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**Key indicators**

Single-crystal X-ray study  
 T = 293 K  
 Mean  $\sigma(C-C)$  = 0.003 Å  
 R factor = 0.023  
 wR factor = 0.054  
 Data-to-parameter ratio = 18.1

For details of how these key indicators were  
 automatically derived from the article, see  
<http://journals.iucr.org/e>.

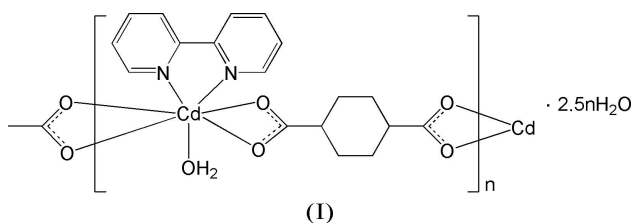
**catena-Poly[[aqua(2,2'-bipyridine-*N,N'*)cadmium(II)-  
 $\mu$ -cyclohexane-1,4-dicarboxylato- $\kappa^4 O, O': O'', O'''$ ]  
 2.5-hydrate]**

The title complex,  $\{[Cd(C_8H_{10}O_4)(C_{10}H_8N_2)(H_2O)] \cdot 2.5H_2O\}_n$ , shows a one-dimensional zigzag chain structure, in which the Cd atom is coordinated by two N atoms from a 2,2'-bipyridine ligand, a water molecule and four carboxylate O atoms from two different cyclohexane-1,4-dicarboxylate ligands.

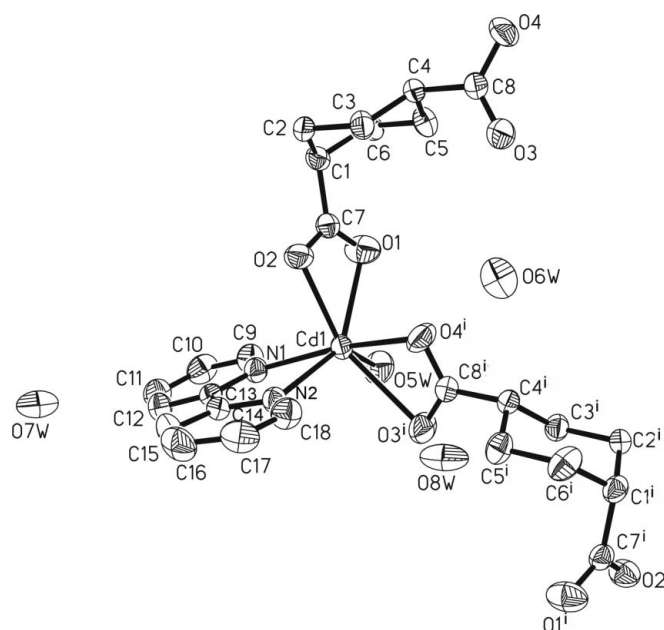
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**Comment**

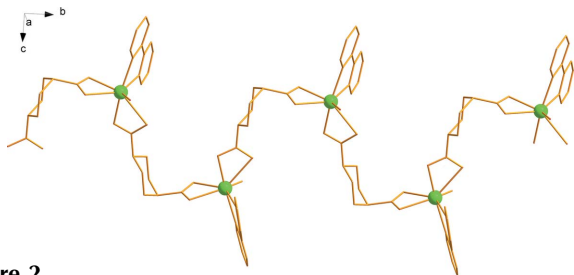
In recent years, ligands containing carboxylate groups have been known to be good building blocks for the construction of polymeric structures (Wang *et al.*, 2006). Here we report a new coordination polymer, (I), constructed from the flexible ligand cyclohexane-1,4-dicarboxylate (chdc) and a Cd<sup>II</sup> ion.



In the asymmetric unit of (I), there are one Cd<sup>II</sup> atom, one chdc ligand, one 2,2'-bipyridine (2,2'-bpy), one coordinated water molecule and two and a half uncoordinated water



**Figure 1**  
 Part of the chain structure of (I). Displacement ellipsoids are drawn at the 50% probability level. H atoms have been omitted for clarity. [Symmetry code: (i)  $\frac{1}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$ .]



**Figure 2**

The one-dimensional zigzag chain in (I). H atoms and uncoordinated water molecules have been omitted for clarity.

molecules (Fig. 1). The Cd<sup>II</sup> atom is seven-coordinated by four O atoms of two chelating carboxylate groups from two different chdc ligands, two N atoms from a 2,2'-bpy ligand and one water molecule. The chdc ligand connects the metal centers, forming an infinite zigzag chain along the *b* axis (Fig. 2). The 2,2'-bpy ligands decorate the chain on each side. The chains are linked to each other by hydrogen bonds between water molecules and carboxylate groups (Table 1). In the chdc ligand, the cyclohexane ring adopts a chair conformation. One carboxylate group lies in the *e*-bond position and the other in the *p*-bond position.

