

## **SUPPLEMENTARY MATERIAL**

### **Antibacterial Secondary Metabolites from Marine-derived Fungus *Aspergillus* sp. IMCASMF180035**

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Hongkai Bi<sup>5</sup> and Xiuli Xu<sup>2,\*</sup>**

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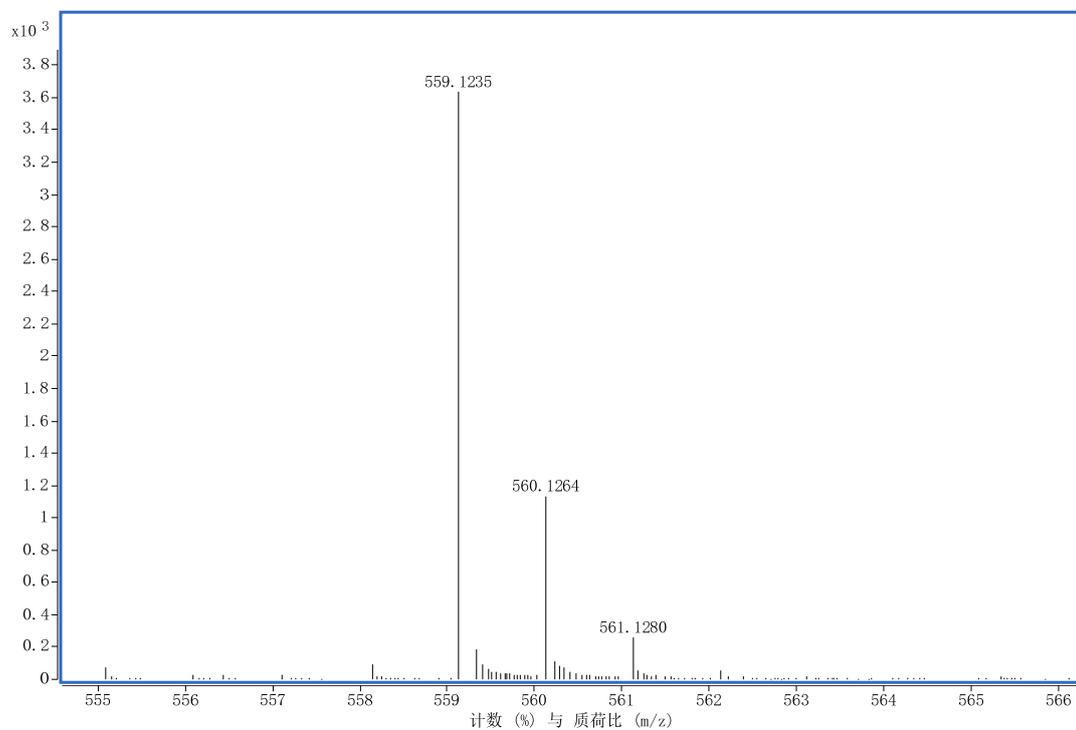
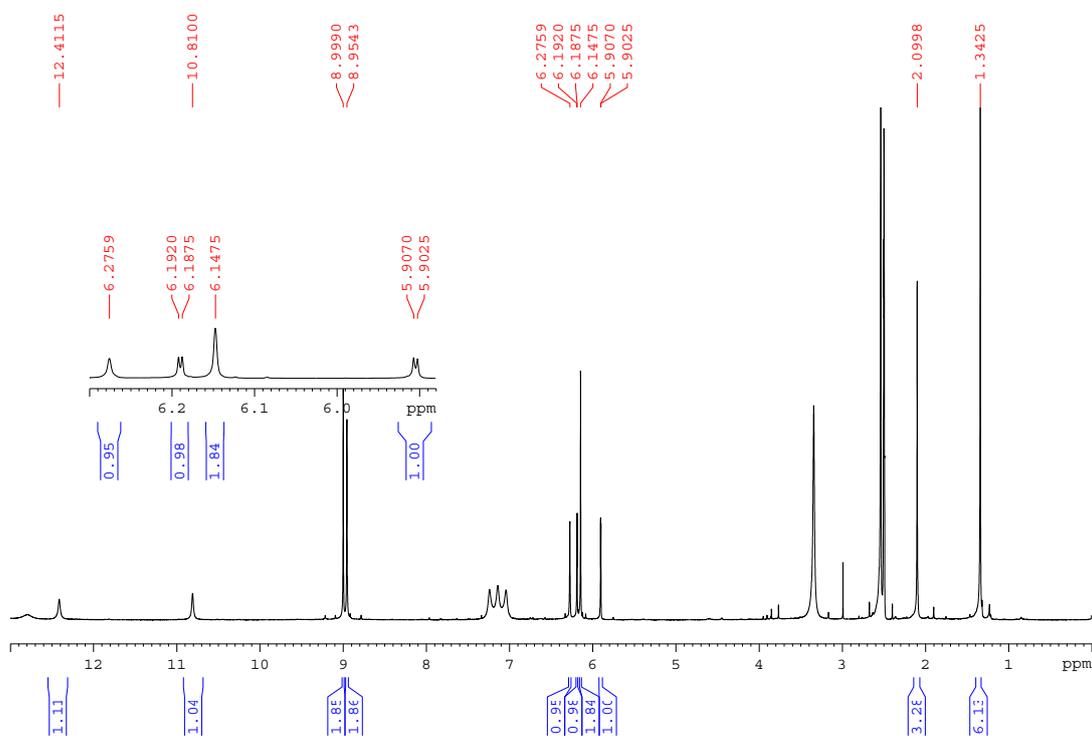
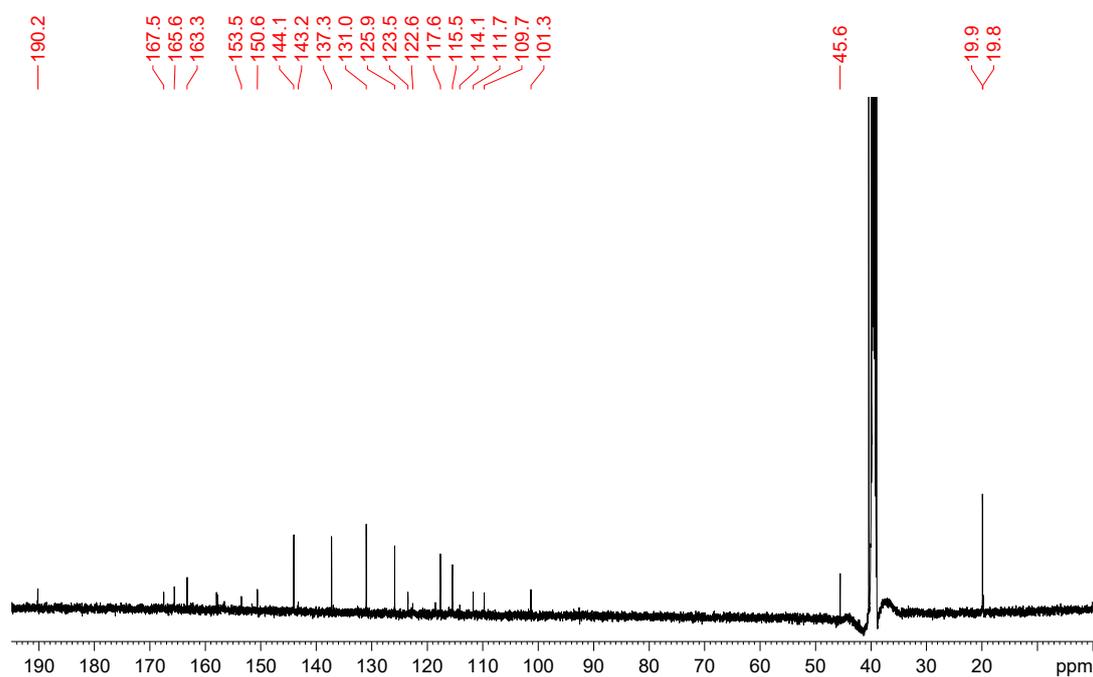
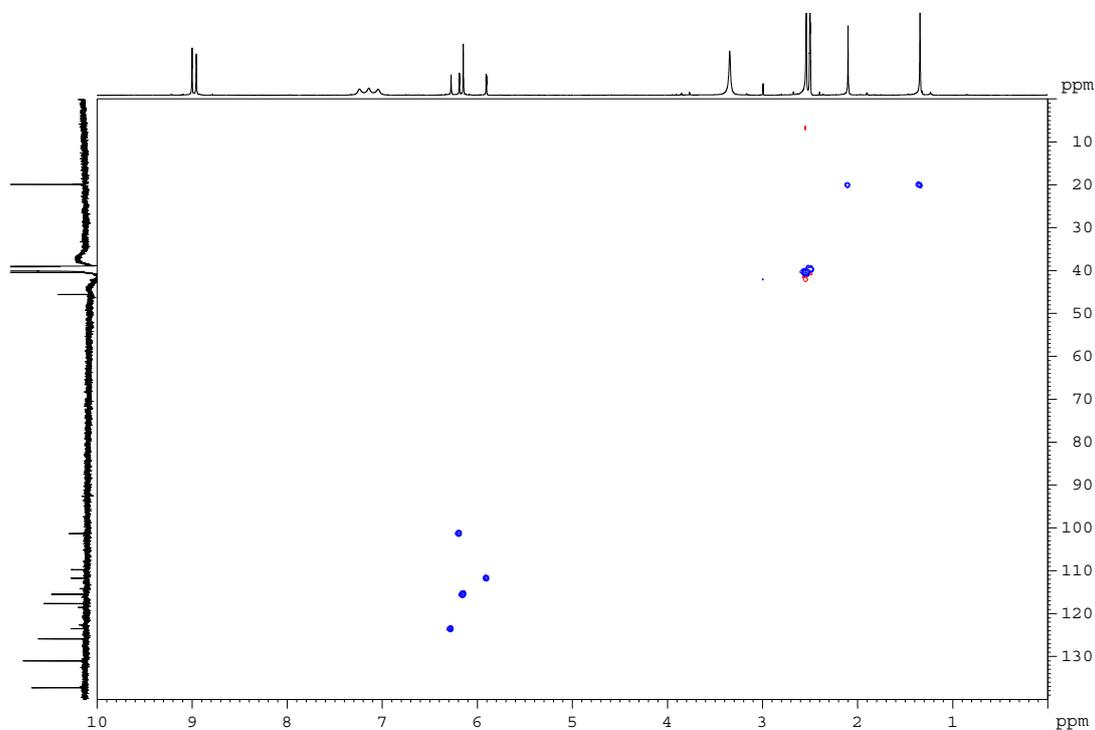
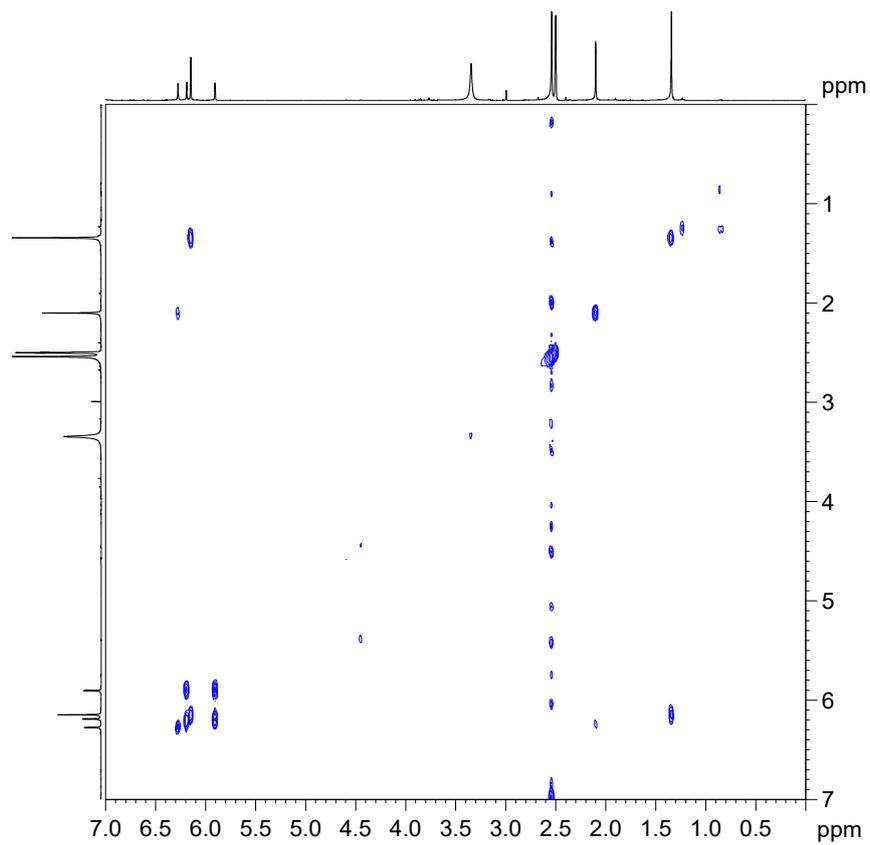


