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**CHARACTERIZATION AND CYTOTOXIC PROPERTY OF A LADDER-LIKE
POLYMERIC SILVER(I) COMPLEX DERIVED FROM
3-AMINOPYRAZINE-2-CARBOXYLIC ACID****G.-S. Li, H.-L. Zhang***Key Laboratory of Surface & Interface Science of Henan, School of Material & Chemical Engineering, Zhengzhou University of Light Industry, Zhengzhou 450002, P. R. China*

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A novel ladder-like polymeric silver(I) complex, $[\text{Ag}_2\text{L}_2]_n \cdot 2n\text{H}_2\text{O}$, where L is 3-aminopyrazine-2-carboxylate, was obtained by the reaction of 3-aminopyrazine-2-carboxylic acid and silver oxide in aqueous ammonia. The complex was characterized by elemental analysis, IR spectra and single crystal X-ray determination. The smallest repeat unit contains a $[\text{Ag}_2\text{L}_2]$ moiety and two water molecules. The $\text{Ag} \cdots \text{Ag}$ distance is 3.176(1) nmol. Each Ag atom is in a Y-shaped coordination, with one carboxylate O and two pyrazine N atoms from two ligands. In the crystal structure of the complex, the dinuclear silver moieties are linked through 3-aminopyrazine-2-carboxylate ligands, to form 1D ladder-like chain along the *b* axis. The water molecules are linked to the silver chain through hydrogen bonds. There are $\pi \cdots \pi$ interactions between the chains. The complex showed effective cytotoxic property on human lung cancer cell line A549.

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Keywords: silver complex, polymeric complex, 3-aminopyrazine-2-carboxylate, crystal structure, cytotoxicity.**INTRODUCTION**

Silver complexes with carboxylate ligands have received much attention in coordination and supramolecular chemistry [1–5]. Owing to the versatile coordination geometry of silver, coordination numbers from two to six are possible [6–10], and because of the relatively weak nature of many Ag-ligand interactions, such complexes are particularly susceptible to the influence of weaker supramolecular forces. Thus, one can not precisely predict what structures will be finally formed for the silver(I) complexes, and more work needs to be carried out to understand the influence effects of such complexes, which has becoming an interesting topic in supramolecular chemistry. In recent years, the self-assembly and structures of a number of silver complexes have been studied. Most of the complexes show interesting luminescence property [11–13] and biological activities [14–17]. Compared to other applications, the cytotoxicity property of the silver complexes received particular attention [18–20]. 3-Aminopyrazine-2-carboxylic acid (HL) is a hopeful ligand containing carboxylic group, amine and pyrazine nitrogen atoms. Some nickel [21], cobalt [22], mercury [23], zinc [24], manganese [25] and lanthanide [26] complexes derived from HL have been reported. However, no silver complex has been reported so far. In the present work, a novel ladder-like polymeric silver(I) complex, $[\text{Ag}_2\text{L}_2]_n \cdot 2n\text{H}_2\text{O}$, is presented.

EXPERIMENTAL

Materials and measurements. 3-Aminopyrazine-2-carboxylic acid and silver oxide were purchased from Fluka and used as received. The solvents used were of reagent grade. Elemental analyses were carried out using a Perkin-Elmer 2400 II elemental analyser. The infrared spectrum was recorded on a Perkin-Elmer FT-IR spectrophotometer with a KBr disc. Thermal analysis was performed with a Pyris 1 thermogravimetric analyser. The X-ray diffraction was carried out on a Bruker SMART 1000 CCD area diffractometer at 298(2) K.

