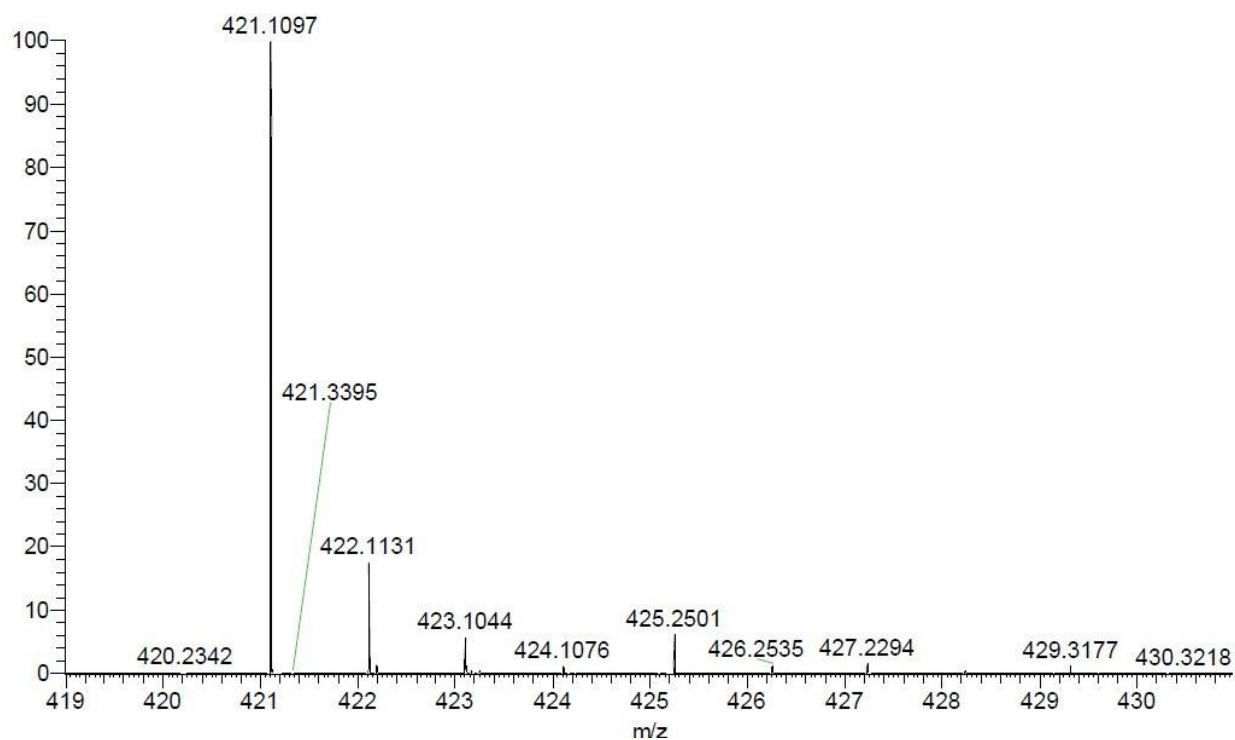
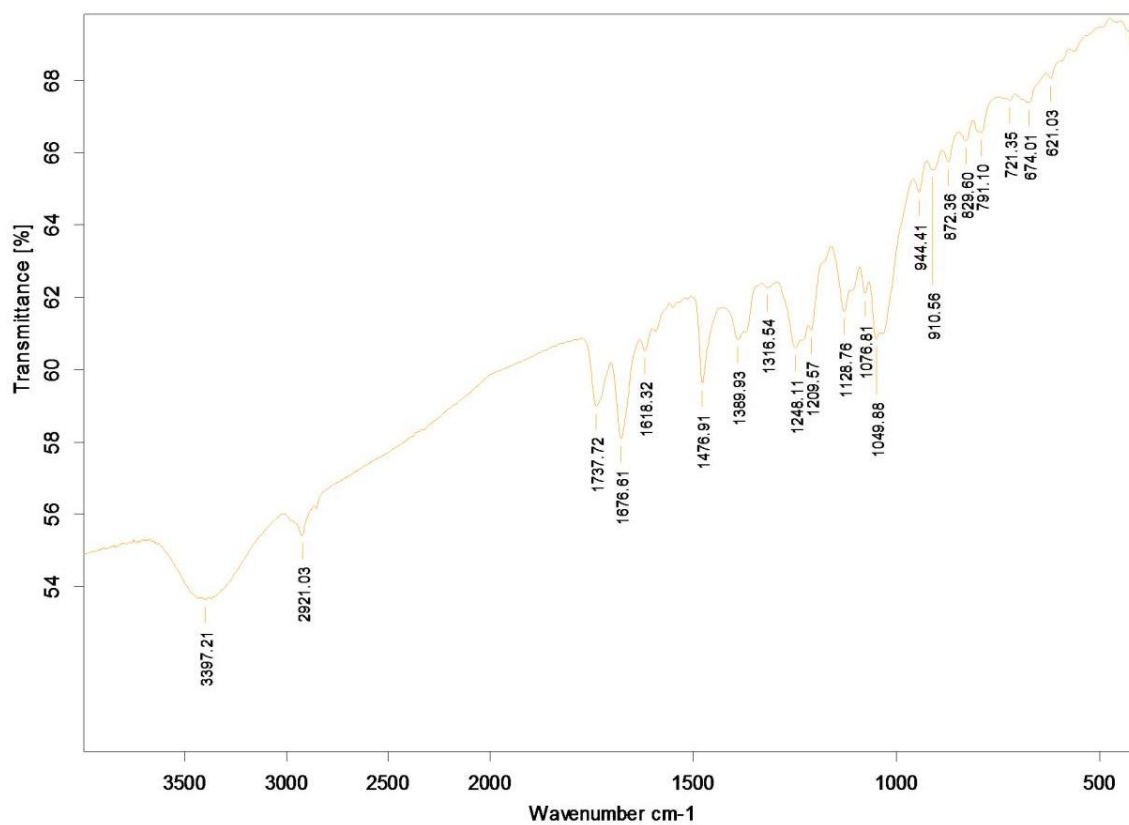


