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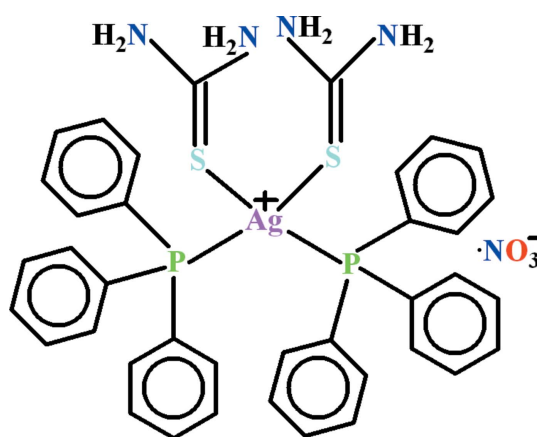
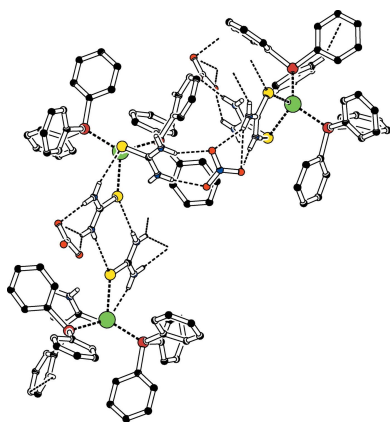
# Crystal structure of bis(thiourea- $\kappa$ S)bis(triphenylphosphane- $\kappa$ P)silver(I) nitrate

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In the title salt,  $[\text{Ag}(\text{CH}_4\text{N}_2\text{S})_2(\text{PPh}_3)_2]\text{NO}_3$ , the  $\text{Ag}^{\text{I}}$  atom is coordinated by two thiourea S atoms and two triphenylphosphane P atoms in a distorted tetrahedral geometry, with bond angles in the range  $102.90(4)$ – $123.29(4)^\circ$ . The  $\text{Ag}-\text{S}=\text{C}$  bond angles are  $101.75(19)$  and  $111.29(18)^\circ$ . In the crystal, the component ions are linked by  $\text{C}-\text{H}\cdots\text{O}$ ,  $\text{C}-\text{H}\cdots\text{S}$ ,  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{N}-\text{H}\cdots\text{S}$  hydrogen bonds, generating  $(10\bar{1})$  sheets.

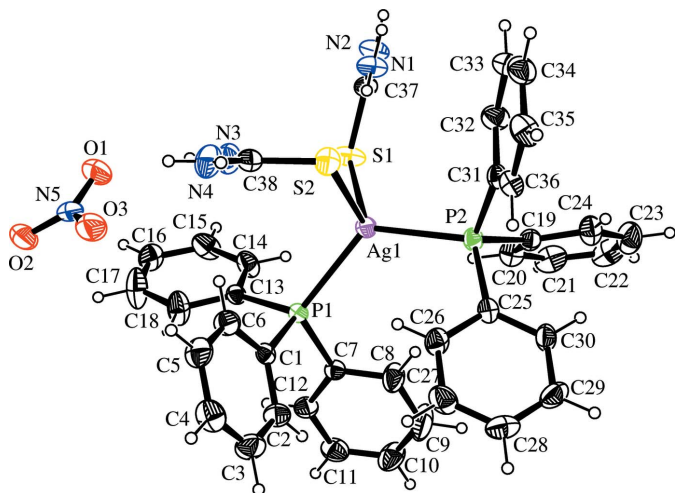
## 1. Chemical context

Silver(I) forms relatively stable compounds with phosphanes and sulfur donor thione ligands due to favorable soft acid–soft base interactions (Ferrari *et al.*, 2007; Isab *et al.*, 2010; Karagiannidis *et al.*, 1990; Nawaz *et al.*, 2011; Ruffer *et al.*, 2011). Interest in these complexes arises from their luminescent (Ferrari *et al.*, 2007), antimicrobial (Ruan *et al.*, 2009) and antitumor properties (Liu *et al.*, 2008). In the light of this, the crystal structures of several silver(I) complexes of phosphanes and thiones have been reported in the literature (Ferrari *et al.*, 2007; Isab *et al.*, 2010; Karagiannidis *et al.*, 1990; Nawaz *et al.*, 2011; Ruffer *et al.*, 2011). Here, we report the crystal structure of a new silver(I) complex of triphenylphosphane ( $\text{PPh}_3$ ) and thiourea (tu), (I) (Fig. 1).



## 2. Structural commentary

The crystal structure of the title complex consists of  $[\text{Ag}(\text{PPh}_3)_2(\text{tu})_2]^+$  cations and  $\text{NO}_3^-$  counter-ions. In the cationic complex,  $[\text{Ag}(\text{PPh}_3)_2(\text{tu})_2]^+$ , the silver(I) atom is bound to two P atoms of  $\text{PPh}_3$  and two sulfur atoms of thiourea, assuming a slightly distorted tetrahedral geometry



**Figure 1**  
View of the title compound with displacement ellipsoids drawn at the 50% probability level.

(Fig. 1). The spread of bond angles around the Ag atom is 102.90 (4)–123.29 (4)°. The high value of the P1–Ag1–P2 angle [123.29 (4)°] is counterbalanced by the smaller S1–Ag1–S2 bond angle [102.90 (4)°]. The deviation from a tetrahedral geometry is apparently due to steric interaction between the bulky phosphane ligands. The Ag–S, Ag–P and other bond lengths (Table 1) are in agreement with those observed in other reported complexes (Ferrari *et al.*, 2007; Isab *et al.*, 2010; Karagiannidis *et al.*, 1990; Nawaz *et al.*, 2011; Rüffer *et al.*, 2011). The nitrate ion is planar, but exhibits low symmetry due to rather strong hydrogen-bonding interactions with the NH group of the tu ligand.

In (I), the dihedral angle between the phenyl rings *A* (C1–C6), *B* (C7–C12), *C* (C13–C18), *D* (C19–C24), *E* (C25–C30) and *F* (C31–C36) are as follows: *A/B*, *A/C*, *B/C*, *D/E*, *D/F* and *E/F* = 82.67 (15), 62.77 (17), 86.59 (14), 73.72 (14), 85.01 (16) and 84.06 (17)°, respectively. The thiourea units *G* (S1/C37/N1/N2) and *H* (S2/C38/N3/N4) are almost planar with r.m.s. deviations of 0.0031 and 0.0007 Å, respectively, and are oriented at a dihedral angle of 76.82 (11)° to each other.

