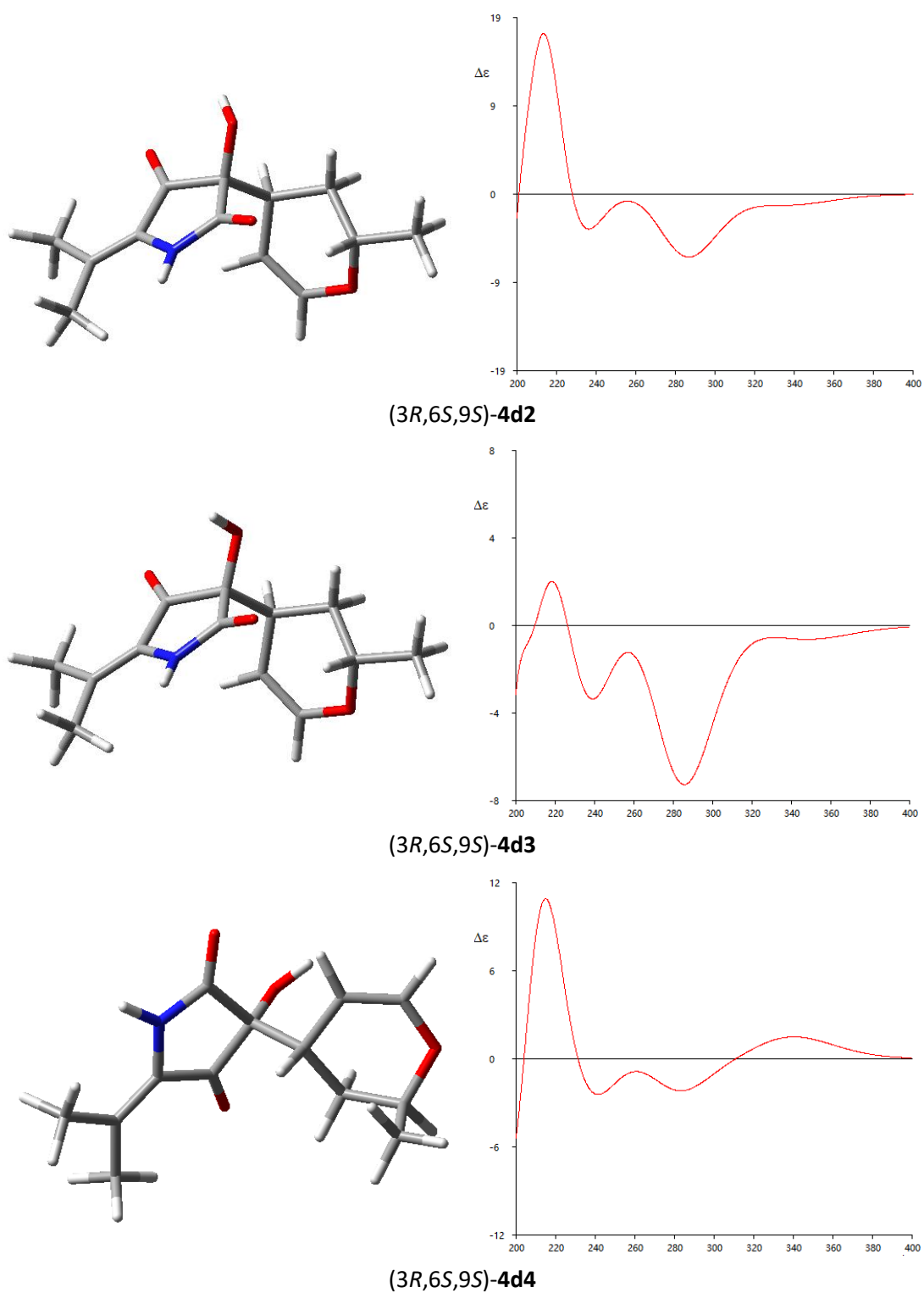


Figure S64. The optimized structures (left) and the calculated CD spectra for conformers (3R,6S,9S)-4d in MeOH at M06-2X/def2TZVP level (right). $\sigma=0.30$ eV.

