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Key indicators

Single-crystal X-ray study
 $T = 150$ K
 Mean $\sigma(\text{C}-\text{C}) = 0.008$ Å
 R factor = 0.050
 wR factor = 0.131
 Data-to-parameter ratio = 15.5

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

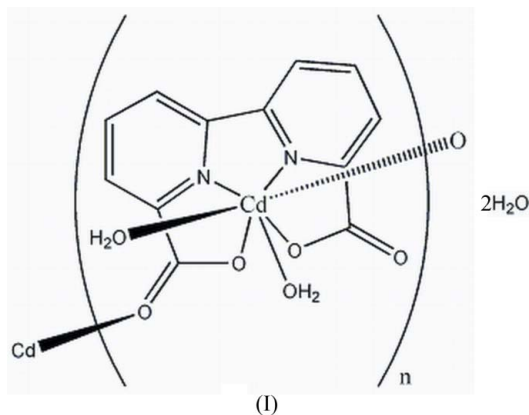
catena-Poly[[[diaquacadmium(II)]- μ -2,2'-bipyridine-6,6'-dicarboxylato] dihydrate]

The title hydrated 2,2'-bipyridine-6,6'-dicarboxylate-cadmium(II) complex, $\{[\text{Cd}(\text{C}_{12}\text{H}_{12}\text{N}_2\text{O}_4)(\text{H}_2\text{O})_2] \cdot 2\text{H}_2\text{O}\}_n$, crystallizes with eight oxygen-bridged monomer units in the unit cell. The divalent cadmium ion is seven-coordinate, and the coordination polyhedron can best be described as slightly distorted pentagonal-bipyramidal.

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Comment

The title compound, (I), has been synthesized as part of a study investigating complex stability. Specifically, the research is part of initial investigations towards generating Gd^{III} -based contrast agents for applications in magnetic resonance imaging (MRI), where overall complex stability is crucial in preventing dissociation *in vivo*, due to the well documented interference of Gd^{3+} in biological processes (Cacheris, 1990).



The research aims to elucidate structural features that contribute significantly to overall stability. A comprehensive series of transition metals has been considered, providing an insight into competition reactions which may compromise the effectiveness of the ability of the ligands to coordinate to the Gd^{3+} ion and neutralize its potential toxicity (Caravan, 1999).

In this complex, the Cd atom is located at the centre of a distorted pentagonal bipyramid of seven coordinating atoms (five O atoms and two N atoms; Fig. 1). One of these donor atoms ($\text{O}4^i$) originates from another symmetry-related complex ($x - \frac{1}{2}, \frac{1}{2} - y, z$). The bond lengths are comparable to those of similar polymeric cadmium-based complexes (Deloume & Loiseleur, 1974). The bridging behaviour results in the formation of a polymer, which extends in a zigzag fashion (see Fig. 2).

The donor atoms $\text{O}4^i$ and $\text{O}5$ are located in the axial positions, with $\text{N}1$, $\text{N}2$, $\text{O}1$, $\text{O}3$ and $\text{O}6$ in the equatorial plane. The deviation of the equatorial angles $\text{N}1-\text{Cd}1-\text{N}2$, $\text{N}2-$

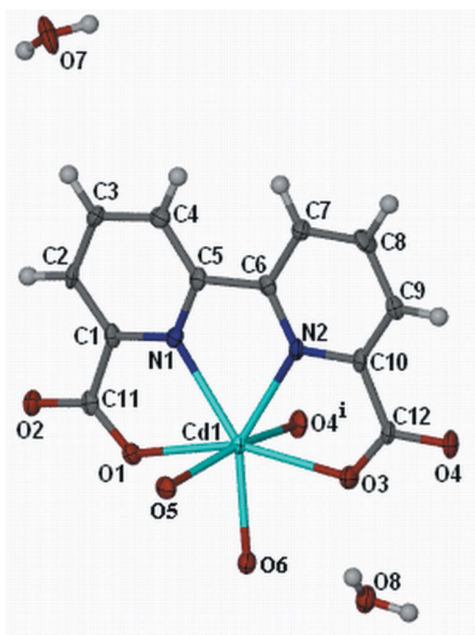


Figure 1

Perspective view of the asymmetric unit, expanded to complete the Cd coordination, showing the atom numbering. Displacement ellipsoids are shown at the 50% probability level. H atoms are represented by circles of arbitrary size. [Symmetry code: (i) $x - \frac{1}{2}, \frac{1}{2} - y, z$.]