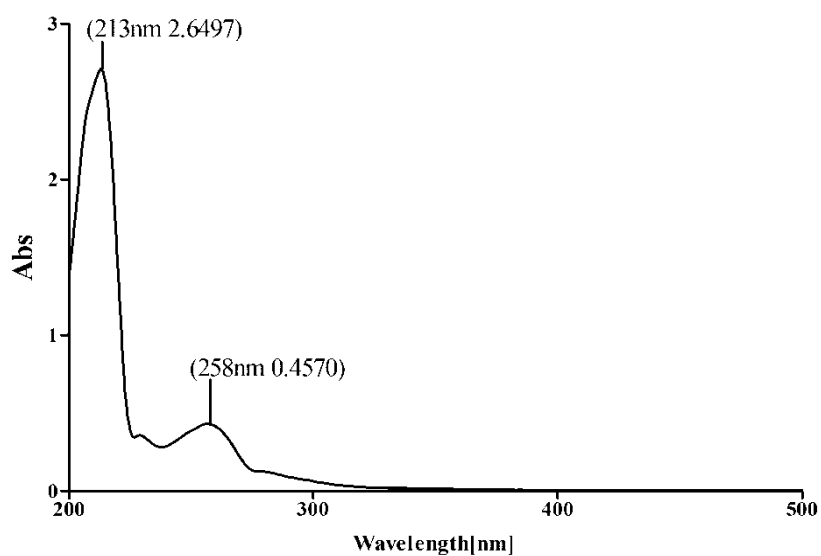
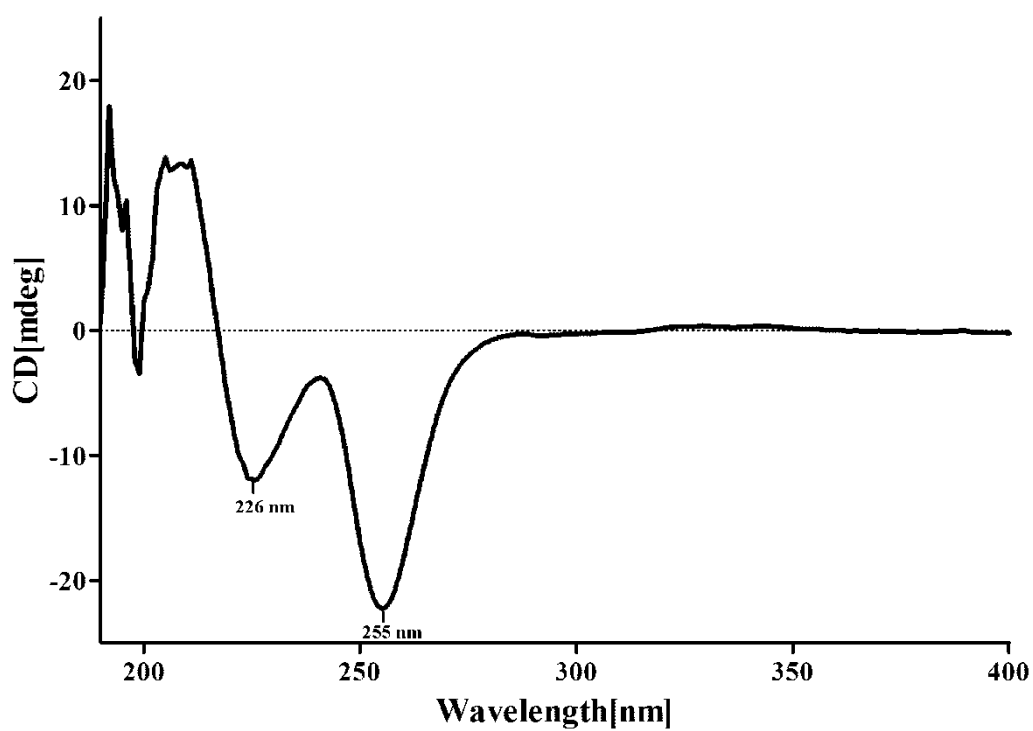
# Supplementary Materials

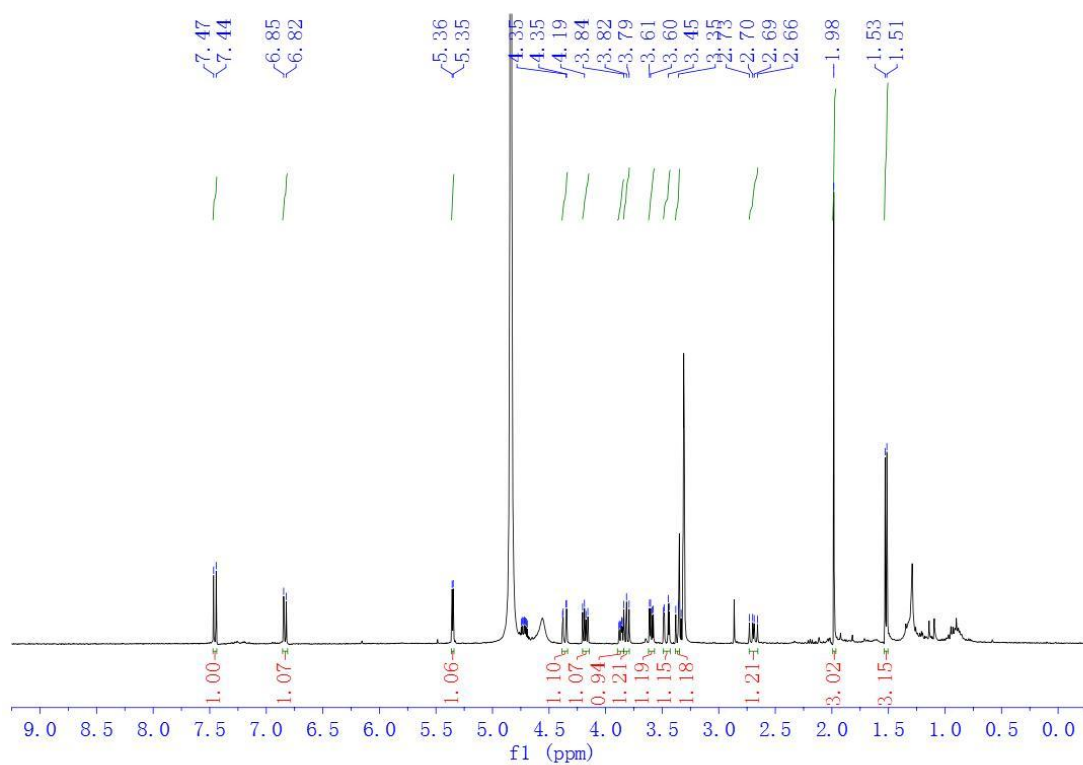
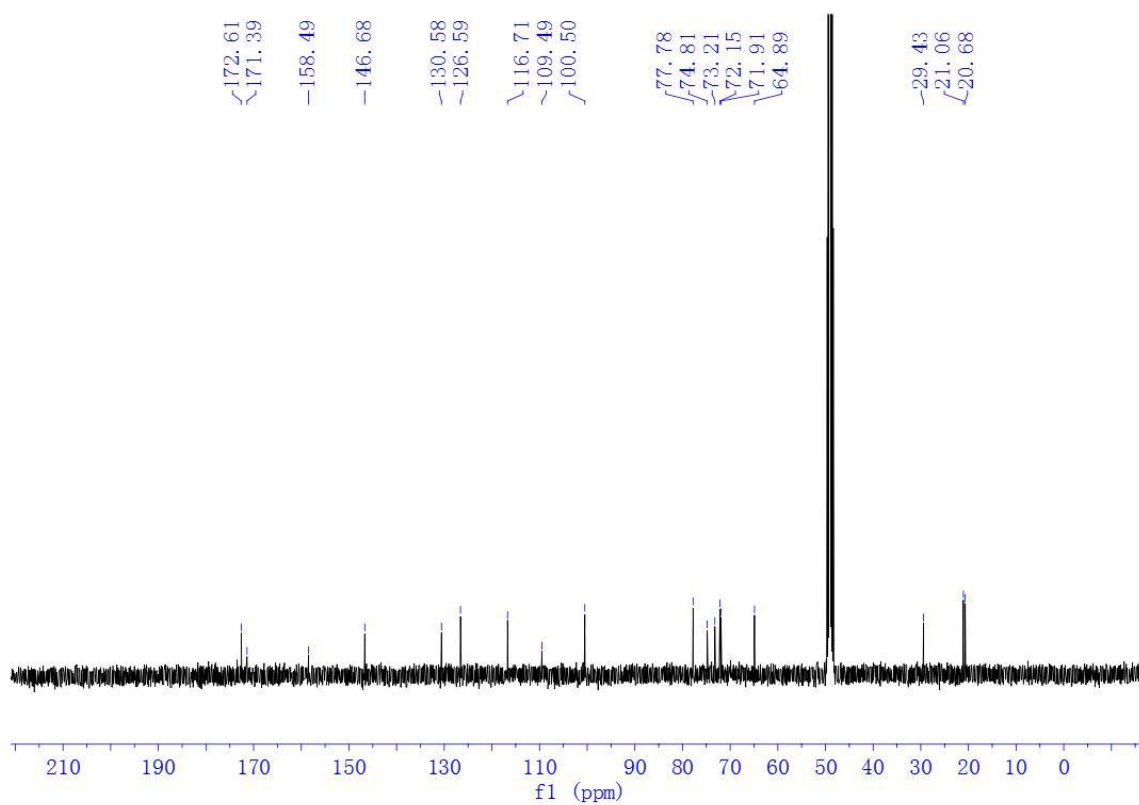
**Figure S1.** HRESIMS spectrum of compound **1**.

