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Synthesis and Reactivity in Inorganic, Metal-Organic, and Nano-Metal Chemistry

Publication details, including instructions for authors and subscription information:

<http://www.tandfonline.com/loi/lsrt20>

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Accepted author version posted online: 18 Feb 2015.



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To cite this article: Chao Li, Xu-Feng Meng, Wei-Ning Li, Xin Zhou & Jing-Jun Ma (2015): Synthesis, Crystal Structure and Biological Property of a Novel Dinuclear Copper(II) Complex, *Synthesis and Reactivity in Inorganic, Metal-Organic, and Nano-Metal Chemistry*, DOI: [10.1080/15533174.2014.963238](https://doi.org/10.1080/15533174.2014.963238)

To link to this article: <http://dx.doi.org/10.1080/15533174.2014.963238>

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Synthesis, Crystal Structure and Biological Property of a Novel Dinuclear Copper(II) Complex

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Abstract

The reaction of 2-acetylpyridine, cyclohexylamine and copper(II) perchlorate in methanol resulted in the dinuclear copper(II) complex $[\text{Cu}_2\text{L}_2]\cdot 2\text{ClO}_4$, where L is the monoanionic form of 1-methoxy-1,3-dipyridin-2-ylbutane-1,3-diol. The complex was characterized by elemental analysis and IR spectroscopic method in the solid state. Single crystal X-ray analysis was performed, which indicated that the complex possesses a crystallographic inversion center symmetry. The Cu atom in the complex is in a square pyramidal geometry. Thermal behavior and biological property on the bacteria strains *B. subtilis*, *E. coli*, and *S. aureus* of the complex were studied.

Keywords Copper complex, dinuclear complex, crystal structure, thermal behavior, antibacterial activity.

INTRODUCTION

Transition metal complexes with various organic ligands have been reported to possess interesting biological properties, such as bacterial, antifungal and antitumor activities.^[1-4] In general, the complexation of organic ligands with most transition metal atoms influences their antibacterial activities.^[5-7] In recent years, copper complexes have been extensively used in antibacterial agents.^[8-10] At this point of view and in continuation of work on the study of metal complexes, herein we report a novel dinuclear copper complex, $[\text{Cu}_2\text{L}_2]\cdot 2\text{ClO}_4$, where L is the monoanionic form of 1-methoxy-1,3-dipyridin-2-ylbutane-1,3-diol.

EXPERIMENTAL

Materials and Methods

2-Acetylpyridine and cyclohexylamine were purchased from Merck and Fluka, and used as received. Copper perchlorate was prepared by the reaction of basic cupric carbonate with perchloric acid in aqueous solution. All other chemicals and solvents used in this work are of analytical grade available commercially and used without further purification. Elemental analyses (carbon, hydrogen and nitrogen) of the complex were obtained from a Carlo ERBA Model EA 1108 analyzer. Infrared spectrum was performed by using a KBr pellet on a Jasco-5300 FT-IR spectrophotometer. Solution electrical conductivity was measured with a DDS-11A conductivity meter.

Synthesis of $[\text{Cu}_2\text{L}_2]\cdot 2\text{ClO}_4$

2-Acetylpyridine (0.01 mol, 1.21 g) and cyclohexylamine (0.01 mol, 0.99 g) were reacted in methanol (15 mL) at room temperature for 30 min. Then, copper perchlorate (0.01 mol, 3.88 g)

and NH_4BF_4 (0.01 mol, 1.05 g) in methanol (10 mL) were added slowly to the above stirred solution. The final mixture was stirred at room temperature for another 30 min to give a blue solution. The solution was filtered to remove some microdirty, and left to slow evaporate in air. A few days later, small block-shaped single crystals were obtained. Yield: 2.75 g (63%). Anal. Calcd. for $\text{C}_{15}\text{H}_{17}\text{ClCuN}_2\text{O}_7$ (%): C, 41.29; H, 3.93; N, 6.42. Found: C, 41.13; H, 4.02; N, 6.55. IR data (cm^{-1} , KBr): 3391, 1609, 1571, 1491, 1474, 1445, 1376, 1218, 1097, 863, 783, 757, 623, 553, 525, 479, 429, 345.

