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Key indicators

Single-crystal X-ray study T = 273 K Mean σ (C–C) = 0.003 Å R factor = 0.023 wR factor = 0.063 Data-to-parameter ratio = 9.6

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

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Aqua(L-aspartato- $\kappa^2 N$,O)(2,2'-bipyridine- $\kappa^2 N$,N)copper(II) pentahydrate

In the title compound, $[Cu(C_4H_5NO_4)(C_{10}H_8N_2)(H_2O)]$. 5H₂O, the Cu^{II} atom is coordinated by an *N*,*O*-bidentate Laspartate dianion, a bipyridine ligand and a water molecule in a distorted square-pyramidal geometry. Intermolecular hydrogen bonds involving the non-coordinated water molecules help to consolidate the crystal packing.

Comment

There is great interest in the ability of amino acids to coordinate to metal ions containing hydrogen bonds in their complexes (Hu *et al.*, 2004). As a binary amino acid, aspartic acid is especially important from both a biological and a simple coordination point of view (Casellato *et al.*, 1991; Ma *et al.*, 2004). By hydrothermal synthesis, an L-aspartate–copper complex [Cu(L-asp)(bipy)(H₂O)]·5H₂O (L-asp is L-aspartate and bipy is 2,2'-bipyridine), (I), was prepared and its crystal structure is presented here. The asymmetric unit of (I) consists of a Cu^{II} ion, an L-aspartate dianion, a bipyridine ligand, a coordinated water and five uncoordinated water molecules.



The molecular structure of (I) is shown in Fig. 1. The coordination geometry around the Cu^{II} atom is five-coordinate distorted square-pyramidal. The L-aspartate dianion (O1/N3) and bipyridine ligand (N1/N2) lie on the basal plane and coordinate to the Cu^{II} ion, forming two five-membered chelate rings. The apical position is occupied by a coordinated water molecule (O5), with a Cu1-O5 distance of 2.2840 (15) Å (Table 1), slightly shorter than the coordinated water bond length (2.368 Å) of the analogous complex [Cu(L-asp)(bipy)(H₂O)]·3H₂O (Brondino *et al.*, 1995; Antolini *et al.*, 1983).

An intramolecular hydrogen bond exists between O3 and N3 (Table 2), which causes the uncoordinated carboxylate group to bend markedly towards the amino group. The uncoordinated carboxylate O2 atom links to an uncoordinated water molecule, while the other uncoordinated carboxylate O atom (O4) links to a neighboring L-aspartate ligand *via* an N3-H12 \cdots O4 hydrogen bond (Fig. 2).

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Figure 1

The structure of (I). Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

A crystal packing diagram of (I), with intermolecular hydrogen bonds involving the uncoordinated water molecules shown as dashed lines.

Experimental

The title complex was synthesized by a hydrothermal synthesis method from a mixture of L-aspartic acid (0.399 g, 3.0 mmol), CuCO₃ (0.240 g, 2.0 mmol), 2,2'-bipyridine (0.156 g, 1.0 mmol) and water (30 ml) in a 40 ml Teflon-lined stainless steel reactor. The mixture was heated at 393 K for 2 d. After cooling to room temperature, darkblue single crystals of (I) were obtained.

Crystal data

10240 (1) 13
$V = 1934.9(1) \text{ A}^3$
Z = 4
$D_x = 1.575 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
$\mu = 1.19 \text{ mm}^{-1}$
T = 273 (2) K
Block, blue
$0.20 \times 0.10 \times 0.08 \text{ mm}$

21692 measured reflections

 $R_{\rm int} = 0.022$

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

3404 independent reflections

 $w = 1/[\sigma^2(F_0^2) + (0.0343P)^2]$

+ 0.7615P] where $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.25 \text{ e } \text{\AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.24 \text{ e } \text{\AA}^{-3}$

3100 reflections with $I > 2\sigma(I)$

Data collection

Bruker SMART CCD area-detector diffractometer ω scans

Absorption correction: multi-scan (SADABS; Sheldrick, 1996) $T_{\min} = 0.797, T_{\max} = 0.911$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.023$ wR(F²) = 0.063 S = 1.063404 reflections 353 parameters All H-atom parameters refined

Table 1

Selected geometric parameters (Å, °).

C11-O2 1.237 (2) Cu1-O1 1.948 C11-O1 1.274 (2) Cu1-N3 1.972 C12-N3 1.478 (2) Cu1-N1 1.985 C14-O3 1.242 (2) Cu1-N2 2.013	
C11-O1 1.274 (2) Cu1-N3 1.972 C12-N3 1.478 (2) Cu1-N1 1.9850 C14-O3 1.242 (2) Cu1-N2 2.0130	4 (12)
C12-N3 1.478 (2) Cu1-N1 1.9850 C14-O3 1.242 (2) Cu1-N2 2.0130	2 (15)
$C_{14}-O_{3}$ 1.242 (2) $C_{11}-N_{2}$ 2.0130	5 (14)
) (14)
C14–O4 1.259 (2) Cu1–O5 2.2840) (15)
O2-C11-O1 123.55 (16) N3-Cu1-N2 98.88	3 (6)
O1-Cu1-N3 83.98 (6) N1-Cu1-N2 81.30) (6)
O1-Cu1-N1 93.66 (5) O1-Cu1-O5 96.40) (6)
N3-Cu1-N1 171.65 (6) C12-N3-Cu1 108.3	(10)
O1-Cu1-N2 164.41 (6) C11-O1-Cu1 115.18	3 (11)

Table 2		
Hydrogen-bond geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N3-H12···O4 ⁱ	0.87 (2)	2.02 (2)	2.890 (2)	176.1 (19)
$O7-H24 \cdot \cdot \cdot O6^{i}$	0.82 (3)	1.98 (4)	2.799 (3)	173 (3)
$O7-H25\cdots O4^{ii}$	0.77 (3)	1.89 (3)	2.652 (2)	173 (3)
O10-H23···O8	0.84 (3)	1.96 (3)	2.762 (3)	159 (3)
O10-H22···O3 ⁱⁱⁱ	0.79 (3)	1.95 (3)	2.738 (2)	170 (3)
O8−H21···O7	0.63 (3)	2.21 (3)	2.836 (3)	170 (4)
O8-H20··· $O2$ ⁱⁱ	0.82(4)	1.99 (4)	2.814 (3)	178 (4)
O9−H19···O10	0.73 (4)	2.05 (4)	2.779 (3)	172 (4)
$O9-H18 \cdot \cdot \cdot O8^{iv}$	0.85(1)	2.51 (4)	3.287 (5)	153 (7)
O6−H17···O9	0.87(5)	1.93 (5)	2.789 (4)	166 (5)
$O6-H16\cdots O2^{ii}$	0.67 (3)	2.14 (3)	2.799 (2)	170 (3)
$N3-H11\cdots O10^{v}$	0.81(2)	2.52 (2)	3.111 (2)	131.0 (17)
N3-H11O3	0.81(2)	2.35 (2)	2.917 (2)	127.4 (17)
$O5-H10$ ··· $O7^{iv}$	0.79 (3)	2.01 (3)	2.784 (2)	166 (2)
O5−H9···O6 ^{vi}	0.80 (3)	2.02 (3)	2.815 (3)	177 (3)

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

The H atoms were identified in difference Fourier syntheses and refined freely.

Data collection: *APEX2* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000); software used to prepare material for publication: *SHELXTL*.

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