

catena-Poly[bis[aqua(2,2-bipyridine- κ^2N,N')cobalt(II)]- μ -1,2,4,5-benzenetetracarboxylato- $\kappa^4O^1:O^2:O^3:O^4$]

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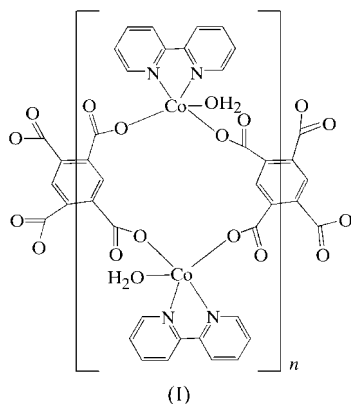
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In the title complex, $[\text{Co}_2(\text{C}_{10}\text{H}_2\text{O}_8)(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_2]$, the four carboxylate groups are fully deprotonated and coordinate to four Co^{II} cations in a monodentate fashion, forming a one-dimensional ribbon-like double-chain structure, with centrosymmetric $[\text{Co}_2(\text{C}_{10}\text{H}_2\text{O}_8)(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_2]$ repeating units and a cavity of approximately $6.8 \times 6.6 \text{ \AA}$. Moreover, a three-dimensional supramolecular structure is formed by face-to-face π - π interactions between the aromatic rings of the 2,2'-bipyridine moieties of two adjacent chains, and by hydrogen-bonding interactions between the coordinated aqua O atom and the coordinated carboxyl O atom from different chains.

Comment

Coordinate polymer solids with a variety of cavities or channels are currently under intense study because of the scope they offer for the generation of new materials with a range of potentially useful properties (Yaghi *et al.*, 1998; Stein *et al.*,



1993). Popular bridging ligands in this field are 1,2,4,5-benzenetetracarboxylic acid (H_4TCB) and 4,4'-bipyridine, and their derivatives (Eddaoudi *et al.*, 2001). We have studied some transition metal complexes bridged by TCB^{4-} or $\text{H}_2\text{TCB}^{2-}$ anions, such as those with Co^{II} (Wang *et al.*, 2000),

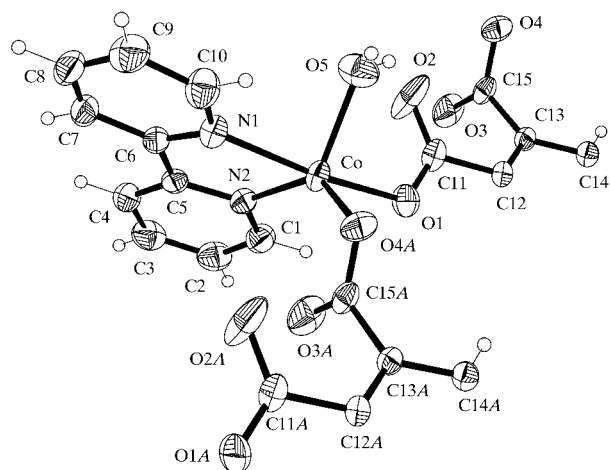


Figure 1

The coordination environment of the Co^{II} ion in (I), with the atom-numbering scheme, showing displacement ellipsoids at the 50% probability level.

Ni^{II} (Cheng *et al.*, 2001) and Cu^{II} (Hu *et al.*, 2003). In the course of our studies, the title novel infinite one-dimensional coordination polymer, $[\text{Co}_2(\text{TCB})(2,2'\text{-bipy})_2(\text{H}_2\text{O})_2]_n$ (2,2'-bipy is 2,2'-bipyridine), (I), was prepared and its structure is reported here. It should be noted that the synthetic conditions, such as H-atom receptors, temperature and solvents, *etc.*, play an important role in determining the compositions of these complexes.

In (I), each Co^{II} cation has a five-coordination environment, completed by two carboxyl O atoms belonging to two TCB^{4-} anions, one aqua O atom and two N atoms from one 2,2'-bipy ligand (Fig. 1). The Co—O bond lengths are all about 2.05 Å, while the Co—N1 bond [2.1590 (17) Å] is longer than the Co—N2 bond [2.0640 (17) Å]. An infinite one-dimensional polymer with a double-chain structure is formed by the Co^{II} cations, the μ_4 -bridging TCB^{4-} anions, the aqua molecules and the terminal 2,2'-bipy ligands (Fig. 2).