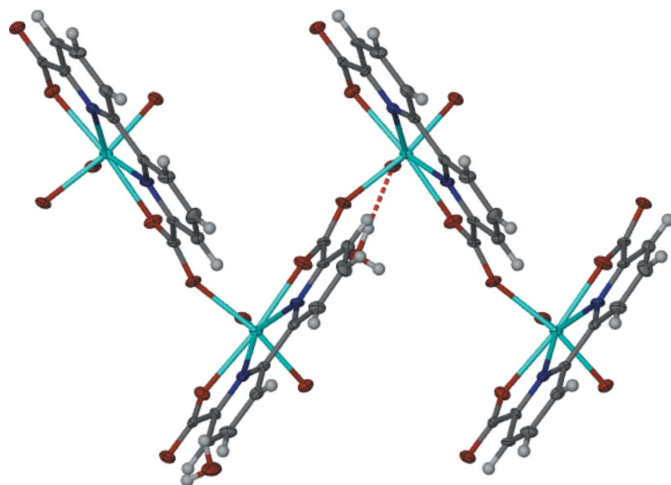


Figure 2

Fragment of the title structure, showing an oxygen-bridged chain extended in a zigzag fashion along the c axis. The dashed line indicates a hydrogen bond.

$\text{Cd1}-\text{O3}$, $\text{O3}-\text{Cd1}-\text{O6}$, $\text{O6}-\text{Cd1}-\text{O1}$, $\text{O1}-\text{Cd1}-\text{N1}$ (Table 1) from the theoretical average angle of 72° can be explained by the narrow bite angle resulting from the coordination of the rigid tetradentate ligand. While the bridging carboxylate group lies almost perpendicular to the tetradentate ligand at $88.75(13)^\circ$, the axial water molecule appears to lean away from the more sterically demanding ligand, tending towards the equatorial water donor at a more acute angle of $82.08(12)^\circ$.

The oxygen-bridged polymer is formed *via* coordination of one of the carbonyl O atoms (O4^i) to the cadmium in an axial orientation at an average angle of 90.10° to the pentagonal plane. The molecular packing is layered, with intermolecular

π -stacking between bipyridine rings ($\text{N1}/\text{C1}-\text{C5}$ and $\text{N2}/\text{C6}-\text{C10}$) at a centroid-centroid distance of 3.267 \AA . The equatorially coordinating water molecule (O6) is hydrogen bonded to a solvent water molecule (O8), which in turn is bonded to a second solvent water molecule (O7) which is hydrogen bonded to the axially coordinating water (O5) (see Fig. 3). All significant hydrogen bonds are listed in Table 2.

Experimental

To an aqueous solution (5 ml water) of 2,2-bipyridine-6,6-dicarboxylate disodium salt (0.05 mol) was added an aqueous solution (5 ml water) of cadmium(II) perchlorate (0.05 mol) and the solution was stirred continuously at room temperature for 5 h. A white precipitate was collected *via* filtration and redissolved in the minimum amount of hot deionized water. Slow evaporation of the solution yielded well defined crystals suitable for X-ray diffraction. ^1H NMR (D_2O): 8.48 (*d*, 2H, $J = 9.63 \text{ Hz}$, py-H1), 8.16 (*t*, 2H, $J = 9.75 \text{ Hz}$, py-H2), 8.08 (*d*, 2H, $J = 7.60 \text{ Hz}$, py-H3).

Crystal data

$[\text{Cd}(\text{C}_{12}\text{H}_8\text{N}_2\text{O}_4)(\text{H}_2\text{O})_2] \cdot 2\text{H}_2\text{O}$

$M_r = 422.63$

Orthorhombic, $Pcab$

$a = 8.665(5) \text{ \AA}$

$b = 17.392(5) \text{ \AA}$

$c = 18.812(5) \text{ \AA}$

$V = 2835(2) \text{ \AA}^3$

$Z = 8$

$D_x = 1.999 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

$\mu = 1.59 \text{ mm}^{-1}$

$T = 150(2) \text{ K}$

Block, colourless

$0.52 \times 0.4 \times 0.22 \text{ mm}$

Data collection

Bruker-Nonius KappaCCD diffractometer

φ and ω scans

Absorption correction: multi-scan

(*SORTAV*; Blessing, 1995)

$T_{\min} = 0.492$, $T_{\max} = 0.722$

13837 measured reflections

3234 independent reflections

2349 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.092$

$\theta_{\max} = 27.5^\circ$

Refinement

Refinement on F^2

$R[F^2 > 2\sigma(F^2)] = 0.050$

$wR(F^2) = 0.131$

$S = 1.04$

3234 reflections

208 parameters

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0704P)^2 + 1.3489P]$

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

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

$\Delta\rho_{\max} = 0.49 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.53 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

