

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

(anti-Chloridothiosemicarbazide- κ S)bis-(triphenylphosphane- κ P)copper(I) 0.48-hydrate

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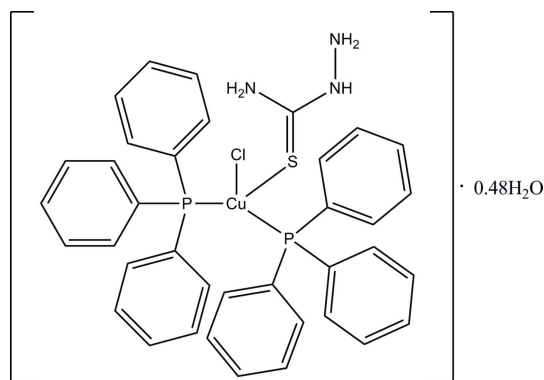
Received 21 March 2013; accepted 28 March 2013

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; disorder in solvent or counterion; R factor = 0.047; wR factor = 0.110; data-to-parameter ratio = 20.1.

In the mononuclear title complex, $[\text{CuCl}(\text{CH}_5\text{N}_3\text{S})(\text{C}_{18}\text{H}_{15}\text{P})_2] \cdot 0.48\text{H}_2\text{O}$, the Cu^{I} ion is in a slightly distorted tetrahedral coordination geometry formed by two P atoms from two triphenylphosphane ligands, one S atom from a thiosemicarbazide ligand and one chloride anion. An intramolecular $\text{N}-\text{H} \cdots \text{N}$ hydrogen bond [graph-set motif $S(5)$] stabilizes the thiosemicarbazide ligand in its *anti* conformation, and an intramolecular $\text{N}-\text{H} \cdots \text{Cl}$ hydrogen bond between the hydrazine $\text{N}-\text{H}$ group and the chloride anion influences the arrangement and orientation of the ligands around the metal center. A weak intramolecular $\text{C}-\text{H} \cdots \text{Cl}$ hydrogen bond is also present. In the crystal, complex molecules are connected through $\text{N}-\text{H} \cdots \text{Cl}$ hydrogen bonds originating from the amide $-\text{NH}_2$ group, and through $\text{O}-\text{H} \cdots \text{S}$ and $\text{O}-\text{H} \cdots \text{Cl}$ hydrogen bonds involving the solvent water molecule. Both the direct $\text{N}-\text{H} \cdots \text{Cl}$ hydrogen bonds as well as the bridging hydrogen bonds mediated by the water molecule connect the complex molecules into zigzag chains that propagate along $[010]$. The solvent water molecule is partially occupied, with a refined occupancy of 0.479 (7).

Related literature

For the coordination of thiosemicarbazide and thiosemicarbazones with metal complexes, see: Andreotti *et al.* (1970); Chattopadhyay *et al.* (1991); Jia *et al.* (2008*a,b*); Villa *et al.* (1972*a,b*); Qirong *et al.* (1987). For potential applications of related complexes, see: Alagarsamy & Parthiban (2011); Kowol *et al.* (2007); Pelosi (2010); Yu *et al.* (2009); Wattanakanjana *et al.* (2012). For hydrogen-bond graph-set motifs, see: Bernstein, *et al.* (1995). For a description of the Cambridge Structural Database (CSD), see: Allen (2002).



Experimental

Crystal data

$[\text{CuCl}(\text{CH}_5\text{N}_3\text{S})(\text{C}_{18}\text{H}_{15}\text{P})_2] \cdot 0.48\text{H}_2\text{O}$
 $M_r = 723.29$
 Monoclinic, $P2_1/n$
 $a = 14.8723$ (7) Å
 $b = 12.4829$ (6) Å
 $c = 19.2103$ (9) Å

$\beta = 96.126$ (1)°
 $V = 3546.0$ (3) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.87$ mm⁻¹
 $T = 293$ K
 $0.34 \times 0.11 \times 0.07$ mm

Data collection

Bruker SMART APEX CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2003)
 $T_{\text{min}} = 0.880$, $T_{\text{max}} = 1$

48021 measured reflections
 8591 independent reflections
 6691 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.110$
 $S = 1.06$
 8591 reflections
 428 parameters
 4 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.42$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{O1}-\text{H1C} \cdots \text{Cl1}$	0.82 (2)	2.37 (2)	3.186 (5)	179 (12)
$\text{O1}-\text{H1D} \cdots \text{S1}^{\text{i}}$	0.82 (2)	2.70 (10)	3.218 (5)	123 (9)
$\text{N1}-\text{H1B} \cdots \text{Cl1}^{\text{ii}}$	0.86	2.48	3.302 (3)	161
$\text{N1}-\text{H1A} \cdots \text{N3}$	0.86	2.27	2.628 (5)	105
$\text{N2}-\text{H2} \cdots \text{Cl1}$	0.86	2.35	3.202 (2)	170
$\text{C42}-\text{H42} \cdots \text{Cl1}$	0.93	2.72	3.626 (3)	164

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

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) and SHELXLE Rev609 (Hübschle *et al.*, 2011); molecular graphics: Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: SHELXL97 and publCIF (Westrip, 2010).

Financial support from the Center of Excellence for Innovation in Chemistry (PERCH-CIC), the Office of the Higher Education Commission, Ministry of Education and Depart-

ment of Chemistry, Prince of Songkla University, is gratefully acknowledged. RN would like to thank Dr Matthias Zeller for valuable suggestions and assistance with X-ray structure determination and use of structure refinement programs.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5600).

