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## Key indicators

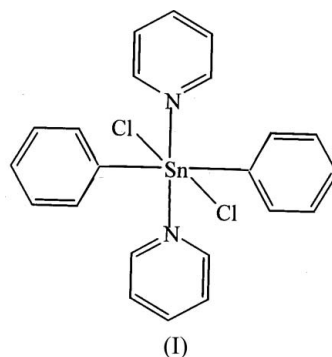
Single-crystal X-ray study  
 $T = 298$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.008$  Å  
 $R$  factor = 0.038  
 $wR$  factor = 0.101  
Data-to-parameter ratio = 21.1For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Dichlorodiphenylbis(pyridine- $\kappa$ N)tin(IV)

In the mononuclear title complex,  $[\text{Sn}(\text{C}_6\text{H}_5)_2\text{Cl}_2(\text{C}_5\text{H}_5\text{N})_2]$ , the central  $\text{Sn}^{\text{IV}}$  atom is octahedrally coordinated by two Cl atoms, two pyridine N atoms, and two benzene C atoms. The dihedral angle between the pyridine ring and the  $\text{SnN}_2\text{Cl}_2$  plane is  $19.2$  ( $9$ )°.

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## Comment

The molecular structure of the title compound, (I), is shown in Fig. 1. The Sn atom is octahedrally coordinated (Table 1) by two pyridine (py) N atoms, two benzene C atoms and two chloride ions in an all-*trans* configuration.



The  $\text{SnC}_2\text{N}_2\text{Cl}_2$  octahedron is relatively undistorted. The Sn—C, Sn—Cl and Sn—N bond lengths in (I) are in good agreement with those in  $\text{py}_2\text{Et}_2\text{SnCl}_2$  (Casas *et al.*, 2000) and  $\text{py}_2\text{Me}_2\text{SnCl}_2$  (Aslanov *et al.*, 1978). The bond angles around Sn in (I) are all very close to  $90^\circ$  [range =  $88.37$  ( $14$ )– $91.84$  ( $14$ )°].

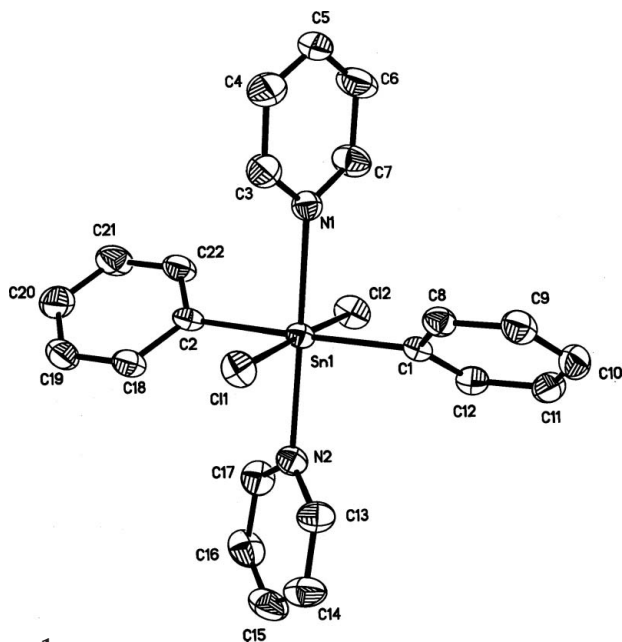
## Experimental

To a benzene suspension of  $(\text{C}_6\text{H}_5)_2\text{SnCl}_2$  (5 mmol) was added a benzene (30 ml) solution of pyridine (10 mmol). The mixture was heated under reflux with stirring for 7 h, and the resulting clear solution was evaporated under vacuum to leave a colourless solid. This was recrystallized from dichloromethane–hexane (3:1 *v/v*), resulting in crystals of (I) suitable for X-ray analysis (m.p. 439 K) after 7 d. Analysis for  $\text{C}_{22}\text{H}_{20}\text{Cl}_2\text{N}_2\text{Sn}$ , calculated: C 52.63, H 4.02%; found: C 52.45, H 4.13%.

