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Aqua[4,4'-dibromo-6,6'-dimethoxy-2,2'-[ethane-1,2-diylbis(nitrilomethylidene)]-diphenolato]copper(II)

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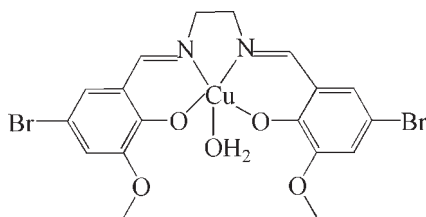
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.009$ Å; disorder in main residue; R factor = 0.053; wR factor = 0.138; data-to-parameter ratio = 12.6.

The title complex, $[\text{Cu}(\text{C}_{18}\text{H}_{16}\text{Br}_2\text{N}_2\text{O}_4)(\text{H}_2\text{O})]$, lies on a crystallographic mirror plane with the Cu^{II} ion coordinated by two N atoms and two O atoms of a tetradentate Schiff base ligand and one O atom from a water ligand in a slightly distorted square-pyramidal environment. The mirror plane, which coincides with the $\text{Cu}-\text{O}_{\text{water}}$ bond, imposes disorder of the atoms of the ethylene group. In the crystal structure, intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds link complex molecules into extended chains along [100].

Related literature

For related structures, see: Nathan *et al.* (2003); Saha *et al.* (2007); Xing (2009). For general background to Schiff base compounds, see: Yu *et al.* (2007); Ghosh *et al.* (2006); Singh *et al.* (2007); Nayka *et al.* (2006).



Experimental

Crystal data

 $[\text{Cu}(\text{C}_{18}\text{H}_{16}\text{Br}_2\text{N}_2\text{O}_4)(\text{H}_2\text{O})]$
 $M_r = 565.70$

 Orthorhombic, $Pnma$
 $a = 8.7299$ (13) Å

 $b = 27.968$ (4) Å

 $c = 7.9900$ (12) Å

 $V = 1950.8$ (5) Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 5.25$ mm⁻¹
 $T = 293$ K

 $0.23 \times 0.20 \times 0.18$ mm

Data collection

Bruker APEXII CCD diffractometer

Absorption correction: multi-scan

(SADABS; Sheldrick, 1996)

 $T_{\text{min}} = 0.378$, $T_{\text{max}} = 0.452$

8970 measured reflections

1759 independent reflections

 1498 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.138$
 $S = 1.13$

1759 reflections

140 parameters

1 restraint

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 1.48$ e Å⁻³
 $\Delta\rho_{\text{min}} = -1.02$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O3}-\text{H3A}\cdots\text{O2}^{\text{i}}$	0.82	2.23	2.963 (5)	150
$\text{O3}-\text{H3A}\cdots\text{O1}^{\text{i}}$	0.82	2.27	2.936 (7)	139

 Symmetry code: (i) $x + \frac{1}{2}, y, -z + \frac{1}{2}$

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2938).

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supporting information

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Aqua{4,4'-dibromo-6,6'-dimethoxy-2,2'-[ethane-1,2-diylbis(nitrilomethylidene)]diphenolato}copper(II)

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S1. Comment

Schiff-bases can readily form stable complexes with most transition metals, in which some may exhibit interesting properties (Yu *et al.*, 2007; Ghosh *et al.*, 2006; Singh *et al.*, 2007; Nayka *et al.*, 2006). Here, we report a Cu(II) complex based on the tetradentate Schiff-base ligand *N,N'*-ethylenebis(5-bromo-3-methoxysalicylalimine).

