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#### **Key indicators**

Single-crystal X-ray study T = 292 KMean  $\sigma(C-C) = 0.007 \text{ Å}$ Disorder in solvent or counterion R factor = 0.043 wR factor = 0.127 Data-to-parameter ratio = 13.1

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

# *catena*-Poly[[[(acetonitrile- $\kappa N$ )silver(I)]- $\mu$ -bis-(pyrimidin-2-ylsulfanyl)methane- $\kappa^2 N^1$ : $N^{1'}$ ] tetrafluoroborate]

In the title complex,  $\{[Ag(C_9H_8N_4S_2)(C_2H_3N)]BF_4\}_n$ , the Ag<sup>I</sup> atom shows trigonal–planar coordination, as it is coordinated by the N atoms from two bis(pyrimidin-2-ylsulfanyl)methane molecules and the N-atom donor of an acetonitrile molecule. The manner in which the donor molecules connect to the Ag atoms leads to a linear chain structure. Adjacent chains are connected into a layer *via* weak Ag···S interactions.

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#### Comment

Following a report on silver (I) *N*-heterocyclic thioether complexes featuring unusual coordination motifs (Hong *et al.*, 2000), we report the silver tetrafluoroborate complex, (I), of bis(pyrimidin-2-ylsulfanyl)methane. The Ag<sup>I</sup> atom exists in a distorted trigonal-planar geometry (Fig. 1) which comprises two pyrimidine N-atom donors from different donor molecules and the N-atom donor from an acetonitrile molecule.



The bond dimensions are within the range reported in similar complexes (Carlucci *et al.*, 1998; Constable *et al.*, 1998; Hou *et al.*, 2004). In the donor molecule, the two pyrimidine



© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved *ORTEPII* (Johnson, 1976) view of the title compound with 30% probability displacement ellipsoids. [Symmetry codes: (i)  $\frac{3}{2} - x, \frac{1}{2} + y, z$ ].

rings are nearly perpendicular to each other [dihedral angle = 87.7 (1)°], the twist being necessary for the pyrimidine rings to engage in coordination to give rise to a linear chain running along the b axis. As the Ag atom in one chain interacts weakly with the S atoms of adjacent chains  $[Ag \cdot \cdot S = 3.050 (2) \text{ Å}]$ , the geometry is distorted towards trigonal bipyramidal; the weak interactions lead to a layer structure (Fig. 2). The anions are found between the layers. The analogous nitrate complex (Zheng et al., 2003) adopts a paired-chain structure.

### **Experimental**

Bis(pyrimidin-2-ylsulfanyl)methane (L) was prepared according to a reported procedure (Zheng et al., 2003) and the product was characterized by NMR and IR spectroscopy. A solution containing a 1:1 molar ratio of AgBF<sub>4</sub> (0.2 mmol, 0.08 g) and the ligand (0.2 mmol, 0.04 g) in acetonitrile-chloroform (1:1) was stirred for 30 min. The mixture was filtered and the solvent allowed to evaporate to afford colourless crystals. CHN elemental analyses confirmed the formulation determined from the diffraction study. FT-IR data (KBr pellet, cm<sup>-1</sup>): 3073 (*m*), 3011 (*w*), 2933 (*w*), 1560 (*s*), 1582 (*s*), 1057 (*s*), 752 (*m*), 693 (*m*), 494 (*w*).

#### Crystal data

$[Ag(C_9H_8N_4S_2)(C_2H_3N)]BF_4$ $M_r = 472.05$ Orthorhombic, <i>Pbca</i> a = 8.7063 (12) Å b = 14.7402 (16) Å c = 24.965 (2) Å V = 3203.9 (6) Å <sup>3</sup>	Mo K $\alpha$ radiation Cell parameters from 1758 reflections $\theta = 2.9-26.4^{\circ}$ $\mu = 1.57 \text{ mm}^{-1}$ T = 292 (1)  K Block colourless
Z = 8 $D_x = 1.957 \text{ Mg m}^{-3}$	$0.20 \times 0.18 \times 0.14 \text{ mm}$
Data collection	
Bruker SMART 1000 CCD diffractometer $\varphi$ and $\omega$ scans Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 1996) $T_{min} = 0.706, T_{max} = 0.804$ 17366 measured reflections	3322 independent reflections 2441 reflections with $I > 2\sigma(R_{int} = 0.041)$ $\theta_{max} = 26.5^{\circ}$ $h = -8 \rightarrow 10$ $k = -17 \rightarrow 18$ $l = -31 \rightarrow 23$
Refinement	
Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.127$ S = 1.18 3322 reflections 253 parameters	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0613P)^{2} + 2.423P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.79 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.68 \text{ e} \text{ Å}^{-3}$

H-atom	parameters	constrained

#### Table 1

Selected geometric parameters (°).

N1-Ag1-N3 <sup>i</sup>	141.32 (13)	N3 <sup>i</sup> -Ag1-N5	92.35 (14)
N1-Ag1-N5	124.56 (15)	0	
Symmetry code: (i)	$x + \frac{3}{2}, y + \frac{1}{2}, z$		





View of the zigzag chains linked through weak Ag...S interactions, shown as dashed lines. H atoms have been omitted.

All H atoms were positioned geometrically, with  $Csp^2 - H = 0.93$  Å and  $Csp^3 - H = 0.97$  Å; they were constrained to ride on their parent atoms, with  $U_{iso}(H) = 1.2U_{eq}(C)$ . The BF<sub>4</sub> anion was disordered and was modelled as two components with site occupancies of 0.86(1)and 0.14 (1). The B-F distances were restrained as 1.35 (1) Å.

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

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 $I > 2\sigma(I)$ 

Extinction correction: SHELXL97

Extinction coefficient: 0.0078 (5)

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