

Supplementary Information

**3,8-connected Cd(II)-based metal-organic framework as an apt  
luminescent sensor for antibiotic sulfasalazine**

**Table of Contents**

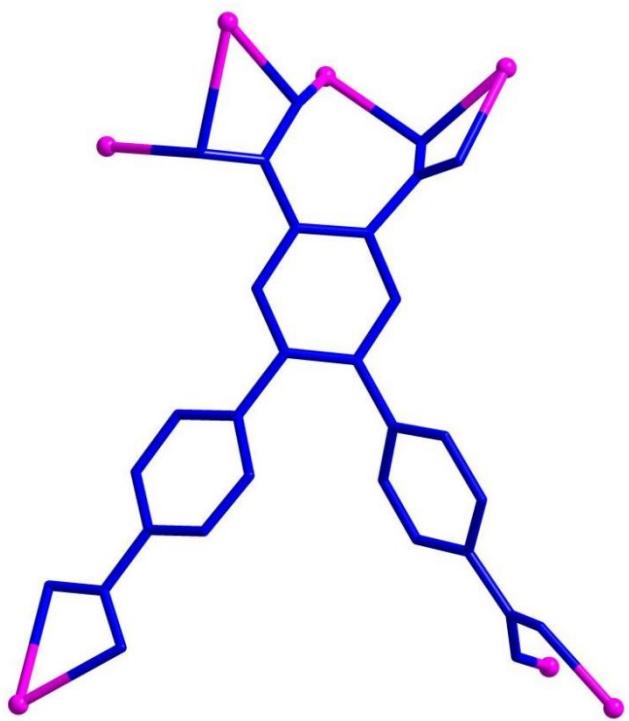
|  |   |
|--|---|
| <b>Table S1.</b> Crystal data and structure refinement parameters of <b>1</b> .....  | 2 |
| <b>Table S2.</b> Partial bond lengths (Å) and bond angles (°) of <b>1</b> .....  | 3 |
| <b>Figure S1.</b> Coordination mode of L <sup>4-</sup> ligand.....   | 3 |
| <b>Figure S2.</b> PXRD simulation pattern of <b>1</b> , PXRD pattern of <b>1</b> after sensing SLA and PXRD pattern of <b>1</b> after four sensing cycles..... | 4 |
| <b>Figure S3.</b> TGA analysis of <b>1</b> .....   | 4 |
| <b>Figure S4.</b> N <sub>2</sub> adsorption and desorption isotherms at 77 K.....  | 5 |
| <b>Figure S5.</b> view of the PL intensity of sample <b>1</b> .....  | 5 |
| <b>Figure S6.</b> The SEM pattern of morphology of <b>1</b> before and after the fluorescent probe test.....   | 6 |
| <b>Figure S7.</b> UV-spectra of antibiotic SLA and <b>1</b> .....  | 6 |
| <b>Figure S8.</b> FT-IR of ligand H <sub>4</sub> L, and <b>1</b> .....   | 7 |

