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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.029 wR factor = 0.075 Data-to-parameter ratio = 17.5

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

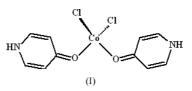
Dichlorobis(4-pyridone-κO)cobalt(II)

The Co atom in the title complex, $[CoCl_2(C_5H_5NO)_2]$, has a distorted tetrahedral coordination involving two Cl⁻ ions and two O atoms of two 4-pyridone ligands. The Co atom occupies a special position with twofold rotation symmetry. A layer structure is formed by N-H···Cl intermolecular hydrogen bonds and π - π stacking interactions of adjacent pyridone rings.

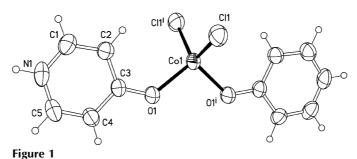
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Comment

4-Hydroxypyridine has been widely used in pharmaceutical synthesis. It usually exists in equilibrium with the form 4-pyridone. However, in contrast to the metal complexes of 2-hydroxypyridine (Goodgame *et al.*, 1989; Blake *et al.*, 1991), there are few reports of structures of complexes of 4-hydroxypyridine or 4-pyridone (LaliaKantouri, 1996). Recently, we synthesized a cobalt complex of 4-pyridone, *viz.* dichlorobis(4-pyridone)cobalt(II), (I) (Fig. 1), by the reaction of 4-hydroxypyridine and CoCl₂·6H₂O in an ethanol solution. Compound (I) is isostructural with [ZnCl₂(C₅H₅NO)₂] (Masse & Le Fur, 1998).



The Co atom in (I) exists in a tetrahedral coordination environment, defined by two Cl⁻ anions and two O atoms of 4pyridone [Co-Cl = 2.2763 (8) Å and Co-O = 1.9550 (12) Å], and the Co atom occupies a special position with twofold rotation symmetry. The C1-C2, C4-C5 and C3-O1 bond lengths are 1.362 (2), 1.359 (3) and 1.2877 (19) Å, respectively. Interestingly, four N-H···Cl intermolecular hydrogen bonds generate a four-membered ring from four adjacent molecules. The four-membered ring has hydrogen-bonding distances (N···Cl) and angles (N-H···Cl) in the ranges 3.3481 (19)-



View of (I) with 50% probability ellipsoids. [Symmetry code: (i) 1 - x, y, $\frac{1}{2} - z$.]

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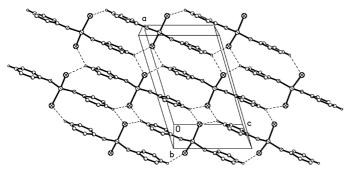


Figure 2

Packing diagram of (I), with hydrogen bonds indicated by dashed lines.

3.3574(18) Å and $127(2)-135(2)^{\circ}$, respectively (Table 1). A two-dimensional layer structure is formed by these hydrogen bonds, and the Co $\cdot \cdot \cdot$ Co distances are 6.7933 (15) and 8.901 (2) Å. Furthermore, the layer architecture is stabilized by $\pi - \pi$ interactions between adjacent pyridone ligands, with $Cg \cdots Cg$ (Cg is the centroid of the N1-containing ring) distances of 3.4429 (13) and 3.5372 (13) Å (Fig. 2).

Experimental

Compound (I) was prepared by the addition of CoCl₂·6H₂O (0.12 g, 0.5 mmol) to an ethanol solution of 4-hydroxypyridine (0.14 g, 1.5 mmol). The mixed solution was allowed to evaporate at room temperature and blue prism-shaped crystals were obtained after three days. Analysis calculated for $C_{10}H_{10}Cl_2CoN_2O_2$: C 37.53, H 3.15, N 8.75%; found: C 37.42, H 3.19, N 8.82%.

Crystal data

 $T_{\min} = 0.551, T_{\max} = 0.700$

5774 measured reflections

$\begin{bmatrix} \text{CoCl}_2(\text{C}_5\text{H}_5\text{NO})_2 \end{bmatrix}$	$D_x = 1.717 \text{ Mg m}^{-3}$
$M_r = 320.03$	Mo K\alpha radiation
Monoclinic, $C2/c$	Cell parameters from 2929
a = 13.185 (3) Å	reflections
b = 11.229 (2) Å	$\theta = 3.1-27.5^{\circ}$
c = 8.7300 (17) Å	$\mu = 1.81 \text{ mm}^{-1}$
$\beta = 106.70$ (3)°	T = 293 (2) K
V = 1238.0 (5) Å ³	Prism, blue
Z = 4	$0.36 \times 0.24 \times 0.20 \text{ mm}$
Data collection	
Rigaku R-AXIS RAPID	1420 independent reflections
diffractometer	1323 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.023$
Absorption correction: multi-scan	$\theta_{\text{max}} = 27.5^{\circ}$
(ABSCOR; Higashi, 1995)	$h = -17 \rightarrow 14$

 $h = -17 \rightarrow 14$ $k = -14 \rightarrow 14$ $-11 \rightarrow 11$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0453P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.029$	+ 0.7656P]
$wR(F^2) = 0.075$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} < 0.001$
1420 reflections	$\Delta \rho_{\rm max} = 0.39 \ {\rm e} \ {\rm \AA}^{-3}$
81 parameters	$\Delta \rho_{\rm min} = -0.25 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Hydrogen-bonding	geometry (Å, °).
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$\overline{D-\mathrm{H}\cdots A}$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} N1 - H6 \cdots Cl1^{ii} \\ N1 - H6 \cdots Cl1^{iii} \end{array}$	0.90 (3)	2.67 (2)	3.3574 (18)	135 (2)
	0.90 (3)	2.74 (2)	3.3481 (19)	127 (2)

Symmetry codes: (ii) x, y, 1 + z; (iii) $\frac{1}{2} - x, \frac{1}{2} - y, 1 - z$.

The H atom on the N atom was freely refined. The H atoms on C atoms were placed in calculated positions and treated as riding [C-H = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(\text{carrier atom})]$.

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: SHELXL97; software used to prepare material for publication: SHELXL97.

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