

# Crystal structure of a one-dimensional helical-type silver(I) coordination polymer: *catena*-poly-[[silver(I)- $\mu$ -*N*-(pyridin-4-ylmethyl)pyridine-3-amine- $\kappa^2$ N:N'] nitrate dimethyl sulfoxide disolvate]

Bokhee Moon,<sup>a</sup> Youngeun Jeon,<sup>b</sup> Suk-Hee Moon<sup>c</sup> and Ki-Min Park<sup>d\*</sup>

Received 10 November 2014

Accepted 12 November 2014

Edited by P. C. Healy, Griffith University, Australia

<sup>a</sup>Busan International High School, Busan 614-100, Republic of Korea, <sup>b</sup>Department of Chemistry, Gyeongsang National University, Jinju 660-701, Republic of Korea, <sup>c</sup>Department of Food & Nutrition, Kyungnam College of Information and Technology, Busan 617-701, Republic of Korea, and <sup>d</sup>Research Institute of Natural Sciences, Gyeongsang National University, Jinju 660-701, Republic of Korea. \*Correspondence e-mail: kmpark@gnu.ac.kr

**Keywords:** crystal structure; silver(I) nitrate; unsymmetrical dipyrindyl ligand; helical chain coordination polymer; hydrogen bonding; Ag $\cdots$ O interactions

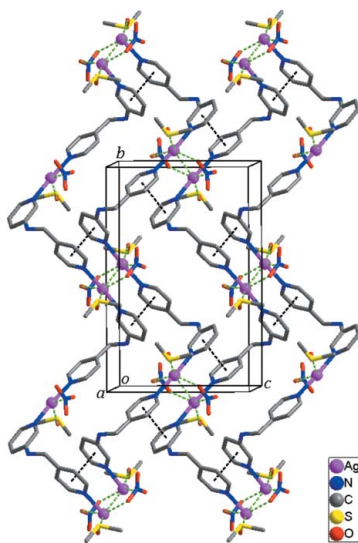
**CCDC reference:** 1033712

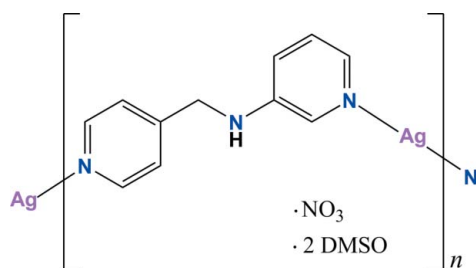
**Supporting information:** this article has supporting information at journals.iucr.org/e

The asymmetric unit of the title compound,  $\{[\text{Ag}(\text{C}_{11}\text{H}_{11}\text{N}_3)]\text{NO}_3 \cdot 2(\text{CH}_3)_2\text{SO}\}_n$ , comprises one Ag<sup>I</sup> atom, one *N*-(pyridine-4-ylmethyl)pyridine-3-amine ligand, one nitrate anion and two dimethyl sulfoxide molecules. The Ag<sup>I</sup> atoms are bridged by two pyridine N atoms from two symmetry-related ligands, forming a helical chain and adopting a slightly distorted linear coordination geometry [ $\text{N}-\text{Ag}-\text{N} = 175.37(8)^\circ$ ]. The helical chain, with a pitch length of 16.7871(8) Å, propagates along the *b*-axis direction. In the crystal, symmetry-related right- and left-handed helical chains are alternately arranged *via* Ag $\cdots$ Ag interactions [3.4145(4) Å] and  $\pi$ - $\pi$  stacking interactions [centroid-centroid distance = 3.650(2) Å], resulting in the formation of a two-dimensional supramolecular network extending parallel to (100). Weak Ag $\cdots$ O [2.775(2), 3.169(4) and 2.690(2) Å] interactions, as well as several N-H $\cdots$ O and C-H $\cdots$ O hydrogen-bonding interactions, contribute to the stabilization of the crystal structure. Parts of the dimethyl sulfoxide solvent molecule are disordered over two sets of sites in a 0.937(3):0.063(3) ratio.

## 1. Chemical context

Self-assembled supramolecular architectures based on the reaction of the silver ion with dipyrindyl-type ligands continue to attract attention not only because of the fascinating structures caused by a variety of coordination geometries for the Ag<sup>I</sup> cation, but also their potential applications as functional materials (Lee *et al.*, 2012; Leong & Vittal, 2011; Park *et al.*, 2010; Zhang *et al.*, 2009, 2013). However, although there has been rapid growth in Ag<sup>I</sup> coordination chemistry based on symmetrical dipyrindyl ligands with nitrogen donor atoms in the same positions on two terminal pyridines, investigations based on unsymmetrical dipyrindyl ligands with nitrogen donor atoms in different positions on two terminal pyridines are still rare (Moon & Park, 2013, 2014; Zhang *et al.*, 2013). Therefore, the development of Ag<sup>I</sup> coordination polymers using unsymmetrical dipyrindyl ligands is a challenging project and deserves exploration. Herein, we report the crystal structure of the title compound prepared by the reaction of silver nitrate with the unsymmetrical dipyrindyl ligand, *N*-(pyridin-4-ylmethyl)pyridine-3-amine, which was synthesized by the reaction of 3-aminopyridine and pyridine-4-carboxaldehyde according to literature methods (Foxon *et al.*, 2002; Lee *et al.*, 2013). The structure of the title compound is related to that of the monohydrated Ag<sup>I</sup> coordination polymer with the same ligand (Zhang *et al.*, 2013).



