

Crystal structure of (benzenecarbothioamide- κ S)chloridobis(triphenylphosphane- κ P)silver(I)

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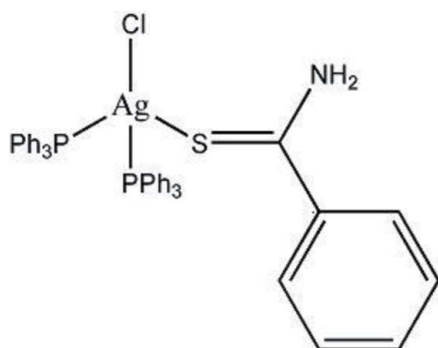
In the mononuclear mixed-ligand title complex, $[\text{AgCl}(\text{C}_7\text{H}_7\text{NS})(\text{C}_{18}\text{H}_{15}\text{P})_2]$, the Ag^{I} ion is four coordinated by one S atom of a benzenecarbothioamide ligand, two P atoms of two triphenylphosphane ligands and one chloride ion, displaying a distorted tetrahedral coordination geometry. In the crystal, pairs of $\text{N}-\text{H}\cdots\text{Cl}$ hydrogen bonds form inversion dimers. An intramolecular $\text{N}-\text{H}\cdots\text{Cl}$ hydrogen bond is also observed.

Keywords: crystal structure; Ag^{I} complex; benzenecarbothioamide; hydrogen bonds.

CCDC reference: 1012936

1. Related literature

For relevant examples of structures of Ag^{I} complexes, see: Aslanidis *et al.* (1997); McFarlane *et al.* (1998); Cox *et al.* (2000); Dennehy *et al.* (2007); Nimthong *et al.* (2008). For potential applications of related complexes, see: Isab *et al.* (2010); Nawaz *et al.* (2011).



2. Experimental

2.1. Crystal data

$[\text{AgCl}(\text{C}_7\text{H}_7\text{NS})(\text{C}_{18}\text{H}_{15}\text{P})_2]$
 $M_r = 805.05$
 Monoclinic, $P2_1/n$
 $a = 14.2405$ (7) Å
 $b = 13.9271$ (7) Å
 $c = 19.0774$ (10) Å
 $\beta = 98.711$ (1)°

$V = 3740.0$ (3) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.78$ mm⁻¹
 $T = 293$ K
 $0.30 \times 0.16 \times 0.10$ mm

2.2. Data collection

Bruker SMART APEX CCD diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 2003)
 $T_{\text{min}} = 0.861$, $T_{\text{max}} = 0.923$

51284 measured reflections
 9308 independent reflections
 8544 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.031$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.033$
 $wR(F^2) = 0.075$
 $S = 1.13$
 9308 reflections

442 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.72$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.39$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1A}\cdots\text{Cl1}^{\text{i}}$	0.86	2.43	3.1798 (17)	146
$\text{N1}-\text{H1B}\cdots\text{Cl1}$	0.86	2.39	3.2434 (17)	173

Symmetry code: (i) $-x + 2, -y, -z$.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 2003); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL2012 (Sheldrick, 2008), SHELXLE (Hübschle *et al.*, 2011); molecular graphics: Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: SHELXL97 and publCIF (Westrip, 2010).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: LH5718).

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supporting information

Acta Cryst. (2014). E70, m310–m311 [doi:10.1107/S1600536814015992]

Crystal structure of (benzenecarbothioamide- κ S)chloridobis(triphenylphosphane- κ P)silver(I)

Wattana Ruangwut and Chaveng Pakawatchai

S1. Experimental

Triphenylphosphine, (0.37 g, 0.14 mmol) was dissolved in 30 cm³ of acetonitrile at 343 K. AgCl (0.10 g, 0.70 mmol) was added and then the mixture was stirred for 3 hours. At that time, a white precipitate was deposited and then benzenecarbothioamide (0.09 g, 0.70 mmol) was added. The new reaction mixture was continually heated under reflux for 5 hours. The resulting clear solution was filtered and the filtrate was kept to evaporate at room temperature. X-ray quality crystals were obtained after two days and were filtered and dried *in vacuo*.

S1.1. Refinement

All H atoms were positioned geometrically and refined using a riding-model, approximation with C—H = 0.93 with $U_{iso}(\text{H}) = 1.2 U_{eq}(\text{C})$, N—H = 0.86 with $U_{iso}(\text{H}) = 1.2 U_{eq}(\text{N})$.

S2. Results and discussion

The coordination compounds of Ag^I complexes with phosphine and thione ligands have gained interest in recent years because of their participation in biological process as antibacterial agents (Isab *et al.*, 2010; Nawaz *et al.*, 2011). A large number of crystal structures of Ag^I complexes containing phosphine and thione as ligands have presented that these complexes are either mononuclear (McFarlane *et al.*, 1998) or dinuclear (Cox *et al.*, 2000) complexes with the silver(I) center exhibiting distorted tetrahedral or trigonal coordination geometry (Aslanidis *et al.*, 1997).