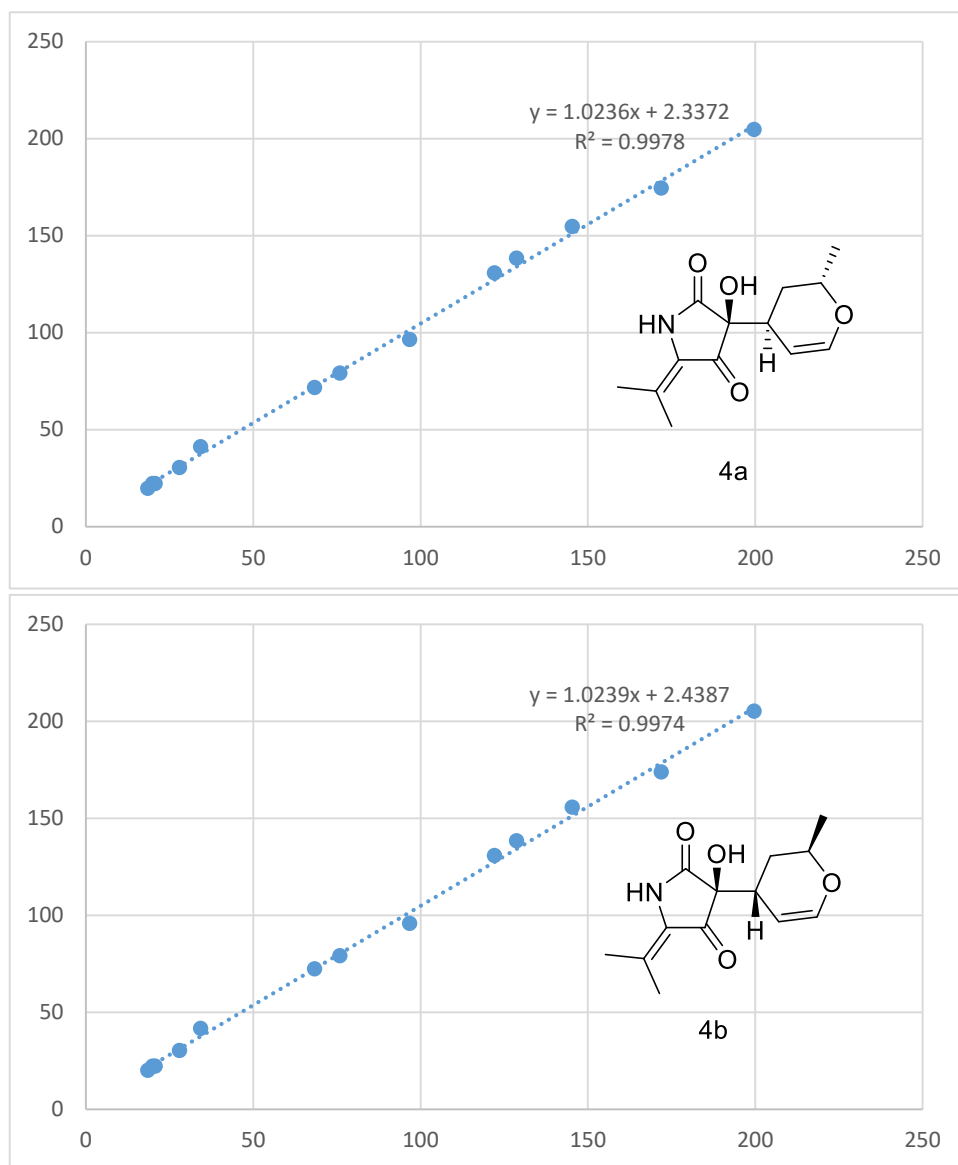


Figure S65. Correlation plots of experimental ^{13}C NMR chemical shifts versus the corresponding calculated data for (3*S*, 6*S*, 9*S*)-**4a** and (3*S*, 6*R*, 9*R*)-**4b**