### 3. Supramolecular features

In the asymmetric unit, strong N–H···S, N–H···O hydrogen bonds complete distorted *S*(6) and *R*<sub>2</sub><sup>2</sup>(8) loops. The other hydrogen-bonding interactions are of the C–H···O, C–H···S, N–H···O and N–H···S types (Table 2, Fig. 2) and lead to a two-dimensional polymeric network in the (10 $\bar{1}$ ) plane.

### 4. Synthesis and crystallization

The title complex was prepared by adding one equivalent of thiourea dissolved in 10 ml methanol to a 1:1 mixture of AgNO<sub>3</sub> and PPh<sub>3</sub> in a methanol–acetonitrile medium (10 ml and 15 ml, respectively). Mixing resulted in the formation of a white precipitate. After stirring for half an hour, the mixture

**Table 1**  
Selected bond lengths (Å).

|        |             |        |             |
|--------|-------------|--------|-------------|
| Ag1–P2 | 2.4888 (13) | Ag1–S1 | 2.6263 (13) |
| Ag1–P1 | 2.5078 (12) | Ag1–S2 | 2.6683 (13) |

**Table 2**  
Hydrogen-bond geometry (Å, °).

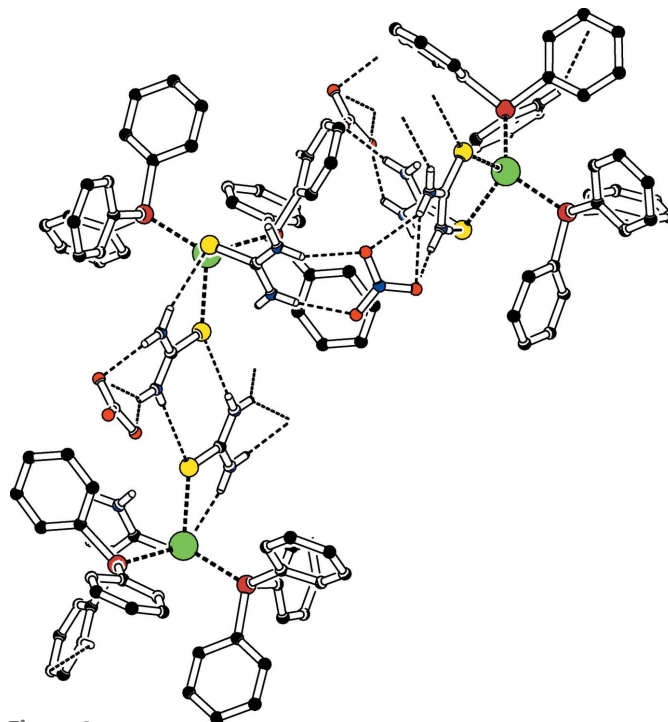
| <i>D</i> –H··· <i>A</i>   | <i>D</i> –H | H··· <i>A</i> | <i>D</i> ··· <i>A</i> | <i>D</i> –H··· <i>A</i> |
|---------------------------|-------------|---------------|-----------------------|-------------------------|
| N1–H1A···O2 <sup>i</sup>  | 0.86        | 2.07          | 2.899 (5)             | 161                     |
| N1–H1B···S2               | 0.86        | 2.57          | 3.417 (4)             | 169                     |
| N2–H2A···O2 <sup>i</sup>  | 0.86        | 2.54          | 3.255 (5)             | 141                     |
| N2–H2A···O3 <sup>i</sup>  | 0.86        | 2.24          | 2.992 (5)             | 147                     |
| N2–H2B···S1 <sup>ii</sup> | 0.86        | 2.66          | 3.453 (4)             | 154                     |
| N3–H3A···O1               | 0.86        | 2.09          | 2.946 (6)             | 171                     |
| N3–H3B···S1               | 0.86        | 2.91          | 3.759 (5)             | 169                     |
| N4–H4A···O3               | 0.86        | 2.26          | 2.976 (6)             | 140                     |
| C2–H2···O1 <sup>iii</sup> | 0.93        | 2.53          | 3.174 (6)             | 126                     |
| C14–H14···S1              | 0.93        | 2.92          | 3.520 (5)             | 124                     |

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

was filtered and the filtrate was left for crystallization. Colorless crystals of (I) were isolated from the filtrate. The crystal structure of the product obtained by adding two equivalents of thiourea has already been reported (Isab *et al.*, 2010).

### 5. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. H atoms were positioned geome-



**Figure 2**  
A partial packing diagram (PLATON; Spek, 2009) illustrating the formation of sheets of molecules with various loops *via* hydrogen-bonding interactions (shown as dashed lines).

**Table 3**  
Experimental details.

|  |  |
|--|--|
| Crystal data   |  |
| Chemical formula   | [Ag(CH <sub>4</sub> N <sub>2</sub> S) <sub>2</sub> (C <sub>18</sub> H <sub>15</sub> P) <sub>2</sub> ] <sub>2</sub> NO <sub>3</sub> |
| <i>M<sub>r</sub></i>   | 846.66   |
| Crystal system, space group  | Monoclinic, <i>P</i> 2 <sub>1</sub> / <i>n</i>   |
| Temperature (K)  | 296  |
| <i>a</i> , <i>b</i> , <i>c</i> (Å)   | 15.0519 (6), 15.1758 (5),<br>17.9186 (8)   |
| $\beta$ (°)  | 107.886 (2)  |
| <i>V</i> (Å <sup>3</sup> )   | 3895.2 (3)   |
| <i>Z</i>   | 4  |
| Radiation type   | Mo <i>K</i> α  |
| $\mu$ (mm <sup>-1</sup> )  | 0.75   |
| Crystal size (mm)  | 0.32 × 0.26 × 0.16   |
| Data collection  |  |
| Diffractometer   | Bruker Kappa APEXII CCD  |
| Absorption correction  | Multi-scan ( <i>SADABS</i> ; Bruker, 2005)   |
| <i>T</i> <sub>min</sub> , <i>T</i> <sub>max</sub>  | 0.798, 0.892   |
| No. of measured, independent and observed [ <i>I</i> > 2σ( <i>I</i> )] reflections                             | 30110, 7659, 3813  |
| <i>R</i> <sub>int</sub>  | 0.100  |
| (sin $\theta/\lambda$ ) <sub>max</sub> (Å <sup>-1</sup> )  | 0.617  |
| Refinement   |  |
| <i>R</i> [ <i>F</i> <sup>2</sup> > 2σ( <i>F</i> <sup>2</sup> )], <i>wR</i> ( <i>F</i> <sup>2</sup> ), <i>S</i> | 0.051, 0.088, 0.98   |
| No. of reflections   | 7659   |
| No. of parameters  | 460  |
| H-atom treatment   | H-atom parameters constrained  |
| $\Delta\rho_{\max}$ , $\Delta\rho_{\min}$ (e Å <sup>-3</sup> )   | 0.47, -0.49  |

Computer programs: *APEX2* and *SAINT* (Bruker, 2007), *SHELXS97* (Sheldrick, 2008), *SHELXL97* (Sheldrick, 2015), *ORTEP-3 for Windows* and *WinGX* (Farrugia, 2012) and *PLATON* (Spek, 2009).

trically (C–H = 0.93, N–H = 0.86 Å) and refined as riding with *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C, N).

