

# Synthesis, Crystal Structure, and Antibacterial Evaluation of a Copper(II) Compound Derived From 1,2-Diaminobenzene-*N*,*N*'-bis(3-methoxysalicylideneimine)

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A copper(II) compound, [CuL][CuL(OH<sub>2</sub>)]·ClO<sub>4</sub>·0.5CH<sub>3</sub>OH, where L = 1,2-diaminobenzene-*N*,*N'*-bis(3-methoxysalicylidenei mine), has been synthesized and characterized by means of spectroscopic methods and single-crystal X-ray structure determination. The compound crystallizes in the orthorhombic space group *Cmc*2<sub>1</sub> with unit cell dimensions a = 17.981(3) Å, b = 17.776(3) Å, c = 14.047(3) Å, V = 4489.8(14) Å<sup>3</sup>, Z = 4,  $R_1 = 0.0511$ , and  $wR_2 = 0.1267$ . The asymmetric unit of the compound contains one mononuclear [CuL] unit, in which the Cu atom adopts a squareplanar geometry, one mononuclear [CuL(OH<sub>2</sub>)] unit, in which the Cu atom adopts a square-pyramidal geometry, one perchlorate anion, and one half methanol solvate. The compound was tested for its antibacterial activities to assess its inhibiting potential.

Keywords antibacterial activity, copper compound, crystal structure, Schiff base, X-ray diffraction

## INTRODUCTION

Schiff bases have often been used as versatile chelating ligands in coordination chemistry. Schiff bases with donors such as O, N, and S have structure similarities with neutral biological systems and due to presence of imine groups are utilized in elucidating the mechanism of transformation of rasemination reaction in biological system.<sup>[1–3]</sup> The Schiff bases prepared from salicylaldehyde and its derivatives are interesting and have received considerable attention not only for their application in coordination chemistry<sup>[4–6]</sup> but also for their importance in medicinal and pharmaceutical fields. Most Schiff bases and their complexes show biological activities including antibacterial,<sup>[7–9]</sup> antifungal,<sup>[9,10]</sup> antitumor,<sup>[11]</sup> anticancer,<sup>[10]</sup> anticorrosion, and anti-inflammatory activities. As a continuation of the biological activities of Schiff base complexes, in the present article, I report a new Schiff base copper compound with the ligand 1,2-diaminobenzene-N,N'-bis(3-methoxysalicylideneimine).

# **EXPERIMENTAL**

### General Considerations

All materials were obtained from commercial sources and used as received. Copper perchlorate was prepared by the reaction of basic cupric carbonate with perchloric acid in distilled water. The Schiff base ligand was prepared according to the literature procedure.<sup>[12]</sup> Infrared spectra were collected on a Nicolet 5SXC FT-IR spectrometer (Linyi University, China) as KBr pellets. Electronic absorption (UV-Vis) spectra were collected on a Cary 500 UV-Vis-NIR spectrophotometer (Linyi University, China). Microanalyses for C, H, and N were performed on a Perkin-Elmer 2400 CHNS/O elemental analyzer (Linyi University, China).

# Preparation of the Copper Compound

To a methanolic solution (20 mL) of the Schiff base ligand (0.38 g, 1 mmol) was added a methanolic solution (10 mL) of copper perchlorate (0.37 g, 1 mmol). The reaction mixture was gently heated at 50°C for 30 min while stirring and the complex precipitated was recrystallized from methanol, yielding well-shaped single crystals suitable for X-ray diffraction. Yield: 45%. Elemental Anal. Calcd. for C<sub>46</sub>H<sub>44</sub>ClCu<sub>2</sub>N<sub>4</sub>O<sub>14</sub>: C, 53.15; H, 4.27; N, 5.39. Found: C, 52.93; H, 4.40; N, 5.51%. IR (KBr, cm<sup>-1</sup>): 1605 (C=N). UV-Vis (toluene,  $\lambda$ , nm, ( $\varepsilon$ , M<sup>-1</sup> cm<sup>-1</sup>)): 325 (21200), 339 (17900), 357 (14500), 403 (11300), 458 (14500), 693 (2730).