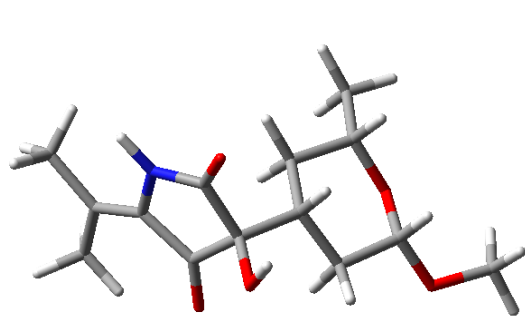
		DP4+	91.53%	8.47%
Nuclei	sp2?	Experimental	Isomer 1	Isomer 2
C	x	122.07	130.833	130.781
C	x	199.64	204.715	205.3
C		75.86	79.184	79.225
C	x	171.89	174.492	173.923
C	x	128.6	138.448	138.369
C		18.44	19.821	20.057
C		19.91	22.225	22.203
C		34.26	41.124	41.651
C	x	96.65	96.549	95.898
C	x	145.25	154.707	155.726
C		68.28	71.742	72.405
C		27.95	30.437	30.371
C		20.63	22.192	22.205

Figure S66. DP4+ probabilities (%) for conformers of (3*S*, 6*S*, 9*S*)-**4a** (isomer 1) and (3*S*, 6*R*, 9*R*)-**4b** (isomer 2).

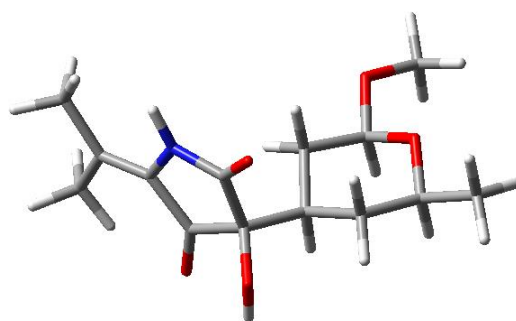
Table S10. Relative free energies^a and equilibrium populations^b of conformers for (3*S*, 6*R*, 8*S*, 9*S*)-**5a**^c

Conformer	ΔG	P (%)
5a1	0.00	24.26
5a2	0.10	20.40
5a3	0.15	18.97
5a4	0.17	18.20
5a5	0.17	18.17

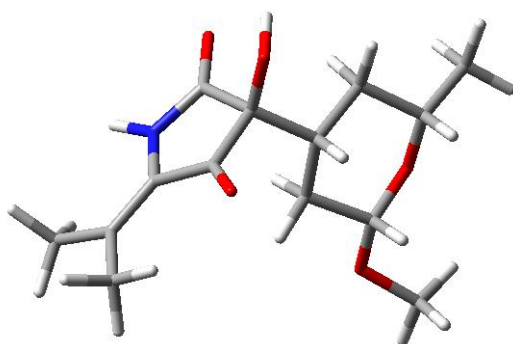
^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