X-ray Crystallography

A well-shaped X-ray quality crystal of the complex was pick up under a microscope and investigated in a diffraction experiment at 298(2) K on a Bruker Apex II diffractometer with monochromated Mo K radiation ($\lambda = 0.71073 \text{ \AA}$) obtained from a graded multilayer X-ray optics. The structure was solved by direct method with SHELXS-97,^[11] and refined with full-matrix least-squares techniques on F^2 with SHELXL-97.^[11] The C- and O-bonded hydrogen atoms were calculated in an idealized geometry, riding on their parent atoms, with distances restrained to 0.9360.97 \AA for C6H and 0.93 \AA for O6H. The perchlorate anion is disordered over two sites with occupancies of 0.64(2) and 0.36(2). The crystal data and refinement parameters are listed in Table 1.

Antibacterial Assay

The antibacterial activities were tested against *B. subtilis*, *E. coli*, and *S. aureus* using Mueller-Hinton medium. The MICs (minimum inhibitory concentrations) of the test compounds were determined by a colorimetric method using the dye MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide]. A stock solution of the synthesized compound (50 g mL^{-1}) in

DMSO was prepared and quantities of the test compounds were incorporated in specified quantity of sterilized liquid Mueller-Hinton medium. A specified quantity of the medium containing the compound was poured into microtitration plates. A suspension of the microorganism was prepared to contain approximately 10^5 cfu mL⁻¹ and applied to microtitration plates with serially diluted compounds in DMSO to be tested and incubated at 37 °C for 24 h. After the MICs were visually determined on each of the microtitration plates, 50 μ L of PBS (phosphate buffered saline 0.01 mol L⁻¹, pH 7.4: Na₂HPO₄·12H₂O 2.9 g, KH₂PO₄ 0.2 g, NaCl 8.0 g, KCl 0.2 g, distilled water 1000 mL) containing 2 mg of MTT/mL was added to each well. Incubation was continued at room temperature for 4-5 h. The content of each well was removed and 100 μ L of isopropanol containing 5% 1 mol L⁻¹ HCl was added to extract the dye. After 12 h of incubation at room temperature, the optical density (OD) was measured with a microplate reader at 550 nm. The antibiotics kanamycin and penicillin were used as standard drugs. The observed MIC values are given in Table 3.

RESULTS AND DISCUSSION

Synthesis and Characterization

Massa and coworkers reported a mononuclear copper(II) complex with the Schiff base ligand derived from pyridine-2-carbaldehyde with cyclohexylamine.^[12] In the present work, we used 2-acetylpyridine and cyclohexylamine as the starting material, to form a Schiff base ligand cyclohexyl-(1-pyridin-2-ylethylidene)amine. The Schiff base ligand was further reacted with copper perchlorate in the presence of NH₄BF₄. To our surprise, the ligand transferred to 1-methoxy-1,3-dipyridin-2-ylbutane-1,3-diol during the reaction and crystallization. The complex was obtained as single crystals, stable in air and soluble in polar organic solvents such as ethanol,

methanol, DMF, and DMSO. Elemental analyses of the complex are consistent with the general molecular formula proposed by single crystal X-ray determination. The molar conductivity of the complex in absolute methanolic solution confirms the 1:1 electrically nature.^[13]