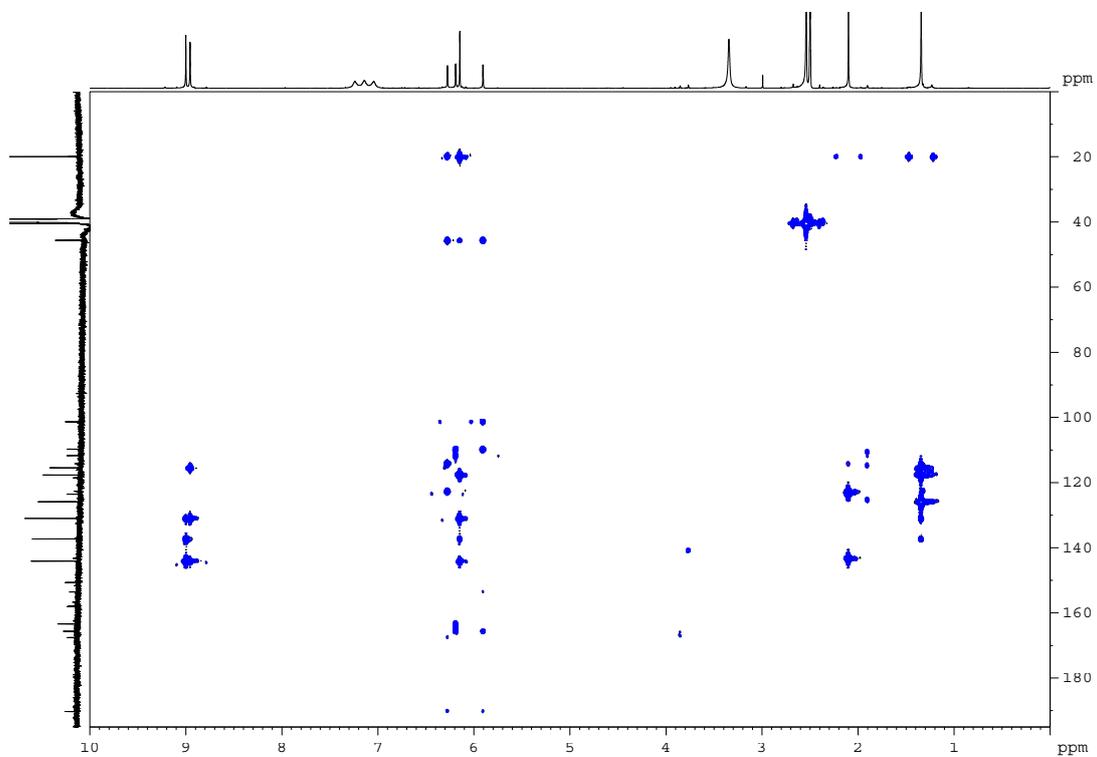
Figure S1. HRESIMS spectrum for compound 1



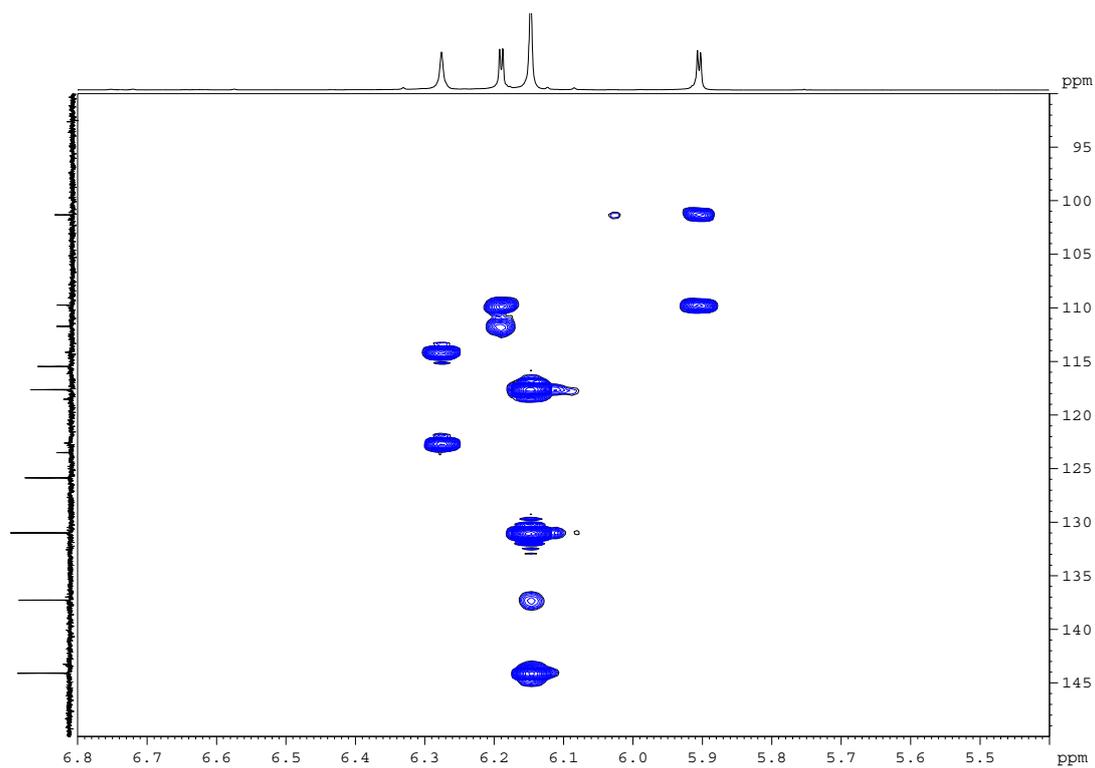
**Figure S2.**  $^1\text{H}$  NMR spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **1****Figure S3.**  $^{13}\text{C}$  NMR spectrum (125 MHz,  $\text{DMSO-}d_6$ ) of **1****Figure S4.** HSQC spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **1**



