

Bis(pyridine-2,4-dicarboxylato)copper(II) dihydrate

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Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$
 R factor = 0.036
 wR factor = 0.098
Data-to-parameter ratio = 10.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

A mononuclear copper(II) coordination complex, $[\text{Cu}(\text{Hpydc})_2] \cdot 2\text{H}_2\text{O}$ (pydc = 2,4-pyridinedicarboxylate, $\text{C}_7\text{H}_4\text{NO}_4$), has been prepared from the hydrothermal reaction of CuO and H_2pydc in H_2O . The complex consists of discrete $[\text{Cu}(\text{Hpydc})_2]$ and two uncoordinated water molecules, the copper(II) displaying a typical planar four-coordinate geometry. All $\text{Cu}-\text{O}$ and $\text{Cu}-\text{N}$ distances range from 1.945 (2) to 1.975 (2) \AA . The discrete structure is further extended into a three-dimensional structure by weak hydrogen bonds.

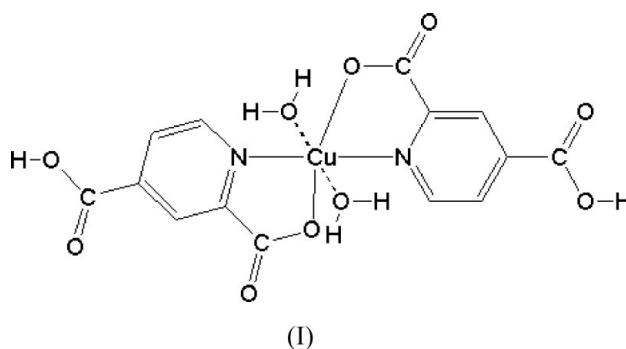
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Comment

Research into transition metal complexes has been rapidly expanding because of their fascinating structural diversity and potential applications as functional materials and enzymes. The key step in the design of polymeric transition metal complexes is to select suitable multidentate bridging ligands containing certain features, such as flexibility, versatile binding modes and the ability to form hydrogen bonds. With these criteria in mind, we chose pyridine-2,5-dicarboxylic acid as ligand. It reacts with transition metals or rare earths; a series of novel complexes having infinite or discrete structures has been obtained and reported (Liang *et al.*, 2000, 2001). In this paper, we report the synthesis and crystal structure of a mononuclear copper(II) compound, (I), *i.e.* $[\text{Cu}(\text{Hpydc})_2] \cdot 2\text{H}_2\text{O}$, (I), where pydc is pyridine-2,4-dicarboxylate.



The coordination complex $[\text{Cu}(\text{Hpydc})_2] \cdot 2\text{H}_2\text{O}$ was prepared by the hydrothermal reaction of CuO and H_2pydc in H_2O . The crystallographic analysis reveals that the compound is a discrete $[\text{Cu}(\text{Hpydc})_2]$ molecule, in which the Cu^{II} atom is coordinated by the two N and two O atoms of two Hpydc ligands to form a distorted planar four-coordinate geometry, as shown in Fig. 1. There is a weak interaction between the Cu and hydrate O atoms, with an average $\text{Cu}-\text{O}$ distance of 2.602 \AA . Other $\text{Cu}-\text{O}$ and $\text{Cu}-\text{N}$ distances are listed in Table

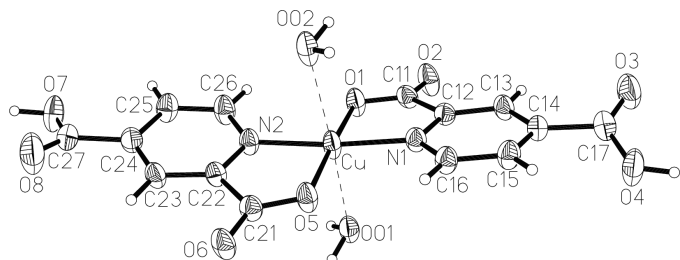


Figure 1
The structure of $[\text{Cu}(\text{Hpydc})_2] \cdot 2\text{H}_2\text{O}$. Displacement ellipsoids are plotted at the 50% probability level.

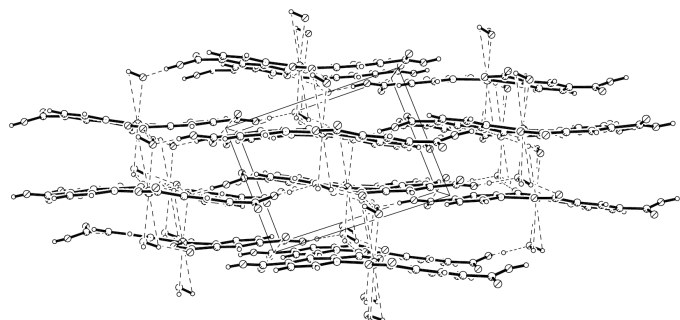


Figure 2
Packing diagram of $[\text{Cu}(\text{Hpydc})_2] \cdot 2\text{H}_2\text{O}$. Dashed lines indicate the hydrogen bonds.

