

Discovery of GOT1 Inhibitors from a Marine-derived *Aspergillus terreus* that Against Pancreatic Ductal Adenocarcinoma

Shan Yan,^{1,†} Changxing Qi,^{1,†} Wei Song,² Qianqian Xu,¹ Weiguang Sun,^{1,*} Yonghui Zhang^{1,*}

¹Hubei Key Laboratory of Natural Medicinal Chemistry and Resource Evaluation, School of Pharmacy, Tongji

Medical College, Huazhong University of Science and Technology, Wuhan 430030, People's Republic of China;

yanshanwhu@163.com (S.Y.); qichangxing@hust.edu.cn (C.Q.); Xuxuqq@hust.edu.cn (Q.X.).

²Guangdong Provincial Key Laboratory of Microbial Culture Collection and Application, State Key Laboratory of

Applied Microbiology Southern China, Institute of Microbiology, Guangdong Academy of Sciences, Guangzhou

510000, China; 20191103@zcmu.edu.cn (W.S.).

*Correspondence: zhangyh@mails.tjmu.edu.cn (Y.Z.); zhuhucheng@hust.edu.cn (H.Z.).

†S.Y. and C.Q. contributed equally to this work.

ITS sequence of the fungus

CTTCCGGTAGGGTGAACCTCGGAAAGGATCATTACCGAGTGCGGGTCTT
TATGGCCAACCTCCCACCGTGACTATTGTACCTTGTGCTTCGGCGGCC
CGCCAGCGTTGCTGGCCGCCGGGGCGACTCGCCCCCGGGCCGTGCC
GCCGGAGACCCAACATGAACCTGTTCTGAAAGCTGCAGTCTGAGTGT
GATTCTTGCAATCAGTAAAATTCACAACATGGATCTTGGTCCGGCA
TCGATGAAGAACGCGAGCAAATGCGATAACTAATGTGAATTGCAGAATTCA
GTGAATCATCGAGTCTTGAACGCACATTGCGCCCCCTGGTATTCCGGGGG
GCATGCCTGTCCGAGCGTCATTGCTGCCCTCAAGCCCGGCTGTGTTGG
GCCCTCGTCCCCCGGCTCCGGGGACGGGCCGAAAGGCAGCGGCC
ACCGCGTCCGGTCTCGAGCGTATGGGCTTCGTCTCCGCTCCGTAGGCC
CGGCCGGCGCCCGCCGACGCATTATTGCAACTTGTGTTCCAGGTTGA
CCTCGGATCAGGTAGGGATAACCGCTGAACTTAACATATCAATAAGCCGG
GAGGAAG

The optical rotation values for compounds 1–18.

1	$[\alpha]_D^{25} : -33.5$ (<i>c</i> 0.39, MeOH)	10	$[\alpha]_D^{25} : +129.3$ (<i>c</i> 0.67, MeOH)
2	$[\alpha]_D^{25} : -46.1$ (<i>c</i> 0.43, MeOH)	11	$[\alpha]_D^{25} : +89.1$ (<i>c</i> 0.24, MeOH)
3	$[\alpha]_D^{25} : -31.1$ (<i>c</i> 0.33, MeOH)	12	$[\alpha]_D^{25} : +31.3$ (<i>c</i> 0.17, MeOH)
4	$[\alpha]_D^{25} : +156.1$ (<i>c</i> 0.15, MeOH)	13	$[\alpha]_D^{25} : +112.3$ (<i>c</i> 0.35, MeOH)
5	$[\alpha]_D^{25} : +179.5$ (<i>c</i> 0.16, MeOH)	14	$[\alpha]_D^{25} : +31.2$ (<i>c</i> 0.21, MeOH)
6	$[\alpha]_D^{25} : 0$ (<i>c</i> 0.15, MeOH) Plane structure without optical rotation value	15	$[\alpha]_D^{25} : +56.8$ (<i>c</i> 0.18, MeOH)
7	$[\alpha]_D^{25} : +17.9$ (<i>c</i> 0.13, MeOH)	16	$[\alpha]_D^{25} : +21.1$ (<i>c</i> 0.12, MeOH)
8	$[\alpha]_D^{25} : +115.3$ (<i>c</i> 0.51, MeOH)	17	$[\alpha]_D^{25} : +51.7$ (<i>c</i> 0.24, MeOH)
9	$[\alpha]_D^{25} : +25.1$ (<i>c</i> 0.17, MeOH)	18	$[\alpha]_D^{25} : +112.3$ (<i>c</i> 0.81, MeOH)

Figure S1. (A) Position of AH (left) and PLP (right) into the active site of GOT1. (B) Surface representation of GOT1 around the active site pocket. (C) Overlay of GOT1-AH (cyan) and WT GOT1 (magenta) crystal structures. (D) Comparison of the center conformations of core molecules in chain A of GOT1-AH (cyan) and wild-type GOT1 (gray).

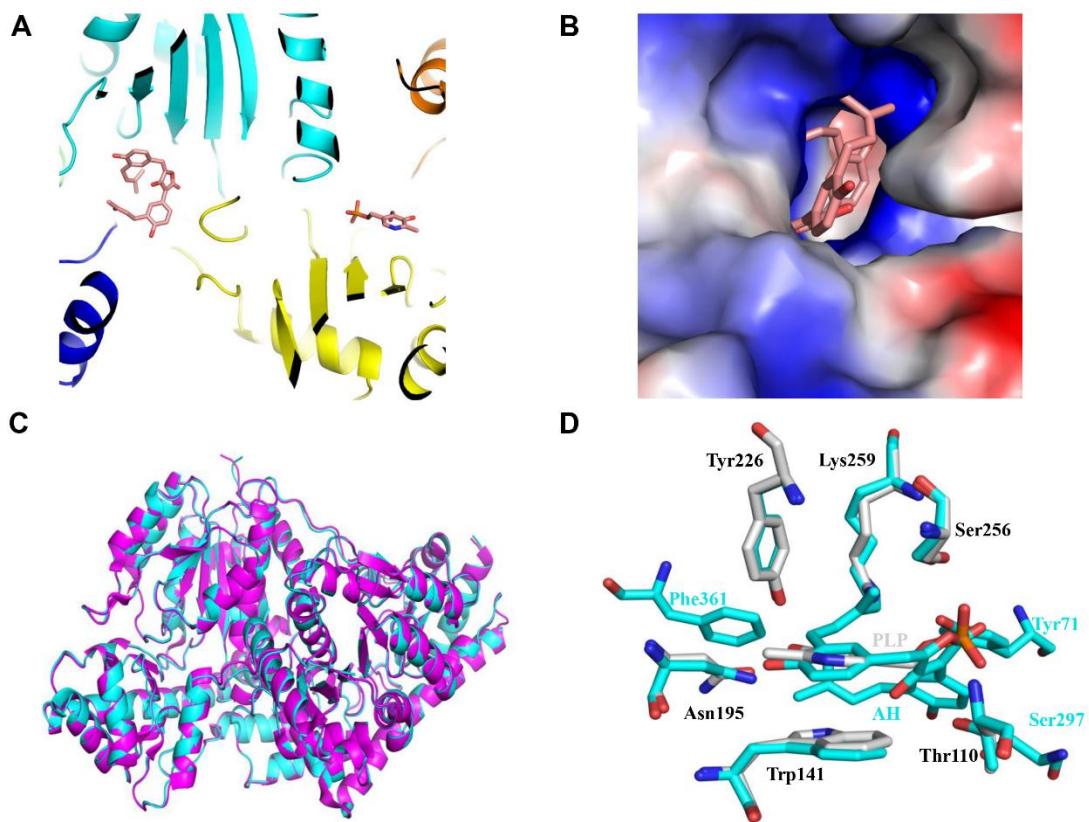


Figure S2. The 2Fo-Fc electron density of PLP bound to chain B is shown as a gray mesh. The map is contoured at the 1σ level.

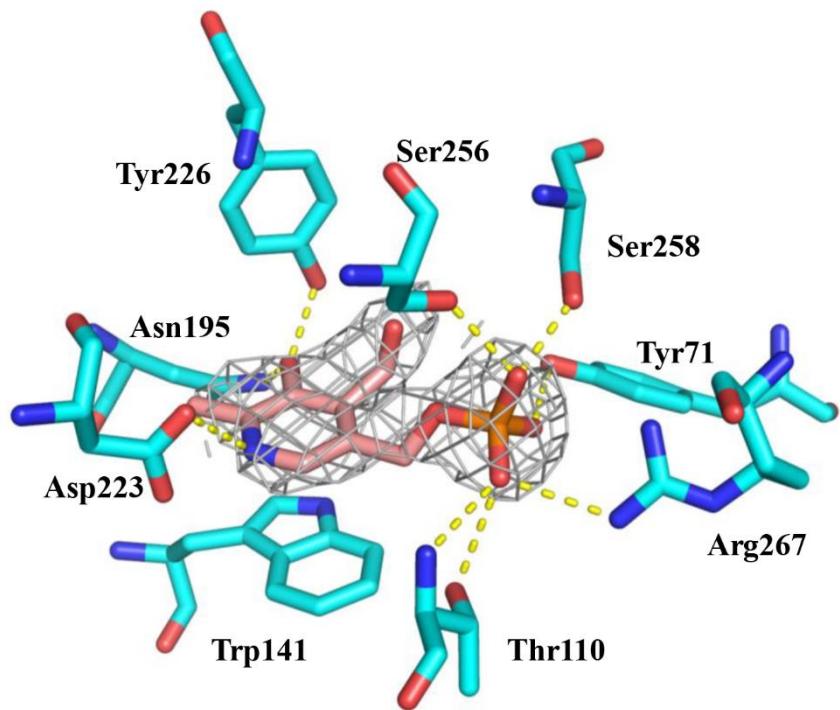


Table S1. Data collection and refinement statistics.

parameter	Enzyme in complex with AH
Resolution (Å)	49.58–2.62 (2.714–2.62)
Space group	P 1 2 ₁ 1
Unit cell dimensions	a=64.87 Å, b=90.42 Å, c=74.05 Å; α=90.0°, β=91.87°, γ=90.0°
Unique reflections	25267 (2557)
Multiplicity	2.0 (2.0)
Completeness (%)	97.67 (97.93)
$I/\sigma(I)$	9.34 (2.99)
$R\text{-merge}^{\dagger}$ (%)	0.07031 (0.2735)
$R\text{-work}$	0.2127 (0.2690)
$R\text{-free}$	0.2661 (0.3291)
Number of non-hydrogen atoms	6,703
Protein residues	820
r.m.s.d. from ideal geometry	
Bonds lengths (Å)	0.002
Angles (°)	0.46
Ramachandran plot (%)	
Favored	96.94
Outliers	0.12
Clashscore	4.51
Average B -factor	29.95
PDB code	6LIG

[†] $R\text{-merge} = \frac{\sum_{hkl} \sum_i |I_i(hkl) - \langle I(hkl) \rangle|}{\sum_{hkl} \sum_i I_i(hkl)}$, where $I(hkl)$ is the intensity of reflection hkl , Σ_{hkl} is the sum over all reflections and Σ_i is the sum over i measurements of reflection hkl . Statistics for the highest-resolution shell are shown in parentheses.

Figure S3. ^1H NMR spectrum of **1**.

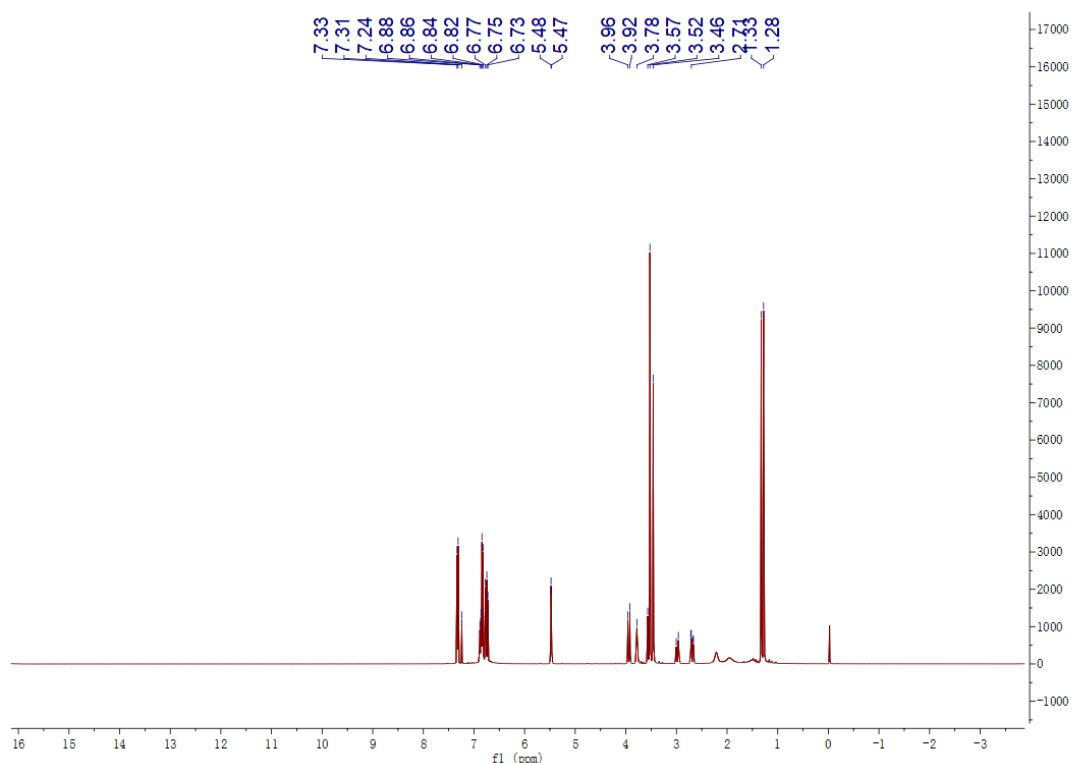


Figure S4. ^{13}C NMR spectrum of **1**.