The molecular structure of the title mononuclear complex displays a distorted tetrahedral coordination geometry (Fig. 1). The deviation from ideal tetrahedral geometry can be described by the P2—Ag1—P1 angle which has value of 120.05 (2)° and this value is much larger than the perfect tetrahedral value of 109.5°. This distortion may be due to the steric imposition induced by the two bulky triphenylphosphine ligands. This has also been observed in previous reports e.g. the P2—Cu1—P1 angle in [CuI(C₇H₈N₂S)(C₁₈H₁₅P)₂] is 118.63 (5)° (Nimthong *et al.*, 2008). The Ag1—S1 bond distance in the title complex 2.5880 (5) Å lies in the range of other Ag^I complexes such as the Ag—S bond in [Ag(C₇H₄NS₂O₂)(C₁₈H₁₅P)₃] of 2.5939 (10) Å (Dennehy *et al.*, 2007). In the crystal, pairs of N—H⋯Cl hydrogen bonds form inversion dimers. An intramolecular N—H⋯Cl hydrogen bond is also observed (Fig. 2).

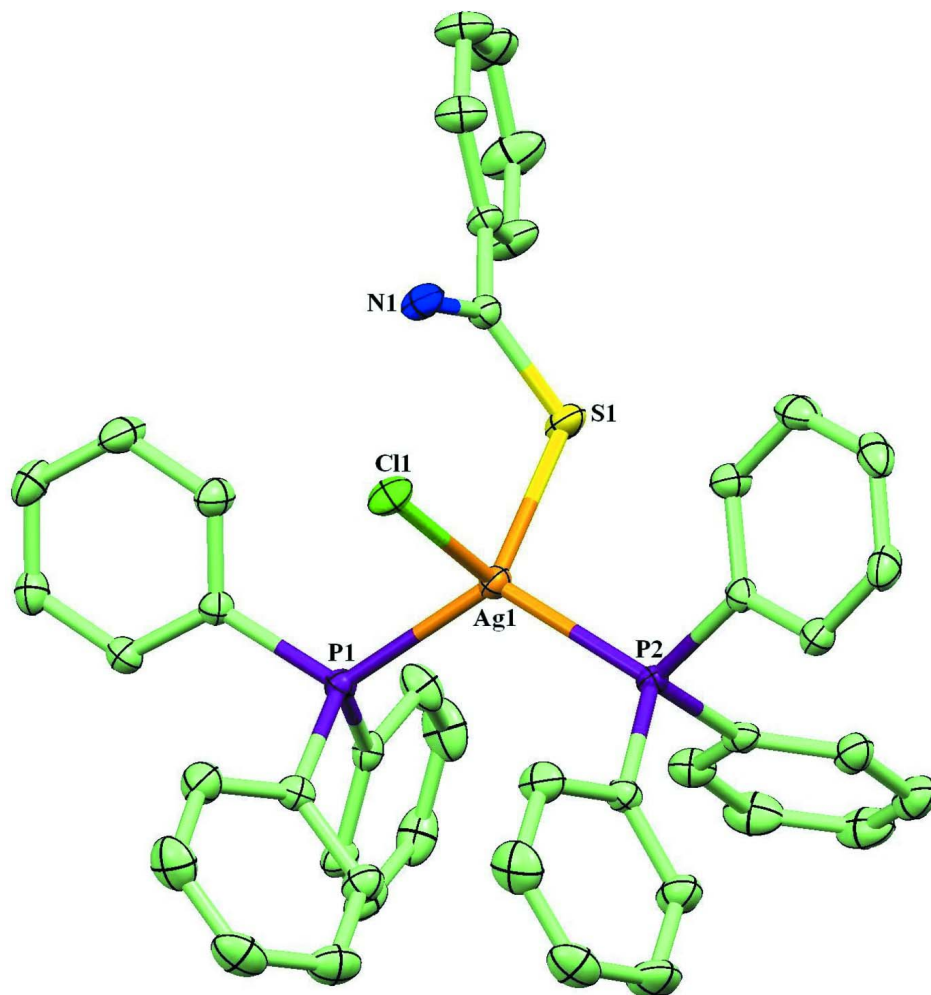
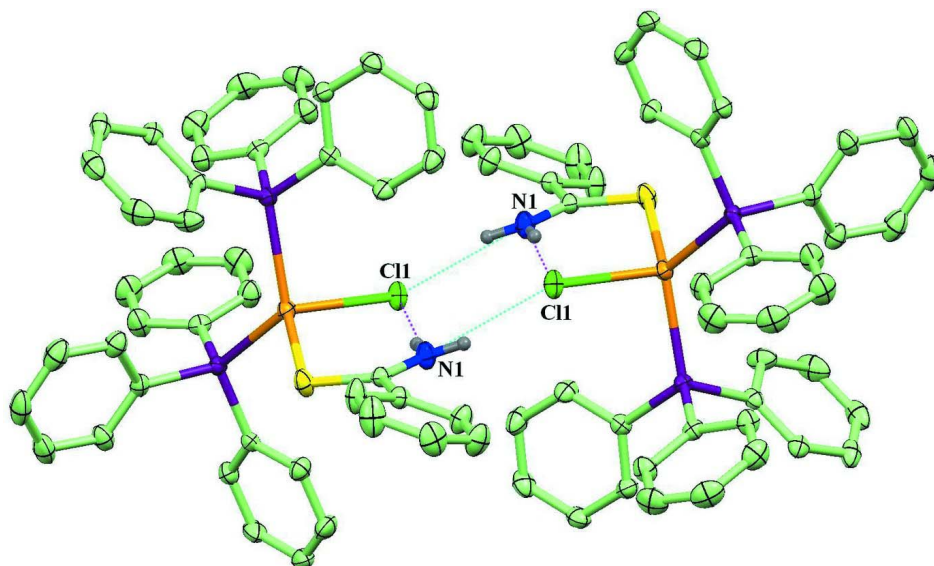


Figure 1

The molecular structure with displacement ellipsoids drawn at the 50% probability level and H atoms are omitted for clarity.