1, as are N—Cu—N, O—Cu—O and O—Cu—N angles. The molecules are connected by hydrogen-bonding interactions between the carboxylate groups and H_2O , with an average O...O distance of 2.675 Å, to form a three-dimensional network, as shown in Fig. 2.

Experimental

A mixture of CuO (0.080 g, 2.0 mmol), $[\text{H}_2\text{pydc}] \cdot \text{H}_2\text{O}$ (0.185 g, 1.0 mmol) and H_2O (16 ml) in a molar ratio of ca 2:1:890 was sealed in a 25 ml stainless-steel reactor with Teflon liner. The reaction system was heated at 443 K for 72 h. Slowly cooling the system to room temperature yielded needle-like green crystals of the complex and some blue-green precipitate. The organic H atoms were positioned geometrically (C—H bond fixed at 0.96 Å) and allowed to ride on their parent C atoms before the final cycle of refinement. The hydrate H atoms were located from difference maps and refined with isotropic displacement parameters.

Crystal data

$[\text{Cu}(\text{C}_7\text{H}_4\text{NO}_4)_2] \cdot 2\text{H}_2\text{O}$

$M_r = 431.80$

Triclinic, $P\bar{1}$

$a = 7.212(3) \text{ \AA}$

$b = 10.139(3) \text{ \AA}$

$c = 11.337(5) \text{ \AA}$

$\alpha = 73.67(3)^\circ$

$\beta = 82.67(3)^\circ$

$\gamma = 85.04(3)^\circ$

$V = 787.8(5) \text{ \AA}^3$

$Z = 2$

$D_x = 1.820 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation

Cell parameters from 4087

reflections

$\theta = 1.9\text{--}25.0^\circ$

$\mu = 1.45 \text{ mm}^{-1}$

$T = 293(2) \text{ K}$

Needle, green

$0.42 \times 0.10 \times 0.04 \text{ mm}$

Data collection

SMART CCD diffractometer

ω scans

Absorption correction: empirical
(SADABS; Sheldrick, 1996)

$T_{\min} = 0.586$, $T_{\max} = 0.944$

4087 measured reflections

2754 independent reflections

2345 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.017$

$\theta_{\max} = 25.0^\circ$

$h = -8 \rightarrow 6$

$k = -11 \rightarrow 12$

$l = -13 \rightarrow 13$

Refinement

Refinement on F^2

$R[F^2 > 2\sigma(F^2)] = 0.036$

$wR(F^2) = 0.098$

$S = 1.04$

2754 reflections

268 parameters

H atoms treated by a mixture of
independent and constrained
refinement

$w = 1/[\sigma^2(F_o^2) + (0.0527P)^2$
 $+ 0.4456P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.61 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.33 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cu—O5	1.945 (2)	O2—O01 ⁱⁱ	2.736 (3)
Cu—O1	1.948 (2)	O4—O02 ⁱⁱⁱ	2.575 (3)
Cu—N2	1.972 (2)	O6—O02 ^{iv}	2.721 (3)
Cu—N1	1.975 (2)	O6—O01 ^v	2.764 (3)
O2—O02 ⁱ	2.716 (3)	O7—O01 ^{vi}	2.600 (3)
O5—Cu—O1	174.94 (9)	C11—O1—Cu	114.43 (17)
O5—Cu—N2	83.63 (9)	C21—O5—Cu	114.81 (17)
O1—Cu—N2	96.47 (9)	C16—N1—Cu	128.61 (19)
O5—Cu—N1	97.05 (9)	C12—N1—Cu	111.75 (18)
O1—Cu—N1	83.40 (9)	C26—N2—Cu	128.11 (19)
N2—Cu—N1	173.73 (10)	C22—N2—Cu	111.84 (18)

Symmetry codes: (i) $3-x, 3-y, 1-z$; (ii) $2-x, 3-y, 1-z$; (iii) $x, 1+y, z$; (iv) $3-x, 3-y, -z$; (v) $2-x, 3-y, -z$; (vi) $x, y-1, z$.

The organic H atoms were positioned geometrically (C—H bond fixed at 0.96 Å) and allowed to ride on their parent C atoms before the final cycle of refinement. The hydrate H atoms were located from difference maps and refined with isotropic displacement parameters.

Data collection: SMART (Siemens, 1994); cell refinement: SMART and SAINT (Siemens, 1994); data reduction: XPREP (Siemens, 1994); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXL97; software used to prepare material for publication: SHELXL97.

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