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Key indicators

Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.043
 wR factor = 0.124
Data-to-parameter ratio = 13.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Diaquadiisothiocyanatobis(pyridine *N*-oxide)-cobalt(II)

In the title mononuclear complex, $[\text{Co}(\text{NCS})_2(\text{C}_5\text{H}_5\text{NO})_2(\text{H}_2\text{O})_2]$, cobalt(II) has a distorted octahedral coordination formed by two N atoms from two thiocyanate anions and four O atoms from two water molecules and two pyridine *N*-oxide molecules. The complexes are connected to each other by $\text{O}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{S}$ hydrogen bonds.

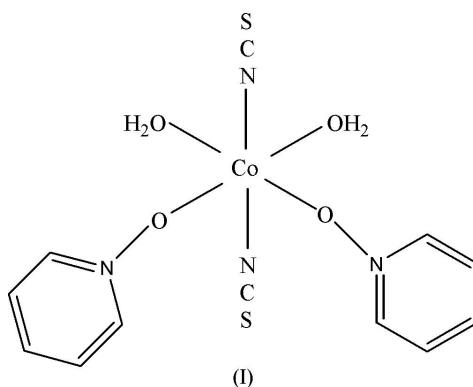
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Comment

Fig. 1 shows the title complex, (I), in which atom Co1 is coordinated by two N atoms from thiocyanate anions and four O atoms from two water molecules and two pyridine *N*-oxide molecules. As indicated in Fig. 1, atom Co1 has a distorted octahedral coordination. Various hydrogen bonds (Table 1 and Fig. 2) help to stabilize the crystal packing. The coordinated water molecule O2 forms two $\text{O}-\text{H}\cdots\text{O}$ bonds to *N*-oxide O-atom acceptors. The coordinated water molecule O1 forms $\text{O}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{S}$ bonds, the acceptor O atom being the O atom of the pyridine *N*-oxide.



Experimental

Pyridine *N*-oxide (0.0639 g, 0.672 mmol) was added to an aqueous solution (12 ml) containing $\text{Co}(\text{ClO}_4)_2 \cdot 6\text{H}_2\text{O}$ (0.1253 g, 0.342 mmol) and sodium thiocyanate (0.0562 g, 0.693 mmol), and the solution was stirred for a few minutes. Pink single crystals of (I) were obtained after the solution had been allowed to stand at room temperature for three weeks.

Crystal data

 $[\text{Co}(\text{NCS})_2(\text{C}_5\text{H}_5\text{NO})_2(\text{H}_2\text{O})_2]$ $M_r = 401.32$ Triclinic, $P\bar{1}$ $a = 5.4773$ (15) Å $b = 11.326$ (3) Å $c = 13.726$ (4) Å $\alpha = 86.366$ (4)° $\beta = 78.985$ (4)° $\gamma = 88.921$ (4)° $V = 834.1$ (4) Å³ $Z = 2$ $D_x = 1.598$ Mg m⁻³Mo $K\alpha$ radiation