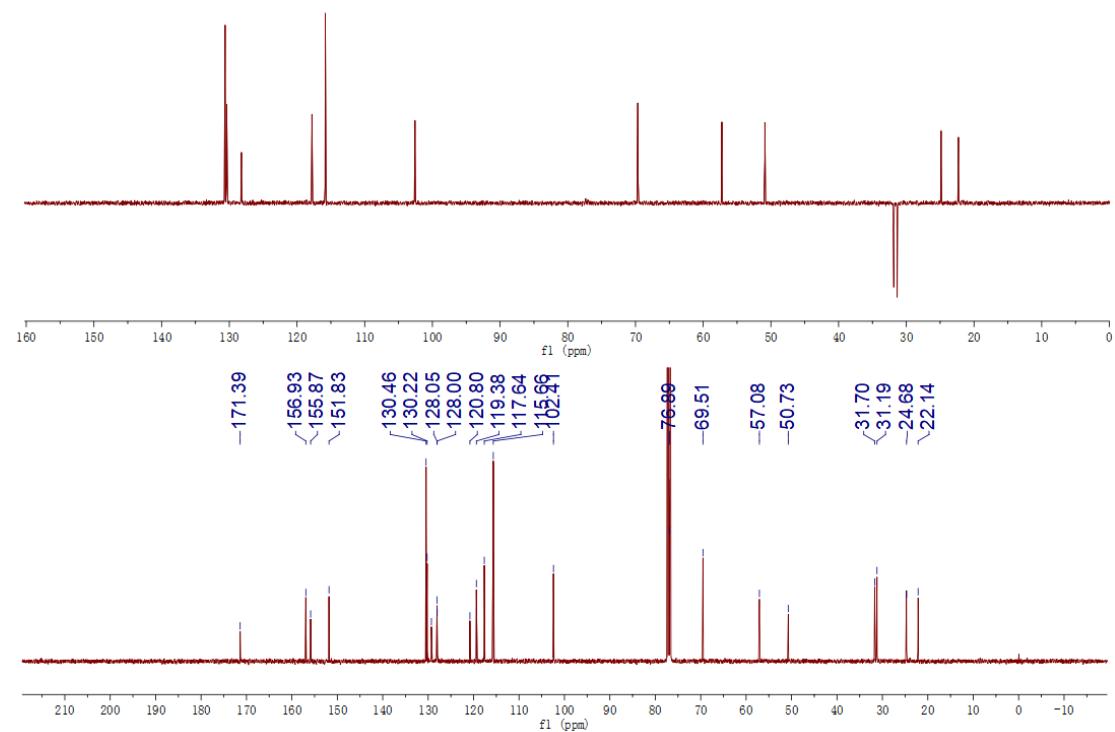


Figure S5. ^1H NMR spectrum of **2**.

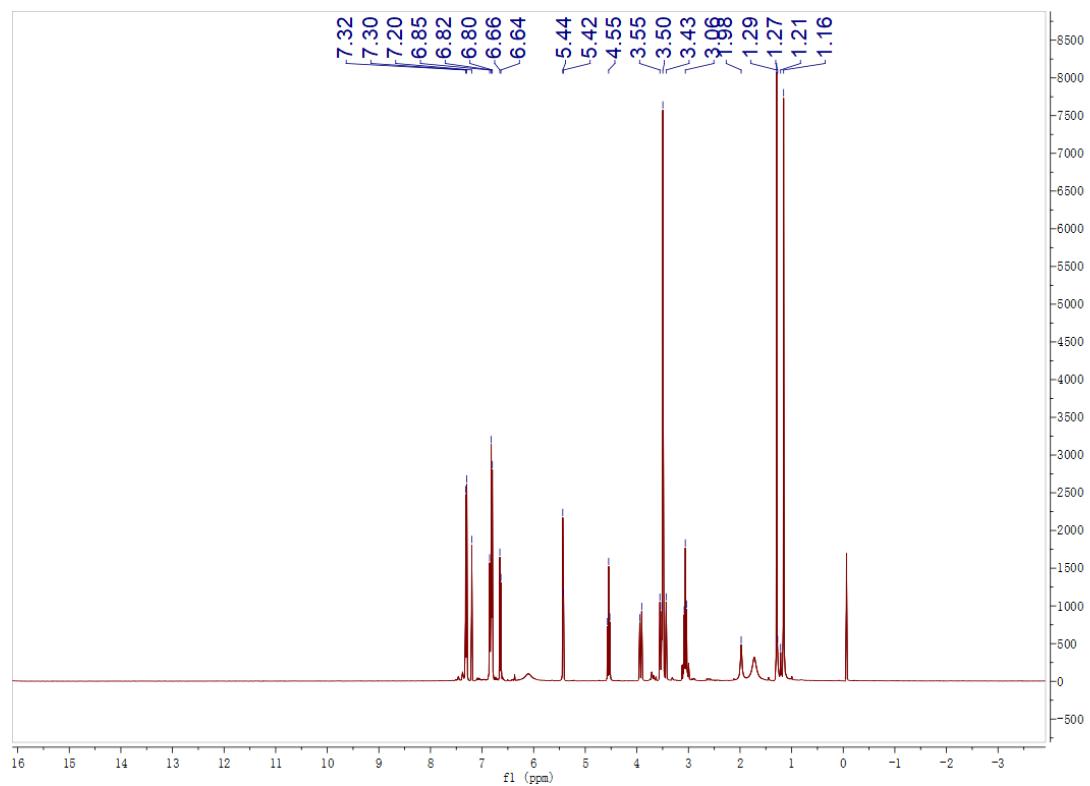


Figure S6. ^{13}C NMR spectrum of **2**.

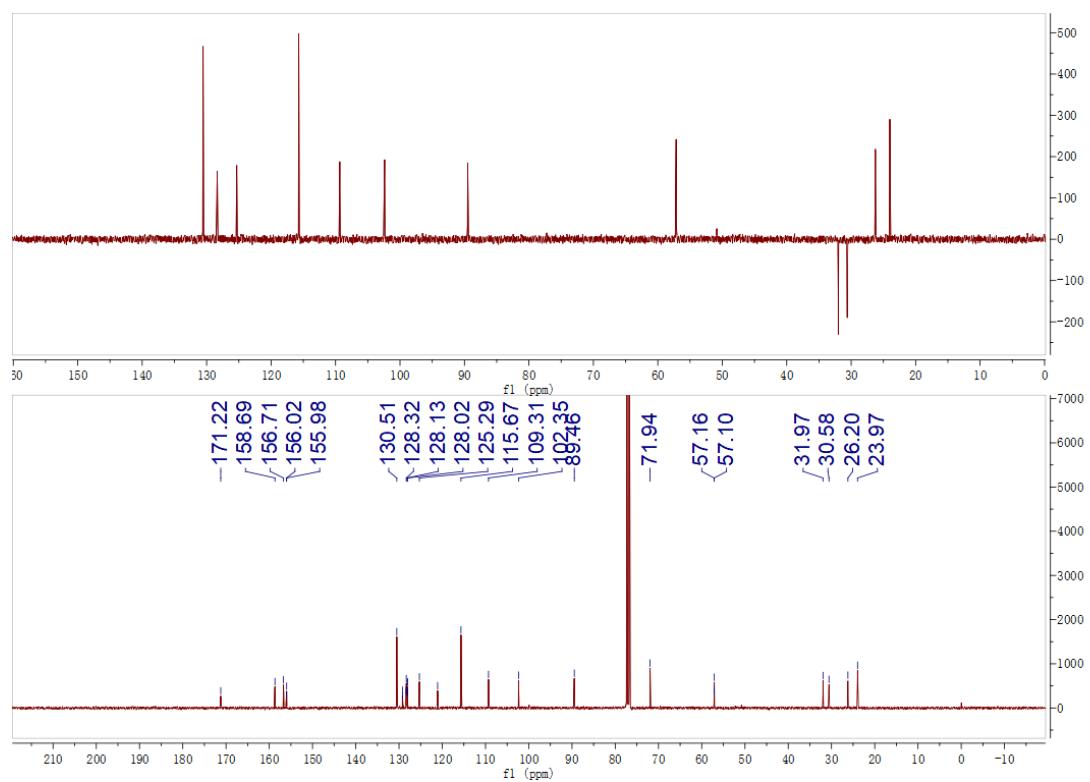


Figure S7. ^1H NMR spectrum of **3**.

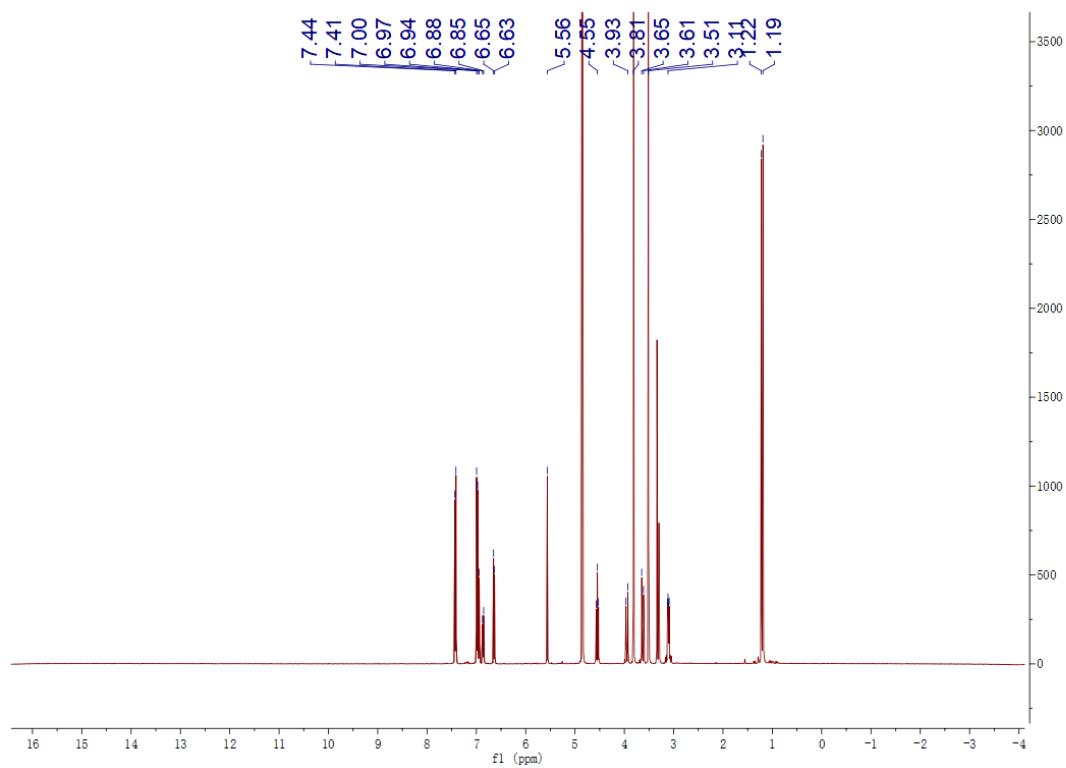


Figure S8. ^{13}C NMR spectrum of **3**.

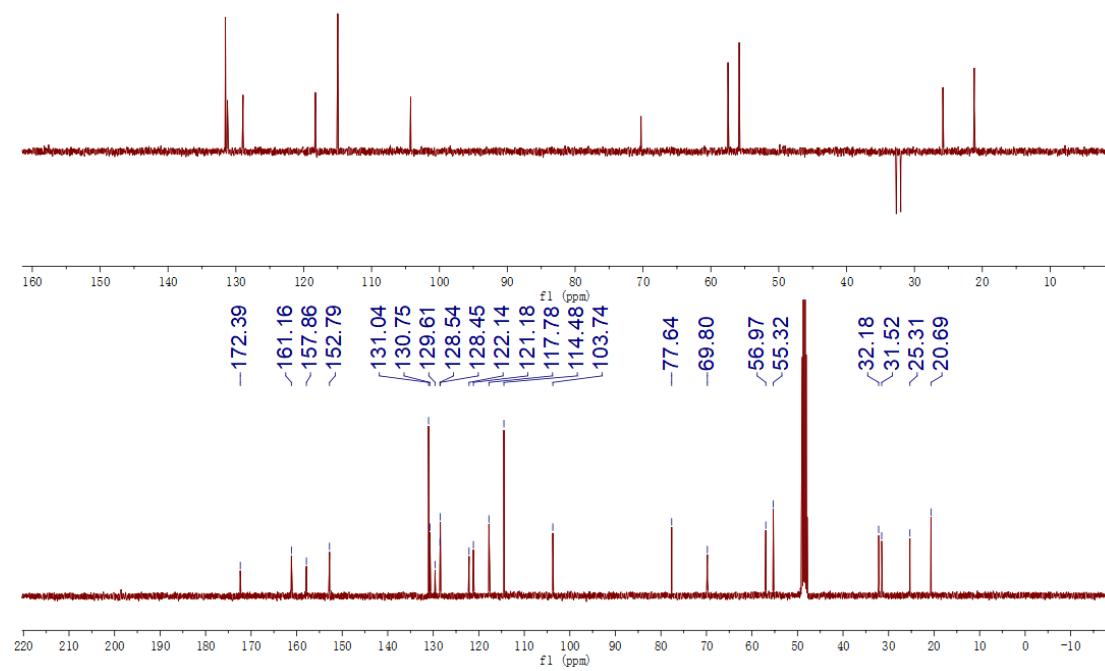


Figure S9. ^1H NMR spectrum of **4**.