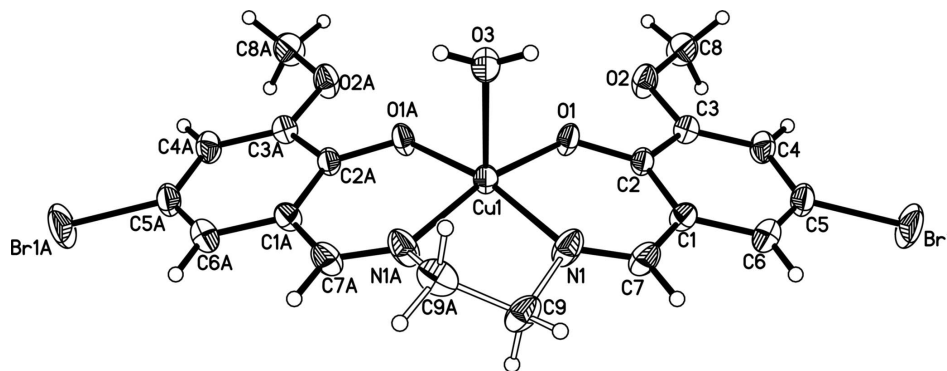
The molecular structure of the title compound is shown in Fig. 1. The complex lies on a crystallographic mirror plane with the Cu^{II} ion coordinated in a slightly distorted square-pyramidal environment. The basal plane is occupied by two N atoms and two O atoms of the Schiff-base ligand, and the apical site is occupied by the O atom of the coordinated water molecule. The Cu^{II} ion is displaced towards the Cu—O_{water} bond from the plane formed by the two N atoms and two O atoms by 0.224 (4) Å. The Cu—N and Cu—O bond lengths are consistent with the corresponding distances found in other Cu Schiff base complexes (Nathan, *et al.*, 2003; Saha, *et al.*, 2007; Xing, 2009).

S2. Experimental

Condensation of ethyl diamine and 5-bromo-3-methoxy-2-hydroxy-benzaldehyde with the ratio 1:2 in ethanol gave the Schiff base ligand. The title compound was synthesized by treatment Cu(ClO₄)₂·6H₂O and the schiff-base ligand (1:1, molar ratio) in methanol. After the mixture was stirred for for about 30 min at room temperature, it was filtered and the filtrate was allowed to partial evaporate in air for one week to produce crystals suitable for X-ray diffraction with a yield about 52%.

S3. Refinement

H atoms were included using the HFIX command in *SHELXL-97* (Sheldrick, 2008), with C—H = 0.96 and 0.93 Å; O—H = 0.82 Å and were allowed for as riding atoms with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C}_{\text{methyl}})$ and ($U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{O})$).

**Figure 1**

The molecular structure of the title compound with the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. All H-atoms are omitted for clarity. Only one disorder component is shown with open bonds [symmetry code (A): (x, -y+1/2, z).

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Crystal data

[Cu(C₁₈H₁₆Br₂N₂O₄)(H₂O)]

M_r = 565.70

Orthorhombic, *Pnma*

Hall symbol: -P 2ac 2n

a = 8.7299 (13) Å

b = 27.968 (4) Å

c = 7.9900 (12) Å

V = 1950.8 (5) Å³

Z = 4

F(000) = 1116

D_x = 1.926 Mg m⁻³

Mo *Kα* radiation, λ = 0.71073 Å

Cell parameters from 2580 reflections

θ = 2.9–26.7°

μ = 5.25 mm⁻¹

T = 293 K

Block, blue

0.23 × 0.20 × 0.18 mm

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

T_{min} = 0.378, *T_{max}* = 0.452

8970 measured reflections

1759 independent reflections

1498 reflections with *I* > 2σ(*I*)

R_{int} = 0.034

θ_{max} = 25.0°, θ_{min} = 2.7°

h = -7→10

k = -33→27

l = -8→9

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.053

wR(*F*²) = 0.138

S = 1.13

1759 reflections

140 parameters

1 restraint

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.0394*P*)² + 13.8845*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.001

