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#### Key indicators

Single-crystal X-ray study T = 295 KMean  $\sigma$ (C–C) = 0.005 Å R factor = 0.052 wR factor = 0.135 Data-to-parameter ratio = 15.2

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

# catena-Poly[[[diaquabis(4-carboxyphenoxyacetato- $\kappa$ O)cobalt(II)]- $\mu$ -4,4'-bipyridine- $\kappa^2$ N,N'] dihydrate]

The title compound,  $\{[Co(C_9H_7O_5)_2(C_{10}H_8N_2)(H_2O)_2]$ -2 $H_2O\}_n$ , has an infinite one-dimensional chain structure along the *a* axis. The Co atom and the mid-point of the bipyridine group are located on inversion centres. A three-dimensional supramolecular network is built up by  $O-H\cdots O$  hydrogenbonding interactions between water molecules, carboxyl hydroxy groups and carboxylate groups.

#### Comment

In recent years, much interest has been focused on using aromatic carboxylic acid ligands to construct supramolecular architectures by self-assembly (Zhang et al., 2004). 4-Carboxyphenoxyacetic acid is a multidentate flexible ligand with versatile binding abilities and the capability of participating in hydrogen bonds as both a donor and an acceptor, thus representing an excellent candidate for the construction of supramolecular complexes. Recently, four one-dimensional Co<sup>II</sup> polymers based on the 4-carboxyphenoxyacetic acid ligand were reported, namely {[Co(4-carboxyphenoxyacetate)(3-hydroxypyridine)<sub>2</sub>(H<sub>2</sub>O)<sub>2</sub>](H<sub>2</sub>O)<sub>n</sub> (Gao *et al.*, [Co(4-carboxyphenoxyacetate)(2,2'-bipyridine)-2004), (H<sub>2</sub>O)]<sub>n</sub> (Gao, Huo, Gu, Liu & Zhao, 2005), [Co(4-carboxyphenoxyacetate)(1,10-phenanthroline)(H<sub>2</sub>O)]<sub>n</sub> (Gao, Huo, Gu, Zhao & Zhao, 2005) and [Co(4-carboxyphenoxyacetate)<sub>2</sub>(4,4'-bipyridine)<sub>2</sub>(H<sub>2</sub>O)<sub>2</sub>]·4H<sub>2</sub>O (Deng *et al.*, 2005). In this paper, we report the new title compound, (I), synthesized by the reaction of 4-carboxyphenoxyacetic acid, 4,4'bipyridine and cobalt dinitrate hexahydrate in an aqueous solution, which forms an infinite one-dimensional chain structure.



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#### Figure 1

Part of the polymeric structure of the title complex, with the atomlabelling scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are represented as spheres of arbitrary radii. [Symmetry codes: (i) -x, 1 - y, 1 - z; (ii) 1 - x, 1 - y, 1 - z].

Complex (I) consists of an infinite one-dimensional chain and uncoordinated water molecules (Fig. 1). The Co atom is located on an inversion centre and is hexacoordinated by two N atoms of the 4,4'-bipyridine ligands, two O atoms of the 4carboxyphenoxyacetate groups and two water molecules. There is another inversion centre at the mid-point of the central C-C bond of the 4,4'-bipyridine unit.

In the crystal structure, a three-dimensional supramolecular network is constructed via  $O-H\cdots O$  hydrogen-bonding interactions between water molecules, carboxylic acid hydroxy groups and carboxylate O atoms (Table 1, Fig. 2).

## **Experimental**

4-Carboxyphenoxyacetic acid was prepared by the reaction of chloroacetic acid with 4-hydroxybenzoic acid, and all other chemicals were of analytical grade and used without further purification. 4-Carboxyphenoxyacetic acid (0.98 g, 5 mmol) and 4,4'-bipyridine (0.78 g, 5 mmol) were dissolved in hot water (10 ml), and the pH was adjusted to about 6 using 0.2 M NaOH solution with stirring. Cobalt(II) nitrate hexahydrate (1.46 g, 5 mmol) was added to the mixed solution. The resulting solution was filtered, and the filtrate was allowed to stand in a desiccator at room temperature for several days. Pink block crystals of (I) were obtained.

