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Key indicators

Single-crystal X-ray study
T = 293 K
Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$
R factor = 0.053
wR factor = 0.149
Data-to-parameter ratio = 14.4

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

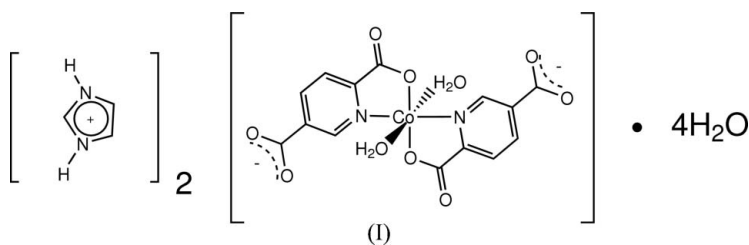
Bis(imidazolium) diaquabis(pyridine-2,5-dicarboxylato- κ^2N,O)cobaltate(II) tetrahydrate

The title complex, $(\text{C}_3\text{H}_5\text{N}_2)_2[\text{Co}(\text{C}_7\text{H}_3\text{NO}_4)_2(\text{H}_2\text{O})_2] \cdot 4\text{H}_2\text{O}$, was prepared by a hydrothermal reaction. Each Co^{II} atom is located on a crystallographic inversion centre and displays a distorted octahedral coordination geometry. The face-to-face distance of $3.529(7) \text{ \AA}$ between partially overlapped parallel pyridine rings reflects a $\pi-\pi$ stacking interaction between neighbouring Co^{II} complex molecules. A network of $\text{O}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds helps to stabilize the crystal packing.

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Comment

The unique strength and direction of hydrogen bonding play a dominant role in the generation of a variety of molecular architectures and molecular recognition in chemical and biological sciences (Lehn *et al.*, 1996; Steiner, 2002). The ligand pyridine-2,5-dicarboxylic acid, known both as a multiple proton donor and acceptor (MacDonald *et al.*, 2000), can use its carboxylate O and pyridine N atoms, to form both monodentate and/or multidentate $M-\text{O}$ and $M-\text{N}$ metal-ligand bonds (Zeng *et al.*, 2004). Protonated imidazole (Him) readily takes part in forming hydrogen-bonded networks (MacDonald *et al.*, 2000). We report here the hydrothermal synthesis and crystal structure of the title complex, (I).



In (I), each Co^{II} atom is coordinated by two N atoms and two O atoms from two chelating pyridine-2,5-dicarboxylate (pdc) ligands, and two O atoms from two aqua ligands to furnish a distorted octahedral geometry. The aqua ligands are located at the axial positions (Table 1 and Fig. 1).

The face-to-face distance of $3.529(7) \text{ \AA}$ between partially overlapped parallel pyridine rings reflects a $\pi-\pi$ stacking interaction between neighbouring Co^{II} complex anions. In addition, neighbouring Co^{II} complex anions are linked to each other by $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds. With the help of $\pi-\pi$ stacking interactions and hydrogen bonds, these complex anions form one-dimensional linear chains running along the *a* axis. Protonated imidazoles link to the chains by $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds (Fig. 2 and Table 2). These chains are linked into a complex network by $\text{O}-\text{H} \cdots \text{O}$ (uncoordinated water and carboxylate, aqua ligands and uncoordinated water) and

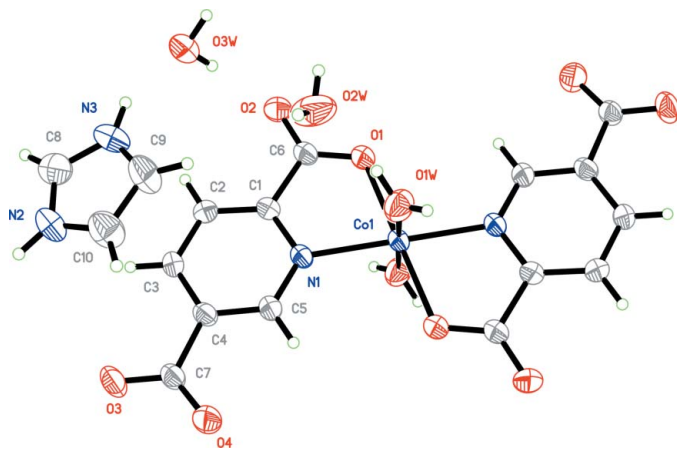


Figure 1
The molecular structure of (I), shown with 50% probability displacement ellipsoids. Unlabelled atoms are related to labelled atoms by the symmetry operator $(-x, 2 - y, -z)$.

$N-H \cdots O$ (protonated imidazoles and uncoordinated water) hydrogen bonds (Fig. 3).

Experimental

Pyridine-2,5-dicarboxylic acid (0.5 mmol), imidazole (0.5 mmol), $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (0.5 mmol) and water (8 ml) were stirred for 15 min in air, then transferred and sealed in a 23 ml Teflon reactor, which was heated at 413 K for 2 d and then cooled to room temperature at a rate of 5 K h^{-1} . Red block-shaped crystals were obtained, and were washed with deionized water and absolute ethanol (yield > 20%, based on Co).

