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Key indicators

Single-crystal X-ray study  
T = 298 K  
Mean  $\sigma(C-C)$  = 0.007 Å  
R factor = 0.044  
wR factor = 0.118  
Data-to-parameter ratio = 12.3

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

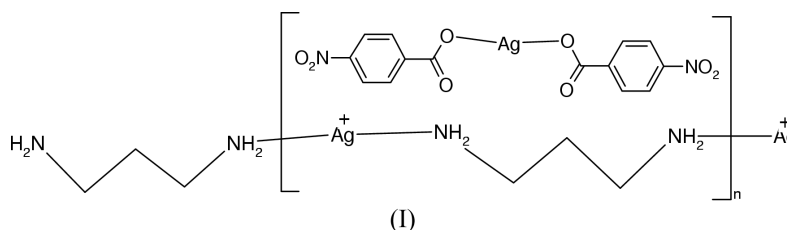
catena-Poly[[silver(I)- $\mu$ -propane-1,3-diamine- $\kappa^2N:N'$ ] bis(4-nitrobenzoato- $\kappa O$ )argentate(I)]

In the title compound,  $\{[Ag(C_3H_{10}N_2)][Ag(C_7H_4NO_4)_2]\}_n$ , one of three independent  $Ag^I$  atoms forms a mononuclear complex with a nearly linear environment, coordinated by two O atoms from two 4-nitrobenzoate anions. The other two Ag atoms lie on inversion centers and are coordinated in a linear configuration by two N atoms from two propane-1,3-diamine ligands, giving zigzag polymeric chains with an  $[-Ag-N-C-C-C-N-]_n$  backbone running along the [011] direction.

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Comment

Silver(I) complexes with carboxylate anions as counter-ions or ligands are a group of metal compounds which, due to their wide usage in many fields, have been structurally characterized for many years (Nomiya *et al.*, 2000; Kristiansson, 2001). Recently, we have reported a few silver(I)-carboxylate complexes with various amines and imines, all of which were structurally characterized (Zhu *et al.*, 1999, 2000; Zheng *et al.*, 2001; Usman *et al.*, 2003). As an extension of our work on the characterization of silver compounds, the structure of the title compound, (I), is reported here.



The title compound, (I), is a polymeric silver(I) complex (Fig. 1). The smallest repeat unit for the complex contains a propane-1,3-diamine-silver(I) cation segment and a bis(4-nitrobenzoato)silver(I) anion. Atom Ag1 in the anion is in a nearly linear coordination environment and is coordinated by two O atoms from two 4-nitrobenzoate anions. Each 4-nitrobenzoate anion coordinates to the Ag atom through one O

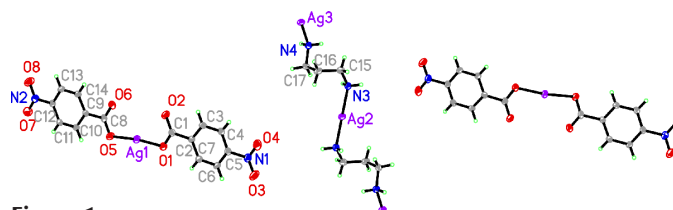
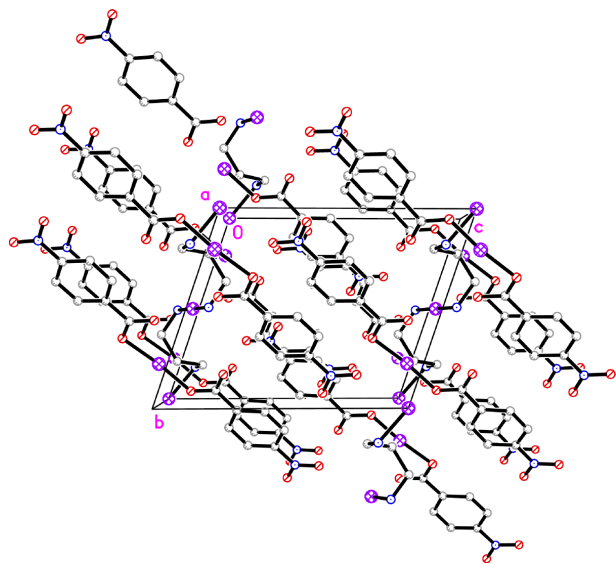


Figure 1 The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. Unlabelled atoms are at the symmetry position  $-x, 2 - y, 2 - z$ .



