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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
Disorder in main residue
$R$ factor $=0.029$
$w R$ factor $=0.080$
Data-to-parameter ratio $=13.1$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Hexaaquacobalt(II) bis(4-formylbenzoate) dihydrate

The crystal structure of the title compound, $\left[\mathrm{Co}\left(\mathrm{H}_{2} \mathrm{O}\right)_{6}\right]$ $\left(\mathrm{C}_{8} \mathrm{H}_{5} \mathrm{O}_{3}\right)_{2} \cdot 2 \mathrm{H}_{2} \mathrm{O}$, consists of octahedral $\mathrm{Co}^{\text {II }}$ complex cations that interact with the anions and uncoordinated water molecules, forming a tightly held, hydrogen-bonded layer. The $\mathrm{Co}^{\text {II }}$ complex cation lies on a special position of site symmetry $\overline{1}$.

## Comment

The known transition metal derivatives of 4-formylbenzoic acid are water-coordinated compounds showing covalent metal-carboxylate bonds, for example the nickel(II) (Deng et al., 2006a) and copper(II) (Deng et al., 2006b) derivatives. However, the cobalt(II) derivative does not have this feature; the title compound (I) has a hexaaquacobalt cation (Table 1), and its charge is balanced by two uncoordinated 4 -formylbenzoate ions. There are one cation and two anions in the asymmetric unit along with two uncoordinated water molecules (Fig. 1). These are linked by hydrogen bonds (Table 2) into a layer structure. The $\mathrm{Co}^{\mathrm{II}}$ complex cation lies on a special position of site symmetry $\overline{1}$.


## Experimental

Cobalt diacetate trihydrate $(0.116 \mathrm{~g}, 0.5 \mathrm{mmol})$ was added to an aqueous solution ( 20 ml ) of 4 -formylbenzoic acid ( $0.15 \mathrm{~g}, 1 \mathrm{mmol}$ ). Sodium hydroxide solution was added to obtain a pH 5 . The filtered solution was allowed to stand for a few days for the pink prismatic crystals to separate. Elemental analysis calculated for $\mathrm{C}_{16} \mathrm{H}_{26} \mathrm{CoO}_{14}$ : C 38.33 , H 5.23\%; found: C 38.31 , H $5.24 \%$.

## Crystal data

| $\left[\mathrm{Co}\left(\mathrm{H}_{2} \mathrm{O}\right)_{6}\right]\left(\mathrm{C}_{8} \mathrm{H}_{5} \mathrm{O}_{3}\right)_{2} \cdot 2 \mathrm{H}_{2} \mathrm{O}$ | $V=538.20(5) \AA^{3}$ |
| :--- | :--- |
| $M_{r}=501.30$ | $Z=1$ |
| Triclinic, $P \overline{1}$ | $D_{x}=1.547 \mathrm{Mg} \mathrm{m}^{-3}$ |
| $a=6.3470(3) \AA$ | Mo $K \alpha$ radiation |
| $b=7.4194(4) \AA$ | $\mu=0.87 \mathrm{~mm}^{-1}$ |
| $c=12.5041(7) \AA$ | $T=295(2) \mathrm{K}$ |
| $\alpha=92.353(2)^{\circ}$ | Prism, pink |
| $\beta=90.029(3)^{\circ}$ | $0.36 \times 0.28 \times 0.18 \mathrm{~mm}$ |
| $\gamma=113.801(6)^{\circ}$ |  |

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## Data collection

Rigaku RAXIS-RAPID IP
diffractometer
$\omega$ scans
Absorption correction: multi-scan
$\quad A B S C O R$ (Higashi, 1995)
$T_{\min }=0.614, T_{\max }=0.860$

5333 measured reflections
2451 independent reflections 2325 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.017$
$\theta_{\text {max }}=27.5^{\circ}$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.029$

$$
S=1.00
$$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0485 P)^{2}\right. \\
& \quad+0.2223 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.55 \text { e } \AA^{-3}
\end{aligned}
$$

2451 reflections
187 parameters
H atoms treated by a mixture of independent and constrained refinement


Figure 1
Displacement ellipsoid plot of the unit-cell contents. Displacement ellipsoids are drawn at the $70 \%$ probability level, and H atoms are shown as spheres of arbitrary radii. Only one disorder component is shown. [Symmetry code: (i) $1-x, 1-y, 1-z$.]

Carbon-bound H atoms were positioned geometrically $(\mathrm{C}-\mathrm{H}$ $0.93 \AA$ ) and were included in the refinement in the riding-model approximation, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$. The water H atoms were located in a difference Fourier map, and were refined with a distance restraint of $\mathrm{O}-\mathrm{H} 0.85$ (1) $\AA$; their displacement parameters were freely refined.

Data collection: RAPID-AUTO (Rigaku, 1998); cell refinement: RAPID-AUTO; data reduction: CrystalStructure (Rigaku/MSC, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: X-SEED (Barbour, 2001); software used to prepare material for publication: publCIF (Westrip, 2006).

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