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#### Key indicators

Single-crystal X-ray study  
T = 293 K  
Mean  $\sigma(\text{C}-\text{C}) = 0.005 \text{ \AA}$   
R factor = 0.051  
wR factor = 0.124  
Data-to-parameter ratio = 13.8

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

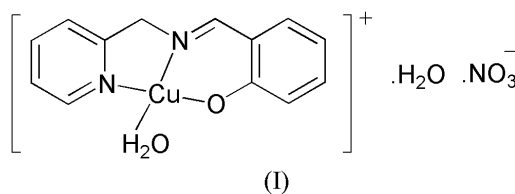
## Aqua[2-(pyridin-2-ylmethyliminomethyl)phenolato- $\kappa^3\text{N},\text{N}',\text{O}$ ]copper(II) nitrate monohydrate

The title compound,  $[\text{Cu}(\text{C}_{13}\text{H}_{11}\text{N}_2\text{O})(\text{H}_2\text{O})](\text{NO}_3)\cdot\text{H}_2\text{O}$ , is a mononuclear copper(II) compound. The Cu atom is coordinated by two N atoms and one O atom from the Schiff base ligand, and another O atom from a water molecule. The four atoms around the metal constitute a slightly distorted square-planar geometry. All O atoms in the nitrate anions and all amine N atoms in the 2-aminomethylpyridine ligands contribute to hydrogen bonds, leading to the formation of sheets parallel to the *ab* plane.

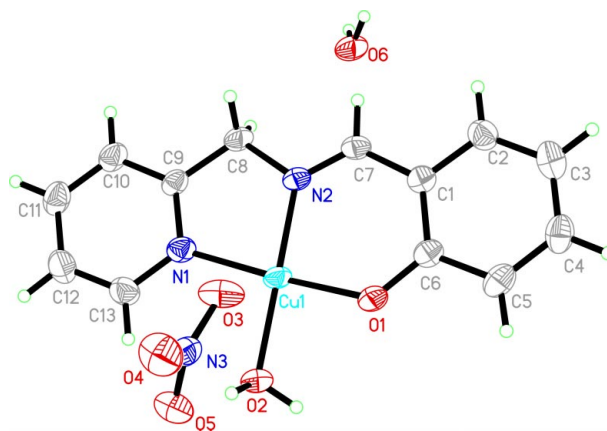
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#### Comment

Recently, we have reported a few Schiff base complexes (You, Lin *et al.*, 2003; You, Qu *et al.*, 2003; You, Xiong *et al.*, 2004; You, Zhu & Liu, 2004). As an extension of our work on the structural characterization of Schiff base complexes, the title mononuclear copper(II) complex, (I), is reported here.

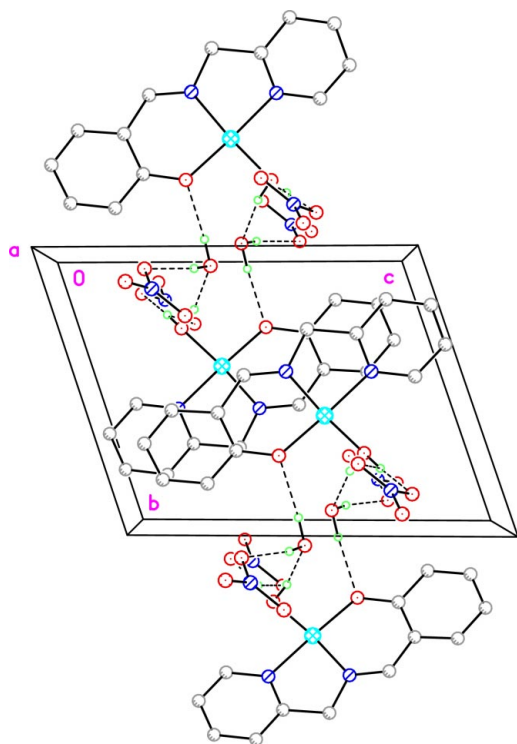


The structure of compound (I) (Fig. 1) consists of a mononuclear  $[\text{Cu}(\text{C}_{13}\text{H}_{11}\text{N}_2\text{O})(\text{H}_2\text{O})]^+$  cation, a nitrate anion and an uncoordinated water molecule. The Cu atom is in a square-planar geometry and is four-coordinated by one O and two N atoms from the Schiff base ligand, and one O atom from the coordinated water molecule. The four coordinating atoms



**Figure 1**

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.



**Figure 2**  
The crystal packing of (I), viewed along the *a* axis. Hydrogen bonds are shown as dashed lines.

around Cu are approximately coplanar, giving a square-planar geometry with an average deviation of 0.091 (4) Å, with the Cu atom 0.064 (3) Å above this plane.

The C7–N2 bond distance of 1.282 (4) Å conforms to the value for a double bond, while the C8–N2 bond distance of 1.469 (4) Å conforms to the value for a single bond. The Cu1–O1 bond length of 1.889 (2) Å (Table 1) is a little shorter than the value of 1.914 (2) Å observed in another Schiff base complex (Langer *et al.*, 2003). The Cu1–N2 bond distance of 1.927 (3) Å is also a little shorter than the value of 1.956 (2) Å observed in the same complex. The Cu1–N1 and Cu1–O2 distances are also comparable with the values found in most copper(II) complexes (Butcher *et al.*, 2003). The bond angles around the Cu<sup>II</sup> centre show some deviations from ideal square-planar geometry.