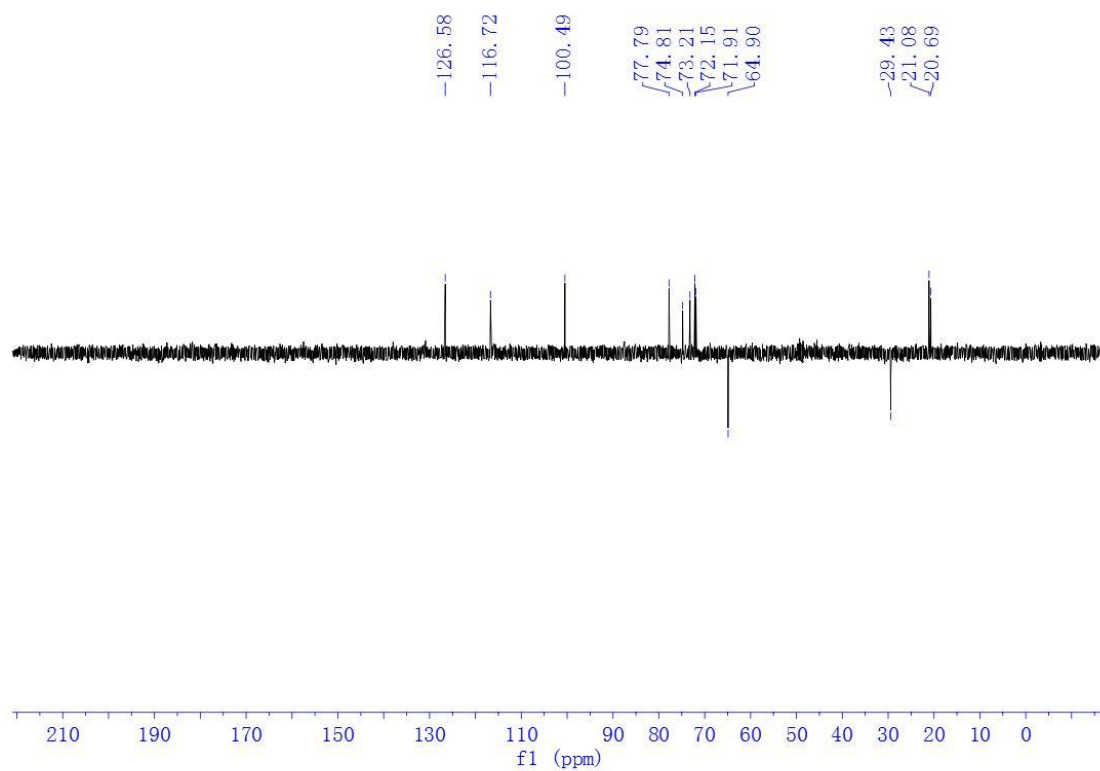
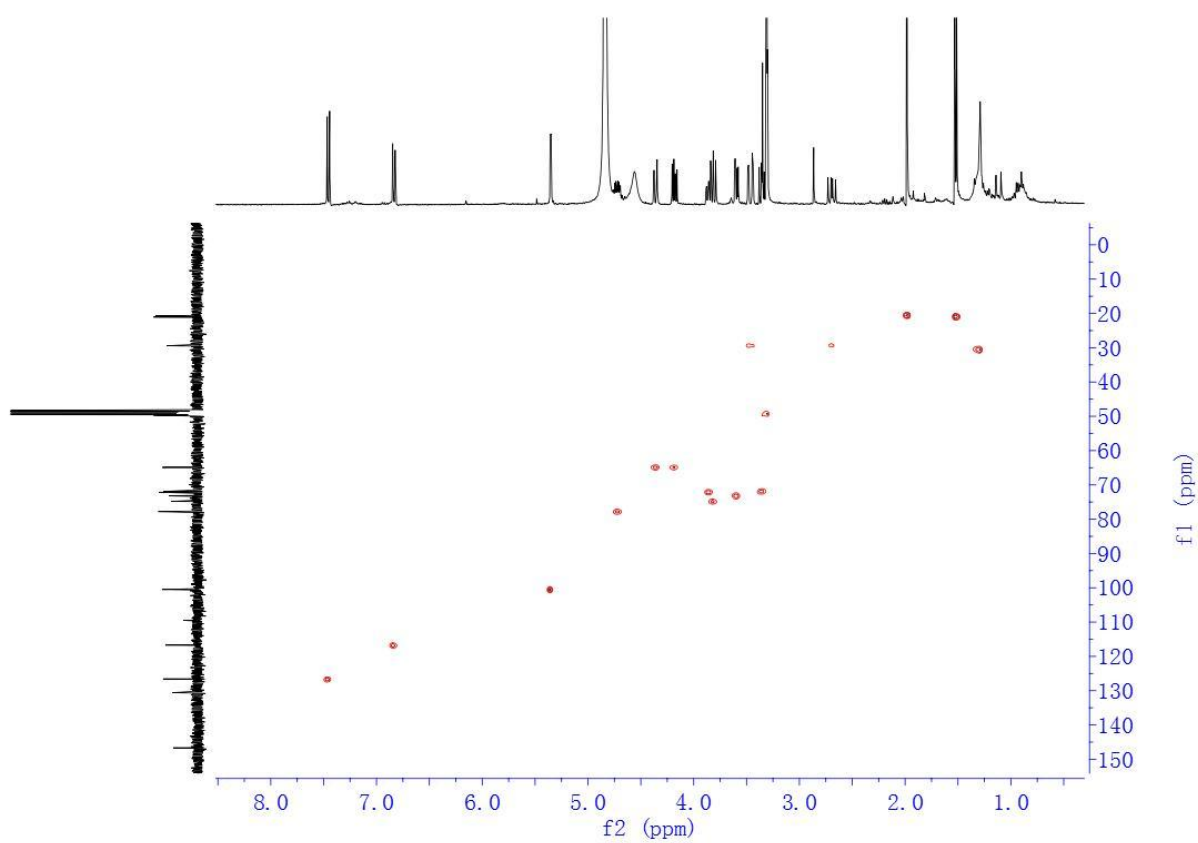