## **Crystal Structure Determination and Refinement**

Crystal data, data collection and refinement parameters for the compound are listed in Table 1. Data were obtained on a Bruker Apex II diffractometer (Shandong University, China) equipped with graphite monochromated Mo K $\alpha$  ( $\lambda = 0.71073$ Å) radiation. The structure of the compound was solved by direct methods and refined on  $F^2$  by full-matrix least-squares using

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Crystal data and structure refinement for the compound		Selected bond lengths (A) and angles (°) for the			
Empirical formula	$C_{46}H_{44}ClCu_2N_4O_{14}$	Bond lengths			
$M/g \text{ mol}^{-1}$	1039.4	Cu1-O3	1.896(4)	Cu1-N2	
Crystal size/mm	$0.30 \times 0.27 \times 0.27$	Cu2-O1	1.905(3)	Cu2-N1	
$2\theta$ range/deg	1.61-27.00	Cu2-O9	2.431(6)		
Completeness to $2\theta$ (%)	100	Bond angles			
<i>h</i> ; <i>k</i> ; <i>l</i> range	-22, 22; -14, 22; -17, 17	O3-Cu1-O3A	89.2(2)	O3-Cu1-N2A	
Temperature/K	298(2)	O3-Cu1-N2	93.6(2)	N2-Cu1-N2A	
Crystal system	Orthorhombic	01-Cu2-O1A	87.8(2)	O1-Cu2-N1A	
Space group	$Cmc2_1$	O1-Cu2-N1	93.6(2)	N1-Cu2-N1A	
a/Å	17.981(3)	O1-Cu2-O9	96.2(2)	N1-Cu2-O9	
b/Å	17.776(3)	Symmetry operation for $A:1 \rightarrow 2$			
c/Å	14.047(3)	Symmetry operation for A. $1 - x$ , y, z.			
V/Å <sup>3</sup>	4489.8(14)	SHELX-97. <sup>[13]</sup> All non-hydrogen atoms were re ically. All hydrogen atoms were placed in calcu			
Ζ	4				
$D_{\rm calc}/{\rm g}~{\rm cm}^{-3}$	1.538				
<i>F</i> (000)	2140	assigned fixed isotropic thermal parameters at			
$\mu/\mathrm{mm}^{-1}$	1.080	the equivalent isotropic $U$ of the atoms to wh			
Max/min transmission	0.759/0.738	The contributions of these hydrogen atoms were structure factor calculations. Selected bond leng			
Reflections collected $(R_{int})$	12853 (0.0438)				
Independent reflections	4776				
Observed reflections	3718	are listed in Table	e 2.		
Parameters refined	320	Antibacterial A	ctivities		
Restraints	43	Antibacterial	activities of th	e compound w	
Max/min $\Delta \rho$ /e Å <sup>-3</sup>	0.813/-0.384	using agar well diffusion method. <sup>[14]</sup> The ac free Schiff base ligand and standard drug in studied against the <i>Staphylococcus aureus</i> and <i>tilis</i> (as gram-positive bacteria) and <i>Pseudon</i>			
$R_1, wR_2 \left[ I > 2\sigma(I) \right]$	0.0511, 0.1267				
$R_1, wR_2$ (all data)	0.0703, 0.1382				
Goodness-of-fit on $F^2$	1.018				

TABLE 1

TABLE 2 (°) for the compound

1.926(4)

1.930(4)

177.2(2)

83.7(3)

172.7(2)

84.2(3)

90.8(2)

were refined anisotropin calculated positions, eters at 1.2 or 1.5 times is to which they are atespective parent atoms. ms were included in the ond lengths and angles

ound were investigated The activities of the drug imipenem were reus and Bacillus sub-Pseudomonas aeruginosa, Escherichia coli, and Salmonella typhi (as gram-negative



FIG. 1. Molecular structure of the complex. Hydrogen atoms have been omitted for clarity. Symmetry operation for A: 1 - x, y, z.

bacteria). Strains were obtained from Dalian Medical University. The solution of 2  $\mu$ g/mL of each compound (free Schiff base ligand and Imipenem) in DMSO was prepared for testing against bacteria. Centrifuged pellets of bacteria from a 24 h old culture containing approximately  $10^4$ – $10^6$  CFU (colony forming unit) per milliliter were spread on the surface of Muller Hinton Agar plates. Wells were created in medium with the help of a sterile metallic bores and nutrients agar media (agar 20 g + beef extract 3 g + peptones 5 g) in 1000 mL of distilled water (pH 7.0), autoclaved and cooled down to  $45^{\circ}$ C. Then it was seeded with 10 mL of prepared inocula to have  $10^6$  CFU/mL. Petri plates were prepared by pouring 75 mL of seeded nutrient agar. The activities were determined by measuring the diameters of the inhibition zones (in mm). The growth inhibition was calculated according to Rahman et al.<sup>[14]</sup>

# **RESULTS AND DISCUSSION**

The Schiff base ligand was prepared by a 1:2 condensation of benzene-1,2-diamine with 3-methoxysalicylaldehyde in methanol. Reaction of the ligand with copper perchlorate in methanol gave the copper compound.

