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Key indicators

Single-crystal X-ray study T = 298 K Mean σ (C–C) = 0.004 Å R factor = 0.032 wR factor = 0.083 Data-to-parameter ratio = 12.0

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

Tetraaquadithiocyanatocobalt(II) bis(2-methylpyrazine 1,4-dioxide)

In the crystal structure of the title compound, $[Co(NCS)_2(H_2O)_4]\cdot 2C_5H_6N_2O_2$, the six-coordinated Co^{II} complex lies on a special position of site symmetry 2/m. The methylpyrazine dioxide lies on a mirror plane and links with the Co^{II} complex *via* O-H···O hydrogen bonding. Received 6 June 2005 Accepted 4 July 2005 Online 9 July 2005

Comment

The thiocyanate anion and pyrazine 1,4-dioxide are common bridging ligands for preparing polynuclear complexes (Shi *et al.*, 2005; Sun *et al.*, 2001). In order to understand the relationship between reaction conditions and the products, the title complex, (I), was synthesized and its crystal structure is reported here.



The molecular structure of (I) is shown in Fig. 1. The Co^{II} atom lies on a special position of site symmetry 2/m. Four water O and two thiocyanate N atoms coordinate to the Co^{II} atom in an octahedral geometry (Table 1). The 2-methyl-



Figure 1

The molecular structure of (I) shown with 30% probability displacement ellipsoids (arbitary spheres for H atoms). [Symmetry codes: (i) -x + 1, -y, -z + 2; (ii) -x + 1, y, -z + 2; (iii) x, -y, z.]

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metal-organic papers

pyrazine 1,4-dioxide, except for two methyl H atoms, lies on a mirror plane and links with the Co^{II} complex via $O-H \cdots O$ hydrogen bonding (Table 2). The structure is similar to the Mn^{II} analogue (Xu et al., 2005).

Experimental

2-Methylpyrazine 1,4-dioxide (0.0436 g, 0.35 mmol) was added to an aqueous solution (15 ml) containing $Co(ClO_4)_2 \cdot 6H_2O$ (0.124 g, 0.34 mmol) and sodium thiocyanate (0.057 g, 0.70 mmol). The mixture was stirred for a few minutes and allowed to stand at room temperature. Pink single crystals of (I) were obtained after three weeks.

Crystal data

[Co(NCS)2(H2O)4]-2C2H6N2O2 $M_r = 499.39$ Monoclinic, C2/m a = 16.982 (4) Å b = 6.7616 (14) Å c = 10.120 (2) Å $\beta = 111.718 (4)^{\circ}$ V = 1079.5 (4) Å³ Z = 2

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\min} = 0.815, T_{\max} = 0.898$ 2873 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.032$ $wR(F^2) = 0.083$ S = 1.121095 reflections 91 parameters H-atom parameters constrained Cell parameters from 2566 reflections $\theta = 2.6 - 28.1^{\circ}$ $\mu = 1.04~\mathrm{mm}^{-1}$ T = 298 (2) K Prism, pink $0.20\,\times\,0.16\,\times\,0.11$ mm

 $D_x = 1.536 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

1095 independent reflections	
1045 reflections with $I > 2\sigma(I)$)
$R_{\rm int} = 0.048$	
$\theta_{\rm max} = 25.5^{\circ}$	
$h = -17 \rightarrow 20$	
$k = -8 \rightarrow 8$	
$l = -12 \rightarrow 10$	

$w = 1/[\sigma^2(F_0^2) + (0.0418P)^2]$
+ 0.4739P]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} < 0.001$
$\Delta \rho_{\rm max} = 0.40 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -0.38 \text{ e } \text{\AA}^{-3}$
Extinction correction: SHELXL97
Extinction coefficient: 0.071 (4)

Table 1

Selected geometric parameters (Å, °).

Co1-N1	2.094 (2)	Co1-O3	2.1102 (12)
N1-Co1-O3	86.46 (6)		

Table 2	
Hydrogen-bond geometry (Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$O3-H2\cdots O1^{i}$	0.85	1.94	2.746 (2)	159
$O3-H1\cdots O2^{ii}$	0.86	1.87	2.7175 (19)	166

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

Pyrazine H atoms were placed in calculated positions. Other H atoms were located in a difference Fourier map. All H atoms were refined using a riding model, with C-H = 0.91-0.96 Å, O-H = 0.85-0.86 Å and $U_{iso}(H) = 1.2U_{eq}(carrier)$.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); 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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