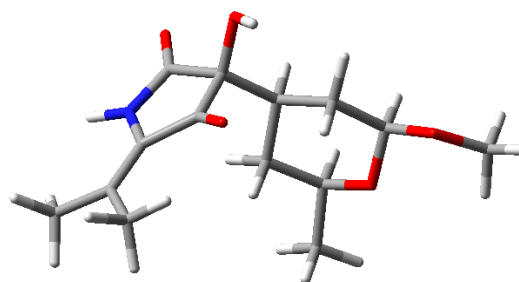
(3*S*,6*R*,8*S*,9*S*)-**5a1**



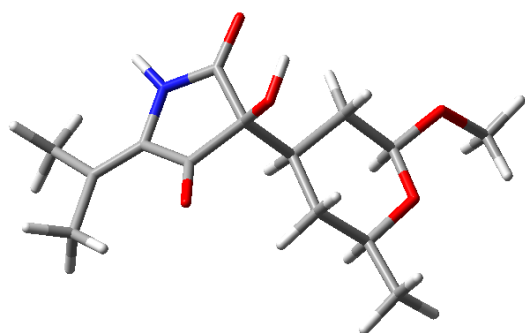
(3*S*,6*R*,8*S*,9*S*)-**5a2**



(3*S*,6*R*,8*S*,9*S*)-**5a3**



(3*S*,6*R*,8*S*,9*S*)-**5a4**



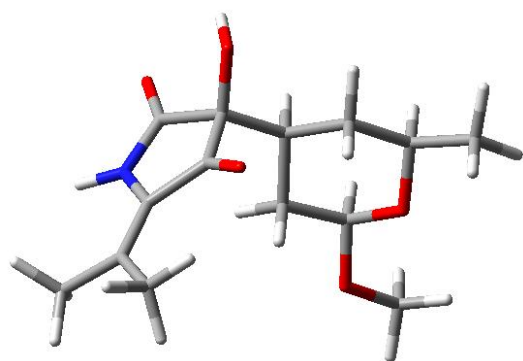
(3*S*,6*R*,8*S*,9*S*)-**5a5**

Figure S67. The optimized structure of conformers for (3*S*,6*R*,8*S*,9*S*)-**5a** in MeOH at B3LYP/6-31G (d) level

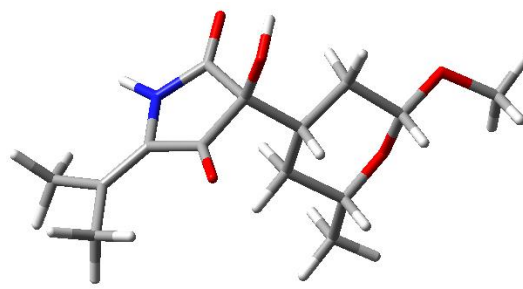
Table S11. Relative free energies^a and equilibrium populations^b of conformers for (3*S*,6*S*,8*R*,9*R*)-**5b**^c

Conformer	ΔG	P (%)
5b1	0.00	48.08
5b2	0.32	28.22
5b3	0.65	16.14
5b4	1.10	7.56

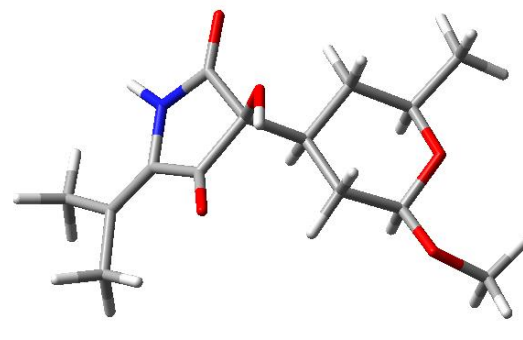
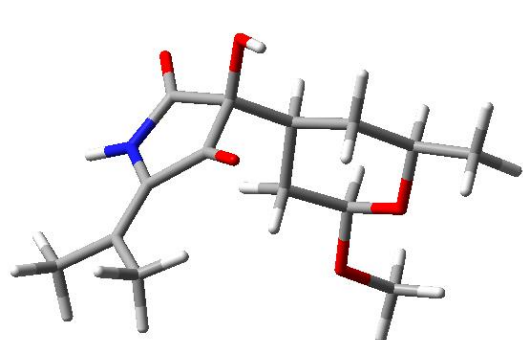
^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



(3*S*,6*S*,8*R*,9*R*)-**5b1**



(3*S*,6*S*,8*R*,9*R*)-**5b2**



(3*S*,6*S*,8*R*,9*R*)-**5b3**

(3*S*,6*S*,8*R*,9*R*)-**5b4**

Figure S68. The optimized structure of conformers for (3*S*,6*S*,8*R*,9*R*)-**5b** in MeOH at B3LYP/6-31G (d) level