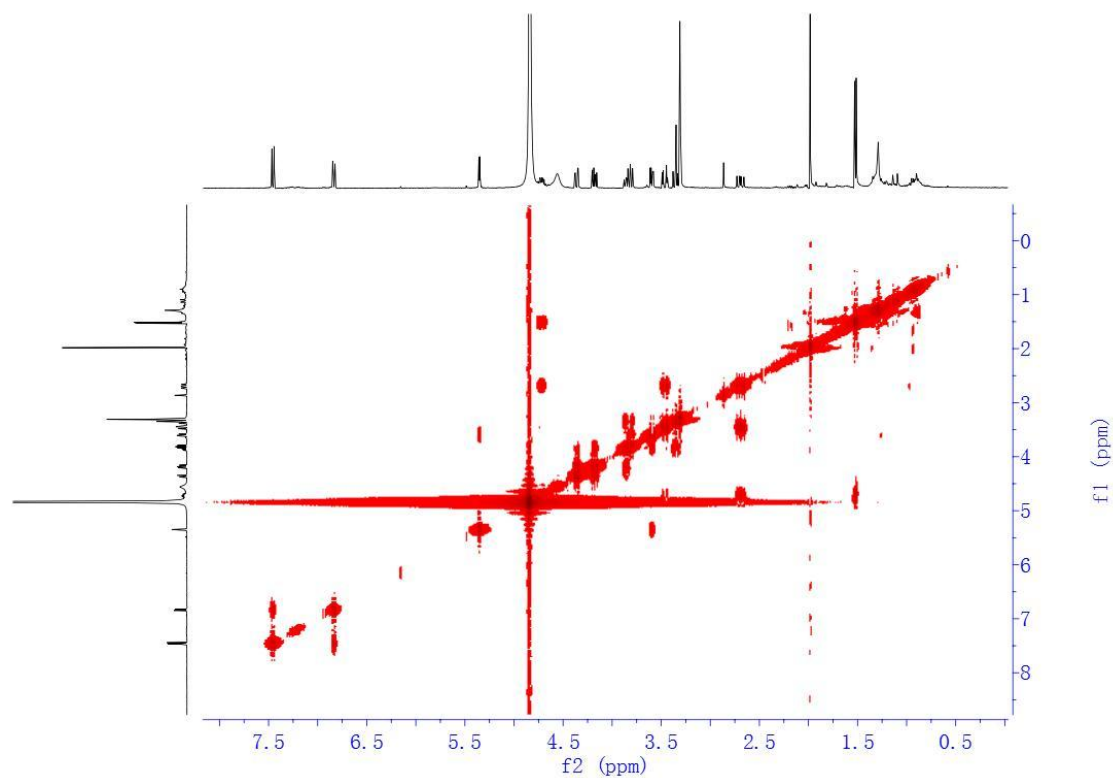
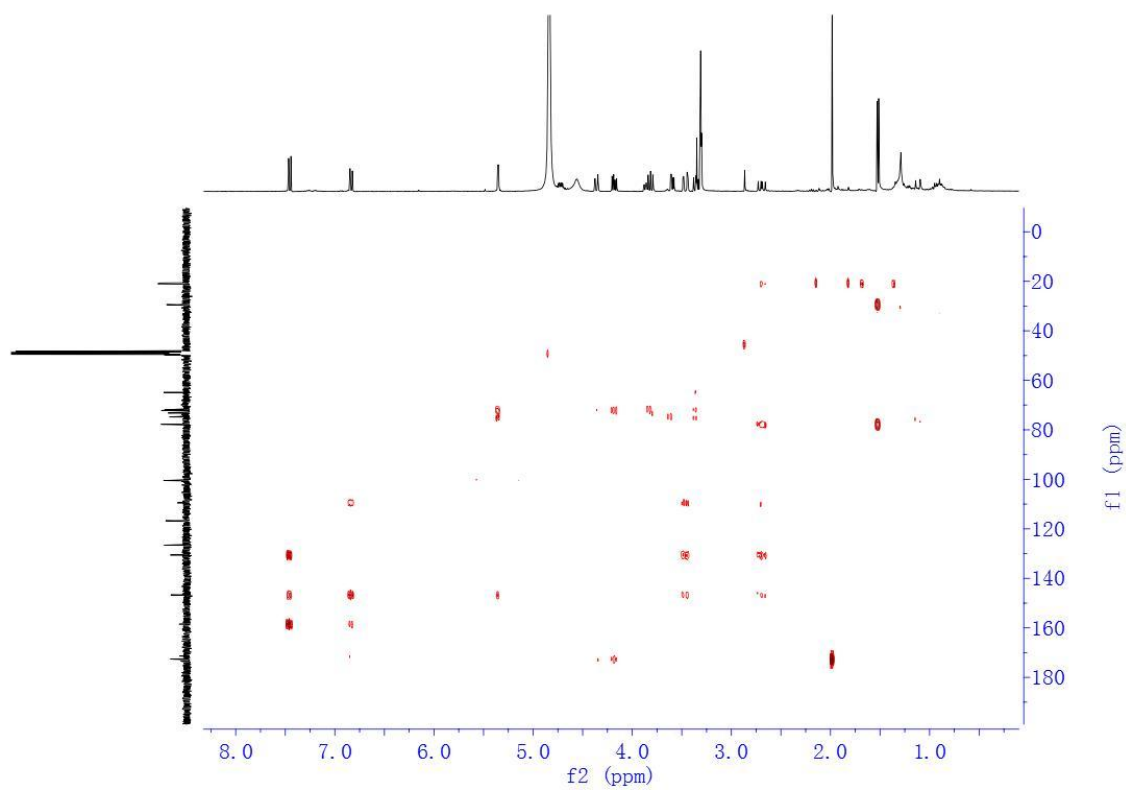
**Figure S2.** IR spectrum of compound **1**.



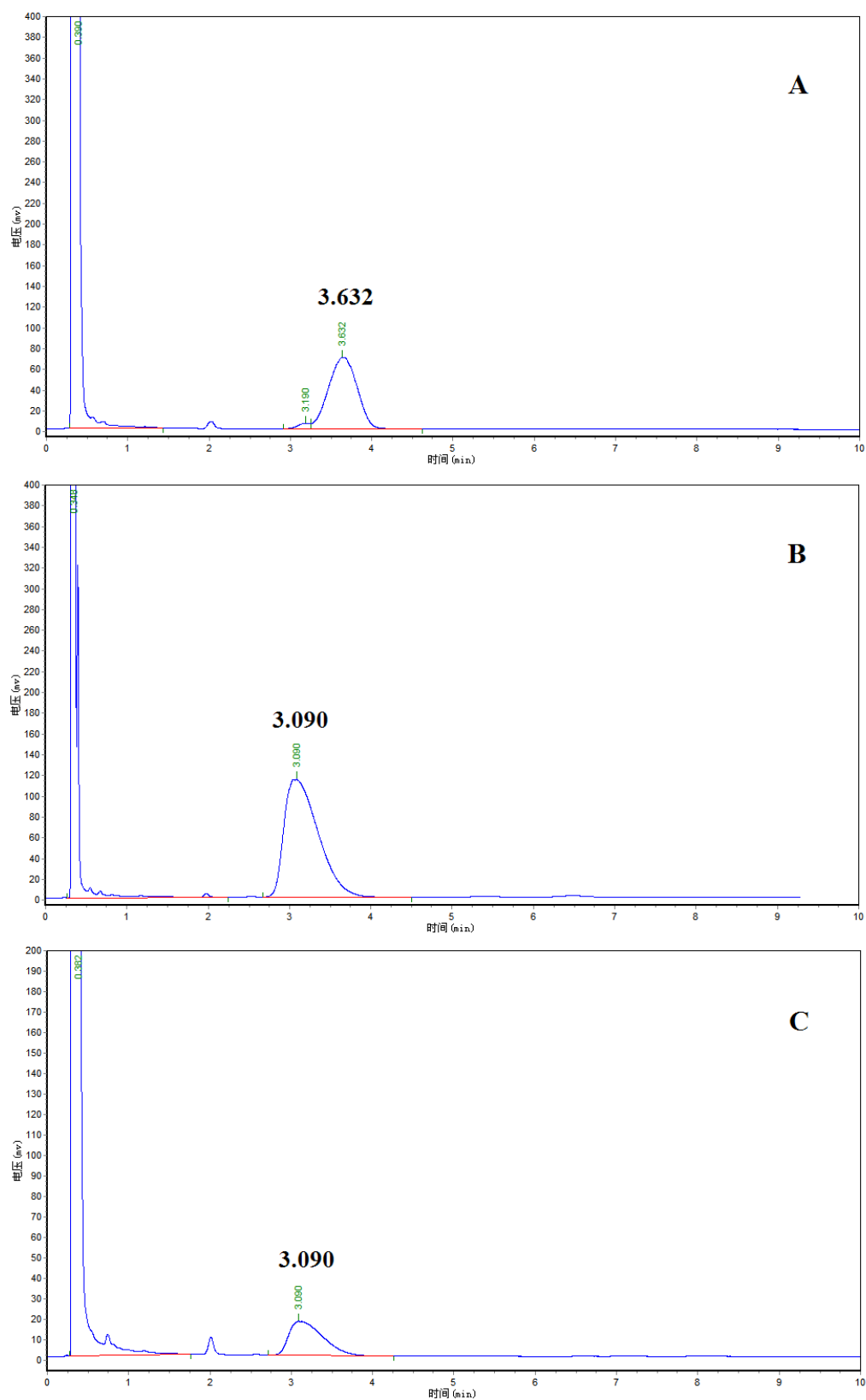
**Figure S3.** UV spectrum of compound 1.**Figure S4.** CD spectrum of compound 1.

