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Key indicators

Single-crystal X-ray study  
 T = 193 K  
 Mean  $\sigma(\text{C}-\text{C}) = 0.010 \text{ \AA}$   
 H-atom completeness 93%  
 Disorder in solvent or counterion  
 R factor = 0.070  
 wR factor = 0.136  
 Data-to-parameter ratio = 11.3

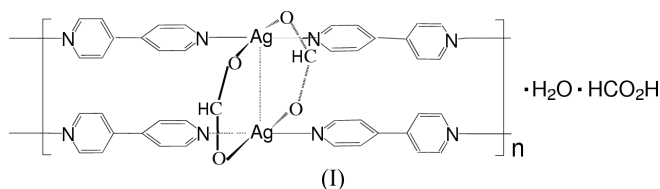
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Poly[[silver(I)- $\mu$ -4,4'-bipyridine- $\mu$ -formate] monohydrate formic acid solvate]

The reaction of silver formate with HCO<sub>2</sub>H and 4,4'-bipyridine in CH<sub>3</sub>CN produced the title compound, [Ag(CHO<sub>2</sub>)(C<sub>10</sub>H<sub>8</sub>N<sub>2</sub>)]·H<sub>2</sub>O·CH<sub>2</sub>O<sub>2</sub>. In the structure, each 4,4'-bipyridine moiety acts as a bidentate ligand that binds two Ag atoms, thus forming a linear chain. The formate ions bridge the adjacent chains through Ag–O bonds [Ag–O = 2.709 (6)–3.089 (5) Å] to form molecular ladders. The molecular ladders are stacked into layers and are interconnected by O–H···O hydrogen bonds.

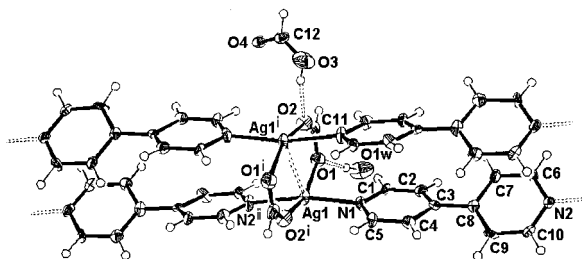
Comment

A large number of silver salts furnish polydimensional supramolecular compounds by the direct reaction of these salts with symmetrically bridging ligands (Batten & Robson, 1998; Hagrman *et al.*, 1999; Tong, Zheng & Chen, 2000), and some of the anionic groups themselves are also able to function as a bridge (Bertelli *et al.*, 1997; Tong *et al.*, 1999; Tong, Chen & Ng, 2000; Robinson & Zaworotko, 1995; Sharma *et al.*, 1998; Yaghi & Li, 1996). The carboxylate system has also been well explored, except for the formate homolog. However, only a few complexes containing metal/formate/4,4'-bipyridine have been reported so far. We report here the structure of the title silver–4,4'-bipyridine–formate complex, (I).



In (I), each Ag atom adopts a distorted square-planar coordination, a geometry that is relatively uncommon (Lippert & Neugebauer, 1982; Villanneau *et al.*, 1998; Tong, Chen & Ng, 2000). Each Ag atom is bonded to two symmetry-related 4,4'-bipyridine units [Ag–N = 2.167 (5) Å], with the donor ends of the ligand occupying *trans* positions of the square [N–Ag–N = 168.1 (2)°], thus generating linear chains. In adjacent chains, two neighboring Ag atoms are bridged by two formate ions through Ag–O coordination, resulting in the formation of a molecular ladder. The Ag–O distances lie in the range 2.709 (6)–3.089 (5) Å (Table 1) and are much longer than those of dinuclear Ag–carboxylate complexes (Bowmaker *et al.*, 2000; Chen & Mak, 1991; Huang *et al.*, 1991). The Ag atoms are also linked by formate-supported Ag–Ag interactions with a separation of 3.1748 (14) Å, which is similar to those found in dinuclear Ag<sup>I</sup>–carboxylate complexes (Chen & Mak, 1991; Huang *et al.*, 1991). The

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**Figure 1**

A part of the polymeric chain, illustrating the coordination environment of the Ag atom in (I). Displacement ellipsoids are shown at the 50% probability level. Only one component of the disordered formic acid molecule is shown. H atoms are drawn as small spheres of arbitrary radii. [Symmetry codes: (i)  $\frac{1}{2} - x, \frac{1}{2} - y, 1 - z$ ; (ii)  $\frac{1}{2} + x, \frac{1}{2} - y, z - \frac{1}{2}$ ]

geometry of the  $\text{HCO}_2^-$  and bipyridine ligands agrees with those observed in other systems (Manson *et al.*, 2001; Han *et al.*, 2000).