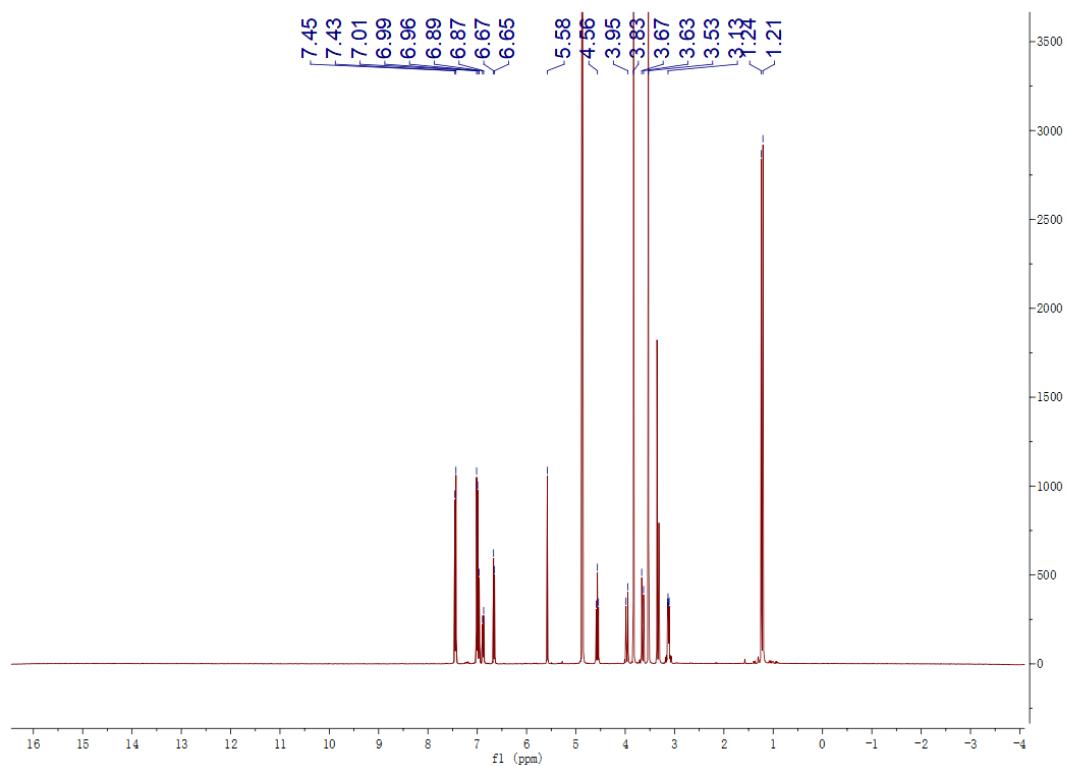


Figure S10. ^{13}C NMR spectrum of **4**.

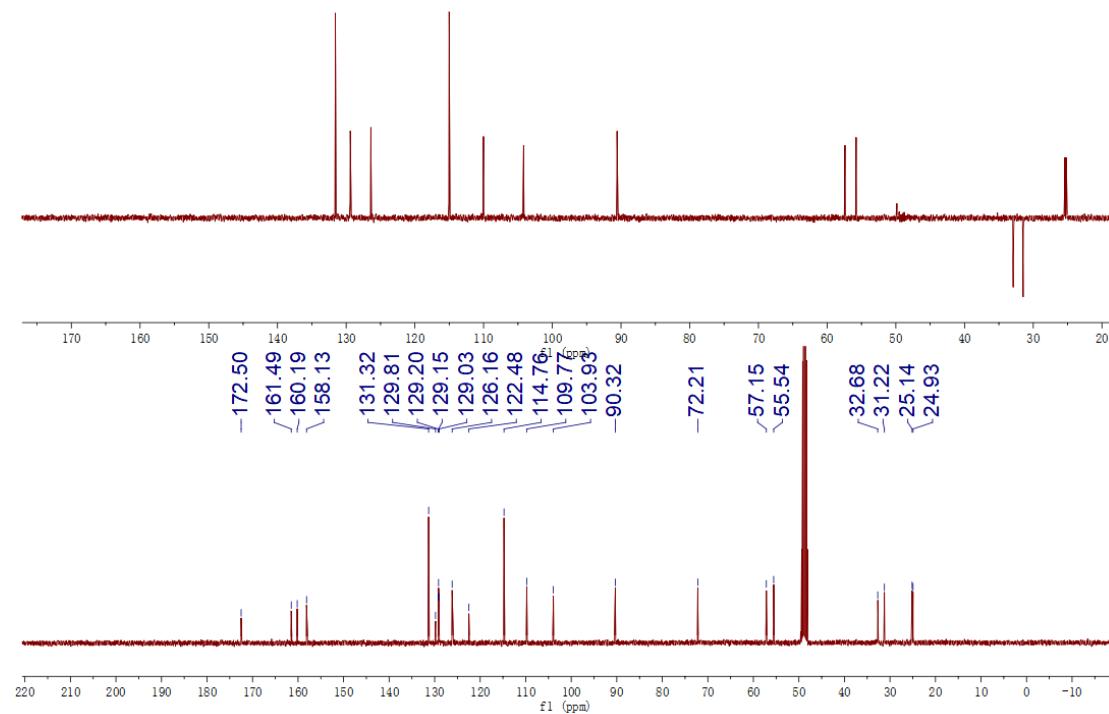


Figure S11. ^1H NMR spectrum of **5**.

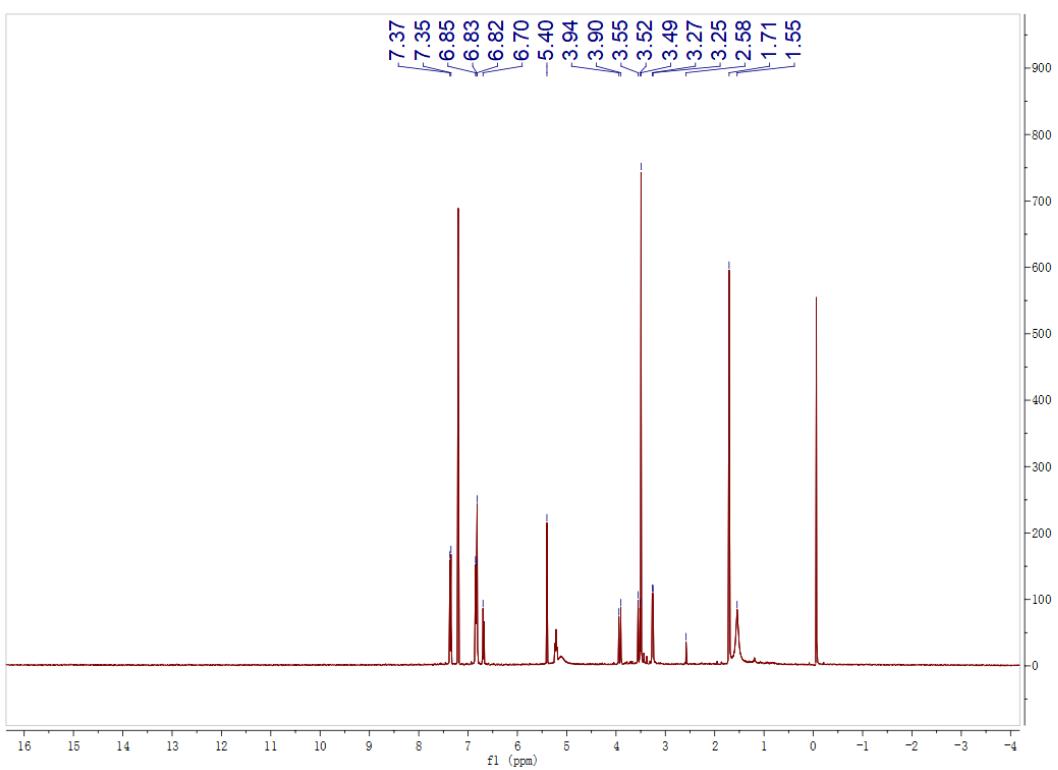


Figure S12. ^{13}C NMR spectrum of **5**.

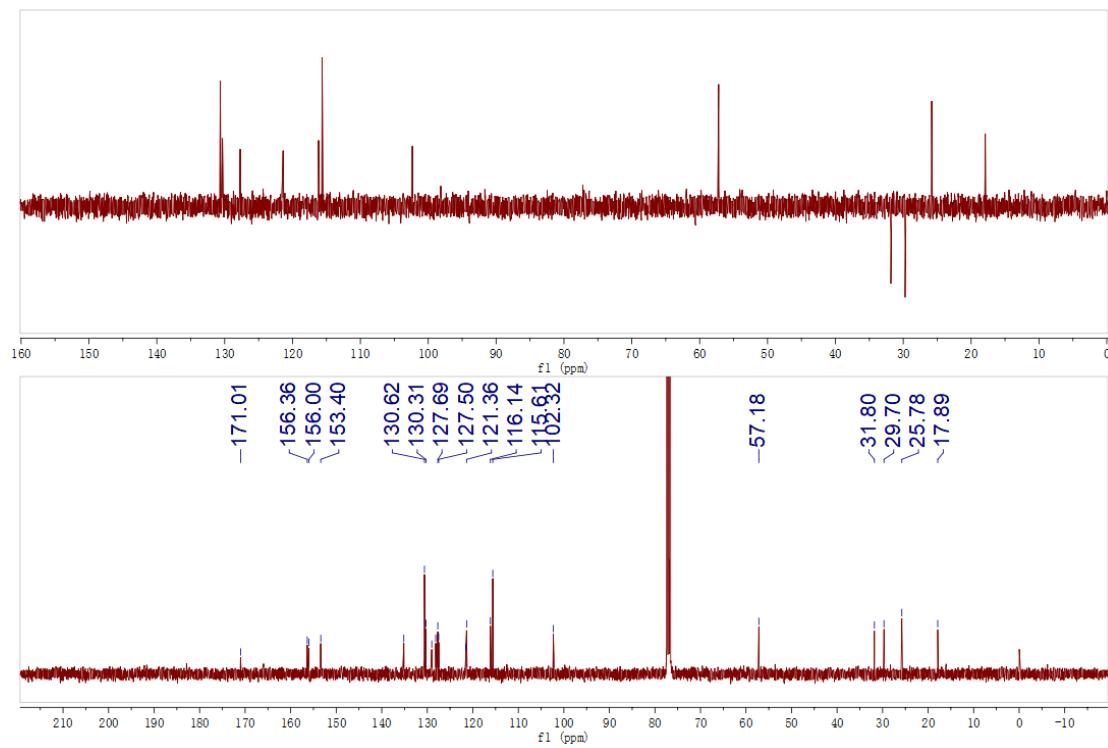


Figure S13. ^1H NMR spectrum of **6**.

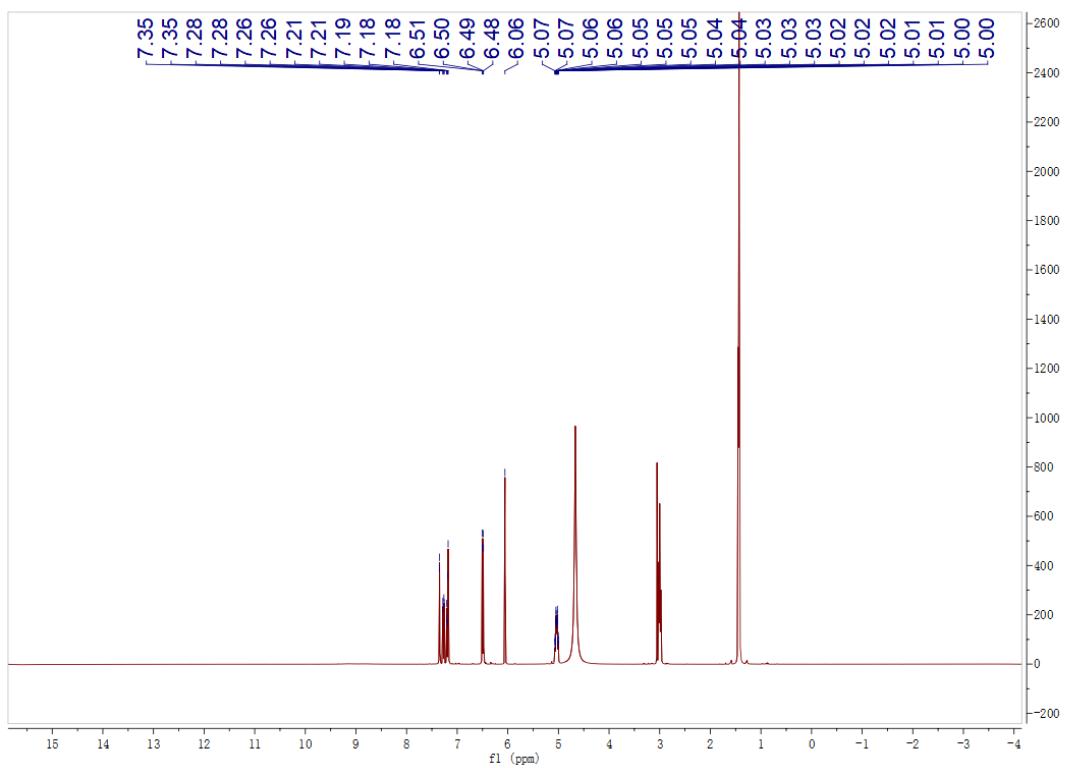


Figure S14. ^{13}C NMR spectrum of **6**.