## Acknowledgements

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## supporting information

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## Crystal structure of bis(thiourea- $\kappa$ S)bis(triphenylphosphane- $\kappa$ P)silver(I) nitrate

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### Computing details

Data collection: *APEX2* (Bruker, 2007); cell refinement: *S SAINT* (Bruker, 2007); data reduction: *S SAINT* (Bruker, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2015); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 2012) and *PLATON* (Spek, 2009).

### Bis(thiourea- $\kappa$ S)bis(triphenylphosphane- $\kappa$ P)silver(I) nitrate

#### Crystal data

$[\text{Ag}(\text{CH}_4\text{N}_2\text{S})_2(\text{C}_{18}\text{H}_{15}\text{P})_2]\text{NO}_3$

$M_r = 846.66$

Monoclinic,  $P2_1/n$

$a = 15.0519$  (6) Å

$b = 15.1758$  (5) Å

$c = 17.9186$  (8) Å

$\beta = 107.886$  (2)°

$V = 3895.2$  (3) Å<sup>3</sup>

$Z = 4$

$F(000) = 1736$

$D_x = 1.444$  Mg m<sup>-3</sup>

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

Cell parameters from 3813 reflections

$\theta = 2.4\text{--}26.0^\circ$

$\mu = 0.75$  mm<sup>-1</sup>

$T = 296$  K

Plate, colorless

$0.32 \times 0.26 \times 0.16$  mm

#### Data collection

Bruker Kappa APEXII CCD  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 8.00 pixels mm<sup>-1</sup>

$\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Bruker, 2005)

$T_{\min} = 0.798$ ,  $T_{\max} = 0.892$

30110 measured reflections

7659 independent reflections

3813 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.100$

$\theta_{\max} = 26.0^\circ$ ,  $\theta_{\min} = 2.4^\circ$

$h = -18 \rightarrow 18$

$k = -18 \rightarrow 18$

$l = -22 \rightarrow 18$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.051$

$wR(F^2) = 0.088$

$S = 0.98$

7659 reflections

460 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

$w = 1/[\sigma^2(F_o^2) + (0.0192P)^2]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.47$  e Å<sup>-3</sup>

$\Delta\rho_{\min} = -0.49$  e Å<sup>-3</sup>

*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 ( $\text{\AA}^2$ )*

|     | <i>x</i>     | <i>y</i>    | <i>z</i>    | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|-------------|-------------|----------------------------------|
| Ag1 | 0.49184 (3)  | 0.18428 (2) | 0.27125 (2) | 0.04076 (12)                     |
| P1  | 0.41457 (8)  | 0.32439 (7) | 0.29245 (7) | 0.0352 (3)                       |
| P2  | 0.63642 (9)  | 0.18065 (8) | 0.23159 (7) | 0.0387 (3)                       |
| S1  | 0.49005 (11) | 0.08173 (7) | 0.38885 (7) | 0.0603 (5)                       |
| S2  | 0.38377 (10) | 0.09742 (8) | 0.14694 (8) | 0.0520 (4)                       |
| N1  | 0.4763 (3)   | -0.0525 (2) | 0.2923 (2)  | 0.0533 (12)                      |
| H1A | 0.4799       | -0.1075     | 0.2820      | 0.064*                           |
| H1B | 0.4609       | -0.0147     | 0.2547      | 0.064*                           |
| N2  | 0.5177 (3)   | -0.0867 (2) | 0.4206 (2)  | 0.0532 (12)                      |
| H2A | 0.5206       | -0.1412     | 0.4083      | 0.064*                           |
| H2B | 0.5300       | -0.0719     | 0.4691      | 0.064*                           |
| N3  | 0.2637 (3)   | 0.1110 (3)  | 0.2272 (3)  | 0.0677 (14)                      |
| H3A | 0.2084       | 0.1177      | 0.2308      | 0.081*                           |
| H3B | 0.3104       | 0.1049      | 0.2689      | 0.081*                           |
| N4  | 0.2023 (3)   | 0.1198 (3)  | 0.0959 (3)  | 0.0744 (15)                      |
| H4A | 0.1480       | 0.1263      | 0.1017      | 0.089*                           |
| H4B | 0.2082       | 0.1195      | 0.0496      | 0.089*                           |
| C1  | 0.3488 (3)   | 0.3837 (3)  | 0.2050 (3)  | 0.0334 (12)                      |
| C2  | 0.3714 (4)   | 0.4676 (3)  | 0.1868 (3)  | 0.0495 (14)                      |
| H2  | 0.4220       | 0.4968      | 0.2209      | 0.059*                           |
| C3  | 0.3192 (4)   | 0.5089 (3)  | 0.1181 (3)  | 0.0651 (17)                      |
| H3  | 0.3353       | 0.5652      | 0.1065      | 0.078*                           |
| C4  | 0.2445 (4)   | 0.4671 (4)  | 0.0676 (3)  | 0.0652 (17)                      |
| H4  | 0.2096       | 0.4953      | 0.0220      | 0.078*                           |
| C5  | 0.2209 (4)   | 0.3842 (4)  | 0.0839 (3)  | 0.0628 (17)                      |
| H5  | 0.1700       | 0.3557      | 0.0494      | 0.075*                           |
| C6  | 0.2728 (3)   | 0.3430 (3)  | 0.1515 (3)  | 0.0525 (15)                      |
| H6  | 0.2565       | 0.2862      | 0.1619      | 0.063*                           |
| C7  | 0.4982 (3)   | 0.4059 (3)  | 0.3462 (3)  | 0.0362 (12)                      |
| C8  | 0.5859 (4)   | 0.4048 (3)  | 0.3392 (3)  | 0.0705 (18)                      |
| H8  | 0.6027       | 0.3600      | 0.3108      | 0.085*                           |
| C9  | 0.6508 (4)   | 0.4701 (4)  | 0.3740 (4)  | 0.096 (2)                        |
| H9  | 0.7097       | 0.4699      | 0.3675      | 0.116*                           |
| C10 | 0.6269 (4)   | 0.5338 (4)  | 0.4176 (4)  | 0.0743 (19)                      |
| H10 | 0.6706       | 0.5761      | 0.4427      | 0.089*                           |