$\text{N1}-\text{Cd1}$	2.388 (4)	$\text{O4}-\text{Cd1}$	2.332 (4)
$\text{N2}-\text{Cd1}$	2.418 (4)	$\text{O5}-\text{Cd1}$	2.287 (3)
$\text{O1}-\text{Cd1}$	2.413 (3)	$\text{O6}-\text{Cd1}$	2.336 (4)
$\text{O3}-\text{Cd1}$	2.423 (4)		
$\text{O5}-\text{Cd1}-\text{O4}$	164.37 (12)	$\text{O4}-\text{Cd1}-\text{N2}$	93.18 (13)
$\text{O5}-\text{Cd1}-\text{O6}$	82.08 (12)	$\text{O6}-\text{Cd1}-\text{N2}$	144.24 (13)
$\text{O4}-\text{Cd1}-\text{O6}$	85.52 (13)	$\text{N1}-\text{Cd1}-\text{N2}$	66.47 (14)
$\text{O5}-\text{Cd1}-\text{N1}$	91.00 (13)	$\text{O1}-\text{Cd1}-\text{N2}$	133.86 (12)
$\text{O4}-\text{Cd1}-\text{N1}$	94.87 (13)	$\text{O5}-\text{Cd1}-\text{O3}$	98.34 (13)
$\text{O6}-\text{Cd1}-\text{N1}$	149.28 (13)	$\text{O4}-\text{Cd1}-\text{O3}$	88.24 (13)
$\text{O5}-\text{Cd1}-\text{O1}$	80.18 (12)	$\text{O6}-\text{Cd1}-\text{O3}$	77.80 (12)
$\text{O4}-\text{Cd1}-\text{O1}$	88.70 (13)	$\text{N1}-\text{Cd1}-\text{O3}$	132.91 (13)
$\text{O6}-\text{Cd1}-\text{O1}$	81.88 (12)	$\text{O1}-\text{Cd1}-\text{O3}$	159.62 (12)
$\text{N1}-\text{Cd1}-\text{O1}$	67.44 (13)	$\text{N2}-\text{Cd1}-\text{O3}$	66.44 (12)
$\text{O5}-\text{Cd1}-\text{N2}$	102.44 (13)		

Table 2
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O7-H7A\cdots O4^i$	0.85	2.26	3.055 (5)	156
$O7-H7B\cdots O5^{ii}$	0.87	1.88	2.729 (5)	167
$O8-H8A\cdots O6^{iii}$	0.88	2.02	2.814 (5)	150
$O8-H8B\cdots O7^{iv}$	0.86	2.00	2.816 (6)	158

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

The carbon bound H atoms were placed in calculated positions using a riding model with $U_{iso}(H) = 1.2U_{eq}(C)$ and $C-H = 0.95$. The solvent water H atoms were located and refined with $U_{iso}(H) = 1.2U_{eq}(O)$; however, the H atoms on the coordinating water molecules could not be found at geometrically sensible positions and were not included in the refinement.

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK* and *DENZO* (Otwinowski & Minor, 1997); program(s) used to solve structure: *DIRDIF99* (Beurskens *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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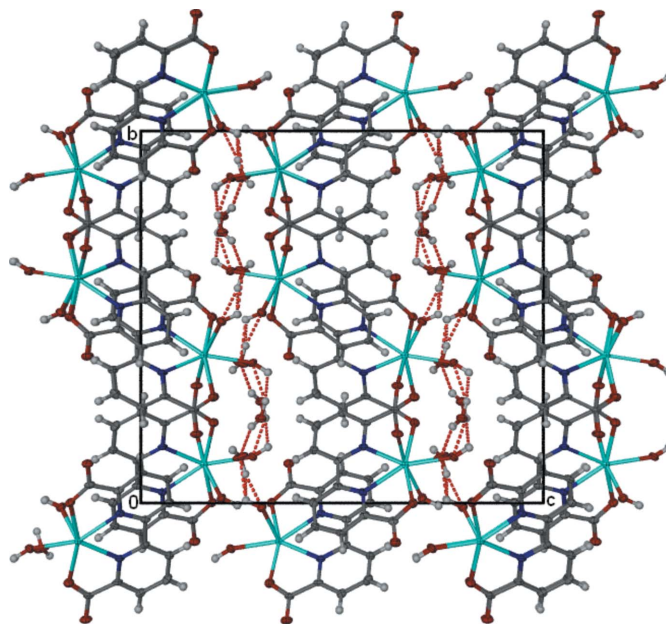


Figure 3
The molecular packing projected on the *bc* plane showing hydrogen bonds (dashed lines) holding the layered chains together in the crystal structure.

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