Δρ_{max} = 1.48 e Å⁻³

Δρ_{min} = -1.02 e Å⁻³

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F, with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Br1	0.28985 (13)	0.50680 (3)	-0.14560 (14)	0.0802 (4)	
Cu1	0.43677 (12)	0.2500	0.01791 (14)	0.0336 (3)	
O1	0.2831 (5)	0.29928 (14)	0.0459 (6)	0.0394 (11)	
O2	0.0393 (5)	0.35017 (16)	0.0910 (7)	0.0492 (12)	
O3	0.5275 (7)	0.2500	0.2889 (8)	0.0446 (16)	
H3A	0.5656	0.2746	0.3257	0.067*	
N1	0.5856 (6)	0.2962 (2)	-0.0675 (8)	0.0515 (16)	
C1	0.4226 (7)	0.3656 (2)	-0.0697 (8)	0.0385 (15)	
C2	0.2929 (7)	0.3439 (2)	0.0005 (8)	0.0334 (13)	
C3	0.1626 (7)	0.3736 (2)	0.0235 (8)	0.0387 (15)	
C4	0.1622 (8)	0.4211 (2)	-0.0169 (9)	0.0436 (16)	
H4	0.0754	0.4397	0.0006	0.052*	
C5	0.2941 (9)	0.4410 (2)	-0.0848 (9)	0.0472 (17)	
C6	0.4200 (9)	0.4150 (2)	-0.1124 (9)	0.0484 (18)	
H6	0.5062	0.4291	-0.1596	0.058*	
C7	0.5621 (8)	0.3402 (2)	-0.1013 (9)	0.0460 (17)	
H7	0.6421	0.3571	-0.1505	0.055*	
C8	-0.0958 (9)	0.3769 (3)	0.1200 (10)	0.058 (2)	
H8A	-0.0753	0.4015	0.2011	0.087*	
H8B	-0.1746	0.3561	0.1615	0.087*	
H8C	-0.1290	0.3913	0.0172	0.087*	
C9	0.7175 (16)	0.2701 (7)	-0.147 (2)	0.054 (5)	0.50
H9A	0.6948	0.2643	-0.2645	0.065*	0.50
H9B	0.8089	0.2897	-0.1413	0.065*	0.50
C9A	0.7457 (13)	0.2236 (7)	-0.061 (3)	0.052 (5)	0.50
H9A1	0.7812	0.2280	0.0527	0.063*	0.50
H9A2	0.8177	0.2037	-0.1222	0.063*	0.50

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.1002 (8)	0.0319 (4)	0.1084 (8)	0.0054 (4)	0.0121 (6)	0.0259 (4)
Cu1	0.0301 (5)	0.0275 (5)	0.0433 (6)	0.000	0.0040 (5)	0.000
O1	0.038 (2)	0.024 (2)	0.056 (3)	0.0025 (18)	0.006 (2)	0.010 (2)
O2	0.038 (3)	0.034 (2)	0.076 (3)	0.006 (2)	0.014 (2)	0.007 (2)

O3	0.053 (4)	0.032 (3)	0.049 (4)	0.000	-0.011 (3)	0.000
N1	0.035 (3)	0.044 (3)	0.076 (4)	0.000 (3)	0.013 (3)	0.020 (3)
C1	0.041 (4)	0.037 (4)	0.038 (3)	-0.002 (3)	0.001 (3)	0.008 (3)
C2	0.039 (3)	0.026 (3)	0.035 (3)	0.000 (3)	0.000 (3)	0.002 (3)
C3	0.041 (3)	0.035 (3)	0.040 (3)	-0.002 (3)	-0.001 (3)	0.002 (3)
C4	0.049 (4)	0.030 (3)	0.051 (4)	0.005 (3)	-0.004 (3)	0.007 (3)
C5	0.065 (5)	0.029 (3)	0.048 (4)	-0.001 (3)	-0.005 (4)	0.005 (3)
C6	0.054 (4)	0.039 (4)	0.052 (4)	-0.004 (3)	0.000 (4)	0.009 (3)
C7	0.038 (4)	0.043 (4)	0.057 (4)	-0.005 (3)	0.008 (3)	0.013 (3)
C8	0.052 (5)	0.053 (5)	0.069 (5)	0.015 (4)	0.011 (4)	0.009 (4)
C9	0.031 (8)	0.053 (9)	0.078 (14)	-0.001 (7)	0.009 (9)	0.021 (10)
C9A	0.023 (8)	0.067 (11)	0.066 (13)	0.004 (7)	-0.006 (8)	-0.011 (11)

Geometric parameters (Å, °)