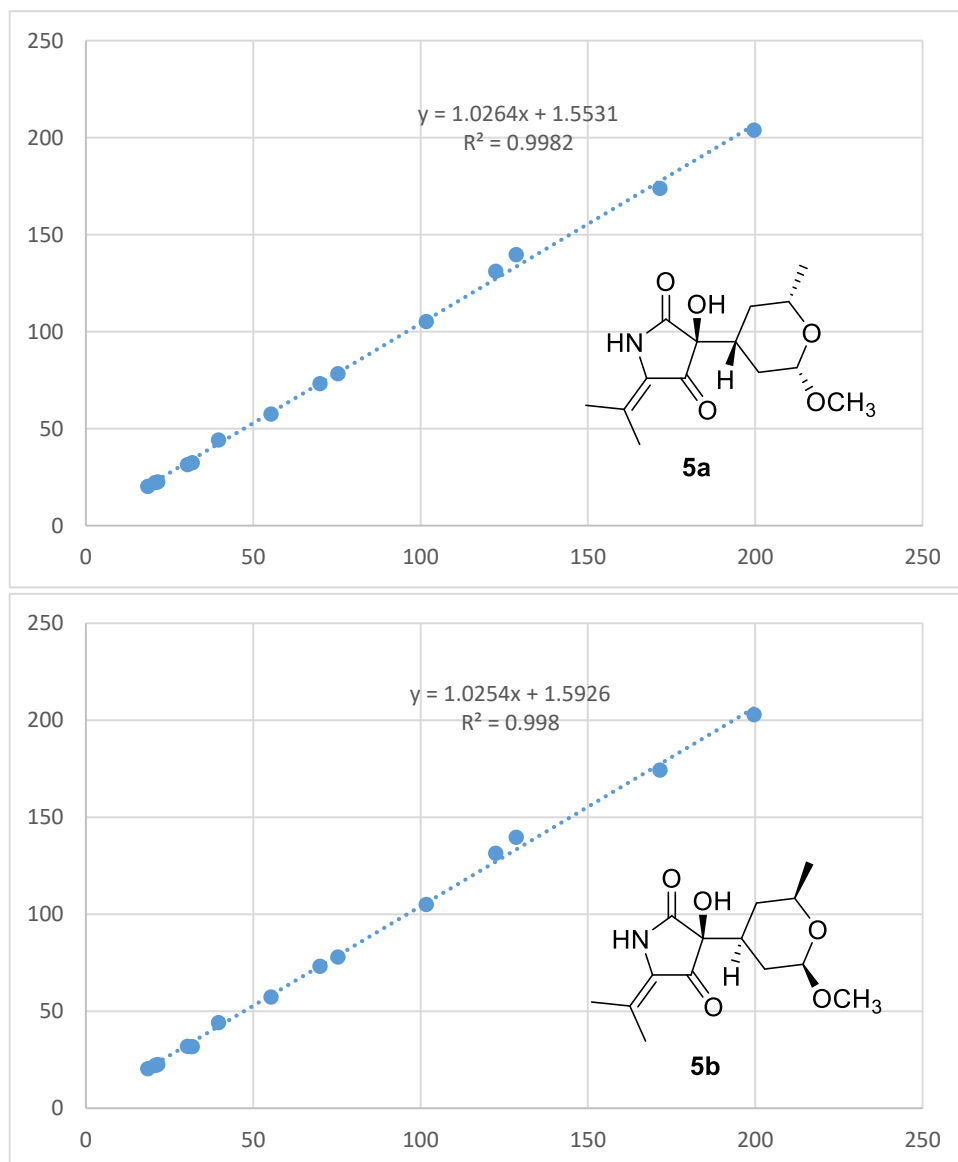


Figure S69. Correlation plots of experimental ¹³C NMR chemical shifts versus the corresponding calculated data for (3*S*,6*R*,8*S*,9*S*)-**5a** and (3*S*,6*S*,8*R*,9*R*)-**5b**.