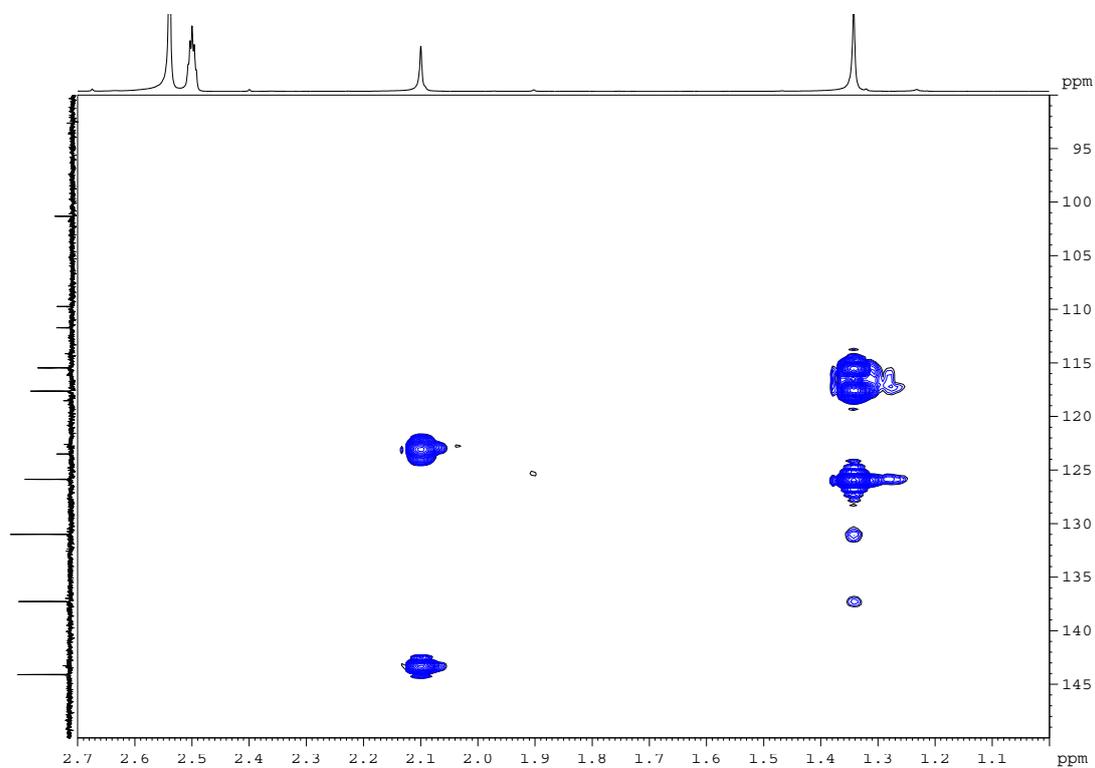
**Figure S5.**  $^1\text{H}$ - $^1\text{H}$  COSY spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **1**



