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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.032$
$w R$ factor $=0.084$
Data-to-parameter ratio $=17.2$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Redetermination of diaquabis(vanillinato- $\kappa^{2} O, O^{\prime}$ )cobalt(II)

The structure of the title complex, $\left[\mathrm{Co}\left(\mathrm{C}_{8} \mathrm{H}_{7} \mathrm{O}_{3}\right)_{2}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right]$, has been redetermined in the space group $C 2 / c$. The original report [Xiao, Lan, Zhang \& Jiang (2002). Guangxi Shifan Daxue Xuebao Ziran Kexueban (J. Guangxi Normal Univ.), 20, 81-83] declared the space group to be $C c$. The molecule lies on a twofold axis; the vanillinate anion chelates to the $\mathrm{Co}^{\mathrm{II}}$ atom through the methoxy and hydroxy groups. The Co$\mathrm{O}_{\text {methoxy }}$ bond is longer than the $\mathrm{Co}-\mathrm{O}_{\text {hydroxy }}$ bond by 0.2673 (18) Å.

## Comment

The title complex, (I), was previously refined in the lower symmetry space group Cc (Xiao et al., 2002). A PLATON check (Spek, 2003) suggests additional symmetry, as shown in this report.

(I)

The molecular structure of (I) is illustrated in Fig. 1. In C2/c, the molecule has a twofold axis of symmetry on which the $\mathrm{Co}^{\mathrm{II}}$ atom lies. The twofold symmetry element relates one vanillinate ligand and one coordinated water molecule to the other. The vanillinate anion chelates to the $\mathrm{Co}^{\mathrm{II}}$ atom through the methoxy and hydroxy groups, the $\mathrm{Co}-\mathrm{O}_{\text {methoxy }}$ bond being longer than the $\mathrm{Co}-\mathrm{O}_{\text {hydroxy }}$ bond by 0.2673 (18) $\AA$. Two water molecules coordinate in a cis manner to the $\mathrm{Co}^{\mathrm{II}}$ atom to complete the distorted octahedral coordination geometry (Table 1). The vanillinate ligand is planar, the maximum deviation being 0.0702 (14) $\AA$ (O1 atom); the Co atom is out-of-plane by 0.6368 (14) $\AA$.

## Experimental

An ethanol solution ( 5 ml ) of vanillin ( 2 mmol ) and $\mathrm{Na}_{2} \mathrm{CO}_{3}$ ( 1 mmol ) was mixed with an aqueous solution $(5 \mathrm{ml})$ of $\mathrm{CoCl}_{2} \cdot 6 \mathrm{H}_{2} \mathrm{O}$ ( 1 mmol ), and the mixture was refluxed for 1 h . After cooling to room

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temperature, the solution was filtered. Red crystals of (I) were obtained after 3 d .

## Crystal data

$\left[\mathrm{Co}\left(\mathrm{C}_{8} \mathrm{H}_{7} \mathrm{O}_{3}\right)_{2}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right]$
$M_{r}=397.23$
Monoclinic, $C 2 / c$
$a=22.270(2) \AA$
$b=10.4487(12) \AA$
$c=7.7771(9) \AA$
$\beta=107.249(12)^{\circ}$
$V=1728.3(3) \AA^{3}$
$Z=4$
$D_{x}=1.527 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 6807 reflections
$\theta=3.2-26.0^{\circ}$
$\mu=1.03 \mathrm{~mm}^{-1}$
$T=295$ (2) K
Prism, red
$0.40 \times 0.38 \times 0.30 \mathrm{~mm}$

## Data collection

Rigaku R-AXIS RAPID
diffractometer

## $\omega$ scans

Absorption correction: multi-scan
(ABSCOR; Higashi, 1995)
$T_{\text {min }}=0.672, T_{\text {max }}=0.738$
7677 measured reflections

## Refinement

Refinement on $F^{2}$
1973 independent reflections
1798 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.024$
$\theta_{\text {max }}=27.5^{\circ}$
$h=-27 \rightarrow 28$
$k=-13 \rightarrow 13$
$l=-10 \rightarrow 10$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.032$
$w R\left(F^{2}\right)=0.084$
$S=1.09$
1973 reflections
115 parameters
H -atom parameters constrained

$$
\begin{aligned}
& w= 1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0446 P)^{2}\right. \\
&+1.1767 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.29 \mathrm{e}^{2} \AA^{-3} \\
& \Delta \rho_{\min }=-0.24 \mathrm{e}^{-3}
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\AA \mathrm{A}^{\circ}$ ).

| $\mathrm{Co}-\mathrm{O} 2$ | $2.0415(13)$ | $\mathrm{O} 3-\mathrm{C} 3$ | $1.382(2)$ |
| :--- | ---: | :--- | ---: |
| $\mathrm{Co}-\mathrm{O} 3$ | $2.2631(13)$ | $\mathrm{O} 3-\mathrm{C} 8$ | $1.408(3)$ |
| $\mathrm{Co}-\mathrm{O} 4$ | $1.9958(12)$ | $\mathrm{O} 4-\mathrm{C} 4$ | $1.311(2)$ |
| $\mathrm{O} 1-\mathrm{C} 7$ | $1.222(3)$ |  |  |
| $\mathrm{O} 2-\mathrm{Co}-\mathrm{O} 2^{\mathrm{i}}$ | $92.02(8)$ | $\mathrm{O} 2-\mathrm{Co}-\mathrm{O} 4$ | $97.28(5)$ |
| $\mathrm{O} 2-\mathrm{Co}-\mathrm{O} 3$ | $172.33(5)$ | $\mathrm{O} 3-\mathrm{Co}-\mathrm{O} 3^{\mathrm{i}}$ | $92.27(8)$ |
| $\mathrm{O} 2-\mathrm{Co}-\mathrm{O} 3^{\mathrm{i}}$ | $88.37(6)$ | $\mathrm{O} 4-\mathrm{Co}-\mathrm{O} 4^{\mathrm{i}}$ | $158.32(8)$ |

Symmetry code: (i) $-x+1, y,-z+\frac{1}{2}$.

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| O2-H2A $\cdots \mathrm{O} 1^{\mathrm{ii}}$ | 0.85 | 1.88 | $2.733(2)$ | 173 |
| $\mathrm{O}^{\mathrm{iii}}-\mathrm{H} 2 B \cdots \mathrm{O} 4^{2}$ | 0.88 | 1.85 | $2.7158(19)$ | 165 |

Symmetry codes: (ii) $x-\frac{1}{2},-y+\frac{1}{2}, z-\frac{1}{2}$; (iii) $-x+1,-y,-z+1$.
H atoms on the water molecule were located in a difference Fourier map and refined as riding in their as-found positions relative to the O atom, with $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{O})$. Methyl H atoms were placed in calculated positions and refined with $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$. Other H atoms were placed in calculated positions with $\mathrm{C}-\mathrm{H}=$


Figure 1
The molecular structure of (I), showing $30 \%$ probability displacement ellipsoids (arbitrary spheres for H atoms). [Symmetry code: (i) $1-x, y$, $\frac{1}{2}-z$.]
$0.93 \AA$ and included in the final cycles of refinement in the riding model, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}$ of the carrier atoms.

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/ MSC, 2002); method used to solve structure: averaging the coordinates of the published $C c$ structure (Xiao et al., 2002); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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