Synthesis of the complex. 3-Aminopyrazine-2-carboxylic acid (0.28 g, 2 mmol) and Ag₂O (0.23 g, 1 mmol) were mixed and stirred in a 30 % aqueous ammonia (30 mL). The mixture was stirred at room temperature until all solid dissolved. The clear colorless solution was kept still at room temperature in dark for several days, to give well-shaped block-like single crystals. Yield: 38 %. Anal. Calcd. for C₁₀H₁₂Ag₂N₆O₆ (%): C 22.7, H 2.3, N 15.9. Found (%): C 22.9, H 2.3, N 15.8. IR (KBr, cm⁻¹): 3407m, 3265w, 1604s, 1555m, 1468w, 1430w, 1381m, 1326w, 1232w, 1165m, 910w, 823w, 557w.

X-ray crystallography. A suitable single crystal of the complex was mounted on the top of a glass fiber. Graphite-monochromatized MoK_α radiation ($\lambda = 0.71073$ nmol) and the ω scan technique were used to collect the diffraction data. Absorption correction was applied with SADABS [27]. The structure of the complex was solved with direct method and refined with a full-matrix least-squares technique with SHELXTL [28]. Anisotropic thermal parameters were applied to all non-hydrogen atoms. The hydrogen atoms of the water molecules were located from electronic density maps, with O \cdots H and H \cdots H distances restrained to 0.85(1) mmol and 1.37(2) mmol, respectively. The other hydrogen atoms were generated geometrically. The crystallographic data and the details of the data collection and refinement for the complex are listed in Table 1. Selected bond lengths and angles are given in Table 2. Hydrogen bonding information is given in Table 3. Crystallographic data for the complex has been deposited with the Cambridge Crystallographic Data Centre (CCDC 1008305).

Table 1

Crystallographic data for the complex

Empirical formula	C ₁₀ H ₁₂ Ag ₂ N ₆ O ₆
<i>FW</i>	528.0
Crystal shape, colour	Block, colorless
Crystal size, mm	0.18×0.17×0.12
Crystal system	Monoclinic
Space group	<i>P</i> 2 ₁ / <i>c</i>
$\lambda(\text{MoK}_\alpha)$, Å	0.71073
<i>T</i> , K	298(2)
μ , mm ⁻¹ (MoK _α)	2.850
Unit cell dimensions <i>a</i> , <i>b</i> , <i>c</i> , Å; β , deg.	13.017(2), 7.1466(9), 15.064(2); 94.518(2)
<i>V</i> , Å ³ ; <i>Z</i>	1397.0(3); 4
<i>T</i> _{min} / <i>T</i> _{max}	0.6279 / 0.7260
No. of measured / unique reflections	7709 / 2623
No. of observed reflections [<i>I</i> ≥ 2σ(<i>I</i>)]	2305
Data / restraints / parameters	2623 / 9 / 229
<i>R</i> _{int}	0.0336
<i>F</i> (000)	1024
Goodness of fit on <i>F</i> ²	1.305
<i>R</i> ₁ , <i>wR</i> ₂ [<i>I</i> ≥ 2σ(<i>I</i>)]	0.0822, 0.1761
<i>R</i> ₁ , <i>wR</i> ₂ (all data)	0.0908, 0.1795
Largest and deepest diff. peak and hole, e/Å ³	1.349 and -1.272

Table 2

Selected bond lengths (Å) and angles (deg.) for the complex

Bond lengths				Bond angles			
Ag1—N1	2.212(10)	Ag1—O1	2.500(10)	N2—Ag1—N1A	171.0(4)	N2—Ag1—O1A	107.2(3)
Ag1—N2A	2.183(10)	Ag2—N5A	2.202(10)	N1—Ag1—O1	70.2(3)	N5—Ag2—N4A	169.1(4)
Ag2—N4	2.241(9)	Ag2—O3	2.573(10)	N5—Ag2—O3A	105.7(3)	N4—Ag2—O3	69.7(3)

Symmetry code for A: $x, 1+y, z$.

