

# **Different coordination abilities of 1,7- and 4,7-phenanthroline in the reactions with copper(II) salts: structural characterization and biological evaluation of the reaction products**

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## Abstract

The reactions between equimolar amounts of CuX<sub>2</sub> (X = NO<sub>3</sub><sup>-</sup> and CF<sub>3</sub>SO<sub>3</sub><sup>-</sup>) and two aromatic nitrogen-containing heterocycles differing in the position of nitrogen atoms, 1,7- and 4,7-phenanthroline (1,7- and 4,7-phen), were performed in ethanol/methanol at room temperature. When CuX<sub>2</sub> salts were mixed with 4,7-phen, two copper(II) complexes, [Cu(NO<sub>3</sub>)<sub>2</sub>(4,7-Hphen)<sub>2</sub>](NO<sub>3</sub>)<sub>2</sub> (**1**) and [Cu(CF<sub>3</sub>SO<sub>3</sub>)(4,7-phen)<sub>2</sub>(H<sub>2</sub>O)<sub>2</sub>]CF<sub>3</sub>SO<sub>3</sub> (**2**), were formed. On the other hand, in the reaction of CuX<sub>2</sub> salts with 1,7-phen, only 1,7-HphenNO<sub>3</sub> (**3a/b**) and 1,7-HphenCF<sub>3</sub>SO<sub>3</sub> (**4**) were obtained as the final products. The obtained products **1** – **4** were characterized by spectroscopic and X-ray diffraction techniques. In the copper(II) complexes **1** and **2**, the coordination geometry around the Cu(II) ion is distorted octahedral and square pyramidal, respectively. The antimicrobial potential of the copper(II) complexes **1** and **2** and corresponding compounds used for their synthesis were assessed against four different bacterial species and *Candida albicans*, displaying moderate growth inhibiting activity. The cytotoxic properties of the investigated complexes were also evaluated against the normal human lung fibroblast cell line (MRC-5) indicating moderate, yet more pronounced cytotoxicity than antimicrobial properties.

**Keywords:** Copper(II), Phenanthroline, Structural characterization, Antimicrobial activity, Cytotoxicity

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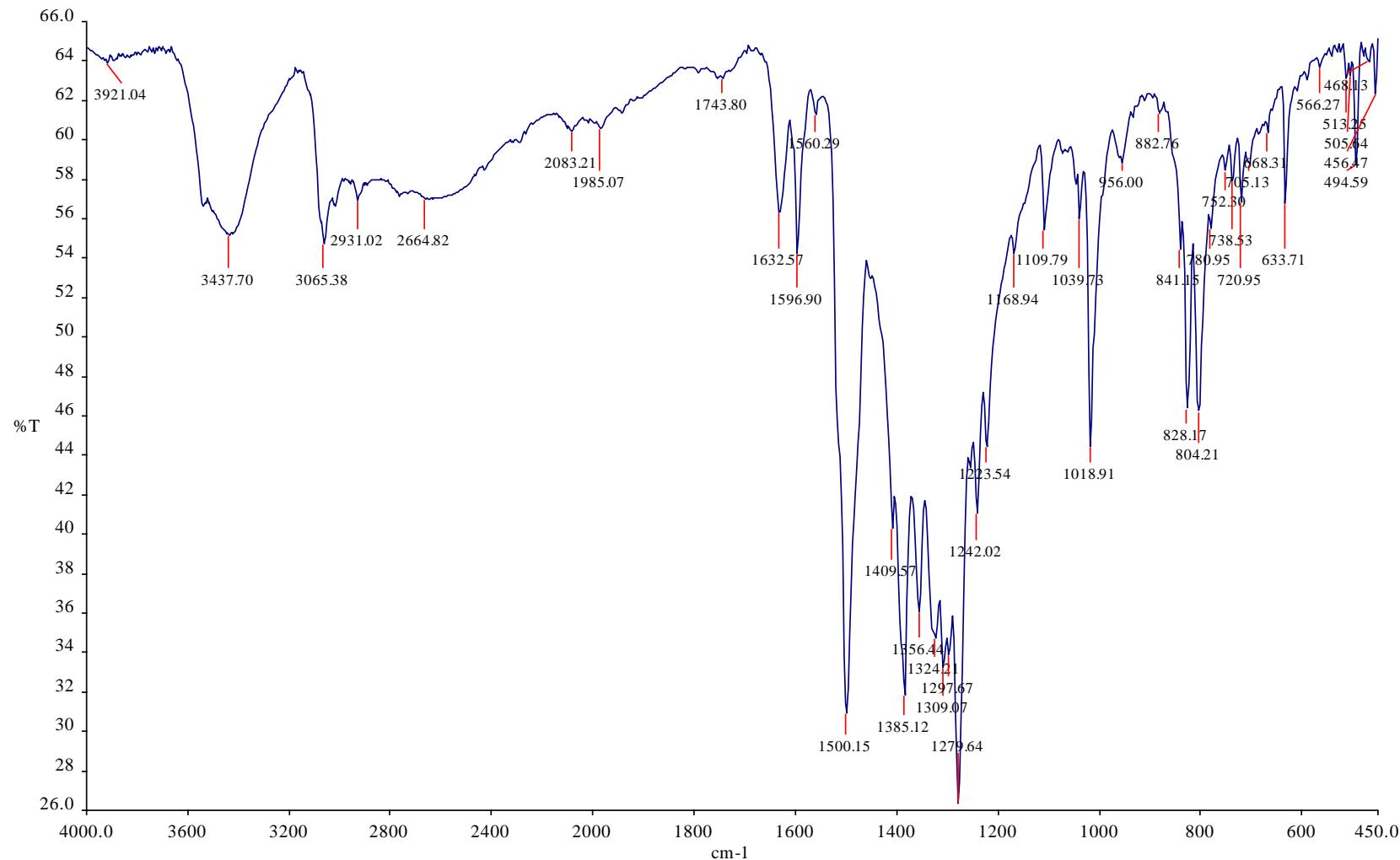
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| UV-Vis spectrum of <b>1</b>              | S5  |
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**Fig. S1.** Molecular structure of **2**. Some hydrogen atoms and partitioning of counter-ions are omitted for clarity. Hydrogen bonds are drawn as dashed blue lines, #1:  $-x+1, -y+1, -z+1$ .