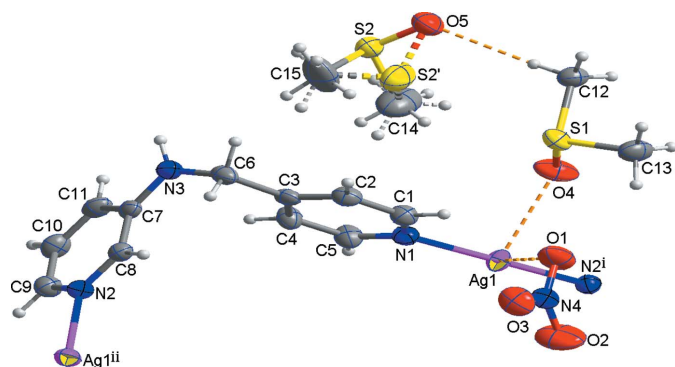


## 2. Structural commentary

The molecular components of the title structure are shown in Fig. 1. The asymmetric unit consists of one  $\text{Ag}^{\text{I}}$  atom, one *N*-(pyridin-4-ylmethyl)pyridine-3-amine ligand, one nitrate anion and two DMSO molecules. The S atom of one of the DMSO molecules is disordered over two sites [site-occupancy factors of 0.937 (3) for S2 and 0.063 (3) for S2']. The Ag atom links two pyridine N atoms from two symmetry-related ligands, forming a helical chain. Thus the  $\text{Ag}^{\text{I}}$  atom is two-coordinate in a slightly distorted linear coordination geometry [ $\text{N}-\text{Ag}-\text{N} = 175.37(8)^\circ$ ], with the  $\text{Ag}-\text{N}$  bond lengths of 2.158 (2) and 2.162 (2) Å. The helical chain propagates along the *b*-axis direction (Fig. 2) and its pitch length is 16.7871 (8) Å, much longer than that [10.135 (2) Å] of the monohydrated  $\text{Ag}^{\text{I}}$  coordination polymer reported by Zhang *et al.* (2013). The two pyridine rings coordinating to the Ag atom are tilted by 9.77 (16)° with respect to each other. In the *N*-(pyridin-4-ylmethyl)pyridine-3-amine ligand, the two pyridine rings are almost perpendicular, the dihedral angle between their mean planes being 86.28 (7)°.

## 3. Supramolecular features

In the crystal structure, the symmetry-related right- and left-handed helical chains are alternately arranged in the structure via  $\text{Ag}\cdots\text{Ag}$  interactions [3.4145 (4) Å], resulting in the



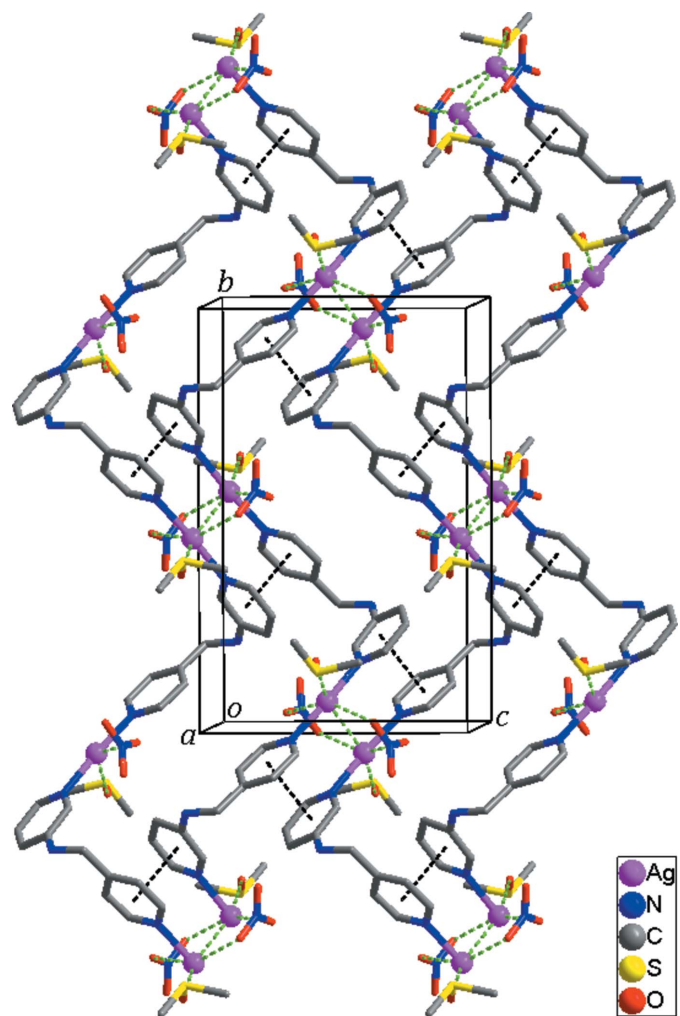
**Figure 1**  
A view of the molecular structure of the title compound, with the atom numbering. Displacement ellipsoids are drawn at the 50% probability level and two-coloured dashed lines indicate the disordered part of DMSO.  $\text{Ag}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{O}$  interactions are shown as yellow dashed lines. [Symmetry codes: (i)  $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$ ; (ii)  $-x + 1, y - \frac{1}{2}, -z + \frac{1}{2}$ ]