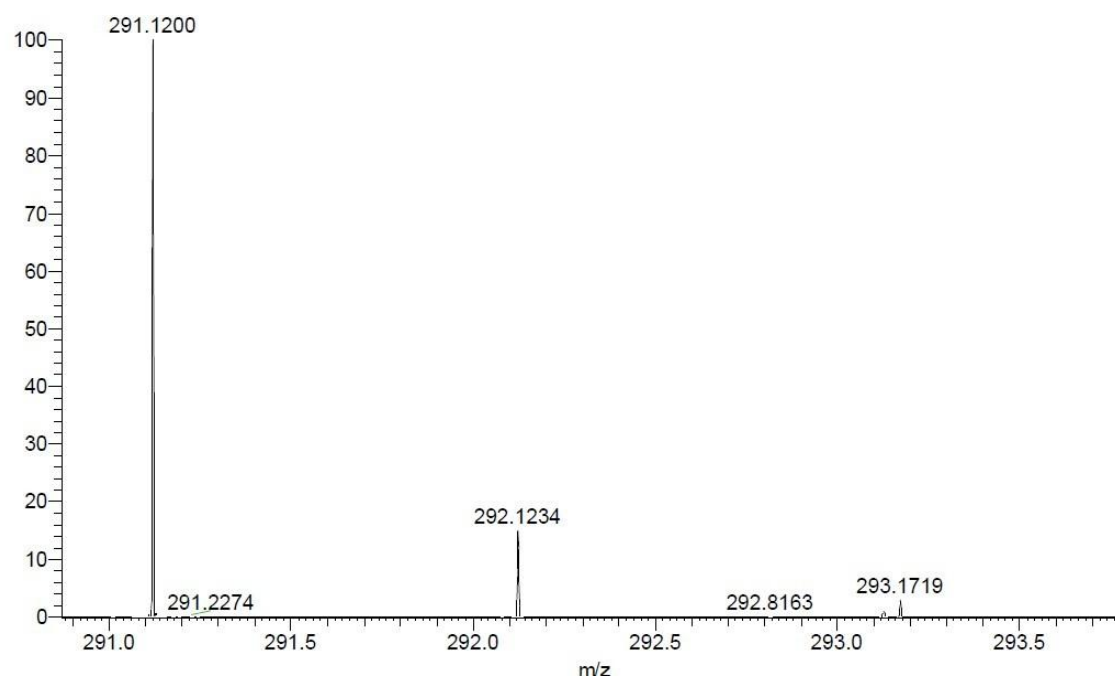
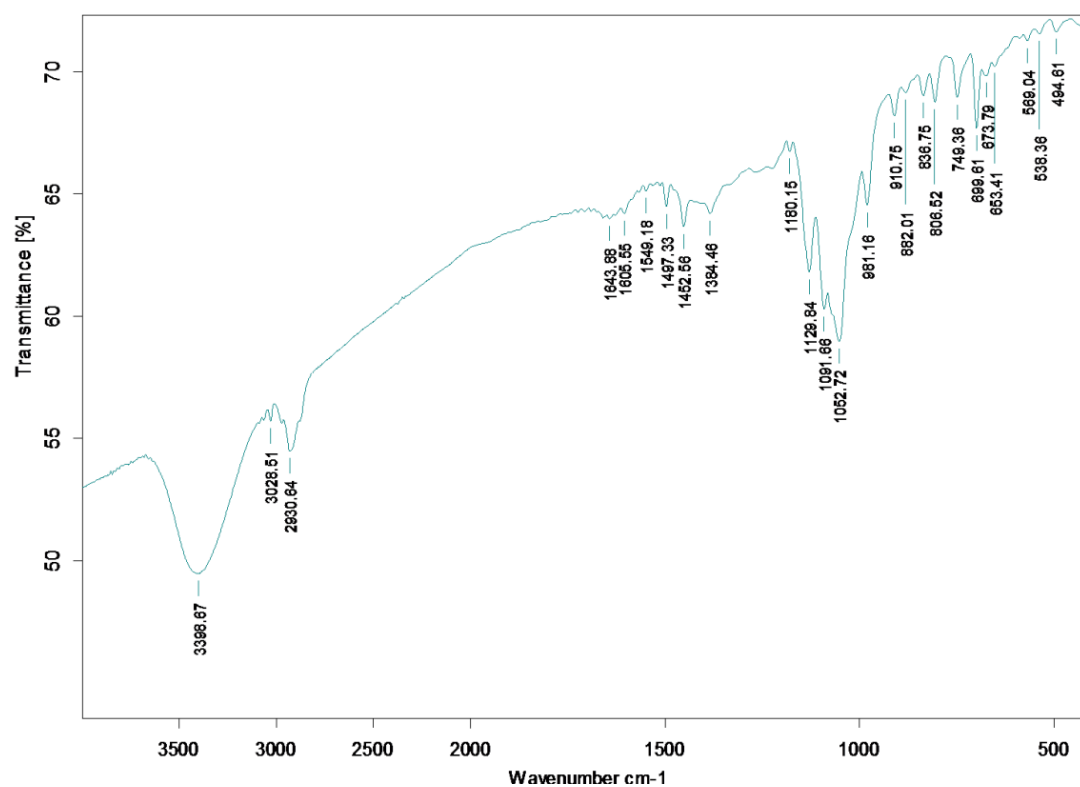
**Figure S5.**  $^1\text{H}$ -NMR spectrum of compound **1**.**Figure S6.**  $^{13}\text{C}$ -NMR spectra of compound **1**.

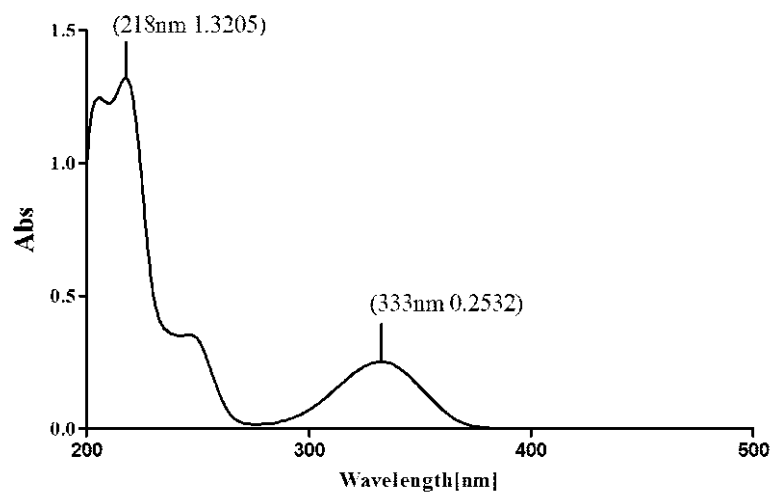
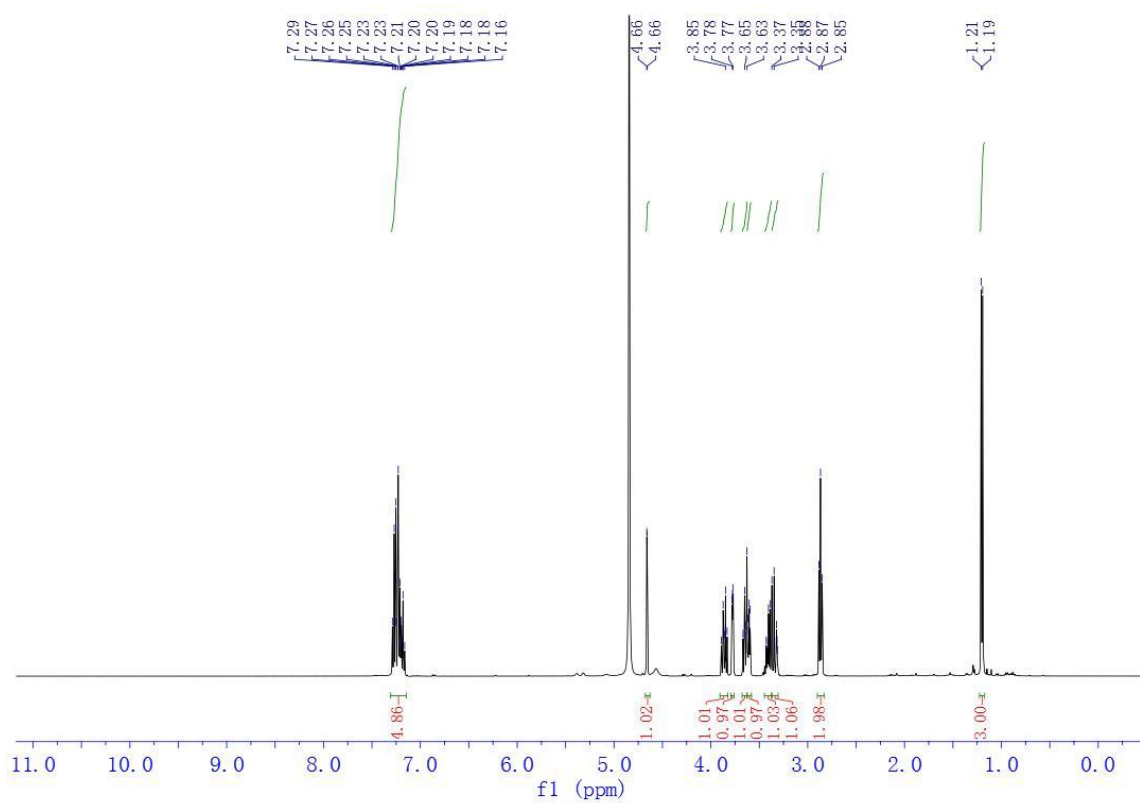
**Figure S7.** DEPT spectra of compound **1**.**Figure S8.** HSQC spectrum of compound **1**.

