

**[2,2'-[4,5-Dibromo-o-phenylene-bis(nitrilodimethylidyne)]diphenolato- $\kappa^4O,N,N',O'$ ](methanol- $\kappa O$ )copper(II)**

Jianxin Xing

Department of Biology, Dezhou University, Dezhou 253023, People's Republic of China

Correspondence e-mail: jianxin\_xing@163.com

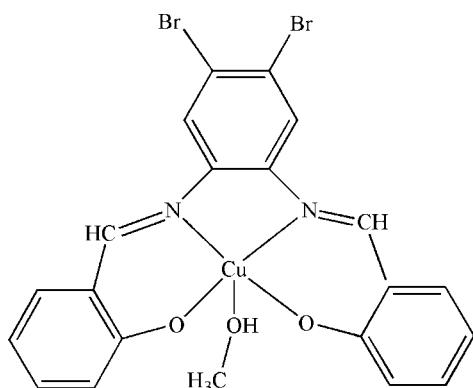
Received 21 March 2009; accepted 26 March 2009

Key indicators: single-crystal X-ray study;  $T = 273\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.008\text{ \AA}$ ;  $R$  factor = 0.043;  $wR$  factor = 0.111; data-to-parameter ratio = 14.6.

In the title compound,  $[\text{Cu}(\text{C}_{20}\text{H}_{12}\text{Br}_2\text{N}_2\text{O}_2)(\text{CH}_3\text{OH})]$ , the  $\text{Cu}^{II}$  ion, and the C, O and hydroxy H atoms of the coordinated methanol molecule are located on a twofold rotation axis, while the methyl H atoms are disordered over two sites about the rotation axis. The  $\text{Cu}^{II}$  ion is coordinated by two N atoms [ $\text{Cu}-\text{N} = 1.960(4)\text{ \AA}$ ] and two O atoms [ $\text{Cu}-\text{O} = 1.908(4)\text{ \AA}$ ] from the tetradeinate Schiff base ligand and by one O atom [ $\text{Cu}-\text{O} = 2.324(6)\text{ \AA}$ ] of the methanol molecule in a square-pyramidal geometry. In the crystal structure, intermolecular  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds link complex molecules into extended chains along [001].

## Related literature

For a related crystal structure, see Saha *et al.* (2007). For general background related to Schiff base compounds, see: Ghosh *et al.* (2006); Nayka *et al.* (2006); Singh *et al.* (2007); Yu *et al.* (2007).



## Experimental

### Crystal data

$[\text{Cu}(\text{C}_{20}\text{H}_{12}\text{Br}_2\text{N}_2\text{O}_2)(\text{CH}_3\text{O})]$	$V = 1982.7(6)\text{ \AA}^3$
$M_r = 567.72$	$Z = 4$
Orthorhombic, $Pnma$	Mo $K\alpha$ radiation
$a = 19.164(4)\text{ \AA}$	$\mu = 5.16\text{ mm}^{-1}$
$b = 19.416(4)\text{ \AA}$	$T = 273\text{ K}$
$c = 5.3287(10)\text{ \AA}$	$0.21 \times 0.15 \times 0.13\text{ mm}$

### Data collection

Bruker APEXII CCD area-detector diffractometer	9881 measured reflections
Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 1996)	2004 independent reflections
$T_{\min} = 0.411$ , $T_{\max} = 0.554$	1517 reflections with $I > 2\sigma(I)$
(expected range = 0.379–0.511)	$R_{\text{int}} = 0.046$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$	242 restraints
$wR(F^2) = 0.111$	H-atom parameters constrained
$S = 1.06$	$\Delta\rho_{\max} = 1.23\text{ e \AA}^{-3}$
2004 reflections	$\Delta\rho_{\min} = -1.82\text{ e \AA}^{-3}$
137 parameters	

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O2—H2A $\cdots$ O1 <sup>i</sup>	0.82	2.30	3.009 (6)	145

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

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*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2794).

## References

- Bruker (2001). *SAINT-Plus*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2004). *APEX2*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Ghosh, R., Rahaman, S. H., Lin, C. N., Lu, T. H. & Ghosh, B. K. (2006). *Polyhedron*, **25**, 3104–3112.
- Nayka, M., Koner, R., Lin, H. H., Flörke, U., Wei, H. H. & Mohanta, S. (2006). *Inorg. Chem.* **45**, 10764–10773.
- Saha, P. K., Dutta, B., Jana, S., Bera, R., Saha, S., Okamoto, K. & Koner, S. (2007). *Polyhedron*, **26**, 563–571.
- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Singh, K., Barwa, M. S. & Tyagi, P. (2007). *Eur. J. Med. Chem.* **42**, 394–402.
- Yu, T. Z., Zhang, K., Zhao, Y. L., Yang, C. H., Zhang, H., Fan, D. W. & Dong, W. K. (2007). *Inorg. Chem. Commun.* **10**, 401–403.

