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## Key indicators

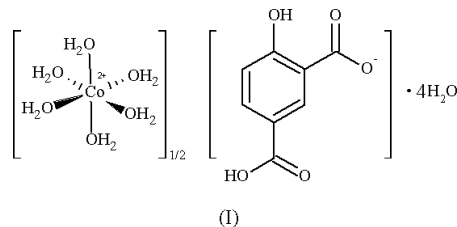
Single-crystal X-ray study  
T = 293 K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$   
R factor = 0.035  
wR factor = 0.095  
Data-to-parameter ratio = 10.2For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Hexaaquacobalt(II) bis(4-hydroxyisophthalate)  
tetrahydrate

Hydrothermal reaction of 4-hydroxyisophthalic acid and  $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$  in water resulted in a new mononuclear cobalt compound,  $[\text{Co}(\text{H}_2\text{O})_6](\text{C}_8\text{H}_5\text{O}_5)_2 \cdot 4\text{H}_2\text{O}$ . The occurrence of an intricate hydrogen-bond network leads to the formation of a three-dimensional structure. The cation lies on a crystallographic inversion center, while the anion and water molecules are in general positions.

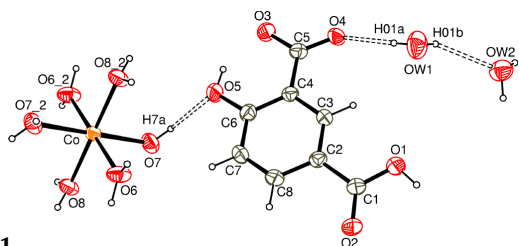
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## Comment

The design and synthesis of new compounds with structures defined through non-covalent connections is currently of great interest because of their potential application in certain areas, such as gas storage (Atwood *et al.*, 2002), catalysis (Thomas, 1998), molecular transport (Gardner *et al.*, 1995) and biosynthesis (Lippert, 1997). In the fields of supramolecular chemistry and crystal engineering, hydrogen bonding plays an important role. Many examples of compounds that assemble by hydrogen-bonding interactions have been reported in previous publications (Yuan *et al.*, 2002). The great structural diversity of crystal packing results not only from the variability of the organic ligands and the ligand-to-metal connections, but also from the types and fashions of hydrogen bonding and other weak intermolecular interactions (Naumann *et al.*, 2002). Although 4-hydroxyisophthalic acid, which possesses four carboxylic O atoms and one hydroxyl O atom, is a good candidate for the generation of many linking modes, including hydrogen bonding, it has rarely been reported in recent years. Here we report the preparation and characterization of the first 4-hydroxyisophthalate-cobalt(II) complex,  $[\text{Co}(\text{H}_2\text{O})_6](\text{C}_8\text{H}_5\text{O}_5)_2 \cdot 4\text{H}_2\text{O}$ , (I).



An X-ray analysis reveals that (I) possesses a mononuclear structure, with the cation on a crystallographic inversion center. A view of the cobalt ion coordination is shown in Fig. 1; the metal center is coordinated in an octahedral geometry by six water molecules with  $\text{Co}-\text{O}$  distances ranging from 2.0386 (18) to 2.1371 (18)  $\text{\AA}$ . The 4-hydroxyisophthalate ligand, in which only one carboxylic acid group is deprotonated, is not directly coordinated to the cobalt ion but is involved in a hydrogen-bond interaction with a coordinated water molecule. Different types of hydrogen-bonding interactions are observed (Table 2) in which O atoms may act as



**Figure 1**

The cation, anion and water molecules, with their hydrogen-bond interactions. Displacement ellipsoids are drawn at the 50% probability level. The suffix 2 corresponds to symmetry code (i) in Table 1.

donors or acceptors: (a) hydrogen bonds between coordinated water and isolated water molecules, with O...O distances ranging from 2.627 (3) to 2.982 (3) Å; (b) hydrogen bonds between coordinated water molecules and carboxylic acid O atoms, with O...O distances ranging from 2.744 (2) to 3.031 (3) Å; (c) hydrogen bond between coordinated water and hydroxyl atom O5, with an O...O distance of 2.886 (3) Å; (d) hydrogen bonds between isolated water molecules and carboxylic acid oxygen atoms, with O...O distances of 2.754 (3) and 2.865 (3) Å; (e) hydrogen bond between the hydroxyl and carboxylic acid groups, with an O...O distance of 2.675 (2) Å. These hydrogen-bonding interactions result in a three-dimensional network (Fig. 2).

## Experimental

A mixture of 4-hydroxyisophthalic acid (0.25 mmol, 0.045 g),  $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$  (0.2 mmol, 0.05 g) and  $\text{H}_2\text{O}$  (20 ml) was sealed in a 25 ml Teflon-lined stainless-steel reactor and heated at 433 K for 72 h, and yielding a red solution. After cooling and evaporation of the red solution over two weeks, red crystals were isolated by filtration (yield, 50%).