#### Crystal data

$[Co(C_9H_7O_5)_2(C_{10}H_8N_2)(H_2O)_2]$	V = 1460.6 (6) Å <sup>3</sup>
$2H_2O$	Z = 2
$M_r = 677.47$	$D_x = 1.540 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
a = 11.726 (2) Å	$\mu = 0.66 \text{ mm}^{-1}$
b = 11.135 (2) Å	T = 295 (2) K
c = 12.485 (3) Å	Block, pink
$\beta = 116.37 \ (3)^{\circ}$	$0.39 \times 0.36 \times 0.26 \text{ mm}$
Data collection	
Rigaku R-AXIS RAPID	13737 measured reflections

diffractometer  $\omega$  scans Absorption correction: multi-scan (Higashi, 1995)  $T_{\min} = 0.780, T_{\max} = 0.844$  13737 measured reflections 3324 independent reflections 2475 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.054$  $\theta_{\text{max}} = 27.5^{\circ}$ 



#### Figure 2

A partial packing view, showing the three-dimensional hydrogen-bonding network. Dashed lines indicate the hydrogen-bonding interactions. H atoms not involved in hydrogen bonds have been omitted for clarity. [Symmetry code: (i) 1 + x, y - 1, z].

#### Refinement

Refinement on $F^2$	w =
$R[F^2 > 2\sigma(F^2)] = 0.052$	
$wR(F^2) = 0.135$	w
S = 1.06	$(\Delta/c$
3324 reflections	$\Delta \rho_{\rm n}$
218 parameters	$\Delta \rho_n$
H atoms treated by a mixture of	
independent and constrained	
refinement	

 $v = 1/[\sigma^{2}(F_{o}^{2}) + (0.074P)^{2} + 0.3963P]$ where  $P = (F_{o}^{2} + 2F_{c}^{2})/3$  $\Delta/\sigma)_{max} < 0.001$  $\Delta\rho_{max} = 0.45 \text{ e} \text{ Å}^{-3}$  $\Delta\rho_{min} = -0.51 \text{ e} \text{ Å}^{-3}$ 

#### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O5-H5A\cdots O7^{i}$	0.82	1.77	2.586 (3)	175
O6−H61···O2 <sup>ii</sup>	0.835 (15)	1.860 (14)	2.648 (3)	157 (3)
O6−H62···O4 <sup>iii</sup>	0.82 (4)	1.879 (11)	2.695 (3)	171 (3)
O7−H71···O2 <sup>ii</sup>	0.83 (4)	1.93 (4)	2.751 (3)	169 (5)
$O7-H72 \cdot \cdot \cdot O1^{iv}$	0.83 (4)	2.044 (17)	2.838 (3)	159 (4)
Symmetry codes:	(i) $x + 1, y$	-1, z; (ii)	-x + 1, -y + 1,	-z + 1; (iii)

Symmetry codes: (1) x + 1, y - 1, z; (11) -x + 1, -y + 1, -z + 1; (11)  $-x + 2, y + \frac{1}{2}, -z + \frac{3}{2};$  (iv)  $-x + 1, y + \frac{1}{2}, -z + \frac{3}{2}.$ 

H atoms bound to C atoms and to the hydroxy group were placed in calculated positions and treated as riding on their parent atoms, with C-H = 0.93 (C<sub>aromatic</sub>) or 0.97 Å (C<sub>methylene</sub>) and O-H = 0.82 Å, and with  $U_{iso}(H) = 1.2U_{eq}(C,O)$ . Water H atoms were located in a difference Fourier map and refined using restrained distances O-H = 0.82 (1) Å and H···H = 1.39 (2) Å, with  $U_{iso}(H) = 1.5U_{eq}(O)$ .

Data collection: *PROCESS-AUTO* (Rigaku, 2002); cell refinement: *PROCESS-AUTO*; data reduction: *PROCESS-AUTO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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