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## Hexaaquacobalt(II) bis(6-hydroxypyridine-3-carboxylate)

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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{o}-\mathrm{O})=0.004 \AA$
Disorder in main residue
$R$ factor $=0.054$
$w R$ factor $=0.159$
Data-to-parameter ratio $=8.6$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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In the title compound, $\left[\mathrm{Co}\left(\mathrm{H}_{2} \mathrm{O}\right)_{6}\right]\left(\mathrm{C}_{6} \mathrm{H}_{4} \mathrm{NO}_{3}\right)_{2}$, the $\mathrm{Co}^{\text {II }}$ atom lies on a special position of $2 / m$ site symmetry in an octahedron made up of water molecules. The anions show orientational disorder over mirror planes and are linked together by a pair of $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds into a dianion. The complex cations and dianions are connected through $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds to form a three-dimensional network.

## Comment

The title compound, (I), is isostructural with the $\mathrm{Zn}^{\mathrm{II}}$ (Zhang et al., 2005) and $\mathrm{Ni}^{\mathrm{II}}$ analogues (Zhang \& $\mathrm{Ng}, 2005$ ).

(I)

The crystal structure of (I) consists of octahedral cations and hydrogen-bonded dianions (Fig. 1). Atom Co1 lies on a position of $2 / m$ site symmetry and atom $\mathrm{O} 2 w$ also lies on the mirror plane, which bisects the $\mathrm{O} 1 w-\mathrm{Co} 1-\mathrm{O} 1 w^{\mathrm{iii}}$ bond angle


A plot of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the $50 \%$ probability level and H atoms are shown as small spheres of arbitrary radii. Dashed lines indicate hydrogen bonds. [Symmetry codes: (i) $1-x, y, 1-z$; (ii) $1-x, 1-y, 1-z$; (iii) $x, 1-y, z$; (iv) $\frac{5}{2}-x, \frac{3}{2}-y, 2-z$.]

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[Fig. 1; symmetry code: (iii) $x, 1-y, z$ ]. In the anion, a crystallographic mirror plane passes through atom C6 perpendicular to the carboxylate group. As a result, the hydroxypyridyl group of the anion shows orientational disorder.

The cations and dianions of (I) are linked by hydrogen bonds (Table 2) to form a three-dimensional network.

## Experimental

A mixture of cobalt(II) chloride hexahydrate $(0.238 \mathrm{~g}, 1 \mathrm{mmol})$, 6-hydroxypyridyl-3-carboxylic acid $(0.139 \mathrm{~g}, \quad 1 \mathrm{mmol})$, sodium hydroxide ( $0.040 \mathrm{~g}, 1 \mathrm{mmol}$ ) and water ( 10 ml ) were sealed in a 23 ml Teflon-lined stainless steel Parr bomb. The bomb was heated to 433 K for 2 d . It was then cooled to room temperature at $10 \mathrm{~K} \mathrm{~h}^{-1}$ to yield red crystals of (I).

## Crystal data

$\left[\mathrm{Co}\left(\mathrm{H}_{2} \mathrm{O}\right)_{6}\right]\left(\mathrm{C}_{6} \mathrm{H}_{4} \mathrm{NO}_{3}\right)_{2}$
$M_{r}=443.23$
Monoclinic, $C 2 / \mathrm{m}$
$a=11.609$ (1) $\AA$
$b=9.754$ (1) A
$c=7.6157(8) \AA$
$\beta=91.448$ (2) ${ }^{\circ}$
$V=862.1$ (2) $\AA^{3}$
$Z=2$
$D_{x}=1.707 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation

Cell parameters from 1422 reflections
$\theta=2.7-27.4^{\circ}$
$\mu=1.07 \mathrm{~mm}^{-1}$
$T=295$ (2) K
Plate, red
$0.34 \times 0.20 \times 0.08 \mathrm{~mm}$
Data collection
Bruker APEX area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\text {min }}=0.499, T_{\text {max }}=0.920$
2296 measured reflections
968 independent reflections 956 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.020$
$\theta_{\text {max }}=27.4^{\circ}$
$h=-14 \rightarrow 15$
$k=-12 \rightarrow 12$
$l=-9 \rightarrow 8$

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /[ \sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.053 P)^{2} \\
&+7.1943 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.54 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.50 \mathrm{e}^{-3}
\end{aligned}
$$

113 reflections
H atoms treated by a mixture of independent and constrained refinement

Table 2
Hydrogen-bond geometry ( $\mathrm{A},{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{O} 1 w-\mathrm{H} 1 w 1 \cdots \mathrm{O} 1$ | 0.85 (1) | 1.83 (2) | 2.659 (4) | 168 (4) |
| $\mathrm{O} 1 w-\mathrm{H} 1 w 2 \cdots \mathrm{O} 1^{\text {i }}$ | 0.84 (1) | 1.87 (1) | 2.710 (5) | 171 (5) |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 1 \cdots \mathrm{O} 2^{\text {ii }}$ | 0.85 (1) | 2.02 (3) | 2.791 (8) | 151 (4) |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 1 \cdots \mathrm{O} 2^{\text {ii }}$ | 0.85 (1) | 2.02 (3) | 2.791 (8) | 151 (4) |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 2 \cdots \mathrm{O} 2^{\text {iii }}$ | 0.85 (1) | 1.97 (1) | 2.769 (9) | 157 (2) |
| $\mathrm{O} 2 w-\mathrm{H} 2 w 2 \cdots \mathrm{O} 2{ }^{\text {iv }}$ | 0.85 (1) | 1.97 (1) | 2.769 (9) | 157 (2) |
| $\mathrm{O} 2-\mathrm{H} 2 \mathrm{o} \cdots{ }^{\text {c }}{ }^{\text {v }}$ | 0.85 | 2.04 | 2.86 (1) | 159 |

Symmetry codes: (i) $\frac{3}{2}-x, \frac{3}{2}-y, 1-z$; (ii) $x-1, y, z-1$; (iii) $2-x, 1-y, 1-z$; (iv) $2-x, y, 1-z$; (v) $\frac{5}{2}-x, \frac{3}{2}-y, 2-z$.

Atoms $\mathrm{C} 1-\mathrm{C} 5 / \mathrm{N} 1 / \mathrm{O} 2$ in the anion are disordered over two possible positions related by mirror symmetry; the $\mathrm{C}-\mathrm{C}$ distances were restrained to 1.39 (1) $\AA$, and the two $\mathrm{N}-\mathrm{C}$ distances were restrained to within $0.01 \AA$ of each other. Additionally, the ring was restrained to near planarity. Water H atoms were located in difference Fourier maps and were refined with a distance restraint of $\mathrm{O}-$ $\mathrm{H}=0.85$ (1) $\AA$. OH groups were allowed to rotate about the $\mathrm{C}-\mathrm{O}$ bond to fit the electron density, with $\mathrm{O}-\mathrm{H}$ constrained to $0.85 \AA$ and $\mathrm{C}-\mathrm{O}-\mathrm{H}=109.5^{\circ}$. Carbon-bound H atoms were placed in calculated positions $(\mathrm{C}-\mathrm{H}=0.93 \AA)$ and included in the refinement in the riding-model approximation, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: atomic coordinates taken from the isostructural Zn analogue (Zhang et al., 2005); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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