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|     |            |             |            |             |
|-----|------------|-------------|------------|-------------|
| C11 | 0.5400 (4) | 0.5362 (3)  | 0.4245 (3) | 0.0619 (17) |
| H11 | 0.5239     | 0.5807      | 0.4535     | 0.074*      |
| C12 | 0.4752 (4) | 0.4727 (3)  | 0.3887 (3) | 0.0519 (15) |
| H12 | 0.4154     | 0.4753      | 0.3934     | 0.062*      |
| C13 | 0.3344 (3) | 0.3126 (3)  | 0.3499 (3) | 0.0355 (11) |
| C14 | 0.3658 (4) | 0.2674 (3)  | 0.4199 (3) | 0.0581 (16) |
| H14 | 0.4274     | 0.2482      | 0.4371     | 0.070*      |
| C15 | 0.3088 (5) | 0.2502 (3)  | 0.4649 (3) | 0.0699 (18) |
| H15 | 0.3321     | 0.2196      | 0.5118     | 0.084*      |
| C16 | 0.2181 (4) | 0.2777 (3)  | 0.4412 (3) | 0.0617 (16) |
| H16 | 0.1784     | 0.2646      | 0.4705     | 0.074*      |
| C17 | 0.1874 (4) | 0.3246 (4)  | 0.3739 (4) | 0.090 (2)   |
| H17 | 0.1261     | 0.3449      | 0.3579     | 0.108*      |
| C18 | 0.2442 (4) | 0.3433 (4)  | 0.3282 (3) | 0.0687 (18) |
| H18 | 0.2214     | 0.3767      | 0.2828     | 0.082*      |
| C19 | 0.7498 (3) | 0.1714 (3)  | 0.3064 (3) | 0.0412 (13) |
| C20 | 0.7566 (4) | 0.1962 (3)  | 0.3825 (3) | 0.0548 (15) |
| H20 | 0.7029     | 0.2116      | 0.3948     | 0.066*      |
| C21 | 0.8416 (5) | 0.1982 (4)  | 0.4402 (4) | 0.0734 (19) |
| H21 | 0.8455     | 0.2146      | 0.4911     | 0.088*      |
| C22 | 0.9209 (5) | 0.1758 (4)  | 0.4214 (4) | 0.082 (2)   |
| H22 | 0.9788     | 0.1788      | 0.4597     | 0.098*      |
| C23 | 0.9158 (4) | 0.1494 (4)  | 0.3484 (4) | 0.080 (2)   |
| H23 | 0.9699     | 0.1335      | 0.3369     | 0.096*      |
| C24 | 0.8308 (4) | 0.1458 (3)  | 0.2907 (3) | 0.0594 (16) |
| H24 | 0.8275     | 0.1261      | 0.2408     | 0.071*      |
| C25 | 0.6492 (4) | 0.2776 (3)  | 0.1747 (3) | 0.0388 (13) |
| C26 | 0.5705 (4) | 0.3294 (3)  | 0.1437 (3) | 0.0503 (14) |
| H26 | 0.5141     | 0.3125      | 0.1505     | 0.060*      |
| C27 | 0.5749 (4) | 0.4057 (3)  | 0.1031 (3) | 0.0648 (17) |
| H27 | 0.5216     | 0.4396      | 0.0824     | 0.078*      |
| C28 | 0.6582 (5) | 0.4316 (3)  | 0.0931 (3) | 0.0632 (18) |
| H28 | 0.6616     | 0.4837      | 0.0668     | 0.076*      |
| C29 | 0.7358 (4) | 0.3803 (4)  | 0.1219 (3) | 0.0628 (17) |
| H29 | 0.7917     | 0.3966      | 0.1138     | 0.075*      |
| C30 | 0.7314 (3) | 0.3043 (3)  | 0.1632 (3) | 0.0500 (14) |
| H30 | 0.7851     | 0.2706      | 0.1836     | 0.060*      |
| C31 | 0.6328 (3) | 0.0869 (3)  | 0.1666 (3) | 0.0395 (13) |
| C32 | 0.6316 (4) | 0.0032 (3)  | 0.1975 (3) | 0.0581 (16) |
| H32 | 0.6392     | -0.0035     | 0.2507     | 0.070*      |
| C33 | 0.6189 (4) | -0.0706 (3) | 0.1495 (4) | 0.0638 (17) |
| H33 | 0.6167     | -0.1266     | 0.1699     | 0.077*      |
| C34 | 0.6098 (4) | -0.0596 (4) | 0.0717 (4) | 0.0653 (18) |
| H34 | 0.6014     | -0.1087     | 0.0392     | 0.078*      |
| C35 | 0.6127 (4) | 0.0224 (4)  | 0.0410 (3) | 0.0636 (17) |
| H35 | 0.6077     | 0.0288      | -0.0118    | 0.076*      |
| C36 | 0.6233 (3) | 0.0960 (3)  | 0.0886 (3) | 0.0486 (14) |
| H36 | 0.6240     | 0.1519      | 0.0674     | 0.058*      |

|     |             |             |            |             |
|-----|-------------|-------------|------------|-------------|
| C37 | 0.4944 (3)  | -0.0262 (3) | 0.3655 (3) | 0.0411 (13) |
| C38 | 0.2767 (4)  | 0.1101 (3)  | 0.1581 (4) | 0.0501 (15) |
| N5  | 0.0359 (4)  | 0.2164 (3)  | 0.1905 (3) | 0.0634 (15) |
| O1  | 0.0756 (3)  | 0.1552 (2)  | 0.2336 (2) | 0.0767 (13) |
| O2  | -0.0238 (3) | 0.2622 (2)  | 0.2067 (3) | 0.0906 (15) |
| O3  | 0.0575 (3)  | 0.2328 (2)  | 0.1292 (3) | 0.0802 (13) |

