

catena-Poly[[[tris(pyridine- κN)copper(II)]- μ -3-carboxylatophenoxyacetato- $\kappa^2 O^3:O'$] trihydrate]

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Key indicators

Single-crystal X-ray study
 T = 295 K
 Mean $\sigma(C-C) = 0.009 \text{ \AA}$
 R factor = 0.073
 wR factor = 0.146
 Data-to-parameter ratio = 17.0

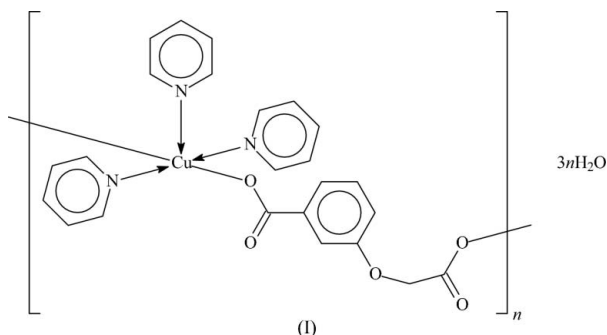
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The 3-carboxyphenoxyacetate ligand in the title compound, $[Cu(C_9H_6O_5)(C_5H_5N)_3] \cdot 3H_2O$, links adjacent tripyridine-copper(II) groups into a helical chain, which runs along the *c* axis of the hexagonal unit cell. The covalently bonded O atoms occupy *trans* sites in the basal plane of the square-pyramidal coordination of the Cu atom. The uncoordinated water molecules connect the chains into a three-dimensional network.

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Comment

We are interested in metal complexes with carboxyphenoxyacetate because the dianion is a multidentate ligand with both rigid and flexible parts. We present here the crystal structure of the title Cu^{II} complex, (I), in which 3-carboxyphenoxyacetate plays the role of bridging ligand.



The Cu^{II} atom has a square-pyramidal coordination geometry (Fig. 1 and Table 1). The $O_2C-C_6H_4O-CH_2-CO_2$ dianion links adjacent tripyridinecopper(II) cations into a helical chain, which runs along the *c* axis of the hexagonal unit cell (Fig. 2). The uncoordinated water molecules connect neighboring chains into a three-dimensional network through hydrogen bonds (Table 2). In the chain, the copper shows square-pyramidal coordination, with the covalently bonded O atoms occupying *trans* sites of the basal plane.

Other $O_2CC_6H_4OCH_2-CO_2Cu$ adducts that have been characterized include the imidazole adduct, in which the ether O atom is also involved in coordination (Gao *et al.*, 2004a), and the benzimidazole adduct, in which the dianion functions in the μ_4 -bridging mode (Gao *et al.*, 2004b). In this pyridine adduct, each carboxylate end of the dianion is only monodentate to copper.

Experimental

Copper dinitrate hexahydrate (1.48 g, 5 mmol) and a slight excess of pyridine (1 ml) were added to an aqueous solution of 3-carboxy-

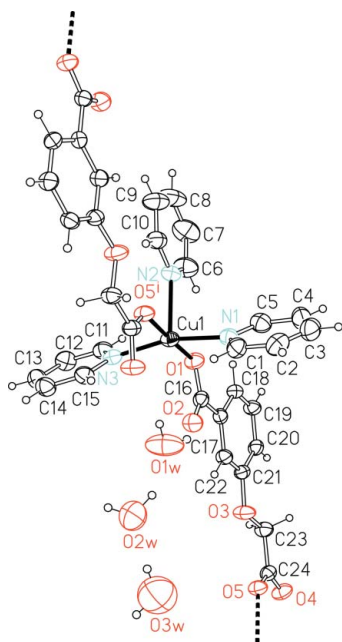


Figure 1
ORTEP (Johnson, 1976) plot of (I). Displacement ellipsoids are drawn at the 30% probability level, and H atoms are drawn as spheres of arbitrary radii. [Symmetry code: (i) $1 - x, -y, -\frac{1}{2} + z$.]