**Figure S9.** COSY spectrum of compound **1**.**Figure S10.** HMBC spectrum of compound **1**.

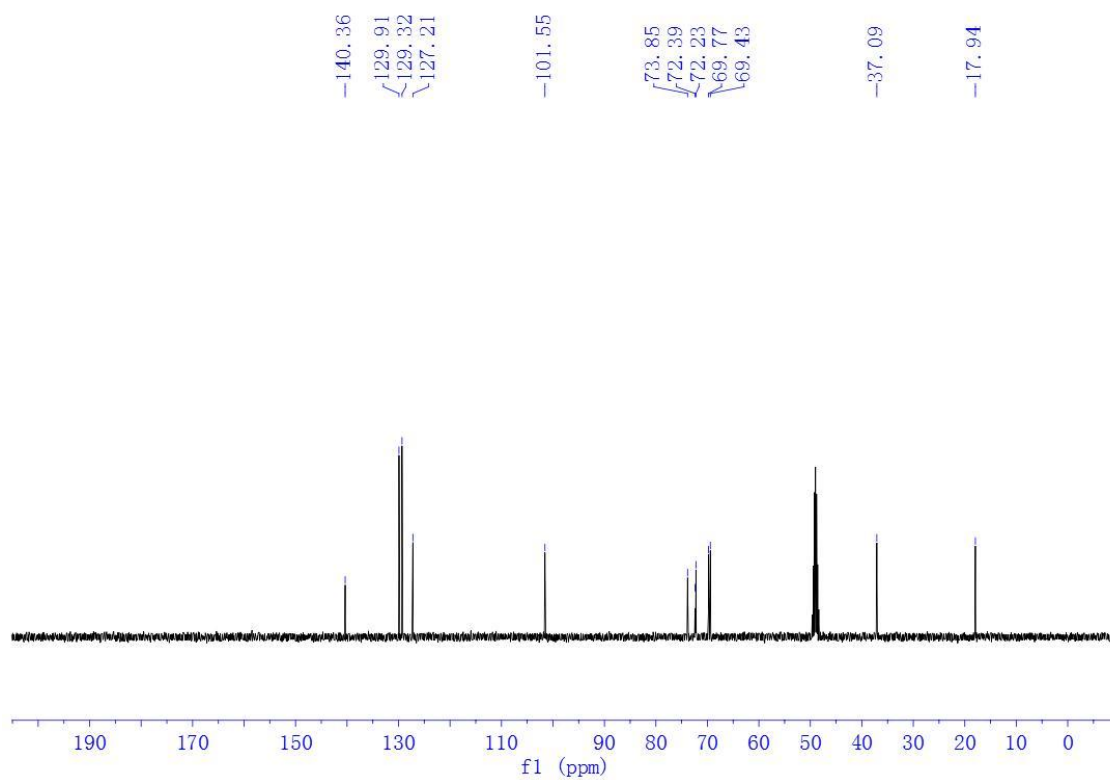
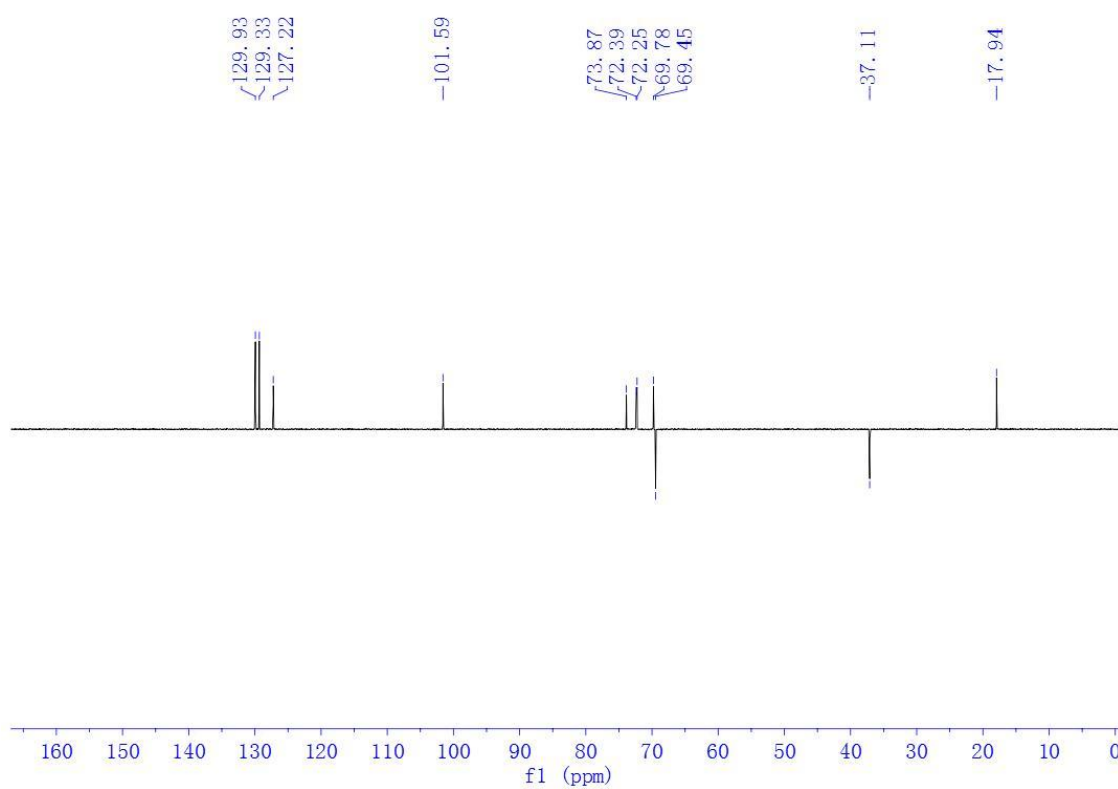
**Figure S11.** GS analysis of the sugar of compound **1** (**A**: L-glucose,  $t_R = 3.632$  min; **B**: D-Glucose,  $t_R = 3.090$  min; **C**: sugar of compound **1**,  $t_R = 3.090$  min).

