metal-organic papers

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Key indicators

Single-crystal X-ray study T = 294 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.024 wR factor = 0.063 Data-to-parameter ratio = 12.6

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

Bis(2-aminonaphthalene-1-sulfonato- κO)bis(1,10-phenanthroline- $\kappa^2 N, N'$)cobalt(II)

The title molecule, $[Co(C_{10}H_8NO_3S)_2(C_{12}H_{10}N_2)_2]$, lies on a crystallographic twofold axis with the Co^{II} atom coordinated by four N atoms from two chelating 1,10-phenanthroline ligands and two sulfonate O atoms from two 2-aminonaph-thalene-1-sulfonate ligands in a distorted octahedral geometry. In the crystal structure, intermolecular $C-H\cdots O$ and $N-H\cdots O$ hydrogen bonds link the mononuclear units into a three-dimensional network. There is also a single $N-H\cdots O$ hydrogen bond and an intramolecular $\pi-\pi$ stacking interaction.

Comment

The design and synthesis of metal-organic frameworks (MOFs) by the mixed-ligand strategy is currently an active research area due to their intriguing structural architectures as well as their potential applications in catalysis, gas adsorption, and separation (Kitagawa *et al.*, 2004). The mixed-ligand strategy has been proven to be an effective method for the preparation of metal–sulfonate complexes (Cai, Chen, Liao, Feng *et al.*, 2001; Cai, Chen, Liao, Yao *et al.*, 2001; Chen *et al.*, 2002; Dai *et al.*, 2006).



As part of our further investigation of the coordination chemistry of aromatic sulfonate anions, we report here the crystal structure of a Co^{II} complex, (I), with mixed ligands of 1,10-phenanthroline and 2-aminonaphthalene-1-sulfonate. The molecular structure of (I) is shown in Fig. 1 and selected bond lengths and angles are listed in Table 1. The Co^{II} atom exhibits a distorted octahedral coordination geometry involving four N-atom donors from two chelating 1,10phenanthroline ligands, and two sulfonate O atoms from two

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Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level. Only H atoms involved in hydrogen bonds have been included. The short-dashed lines indicate intramolecular hydrogen bonds and the long-dashed lines indicate π - π stacking interactions [symmetry code: (A) y, x, -z].

monodentate 2-aminonaphthalene-1-sulfonate ligands. The molecule lies on a crystallographic twofold axis.

There is a single intramolecular π - π stacking interaction between the naphthalene ring system and the quinoline ring system of the 1,10-phenanthroline ligand, with a centroidcentroid distance of 3.670 (1) Å and a dihedral angle between the rings of 5.20 (4)°. In the crystal structure, intermolecular N-H ···O and C-H ···O hydrogen bonds link complex molecules, forming a three-dimensional network (Table 2 and Fig. 2).

Experimental

The title complex was synthesized by dissolving a reaction mixture of $Co(CH_3COO)_2 \cdot 4H_2O$ (99.6 mg, 0.4 mmol), 1,10-phenanthroline (39.6 mg, 0.2 mmol), 2-aminonaphthalene-1-sulfonic acid (89.3 mg, 0.4 mmol), NaOH (0.6 mmol), and H₂O (12 ml) in a 23 ml Teflonlined autoclave under autogenous pressure at 413 K for 2 d. After the reaction, the mixture was cooled slowly to room temperature at a rate of 5 K h⁻¹; red block-shaped crystals suitable for X-ray analysis were obtained in 30% yield. Analysis calculated for C₄₄H₃₂CON₆O₆S₂: C 61.18, H 3.73, N 9.73%; found: C 61.35, H 3.89, N 9.89%.

Crystal data

[Co(C10H8NO3S)2(C12H8N2)2]
$M_r = 863.81$
Tetragonal, $P4_{1}2_{1}2$
a = 12.468 (3) Å
c = 24.510 (9) Å
$V = 3810.3 (19) \text{ Å}^3$
Z = 4

Data collection

Bruker APEX-II CCD diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.812, T_{\max} = 0.883$ 20658 measured reflections 3363 independent reflections 3105 reflections with $I > 2\sigma(I)$ $R_{int} = 0.038$ $\theta_{max} = 25.0^{\circ}$

 $D_x = 1.506 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.62 \text{ mm}^{-1}$ T = 294 (2) K Block, red

0.44 \times 0.28 \times 0.20 mm



Figure 2

Part of the crystal structure of (I), showing $N-H\cdots O$ and $C-H\cdots O$ hydrogen bonds as dashed lines. Only H atoms involved in hydrogen bonds are shown.

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0406P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.024$	+ 0.043P]
$wR(F^2) = 0.063$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.04	$(\Delta/\sigma)_{\rm max} = 0.001$
3363 reflections	$\Delta \rho_{\rm max} = 0.13 \text{ e} \text{ \AA}^{-3}$
267 parameters	$\Delta \rho_{\rm min} = -0.26 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	Absolute structure: Flack (1983)
	1364 Friedel pairs

Flack parameter: -0.034 (12)

Table 1

Selected geometric parameters (Å, °).

Co1-O1	2.0716 (14)	Co1-N2	2.1523 (16)
Co1-N1	2.1520 (16)		
O1 ⁱ -Co1-O1	90.19 (8)	O1-Co1-N2	93.65 (6)
O1 ⁱ -Co1-N1	86.50 (6)	N1 ⁱ -Co1-N2	98.66 (6)
O1-Co1-N1	97.47 (6)	N1-Co1-N2	77.21 (6)
N1 ⁱ -Co1-N1	174.40 (9)	N2-Co1-N2 ⁱ	87.12 (8)
O1 ⁱ -Co1-N2	163.61 (6)		· · · ·

Symmetry code: (i) y, x, -z.

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

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N3-H3B\cdots O3^{ii}$	0.86	2.22	2.901 (3)	137
$N3-H3A\cdots O2$	0.86	1.95	2.650 (3)	137
C21-H21···O1	0.93	2.47	3.027 (3)	119
C21-H21···O3	0.93	2.50	3.063 (3)	119
C7−H7···O3 ⁱⁱⁱ	0.93	2.57	3.471 (3)	163
C8−H8···O2 ^{iv}	0.93	2.53	3.385 (3)	154
$C12{-}H12{\cdot}{\cdot}{\cdot}O1^i$	0.93	2.48	3.025 (3)	117
Symmetry codes:	(i) $v_1 x_2 - z_2$ (ii) $v + \frac{1}{2} - x + \frac{1}{2} - \frac{1}{2} + \frac{1}{2} - \frac{1}{2} + \frac{1}{2} - \frac{1}{2} + $	$\frac{1}{2}$, $z = \frac{1}{2}$; (iii) v	+1.x7; (iv)

Symmetry codes: (1) y, x, -z; (1) $y + \frac{1}{2}, -x + \frac{1}{2}, z - \frac{1}{4};$ (11) y + 1, x, -z; (1V) $x + \frac{1}{2}, -y + \frac{1}{2}, -z - \frac{1}{4}.$

H atoms were located in difference maps, but were subsequently placed in calculated positions and treated as riding, with C–H = 0.93 Å and N–H = 0.86 Å, with $U_{iso}(H) = 1.2U_{eq}(C, N)$.

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Data collection: *APEX2* (Bruker, 2003); 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: *SHELXTL* (Bruker, 2001) and *DIAMOND* (Brandenburg & Berndt, 1999); software used to prepare material for publication: *SHELXTL*.

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