**Figure S6.** HMBC spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **1**



**Figure S7.** Expansion of HMBC spectrum (500 MHz, DMSO-*d*<sub>6</sub>) of **1**



**Figure S8.** Expansion of HMBC spectrum (500 MHz, DMSO-*d*<sub>6</sub>) of **1**

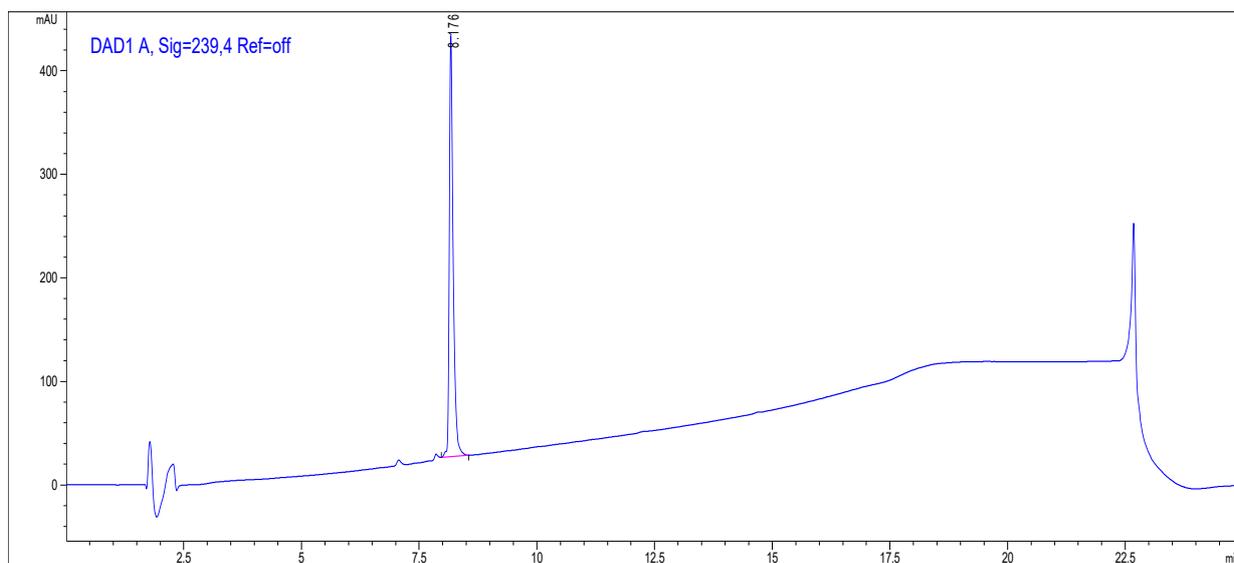


Figure S9. HPLC profile of 1

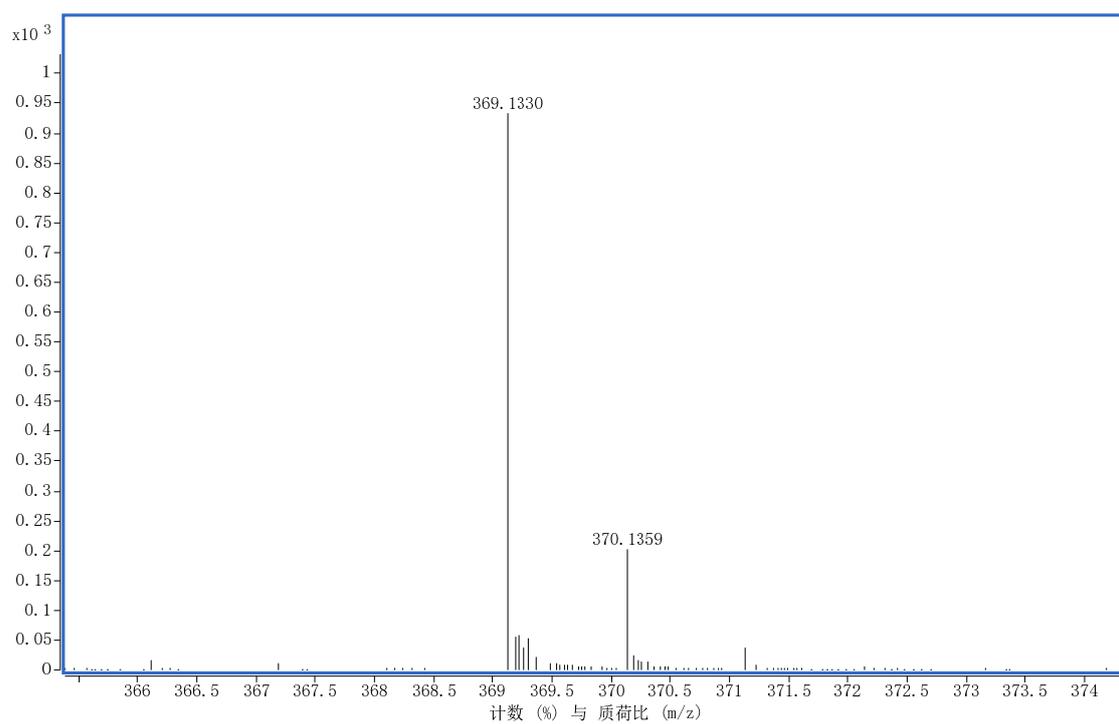


Figure S10. HRESIMS spectrum of for 2

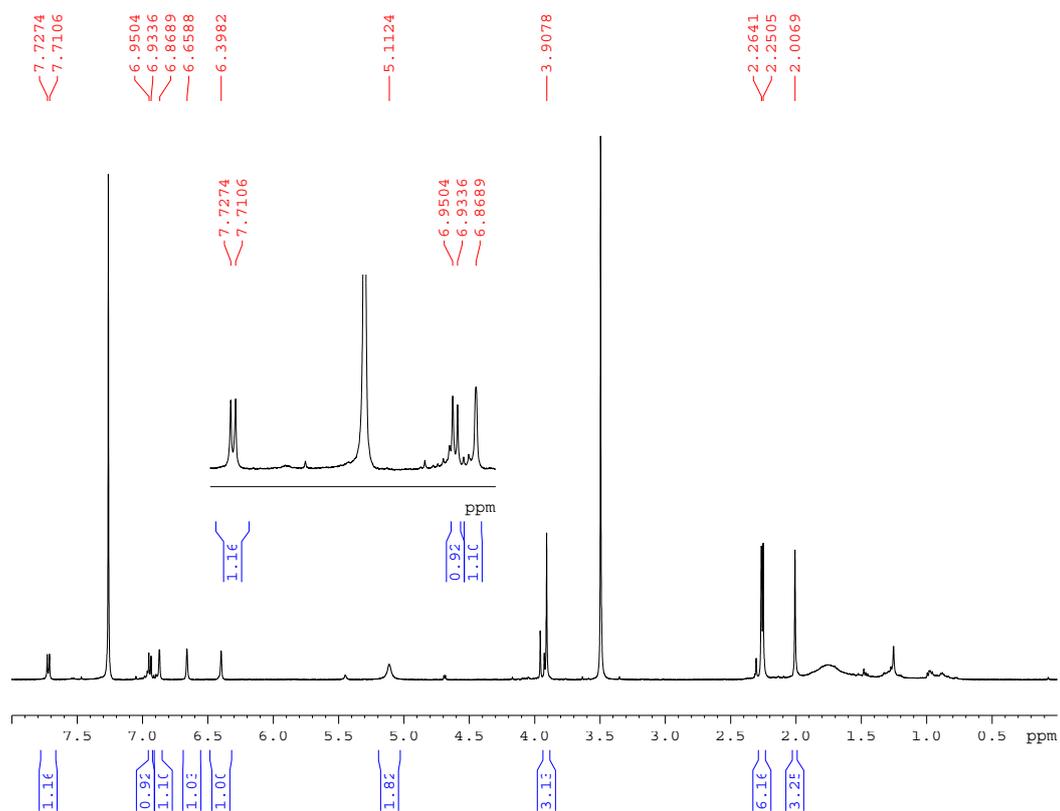


Figure S11.  $^1\text{H}$  NMR spectrum (500 MHz,  $\text{CDCl}_3$ ) of **2**

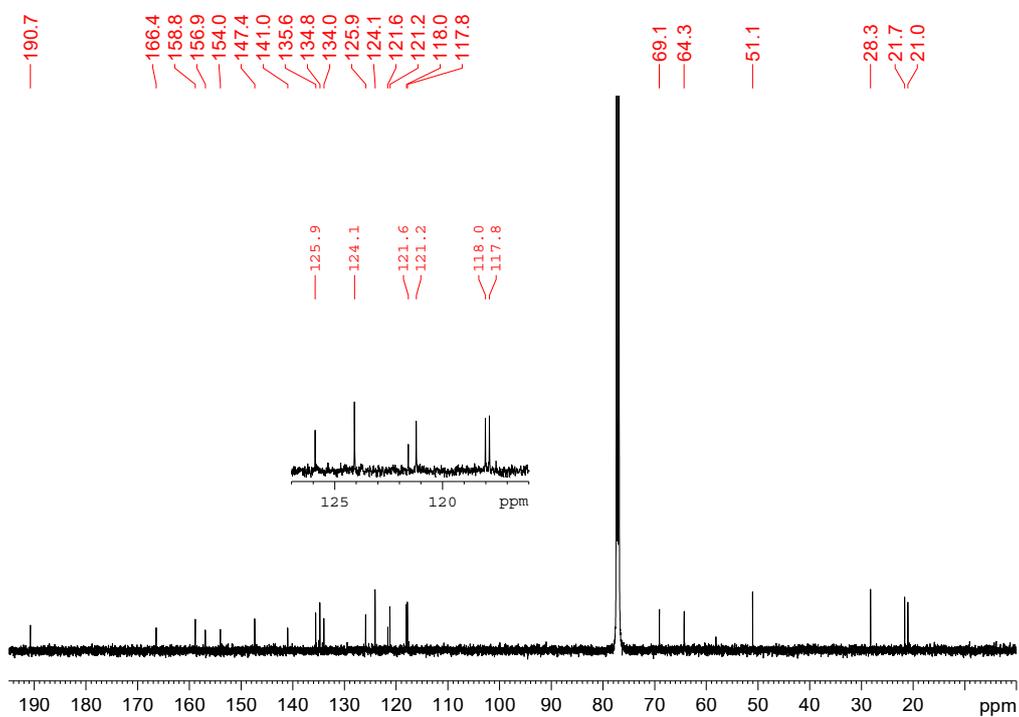
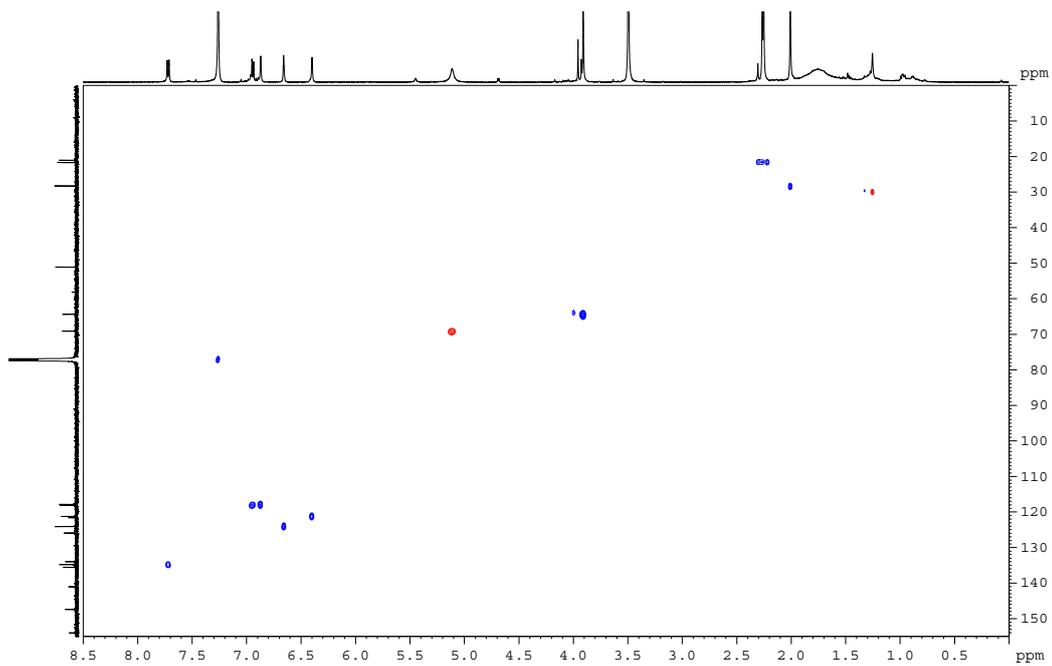
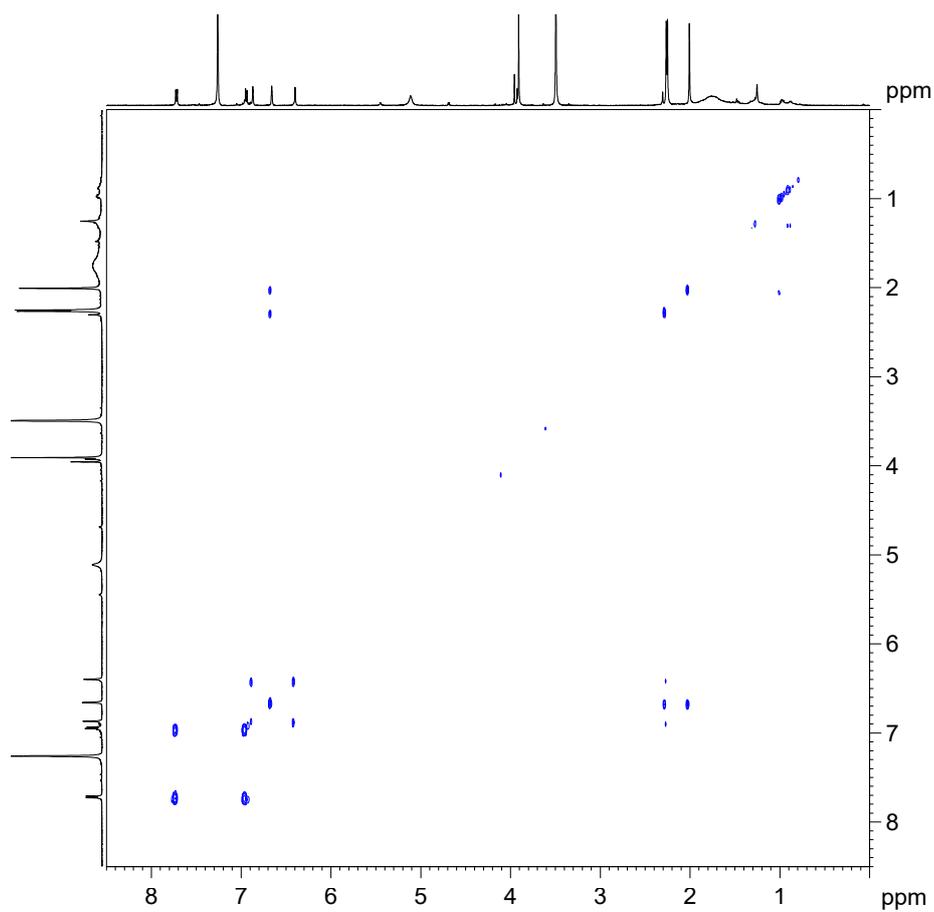


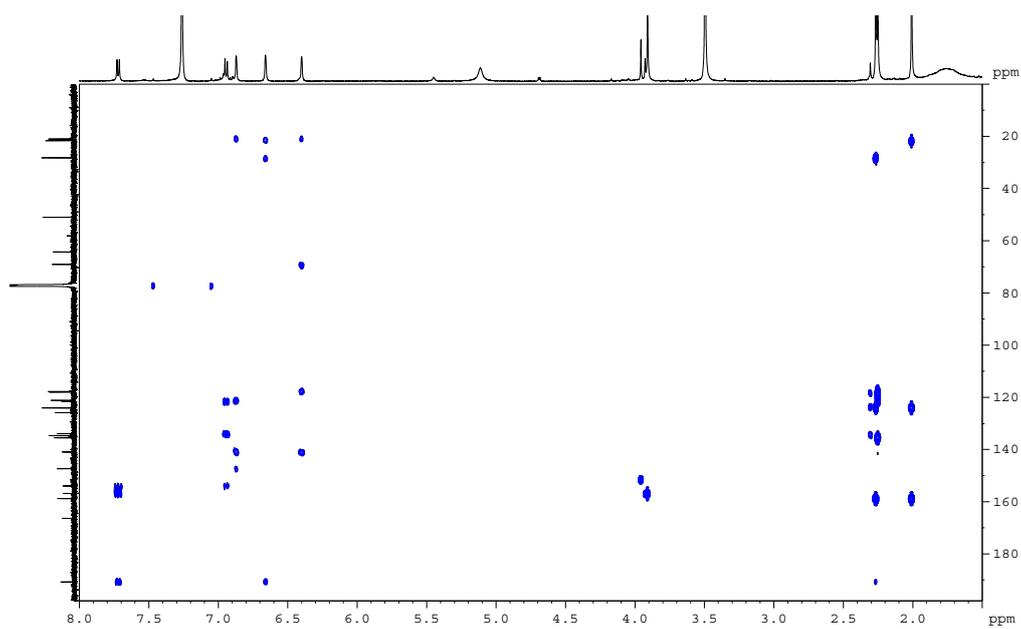
Figure S12.  $^{13}\text{C}$  NMR spectrum (125 MHz,  $\text{CDCl}_3$ ) of **2**