# Structure Description of the Compound

The crystal structure analysis of the compound reveals that it contains a [CuL] unit, a [CuL(OH<sub>2</sub>)] unit, a perchlorate anion, and one half methanol solvate in the asymmetric unit (Figure 1). In [CuL], the Cu atom is in a square-planar geometry, coordinated by two imine N and two phenolate O atoms of L. In [CuL(OH<sub>2</sub>)], the Cu atom is in a square-pyramidal geometry, coordinated by two imine N and two phenolate O atoms of L in the basal plane, and weakly coordinated by one water O atom at the apical position. In [CuL(OH<sub>2</sub>)], the C8-C9-C10-C10A-C9A-C8A (symmetry code for A: 1 - x, y, z) benzene ring forms a dihedral angle of  $1.5(3)^{\circ}$  with the C1-C6 benzene ring. In [CuL], the C19-C20-C21-C21A-C20A-C19A benzene ring forms a dihedral angle of 7.7(3)° with the C12-C17 benzene ring. The Cu-O and Cu-N bonds in the basal plane of [CuL(OH<sub>2</sub>)] are slightly longer than those in [CuL], and all of them are in the range of similar Schiff base copper(II) complexes with square-planar and square-pyramidal geometries.<sup>[15-18]</sup> In the three-dimensional crystal structure of the compound, the [CuL] and [CuL(OH<sub>2</sub>)] molecules are linked together by four intermolecular O-H···O hydrogen bonds (Table 3, Figure 2).

TABLE 3Hydrogen-bond geometry for the compound (Å, °)

D-H···A	D-H	H···A	D····A	D−H…A
09–H9W…O3	0.94(1)	2.30(2)	3.01(3)	132(3)
09–H9W…O4	0.94(1)	2.02(2)	2.86(3)	147(3)



FIG. 2. The molecular packing structure of the complex, viewed along the *y* axis. Hydrogen bonds are drawn as dashed lines.

## **Antibacterial Activity**

The copper compound, the free Schiff base ligand, and the standard drug imipenem were screened separately for their antibacterial activities against the bacteria *Staphylococcus aureus*, *Bacillus subtilis*, *Pseudomonas aeruginosa*, *Escherichia coli*, and *Salmonella typhi*. The diffusion agar technique was used to evaluate the antibacterial activities of the compounds.<sup>[19]</sup> The results are summarized in Table 4. From the results, it is obvious that the copper compound is more active toward gram-positive

 TABLE 4

 Bactericidal screening data of the tested materials (inhibition zone in mm)

Microorganism	Copper compound	Free Schiff base ligand	Imipenem
Staphylococcus aureus	++++	++	++++
Bacillus subtilis	+++	++	++++
Pseudomonas aeruginoca	+	_	++++
Escherichia coli	++	_	++++
Salmonella typhi	+	_	++++

++++ = excellent activity (>90% inhibition); +++ = significant activity (70–90% inhibition); ++ = moderate activity (40–70% inhibition); + = weak activity (<40% inhibition), - = no activity.

strains than gram-negative strains. This may be caused by the difference in the structures of the cell walls. The walls of gram-negative cells are more complex than those of gram-positive cells. The copper compound showed excellent activity against *Staphylococcus aureus* and significant activity against *Bacillus subtilis*, but weak or moderate activities against the gram-negative strains. The free Schiff base ligand has moderate activities against *Staphylococcus aureus* and *Bacillus subtilis* but no activity against *Pseudomonas aeruginoca, Escherichia coli*, and *Salmonella typhi*. In general, the copper compound showed stronger antibacterial activities than the free Schiff base ligand.

## CONCLUSION

A copper(II) compound with the Schiff base ligand 1,2diaminobenzene-N,N'-bis(3-methoxysalicylideneimine) were prepared and characterized. The antibacterial evaluation indicates that the compound may be used as an efficient antibacterial material. This is in accordance with those reported previously that metal complexes showed stronger antibacterial activities than the free Schiff base ligands.<sup>[20–22]</sup>

# SUPPLEMENTARY MATERIALS

Crystallographic data for the structural analyses has been deposited with the Cambridge Crystallographic Data Centre, CCDC No. 866129. Copy of this information maybe obtained free of charge from The Director, CCDC, 12 Union Road, Cambridge, CB2 1EZ, UK, fax: +44 1223 366 033, e-mail: deposit@ ccdc.cam.ac.uk or on the web www: http://www.ccdc.cam.ac. uk.

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