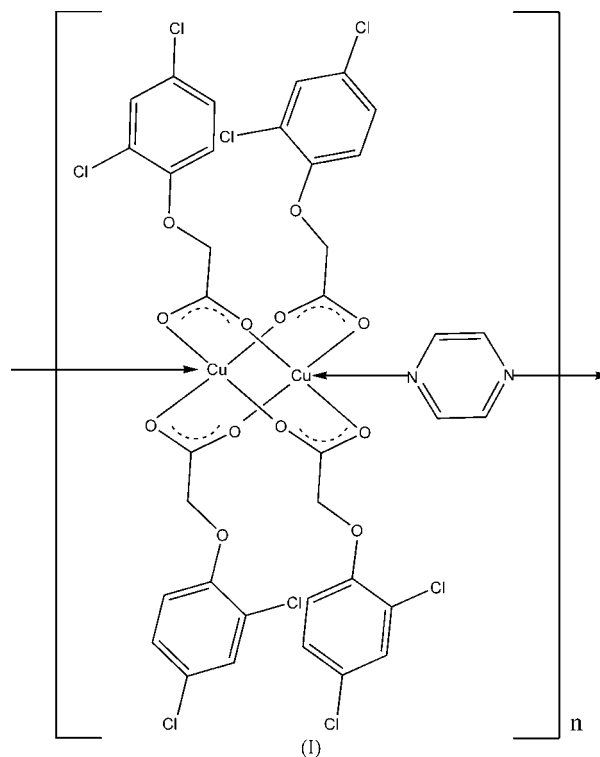


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Key indicators

Single-crystal X-ray study
T = 296 K
Mean $\sigma(\text{C}-\text{C}) = 0.004 \text{ \AA}$
R factor = 0.030
wR factor = 0.085
Data-to-parameter ratio = 12.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.**catena-Poly[[tetrakis(μ -2,4-dichlorophenoxyacetato- $\kappa^2\text{O}:\text{O}'$)dicopper(II)]- μ -pyrazine- $\kappa^2\text{N}:N'$]**The title complex, $[\text{Cu}_2(\text{C}_8\text{H}_5\text{Cl}_2\text{O}_3)_4(\text{C}_4\text{H}_4\text{N}_2)]_n$, crystallizes as a chain of alternating tetrakis(2,4-dichlorophenoxyacetato)dicopper(II) and pyrazine units in which square-pyramidal copper(II) centers are alternately linked by four bridging 2,4-dichlorophenoxyacetate anions and a single bridging pyrazine molecule. Both the dinuclear copper unit and the pyrazine ligand have a center of symmetry.Received 30 November 2006
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Comment

Phenoxyacetic acid and its derivatives are biologically active compounds which are widely used as herbicides and plant-growth substances. Due to their versatile bonding modes with metal ions, they have also been used in the synthesis of metal-organic frameworks (Psomas *et al.*, 2000; Liu *et al.*, 2004; Gao *et al.*, 2005). We report here the crystal structure of a new copper(II) complex, (I), obtained by reacting a copper salt with 2,4-dichlorophenoxyacetic acid and a pyrazine ligand.

As can be seen from Fig. 1, the asymmetric unit comprises one half of the tetrakis(2,4-dichlorophenoxyacetato)dicopper(II) unit, and one half-pyrazine ligand; the complete dinuclear unit and pyrazine are generated by an inversion center. The square-pyramidal coordination geometry of the

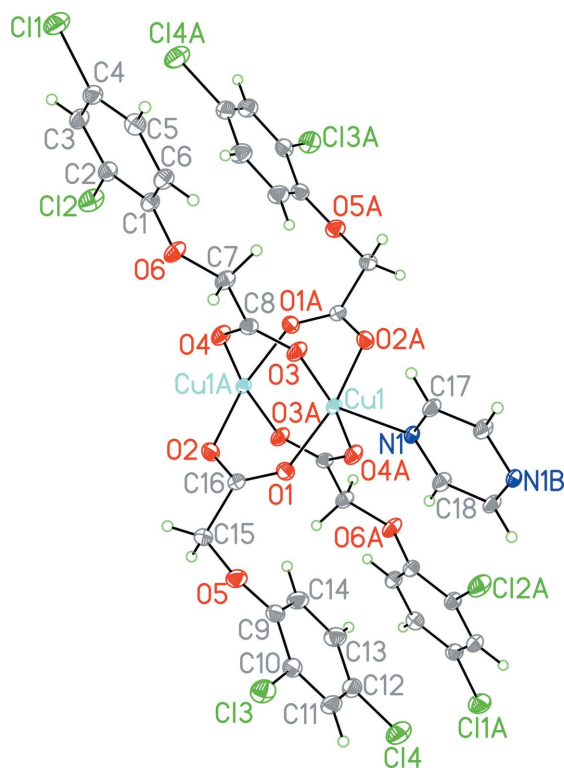


Figure 1
Part of the polymeric structure of the title complex with the atom-labeling scheme. Displacement ellipsoids are drawn at the 35% probability level. [Symmetry codes: (A) $1 - x, 1 - y, 1 - z$; (B) $-x, 1 - y, 1 - z$.]

