

Chuan-Bi Li* and Ying-Chun Gao

College of Science, Changchun University,
Changchun 130022, People's Republic of China

Correspondence e-mail: chuanbl@gmail.com

Key indicators

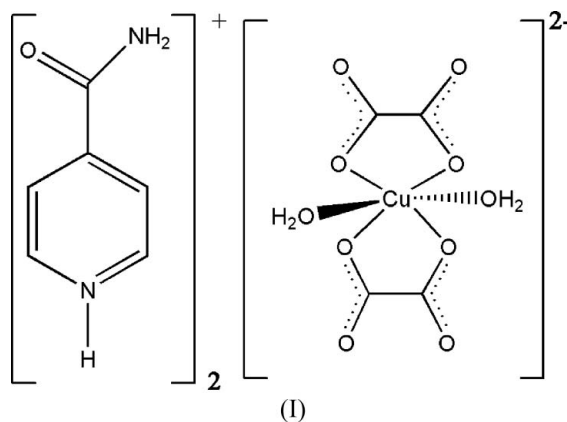
Single-crystal X-ray study
 $T = 295$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.025
 wR factor = 0.067
Data-to-parameter ratio = 12.5For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis(4-carboxamidopyridinium) diaqua-
dioxalatocuprate(II)

In the title compound, $(\text{C}_6\text{H}_7\text{N}_2\text{O})_2[\text{Cu}(\text{C}_2\text{O}_4)_2(\text{H}_2\text{O})_2]$, the Cu^{II} center (site symmetry $\bar{1}$) exhibits an octahedral coordination geometry arising from four O atoms of two oxalate dianions and two O atoms of water molecules. The protonated isonicotinamide cation interacts with the anion by $\text{O}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds, generating a three-dimensional network.

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Comment

To extend our work (Gao *et al.*, 2006) on hydrogen-bonded supramolecular architectures (Eddaoudi *et al.*, 2002), we obtained the title compound, (I), by choosing oxalate dianions, isonicotinamide and Cu^{2+} cations as the starting materials.



Compound (I) contains $[\text{Cu}(\text{C}_2\text{O}_4)_2(\text{H}_2\text{O})_2]^{2-}$ dianions and $[\text{C}_6\text{H}_7\text{N}_2\text{O}]^+$ cations, as shown in Fig. 1. The Cu^{II} atom (site symmetry $\bar{1}$) shows a distorted octahedral coordination geometry, being coordinated by four O atoms from two oxalate ligands and two O atoms from water molecules, with the $\text{Cu}-\text{O}$ distances showing a typical Jahn–Teller distortion (Table 1).

The monoprotonated (at the pyridyl N atom) isonicotinamide cations balance the charge of the anionic complex ion. $\text{O}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen-bonding interactions (Table 2) play an important role in stabilizing the network in the crystal structure of (I) (Fig. 2), resulting in a three-dimensional supramolecular network.

Experimental

A mixture of CuSO_4 (0.5 mmol), $\text{K}_2\text{C}_2\text{O}_4\cdot\text{H}_2\text{O}$ (0.5 mmol) and isonicotinamide (1 mmol) was heated in 20 ml of water/ethanol (1:1 v/v) solution and continually stirred for about 30 min at 333 K. The pH value was adjusted to about 4.0 with 4 M HCl. The mixture was

then cooled to room temperature and filtered. The filtrate was allowed to evaporate at room temperature. One week later, aquamarine-blue crystals of (I) were obtained.

