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CRYSTAL STRUCTURE AND ANTITUMOR ACTIVITIES OF THE DICHLORIDE 2,6-BIS(1-PHENYLBENZIMIDAZOL-2-YL)PYRIDINE COPPER(II) COMPLEX

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A new complex Cu(bpbp)Cl₂·DMF·H₂O is synthesized by treatment of CuCl₂ with 2,6-bis(1phenylbenzimidazol-2-yl)pyidine in DMF. Its structure is characterized by elemental, IR, UV, and single crystal X-ray structure analyses. For the complex: crystal system, triclinic, space group $P\overline{1}$, a = 1.0537(1) nm, b = 1.1735(1) nm, c = 1.3749(2) nm, $\alpha = 112.275(2)^{\circ}$, $\beta = 91.531(2)^{\circ}$, $\gamma = 97.700(2)^{\circ}$, V = 1.553(4) nm³, Z = 2. In a distorted trigonal bipyramidal geometry the Cu(II) ion is coordinated by three nitrogen atoms from 2,6-bis(1-phenylbenzimidazol-2-yl)pyridine and two chloride ions. The antiproliferative activities of the complex are screened by MTT assay against HepG2, Huh7, Eca109, and Eca9706 cancer cells. The complex exhibits specific inhibition on Eca9706 cancer cells with the IC₅₀ value of 28 μ M after 48 h treatment. CCDC: 968927.

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K e y w o r d s : benzimidazole, copper complex, crystal structure, antitumor activities.

The widespread success of cisplatin in the clinical treatment of various types of neoplasias has placed coordination chemistry of metal-based drugs in the frontline in the fight against cancer [1-4]. The cure with cisplatin is still limited by dose-limiting side effects and inherited or acquired resistance phenomena. These problems have stimulated an extensive search and prompted chemists to develop alternative strategies based on different metals with improved pharmacological properties and aimed at different targets. However, the accumulation of metal ions in the body can lead to deleterious effects and unavoidable toxicity. Thus, biodistribution and clearance of the metal complex as well as its pharmacological specificity need to be considered [5]. Benzimidazole derivatives are important pharmacophores in drugs that display a diversity of pharmacological activities, such as anti-inflammatory, antioxidant, gastroprotective, and antiparasitic activities [6]. Many DNA minor groove binders containing one or more benzimidazole heterocycles endowed with promising antitumor and antiparasitic activities have been reported to date [7, 8]. Studies have showed that the intercalating abilities of the complexes depended on the types of metal ions, the ligand donor atoms, the planarity of ligands, and the coordination geometry [9]. In our previous study, two zinc complexes based on 2,6-bis(1phenylbenzimidazol-2-yl)pyridine have been synthesized and evaluated for in vitro anticancer activities [10].

Copper is an essential element for most aerobic organisms employed as a structural and catalytic cofactor, and consequently, it is involved in many biological pathways [11–13]. Several strategies aimed at the development of new anticancer therapeutics targeting the elevated tumor-specific copper

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level have been proposed [14, 15]. In this article, we synthesized a new copper(II) complex based on 2,6-bis(1-phenylbenzimidazol-2-yl)pyridine and investigated its antitumor activities. The result demonstrates that the dichloride 2,6-bis(1-phenylbenzimidazol-2-yl)pyridine copper(II) complex has high proliferation inhibition toward Eca9706 cancer cells *in vitro*.

Experimental. General. *o*-Phenylenediamine, pyridine-2,6-dicaboxyl acid, bromobenzene, and CuCl₂ were purchased from Shanghai Aladdin Reagent Company. All the chemicals and solvents were analytically pure and used without further purification. The analyses (C, H, and N) were made on a Perkin-Elmer 240C elemental analyzer. The solid infrared spectra (IR) were obtained from a Bruker IFS66V vacuum-type FT-IR spectrophotometer in KBr pellets. The UV absorption spectra were recorded on a model UV-240 spectrophotometer (Shimadzu, Japan). The synthesis of 2,6-bis(1-phenylbenzimidazol-2-yl)pyridine (bpbp) was accorded to the literature method [16].