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supporting information

Acta Cryst. (2013). E69, m244–m245 [https://doi.org/10.1107/S1600536813008556]

(*anti*-Chloridothiosemicabazide- κ S)bis(triphenylphosphane- κ P)copper(I) 0.48-hydrate

Ruthairat Nimthong, Chaveng Pakawatchai, Nuchanat Phongphayak and Yupa Wattanakanjana

S1. Comment

Thiosemicarbazonones, the condensation products of thiosemicarbazide and aldehydes, are compounds of considerable interest due to their remarkable number and variety of biological properties and the potential medical and pharmaceutical applications that result (Pelosi, 2010; Yu *et al.*, 2009). Biological applications are not limited to the thiosemicarbazonones, but their metal complexes have been described to also have similar effects, and are often even more active than the uncoordinated thiosemicarbazonones. Metal complexes of thiosemicarbazonones have been described to be active antimicrobial agents (Alagarsamy *et al.*, 2011), are highly effective anticancer agents (Yu *et al.*, 2009), they exhibit cytotoxicity and do interact with ribonucleotide reductase (Kowol *et al.*, 2007), to name just a few of the properties they have been investigated for. The parent compound of all thiosemicarbazonones, thiosemicarbazide, does also exhibit a rich metal coordination chemistry. Having no substituent at the hydrazide NH₂ group, thiosemicarbazide is able to coordinate to metal ions in an N,*S*-chelating mode, and this is realised in most of its complexes. Of 79 complexes of thiosemicarbazide reported in the Cambridge Structural Database (2013 release, Allen, 2002), 61 featured N,*S*-chelation of the ligand, 5 displayed N,*S*-chelation and additional coordination *via* the sulfur atom, and 13 displayed mono- or bidentate coordination *via* the sulfur only. Coordination *via* only nitrogen was not observed at all. The coordination mode towards the metal dictates the conformation of the thiosemicarbazide ligand. In N,*S*-chelating complexes, the ligand necessarily features a *syn*-conformation of the sulfur atom *versus* the hydrazide NH₂ group. In the complexes with only sulfur coordination both *syn* and *anti* conformation can be imagined. However, all reported examples feature the *anti*-conformation which is also observed in the parent compound. In salts of with protonated thiosemicarbazide cations, on the other hand, the *anti* conformation is again the only one realised (Andreotti *et al.*, 1970).

The reasons for the often increased biological activity of metal complexes, when compared to those of only the ligands by themselves, is manifold and depend highly on the individual case and type of biological activity under investigation. Common to many mechanisms is the need for the active compounds to pass through the cell membrane, a barrier difficult to penetrate for polar compounds such as thiosemicarbazonone. Coordination to metal complexes, especially when paired with other more lipophilic ligands, reduces the polarity of the thiosemicarbazonone compound. In an effort to investigate the influence of metal coordination and use of various phenyl phosphane ligands on the antibacterial properties of copper(I) and silver(I) complexes of thiosemicarbazonone we reacted thiosemicarbazonone with the triphenylphosphine adduct of copper(I) chloride

The use of monodentate triphenylphosphane lead to the formation of the mononuclear complex have prepared the title compound of this study, (*anti*-thiosemicabazide- κ S)chloridobis(triphenylphosphane- κ P)copper(I), which crystallized from acetonitrile in the form of its hemihydrate (with a refined occupancy of 0.479 (7)). As is typical for soft metal ions such as Cu(I), thiosemicarbazonone coordination is *via* the soft sulfur donor only, as had been observed for the other two Cu(I)

complexes of thiosemicarbazide (Jia *et al.*, 2008*a,b*). Copper(II), on the other hand shows N,*S*-chelation with thiosemicarbazide (Villa *et al.*, 1972*a,b*; Qirong *et al.*, 1987; Chattopadhyay *et al.*, 1991). In the title compound, the metal center is coordinated to one S donor of the thiosemicarbazide ligand, two P atoms of two triphenylphosphane ligands one chlorine atom. In agreement with expectation for a Cu(I) center with soft donor atoms, It displays a distorted tetrahedral coordination of the Cu^I center. (Fig. 1). The Cu—S bond distance of 2.3841 (7) Å is slightly longer than in the other two Cu(I) thiosemicarbazides (2.2401 (10) to 2.3192 (13) Å, Jia *et al.*, 2008*a,b*), but very similar to several other similar complexes such as the thiosemicarbazone complex [CuI(C₄H₉N₃S)(C₁₈H₁₅P)₂] with a Cu—S bond length of 2.3866 (7) Å (Wattanakanjana *et al.*, 2012). As in all other complexes with S-only coordination of thiosemicarbazides the ligand is in the *anti*-conformation, which is stabilized by an intramolecular N—H⋯N hydrogen bond between N1—H1A and N3 (graph set designator as S(5) (Bernstein *et al.*, 1995)). Another intramolecular hydrogen bond, between the hydrazine N2—H2 group and the chloride anion, orients the hydrazide N—H group towards the chlorine atom and, through this, influences the arrangement and orientation of the ligands around the Cu^I center. Neighboring complexes are connected with each other through N1—H1B⋯Cl1 hydrogen bonds originating from the amide NH₂ group, and through O1—H1D⋯S1 and O1—H1C⋯Cl1 hydrogen bonds involving the solvate water molecule. Both the direct N2—H2⋯Cl1 hydrogen bonds as well as the bridging H-bonds mediated by the water molecule are linking molecules with each other leading to formation of a one-dimensional zigzag chain parallel to the *b* axis (see Table and Fig.2).

