

## A new chain-like cadmium(II) coordination polymer containing chains of water molecules

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Received 31 January 2005

Accepted 7 March 2005

Online 13 May 2005

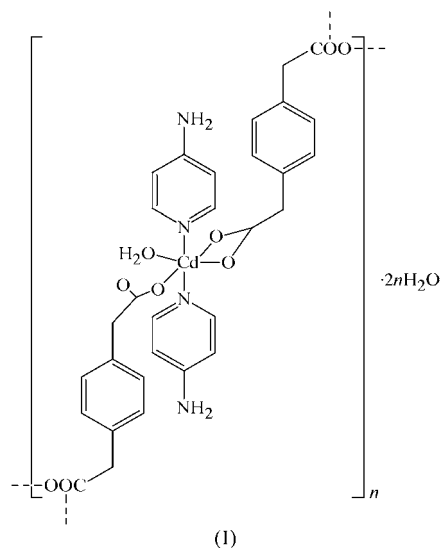
The title cadmium(II) polymer, *catena*-poly[[[bis(4-aminopyridine- $\kappa$ N)aquacadmium(II)]- $\mu$ -1,4-phenylenediacetate- $\kappa^4$ O,O':O'',O'''] dihydrate],  $[[\text{Cd}(\text{C}_{10}\text{H}_8\text{O}_4)(\text{C}_5\text{H}_6\text{N}_2)_2(\text{H}_2\text{O})] \cdot 2\text{H}_2\text{O}]_n$ , comprises one-dimensional wave-like chains, in which the Cd atom is coordinated by 1,4-phenylenediacetate and 4-aminopyridine molecules. The 1,4-phenylenediacetate ligands lie about inversion centres. Extensive hydrogen-bonding interactions between the chains lead to a three-dimensional structure. Free water molecules form chains in the structure.

### Comment

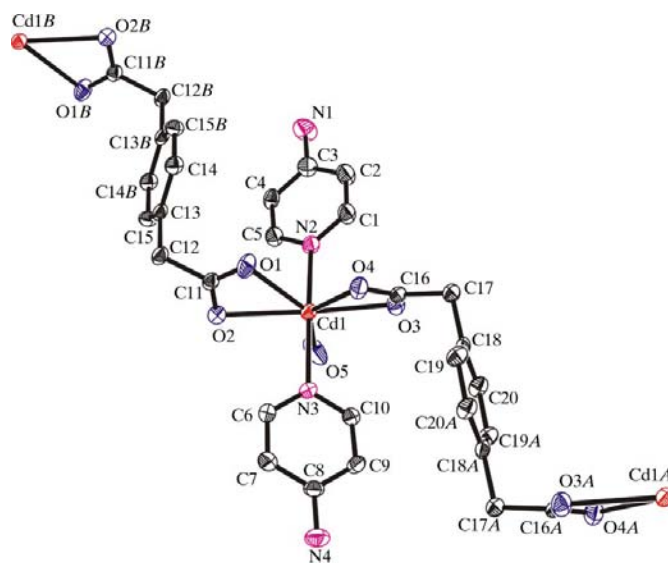
Various rigid benzenedicarboxylic acids, such as 1,2-benzenedicarboxylic acid and 1,4-benzenedicarboxylic acid, which have been demonstrated to be excellent candidates for bridging metal centers and metal clusters and which form architectures of various dimensions, have been used in numerous reactions to produce metal-organic frameworks (Li *et al.*, 2004). However, phenylenediacetic acids, for example, 1,4-phenylenediacetic acid ( $\text{H}_2\text{PDA}$ ), derived from benzenedicarboxylic acid, have more flexible carboxylic acid groups, which do not tend to take part in reactions; hence, there are only a few examples of polymeric compounds containing these acids (Pan *et al.*, 2003). In this context, we introduced a secondary building unit (4-aminopyridine) for the design of metal-organic carboxylate frameworks, to improve the activity of the carboxylate group coordination with transition metal ions. Remarkably and fortunately, a three-dimensional complex was obtained containing the two ligands. The other significant observation is that free water molecules form hydrogen-bonded chains in the structure, a phenomenon that is of current interest (Ghosh & Bharadwaj, 2004).