### Crystal data

$[\text{Co}(\text{H}_2\text{O})_6](\text{C}_8\text{H}_5\text{O}_5)_2 \cdot 4\text{H}_2\text{O}$   
 $M_r = 601.33$   
 Triclinic,  $P\bar{1}$   
 $a = 7.1863$  (6) Å  
 $b = 8.5173$  (7) Å  
 $c = 9.8959$  (7) Å  
 $\alpha = 92.088$  (1)°  
 $\beta = 91.424$  (1)°  
 $\gamma = 90.548$  (1)°  
 $V = 605.08$  (8) Å<sup>3</sup>

$Z = 1$   
 $D_x = 1.650$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 34 reflections  
 $\theta = 2.1$ – $25.0^\circ$   
 $\mu = 0.80$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 Block, red  
 $0.28 \times 0.16 \times 0.10$  mm  
 2076 independent reflections  
 1847 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.019$   
 $\theta_{\text{max}} = 25.0^\circ$   
 $h = -7 \rightarrow 8$   
 $k = -10 \rightarrow 10$   
 $l = -11 \rightarrow 11$

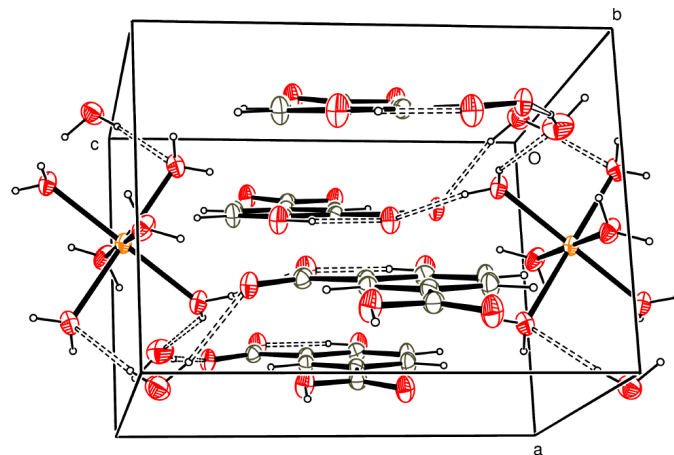
### Data collection

Siemens SMART CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\text{min}} = 0.857$ ,  $T_{\text{max}} = 0.923$   
 3102 measured reflections

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.095$   
 $S = 1.07$   
 2076 reflections  
 203 parameters  
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0524P)^2 + 0.2322P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.54$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.43$  e Å<sup>-3</sup>



**Figure 2**

Packing of the title complex, showing the three-dimensional hydrogen-bonding network.

**Table 1**

Selected geometric parameters (Å, °).

Co—O6	2.0386 (18)	O2—C1	1.230 (3)
Co—O8	2.0873 (17)	O3—C5	1.270 (3)
Co—O7	2.1371 (18)	O4—C5	1.263 (3)
O1—C1	1.310 (3)	O5—C6	1.344 (3)
O6—Co—O8 <sup>i</sup>	90.38 (7)	O2—C1—O1	122.6 (2)
O6—Co—O8	89.62 (7)	O2—C1—C2	122.5 (2)
O6—Co—O7	88.04 (8)	O1—C1—C2	114.8 (2)
O6 <sup>i</sup> —Co—O7	91.96 (8)	O4—C5—O3	123.1 (2)
O8 <sup>i</sup> —Co—O7	93.09 (8)	O4—C5—C4	119.2 (2)
O8—Co—O7	86.91 (8)	O3—C5—C4	117.7 (2)

Symmetry code: (i)  $1 - x, 1 - y, 2 - z$ .

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1—H1...O2 <sup>i</sup>	0.82	1.86	2.675 (2)	173
O5—H5...O3	0.82	1.80	2.527 (2)	148
O6—H6A...OW1 <sup>ii</sup>	0.84 (3)	1.82 (3)	2.627 (3)	161 (3)
O6—H6B...O4 <sup>iii</sup>	0.84 (1)	1.86 (1)	2.674 (2)	164 (3)
O7—H7A...O5	0.84 (1)	2.10 (2)	2.886 (3)	156 (3)
O7—H7A...O2 <sup>iv</sup>	0.84 (1)	2.59 (3)	3.031 (3)	115 (3)
O7—H7B...OW2 <sup>v</sup>	0.81 (3)	2.05 (3)	2.847 (3)	167 (3)
O8—H8A...OW1 <sup>v</sup>	0.84 (1)	2.25 (2)	2.982 (3)	147 (2)
O8—H8B...O3 <sup>iii</sup>	0.84 (1)	1.92 (1)	2.744 (2)	169 (3)
OW2—HW2A...O7 <sup>vi</sup>	0.84 (3)	2.12 (3)	2.952 (3)	175 (3)
OW2—HW2B...O4 <sup>vii</sup>	0.84 (1)	2.04 (3)	2.865 (3)	167 (4)
OW1—HW1A...O4	0.85 (2)	1.90 (2)	2.754 (3)	174 (4)
OW1—HW1B...OW2	0.85 (3)	2.07 (3)	2.759 (3)	138 (3)

Symmetry codes: (i)  $-x, -1 - y, 1 - z$ ; (ii)  $1 - x, -y, 1 - z$ ; (iii)  $x, y, 1 + z$ ; (iv)  $x, 1 + y, z$ ; (v)  $x, 1 + y, 1 + z$ ; (vi)  $-x, -y, 1 - z$ ; (vii)  $-x, -y, -z$ .

The H atoms were located in an electron-density difference map. The H atoms of C—H and hydroxyl O—H groups were placed in calculated positions (C—H = 0.96 Å and O—H = 0.82 Å), and were allowed to refine as riding, with displacement parameters fixed at 120% of those of their parent atoms. The H atoms of the water molecules (free and coordinated) were refined with O—H distances restrained to 0.84 (1) Å and H...H distances restrained to 1.37 (1) Å with displacement parameters fixed at 150% of the parent O atoms.

These restraints ensure a reasonable geometry for the water molecules.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SMART*; data reduction: *SAINT* (Siemens, 1994; program(s) used to solve structure: *SHELXTL* (Siemens, 1994); program(s) used to refine structure: *SHELXTL*; molecular graphics: *ORTEP-3* (Farrugia, 1997) and *ORTEP-III* (Burnett & Johnson, 1996); software used to prepare material for publication: *SHELXTL*.

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