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Crystal structure of *catena*-poly[[silver(I)- μ -N-(pyridin-2-ylmethyl)pyridine-3-amine- κ^2 N:N'] trifluoromethanesulfonate]

Suk-Hee Moon,^a Seonghwa Cho^b and Ki-Min Park^{c*}

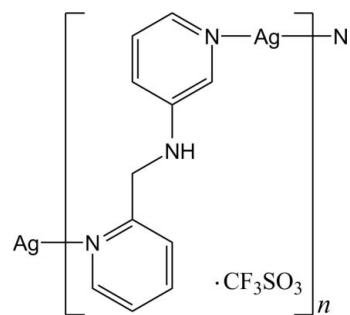
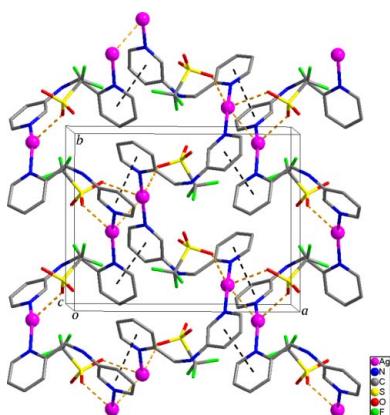
^aDepartment of Food & Nutrition, Kyungnam College of Information and Technology, Busan 617-701, Republic of Korea,

^bDepartment of Chemistry, Gyeongsang National University, Jinju 660-701, Republic of Korea, and ^cResearch Institute of Natural Sciences, Gyeongsang National University, Jinju 660-701, Republic of Korea. *Correspondence e-mail: kmpark@gnu.ac.kr

In the asymmetric unit of the title compound, $\{[\text{Ag}(\text{C}_{11}\text{H}_{11}\text{N}_3)]\text{CF}_3\text{SO}_3\}_n$, there are two Ag^{I} atoms, two *N*-(pyridine-2-ylmethyl)pyridine-3-amine ligands (*A* and *B*) and two CF_3SO_3^- anions. Both Ag^{I} atoms are bridged by two pyridine N atoms from two symmetry-related *A* or *B* ligands, forming right- or left-handed helical chains, respectively. The Ag^{I} atom of the right-handed helical chain adopts a slightly distorted linear coordination geometry [$\text{N}-\text{Ag}-\text{N} = 170.69 (14)$ °], while that of the left-handed helical chain adopts a bent geometry [$\text{N}-\text{Ag}-\text{N} = 149.42 (14)$ °]. Both helical chains have the same pitch length [10.8437 (5) Å], propagate along the *b*-axial direction and are alternately arranged via $\text{Ag}\cdots\text{Ag}$ [3.0814 (5) Å] and $\pi-\pi$ stacking interactions [centroid-centroid distances = 3.514 (3) and 3.487 (3) Å], resulting in the formation of a two-dimensional supramolecular network extending parallel to the *ab* plane. Weak $\text{Ag}\cdots\text{O}$ [2.861 (4), 2.617 (3), and 2.624 (4) Å] and $\text{Ag}\cdots\text{F}$ [3.017 (3) Å] interactions as well as $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{F}$ hydrogen-bonding interactions occur between the helical chains and the anions.

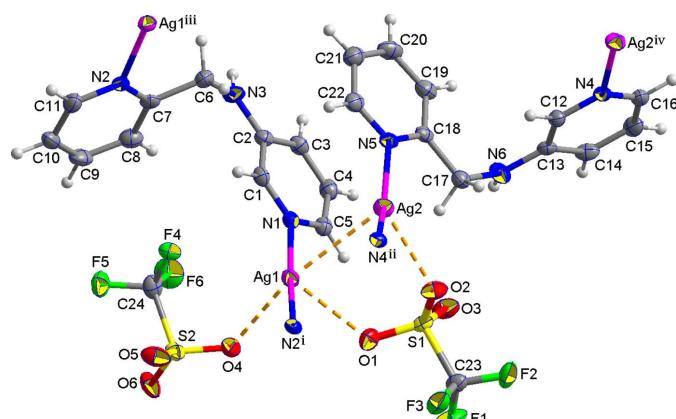
1. Chemical context

A few silver coordination polymers based on unsymmetrical dipyridyl ligands composed of two terminal pyridines with different substituted-nitrogen positions have been reported (Moon & Park, 2013, 2014; Zhang *et al.*, 2013). In an extension of investigations on Ag^{I} coordination polymers with unsymmetrical dipyridyl ligands, the title compound was prepared by the reaction of silver trifluoromethanesulfonate with *N*-(pyridine-2-ylmethyl)pyridine-3-amine. The structure of title compound is related to that of the perchlorate salt (Moon & Park, 2014; Zhang *et al.*, 2013).