S2. Experimental

Triphenylphosphane (0.53 g, 2 mmol) was dissolved in 30 cm³ of acetonitrile at 335 K. CuCl (0.10 g, 1 mmol) was added and the mixture was stirred for 1.5 h. Thiosemicarbazide (0.09 g, 1 mmol) was added and the reaction mixture was heated to reflux of the solvent for 5.5 h. The resulting clear solution was filtered and left to evaporate at room temperature. The crystalline solid, which precipitated upon standing for three days, was filtered off and dried under reduced pressure (0.38 g, yield 52%). Mp = 439–441 K. Main IR peaks (KBr, cm⁻¹): ν (N—H) 3420, 3248, 3126 m, ν (C—N) + δ (N—H) 1620, 1598, 1476 m, ν (C=S) + ν (C=N) + ν (C—N) 1326, 1308 m, ν (C—N) + ν (C—S) 1082 s, ν (C—S) 775 s.

S3. Refinement

The two hydrazide NH₂ H atoms were located in a difference Fourier map and refined isotropically, with N—H distances restrained to 0.86 (2) Å, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$. The remaining H atoms bonded to N or C atoms were positioned geometrically and refined using a riding model, with C—H = 0.93, N—H = 0.86 Å with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C/N})$. A solvate water molecule is partially occupied, the occupancy of the water molecule refined to 0.479 (7). The H atoms attached to the water molecules were located in a difference Fourier map and refined isotropically, with O—H distances restrained to 0.82 (2) Å, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$.

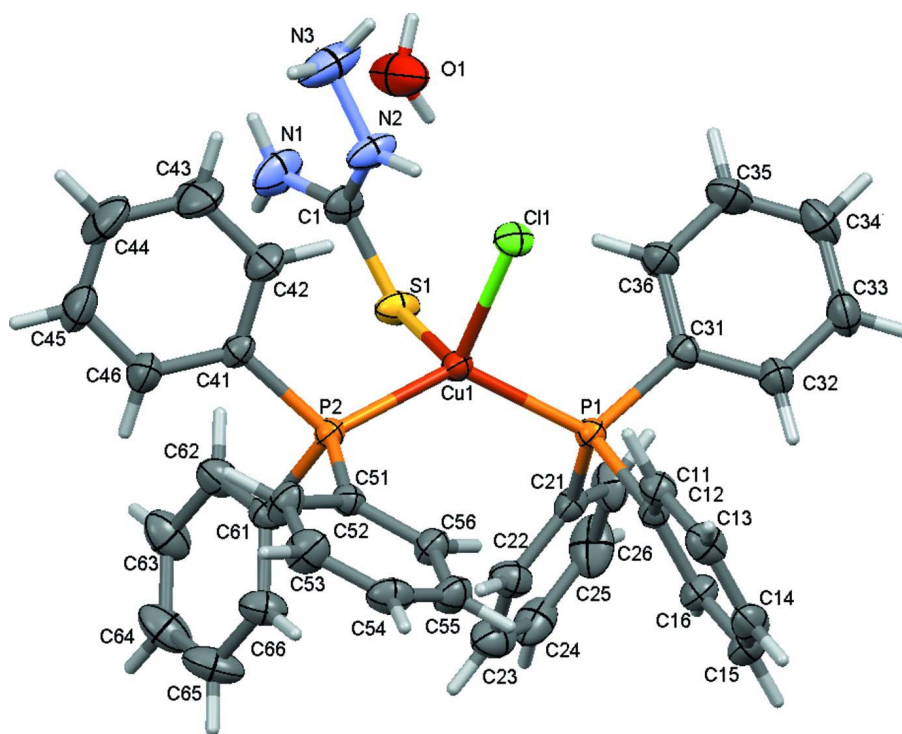


Figure 1

The molecular structure with displacement ellipsoids drawn at the 50% probability level.

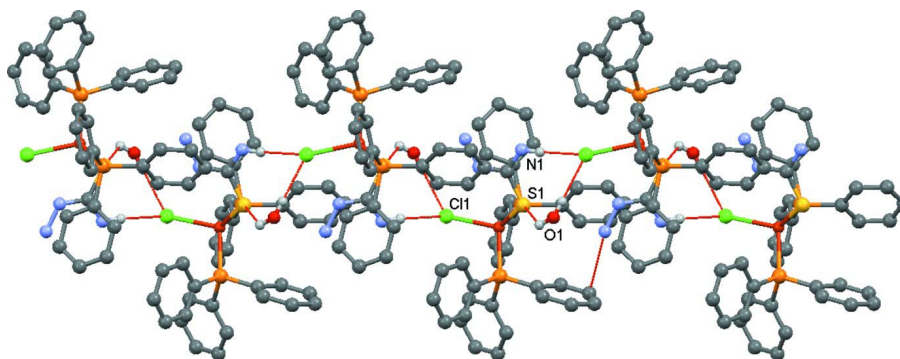


Figure 2

Part of the crystal structure with O—H...S and N—H...Cl hydrogen bonds, shown as dashed lines, which link molecules into one dimensional zigzag chain parallel to the *b* axis.

