

catena-Poly[[[tris(pyridine- κ N)cadmium(II)]- μ -{2-[3-(1-carboxylatomethyl)-4-nitrophenyl]-propionato- κ^2 O,O'}] dihydrate]

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Key indicators

Single-crystal X-ray study
 T = 295 K
 Mean σ (C–C) = 0.006 Å
 Disorder in main residue
 R factor = 0.037
 wR factor = 0.094
 Data-to-parameter ratio = 17.3

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

The dicarboxylate dianion in the title compound, [Cd(C₅H₅N)₃(C₁₁H₉NO₆)₂·2H₂O], uses its carboxylatomethyl arm to chelate to one Cd atom and the 1-carboxylatoethyl arm to bind to another Cd atom through only one terminal O atom, giving rise to a polymeric chain that runs along [10 $\bar{1}$]. The octahedral geometry of the Cd atom is completed by the N atoms of three pyridine N-atom donors.

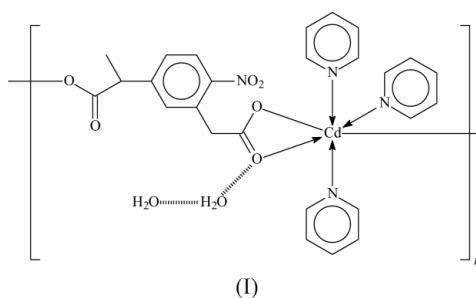
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Comment

We have recently synthesized 2-(3-carboxymethyl-4-nitrophenyl)propionic acid (Huo *et al.*, 2004) for complexation studies with metal systems. In the title cadmium derivative, the dicarboxylate dianion uses its acetate arm to chelate to one Cd atom and the propionate arm to bind to another Cd atom through only one terminal O. The six-coordinate geometry is distorted owing to the proximity of the uncoordinated O atom (Fig. 1). In the acetate arm, the C–O single and double bonds are not significantly different [1.254 (3) and 1.241 (4) Å], as are those in the propionate arm [1.249 (4) and 1.236 (4) Å]. Nevertheless, if the shorter of the pair of bonds is regarded as the double bond, then the water molecule O1 w forms hydrogen bonds to the carbonyl O atom of the same dianion. On the other hand, the O2 w water molecule forms hydrogen bonds to the O1 w O atom and also to the uncoordinated carbonyl O atom (Table 2). The resulting structure assumes a chain motif along [10 $\bar{1}$]. The monodentate/bidentate nature of the dianion contrasts with the monodentate nature of both arms in the tripyridinecadmium benzene-1,3-dioxyacetate complex; the six-coordinate geometry is completed by a water ligand that engages in hydrogen-bonding interactions (Gao *et al.*, 2004).



The tripyridine adduct of cadmium bis(4-hydroxybenzoate) also displays a chelating carboxylate anion and a unidentate carboxylate anion; the compound crystallizes as a pyridine solvate (Shnulin *et al.*, 1981).

Experimental

The title complex was prepared by the addition of 2-(3-carboxymethyl-4-nitrophenyl)propionic acid (5.06 g, 20 mmol) (Gao *et al.*, 2004) and pyridine (1.5 ml) to an aqueous solution of cadmium nitrate tetrahydrate (6.17 g, 20 mmol). The resulting mixture was heated for 30 min. Yellow crystals separated from the solution after several days. Analysis calculated for $C_{26}H_{28}CdN_4O_8$: C 49.03, H 4.43, N 8.80%; found: C 49.08, H 4.51 N 8.84%.

Crystal data

$[Cd(C_5H_5N)_3(C_{11}H_9NO_6)] \cdot 2H_2O$
 $M_r = 636.92$
 Monoclinic, $P2_1/n$
 $a = 9.748$ (2) Å
 $b = 19.923$ (4) Å
 $c = 14.598$ (3) Å
 $\beta = 95.92$ (3)°
 $V = 2820$ (1) Å³
 $Z = 4$

$D_x = 1.500$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 23092 reflections
 $\theta = 3.2$ – 27.5 °
 $\mu = 0.83$ mm⁻¹
 $T = 295$ (2) K
 Prism, yellow
 $0.35 \times 0.24 \times 0.18$ mm