## Experimental

Cd(NO<sub>3</sub>)<sub>2</sub>·4H<sub>2</sub>O (0.062 g, 0.2 mmol), H<sub>2</sub>chdc (0.035 g, 0.2 mmol) and 2,2'-bpy (0.031 g, 0.2 mmol) were dissolved in water (20 ml). To this solution, 1 M NaOH was added dropwise to give a pH of 6.0. The final solution was sealed in a 25 ml Teflon-lined stainless steel vessel, which was heated to 453 K for 3 d and then cooled to room temperature. Colorless prismatic crystals of (I) were collected (yield 42%).

### Crystal data

|  |   |
|--|---|
| [Cd(C <sub>8</sub> H <sub>10</sub> O <sub>4</sub> )(C <sub>10</sub> H <sub>8</sub> N <sub>2</sub> )(H <sub>2</sub> O)]·2.5H <sub>2</sub> O | $\beta = 111.129 (7)^\circ$               |
| $M_r = 501.81$   | $V = 4085.2 (11) \text{ \AA}^3$           |
| Monoclinic, <i>C2/c</i>  | $Z = 8$                                   |
| $a = 15.657 (2) \text{ \AA}$   | Mo $K\alpha$ radiation                    |
| $b = 13.140 (2) \text{ \AA}$   | $\mu = 1.11 \text{ mm}^{-1}$              |
| $c = 21.287 (3) \text{ \AA}$   | $T = 293 (2) \text{ K}$                   |
|  | $0.35 \times 0.25 \times 0.15 \text{ mm}$ |

### Data collection

|  |  |
|--|--|
| Rigaku Mercury70 CCD diffractometer                                    | 16648 measured reflections             |
| Absorption correction: multi-scan (SADABS; Sheldrick, 1996)            | 4681 independent reflections           |
| $T_{\min} = 0.789$ , $T_{\max} = 0.893$ (expected range = 0.748–0.846) | 4397 reflections with $I > 2\sigma(I)$ |
|  | $R_{\text{int}} = 0.013$               |

### Refinement

|                                 |  |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.023$ | 258 parameters                                 |
| $wR(F^2) = 0.054$               | H-atom parameters constrained                  |
| $S = 1.07$                      | $\Delta\rho_{\max} = 0.60 \text{ e \AA}^{-3}$  |
| 4681 reflections                | $\Delta\rho_{\min} = -0.42 \text{ e \AA}^{-3}$ |

**Table 1**

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

| <i>D</i> —H··· <i>A</i>      | <i>D</i> —H | H··· <i>A</i> | <i>D</i> ··· <i>A</i> | <i>D</i> —H··· <i>A</i> |
|------------------------------|-------------|---------------|-----------------------|-------------------------|
| O5W—H5WA···O8W               | 0.85        | 1.93          | 2.763 (3)             | 168                     |
| O5W—H5WB···O6W <sup>†</sup>  | 0.85        | 2.02          | 2.797 (3)             | 152                     |
| O6W—H6WA···O1                | 0.85        | 2.44          | 3.251 (3)             | 159                     |
| O6W—H6WB···O1 <sup>†</sup>   | 0.85        | 2.00          | 2.770 (2)             | 150                     |
| O7W—H7WA···O4 <sup>ii</sup>  | 0.85        | 1.88          | 2.728 (2)             | 174                     |
| O7W—H7WB···O3 <sup>iii</sup> | 0.85        | 1.95          | 2.760 (2)             | 159                     |
| O8W—H8WA···O7W <sup>iv</sup> | 0.85        | 1.84          | 2.629 (2)             | 153                     |

Symmetry codes: (i)  $-x, y, -z + \frac{1}{2}$ ; (ii)  $-x, -y, -z$ ; (iii)  $x - \frac{1}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$ ; (iv)  $-x, -y + 1, -z$ .

All H atoms on C atoms were positioned geometrically and refined as riding, with C—H = 0.95 (aromatic ring), 0.99 (CH<sub>2</sub>) or 1.00  $\text{\AA}$  (CH) and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ . H atoms on water molecules were located in a difference Fourier map and refined as riding (O—H = 0.85  $\text{\AA}$ ), with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$ .

Data collection: *CrystalClear* (Rigaku, 2000); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Siemens, 1994); software used to prepare material for publication: *SHELXTL*.

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