

V = 1319.6 (3) Å<sup>3</sup>

Mo  $K\alpha$  radiation

 $0.16 \times 0.14 \times 0.10 \text{ mm}$ 

6681 measured reflections

1584 independent reflections 1562 reflections with  $I > 2\sigma(I)$ 

 $\mu = 2.63 \text{ mm}^-$ 

T = 100 K

 $R_{\rm int} = 0.024$ 

Z = 4

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### Dichloridobis(pyridine-2-thiolato- $\kappa^2 N$ ,S)tin(IV): a new polymorph

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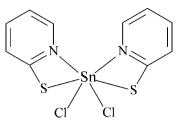
Received 26 April 2012; accepted 25 May 2012

Key indicators: single-crystal X-ray study; T = 100 K; mean  $\sigma$ (C–C) = 0.005 Å; R factor = 0.020; wR factor = 0.050; data-to-parameter ratio = 20.1.

The title compound,  $[SnCl_2(C_5H_4NS)_2]$ , is the product of reaction of 2,2'-dipyridyl disulfide with tin tetrachloride. The Sn<sup>IV</sup> atom adopts a distorted octahedral geometry, with the two bidentate pyridine-2-thiolate ligands forming two planar four-membered chelate rings. The two Sn-Cl, two Sn-N and two Sn-S bonds are in *cis*, *cis* and *trans* configurations, respectively. The crystal grown from acetonitrile represents a new monoclinic polymorph in space group C2/c with the molecule having twofold rotational symmetry, the Sn<sup>IV</sup> atom lying on the twofold axis. The molecular structure of the monoclinic polymorph is very close to that of the triclinic polymorph studied previously in space group  $P\overline{1}$ , the molecule occupying a general position [Masaki & Matsunami (1976). Bull. Chem. Soc. Jpn, 49, 3274–3279; Masaki et al. (1978). Bull. Chem. Soc. Jpn, 51, 3298-3301]. Apparently, the formation of the two polymorphs is determined by the different systems of intermolecular interactions. In the crystal of the monoclinic polymorph, molecules are bound into ribbons along the c axis by  $C-H \cdots Cl$  hydrogen bonds, whereas in the crystal of the triclinic polymorph, molecules form chains along the *a* axis by attractive  $S \cdot \cdot S$  interactions. The crystal studied was a pseudomerohedral twin; the refined BASF value is 0.221 (1).

#### **Related literature**

For metal complexes with 2,2'-dipyridyl dichalcogenides, see: Kadooka et al. (1976a,b); Cheng et al. (1996); Kienitz et al. (1996); Bell et al. (2000); Kita et al. (2001); Kedarnath et al. (2009). For the triclinic polymorph, see: Masaki & Matsunami (1976); Masaki et al. (1978).



#### **Experimental**

Crystal data
$[SnCl_2(C_5H_4NS)_2]$
$M_r = 409.93$
Monoclinic, $C2/c$
a = 6.3240 (7)  Å
b = 12.9391 (14)  Å
c = 16.4240 (18)  Å
$\beta = 100.922 (2)^{\circ}$
Data collection
Duta concention
Bruker SMART 1K CCD
diffractometer
Absorption correction: multi-scan

Absorption correction: multi-scan	
(SADABS; Sheldrick, 1998)	
$T_{\min} = 0.678, \ T_{\max} = 0.779$	

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.020$	79 parameters
$wR(F^2) = 0.050$	H-atom parameters constrained
S = 1.00	$\Delta \rho_{\rm max} = 0.81 \ {\rm e} \ {\rm \AA}^{-3}$
1584 reflections	$\Delta \rho_{\min} = -0.53 \text{ e} \text{ Å}^{-3}$

#### Table 1

Hydrogen-bond geometry (Å, °).

 $\overline{D - \mathbf{H} \cdot \cdot \cdot A}$ D-H $H \cdot \cdot \cdot A$  $D \cdots A$  $D - H \cdot \cdot \cdot A$ C3-H3···Cl1<sup>i</sup> 0.95 2.80 3.673 (3) 154

Symmetry code: (i) -x + 1, -y + 1, -z.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

We thank Professor Abel M. Maharramov for fruitful discussions and help in this work.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2356).

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Dichloridobis(pyridine-2-thiolato- $\kappa^2 N$ , S)tin(IV): a new polymorph

# Sheyda R. Ismaylova, Zhanna V. Matsulevich, Galina N. Borisova, Alexander V. Borisov and Victor N. Khrustalev

#### S1. Comment

The coordination chemistry of 2,2'-dipyridyl dichalcogenides to metal ions is a topic of current research interest owing to the application of these complexes as potential precursors for the generation of semiconducting materials (Kadooka *et al.*, 1976*a*, 1976*b*; Cheng *et al.*, 1996; Kienitz *et al.*, 1996; Bell *et al.*, 2000; Kita *et al.*, 2001; Kedarnath *et al.*, 2009).

