Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma(C-C) = 0.010 \text{ Å}$ Disorder in main residue R factor = 0.042 wR factor = 0.117 Data-to-parameter ratio = 16.9

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

# Tri-*n*-butyl(5-chloro-6-hydroxypyridine-3-carboxylato)tin(IV)

The crystal structure of the title complex,  $[Sn(C_4H_9)_3(C_6H_3-CINO_3)]$ , possesses an infinite two-dimensional polymeric chain structure in the solid state, resulting from strong intermolecular N-H···O hydrogen bonds. The Sn atom is five-coordinate with a distorted trigonal bipyramidal geometry.

# Comment

The title complex, (I), consists of three butyl and one 5-chloro-6-hydroxypyridine-3-carboxylate group bonded to the Sn atom (Fig. 1) and has an infinite one-dimensional polymeric chain structure (Fig. 2), due to the Sn1 $\cdots$ O3<sup>i</sup> [symmetry code: (i) x, y + 1, z] distance of 2.537 (4)Å, compared to the van der Waals sum of 4.0Å (Vollano *et al.*, 1984).



The Sn atom has a distorted trigonal-bipyramidal geometry, with atoms O1 and O3<sup>i</sup> in axial positions and C atoms of the three butyl groups in equatorial positions. The distortions away from the ideal trigonal-bipyramidal geometry are not along the Berry coordinate (Hämäläinen & Turpeinen, 1987) and are reflected by the bond angles around the Sn atom (Table 1). The angle O1-Sn1···O3<sup>i</sup> [178.21 (19)°] deviates from linearity by  $1.87^{\circ}$ . The sum of the C15-Sn1-C7 [126.7 (3)°], C15-Sn1-C11 [115.1 (3)°] and C7-Sn1-C11 [115.1 (3)°] angles is 356.9°, which shows that these atoms are not coplanar.

The coordination geometry of the Sn atom and the molecular structure of (I) are normal (Xie *et al.*, 1991). The central Sn atom is not coordinated by the N atom of the 5-chloro-6hydroxynicotinic acid, which is different from the triphenyltin 4-picolinate (Yin *et al.*, 2002).

Strong intermolecular  $N-H\cdots O$  hydrogen bonds (Table 2) between O atoms of the carboxylate groups and H(N) atoms of the pyridine rings result in the formation of two-dimensional polymeric sheets lying in the *bc* plane (Fig. 2).

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Received 23 December 2005 Accepted 22 February 2006

# Experimental

A mixture of tri-*n*-butyltin oxide (596.1 mg, 2.0 mmol) and 5-chloro-6-hydroxynicotinic acid (347.1 mg, 2.0 mmol) in benzene (80 ml) was heated under reflux for 8 h. The resulting clear solution was evaporated under vacuum. The product was crystallized from a mixture of dichloromethane/methanol (1:1) (yield 712.2 mg, 76%; m.p. 430 K). Analysis calculated for  $C_{18}H_{29}$ CINSn: C 46.84, H 6.33, N 3.30%; found: C 46.82, H 6.36, N 3.38%.

Mo  $K\alpha$  radiation

reflections

 $\theta = 2.5 - 26.3^{\circ}$  $\mu = 1.32 \text{ mm}^{-1}$ 

T = 298 (2) K

 $R_{\rm int} = 0.055$ 

 $\theta_{\rm max} = 25.0^{\circ}$ 

 $h = -10 \rightarrow 10$ 

 $k = -10 \rightarrow 12$  $l = -23 \rightarrow 26$ 

Block, colorless

 $0.38 \times 0.24 \times 0.17 \text{ mm}$ 

3815 independent reflections 2876 reflections with  $I > 2\sigma(I)$ 

Flack parameter: -0.05(5)

Cell parameters from 3993

# Crystal data

 $[Sn(C_4H_9)_3(C_6H_3CINO_3)]$   $M_r = 462.57$ Orthorhombic,  $P2_12_12_1$  a = 8.825 (2) Å b = 10.823 (3) Å c = 22.592 (6) Å V = 2157.9 (10) Å<sup>3</sup> Z = 4 $D_x = 1.424$  Mg m<sup>-3</sup>

#### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{min} = 0.634, T_{max} = 0.807$ 11314 measured reflections

#### Refinement

 $\begin{array}{ll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.0634P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.042 & + 1.3585P] \\ wR(F^2) = 0.117 & where \ P = (F_o^2 + 2F_c^2)/3 \\ S = 1.00 & (\Delta/\sigma)_{max} = 0.002 \\ 3815 \ reflections & \Delta\rho_{max} = 1.30 \ e^{-3} \\ 226 \ parameters & \Delta\rho_{min} = -0.65 \ e^{-3} \\ H-atom \ parameters \ constrained & 2193 \ Friedel \ pairs \\ \end{array}$ 

# Table 1

Selected geometric parameters (Å, °).

Sn1-C15	2.119 (7)	Sn1-O1	2.146 (4)
Sn1-C7	2.133 (6)	Sn1-O3 <sup>i</sup>	2.537 (4)
Sn1-C11	2.143 (7)	O3-Sn1 <sup>ii</sup>	2.537 (4)
C15-Sn1-C7	126.7 (3)	C11-Sn1-O1	92.7 (3)
C15-Sn1-C11	115.1 (3)	C15-Sn1-O3 <sup>i</sup>	81.9 (2)
C7-Sn1-C11	115.1 (3)	C7-Sn1-O3 <sup>i</sup>	82.4 (2)
C15-Sn1-O1	98.1 (2)	C11-Sn1-O3 <sup>i</sup>	88.9 (3)
C7-Sn1-O1	96.2 (2)	$O1-Sn1-O3^i$	178.21 (19)

Symmetry codes: (i) x, y - 1, z; (ii) x, y + 1, z.

# Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N1-H1···O2 <sup>iii</sup>	0.86	1.95	2.781 (8)	167
Symmetry code: (iii)	$-x, y + \frac{1}{2}, -z +$	- <u>3</u> -		

When the crystal structure was solved, atoms C17 and C18 were found to be disordered. During the refinement process with isotropic displacement parameters, the occupancies of the disordered atoms









The packing, viewed approximately down the c axis. Hydrogen bonds are indicated by dashed lines. H atoms and the minor disorder component have been omitted for clarity.

were refined. These site-occupancy factors were fixed as C17 = 0.80, C17' = 0.20, C18 = 0.80 and C18' = 0.20 during the anisotropic refinement. H atoms were positioned geometrically, with N-H = 0.86Å and C-H = 0.93, 0.96 and 0.97Å for aromatic, methyl and methylene H atoms, respectively, and constrained to ride on their parent atoms, with  $U_{iso}(H) = xU_{cq}(C,N)$  where x = 1.5 for methyl H and x = 1.2 for all other H atoms.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

We acknowledge financial support from the Shandong Province Science Foundation and the State Key Laboratory of Crystal Material, Shandong University, China.

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