**Table 1**  
Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N3}-\text{H3}\cdots\text{O3}^{\text{i}}$	0.88	2.17	3.042 (3)	173
$\text{C1}-\text{H1}\cdots\text{O1}$	0.95	2.55	3.306 (4)	136
$\text{C5}-\text{H5}\cdots\text{O2}^{\text{ii}}$	0.95	2.32	3.151 (3)	145
$\text{C6}-\text{H6A}\cdots\text{O3}^{\text{iii}}$	0.99	2.42	3.405 (4)	175
$\text{C8}-\text{H8}\cdots\text{O3}^{\text{iii}}$	0.95	2.55	3.480 (4)	168
$\text{C10}-\text{H10}\cdots\text{O4}^{\text{iv}}$	0.95	2.44	3.309 (4)	152
$\text{C12}-\text{H12A}\cdots\text{O5}$	0.98	2.43	3.377 (4)	161
$\text{C12}-\text{H12B}\cdots\text{O5}^{\text{v}}$	0.98	2.55	3.478 (4)	159
$\text{C12}-\text{H12C}\cdots\text{O3}^{\text{vi}}$	0.98	2.35	3.270 (4)	156
$\text{C13}-\text{H13A}\cdots\text{O1}$	0.98	2.53	3.292 (4)	134
$\text{C15}-\text{H15C}\cdots\text{O2}^{\text{i}}$	0.98	2.59	3.470 (6)	149

Symmetry codes: (i)  $x, y, z + 1$ ; (ii)  $-x + 1, -y + 1, -z$ ; (iii)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (iv)  $-x + 1, -y + 1, -z + 1$ ; (v)  $x, -y + \frac{3}{2}, z - \frac{1}{2}$ ; (vi)  $-x + 2, -y + 1, -z$ .

formation of a two-dimensional supramolecular network extending parallel to (100) (Fig. 2).  $\pi-\pi$  stacking interactions [centroid-centroid distance = 3.650 (2) Å] between the pyridine rings of both helical chains contribute to the stabi-



**Figure 2**  
The two-dimensional supramolecular network formed through  $\text{Ag}\cdots\text{Ag}$  and  $\text{Ag}\cdots\text{O}$  interactions (green dashed lines) as well as  $\pi-\pi$  stacking interactions (black dashed lines).

**Table 2**  
Experimental details.

Crystal data	
Chemical formula	[Ag(C <sub>11</sub> H <sub>11</sub> N <sub>3</sub> )]NO <sub>3</sub> ·2C <sub>2</sub> H <sub>6</sub> OS
<i>M<sub>r</sub></i>	511.36
Crystal system, space group	Monoclinic, <i>P</i> 2 <sub>1</sub> / <i>c</i>
Temperature (K)	173
<i>a</i> , <i>b</i> , <i>c</i> (Å)	11.7046 (6), 16.7871 (8), 10.4922 (5)
$\beta$ (°)	91.950 (1)
<i>V</i> (Å <sup>3</sup> )	2060.38 (17)
<i>Z</i>	4
Radiation type	Mo <i>K</i> $\alpha$
$\mu$ (mm <sup>-1</sup> )	1.21
Crystal size (mm)	0.31 × 0.24 × 0.12
Data collection	
Diffraction	Bruker SMART CCD
Absorption correction	Multi-scan ( <i>SADABS</i> ; Bruker, 2000)
<i>T<sub>min</sub></i> , <i>T<sub>max</sub></i>	0.705, 0.868
No. of measured, independent and observed [ <i>I</i> > 2 $\sigma$ ( <i>I</i> )] reflections	11476, 4039, 3485
<i>R<sub>int</sub></i>	0.063
( <i>sin</i> $\theta$ / $\lambda$ ) <sub>max</sub> (Å <sup>-1</sup> )	0.617
Refinement	
<i>R</i> [ <i>F</i> <sup>2</sup> > 2 $\sigma$ ( <i>F</i> <sup>2</sup> )], <i>wR</i> ( <i>F</i> <sup>2</sup> ), <i>S</i>	0.028, 0.078, 1.08
No. of reflections	4039
No. of parameters	255
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{max}$ , $\Delta\rho_{min}$ (e Å <sup>-3</sup> )	0.58, -0.63

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

zation of the two-dimensional network. The two-dimensional networks are further stabilized by Ag $\cdots$ O interactions [Ag1 $\cdots$ O1 = 2.775 (2), Ag1 $\cdots$ O2<sup>i</sup> = 3.169 (4) and Ag1 $\cdots$ O4 = 2.690 (2) Å; symmetry code: (i)  $-x + 1, -y + 1, -z$ ] (Fig. 2), as well as N–H $\cdots$ O and C–H $\cdots$ O hydrogen bonds between the helical chains and the nitrate anions or the DMSO molecules (Table 1). In addition, several C–H $\cdots$ O hydrogen bonds between the DMSO molecules, and between the DMSO molecules and the nitrate anions are also observed.