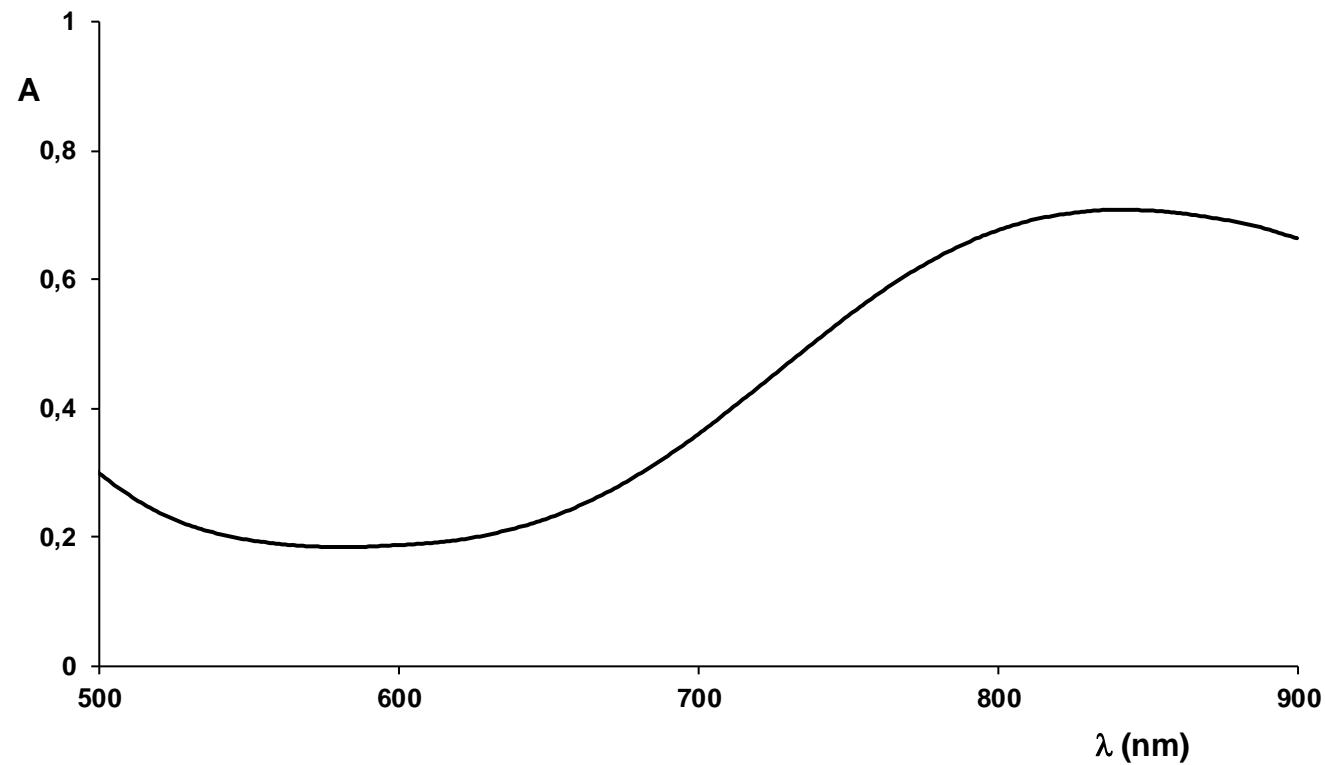
**Fig. S2.** Molecular structure of **3b**. Displacement ellipsoids are drawn at 50% probability level and H atoms are represented by spheres of arbitrary size.

**Fig. S3.** The overlay of the packing of the **3a** (red) and **3b** (blue) along the *a* (**a**), *b* (**b**) and *c* (**c**) axis.