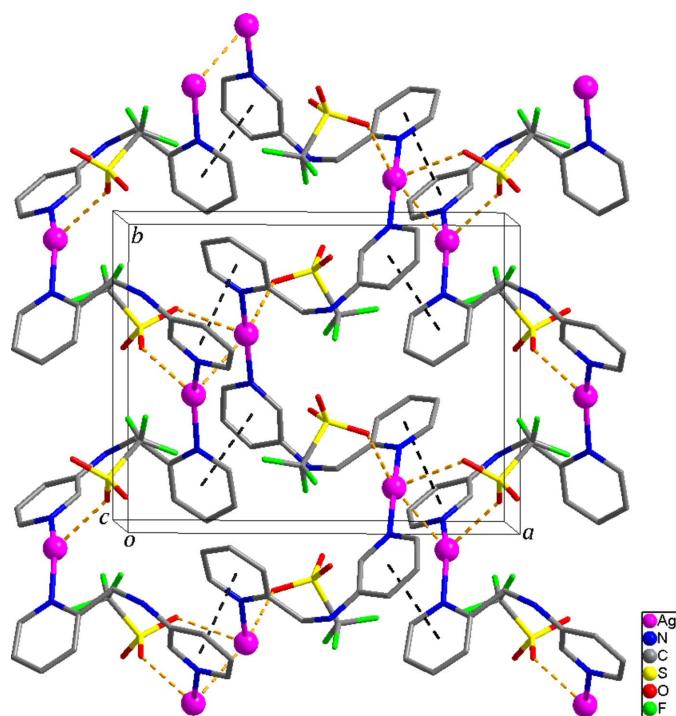
2. Structural commentary

The molecular components of the title structure are shown in Fig. 1. The asymmetric unit contains two Ag^{I} atoms (Ag1 and Ag2), two *N*-(pyridine-2-ylmethyl)pyridine-3-amine (Lee *et*

**Figure 1**

A view of the molecular structure of the title compound, with the atom numbering. Displacement ellipsoids are drawn at the 50% probability level and dashed lines represent $\text{Ag}\cdots\text{Ag}$ and $\text{Ag}\cdots\text{O}$ interactions [symmetry codes: (i) $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$; (ii) $-x, y + \frac{1}{2}, -z + \frac{1}{2}$; (iii) $-x + 1, y - \frac{1}{2}, -z + \frac{1}{2}$; (iv) $-x, y - \frac{1}{2}, -z + \frac{1}{2}$].

al., 2013) ligands (*A* and *B*) and two trifluoromethanesulfonate anions. The $\text{Ag}1$ atom is coordinated by two pyridine N atoms from two symmetry-related *A* ligands giving a geometry which is slightly distorted from linear [$\text{N}1-\text{Ag}1-\text{N}2 = 170.69 (14)^\circ$], forming a right-handed helical chain, while the $\text{Ag}2$ atom is coordinated by two pyridine N atoms from two symmetry-related *B* ligands in a bent arrangement [$\text{N}4-\text{Ag}2-\text{N}5 = 149.42 (14)^\circ$], forming a left-handed helical chain. Two pyridine rings coordinating to the $\text{Ag}1$ and $\text{Ag}2$ atoms are tilted by 14.1 (3) and 28.9 (2) $^\circ$, respectively, with respect to each other.

**Figure 2**

The two-dimensional supramolecular network formed through $\text{Ag}\cdots\text{Ag}$ and $\text{Ag}\cdots\text{O}$ interactions (yellow dashed lines) and $\pi\cdots\pi$ stacking interactions (black dashed lines).

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N}3-\text{H}3\cdots\text{O}6^i$	0.88	2.51	3.206 (6)	136
$\text{N}6-\text{H}6\cdots\text{O}3^{ii}$	0.88	2.43	3.217 (6)	149
$\text{C}1-\text{H}1\cdots\text{O}5^{iii}$	0.95	2.55	3.339 (6)	141
$\text{C}4-\text{H}4\cdots\text{O}3^{ii}$	0.95	2.54	3.383 (6)	147
$\text{C}6-\text{H}6A\cdots\text{O}5^i$	0.99	2.57	3.471 (6)	151
$\text{C}11-\text{H}11\cdots\text{F}3^{iii}$	0.95	2.49	3.311 (6)	145
$\text{C}11-\text{H}11\cdots\text{O}1^{iii}$	0.95	2.54	3.350 (7)	143
$\text{C}15-\text{H}15\cdots\text{O}6^{ii}$	0.95	2.53	3.450 (6)	164
$\text{C}16-\text{H}16\cdots\text{O}6^{iv}$	0.95	2.58	3.316 (6)	135
$\text{C}17-\text{H}17B\cdots\text{O}3$	0.99	2.58	3.422 (6)	143

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

3. Supramolecular features

Both helical chains in the structure have the same pitch length [10.8437 (5) \AA], propagate along the *b*-axial direction and are alternately arranged via $\text{Ag}1\cdots\text{Ag}2$ interactions [3.0814 (5) \AA], resulting in the formation of a two-dimensional supramolecular network extending parallel to the *ab* plane (Fig. 2). Furthermore, $\pi\cdots\pi$ stacking interactions [centroid–centroid distances = 3.514 (3) and 3.487 (3) \AA] between pyridine rings of both helical chains contribute to the stabilization of the two-dimensional network. In the crystal structure, the two-dimensional networks are further stabilized by $\text{Ag}\cdots\text{O}$ and $\text{Ag}\cdots\text{F}$ interactions [$\text{Ag}1\cdots\text{O}1$ 2.861 (4), $\text{Ag}1\cdots\text{O}4$

Table 2
Experimental details.