*Atomic displacement parameters (Å<sup>2</sup>)*

|     | $U^{11}$    | $U^{22}$     | $U^{33}$   | $U^{12}$    | $U^{13}$   | $U^{23}$    |
|-----|-------------|--------------|------------|-------------|------------|-------------|
| Ag1 | 0.0430 (2)  | 0.03710 (19) | 0.0476 (2) | 0.0009 (2)  | 0.0218 (2) | -0.0015 (2) |
| P1  | 0.0355 (8)  | 0.0314 (7)   | 0.0395 (8) | 0.0020 (6)  | 0.0128 (7) | 0.0002 (6)  |
| P2  | 0.0383 (8)  | 0.0434 (7)   | 0.0382 (8) | 0.0034 (7)  | 0.0173 (7) | 0.0006 (7)  |
| S1  | 0.1117 (14) | 0.0337 (7)   | 0.0409 (8) | 0.0012 (8)  | 0.0313 (9) | 0.0007 (6)  |
| S2  | 0.0517 (10) | 0.0620 (9)   | 0.0426 (9) | -0.0022 (7) | 0.0148 (8) | -0.0110 (7) |
| N1  | 0.089 (4)   | 0.037 (2)    | 0.041 (3)  | -0.001 (2)  | 0.030 (3)  | 0.000 (2)   |
| N2  | 0.083 (4)   | 0.036 (2)    | 0.038 (3)  | 0.007 (2)   | 0.013 (3)  | 0.000 (2)   |
| N3  | 0.056 (3)   | 0.078 (3)    | 0.077 (4)  | -0.002 (2)  | 0.032 (3)  | -0.007 (3)  |
| N4  | 0.045 (3)   | 0.087 (3)    | 0.079 (4)  | -0.007 (3)  | 0.001 (3)  | -0.003 (3)  |
| C1  | 0.037 (3)   | 0.036 (3)    | 0.032 (3)  | 0.005 (2)   | 0.018 (3)  | -0.001 (2)  |
| C2  | 0.058 (4)   | 0.045 (3)    | 0.044 (3)  | -0.002 (3)  | 0.014 (3)  | 0.005 (3)   |
| C3  | 0.072 (5)   | 0.057 (4)    | 0.067 (4)  | -0.004 (3)  | 0.021 (4)  | 0.020 (3)   |
| C4  | 0.069 (5)   | 0.079 (4)    | 0.043 (4)  | 0.016 (4)   | 0.011 (4)  | 0.018 (3)   |
| C5  | 0.048 (4)   | 0.071 (4)    | 0.057 (4)  | -0.007 (3)  | -0.004 (3) | -0.001 (3)  |
| C6  | 0.050 (4)   | 0.051 (3)    | 0.049 (4)  | -0.001 (3)  | 0.004 (3)  | 0.007 (3)   |
| C7  | 0.037 (3)   | 0.035 (3)    | 0.034 (3)  | -0.001 (2)  | 0.008 (3)  | -0.001 (2)  |
| C8  | 0.050 (4)   | 0.075 (4)    | 0.095 (5)  | -0.014 (3)  | 0.034 (4)  | -0.042 (4)  |
| C9  | 0.049 (4)   | 0.122 (6)    | 0.128 (6)  | -0.028 (4)  | 0.042 (5)  | -0.055 (5)  |
| C10 | 0.065 (5)   | 0.078 (4)    | 0.081 (5)  | -0.028 (4)  | 0.023 (4)  | -0.028 (4)  |
| C11 | 0.073 (5)   | 0.047 (3)    | 0.060 (4)  | 0.001 (3)   | 0.012 (4)  | -0.015 (3)  |
| C12 | 0.051 (4)   | 0.040 (3)    | 0.061 (4)  | -0.006 (3)  | 0.012 (3)  | -0.012 (3)  |
| C13 | 0.038 (3)   | 0.033 (2)    | 0.038 (3)  | 0.003 (2)   | 0.017 (3)  | -0.001 (2)  |
| C14 | 0.060 (4)   | 0.063 (3)    | 0.061 (4)  | 0.021 (3)   | 0.032 (4)  | 0.012 (3)   |
| C15 | 0.098 (6)   | 0.063 (4)    | 0.066 (4)  | 0.023 (4)   | 0.050 (5)  | 0.024 (3)   |
| C16 | 0.068 (5)   | 0.071 (4)    | 0.062 (4)  | -0.004 (3)  | 0.042 (4)  | -0.006 (3)  |
| C17 | 0.048 (4)   | 0.159 (6)    | 0.068 (5)  | 0.022 (5)   | 0.026 (4)  | 0.015 (5)   |
| C18 | 0.051 (4)   | 0.109 (5)    | 0.052 (4)  | 0.020 (4)   | 0.025 (4)  | 0.027 (3)   |
| C19 | 0.040 (3)   | 0.041 (3)    | 0.043 (3)  | 0.001 (3)   | 0.013 (3)  | 0.007 (2)   |
| C20 | 0.056 (4)   | 0.063 (4)    | 0.044 (4)  | 0.001 (3)   | 0.014 (3)  | 0.004 (3)   |
| C21 | 0.086 (5)   | 0.074 (4)    | 0.053 (4)  | -0.016 (4)  | 0.011 (4)  | -0.001 (3)  |
| C22 | 0.065 (5)   | 0.085 (5)    | 0.071 (5)  | -0.020 (4)  | -0.016 (5) | 0.032 (4)   |
| C23 | 0.048 (5)   | 0.095 (5)    | 0.093 (6)  | 0.016 (4)   | 0.015 (5)  | 0.040 (4)   |
| C24 | 0.048 (4)   | 0.075 (4)    | 0.058 (4)  | 0.011 (3)   | 0.019 (4)  | 0.013 (3)   |
| C25 | 0.036 (3)   | 0.049 (3)    | 0.031 (3)  | 0.004 (3)   | 0.010 (3)  | -0.001 (2)  |
| C26 | 0.052 (4)   | 0.061 (3)    | 0.045 (3)  | 0.002 (3)   | 0.026 (3)  | 0.004 (3)   |
| C27 | 0.072 (5)   | 0.064 (4)    | 0.062 (4)  | 0.022 (3)   | 0.025 (4)  | 0.025 (3)   |
| C28 | 0.097 (6)   | 0.044 (3)    | 0.055 (4)  | -0.013 (4)  | 0.033 (4)  | -0.001 (3)  |
| C29 | 0.074 (5)   | 0.058 (4)    | 0.070 (4)  | -0.019 (3)  | 0.043 (4)  | 0.000 (3)   |

