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## Structure Reports

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catena-Poly[[triaquacadmium(II)]- $\mu$ -pyridine-2,3-dicarboxylato- $\kappa^3$ N,O<sup>2</sup>:O<sup>3</sup>]Hossein Aghabozorg,<sup>a\*</sup> Elham Motyeian,<sup>a</sup> Ramona Khadivi,<sup>b</sup> Mohammad Ghadermazi<sup>c</sup> and Faranak Manteghi<sup>a</sup>

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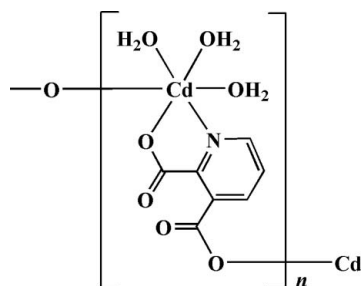
Received 17 November 2007; accepted 27 December 2007

Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(\text{C}-\text{C}) = 0.007$  Å;  $R$  factor = 0.029;  $wR$  factor = 0.057; data-to-parameter ratio = 14.9.

The title polymeric compound,  $[\text{Cd}(\text{C}_7\text{H}_3\text{NO}_4)(\text{H}_2\text{O})_3]_n$  or  $[\text{Cd}(\text{py-2,3-dc})(\text{H}_2\text{O})_3]_n$ , where py-2,3-dcH<sub>2</sub> is pyridine-2,3-dicarboxylic acid, was obtained by the reaction of cadmium(II) nitrate hexahydrate with (pipzH<sub>2</sub>)(py-2,3-dc) as a proton-transfer compound in aqueous solution (pipz is piperazine). The molecular structure shows that only the anionic fragment of the starting proton-transfer compound is present in the complex, while the (pipzH<sub>2</sub>)<sup>2+</sup> dication has been lost. Each (py-2,3-dc)<sup>2-</sup> ligand bridges two Cd<sup>II</sup> atoms in two different coordination modes, *i.e.* one end acts as a monodentate and the other end as a bidentate ligand. The three remaining coordination sites on the metal center are occupied by water molecules. The geometric arrangement of the six donor atoms around the Cd<sup>II</sup> atom is distorted octahedral. In the crystal structure, O—H...O and C—H...O hydrogen bonds play an important role in stabilizing the supramolecular structure.

## Related literature

For related ion pairs or complexes, see: Aghabozorg, Daneshvar, Motyeian *et al.* (2007); Aghabozorg *et al.* (2008); Manteghi *et al.* (2007); Aghabozorg, Sadr-khanlou *et al.* (2007); Li *et al.* (2004).



## Experimental

## Crystal data

$[\text{Cd}(\text{C}_7\text{H}_3\text{NO}_4)(\text{H}_2\text{O})_3]$   
 $M_r = 331.55$   
Orthorhombic,  $Pca2_1$   
 $a = 16.820$  (3) Å  
 $b = 6.8076$  (14) Å  
 $c = 8.6658$  (17) Å

$V = 992.3$  (3) Å<sup>3</sup>  
 $Z = 4$   
Mo  $K\alpha$  radiation  
 $\mu = 2.22$  mm<sup>-1</sup>  
 $T = 100$  (2) K  
 $0.40 \times 0.08 \times 0.05$  mm

## Data collection

Bruker APEX 1000 CCD area-detector diffractometer  
Absorption correction: multi-scan (APEX2; Bruker, 2005)  
 $T_{\min} = 0.810$ ,  $T_{\max} = 0.901$

8990 measured reflections  
2255 independent reflections  
1848 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.060$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.029$   
 $wR(F^2) = 0.056$   
 $S = 1.01$   
2255 reflections  
151 parameters  
7 restraints

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.61$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.84$  e Å<sup>-3</sup>  
Absolute structure: Flack (1983), 1042 Friedel pairs  
Flack parameter: 0.05 (4)

Table 1

Selected geometric parameters (Å, °).

