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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.027$
$w R$ factor $=0.067$
Data-to-parameter ratio $=15.6$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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# catena-Poly[[diaquabis(1H-benzimidazole- $\kappa \mathrm{N}^{3}$ )-cobalt(II)]- $\mu$-succinato-кO] 

The crystal structure of the title compound $\left[\mathrm{Co}\left(\mathrm{C}_{4} \mathrm{H}_{4} \mathrm{O}_{4}\right)\right.$ $\left.\left(\mathrm{C}_{7} \mathrm{H}_{6} \mathrm{~N}_{2}\right)_{2}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right]_{n}$, consists of a polymeric $\mathrm{Co}^{\mathrm{II}}$ complex bridged by a succinate dianion. Neighboring polymeric chains linked via hydrogen bonds. The overlapped arrangement and short separation of 3.365 (3) $\AA$ between parallel benzimidazole rings suggest the existence of $\pi-\pi$ stacking. The centrosymmetric complex has an octahedral coordination geometry.

## Comment

$\pi-\pi$ stacking between aromatic rings has been shown to be correlated with the electron-transfer process in some biological systems (Deisenhofer \& Michel, 1989). As a part of an investigation of $\pi-\pi$ stacking in metal complexes (Chen et al., 2003), the title polymeric $\mathrm{Co}^{\mathrm{II}}$ complex, (I), has recently been prepared in our laboratory and its X-ray structure is presented here.


The title $\mathrm{Co}^{\text {II }}$ complex is isomorphous with the $\mathrm{Ni}^{\mathrm{II}}$ complex reported recently (Liu et al., 2003). The crystal structure consists of polymeric chains of the $\mathrm{Co}^{\mathrm{II}}$ complex. The octahedral coordination environment around the $\mathrm{Co}^{\mathrm{II}}$ atom, located on an inversion center, is illustrated in Fig. 1. The CoO (water) bond $[2.1404(11) \AA]$ is longer than the Co O (carboxyl) bond [2.0898 (11) Å] (Table 1). The carboxyl group of the succinate is monodentate; the uncoordinated carboxyl atom O 2 is hydrogen-bonded to the adjacent coordinated water and benzimidazole molecules, as shown in Fig. 1.

The succinate dianions bridge the $\mathrm{Co}^{\mathrm{II}}$ atoms through both terminal carboxyl groups to form one-dimensional polymeric chains (Fig. 2). Adjacent chains are linked via $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds between benzimidazole and carboxyl groups and $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds between coordinated water molecules and carboxyl groups (Table 2). Fig. 2 also shows the overlapped arrangement of parallel benzimidazole groups from adjacent polymeric chains. Neighboring benzimidazole rings related by $(-x, 1-y,-z)$ are 3.365 (3) A apart. These findings suggest the existence of $\pi-\pi$ stacking between adjacent polymeric chains.

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Figure 1
The coordination environment around the $\mathrm{Co}^{\mathrm{II}}$ atom in (I), with $50 \%$ probability displacement ellipsoids. Dashed lines indicate hydrogen bonding. [Symmetry codes: (i) $-x,-y, 1-z$; (ii) $1-x, 1-y, 1-z$; (iii) $x, y, 1+z$; (iv) $x-1, y, z]$.


Figure 2
The molecular packing diagram, showing the $\pi-\pi$ stacking between benzimidazole rings from neighboring polymeric chains. Dashed lines indicate hydrogen bonds.

## Experimental

$\mathrm{CoCl}_{2} \cdot 6 \mathrm{H}_{2} \mathrm{O}(0.48 \mathrm{~g}, 2 \mathrm{mmol})$ was added to an aqueous solution $(10 \mathrm{ml})$ containing succinic acid $(0.24 \mathrm{~g}, 2 \mathrm{mmol})$ and $\mathrm{NaOH}(0.16 \mathrm{~g}$, 4 mmol ). After the mixture was refluxed for 1 h , an ethanol solution $(10 \mathrm{ml})$ of benzimidazole $(0.24 \mathrm{~g}, 2 \mathrm{mmol})$ was added with continuous stirring. The solution was refluxed for 2 h , then was cooled to room temperature and filtered. Pink single crystals were obtained from the filtrate after 3 d .