**Table S1.** Crystal data and structure refinement parameters of **1**

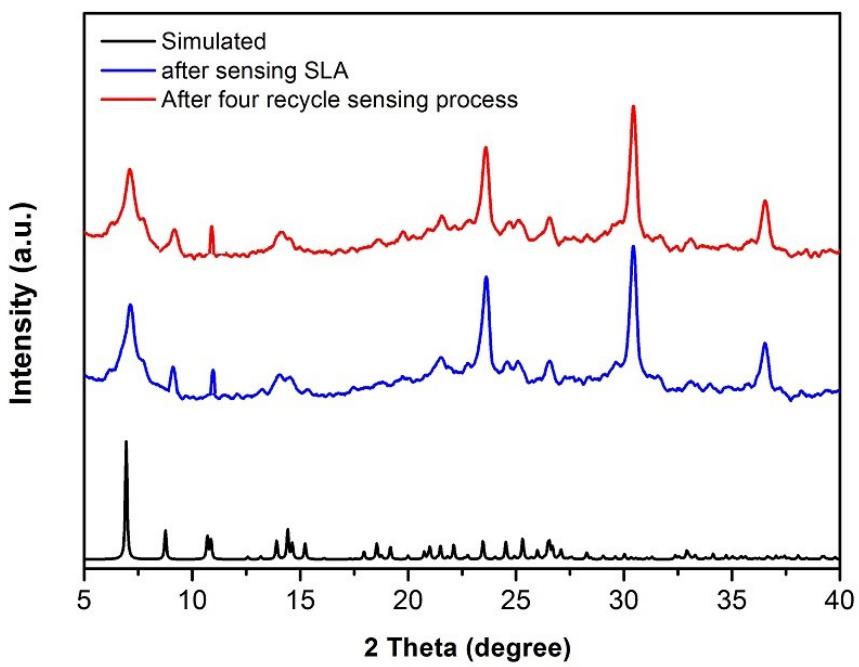
| Compound   | <b>1</b>  |
|--|---|
| Formula  | C <sub>44</sub> H <sub>26</sub> Cd <sub>4</sub> O <sub>19</sub>   |
| Formula weight   | 1308.5  |
| Temperature (K)  | 293(2)  |
| Crystal system   | Monoclinic  |
| Space group  | C2/c  |
| <i>a</i> (Å), <i>b</i> (Å), <i>c</i> (Å)   | 25.4577(11), 11.0140(7), 14.1461(12)                              |
| $\alpha$ (°), $\beta$ (°), $\gamma$ (°)  | 90, 92.376(5), 90   |
| <i>V</i> (Å <sup>3</sup> )   | 3963.0(5)   |
| <i>Z</i>   | 4   |
| <i>D</i> calc (Mg m <sup>-3</sup> )  | 2.193   |
| $\mu$ (mm <sup>-1</sup> )  | 17.739  |
| <i>F</i> (000)   | 2536.0  |
| Crystal size (mm)  | 0.05× 0.02× 0.02  |
| $\theta$ range (°)   | 3.475 to 67.06  |
| Index ranges   | -30 ≤ <i>h</i> ≤ 23<br>-11 ≤ <i>K</i> ≤ 13<br>-16 ≤ <i>L</i> ≤ 16 |
| Reflections collected  | 12248   |
| Independent reflection   | 3507 [ <i>R</i> <sub>int</sub> = 0.2003]                          |
| Data/restraints/parameters   | 3507/229/342  |
| Final <i>R</i> <sub>1</sub> , <i>wR</i> <sub>2</sub> indices [ <i>I</i> > 2σ ( <i>I</i> )] | 0.1633, 0.3734  |
| <i>R</i> <sub>1</sub> , <i>wR</i> <sub>2</sub> indices (all data)                          | 0.1784, 0.3825  |
| <i>GOF</i>   | 1.073   |
| $\Delta r_{\text{max},\text{min}}$ (e Å <sup>-3</sup> )                                    | 1.60/-2.1   |