**Figure 2**

Part of the crystal structure showing the intermolecular and intramolecular hydrogen bonds as dashed lines are drawn between the non-hydrogen atoms. The symmetry related molecule is generated by the operator $(-x+2, -y, -z)$.

(Benzenecarbothioamide- κ S)chloridobis(triphenylphosphane- κ P)silver(I)

Crystal data

$[\text{AgCl}(\text{C}_7\text{H}_7\text{NS})(\text{C}_{18}\text{H}_{15}\text{P})_2]$

$M_r = 805.05$

Monoclinic, $P2_1/n$

$a = 14.2405 (7) \text{ \AA}$

$b = 13.9271 (7) \text{ \AA}$

$c = 19.0774 (10) \text{ \AA}$

$\beta = 98.711 (1)^\circ$

$V = 3740.0 (3) \text{ \AA}^3$

$Z = 4$

$F(000) = 1648$

$D_x = 1.430 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 7876 reflections

$\theta = 2.4\text{--}28.3^\circ$

$\mu = 0.78 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Block, colorless

$0.30 \times 0.16 \times 0.10 \text{ mm}$

Data collection

Bruker SMART APEX CCD
diffractometer

Radiation source: sealed tube

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 2003)

$T_{\min} = 0.861$, $T_{\max} = 0.923$

51284 measured reflections

9308 independent reflections

8544 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.031$

$\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 1.7^\circ$

$h = -19 \rightarrow 18$

$k = -18 \rightarrow 18$

$l = -25 \rightarrow 25$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.033$

$wR(F^2) = 0.075$

$S = 1.13$

9308 reflections

442 parameters

0 restraints

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0351P)^2 + 1.9067P]$

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

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

$\Delta\rho_{\max} = 0.72 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.39 \text{ e \AA}^{-3}$

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.

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	0.96337 (2)	0.11780 (2)	0.18826 (2)	0.01933 (5)
Cl1	1.07616 (3)	0.00745 (4)	0.12713 (2)	0.02523 (10)
S1	0.79118 (3)	0.07930 (4)	0.13044 (2)	0.02617 (11)
P1	1.02572 (3)	0.27274 (3)	0.15422 (2)	0.01705 (9)
P2	0.98356 (3)	0.07020 (3)	0.31343 (2)	0.01641 (9)
N1	0.86973 (12)	-0.00117 (13)	0.02851 (9)	0.0257 (4)
H1A	0.8678	-0.0242	-0.0135	0.031*
H1B	0.9221	-0.0018	0.0576	0.031*
C1	1.02873 (12)	0.29144 (13)	0.05997 (9)	0.0187 (3)
C2	1.05820 (14)	0.37795 (14)	0.03329 (10)	0.0236 (4)
H2	1.0772	0.4285	0.0640	0.028*
C3	1.05912 (15)	0.38854 (15)	-0.03897 (11)	0.0268 (4)
H3	1.0791	0.4460	-0.0566	0.032*
C4	1.03040 (16)	0.31370 (16)	-0.08444 (11)	0.0296 (4)
H4	1.0312	0.3210	-0.1328	0.036*
C5	1.00051 (17)	0.22818 (16)	-0.05892 (11)	0.0324 (5)
H5	0.9810	0.1781	-0.0899	0.039*
C6	0.99972 (15)	0.21714 (14)	0.01362 (10)	0.0258 (4)
H6	0.9796	0.1595	0.0309	0.031*
C7	1.14768 (13)	0.29086 (13)	0.19631 (10)	0.0200 (4)
C8	1.16664 (14)	0.30735 (15)	0.26924 (10)	0.0263 (4)
H8	1.1166	0.3123	0.2952	0.032*
C9	1.25915 (16)	0.31638 (17)	0.30335 (12)	0.0352 (5)
H9	1.2711	0.3289	0.3518	0.042*
C10	1.33373 (16)	0.30669 (18)	0.26532 (13)	0.0374 (5)
H10	1.3960	0.3129	0.2881	0.045*
C11	1.31608 (15)	0.28778 (18)	0.19375 (13)	0.0377 (5)
H11	1.3665	0.2800	0.1685	0.045*
C12	1.22346 (14)	0.28022 (16)	0.15898 (11)	0.0286 (4)
H12	1.2121	0.2680	0.1105	0.034*
C13	0.96006 (13)	0.37748 (13)	0.17718 (10)	0.0197 (4)
C14	0.86150 (15)	0.37148 (15)	0.16127 (13)	0.0305 (5)
H14	0.8335	0.3157	0.1411	0.037*
C15	0.80496 (16)	0.44786 (18)	0.17519 (14)	0.0390 (6)
H15	0.7393	0.4438	0.1632	0.047*
C16	0.84564 (17)	0.53014 (17)	0.20690 (12)	0.0348 (5)
H16	0.8075	0.5811	0.2166	0.042*
C17	0.94290 (16)	0.53631 (15)	0.22403 (10)	0.0286 (4)
H17	0.9702	0.5913	0.2460	0.034*