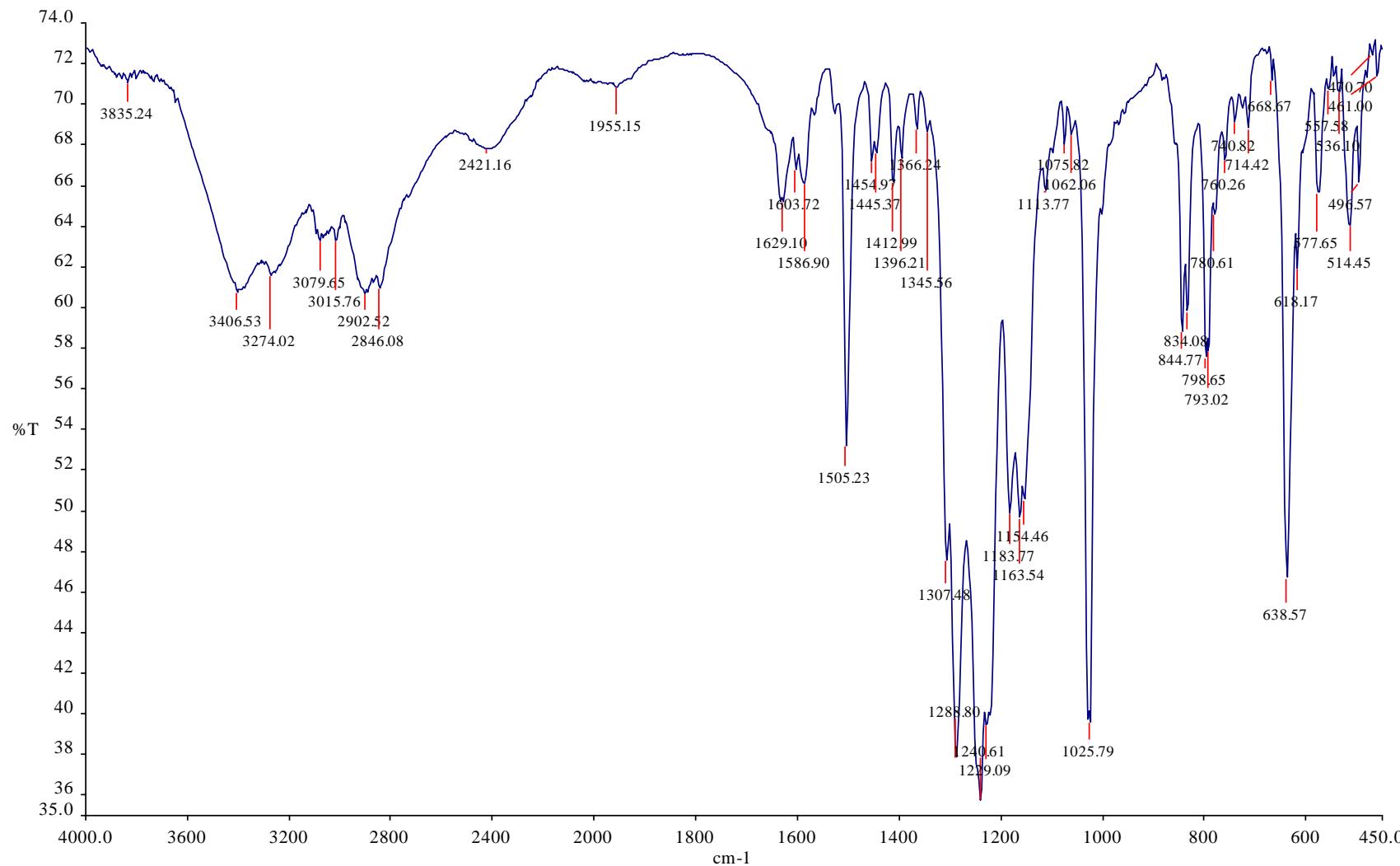
IR spectrum of **1**



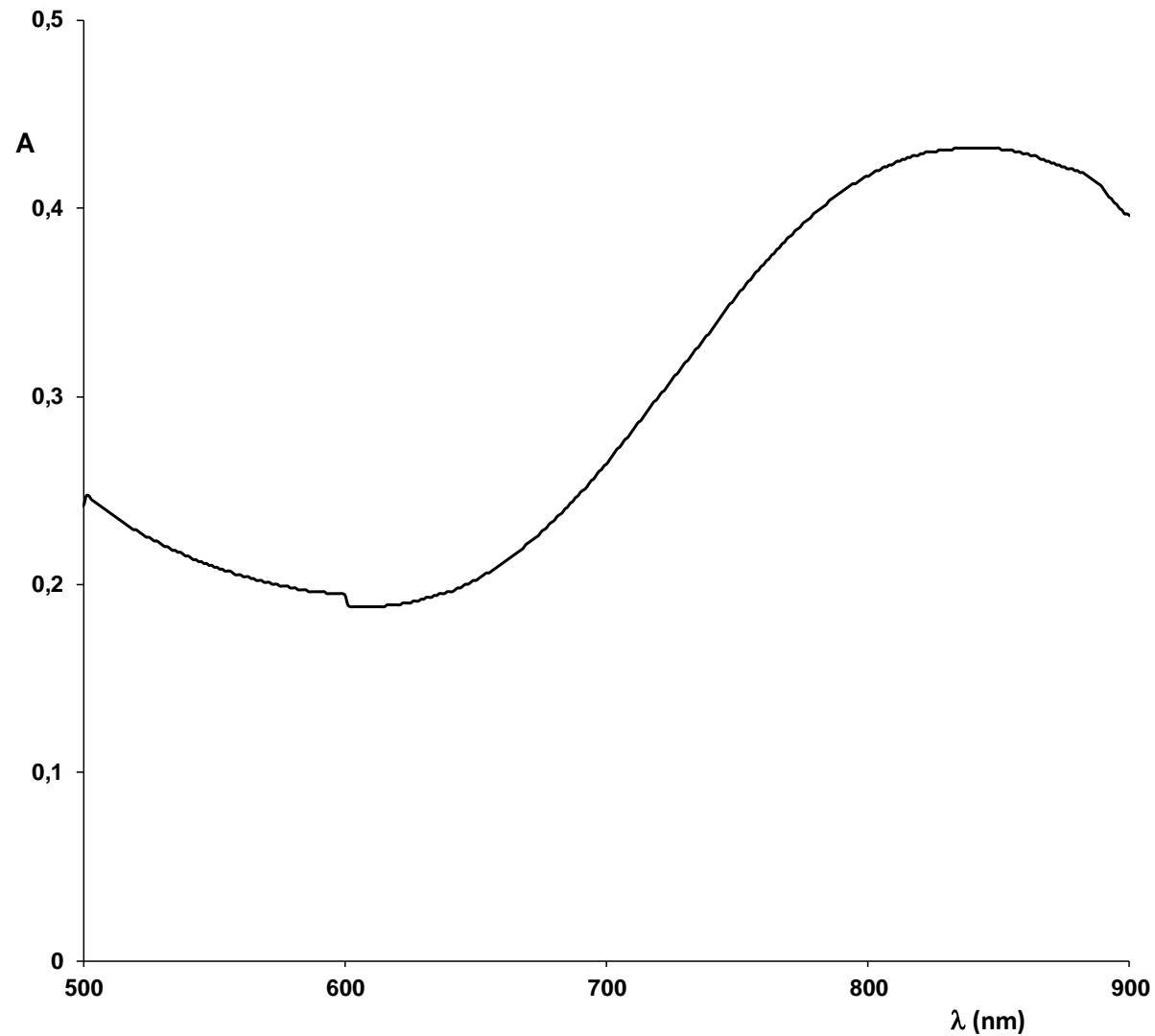
UV-Vis spectrum of **1** (DMSO)