As shown in Fig. 1, the chain consists of  $[\text{Cd}^{\text{II}}(\text{PDA})(4\text{-aminopyridine})(\text{H}_2\text{O})]$  building blocks that are linked head-

to-tail by  $\text{PDA}^{2-}$  ligands through  $M\text{—O}$  bonds. The Cd atom has a distorted decahedral environment, with the equatorial

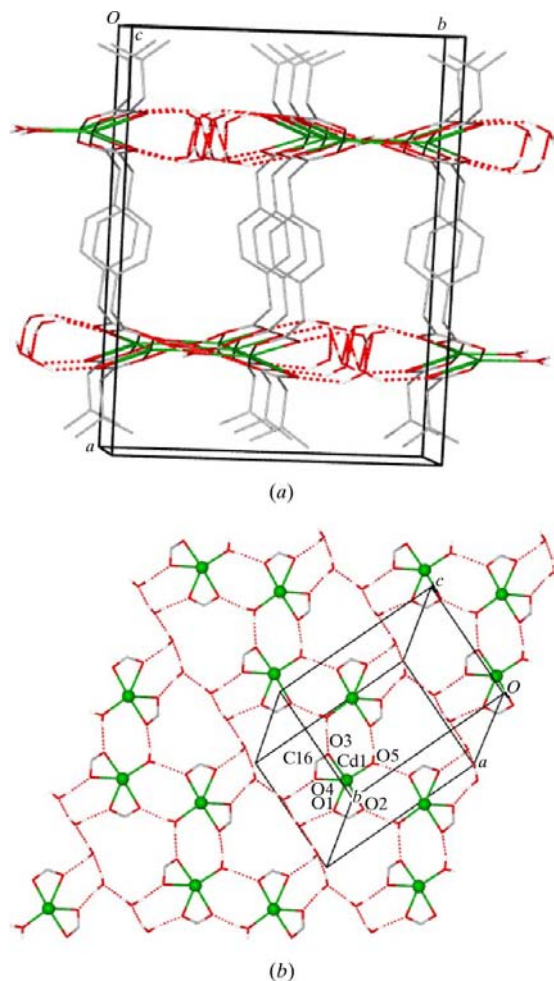


plane occupied by five O atoms, *viz.* four (O1, O2, O3 and O4) from two deprotonated  $\text{PDA}^{2-}$  molecules and one (O5) from the coordinated water molecule (Table 1). Two N atoms from two 4-aminopyridine molecules occupy the two axial positions, with bond lengths of 2.276 (3) (Cd—N3) and 2.296 (3) Å (Cd—N2). The 4-aminopyridine ligand uses its pyridine ring N atom to coordinate with the metal ion and its amine N atom is uncoordinated. The deprotonated  $\text{PDA}^{2-}$  ligands both lie about inversion centres and act as bridges, linking three  $\text{Cd}^{\text{II}}$  ions through carboxylate groups. The  $M\text{—O}_{\text{PDA}^{2-}}$  distances range from 2.378 (3) to 2.651 (3) Å, and the  $M\text{—O}_{\text{water}}$  distance is 2.334 (4) Å. Hence, each metal centre connects two  $\text{PDA}^{2-}$  ligands to form a one-dimensional chain in a step-like arrangement, with metal–metal distances of 11.644 (5) Å.

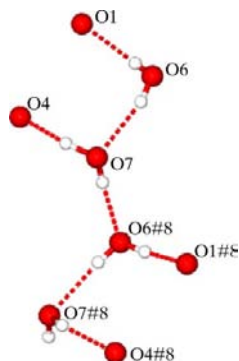


**Figure 1**  
 The local environment of the  $\text{Cd}^{\text{II}}$  ions in (I). H atoms have been omitted for clarity. Displacement ellipsoids are shown at the 50% probability level. [Symmetry codes: (A)  $-x, 2 - y, 1 - z$ ; (B)  $1 - x, 2 - y, -z$ .]