(anti-Chloridothiosemicabazide- κ S)bis(triphenylphosphane- κ P)copper(I) 0.48-hydrate

Crystal data

[CuCl(CH₅N₃S)(C₁₈H₁₅P)₂] \cdot 0.48H₂O

M_r = 723.29

Monoclinic, *P*2₁/*n*

a = 14.8723 (7) Å

b = 12.4829 (6) Å

c = 19.2103 (9) Å

β = 96.126 (1)°

V = 3546.0 (3) Å³

Z = 4

F(000) = 1499.2

D_x = 1.355 Mg m⁻³

Mo *K* α radiation, λ = 0.71073 Å

Cell parameters from 7520 reflections

θ = 2.3–23.0°

$\mu = 0.87 \text{ mm}^{-1}$
 $T = 293 \text{ K}$

Block, colorless
 $0.34 \times 0.11 \times 0.07 \text{ mm}$

Data collection

Bruker SMART APEX CCD
 diffractometer
 Radiation source: sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2003)
 $T_{\min} = 0.880$, $T_{\max} = 1$

48021 measured reflections
 8591 independent reflections
 6691 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$
 $\theta_{\max} = 28.1^\circ$, $\theta_{\min} = 1.7^\circ$
 $h = -19 \rightarrow 19$
 $k = -16 \rightarrow 16$
 $l = -25 \rightarrow 25$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.110$
 $S = 1.06$
 8591 reflections
 428 parameters
 4 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: mixed
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.050P)^2 + 1.0259P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.42 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$

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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Cu1	0.87688 (2)	0.69145 (2)	0.14512 (2)	0.03859 (9)	
O1	0.6525 (4)	0.3863 (6)	0.1104 (3)	0.106 (3)	0.479 (7)
H1C	0.701 (4)	0.417 (8)	0.120 (6)	0.159*	0.479 (7)
H1D	0.631 (7)	0.336 (6)	0.130 (5)	0.159*	0.479 (7)
C11	0.84280 (4)	0.50261 (5)	0.14456 (3)	0.04861 (15)	
S1	0.80953 (5)	0.77828 (5)	0.23718 (4)	0.05356 (18)	
P1	1.02443 (4)	0.69509 (5)	0.19010 (3)	0.03600 (13)	
P2	0.80809 (4)	0.77112 (5)	0.04621 (3)	0.03623 (14)	
N1	0.64208 (18)	0.7719 (2)	0.27084 (16)	0.0810 (8)	
H1A	0.5922	0.7396	0.2761	0.097*	
H1B	0.6491	0.8380	0.2829	0.097*	
N2	0.69393 (15)	0.61907 (19)	0.22675 (13)	0.0648 (7)	
H2	0.7356	0.5829	0.2096	0.078*	
N3	0.6107 (2)	0.5707 (3)	0.2361 (2)	0.1044 (13)	
H3A	0.583 (3)	0.566 (4)	0.1932 (13)	0.125*	
H3B	0.618 (3)	0.5052 (19)	0.248 (3)	0.125*	
C1	0.70773 (18)	0.7203 (2)	0.24444 (13)	0.0514 (6)	

C11	1.11233 (14)	0.66548 (18)	0.13309 (11)	0.0365 (5)
C12	1.09407 (17)	0.58379 (19)	0.08457 (13)	0.0469 (6)
H12	1.0388	0.5483	0.0817	0.056*
C13	1.1577 (2)	0.5551 (2)	0.04057 (14)	0.0567 (7)
H13	1.1452	0.5004	0.0081	0.068*
C14	1.2390 (2)	0.6069 (2)	0.04457 (15)	0.0585 (7)
H14	1.2815	0.5874	0.0147	0.070*
C15	1.25819 (18)	0.6875 (2)	0.09241 (16)	0.0583 (7)
H15	1.3138	0.7222	0.0950	0.070*
C16	1.19517 (16)	0.7173 (2)	0.13669 (13)	0.0477 (6)
H16	1.2083	0.7722	0.1689	0.057*
C21	1.05730 (15)	0.82700 (19)	0.22479 (13)	0.0418 (5)
C22	1.0509 (2)	0.9125 (2)	0.17833 (15)	0.0582 (7)
H22	1.0288	0.9006	0.1318	0.070*
C23	1.0763 (2)	1.0143 (2)	0.1993 (2)	0.0718 (9)
H23	1.0726	1.0702	0.1671	0.086*
C24	1.1072 (2)	1.0327 (3)	0.2678 (2)	0.0782 (10)
H24	1.1258	1.1011	0.2822	0.094*
C25	1.1109 (2)	0.9510 (3)	0.3150 (2)	0.0877 (12)
H25	1.1304	0.9644	0.3618	0.105*
C26	1.0856 (2)	0.8474 (3)	0.29384 (16)	0.0646 (8)
H26	1.0879	0.7923	0.3265	0.077*
C31	1.05437 (16)	0.60214 (19)	0.26296 (12)	0.0419 (5)
C32	1.14238 (18)	0.5696 (2)	0.28240 (14)	0.0559 (7)
H32	1.1889	0.5962	0.2586	0.067*
C33	1.1621 (2)	0.4978 (3)	0.33684 (16)	0.0677 (8)
H33	1.2217	0.4774	0.3498	0.081*
C34	1.0941 (2)	0.4567 (3)	0.37161 (16)	0.0727 (9)
H34	1.1074	0.4085	0.4082	0.087*
C35	1.0072 (2)	0.4869 (3)	0.35232 (16)	0.0747 (9)
H35	0.9611	0.4581	0.3755	0.090*
C36	0.98627 (19)	0.5600 (2)	0.29870 (14)	0.0573 (7)
H36	0.9266	0.5807	0.2867	0.069*
C41	0.68669 (15)	0.7430 (2)	0.02893 (12)	0.0422 (5)
C42	0.65597 (19)	0.6448 (3)	0.04902 (16)	0.0626 (7)
H42	0.6965	0.5956	0.0710	0.075*
C43	0.5651 (2)	0.6188 (3)	0.0366 (2)	0.0890 (11)
H43	0.5449	0.5524	0.0503	0.107*
C44	0.5049 (2)	0.6905 (4)	0.0044 (2)	0.0912 (12)
H44	0.4439	0.6731	-0.0035	0.109*
C45	0.5344 (2)	0.7876 (3)	-0.01622 (19)	0.0780 (10)
H45	0.4935	0.8360	-0.0386	0.094*
C46	0.62496 (18)	0.8145 (3)	-0.00402 (16)	0.0610 (7)
H46	0.6445	0.8812	-0.0180	0.073*
C51	0.85129 (15)	0.73927 (18)	-0.03696 (12)	0.0389 (5)
C52	0.79750 (19)	0.7307 (3)	-0.10006 (14)	0.0580 (7)
H52	0.7360	0.7458	-0.1022	0.070*
C53	0.8345 (2)	0.6999 (3)	-0.16006 (15)	0.0660 (8)