#### 4. Database survey

The structures of the silver(I) nitrate and perchlorate complexes of the same ligand have been reported as their monohydrated and non-solvated forms, respectively, by Zhang *et al.* (2013). These complexes have been also studied for their luminescent properties in the solid state.

#### 5. Synthesis and crystallization

*N*-(Pyridin-4-ylmethyl)pyridine-3-amine was prepared according to the procedure described by Lee *et al.* (2013) and Foxon *et al.* (2002). 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) nitrate in the molar ratio 1:1 in methanol.

#### 6. Refinement details

Crystal data, data collection and structure refinement details are summarized in Table 2. Atoms S2 and S2' of one DMSO molecule are disordered over two sites with site-occupation factors of 0.937 (3) and 0.063 (3), respectively. All H atoms were positioned geometrically and refined using a riding model, with C–H = 0.95 Å for Csp<sup>2</sup>–H, 0.88 Å for amine N–H and 0.99 Å for methylene C–H. For all H atoms, *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C,N).

#### Acknowledgements

This work was supported by NRF (2010-0022675) projects.

#### References

- Brandenburg, K. (2005). *DIAMOND*. Crystal Impact GbR, Germany.
- Bruker. (2000). *SMART*, *SAINT-Plus* and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Foxon, S. P., Walter, O. & Schindler, S. (2002). *Eur. J. Inorg. Chem.* pp. 111–121.
- Lee, E., Ryu, H., Moon, S.-H. & Park, K.-M. (2013). *Bull. Korean Chem. Soc.* **34**, 3477–3480.
- Lee, E., Seo, J., Lee, S. S. & Park, K.-M. (2012). *Cryst. Growth Des.* **12**, 3834–3837.
- Leong, W. L. & Vittal, J. J. (2011). *Chem. Rev.* **111**, 688–764.
- Moon, S.-H. & Park, K.-M. (2013). *Acta Cryst.* **E69**, m414–m415.
- Moon, S.-H. & Park, K.-M. (2014). *Acta Cryst.* **E70**, m233.
- Park, K.-M., Seo, J., Moon, S.-H. & Lee, S. S. (2010). *Cryst. Growth Des.* **10**, 4148–4154.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Zhang, Z.-Y., Deng, Z.-P., Huo, L.-H., Zhao, H. & Gao, S. (2013). *Inorg. Chem.* **52**, 5914–5923.
- Zhang, W., Ye, H. & Xiong, R. G. (2009). *Coord. Chem. Rev.* **253**, 2980–2997.

## supporting information

*Acta Cryst.* (2014). E70, 507-509 [doi:10.1107/S1600536814024817]

**Crystal structure of a one-dimensional helical-type silver(I) coordination polymer: *catena*-poly[[silver(I)- $\mu$ -*N*-(pyridin-4-ylmethyl)pyridine-3-amine- $\kappa^2$ *N:N'*] nitrate dimethyl sulfoxide disolvate]**

**Bokhee Moon, Youngeun Jeon, Suk-Hee Moon and Ki-Min Park**

**Computing details**

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE-Plus* (Bruker, 2000); data reduction: *SAINTE-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)- $\mu$ -*N*-(pyridin-4-ylmethyl)pyridine-3-amine- $\kappa^2$ *N:N'*] nitrate dimethyl sulfoxide disolvate]**

*Crystal data*

[Ag(C<sub>11</sub>H<sub>11</sub>N<sub>3</sub>)]NO<sub>3</sub>·2C<sub>2</sub>H<sub>6</sub>OS

$M_r = 511.36$

Monoclinic, *P2<sub>1</sub>/c*

Hall symbol: -P 2ybc

$a = 11.7046$  (6) Å

$b = 16.7871$  (8) Å

$c = 10.4922$  (5) Å

$\beta = 91.950$  (1)°

$V = 2060.38$  (17) Å<sup>3</sup>

$Z = 4$

$F(000) = 1040$

$D_x = 1.649$  Mg m<sup>-3</sup>

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

Cell parameters from 7555 reflections

$\theta = 2.3$ – $28.3$ °

$\mu = 1.21$  mm<sup>-1</sup>

$T = 173$  K

Plate, colorless

$0.31 \times 0.24 \times 0.12$  mm

*Data collection*

Bruker SMART CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

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

$T_{\min} = 0.705$ ,  $T_{\max} = 0.868$

11476 measured reflections

4039 independent reflections

3485 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.063$

$\theta_{\max} = 26.0$ °,  $\theta_{\min} = 1.7$ °

$h = -12 \rightarrow 14$

$k = -20 \rightarrow 19$

$l = -12 \rightarrow 12$

*Refinement*

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.028$

$wR(F^2) = 0.078$

$S = 1.08$

4039 reflections

255 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0348P)^2 + 1.0473P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.58 \text{ e } \text{Å}^{-3}$   
 $\Delta\rho_{\min} = -0.63 \text{ e } \text{Å}^{-3}$

