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# Soo Yei Ho and Edward R. T. Tiekink\*

Department of Chemistry, National University of Singapore, Singapore 117543

Correspondence e-mail: chmtert@nus.edu.sg

#### Key indicators

Single-crystal X-ray study T = 223 KMean  $\sigma(C-C) = 0.008 \text{ Å}$  R factor = 0.042 wR factor = 0.115 Data-to-parameter ratio = 28.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

© 2001 International Union of Crystallography Printed in Great Britain – all rights reserved The Au atom in the title compound,  $(c-C_6H_{11})_3PAu(S_2CNEt_2)$ or  $[Au(C_{18}H_{33}P)(C_5H_{10}NS_2)]$ , exists in a linear geometry so that Au-S is 2.3340 (11), Au-P is 2.2599 (10) Å and the angle at gold is 171.61 (4)°. The distortion from linearity may be traced to the close approach of the non-coordinating S atom which is separated from the Au atom by 3.0859 (13) Å.

Comment

In connection with previous studies on compounds of the general formula  $R_3$ PAu(S<sub>2</sub>COR'), R and R' = alkyl (Siasios & Tiekink, 1993*a*,*b*), which demonstrated interesting flexibility in the mode of coordination of the thiolate ligands, the structure of the title compound,  $(c-C_6H_{11})_3$ PAu(S<sub>2</sub>CNEt<sub>2</sub>), (I), was investigated.



The Au atom in (I) exists in an approximately linear geometry defined by the S atom of the monodentate dithiocarbamate ligand and the P atom of the phosphine ligand (Fig. 1 and Table 1). There is a significant distortion from the ideal 180° angle for P-Au-S that may be correlated with the close approach of atom S2 to the Au centre. Thus, the Au···S2 separation is 3.0859(13) Å, a distance that is well within the sum of the van der Waals radii, 3.50 Å, for these atoms (Bondi, 1964). Although there are a number of related gold-phosphine-dithiocarbamate structures in the literature, there is, surprisingly, only one directly comparable mononuclear structure available for comparison, namely that of the triphenylphosphine analogue (Wijnhoven et al., 1972). The respective Au-S bond distance in the present structure and that in  $(C_6H_5)_3PAu(S_2CNEt_2)$  of 2.3340 (11) and 2.338 (3) Å, respectively, are equal within experimental error. Likewise, the Au-P bond distances of 2.2599 (10) and 2.251 (3) Å, respectively, are also equivalent. However, the distortion from linearity in the cyclohexyl derivative is somewhat greater as the P-Au-S angle in the triphenylphosphine analogue was determined to be 175.7  $(1)^{\circ}$ . These results show that in the uncongested Au atom environment, bulky phosphines may be accommodated without significant deviations in the geometric parameters about Au. A similar conclusion has been reported

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Figure 1

The molecular structure and crystallographic numbering scheme for (I); displacement ellipsoids are shown at the 50% probability level (Johnson, 1976).

previously for the closely related dithiocarbonate structures, *i.e.*  $R_3$ PAu(S<sub>2</sub>COR') (Siasios & Tiekink, 1993*a*,*b*). The crystal structure is stabilized by hydrophobic interactions.

# **Experimental**

To a dichloromethane solution (4 ml) of  $(C_6H_{11})_3$ PAuCl (0.20 g, 0.39 mmol) was added NaS<sub>2</sub>CNEt<sub>2</sub> (67 mg, 0.39 mmol). The colourless solution immediately turned yellow, indicating the formation of the product, and was stirred for 2 h. The yellow solution was filtered through Celite and concentrated to approximately 1 ml to yield the product. The product was recrystallized by the vapour diffusion of hexane into a dichloromethane solution to yield X-ray quality yellow crystals of (I). <sup>1</sup>H NMR (CDCl<sub>3</sub>):  $\delta$  3.91 (*q*, 4 H), 2.02–1.54 (*m*, 33 H), 1.30 (*t*, 6 H). <sup>13</sup>C{<sup>1</sup>H} NMR (CDCl<sub>3</sub>):  $\delta$  216.3 (*s*, CS<sub>2</sub>), 48.9 (*s*, ethyl-CH<sub>2</sub>), 33.5–25.9 (*m*, P(C<sub>6</sub>H<sub>11</sub>)<sub>3</sub>), 12.1 (*s*, CH<sub>3</sub>). <sup>31</sup>P{<sup>1</sup>H} (CDCl<sub>3</sub>):  $\delta$  55.3 p.p.m. ESI–MS: *m*/z 626 (*M*<sup>+</sup>). IR (KBr): 1476  $\nu$ (C–N), 1083 and 991  $\nu$ (C–S) cm<sup>-1</sup>.