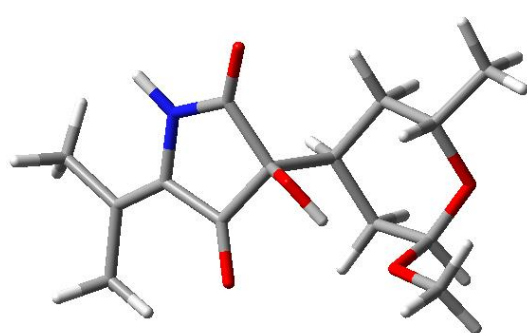
		DP4+	86.42%	13.58%
Nuclei	sp2?	experimental	Isomer 1	Isomer 2
C	x	122.44	131.238	131.53
C	x	199.59	203.942	203.056
C		75.29	78.338	78.149
C	x	171.51	173.932	174.402
C	x	128.5	139.759	139.848
C		18.46	20.382	20.534
C		20.69	22.232	22.16
C		39.6	44.243	44.288
C		30.31	31.56	31.994
C		101.66	105.23	105.176
C		69.93	73.333	73.261
C		31.72	32.473	31.823
C		21.45	22.653	22.652
C		55.32	57.549	57.509

Figure S70. DP4+ probabilities (%) for conformers of (3*S*,6*R*,8*S*,9*S*)-**5a** (isomer 1) and (3*S*,6*S*,8*R*,9*R*)-**5b** (isomer 2).

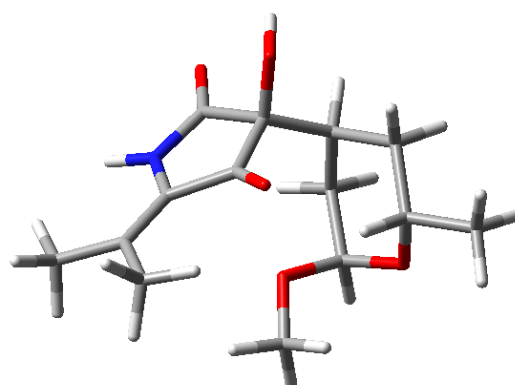
Table S12. Relative free energies^a and equilibrium populations^b of conformers for (3*S*,6*S*,8*R*,9*S*)-**6a**^c

Conformer	ΔG	P (%)
6a1	0.00	68.40
6a2	0.72	20.24
6a3	1.06	11.35

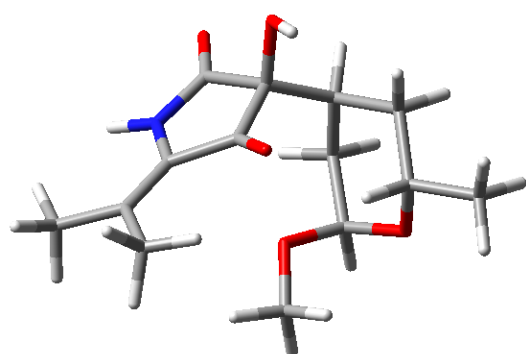
^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



(3*S*,6*S*,8*R*,9*S*)-**6a1**



(3*S*,6*S*,8*R*,9*S*)-**6a2**



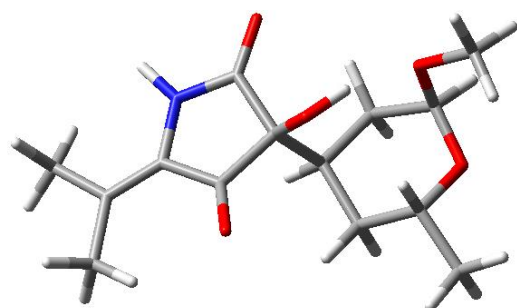
(3*S*,6*S*,8*R*,9*S*)-**6a3**

Figure S71. The optimized structure of conformers for (3*S*,6*S*,8*R*,9*S*)-**6a** in MeOH at B3LYP/6-31G (d) level.

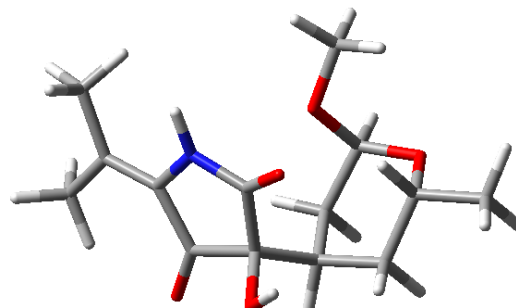
Table S13. Relative free energies^a and equilibrium populations^b of conformers for (3*S*,6*R*,8*S*,9*R*)-**6b**^c

