## metal-organic papers

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### Normah Awang, Ibrahim Baba, M. Sukeri M. Yusof and Bohari M. Yamin\*

School of Chemical Sciences and Food Technology, Universiti Kebangsaan Malaysia, 43600 Bangi, Selangor, Malaysia

Correspondence e-mail: bohari@pkrisc.cc.ukm.my

#### Key indicators

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.025 wR factor = 0.071 Data-to-parameter ratio = 20.5

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

# (*N*-Cyclohexyl-*N*-ethyldithiocarbamato)triphenyltin(IV)

The title compound,  $[Sn(C_6H_5)_3[S_2CN(Et)C_6H_{11}]]$ , has a similar geometry to its methyl analogue, between tetrahedral and distorted trigonal bipyramidal but closer to the latter. The long intramolecular  $Sn \cdots S$  interaction of 2.9426 (10) Å is slightly shorter than in its methyl analogue, 3.0134 (8) Å.

### Comment

The structural dimensions of the title compound, (I), are in agreement with the analogues (*N*-cyclohexyl-*N*-methyldithiocarbamato)triphenyltin(IV) (Awang *et al.*, 2003), (*N*-butyl-*N*-methyldithiocarbamato)triphenyltin (Kana *et al.*, 2001) and (diethyldithiocarbamato)triphenyltin(IV) (Lindley & Carr, 1974). The presence of an ethyl group results in only a slightly shorter weak intermolecular interaction [Sn1 $\cdots$ S2 = 2.9426 (10) Å] compared with 3.0134 (8) Å in *N*-cyclohexyl-*N*methyldithiocarbamato)triphenyltin(IV). The geometry of the central Sn atom is also between tetrahedral and trigonal bipyramidal, but closer to the latter with C15–Sn1–C22, C15–Sn1–S1 and C22–Sn1–S1 angles of 114.20 (8), 117.41 (6) and 118.88 (5)°, respectively, in the equatorial positions; the C21–Sn1–S2 angle for the axial positions is 157.11 (6)°.



### **Experimental**

The title compound was synthesized by addition of carbon disulfide (1.8 ml, 0.03 mol) to an ethanolic solution of *N*-ethylcyclohexylamine (4 ml, 0.03 mol) and stirring for 1 h at 269 K. After stirring, triphenyltin(IV) chloride (11.6 g, 0.03 mol) solution was added and the solution mixture was further stirred for another 1 h. The white precipitate was filtered off and washed with cold ethanol and dried in vacuum. Some good quality crystals suitable for X-ray analysis were obtained by recrystallization from a 1:1 mixture of ethanol and chloroform.

Crystal data

[Sn(C<sub>6</sub>H<sub>5</sub>)<sub>3</sub>(C<sub>9</sub>H<sub>16</sub>NS<sub>2</sub>)] Z = 2 $M_{\rm m} = 552.34$  $D_r = 1.442 \text{ Mg m}^{-3}$ Triclinic, P1 Mo Ka radiation a = 10.036 (2) Å Cell parameters from 6453 reflections b = 11.435(3) Å c = 11.989 (3) Å  $\theta = 1.8\text{--}27.5^\circ$  $\mu = 1.18~\mathrm{mm}^{-1}$  $\alpha = 105.586 \ (4)^{\circ}$  $\beta = 105.443 \ (4)^{\circ}$ T = 293 (2) K  $\gamma = 90.646 (4)^{\circ}$ Block, colourless  $V = 1272.4 (5) \text{ Å}^3$  $0.55 \times 0.35 \times 0.26 \text{ mm}$ 

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Bruker SMART APEX CCD area- detector diffractometer $\omega$ scans	5793 independent reflections 5492 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.022$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -13 \rightarrow 13$
$T_{\min} = 0.562, \ T_{\max} = 0.748$	$k = -14 \rightarrow 14$
16 186 measured reflections	$l = -15 \rightarrow 15$
Refinement	
Refinement on $F^2$	$w = 1/[\sigma^2(F_a^2) + (0.0385P)^2$
$R[F^2 > 2\sigma(F^2)] = 0.025$	+ 0.5044P]
$wR(F^2) = 0.071$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.05	$(\Delta/\sigma)_{\rm max} < 0.001$
5793 reflections	$\Delta \rho_{\rm max} = 0.73 \ {\rm e} \ {\rm \AA}^{-3}$
282 parameters	$\Delta \rho_{\rm min} = -0.42 \ {\rm e} \ {\rm \AA}^{-3}$

282 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

Sn1-S1	2.4758 (7)	Sn1-C21	2.170 (2)
Sn1-S2	2.9426 (10)	S1-C9	1.750 (2)
Sn1-C15 Sn1-C22	2.144 (2) 2.148 (2)	S2-C9	1.690 (2)
C15-Sn1-C22 C15-Sn1-S1 C22-Sn1-S1 C21-Sn1-S2	114.20 (8) 117.41 (6) 118.88 (5) 157.11 (6)	C15-Sn1-C21 C22-Sn1-C21 C21-Sn1-S1	106.21 (8) 103.04 (8) 92.04 (6)

Extinction correction: SHELXL97

Extinction coefficient: 0.0054 (7)

After their location in a difference Fourier map, all H atoms were included in the refinement in geometrically calculated positions, and allowed to ride on the parent C atoms with C-H distances in the range 0.93–0.97 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$  or  $1.5U_{eq}(C)$ .

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for



The molecular structure of (I), with 50% probability displacement ellipsoids.

publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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