Crystal data	[$\text{Ag}(\text{C}_{11}\text{H}_{11}\text{N}_3)\text{-CF}_3\text{SO}_3$
Chemical formula	
M_r	442.17
Crystal system, space group	Monoclinic, $P2_1/c$
Temperature (K)	173
a, b, c (\AA)	13.7529 (6), 10.8437 (5), 19.5795 (9)
β ($^\circ$)	99.826 (1)
V (\AA^3)	2877.1 (2)
Z	8
Radiation type	Mo $K\alpha$
μ (mm^{-1})	1.60
Crystal size (mm)	0.31 \times 0.22 \times 0.10
Data collection	Bruker SMART CCD area detector
Diffractometer	Multi-scan (SADABS; Bruker, 2000)
Absorption correction	
T_{\min}, T_{\max}	0.637, 0.857
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	15852, 5629, 4286
R_{int}	0.038
($\sin \theta/\lambda$) _{max} (\AA^{-1})	0.617
Refinement	
$R[F^2 > 2\sigma(F^2)]$, $wR(F^2)$, S	0.039, 0.093, 1.06
No. of reflections	5629
No. of parameters	415
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ ($e \text{\AA}^{-3}$)	1.04, -0.67

Computer programs: SMART and SAINT-Plus (Bruker, 2000), SHELXS97, SHELXL97 and SHELXTL (Sheldrick, 2008) and DIAMOND (Brandenburg, 2005).

2.624 (4), $\text{Ag}_2\cdots\text{O}_2$ 2.617 (3), $\text{Ag}_2\cdots\text{F}_3^{\text{iv}}$ 3.017 (3) Å; symmetry code: (iv) $-x, y - \frac{1}{2}, -z + \frac{1}{2}$] (Figs. 1 and 2) as well as N—H···O and N—H···O and C—H···O and C—H···F hydrogen-bonds (Table 1) between the helical chains and CF_3SO_3^- anions.

4. Synthesis and crystallization

The ligand (*N*-(pyridin-2-ylmethyl)pyridine-3-amine) was prepared according to a procedure described by Lee *et al.* (2013). Crystals of the title compound suitable for X-ray analysis were obtained by vapour diffusion of diethyl ether into a DMSO solution of the white precipitate afforded by the reaction of the ligand with silver(I) hexafluoridophosphate in the molar ratio 1:1 in methanol.

5. Refinement details

Crystal data, data collection and structure refinement details are summarized in Table 2. All H atoms were positioned

geometrically and refined using a riding model, with $d(\text{C—H}) = 0.95$ Å for Csp^2 —H, 0.88 Å for amine N—H and 0.99 Å for methylene C—H. For all H atoms $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C,N})$.

Acknowledgements

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supporting information

Acta Cryst. (2014). E70, 389-391 [doi:10.1107/S1600536814022922]

Crystal structure of *catena-poly[[silver(I)- μ -N-(pyridin-2-ylmethyl)pyridine-3-amine- κ^2 N:N'] trifluoromethanesulfonate]*

Suk-Hee Moon, Seonghwa Cho and Ki-Min Park

Computing details

Data collection: SMART (Bruker, 2000); cell refinement: SAINT-Plus (Bruker, 2000); data reduction: SAINT-Plus (Bruker, 2000); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: DIAMOND (Brandenburg, 2005); software used to prepare material for publication: SHELXTL (Sheldrick, 2008).

catena-Poly[[silver(I)- μ -N-(pyridin-2-ylmethyl)pyridine-3-amine- κ^2 N:N'] trifluoromethanesulfonate]

Crystal data



$M_r = 442.17$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 13.7529 (6)$ Å

$b = 10.8437 (5)$ Å

$c = 19.5795 (9)$ Å

$\beta = 99.826 (1)^\circ$

$V = 2877.1 (2)$ Å³

$Z = 8$

$F(000) = 1744$

$D_x = 2.042$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 5639 reflections

$\theta = 2.4\text{--}28.2^\circ$

$\mu = 1.60$ mm⁻¹

$T = 173$ K

Plate, colorless

$0.31 \times 0.22 \times 0.10$ mm

Data collection

Bruker SMART CCD area detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2000)

$T_{\min} = 0.637$, $T_{\max} = 0.857$

15852 measured reflections

5629 independent reflections

4286 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.038$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 1.5^\circ$

$h = -16 \rightarrow 16$

$k = -13 \rightarrow 9$

$l = -23 \rightarrow 24$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.039$