**Table S2.** Bond length ( $\text{\AA}$ ) and bond angle ( $^\circ$ ) of **1**

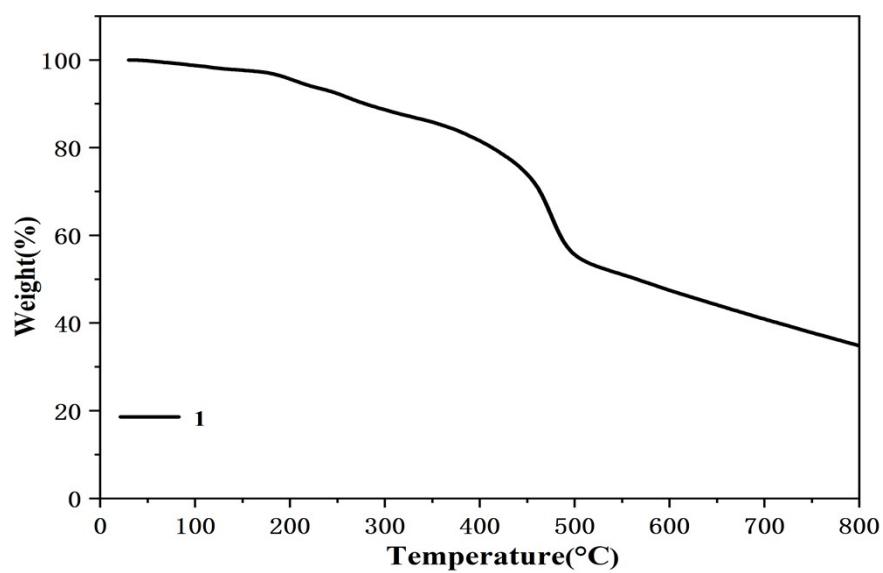
| <b>1</b>   |           |                      |           |            |           |
|--|-----------|----------------------|-----------|------------|-----------|
| Cd1-O(10)  | 2.342(17) | Cd1#2-O(2)           | 2.36(2)   | Cd1#4-O(1) | 2.08(2)   |
| Cd2-O(8)#1   | 2.360(17) | Cd2-O(3)#2           | 2.145(19) | Cd1#5-O11  | 2.425(17) |
| Cd2#6-O(3)   | 2.145(19) | Cd3-O(6)#1           | 2.333(18) | Cd3-O(5)   | 1.70(5)   |
| O(10)-Cd(1)-O(4)#9   | 88.0(6)   | O(10)-Cd(1)-O(11)#9  |           |            | 54.6(6)   |
| O(10)-Cd(1)-O(2)#7   | 100.4(7)  | O(2)#7Cd(1)-O(2)     |           |            | 78.2(6)   |
| O(8)-Cd(2)-O8)#1   | 76.5(8)   | O(10)-Cd(2)-O(8)#1   |           |            | 76.5(6)   |
| O(3)#2Cd(2)-O(8)   | 164.6(7)  | O(3)#3-Cd(2)-O(8)    |           |            | 88.1(7)   |
| O(3)#3-Cd(2)-O(10)   | 100.1(7)  | O(3)#2-Cd(2)-O(10)#1 |           |            | 95.9(7)   |
| O(6)#1-Cd(3)-O(8)#1  | 52.3(6)   | O(6)#1-Cd(3)-O(5)#1  |           |            | 93.1(10)  |
| O(7)-Cd(3)-O(6)  | 113.4(12) | O(7)-Cd(3)-O(8)      |           |            | 92.0(12)  |
| O(9) -Cd(3)-O(6)   | 102.0(18) |                      |           |            |           |
| Symmetry codes for #1 1-x,+y,-1/2-z; #2 1/2+x,1/2+y,+z; #3 1/2-x,1/2+y,-1/2-z; #4<br>-1/2+x,1/2-y,-1/2+z; #5 -1/2+x,1/2+y,+z; #6-1/2+x,-1/2+y,+z; #7 1-x,1-y,-z; |           |                      |           |            |           |



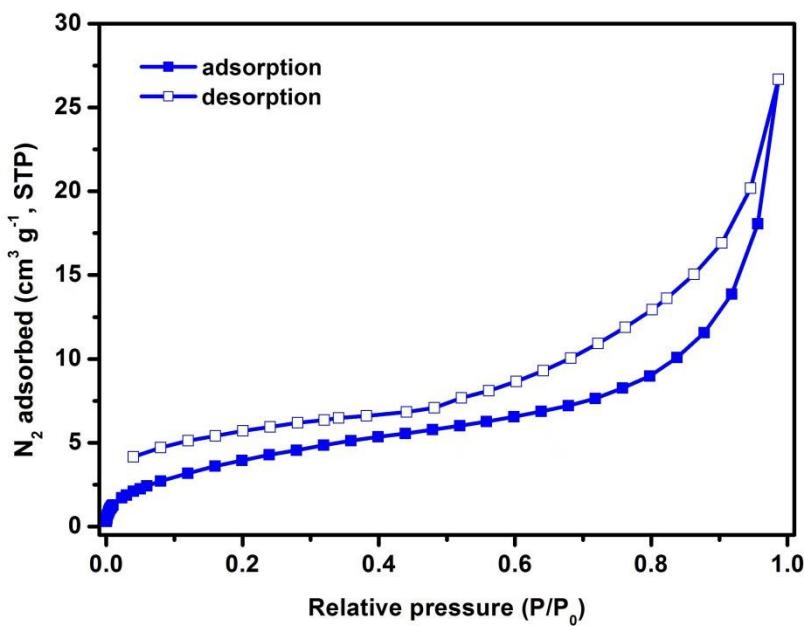
**Figure S1.** Coordination mode of  $\text{L}^4$  ligand