Br1—C5	1.903 (7)	C2—C3	1.420 (9)
Cu1—O1 ⁱ	1.937 (4)	C3—C4	1.367 (9)
Cu1—O1	1.937 (4)	C4—C5	1.389 (10)
Cu1—N1	1.954 (5)	C4—H4	0.9300
Cu1—N1 ⁱ	1.954 (5)	C5—C6	1.338 (10)
Cu1—O3	2.305 (6)	C6—H6	0.9300
O1—C2	1.303 (7)	C7—H7	0.9300
O2—C3	1.371 (8)	C8—H8A	0.9600
O2—C8	1.416 (8)	C8—H8B	0.9600
O3—H3A	0.8188	C8—H8C	0.9600
N1—C7	1.277 (9)	C9—C9A	1.491 (19)
N1—C9A ⁱ	1.504 (10)	C9—H9A	0.9700
N1—C9	1.505 (10)	C9—H9B	0.9700
C1—C2	1.402 (9)	C9A—N1 ⁱ	1.504 (10)
C1—C6	1.421 (9)	C9A—H9A1	0.9700
C1—C7	1.433 (10)	C9A—H9A2	0.9700
O1 ⁱ —Cu1—O1	90.8 (2)	C5—C4—H4	120.6
O1 ⁱ —Cu1—N1	166.2 (3)	C6—C5—C4	121.8 (6)
O1—Cu1—N1	91.8 (2)	C6—C5—Br1	120.1 (6)
O1 ⁱ —Cu1—N1 ⁱ	91.8 (2)	C4—C5—Br1	118.1 (5)
O1—Cu1—N1 ⁱ	166.2 (3)	C5—C6—C1	120.2 (7)
N1—Cu1—N1 ⁱ	82.7 (4)	C5—C6—H6	119.9
O1 ⁱ —Cu1—O3	97.45 (18)	C1—C6—H6	119.9
O1—Cu1—O3	97.45 (18)	N1—C7—C1	125.3 (6)
N1—Cu1—O3	95.7 (2)	N1—C7—H7	117.4
N1 ⁱ —Cu1—O3	95.7 (2)	C1—C7—H7	117.4
C2—O1—Cu1	127.2 (4)	O2—C8—H8A	109.5
C3—O2—C8	117.8 (5)	O2—C8—H8B	109.5
Cu1—O3—H3A	118.5	H8A—C8—H8B	109.5
C7—N1—C9A ⁱ	120.7 (10)	O2—C8—H8C	109.5
C7—N1—C9	120.0 (9)	H8A—C8—H8C	109.5
C7—N1—Cu1	127.2 (5)	H8B—C8—H8C	109.5

C9A ⁱ —N1—Cu1	111.3 (9)	C9A—C9—N1	110.7 (14)
C9—N1—Cu1	109.7 (8)	C9A—C9—H9A	109.5
C2—C1—C6	120.2 (6)	N1—C9—H9A	109.5
C2—C1—C7	122.8 (6)	C9A—C9—H9B	109.5
C6—C1—C7	117.0 (6)	N1—C9—H9B	109.5
O1—C2—C1	125.4 (6)	H9A—C9—H9B	108.1
O1—C2—C3	118.1 (6)	C9—C9A—N1 ⁱ	98.7 (12)
C1—C2—C3	116.5 (6)	C9—C9A—H9A1	112.0
C4—C3—O2	123.7 (6)	N1 ⁱ —C9A—H9A1	112.0
C4—C3—C2	122.6 (6)	C9—C9A—H9A2	112.0
O2—C3—C2	113.7 (5)	N1 ⁱ —C9A—H9A2	112.0
C3—C4—C5	118.7 (7)	H9A1—C9A—H9A2	109.7
C3—C4—H4	120.6		

Symmetry code: (i) $x, -y+1/2, z$.

Hydrogen-bond geometry (Å, °)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O3—H3A \cdots O2 ⁱⁱ	0.82	2.23	2.963 (5)	150
O3—H3A \cdots O1 ⁱⁱ	0.82	2.27	2.936 (7)	139

Symmetry code: (ii) $x+1/2, y, -z+1/2$.