Cell parameters from 1753

reflections

 $\theta = 2.4$ – 26.3 ° $\mu = 1.30$ mm⁻¹ $T = 298$ (2) K

Prism, pink

 $0.20 \times 0.09 \times 0.08$ mm

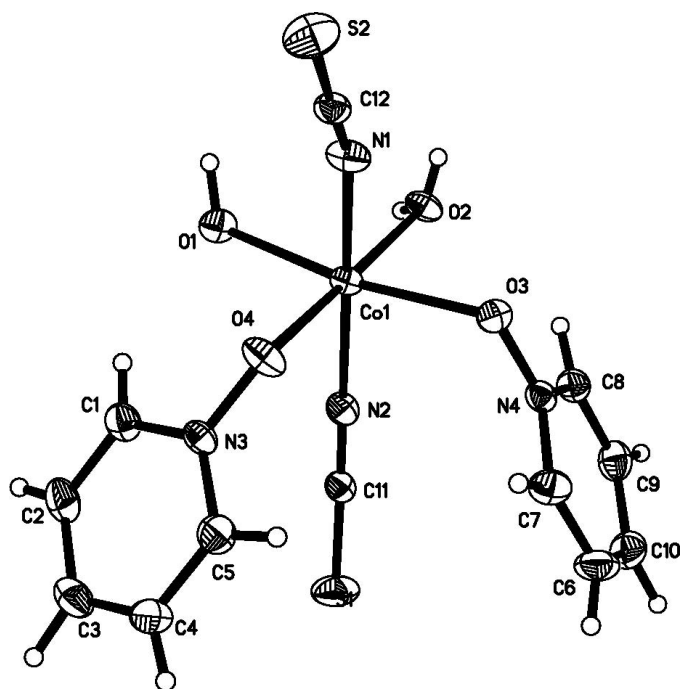


Figure 1
View of (I), showing 30% displacement ellipsoids. H atoms are shown as spheres of arbitrary radius.

Data collection

Bruker SMART CCD area-detector diffractometer	2892 independent reflections
φ and ω scans	2582 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{int} = 0.022$
$T_{min} = 0.781, T_{max} = 0.903$	$\theta_{max} = 25.0^\circ$
4291 measured reflections	$h = -6 \rightarrow 6$
	$k = -11 \rightarrow 13$
	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.046P)^2 + 1.1096P]$
$R[F^2 > 2\sigma(F^2)] = 0.043$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.124$	$(\Delta/\sigma)_{max} = 0.001$
$S = 1.14$	$\Delta\rho_{max} = 0.35 \text{ e } \text{\AA}^{-3}$
2892 reflections	$\Delta\rho_{min} = -0.31 \text{ e } \text{\AA}^{-3}$
212 parameters	
H-atom parameters constrained	

Table 1
Hydrogen-bonding geometry ($\text{\AA}, ^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O1-H11 \cdots S2^i$	0.87	2.59	3.433 (3)	164
$O1-H12 \cdots O4^i$	0.89	1.87	2.753 (4)	175
$O2-H14 \cdots O3^i$	0.99	1.86	2.833 (4)	165
$O2-H13 \cdots O3^{ii}$	0.71	2.23	2.901 (4)	157

Symmetry codes: (i) $1+x, y, z$; (ii) $1-x, 1-y, 1-z$.

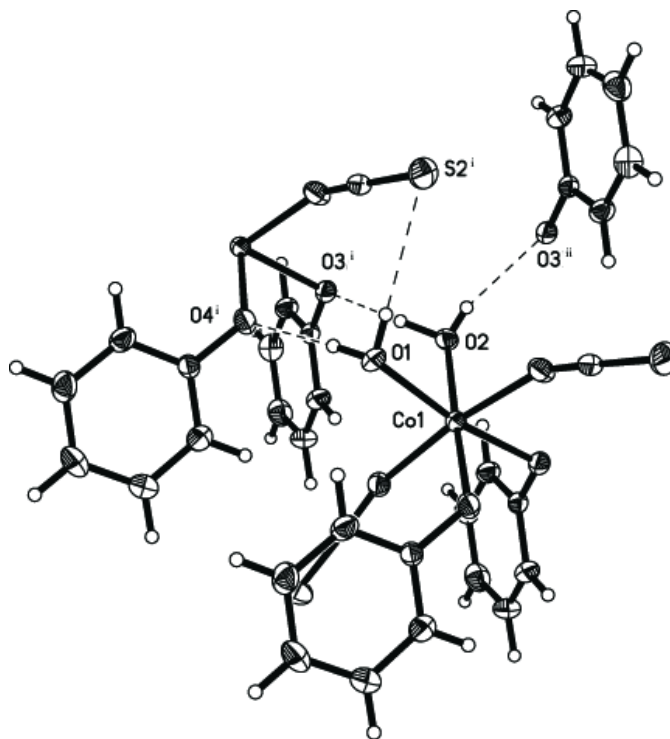


Figure 2
Hydrogen bonds (shown as broken lines) between the complex molecules. Symmetry codes are as in Table 1.

H atoms bonded to C atoms were included in calculated positions. Other H atoms were located in a difference map. All H atoms were refined using a riding model [$C-H = 0.93 \text{ \AA}$ and $U_{iso}(H) = 1.2U_{eq}(C)$; $O-H = 0.71-0.99 \text{ \AA}$ and $U_{iso}(H) = 1.2U_{eq}(O)$].

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 2001; software used to prepare material for publication: SHELXTL).

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