Description of the Crystal Structure of the Complex

Figure 1 gives the ORTEP diagram with atomic labeling scheme of the complex. Selected bond lengths and angles are given in Table 2. The complex possesses a crystallographic inversion center symmetry, and the asymmetric unit of the complex contains a dinuclear copper complex cation and two disordered perchlorate anions. The Cu...Cu distance is 2.985(1) Å. In the complex cation, two of the monoanionic four-dentate ligands wrap the Cu atoms with coordination of two pyridine nitrogen atoms and two oxygen atoms in a distorted square pyramidal coordination, which is proved by the τ reference ($\tau = 0.24$).^[14] One phenolic hydroxyl is deprotonated on complexation, and acts as a bridging group, while the other one is kept in neutral. The distortion of the square pyramidal coordination mainly comes from the tensile force of the four-membered chelate ring Cu1-O6-Cu1A-O6A and five-membered chelate rings Cu1-N2-C1-C6-O7 and Cu1-N3A-C10A-C9A-O6A. The apical O7 atom forms coordinate bond angles in the range of 76.0(1)-108.5(2)° with the basal donor atoms N2, N3, O6 and O6A. The two *trans* angles in the basal plane of the square pyramidal coordination differ by 8.0(2) and 22.5(2)° from the ideal values. The coordinate bond values are with normal ranges as compared to other similar copper(II) complexes.^[15-18]

The molecular packing structure of the complex is shown in Figure 2. The perchlorate anions are linked to the complex cations through intermolecular O7 δ H7D...O1 hydrogen bonds [O7 δ

H7D = 0.93 Å, H7D...O1^{#1} = 2.28 Å, O7...O1^{#1} = 2.89(1) Å, O7...H7D...O1^{#1} = 123°; symmetry code for #1: 1 ó x, 6 y, 1 6 z].

Thermal Behavior

Thermogravimetric analysis of the complex was carried out in the temperature range 50-800 °C at a heating rate of 10 °C min⁻¹. The TG curve of the complex is given following degradation scheme (Figure 3). The complex decomposes from 180 °C to 520 °C. The final residue is CuO. The observed residue of 16.5% is close to the calculated value of 18.0%.

Antibacterial Activity

From Table 3, the copper complex has medium activity against *E. coli*, and strong activities against *B. subtilis* and *S. aureus*. Especially for *S. aureus*, the complex has even stronger activity than the reference drugs Penicillin and Kanamycin. The complex has stronger activity against *E. coli* than Penicillin, but weaker than Kanamycin. As for *B. subtilis*, the activity of the complex is slightly weaker than the two reference drugs.

CONCLUSIONS

A novel dinuclear copper(II) complex was synthesized by the reaction of 2-acetylpyridine, cyclohexylamine, copper perchlorate and NH₄BF₄ in methanol at room temperature. The ligand underwent an interesting transformation. The reaction mechanism may be further studied by organic and catalytic researchers. The complex has effective antibacterial activities on the strains *B. subtilis*, *E. coli*, and *S. aureus*.

ACKNOWLEDGMENTS

The authors are grateful to Hebei Key Laboratory of Bioinorganic Chemistry and College of Sciences of Agricultural University of Hebei for financial support.

SUPPLEMENTARY MATERIAL

CCDC 980234 contains the supplementary crystallographic data for the complex. The data can be obtained free of charge *via* <http://www.ccdc.cam.ac.uk/conts/retrieving.html>, or from the Cambridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: (+44) 1223-336-033; or e-mail: deposit@ccdc.cam.ac.uk.

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Table 1. Crystallographic data for the complex

Chemical formula	$C_{15}H_{17}ClCuN_2O_7$
Formula weight	436.3
Crystal system	Monoclinic
Space group	$P2_1/n$
Unit cell dimensions	
a (Å)	10.315(2)
b (Å)	12.416(2)
c (Å)	13.964(2)
β (°)	97.494(2)
V (Å ³)	1773.2(6)
Z	4
ρ (g cm ⁻³)	1.634
μ (mm ⁻¹)	1.423
T_{\min}, T_{\max}	0.7838, 0.7940
Reflections collected	9371
Reflections unique	3466
Reflections observed [$I > 2\sigma(I)$]	2339
Parameters	274
Restraints	101
R_1, wR_2 [$I > 2\sigma(I)$]	0.0518, 0.1383
R_1, wR_2 (all data)	0.0847, 0.1549

