Acta Crystallographica Section C Crystal Structure Communications

ISSN 0108-2701

# catena-Poly[bis[(2,2-bipyridine- $\kappa^2 N, N'$ )copper(II)]- $\mu_4$ -1,2,4,5-benzene-tetracarboxylato- $\kappa^4 O^1$ : $O^2$ : $O^3$ : $O^4$ ]

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Received 2 December 2003 Accepted 15 December 2003 Online 17 January 2004

In the title complex,  $[Cu_2(C_{10}H_2O_8)(C_{10}H_8N_2)_2]_n$ , the Cu<sup>II</sup> cation has a four-coordinated environment, completed by two carboxyl O atoms belonging to two 1,2,4,5-benzenetetra-carboxylate anions (TCB<sup>4-</sup>) and two N atoms from one 2,2'-bipyridine (2,2'-bipy) ligand, forming a distorted square-planar geometry. The  $[Cu(2,2'-bipy)]^{2+}$  moieties are bridged by TCB<sup>4-</sup> anions, which lie about inversion centres, forming an infinite one-dimensional coordination polymer with a double-chain structure along the *a* axis. A two-dimensional network structure is formed *via* a face-to-face  $\pi$ - $\pi$  interaction between the 2,2'-bipy rings belonging to two adjacent double chains, at a distance of approximately 3.56 Å.

# Comment

In recent years, the design and construction of metal-organic coordination polymers have been extensively studied, due to the crystallographic diversity of these compounds and their potential applications in catalysis, non-linear optics and gas



adsorption (Chen *et al.*, 2003; Eddaoudi *et al.*, 2001; Shi *et al.*, 2003; Stein *et al.*, 1993). In this field, 1,2,4,5-benzenetetracarboxylic acid (H<sub>4</sub>TCB) is a good bridging ligand, and numerous complexes with TCB<sup>4–</sup> anions have been prepared.



#### Figure 1

The coordination environment of the Cu<sup>II</sup> cation in (I), with the atomnumbering scheme, showing the double-chain structure with cavities of approximately 4.5 × 6.1 Å, viewed along the *b* axis [symmetry codes: (i) -x, 1 - y, -z; (ii) x - 1, y, z]. Displacement ellipsoids are drawn at the 50% probability level and H atoms have been omitted for clarity.

The TCB<sup>4-</sup> anions usually coordinate to metal ions in a  $\mu_2$ bridging mode (Hu *et al.*, 2003) and occasionally in a  $\mu_4$ bridging mode (Shi *et al.*, 2001), while in the title complex, [Cu(TCB)<sub>0.5</sub>(2,2'-bpy)]<sub>n</sub>, (I), each TCB<sup>4-</sup> anion binds four Cu<sup>II</sup> ions in a tetra-monodentate mode. Moreover, it should be pointed out that new complexes are constantly being obtained *via* different reaction conditions, such as the use of different solvents, synthesis conditions or H-atom acceptors.

In (I), each Cu<sup>II</sup> cation has a four-coordinated environment, completed by two carboxyl O atoms belonging to two TCB<sup>4–</sup> anions and two N atoms from one 2,2'-bipy ligand (Fig. 1). The bond lengths around the Cu<sup>II</sup> centres are 1.956 (2) Å for Cu1-O3, 1.972 (2) Å for Cu1-O2<sup>i</sup>, 1.978 (2) Å for Cu1-N2 and 1.987 (2) Å for Cu1-N1 [symmetry code: (i) -x, 1 - y, -z]. The N2-Cu1-N1, O3-Cu1-O2<sup>i</sup>, O3-Cu1-N1 and O2<sup>i</sup>-Cu1-N2 bond angles are in the range 81.80 (10)– 98.33 (10)°, significantly different from the ideal value.

From these bond lengths and angles, we can best describe the coordination of the Cu<sup>II</sup> cation as a distorted squareplanar geometry. Moreover, the coordination mode of the TCB<sup>4-</sup> anion is similar to that in [Cu<sub>2</sub>(TCB)(phen)<sub>2</sub>]<sub>n</sub>·nH<sub>2</sub>O (Shi *et al.*, 2001); the four carboxylate groups are all deprotonated and coordinate to four Cu<sup>II</sup> cations in a monodentate fashion, forming an infinite one-dimensional ribbon-like double-chain structure with cavities of approximately  $4.5 \times 6.1$  Å along the *a* axis (Fig. 1). The Cu1···Cu1<sup>i</sup> and Cu1<sup>i</sup>···Cu1<sup>ii</sup> separation distances are 7.931 and 7.279 Å, respectively [symmetry code: (ii) x - 1, y, z]. Furthermore, there are  $\pi$ - $\pi$  interactions between the aromatic rings of 2,2'bipy ligands belonging to two adjacent double chains, at a distance of approximately 3.56 Å, resulting in a two-dimensional network structure (Fig. 2).



Figure 2

A view of the two-dimensional network structure in (I), formed by  $\pi$ - $\pi$  interactions.

# **Experimental**

A solution (10 ml) of dimethylformamide containing CuCl<sub>2</sub>·2H<sub>2</sub>O (0.5 mmol, 0.085 g) and H<sub>4</sub>TCB (0.5 mmol, 0.127 g) was added slowly to a solution (10 ml) of dimethylformamide containing 2,2'-bipy (0.5 mmol, 0.078 g). The mixture was stirred for 30 min and then left to stand at room temperature for about a month. Light-blue prism-shaped crystals of (I) were obtained.

#### Crystal data

 $\begin{bmatrix} Cu_2(C_{10}H_2O_8)(C_{10}H_8N_2)_2 \end{bmatrix}$   $M_r = 689.56$ Monoclinic,  $P2_1/n$  a = 7.2794 (7) Å b = 12.5009 (12) Å c = 14.4436 (14) Å  $\beta = 104.021$  (2)° V = 1275.2 (2) Å<sup>3</sup> Z = 2

#### Cell parameters from 532 reflections $\theta = 2.4-22.0^{\circ}$ $\mu = 1.73 \text{ mm}^{-1}$ T = 273 (2) K Prism, blue $0.32 \times 0.29 \times 0.24 \text{ mm}$

 $D_{\rm r} = 1.796 {\rm Mg m}^{-3}$ 

Mo  $K\alpha$  radiation

#### Data collection

Bruker SMART CCD area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2000)
$T_{\min} = 0.582, \ T_{\max} = 0.663$
9055 measured reflections

2263 independent reflections 2179 reflections with  $I > 2\sigma(I)$  $R_{int} = 0.024$  $\theta_{max} = 25.1^{\circ}$  $h = -8 \rightarrow 8$  $k = -14 \rightarrow 14$  $l = -17 \rightarrow 17$ 

#### Refinement

Refinement on $F^2$
$R[F^2 > 2\sigma(F^2)] = 0.036$
$wR(F^2) = 0.087$
S = 1.16
2263 reflections
200 parameters
H-atom parameters constrained

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0367P)^2 \\ &+ 1.7086P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.48 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.29 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

All H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of 0.93 Å, with  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$  (parent atom).

Data collection: *SMART* (Bruker, 2000); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve structure: *SHELXTL* (Bruker, 2000); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

The authors acknowledge financial support from the Zhejiang Provincial Natural Science Foundation of China (grant No. 202137).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SQ1144). Services for accessing these data are described at the back of the journal.

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