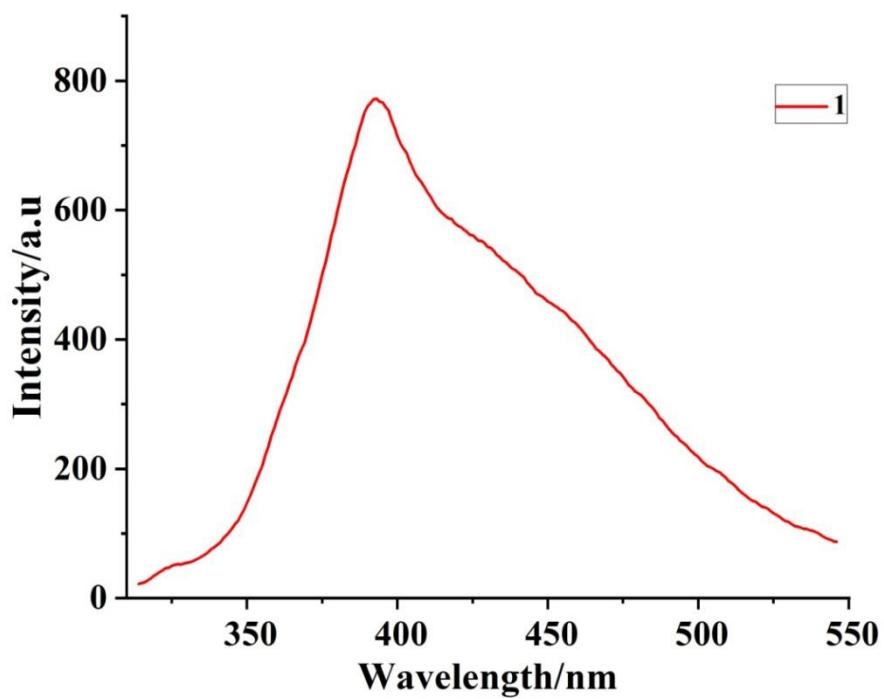
**Figure S2.** PXRD simulation pattern of **1**, PXRD pattern of **1** after sensing SLA and PXRD pattern of **1** after four sensing cycles.