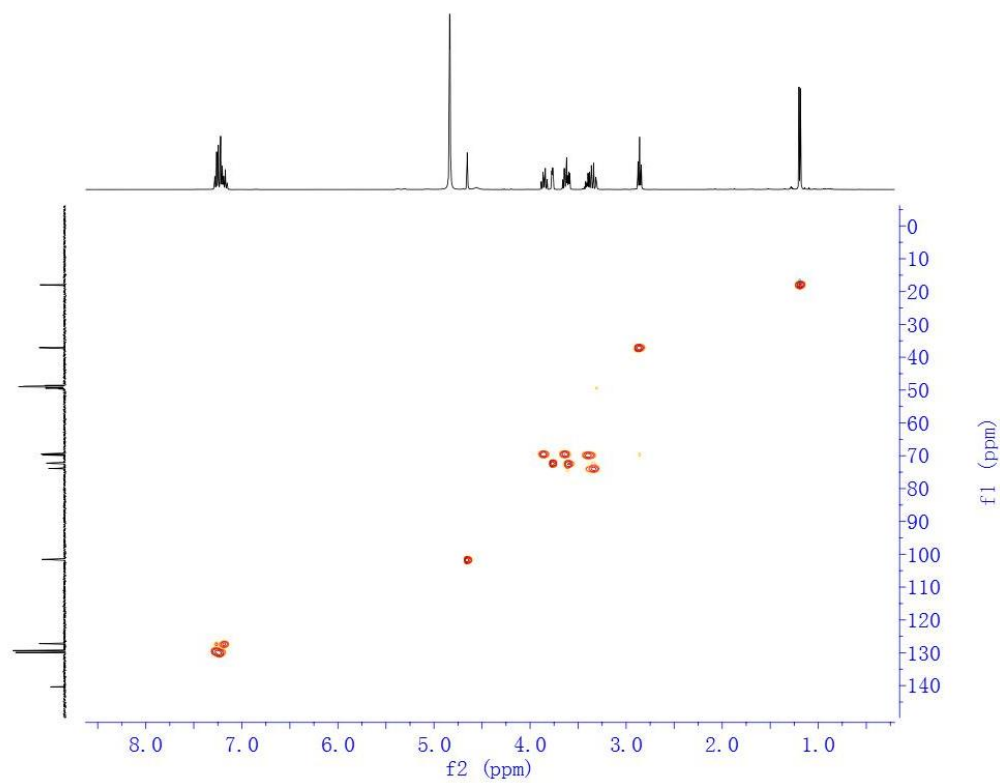
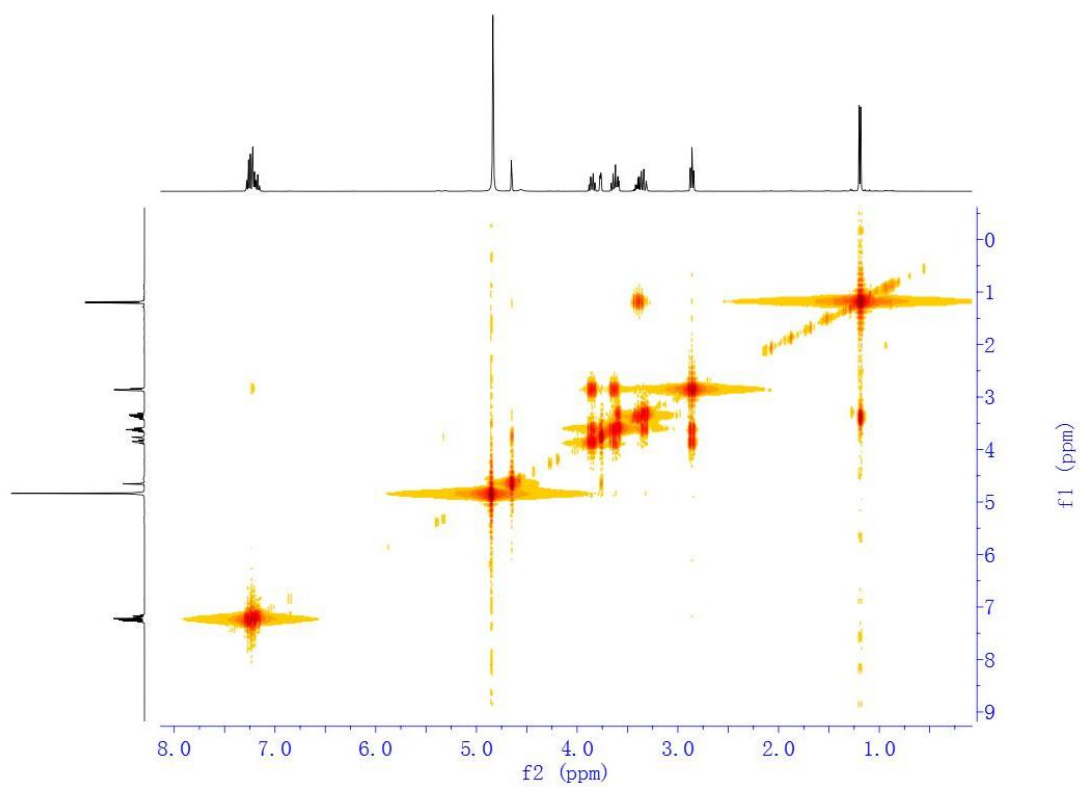