C18	1.00066 (14)	0.46105 (14)	0.20872 (10)	0.0223 (4)
H18	1.0664	0.4664	0.2195	0.027*
C19	1.10059 (13)	0.08806 (13)	0.36558 (10)	0.0193 (4)
C20	1.11456 (15)	0.12565 (17)	0.43381 (11)	0.0319 (5)
H20	1.0625	0.1439	0.4548	0.038*
C21	1.20579 (16)	0.13612 (19)	0.47091 (13)	0.0386 (5)
H21	1.2144	0.1611	0.5167	0.046*
C22	1.28319 (15)	0.10983 (17)	0.44038 (13)	0.0358 (5)
H22	1.3442	0.1164	0.4654	0.043*
C23	1.26999 (16)	0.07346 (19)	0.37215 (14)	0.0385 (5)
H23	1.3224	0.0563	0.3511	0.046*
C24	1.17911 (14)	0.06228 (16)	0.33477 (12)	0.0297 (4)
H24	1.1709	0.0375	0.2890	0.036*
C25	0.90286 (13)	0.13427 (14)	0.36321 (10)	0.0201 (4)
C26	0.86802 (14)	0.09695 (16)	0.42206 (11)	0.0272 (4)
H26	0.8856	0.0355	0.4380	0.033*
C27	0.80725 (15)	0.15072 (19)	0.45705 (12)	0.0362 (5)
H27	0.7838	0.1249	0.4959	0.043*
C28	0.78167 (16)	0.2422 (2)	0.43432 (13)	0.0413 (6)
H28	0.7413	0.2781	0.4580	0.050*
C29	0.81590 (16)	0.28085 (17)	0.37632 (14)	0.0381 (5)
H29	0.7990	0.3429	0.3613	0.046*
C30	0.87564 (14)	0.22677 (15)	0.34051 (11)	0.0280 (4)
H30	0.8977	0.2525	0.3011	0.034*
C31	0.96144 (13)	-0.05724 (13)	0.32671 (9)	0.0185 (3)
C32	0.91259 (14)	-0.10887 (14)	0.26999 (11)	0.0246 (4)
H32	0.8938	-0.0785	0.2267	0.030*
C33	0.89178 (15)	-0.20533 (15)	0.27774 (12)	0.0297 (4)
H33	0.8587	-0.2391	0.2398	0.036*
C34	0.92006 (16)	-0.25143 (15)	0.34174 (12)	0.0295 (4)
H34	0.9052	-0.3158	0.3471	0.035*
C35	0.97056 (16)	-0.20123 (15)	0.39764 (11)	0.0288 (4)
H35	0.9907	-0.2323	0.4404	0.035*
C36	0.99138 (14)	-0.10483 (14)	0.39031 (10)	0.0243 (4)
H36	1.0256	-0.0717	0.4282	0.029*
C37	0.79299 (13)	0.03483 (13)	0.04814 (9)	0.0200 (4)
C38	0.70443 (14)	0.03396 (14)	-0.00449 (9)	0.0219 (4)
C39	0.69078 (16)	-0.03464 (16)	-0.05811 (11)	0.0307 (5)
H39	0.7379	-0.0797	-0.0620	0.037*
C40	0.60665 (17)	-0.03594 (18)	-0.10586 (12)	0.0379 (5)
H40	0.5977	-0.0820	-0.1415	0.045*
C41	0.53653 (16)	0.03057 (19)	-0.10061 (12)	0.0386 (5)
H41	0.4800	0.0289	-0.1323	0.046*
C42	0.55038 (18)	0.0996 (2)	-0.04827 (13)	0.0444 (6)
H42	0.5035	0.1451	-0.0450	0.053*
C43	0.63401 (16)	0.10130 (18)	-0.00047 (12)	0.0348 (5)
H43	0.6429	0.1482	0.0346	0.042*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Ag1	0.02407 (8)	0.01923 (7)	0.01454 (7)	-0.00135 (5)	0.00242 (5)	0.00012 (5)
Cl1	0.0264 (2)	0.0283 (2)	0.0204 (2)	0.00856 (19)	0.00159 (17)	-0.00369 (18)
S1	0.0203 (2)	0.0411 (3)	0.0165 (2)	0.0002 (2)	0.00093 (17)	-0.0062 (2)
P1	0.0174 (2)	0.0171 (2)	0.0162 (2)	-0.00025 (17)	0.00140 (16)	0.00061 (16)
P2	0.0171 (2)	0.0186 (2)	0.0133 (2)	-0.00141 (17)	0.00131 (16)	-0.00025 (16)
N1	0.0225 (8)	0.0345 (9)	0.0191 (8)	0.0021 (7)	-0.0006 (6)	-0.0072 (7)
C1	0.0167 (8)	0.0210 (9)	0.0179 (8)	0.0013 (7)	0.0012 (6)	0.0017 (7)
C2	0.0261 (10)	0.0208 (9)	0.0237 (9)	-0.0039 (7)	0.0035 (7)	-0.0005 (7)