This article describes the new monoclinic polymorph of dichlorobis(2-pyridinethiolato)tin(IV),  $C_{10}H_8Cl_2N_2S_2Sn$  (I), which was obtained by the reaction of 2,2'-dipyridyl disulfide with tin tetrachloride (Fig. 1). The synthesis of the title compound by the reaction of 2,2'-dipyridyl disulfide with tin dichloride and its triclinic polymorph were reported previously (Masaki & Matsunami, 1976; Masaki *et al.*, 1978).

The molecule of I possesses overall intrinsic  $C_2$  symmetry. In contrast to the triclinic polymorph (the space group  $P\overline{1}$ , the molecule occupies a common position), this symmetry is realised in the crystal of the monoclinic polymorph (the space group C2/c, the molecule occupies a special position on the twofold axis). The tin atom adopts a distorted octahedral geometry, with the two bidentate 2-pyridinethiolato ligands forming two planar four-membered chelate rings (Fig. 2). The two Sn–Cl, two Sn–N and two Sn–S bonds are in *cis-*, *cis-* and *trans-*configurations, respectively. Generally, the molecular structure of the monoclinic polymorph of I is very close to that of the triclinic polymorph.

Apparently, the formation of the two polymorphs of **I** is determined by the different systems of intermolecular nonvalent interactions. In the crystal of the monoclinic polymorph, the molecules are bound into the ribbons along the *c* axis by the weak intermolecular C3–H3···Cl1<sup>i</sup> hydrogen bonds (Fig. 3, Table 1), whereas, in the crystal of the triclinic polymorph, the molecules form the chains along the *a* axis by the weak attractive intermolecular S···S (3.544 (3)Å) interactions (Fig. 4). Symmetry code: (i) -*x*+1, -*y*+1, -*z*.

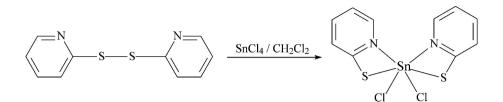
#### **S2. Experimental**

A solution of SnCl<sub>4</sub> (0.13 g, 0.5 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (25 ml) was added to a solution of 2,2'-dipyridyl disulfide (0.11 g, 0.5 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (25 ml) with stirring at room temperature. After 1 h, the powder of complex (C<sub>5</sub>H<sub>4</sub>NS)<sub>2</sub>SnCl<sub>6</sub> was separated by filtration. The filtrate was concentrated in *vacuo*. The solid was re-crystallized from CH<sub>3</sub>CN to give **I** as colourless crystals. Yield is 43%. M.p. = 546-548 K. <sup>1</sup>H NMR (*DMSO-d*<sub>6</sub>, 300 MHz, 302 K):  $\delta$  = 8.48 (d, 2H, H6, J = 4.4), 7.81 (t, 2H, H4, J = 7.3), 7.62 (d, 2H, H3, J = 7.3), 7.28 (dd, 2H, H5, J = 7.3, J = 4.4). Anal. Calcd. for C<sub>10</sub>H<sub>8</sub>Cl<sub>2</sub>N<sub>2</sub>S<sub>2</sub>Sn: C, 29.29; H, 1.97; N, 6.83. Found: C, 29.21; H, 1.92; N, 6.79.

#### **S3. Refinement**

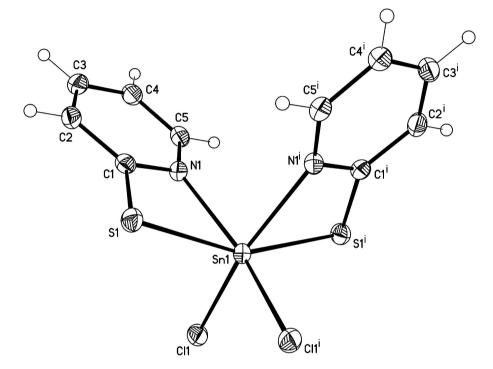
The crystal of I was a pseudo-merohedral twin. The twin matrix is (1 0 0 0 -1 0 -1 0 -1), and BASF is equal to 0.221 (1).

The hydrogen atoms were placed in calculated positions with C–H = 0.95Å and refined in the riding model with fixed isotropic displacement parameters  $U_{iso}(H) = 1.2U_{eq}(C)$ .



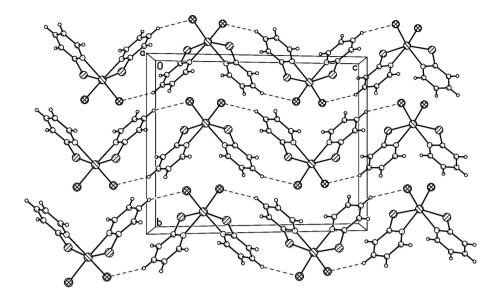
#### Figure 1

Reaction of 2,2'-dipyridyl disulfide with tin tetrachloride.