|     |           |           |           |            |           |            |
|-----|-----------|-----------|-----------|------------|-----------|------------|
| C30 | 0.040 (3) | 0.060 (4) | 0.057 (4) | 0.007 (3)  | 0.025 (3) | 0.008 (3)  |
| C31 | 0.037 (3) | 0.048 (3) | 0.040 (3) | 0.002 (2)  | 0.022 (3) | 0.000 (3)  |
| C32 | 0.074 (5) | 0.050 (3) | 0.058 (4) | 0.004 (3)  | 0.031 (4) | 0.007 (3)  |
| C33 | 0.071 (5) | 0.040 (3) | 0.085 (5) | 0.003 (3)  | 0.030 (4) | -0.003 (3) |
| C34 | 0.059 (4) | 0.059 (4) | 0.074 (5) | 0.002 (3)  | 0.016 (4) | -0.021 (4) |
| C35 | 0.068 (5) | 0.074 (4) | 0.047 (4) | 0.014 (3)  | 0.015 (3) | -0.008 (3) |
| C36 | 0.046 (4) | 0.050 (3) | 0.050 (4) | 0.008 (3)  | 0.015 (3) | -0.002 (3) |
| C37 | 0.050 (4) | 0.036 (3) | 0.039 (3) | -0.002 (2) | 0.016 (3) | 0.008 (3)  |
| C38 | 0.049 (4) | 0.039 (3) | 0.061 (4) | -0.001 (3) | 0.015 (4) | -0.007 (3) |
| N5  | 0.059 (4) | 0.033 (3) | 0.106 (5) | -0.008 (3) | 0.036 (4) | -0.002 (3) |
| O1  | 0.079 (3) | 0.051 (2) | 0.103 (3) | 0.016 (2)  | 0.032 (3) | 0.020 (2)  |
| O2  | 0.079 (3) | 0.051 (2) | 0.164 (4) | 0.017 (2)  | 0.070 (3) | 0.019 (3)  |
| O3  | 0.084 (3) | 0.060 (2) | 0.109 (4) | 0.007 (2)  | 0.047 (3) | 0.020 (2)  |

*Geometric parameters (Å, °)*

|        |             |         |           |
|--------|-------------|---------|-----------|
| Ag1—P2 | 2.4888 (13) | C13—C14 | 1.379 (6) |
| Ag1—P1 | 2.5078 (12) | C14—C15 | 1.371 (7) |
| Ag1—S1 | 2.6263 (13) | C14—H14 | 0.9300    |
| Ag1—S2 | 2.6683 (13) | C15—C16 | 1.365 (7) |
| P1—C1  | 1.815 (4)   | C15—H15 | 0.9300    |
| P1—C7  | 1.816 (4)   | C16—C17 | 1.353 (7) |
| P1—C13 | 1.821 (5)   | C16—H16 | 0.9300    |
| P2—C19 | 1.823 (5)   | C17—C18 | 1.382 (7) |
| P2—C31 | 1.828 (5)   | C17—H17 | 0.9300    |
| P2—C25 | 1.834 (5)   | C18—H18 | 0.9300    |
| S1—C37 | 1.698 (4)   | C19—C20 | 1.387 (6) |
| S2—C38 | 1.695 (6)   | C19—C24 | 1.388 (6) |
| N1—C37 | 1.317 (5)   | C20—C21 | 1.376 (6) |
| N1—H1A | 0.8600      | C20—H20 | 0.9300    |
| N1—H1B | 0.8600      | C21—C22 | 1.378 (8) |
| N2—C37 | 1.315 (5)   | C21—H21 | 0.9300    |
| N2—H2A | 0.8600      | C22—C23 | 1.348 (8) |
| N2—H2B | 0.8600      | C22—H22 | 0.9300    |
| N3—C38 | 1.312 (6)   | C23—C24 | 1.377 (7) |
| N3—H3A | 0.8600      | C23—H23 | 0.9300    |
| N3—H3B | 0.8600      | C24—H24 | 0.9300    |
| N4—C38 | 1.324 (6)   | C25—C30 | 1.376 (6) |
| N4—H4A | 0.8600      | C25—C26 | 1.387 (6) |
| N4—H4B | 0.8600      | C26—C27 | 1.380 (6) |
| C1—C2  | 1.382 (5)   | C26—H26 | 0.9300    |
| C1—C6  | 1.391 (6)   | C27—C28 | 1.377 (7) |
| C2—C3  | 1.389 (6)   | C27—H27 | 0.9300    |
| C2—H2  | 0.9300      | C28—C29 | 1.366 (7) |
| C3—C4  | 1.363 (6)   | C28—H28 | 0.9300    |
| C3—H3  | 0.9300      | C29—C30 | 1.382 (6) |
| C4—C5  | 1.363 (6)   | C29—H29 | 0.9300    |
| C4—H4  | 0.9300      | C30—H30 | 0.9300    |



