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## Key indicators

Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
Disorder in solvent or counterion
$R$ factor $=0.030$
$w R$ factor $=0.084$
Data-to-parameter ratio $=15.6$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## catena-Poly[[bromatosilver(I)- $\mu$-4,4'-bipyridine] 2.5-hydrate]

In the crystal structure of $\left\{\left[\mathrm{Ag}\left(\mathrm{BrO}_{3}\right)\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right)\right] \cdot 2.5 \mathrm{H}_{2} \mathrm{O}\right\}_{n}$, the Ag atom is coordinated by the N atoms of two different spacer donor ligands $[\mathrm{Ag}-\mathrm{N}=2.147$ (2) and 2.152 (2) $\AA$, and $\left.\mathrm{N}-\mathrm{Ag}-\mathrm{N}=172.0(1)^{\circ}\right]$, but is connected only weakly to the O atoms of two different bromate counter-ions $[\mathrm{Ag}-\mathrm{O}=$ 2.877 (3) and 2.972 (3) $\AA$, and $\mathrm{O}-\mathrm{Ag}-\mathrm{O}=162.3$ (1) $\left.{ }^{\circ}\right]$; the coordination geometry of the Ag atom is approximately planar. The $\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right) \mathrm{Ag}$ chains, which propagate along [10 $\left.\overline{1}\right]$, are cross-linked in the $b$ direction by bromate anions and $\mathrm{Ag} \cdots \mathrm{Ag}[3.3197$ (6) $\AA$ A ] interactions, thus forming sheets.

## Comment

The 4,4'-bipyridine spacer ligand forms a large number of complexes with the silver(I) cation; in some of these complexes, the $\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right) \mathrm{Ag}$ units are assembled into linear chains, and the inorganic counter-ions interact, although somewhat weakly, with the metal atoms, e.g. $\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right) \mathrm{Ag}\left(\mathrm{NO}_{3}\right)$, which crystallizes in Pmma ( Bi et al., 2002) and Fdd2 (Yaghi \& Li, 1996) modifications. For the Pmma structure, if two other Ag...O contacts [2.835 (6) and 2.747 (6) $\AA$ ] are considered as bonding interactions, the geometry approximates to that of a rectangular plane. The nitrite salt, which is also water-free, (Blake et al., 1999), exhibits shorter and stronger Ag...O interactions [Ag. $\cdot \mathrm{O}=$ 2.487 (6) Å]. The present study of the bromate salt continues studies on these complexes with other inorganic anions; an earlier study of the dihydrogen orthophosphate reveals the existence of a phosphoric acid molecule in the formula unit (Tong et al., 2000).

In the title compound, $\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right) \mathrm{Ag}\left(\mathrm{BrO}_{3}\right) \cdot 2.5 \mathrm{H}_{2} \mathrm{O}$, (I) (Fig. 1), the Ag atom is coordinated by the N atoms of two donor ligands $[\mathrm{Ag}-\mathrm{N}=2.147$ (2) and 2.152 (2) $\AA$, and $\mathrm{N}-$ $\left.\mathrm{Ag}-\mathrm{N}=172.0(1)^{\circ}\right]$ in a nearly linear geometry. The rings are twisted by $30.9(1)^{\circ}$ with respect to each other. The Ag atom is linked to the O atoms of two bromate counter-ions so that the geometry approximates to that of a rectangular plane. The


Figure 1
A view of (I), showing $75 \%$ probability displacement ellipsoids. H atoms are drawn as spheres of arbitrary radii. [Symmetry codes: (i) $\frac{1}{2}+x, \frac{5}{2}-y$, $z-\frac{1}{2}$; (ii) $-x, 2-y, 1-z$.]

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chains propagate along [10 $\overline{1}]$; they are cross-linked by the bromate anions and by $\operatorname{Ag} \cdots \operatorname{Ag}(-x, \quad 2-y, \quad 1-z)$ [3.3197 (6) Å] interactions, thus forming layers (Fig. 2). The metal-metal interaction in (I) is longer than that [2.78 (1) A] found in the $F d d 2$ modification of the nitrate (Yaghi \& Li, 1996) but similar to that [ 3.286 (2) $\AA$ ] found in the dihydrogen orthophosphate (Tong et al., 2000). The latter study lists the $\mathrm{Ag} \cdots \mathrm{Ag}$ distances in several related systems. The present structure is consolidated by hydrogen bonds (Table 2); the water molecules can be envisaged as occupying the spaces between the layers. One of the water molecules lies on a special position with $\overline{1}$ site symmetry, and thus its H atoms are necessarily disordered.


## Experimental

Silver bromate ( $0.24 \mathrm{~g}, 1 \mathrm{mmol}$ ), 4,4'-bipyridine ( $0.16 \mathrm{~g}, 1 \mathrm{mmol}$ ) and water ( 10 ml ) were placed in a 23 ml Teflon-lined, stainless-steel bomb, which was heated at 413 K for 6 d . The bomb was then cooled at $5 \mathrm{~K} \mathrm{~h}^{-1}$. Needle-shaped crystals were isolated in about $70 \%$ yield. A needle was cut into a block to provide the crystal for measurement. Analysis calculated for $\mathrm{C}_{10} \mathrm{H}_{12} \mathrm{~N}_{2} \mathrm{O}_{5.5}$ : C $27.55, \mathrm{H} 2.77, \mathrm{~N} 6.45 \%$; found: C $27.54, \mathrm{H} 3.02, \mathrm{~N} 6.37 \%$.