## Crystal data

$[\text{Sn}(\text{C}_6\text{H}_5)_2\text{Cl}_2(\text{C}_5\text{H}_5\text{N})_2]$   
 $M_r = 501.99$   
Monoclinic,  $P2_1/n$   
 $a = 9.445$  (3) Å  
 $b = 16.720$  (5) Å  
 $c = 13.474$  (4) Å  
 $\beta = 91.584$  (4)°  
 $V = 2127.0$  (10) Å<sup>3</sup>  
 $Z = 4$

$D_x = 1.568$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 3361  
reflections  
 $\theta = 2.4$ – $25.7^\circ$   
 $\mu = 1.46$  mm<sup>-1</sup>  
 $T = 298$  (2) K  
Block, colourless  
 $0.48 \times 0.39 \times 0.36$  mm



**Figure 1**  
The structure of (I), showing 30% probability displacement ellipsoids. H atoms have been omitted for clarity.

#### Data collection

Siemens SMART CCD diffractometer	5142 independent reflections
$\varphi$ and $\omega$ scans	2752 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.042$
$T_{\text{min}} = 0.541$ , $T_{\text{max}} = 0.621$	$\theta_{\text{max}} = 28.4^\circ$
13388 measured reflections	$h = -11 \rightarrow 12$
	$k = -16 \rightarrow 22$
	$l = -14 \rightarrow 17$

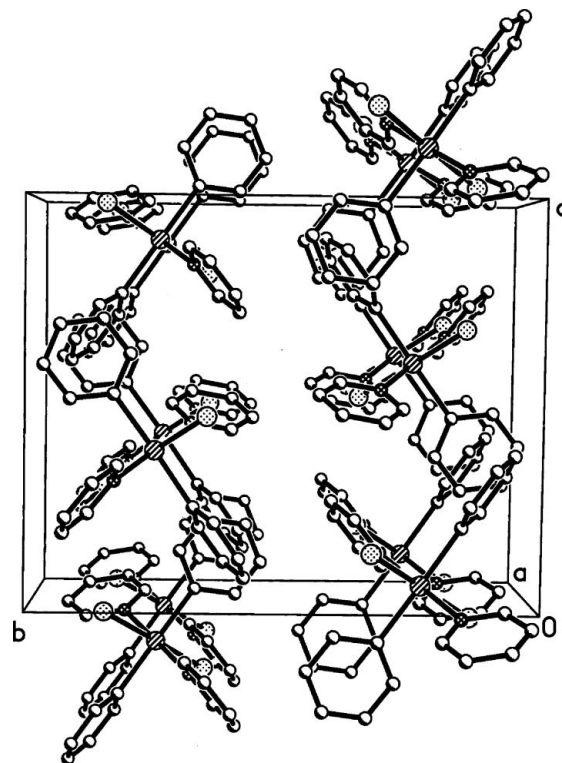
#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0399P)^2 + 0.5683P]$
$R[F^2 > 2\sigma(F^2)] = 0.038$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.101$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.00$	$\Delta\rho_{\text{max}} = 1.12 \text{ e } \text{\AA}^{-3}$
5142 reflections	$\Delta\rho_{\text{min}} = -0.49 \text{ e } \text{\AA}^{-3}$
244 parameters	
H-atom parameters constrained	

**Table 1**  
Selected bond lengths ( $\text{\AA}$ ).

Sn1—C2	2.192 (4)	Sn1—N1	2.331 (4)
Sn1—C1	2.194 (4)	Sn1—Cl2	2.5310 (12)
Sn1—N2	2.314 (4)	Sn1—Cl1	2.5331 (12)

All H atoms were positioned geometrically and refined as riding, with C—H = 0.93  $\text{\AA}$  and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ . The maximum and minimum electron-density peaks are located 1.38  $\text{\AA}$  from atom H18 and 1.40  $\text{\AA}$  from atom Cl2, respectively.



**Figure 2**  
The packing of (I). H atoms have been omitted.

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 publication: *SHELXTL*.

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