Data collection

Rigaku R-AXIS RAPID diffractometer
 ω scans
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
 $T_{min} = 0.594$, $T_{max} = 0.865$
 25939 measured reflections

6413 independent reflections
 4950 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.032$
 $\theta_{max} = 27.5$ °
 $h = -10 \rightarrow 12$
 $k = -25 \rightarrow 25$
 $l = -18 \rightarrow 18$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.094$
 $S = 1.04$
 6413 reflections
 371 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0464P)^2 + 1.0608P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.001$
 $\Delta\rho_{max} = 0.63$ e Å⁻³
 $\Delta\rho_{min} = -0.52$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Cd1—O1	2.328 (2)	Cd1—N1	2.364 (3)
Cd1—O2	2.579 (2)	Cd1—N2	2.403 (3)
Cd1—O3 ⁱ	2.293 (2)	Cd1—N3	2.330 (3)
O1—Cd1—O2	52.75 (7)	O2—Cd1—N3	101.64 (9)
O1—Cd1—O3 ⁱ	137.18 (7)	O3 ⁱ —Cd1—N1	86.50 (9)
O1—Cd1—N1	97.66 (9)	O3 ⁱ —Cd1—N2	137.08 (8)
O1—Cd1—N2	85.65 (8)	O3 ⁱ —Cd1—N3	94.37 (9)
O1—Cd1—N3	90.01 (9)	N1—Cd1—N2	84.35 (9)
O2—Cd1—O3 ⁱ	84.75 (7)	N1—Cd1—N3	167.75 (9)
O2—Cd1—N1	90.61 (9)	N2—Cd1—N3	86.74 (9)
O2—Cd1—N2	137.06 (8)		

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

Table 2

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1w—H1w1 \cdots O2	0.85 (1)	2.08 (2)	2.882 (4)	159 (4)
O1w—H1w2 \cdots O4	0.85 (1)	2.04 (1)	2.851 (3)	160 (3)
O2w—H2w1 \cdots O1w	0.86 (1)	2.00 (1)	2.860 (4)	176 (4)
O2w—H2w2 \cdots O3 ⁱ	0.86 (1)	1.98 (2)	2.812 (4)	165 (4)

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

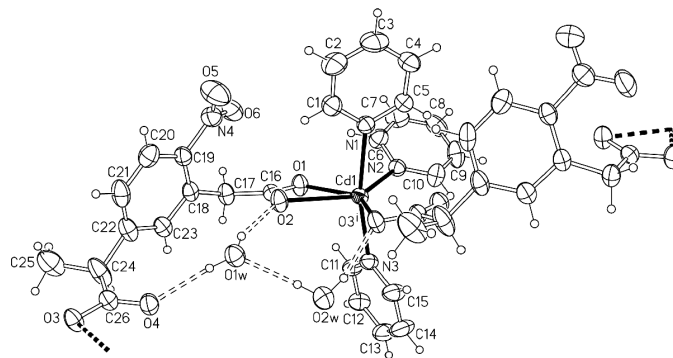


Figure 1

ORTEPII (Johnson, 1976) plot of a portion of the polymeric structure of (I) with ellipsoids shown at the 30% probability level. H atoms are drawn as spheres of arbitrary radii. Broken lines indicate hydrogen bonds.

The structure is disordered in the methyl substituent of the 1-carboxylatoethyl arm of the dicarboxylate ligand. The methyl C atom (C25) was refined as two atoms of 0.5 site occupancy, as the occupancy could not be refined; the C24—C25 and C24—C25' distances were restrained to 1.50 (1) Å and the displacement parameters of the unprimed and primed atoms were made equal to each other. The vibration of the C24 and C25 atoms was restrained to be approximately isotropic. The H atoms were placed in calculated positions [C—H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ (aromatic); C—H 0.98 Å and $U_{iso} = 1.2U_{eq}(C)$ (methine); C—H = 0.97 Å and $U_{iso} = 1.2U_{eq}(C)$ (methylene); C—H = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ (methyl)], and were included in the refinement in the riding-model approximation. The water H atoms were located and refined with distance restraints of O—H = 0.85 (1) Å and H \cdots H = 1.39 (1) Å.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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