|            |             |             |           |
|------------|-------------|-------------|-----------|
| C5—C6      | 1.374 (6)   | C31—C36     | 1.369 (6) |
| C5—H5      | 0.9300      | C31—C32     | 1.387 (6) |
| C6—H6      | 0.9300      | C32—C33     | 1.390 (6) |
| C7—C8      | 1.364 (6)   | C32—H32     | 0.9300    |
| C7—C12     | 1.375 (6)   | C33—C34     | 1.368 (7) |
| C8—C9      | 1.396 (7)   | C33—H33     | 0.9300    |
| C8—H8      | 0.9300      | C34—C35     | 1.367 (7) |
| C9—C10     | 1.359 (7)   | C34—H34     | 0.9300    |
| C9—H9      | 0.9300      | C35—C36     | 1.384 (6) |
| C10—C11    | 1.352 (7)   | C35—H35     | 0.9300    |
| C10—H10    | 0.9300      | C36—H36     | 0.9300    |
| C11—C12    | 1.381 (6)   | N5—O1       | 1.237 (5) |
| C11—H11    | 0.9300      | N5—O2       | 1.241 (5) |
| C12—H12    | 0.9300      | N5—O3       | 1.261 (5) |
| C13—C18    | 1.373 (6)   |             |           |
| P2—Ag1—P1  | 123.29 (4)  | C16—C15—H15 | 119.8     |
| P2—Ag1—S1  | 116.19 (5)  | C14—C15—H15 | 119.8     |
| P1—Ag1—S1  | 105.15 (4)  | C17—C16—C15 | 118.2 (6) |
| P2—Ag1—S2  | 96.49 (4)   | C17—C16—H16 | 120.9     |
| P1—Ag1—S2  | 110.67 (4)  | C15—C16—H16 | 120.9     |
| S1—Ag1—S2  | 102.90 (4)  | C16—C17—C18 | 122.1 (6) |
| C1—P1—C7   | 103.0 (2)   | C16—C17—H17 | 118.9     |
| C1—P1—C13  | 104.9 (2)   | C18—C17—H17 | 118.9     |
| C7—P1—C13  | 103.8 (2)   | C13—C18—C17 | 120.0 (5) |
| C1—P1—Ag1  | 116.41 (14) | C13—C18—H18 | 120.0     |
| C7—P1—Ag1  | 112.30 (16) | C17—C18—H18 | 120.0     |
| C13—P1—Ag1 | 114.92 (14) | C20—C19—C24 | 118.0 (5) |
| C19—P2—C31 | 104.3 (2)   | C20—C19—P2  | 118.2 (4) |
| C19—P2—C25 | 103.7 (2)   | C24—C19—P2  | 123.7 (4) |
| C31—P2—C25 | 105.0 (2)   | C21—C20—C19 | 121.0 (5) |
| C19—P2—Ag1 | 119.64 (17) | C21—C20—H20 | 119.5     |
| C31—P2—Ag1 | 109.93 (16) | C19—C20—H20 | 119.5     |
| C25—P2—Ag1 | 113.04 (17) | C20—C21—C22 | 119.1 (6) |
| C37—S1—Ag1 | 111.29 (18) | C20—C21—H21 | 120.4     |
| C38—S2—Ag1 | 101.75 (19) | C22—C21—H21 | 120.4     |
| C37—N1—H1A | 120.0       | C23—C22—C21 | 121.0 (6) |
| C37—N1—H1B | 120.0       | C23—C22—H22 | 119.5     |
| H1A—N1—H1B | 120.0       | C21—C22—H22 | 119.5     |
| C37—N2—H2A | 120.0       | C22—C23—C24 | 120.1 (7) |
| C37—N2—H2B | 120.0       | C22—C23—H23 | 119.9     |
| H2A—N2—H2B | 120.0       | C24—C23—H23 | 119.9     |
| C38—N3—H3A | 120.0       | C23—C24—C19 | 120.6 (5) |
| C38—N3—H3B | 120.0       | C23—C24—H24 | 119.7     |
| H3A—N3—H3B | 120.0       | C19—C24—H24 | 119.7     |
| C38—N4—H4A | 120.0       | C30—C25—C26 | 118.1 (4) |
| C38—N4—H4B | 120.0       | C30—C25—P2  | 124.9 (4) |
| H4A—N4—H4B | 120.0       | C26—C25—P2  | 117.0 (4) |

|              |            |                |           |
|--------------|------------|----------------|-----------|
| C2—C1—C6     | 117.2 (4)  | C27—C26—C25    | 120.8 (5) |
| C2—C1—P1     | 123.5 (4)  | C27—C26—H26    | 119.6     |
| C6—C1—P1     | 119.3 (3)  | C25—C26—H26    | 119.6     |
| C1—C2—C3     | 120.7 (5)  | C28—C27—C26    | 120.1 (5) |
| C1—C2—H2     | 119.6      | C28—C27—H27    | 120.0     |
| C3—C2—H2     | 119.6      | C26—C27—H27    | 120.0     |
| C4—C3—C2     | 120.3 (5)  | C29—C28—C27    | 119.6 (5) |
| C4—C3—H3     | 119.8      | C29—C28—H28    | 120.2     |
| C2—C3—H3     | 119.8      | C27—C28—H28    | 120.2     |
| C3—C4—C5     | 120.2 (5)  | C28—C29—C30    | 120.2 (5) |
| C3—C4—H4     | 119.9      | C28—C29—H29    | 119.9     |
| C5—C4—H4     | 119.9      | C30—C29—H29    | 119.9     |
| C4—C5—C6     | 119.5 (5)  | C25—C30—C29    | 121.2 (5) |
| C4—C5—H5     | 120.2      | C25—C30—H30    | 119.4     |
| C6—C5—H5     | 120.2      | C29—C30—H30    | 119.4     |
| C5—C6—C1     | 122.0 (5)  | C36—C31—C32    | 119.4 (4) |
| C5—C6—H6     | 119.0      | C36—C31—P2     | 123.0 (4) |
| C1—C6—H6     | 119.0      | C32—C31—P2     | 117.4 (4) |
| C8—C7—C12    | 118.4 (4)  | C31—C32—C33    | 120.4 (5) |
| C8—C7—P1     | 118.4 (4)  | C31—C32—H32    | 119.8     |
| C12—C7—P1    | 123.0 (4)  | C33—C32—H32    | 119.8     |
| C7—C8—C9     | 121.0 (5)  | C34—C33—C32    | 119.0 (5) |
| C7—C8—H8     | 119.5      | C34—C33—H33    | 120.5     |
| C9—C8—H8     | 119.5      | C32—C33—H33    | 120.5     |
| C10—C9—C8    | 119.3 (6)  | C35—C34—C33    | 121.0 (5) |
| C10—C9—H9    | 120.3      | C35—C34—H34    | 119.5     |
| C8—C9—H9     | 120.3      | C33—C34—H34    | 119.5     |
| C11—C10—C9   | 120.4 (6)  | C34—C35—C36    | 119.9 (5) |
| C11—C10—H10  | 119.8      | C34—C35—H35    | 120.0     |
| C9—C10—H10   | 119.8      | C36—C35—H35    | 120.0     |
| C10—C11—C12  | 120.3 (5)  | C31—C36—C35    | 120.3 (5) |
| C10—C11—H11  | 119.8      | C31—C36—H36    | 119.9     |
| C12—C11—H11  | 119.8      | C35—C36—H36    | 119.9     |
| C7—C12—C11   | 120.6 (5)  | N2—C37—N1      | 117.6 (4) |
| C7—C12—H12   | 119.7      | N2—C37—S1      | 120.7 (4) |
| C11—C12—H12  | 119.7      | N1—C37—S1      | 121.8 (4) |
| C18—C13—C14  | 117.3 (5)  | N3—C38—N4      | 117.5 (6) |
| C18—C13—P1   | 125.1 (4)  | N3—C38—S2      | 122.4 (5) |
| C14—C13—P1   | 117.5 (4)  | N4—C38—S2      | 120.1 (5) |
| C15—C14—C13  | 121.9 (5)  | O1—N5—O2       | 121.3 (6) |
| C15—C14—H14  | 119.1      | O1—N5—O3       | 119.3 (5) |
| C13—C14—H14  | 119.1      | O2—N5—O3       | 119.4 (5) |
| C16—C15—C14  | 120.4 (5)  |                |           |
| C7—P1—C1—C2  | 6.2 (5)    | Ag1—P2—C19—C20 | -21.5 (4) |
| C13—P1—C1—C2 | 114.6 (4)  | C31—P2—C19—C24 | 39.0 (5)  |
| Ag1—P1—C1—C2 | -117.2 (4) | C25—P2—C19—C24 | -70.7 (4) |
| C7—P1—C1—C6  | -175.4 (4) | Ag1—P2—C19—C24 | 162.3 (3) |

