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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.009 Å Disorder in main residue R factor = 0.064 wR factor = 0.207 Data-to-parameter ratio = 23.3

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title compound, $[Co_2(CH_3O)_2(C_4H_5N_3)_4](BF_4)_2$, consists of a discrete dimeric Co^{II} complex. The asymmetric unit consists of one pyrimidin-2-amine ligand, which occupies a general position, and one methanolate group, as well as one Co^{II} cation and one tetrafluoroborate anion, which are located in special positions. The Co^{II} cation lies on a twofold rotation axis, whereas the tetrafluoroborate anion and the methanolate group lie on mirror planes. The N atoms of the pyrimidin-2amine ligand are involved in intermolecular hydrogen bonding. From this arrangement, large channels are formed, which elongate in the direction of the crystallographic *c* axis.

 κN^{1})cobalt(II)] bis(tetrafluoroborate)

Di-µ2-methanolato-bis[tetrakis(pyrimidin-2-amine-

Comment

Cobalt complexes are of great interest in coordination chemistry in relation to catalysis and enzymatic reactions, magnetism, and molecular architectures (Billson *et al.*, 2000; Fritsky *et al.*, 2003; Kotera *et al.*, 2003). Until recently, only a few cobalt complexes with pyrimidin-2-amine as ligand have been reported. In all of these compounds, the Co atom is coordinated by the N atom of the pyrimidine ring. Most of these compounds represent discrete mononuclear complexes (Kistenmacher *et al.*, 1978; Masaki *et al.*, 2002; Song *et al.*, 2005) but in some of them the Co atoms are connected into polymers, $[C_{16}H_{10}CO_2N_{18}]_n$ (Jensen *et al.*, 2000). To date, there is one dinuclear compound, $[C_8H_{22}Co_2N_6O_{14}S_2\cdot 2H_2O]$, in which the two Co atoms are connected by two sulfate anions (Lumme *et al.*, 1996). We report here the structure of another dinuclear Co compound with pyrimidin-2-amine, (I).



The structure of compound (I), is presented in Fig. 1. The Co atom lies on a twofold rotation axis, whereas the tetra-fluoroborate anion and the methanolate group lie on mirror

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Figure 1

Crystal structure of (I), with displacement ellipsoids drawn at the 30% probability level. Only one disorder component is shown. [Symmetry codes: (i) -x, y, -z; (ii) -x, -y, -z; (iii) x, -y, z.]



Figure 2 Packing diagram of (I), viewed along the c axis (hydrogen bonding is shown as dashed lines).

planes. The cobalt centres are four-coordinated by two O atoms of the methanolate group and two N atoms of two symmetry-related pyrimidin-2-amine ligands. The Co-O bond lengths of 1.920 (2) Å and the Co–N bond lengths of 2.014 (3) Å are comparable to those of similar Co complexes. The *cis*-angles at the Co^{II} centre range from 79.86 (17) to 95.33 (14)°, which indicates a slightly distorted square geometry of Co atoms. Two Co atoms are connected by two methanolate groups via μ -O,O' coordination into dimers, which are located on site 2/m. The tetrafluoroborate anions are located 3.4471 Å above the plane formed by the two Co and the two O atoms. The complexes are connected via N- $H \cdots N$ and $N - H \cdots F$ hydrogen bonding. From this arrangement, large channels are formed, which elongate in the direction of the c axis (Fig. 2). The channel volume was calculated using PLATON (Spek, 2003) to be 41.8% of the unit-cell volume.

Experimental

All chemicals were of AR grade. Co(CH₃COO)₂·4H₂O (0.2 mmol, 49.8 mg) and pyrimidin-2-amine (0.2 mmol, 37.6 mg) were dissolved in methanol (20 ml). A solution of sodium fluoroborate (0.1 mmol, 10.9 mg) in water (3 ml) was then added. The mixture was stirred for 30 min at 353 K and the remaining solid was filtered off. After slow evaporation of the solvent from the filtrate red block-shaped crystals of (I) grew within 10 d. Analysis found: C 29.23, H 3.62, N 22.85%; calculated for C₁₈H₂₆B₂Co₂F₈N₁₂O₂: C 29.46, H 3.57, N 22.90%.

Z = 2

 $D_x = 1.140 \text{ Mg m}^{-3}$

 $0.32 \times 0.28 \times 0.20 \ \text{mm}$

2626 measured reflections

2541 independent reflections

2010 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 0.84 \text{ mm}^{-1}$

T = 293 (2) K

Block, red

 $R_{\rm int}=0.032$ $\theta_{\rm max} = 27.5^{\circ}$

Crystal data

[Co2(CH3O)2(C4H5N3)4](BF4)2 $M_r = 733.99$ Monoclinic, C2/m a = 16.338 (6) Å b = 14.704 (2) Å c = 12.040 (4) Å $\beta = 132.34 (1)^{\circ}$ V = 2138.0 (11) Å³

Data collection

Bruker APEX area-detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Bruker, 2001) $T_{\rm min}=0.775,\ T_{\rm max}=0.850$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.1252P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.064$	+ 2.7158 <i>P</i>]
$wR(F^2) = 0.207$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.11	$(\Delta/\sigma)_{\rm max} = 0.003$
2541 reflections	$\Delta \rho_{\rm max} = 0.86 \text{ e } \text{\AA}^{-3}$
109 parameters	$\Delta \rho_{\rm min} = -0.60 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N2 - H2B \cdots N3^{iv}$ $N2 - H2A \cdots F3^{v}$ $N2 - H2A \cdots F1^{vi}$	0.90 0.90 0.90	2.10 2.15 2.40	3.001 (6) 3.022 (6) 3.091 (6)	176 163 134

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

All C-bound H atoms were placed in idealized positions and were refined using a riding model, with C-H = 0.93 Å and $U_{iso}(H)$ = $1.2U_{eq}(C)$ for aromatic H atoms, and C-H = 0.96 Å and $U_{iso}(H) =$ $1.5U_{eq}(C)$ for methyl H atoms. The N-bound H atoms were located in a difference map, their bond lengths were set to 0.90 Å and they were refined using a riding model $[U_{iso}(H) = 1.5U_{eq}(N)]$. The methyl H atoms are disordered due to symmetry; site occupancy factors are 0.5. There are no indications that the channels are filled with solvent molecules.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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