H53	0.7975	0.6944	-0.2021	0.079*
C54	0.9242 (2)	0.6776 (2)	-0.15837 (15)	0.0587 (7)
H54	0.9482	0.6560	-0.1989	0.070*
C55	0.97882 (19)	0.6872 (2)	-0.09687 (15)	0.0576 (7)
H55	1.0404	0.6735	-0.0956	0.069*
C56	0.94248 (17)	0.7172 (2)	-0.03655 (14)	0.0486 (6)
H56	0.9801	0.7226	0.0052	0.058*
C61	0.80914 (16)	0.91750 (19)	0.04859 (12)	0.0429 (5)
C62	0.7576 (2)	0.9697 (2)	0.09473 (15)	0.0605 (7)
H62	0.7245	0.9294	0.1238	0.073*
C63	0.7549 (3)	1.0802 (2)	0.09799 (17)	0.0766 (9)
H63	0.7194	1.1138	0.1286	0.092*
C64	0.8040 (3)	1.1398 (3)	0.05644 (19)	0.0857 (11)
H64	0.8020	1.2142	0.0584	0.103*
C65	0.8562 (3)	1.0901 (3)	0.01195 (19)	0.0883 (11)
H65	0.8905	1.1310	-0.0158	0.106*
C66	0.8586 (2)	0.9793 (2)	0.00755 (15)	0.0623 (7)
H66	0.8941	0.9467	-0.0234	0.075*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cu1	0.03296 (15)	0.04514 (17)	0.03791 (16)	-0.00235 (12)	0.00491 (11)	0.00367 (12)
O1	0.104 (5)	0.132 (6)	0.074 (4)	-0.054 (4)	-0.025 (3)	0.031 (3)
Cl1	0.0553 (4)	0.0380 (3)	0.0546 (4)	-0.0016 (3)	0.0157 (3)	-0.0045 (3)
S1	0.0600 (4)	0.0470 (4)	0.0581 (4)	-0.0062 (3)	0.0267 (3)	-0.0090 (3)
P1	0.0326 (3)	0.0408 (3)	0.0345 (3)	-0.0015 (2)	0.0031 (2)	0.0009 (2)
P2	0.0323 (3)	0.0405 (3)	0.0359 (3)	-0.0013 (2)	0.0036 (2)	0.0022 (2)
N1	0.0616 (16)	0.0768 (18)	0.111 (2)	0.0057 (14)	0.0397 (16)	-0.0275 (16)
N2	0.0507 (13)	0.0628 (15)	0.0868 (17)	-0.0123 (11)	0.0347 (12)	-0.0253 (13)
N3	0.068 (2)	0.099 (2)	0.156 (3)	-0.0340 (19)	0.055 (2)	-0.046 (3)
C1	0.0513 (15)	0.0576 (16)	0.0480 (14)	0.0054 (12)	0.0178 (12)	-0.0065 (12)
C11	0.0350 (11)	0.0408 (12)	0.0339 (11)	0.0038 (9)	0.0047 (9)	0.0038 (9)
C12	0.0510 (14)	0.0423 (13)	0.0475 (14)	-0.0025 (11)	0.0055 (11)	-0.0006 (11)
C13	0.0751 (19)	0.0470 (15)	0.0499 (15)	0.0075 (14)	0.0155 (13)	-0.0055 (12)
C14	0.0648 (18)	0.0579 (17)	0.0574 (16)	0.0172 (14)	0.0276 (14)	0.0085 (13)
C15	0.0424 (14)	0.0652 (18)	0.0700 (18)	-0.0009 (12)	0.0180 (13)	0.0062 (15)
C16	0.0407 (13)	0.0538 (15)	0.0491 (14)	-0.0042 (11)	0.0061 (11)	-0.0041 (11)
C21	0.0315 (11)	0.0457 (13)	0.0484 (13)	-0.0006 (9)	0.0057 (10)	-0.0063 (10)
C22	0.0693 (18)	0.0483 (15)	0.0594 (17)	-0.0014 (13)	0.0173 (14)	-0.0006 (13)
C23	0.074 (2)	0.0436 (16)	0.102 (3)	-0.0007 (14)	0.0286 (19)	-0.0052 (16)
C24	0.0532 (18)	0.0538 (18)	0.126 (3)	-0.0037 (14)	0.0035 (19)	-0.031 (2)
C25	0.086 (2)	0.080 (2)	0.088 (3)	0.009 (2)	-0.030 (2)	-0.039 (2)
C26	0.0679 (19)	0.0608 (17)	0.0601 (17)	0.0080 (15)	-0.0159 (14)	-0.0101 (14)
C31	0.0436 (13)	0.0444 (13)	0.0368 (12)	0.0000 (10)	0.0006 (10)	0.0007 (10)
C32	0.0508 (15)	0.0623 (17)	0.0540 (15)	0.0066 (13)	0.0028 (12)	0.0120 (13)
C33	0.0650 (19)	0.072 (2)	0.0642 (18)	0.0212 (16)	-0.0008 (15)	0.0146 (15)
C34	0.096 (3)	0.0641 (19)	0.0569 (18)	0.0155 (18)	0.0017 (17)	0.0196 (15)