Goodness-of-fit on F^2	1.045
Highest peak and deepest hole ($e \text{ \AA}^{-3}$)	0.667, 60.612

Table 2. Selected bond lengths (Å) and angles (°) for the complex

Bond lengths (Å)			
Cu1óO6A	1.904(3)	Cu1óO6	1.948(3)
Cu1óN3A	1.973(4)	Cu1óN2	1.978(4)
Cu1óO7	2.238(3)		
Bond angles (°)			
O6óCu1óO6A	78.43(14)	O6AóCu1óN3A	81.87(15)
O6óCu1óN3A	157.55(15)	O6óCu1óN2A	172.04(16)
O6óCu1óN2	93.85(15)	N3óCu1óN2A	106.07(17)
O6óCu1óO7A	101.24(13)	O6óCu1óO7	85.99(13)
N3óCu1óO7A	108.48(15)	N2óCu1óO7	76.04(14)

Symmetry code for A: 1 ó x, ó y, 1 ó z.

Table 3. MIC values ($\mu\text{g mL}^{-1}$) of the tested compounds

	<i>B. subtilis</i>	<i>E. coli</i>	<i>S. aureus</i>
The complex	1.56	12.5	0.78
Penicillin	0.78	>100	3.13
Kanamycin	0.39	6.25	1.56

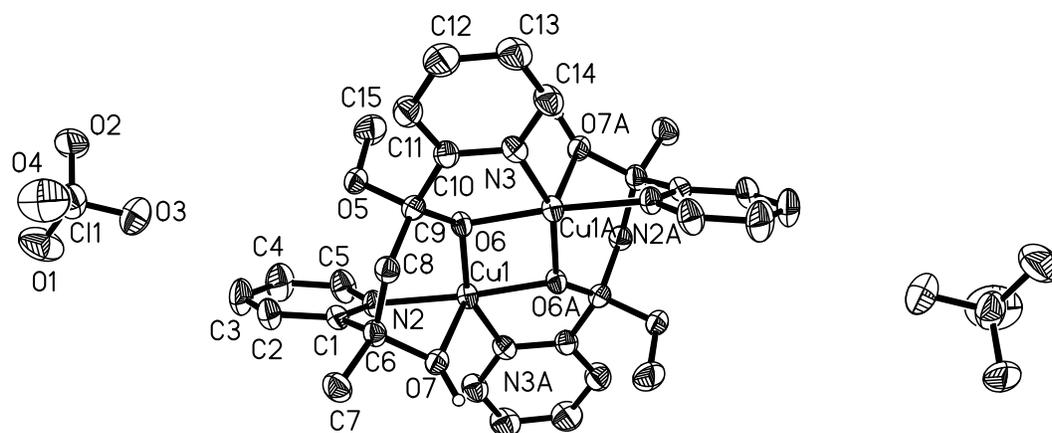


Figure 1. Molecular structure of the complex with the atom labeling scheme. All non-hydrogen atoms are represented at 30% probability thermal ellipsoids. Only the major components of the disordered perchlorate anions are shown.

