metal-organic papers

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Key indicators

Single-crystal X-ray study T = 298 K Mean σ (N–C) = 0.005 Å R factor = 0.025 wR factor = 0.065 Data-to-parameter ratio = 21.9

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

trans-Tetrabromobis(dimethylamine)tin

The title compound, $[SnBr_4(C_2H_7N)_2]$, consists of separate centrosymmetric molecules with nearly octahedral geometry, in which the central Sn metal is surrounded by four Br atoms and two dimethylamine ligands. The unique Sn-N bond length is 2.244 (3) Å and the two Sn-Br lengths are 2.5867 (11) and 2.5707 (12) Å. The C-N bond lengths of the amine ligand are 1.482 (6) and 1.492 (6) Å. The unique angle in the SnBr₄ plane is 90.82 (3)°, and the axial to equatorial angles are 86.18 (10) and 85.46 (10)°.

Comment

Recently, we have utilized reactions between common germanium hydrides and SnD4 to grow epitaxic layers of $Ge_{1-x}Sn_x$ semiconductors on Si(100) substrates. The bandgaps of these materials are intermediate between those of Ge (Eg =0.066 eV) and Sn (Eg = 0.1 eV), and decrease monotonically with increasing Sn concentration in the alloy, indicating that this material will have important application in Si-based devices, such as IR photodetectors. The application of pure SnD₄ is the essential component in the preparation of these metastable and technologically important materials. SnD₄ is synthesized by reduction of SnBr₄ with LiAlD₄ and it is typically isolated as a volatile liquid, which is unstable at room temperature with respect to Sn and D₂. Nevertheless, mixtures of the compound with high-purity H₂ have the necessary stability at 295 K to be used as viable CVD sources in deposition of $\text{Ge}_{1-x}\text{Sn}_x$ (Bauer, Taraci *et al.*, 2002). Ongoing efforts to further stabilize SnD₄ by adduct formation with simple Lewis bases, such as dimethylamine, have led to the synthesis of the previously unknown complex SnBr₄[HN(CH₃)₂]₂. This compound is currently utilized to prepare the corresponding SnH₄ and SnD₄ derivatives as viable low-temperature CVD precursors of Sn.



The central Sn metal of the title compound is sixfold coordinated, with four Br atoms arranged in square-planar fashion and the two amine ligands nearly perpendicular to the $SnBr_4$ plane (Fig. 1). The Sn-Br bond distances are

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Figure 1

Displacement ellipsoid plot shown at the 50% probability level [symmetry code: (i) -x, -y, -z].

2.5867 (11) (Sn1-Br2) and 2.5707 (12) Å (Sn1-Br3). These values are significantly longer than those reported for a related SnBr₄(dioxane) complex (Bauer, Groy & Kouvetakis, 2002), which are in the range 2.5005 (7)–2.5045 (6) Å. The angles within the SnBr₄ square plane are nearly 90° [90.82 (3)° for Br2-Sn1-Br3] and the angles between the plane and the axis of the HN(CH₃)₂ ligands are 86.18 (10) (N4-Sn1-Br2) and 85.46 (10)° (N4-Sn1-Br3).

All H-atom positions were calculated based on idealized geometry. The H atoms were then allowed to ride on their bonding partners during the final stages of refinement. There is an indication of intermolecular interaction between one amine H atom and the nearest neighbor Br atom, since the Br2 \cdots H4A distance is 2.63 Å and the N4-H4A \cdots Br2 angle is 150°.

Experimental

 $SnBr_4[HN(CH_3)_2]_2$ is readily synthesized in toluene by direct combination of $SnBr_4$ with purified $HN(CH_3)_2$. This compound has been characterized by spectroscopic methods and combustion analysis. Single crystals have been grown by slowly cooling concentrated toluene solutions of the compound.

Crystal data

```
[SnBr_4(C_2H_7N)_2]
                                                      D_x = 2.674 \text{ Mg m}^{-3}
M_r = 528.50
                                                      Mo K\alpha radiation
Monoclinic, P2_1/c
                                                      Cell parameters from 3041
                                                         reflections
a = 6.592 (4) \text{ Å}
b = 12.029(7) Å
                                                      \theta = 3.0-25.1^{\circ}
                                                      \mu = 14.08 \text{ mm}^{-1}
c = 8.326(5) Å
\beta = 96.114 (11)^{\circ}
                                                      T = 298 (2) \text{ K}
V = 656.5 (7) \text{ Å}^3
                                                      Block, colorless
Z = 2
                                                      0.28 \times 0.21 \times 0.14 \text{ mm}
Data collection
Bruker SMART APEX
                                                      1161 independent reflections
   diffractometer
                                                      1028 reflections with I > 2\sigma(I)
                                                      R_{\rm int} = 0.042
(i) scans
Absorption correction: multi-scan
                                                      \theta_{\rm max} = 25.1^{\circ}
                                                      h = -7 \rightarrow 7
   (SADABS; Bruker, 2001)
   T_{\min} = 0.034, \ T_{\max} = 0.135
                                                      k = -14 \rightarrow 14
5130 measured reflections
                                                      l = -9 \rightarrow 9
Refinement
Refinement on F^2
                                                      w = 1/[\sigma^2(F_o^2) + (0.0306P)^2]
R[F^2 > 2\sigma(F^2)] = 0.026
wR(F<sup>2</sup>) = 0.065
                                                           + 0.2351P]
                                                          where P = (F_o^2 + 2F_c^2)/3
                                                      (\Delta/\sigma)_{\rm max} < 0.001
S = 1.05
                                                      \Delta \rho_{\rm max} = 0.71 \text{ e } \text{\AA}^{-3}
1161 reflections
                                                      \Delta \rho_{\rm min} = -0.73 \ {\rm e} \ {\rm \AA}^{-3}
53 parameters
                                                      Extinction correction: SHELXTL
H-atom parameters constrained
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Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

Extinction coefficient: 0.0021 (5)

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