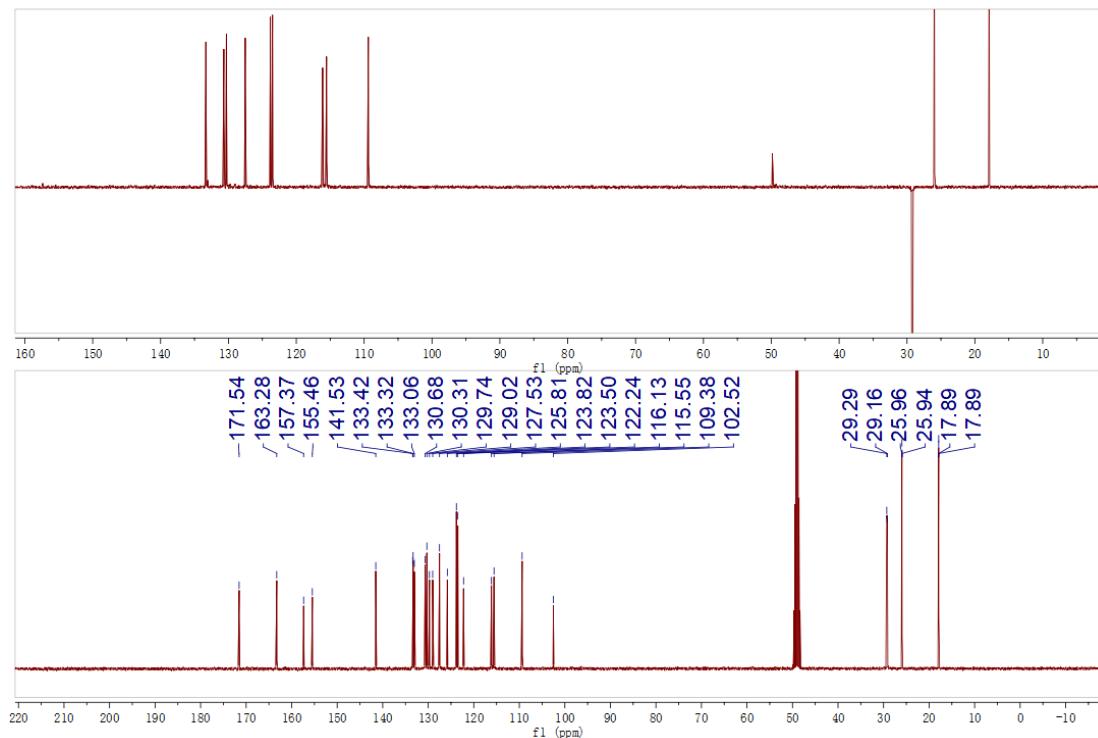


Figure S15. ^1H NMR spectrum of **7**.

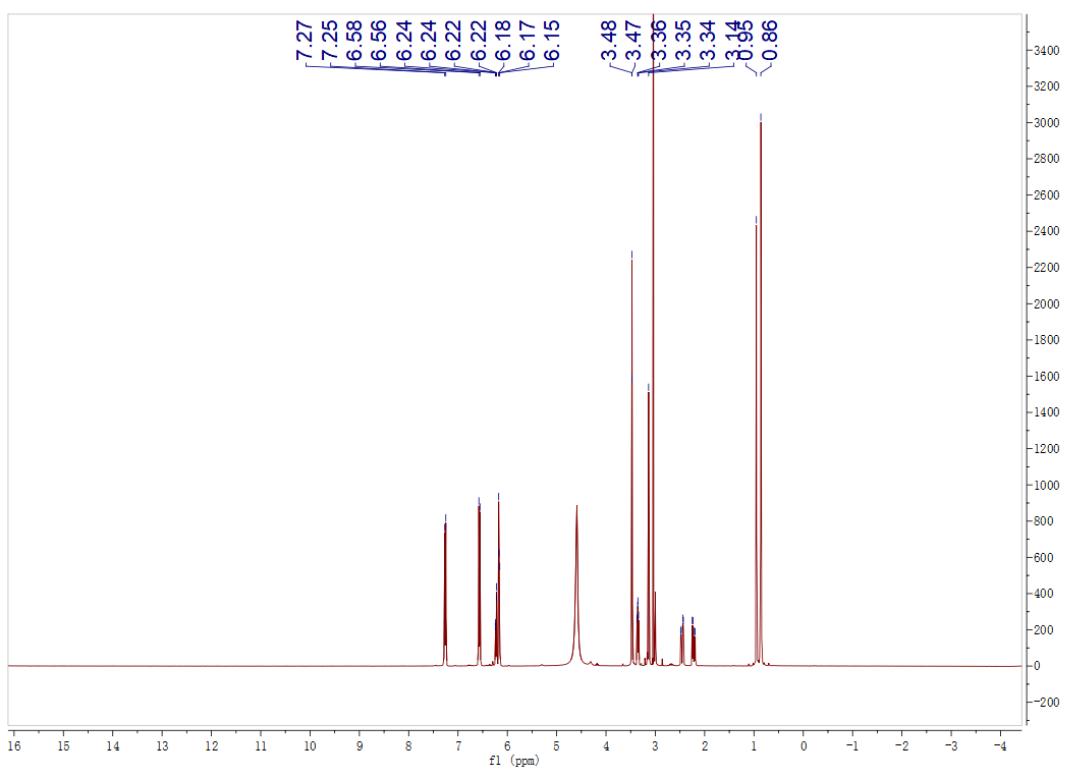


Figure S16. ^{13}C NMR spectrum of **7**.

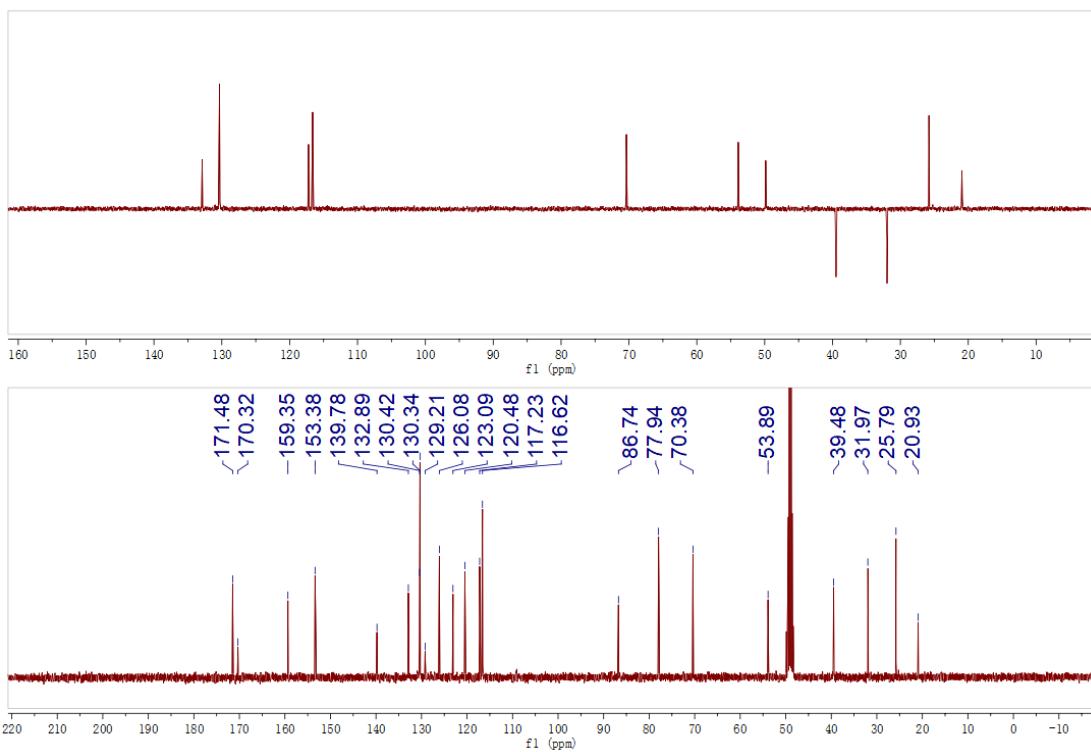


Figure S17. ^1H NMR spectrum of **8**.

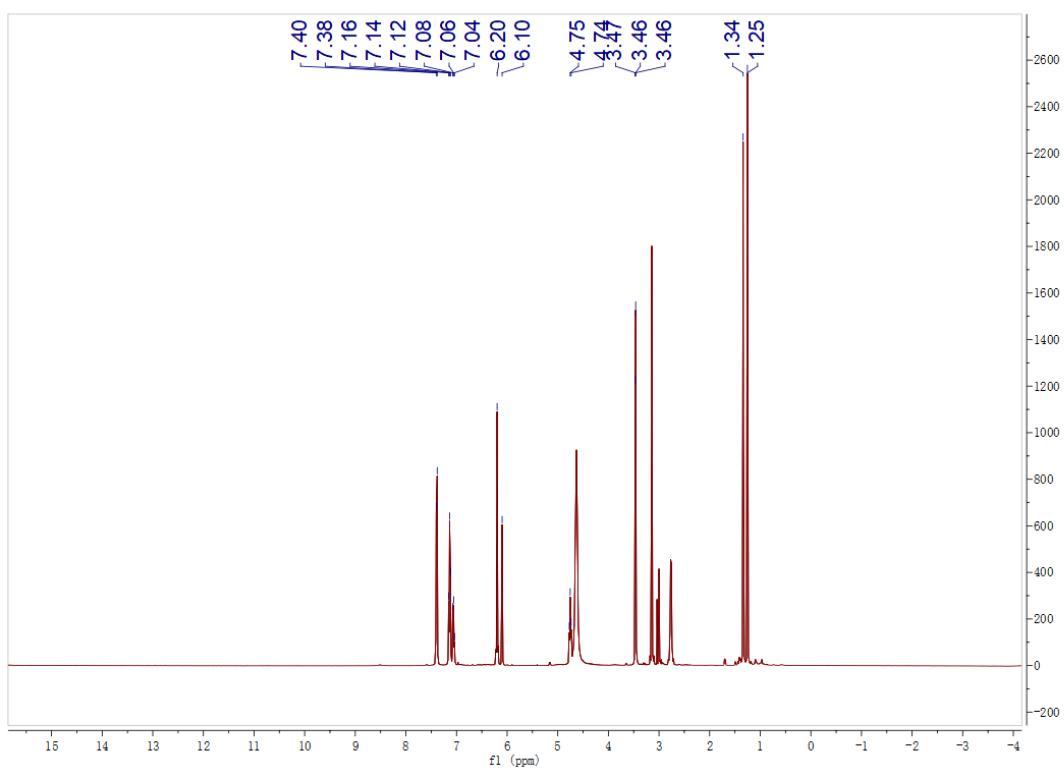


Figure S18. ^{13}C NMR spectrum of **8**.

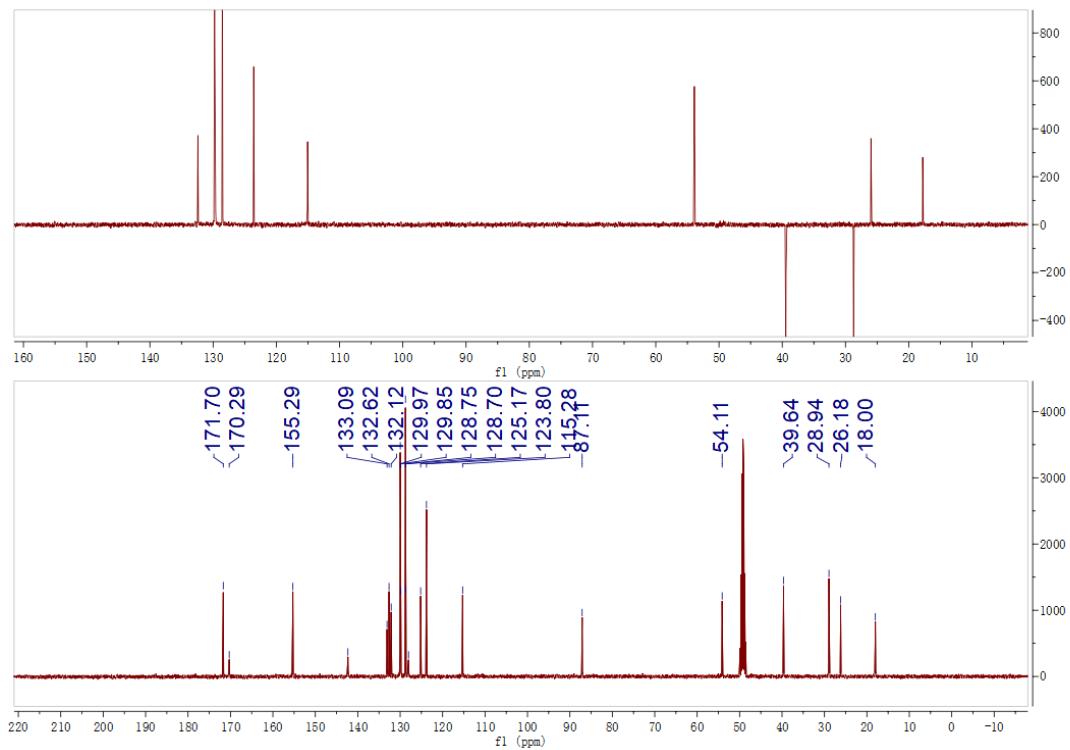


Figure S19. ^1H NMR spectrum of **9**.