#### 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 ( $\text{Å}^2$ )

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Ag1	0.599664 (17)	0.557957 (11)	0.078155 (18)	0.03159 (9)	
N1	0.64574 (18)	0.46711 (13)	0.2167 (2)	0.0311 (5)	
N2	0.45491 (18)	0.15285 (12)	0.5486 (2)	0.0287 (5)	
N3	0.6908 (2)	0.29282 (13)	0.6162 (2)	0.0346 (5)	
H3	0.7182	0.3160	0.6859	0.092 (15)*	
C1	0.7468 (2)	0.42880 (15)	0.2103 (3)	0.0328 (6)	
H1	0.7945	0.4404	0.1412	0.039*	
C2	0.7836 (2)	0.37356 (15)	0.2997 (3)	0.0321 (6)	
H2	0.8551	0.3477	0.2911	0.039*	
C3	0.7164 (2)	0.35560 (14)	0.4024 (2)	0.0279 (5)	
C4	0.6126 (2)	0.39512 (16)	0.4087 (3)	0.0355 (6)	
H4	0.5635	0.3849	0.4772	0.043*	
C5	0.5808 (2)	0.44944 (16)	0.3151 (3)	0.0373 (6)	
H5	0.5092	0.4755	0.3211	0.045*	
C6	0.7578 (2)	0.29614 (15)	0.5034 (3)	0.0336 (6)	
H6A	0.7585	0.2425	0.4644	0.040*	
H6B	0.8375	0.3096	0.5294	0.040*	
C7	0.5858 (2)	0.25525 (14)	0.6209 (2)	0.0284 (5)	
C8	0.5553 (2)	0.19125 (14)	0.5408 (2)	0.0270 (5)	
H8	0.6074	0.1744	0.4788	0.032*	
C9	0.3803 (2)	0.17604 (17)	0.6358 (3)	0.0350 (6)	
H9	0.3100	0.1483	0.6420	0.042*	
C10	0.4037 (3)	0.23941 (17)	0.7165 (3)	0.0388 (6)	
H10	0.3495	0.2553	0.7769	0.047*	
C11	0.5062 (2)	0.27946 (16)	0.7091 (2)	0.0356 (6)	
H11	0.5226	0.3234	0.7639	0.043*	
S1	0.90399 (6)	0.61772 (4)	0.14218 (7)	0.03541 (16)	
O4	0.78359 (17)	0.64866 (14)	0.1364 (2)	0.0545 (6)	
C12	0.9849 (2)	0.68750 (17)	0.2364 (3)	0.0392 (7)	
H12A	0.9627	0.6838	0.3254	0.047*	
H12B	0.9699	0.7415	0.2046	0.047*	
H12C	1.0665	0.6755	0.2309	0.047*	

C13	0.9641 (3)	0.6403 (2)	-0.0077 (3)	0.0470 (7)	
H13A	0.9288	0.6065	-0.0741	0.071*	
H13B	1.0467	0.6305	-0.0025	0.071*	
H13C	0.9500	0.6964	-0.0287	0.071*	
S2	0.85523 (7)	0.61054 (5)	0.61168 (7)	0.0404 (3)	0.937 (3)
S2'	0.8683 (14)	0.5787 (10)	0.4964 (13)	0.060 (5)	0.063 (3)
O5	0.95718 (18)	0.64284 (13)	0.5476 (2)	0.0486 (5)	
C14	0.7356 (3)	0.6311 (3)	0.5101 (4)	0.0866 (14)	
H14A	0.7204	0.6885	0.5099	0.104*	0.937 (3)
H14B	0.6687	0.6026	0.5407	0.104*	0.937 (3)
H14C	0.7512	0.6135	0.4234	0.104*	0.937 (3)
H14D	0.7343	0.6771	0.4527	0.104*	0.063 (3)
H14E	0.7282	0.6493	0.5982	0.104*	0.063 (3)
H14F	0.6719	0.5954	0.4870	0.104*	0.063 (3)
C15	0.8616 (4)	0.5055 (2)	0.5930 (5)	0.0829 (14)	
H15A	0.9245	0.4841	0.6466	0.100*	0.937 (3)
H15B	0.8743	0.4926	0.5035	0.100*	0.937 (3)
H15C	0.7893	0.4818	0.6185	0.100*	0.937 (3)
H15D	0.9322	0.4742	0.5905	0.100*	0.063 (3)
H15E	0.7963	0.4716	0.5681	0.100*	0.063 (3)
H15F	0.8519	0.5256	0.6796	0.100*	0.063 (3)
N4	0.72721 (19)	0.45059 (12)	-0.1442 (2)	0.0309 (5)	
O1	0.7696 (2)	0.50433 (13)	-0.0788 (2)	0.0544 (6)	
O2	0.6392 (2)	0.46146 (17)	-0.2082 (3)	0.0807 (10)	
O3	0.77059 (19)	0.38352 (12)	-0.1461 (2)	0.0521 (6)	