The coordination mode of the TCB^{4-} anion in (I) is similar to that in $[\text{Cu}_2(\text{TCB})(\text{phen})_2]_n \cdot n\text{H}_2\text{O}$ (phen is 1,10-phenanthroline; Shi *et al.*, 2001). The four carboxylate groups are fully deprotonated and coordinate to four Co^{II} cations in a monodentate fashion, forming a one-dimensional ribbon-like

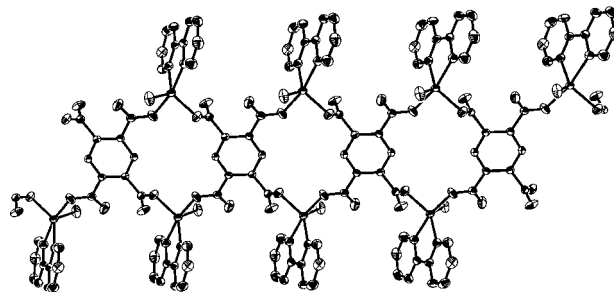


Figure 2

The double-chain structure of (I), showing the cavities.

double-chain structure with cavities of approximately 6.8×6.6 Å. Intermolecular hydrogen-bonding interactions of 2.623 (2) Å are found between the coordinated aqua O atom and the coordinated carboxyl O atom from different chains, resulting in a two-dimensional supramolecular structure.

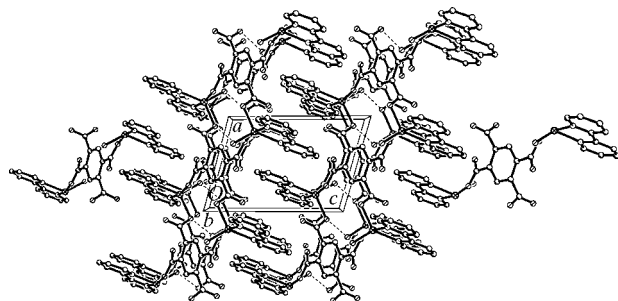


Figure 3
The three-dimensional network structure of (I).

Furthermore, a three-dimensional network structure is formed by face-to-face π - π interactions of approximately 3.78 Å between the aromatic rings of the 2,2'-bipy ligands of two adjacent chains (Fig. 3).

Experimental

A solution of dimethylformamide (10 ml) containing $\text{CoCl}_2 \cdot 6\text{H}_2\text{O}$ (0.5 mmol, 0.119 g) and H_4TCB (0.5 mmol, 0.127 g) was added slowly to a solution of dimethylformamide (10 ml) containing 2,2'-bipyridine (0.5 mmol, 0.078 g). The mixture was stirred for 30 min and left to stand at room temperature for about three weeks. Deep-red prism-shaped crystals of (I) were obtained.

Crystal data

$[\text{Co}_2(\text{C}_{10}\text{H}_2\text{O}_8)(\text{C}_{10}\text{H}_8\text{N}_2)_2(\text{H}_2\text{O})_2]$	$Z = 1$
$M_r = 716.38$	$D_x = 1.766 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 7.5859$ (8) Å	Cell parameters from 534 reflections
$b = 8.9838$ (10) Å	$\theta = 2.4$ – 23.0°
$c = 10.6729$ (11) Å	$\mu = 1.30 \text{ mm}^{-1}$
$\alpha = 80.441$ (2) $^\circ$	$T = 273$ (2) K
$\beta = 72.913$ (2) $^\circ$	Prism, red
$\gamma = 77.096$ (2) $^\circ$	$0.31 \times 0.21 \times 0.12 \text{ mm}$
$V = 673.75$ (12) Å ³	

Data collection

Bruker SMART CCD area-detector diffractometer	2406 independent reflections
φ and ω scans	2273 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 2000)	$R_{\text{int}} = 0.015$
$T_{\text{min}} = 0.727$, $T_{\text{max}} = 0.855$	$\theta_{\text{max}} = 25.2^\circ$
4962 measured reflections	$h = -9 \rightarrow 9$
	$k = -10 \rightarrow 10$
	$l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.027$
 $wR(F^2) = 0.071$
 $S = 1.08$
 2406 reflections
 217 parameters
 H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0362P)^2 + 0.3213P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.023$
 $\Delta\rho_{\text{max}} = 0.31 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.22 \text{ e \AA}^{-3}$

Table 1
Hydrogen-bonding geometry (Å, $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O5–H12 \cdots O4 ⁱ	0.854 (9)	1.770 (10)	2.623 (2)	177 (3)
O5–H13 \cdots O2	0.855 (10)	2.010 (17)	2.715 (2)	139 (2)

Symmetry code: (i) $1 - x, 1 - y, -z$.

The positions and isotropic displacement parameters of the water H atoms, H12 and H13, were refined subject to $O-H = 0.85$ (1) Å. The other H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of 0.93 Å, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: SMART (Bruker, 2000); cell refinement: SMART; data reduction: SAINT (Bruker, 2000); program(s) used to solve structure: SHELXTL (Bruker, 2000); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: TA1428). Services for accessing these data are described at the back of the journal.

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