$wR(F^2) = 0.093$

$S = 1.06$

5629 reflections

415 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0369P)^2 + 6.3774P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$

$$\Delta\rho_{\max} = 1.04 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.67 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Ag1	0.30869 (3)	0.63691 (3)	0.192614 (19)	0.02491 (11)
Ag2	0.17632 (3)	0.43820 (3)	0.24204 (2)	0.02787 (11)
N1	0.3183 (3)	0.4818 (4)	0.1251 (2)	0.0220 (9)
N2	0.7006 (3)	0.2705 (3)	0.2265 (2)	0.0223 (9)
N3	0.4684 (3)	0.2087 (4)	0.1066 (2)	0.0284 (10)
H3	0.4727	0.1538	0.0742	0.034*
C1	0.3913 (3)	0.3969 (4)	0.1368 (2)	0.0231 (11)
H1	0.4428	0.4078	0.1753	0.028*
C2	0.3940 (3)	0.2938 (4)	0.0943 (2)	0.0228 (10)
C3	0.3168 (3)	0.2794 (5)	0.0383 (2)	0.0252 (11)
H3A	0.3152	0.2103	0.0083	0.030*
C4	0.2434 (4)	0.3662 (5)	0.0272 (3)	0.0275 (11)
H4	0.1904	0.3569	-0.0104	0.033*
C5	0.2462 (4)	0.4662 (5)	0.0701 (2)	0.0237 (11)
H5	0.1956	0.5265	0.0607	0.028*
C6	0.5398 (4)	0.2033 (5)	0.1695 (3)	0.0288 (12)
H6A	0.5618	0.1167	0.1774	0.035*
H6B	0.5071	0.2279	0.2087	0.035*
C7	0.6299 (4)	0.2836 (4)	0.1702 (3)	0.0234 (11)
C8	0.6403 (4)	0.3658 (5)	0.1181 (3)	0.0327 (12)
H8	0.5897	0.3735	0.0787	0.039*
C9	0.7247 (4)	0.4367 (5)	0.1237 (3)	0.0346 (13)
H9	0.7330	0.4928	0.0879	0.042*
C10	0.7964 (4)	0.4258 (5)	0.1812 (3)	0.0324 (12)
H10	0.8543	0.4751	0.1868	0.039*
C11	0.7815 (4)	0.3405 (5)	0.2307 (3)	0.0314 (12)
H11	0.8318	0.3310	0.2701	0.038*
N4	-0.1913 (3)	0.0415 (3)	0.1616 (2)	0.0218 (9)
N5	0.1834 (3)	0.2634 (4)	0.1887 (2)	0.0224 (9)
N6	-0.0542 (3)	0.2536 (4)	0.0675 (2)	0.0295 (10)
H6	-0.0559	0.2757	0.0241	0.035*
C12	-0.1212 (3)	0.1180 (4)	0.1464 (2)	0.0211 (10)

H12	-0.0651	0.1334	0.1810	0.025*
C13	-0.1267 (3)	0.1755 (4)	0.0824 (2)	0.0221 (10)
C14	-0.2106 (4)	0.1521 (4)	0.0325 (2)	0.0273 (11)
H14	-0.2181	0.1902	-0.0118	0.033*
C15	-0.2815 (4)	0.0739 (5)	0.0483 (3)	0.0306 (12)
H15	-0.3385	0.0570	0.0148	0.037*
C16	-0.2702 (4)	0.0194 (4)	0.1130 (3)	0.0261 (11)
H16	-0.3197	-0.0352	0.1233	0.031*
C17	0.0244 (3)	0.3012 (5)	0.1190 (3)	0.0271 (11)
H17A	-0.0031	0.3203	0.1614	0.033*
H17B	0.0475	0.3798	0.1016	0.033*
C18	0.1121 (4)	0.2185 (4)	0.1389 (2)	0.0227 (10)
C19	0.1216 (4)	0.1044 (5)	0.1089 (3)	0.0313 (12)
H19	0.0706	0.0738	0.0742	0.038*
C20	0.2063 (4)	0.0352 (5)	0.1301 (3)	0.0351 (13)
H20	0.2139	-0.0430	0.1098	0.042*
C21	0.2788 (4)	0.0807 (5)	0.1806 (3)	0.0340 (13)
H21	0.3372	0.0348	0.1961	0.041*
C22	0.2647 (4)	0.1953 (5)	0.2082 (3)	0.0281 (11)
H22	0.3152	0.2273	0.2428	0.034*
S1	0.03658 (9)	0.65963 (11)	0.12069 (6)	0.0230 (3)
O1	0.1266 (3)	0.7218 (3)	0.1137 (2)	0.0390 (9)
O2	0.0406 (3)	0.5931 (3)	0.18476 (18)	0.0343 (9)
O3	-0.0106 (3)	0.5974 (3)	0.05959 (18)	0.0369 (9)
C23	-0.0469 (4)	0.7861 (5)	0.1303 (3)	0.0289 (12)
F1	-0.0609 (3)	0.8570 (3)	0.07437 (17)	0.0480 (9)
F2	-0.1350 (2)	0.7440 (3)	0.13945 (17)	0.0439 (8)
F3	-0.0113 (2)	0.8558 (3)	0.18486 (16)	0.0381 (8)
S2	0.47697 (9)	0.84803 (12)	0.12529 (6)	0.0269 (3)
O4	0.3758 (3)	0.8093 (3)	0.1188 (2)	0.0380 (9)
O5	0.5228 (3)	0.8873 (3)	0.19320 (19)	0.0403 (10)
O6	0.4983 (3)	0.9245 (4)	0.07004 (19)	0.0410 (10)
C24	0.5413 (4)	0.7040 (5)	0.1142 (3)	0.0311 (12)
F4	0.5337 (2)	0.6255 (3)	0.16634 (16)	0.0347 (7)
F5	0.6375 (2)	0.7235 (3)	0.11623 (17)	0.0428 (8)
F6	0.5059 (3)	0.6481 (3)	0.05552 (17)	0.0539 (10)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Ag1	0.0246 (2)	0.0220 (2)	0.0286 (2)	-0.00393 (16)	0.00586 (15)	-0.00514 (16)
Ag2	0.0321 (2)	0.0252 (2)	0.0282 (2)	-0.00139 (17)	0.01060 (17)	-0.00680 (17)
N1	0.020 (2)	0.021 (2)	0.025 (2)	-0.0021 (17)	0.0048 (17)	0.0009 (17)
N2	0.022 (2)	0.016 (2)	0.029 (2)	0.0018 (16)	0.0050 (17)	-0.0032 (17)
N3	0.033 (2)	0.020 (2)	0.031 (2)	0.0039 (18)	0.0014 (19)	-0.0082 (18)
C1	0.022 (3)	0.027 (3)	0.021 (2)	-0.005 (2)	0.002 (2)	-0.001 (2)
C2	0.027 (3)	0.019 (2)	0.023 (3)	-0.002 (2)	0.006 (2)	0.001 (2)
C3	0.029 (3)	0.025 (3)	0.022 (3)	-0.005 (2)	0.005 (2)	-0.004 (2)