Cd1—O1W	2.254 (4)	Cd1—O4	2.279 (3)
Cd1—O2 <sup>i</sup>	2.259 (3)	Cd1—N1 <sup>i</sup>	2.302 (4)
Cd1—O3W	2.274 (3)	Cd1—O2W	2.385 (3)
O3W—Cd1—O4	156.89 (12)	O2 <sup>i</sup> —Cd1—O2W	171.30 (12)
O1W—Cd1—N1 <sup>i</sup>	163.70 (14)		

Symmetry code: (i)  $-x + \frac{1}{2}, y, z - \frac{1}{2}$ .

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1W—H1...O2W <sup>ii</sup>	0.85 (2)	1.99 (2)	2.782 (5)	155.61
O1W—H2...O3 <sup>iii</sup>	0.87 (2)	1.86 (2)	2.702 (5)	163.55
O2W—H3...O1 <sup>iii</sup>	0.88 (2)	1.85 (2)	2.731 (5)	177.88
O2W—H4...O3 <sup>iv</sup>	0.87 (2)	1.84 (2)	2.687 (5)	164.45
O3W—H5...O4 <sup>iii</sup>	0.86 (2)	1.86 (2)	2.712 (4)	171.29
O3W—H6...O1 <sup>v</sup>	0.87 (2)	1.89 (2)	2.735 (8)	164.32
C3—H3A...O1W <sup>vi</sup>	0.95 (2)	2.49 (2)	3.420 (6)	165

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

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: APEX2; program(s) used to solve structure: SHELXTL (Sheldrick, 1998); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2028).

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## supporting information

*Acta Cryst.* (2008). E64, m320–m321 [doi:10.1107/S160053680706850X]

**catena-Poly[[triaquacadmium(II)]- $\mu$ -pyridine-2,3-dicarboxylato- $\kappa^3$ N,O<sup>2</sup>:O<sup>3</sup>]**

Hossein Aghabozorg, Elham Motyeian, Ramona Khadivi, Mohammad Ghadermazi and Faranak Manteghi