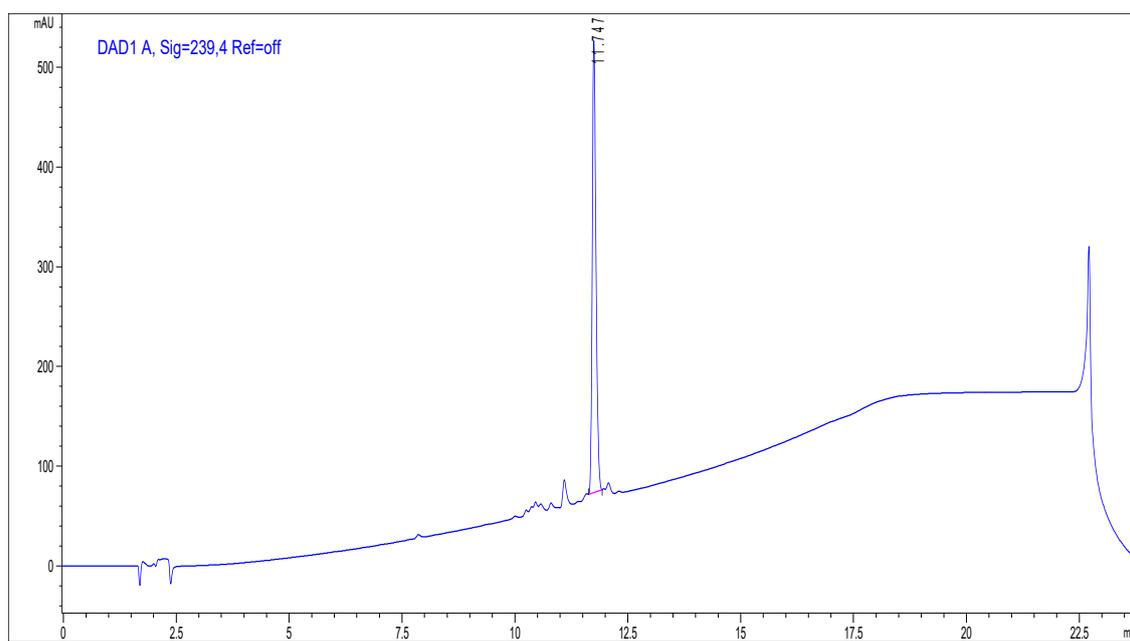
**Figure S13.** HSQC spectrum (500 MHz,  $\text{CDCl}_3$ ) of **2**



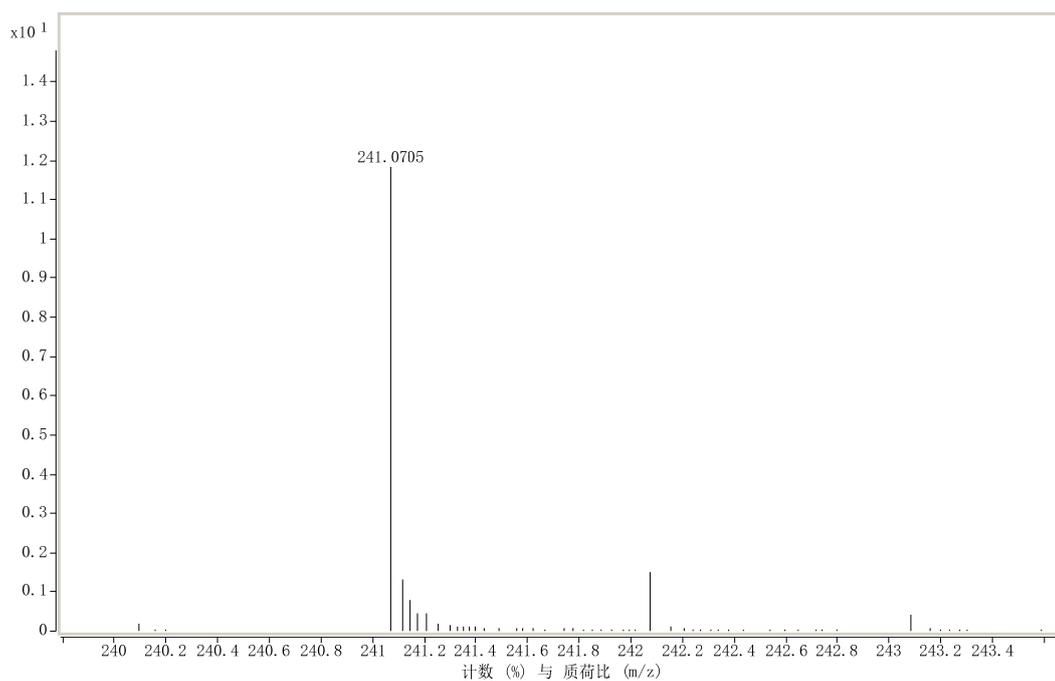
**Figure S14.**  $^1\text{H}$ - $^1\text{H}$  COSY spectrum (500 MHz,  $\text{CDCl}_3$ ) of **2**



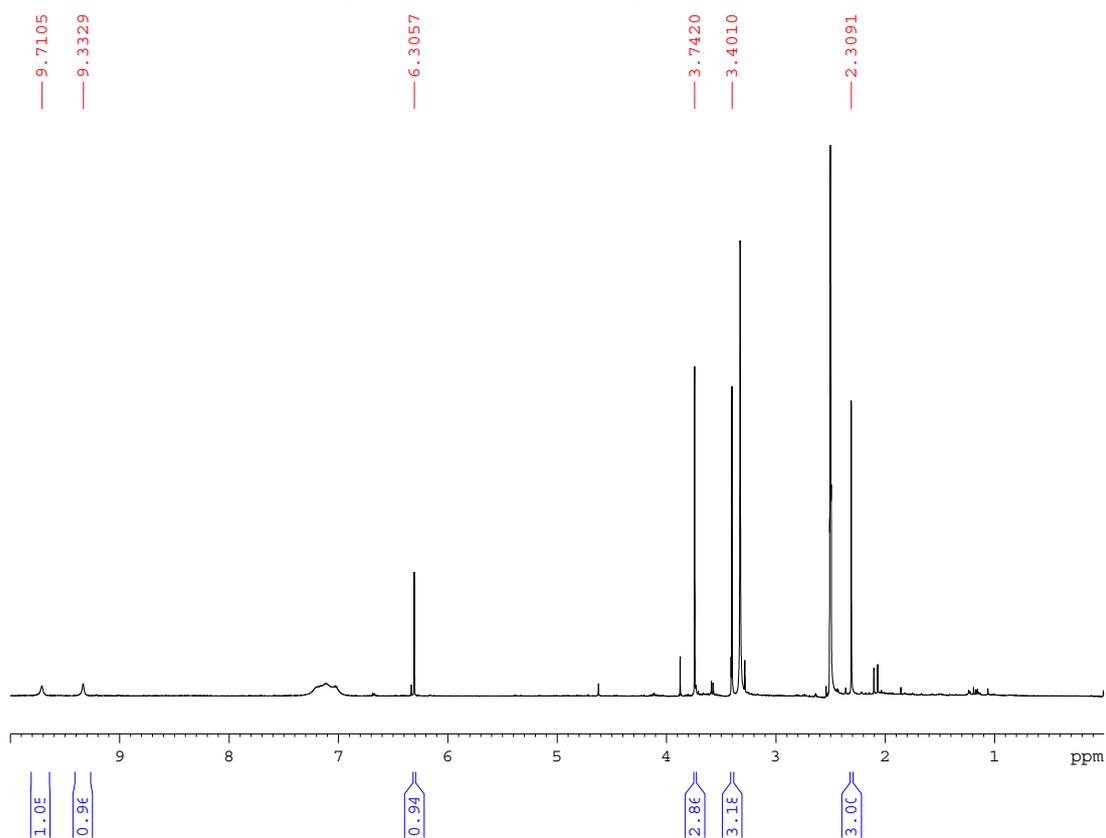
**Figure S15.** HMBC spectrum (500 MHz, CDCl<sub>3</sub>) of **2**



