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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.023 wR factor = 0.062 Data-to-parameter ratio = 10.9

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

catena-Poly[[aqua(diformato- κO)copper(II)]- μ -1,4-diazabicyclo[2.2.2]octane- $\kappa^2 N:N'$]

The title compound, $[Cu(CHO_2)_2(C_6H_{12}N_2)(H_2O)]_n$, forms a polymeric chain, $[Cu(HCOO)_2(dabco)(H_2O)]_{\infty}$ (dabco is 1,4diazabicyclo[2.2.2]octane). Both formate ligands are Omonodentate anions and dabco acts as a bridging ligand, creating a linear polymeric arrangement interconnected by $O_{water} - H \cdots O_{carboxy}$ hydrogen bonds. The deformed squarepyramidal Cu^{II} coordination comprises two N and two O atoms as the base, and a water molecule in the apical position. The point symmetry of the Cu^{II} polyhedron and the dabco ligand is *mm*, and the formate anions lie on the mirror planes $\frac{1}{4}$, *y*, *z* and $\frac{3}{4}$, *y*, *z*.

Comment

So far, Cu^{II} -dabco coordination compounds has not been intensively investigated. Only one mononuclear structure (Karan *et al.*, 1999), two dinuclear structures (Durley *et al.*, 1980; Maverick *et al.*, 1986) and one polymeric structure (Rao *et al.*, 1983) of Cu^{II} compounds containing the dabco ligand have been reported. The present paper reports the first example of the coordination of the dabco ligand in a basal position of a square-pyramidal Cu^{II} polyhedron. The structure of the title complex, (I), is polymeric, with $[Cu(HCOO)_2(dabco)(H_2O)]_{\infty}$ chains running along the *a* axis. In Fig. 1, the labelled atoms indicate the independent fragment of the chain.



The chain consists of pentacoordinated Cu^{II} ions in a distorted square-pyramidal (SQP) geometry, with two Cu–N bonds of 2.093 (2) Å and two Cu–O(formate) bonds of 1.962 (2) Å in the basal plane. The apical position is occupied by the water molecule [Cu–OH₂ = 2.238 (2) Å]. The Cu atom is displaced from the basal plane by 0.124 (1) Å towards atom O1. The point symmetry of the Cu^{II} polyhedron and the dabco ligand is *mm*, and the formate anions lie on the mirror planes $\frac{1}{4}$, *y*, *z* and $\frac{3}{4}$, *y*, *z*.

The observed SQP coordination is distinctly deformed in the direction of trigonal-bipyramidal (TBP) coordination, with

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Figure 1

A fragment of the polymeric structure of the title compound. Displacement ellipsoids for non-H atoms are drawn at the 40% probability level.



Figure 2

Perspective view of the crystal packing in the unit cell, showing the linkage of the polymeric chains by hydrogen bonding as dashed lines.

the trigonality parameter $\tau = 0.24$ [τ is defined by Addison *et al.* (1984)]; for the regular SQP structure, the trigonality parameter is 0 and for TBP distortion it increases to 1.

The formate group acts as a monodentate ligand, the distance between the Cu^{II} ion and uncoordinated atom O3 being 3.287 (3) Å. Such behaviour may be caused by the participation of this atom in a strong hydrogen bond with the water molecule. These interchain interactions, running along the *c* axis, are shown in Fig. 2. This strong hydrogen bond does not cause a delocalization of the π bond in the carboxyl group as the C2–O2 and C2–O3 bonds are distinctly different [1.257 (3) and 1.214 (4) Å, respectively].

The intrachain Cu···Cu distance of 6.808 (1) Å is longer than the shortest interchain Cu···Cu distance of 6.422 (2) Å along the *c* axis. Other short interchain Cu···Cu distances of 7.246 (2) and 8.157 (2) Å are between the two Cu^{II} ions related by the 2_1 screw axis, and with no spacer between them.

Experimental

The title complex was prepared by dissolving cupric formate $[Cu(HCOO)_2 \cdot 2H_2O, 2 \text{ mmol}]$ in 50 ml of water with dabco $(C_6H_{12}N_2, 2 \text{ mmol})$. After heating to boiling, a few drops of formic acid were added to clear the solution. The solution was filtered and allowed to cool. After several days, turquoise crystals were obtained.

Crystal data

 $[Cu(CHO_2)_2(C_6H_{12}N_2)(H_2O)]$ Mo Ka radiation Cell parameters from 38 $M_r = 283.77$ Orthorhombic, Pmmn reflections a = 6.8084 (13) Å $\theta = 5 - 18^{\circ}$ $\mu = 2.08 \text{ mm}^{-1}$ b = 12.071 (2) Åc = 6.4224 (15) ÅT = 293 K $V = 527.80 (19) \text{ Å}^3$ Prism, turquoise Z = 2 $0.30 \times 0.20 \times 0.20 \mbox{ mm}$ $D_x = 1.786 \text{ Mg m}^{-3}$ Data collection Siemens P3 diffractometer $\theta_{\rm max} = 28.0^\circ$ $h = 0 \rightarrow 8$ ω -2 θ scans $k = 0 \rightarrow 15$ Absorption correction: ψ scan (North et al., 1968) $l = 0 \rightarrow 8$ $T_{\min} = 0.557, \ T_{\max} = 0.661$ 3 standard reflections 719 measured reflections every 100 reflections 719 independent reflections intensity decay: none 710 reflections with $I > 2\sigma(I)$ Refinement Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0345P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.023$ + 0.21P] where $P = (F_o^2 + 2F_c^2)/3$ $wR(F^2) = 0.062$ $(\Delta/\sigma)_{\rm max} < 0.001$ S = 1.23719 reflections $\Delta \rho_{\rm max} = 0.39 \, {\rm e \, \AA}$ $\Delta \rho_{\rm min} = -0.49 \,\mathrm{e} \,\mathrm{\AA}^{-3}$ 66 parameters All H-atom parameters refined

Table 1

Selected geometric parameters (Å, °).

Cu-O1	2.238 (2)	O2-C2	1.257 (3)
Cu-O2 Cu-N1	1.962 (2) 2.093 (2)	O3-C2	1.214 (4)
O1-Cu-O2 O1-Cu-N1	98.42 (6) 88.90 (5)	$O2-Cu-O2^i$ N1-Cu-N1 ⁱ	163.16 (11) 177.81 (9)
O2-Cu-N1	90.16 (1)		

Symmetry code: (i) $\frac{1}{2} - x, \frac{1}{2} - y, z$.

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
$O1-H1\cdots O3^{ii}$	0.94 (4)	1.77 (4)	2.699 (3)	170 (4)
Symmetry code: (ii) $\frac{1}{2}$ -	$-x, \frac{1}{2} - y, 1 + z$			

All H atoms were located from a difference synthesis and refined isotropically. The C–H distances range from 0.92 (3) to 1.00 (4) Å and the O–H distance refined to 0.94 (4) Å.

Data collection: *P3* (Siemens, 1993); cell refinement: *P3*; data reduction: *XDISK* in *SHELXTL/PC* (Sheldrick, 1990); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL/PC*; software used to prepare material for publication: *PLATON* (Spek, 1990).

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