Synthesis of the Cu(bpbp)Cl₂·DMF·2H₂O complex. A solution of CuCl₂ (0.135 g, 1.0 mmol) in 5 ml of water was added to a hot stirred solution of bpbp (0.464 g, 1.0 mmol) in 30 ml of DMF. The reaction mixture was stirred for 20 min at room temperature. After two weeks, blue crystals were obtained from the DMF solution. Yield 0.239 g (80 %). Calculated for $C_{34}H_{30}Cl_2CuN_6O_2$ [Cu(bpbp)Cl₂·DMF·2H₂O] C, 59.26, H, 4.39 % N, 12.20 %: found: C, 59.39 %, 4.26, N, 12.10 %. Selected IR data (KBr, cm⁻¹): 3415, 3342, 3069, 2924, 1668, 1590, 1501, 1456, 1384, 1334, 1300, 1149, 1093, 994, 887,814,753, 680, 647, 613.

X-ray crystallography. The single crystal structure determination of the complex was performed on a Bruker SMART APEX CCD diffractometer equipped with a normal focus, 3 kW sealed tube X-ray source and graphite monochromated Mo K_{α} radiation ($\lambda = 0.71073$ Å) at 173 K, operating at 50 kV and 30 mA. The structures were solved by direct methods using the SHELXTL program. Absorption correction was semi-empirical from the equivalents using the Fourier difference techniques and refined by full-matrix least-squares. All non-hydrogen atoms in both structures were refined by means of anisotropic displacement parameters. All hydrogen atoms were added theoretically. CIF file containing complete information on the studied structure was deposited with CCDC, deposition num-

Table 1

Empirical formula	$C_{34}H_{30}CuN_6O_2$
Formula weight	689.07
Crystal system	Triclinic
Space group	Pī
<i>a</i> , <i>b</i> , <i>c</i> , nm	1.0537(1), 1.1735(1), 1.3749(2)
α , β , γ , deg.	112.275(2), 91.531(2), 97.700(2)
<i>V</i> , nm ³	1.553 (4)
Ζ	2
$d_{\rm c},{\rm g/cm}^{-3}$	1.471
Absorption coefficient, mm ⁻¹	0.918
<i>F</i> (000)	708
Theta range for data collection	1.61 to 25.01
Index ranges	$-12 \le h \le 12, -13 \le k \le 10, -16 \le l \le 14$
Independent reflections	6878 $[R_{int} = 0.0202]$
Reflections collected	5482
Data / restraints / parameters	5294 / 0 / 418
Goodness-of-fit on F^2	1.069
Final <i>R</i> indices $[I \ge 2\sigma(I)]$	$R_1 = 0.0451, \ wR_2 = 0.1330$
<i>R</i> indices (all data)	$R_1 = 0.0671, \ wR_2 = 0.1625$
Largest diff. peak and hole, e/nm^{-3}	0.662, -0.830

Crystal data and structure refinement for the complex

Fig. 1	!.	UV-vis	absorption	spectra	of t	he	free	ligand	and	the	
complex											

ber 968927, and is freely available upon request from the following website: www.ccdc.cam.ac.uk/data_ request/cif. The crystal data are summarized in Table 1.

Cell culture. The cell lines were obtained from American Type Culture Collection (ATCC, Manassas, VA) and maintained in the DMEM medium supplemented with fetal bovine serum (10 %), penicillin (100 units \cdot ml⁻¹), and streptomycin (50 units \cdot ml⁻¹) at 37 °C in a humidified incubator with 5 % CO₂ in the atmosphere.



MTT assay. The effects of the complex on cell proliferation were determined by the MTT assay. Briefly, the cells were seeded in 96-well culture plates at different densities. After 24 h, different concentrations of compounds were added and incubated for the indicated time. Then, 20 μ l per well of the MTT solution (5 mg·ml⁻¹ phosphate buffered saline) was added and incubated for 5 h. The medium was aspirated and replaced with 200 μ l per well of DMSO to dissolve the formazan salt formed. The color intensity of the formazan solution, which reflects the cell growth condition, was measured at 570 nm using a microplate spectrophotometer (VERSA max).