C4	0.025 (3)	0.032 (3)	0.024 (3)	-0.004 (2)	-0.001 (2)	0.002 (2)
C5	0.022 (3)	0.027 (3)	0.022 (3)	0.000 (2)	0.004 (2)	0.003 (2)
C6	0.024 (3)	0.024 (3)	0.038 (3)	0.004 (2)	0.007 (2)	0.002 (2)
C7	0.027 (3)	0.015 (2)	0.029 (3)	0.003 (2)	0.006 (2)	-0.004 (2)
C8	0.040 (3)	0.030 (3)	0.029 (3)	0.003 (2)	0.005 (2)	0.001 (2)
C9	0.046 (3)	0.023 (3)	0.038 (3)	0.004 (2)	0.018 (3)	0.009 (2)
C10	0.032 (3)	0.021 (3)	0.049 (3)	-0.002 (2)	0.018 (3)	-0.001 (2)
C11	0.024 (3)	0.029 (3)	0.040 (3)	0.001 (2)	0.002 (2)	-0.004 (2)
N4	0.025 (2)	0.016 (2)	0.025 (2)	0.0020 (16)	0.0047 (17)	-0.0005 (17)
N5	0.029 (2)	0.019 (2)	0.021 (2)	0.0012 (17)	0.0104 (17)	0.0035 (17)
N6	0.035 (2)	0.033 (2)	0.020 (2)	-0.006 (2)	0.0030 (18)	0.0102 (19)
C12	0.021 (2)	0.020 (2)	0.021 (2)	-0.0001 (19)	0.001 (2)	-0.0012 (19)
C13	0.026 (3)	0.016 (2)	0.025 (3)	0.0041 (19)	0.004 (2)	0.000 (2)
C14	0.038 (3)	0.023 (3)	0.017 (2)	0.005 (2)	-0.003 (2)	0.000 (2)
C15	0.027 (3)	0.028 (3)	0.033 (3)	0.000 (2)	-0.007 (2)	-0.009 (2)
C16	0.022 (3)	0.020 (3)	0.036 (3)	0.001 (2)	0.006 (2)	-0.001 (2)
C17	0.026 (3)	0.024 (3)	0.031 (3)	-0.003 (2)	0.005 (2)	0.003 (2)
C18	0.029 (3)	0.018 (2)	0.025 (3)	-0.003 (2)	0.013 (2)	0.001 (2)
C19	0.039 (3)	0.024 (3)	0.032 (3)	-0.006 (2)	0.011 (2)	-0.001 (2)
C20	0.051 (4)	0.023 (3)	0.035 (3)	0.002 (3)	0.019 (3)	-0.004 (2)
C21	0.035 (3)	0.031 (3)	0.038 (3)	0.004 (2)	0.011 (3)	0.004 (3)
C22	0.030 (3)	0.027 (3)	0.029 (3)	0.000 (2)	0.011 (2)	0.004 (2)
S1	0.0249 (6)	0.0222 (6)	0.0208 (6)	0.0063 (5)	0.0006 (5)	-0.0015 (5)
O1	0.029 (2)	0.037 (2)	0.054 (2)	0.0034 (17)	0.0158 (18)	-0.0033 (19)
O2	0.046 (2)	0.031 (2)	0.0251 (19)	0.0098 (17)	0.0012 (17)	0.0060 (16)
O3	0.044 (2)	0.033 (2)	0.030 (2)	0.0092 (17)	-0.0026 (17)	-0.0112 (17)
C23	0.029 (3)	0.028 (3)	0.028 (3)	0.004 (2)	-0.001 (2)	-0.006 (2)
F1	0.062 (2)	0.0397 (19)	0.0404 (19)	0.0274 (17)	0.0022 (16)	0.0099 (16)
F2	0.0218 (16)	0.052 (2)	0.058 (2)	0.0028 (15)	0.0059 (15)	-0.0139 (17)
F3	0.0315 (16)	0.0381 (18)	0.0420 (18)	0.0053 (14)	-0.0013 (14)	-0.0205 (15)
S2	0.0274 (7)	0.0256 (7)	0.0266 (7)	-0.0033 (5)	0.0018 (5)	0.0070 (5)
O4	0.030 (2)	0.031 (2)	0.055 (3)	0.0000 (16)	0.0120 (18)	0.0148 (19)
O5	0.061 (3)	0.028 (2)	0.030 (2)	-0.0019 (19)	0.0017 (19)	-0.0031 (17)
O6	0.035 (2)	0.048 (2)	0.036 (2)	-0.0173 (18)	-0.0061 (17)	0.0191 (19)
C24	0.031 (3)	0.038 (3)	0.027 (3)	-0.005 (2)	0.011 (2)	-0.005 (2)
F4	0.0365 (17)	0.0227 (15)	0.0456 (18)	0.0006 (13)	0.0087 (14)	0.0061 (14)
F5	0.0282 (17)	0.049 (2)	0.055 (2)	-0.0043 (15)	0.0179 (15)	-0.0013 (17)
F6	0.061 (2)	0.061 (2)	0.0378 (19)	-0.0073 (19)	0.0040 (17)	-0.0228 (18)

