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Key indicators

Single-crystal X-ray study T = 295 K Mean σ (C–C) = 0.003 Å R factor = 0.037 wR factor = 0.099 Data-to-parameter ratio = 15.2

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

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Diaquabis(3-hydroxybenzoato- κO)bis(5nitro-1*H*-benzimidazole- κN^3)cobalt(II) bis(5-nitro-1*H*-benzimidazole) dihydrate

The title compound, $[Co(C_7H_5O_3)_2(C_7H_5N_3O_2)_2(H_2O)_2]$ ·-2C₇H₅N₃O₂·2H₂O, is composed of neutral Co^{II}-containing complexes, accompanied by uncoordinated nitrobenzimidazole (Nbzim) molecules and water molecules. The Co^{II} atom is located on a twofold axis and coordinated by two Nbzim molecules, two 3-hydroxybenzoate anions and two water molecules in a slightly distorted octahedral geometry. The crystal packing is stabilized by extensive hydrogen bonding and π - π stacking between coordinated and uncoordinated Nbzim molecules.

Comment

As part of our ongoing investigations of the nature of π - π stacking in metal complexes (Li *et al.*, 2005; Zhang *et al.*, 2005), the title Co^{II} compound, (I), incorporating 5-nitrobenzimidazole (Nbzim) ligands, has been prepared and its crystal structure is presented here.



Compound (I) is composed of Co^{II} complexes, accompanied by non-coordinated Nbzim molecules and water molecules (Fig. 1). The Co^{II} atom is located on a twofold axis and is coordinated by two 3-hydroxybenzoate (HBA) anions, two Nbzim molecules and two water molecules in a slightly distorted octahedral geometry (Table 1). The cis bond angles about Co range from 85.88 (8) to 93.88 (6) $^{\circ}$ and the *trans* angles lie between 175.22 (7) and 175.97 (5)°. The monodentate HBA anions coordinate to the Co atom in a cis configuration, while the Nbzim molecules coordinate to the Co atom in a trans configuration. Within both the coordinated and uncoordinated Nbzim species, the nitro groups are almost coplanar with the benzimidazole skeleton, the dihedral angles being 2.9 (2) (N12-nitro) and 1.89 (19)° (N22-nitro). The coordinated N11-Nbzim molecule is approximately parallel to the uncoordinated Nbzim molecule, the dihedral angle being 10.87 (10)°.

A partially overlapped arrangement of the coordinated Nbzim and uncoordinated Nbzim molecules is observed (Fig. 2). The distances of atoms of N11-Nbzim to the N21Received 18 May 2006 Accepted 18 May 2006

V = 4372 (2) Å³ Z = 4

 $D_x = 1.607 \text{ Mg m}^{-3}$

 $0.31 \times 0.18 \times 0.11 \text{ mm}$

20853 measured reflections

 $w = 1/[\sigma^2(F_0^2) + (0.0541P)^2]$

+ 1.4439P] where $P = (F_0^2 + 2F_c^2)/3$

 $\Delta \rho_{\rm min} = -0.31 \text{ e } \text{\AA}^{-3}$

 $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.33 \text{ e } \text{\AA}^{-3}$

5000 independent reflections

3990 reflections with $I > 2\sigma(I)$)

Mo $K\alpha$ radiation

 $\mu = 0.49 \text{ mm}^{-1}$

T = 295 (2) K

Prism, pink

 $R_{\rm int} = 0.035$

 $\theta_{\rm max} = 27.5^{\circ}$



Figure 1

The molecular structure of (I), shown with 30% probability displacement ellipsoids (arbitrary spheres for H atoms) [symmetry code: (i) 1 - x, y, $\frac{1}{2} - z$].



Figure 2 A diagram showing the π - π stacking in (I).

Nbzim mean plane are 3.505 (2) (N11), 3.277 (3) (C12) and 3.493 (2) Å (C18). These findings suggest the existence of $\pi - \pi$ stacking between coordinated Nbzim and uncoordinated Nbzim.

An extensive network of $O-H \cdots O$, $O-H \cdots N$ and N- $H \cdots O$ hydrogen bonds helps to stabilize the crystal structure of (I) (Table 2).

Experimental

CoCl₂·6H₂O (0.24 g, 1 mmol), Na₂CO₃ (0.053 g, 0.5 mmol) and sodium 3-hydroxybenzoate (0.16 g, 1 mmol) were dissolved in a water-ethanol solution (15 ml, 1:2). 6-Nitrobenzimidazole (6-Nbzim) (0.33 g, 1 mmol) was then added to the above solution. The mixture was refluxed for 2.5 h, and then filtered after cooling to room

Crystal data

[Co(C7H5O3)2(C7H5N3O2)2- $(H_2O)_2] \cdot 2C_7H_5N_3O_2 \cdot 2H_2O$ $M_r = 1057.77$ Monoclinic C_2/c a = 26.536 (8) Å b = 9.547 (3) Å c = 18.026 (4) Å $\beta = 106.79 (1)^{\circ}$

Data collection

Rigaku R-AXIS RAPID diffractometer (i) scans Absorption correction: multi-scan (ABSCOR; Higashi, 1995) $T_{\min} = 0.857, \ T_{\max} = 0.945$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.037$ wR(F²) = 0.099 S = 1.055000 reflections 330 parameters H-atom parameters constrained

Table 1

S

Selected bond lengths (Å).					
Co-O1	2.1375 (14)	C12-N13	1.313 (2)		
Co-O31	2.0821 (12)	C12-N11	1.348 (2)		
Co-N13	2.1400 (15)	C22-N23	1.311 (3)		
C31-O31	1.262 (2)	C22-N21	1.349 (3)		
C31-O32	1.265 (2)				

Table 2

Hydrogen-bond geometry (Å, °).

$\overline{D-\mathrm{H}\cdots A}$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1-H1A\cdots N23$	0.93	2.13	3.041 (3)	166
$O1 - H1B \cdots O32$	0.95	1.78	2.736 (2)	175
$O33-H3\cdots O1W^{i}$	0.95	1.76	2.704 (2)	172
$O1W-H1\cdots N23$	0.93	2.08	2.994 (2)	168
$O1W - H2 \cdot \cdot \cdot O32$	0.89	2.00	2.884 (2)	173
$N11-H11\cdots O32^{ii}$	0.86	2.17	2.934 (2)	147
$N21 - H21 \cdots O33^{ii}$	0.86	2.10	2.887 (2)	153

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

Water H atoms and the hydroxy H atom were located in a difference Fourier map and refined as riding in their as-found relative positions (O-H = 0.89–0.95 Å), with $U_{iso}(H) = 1.5U_{ea}(O)$. Other H atoms were placed in calculated positions, C-H = 0.93 Å and N-H= 0.86 Å, and refined as riding, with $U_{iso}(H) = 1.2U_{eq}(C,N)$.

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/ MSC, 2002); program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia,

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1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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