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Ag1	0.03243 (13)	0.03166 (13)	0.03064 (12)	0.00682 (8)	0.00031 (9)	0.00448 (8)
N1	0.0263 (11)	0.0307 (11)	0.0361 (12)	0.0013 (9)	-0.0021 (9)	0.0028 (9)
N2	0.0270 (11)	0.0292 (11)	0.0298 (11)	-0.0012 (8)	0.0004 (9)	-0.0015 (9)
N3	0.0403 (13)	0.0306 (11)	0.0324 (12)	-0.0062 (9)	-0.0075 (10)	0.0020 (9)
C1	0.0271 (13)	0.0353 (14)	0.0360 (14)	0.0018 (10)	0.0035 (11)	-0.0005 (11)
C2	0.0249 (12)	0.0312 (13)	0.0402 (14)	0.0039 (10)	-0.0003 (11)	-0.0028 (11)
C3	0.0261 (12)	0.0207 (11)	0.0365 (13)	-0.0034 (9)	-0.0055 (11)	-0.0014 (10)
C4	0.0284 (13)	0.0354 (14)	0.0427 (15)	0.0014 (11)	0.0040 (12)	0.0086 (12)
C5	0.0266 (13)	0.0375 (15)	0.0480 (17)	0.0055 (11)	0.0022 (12)	0.0085 (12)
C6	0.0279 (13)	0.0272 (13)	0.0451 (15)	-0.0028 (10)	-0.0075 (12)	0.0051 (11)
C7	0.0359 (14)	0.0239 (12)	0.0249 (11)	-0.0003 (10)	-0.0053 (10)	0.0037 (10)
C8	0.0285 (12)	0.0272 (12)	0.0253 (12)	0.0015 (10)	-0.0010 (10)	-0.0009 (9)
C9	0.0301 (13)	0.0391 (15)	0.0358 (14)	0.0001 (11)	0.0030 (11)	-0.0003 (12)
C10	0.0436 (16)	0.0423 (16)	0.0309 (13)	0.0084 (13)	0.0071 (12)	-0.0037 (12)
C11	0.0474 (16)	0.0306 (13)	0.0283 (12)	0.0044 (12)	-0.0044 (12)	-0.0053 (11)
S1	0.0319 (3)	0.0323 (3)	0.0423 (4)	-0.0020 (3)	0.0062 (3)	-0.0029 (3)
O4	0.0266 (10)	0.0654 (14)	0.0718 (15)	-0.0028 (10)	0.0073 (10)	-0.0272 (13)
C12	0.0343 (15)	0.0469 (17)	0.0360 (14)	-0.0019 (12)	-0.0018 (12)	-0.0044 (13)
C13	0.0429 (17)	0.063 (2)	0.0353 (15)	-0.0057 (15)	0.0019 (13)	-0.0093 (14)

S2	0.0469 (5)	0.0423 (5)	0.0318 (4)	-0.0074 (3)	0.0008 (3)	-0.0013 (3)
S2'	0.057 (9)	0.081 (11)	0.040 (7)	-0.001 (7)	0.004 (6)	0.001 (7)
O5	0.0421 (12)	0.0470 (12)	0.0565 (13)	-0.0080 (9)	0.0009 (10)	0.0018 (10)
C14	0.043 (2)	0.141 (4)	0.075 (3)	-0.001 (2)	-0.001 (2)	0.009 (3)
C15	0.113 (4)	0.042 (2)	0.095 (3)	-0.023 (2)	0.024 (3)	-0.003 (2)
N4	0.0269 (11)	0.0321 (12)	0.0340 (12)	0.0015 (9)	0.0039 (9)	-0.0025 (9)
O1	0.0673 (15)	0.0464 (12)	0.0498 (13)	-0.0149 (11)	0.0055 (11)	-0.0193 (11)
O2	0.0514 (15)	0.0690 (17)	0.119 (3)	0.0119 (13)	-0.0377 (17)	0.0006 (17)
O3	0.0473 (13)	0.0342 (11)	0.0748 (16)	0.0130 (9)	0.0028 (12)	0.0001 (11)

