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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.009 \AA$
Disorder in main residue
$R$ factor $=0.052$
$w R$ factor $=0.108$
Data-to-parameter ratio $=12.0$

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## Poly[[[[(2,2'-bipyridine)copper(II)]-hemi- $\mu_{2}$-2-nitro-terephthalato-hemi- $\mu_{4}$-2-nitroterephthalato] monohydrate]

The polymeric title compound, $\left\{\left[\mathrm{Cu}\left(\mathrm{C}_{8} \mathrm{H}_{3} \mathrm{NO}_{6}\right)\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right)\right]\right.$-$\left.\mathrm{H}_{2} \mathrm{O}\right\}_{n}$, has two disordered nitroterephthalate units located around centres of inversion; one functions in a $\mu_{2}$-bridging mode and the other in a $\mu_{4}$-bridging mode to surround the five-coordinated Cu atom in a square-pyramidal geometry. The crystal structure reveals a two-dimensional hydrogenbonded network involving a water molecule of crystallization.

## Comment

A previous study of copper 2-nitroterephthalate reported the inorganic-organic hybrid structure (Ma \& Zhu, 2004); the compound exhibits a square-pyramidal coordination involving bridging dicarboxylate and hydroxyl groups, with the apical site of the pyramid occupied by a water molecule. The crystal structure is a three-dimensional network.

(I)

However, with the chelating $2,2^{\prime}$-bipyridine heterocycle, copper 2-nitroterephthalate adopts a only two-dimensional structure and does not include a coordinated water molecule, (I) (Fig. 1). The two symmetry-independent dianionic groups are each disordered over an inversion centre. One of the groups functions in the $\mu_{2}$-bridging mode and the other in the $\mu_{4}$-bridging mode to surround the five-coordinated Cu atom. The apical site of the square pyramid is occupied by the O atom of the $\mu_{4}$-bridging dianion.

## Experimental

A mixture of copper chloride dihydrate $(0.035 \mathrm{~g}, 0.2 \mathrm{mmol})$, 2-nitroterephthalic acid $(0.021 \mathrm{~g}, 0.1 \mathrm{mmol}), 2,2^{\prime}$-bipyridine ( 0.032 g , $0.2 \mathrm{mmol})$, sodium hydroxide $(0.008 \mathrm{~g}, 0.2 \mathrm{mmol})$ and water $(8 \mathrm{ml})$ was sealed in a 25 ml stainless steel Teflon-lined bomb. The bomb was heated at 423 K for 3 d . After cooling, deep-blue block-shaped crystals of (I) were obtained.

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## Crystal data

$\left[\mathrm{Cu}\left(\mathrm{C}_{8} \mathrm{H}_{3} \mathrm{NO}_{6}\right)\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right)\right] \cdot \mathrm{H}_{2} \mathrm{O}$
$M_{r}=446.85$
Triclinic, $P \overline{1}$
$a=9.537$ (1) $\AA$
$b=9.950$ (1) $\AA$
$c=11.363$ (1) $\AA$
$\alpha=94.156(2)^{\circ}$
$\beta=107.839(2)^{\circ}$
$\gamma=113.535(2)^{\circ}$
$V=917.31(16) \AA^{3}$

## Data collection

Bruker SMART APEX areadetector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\text {min }}=0.764, T_{\text {max }}=0.817$
5077 measured reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.052$
$w R\left(F^{2}\right)=0.108$
$S=0.91$
3534 reflections
295 parameters
$Z=2$
$D_{x}=1.618 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 963
reflections
$\theta=2.3-20.4^{\circ}$
$\mu=1.24 \mathrm{~mm}^{-1}$
$T=295$ (2) K
Block, blue
$0.23 \times 0.19 \times 0.17 \mathrm{~mm}$

3534 independent reflections
2227 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.048$
$\theta_{\text {max }}=26.0^{\circ}$
$h=-11 \rightarrow 11$
$k=-12 \rightarrow 12$
$l=-14 \rightarrow 11$

