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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.005 \text{ Å}$ H-atom completeness 82% Disorder in solvent or counterion R factor = 0.042 wR factor = 0.130 Data-to-parameter ratio = 16.8

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

© 2003 International Union of Crystallography Printed in Great Britain – all rights reserved The title cobalt(III) complex, $[Co(C_9H_6NO)_3]$ ·CH₃OH, features three 8-quinolinolate ligands that chelate the cobalt(III) atom with an octahedral coordination geometry. The methanol solvent molecule is disordered in the structure. The O···O distances 2.726 (7) and 2.871 (8) Å between the quinolinol hydroxyl and disordered methanol suggest the existence of hydrogen bonding. The separation distances of 3.404 (15) and 3.386 (7) Å between neighboring complex molecules.

Comment

As part of an investigation into π - π stacking interactions in metal complexes, several 8-quinolinolate metal complexes have been prepared in our laboratory (Li *et al.*, 2003). Compound (I) is a representative of this group.



The structure of (I) is shown in Fig. 1. Three 8-quinolinolate monoanions chelate to the Co^{III} atom with an octahedral coordination geometry. The planar 8-quinolinolate monoanions are nearly perpendicular to each other, with dihedral angles of 93.80 (7), 95.34 (6) and 91.49 (7)°. The overlapped arrangement of neighboring parallel quinoline rings is shown in Fig. 2. The quinoline plane containing N11(1 - x, 1 - y, 1 - z) by 3.404 (15) Å. Likewise, the quinoline plane containing N31(1 - x, 1 - y, -z) by 3.386 (7) Å. These findings indicate the existence of π - π stacking between the neighboring complex molecules.

The methanol solvent molecule is disordered, both C and O atoms being located in two sites with 0.5 occupancy factors. Although H atoms of the disordered methanol were not located, the O1a···O11 distance of 2.726 (7) Å and the O1b···O11 distance of 2.871 (8) Å suggest the existence of hydrogen bonding between the quinolinol hydroxyl group and the disordered methanol molecule. The C24–H24···O1b($\frac{1}{2} - x$, $y - \frac{1}{2}$, $\frac{1}{2} - z$) angle of 133° and the

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The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level. Dashed lines indicate hydrogen bonds. Both disorder components of methanol are shown.



A packing diagram, showing the π - π interaction between neighboring quinoline rings.

 $C24 \cdots O1b$ distance of 3.154 (8) Å suggest that the disordered methanol molecule is also involved in weak C-H···O hydrogen bonding.

Compound (I) is isomorphous with the manganese(III) compound (Hems & Mackay, 1975). The major difference between the Co and Mn complexes is the length of the coordination bonds: Mn-O distances range from 1.905 to 1.924 Å and Co-O distances range from 1.8975 (19) to 1.9066 (19) Å; Mn–N distances range from 2.059 to 2.266 Å, while Co-N distances range from 1.921 (2) to 1.936 (2) Å. The other difference between the two structures is that the solvent methanol molecule in the Mn structure was not treated with a disorder model, even though the C-O distance of 1.352 Å is shorter than the expected value.

Experimental

The complex was prepared by refluxing a methanol solution (15 ml) containing CoCl₂·6H₂O (0.24 g, 1 mmol) and 8-quinolinol (0.15 g, 1 mmol) for 2 h. The resulting solution was filtered at room temperature. Brown single crystals were obtained from the filtrate after two weeks.

Crystal data

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1

$Co(C_9H_6NO)_3]$ ·CH ₄ O	$D_x = 1.466 \text{ Mg m}^{-3}$	
$A_r = 523.42$	Mo $K\alpha$ radiation	
Aonoclinic, $P2_1/n$	Cell parameters from 9762	
= 10.9235 (12) Å	reflections	
= 13.1172 (14) Å	$\theta = 3.0-27.0^{\circ}$	
= 16.6861 (14) Å	$\mu = 0.77 \text{ mm}^{-1}$	
$B = 97.325 \ (8)^{\circ}$	T = 298 (2) K	
$V = 2371.4 (4) Å^3$	Prism, brown	
Z = 4	$0.36 \times 0.20 \times 0.10 \text{ mm}$	
Data collection		
Rigaku R-AXIS-RAPID	5432 independent reflections	

Kigaku K-AXIS-KAPID	5452 independ
diffractometer	4009 reflection
v scans	$R_{\rm int} = 0.020$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(ABSCOR; Higashi, 1995)	$h = -14 \rightarrow 14$
$T_{\min} = 0.750, \ T_{\max} = 0.918$	$k = -17 \rightarrow 17$
0460 measured reflections	$l = -21 \rightarrow 21$

Refinement

F

S 5

3

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0805P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 0.1393P]
$\nu R(F^2) = 0.130$	where $P = (F_o^2 + 2F_c^2)/3$
1 = 1.04	$(\Delta/\sigma)_{\rm max} = 0.003$
432 reflections	$\Delta \rho_{\rm max} = 0.59 \ {\rm e} \ {\rm \AA}^{-3}$
23 parameters	$\Delta \rho_{\rm min} = -0.42 \text{ e } \text{\AA}^{-3}$
I-atom parameters constrained	

Table 1

Selected geometric parameters (Å).

Co-O31	1.8975 (19)	Co-N11	1.921 (2)
Co-O21	1.8994 (18)	Co-N21	1.924 (2)
Co-O11	1.9066 (19)	Co-N31	1.936 (2)

The disordered methanol molecule was refined isotropically and its H atoms were not located. Other H atoms were placed in calculated positions with C-H = 0.93 Å, and included in the final cycles of refinement using a riding model, with $U_{iso}(H) = 1.2U_{eq}$ of the carrier atoms.

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

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

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