C3	0.0301 (10)	0.0256 (10)	0.0261 (10)	-0.0010 (8)	0.0083 (8)	0.0060 (8)
C4	0.0378 (12)	0.0331 (11)	0.0185 (9)	0.0057 (9)	0.0058 (8)	0.0053 (8)
C5	0.0494 (14)	0.0257 (10)	0.0201 (9)	-0.0008 (9)	-0.0013 (9)	-0.0028 (8)
C6	0.0353 (11)	0.0210 (9)	0.0199 (9)	-0.0015 (8)	0.0000 (8)	0.0034 (7)
C7	0.0192 (8)	0.0178 (8)	0.0220 (9)	0.0003 (7)	0.0000 (7)	0.0005 (7)
C8	0.0250 (10)	0.0304 (10)	0.0226 (9)	0.0018 (8)	0.0007 (8)	-0.0007 (8)
C9	0.0344 (12)	0.0376 (12)	0.0295 (11)	0.0001 (10)	-0.0083 (9)	-0.0021 (9)
C10	0.0222 (10)	0.0395 (13)	0.0463 (13)	-0.0031 (9)	-0.0088 (9)	0.0019 (10)
C11	0.0198 (10)	0.0463 (14)	0.0478 (14)	0.0014 (9)	0.0072 (9)	0.0036 (11)
C12	0.0241 (10)	0.0350 (11)	0.0265 (10)	0.0017 (8)	0.0036 (8)	-0.0010 (8)
C13	0.0207 (9)	0.0209 (9)	0.0183 (8)	0.0029 (7)	0.0052 (7)	0.0039 (7)
C14	0.0216 (10)	0.0261 (10)	0.0439 (13)	0.0003 (8)	0.0050 (9)	0.0054 (9)
C15	0.0228 (10)	0.0388 (13)	0.0580 (15)	0.0107 (9)	0.0139 (10)	0.0135 (11)
C16	0.0421 (13)	0.0315 (11)	0.0346 (11)	0.0170 (10)	0.0180 (10)	0.0103 (9)
C17	0.0449 (12)	0.0240 (10)	0.0170 (9)	0.0079 (9)	0.0050 (8)	0.0014 (7)
C18	0.0265 (9)	0.0214 (9)	0.0187 (8)	0.0022 (7)	0.0019 (7)	0.0017 (7)
C19	0.0175 (8)	0.0193 (8)	0.0200 (8)	-0.0007 (7)	-0.0007 (7)	0.0022 (7)
C20	0.0225 (10)	0.0473 (13)	0.0255 (10)	-0.0013 (9)	0.0019 (8)	-0.0068 (9)
C21	0.0291 (11)	0.0536 (15)	0.0298 (11)	-0.0051 (10)	-0.0064 (9)	-0.0102 (10)
C22	0.0199 (10)	0.0411 (13)	0.0429 (13)	-0.0012 (9)	-0.0070 (9)	0.0014 (10)
C23	0.0201 (10)	0.0462 (14)	0.0494 (14)	0.0062 (9)	0.0065 (9)	-0.0019 (11)
C24	0.0239 (10)	0.0342 (11)	0.0308 (11)	0.0024 (8)	0.0037 (8)	-0.0061 (9)
C25	0.0151 (8)	0.0247 (9)	0.0196 (8)	-0.0004 (7)	-0.0001 (7)	-0.0053 (7)
C26	0.0212 (9)	0.0377 (12)	0.0229 (9)	-0.0005 (8)	0.0041 (7)	-0.0030 (8)
C27	0.0250 (11)	0.0556 (15)	0.0297 (11)	-0.0022 (10)	0.0096 (9)	-0.0119 (10)
C28	0.0259 (11)	0.0541 (16)	0.0441 (14)	0.0043 (10)	0.0062 (10)	-0.0252 (12)
C29	0.0311 (11)	0.0312 (12)	0.0499 (14)	0.0070 (9)	-0.0004 (10)	-0.0121 (10)
C30	0.0256 (10)	0.0264 (10)	0.0307 (10)	0.0014 (8)	0.0001 (8)	-0.0041 (8)
C31	0.0173 (8)	0.0197 (9)	0.0194 (8)	-0.0008 (7)	0.0060 (7)	-0.0005 (7)
C32	0.0263 (10)	0.0244 (10)	0.0220 (9)	-0.0008 (8)	0.0003 (8)	-0.0021 (7)
C33	0.0325 (11)	0.0233 (10)	0.0327 (11)	-0.0047 (8)	0.0029 (9)	-0.0080 (8)
C34	0.0345 (11)	0.0193 (9)	0.0384 (11)	-0.0029 (8)	0.0170 (9)	-0.0016 (8)
C35	0.0393 (12)	0.0250 (10)	0.0245 (10)	0.0002 (9)	0.0123 (9)	0.0053 (8)
C36	0.0292 (10)	0.0254 (10)	0.0186 (9)	-0.0019 (8)	0.0049 (8)	-0.0005 (7)
C37	0.0227 (9)	0.0198 (9)	0.0172 (8)	-0.0031 (7)	0.0017 (7)	0.0003 (7)
C38	0.0231 (9)	0.0255 (9)	0.0162 (8)	-0.0035 (7)	-0.0003 (7)	0.0003 (7)
C39	0.0307 (11)	0.0296 (11)	0.0292 (10)	0.0013 (9)	-0.0034 (8)	-0.0070 (9)