Results and discussion. IR, UV for the complex. The purity of the complex was carefully checked by the elemental analysis. Single crystals suitable for the X-ray crystallographic analysis were obtained by recrystallization from DMF. The IR spectra of the free bpbp ligand and the complex show all absorption bands resulting from the skeletal vibration of benzimidazole. In the complex there is a strong peak at 1668 cm⁻¹, which shows DMF in this complex. The UV-vis absorption was recorded at a concentration of 1.0×10^{-5} mol/l in DMF at room temperature. The UV-vis absorption spectra of the free ligand and the complex are shown in Fig. 1. The ligand has absorption in the range of 270—350 nm and the complex has absorption in the range of 270—400 nm. The maximum ligand absorption is located at about 314 nm, which can be assigned to intraligand $\pi \rightarrow \pi^*$ transitions. There is one more peak observed at 365 nm for the complex which probably arises from the metal—ligand charge transfer.

Crystal structure. The ORTEP drawing for the complex with atom numbering is shown in Fig. 2. The central $Cu_1(II)$ atom has an N_3Cl_2 distorted trigonal bipyramidal geometry. The benzimida-



Fig. 2. Molecular structure of Cu(bpbp)Cl₂ at a 30 % probability. H atoms and solvents were omitted for clarity



Table 2

Selected bong lengths (nm) and bond angles (deg.)

Cu1—N3	0.1999(3)	N3—Cu1—N4	78.44(13)	N3—Cu1—N1	78.16(13)
Cu1—N(1)	0.2038(3)	N4—Cu1—N1	155.05(14)	N3—Cu1—Cl1	160.82(10)
Cu1—Cl2	2.4837(12)	N4—Cu1—Cl1	100.22(10)	N1—Cu1—Cl1	99.03(10)
Cu1—N4	0.2034(3)	N3—Cu1—Cl2	99.17(10)	N4—Cu1—Cl2	97.45(10)
Cu1—Cl1	0.2244(1)	N1—Cu1—Cl2	94.69(10)	Cl1—Cu1—Cl2	99.97(4)

zole rings and the central pyridine ring of the ligand together with Cu_1^{2+} form a plane (r.m.s. = = 0.0645°). The two Cl⁻ ions are located on two sides of the plane formed by N₁, N₃, N₄, and Cu₁. The two substituted phenyl rings (C₇—C₁₂, C₂₀—C₂₅) are inclined with their attached planar benzimidazole rings with the dihedral angles of 88.5° and 87.1° respectively. Selected bond lengths and angles for the complex are listed in Table 2. The average Cu—N bond length is 2.021 Å, which falls into the range of normal Cu—N distances.

The antiproliferative activities of the complex were screened by MTT assay against HepG2, Huh7, Eca109, and Eca9706 cancer cells. As shown in Fig. 3, the copper complex exhibits specific inhibition on Eca9706 cancer cells with IC_{50} of 28 μ M after 48 h treatment, with cisplatin (12 μ M) used as a positive control. The complex has no antitumor activities on HepG2, Huh7, and Eca109 cancer cells. The copper complex is a promising novel complex with the application potential in treatment of Eca9706 cancer.

Conclusions. A new complex $Cu(bpbp)Cl_2 \cdot DMF \cdot H_2O$ was synthesized by treatment of $CuCl_2$ with 2,6-bis(1-phenylbenzimidazol-2-yl)pyridine (bpbp) in DMF at room temperature. The complex was characterized by single X-ray crystal structure analyses. In the complex, the distorted trigonal bipyramidal geometry of the Cu(II) ion is coordinated by three nitrogen atoms from 2,6-bis(1-phenylbenzimidazol-2-yl)pyridine and two chloride ions. The antiproliferative activities of the complex were screened by MTT assay against HepG2, Huh7, Eca109, and Eca9706 cancer cells. The copper complex exhibits specific inhibition on Eca9706 cancer cells.

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