**Figure S16.** HPLC profile of **2**



**Figure S17.** HRESIMS spectrum for **3**



**Figure S18.**  $^1\text{H}$  NMR spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **3**

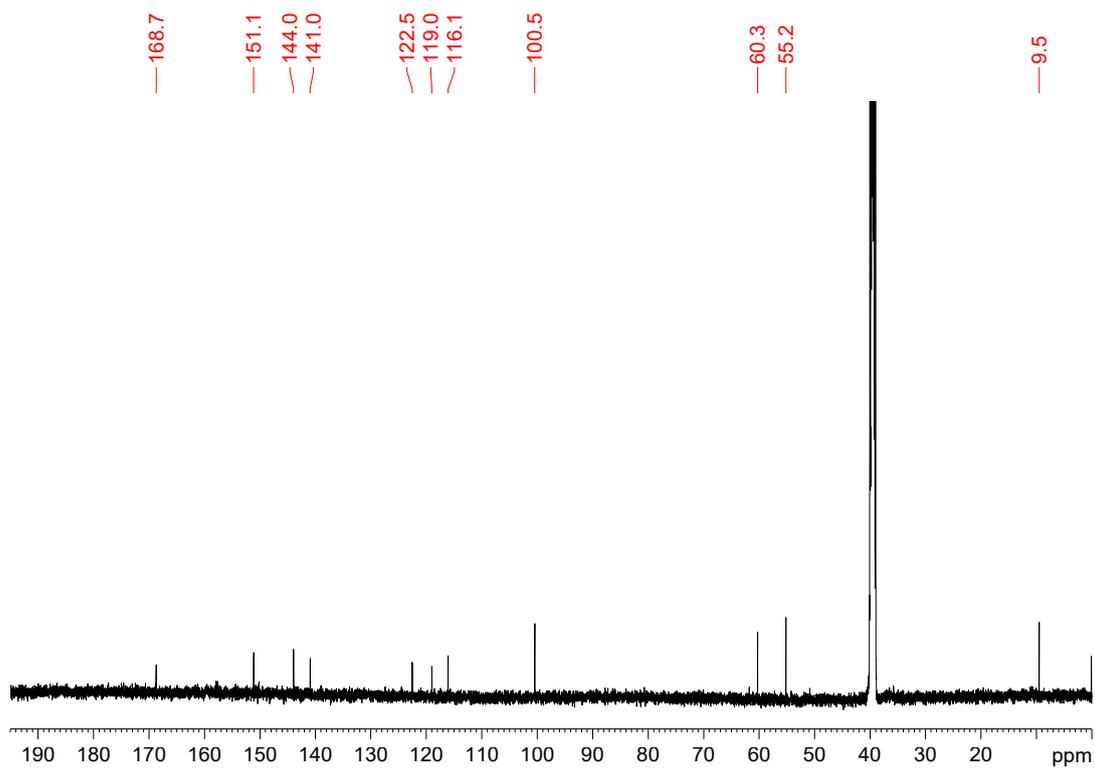


Figure S19.  $^{13}\text{C}$  NMR spectrum (125 MHz,  $\text{DMSO-}d_6$ ) of **3**

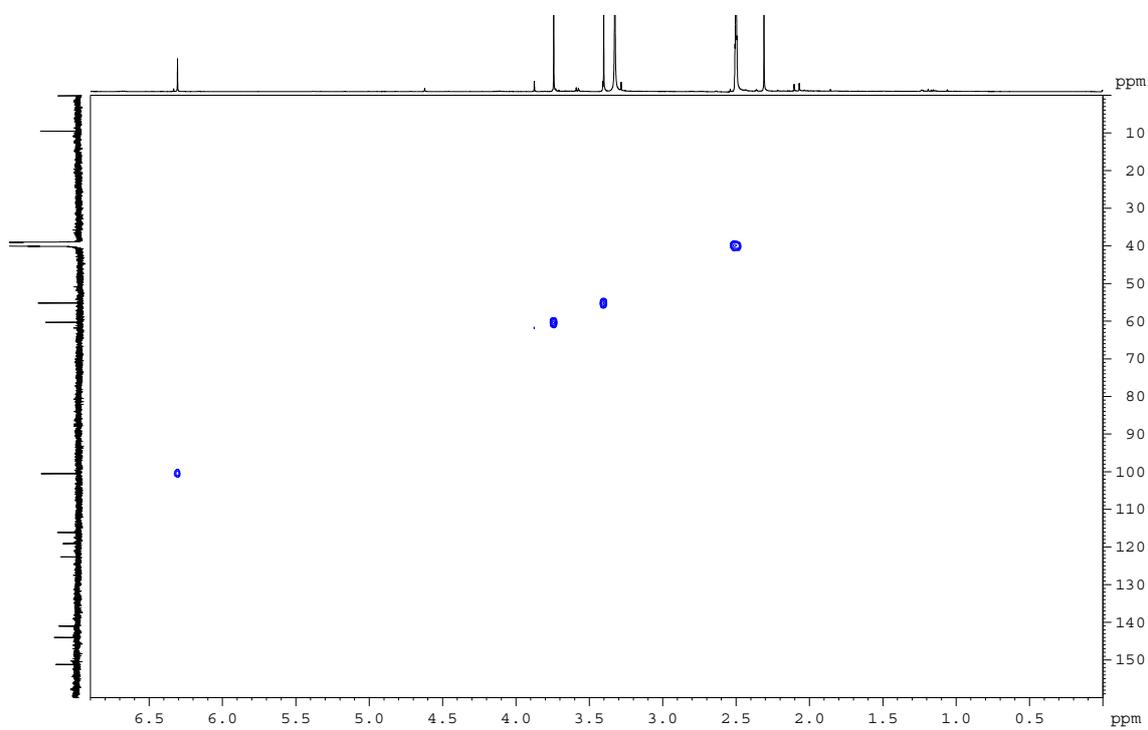
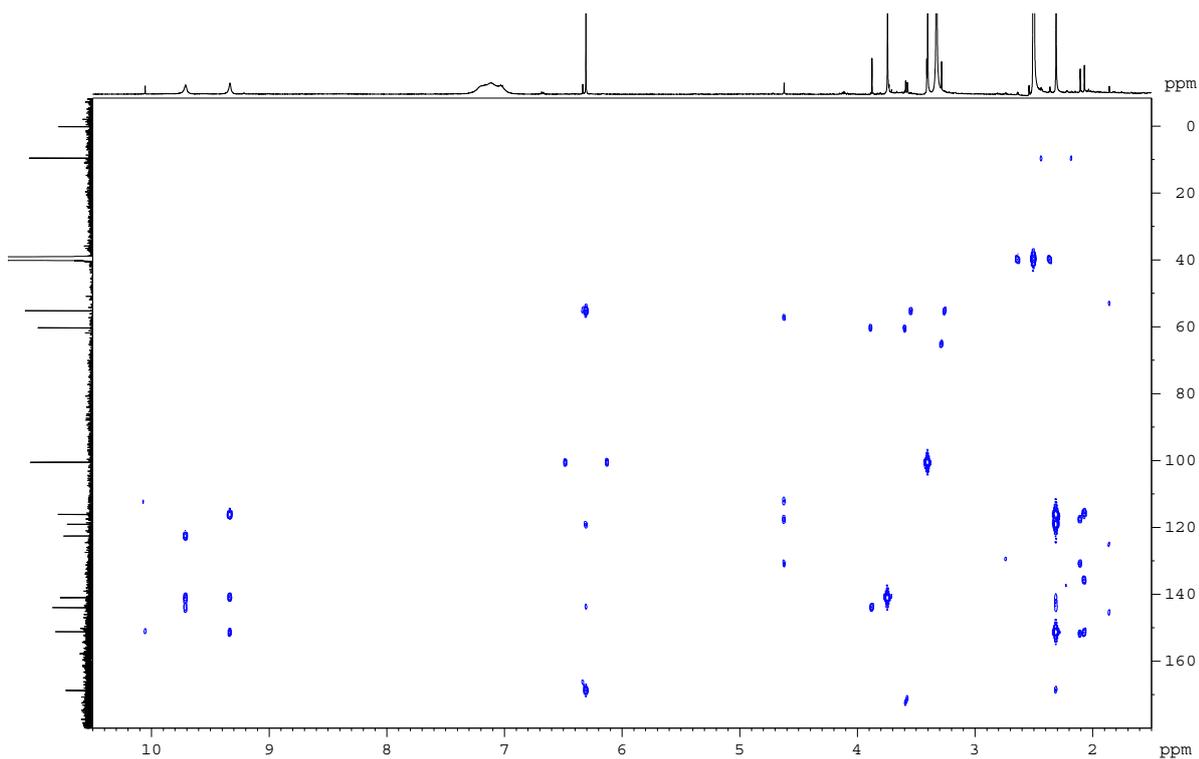
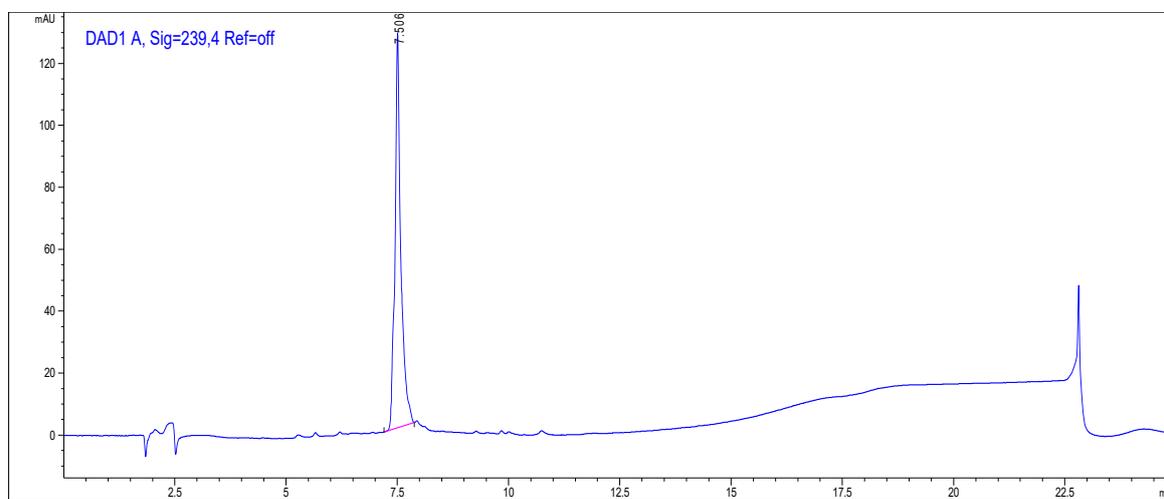


