Acta Cryst. (2003). E59, m809–m811 DOI: 10.1107/S1600536803018762 Wu, Xue and Xu • [Co(C₄H₄O₅)(C₁₂H₈N₂)(H₂O)]·1.5H₂O **m809**

Figure 1

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.005 \text{ Å}$ R factor = 0.038 wR factor = 0.109 Data-to-parameter ratio = 16.6

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

Aqua(oxydiacetato- $\kappa^3 O, O', O''$)(1,10phenanthroline- $\kappa^2 N, N'$)cobalt(II) sesquihydrate

The title compound, $[Co(C_4H_4O_5)(C_{12}H_8N_2)H_2O] \cdot 1.5H_2O$, consists of the octahedral complex and lattice water. The asymmetric unit contains two molecules of the complex, which form a hydrogen-bonded dimer. The overlapped arrangement and shorter separations of 3.532 (6) and 3.385 (12) Å between neighboring parallel phenanthroline rings suggest the existence of π - π -stacking interactions.

Comment

Aromatic π - π stacking interactions are important for some biological systems (Deisenhofer & Michel, 1989). As part of our investigation of π - π stacking in metal complexes (Chen *et al.*, 2003), the title Co^{II} complex, (I), was prepared.

 Co^{II} compound (I) is isomorphous with the Ni^{II} (Baggio *et al.*, 2000) and Zn^{II} analogues (Baggio *et al.*, 1996). The asymmetric unit of (I) contains two molecules of the complex and three molecules of lattice water, as shown in Fig. 1. The two independent complex molecules display a similar octa-



© 2003 International Union of Crystallography Printed in Great Britain – all rights reserved The molecular structure of (I), with 30% probability displacement ellipsoids. Dashed lines indicate hydrogen bonding.



Received 15 August 2003 Accepted 26 August 2003 Online 30 August 2003



Figure 2

The molecular packing diagram, showing the hydrophilic and hydrophobic layered structure. H atoms have been omitted for clarity.



Figure 3

A diagram showing the π - π stacking between (a) N1-phen and N1ⁱⁱⁱ-phen rings, and (b) N3-phen and N3^{iv}-phen rings. [Symmetry codes: (iii) 1 - x, -y, 1-z; (iv) -x, 1-y, 1-z.]

(*b*)

hedral coordination geometry (Table 1), formed by a phenanthroline (phen), a meridional oxydiacetate and a coordinated water molecule. The lattice water molecules are hydrogen bonded to the adjacent carboxyl groups (Table 2). Through hydrogen bonding between carboxyl groups and coordinated water molecules, two independent complex molecules link to each other to form a dimer (Fig. 1). All lattice water molecules are ordered in the Co^{II} compound, as is the case in the isomorphous Zn^{II} complex but different from the disorder found in the isomorphous Ni^{II} complex.

The molecular packing in the unit cell is illustrated in Fig. 2. The crystal structure consists of the hydrophilic layers formed by carboxylate and water molecules and hydrophobic layers formed by phen ligands. Parallel disposition of phen ligands is observed in the crystal. Fig. 3 shows the overlapped arrangement of neighboring parallel phen ligands. The separations of 3.532 (6) Å between N1-phen and N1ⁱⁱⁱ-phen and 3.385 (12) Å between the N3-phen and N3^{iv}-phen [symmetry codes: (iii) 1 - x, -y, 1 - z; (iv) -x, 1 - y, 1 - z] suggest the existence of aromatic π - π stacking between phen ligands.

Experimental

An aqueous solution (10 ml) containing CoCl₂·6H₂O (0.24 g, 1 mmol) and 1,10-phenanthroline (0.18 g, 1 mmol) was mixed with an aqueous solution (10 ml) containing oxydiacetic acid hydrate (0.15 g, 1 mmol) and NaOH (0.08 g, 2 mmol) at room temperature. The mixture was refluxed for 3 h. Pink single crystals of the title compound were obtained from the filtrate after 2 weeks.

Crystal data

$[Co(C_4H_4O_5)(C_{12}H_8N_2)-$	$D_x = 1.591 \text{ Mg m}^{-3}$
$(H_2O)] \cdot 1.5H_2O$	Mo $K\alpha$ radiation
$M_r = 416.25$	Cell parameters from 16878
Monoclinic, $P2_1/c$	reflections
a = 10.4722 (16)Å	$\theta = 2.8-24.0^{\circ}$
b = 24.3161 (9) Å	$\mu = 1.03 \text{ mm}^{-1}$
c = 13.6729 (14) Å	T = 293 (2) K
$\beta = 93.627 \ (1)^{\circ}$	Prism, pink
$V = 3474.7 (7) \text{ Å}^3$	$0.32 \times 0.28 \times 0.20 \text{ mm}$
Z = 8	

Data collection

Bruker SMART CCD diffractometer ω and φ scans Absorption correction: multi-scan (SADABS; Bruker, 1999)

 $h = -13 \rightarrow 13$ $T_{\min} = 0.715, \ T_{\max} = 0.810$ $k = -31 \rightarrow 30$ 21949 measured reflections $l = -11 \rightarrow 17$

Refinement

Refinement on F^2 H-atom parameters constrained $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0452P)^{2}]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.109$ S = 0.93 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.56 \text{ e } \text{\AA}^{-3}$ 7932 reflections 479 parameters $\Delta \rho_{\rm min} = -0.41 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Selected geometric parameters (Å).

Co1-O1	2.070 (2)	Co2-O12	2.060 (2)
Co1-O6	2.074 (2)	Co2-O10	2.096 (2)
Co1-O4	2.084 (2)	Co2-O7	2.117 (2)
Co1-N2	2.098 (2)	Co2-N3	2.118 (2)
Co1-N1	2.116 (2)	Co2-O9	2.118 (2)
Co1-O3	2.130 (2)	Co2-N4	2.119 (2)

7932 independent reflections

 $R_{\rm int}=0.039$

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

4856 reflections with $I > 2\sigma(I)$

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

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O6-H61···O8	0.87	1.79	2.647 (3)	169
$O6-H62\cdots O11^{i}$	0.92	1.77	2.685 (3)	178
O12-H121O5	0.91	1.77	2.676 (3)	175
$O12-H122\cdots O2^{ii}$	0.96	1.64	2.596 (3)	171
O21-H211···O1	0.91	2.04	2.933 (3)	166
$O21 - H212 \cdot \cdot \cdot O5^i$	0.95	1.95	2.890 (3)	168
O22-H221···O7	0.95	1.94	2.867 (3)	163
$O22-H222\cdots O11^{i}$	0.88	1.95	2.811 (3)	168
O23-H231···O10	0.87	2.07	2.915 (3)	165
$O23\!-\!H232\!\cdots\!O22^{ii}$	0.87	2.24	3.094 (3)	168

Symmetry code: (i) 1 + x, y, z; (ii) x - 1, y, z.

H atoms of the water molecules were located in a difference Fourier map and were included in the structure-factor calculations with fixed coordinates and isotropic displacement parameters of 0.05 Å^2 . Other H atoms were placed in calculated positions, with C– H = 0.93 Å (phen) or 0.97 Å (oxydiacetate) and were included in the final cycles of refinement in a riding model, with $U_{iso}(H) = 1.2U_{eq}$ of the carrier atoms.

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1993); program(s) used to refine

structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

This project was supported by the National Natural Science Foundation of China (Nos. 29973036 and 20240430654). The authors thank Dr Chen-Hsiung Hung for assistance in the data collection.

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