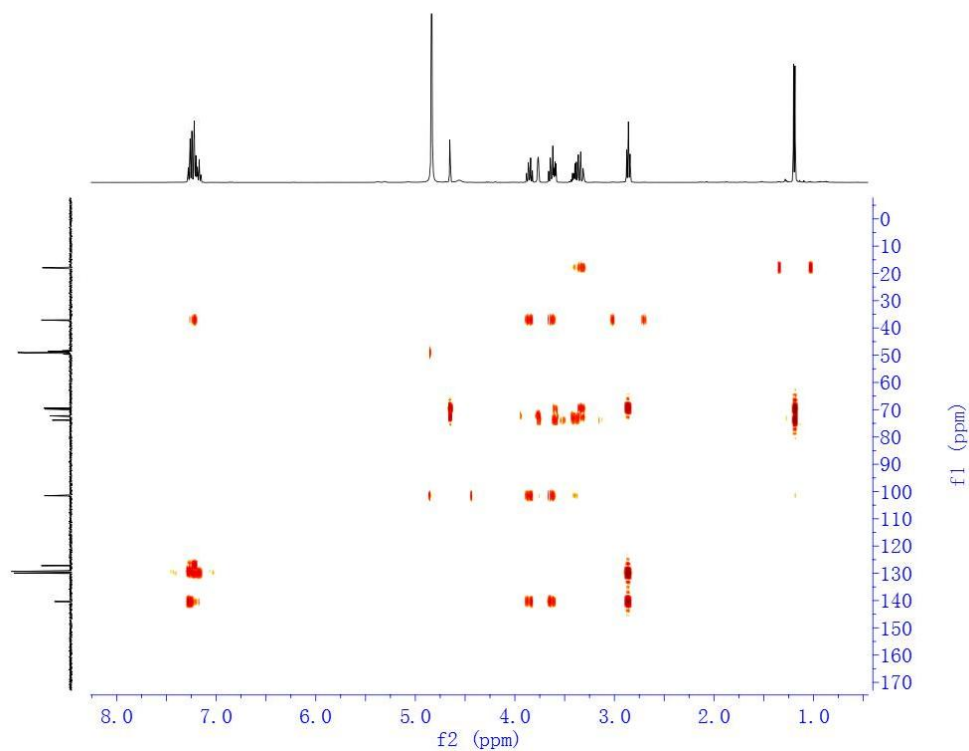
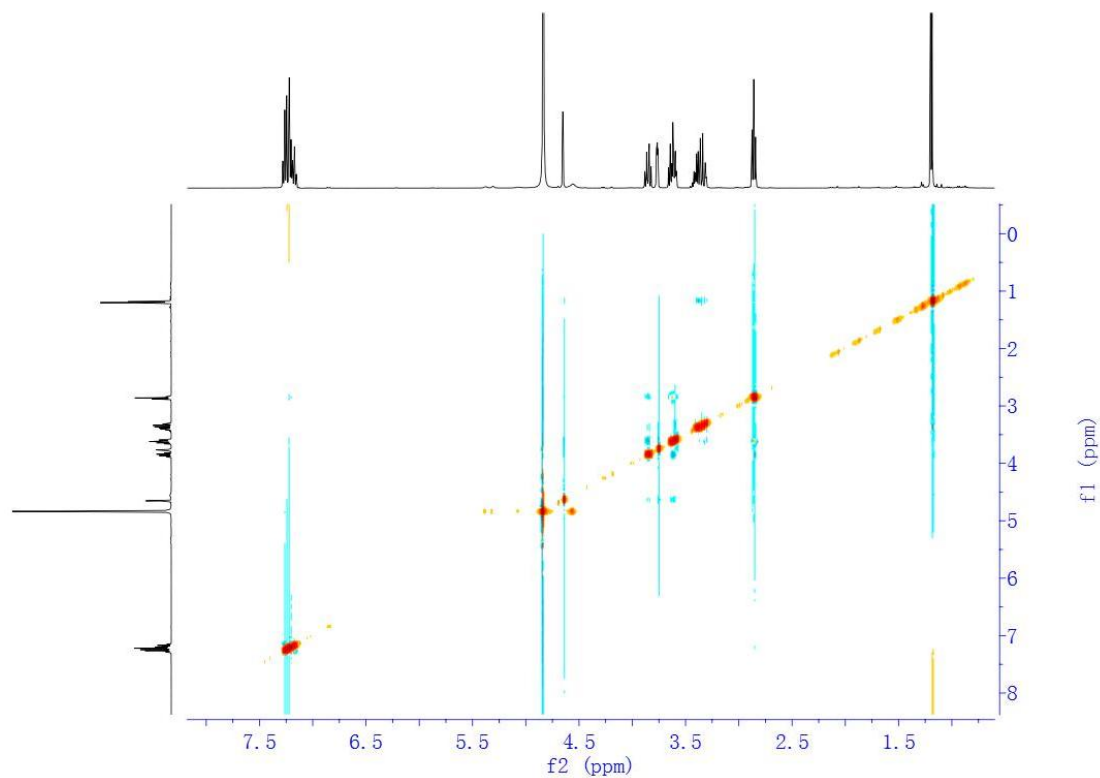
**Figure S12.** HRESIMS spectrum of compound 2.**Figure S13.** IR spectrum of compound 2.

**Figure S14.** UV spectrum of compound 2.**Figure S15.**  $^1\text{H}$ -NMR spectrum of compound 2.

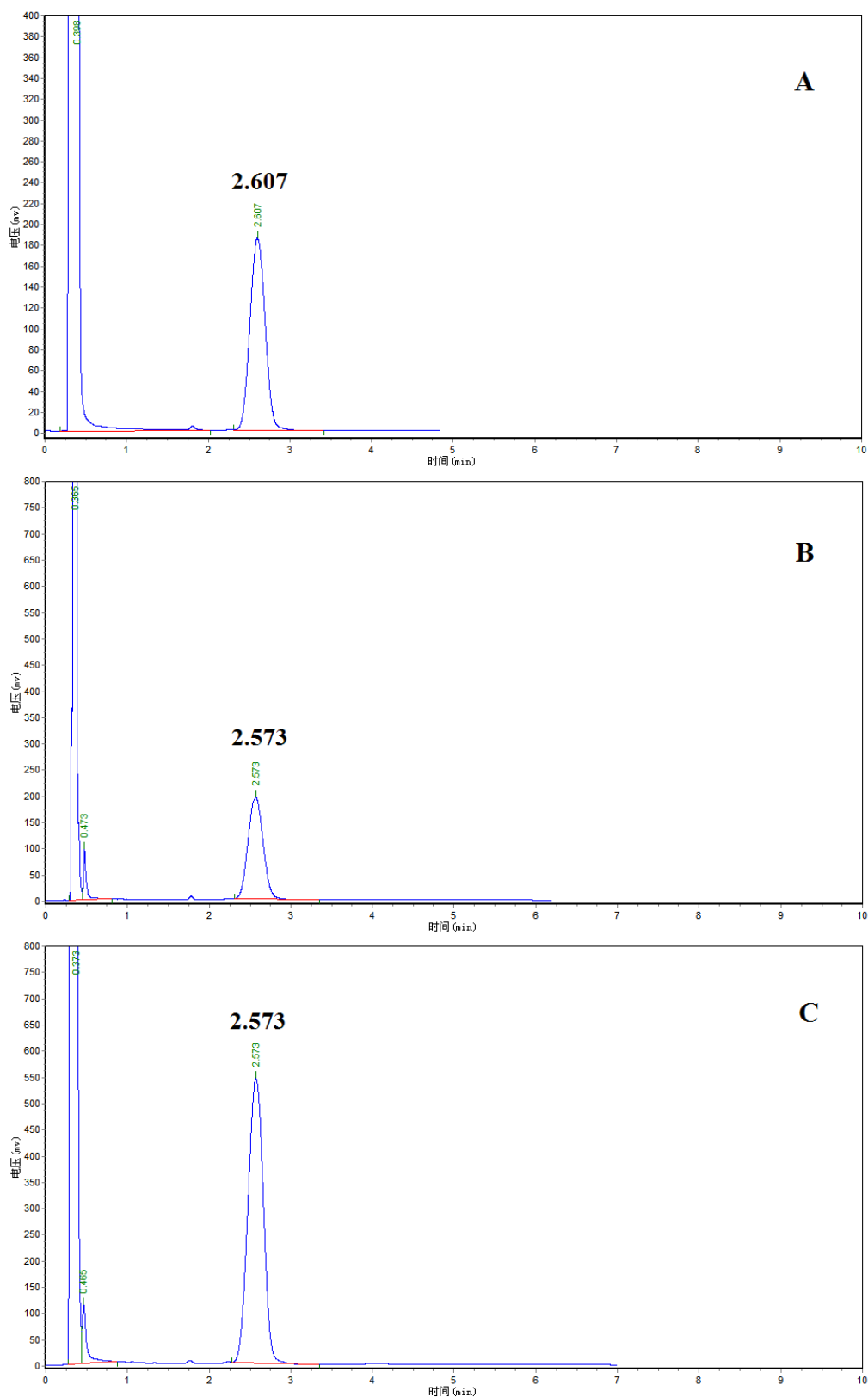


**Figure S16.**  $^{13}\text{C}$ -NMR spectra of compound 2.**Figure S17.** DEPT spectra of compound 2.

**Figure S18.** HSQC spectrum of compound 2.**Figure S19.** COSY spectrum of compound 2.

**Figure S20.** HMBC spectrum of compound 2.**Figure S21.** NOSEY spectrum of compound 2.

**Figure S22.** GS analysis of the sugar of compound **2** (**A**: L-rhamnose,  $t_R = 2.607$  min; **B**: sugar of compound **2**,  $t_R = 2.573$  min; **C**: A mixture 1:1 of L-rhamnose and sugar of compound **2**,  $t_R = 2.573$  min).



**Figure S23.** Pictures of the colony and the culture of *T. marneffeii*.**Figure S24.** Phylogenetic analysis of ITS1-4 gene sequence.

GATGATGTATACTCCAACCCATGTGACATACCTCATGTTGCCTCGGCAGGTCGTGCCTCCCT  
 CGTAGGTCCTACCCTGTAGGCTCCTACCCGGAAGGCGCGGGTACCCCTGCCGGTGGCCCAG  
 GAAACTCTGTCTCATCGTTGAATTCTGAACCTATAACTAAATAAGTTAAAACCTTTCAACAAC  
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