# supporting information

*Acta Cryst.* (2009). E65, m469 [doi:10.1107/S1600536809011179]

## {2,2'-[4,5-Dibromo-o-phenylenebis(nitrilodimethylidyne)]diphenolato- $\kappa^4O,N,N',O'$ }(methanol- $\kappa O$ )copper(II)

Jianxin Xing

### S1. Comment

Schiff-bases have played an important role in the development of coordination chemistry as they readily form stable complexes with most of the 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 new Cu(II) complex based on the tetradentate Schiff-base ligand 4,5-dibromo-1,2-diaminobenzene-*N,N'*-bis (salicylideneimine).

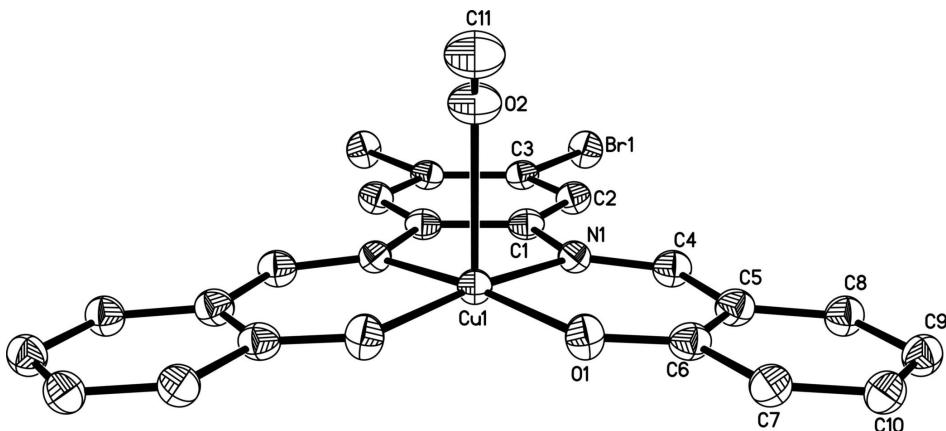
The molecular structure of the title compound is shown in Fig. 1. The Cu<sup>II</sup> ion is pentacoordinated with the four basal sites occupied by two N atoms and two O atoms of the Schiff-base ligand, while the apical site is occupied by the O atom of the coordinated methanol molecule. The Cu<sup>II</sup> ion is displaced towards the Cu—O<sub>methanol</sub> bond from the plane formed by the two N atoms and two O atoms by 0.1017 Å. The coordination geometry of the Cu<sup>II</sup> ion is square-pyramidal. The Cu—N, Cu—O and Cu—O<sub>methanol</sub> bond lengths are consistent with the corresponding distances in aqua-(*N,N'*-ethylenebis(3-methoxysalicylaldiminato)-*N,N',O,O'*)copper(II) (Saha, *et al.*, 2007).

### S2. Experimental

The Schiff base ligand was synthesized by condensation of 4,5-dibromo-1,2-diaminobenzene and 2-hydroxybenzaldehyde with the ratio 1:2 in ethanol. The synthesis of the title complex was carried out by reacting Cu(ClO<sub>4</sub>)<sub>2</sub>·6H<sub>2</sub>O, and the schiff-base ligand (1:1, molar ratio) in methanol. After the stirring process was continued for about 10 min at room temperature, the mixture was filtered and the filtrate was allowed to partial evaporate in air for several days to produce crystals suitable for X-ray diffraction with a yield about 55%.

### S3. Refinement

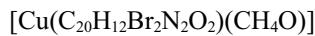
H atoms were included using the HFIX command in *SHELXL-97* (Sheldrick, 2008), with C—H = 0.96 and 0.93 Å; O—H = 0.82 Å and were allowed for as riding atoms with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C}_{\text{methyl}})$  and ( $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{O})$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

**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. Unlabelled atoms are related by the symmetry operator ( $x, -y+3/2, z$ ).

### {2,2'-[4,5-Dibromo-*o*-phenylenebis(nitrilodimethylidyne)]diphenolato- $\kappa^4O,N,N',O'$ }(methanol- $\kappa O$ )copper(II)