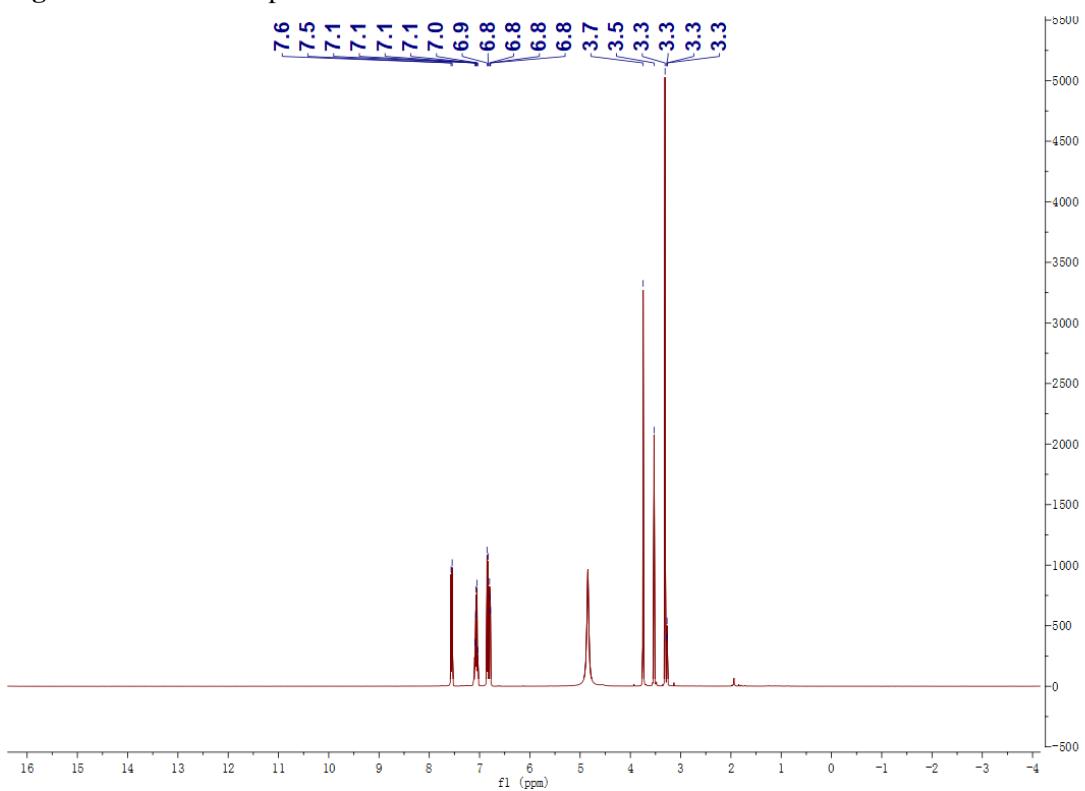


Figure S20. ^{13}C NMR spectrum of **9**.

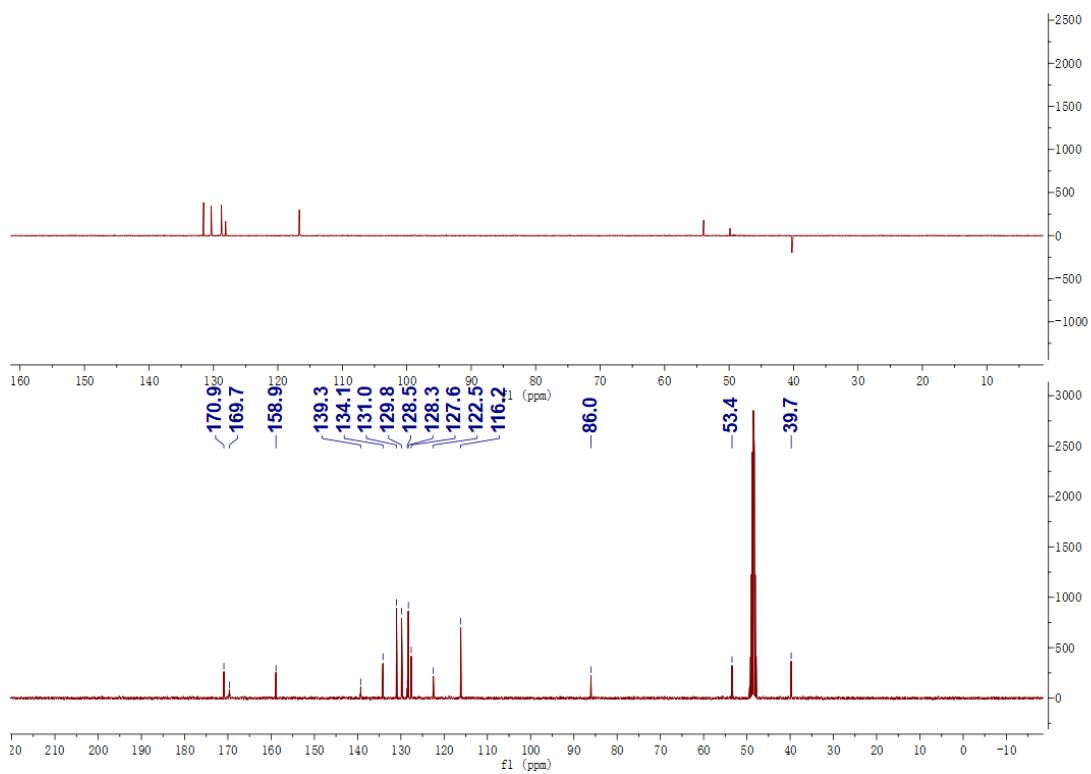


Figure S21. ^1H NMR spectrum of **10**.

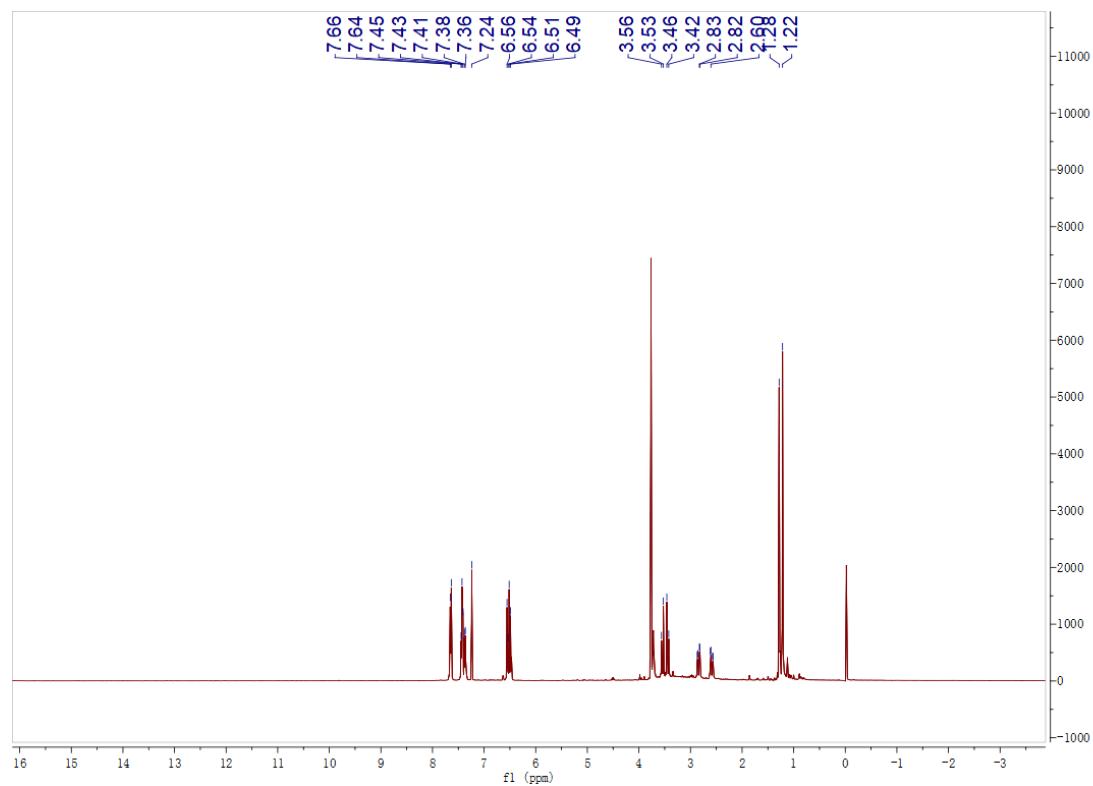


Figure S22. ^{13}C NMR spectrum of **10**.

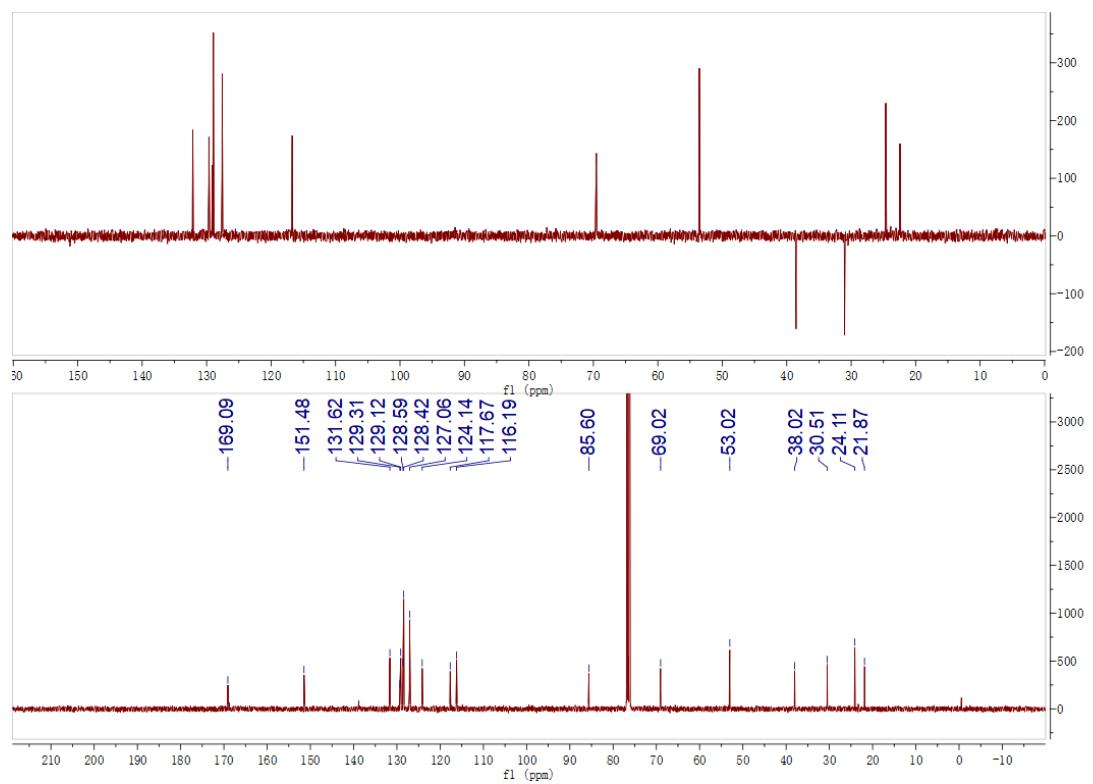


Figure S23. ^1H NMR spectrum of **11**.

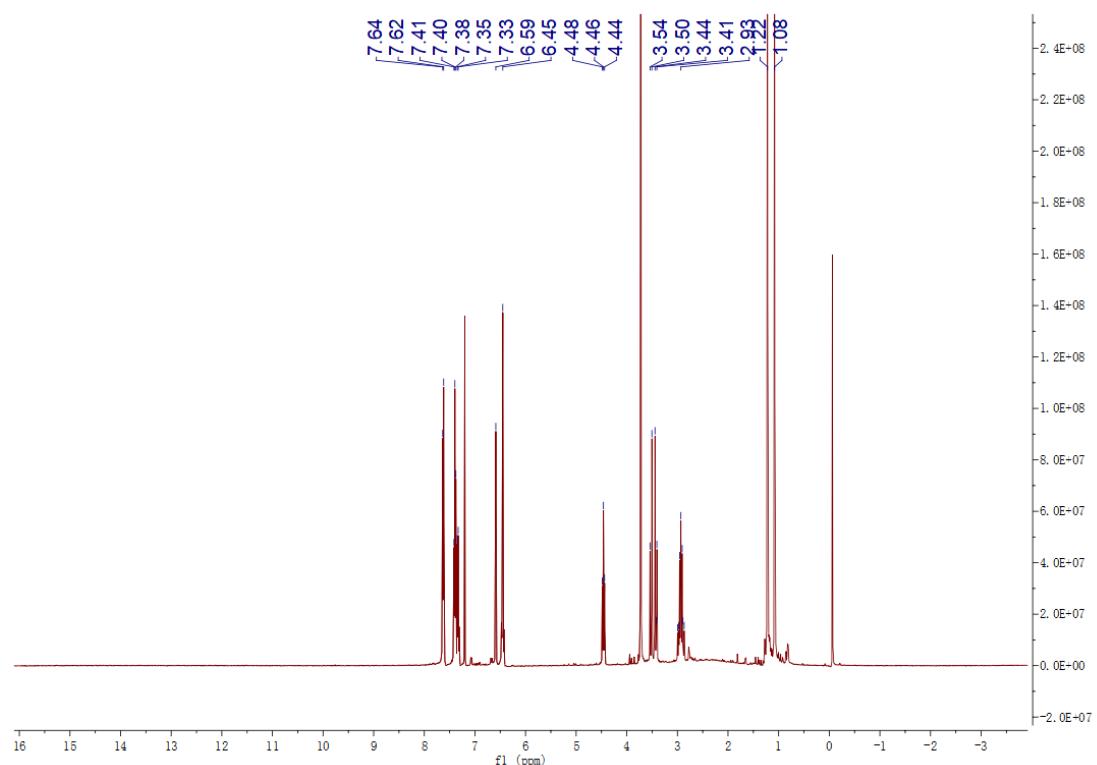


Figure S24. ^{13}C NMR spectrum of **11**.