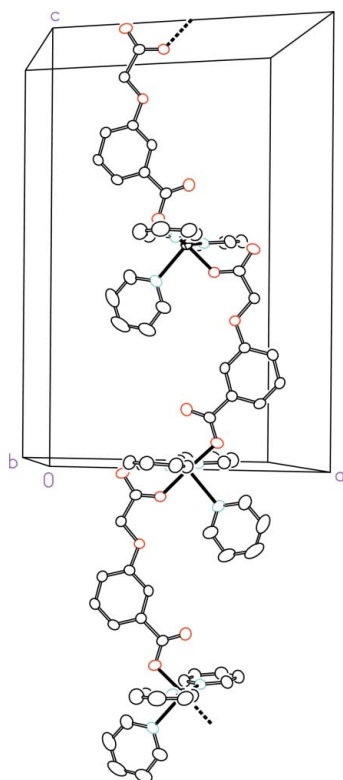


Figure 2
ORTEP (Johnson, 1976) plot of the polymeric chain; the water molecules and H atoms are not shown.

phenoxyacetic acid (0.97 g, 5 mmol). Drops of 0.2 M sodium hydroxide solution were added until the pH of the solution was approximately 6. Blue block-shaped crystals of (I) were obtained after a week. Analysis calculated for $C_{24}H_{27}CuN_3O_8$: C 52.54, H 4.96, N 7.66%; found: C 52.52, H 4.98, N 6.69%.

Crystal data

$[Cu(C_9H_6O_5)(C_5H_5N)_3] \cdot 3H_2O$
 $M_r = 549.04$
 Hexagonal, $P6_3$
 $a = 14.408 (2) \text{ \AA}$
 $c = 21.630 (4) \text{ \AA}$
 $V = 3888.6 (8) \text{ \AA}^3$
 $Z = 6$
 $D_x = 1.407 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation
 Cell parameters from 26385 reflections
 $\theta = 3.3\text{--}27.5^\circ$
 $\mu = 0.89 \text{ mm}^{-1}$
 $T = 295 (2) \text{ K}$
 Block, blue
 $0.37 \times 0.25 \times 0.19 \text{ mm}$

Data collection

Rigaku R-Axis RAPID IP diffractometer
 ω scans
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.591, T_{\max} = 0.849$
 27636 measured reflections

5518 independent reflections
 4814 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$
 $\theta_{\max} = 27.5^\circ$
 $h = -18 \rightarrow 18$
 $k = -18 \rightarrow 18$
 $l = -28 \rightarrow 26$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.073$
 $wR(F^2) = 0.146$
 $S = 1.24$
 5518 reflections
 325 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0697P)^2 + 0.7075P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.03$
 $\Delta\rho_{\max} = 0.48 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.39 \text{ e \AA}^{-3}$
 Absolute structure: Flack (1983),
 2571 Friedel pairs
 Flack parameter: 0.05 (2)

Table 1

Selected geometric parameters ($\text{\AA}, ^\circ$).

Cu1—O1	1.952 (3)	Cu1—N2	2.332 (4)
Cu1—O5 ⁱ	1.951 (3)	Cu1—N3	2.055 (4)
Cu1—N1	2.057 (4)		
O1—Cu1—O5 ⁱ	177.5 (2)	O5 ⁱ —Cu1—N2	87.9 (2)
O1—Cu1—N1	89.1 (2)	O5 ⁱ —Cu1—N3	90.8 (1)
O1—Cu1—N2	89.7 (1)	N1—Cu1—N2	97.1 (2)
O1—Cu1—N3	89.9 (1)	N1—Cu1—N3	164.6 (2)
O5 ⁱ —Cu1—N1	90.8 (1)	N2—Cu1—N3	98.3 (1)

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

Table 2

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1w—H1w1 \cdots O2	0.85	2.17	2.800 (6)	131
O1w—H1w2 \cdots O4 ⁱ	0.85	2.03	2.835 (6)	158
O2w—H2w1 \cdots O4 ⁱⁱ	0.86	2.25	2.789 (7)	121
O2w—H2w2 \cdots O1w	0.86	2.22	2.766 (8)	121
O3w—H3w1 \cdots O2w	0.86	1.83	2.69 (1)	170

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

C-bound H atoms were placed in calculated positions [$C-H = 0.93$ (aromatic) and 0.97 \AA (methylene)] and included in the refinement as riding [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$]. Water H atoms were placed in chemically sensible positions on the basis of hydrogen-bonding interactions but were not refined [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{O})$].

Data collection and cell refinement: *RAPID-AUTO* (Rigaku, 1998); data reduction: *CrystalStructure* (Rigaku/MSK, 2002); structure solution: *SHELXS97* (Sheldrick, 1997); structure refinement: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP* (Johnson, 1976); publication material: *SHELXL97*.

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