#### Crystal data



$$M_r = 567.72$$

Orthorhombic,  $Pnma$

Hall symbol: -P 2ac 2n

$$a = 19.164 (4) \text{ \AA}$$

$$b = 19.416 (4) \text{ \AA}$$

$$c = 5.3287 (10) \text{ \AA}$$

$$V = 1982.7 (6) \text{ \AA}^3$$

$$Z = 4$$

$$F(000) = 1116$$

$$D_x = 1.902 \text{ Mg m}^{-3}$$

$$D_m = 1.902 \text{ Mg m}^{-3}$$

$D_m$  measured by not measured

Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1834 reflections

$$\theta = 3.0\text{--}22.2^\circ$$

$$\mu = 5.16 \text{ mm}^{-1}$$

$$T = 273 \text{ K}$$

Block, red

$$0.21 \times 0.15 \times 0.13 \text{ mm}$$

#### Data collection

Bruker APEXII CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Sheldrick, 1996)

$$T_{\min} = 0.411, T_{\max} = 0.554$$

$$9881 \text{ measured reflections}$$

$$2004 \text{ independent reflections}$$

$$1517 \text{ reflections with } I > 2\sigma(I)$$

$$R_{\text{int}} = 0.046$$

$$\theta_{\max} = 26.0^\circ, \theta_{\min} = 2.1^\circ$$

$$h = -23 \rightarrow 16$$

$$k = -23 \rightarrow 23$$

$$l = -6 \rightarrow 6$$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.043$$

$$wR(F^2) = 0.111$$

$$S = 1.06$$

$$2004 \text{ reflections}$$

$$137 \text{ parameters}$$

$$242 \text{ restraints}$$

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

$$w = 1/[\sigma^2(F_o^2) + (0.0453P)^2 + 5.5878P]$$

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

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

$$\Delta\rho_{\max} = 1.23 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -1.82 \text{ e \AA}^{-3}$$

*Special details*

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s 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  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*$ / $U_{\text{eq}}$	Occ. (<1)
Cu1	0.12529 (5)	0.7500	1.11061 (15)	0.0314 (2)	
Br1	-0.10466 (3)	0.66291 (3)	0.15521 (10)	0.0450 (2)	
O1	0.1647 (2)	0.68206 (19)	1.3271 (6)	0.0423 (9)	
O2	0.2119 (3)	0.7500	0.8053 (11)	0.0547 (10)	
H2A	0.1990	0.7500	0.6587	0.066*	
N1	0.0729 (2)	0.6827 (2)	0.9116 (7)	0.0301 (9)	
C1	0.0321 (3)	0.7138 (3)	0.7227 (9)	0.0347 (9)	
C2	-0.0080 (3)	0.6782 (3)	0.5479 (9)	0.0367 (9)	
H2	-0.0076	0.6304	0.5456	0.044*	
C3	-0.0485 (3)	0.7142 (3)	0.3777 (9)	0.0327 (10)	
C4	0.0738 (3)	0.6172 (3)	0.9476 (10)	0.0394 (8)	
H4	0.0472	0.5905	0.8391	0.047*	
C5	0.1119 (3)	0.5817 (3)	1.1386 (9)	0.0392 (8)	
C6	0.1546 (3)	0.6165 (3)	1.3181 (9)	0.0397 (9)	
C7	0.1889 (3)	0.5735 (3)	1.4984 (10)	0.0416 (9)	
H7	0.2177	0.5936	1.6182	0.050*	
C8	0.1056 (3)	0.5104 (3)	1.1487 (10)	0.0425 (9)	
H8	0.0772	0.4889	1.0308	0.051*	
C9	0.1388 (3)	0.4700 (3)	1.3222 (10)	0.0441 (10)	
H9	0.1339	0.4224	1.3226	0.053*	
C10	0.1804 (3)	0.5040 (3)	1.4990 (11)	0.0433 (10)	
H10	0.2030	0.4779	1.6207	0.052*	
C11	0.2775 (5)	0.7500	0.8417 (18)	0.067 (2)	
H11A	0.3003	0.7274	0.7040	0.101*	0.50
H11B	0.2878	0.7260	0.9947	0.101*	0.50
H11C	0.2938	0.7966	0.8539	0.101*	0.50

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Cu1	0.0344 (5)	0.0327 (5)	0.0269 (4)	0.000	-0.0054 (4)	0.000
Br1	0.0519 (4)	0.0464 (4)	0.0369 (3)	-0.0063 (3)	-0.0153 (2)	-0.0044 (2)
O1	0.053 (2)	0.039 (2)	0.036 (2)	0.0042 (18)	-0.0152 (17)	0.0006 (16)
O2	0.048 (2)	0.076 (2)	0.040 (2)	0.000	-0.0030 (18)	0.000
N1	0.031 (2)	0.032 (2)	0.027 (2)	0.0034 (18)	-0.0029 (17)	0.0005 (18)