*Geometric parameters (Å, °)*

Ag1—N2 <sup>i</sup>	2.158 (2)	S1—C13	1.785 (3)
Ag1—N1	2.162 (2)	C12—H12A	0.9800
N1—C5	1.337 (4)	C12—H12B	0.9800
N1—C1	1.350 (3)	C12—H12C	0.9800
N2—C9	1.343 (4)	C13—H13A	0.9800
N2—C8	1.345 (3)	C13—H13B	0.9800
N2—Ag1 <sup>ii</sup>	2.158 (2)	C13—H13C	0.9800
N3—C7	1.383 (3)	S2—O5	1.492 (2)
N3—C6	1.443 (4)	S2—C14	1.764 (4)
N3—H3	0.8800	S2—C15	1.777 (4)
C1—C2	1.378 (4)	S2—H14E	1.6245
C1—H1	0.9500	S2—H15F	1.5946
C2—C3	1.389 (4)	S2'—O5	1.579 (16)
C2—H2	0.9500	S2'—C15	1.597 (16)
C3—C4	1.388 (4)	S2'—C14	1.794 (17)
C3—C6	1.523 (3)	C14—H14A	0.9800
C4—C5	1.382 (4)	C14—H14B	0.9800
C4—H4	0.9500	C14—H14C	0.9800
C5—H5	0.9500	C14—H14D	0.9799
C6—H6A	0.9900	C14—H14E	0.9799
C6—H6B	0.9900	C14—H14F	0.9800
C7—C11	1.396 (4)	C15—H15A	0.9800
C7—C8	1.402 (3)	C15—H15B	0.9800
C8—H8	0.9500	C15—H15C	0.9800
C9—C10	1.381 (4)	C15—H15D	0.9800
C9—H9	0.9500	C15—H15E	0.9801
C10—C11	1.380 (4)	C15—H15F	0.9799
C10—H10	0.9500	N4—O2	1.224 (3)
C11—H11	0.9500	N4—O1	1.228 (3)
S1—O4	1.501 (2)	N4—O3	1.236 (3)
S1—C12	1.784 (3)		
N2 <sup>i</sup> —Ag1—N1	175.37 (8)	O5—S2—H14E	124.0
C5—N1—C1	117.0 (2)	C15—S2—H14E	115.4
C5—N1—Ag1	122.74 (17)	O5—S2—H15F	124.0
C1—N1—Ag1	120.11 (19)	C14—S2—H15F	114.4

C9—N2—C8	119.5 (2)	H14E—S2—H15F	111.1
C9—N2—Ag1 <sup>ii</sup>	116.62 (17)	O5—S2'—C15	110.8 (9)
C8—N2—Ag1 <sup>ii</sup>	123.85 (17)	O5—S2'—C14	101.6 (9)
C7—N3—C6	123.7 (2)	C15—S2'—C14	105.3 (9)
C7—N3—H3	118.2	S2—O5—S2'	51.5 (6)
C6—N3—H3	118.2	S2—C14—H14A	109.5
N1—C1—C2	122.7 (3)	S2'—C14—H14A	129.7
N1—C1—H1	118.6	S2—C14—H14B	109.5
C2—C1—H1	118.6	S2'—C14—H14B	119.4
C1—C2—C3	120.2 (2)	H14A—C14—H14B	109.5
C1—C2—H2	119.9	S2—C14—H14C	109.5
C3—C2—H2	119.9	S2'—C14—H14C	65.7
C4—C3—C2	116.9 (2)	H14A—C14—H14C	109.5
C4—C3—C6	122.6 (2)	H14B—C14—H14C	109.5
C2—C3—C6	120.5 (2)	S2—C14—H14D	121.4
C5—C4—C3	119.8 (3)	S2'—C14—H14D	109.4
C5—C4—H4	120.1	H14B—C14—H14D	126.1
C3—C4—H4	120.1	H14C—C14—H14D	70.5
N1—C5—C4	123.3 (3)	S2—C14—H14E	65.5
N1—C5—H5	118.3	S2'—C14—H14E	109.5
C4—C5—H5	118.3	H14A—C14—H14E	70.9
N3—C6—C3	115.3 (2)	H14B—C14—H14E	75.3
N3—C6—H6A	108.5	H14C—C14—H14E	174.3
C3—C6—H6A	108.5	H14D—C14—H14E	109.5
N3—C6—H6B	108.5	S2—C14—H14F	127.7
C3—C6—H6B	108.5	S2'—C14—H14F	109.5
H6A—C6—H6B	107.5	H14A—C14—H14F	117.6
N3—C7—C11	120.3 (2)	H14C—C14—H14F	75.5
N3—C7—C8	122.5 (2)	H14D—C14—H14F	109.5
C11—C7—C8	117.2 (2)	H14E—C14—H14F	109.5
N2—C8—C7	122.4 (2)	S2'—C15—S2	46.3 (6)
N2—C8—H8	118.8	S2'—C15—H15A	126.5
C7—C8—H8	118.8	S2—C15—H15A	109.5
N2—C9—C10	121.4 (3)	S2'—C15—H15B	63.2
N2—C9—H9	119.3	S2—C15—H15B	109.5
C10—C9—H9	119.3	H15A—C15—H15B	109.5
C11—C10—C9	119.7 (3)	S2'—C15—H15C	123.2
C11—C10—H10	120.2	S2—C15—H15C	109.5
C9—C10—H10	120.2	H15A—C15—H15C	109.5
C10—C11—C7	119.8 (2)	H15B—C15—H15C	109.5
C10—C11—H11	120.1	S2'—C15—H15D	109.6
C7—C11—H11	120.1	S2—C15—H15D	124.9
O4—S1—C12	105.96 (13)	H15B—C15—H15D	72.5
O4—S1—C13	106.77 (15)	H15C—C15—H15D	121.8
C12—S1—C13	97.50 (14)	S2'—C15—H15E	109.4
S1—C12—H12A	109.5	S2—C15—H15E	124.8
S1—C12—H12B	109.5	H15A—C15—H15E	120.3
H12A—C12—H12B	109.5	H15B—C15—H15E	76.1