**Figure 2**  
The crystal packing of (I), viewed along the *a* axis. H atoms have been omitted.

atom of the carboxylate group. The O1–Ag1–O5 angle [172.24 (13)°, Table 1] deviates slightly from the ideal value of 180°, which is probably due to the weak interactions involving atoms O2 and O6 of the 4-nitrobenzoate anion, with an average Ag···O distance of 2.972 (5) Å; these pull Ag<sup>I</sup> in towards the direction of atoms O2 and O6. There are intermolecular N–H···O hydrogen bonds (Table 2). Ag2 and Ag3 both lie on inversion centers and are coordinated in a linear configuration by two N atoms from two propane-1,3-diamine ligands, giving zigzag polymeric cation chains with an [–Ag–N–C–C–C–N–]<sub>n</sub> backbone running along the [011] direction (Fig. 2). All the Ag–O and Ag–N bond lengths are comparable to the values observed in a similar silver(I) complex (Zhu *et al.*, 2003).

## Experimental

Silver 4-nitrobenzoate (0.1 mmol, 27.4 mg) and propane-1,3-diamine (0.1 mmol, 7.4 mg) were dissolved in an aqueous ammonia solution (10 ml, 30%). The mixture was stirred for about 10 min at room temperature to obtain a clear colorless solution. The resulting solution was kept in the dark, and after slow evaporation of the solvent over a period of 8 d, crystals of (I) were isolated, washed three times with water and dried in a vacuum desiccator using anhydrous CaCl<sub>2</sub> (yield 73%). Elemental analysis calculated: C 32.8, H 2.9, N 9.0%; found: C 32.6, H 2.9, N 9.1%.

### Crystal data

[Ag(C<sub>3</sub>H<sub>10</sub>N<sub>2</sub>)] [Ag(C<sub>7</sub>H<sub>4</sub>NO<sub>4</sub>)<sub>2</sub>]  
*M<sub>r</sub>* = 622.09  
 Triclinic, *P* $\bar{1}$   
*a* = 9.530 (5) Å  
*b* = 9.561 (6) Å  
*c* = 11.609 (5) Å  
 $\alpha$  = 108.472 (5)°  
 $\beta$  = 93.918 (6)°  
 $\gamma$  = 92.673 (5)°  
*V* = 998.3 (9) Å<sup>3</sup>

*Z* = 2  
*D<sub>x</sub>* = 2.071 Mg m<sup>−3</sup>  
 Mo *K*α radiation  
 Cell parameters from 3715 reflections  
 $\theta$  = 2.3–26.5°  
 $\mu$  = 2.02 mm<sup>−1</sup>  
*T* = 298 (2) K  
 Block, colorless  
 0.32 × 0.26 × 0.18 mm

### Data collection

Bruker SMART CCD area-detector diffractometer  
 $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
*T<sub>min</sub>* = 0.553, *T<sub>max</sub>* = 0.699  
 5306 measured reflections

3493 independent reflections  
 2999 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.021  
 $\theta_{\max}$  = 25.0°  
*h* = −11 → 10  
*k* = −11 → 11  
*l* = −13 → 10

### Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.044  
*wR* (*F*<sup>2</sup>) = 0.118  
*S* = 1.07  
 3493 reflections  
 283 parameters

H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0841P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 3.45 \text{ e } \text{Å}^{-3}$   
 $\Delta\rho_{\min} = -1.03 \text{ e } \text{Å}^{-3}$

**Table 1**

Selected geometric parameters (Å, °).

Ag1–O1	2.087 (5)	Ag2–N3	2.151 (5)
Ag1–O5	2.124 (5)	Ag3–N4	2.150 (6)
O1–Ag1–O5	172.24 (13)	N4 <sup>ii</sup> –Ag3–N4	180 (1)
N3 <sup>i</sup> –Ag2–N3	180 (1)		

Symmetry codes: (i)  $-x, 2-y, 2-z$ ; (ii)  $1-x, 1-y, 2-z$ .

**Table 2**

Hydrogen-bonding geometry (Å, °).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
N4–H4B···O6 <sup>ii</sup>	0.90	2.13	3.008 (7)	167
N4–H4A···O2 <sup>ii</sup>	0.90	2.39	3.203 (9)	150
N3–H3B···O6 <sup>iii</sup>	0.90	2.14	3.025 (7)	167
N3–H3A···O2 <sup>iii</sup>	0.90	2.40	3.204 (8)	149

Symmetry codes: (ii)  $1-x, 1-y, 2-z$ ; (iii)  $x-1, y, z$ .

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C–H distances of 0.93–0.97 Å and N–H distances of 0.90 Å, and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C/N})$ . The maximum residual density is located 1.86 Å from atom Ag1, while the minimum residual density is 0.91 Å from Ag1.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *SHELXTL*.

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