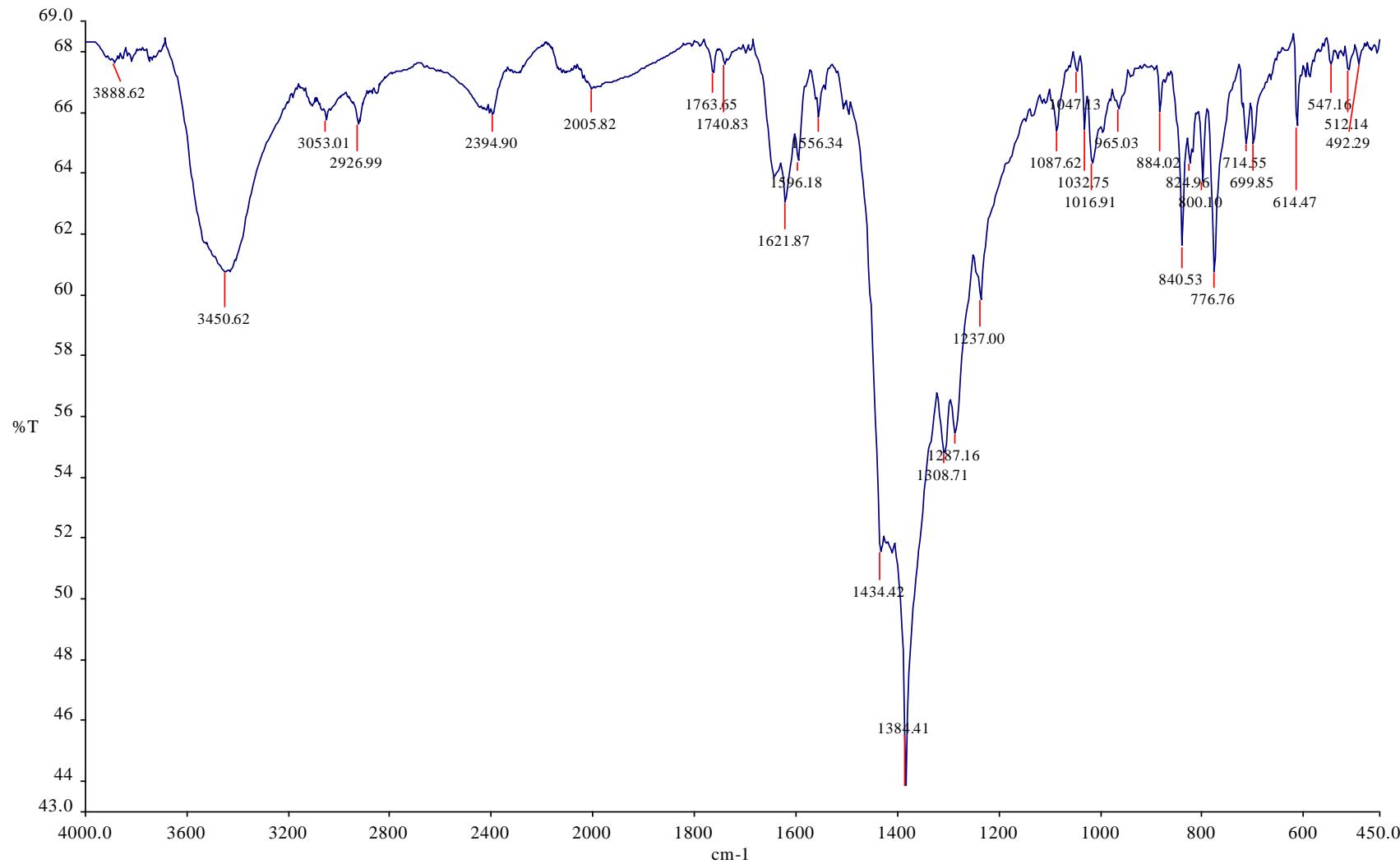
IR spectrum of **2**



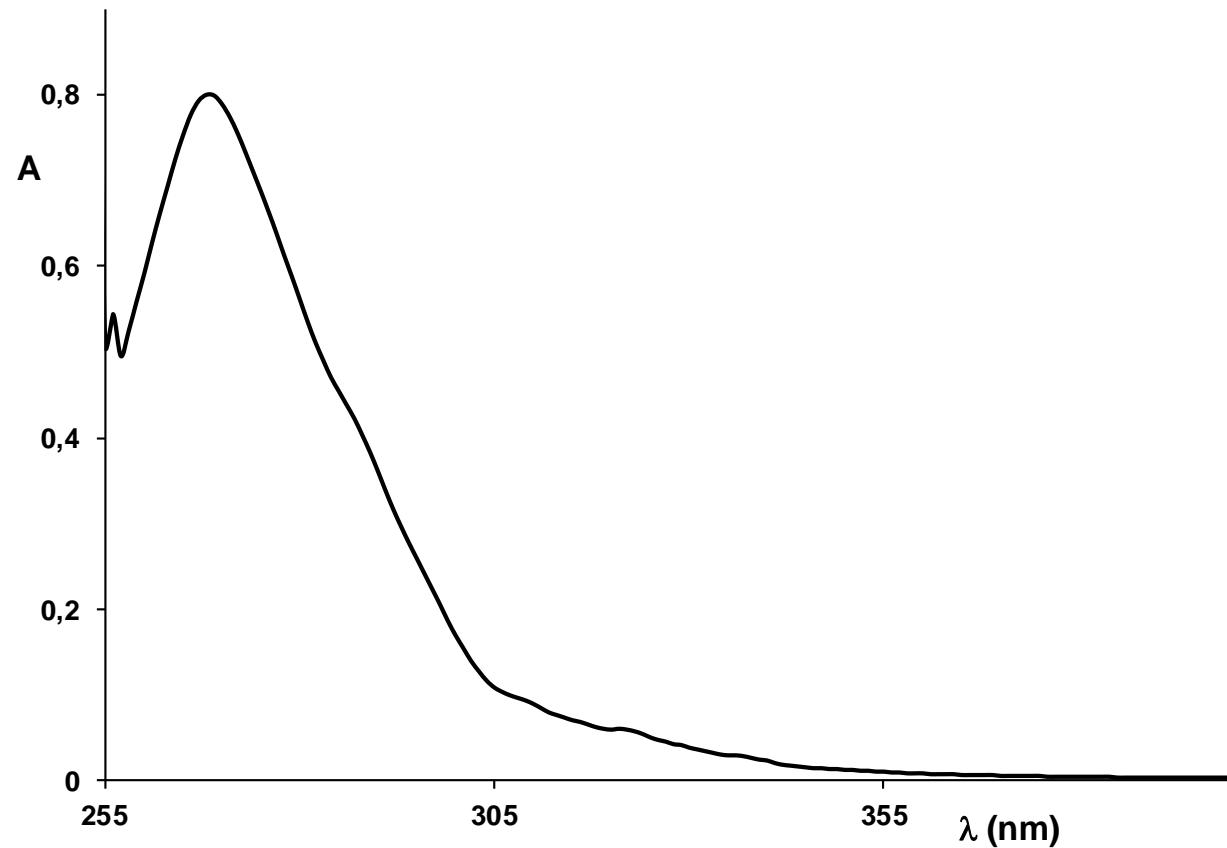
UV-Vis spectrum of **2** (DMSO)