**S1. Comment**

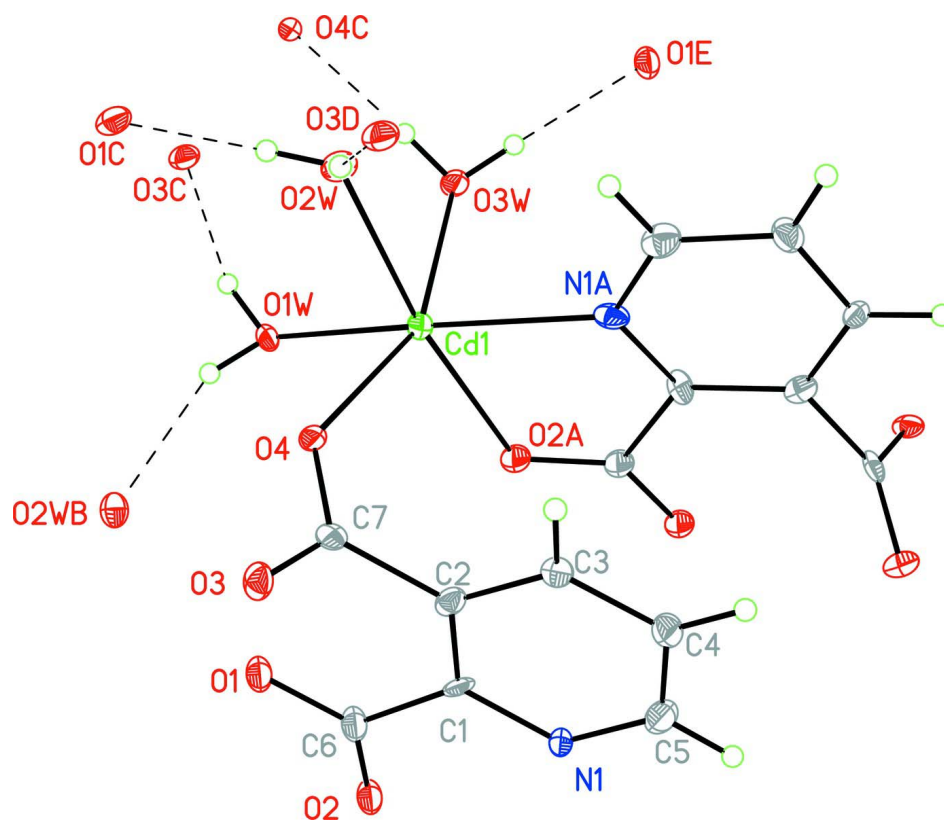
Recently, ion pairs or complexes related to the title compound have been reported (Aghabozorg, Daneshvar, Motyeian *et al.*, 2007). Proton transfer from pyridine-2,3-dicarboxylic acid (py-2,3-dcH<sub>2</sub>) to amines, such as piperazine (pipz) and propane-1,3-diamine (pn), resulted in the formation of novel systems (Aghabozorg, Manteghi *et al.*, 2007 submitted; Manteghi *et al.*, 2007). The resulting compounds, with some remaining sites as electron donors, can coordinate to metallic ions (Aghabozorg, Sadr-khanlou *et al.*, 2007). The molecular structure of the title compound shows that only the anionic fragment of starting proton transfer compound is incorporated into the complex and that the (pipzH<sub>2</sub>)<sup>2+</sup> dication has been lost. Each cadmium(II) atom is coordinated by five O-atoms and one N-atom. The asymmetric unit consists of one cadmium, one bridging (py-2,3-dc)<sup>2-</sup> and three coordinated water molecules (Fig. 1). The (py-2,3-dc)<sup>2-</sup> groups bridge two cadmium ions by adopting two different coordination modes, bidentate and monodentate. The existence of both coordination modes is seldom found in pyridine multicarboxylate coordination polymers (Li *et al.* 2004). The bond lengths and bond angles of the equatorial bonds around the metal center with atoms N1A, O2A, O1W and O2W, and the axial bonds with atoms O4 and O3W, indicate that the geometric arrangement of the six donor atoms around the cadmium(II) atom is distorted octahedral (Table 1). It is noticeable that one of the carboxylate groups is almost coplanar with the pyridine ring and the other is perpendicular to it (Fig. 1). The formation of the polymeric chains along the *c* axis is illustrated in Fig. 2. There are a number of O—H $\cdots$ O hydrogen bonds (Table 2) involving the coordinated water molecules and other O-atoms, [D $\cdots$ A distances ranging from 2.685 (5) to 2.789 (5) Å], and a C—H $\cdots$ O bond [D $\cdots$ A distance of 3.430 (6) Å], that give rise to the formation of a three-dimensional network (Fig. 3).