Figure S20. HSQC spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **3**



**Figure S21.** HMBC spectrum (500 MHz, DMSO-*d*<sub>6</sub>) of **3**



**Figure S22.** HPLC profile of **3**

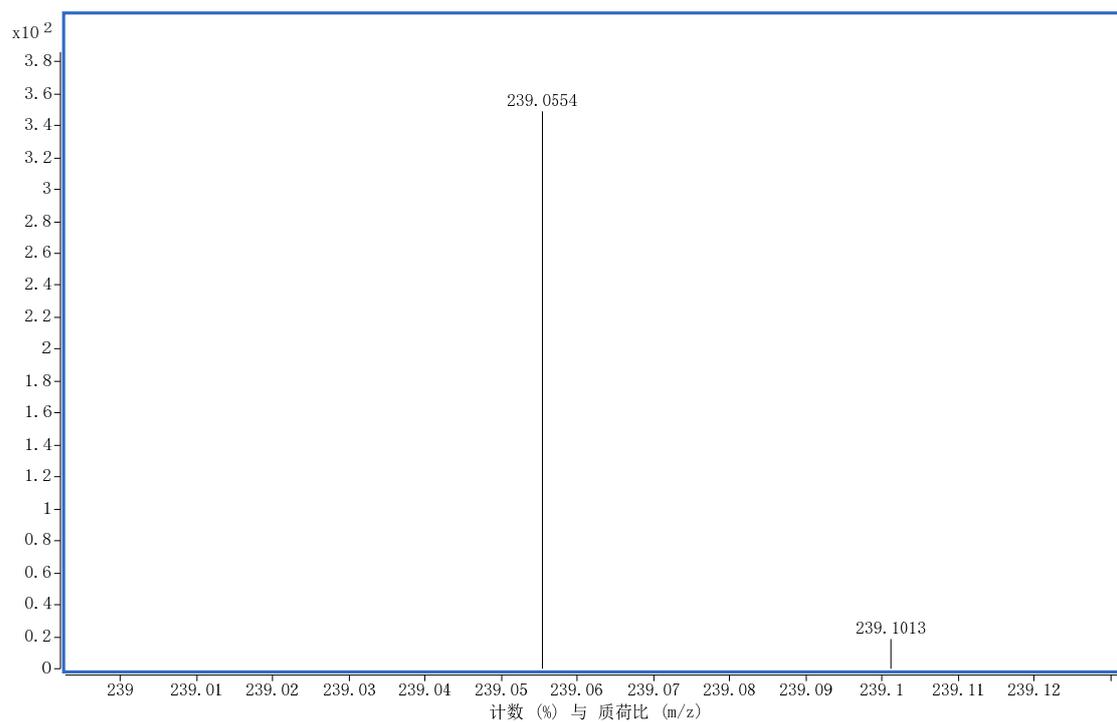
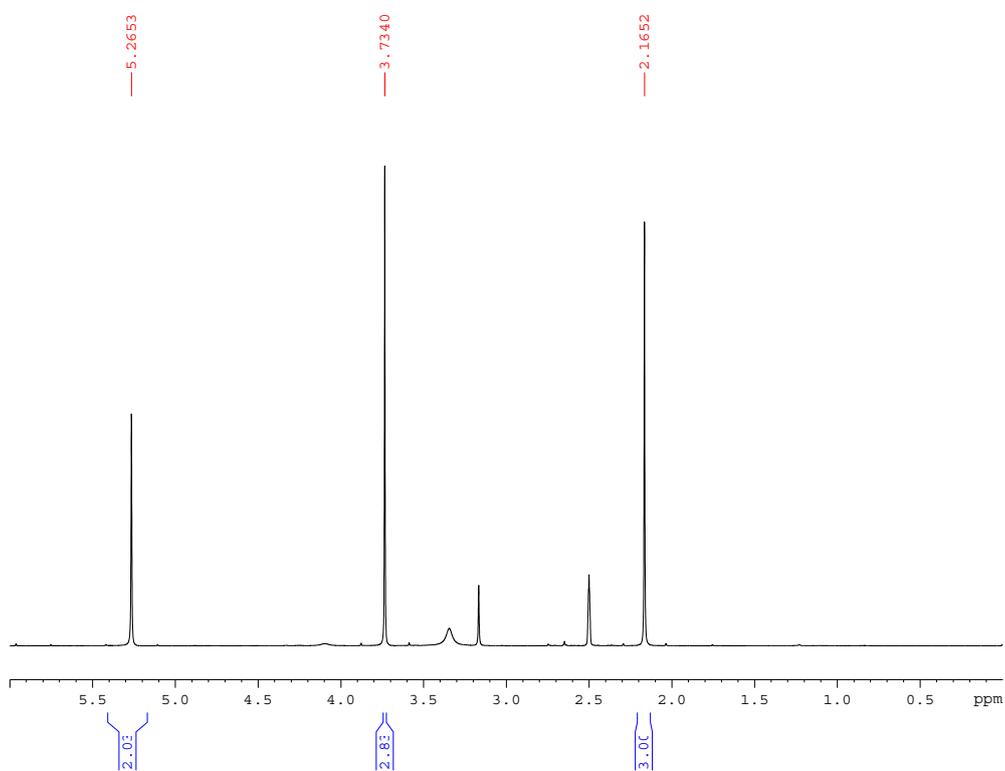
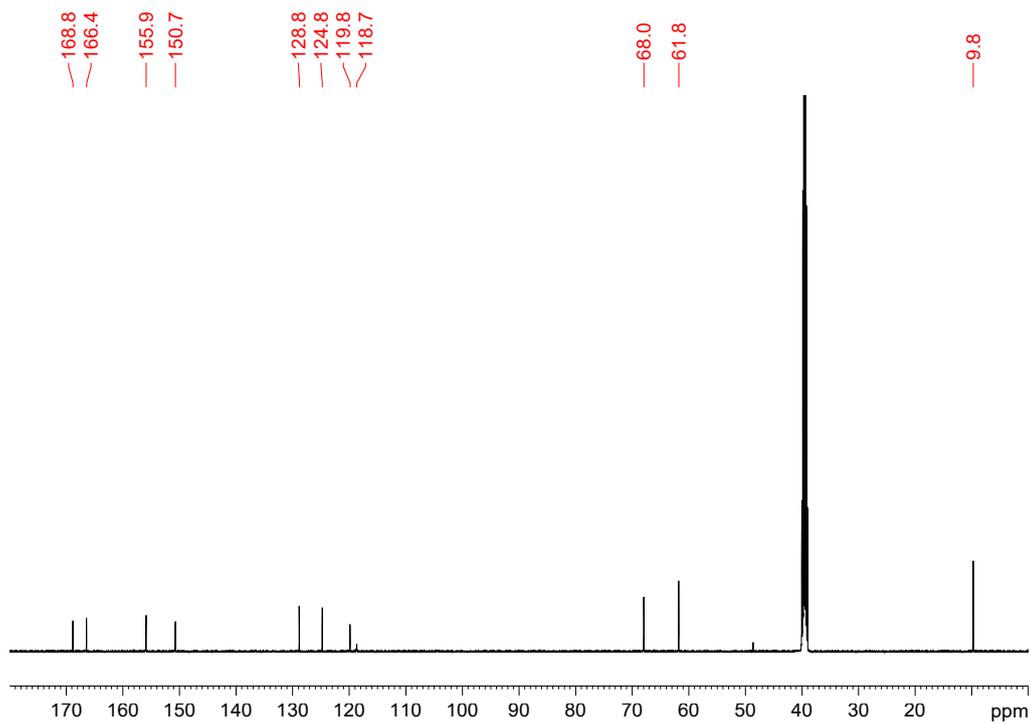
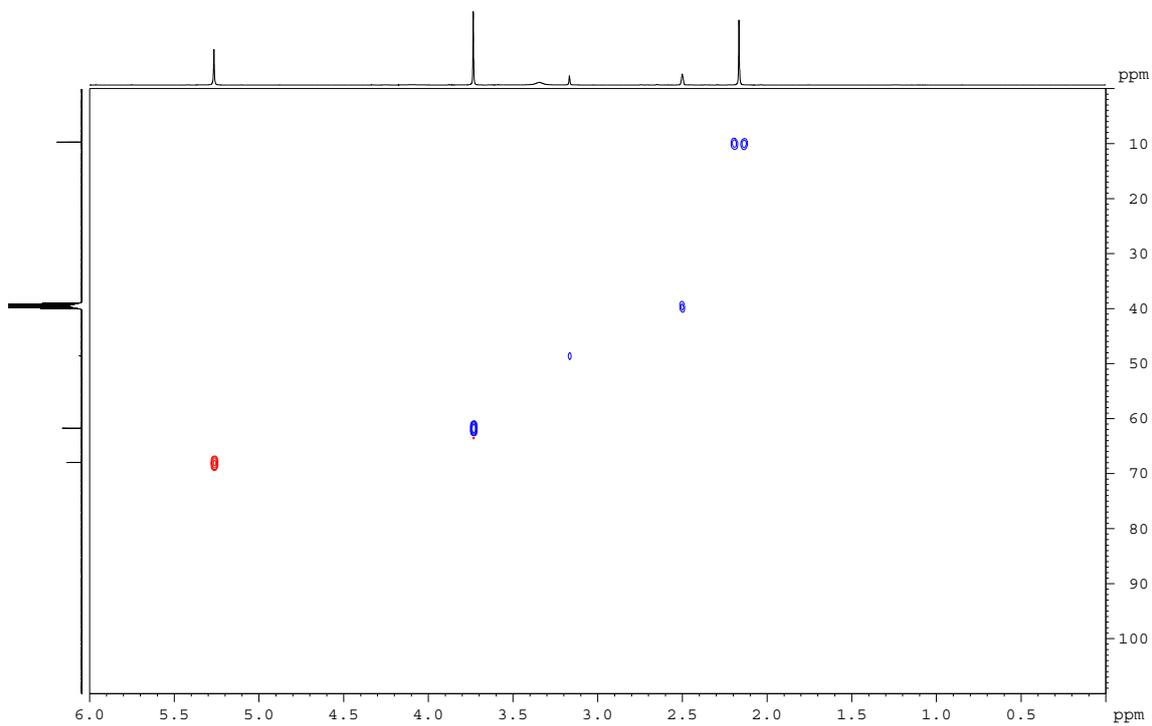


