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## Key indicators

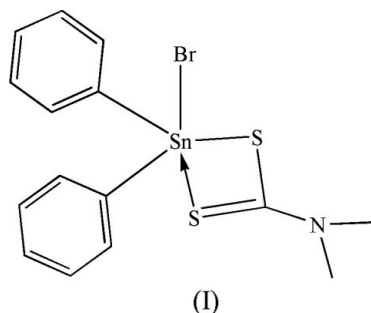
Single-crystal X-ray study  
 $T = 295$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.009$  Å  
 $R$  factor = 0.042  
 $wR$  factor = 0.094  
Data-to-parameter ratio = 19.9For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Bromo(*N,N*-dimethyldithiocarbamato- $\kappa^2\text{S,S}'$ )-  
diphenyltin(IV)The Sn atom in the title compound,  $[\text{SnBr}(\text{C}_6\text{H}_5)_2(\text{C}_3\text{H}_6\text{NS}_2)]$ ,  
adopts a distorted  $\text{SnBrC}_2\text{S}_2$  trigonal-bipyramidal environ-  
ment.

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## Comment

The structures of some chloro-diorgano-tin *N,N*-dialkyl-dithiocarbamates,  $R_2\text{Sn}(\text{S}_2\text{CNR}'_2)\text{Cl}$  have shown that the Sn atom exists in a distorted  $\text{SnClC}_2\text{S}_2$  trigonal-bipyramidal environment (Tiekink, 1992; Tiekink *et al.*, 1999; Yin *et al.*, 2001). In the bromine-containing title compound, (I), the Sn atom is also five-coordinate and possesses a distorted trigonal-bipyramidal geometry (Fig. 1 and Table 1). The equatorial plane is defined by two C atoms (C4 and C10) of the phenyl groups and the more tightly held S1 atom from the asymmetrically chelating dithiocarbamate ligand, and the axial positions are occupied by the less tightly held atom S2 and atom Br1. The tin atom lies 0.1475 (5) Å out of the  $\text{SC}_2$  trigonal plane in the direction of the Br atom. The dithiocarbamate ligand chelates in an asymmetric manner to the Sn atom with a difference in the Sn–S bond lengths of some 0.214 Å; the average Sn–S bond distance is 2.5605 (16) Å, which is similar to those found in  $\text{Ph}_2\text{Sn}(\text{S}_2\text{CNET}_2)\text{Cl}$  (Dakternieks *et al.*, 1992),  $\text{Ph}_2\text{Sn}(\text{S}_2\text{CNCy}_2)\text{Cl}$  (Basu Baul & Tiekink, 1993),  $\text{Ph}_2\text{Sn}(\text{S}_2\text{CNETCy})\text{Cl}$  (Hall & Tiekink, 1995) and  $(\text{C}_6\text{H}_5\text{CH}_2)_2\text{Sn}(\text{S}_2\text{CNMe}_2)\text{Cl}$  (Yin *et al.*, 2001). The thiocarbamate C–S bond lengths suggest that the bonding of this group is localized.



## Experimental

A solution of  $\text{NaS}_2\text{CNMe}_2 \cdot 2\text{H}_2\text{O}$  (0.36 g, 2 mmol) in 30 ml of hot acetone was added dropwise to a solution of diphenyltin dibromide (0.87 g, 2 mmol) in 30 ml of the same solvent. The reaction mixture was stirred for about an hour under reflux, and then cooled to about 273 K. The NaBr formed was removed by filtration, and the solvent was evaporated under reduced pressure. The solid obtained was recrystallized from trichloromethane–hexane (1:1 v/v) and crystals of

(I) were obtained from dichloromethane–hexane (1:1 v/v) by slow evaporation at 298 K (yield 76.3%, m.p. 406–407 K). Analysis found: C 38.12, H 3.27, N 2.83%; calculated for C<sub>15</sub>H<sub>16</sub>BrNS<sub>2</sub>Sn: C 38.09, H 3.41, N 2.96%.

