Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.008 Å R factor = 0.063 wR factor = 0.158 Data-to-parameter ratio = 13.8

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. Received 7 February 2007

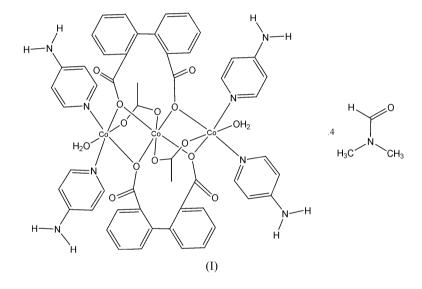
Accepted 1 March 2007

Di- μ_2 -acetato-tetrakis(4-aminopyridine)diaquabis(μ_3 -biphenyl-2,2'-dicarboxylato)tricobalt(II) *N*,*N*'-dimethylformamide tetrasolvate

The title complex, $[Co_3(C_2H_3O_2)_2(C_14H_8O_4)_2(C_5H_6N_2)_4(H_2O)_2]\cdot4C_3H_7NO$, is a linear trinuclear cobalt complex crystallizing with the central Co atom on a centre of symmetry and with with four dimethylformamide solvent molecules per complex. The crystal packing is stabilized by $N-H\cdots O$ and $O-H\cdots O$ hydrogen bonds.

Comment

Linear tricobalt complexes have been investigated extensively because of their potential application in molecular wires, as magnetic materials and as catalysts (Clerac *et al.*, 2000).



In the title complex, (I) (Fig. 1), the outer Co atoms are sixcoordinate in a slightly distorted octahedral mode. Whereas the central Co atom is bonded to six O atoms, the terminal two Co atoms are each bonded to four O and two N atoms. The Co1···Co2 distance is 3.2141 (7) Å, which is slightly shorter than in the complex $[Co_3(C_{18}H_{18}N_2O_2)_2(C_2H_3O_2)_2]$ (You *et al.*, 2004). The crystal packing is stabilized by N-H···O and O-H···O hydrogen bonds (Table 1).

Experimental

When a mixture of Co(CH₃COO)₂·2H₂O (0.0754 g, 0.032 mmol), diphenic acid (0.0772 g, 0.032 mmol) and 4-aminopyridine (0.0570 g, 0.060 mmol) was dissolved in CH₃OH (10 ml) and dimethyl-formamide (10 ml), a red solution was obtained, which was set aside at room temperature and allowed to evaporate slowly. After 10 d, pink block-shaped crystals were collected by filtration. Analysis calculated for C₆₄H₇₈Co₃N₁₂O₁₈: C 51.93, H 5.31, N 11.36%; found: C 51.98, H 5.36, N 11.41%.

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Crystal data

 $\begin{bmatrix} Co_3(C_2H_3O_2)_2(C_{14}H_8O_4)_2 - \\ (C_5H_6N_2)_4(H_2O)_2 \end{bmatrix} \cdot 4C_3H_7NO \\ M_r = 1480.17 \\ Monoclinic, P2_1/c \\ a = 12.2379 (17) \\ b = 11.1660 (15) \\ b = 25.985 (4) \\ A \end{bmatrix}$

Data collection

Bruker SMART CCD diffractometer Absorption correction: multi-scan (*SADABS*; Bruker, 1997) $T_{\rm min} = 0.920, T_{\rm max} = 0.927$

Refinement

 $\begin{array}{ll} R[F^2 > 2\sigma(F^2)] = 0.063 & \text{H atoms treated by a mixture of} \\ wR(F^2) = 0.158 & \text{independent and constrained} \\ S = 1.02 & \text{refinement} \\ 6218 \text{ reflections} & \Delta\rho_{\max} = 0.40 \text{ e} \text{ Å}^{-3} \\ 452 \text{ parameters} & \Delta\rho_{\min} = -0.41 \text{ e} \text{ Å}^{-3} \end{array}$

 $\beta = 96.571 \ (2)^{\circ}$

Z = 2

V = 3527.5 (8) Å³

Mo $K\alpha$ radiation $\mu = 0.77 \text{ mm}^{-1}$

 $0.11 \times 0.10 \times 0.10$ mm

24864 measured reflections

6218 independent reflections

3850 reflections with $I > 2\sigma(I)$

T = 293 (2) K

 $R_{\rm int} = 0.100$

Table '	1
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Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N2-H2A···O8	0.86	2.15	2.966 (8)	157
$N2-H2B\cdots O9$	0.86	2.03	2.832 (7)	155
N4 $-$ H4 A ···O7 ⁱ	0.86	2.10	2.944 (6)	164
$N4 - H4B \cdots O9^{ii}$	0.86	2.13	2.983 (7)	170
$O7-H7A\cdots O2^{iii}$	0.85 (4)	1.80 (4)	2.627 (5)	164 (4)
$O7 - H7B \cdots O3$	0.85 (2)	1.75 (2)	2.605 (6)	175 (4)

Symmetry codes: (i) $-x, y + \frac{1}{2}, -z + \frac{3}{2}$; (ii) x + 1, y, z; (iii) -x, -y + 2, -z + 2.

H atoms bonded to C and N were positioned geometrically and refined using a riding model, with aromatic C-H = 0.93 Å, methyl C-H = 0.96 Å and N-H = 0.86 Å. $U_{iso}(H)$ values were set at $1.2U_{eq}(C,N)$ or $1.5U_{eq}(methyl C)$. The methyl groups were allowed to rotate but not to tip. Water H atoms were found in a difference Fourier map and refined with an O-H distance restraint of 0.89 (2) Å and with free $U_{iso}(H)$ values.

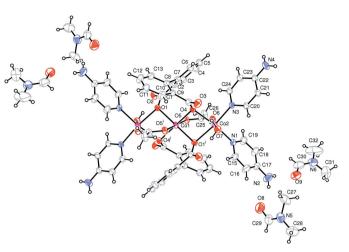


Figure 1

The molecular structure of the title compound, showing the atomnumbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 30% probability level. [Symmetry code: (i) -x, 2 - y, 2 - z.]

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT* and *SHELXTL* (Bruker, 1997); program(s) used to solve structure: *SHELXS86* (Sheldrick, 1985); 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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