Conformer	ΔG	P (%)
6b1	0.00	77.82
6b2	0.98	14.77
6b3	1.39	7.41

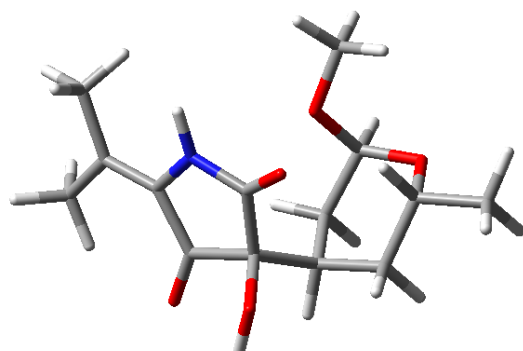
^a B3LYP/6-31G(d), in kcal/mol. ^b From ΔG values at 298.15 K. ^c in MeOH, no imaginary frequency.



(3*S*,6*R*,8*S*,9*R*)-**6b1**



(3*S*,6*R*,8*S*,9*R*)-**6b2**



(3*S*,6*R*,8*S*,9*R*)-**6b3**

Figure S72. The optimized structure of conformers for (3*S*,6*R*,8*S*,9*R*)-**6b** in MeOH at B3LYP/6-31G (d) level.

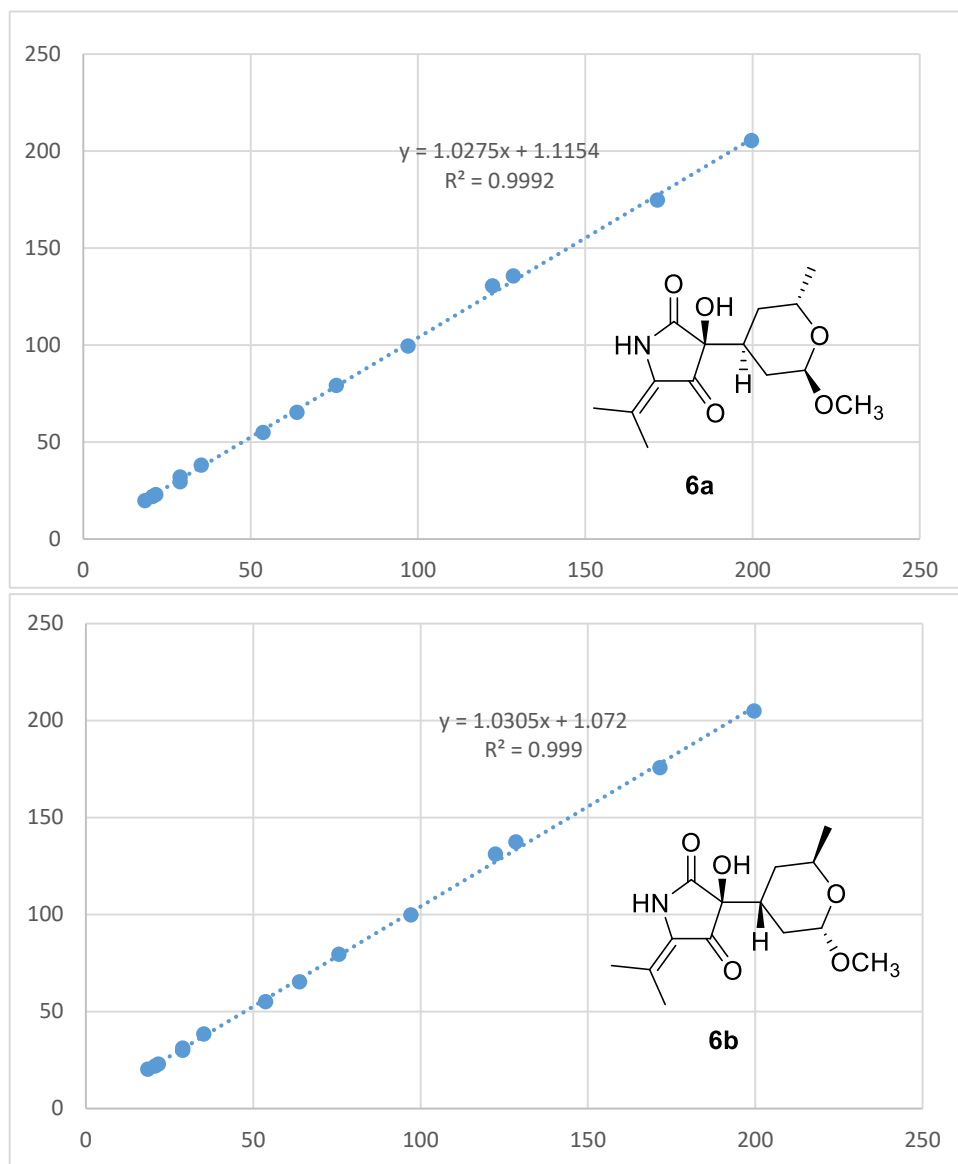


Figure S73. Correlation plots of experimental ^{13}C NMR chemical shifts versus the corresponding calculated data for (3*S*,6*S*,8*R*,9*S*)-**6a** and (3*S*,6*R*,8*S*,9*R*)-**6b**

		DP4+	86.61%	13.39%
Nuclei	sp2?	experimental	Isomer 1	Isomer 2
C	x	122.28	130.626	131.05
C	x	199.65	205.514	204.922
C		75.55	79.267	79.475
C	x	171.53	174.733	175.655
C	x	128.44	135.751	137.446
C		18.43	19.937	20.266