Crystal data

[SnBr(C<sub>6</sub>H<sub>5</sub>)<sub>2</sub>(C<sub>3</sub>H<sub>6</sub>NS<sub>2</sub>)]  
*M<sub>r</sub>* = 473.01  
 Monoclinic, *C*2/*c*  
*a* = 17.560 (8) Å  
*b* = 9.133 (3) Å  
*c* = 21.838 (10) Å  
 β = 91.772 (6)°  
*V* = 3501 (3) Å<sup>3</sup>

*Z* = 8  
*D<sub>x</sub>* = 1.795 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 μ = 3.97 mm<sup>-1</sup>  
*T* = 295 (2) K  
 Block, colorless  
 0.14 × 0.10 × 0.10 mm

Data collection

Bruker APEX area-detector diffractometer  
 φ and ω scans  
 Absorption correction: multi-scan (SADABS; Bruker, 2002)  
*T<sub>min</sub>* = 0.606, *T<sub>max</sub>* = 0.692

9614 measured reflections  
 3634 independent reflections  
 2687 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.042  
 θ<sub>max</sub> = 26.5°

Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.042  
*wR* (*F*<sup>2</sup>) = 0.094  
*S* = 1.04  
 3634 reflections  
 183 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.029P)^2 + 3.0728P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 (Δ/σ)<sub>max</sub> = 0.001  
 Δρ<sub>max</sub> = 0.61 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = -0.64 e Å<sup>-3</sup>

Table 1 Selected geometric parameters (Å, °).

Sn1—C10	2.124 (5)	Sn1—S2	2.6674 (16)
Sn1—C4	2.132 (5)	C1—S1	1.744 (5)
Sn1—S1	2.4537 (15)	C1—S2	1.701 (6)
Sn1—Br1	2.6098 (11)		
C10—Sn1—C4	111.87 (19)	S1—Sn1—Br1	87.02 (5)
C10—Sn1—S1	113.58 (14)	C10—Sn1—S2	97.45 (15)
C4—Sn1—S1	133.21 (14)	C4—Sn1—S2	93.80 (14)
C10—Sn1—Br1	98.05 (15)	S1—Sn1—S2	70.05 (5)
C4—Sn1—Br1	97.21 (14)	Br1—Sn1—S2	156.05 (4)

H atoms were placed at calculated positions and were included in the refinement in the riding-model approximation, with C—H = 0.93 Å and *U*<sub>iso</sub>(H) = 1.2*U*<sub>eq</sub>(C) for aromatic H atoms, and C—H = 0.96 Å and *U*<sub>iso</sub>(H) = 1.5*U*<sub>eq</sub>(C) for methyl H atoms.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine

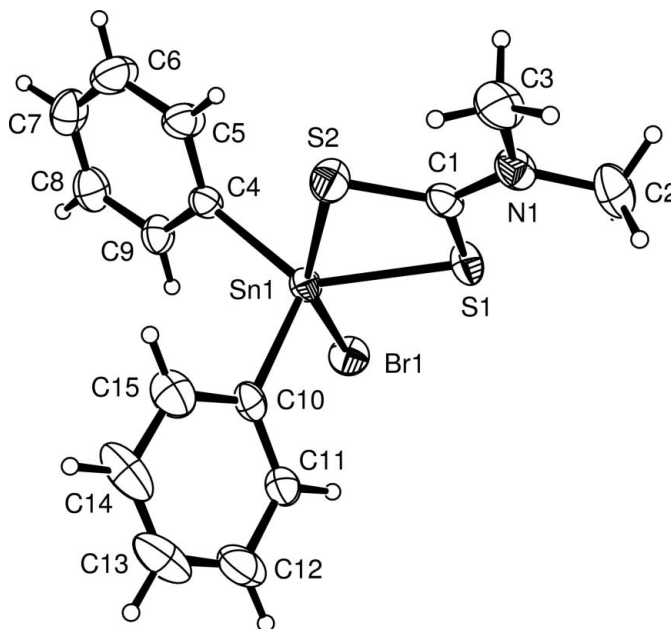


Figure 1 The structure of (I), with displacement ellipsoids drawn at the 30% probability level (arbitrary spheres for the H atoms).

structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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