Geometric parameters (\AA , $^{\circ}$)

Ag1—N1	2.156 (4)	N6—C13	1.377 (6)
Ag1—N2 ⁱ	2.167 (4)	N6—C17	1.443 (6)
Ag1—Ag2	3.0814 (5)	N6—H6	0.8800
Ag2—N4 ⁱⁱ	2.174 (4)	C12—C13	1.390 (7)
Ag2—N5	2.175 (4)	C12—H12	0.9500
N1—C5	1.344 (6)	C13—C14	1.402 (7)
N1—C1	1.353 (6)	C14—C15	1.367 (7)

N2—C11	1.338 (6)	C14—H14	0.9500
N2—C7	1.347 (6)	C15—C16	1.383 (7)
N2—Ag1 ⁱⁱⁱ	2.167 (4)	C15—H15	0.9500
N3—C2	1.369 (6)	C16—H16	0.9500
N3—C6	1.439 (6)	C17—C18	1.500 (7)
N3—H3	0.8800	C17—H17A	0.9900
C1—C2	1.398 (7)	C17—H17B	0.9900
C1—H1	0.9500	C18—C19	1.385 (7)
C2—C3	1.398 (6)	C19—C20	1.389 (8)
C3—C4	1.369 (7)	C19—H19	0.9500
C3—H3A	0.9500	C20—C21	1.370 (8)
C4—C5	1.368 (7)	C20—H20	0.9500
C4—H4	0.9500	C21—C22	1.381 (7)
C5—H5	0.9500	C21—H21	0.9500
C6—C7	1.512 (7)	C22—H22	0.9500
C6—H6A	0.9900	S1—O3	1.429 (4)
C6—H6B	0.9900	S1—O1	1.436 (4)
C7—C8	1.380 (7)	S1—O2	1.440 (4)
C8—C9	1.382 (8)	S1—C23	1.819 (5)
C8—H8	0.9500	C23—F1	1.324 (6)
C9—C10	1.369 (8)	C23—F3	1.332 (5)
C9—H9	0.9500	C23—F2	1.336 (6)
C10—C11	1.379 (7)	S2—O6	1.433 (4)
C10—H10	0.9500	S2—O5	1.435 (4)
C11—H11	0.9500	S2—O4	1.438 (4)
N4—C16	1.336 (6)	S2—C24	1.826 (6)
N4—C12	1.343 (6)	C24—F6	1.317 (6)
N4—Ag2 ^{iv}	2.174 (4)	C24—F5	1.334 (6)
N5—C22	1.341 (6)	C24—F4	1.347 (6)
N5—C18	1.349 (6)		
N1—Ag1—N2 ⁱ	170.69 (14)	N4—C12—C13	123.1 (4)
N1—Ag1—Ag2	75.41 (10)	N4—C12—H12	118.4
N2 ⁱ —Ag1—Ag2	97.25 (10)	C13—C12—H12	118.4
N4 ⁱⁱ —Ag2—N5	149.42 (14)	N6—C13—C12	122.5 (4)
N4 ⁱⁱ —Ag2—Ag1	86.49 (10)	N6—C13—C14	120.2 (4)
N5—Ag2—Ag1	112.41 (10)	C12—C13—C14	117.3 (4)
C5—N1—C1	118.3 (4)	C15—C14—C13	119.3 (5)
C5—N1—Ag1	118.4 (3)	C15—C14—H14	120.4
C1—N1—Ag1	123.3 (3)	C13—C14—H14	120.4
C11—N2—C7	117.9 (4)	C14—C15—C16	120.0 (5)
C11—N2—Ag1 ⁱⁱⁱ	119.1 (3)	C14—C15—H15	120.0
C7—N2—Ag1 ⁱⁱⁱ	122.9 (3)	C16—C15—H15	120.0
C2—N3—C6	124.0 (4)	N4—C16—C15	121.7 (5)
C2—N3—H3	118.0	N4—C16—H16	119.1
C6—N3—H3	118.0	C15—C16—H16	119.1
N1—C1—C2	122.6 (4)	N6—C17—C18	116.2 (4)
N1—C1—H1	118.7	N6—C17—H17A	108.2