All O atoms in the nitrate anions and all amine N atoms in the 2-aminomethylpyridine ligands contribute to hydrogen bonds, leading to the formation of sheets parallel to the *ab* plane (Table 2 and Fig. 2).

## Experimental

All chemicals used (reagent grade) were commercially available. Elemental analyses for C, H and N were performed on a Perkin–Elmer 240 C elemental analyser. 2-Aminomethylpyridine (0.1 mmol, 10.8 mg) and Cu(NO<sub>3</sub>)·3H<sub>2</sub>O (0.1 mmol, 24.2 mg) were dissolved in methanol (10 ml). The mixture was stirred for 1 h to give a clear blue solution, which was allowed to evaporate slowly in the open at room temperature. After 5 d, blue block crystals of (I) were formed at the bottom of the vessel. These crystals were isolated, washed three times with methanol and dried in a vacuum desiccator using anhydrous

CaCl<sub>2</sub> (yield 81.3%). Analysis found: C 41.9, H 4.1, N 11.3%; calculated for C<sub>13</sub>H<sub>15</sub>N<sub>3</sub>O<sub>6</sub>Cu: C 41.7, H 4.2, N 11.1%.

### Crystal data

[Cu(C <sub>13</sub> H <sub>11</sub> N <sub>2</sub> O)(H <sub>2</sub> O)](NO <sub>3</sub> )·H <sub>2</sub> O	<i>Z</i> = 2
<i>M<sub>r</sub></i> = 372.82	<i>D<sub>x</sub></i> = 1.646 Mg m <sup>−3</sup>
Triclinic, <i>P</i> $\bar{1}$	Mo <i>K</i> α radiation
<i>a</i> = 7.777 (2) Å	Cell parameters from 3171 reflections
<i>b</i> = 8.993 (2) Å	<i>θ</i> = 2.4–26.8°
<i>c</i> = 11.389 (2) Å	<i>μ</i> = 1.49 mm <sup>−1</sup>
<i>α</i> = 71.25 (3)°	<i>T</i> = 293 (2) K
<i>β</i> = 86.18 (3)°	Block, blue
<i>γ</i> = 86.88 (3)°	0.27 × 0.23 × 0.15 mm
<i>V</i> = 752.2 (3) Å <sup>3</sup>	

### Data collection

Bruker SMART CCD area-detector diffractometer	3044 independent reflections
<i>φ</i> and <i>ω</i> scans	2816 reflections with <i>I</i> > 2σ( <i>I</i> )
Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 1996; Blessing, 1995)	<i>R<sub>int</sub></i> = 0.022
<i>T<sub>min</sub></i> = 0.690, <i>T<sub>max</sub></i> = 0.808	<i>θ<sub>max</sub></i> = 26.5°
5940 measured reflections	<i>h</i> = −9 → 9
	<i>k</i> = −11 → 11
	<i>l</i> = −14 → 14

### Refinement

Refinement on <i>F</i> <sup>2</sup>	$w = 1/[\sigma^2(F_o^2) + (0.063P)^2 + 0.1843P]$
$R[F^2 > 2\sigma(F^2)] = 0.051$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.124$	( $\Delta/\sigma$ ) <sub>max</sub> < 0.001
<i>S</i> = 1.18	$\Delta\rho_{max} = 0.62 \text{ e \AA}^{-3}$
3044 reflections	$\Delta\rho_{min} = -0.25 \text{ e \AA}^{-3}$
220 parameters	
H atoms treated by a mixture of independent and constrained refinement	

**Table 1**

Selected geometric parameters (Å, °).

Cu1–O1	1.889 (2)	Cu1–O2	1.977 (2)
Cu1–N2	1.927 (3)	Cu1–N1	1.993 (3)
O1–Cu1–N2	93.93 (11)	O1–Cu1–N1	176.23 (10)
O1–Cu1–O2	89.19 (11)	N2–Cu1–N1	82.80 (11)
N2–Cu1–O2	170.39 (10)	O2–Cu1–N1	94.35 (11)

**Table 2**

Hydrogen-bonding geometry (Å, °).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
O2–H2A···O5	0.897 (10)	1.787 (11)	2.680 (4)	173 (4)
O2–H2A···N3	0.897 (10)	2.47 (3)	3.248 (4)	146 (4)
O2–H2A···O3	0.897 (10)	2.51 (4)	3.024 (4)	117 (3)
O6–H6B···O4 <sup>i</sup>	0.896 (10)	1.951 (16)	2.825 (4)	165 (4)
O2–H2B···O6 <sup>ii</sup>	0.892 (10)	1.784 (14)	2.665 (4)	169 (4)
O6–H6A···O1 <sup>iii</sup>	0.900 (10)	2.022 (12)	2.917 (3)	173 (3)

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

Atoms H2A, H2B, H6A and H6B were located in a difference Fourier map and refined isotropically, with *U*<sub>iso</sub>(H) values fixed at 0.08 Å<sup>2</sup>. All remaining H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C–H distances in the range 0.93–0.97 Å and with *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C).

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT* and *SHELXTL* (Sheldrick, 1997a); program(s) used to solve structure: *SHELXS97* (Sheldrick,

1997b); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997b); molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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