C40	0.0370 (12)	0.0416 (13)	0.0309 (11)	-0.0022 (10)	-0.0080 (9)	-0.0118 (10)
C41	0.0292 (11)	0.0586 (16)	0.0242 (10)	0.0023 (11)	-0.0082 (9)	-0.0049 (10)
C42	0.0349 (13)	0.0619 (17)	0.0326 (12)	0.0199 (12)	-0.0072 (10)	-0.0099 (11)
C43	0.0336 (12)	0.0441 (13)	0.0239 (10)	0.0105 (10)	-0.0042 (9)	-0.0115 (9)

Geometric parameters (Å, °)

Ag1—P2	2.4529 (5)	C18—H18	0.9300
Ag1—P1	2.4578 (5)	C19—C24	1.387 (3)
Ag1—S1	2.5880 (5)	C19—C20	1.389 (3)
Ag1—C11	2.6208 (5)	C20—C21	1.390 (3)
S1—C37	1.6918 (19)	C20—H20	0.9300
P1—C7	1.8173 (19)	C21—C22	1.372 (3)
P1—C13	1.8216 (19)	C21—H21	0.9300
P1—C1	1.8238 (18)	C22—C23	1.383 (3)
P2—C19	1.8240 (18)	C22—H22	0.9300
P2—C31	1.8269 (19)	C23—C24	1.389 (3)
P2—C25	1.8300 (19)	C23—H23	0.9300
N1—C37	1.308 (2)	C24—H24	0.9300
N1—H1A	0.8600	C25—C26	1.395 (3)
N1—H1B	0.8600	C25—C30	1.395 (3)
C1—C6	1.383 (3)	C26—C27	1.389 (3)
C1—C2	1.397 (3)	C26—H26	0.9300
C2—C3	1.388 (3)	C27—C28	1.377 (4)
C2—H2	0.9300	C27—H27	0.9300
C3—C4	1.378 (3)	C28—C29	1.384 (4)
C3—H3	0.9300	C28—H28	0.9300
C4—C5	1.378 (3)	C29—C30	1.391 (3)
C4—H4	0.9300	C29—H29	0.9300
C5—C6	1.394 (3)	C30—H30	0.9300
C5—H5	0.9300	C31—C36	1.392 (3)
C6—H6	0.9300	C31—C32	1.394 (3)
C7—C12	1.388 (3)	C32—C33	1.388 (3)
C7—C8	1.396 (3)	C32—H32	0.9300
C8—C9	1.384 (3)	C33—C34	1.384 (3)
C8—H8	0.9300	C33—H33	0.9300
C9—C10	1.381 (3)	C34—C35	1.382 (3)
C9—H9	0.9300	C34—H34	0.9300
C10—C11	1.376 (3)	C35—C36	1.386 (3)
C10—H10	0.9300	C35—H35	0.9300
C11—C12	1.388 (3)	C36—H36	0.9300
C11—H11	0.9300	C37—C38	1.488 (2)
C12—H12	0.9300	C38—C43	1.384 (3)
C13—C14	1.393 (3)	C38—C39	1.392 (3)
C13—C18	1.395 (3)	C39—C40	1.391 (3)
C14—C15	1.384 (3)	C39—H39	0.9300
C14—H14	0.9300	C40—C41	1.377 (3)
C15—C16	1.381 (4)	C40—H40	0.9300