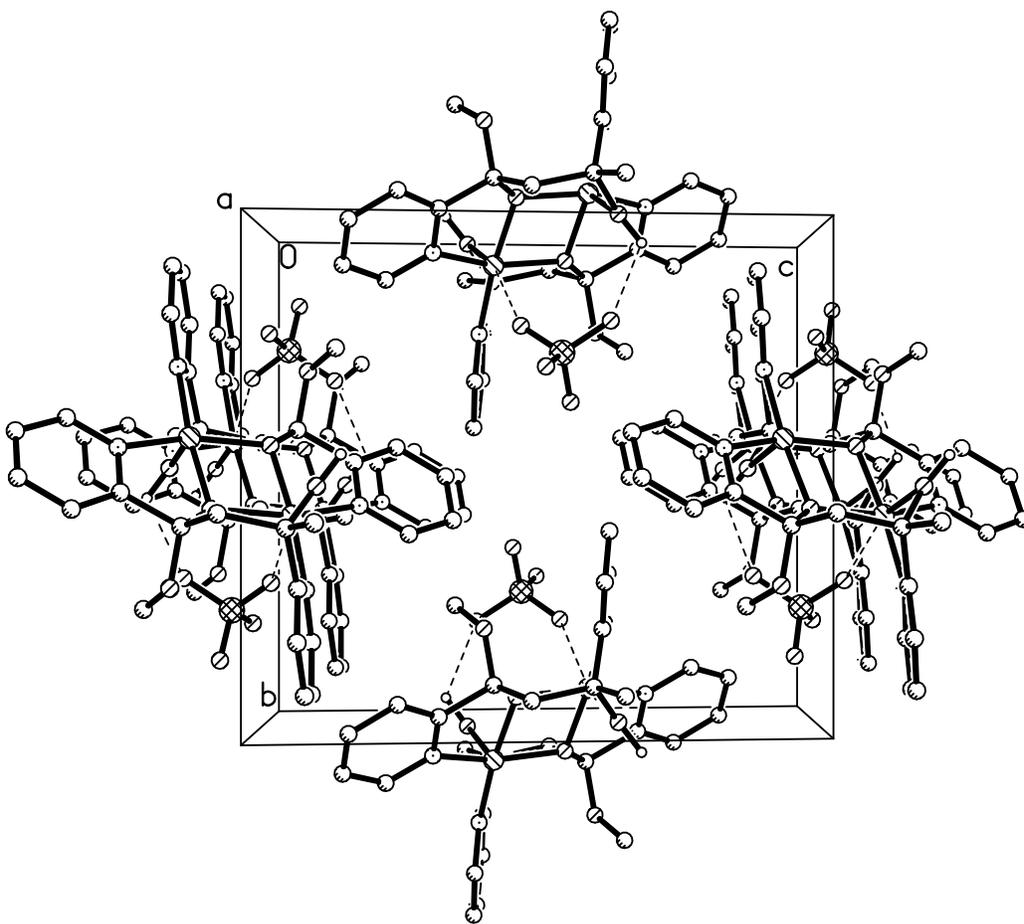


Figure 2. Packing structure of the complex. Hydrogen bonds are shown as thin dashed lines.

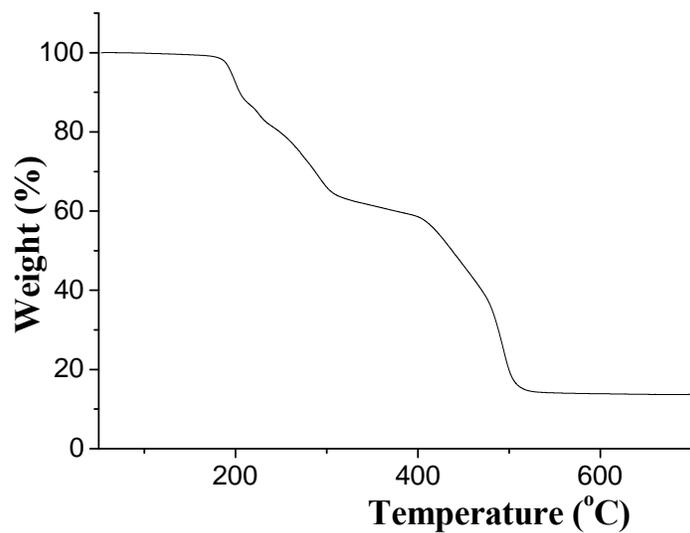


Figure 3. DT curve of the complex.