### Crystal data

[Au(C <sub>5</sub> H <sub>10</sub> NS <sub>2</sub> )(C <sub>18</sub> H <sub>33</sub> P)]
$M_r = 625.64$
Triclinic, P1
a = 10.1697 (5)  Å
b = 11.7231 (6) Å
c = 12.3281 (6) Å
$\alpha = 71.710 \ (1)^{\circ}$
$\beta = 77.917 (1)^{\circ}$
$\gamma = 65.803 (1)^{\circ}$
$V = 1267.36 (11) \text{ Å}^3$
. ,

#### Data collection

Bruker AXS SMART CCD
diffractometer
$\omega$ scans
Absorption correction: empirical
(SADABS; Bruker, 2000)
$T_{\min} = 0.112, \ T_{\max} = 0.297$
10496 measured reflections

Z = 2  $D_x = 1.639 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 10496 reflections  $\theta = 1.8 - 30.0^{\circ}$   $\mu = 6.04 \text{ mm}^{-1}$  T = 223 (2) KBlock, pale yellow  $0.39 \times 0.31 \times 0.21 \text{ mm}$ 

	7203 independent reflections
	6682 reflections with $I > 2\sigma(I)$
	$R_{\rm int} = 0.036$
1	$\theta_{\rm max} = 30.0^{\circ}$
	$h = -10 \rightarrow 14$
	$k = -10 \rightarrow 16$
	$l = -17 \rightarrow 17$

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0897P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	where $P = (F_o^2 + 2F_c^2)/3$
$vR(F^2) = 0.115$	$(\Delta/\sigma)_{\rm max} = 0.001$
S = 1.05	$\Delta \rho_{\rm max} = 2.08 \text{ e } \text{\AA}^{-3}$
7203 reflections	$\Delta \rho_{\rm min} = -2.91 \text{ e } \text{\AA}^{-3}$
254 parameters	
Hatom parameters constrained	

## Table 1

Selected geometric parameters (Å, °).

Au-P1	2.2599 (10)	P1-C12	1.858 (5)
Au-S1	2.3340 (11)	P1-C18	1.874 (4)
S1-C1	1.745 (4)	N1-C1	1.348 (5)
S2-C1	1.699 (4)	N1-C4	1.464 (6)
P1-C6	1.836 (4)	N1-C2	1.464 (6)
P1-Au-S1	171.61 (4)	C18–P1–Au	112.27 (15)
C1-S1-Au	98.38 (14)	C1-N1-C4	121.5 (4)
C6-P1-C12	105.79 (19)	C1-N1-C2	121.2 (4)
C6-P1-C18	106.16 (19)	C4-N1-C2	116.8 (4)
C12-P1-C18	112.0 (2)	N1-C1-S2	122.2 (3)
C6-P1-Au	112.06 (13)	N1-C1-S1	117.4 (3)
C12-P1-Au	108.40 (14)	S2-C1-S1	120.5 (2)

The C-bound H atoms were placed in their geometrically calculated positions and included in the final refinement in the ridingmodel approximation with an overall displacement parameter,  $U_{\rm iso}$ , with  $1.0U_{\rm iso}$  for methine-H,  $1.25U_{\rm iso}$  for methylene-H and  $1.5U_{\rm iso}$  for CH<sub>3</sub>. The residual electron-density peak is located in the vicinity of the Au atom.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SHELXTL* (Bruker, 2000); program(s) used to solve structure: *DIRDIF92 PATTY* (Beurskens *et al.*, 1992); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP*II (Johnson, 1976); software used to prepare material for publication: *SHELXTL*.

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