H atoms treated by a mixture of independent and constrained refinement
$w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0309 P)^{2}\right]$
where $P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }=0.001$
$\Delta \rho_{\text {max }}=0.40 \mathrm{e}^{-3}$
$\Delta \rho_{\min }=-0.41 \mathrm{e}^{-3}$

Table 1
Selected geometric parameters ( $\left({ }^{\circ},{ }^{\circ}\right)$.

| $\mathrm{Cu} 1-\mathrm{O} 5$ | $1.949(3)$ | $\mathrm{Cu} 1-\mathrm{N} 2$ | $1.996(4)$ |
| :--- | ---: | :--- | ---: |
| $\mathrm{Cu} 1-\mathrm{O} 1$ | $1.955(3)$ | $\mathrm{Cu} 1-\mathrm{O} 5^{\mathrm{i}}$ | $2.373(3)$ |
| $\mathrm{Cu} 1-\mathrm{N} 1$ | $1.994(4)$ |  |  |
| $\mathrm{O} 5-\mathrm{Cu} 1-\mathrm{O} 1$ | $88.80(12)$ | $\mathrm{N} 1-\mathrm{Cu} 1-\mathrm{N} 2$ | $80.84(17)$ |
| $\mathrm{O} 5-\mathrm{Cu} 1-\mathrm{N} 1$ | $178.37(15)$ | $\mathrm{O} 5-\mathrm{Cu} 1-\mathrm{O} 5^{\mathrm{i}}$ | $77.27(12)$ |
| $\mathrm{O} 1-\mathrm{Cu} 1-\mathrm{N} 1$ | $91.64(15)$ | $\mathrm{O} 1-\mathrm{Cu} 1-\mathrm{O} 5^{\mathrm{i}}$ | $89.05(13)$ |
| $\mathrm{O} 5-\mathrm{Cu} 1-\mathrm{N} 2$ | $98.57(14)$ | $\mathrm{N} 1-\mathrm{Cu} 1-\mathrm{O} 5^{\mathrm{i}}$ | $104.30(13)$ |
| $\mathrm{O} 1-\mathrm{Cu} 1-\mathrm{N} 2$ | $170.92(15)$ | $\mathrm{N} 2-\mathrm{Cu} 1-\mathrm{O} 5^{\mathrm{i}}$ | $97.70(13)$ |

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

Table 2
Hydrogen-bonding geometry $\left(\AA^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| O1W-H1W1 $\cdots$ O2 | $0.85(6)$ | $1.98(6)$ | $2.815(6)$ | $169(6)$ |
| O1 $W-\mathrm{H} 1 W 2 \cdots 6^{\mathrm{ii}}$ | $0.85(5)$ | $2.09(4)$ | $2.879(7)$ | $155(8)$ |

Symmetry code: (ii) $-x,-y, 1-z$.
The aromatic H atoms were positioned geometrically, and were included in the refinement in the riding-model approximation $[\mathrm{C}-\mathrm{H}$ $=0.93 \AA$ and $\left.U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})\right]$. The water H atoms were located


Figure 1
ORTEPII (Johnson, 1976) plot of a portion of the title compound. Displacement ellipsoids are drawn at the $50 \%$ probability level. The water molecules are not shown. [Symmetry code: (i) $1-x, 1-y, 1-z$.]
in a difference Fourier map and were refined with distance restraints of $\mathrm{O}-\mathrm{H}=0.85(1) \AA$ and $\mathrm{H} \cdots \mathrm{H}=1.39(1) \AA$, and $U_{\text {iso }}(\mathrm{H})=$ $1.2 U_{\text {eq }}(\mathrm{O})$. The two nitro groups are disordered with respect to H atoms; the $\mathrm{C}-\mathrm{N}$ distances were restrained to 1.470 (5) $\AA$, a tighter restraint being used to avoid too large a difference in distances. Additionally, the displacement parameters of the O atoms of the nitro groups were restrained to be nearly isotropic.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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