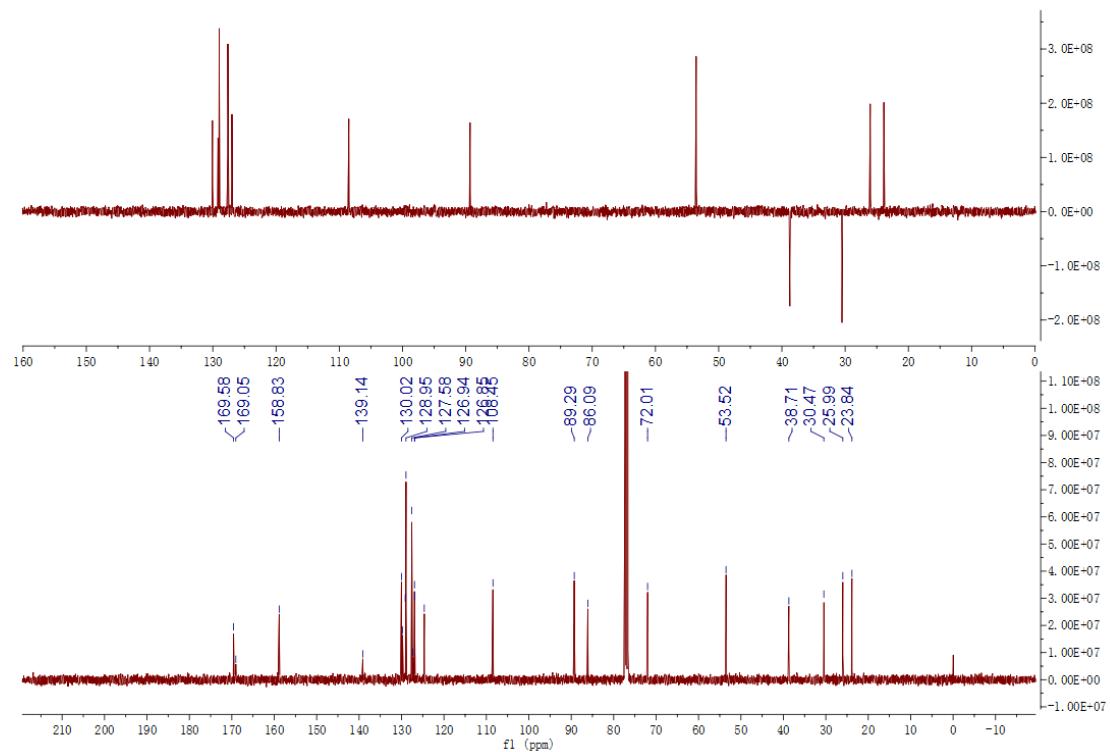


Figure S25. ^1H NMR spectrum of **12**.

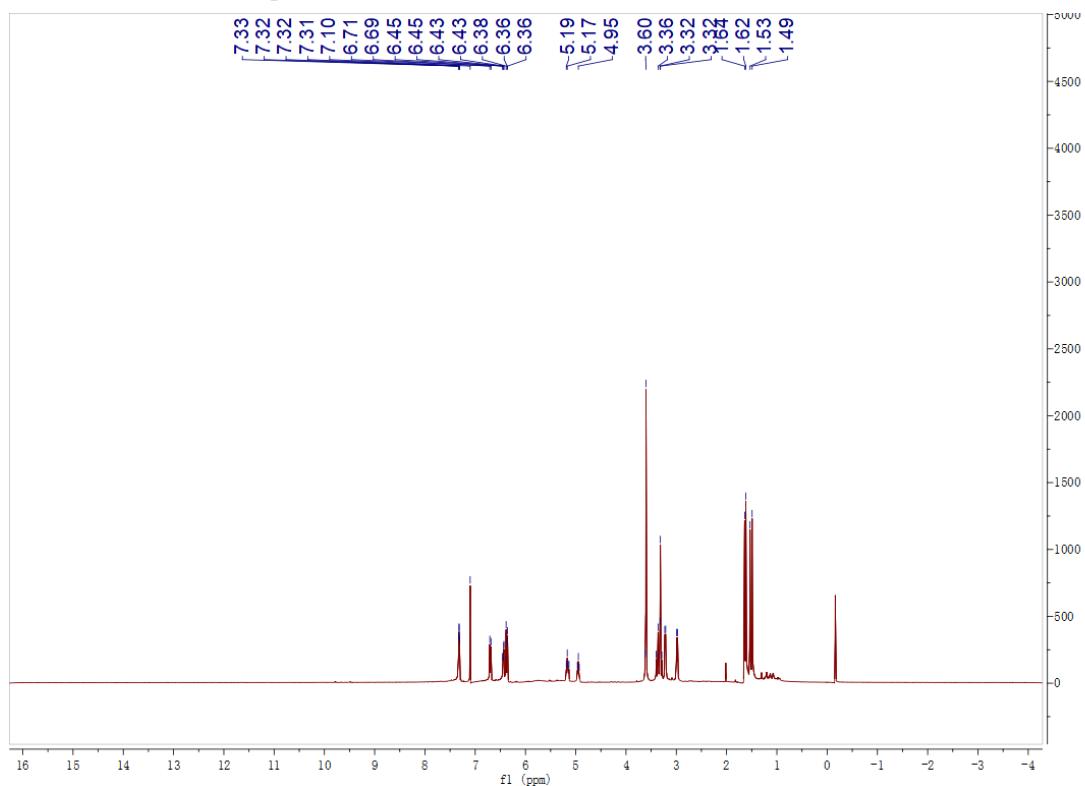


Figure S26. ^{13}C NMR spectrum of **12**.

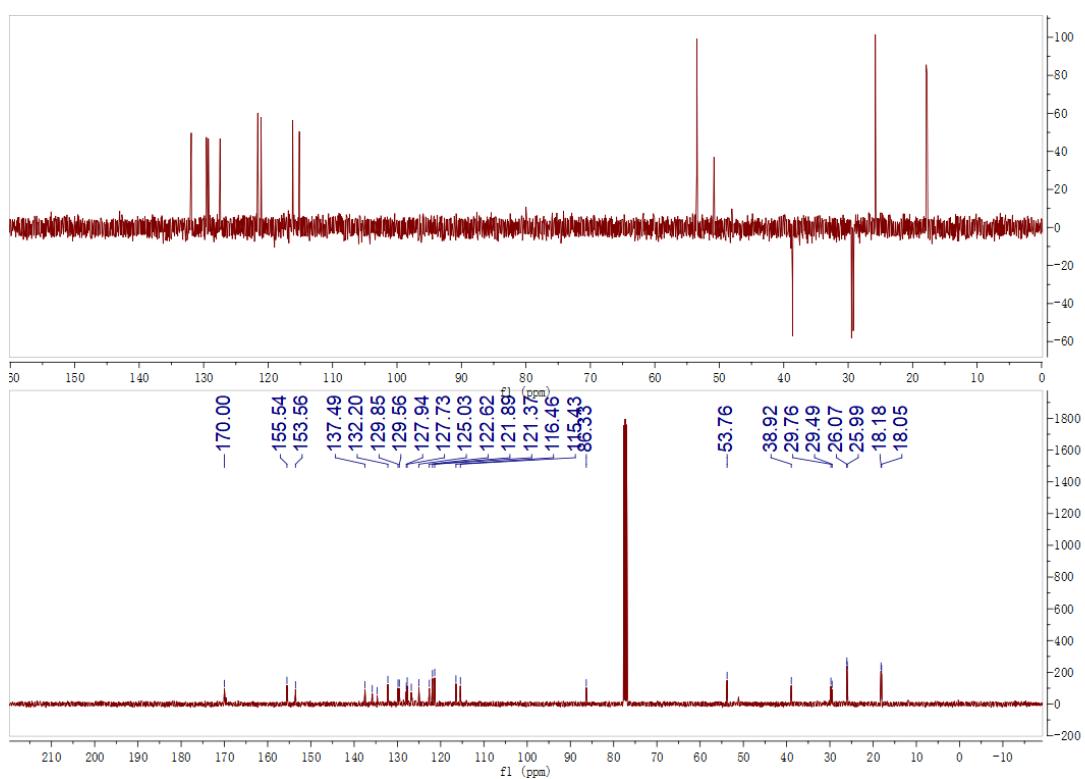


Figure S27. ^1H NMR spectrum of **13**.

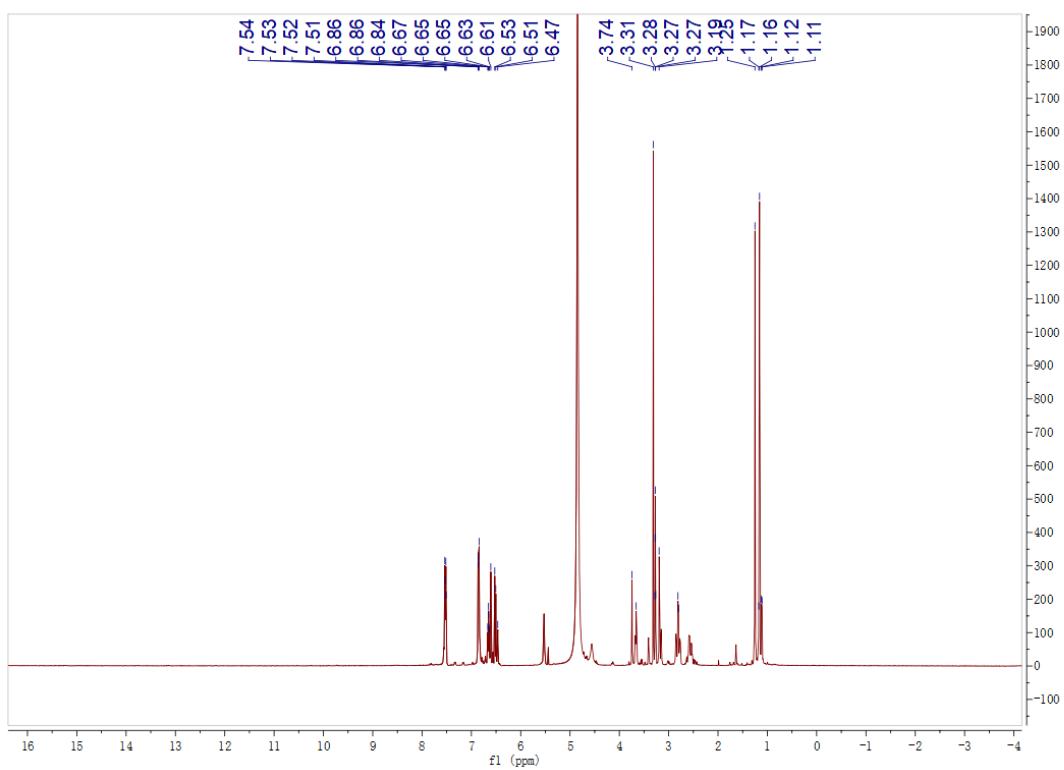


Figure S28. ^{13}C NMR spectrum of **13**.

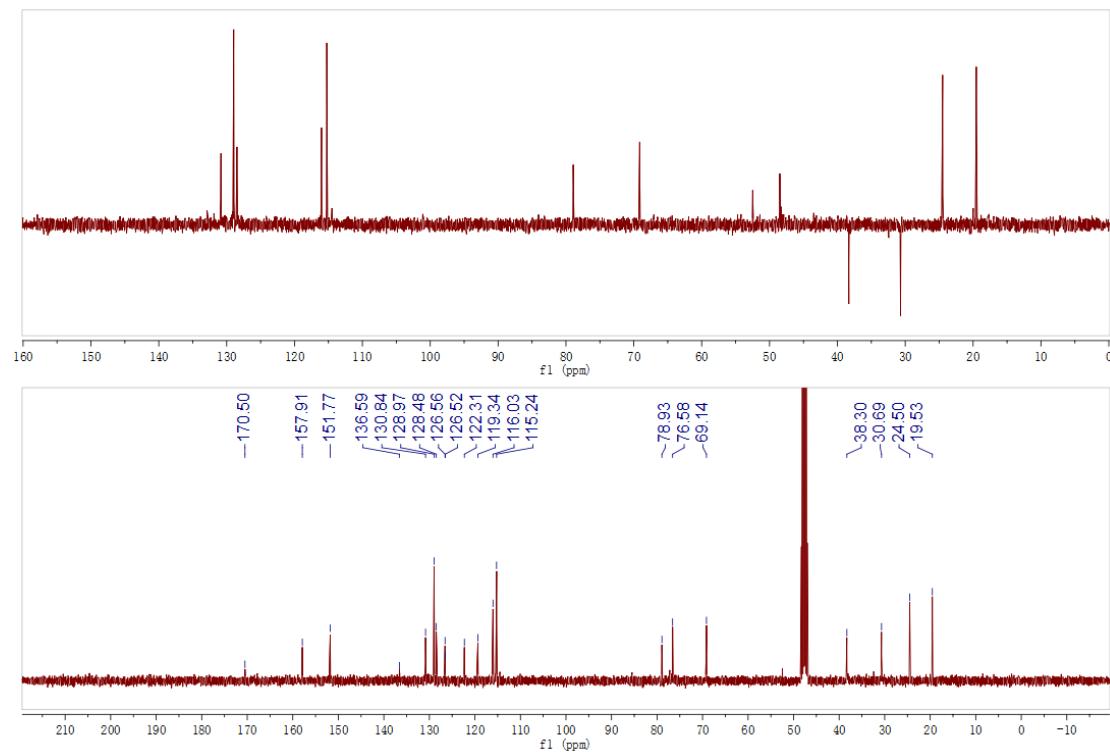


Figure S29. ^1H NMR spectrum of **14**.