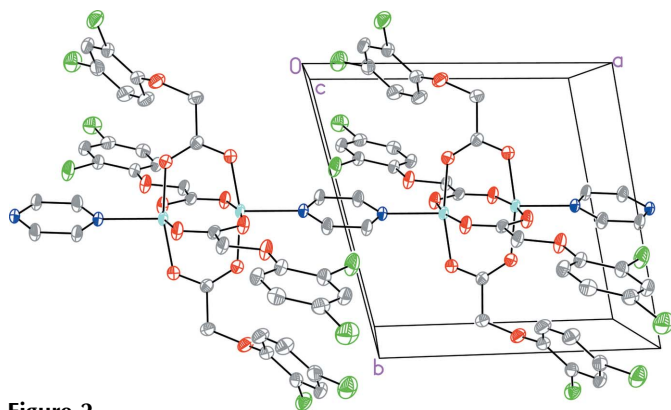


Figure 2
The one-dimensional chain structure of the title complex. H atoms have been omitted for clarity.

copper(II) atom comprises four basal O atoms from four bridging 2,4-dichlorophenoxyacetate anions and an apical N atom from a single bridging pyrazine molecule. The basic architectural motif of the extended structure is a one-dimensional chain of alternating tetrakis(2,4-dichlorophenoxyacetato)dicopper(II) and pyrazine units, as shown in Fig. 2.

Experimental

The title compound was obtained from a solvothermal reaction. Copper(II) nitrate trihydrate (0.06 g, 0.25 mmol), 2,4-dichlorophenoxyacetic acid (0.11 g, 0.5 mmol), pyrazine (0.04 g, 0.5 mmol), and acetonitrile (10 ml) were placed in a Teflon-lined stainless steel

Parr bomb and heated at 353 K for 3 d, and then cooled slowly to room temperature, yielding green single crystals of the title compound.

Crystal data

[Cu₂(C₈H₅Cl₂O₃)₄(C₄H₄N₂)]
 $M_r = 1087.26$
 Triclinic, $P\bar{1}$
 $a = 9.8662(3) \text{ \AA}$
 $b = 10.9610(4) \text{ \AA}$
 $c = 11.0262(3) \text{ \AA}$
 $\alpha = 63.467(2)^\circ$
 $\beta = 78.764(2)^\circ$
 $\gamma = 72.876(2)^\circ$

$V = 1016.73(6) \text{ \AA}^3$
 $Z = 1$
 $D_x = 1.776 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 $\mu = 1.64 \text{ mm}^{-1}$
 $T = 296(2) \text{ K}$
 Block, green
 $0.24 \times 0.14 \times 0.10 \text{ mm}$

Data collection

Bruker SMART APEX-II CCD diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.699, T_{\max} = 0.859$

8012 measured reflections
 3444 independent reflections
 2807 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$
 $\theta_{\max} = 25.0^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.085$
 $S = 1.01$
 3444 reflections
 271 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0528P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.31 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.34 \text{ e \AA}^{-3}$

Table 1

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

Cu1—O2 ⁱ	1.9573 (18)	Cu1—O3	1.9734 (19)
Cu1—O1	1.9713 (17)	Cu1—N1	2.204 (2)
Cu1—O4 ⁱ	1.9734 (19)		
O2 ⁱ —Cu1—O1	166.77 (8)	O4 ⁱ —Cu1—O3	166.87 (8)
O2 ⁱ —Cu1—O4 ⁱ	89.96 (8)	O2 ⁱ —Cu1—N1	101.10 (8)
O1—Cu1—O4 ⁱ	88.11 (8)	O1—Cu1—N1	92.09 (7)
O2 ⁱ —Cu1—O3	89.09 (8)	O4 ⁱ —Cu1—N1	93.73 (8)
O1—Cu1—O3	89.83 (8)	O3—Cu1—N1	99.30 (7)

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

All H atoms were placed in idealized positions and constrained to ride on their parent C atoms, with $C-H = 0.93 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic, and $C-H = 0.97 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for methylene H atoms.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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