C35	0.081 (2)	0.085 (2)	0.0607 (19)	-0.0052 (18)	0.0165 (16)	0.0279 (17)
C36	0.0543 (16)	0.0706 (18)	0.0470 (15)	-0.0003 (14)	0.0062 (12)	0.0146 (13)
C41	0.0337 (11)	0.0549 (15)	0.0376 (12)	-0.0034 (10)	0.0021 (9)	-0.0042 (10)
C42	0.0473 (15)	0.0670 (18)	0.0710 (19)	-0.0138 (13)	-0.0052 (13)	0.0002 (15)
C43	0.062 (2)	0.096 (3)	0.106 (3)	-0.037 (2)	-0.0070 (19)	0.006 (2)
C44	0.0389 (17)	0.141 (4)	0.092 (3)	-0.019 (2)	-0.0038 (16)	-0.006 (2)
C45	0.0416 (16)	0.110 (3)	0.080 (2)	0.0091 (17)	-0.0061 (15)	0.003 (2)
C46	0.0427 (15)	0.0740 (19)	0.0654 (18)	0.0024 (13)	0.0007 (13)	0.0113 (15)
C51	0.0401 (12)	0.0391 (12)	0.0381 (12)	-0.0010 (10)	0.0075 (9)	0.0010 (9)
C52	0.0441 (14)	0.086 (2)	0.0431 (14)	0.0032 (14)	0.0008 (11)	-0.0040 (14)
C53	0.0654 (19)	0.093 (2)	0.0395 (14)	-0.0050 (16)	0.0033 (13)	-0.0071 (14)
C54	0.0719 (19)	0.0564 (17)	0.0520 (16)	-0.0060 (14)	0.0265 (14)	-0.0076 (12)
C55	0.0478 (15)	0.0610 (17)	0.0672 (18)	0.0020 (13)	0.0206 (13)	-0.0021 (14)
C56	0.0417 (13)	0.0567 (16)	0.0477 (14)	0.0002 (11)	0.0063 (11)	-0.0009 (11)
C61	0.0479 (13)	0.0395 (12)	0.0401 (12)	-0.0013 (10)	-0.0011 (10)	0.0033 (10)
C62	0.077 (2)	0.0506 (16)	0.0557 (16)	0.0073 (14)	0.0170 (14)	0.0043 (13)
C63	0.111 (3)	0.0526 (18)	0.067 (2)	0.0171 (18)	0.0143 (19)	-0.0046 (15)
C64	0.140 (3)	0.0417 (17)	0.073 (2)	-0.002 (2)	0.003 (2)	-0.0008 (16)
C65	0.136 (3)	0.0539 (19)	0.079 (2)	-0.027 (2)	0.030 (2)	0.0082 (17)
C66	0.079 (2)	0.0498 (16)	0.0603 (17)	-0.0128 (14)	0.0192 (15)	0.0005 (13)

Geometric parameters (Å, °)

Cu1—P1	2.2714 (6)	C31—C36	1.387 (3)
Cu1—P2	2.2872 (6)	C32—C33	1.385 (4)
Cu1—S1	2.3841 (7)	C32—H32	0.9300
Cu1—C11	2.4110 (7)	C33—C34	1.369 (5)
O1—H1C	0.82 (2)	C33—H33	0.9300
O1—H1D	0.82 (2)	C34—C35	1.359 (5)
S1—C1	1.697 (3)	C34—H34	0.9300
P1—C21	1.823 (2)	C35—C36	1.386 (4)
P1—C11	1.831 (2)	C35—H35	0.9300
P1—C31	1.836 (2)	C36—H36	0.9300
P2—C61	1.828 (2)	C41—C42	1.377 (4)
P2—C51	1.828 (2)	C41—C46	1.385 (4)
P2—C41	1.834 (2)	C42—C43	1.386 (4)
N1—C1	1.316 (3)	C42—H42	0.9300
N1—H1A	0.8600	C43—C44	1.366 (5)
N1—H1B	0.8600	C43—H43	0.9300
N2—C1	1.319 (3)	C44—C45	1.362 (5)
N2—N3	1.406 (4)	C44—H44	0.9300
N2—H2	0.8600	C45—C46	1.383 (4)
N3—H3A	0.879 (19)	C45—H45	0.9300
N3—H3B	0.851 (19)	C46—H46	0.9300
C11—C16	1.387 (3)	C51—C56	1.383 (3)
C11—C12	1.389 (3)	C51—C52	1.383 (3)
C12—C13	1.382 (4)	C52—C53	1.384 (4)
C12—H12	0.9300	C52—H52	0.9300

