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Key indicators

Single-crystal X-ray study  
 T = 293 K  
 Mean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$   
 R factor = 0.021  
 wR factor = 0.064  
 Data-to-parameter ratio = 19.7

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

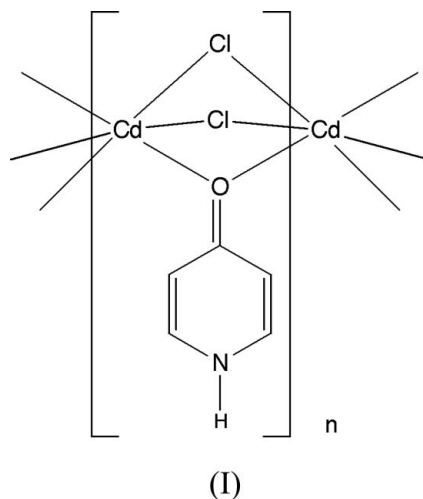
catena-Poly[cadmium(II)-di- $\mu$ -chloro- $\mu$ -4-1H-pyridone- $\kappa\text{O}:\kappa\text{O}$ ]

The crystal structure of the title compound,  $[\text{CdCl}_2(\text{C}_5\text{H}_5\text{NO})]_n$ , is reported. The asymmetric unit consists of one-half of a pyridone ligand which is located on a twofold axis, one cadmium cation which occupies a centre of inversion and one chloro ligand in a general position. This one-dimensional polymer represents a rare example of a structure with an oxygen-bonded bridging 4-pyridone ligand.

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Comment

The title polymer, (I), consists of infinite chains of face-sharing cadmium-centred octahedra with a metal–metal distance of 3.3404 (9) Å. For coordination polymers based on Cd this type of connectivity is much less common than edge-sharing (Hu & Englert, 2002; Hu *et al.*, 2003). A comparable Cd...Cd distance occurs in a chain polymer described by Puget *et al.* (1991). Neighbouring Cd atoms are bridged by two chloro and one oxygen-bonded pyridone ligands. Closest contacts between adjacent polymer strands are due to CH...Cl interactions at a distance of 2.74 Å; no classical hydrogen bonds are involved. The metal centres occupy inversion centres; the bridging O atoms, C3, N1 and the H atom attached to the latter are located on twofold axes in space group *I2/a*. We are aware of only one previous example of a bridging pyridone ligand (Gao *et al.*, 2005).



Experimental

Single crystals were obtained from an aqueous solution. One equivalent of  $\text{CdCl}_2$  was added to two equivalents of 4-hydroxypyridine in water. After stirring for 5 min, the solution was left to stand for 9 d at 315 K. The crystals are colourless platelets with dominant {010} faces.

## Crystal data

[CdCl<sub>2</sub>(C<sub>5</sub>H<sub>5</sub>NO)]  
*M<sub>r</sub>* = 278.42  
 Monoclinic, *I*2/*a*  
*a* = 7.510 (2) Å  
*b* = 15.531 (4) Å  
*c* = 6.6809 (18) Å  
 $\beta$  = 98.057 (5)°  
*V* = 771.6 (4) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 2.397 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 Cell parameters from 3571 reflections  
 $\theta$  = 2.6–28.4°  
 $\mu$  = 3.45 mm<sup>-1</sup>  
*T* = 293 (2) K  
 Platelet, colourless  
 0.47 × 0.40 × 0.07 mm

## Data collection

Bruker SMART CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: analytical (*SHELXTL*; Sheldrick, 1998)  
*T<sub>min</sub>* = 0.23, *T<sub>max</sub>* = 0.79  
 5114 measured reflections

967 independent reflections  
 916 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.032  
 $\theta_{\max}$  = 28.4°  
*h* = -10 → 10  
*k* = -20 → 20  
*l* = -8 → 8

## Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.021  
*wR* [*F*<sup>2</sup>] = 0.064  
*S* = 1.40  
 967 reflections  
 49 parameters

H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0333P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.47 \text{ e } \text{Å}^{-3}$   
 $\Delta\rho_{\min} = -0.57 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cd1—O1	2.3351 (14)	O1—C1	1.286 (3)
Cd1—CH <sup>i</sup>	2.5922 (9)	C1—C2	1.417 (2)
Cd1—Cl1	2.6288 (8)	C2—C3	1.352 (3)
Cd1···Cd1 <sup>i</sup>	3.3404 (10)	C3—N1	1.343 (3)
O1 <sup>ii</sup> —Cd1—Cl1	102.33 (3)	Cd1 <sup>i</sup> —Cl1—Cd1	79.55 (3)
O1—Cd1—Cl1	77.67 (3)	Cd1 <sup>i</sup> —O1—Cd1	91.33 (7)
Cl1 <sup>i</sup> —Cd1—Cl1	85.67 (3)		

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

All H atoms were introduced in idealized positions (C—H = 0.93 Å and N—H = 0.86 Å) and were refined using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ . The setting *I*2/*a* for space group 15 was preferred over *C*2/*c* because the very obtuse monoclinic angle in the latter resulted in correlations between *x* and *z* parameters in the refinement matrix.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 1999); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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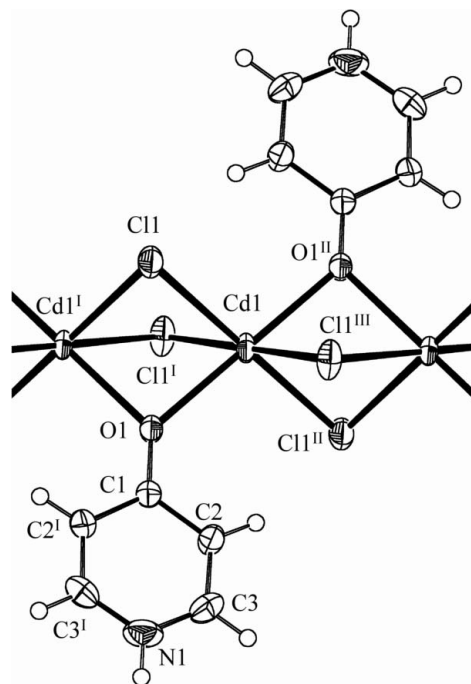


Figure 1

Displacement ellipsoid plot (50% probability level, H atoms with arbitrary radii) of a section of the coordination polymer. The chain extends along *c*. [Symmetry operations: (I)  $\frac{1}{2} - x, y, -z$ ; (II)  $\frac{1}{2} - x, \frac{1}{2} - y, \frac{1}{2} - z$ ; (III)  $x, \frac{1}{2} - y, \frac{1}{2} + z$ .]

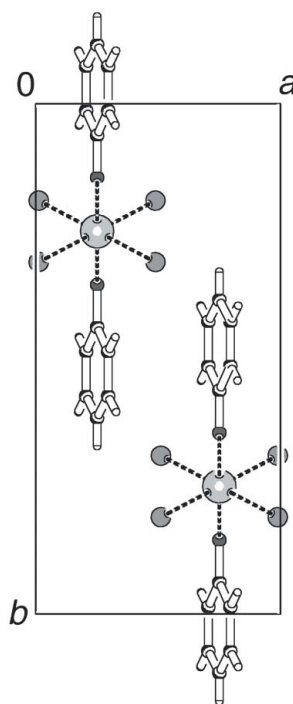


Figure 2

Packing diagram of two polymer chains in the unit cell. The view is in the chain direction, along *c*.

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