**S2. Experimental**

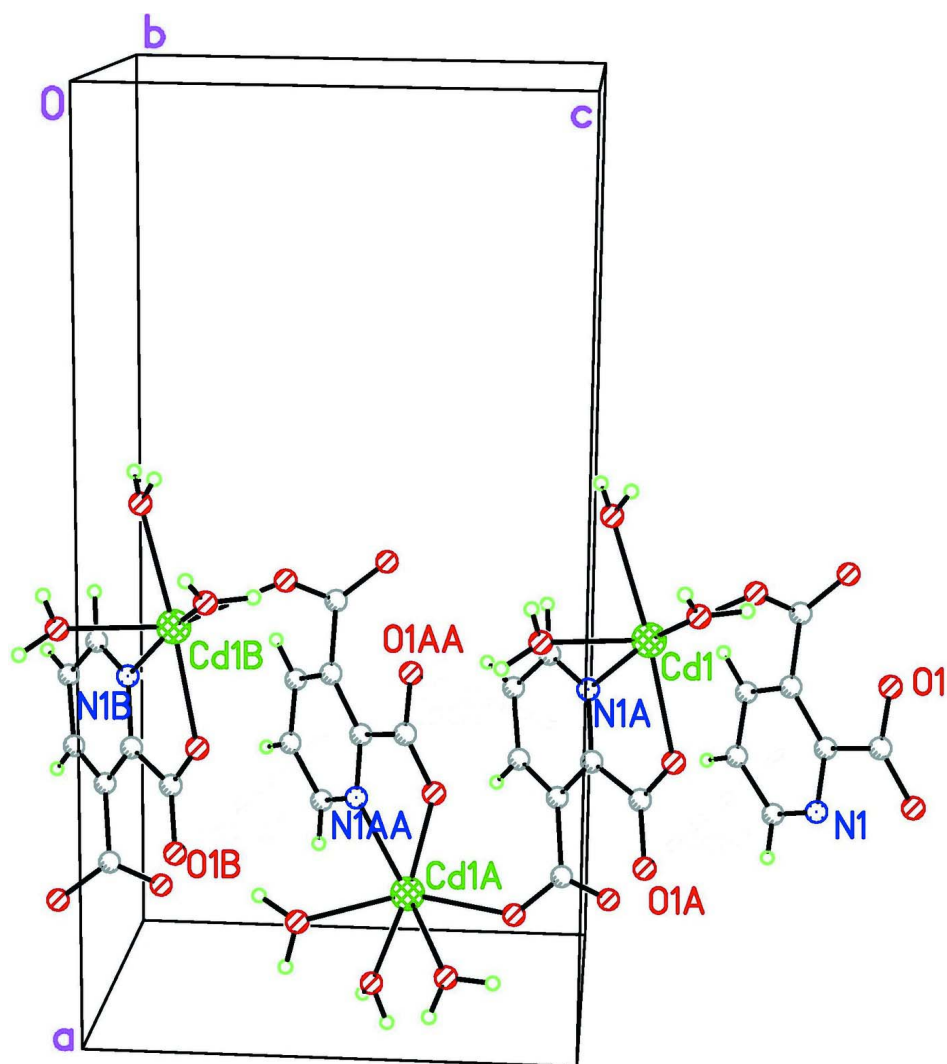
The proton transfer ion pair was prepared according to the literature (Aghabozorg *et al.*, 2008, in press). A solution of Cd(NO<sub>3</sub>)<sub>2</sub> · 6H<sub>2</sub>O (158 mg, 0.5 mmol) in water (20 ml) was added to a solution of (pipzH<sub>2</sub>)(py- 2,3-dc) (253 mg, 1.0 mmol) in water (20 ml), in a 1:2 molar ratio. Colorless crystals of the title compound suitable for X-ray characterization were obtained after a few days at room temperature.

**S3. Refinement**

All the hydrogen atoms could be located from the difference Fourier syntheses. The water H-atoms were refined isotropically with  $U_{\text{iso}}(\text{H}) = 0.022$ . The C-bond H-atoms were included in calculated positions and treated as riding atoms: C—H = 0.95 Å with  $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ .

