Received 2 November 2006

Accepted 15 April 2007

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.006 Å Disorder in solvent or counterion R factor = 0.054 wR factor = 0.108 Data-to-parameter ratio = 14.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. The title compound, $\{[Cu(C_{14}H_{14}N_4)](ClO_4)\}_n$, has been synthesized by the hydrothermal method. The Cu^I atom, lying on a crystallographic twofold rotation axis, is two-coordinate with a linear geometry. The Cl atom also lies on a crystallographic twofold rotation axis. The structure contains one-dimensional chains and a three-dimensional supramolecular network is formed *via* π - π interactions.

methyl)benzene- $\kappa^2 N:N'$] perchlorate]

catena-Poly[[copper(I)-µ-1,4-bis(imidazol-1-yl-

Comment

Copper is an important metal ion in biological systems and it plays diverse roles in living organisms (Holm *et al.*, 1996; Kaim & Rall, 1996). The knowledge of the chemistry of well defined coordinatively unsaturated copper(I) complexes is necessary for the understanding of the reactivity of metals in various biological processes. Hydrothermal/solvothermal reactions under pressure have been used as a synthetic strategy to reduce various metal ions, including copper (Lu & Babb, 2002). The simultaneous reduction of Cu^{2+} to Cu^+ and the formation of a stable linear complex of copper(I) at ambient temperature are thus acomplished. Of these compounds, the two-coordinate copper(I) complexes with nitrogen ligands are scarcely represented in the literature and structurally characterized ones are even rarer (Le Clainche *et al.*, 2000; Liang *et al.*, 2002; Sanyal *et al.*, 1993; Sorell & Jameson, 1983).



Complex (I) consists of chains with alternate Cu^I ions and the bridging ligands. Copper(I) is coordinated by the N atom of each bix ligand unit [bix is 1,4-bis(imidazole-l-ylmethyl)benzene]. The structure is shown in Fig. 1. The geometry of the cationic portion is a linear bix–Cu^I–bix unit. The Cu–N bond length is quite similar to that previously reported for twocoordinate copper(I) complexes with nitrogen donor ligands (Le Clainche et al., 2000; Liang et al., 2002) The dihedral angle between the two imidazole rings coordinated to Cu^I is $73.7 (3)^{\circ}$. All the bix ligands adopt *trans* conformations to form a one-dimensional chain along the b axis, the distance between the Cu ions along the chain being 14.026 (2) Å (Fig. 2). The plane of the imidazole ring is bent relative to the benzene ring, with a dihedral angle between the rings of $80.0 (2)^{\circ}$. It is noteworthy that one imidazole ring interacts with another imidazole ring and a benzene ring belonging to different adjacent chains [symmetry codes: (i) -x, -y, 1 - z;

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metal-organic papers





Figure 1

Part of the polymeric structure of (I). Displacement ellipsoids are drawn at the 30% probability level. H atoms have been omitted for clarity. [Symmetry codes: (A) -x, y, $\frac{3}{2} - z$; (B) $\frac{1}{2} - x$, $-\frac{1}{2} - y$, 1 - z; (C) $\frac{1}{2} + x$, $-\frac{1}{2} - y$, $\frac{1}{2} + z$.]



Figure 2

The one-dimensional chain of (I). H atoms have been omitted.



The three-dimensional supramolecular network formed by π - π stacking interactions (dashed lines).

(ii) $x, -y, \frac{1}{2} + z$] [centroid–centroid distances of 3.8408 (10) and 4.0142 (9) Å, respectively], indicating significant $\pi - \pi$ interaction (Janiak, 2000). As a result, a three-dimensional supra-molecular structure is formed (Fig. 3). Furthermore, packing along the *c* axis, the cavities form channels occupied by perchlorate anions (Fig. 4).

Experimental

Bix dihydrate was prepared as described by Hoskins *et al.* (1997). Compound (I) was hydrothermally synthesized under autogenous pressure. A mixture of Cu(ClO₄)₂·6H₂O (215.2 mg, 0.5 mmol), bix dihydrate (111.2 mg, 0.4 mmol), 2-butynedioic acid (57.1 mg, 0.4 mmol) and H₂O (10 ml) was sealed in a stainless steel reactor with a Teflon liner, which was heated to 433 K for three days. After slow



Figure 4

Packing diagram of (I), showing the channels occupied by perchlorate anions.

cooling to room temperature, white prismatic crystals of (I) were obtained as a major phase by filtration; these were washed with distilled water and finally dried in air.

Crystal data

 $\begin{array}{ll} [\mathrm{Cu}(\mathrm{C}_{14}\mathrm{H}_{14}\mathrm{N}_4)](\mathrm{ClO}_4) & V = 1580.5 \ (7) \ \text{\AA}^3 \\ M_r = 401.28 & Z = 4 \\ \mathrm{Monoclinic}, \ C2/c & \mathrm{Mo} \ K\alpha \ \mathrm{radiation} \\ a = 15.675 \ (3) \ \text{\AA} & \mu = 1.58 \ \mathrm{mm}^{-1} \\ b = 11.791 \ (3) \ \text{\AA} & T = 173 \ (2) \ \mathrm{K} \\ c = 11.316 \ (4) \ \text{\AA} & 0.30 \times 0.25 \times 0.12 \ \mathrm{mm} \\ \beta = 130.913 \ (9)^{\circ} \end{array}$

Data collection

| Rigaku Mercury CCD detector | 5981 measured reflections |
|--|--|
| diffractometer | 1817 independent reflections |
| Absorption correction: multi-scan | 1447 reflections with $I > 2\sigma(I)$ |
| (CrystalClear; Rigaku, 2000) | $R_{\rm int} = 0.025$ |
| $T_{\min} = 0.684, \ T_{\max} = 0.827$ | |

Refinement

| $R[F^2 > 2\sigma(F^2)] = 0.054$ | 128 parameters |
|---------------------------------|--|
| $wR(F^2) = 0.108$ | H-atom parameters constrained |
| S = 1.00 | $\Delta \rho_{\rm max} = 0.42 \ {\rm e} \ {\rm \AA}^{-3}$ |
| 1817 reflections | $\Delta \rho_{\rm min} = -0.36 \text{ e } \text{\AA}^{-3}$ |

Table 1 Selected geometric parameters (Å, °).

| Cu1-N1 | 1.862 (3) | Cu1-N1 ⁱ | 1.862 (3) |
|-------------------------------------|------------------------|-----------------------|------------------------|
| N1-Cu1-N1 ⁱ C2-N1-Cu1 | 178.6 (2) 125.8 (3) | C1-N1-Cu1 N2-C4-C5 | 128.4 (3) 112.4 (3) |
| | | | |

Symmetry code: (i) -x, y, $-z + \frac{3}{2}$.

All H atoms bonded to carbon were placed at calculated positions, refined with isotropic displacement parameters, riding on their parent atoms with C–H = 0.95 and 0.99 Å for the aromatic CH and the CH₂ H atoms, respectively; $U_{iso}(H) = 1.2U_{eq}(C)$. The O atoms are disordered equally about the crystallographic twofold rotation axis passing through Cl.

Data collection: CrystalClear (Rigaku, 2000); cell refinement: CrystalClear; data reduction: CrystalClear; program(s) used to solve

structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000); software used to prepare material for publication: *SHELXTL*.

The work was supported by grants from the State Key Laboratory of Structural Chemistry, Fujian Institute of Research on the Structure of Matter, Chinese Academy of Sciences (CAS), the Ministry of Science and Technology of China (001CB108906), and the National Science Foundation of China (20333070).

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