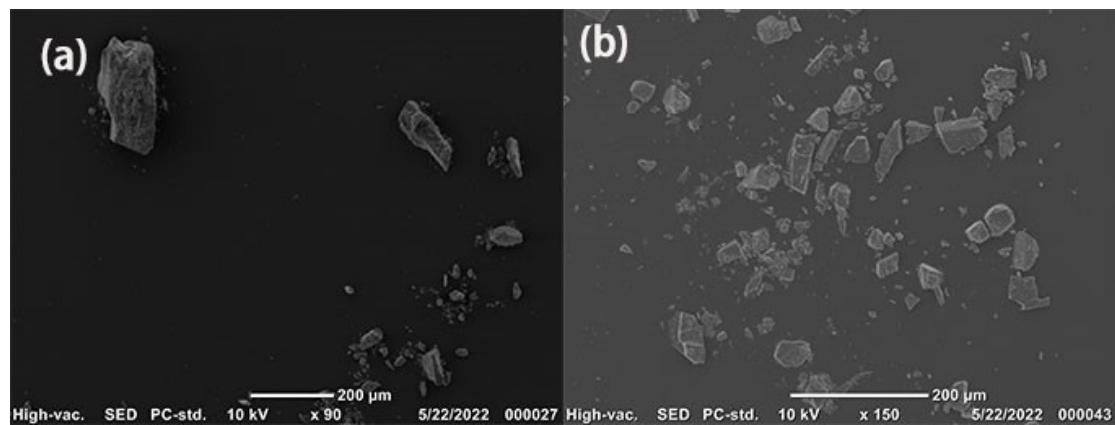
**Figure S3.** TGA analysis of **1**



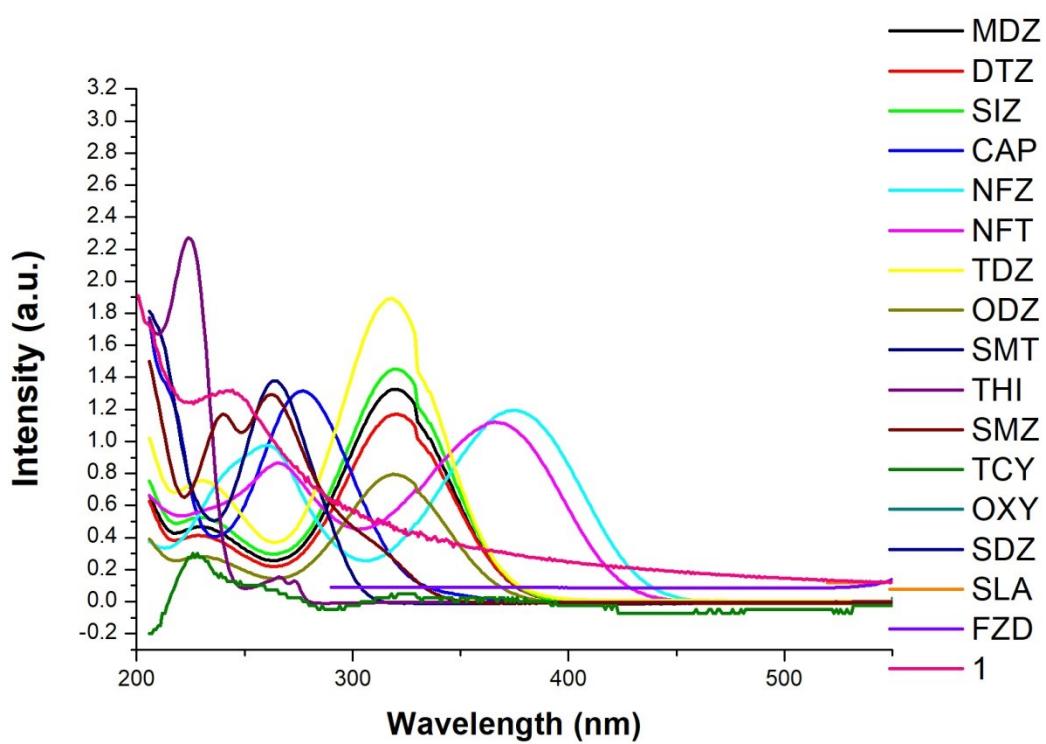
**Figure S4.**  $N_2$  adsorption and desorption isotherms at 77 K.



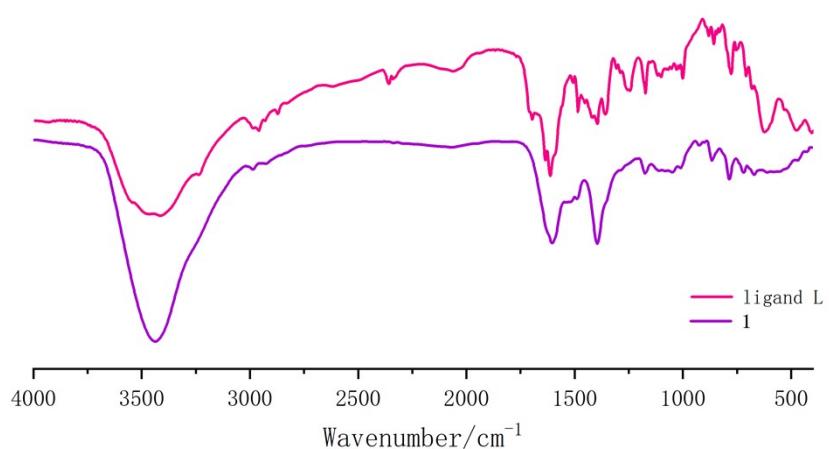
**Figure S5.** view of the PL intensity of sample **1**.



**Figure S6.** (a) The SEM pattern of morphology of **1** before the fluorescent probe test, (b) The SEM pattern of morphology of **1** after the fluorescent probe test.



**Figure S7.** UV-spectra of antibiotics and **1**.



**Figure S8.** FT-IR of ligand L, and **1**.