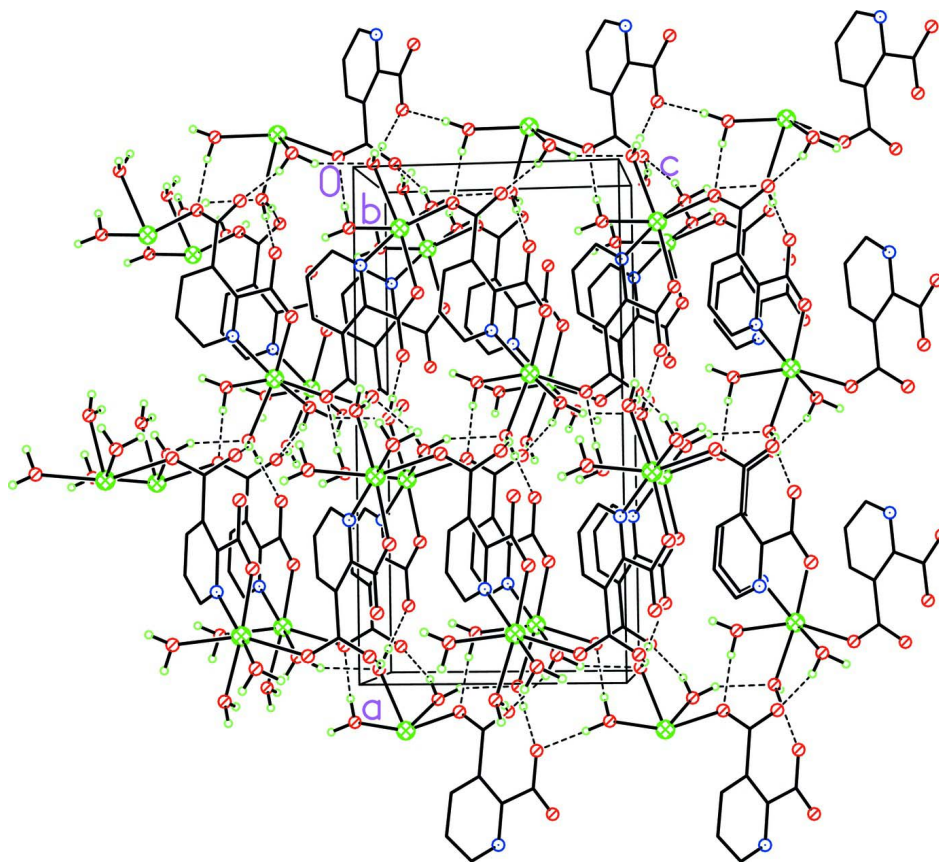
**Figure 1**

Molecular structure of the title compound, showing the atom-numbering scheme and displacement ellipsoids at the 50% probability level. Hydrogen bonds are shown as dashed lines. Atoms marked with suffixes A–E are related by the symmetry codes (A)  $-x + 3/2, y, z - 1/2$ ; (B)  $-x + 1, -y + 1, z + 1/2$ ; (C)  $-x + 1, -y + 1, z - 1/2$ ; (D)  $-x + 1, -y + 2, z - 1/2$ ; (E)  $x, y, z - 1$ ).



**Figure 2**

A view of the polymeric chain of the title compound extending along the *c* axis.



**Figure 3**

A view along the *b* axis of the crystal packing of the title compound [the hydrogen bonds are shown as dashed lines and the hydrogen atoms of the pyridine-2,3-dicarboxylates ligands are omitted for clarity].

**catena-Poly[[triaquacadmium(II)]- $\mu$ -pyridine-2,3-dicarboxylato-  $\kappa^3N,O^2:O^3$ ]**

*Crystal data*

[Cd(C<sub>7</sub>H<sub>3</sub>NO<sub>4</sub>)(H<sub>2</sub>O)<sub>3</sub>]

*M<sub>r</sub>* = 331.55

Orthorhombic, *Pca*2<sub>1</sub>

Hall symbol: P 2c -2ac

*a* = 16.820 (3) Å

*b* = 6.8076 (14) Å

*c* = 8.6658 (17) Å

*V* = 992.3 (3) Å<sup>3</sup>

*Z* = 4

*F*(000) = 648

*D<sub>x</sub>* = 2.219 Mg m<sup>-3</sup>

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 1136 reflections

θ = 3.0–24.6°

μ = 2.22 mm<sup>-1</sup>

*T* = 100 K

Needle, colourless

0.40 × 0.08 × 0.05 mm

*Data collection*

Bruker APEX 1000 CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*APEX2*; Bruker, 2005)

*T<sub>min</sub>* = 0.810, *T<sub>max</sub>* = 0.901

8990 measured reflections

2255 independent reflections

1848 reflections with *I* > 2σ(*I*)

*R<sub>int</sub>* = 0.060

θ<sub>max</sub> = 27.5°, θ<sub>min</sub> = 2.4°

*h* = -21 → 21

*k* = -8 → 8

*l* = -11 → 11

Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.029$

$wR(F^2) = 0.056$

$S = 1.01$

2255 reflections

151 parameters

7 restraints

Primary atom site location: structure-invariant  
direct methods

Secondary atom site location: difference Fourier  
map

Hydrogen site location: difference Fourier map

H atoms treated by a mixture of independent  
and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.02P)^2]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.61 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.84 \text{ e } \text{\AA}^{-3}$