C13—C14	1.365 (4)	C53—C54	1.359 (4)
C13—H13	0.9300	C53—H53	0.9300
C14—C15	1.373 (4)	C54—C55	1.366 (4)
C14—H14	0.9300	C54—H54	0.9300
C15—C16	1.382 (4)	C55—C56	1.382 (4)
C15—H15	0.9300	C55—H55	0.9300
C16—H16	0.9300	C56—H56	0.9300
C21—C26	1.372 (4)	C61—C66	1.372 (4)
C21—C22	1.388 (4)	C61—C62	1.394 (4)
C22—C23	1.373 (4)	C62—C63	1.382 (4)
C22—H22	0.9300	C62—H62	0.9300
C23—C24	1.367 (5)	C63—C64	1.360 (5)
C23—H23	0.9300	C63—H63	0.9300
C24—C25	1.362 (5)	C64—C65	1.364 (5)
C24—H24	0.9300	C64—H64	0.9300
C25—C26	1.395 (5)	C65—C66	1.386 (4)
C25—H25	0.9300	C65—H65	0.9300
C26—H26	0.9300	C66—H66	0.9300
C31—C32	1.383 (3)		
P1—Cu1—P2	129.47 (2)	C36—C31—P1	119.14 (18)
P1—Cu1—S1	100.02 (3)	C31—C32—C33	120.8 (3)
P2—Cu1—S1	103.44 (3)	C31—C32—H32	119.6
P1—Cu1—Cl1	102.44 (2)	C33—C32—H32	119.6
P2—Cu1—Cl1	110.26 (2)	C34—C33—C32	120.2 (3)
S1—Cu1—Cl1	110.06 (2)	C34—C33—H33	119.9
H1C—O1—H1D	129 (10)	C32—C33—H33	119.9
C1—S1—Cu1	108.44 (9)	C35—C34—C33	119.6 (3)
C21—P1—C11	102.72 (10)	C35—C34—H34	120.2
C21—P1—C31	104.98 (11)	C33—C34—H34	120.2
C11—P1—C31	101.42 (10)	C34—C35—C36	121.1 (3)
C21—P1—Cu1	111.56 (8)	C34—C35—H35	119.4
C11—P1—Cu1	119.40 (7)	C36—C35—H35	119.4
C31—P1—Cu1	115.04 (8)	C35—C36—C31	120.0 (3)
C61—P2—C51	103.67 (11)	C35—C36—H36	120.0
C61—P2—C41	101.63 (11)	C31—C36—H36	120.0
C51—P2—C41	103.36 (10)	C42—C41—C46	118.6 (2)
C61—P2—Cu1	114.33 (8)	C42—C41—P2	118.0 (2)
C51—P2—Cu1	117.63 (8)	C46—C41—P2	123.4 (2)
C41—P2—Cu1	114.30 (8)	C41—C42—C43	120.4 (3)
C1—N1—H1A	120.0	C41—C42—H42	119.8
C1—N1—H1B	120.0	C43—C42—H42	119.8
H1A—N1—H1B	120.0	C44—C43—C42	120.3 (3)
C1—N2—N3	119.6 (2)	C44—C43—H43	119.9
C1—N2—H2	120.2	C42—C43—H43	119.9
N3—N2—H2	120.2	C45—C44—C43	119.9 (3)
N2—N3—H3A	104 (3)	C45—C44—H44	120.0
N2—N3—H3B	111 (3)	C43—C44—H44	120.0

H3A—N3—H3B	102 (4)	C44—C45—C46	120.3 (3)
N1—C1—N2	117.8 (3)	C44—C45—H45	119.8
N1—C1—S1	121.9 (2)	C46—C45—H45	119.8
N2—C1—S1	120.19 (19)	C45—C46—C41	120.4 (3)
C16—C11—C12	119.0 (2)	C45—C46—H46	119.8
C16—C11—P1	124.28 (18)	C41—C46—H46	119.8
C12—C11—P1	116.67 (17)	C56—C51—C52	117.7 (2)
C13—C12—C11	120.2 (2)	C56—C51—P2	118.30 (18)
C13—C12—H12	119.9	C52—C51—P2	123.93 (19)
C11—C12—H12	119.9	C51—C52—C53	120.5 (3)
C14—C13—C12	120.2 (3)	C51—C52—H52	119.8
C14—C13—H13	119.9	C53—C52—H52	119.8
C12—C13—H13	119.9	C54—C53—C52	120.9 (3)
C13—C14—C15	120.3 (2)	C54—C53—H53	119.6
C13—C14—H14	119.8	C52—C53—H53	119.6
C15—C14—H14	119.8	C53—C54—C55	119.6 (3)
C14—C15—C16	120.2 (3)	C53—C54—H54	120.2
C14—C15—H15	119.9	C55—C54—H54	120.2
C16—C15—H15	119.9	C54—C55—C56	119.9 (3)
C15—C16—C11	120.1 (2)	C54—C55—H55	120.0
C15—C16—H16	120.0	C56—C55—H55	120.0
C11—C16—H16	120.0	C55—C56—C51	121.4 (2)
C26—C21—C22	118.1 (2)	C55—C56—H56	119.3
C26—C21—P1	124.3 (2)	C51—C56—H56	119.3
C22—C21—P1	117.53 (19)	C66—C61—C62	117.9 (2)
C23—C22—C21	121.6 (3)	C66—C61—P2	123.5 (2)
C23—C22—H22	119.2	C62—C61—P2	118.63 (19)
C21—C22—H22	119.2	C63—C62—C61	121.1 (3)
C24—C23—C22	119.5 (3)	C63—C62—H62	119.5
C24—C23—H23	120.2	C61—C62—H62	119.5
C22—C23—H23	120.2	C64—C63—C62	120.0 (3)
C25—C24—C23	120.0 (3)	C64—C63—H63	120.0
C25—C24—H24	120.0	C62—C63—H63	120.0
C23—C24—H24	120.0	C63—C64—C65	119.7 (3)
C24—C25—C26	120.7 (3)	C63—C64—H64	120.1
C24—C25—H25	119.7	C65—C64—H64	120.1
C26—C25—H25	119.7	C64—C65—C66	120.9 (3)
C21—C26—C25	119.9 (3)	C64—C65—H65	119.6
C21—C26—H26	120.0	C66—C65—H65	119.6
C25—C26—H26	120.0	C61—C66—C65	120.4 (3)
C32—C31—C36	118.3 (2)	C61—C66—H66	119.8
C32—C31—P1	122.49 (19)	C65—C66—H66	119.8
N3—N2—C1—N1	0.5 (5)	C32—C31—C36—C35	-0.1 (4)
N3—N2—C1—S1	177.1 (3)	P1—C31—C36—C35	177.8 (2)
Cu1—S1—C1—N1	-154.9 (2)	C61—P2—C41—C42	-156.0 (2)
Cu1—S1—C1—N2	28.6 (3)	C51—P2—C41—C42	96.8 (2)
C21—P1—C11—C16	18.2 (2)	Cu1—P2—C41—C42	-32.3 (2)