S1—C12—H12C	109.5	H15D—C15—H15E	109.5
H12A—C12—H12C	109.5	S2'—C15—H15F	109.4
H12B—C12—H12C	109.5	S2—C15—H15F	63.1
S1—C13—H13A	109.5	H15A—C15—H15F	72.7
S1—C13—H13B	109.5	H15B—C15—H15F	172.3
H13A—C13—H13B	109.5	H15C—C15—H15F	76.1
S1—C13—H13C	109.5	H15D—C15—H15F	109.5
H13A—C13—H13C	109.5	H15E—C15—H15F	109.5
H13B—C13—H13C	109.5	O2—N4—O1	120.9 (2)
O5—S2—C14	106.70 (18)	O2—N4—O3	117.7 (2)
O5—S2—C15	105.90 (18)	O1—N4—O3	121.3 (2)
C14—S2—C15	99.3 (3)		
C5—N1—C1—C2	0.1 (4)	C8—N2—C9—C10	1.0 (4)
Ag1—N1—C1—C2	176.35 (19)	Ag1 <sup>ii</sup> —N2—C9—C10	-176.0 (2)
N1—C1—C2—C3	-0.5 (4)	N2—C9—C10—C11	-0.8 (4)
C1—C2—C3—C4	0.4 (4)	C9—C10—C11—C7	-0.5 (4)
C1—C2—C3—C6	-178.5 (2)	N3—C7—C11—C10	-177.3 (2)
C2—C3—C4—C5	0.0 (4)	C8—C7—C11—C10	1.5 (4)
C6—C3—C4—C5	178.9 (2)	C14—S2—O5—S2'	-58.3 (8)
C1—N1—C5—C4	0.3 (4)	C15—S2—O5—S2'	46.9 (8)
Ag1—N1—C5—C4	-175.8 (2)	C15—S2'—O5—S2	-56.6 (7)
C3—C4—C5—N1	-0.4 (4)	C14—S2'—O5—S2	54.9 (5)
C7—N3—C6—C3	76.5 (3)	O5—S2—C14—S2'	57.3 (8)
C4—C3—C6—N3	-9.4 (3)	C15—S2—C14—S2'	-52.5 (8)
C2—C3—C6—N3	169.5 (2)	O5—S2'—C14—S2	-51.1 (5)
C6—N3—C7—C11	-154.1 (2)	C15—S2'—C14—S2	64.6 (6)
C6—N3—C7—C8	27.1 (4)	O5—S2'—C15—S2	49.4 (6)
C9—N2—C8—C7	0.0 (4)	C14—S2'—C15—S2	-59.7 (7)
Ag1 <sup>ii</sup> —N2—C8—C7	176.86 (17)	O5—S2—C15—S2'	-51.4 (8)
N3—C7—C8—N2	177.5 (2)	C14—S2—C15—S2'	59.1 (8)
C11—C7—C8—N2	-1.3 (3)		

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

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

<i>D</i> —H $\cdots$ <i>A</i>	<i>D</i> —H	H $\cdots$ <i>A</i>	<i>D</i> $\cdots$ <i>A</i>	<i>D</i> —H $\cdots$ <i>A</i>
N3—H3 $\cdots$ O3 <sup>iii</sup>	0.88	2.17	3.042 (3)	173
C1—H1 $\cdots$ O1	0.95	2.55	3.306 (4)	136
C5—H5 $\cdots$ O2 <sup>iv</sup>	0.95	2.32	3.151 (3)	145
C6—H6A $\cdots$ O3 <sup>v</sup>	0.99	2.42	3.405 (4)	175
C8—H8 $\cdots$ O3 <sup>v</sup>	0.95	2.55	3.480 (4)	168
C10—H10 $\cdots$ O4 <sup>vi</sup>	0.95	2.44	3.309 (4)	152
C12—H12A $\cdots$ O5	0.98	2.43	3.377 (4)	161
C12—H12B $\cdots$ O5 <sup>vii</sup>	0.98	2.55	3.478 (4)	159
C12—H12C $\cdots$ O3 <sup>viii</sup>	0.98	2.35	3.270 (4)	156

C13—H13A···O1	0.98	2.53	3.292 (4)	134
C15—H15C···O2 <sup>iii</sup>	0.98	2.59	3.470 (6)	149

---

Symmetry codes: (iii)  $x, y, z+1$ ; (iv)  $-x+1, -y+1, -z$ ; (v)  $x, -y+1/2, z+1/2$ ; (vi)  $-x+1, -y+1, -z+1$ ; (vii)  $x, -y+3/2, z-1/2$ ; (viii)  $-x+2, -y+1, -z$ .