In the structure, several different kinds of hydrogen bonds (Table 2) are observed: (i) hydrogen bonds between free water molecules [ $O6 \cdots O7 = 2.825(6) \text{ \AA}$ ], (ii) hydrogen bonding of free water and carboxyl O atoms [ $O7 \cdots O4 = 2.770(5) \text{ \AA}$ ], (iii)



**Figure 2**  
(a) The two-dimensional layer in the title compound, showing hydrogen-bonding interactions along the *b* axis. H atoms, except those involved in the interactions shown, and 4-aminopyridine molecules have been omitted. (b) The two-dimensional layer of the title compound, showing hydrogen-bond interactions along the *a* axis. Benzene rings and 4-aminopyridine molecules have been omitted.



**Figure 3**  
The chain of water molecules in the title structure. [Symmetry code: (vi)  $x, -y + \frac{5}{2}, z + \frac{1}{2}$ ]

hydrogen bonding of coordinated water and carboxyl O atoms [ $O5 \cdots O2^{ii} = 2.745(5) \text{ \AA}$  and  $O5 \cdots O3^{iii} = 2.782(5) \text{ \AA}$ ; symmetry codes are as given in Table 2], (iv) hydrogen bonding between the amine N atoms in the 4-aminopyridine molecules and free water molecules [ $N4 \cdots O7^{iv} = 3.168(6) \text{ \AA}$  and  $N1 \cdots O6^v = 3.015(7) \text{ \AA}$ ], and (v) hydrogen bonding between the amine N atoms and carboxyl O atoms [ $N4 \cdots O4^i = 3.241(6) \text{ \AA}$ ]. Through these hydrogen-bonding interactions, each  $Cd^{II}$  polymer chain links to other chains to form a three-dimensional structure (Fig. 2*a*) in a layer-chain grid construction (for clarity, 4-aminopyridine molecules have been omitted from the plot). These central parts are linked by hydrogen bonds to form two-dimensional layers (Fig. 2*b*). The supramolecular layers are connected by the  $PDA^{2-}$  molecules of the step-like chains (Fig. 2*a*) to form the final three-dimensional supramolecular structure. Moreover, the uncoordinated amine N atoms of the 4-aminopyridine molecules are hydrogen bonded to O atoms of water molecules and carboxyl groups to stabilize the conformation.

Free water molecules play an interesting role in the structure, forming infinite chains by supramolecular assembly (Fig. 3). Through hydrogen bonds, these chains further organize the coordination polymer chains to be distributed alternately side by side, thus forming the two-dimensional structure (Fig. 2*b*).

## Experimental

The title compound, (I), was produced under hydrothermal conditions. Cadmium nitrate tetrahydrate (0.119 g), 1,4-phenylenediacetic acid (97%, 0.040 g), 4-aminopyridine (98%, 0.020 g) and water (15 ml) were added to a Teflon-lined steel autoclave vessel. The mixture was heated at 438 K for 3 d and then cooled slowly over a period of 2 d to room temperature. The product consisted of white crystals of the title complex in 40% yield.

### Crystal data

$[Cd(C_{10}H_8O_4)(C_5H_6N_2)_2 \cdot (H_2O)] \cdot 2H_2O$   
 $M_r = 546.85$   
 Monoclinic,  $P2_1/c$   
 $a = 18.1668(12) \text{ \AA}$   
 $b = 13.6273(10) \text{ \AA}$   
 $c = 9.2692(7) \text{ \AA}$   
 $\beta = 102.998(3)^\circ$   
 $V = 2235.9(3) \text{ \AA}^3$   
 $Z = 4$