The molecular ladders are stacked into layers, which are further interconnected by hydrogen-bonding networks formed by formic acid and water molecules, resulting in a three-dimensional network (Table 2).

## Experimental

To a suspension of silver formate (0.046 g, 0.2 mmol) in acetonitrile (2 ml) was added formic acid (0.042 g, 0.8 mmol) to furnish a clear solution. To this solution was added 4,4'-bipyridine (0.063 g, 0.4 mmol) in the same solvent (2 ml). The precipitate that formed was removed by filtration; the filtrate, when set aside at room temperature, afforded colorless prism-shaped crystals, which were collected by filtration, washed with methanol and diethyl ether, and dried in a vacuum. Yield 0.050 g, 64.2%.

### Crystal data

$[\text{Ag}(\text{CHO}_2)(\text{C}_{10}\text{H}_8\text{N}_2)] \cdot \text{H}_2\text{O} \cdot \text{CH}_2\text{O}_2$	$D_x = 1.834 \text{ Mg m}^{-3}$
$M_r = 373.11$	Mo $K\alpha$ radiation
Monoclinic, $C2/c$	Cell parameters from 3186 reflections
$a = 17.848 (4) \text{ \AA}$	$\theta = 3.1\text{--}27.5^\circ$
$b = 16.863 (3) \text{ \AA}$	$\mu = 1.51 \text{ mm}^{-1}$
$c = 9.485 (3) \text{ \AA}$	$T = 193 (2) \text{ K}$
$\beta = 108.79 (1)^\circ$	Prism, colorless
$V = 2702.6 (12) \text{ \AA}^3$	$0.35 \times 0.18 \times 0.06 \text{ mm}$
$Z = 8$	

### Data collection

Rigaku Mercury area-detector diffractometer	2365 independent reflections
$\omega$ scans	2176 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (Blessing, 1995; Jacobson, 1998)	$R_{\text{int}} = 0.054$
$T_{\text{min}} = 0.784, T_{\text{max}} = 0.915$	$\theta_{\text{max}} = 25.0^\circ$
8855 measured reflections	$h = -21 \rightarrow 21$
	$k = -16 \rightarrow 20$
	$l = -11 \rightarrow 9$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0427P)^2 + 20.891P]$
$R[F^2 > 2\sigma(F^2)] = 0.070$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.136$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.37$	$\Delta\rho_{\text{max}} = 1.12 \text{ e \AA}^{-3}$
2365 reflections	$\Delta\rho_{\text{min}} = -0.53 \text{ e \AA}^{-3}$
209 parameters	
H-atom parameters constrained	

**Table 1**

Selected geometric parameters ( $\text{\AA}, ^\circ$ ).

Ag1—N1	2.167 (5)	Ag1—O2 <sup>i</sup>	3.089 (5)
Ag1—O1	2.709 (6)	Ag1—Ag1 <sup>i</sup>	3.1748 (14)
Ag1—O2	2.809 (5)		
N2 <sup>ii</sup> —Ag1—N1	168.1 (2)	N2 <sup>ii</sup> —Ag1—O2 <sup>i</sup>	88.47 (18)
N2 <sup>ii</sup> —Ag1—O1	91.2 (2)	N1—Ag1—O2 <sup>i</sup>	87.80 (18)
N1—Ag1—O1	95.83 (19)	O1—Ag1—O2 <sup>i</sup>	162.27 (16)

Symmetry codes: (i)  $\frac{1}{2} - x, \frac{1}{2} - y, 1 - z$ ; (ii)  $\frac{1}{2} + x, \frac{1}{2} - y, z - \frac{1}{2}$ .

**Table 2**

Hydrogen-bonding geometry ( $\text{\AA}, ^\circ$ ).

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
O3—H3O <sup>o</sup> ···O2	0.84	1.75	2.57 (2)	167
O3'—H3O <sup>o</sup> ···O2	0.84	1.70	2.52 (3)	163
O1W—H1W1···O1	0.83	1.93	2.74 (1)	166

The formic acid molecule is disordered over two positions with populations of 50% each. The disordered atoms were anisotropically refined with their  $U_{ij}$  components approximated to isotropic behavior and with suitable restraints for C—O distances. Only one of the H atoms of the water molecule was located in a difference map, and this atom was allowed to ride on the parent atom with the O—H distance fixed at 0.83  $\text{\AA}$ . All other H atoms were placed at calculated positions (O—H = 0.84  $\text{\AA}$  and C—H = 0.95  $\text{\AA}$ ) and were included in the refinement in the riding-model approximation with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{O})$ . The final difference Fourier map had a peak larger than  $1 \text{ e \AA}^{-3}$  at about 1  $\text{\AA}$  from atom Ag1.

Data collection: *CrystalClear* (Rigaku/MSK, 2001); cell refinement: *CrystalClear*; data reduction: *CrystalStructure* (Rigaku/MSK, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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