Absolute structure: Flack (1983), 1042 Friedel  
pairs

Absolute structure parameter: 0.05 (4)

Special details

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
Cd1	0.602315 (16)	0.57263 (4)	1.12348 (5)	0.01148 (8)
O1	0.64375 (15)	0.5438 (4)	1.6269 (10)	0.0146 (4)
O2	0.77114 (19)	0.4893 (5)	1.6812 (4)	0.0146 (4)
O3	0.55536 (18)	0.9159 (5)	1.5507 (4)	0.0139 (7)
O4	0.56140 (18)	0.7169 (5)	1.3472 (4)	0.0131 (8)
C1	0.7384 (3)	0.7512 (7)	1.5083 (6)	0.0114 (10)
C2	0.6813 (3)	0.8656 (7)	1.4333 (6)	0.0120 (11)
C3	0.7065 (3)	1.0220 (8)	1.3431 (7)	0.0110 (11)
H3A	0.6685	1.1023	1.2920	0.013*
C4	0.7868 (3)	1.0616 (8)	1.3272 (7)	0.0149 (11)
H4A	0.8050	1.1660	1.2636	0.018*
C5	0.8403 (3)	0.9424 (8)	1.4080 (6)	0.0168 (11)
H5A	0.8956	0.9694	1.4001	0.020*
C6	0.7166 (2)	0.5813 (6)	1.6146 (9)	0.0146 (4)
C7	0.5934 (3)	0.8271 (7)	1.4475 (5)	0.0113 (10)
N1	0.8168 (2)	0.7919 (6)	1.4961 (5)	0.0125 (9)
O1W	0.5482 (2)	0.2964 (6)	1.2211 (4)	0.0145 (8)
H1	0.526 (3)	0.302 (9)	1.309 (4)	0.022*
H2	0.509 (2)	0.227 (7)	1.183 (5)	0.022*
O2W	0.47743 (19)	0.6981 (5)	1.0383 (4)	0.0134 (7)
H3	0.439 (2)	0.618 (6)	1.065 (6)	0.020*
H4	0.464 (3)	0.818 (4)	1.060 (6)	0.020*
O3W	0.59058 (19)	0.3963 (5)	0.9018 (4)	0.0147 (8)

H5	0.5445 (19)	0.350 (8)	0.879 (7)	0.022*
H6	0.609 (3)	0.420 (8)	0.811 (3)	0.022*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Cd1	0.00902 (12)	0.01505 (14)	0.01036 (13)	-0.00047 (13)	0.0004 (3)	0.0018 (3)
O1	0.0109 (8)	0.0186 (11)	0.0144 (10)	0.0005 (8)	0.0029 (11)	0.0048 (12)
O2	0.0109 (8)	0.0186 (11)	0.0144 (10)	0.0005 (8)	0.0029 (11)	0.0048 (12)
O3	0.0112 (16)	0.0158 (18)	0.0146 (16)	-0.0006 (14)	0.0041 (13)	-0.0032 (16)
O4	0.0077 (16)	0.023 (2)	0.0084 (17)	-0.0019 (14)	0.0004 (14)	-0.0031 (16)
C1	0.011 (2)	0.012 (3)	0.011 (2)	-0.0042 (19)	-0.0063 (19)	-0.001 (2)
C2	0.010 (2)	0.014 (3)	0.012 (3)	-0.0018 (19)	0.000 (2)	-0.005 (2)
C3	0.009 (3)	0.013 (3)	0.011 (3)	0.004 (2)	0.000 (2)	-0.002 (2)
C4	0.014 (3)	0.016 (3)	0.014 (3)	-0.002 (2)	-0.001 (2)	0.005 (3)
C5	0.012 (3)	0.020 (3)	0.017 (3)	-0.004 (2)	0.000 (2)	-0.004 (3)
C6	0.0109 (8)	0.0186 (11)	0.0144 (10)	0.0005 (8)	0.0029 (11)	0.0048 (12)
C7	0.013 (2)	0.011 (3)	0.010 (2)	-0.001 (2)	-0.003 (2)	0.0057 (19)
N1	0.011 (2)	0.016 (2)	0.010 (2)	-0.0057 (17)	0.0018 (16)	-0.0007 (18)
O1W	0.0142 (18)	0.022 (2)	0.0070 (17)	-0.0081 (15)	0.0020 (14)	-0.0012 (16)
O2W	0.0127 (17)	0.0119 (19)	0.0156 (19)	0.0014 (14)	-0.0030 (14)	-0.0008 (17)
O3W	0.0113 (18)	0.023 (2)	0.0093 (16)	-0.0033 (15)	0.0003 (14)	-0.0040 (16)