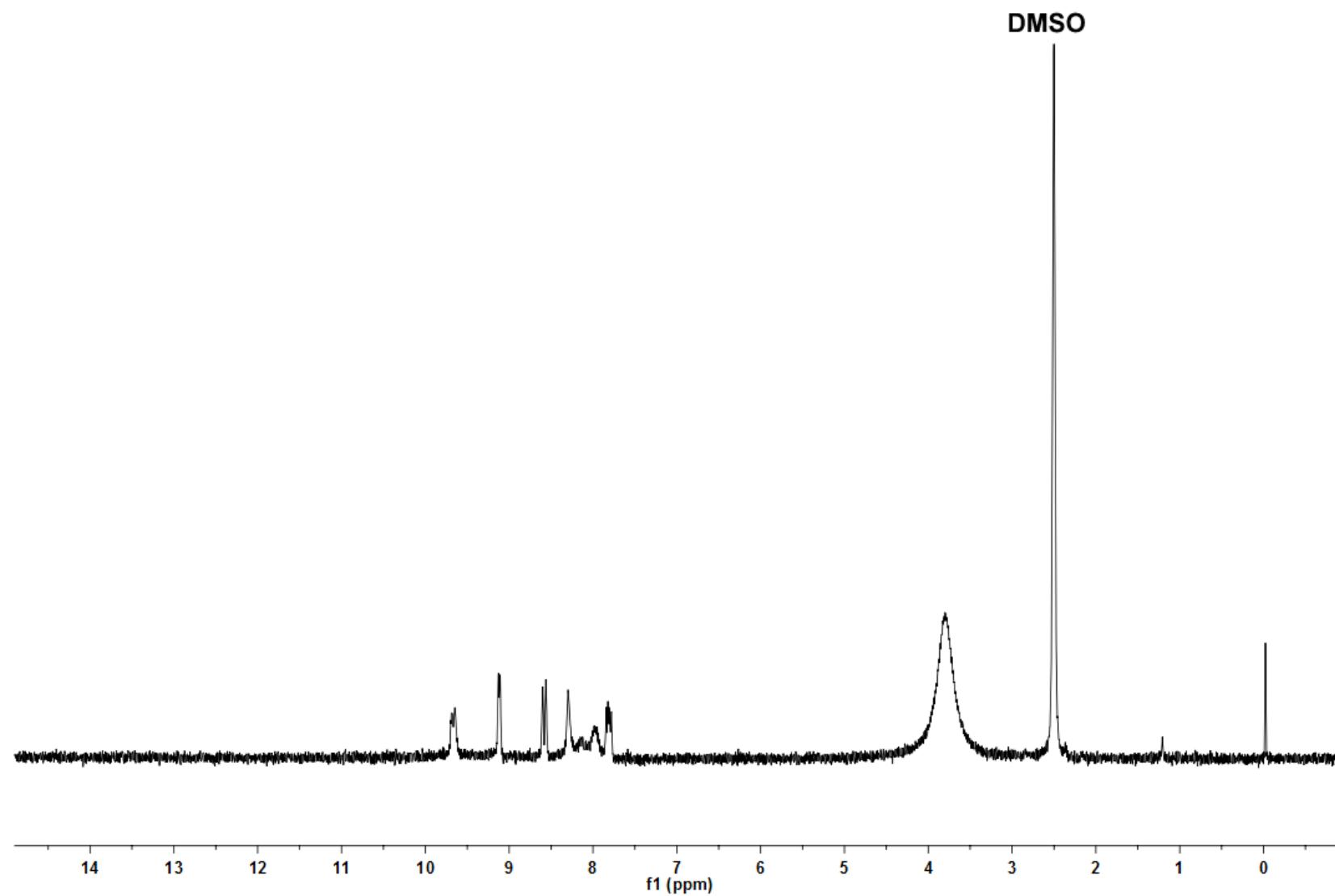
IR spectrum of **3a**



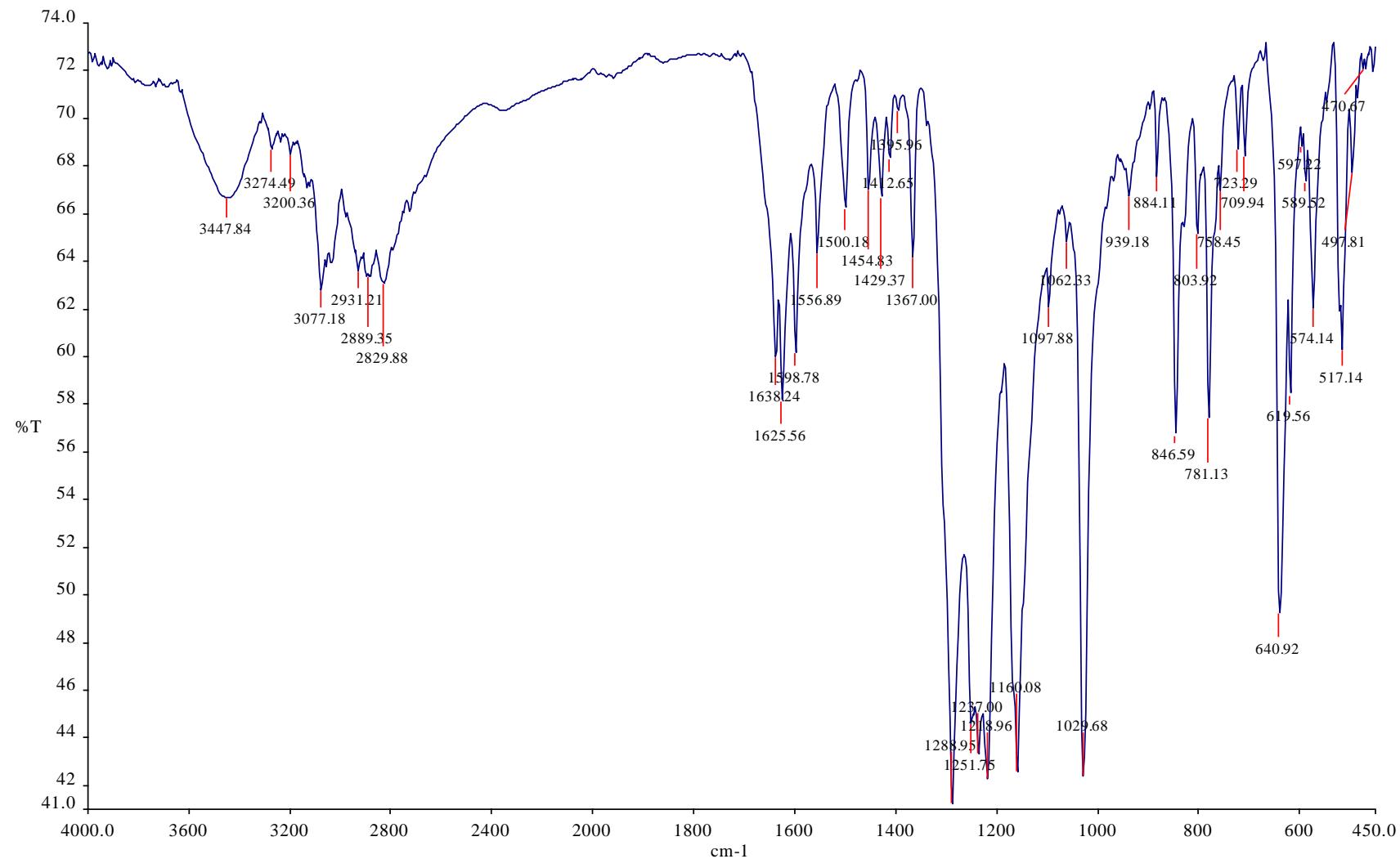
UV-Vis spectrum of **3a** (DMSO)