Cytotoxicity assay. Human lung carcinoma A549 cells were cultured in an F12 K medium supplemented with 10 % heat inactivated fetal bovine serum (FBS), 2 mM glutamine, 100 U/mL penicillin, and 100 µg/mL streptomycin, and maintained at 37 °C in humidified atmosphere of 5 % CO₂. MTT assay was conducted and modified as described in the literature [29]. A549 cells (2.5×10^3 cells) were seeded on 96-well microtitre plates in F12 K medium with 10 % FBS and incubated overnight. The cell culture medium was replaced by the different dose of compounds solution, and then the cells were cultured for another 72 h. The MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] reagent was added to the cell supernatant for a final concentration of 0.5 mg/mL of MTT. After 3 h the cell culture medium was removed. Formazan crystals in adherent cells were dissolved in 200 µL DMSO and the absorbance of the formazan solution was measured. Each compound was tested in triplicate and the experiments were repeated three times.

Table 3

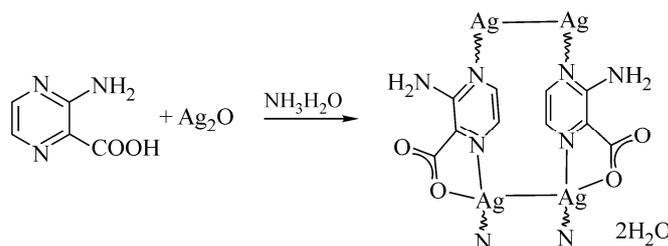
Hydrogen-bonding geometry (Å, deg.)

D—H···A	D—H	H···A	D···A	D—H···A
N3—H3A···O1 ⁱ	0.86	2.06	2.915(14)	177
N6—H6A···O3 ⁱ	0.86	2.24	3.078(15)	165
N3—H3B···O2	0.86	2.01	2.657(15)	131
N6—H6B···O4	0.86	1.99	2.644(16)	132
O5—H5A···O3 ⁱⁱ	0.85(1)	2.02(5)	2.820(14)	156(11)

Symmetry codes: ⁱ $x, -1+y, z$; ⁱⁱ $x, 3/2-y, 1/2+z$.

RESULTS AND DISCUSSION

Synthesis. The synthetic procedure of the complex is described as follows:



In the present complex, the Ag atom combines with the pyrazine N and carboxylate O atoms, instead of the amino N atom. This is due to the strain of the five-membered chelate ring is weaker than the four-membered chelate ring. Elemental analyses of the complex are in good agreement with the expected values. Even though most silver complexes are sensitive to light, this complex is very stable.

Infrared spectrum of the complex. The medium and broad absorption centered at 3407 cm^{-1} can be assigned to the vibration of the water molecules. The weak and sharp absorption at 3265 cm^{-1} comes from the N—H vibrations of the amino groups. The $\nu_{\text{as}}(\text{COO})$ is related to the strong band observed at 1604 cm^{-1} , whereas the $\nu_{\text{s}}(\text{COO})$ is attributed to the medium band observed at 1381 cm^{-1} . This gives rise to a $\Delta\nu$ of 223 cm^{-1} characteristic of the monodentate coordination of the carboxylate group [30].