C31—P1—C11—C16	-90.2 (2)	C61—P2—C41—C46	25.0 (3)
Cu1—P1—C11—C16	142.20 (18)	C51—P2—C41—C46	-82.3 (2)
C21—P1—C11—C12	-163.36 (18)	Cu1—P2—C41—C46	148.6 (2)
C31—P1—C11—C12	88.22 (19)	C46—C41—C42—C43	-0.3 (4)
Cu1—P1—C11—C12	-39.3 (2)	P2—C41—C42—C43	-179.4 (3)
C16—C11—C12—C13	-0.1 (4)	C41—C42—C43—C44	0.0 (6)
P1—C11—C12—C13	-178.7 (2)	C42—C43—C44—C45	0.5 (6)
C11—C12—C13—C14	0.0 (4)	C43—C44—C45—C46	-0.6 (6)
C12—C13—C14—C15	0.2 (4)	C44—C45—C46—C41	0.3 (5)
C13—C14—C15—C16	-0.3 (4)	C42—C41—C46—C45	0.1 (4)
C14—C15—C16—C11	0.2 (4)	P2—C41—C46—C45	179.2 (2)
C12—C11—C16—C15	0.0 (4)	C61—P2—C51—C56	95.5 (2)
P1—C11—C16—C15	178.4 (2)	C41—P2—C51—C56	-158.8 (2)
C11—P1—C21—C26	-114.0 (2)	Cu1—P2—C51—C56	-31.8 (2)
C31—P1—C21—C26	-8.3 (3)	C61—P2—C51—C52	-87.8 (2)
Cu1—P1—C21—C26	117.0 (2)	C41—P2—C51—C52	17.9 (3)
C11—P1—C21—C22	67.8 (2)	Cu1—P2—C51—C52	144.9 (2)
C31—P1—C21—C22	173.51 (19)	C56—C51—C52—C53	0.7 (4)
Cu1—P1—C21—C22	-61.3 (2)	P2—C51—C52—C53	-176.0 (2)
C26—C21—C22—C23	3.4 (4)	C51—C52—C53—C54	-0.1 (5)
P1—C21—C22—C23	-178.3 (2)	C52—C53—C54—C55	-0.9 (5)
C21—C22—C23—C24	-1.3 (5)	C53—C54—C55—C56	1.4 (4)
C22—C23—C24—C25	-1.3 (5)	C54—C55—C56—C51	-0.8 (4)
C23—C24—C25—C26	1.7 (5)	C52—C51—C56—C55	-0.3 (4)
C22—C21—C26—C25	-2.9 (4)	P2—C51—C56—C55	176.6 (2)
P1—C21—C26—C25	178.8 (2)	C51—P2—C61—C66	-18.1 (3)
C24—C25—C26—C21	0.5 (5)	C41—P2—C61—C66	-125.2 (2)
C21—P1—C31—C32	-77.2 (2)	Cu1—P2—C61—C66	111.2 (2)
C11—P1—C31—C32	29.5 (2)	C51—P2—C61—C62	162.3 (2)
Cu1—P1—C31—C32	159.82 (19)	C41—P2—C61—C62	55.3 (2)
C21—P1—C31—C36	104.9 (2)	Cu1—P2—C61—C62	-68.4 (2)
C11—P1—C31—C36	-148.4 (2)	C66—C61—C62—C63	1.4 (4)
Cu1—P1—C31—C36	-18.1 (2)	P2—C61—C62—C63	-179.1 (2)
C36—C31—C32—C33	-0.9 (4)	C61—C62—C63—C64	-0.9 (5)
P1—C31—C32—C33	-178.8 (2)	C62—C63—C64—C65	-0.3 (6)
C31—C32—C33—C34	1.0 (5)	C63—C64—C65—C66	1.2 (6)
C32—C33—C34—C35	0.0 (5)	C62—C61—C66—C65	-0.6 (4)
C33—C34—C35—C36	-1.1 (5)	P2—C61—C66—C65	179.9 (3)
C34—C35—C36—C31	1.1 (5)	C64—C65—C66—C61	-0.7 (6)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1C \cdots C11	0.82 (2)	2.37 (2)	3.186 (5)	179 (12)
O1—H1D \cdots S1 ⁱ	0.82 (2)	2.70 (10)	3.218 (5)	123 (9)
N1—H1B \cdots C11 ⁱⁱ	0.86	2.48	3.302 (3)	161
N1—H1A \cdots N3	0.86	2.27	2.628 (5)	105

N2—H2···C11	0.86	2.35	3.202 (2)	170
C42—H42···C11	0.93	2.72	3.626 (3)	164

Symmetry codes: (i) $-x+3/2, y-1/2, -z+1/2$; (ii) $-x+3/2, y+1/2, -z+1/2$.