C2—C1—H1	118.7	C18—C17—H17A	108.2
N3—C2—C1	121.9 (4)	N6—C17—H17B	108.2
N3—C2—C3	120.5 (4)	C18—C17—H17B	108.2
C1—C2—C3	117.5 (4)	H17A—C17—H17B	107.4
C4—C3—C2	119.2 (5)	N5—C18—C19	121.3 (5)
C4—C3—H3A	120.4	N5—C18—C17	115.1 (4)
C2—C3—H3A	120.4	C19—C18—C17	123.6 (5)
C5—C4—C3	120.3 (5)	C18—C19—C20	119.4 (5)
C5—C4—H4	119.8	C18—C19—H19	120.3
C3—C4—H4	119.8	C20—C19—H19	120.3
N1—C5—C4	122.1 (5)	C21—C20—C19	119.4 (5)
N1—C5—H5	119.0	C21—C20—H20	120.3
C4—C5—H5	119.0	C19—C20—H20	120.3
N3—C6—C7	115.1 (4)	C20—C21—C22	118.3 (5)
N3—C6—H6A	108.5	C20—C21—H21	120.9
C7—C6—H6A	108.5	C22—C21—H21	120.9
N3—C6—H6B	108.5	N5—C22—C21	123.3 (5)
C7—C6—H6B	108.5	N5—C22—H22	118.3
H6A—C6—H6B	107.5	C21—C22—H22	118.3
N2—C7—C8	121.4 (5)	O3—S1—O1	114.7 (2)
N2—C7—C6	115.0 (4)	O3—S1—O2	115.9 (2)
C8—C7—C6	123.6 (5)	O1—S1—O2	114.3 (2)
C7—C8—C9	119.5 (5)	O3—S1—C23	103.8 (2)
C7—C8—H8	120.3	O1—S1—C23	103.0 (2)
C9—C8—H8	120.3	O2—S1—C23	102.7 (2)
C10—C9—C8	119.7 (5)	F1—C23—F3	108.4 (4)
C10—C9—H9	120.2	F1—C23—F2	107.6 (4)
C8—C9—H9	120.2	F3—C23—F2	107.6 (4)
C9—C10—C11	117.6 (5)	F1—C23—S1	111.0 (4)
C9—C10—H10	121.2	F3—C23—S1	111.0 (3)
C11—C10—H10	121.2	F2—C23—S1	111.0 (4)
N2—C11—C10	123.9 (5)	O6—S2—O5	114.4 (2)
N2—C11—H11	118.0	O6—S2—O4	115.0 (2)
C10—C11—H11	118.0	O5—S2—O4	115.7 (2)
C16—N4—C12	118.7 (4)	O6—S2—C24	103.9 (2)
C16—N4—Ag2 ^{iv}	118.0 (3)	O5—S2—C24	102.6 (2)
C12—N4—Ag2 ^{iv}	123.0 (3)	O4—S2—C24	102.7 (2)
C22—N5—C18	118.4 (4)	F6—C24—F5	108.4 (4)
C22—N5—Ag2	116.5 (3)	F6—C24—F4	107.8 (4)
C18—N5—Ag2	125.1 (3)	F5—C24—F4	106.4 (4)
C13—N6—C17	123.7 (4)	F6—C24—S2	112.5 (4)
C13—N6—H6	118.2	F5—C24—S2	111.0 (4)
C17—N6—H6	118.2	F4—C24—S2	110.4 (3)
N1—Ag1—Ag2—N4 ⁱⁱ	160.70 (15)	N4—C12—C13—N6	-179.9 (4)
N2 ⁱ —Ag1—Ag2—N4 ⁱⁱ	-13.47 (14)	N4—C12—C13—C14	0.6 (7)
N1—Ag1—Ag2—N5	5.50 (15)	N6—C13—C14—C15	179.7 (5)
N2 ⁱ —Ag1—Ag2—N5	-168.67 (15)	C12—C13—C14—C15	-0.8 (7)