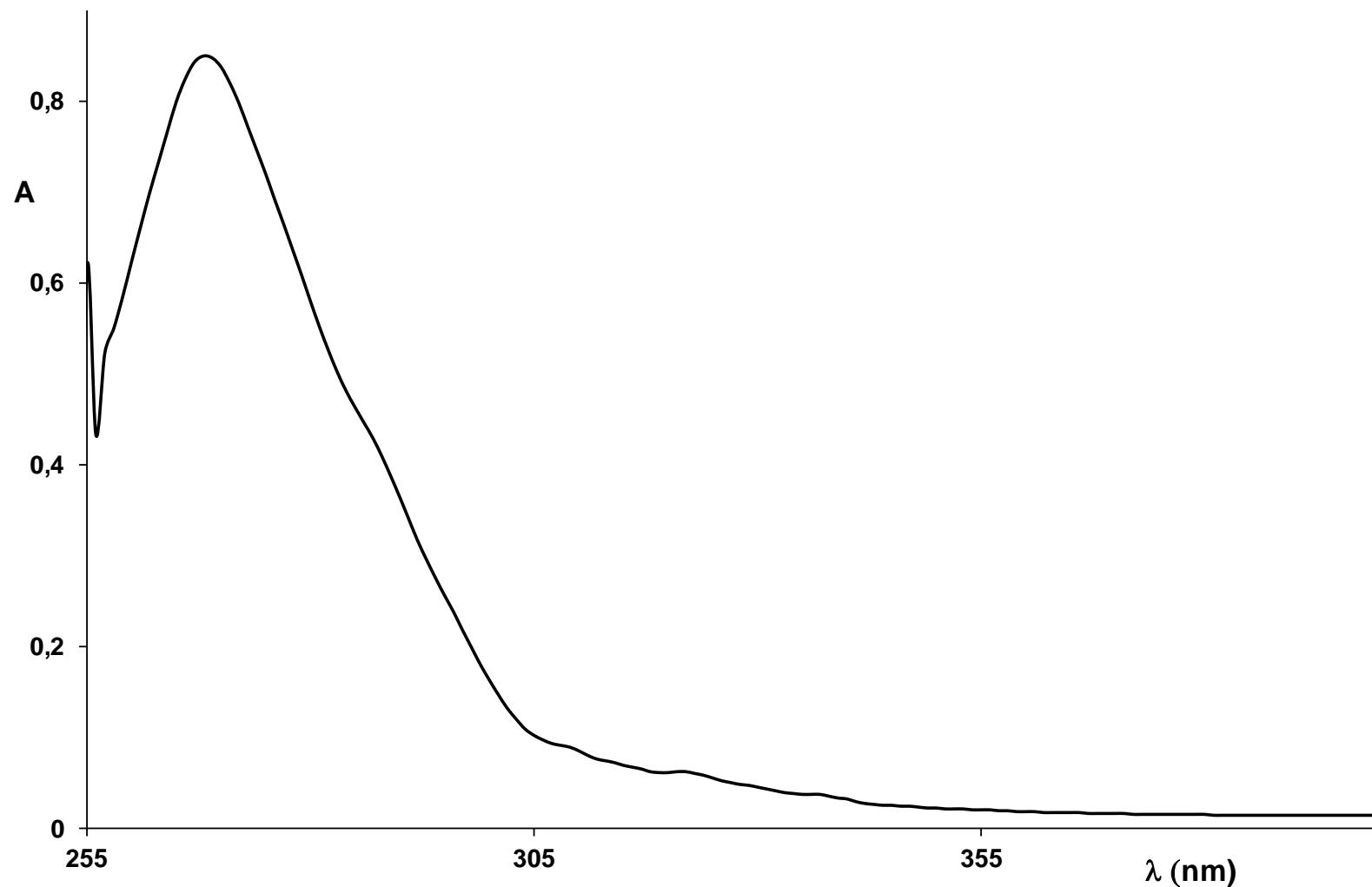
$^1\text{H}$  NMR spectrum of **3a** (200 MHz, DMSO-*d*<sub>6</sub>)