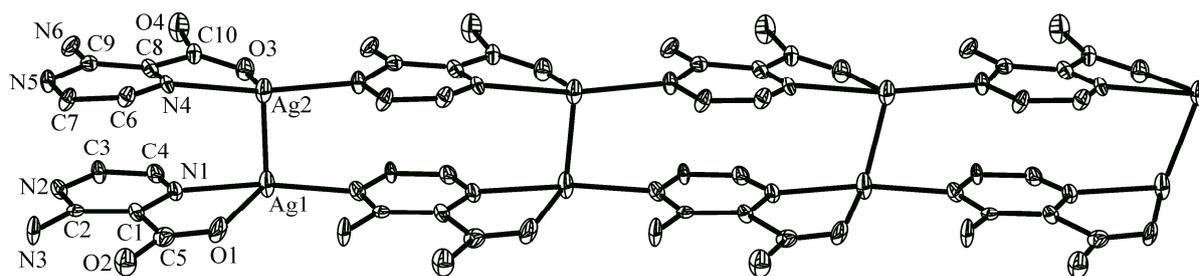


Fig. 1. Molecular structure of the silver complex with 30 % probability level

Structure description of the complex. The molecular structure of the complex is shown in Fig. 1. The smallest repeat unit contains a $[Ag_2L_2]$ moiety and two water molecules. The Ag \cdots Ag distance is 3.176(1) mmol. Each Ag atom is in a Y-shaped coordination, with one carboxylate O and two pyrazine N atoms from two ligands. The Ag1 and Ag2 atoms deviate from the least-squares planes defined by the corresponding three donor atoms by 0.168(2) mmol and 0.198(2) mmol, respectively. The distortion of the coordination can be observed from the bond lengths and angles related to the Ag atom. The Ag—O bonds are much longer than the Ag—N bonds. Due to the strain created by the five-membered chelated rings Ag1—N1—C1—C5—O1 and Ag2—N4—C8—C10—O3, the angles N1—Ag1—O1 and N4—Ag2—O3 are much smaller. Yet, both Ag—O and Ag—N bonds are comparable to those observed in similar silver complexes [31, 32]. The dihedral angle between the least-squares planes of N1—C1—C2—N2—C3—C4 and O1—C5—O2 is 6.4(3) $^\circ$, and that between the least-squares planes of N4—C6—C7—N5—C9—C8 and O3—C10—O4 is 17.4(3) $^\circ$. The dinuclear silver moieties are linked through 3-aminopyrazine-2-carboxylate ligands, to form 1D ladder-like chain along the *b* axis (Fig. 2). The water molecules are linked to the silver chain through intermolecular N—H \cdots O and O—H \cdots O hydrogen bonds (Table 3). There are $\pi\cdots\pi$ interactions between the chains (Table 4).

Thermal analysis. Fig. 3 shows the TG-DT curves of the complex. The complex undergoes three steps of decomposition. The first step, starting from 80 $^\circ$ C to 140 $^\circ$ C, corresponds to the loss of water molecules. The weight loss of 7.3 % is close to the calculated value of 6.9 %. The second step, starting