## Crystal data

$\left[\mathrm{Ag}\left(\mathrm{BrO}_{3}\right)\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right)\right] \cdot 2.5 \mathrm{H}_{2} \mathrm{O}$
$M_{r}=437.00$
Monoclinic, $P 2_{1} / n$
$a=13.1128(8) \AA$
$b=6.9843(4) \AA$
$c=15.3908(9) \AA$
$\beta=104.647(1)^{\circ}$
$V=1363.74(14) \AA^{3}$
$Z=4$
$D_{x}=2.128 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 4053
$\quad$ reflections
$\theta=2.4-28.2^{\circ}$
$\mu=4.43 \mathrm{~mm}^{-1}$
$T=298(2) \mathrm{K}$
Block, colorless
$0.37 \times 0.25 \times 0.24 \mathrm{~mm}$

Data collection
Bruker APEX area-detector diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2001)
$T_{\text {min }}=0.201, T_{\text {max }}=0.416$
7759 measured reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.030$
$w R\left(F^{2}\right)=0.084$
$S=1.05$
3066 reflections
196 parameters
H atoms treated by a mixture of
$\quad$ independent and constrained
$\quad$ refinement


Figure 2
A view of (I), showing the extended chain structure of the $\left(\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2}\right) \mathrm{Ag}$ units linked by $\mathrm{Ag} \cdots \mathrm{Ag}$ interactions (dashed lines) into a layer structure. Water molecules and C-bound H atoms are not shown.

Table 1
Selected geometric parameters ( $\AA \AA^{\circ}$ ).

| $\mathrm{Ag} 1-\mathrm{N} 1$ | $2.147(2)$ | $\mathrm{Ag} 1-\mathrm{O} 1$ | $2.877(3)$ |
| :--- | ---: | :--- | :---: |
| $\mathrm{Ag} 1-\mathrm{N} 2^{\mathrm{i}}$ | $2.152(2)$ | $\mathrm{Ag} 1-\mathrm{O} 2^{\mathrm{ii}}$ | $2.972(3)$ |
|  |  |  |  |
| $\mathrm{N} 1-\mathrm{Ag} 1-\mathrm{N} 2^{\mathrm{i}}$ | $172.0(1)$ | $\mathrm{N}^{\mathrm{i}}-\mathrm{Ag} 1-\mathrm{O} 1$ | $90.3(1)$ |
| $\mathrm{N} 1-\mathrm{Ag} 1-\mathrm{O} 1$ | $88.2(1)$ | $\mathrm{N}^{2}-{\mathrm{Ag} 1-\mathrm{O} 2^{i i}}$ | $92.81(9)$ |
| $\mathrm{N} 1-\mathrm{Ag} 1-\mathrm{O} 2^{\mathrm{ii}}$ | $91.1(1)$ | $\mathrm{O} 1-\mathrm{Ag} 1-\mathrm{O} 2^{i i}$ | $162.3(1)$ |

Symmetry codes: (i) $\frac{1}{2}+x, \frac{5}{2}-y, z-\frac{1}{2}$; (ii) $-x, 2-y, 1-z$.

Table 2
Hydrogen-bonding geometry ( $\mathrm{A},{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| O1W-H1W2 $\cdots$ O2 | $0.86(1)$ | $2.12(3)$ | $2.927(5)$ | $155(6)$ |
| O1 $W-\mathrm{H} 1 W 1 \cdots \mathrm{O} 3 W$ | $0.86(1)$ | $2.46(6)$ | $2.894(5)$ | $112(5)$ |
| O2 $W-\mathrm{H} 2 W 1 \cdots \mathrm{O}^{\text {iii }}$ | $0.87(1)$ | $2.18(5)$ | $2.76(1)$ | $124(5)$ |
| O2 $^{\text {(ii }} W-\mathrm{H} 2 W 2 \cdots \mathrm{O}^{\text {iv }}$ | $0.86(1)$ | $2.13(4)$ | $2.853(5)$ | $141(5)$ |

Symmetry codes: (iii) $1-x,-y, 1-z$; (iv) $1-x, 1-y, 1-z$.
The aromatic H atoms were placed at calculated positions in the riding-model approximation $\left[\mathrm{C}-\mathrm{H}=0.93 \AA\right.$ and $U_{\text {iso }}(\mathrm{H})=$ $\left.1.2 U_{\text {eq }}(\mathrm{C})\right]$. Water H atoms were located and refined with distance restraints $[\mathrm{O}-\mathrm{H}=0.85(1) \AA$ and $\mathrm{H} \cdots \mathrm{H}=1.39(1) \AA$, and their

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displacement parameters were tied to the $U_{\text {eq }}$ values of the O atoms by a factor of 1.2. The O3W water molecule lies on an inversion center, so it should be bonded to only one H atom; however, the inversion symmetry element constrains the $\mathrm{H}-\mathrm{O}-\mathrm{H}$ angle to be $180^{\circ}$. Two H atoms were refined, each with a site occupancy of 0.5 (H3W1 and H3W2). Hydrogen bonds involving these disordered atoms are not listed in Table 2.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; 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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