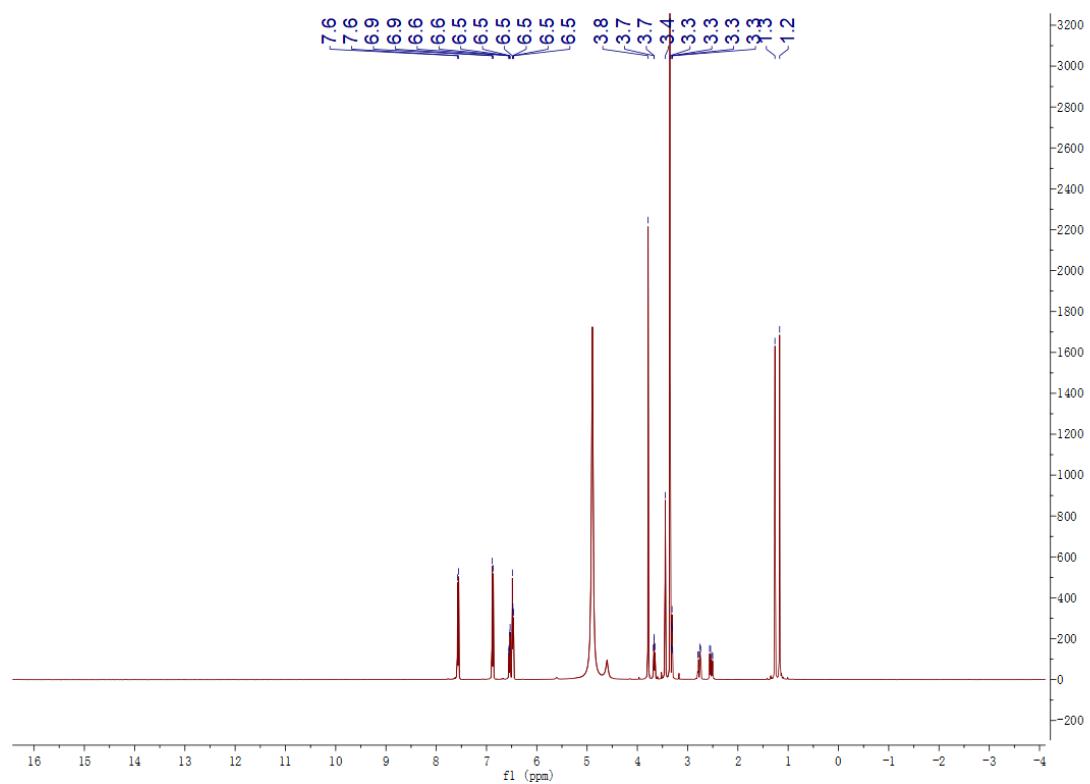


Figure S30. ^{13}C NMR spectrum of **14**.

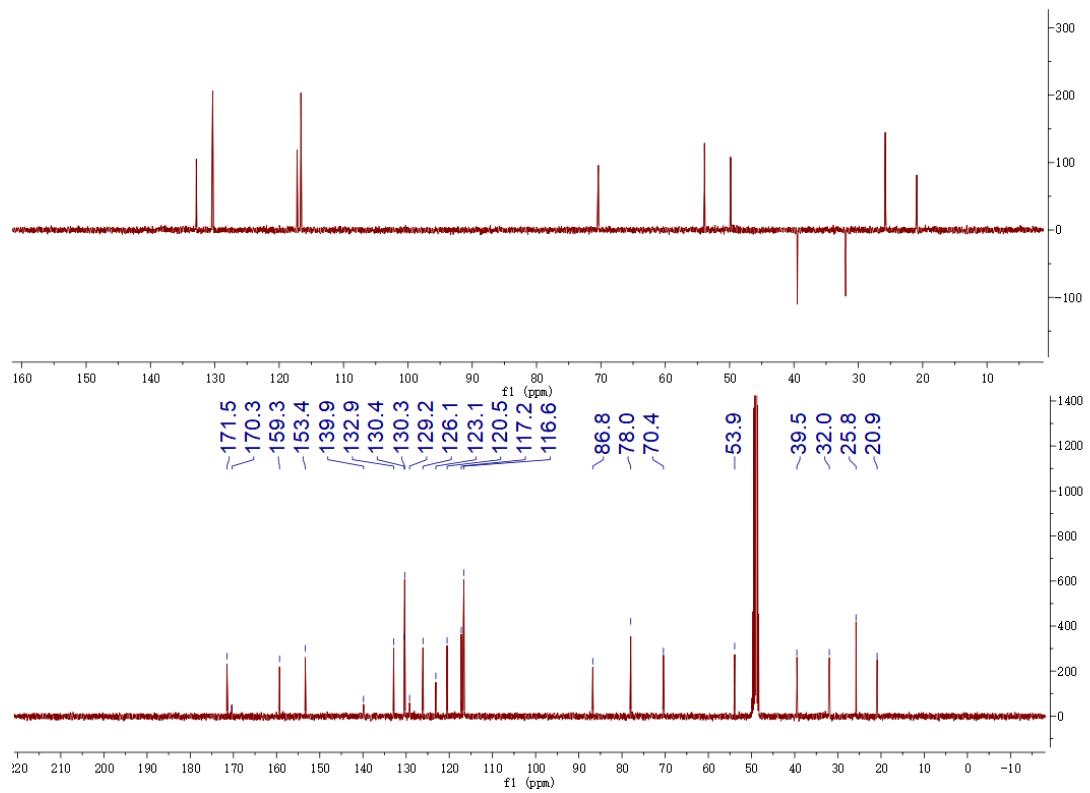


Figure S31. ^1H NMR spectrum of **15**.

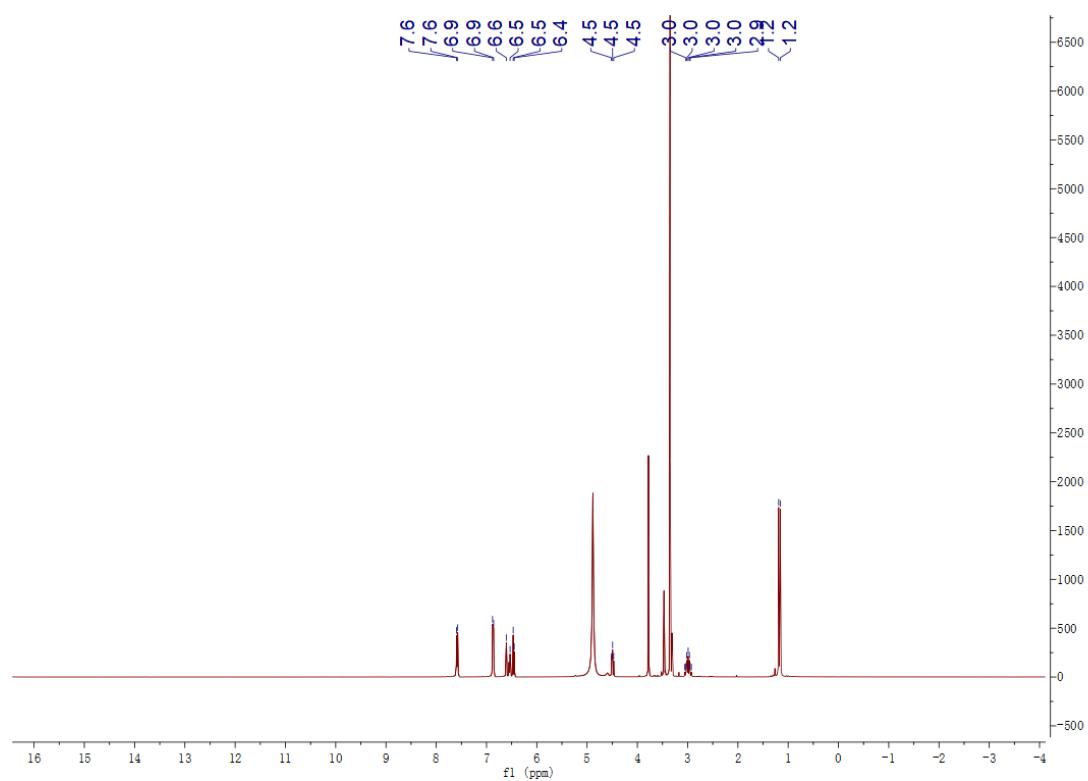


Figure S32. ^{13}C NMR spectrum of **15**.

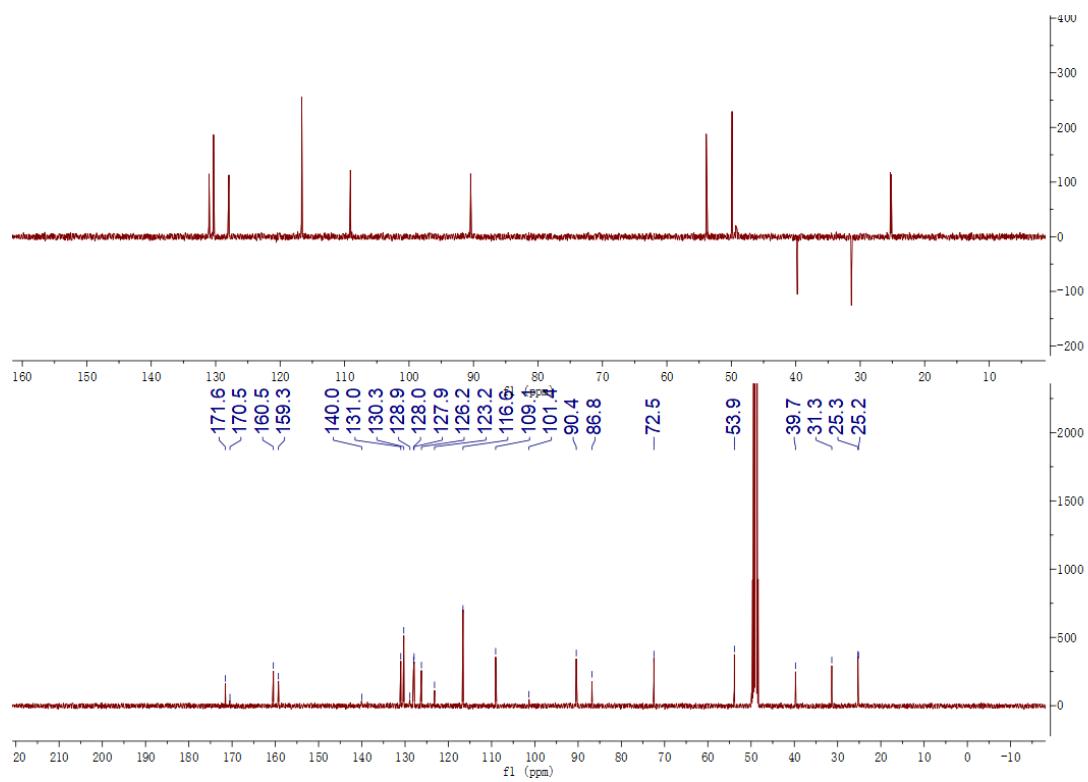


Figure S33. ^1H NMR spectrum of **16**.

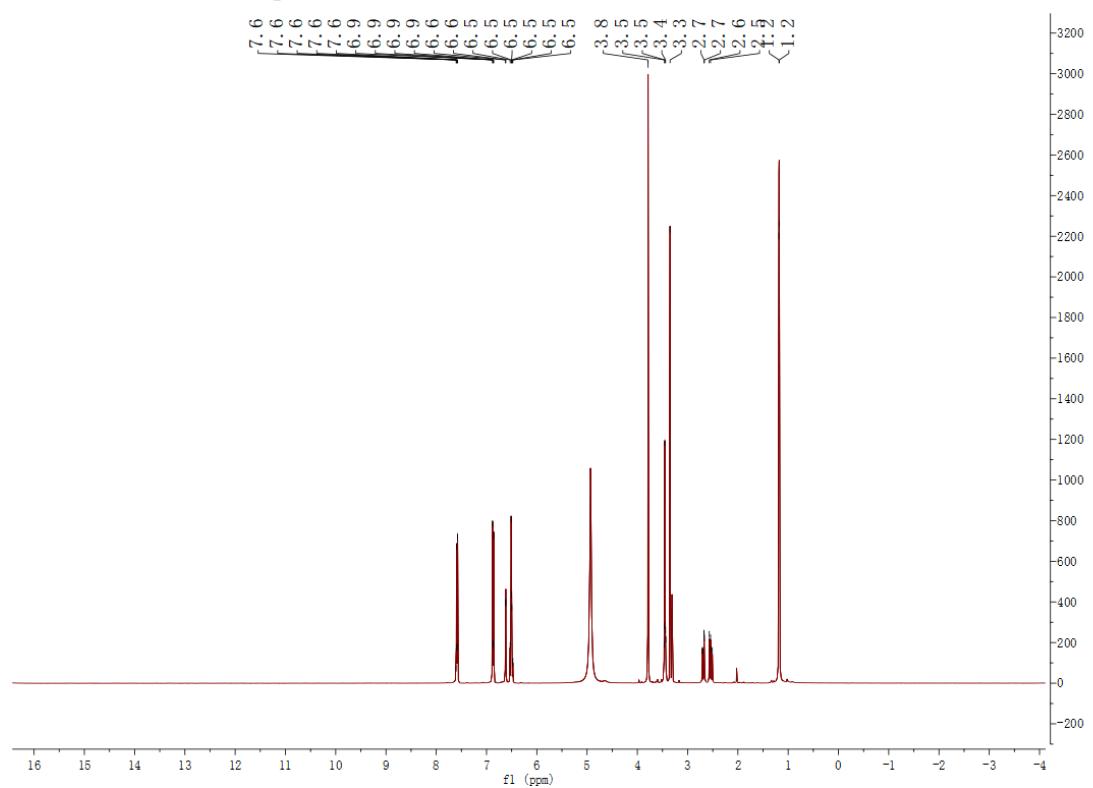


Figure S34. ^{13}C NMR spectrum of **16**.