C1	0.0348 (17)	0.0422 (17)	0.0272 (16)	0.0003 (15)	-0.0009 (15)	0.0008 (15)
C2	0.0377 (19)	0.042 (2)	0.0308 (18)	-0.0010 (17)	-0.0037 (17)	0.0007 (17)
C3	0.033 (2)	0.040 (2)	0.0252 (19)	-0.0030 (18)	-0.0033 (17)	-0.0008 (18)
C4	0.0398 (16)	0.0449 (17)	0.0336 (16)	0.0007 (15)	-0.0047 (14)	0.0008 (14)
C5	0.0406 (17)	0.0443 (17)	0.0327 (16)	0.0021 (15)	-0.0034 (15)	0.0009 (15)
C6	0.0397 (17)	0.0469 (18)	0.0326 (16)	0.0026 (16)	-0.0015 (15)	0.0020 (15)
C7	0.0446 (19)	0.0446 (19)	0.0355 (18)	0.0035 (18)	-0.0059 (17)	0.0017 (17)
C8	0.0452 (18)	0.0444 (18)	0.0380 (18)	0.0005 (17)	-0.0052 (16)	0.0017 (16)
C9	0.048 (2)	0.044 (2)	0.0400 (19)	0.0020 (18)	-0.0039 (17)	0.0039 (17)
C10	0.047 (2)	0.0451 (19)	0.0381 (19)	0.0044 (18)	-0.0044 (17)	0.0055 (17)
C11	0.052 (4)	0.090 (4)	0.060 (4)	0.000	0.002 (4)	0.000

Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )

Cu1—O1 <sup>i</sup>	1.908 (4)	C4—C5	1.430 (7)
Cu1—O1	1.908 (3)	C4—H4	0.9300
Cu1—N1 <sup>i</sup>	1.960 (4)	C5—C8	1.390 (8)
Cu1—N1	1.960 (4)	C5—C6	1.429 (7)
Cu1—O2	2.324 (6)	C6—C7	1.432 (7)
Br1—C3	1.886 (5)	C7—C10	1.358 (8)
O1—C6	1.289 (7)	C7—H7	0.9300
O2—C11	1.271 (9)	C8—C9	1.368 (7)
O2—H2A	0.8199	C8—H8	0.9300
N1—C4	1.285 (7)	C9—C10	1.399 (8)
N1—C1	1.410 (6)	C9—H9	0.9300
C1—C2	1.392 (7)	C10—H10	0.9300
C1—C1 <sup>i</sup>	1.405 (10)	C11—H11A	0.9600
C2—C3	1.383 (7)	C11—H11B	0.9600
C2—H2	0.9300	C11—H11C	0.9600
C3—C3 <sup>i</sup>	1.388 (10)		
O1 <sup>i</sup> —Cu1—O1	87.5 (2)	N1—C4—H4	116.9
O1 <sup>i</sup> —Cu1—N1 <sup>i</sup>	93.95 (16)	C5—C4—H4	116.9
O1—Cu1—N1 <sup>i</sup>	172.22 (18)	C8—C5—C6	119.6 (5)
O1 <sup>i</sup> —Cu1—N1	172.22 (18)	C8—C5—C4	117.7 (5)
O1—Cu1—N1	93.95 (16)	C6—C5—C4	122.7 (5)
N1 <sup>i</sup> —Cu1—N1	83.6 (2)	O1—C6—C5	125.3 (5)
O1 <sup>i</sup> —Cu1—O2	98.09 (16)	O1—C6—C7	118.9 (5)
O1—Cu1—O2	98.09 (16)	C5—C6—C7	115.9 (5)
N1 <sup>i</sup> —Cu1—O2	89.29 (16)	C10—C7—C6	121.7 (5)
N1—Cu1—O2	89.29 (16)	C10—C7—H7	119.1
C6—O1—Cu1	127.0 (3)	C6—C7—H7	119.1
C11—O2—Cu1	126.8 (6)	C9—C8—C5	123.8 (5)
C11—O2—H2A	116.4	C9—C8—H8	118.1
Cu1—O2—H2A	116.8	C5—C8—H8	118.1
C4—N1—C1	122.6 (4)	C8—C9—C10	116.7 (6)
C4—N1—Cu1	124.8 (3)	C8—C9—H9	121.6
C1—N1—Cu1	112.6 (3)	C10—C9—H9	121.6

C2—C1—C1 <sup>i</sup>	119.8 (3)	C7—C10—C9	122.3 (5)
C2—C1—N1	124.8 (5)	C7—C10—H10	119.2
C1 <sup>i</sup> —C1—N1	115.4 (3)	C9—C10—H10	118.5
C3—C2—C1	119.9 (5)	O2—C11—H11A	109.5
C3—C2—H2	120.1	O2—C11—H11B	109.5
C1—C2—H2	120.1	H11A—C11—H11B	109.5
C2—C3—C3 <sup>i</sup>	120.3 (3)	O2—C11—H11C	109.5
C2—C3—Br1	117.7 (4)	H11A—C11—H11C	109.5
C3 <sup>i</sup> —C3—Br1	121.90 (15)	H11B—C11—H11C	109.5
N1—C4—C5	126.3 (5)		

Symmetry code: (i)  $x, -y+3/2, z$ .

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

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O2—H2A $\cdots$ O1 <sup>ii</sup>	0.82	2.30	3.009 (6)	145

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