C15—H15	0.9300	C41—C42	1.378 (3)
C16—C17	1.377 (3)	C41—H41	0.9300
C16—H16	0.9300	C42—C43	1.386 (3)
C17—C18	1.390 (3)	C42—H42	0.9300
C17—H17	0.9300	C43—H43	0.9300
P2—Ag1—P1	120.053 (16)	C24—C19—C20	118.98 (18)
P2—Ag1—S1	108.823 (16)	C24—C19—P2	117.53 (15)
P1—Ag1—S1	115.223 (17)	C20—C19—P2	123.49 (15)
P2—Ag1—C11	106.579 (16)	C19—C20—C21	120.5 (2)
P1—Ag1—C11	97.298 (16)	C19—C20—H20	119.8
S1—Ag1—C11	107.087 (16)	C21—C20—H20	119.8
C37—S1—Ag1	108.95 (7)	C22—C21—C20	120.3 (2)
C7—P1—C13	105.87 (9)	C22—C21—H21	119.8
C7—P1—C1	104.61 (8)	C20—C21—H21	119.8
C13—P1—C1	102.26 (8)	C21—C22—C23	119.6 (2)
C7—P1—Ag1	111.37 (6)	C21—C22—H22	120.2
C13—P1—Ag1	114.86 (6)	C23—C22—H22	120.2
C1—P1—Ag1	116.70 (6)	C22—C23—C24	120.5 (2)
C19—P2—C31	102.67 (8)	C22—C23—H23	119.7
C19—P2—C25	104.13 (8)	C24—C23—H23	119.7
C31—P2—C25	105.47 (9)	C19—C24—C23	120.1 (2)
C19—P2—Ag1	117.18 (6)	C19—C24—H24	119.9
C31—P2—Ag1	113.60 (6)	C23—C24—H24	119.9
C25—P2—Ag1	112.53 (6)	C26—C25—C30	118.54 (18)
C37—N1—H1A	120.0	C26—C25—P2	124.32 (15)
C37—N1—H1B	120.0	C30—C25—P2	117.13 (15)
H1A—N1—H1B	120.0	C27—C26—C25	120.6 (2)
C6—C1—C2	119.31 (17)	C27—C26—H26	119.7
C6—C1—P1	118.31 (14)	C25—C26—H26	119.7
C2—C1—P1	122.38 (14)	C28—C27—C26	120.1 (2)
C3—C2—C1	120.12 (18)	C28—C27—H27	119.9
C3—C2—H2	119.9	C26—C27—H27	119.9
C1—C2—H2	119.9	C27—C28—C29	120.2 (2)
C4—C3—C2	119.88 (18)	C27—C28—H28	119.9
C4—C3—H3	120.1	C29—C28—H28	119.9
C2—C3—H3	120.1	C28—C29—C30	119.9 (2)
C3—C4—C5	120.62 (19)	C28—C29—H29	120.1
C3—C4—H4	119.7	C30—C29—H29	120.1
C5—C4—H4	119.7	C29—C30—C25	120.6 (2)
C4—C5—C6	119.69 (19)	C29—C30—H30	119.7
C4—C5—H5	120.2	C25—C30—H30	119.7
C6—C5—H5	120.2	C36—C31—C32	118.82 (17)
C1—C6—C5	120.38 (19)	C36—C31—P2	123.17 (14)
C1—C6—H6	119.8	C32—C31—P2	118.01 (14)
C5—C6—H6	119.8	C33—C32—C31	120.38 (19)
C12—C7—C8	118.76 (18)	C33—C32—H32	119.8
C12—C7—P1	121.58 (15)	C31—C32—H32	119.8

C8—C7—P1	119.41 (15)	C34—C33—C32	120.30 (19)
C9—C8—C7	120.7 (2)	C34—C33—H33	119.9
C9—C8—H8	119.7	C32—C33—H33	119.9
C7—C8—H8	119.7	C35—C34—C33	119.59 (19)
C10—C9—C8	119.8 (2)	C35—C34—H34	120.2
C10—C9—H9	120.1	C33—C34—H34	120.2
C8—C9—H9	120.1	C34—C35—C36	120.40 (19)
C11—C10—C9	120.0 (2)	C34—C35—H35	119.8
C11—C10—H10	120.0	C36—C35—H35	119.8
C9—C10—H10	120.0	C35—C36—C31	120.48 (19)
C10—C11—C12	120.4 (2)	C35—C36—H36	119.8
C10—C11—H11	119.8	C31—C36—H36	119.8
C12—C11—H11	119.8	N1—C37—C38	117.60 (16)
C11—C12—C7	120.2 (2)	N1—C37—S1	122.48 (14)
C11—C12—H12	119.9	C38—C37—S1	119.91 (14)
C7—C12—H12	119.9	C43—C38—C39	118.97 (18)
C14—C13—C18	118.83 (18)	C43—C38—C37	120.29 (17)
C14—C13—P1	115.88 (15)	C39—C38—C37	120.74 (18)
C18—C13—P1	125.29 (14)	C40—C39—C38	120.1 (2)
C15—C14—C13	120.6 (2)	C40—C39—H39	120.0
C15—C14—H14	119.7	C38—C39—H39	120.0
C13—C14—H14	119.7	C41—C40—C39	120.4 (2)
C16—C15—C14	120.3 (2)	C41—C40—H40	119.8
C16—C15—H15	119.9	C39—C40—H40	119.8
C14—C15—H15	119.9	C40—C41—C42	119.7 (2)
C17—C16—C15	119.7 (2)	C40—C41—H41	120.1
C17—C16—H16	120.1	C42—C41—H41	120.1
C15—C16—H16	120.1	C41—C42—C43	120.2 (2)
C16—C17—C18	120.6 (2)	C41—C42—H42	119.9
C16—C17—H17	119.7	C43—C42—H42	119.9
C18—C17—H17	119.7	C38—C43—C42	120.6 (2)
C17—C18—C13	119.99 (19)	C38—C43—H43	119.7
C17—C18—H18	120.0	C42—C43—H43	119.7
C13—C18—H18	120.0		
C7—P1—C1—C6	121.08 (16)	P2—C19—C20—C21	179.13 (18)
C13—P1—C1—C6	-128.69 (16)	C19—C20—C21—C22	0.3 (4)
Ag1—P1—C1—C6	-2.47 (17)	C20—C21—C22—C23	0.5 (4)
C7—P1—C1—C2	-59.59 (17)	C21—C22—C23—C24	-0.8 (4)
C13—P1—C1—C2	50.64 (17)	C20—C19—C24—C23	0.5 (3)
Ag1—P1—C1—C2	176.86 (13)	P2—C19—C24—C23	-179.44 (18)
C6—C1—C2—C3	-0.6 (3)	C22—C23—C24—C19	0.3 (4)
P1—C1—C2—C3	-179.96 (15)	C19—P2—C25—C26	-81.47 (18)
C1—C2—C3—C4	0.4 (3)	C31—P2—C25—C26	26.25 (18)
C2—C3—C4—C5	0.1 (3)	Ag1—P2—C25—C26	150.62 (15)
C3—C4—C5—C6	-0.3 (3)	C19—P2—C25—C30	97.99 (15)
C2—C1—C6—C5	0.5 (3)	C31—P2—C25—C30	-154.29 (14)
P1—C1—C6—C5	179.81 (16)	Ag1—P2—C25—C30	-29.92 (16)