$D_x = 1.624 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 Cell parameters from 5000 reflections  
 $\theta = 3.1\text{--}27.5^\circ$   
 $\mu = 1.02 \text{ mm}^{-1}$   
 $T = 293(2) \text{ K}$   
 Prism, white  
 $0.30 \times 0.16 \times 0.15 \text{ mm}$

### Data collection

Rigaku Mercury 70 diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan [empirical from equivalent reflections (*XEMP* in *SHELXTL*; Siemens, 1994)]  
 $T_{min} = 0.757$ ,  $T_{max} = 0.857$   
 16 963 measured reflections

5117 independent reflections  
 4579 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.030$   
 $\theta_{max} = 27.5^\circ$   
 $h = -23 \rightarrow 23$   
 $k = -8 \rightarrow 17$   
 $l = -12 \rightarrow 12$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.045$   
 $wR(F^2) = 0.148$   
 $S = 1.07$   
 5117 reflections  
 393 parameters

All H-atom parameters refined  
 $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} < 0.001$   
 $\Delta\rho_{max} = 0.60 \text{ e \AA}^{-3}$   
 $\Delta\rho_{min} = -0.51 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters (Å, °).

Cd1—O1	2.418 (3)	Cd1—N3	2.276 (3)
Cd1—O2	2.511 (3)	O1—C11	1.250 (5)
Cd1—O3	2.378 (3)	O2—C11	1.243 (5)
Cd1—O4	2.651 (3)	O3—C16	1.250 (5)
Cd1—O5	2.334 (4)	O4—C16	1.244 (5)
Cd1—N2	2.296 (3)		
N3—Cd1—O3	91.49 (12)	O5—Cd1—O2	89.13 (12)
N2—Cd1—O3	89.47 (12)	O1—Cd1—O2	52.27 (9)
O5—Cd1—O3	95.40 (12)	O3—Cd1—O4	51.23 (9)

**Table 2**

Hydrogen-bond geometry (Å, °).

<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>
N4—H9...O4 <sup>i</sup>	0.82 (6)	2.44 (6)	3.241 (6)	170 (5)
O5—H21...O2 <sup>ii</sup>	0.90 (6)	1.86 (5)	2.745 (5)	166 (5)
O5—H22...O3 <sup>iii</sup>	0.71 (5)	2.07 (5)	2.782 (5)	172 (6)
N4—H10...O7 <sup>iv</sup>	0.96 (6)	2.25 (6)	3.168 (6)	161 (5)
N1—H4...O6 <sup>v</sup>	0.78 (6)	2.23 (6)	3.015 (7)	178 (6)
O6—H26...O1	0.82 (1)	1.94 (3)	2.703 (5)	156 (8)
O6—H25...O7	0.81 (1)	2.03 (2)	2.825 (6)	166 (5)
O7—H23...O4	0.94 (6)	1.92 (6)	2.770 (5)	149 (6)
O7—H24...O6 <sup>vi</sup>	0.82 (1)	2.01 (1)	2.832 (6)	177 (5)

 Symmetry codes: (i)  $-x, -y + 2, -z$ ; (ii)  $x, -y + \frac{3}{2}, z + \frac{1}{2}$ ; (iii)  $x, -y + \frac{3}{2}, z - \frac{1}{2}$ ; (iv)  $-x, y - \frac{1}{2}, -z + \frac{1}{2}$ ; (v)  $-x + 1, y - \frac{1}{2}, -z + \frac{1}{2}$ ; (vi)  $x, -y + \frac{3}{2}, z + \frac{1}{2}$ .

H atoms were included in calculated positions and their parameters were refined isotropically [C—H = 0.87 (5)–1.07 (5) Å, O—H = 0.71 (5)–0.94 (6) Å and N—H = 0.78 (6)–0.96 (6) Å].

Data collection: *CrystalClear* (Rigaku/MSK, 2001); 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*.

This work was supported by the Natural Science Foundation of China (grant No. 29901005) and the Natural Science Foundation of Fujian Province.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: HJ1044). Services for accessing these data are described at the back of the journal.

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