Ag2—Ag1—N1—C5	82.5 (3)	C13—C14—C15—C16	0.3 (7)
Ag2—Ag1—N1—C1	-95.0 (4)	C12—N4—C16—C15	-0.6 (7)
C5—N1—C1—C2	-0.6 (7)	Ag2 ^{iv} —N4—C16—C15	-174.9 (4)
Ag1—N1—C1—C2	177.0 (3)	C14—C15—C16—N4	0.4 (8)
C6—N3—C2—C1	11.5 (7)	C13—N6—C17—C18	83.2 (6)
C6—N3—C2—C3	-167.8 (5)	C22—N5—C18—C19	0.6 (7)
N1—C1—C2—N3	-180.0 (4)	Ag2—N5—C18—C19	-178.2 (3)
N1—C1—C2—C3	-0.6 (7)	C22—N5—C18—C17	-178.9 (4)
N3—C2—C3—C4	-180.0 (4)	Ag2—N5—C18—C17	2.3 (6)
C1—C2—C3—C4	0.6 (7)	N6—C17—C18—N5	-178.1 (4)
C2—C3—C4—C5	0.5 (7)	N6—C17—C18—C19	2.3 (7)
C1—N1—C5—C4	1.8 (7)	N5—C18—C19—C20	-0.4 (7)
Ag1—N1—C5—C4	-175.9 (4)	C17—C18—C19—C20	179.1 (5)
C3—C4—C5—N1	-1.8 (7)	C18—C19—C20—C21	0.3 (8)
C2—N3—C6—C7	-87.1 (6)	C19—C20—C21—C22	-0.4 (8)
C11—N2—C7—C8	0.5 (7)	C18—N5—C22—C21	-0.7 (7)
Ag1 ⁱⁱⁱ —N2—C7—C8	-177.0 (4)	Ag2—N5—C22—C21	178.1 (4)
C11—N2—C7—C6	-179.0 (4)	C20—C21—C22—N5	0.7 (8)
Ag1 ⁱⁱⁱ —N2—C7—C6	3.6 (6)	O3—S1—C23—F1	-57.7 (4)
N3—C6—C7—N2	-174.5 (4)	O1—S1—C23—F1	62.2 (4)
N3—C6—C7—C8	6.1 (7)	O2—S1—C23—F1	-178.8 (4)
N2—C7—C8—C9	-0.3 (8)	O3—S1—C23—F3	-178.4 (4)
C6—C7—C8—C9	179.1 (5)	O1—S1—C23—F3	-58.5 (4)
C7—C8—C9—C10	-0.8 (8)	O2—S1—C23—F3	60.5 (4)
C8—C9—C10—C11	1.7 (8)	O3—S1—C23—F2	61.9 (4)
C7—N2—C11—C10	0.5 (7)	O1—S1—C23—F2	-178.2 (3)
Ag1 ⁱⁱⁱ —N2—C11—C10	178.1 (4)	O2—S1—C23—F2	-59.2 (4)
C9—C10—C11—N2	-1.6 (8)	O6—S2—C24—F6	-64.1 (4)
N4 ⁱⁱ —Ag2—N5—C22	-48.9 (5)	O5—S2—C24—F6	176.4 (4)
Ag1—Ag2—N5—C22	75.7 (3)	O4—S2—C24—F6	56.0 (4)
N4 ⁱⁱ —Ag2—N5—C18	129.9 (4)	O6—S2—C24—F5	57.6 (4)
Ag1—Ag2—N5—C18	-105.5 (3)	O5—S2—C24—F5	-61.9 (4)
C16—N4—C12—C13	0.1 (7)	O4—S2—C24—F5	177.7 (3)
Ag2 ^{iv} —N4—C12—C13	174.1 (3)	O6—S2—C24—F4	175.4 (3)
C17—N6—C13—C12	-11.8 (7)	O5—S2—C24—F4	56.0 (4)
C17—N6—C13—C14	167.7 (4)	O4—S2—C24—F4	-64.5 (4)

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

Hydrogen-bond geometry (\AA , $^\circ$)

D—H···A	D—H	H···A	D···A	D—H···A
N3—H3···O6 ^v	0.88	2.51	3.206 (6)	136
N6—H6···O3 ^{vi}	0.88	2.43	3.217 (6)	149
C1—H1···O5 ⁱⁱⁱ	0.95	2.55	3.339 (6)	141
C4—H4···O3 ^{vi}	0.95	2.54	3.383 (6)	147
C6—H6A···O5 ^v	0.99	2.57	3.471 (6)	151
C8—H8···N3	0.95	2.57	2.891 (7)	100

C11—H11···F3 ⁱⁱⁱ	0.95	2.49	3.311 (6)	145
C11—H11···O1 ⁱⁱⁱ	0.95	2.54	3.350 (7)	143
C15—H15···O6 ^{vi}	0.95	2.53	3.450 (6)	164
C16—H16···O6 ^{vii}	0.95	2.58	3.316 (6)	135
C17—H17B···O3	0.99	2.58	3.422 (6)	143
C19—H19···N6	0.95	2.59	2.906 (7)	100

Symmetry codes: (iii) $-x+1, y-1/2, -z+1/2$; (v) $x, y-1, z$; (vi) $-x, -y+1, -z$; (vii) $x-1, y-1, z$.