## Crystal data

| $\left[\mathrm{Co}\left(\mathrm{C}_{4} \mathrm{H}_{4} \mathrm{O}_{4}\right)\left(\mathrm{C}_{7} \mathrm{H}_{6} \mathrm{~N}_{2}\right)_{2} \mathrm{H}_{2} \mathrm{O}_{2}\right]$ | $Z=1$ |
| :--- | :--- |
| $M_{r}=447.31$ | $D_{x}=1.628 \mathrm{Mg} \mathrm{m}^{-3}$ |
| Triclinic, $P \overline{1}$ | Mo $K \alpha$ radiation |
| $a=7.0717(9) \AA$ | Cell parameters from 2067 |
| $b=8.6326(11) \AA$ | reflections |
| $c=8.8223(12) \AA$ | $\theta=2.5-25.0^{\circ}$ |
| $\alpha=100.699(7)^{\circ}$ | $\mu=0.99 \mathrm{~mm}^{\circ}$ |
| $\beta=112.489(9)^{\circ}$ | $T=295(2) \mathrm{K}$ |
| $\gamma=104.780(6)^{\circ}$ | Plate, pink |
| $V=456.33(11) \AA^{\circ}$ | $0.28 \times 0.18 \times 0.06 \mathrm{~mm}$ |

## Data collection

Rigaku RAXIS-RAPID diffractometer

## $\omega$ scans

Absorption correction: multi-scan
(ABSCOR; Higashi, 1995)
$T_{\text {min }}=0.756, T_{\text {max }}=0.940$
4322 measured reflections

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.033 P)^{2}\right. \\
& \quad+0.2062 P] \\
& \text { where } P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.28 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.24 \mathrm{e}^{-3}
\end{aligned}
$$

Table 1
Selected geometric parameters $\left(\AA^{\circ},{ }^{\circ}\right)$.

| $\mathrm{Co}-\mathrm{O} 1$ | $2.0894(11)$ | $\mathrm{O} 2-\mathrm{C} 11$ | $1.2695(19)$ |
| :--- | ---: | :--- | :--- |
| $\mathrm{Co}-\mathrm{N} 3$ | $2.1309(13)$ | $\mathrm{C} 11-\mathrm{C} 12$ | $1.517(2)$ |
| $\mathrm{C}-\mathrm{O} 3$ | $2.1407(11)$ | $\mathrm{C} 12-\mathrm{C} 12^{\mathrm{i}}$ | $1.521(3)$ |
| $\mathrm{O} 1-\mathrm{C} 11$ | $1.2568(18)$ |  |  |
| $\mathrm{O} 1-\mathrm{Co}-\mathrm{N} 3$ | $91.30(5)$ | $\mathrm{N} 3-\mathrm{Co}-\mathrm{O} 3$ | $86.21(5)$ |
| $\mathrm{O} 1-\mathrm{Co}-\mathrm{O} 3$ | $90.50(4)$ |  |  |

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

Table 2
Hydrogen-bonding geometry $\left(\AA^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{O} 3-\mathrm{H} 32 \cdots \mathrm{O}^{\mathrm{ii}}$ | 0.87 | 1.85 | $2.7020(18)$ | 167 |
| $\mathrm{~N} 1-\mathrm{H} 1 \cdots \mathrm{O}^{\mathrm{v}}$ | 0.86 | 2.07 | $2.8775(19)$ | 157 |
| $\mathrm{O}^{\mathrm{O}}-\mathrm{H} 31 \cdots \mathrm{O}^{\text {vi }}$ | 0.89 | 2.00 | $2.860(2)$ | 160 |

Symmetry codes: (ii) $1-x, 1-y, 1-z$; (v) $x, y, z-1$; (vi) $1+x, y, z$.
H atoms of water molecules were located in a difference Fourier map and included in the structure-factor calculation, with fixed coordinates and isotropic displacement parameters of $0.05 \AA^{2}$. Other H atoms were placed in calculated positions, with $\mathrm{C}-\mathrm{H}=0.93 \AA$ (benzimidazole) or $0.97 \AA$ (succinate) and $\mathrm{N}-\mathrm{H}=0.86 \AA$, and included in the final cycles of refinement in a riding model, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}$ of the carrier atoms.

Data collection: PROCESS-AUTO (Rigaku Corporation, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/MSC and Rigaku Corporation, 2002); program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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