C		20.66	22.002	21.895
C		35.23	38.188	38.331
C		28.86	29.628	29.974
C		97.05	99.607	99.78
C		63.82	65.342	65.329
C		28.86	32.037	31.086
C		21.6	22.914	22.846
C		53.67	54.999	55.092

Figure S74. DP4+ probabilities (%) for conformers of (3*S*,6*S*,8*R*,9*S*)-**6a** (isomer 1) and (3*S*,6*R*,8*S*,9*R*)-**6b** (isomer 2).

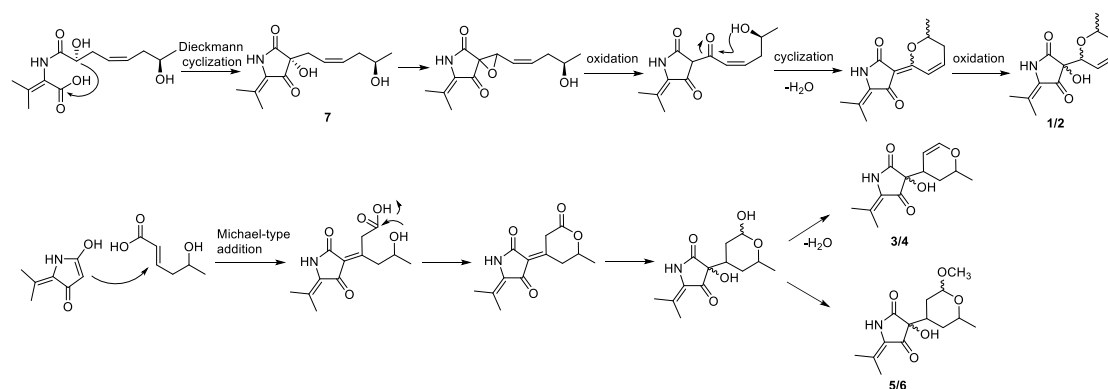


Figure S74. Proposed biosynthetic pathway of compounds **1–7** [1,2].

[1] Rischer, M.; Lee, S. R.; Eom, H. J.; Park, H. B.; Vollmer, J.; Kaster, A. K.; Shin, Y. H.; Oh, D. C.; Kim, K. H.; Beemelmans, C. Spirocyclic cladosporicin A and cladosporiumins I and J from a Hydractinia-associated *Cladosporium sphaerospermum* SW67. *Org. Chem. Front.* **2019**, *6*, 1084-1093.

[2] Mo, X.; Gulder, T. Biosynthetic strategies for tetramic acid formation. *Nat. Prod. Rep.* **2021**, *38*, 1555-1566.