Crystal data

$(\text{C}_3\text{H}_5\text{N}_2)_2[\text{Co}(\text{C}_7\text{H}_3\text{NO}_4)_2 \cdot (\text{H}_2\text{O})_2] \cdot 4\text{H}_2\text{O}$	$Z = 1$
$M_r = 635.41$	$D_x = 1.559 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 8.844 (5) \text{ \AA}$	Cell parameters from 25 reflections
$b = 9.062 (4) \text{ \AA}$	$\theta = 2-7^\circ$
$c = 9.635 (4) \text{ \AA}$	$\mu = 0.71 \text{ mm}^{-1}$
$\alpha = 97.70 (3)^\circ$	$T = 293 (2) \text{ K}$
$\beta = 116.80 (3)^\circ$	Block, red
$\gamma = 93.26 (3)^\circ$	$0.49 \times 0.38 \times 0.29 \text{ mm}$
$V = 677.0 (6) \text{ \AA}^3$	

Data collection

Siemens R3m diffractometer
 ω scans
 Absorption correction: ψ scan
 (North *et al.*, 1968)
 $T_{\text{min}} = 0.721$, $T_{\text{max}} = 0.820$
 3153 measured reflections
 2959 independent reflections
 2532 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.149$
 $S = 1.05$
 2959 reflections
 205 parameters
 H atoms treated by a mixture of independent and constrained refinement

$R_{\text{int}} = 0.042$
 $\theta_{\text{max}} = 27.0^\circ$
 $h = 0 \rightarrow 11$
 $k = -11 \rightarrow 11$
 $l = -12 \rightarrow 10$
 2 standard reflections
 every 200 reflections
 intensity decay: none

$w = 1/[\sigma^2(F_o^2) + (0.0964P)^2 + 0.3143P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.47 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.51 \text{ e \AA}^{-3}$

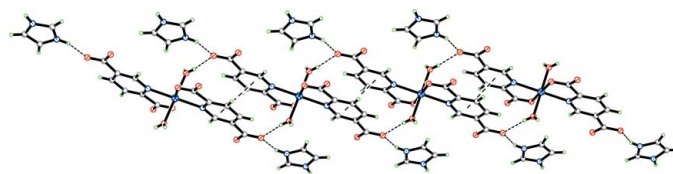


Figure 2
Perspective view of the chain along the a axis. Hydrogen bonds and π - π stacking interactions are shown as dashed lines.

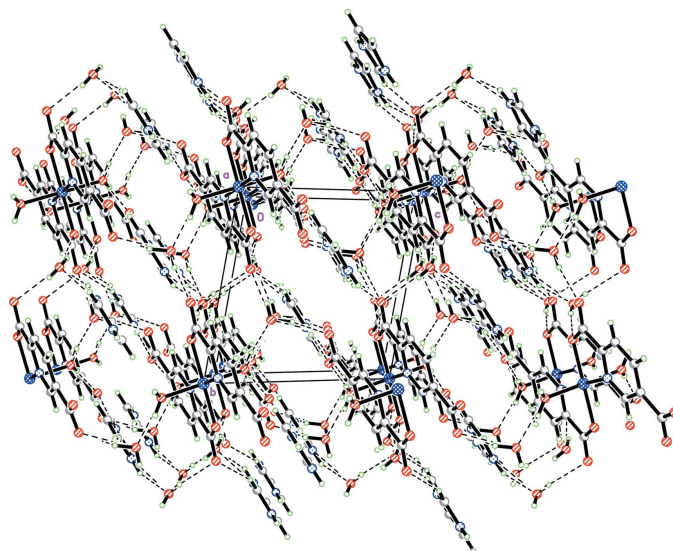


Figure 3
Packing diagram of (I). Dashed lines indicate hydrogen bonds.

Table 1

Selected geometric parameters (\AA , $^\circ$).

Co1—O1W	2.085 (2)	Co1—N1	2.124 (2)
Co1—O1	2.098 (2)		
O1W—Co1—O1 ⁱ	89.90 (9)	O1W—Co1—N1 ⁱ	91.07 (10)
O1W—Co1—O1	90.10 (9)	O1 ⁱ —Co1—N1	101.77 (9)
O1W ⁱ —Co1—N1	91.08 (10)	O1—Co1—N1	78.23 (9)
O1W—Co1—N1	88.92 (10)		

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

Table 2

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1W—H1A \cdots O3 ⁱⁱ	0.70 (5)	2.01 (5)	2.703 (3)	169 (5)
O1W—H1B \cdots O2W	0.86 (5)	1.85 (5)	2.688 (4)	166 (4)
O2W—H2C \cdots O4 ⁱⁱⁱ	0.97 (6)	1.78 (7)	2.731 (4)	165 (5)
O2W—H2D \cdots O2 ^{iv}	0.82 (7)	1.96 (7)	2.777 (4)	169 (6)
O3W—H3C \cdots O1 ^{iv}	0.93 (6)	1.94 (6)	2.855 (4)	167 (5)
O3W—H3D \cdots O2	0.75 (6)	2.14 (6)	2.882 (4)	168 (6)
N2—H2B \cdots O3 ^v	0.86	1.92	2.726 (4)	157
N3—H3A \cdots O3W	0.86	1.99	2.797 (4)	156

Symmetry codes: (ii) $x - 1, y, z$; (iii) $-x + 1, -y + 2, -z + 1$; (iv) $-x, -y + 1, -z$; (v) $-x + 2, -y + 2, -z + 1$.

H atoms attached to C or N atoms were positioned geometrically (C–H = 0.93 Å and N–H = 0.86 Å) and refined as riding on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C or N})$. O-bound H atoms were located in difference Fourier maps and refined with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

Data collection: *R3m Software* (Siemens, 1990); cell refinement: *R3m Software*; data reduction: *R3m Software*; program(s) used to solve structure: *SHELXTL* (Bruker, 1999); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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