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[*N*,*N*-Bis(2-hydroxyethyl)dithiocarbamato-*S*]bis(triphenylphosphine-*P*)phosphine solvate

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In the title compound, $[Cu(C_5H_{10}NO_2S_2)(C_{18}H_{15}P)_2]\cdot C_{18}H_{15}P$, the Cu atom is in a distorted tetrahedral coordination, with two triphenylphosphine P atoms and two S atoms from an *N*,*N*-bis(2-hydroxyethyl)dithiocarbamate ligand occupying the vertices. The crystal structure is characterized by alternate layers of complex and triphenylphosphine molecules.

Comment

Group 12 metals form a wide range of tertiary phosphine (R_3P) complexes of the general stoichiometry $L_nMX(L = R_3P, n = 1-4;$ Attar, Alcock, Frye *et al.*, 1991; Attar, Alcock, Bowmaker *et al.*, 1991; Alyea *et al.*, 1992). The value of *n* and the net electronic charge on the complex are somewhat dependent upon the coordinating ability of *X*. Also, the transition metal dithiolates containing MS_4 chromophores show interesting variations in their reactions with Lewis bases (Cordes & Draganjac, 1988; Venkatachalam *et al.*, 1996).



The asymmetric unit of the title structure, (I), contains one molecule of $[Cu\{(C_2H_4OH)_2S_2CN\}(PPh_3)_2]$ together with a molecule of uncoordinated triphenylphosphine. The Cu1 atom is in a distorted tetrahedral environment, with the P atoms of the triphenylphosphine ligands and the S atoms of the bidentate dithiocarbamate ligand occupying the vertices. The angles around Cu1 range between 74.77 (4) (S1-Cu1-S2) and 124.51 (4)° (P1-Cu1-P2). The small S1-Cu1-S2 bond angle is due to the restricted bite angle of the ligand. The P1-Cu1-P2 bond angle is comparable with values reported for other bis(triphenylphosphine)copper(I) compounds (Kara-

glannidis et al., 1989; Long et al., 1996). The Cu-P distances are comparable to those found in $(PPh_3)_2Cu(C_2H_4N_2S_2)Br$ (Long et al., 1996). The bond lengths of Cu-S1 and S2-C37 are longer than those of Cu-S2 and S1-C37, respectively. One of the S–C bond lengths (S1-C37) is close to the C=S distance of 1.69 Å, while S2–C37 shows a value intermediate between single- and double-bond distances, indicating partial double-bond character. The longer Cu-S bond length (Cu1-S1) lies adjacent to C-S bond which is closer to the C=Sbond distance. The same characteristic is found in the structure of $[Ni(pipdtc)(PPh_3)_2]ClO_4(PPh_3) \cdot H_2O$ (pipdtc is piperidinedithiocarbamate; Venkatachalam et al., 1996), but different from the complex of $[Au(S_2CPh)(PPh_3)_2]$ (Lanfredi et al., 1992), where the shorter Au-S bond lies adjacent to the C-S bond length which is closer to the C=S bond distance. The four-membered CuS₂C is nearly planar with normal bond lengths. The bond length of C37–N1 [1.360(4) Å] has double-bond character, but slightly longer than those observed in a related structure (Jian et al., 1999). All the phenyl rings show normal bond parameters. The P-C bond distances are also normal. The C-P-C angles deviate appreciably from the ideal tetrahedral angle of 109.5°. The crystal structure is characterized by alternate layers of the complexes and the triphenylphosphine ligands which lie parallel to the bc plane. The structure is stabilized by C- $H \cdot \cdot \pi$ and van der Waals interactions (Table 2).

Experimental

Sodium *N*,*N*-bis(2-hydroxyethyl)dithiocarbamate, Na[S₂CN(C₂H₄-OH)₂], was dissolved in H₂O. CuCl₂ was then added at room temperature with stirring. The brown reaction mixture was filtered and the filtrate was washed with water and dried overnight *in vacuo*. The dried brown solid was dissolved in EtOH and triphenyl-phosphine was added at 253–263 K with stirring and the resulting solution refluxed for 4–5 h. Upon cooling, the colourless solution obtained was filtered and left to evaporate. After a few days, colourless solids were separated out and recrystallized from EtOH/ CHCl₃. Single crystals suitable for X-ray analysis were obtained by slow evaporation at room temperature of the EtOH solvent.

Crystal data

$[Cu(C_5H_{10}NO_2S_2)(C_{18}H_{15}P)_2]$	Z = 2
C ₁₈ H ₁₅ P	$D_x = 1.291 \text{ Mg m}^{-3}$
$M_r = 1030.61$	Mo $K\alpha$ radiation
Triclinic, $P\overline{1}$	Cell parameters from 7234
a = 12.2892 (4) Å	reflections
b = 13.1612(5) Å	$\theta = 1.11 - 28.34^{\circ}$
c = 19.3696 (7) Å	$\mu = 0.624 \text{ mm}^{-1}$
$\alpha = 95.512 \ (1)^{\circ}$	T = 293 (2) K
$\beta = 103.177 \ (1)^{\circ}$	Slab, colourless
$\gamma = 116.801 \ (1)^{\circ}$	$0.40 \times 0.30 \times 0.14 \text{ mm}$
$V = 2650.47 (16) \text{ Å}^3$	
Data collection	
Siemens SMART CCD area-	11 791 independent reflections
detector diffractometer	5663 reflections with $I > 2\sigma(I)$
ω scans	$R_{\rm int} = 0.065$
Absorption correction: empirical	$\theta_{\rm max} = 27.5^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -15 \rightarrow 14$
$T_{\min} = 0.788, \ T_{\max} = 0.918$	$k = -17 \rightarrow 17$
18 118 measured reflections	$l = -18 \rightarrow 25$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.057$	$w = 1/[\sigma^2(F_o^2) + (0.0271P)^2]$
$wR(F^2) = 0.139$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.86	$(\Delta/\sigma)_{\rm max} < 0.001$
11791 reflections	$\Delta \rho_{\rm max} = 0.44 \ {\rm e} \ {\rm \AA}^{-3}$
624 parameters	$\Delta \rho_{\rm min} = -0.76 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

Cu1-P2	2.2595 (9)	\$1-C37	1.702 (4)	
Cu1-P1	2.2849 (10)	S2-C37	1.710 (4)	
Cu1-S2	2.3946 (11)	N1-C37	1.360 (5)	
Cu1-S1	2.4287 (10)			
P2-Cu1-P1	124.51 (4)	P2-Cu1-S1	122.04 (4)	
P2-Cu1-S2	112.35 (4)	P1-Cu1-S1	102.49 (4)	
P1-Cu1-S2	109.85 (4)	S2-Cu1-S1	74.77 (4)	
-				

Table 2

Hydrogen-bonding geometry (Å, °).

Cg1 and Cg2 are the centroids of the C19-C24 and C7-C12 rings, respectively.

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} C12-H12A\cdots Cg1\\ C44-H44A\cdots Cg2^{i} \end{array}$	0.93	2.84	3.633	144
	0.93	2.97	3.821	153

Symmetry code: (i) x - 1, y - 1, z.

After checking their presence in the difference map, all H atoms were geometrically fixed and allowed to ride on their attached atoms. The O1 atom in one of the hydroxyethyl groups is disordered and was refined with an occupancy ratio of of 78:22.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 1990).

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