*Geometric parameters (Å, °)*

Cd1—O1W	2.254 (4)	C2—C7	1.506 (6)
Cd1—O2 <sup>i</sup>	2.259 (3)	C3—C4	1.385 (6)
Cd1—O3W	2.274 (3)	C3—H3A	0.9500
Cd1—O4	2.279 (3)	C4—C5	1.400 (7)
Cd1—N1 <sup>i</sup>	2.302 (4)	C4—H4A	0.9500
Cd1—O2W	2.385 (3)	C5—N1	1.338 (7)
O1—C6	1.256 (4)	C5—H5A	0.9500
O2—C6	1.252 (6)	N1—Cd1 <sup>ii</sup>	2.302 (4)
O2—Cd1 <sup>ii</sup>	2.259 (3)	O1W—H1	0.850 (19)
O3—C7	1.255 (6)	O1W—H2	0.869 (19)
O4—C7	1.268 (6)	O2W—H3	0.88 (2)
C1—N1	1.351 (6)	O2W—H4	0.866 (19)
C1—C2	1.397 (7)	O3W—H5	0.858 (19)
C1—C6	1.523 (7)	O3W—H6	0.87 (2)
C2—C3	1.387 (7)		
O1W—Cd1—O2 <sup>i</sup>	95.06 (13)	C2—C3—H3A	119.9
O1W—Cd1—O3W	80.89 (13)	C3—C4—C5	117.7 (5)
O2 <sup>i</sup> —Cd1—O3W	97.84 (12)	C3—C4—H4A	121.1
O1W—Cd1—O4	85.31 (13)	C5—C4—H4A	121.1
O2 <sup>i</sup> —Cd1—O4	101.79 (12)	N1—C5—C4	122.6 (5)
O3W—Cd1—O4	156.89 (12)	N1—C5—H5A	118.7
O1W—Cd1—N1 <sup>i</sup>	163.70 (14)	C4—C5—H5A	118.7