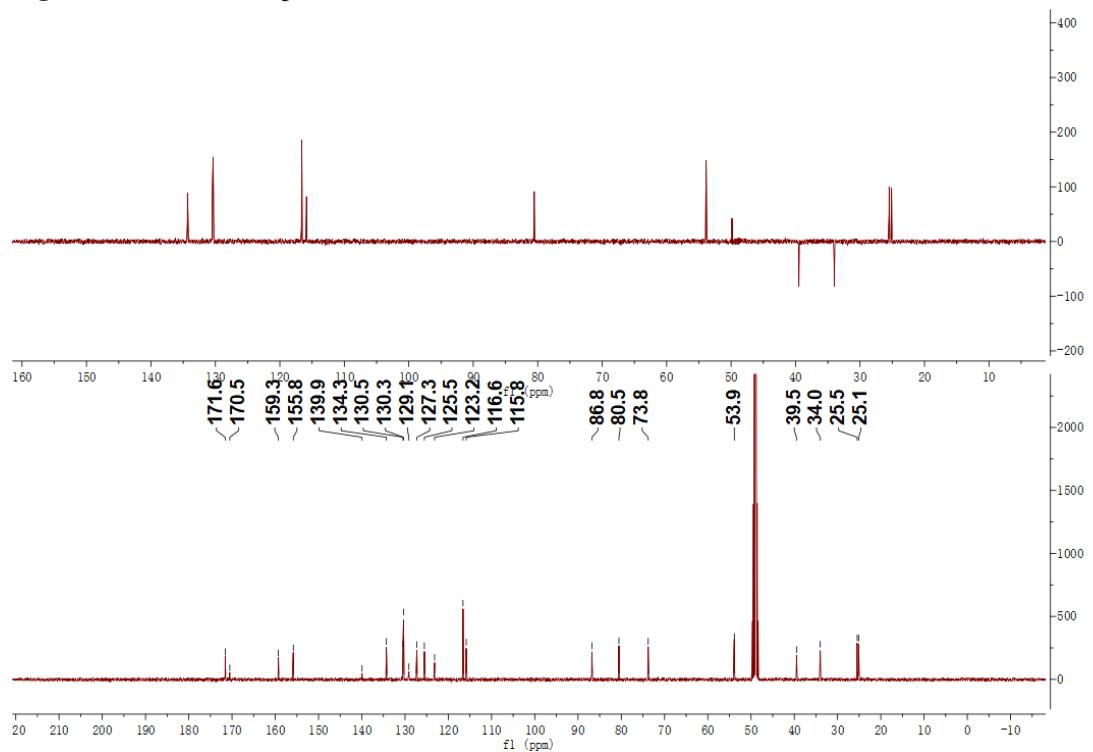


Figure S35. ^{13}C NMR spectrum of **17**.

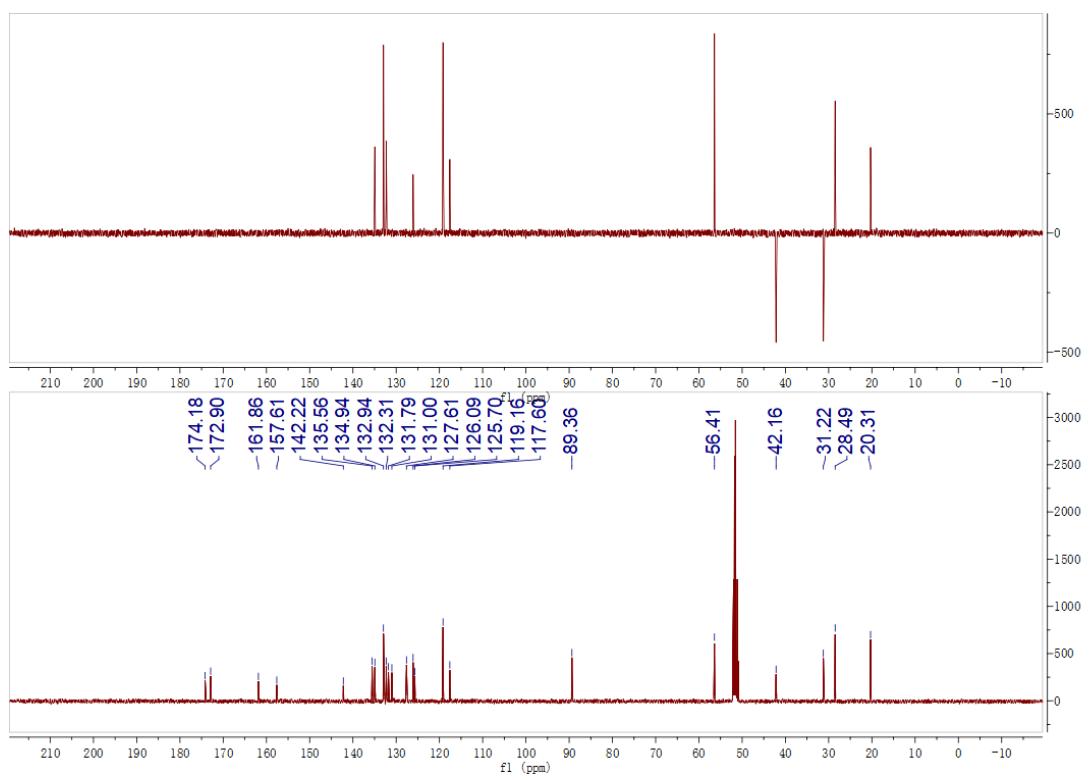


Figure S36. ^1H NMR spectrum of **18**.

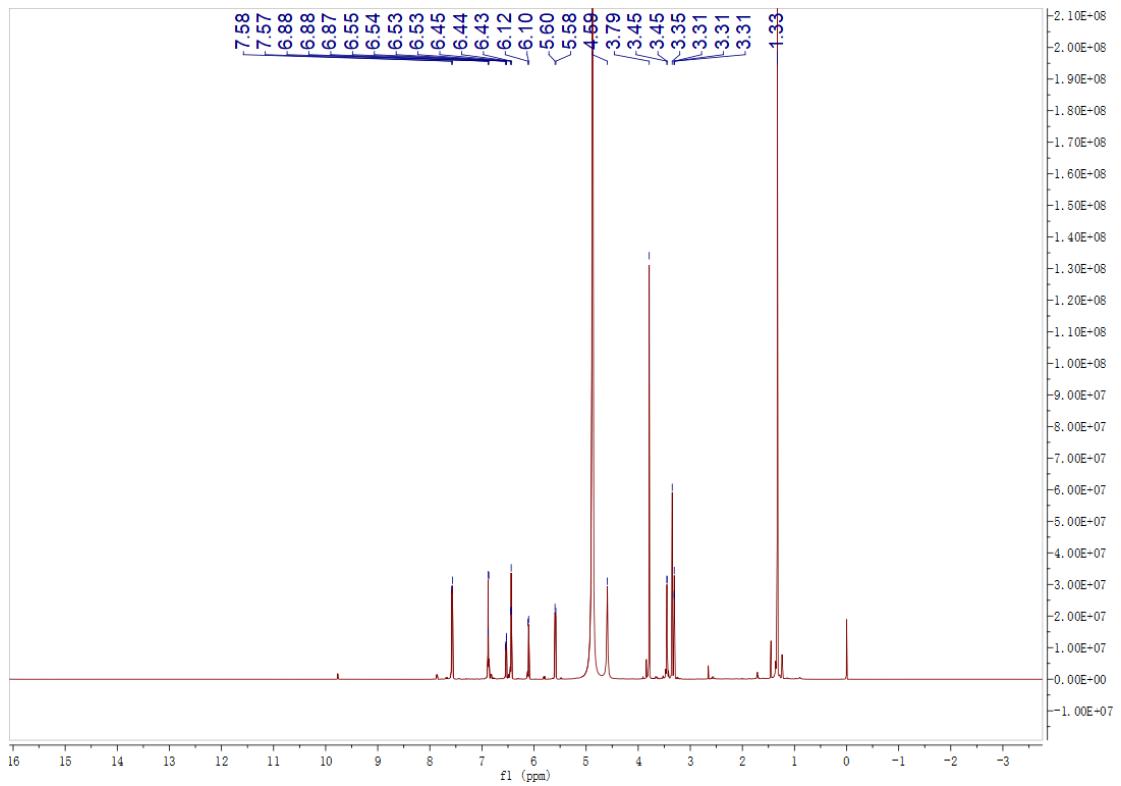


Figure S37. ^{13}C NMR spectrum of **18**.

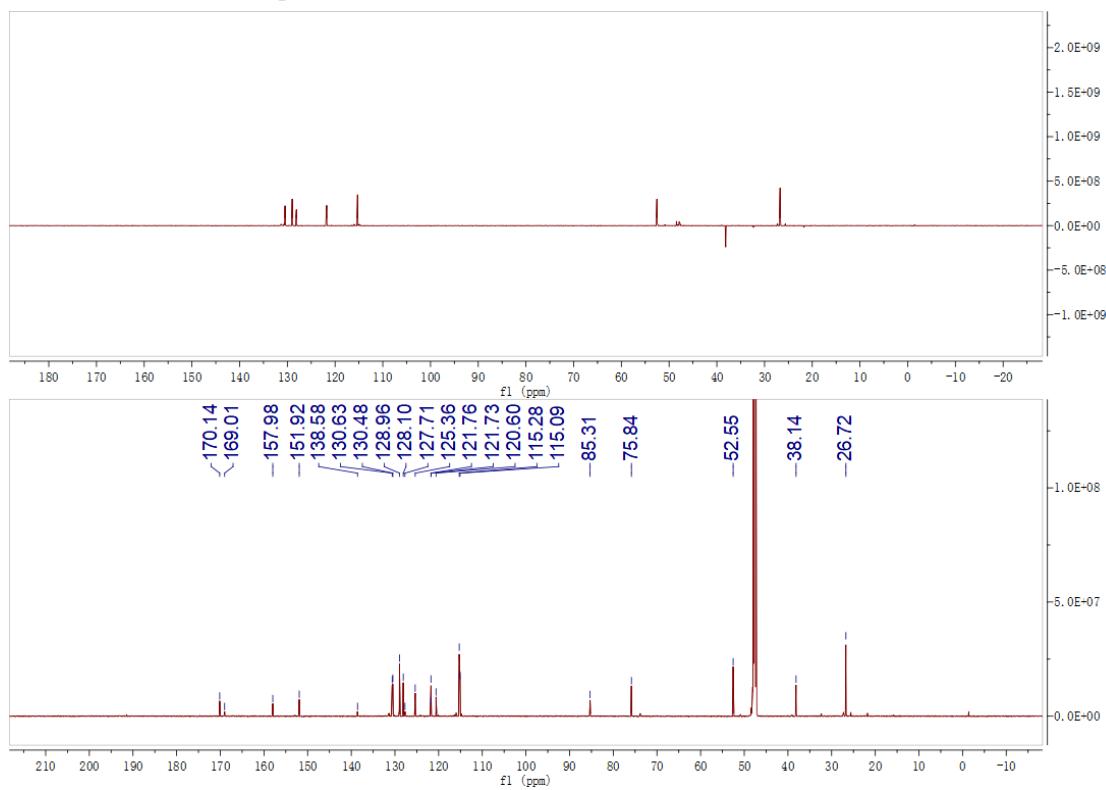


Figure S38. HPLC profile for the extract of the culture of *Aspergillus terreus* [MeOH–H₂O (20:80 to 80:20, 80 min, v/v, 2.0 mL/min)].

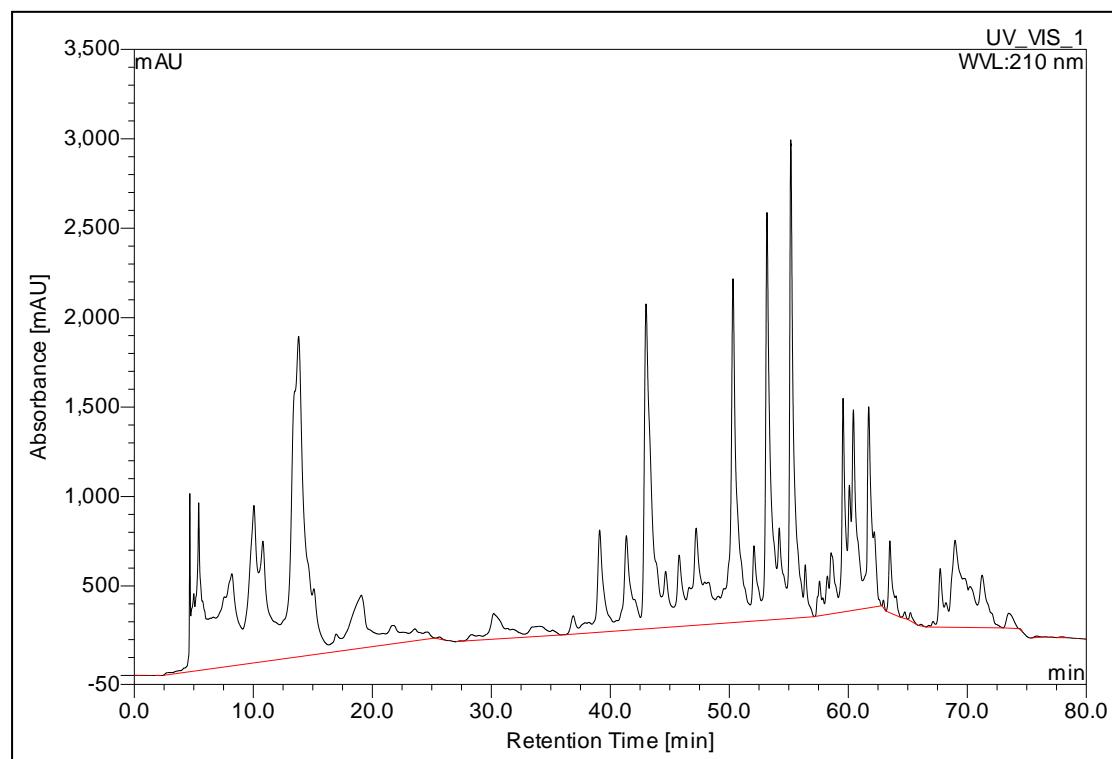


Figure S39. Purity tests of all compounds 1-18.