Figure S23. HRESIMS spectrum for 4

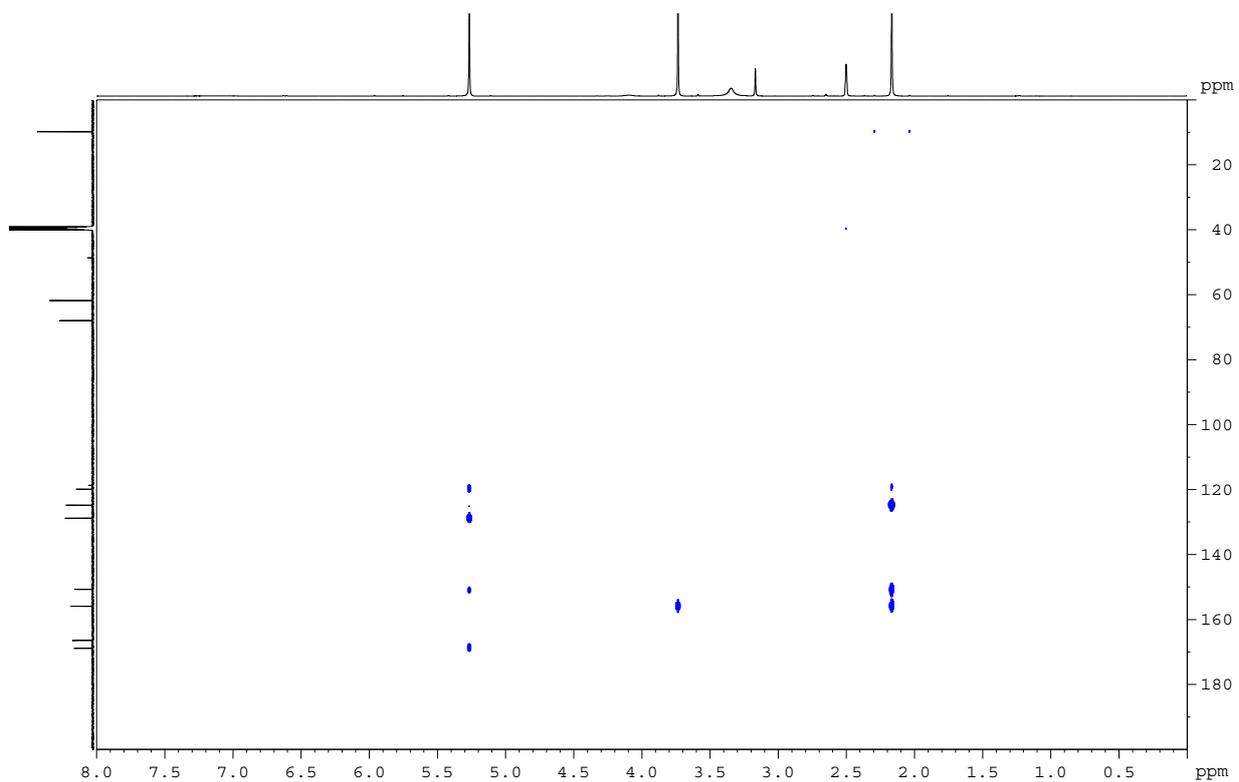
Figure S24. <sup>1</sup>H NMR spectrum (500 MHz, DMSO-*d*<sub>6</sub>) of 4



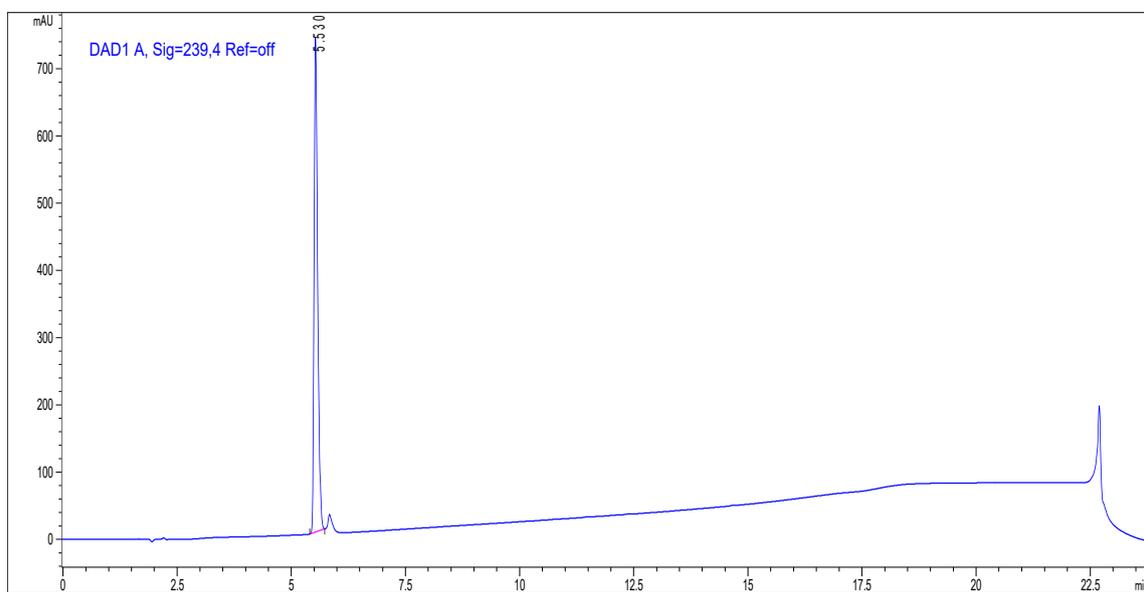
**Figure S25.**  $^{13}\text{C}$  NMR spectrum (125 MHz,  $\text{DMSO-}d_6$ ) of **4**



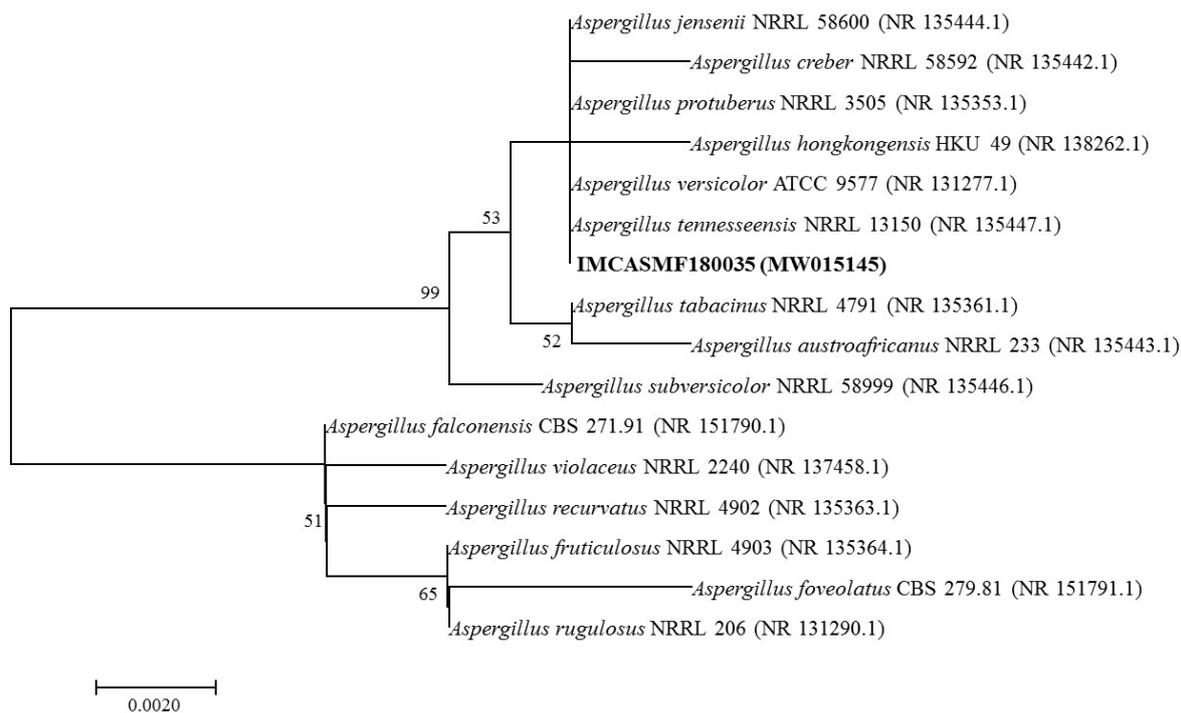
**Figure S26.** HSQC spectrum (500 MHz,  $\text{DMSO-}d_6$ ) of **4**



**Figure S27.** HMBC spectrum (500 MHz, DMSO-*d*<sub>6</sub>) of **4**



**Figure S28.** HPLC profile of **4**



**Figure S29.** Neighbor-joining phylogenetic tree of IMCASM180035 and its most related type strains based on internal transcribed spacer region (ITS) from NCBI ITS database. Numbers at nodes indicate levels of bootstrap support (%) based on a neighbor joining analysis of 1,000 resampled datasets; only values >50 % are given. NCBI accession numbers are given in parentheses. Bar 0.002 nucleotide substitutions per site.