O2 <sup>i</sup> —Cd1—N1 <sup>i</sup>	73.25 (13)	O2—C6—O1	125.0 (6)
O3W—Cd1—N1 <sup>i</sup>	89.34 (14)	O2—C6—C1	118.8 (4)
O4—Cd1—N1 <sup>i</sup>	107.87 (14)	O1—C6—C1	116.2 (5)
O1W—Cd1—O2W	93.38 (13)	O3—C7—O4	123.9 (4)
O2 <sup>i</sup> —Cd1—O2W	171.30 (12)	O3—C7—C2	118.3 (4)
O3W—Cd1—O2W	81.44 (12)	O4—C7—C2	117.6 (4)
O4—Cd1—O2W	80.96 (12)	C5—N1—C1	119.4 (4)
N1 <sup>i</sup> —Cd1—O2W	98.06 (13)	C5—N1—Cd1 <sup>ii</sup>	126.5 (3)
C6—O2—Cd1 <sup>ii</sup>	117.6 (3)	C1—N1—Cd1 <sup>ii</sup>	114.0 (3)
C7—O4—Cd1	135.2 (3)	Cd1—O1W—H1	118 (4)
N1—C1—C2	121.3 (5)	Cd1—O1W—H2	128 (3)
N1—C1—C6	116.0 (4)	H1—O1W—H2	92 (5)
C2—C1—C6	122.6 (4)	Cd1—O2W—H3	110 (3)
C3—C2—C1	118.7 (4)	Cd1—O2W—H4	120 (4)
C3—C2—C7	118.6 (4)	H3—O2W—H4	110 (5)
C1—C2—C7	122.7 (4)	Cd1—O3W—H5	118 (4)
C4—C3—C2	120.2 (5)	Cd1—O3W—H6	130 (4)
C4—C3—H3A	119.9	H5—O3W—H6	101 (5)
O1W—Cd1—O4—C7	131.6 (5)	C2—C1—C6—O2	-177.9 (5)
O2 <sup>i</sup> —Cd1—O4—C7	37.4 (5)	N1—C1—C6—O1	179.6 (6)
O3W—Cd1—O4—C7	-175.1 (4)	C2—C1—C6—O1	2.7 (9)
N1 <sup>i</sup> —Cd1—O4—C7	-38.6 (5)	Cd1—O4—C7—O3	168.0 (3)
O2W—Cd1—O4—C7	-134.3 (5)	Cd1—O4—C7—C2	-7.0 (7)
N1—C1—C2—C3	0.9 (7)	C3—C2—C7—O3	-87.1 (6)
C6—C1—C2—C3	177.6 (5)	C1—C2—C7—O3	92.4 (6)
N1—C1—C2—C7	-178.5 (4)	C3—C2—C7—O4	88.2 (6)
C6—C1—C2—C7	-1.8 (8)	C1—C2—C7—O4	-92.3 (6)
C1—C2—C3—C4	0.6 (9)	C4—C5—N1—C1	0.2 (8)
C7—C2—C3—C4	-179.9 (5)	C4—C5—N1—Cd1 <sup>ii</sup>	176.0 (4)
C2—C3—C4—C5	-1.7 (9)	C2—C1—N1—C5	-1.3 (7)
C3—C4—C5—N1	1.3 (9)	C6—C1—N1—C5	-178.3 (5)
Cd1 <sup>ii</sup> —O2—C6—O1	175.2 (6)	C2—C1—N1—Cd1 <sup>ii</sup>	-177.7 (4)
Cd1 <sup>ii</sup> —O2—C6—C1	-4.1 (7)	C6—C1—N1—Cd1 <sup>ii</sup>	5.4 (6)
N1—C1—C6—O2	-1.0 (8)		

Symmetry codes: (i)  $-x+3/2, y, z-1/2$ ; (ii)  $-x+3/2, y, z+1/2$ .

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1W—H1 $\cdots$ O2W <sup>iii</sup>	0.85 (2)	1.99 (2)	2.782 (5)	156
O1W—H2 $\cdots$ O3 <sup>iv</sup>	0.87 (2)	1.86 (2)	2.702 (5)	164
O2W—H3 $\cdots$ O1 <sup>iv</sup>	0.88 (2)	1.85 (2)	2.731 (5)	178
O2W—H4 $\cdots$ O3 <sup>v</sup>	0.87 (2)	1.84 (2)	2.687 (5)	165
O3W—H5 $\cdots$ O4 <sup>iv</sup>	0.86 (2)	1.86 (2)	2.712 (4)	171

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O3W—H6···O1 <sup>vi</sup>	0.87 (2)	1.89 (2)	2.735 (8)	164
C3—H3A···O1W <sup>vii</sup>	0.95 (2)	2.49 (2)	3.420 (6)	165

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Symmetry codes: (iii)  $-x+1, -y+1, z+1/2$ ; (iv)  $-x+1, -y+1, z-1/2$ ; (v)  $-x+1, -y+2, z-1/2$ ; (vi)  $x, y, z-1$ ; (vii)  $x, y+1, z$ .