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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.007 Å Disorder in solvent or counterion R factor = 0.036 wR factor = 0.088 Data-to-parameter ratio = 13.9

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# catena-Poly[[silver(I)- $\mu$ -1-phenylsulfanyl-2-(2-pyridyl-sulfanyl)ethane- $\kappa^2 S^1$ : $N^2$ ] tetrafluoroborate]

In the title complex,  $\{[Ag(C_{13}H_{13}NS_2)]BF_4\}_n$ , the Ag<sup>I</sup> ions show linear coordination through the N atom from one 1phenylsulfanyl-2-(2-pyridylsulfanyl)ethane (*L*) molecule and the S atom of another. Each *L* ligand bridges two adjacent Ag atoms through the donor molecules, leading to a linear chain structure. Received 28 July 2005 Accepted 5 August 2005 Online 28 September 2005

## Comment

There are many silver(I) complexes with substituted bisthioether compounds featuring unusual coordination motifs (Hong *et al.*, 2000) in which donor atoms have always been the same. We report here the structure of the silver tetrafluoroborate complex, (I), of AgBF<sub>4</sub> with the *N/S*-donor ligand 1-phenylsulfanyl-2-(2-pyridylsulfanyl)ethane (*L*).



In (I), the  $Ag^{I}$  atom exists in a nearly linear geometry (Fig. 1), comprising a pyridyl *N*-atom donor and a phenyl-



#### Figure 1 ORTEPII (Johnson, 1976) view of the title compound, with 30%

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved  $x_1, \frac{1}{2} + y, \frac{3}{2} - z$ .]



A view of the chains linked through  $Ag \cdots S$  weak interactions (dashed lines). H and disordered F atoms have been omitted.

sulfanyl S-atom donor from different *L* molecules. The bond dimensions are within the range reported in similar complexes (Wang *et al.*, 1992; Constable *et al.*, 1998; Hou *et al.*, 2004). The  $BF_4^-$  anion shows a weak interaction with the  $Ag^I$  center with an  $Ag1\cdots$ F1 distance of 2.630 (9) Å, which is longer than the sum of ionic radii (2.31 Å; Shannon & Prewitt, 1969). This interaction causes a small distortion of the linear coordination of the  $Ag^I$  atom [S1-Ag1- $N1^i$  = 174.02 (9)°].

In the ligand, the phenyl and pyridyl rings are inclined with respect to each other at an angle of 73.1 (2)° and the S1···S2 non-bonded distance is 4.417 (2) Å. Each ligand bridges two adjacent Ag atoms through Ag—N and Ag—S coordination to give rise to an infinite chain running along the *b* axis. In the crystal packing of (I), adjacent chains are potentially linked into a two-dimensional network (Fig. 2) through weak Ag··S interactions [Ag1···S2( $x, \frac{3}{2} - y, -\frac{1}{2} + z$ ) = S2···Ag1( $x, \frac{3}{2} - y, \frac{1}{2} + z$ ) = 3.013 (1) Å; Suenaga *et al.*, 1999]. The anions are found between the layers.

# Experimental

1-Phenylsulfanyl-2-(2-pyridylsulfanyl)ethane (*L*) was prepared according to a reported procedure (Chen *et al.*, 2001). A solution containing a 1:1 molar ratio of AgBF<sub>4</sub> (0.2 mmol) and *L* (0.2 mmol) in acetonitrile–chloroform (1:1) was stirred for 30 min at room temperature, and the mixture was filtered. Colorless single crystals suitable for X-ray investigation were obtained from this filtrate by slow evaporation of the solvent.

### Crystal data

$[Ag(C_{13}H_{13}NS_2)]BF_4$	$D_x = 1.836 \text{ Mg m}^{-3}$
$M_r = 442.04$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 1006
a = 11.252 (3)  Å	reflections
b = 13.999 (4) Å	$\theta = 3.6-25.8^{\circ}$
c = 11.315 (3) Å	$\mu = 1.56 \text{ mm}^{-1}$
$\beta = 116.227 \ (4)^{\circ}$	T = 293 (2) K
$V = 1598.8 (8) \text{ Å}^3$	Block, colorless
Z = 4	$0.16 \times 0.12 \times 0.08 \text{ mm}$

#### Data collection

Bruker SMART 1000 CCD	3260 independent reflections
diffractometer	2324 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.036$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.4^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -10 \rightarrow 14$
$T_{\min} = 0.643, \ T_{\max} = 0.883$	$k = -13 \rightarrow 17$
9039 measured reflections	$l = -14 \rightarrow 9$
Refinement	
Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0437P)^2$
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 0.3971P]
$wR(F^2) = 0.088$	where $P = (F_0^2 + 2F_c^2)/3$
S = 0.99	$(\Delta/\sigma)_{\rm max} = 0.004$
3260 reflections	$\Delta \rho_{\rm max} = 0.52 \ {\rm e} \ {\rm \AA}^{-3}$
235 parameters	$\Delta \rho_{\rm min} = -0.39 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	

Table 1
Selected geometric parameters (Å, °).

Ag1-N1 <sup>i</sup>	2.181 (3)	Ag1-S1	2.4127 (11)	
-		-		
N1 <sup>i</sup> -Ag1-S1	174.02 (9)			
Symmetry code: (i) $-x + 1$ , $y + \frac{1}{2}$ , $-z + \frac{3}{2}$ .				

All H atoms were positioned geometrically with  $Csp^2 - H = 0.93 \text{ Å}$ and  $Csp^3 - H = 0.97 \text{ Å}$ ; they were constrained to ride on their parent atoms, with  $U_{iso}(H) = 1.2U_{eq}(C)$ . The BF<sub>4</sub><sup>-</sup> anion was observed to be disordered and was eventually modeled as eight 0.5-occupied F atoms surrounding the central B atom.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 1998); 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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