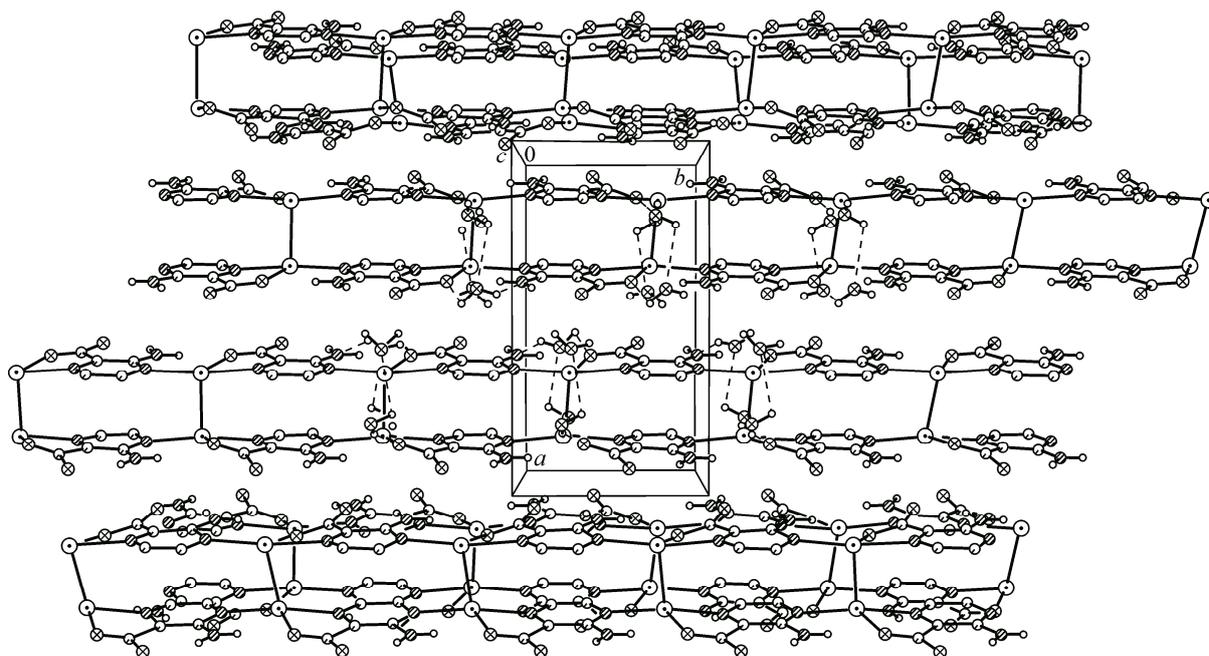


Fig. 2. Molecular packing of the complex, viewed along the *c* axis. Hydrogen bonds are shown as dashed lines

Table 4

Parameters between the planes for the complex

Cg	Distance between ring centroids, Å	Dihedral angle, deg.	Perpendicular distance of Cg(I) on Cg(J), Å	Perpendicular distance of Cg(I) on Cg(J), Å	Slippage
Cg(1)—Cg(1) ⁱⁱⁱ	3.410	0	3.274	3.274	0.954
Cg(1)—Cg(2) ⁱⁱⁱ	4.340	5	3.244	3.339	
Cg(1)—Cg(3)	3.674	6	3.215	-3.344	
Cg(2)—Cg(3)	3.462	2	3.243	-3.276	
Cg(3)—Cg(3) ^{iv}	4.895	0	-3.047	-3.047	3.831
Cg(3)—Cg(3) ^v	4.726	0	-3.162	-3.162	3.512

Symmetry codes: ⁱⁱⁱ 1-x, 1-y, 1-z; ^{iv} -x, -y, 1-z; ^v -x, 1-y, 1-z.

Cg(1), Cg(2), and Cg(3) are the centroids of Ag1—O1—C5—C1—N1, N1—C1—C2—N2—C3—C4, and N4—C6—C7—N5—C9—C8, respectively.

from 250 °C to 380 °C, corresponds to the loss of the ligands, and the formation of Ag₂O. The weight loss of 51 % agrees well with the calculated value of 49 %. Then, Ag₂O decompose to give Ag as the final product.

Cytotoxic property. A549 cells were used by MTT method to study the growth inhibitory effects of the complex on lung cancer cells. The IC₅₀ value for the complex is 10.3±1.5 μM. As a comparison, the silver nitrate and cisplatin showed IC₅₀ values of 8.7±0.9 μM and 6.9±1.1 μM, and the ligand itself showed no activity. Thus, the cytotoxic property of the complex is weaker than the silver nitrate and cisplatin. However, considering the obvious side effects of the silver nitrate and cisplatin, the present silver complex is also a promising drug, and deserves further study.

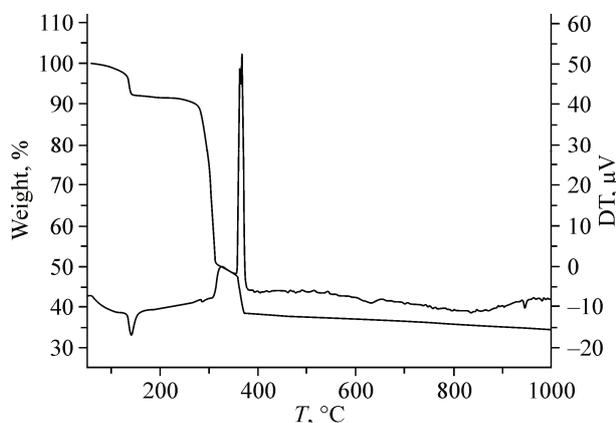


Fig. 3. TG curve of the complex

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