|                 |            |                 |            |
|-----------------|------------|-----------------|------------|
| C13—P1—C1—C6    | -67.0 (4)  | C24—C19—C20—C21 | 2.2 (7)    |
| Ag1—P1—C1—C6    | 61.2 (4)   | P2—C19—C20—C21  | -174.2 (4) |
| C6—C1—C2—C3     | 0.4 (7)    | C19—C20—C21—C22 | 0.4 (8)    |
| P1—C1—C2—C3     | 178.8 (4)  | C20—C21—C22—C23 | -2.1 (9)   |
| C1—C2—C3—C4     | 0.3 (8)    | C21—C22—C23—C24 | 1.0 (9)    |
| C2—C3—C4—C5     | -0.6 (9)   | C22—C23—C24—C19 | 1.7 (8)    |
| C3—C4—C5—C6     | 0.1 (9)    | C20—C19—C24—C23 | -3.2 (7)   |
| C4—C5—C6—C1     | 0.6 (8)    | P2—C19—C24—C23  | 172.9 (4)  |
| C2—C1—C6—C5     | -0.8 (7)   | C19—P2—C25—C30  | 30.9 (5)   |
| P1—C1—C6—C5     | -179.3 (4) | C31—P2—C25—C30  | -78.3 (5)  |
| C1—P1—C7—C8     | -97.0 (4)  | Ag1—P2—C25—C30  | 161.9 (4)  |
| C13—P1—C7—C8    | 153.8 (4)  | C19—P2—C25—C26  | -146.5 (4) |
| Ag1—P1—C7—C8    | 29.0 (5)   | C31—P2—C25—C26  | 104.4 (4)  |
| C1—P1—C7—C12    | 77.7 (4)   | Ag1—P2—C25—C26  | -15.4 (4)  |
| C13—P1—C7—C12   | -31.5 (4)  | C30—C25—C26—C27 | -0.3 (7)   |
| Ag1—P1—C7—C12   | -156.2 (4) | P2—C25—C26—C27  | 177.2 (4)  |
| C12—C7—C8—C9    | -0.2 (9)   | C25—C26—C27—C28 | -0.3 (8)   |
| P1—C7—C8—C9     | 174.8 (5)  | C26—C27—C28—C29 | 1.6 (9)    |
| C7—C8—C9—C10    | 2.1 (10)   | C27—C28—C29—C30 | -2.1 (9)   |
| C8—C9—C10—C11   | -2.7 (10)  | C26—C25—C30—C29 | -0.2 (7)   |
| C9—C10—C11—C12  | 1.3 (10)   | P2—C25—C30—C29  | -177.5 (4) |
| C8—C7—C12—C11   | -1.2 (8)   | C28—C29—C30—C25 | 1.4 (8)    |
| P1—C7—C12—C11   | -175.9 (4) | C19—P2—C31—C36  | -119.9 (5) |
| C10—C11—C12—C7  | 0.6 (8)    | C25—P2—C31—C36  | -11.2 (5)  |
| C1—P1—C13—C18   | 0.2 (5)    | Ag1—P2—C31—C36  | 110.7 (4)  |
| C7—P1—C13—C18   | 108.0 (5)  | C19—P2—C31—C32  | 65.8 (4)   |
| Ag1—P1—C13—C18  | -129.0 (4) | C25—P2—C31—C32  | 174.5 (4)  |
| C1—P1—C13—C14   | 178.6 (4)  | Ag1—P2—C31—C32  | -63.7 (4)  |
| C7—P1—C13—C14   | -73.6 (4)  | C36—C31—C32—C33 | -1.3 (8)   |
| Ag1—P1—C13—C14  | 49.4 (4)   | P2—C31—C32—C33  | 173.2 (4)  |
| C18—C13—C14—C15 | 2.8 (8)    | C31—C32—C33—C34 | 1.5 (9)    |
| P1—C13—C14—C15  | -175.8 (4) | C32—C33—C34—C35 | -0.1 (9)   |
| C13—C14—C15—C16 | 0.0 (8)    | C33—C34—C35—C36 | -1.3 (9)   |
| C14—C15—C16—C17 | -2.2 (9)   | C32—C31—C36—C35 | -0.1 (8)   |
| C15—C16—C17—C18 | 1.5 (9)    | P2—C31—C36—C35  | -174.3 (4) |
| C14—C13—C18—C17 | -3.3 (8)   | C34—C35—C36—C31 | 1.4 (8)    |
| P1—C13—C18—C17  | 175.1 (4)  | Ag1—S1—C37—N2   | 162.8 (4)  |
| C16—C17—C18—C13 | 1.3 (10)   | Ag1—S1—C37—N1   | -16.2 (5)  |
| C31—P2—C19—C20  | -144.9 (4) | Ag1—S2—C38—N3   | -35.5 (4)  |
| C25—P2—C19—C20  | 105.5 (4)  | Ag1—S2—C38—N4   | 144.3 (4)  |

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

| $D-H\cdots A$                   | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------------|-------|-------------|-------------|---------------|
| N1—H1A $\cdots$ O2 <sup>i</sup> | 0.86  | 2.07        | 2.899 (5)   | 161           |
| N1—H1B $\cdots$ S2              | 0.86  | 2.57        | 3.417 (4)   | 169           |
| N2—H2A $\cdots$ O2 <sup>i</sup> | 0.86  | 2.54        | 3.255 (5)   | 141           |
| N2—H2A $\cdots$ O3 <sup>i</sup> | 0.86  | 2.24        | 2.992 (5)   | 147           |

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|                           |      |      |           |     |
|---------------------------|------|------|-----------|-----|
| N2—H2B···S1 <sup>ii</sup> | 0.86 | 2.66 | 3.453 (4) | 154 |
| N3—H3A···O1               | 0.86 | 2.09 | 2.946 (6) | 171 |
| N3—H3B···S1               | 0.86 | 2.91 | 3.759 (5) | 169 |
| N4—H4A···O3               | 0.86 | 2.26 | 2.976 (6) | 140 |
| C2—H2···O1 <sup>iii</sup> | 0.93 | 2.53 | 3.174 (6) | 126 |
| C14—H14···S1              | 0.93 | 2.92 | 3.520 (5) | 124 |

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