Crystal data

$(C_6H_7N_2O)_2[Cu(C_2O_4)_2(H_2O)_2]$	$\gamma = 79.483 (3)^\circ$
$M_r = 521.89$	$V = 476.9 (2) \text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 1$
$a = 6.788 (2) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 6.957 (2) \text{ \AA}$	$\mu = 1.23 \text{ mm}^{-1}$
$c = 10.849 (3) \text{ \AA}$	$T = 295 (2) \text{ K}$
$\alpha = 76.127 (2)^\circ$	$0.43 \times 0.32 \times 0.24 \text{ mm}$
$\beta = 75.198 (2)^\circ$	

Data collection

Bruker SMART CCD APEX II diffractometer	4061 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2002)	1887 independent reflections
$T_{\min} = 0.635, T_{\max} = 0.750$	1822 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.012$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.025$	151 parameters
$wR(F^2) = 0.067$	H-atom parameters constrained
$S = 1.00$	$\Delta\rho_{\max} = 0.25 \text{ e \AA}^{-3}$
1887 reflections	$\Delta\rho_{\min} = -0.46 \text{ e \AA}^{-3}$

Table 1

Selected bond lengths (\AA).

Cu—O1	1.9428 (13)	Cu—OW1	2.4872 (16)
Cu—O2	1.9873 (13)		

Table 2

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
OW1—HW1 \cdots O5 ^v	0.84	1.97	2.7959 (19)	170
OW1—HW2 \cdots O4 ⁱⁱ	0.79	1.94	2.733 (2)	175
N1—H1 \cdots O3 ⁱⁱⁱ	0.86	1.91	2.675 (2)	148
N1—H1 \cdots O4 ⁱⁱⁱ	0.86	2.31	2.963 (2)	132
N2—HN1 \cdots OW1 ^{iv}	0.86	2.04	2.837 (2)	154
N2—HN2 \cdots O2 ⁱ	0.86	2.31	3.078 (2)	149

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

The water H atoms were located in difference maps and refined as riding in their as-found relative positions ($O-H = 0.79-0.84 \text{ \AA}$), with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(O)$. The other H atoms were positioned geometrically and refined with a riding model ($N-H = 0.86 \text{ \AA}$ and $C-H = 0.93 \text{ \AA}$), with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C, N)$.

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:

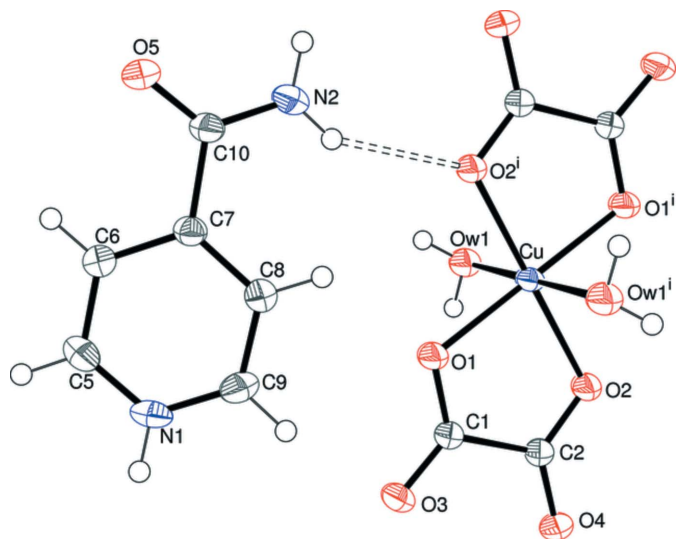


Figure 1

The structure of (I), showing 30% probability displacement ellipsoids (arbitrary spheres for the H atoms). The hydrogen bond is shown as a double dashed line. [Symmetry code: (i) $1-x, 1-y, 1-z$.]

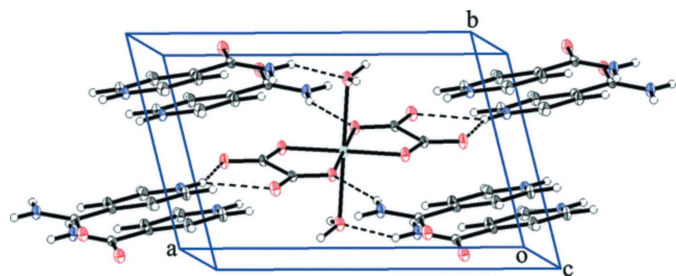


Figure 2

The unit cell packing for (I), showing the hydrogen bonds as dashed lines.

DIAMOND (Brandenburg, 1999); software used to prepare material for publication: SHELXL97.

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