C4—C5—C6—C1	0.0 (3)	C30—C25—C26—C27	0.2 (3)
C13—P1—C7—C12	-130.56 (17)	P2—C25—C26—C27	179.70 (16)
C1—P1—C7—C12	-22.97 (19)	C25—C26—C27—C28	-0.8 (3)
Ag1—P1—C7—C12	103.94 (16)	C26—C27—C28—C29	0.4 (3)
C13—P1—C7—C8	55.30 (17)	C27—C28—C29—C30	0.6 (3)
C1—P1—C7—C8	162.89 (16)	C28—C29—C30—C25	-1.1 (3)
Ag1—P1—C7—C8	-70.20 (16)	C26—C25—C30—C29	0.7 (3)
C12—C7—C8—C9	2.3 (3)	P2—C25—C30—C29	-178.80 (16)
P1—C7—C8—C9	176.59 (17)	C19—P2—C31—C36	35.53 (18)
C7—C8—C9—C10	-1.6 (3)	C25—P2—C31—C36	-73.24 (17)
C8—C9—C10—C11	-0.2 (4)	Ag1—P2—C31—C36	163.06 (14)
C9—C10—C11—C12	1.3 (4)	C19—P2—C31—C32	-144.12 (15)
C10—C11—C12—C7	-0.6 (4)	C25—P2—C31—C32	107.10 (16)
C8—C7—C12—C11	-1.2 (3)	Ag1—P2—C31—C32	-16.59 (17)
P1—C7—C12—C11	-175.38 (17)	C36—C31—C32—C33	2.0 (3)
C7—P1—C13—C14	-168.93 (15)	P2—C31—C32—C33	-178.37 (16)
C1—P1—C13—C14	81.80 (16)	C31—C32—C33—C34	-0.5 (3)
Ag1—P1—C13—C14	-45.60 (17)	C32—C33—C34—C35	-1.1 (3)
C7—P1—C13—C18	11.10 (18)	C33—C34—C35—C36	1.2 (3)
C1—P1—C13—C18	-98.18 (17)	C34—C35—C36—C31	0.2 (3)
Ag1—P1—C13—C18	134.42 (15)	C32—C31—C36—C35	-1.8 (3)
C18—C13—C14—C15	1.3 (3)	P2—C31—C36—C35	178.54 (15)
P1—C13—C14—C15	-178.66 (18)	Ag1—S1—C37—N1	19.52 (18)
C13—C14—C15—C16	-1.8 (4)	Ag1—S1—C37—C38	-161.76 (13)
C14—C15—C16—C17	0.6 (3)	N1—C37—C38—C43	-153.7 (2)
C15—C16—C17—C18	1.0 (3)	S1—C37—C38—C43	27.5 (3)
C16—C17—C18—C13	-1.5 (3)	N1—C37—C38—C39	26.7 (3)
C14—C13—C18—C17	0.3 (3)	S1—C37—C38—C39	-152.13 (17)
P1—C13—C18—C17	-179.73 (14)	C43—C38—C39—C40	-1.1 (3)
C31—P2—C19—C24	81.55 (17)	C37—C38—C39—C40	178.5 (2)
C25—P2—C19—C24	-168.66 (16)	C38—C39—C40—C41	0.2 (4)
Ag1—P2—C19—C24	-43.67 (18)	C39—C40—C41—C42	0.8 (4)
C31—P2—C19—C20	-98.36 (18)	C40—C41—C42—C43	-0.9 (4)
C25—P2—C19—C20	11.4 (2)	C39—C38—C43—C42	1.1 (4)
Ag1—P2—C19—C20	136.42 (16)	C37—C38—C43—C42	-178.5 (2)
C24—C19—C20—C21	-0.8 (3)	C41—C42—C43—C38	-0.1 (4)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots C11 ⁱ	0.86	2.43	3.1798 (17)	146
N1—H1B \cdots C11	0.86	2.39	3.2434 (17)	173

Symmetry code: (i) $-x+2, -y, -z$.