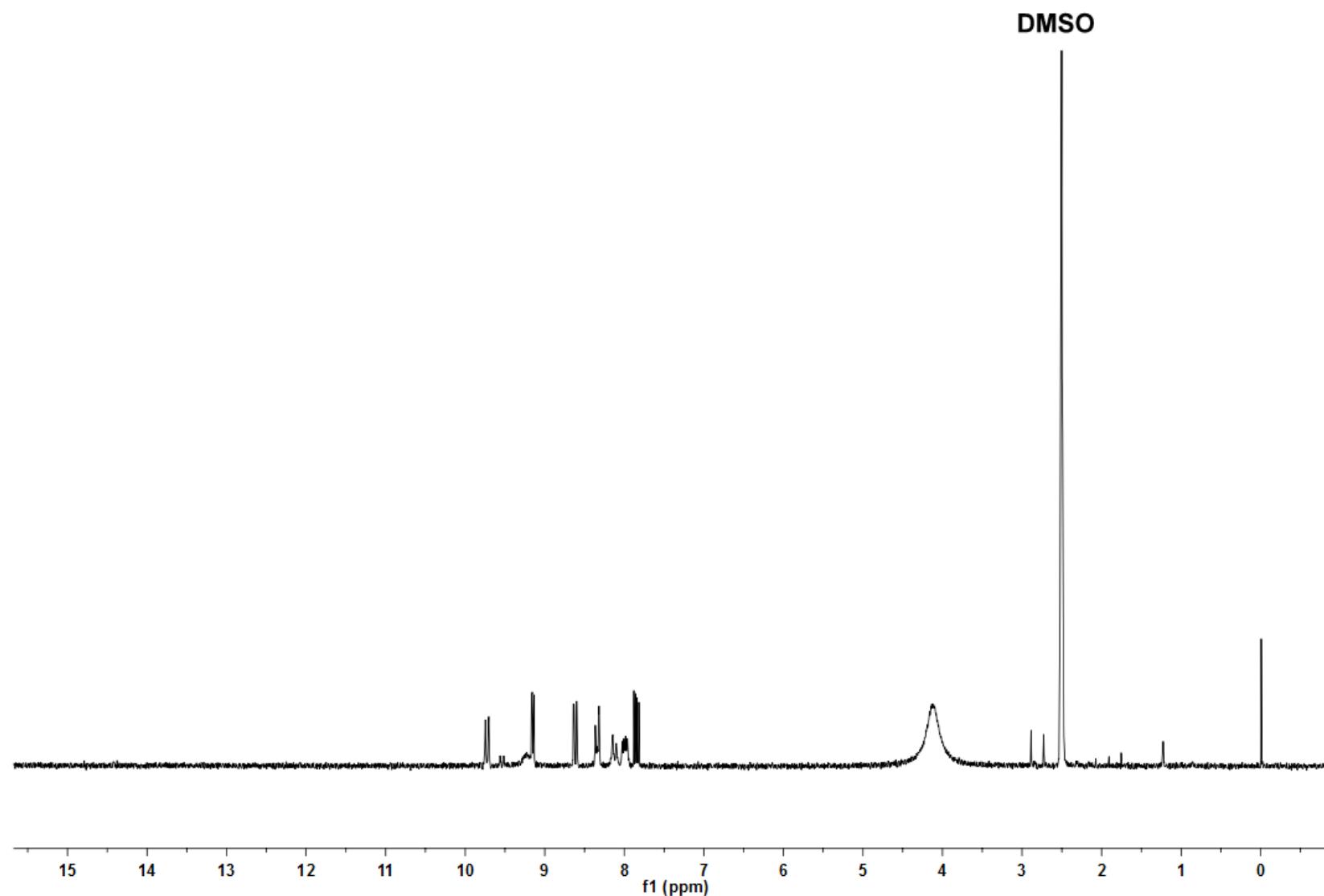
IR spectrum of 4

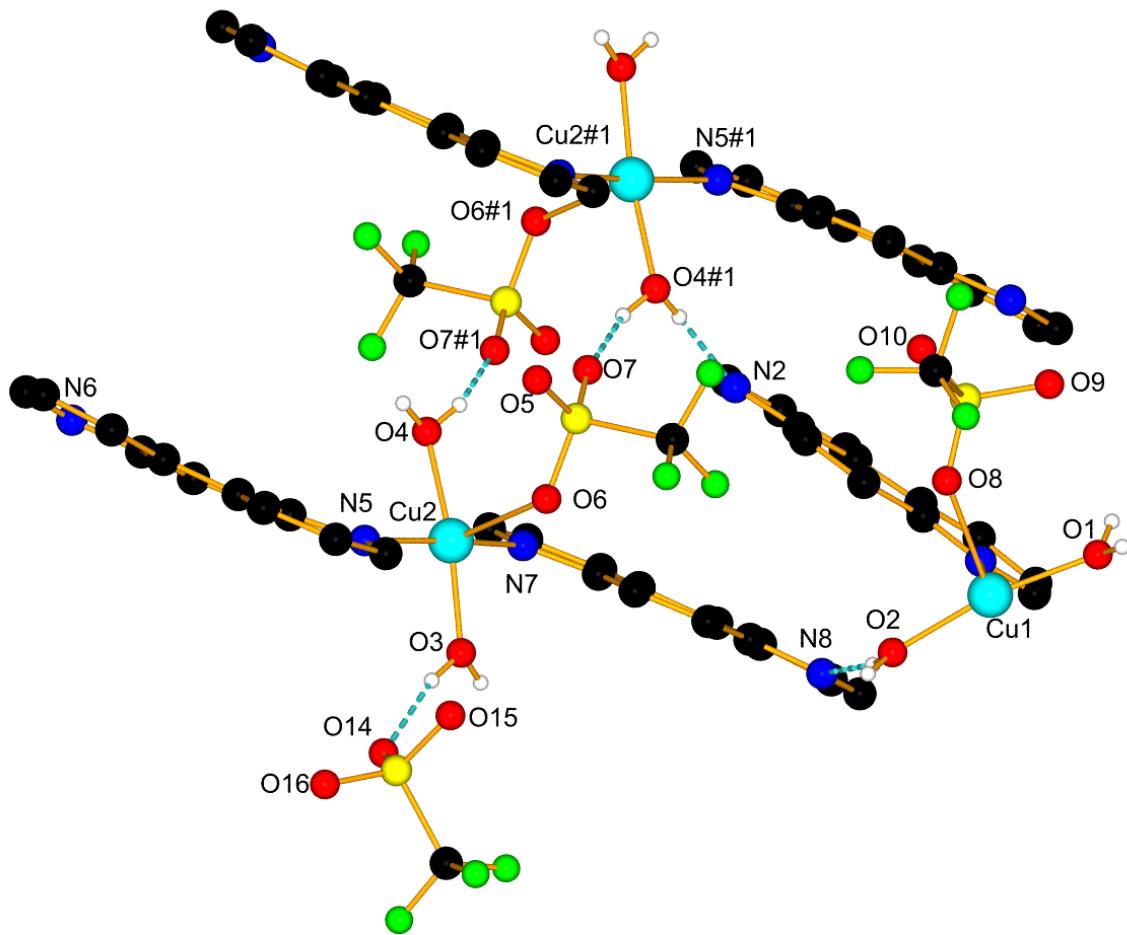


UV-Vis spectrum of **4** (DMSO)

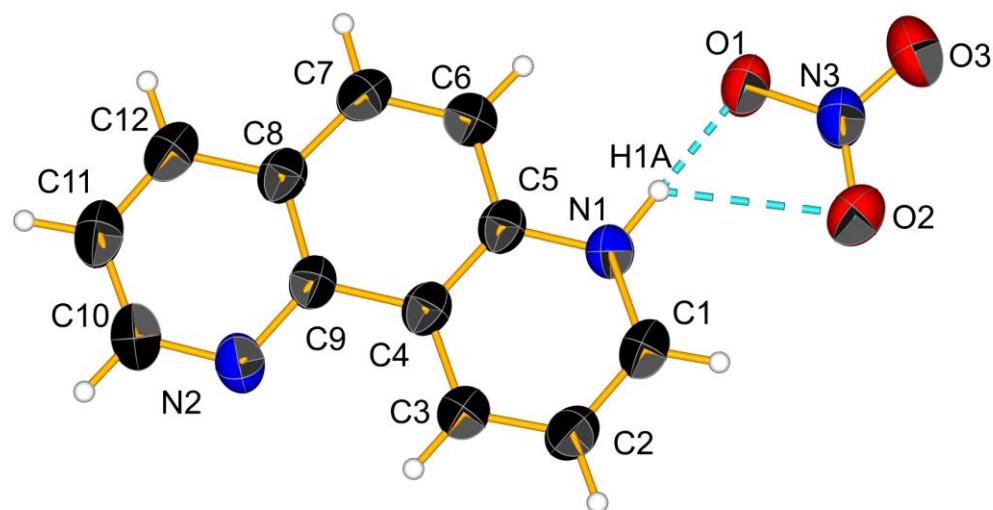


$^1\text{H}$  NMR spectrum of **4** (200 MHz, DMSO- $d_6$ )



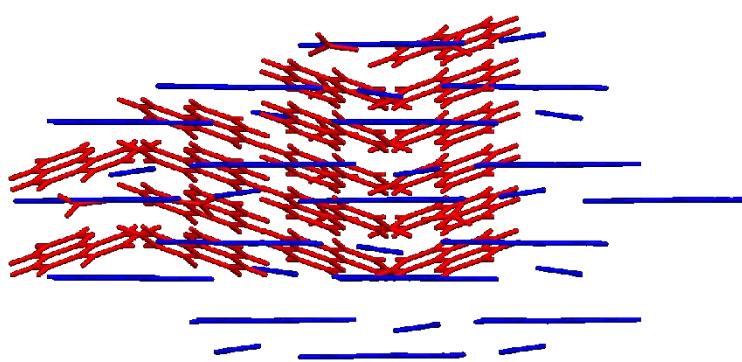


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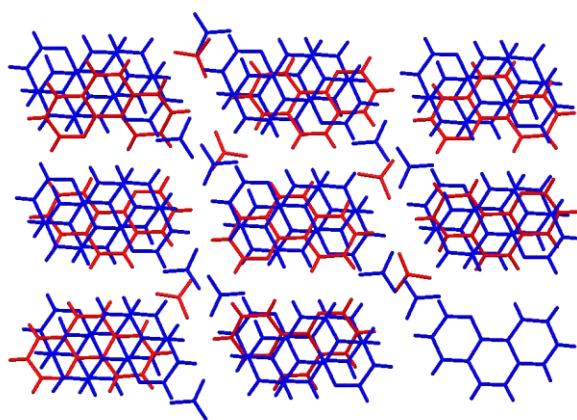


**Fig. S2.** Molecular structure of **3b**. Displacement ellipsoids are drawn at 50% probability level and H atoms are represented by spheres of arbitrary size.

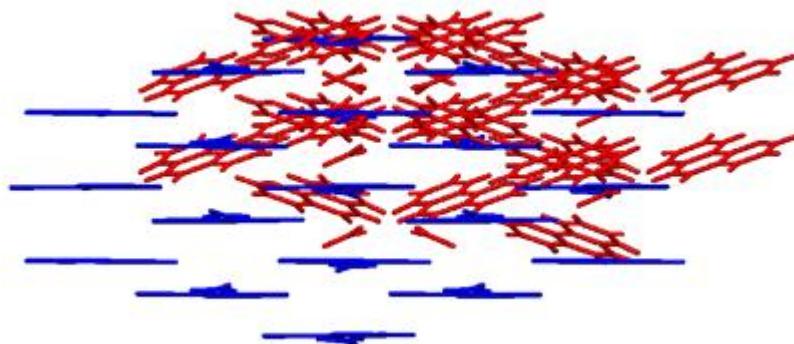
a)



b)



c)



**Fig. S3.** The overlay of the packing of the **3a** (red) and **3b** (blue) along the *a* (a), *b* (b) and *c* (c) axis.