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Key indicators

Single-crystal X-ray study T = 295 K Mean σ (C–C) = 0.003 Å R factor = 0.043 wR factor = 0.127 Data-to-parameter ratio = 15.6

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

catena-Poly[[[(4'-phenyl-2,2':6',2''-terpyridine- $\kappa^3 N$)copper(II)]- μ -terephthalato- $\kappa^2 O$:O'] terephthalic acid]

The title compound, $[Cu(C_8H_4O_4)(C_{21}H_{15}N_3)]\cdot C_8H_6O_4$, consists of polymeric chains in which the terpyridine heterocycle chelates to the Cu atom in an N,N',N''-tridentate manner; the Cu atoms are also covalently bonded through the two singly-bonded carboxyl O atoms of the dicarboxylate dianions to give rise to a chain structure. An additional terephthalic acid molecule is hydrogen bonded to the chains. The asymmetric unit contains halves of two different halfmolecule terephthalate ions; in each case, the complete ions are generated by inversion symmetry.

Comment

The terephthalate (tp) ion, $C_8H_4O_4^{2-}$, has been widely used as a bridging ligand in the syntheses of coordination polymers (Xanthopoulos *et al.*, 1993; Bakalbassis *et al.*, 1986, 1988; Verdaguer *et al.*, 1984). In most cases, the dianion acts as a monodentate or bidentate ligand (Cano *et al.*, 1997, and references therein; Fun *et al.*, 1999). It has seldom been found as a neutral (protonated) filler in the crystal packing.



We describe here a zigzag polymeric complex, $[Cu(C_8H_4O_4)(C_{21}H_{15}N_3)]\cdot C_8H_6O_4$, (I), that was obtained by the hydrothermal reaction of copper(II) chloride with terpyridine and terephthalic acid (H₂tp), instead of a binuclear complex as reported by Sun *et al.* (2001). The Cu ions in (I) possess N₃O₂ coordination spheres (Table 1). If a very long bond to O2 [d = 2.801 (2) Å] was considered to be significant, a distorted CuN₃O₃ octahedron would result. According to the structural index τ (Atakol *et al..*, 1999), the coordination geometry about Cu approximates to distorted square-pyramidal, with atom O3 occupying the apex and atoms N1, N2, N3 and O1 forming the basal plane (Fig. 1). In the crystal packing,

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metal-organic papers



Figure 1

View of a fragment of (I), with displacement ellipsoids drawn at the 50% probability level and H atoms drawn as spheres of arbitrary radii. The complete tp species containing O1 is generated by the symmetry code (1 - x, 1 - y, 1 - z), and that containing O3 by (1 - x, -y, -z).



Figure 2 The polymeric chain in (I), with Cu atoms coloured light blue.

each tp ions acts as a doubly monodentate bridging ligand and binds with two copper ions to form a zigzag chain, \cdots Cu-tp-Cu-tp \cdots , with the O1 and O3 tp species alternating. Terpyridine acts as a tridentate ligand to complete the coordination about each Cu atom (Fig. 2). Interestingly, a molecule of H2tp was also found in the structure of (I). It acts as a bridge to support the zigzag chains with hydrogen bonds (Table 2) to carbonyl O atoms of coordinated tp ions, which gives rise to a sheet-like network (Fig. 3).

Experimental

To an aqueous solution (15 ml) of copper(II) chloride dihydrate (0.10 g, 0.50 mmol) was added 4'-phenyl-2,2':6',2"-terpyridine [0.16 g, 0.50 mmol, synthesized according to a literature procedure (Constable *et al.*, 1990)], followed by terephthalic acid (0.04 g, 0.25 mmol) and piperidine (0.06 g, 0.5 mmol). The mixture was placed in a 25 ml Teflon-lined hydrothermal bomb and heated at 423 K for 40 h. The bomb was then cooled to room temperature at about 5 K h⁻¹. Green prismatic crystals of (I) were obtained from the cooled solution. Analysis calculated for $C_{37}H_{25}CuN_3O_8$: C 63.20, N



Figure 3

Detail of the hydrogen bonding between terephthalic acid and the terephthalate dianions in (I).

3.58, H 5.98%; found: C 63.21, H 3.32, N 6.01%. IR (KBr, cm⁻¹): 3432 (*br s*), 3060 (*w*), 2349 (*w*), 1699 (*w*), 1601 (*vs*), 1474 (s), 1417 (s), 1352 (s), 1247 (*w*), 1021 (*m*), 824 (*w*), 771 (*m*).

Crystal data

Data collection

Bruker APEX area-dectector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{\min} = 0.707, T_{\max} = 0.920$ 13 572 measured reflections

Refinement

Refinement on F^2

 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.127$ S = 1.087024 reflections 450 parameters

H atoms treated by a mixture of independent and constrained refinement

Z = 2 $D_x = 1.510 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 4160 reflections $\theta = 2.2 - 26.0^{\circ}$ $\mu = 0.77 \text{ mm}^{-1}$ T = 295 (2) KPrism, green $0.26 \times 0.18 \times 0.11 \text{ mm}$

7024 independent reflections 5819 reflections with $I > 2\sigma(I)$ $R_{int} = 0.020$ $\theta_{max} = 27.9^{\circ}$ $h = -10 \rightarrow 10$ $k = -18 \rightarrow 18$ $l = -17 \rightarrow 18$

$$\begin{split} &w = 1/[\sigma^2(F_{\rm o}^2) + (0.0744P)^2 \\ &+ 0.1147P] \\ &where \ P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} = 0.001 \\ \Delta\rho_{\rm max} = 0.61 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.47 \ {\rm e} \ {\rm \AA}^{-3} \end{split}$$

Ta	ab	le	1			
0						

Selected geometri	c parameters	(A,	°).	
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Cu1-O1	1.901 (2)	Cu1-N2	1.927 (2)
Cu1-O3	2.280 (2)	Cu1-N3	2.018 (2)
Cu1-N1	2.047 (2)		
O1-Cu1-O3	89.94 (6)	O3-Cu1-N2	97.43 (7)
O1-Cu1-N1	99.11 (8)	O3-Cu1-N3	101.00 (7)
O1-Cu1-N2	172.45 (7)	N1-Cu1-N2	79.65 (7)
O1-Cu1-N3	100.18 (8)	N1-Cu1-N3	158.81 (8)
O3-Cu1-N1	87.69 (7)	N2-Cu1-N3	80.07 (7)

Table 2

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

	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{matrix} O5-H5O\cdots O4\\ O7-H7O\cdots O3^i \end{matrix}$	0.85(1)	1.75 (1)	2.594 (2)	176 (4)
	0.84(1)	1.82 (1)	2.666 (2)	177 (4)

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

Hydrogen atoms bonded to C atoms were placed in calculated positions (C-H = 0.93 Å) and were included in the refinement as riding $[U_{iso}(H) = 1.2U_{eq}C)]$. H atoms of the carboxylic acid (H5O and H7O) were located in a difference map and refined freely.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine

structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL*97.

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