#### Figure 2

Molecular structure of **I** with the atom numbering scheme. Displacement ellipsoids are shown at the 50% probability level. H atoms are presented as a small spheres of arbitrary radius. Symmetry code: (i) -x+1, y, -z+1/2.





The H-bonded ribbons along the c axis in the monoclinic polymorph of I.

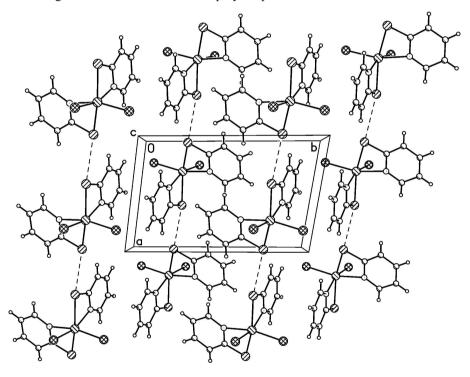


Figure 4

The S $\cdots$ S bonded chains along the *a* axis in the triclinic polymorph of **I**. Dashed lines indicate the intermolecular non-valent interactions.

Dichloridobis(pyridine-2-thiolato-κ<sup>2</sup>N,S)tin(IV)

Crystal data  $[SnCl_2(C_5H_4NS)_2]$  $M_r = 409.93$ 

Monoclinic, *C*2/*c* Hall symbol: -C 2yc Mo *K* $\alpha$  radiation,  $\lambda = 0.71073$  Å

 $\theta = 2.5 - 30.0^{\circ}$ 

 $\mu = 2.63 \text{ mm}^{-1}$ 

Prism, colourless

 $0.16 \times 0.14 \times 0.10 \text{ mm}$ 

T = 100 K

Cell parameters from 6340 reflections

a = 6.3240 (7) Å b = 12.9391 (14) Å c = 16.4240 (18) Å  $\beta = 100.922 (2)^{\circ}$   $V = 1319.6 (3) \text{ Å}^{3}$  Z = 4 F(000) = 792 $D_{x} = 2.063 \text{ Mg m}^{-3}$ 

#### Data collection

Dura concention	
Bruker SMART 1K CCD	6681 measured reflections
diffractometer	1584 independent reflections
Radiation source: fine-focus sealed tube	1562 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\rm int} = 0.024$
$\varphi$ - and $\omega$ -scans	$\theta_{\text{max}} = 28.0^{\circ}, \ \theta_{\text{min}} = 1.3^{\circ}$
Absorption correction: multi-scan	$h = -8 \rightarrow 8$
(SADABS; Sheldrick, 1998)	$k = -16 \rightarrow 16$
$T_{\min} = 0.678, \ T_{\max} = 0.779$	$l = -21 \rightarrow 21$

#### Refinement

Refinement on $F^2$	Secondary atom site location: difference Fourier
Least-squares matrix: full	map
$R[F^2 > 2\sigma(F^2)] = 0.020$	Hydrogen site location: inferred from
$wR(F^2) = 0.050$	neighbouring sites
S = 1.00	H-atom parameters constrained
1584 reflections	$w = 1/[\sigma^2(F_o^2) + (0.014P)^2 + 7.75P]$
79 parameters	where $P = (F_o^2 + 2F_c^2)/3$
0 restraints	$(\Delta/\sigma)_{\rm max} < 0.001$
Primary atom site location: structure-invariant	$\Delta \rho_{\rm max} = 0.81 \text{ e } \text{\AA}^{-3}$
direct methods	$\Delta \rho_{\rm min} = -0.53 \text{ e} \text{ Å}^{-3}$

#### Special details

**Geometry**. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'s involving l.s. planes.

**Refinement**. Refinement of  $F^2$  against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on  $F^2$ , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating *R*-factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. *R*-factors based on  $F^2$  are statistically about twice as large as those based on *F*, and *R*-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters  $(Å^2)$ 

	x	у	Ζ	$U_{ m iso}$ */ $U_{ m eq}$
Sn1	0.5000	0.38034 (2)	0.2500	0.01554 (7)
C11	0.68565 (13)	0.25722 (5)	0.18092 (4)	0.02082 (15)
S1	0.17981 (13)	0.41891 (5)	0.14189 (4)	0.01907 (14)
N1	0.5510 (4)	0.50833 (19)	0.16215 (15)	0.0166 (5)
C1	0.3563 (5)	0.5095 (2)	0.11252 (18)	0.0170 (6)
C2	0.3100 (5)	0.5780 (2)	0.04557 (18)	0.0201 (6)
H2	0.1720	0.5786	0.0105	0.024*
C3	0.4699 (6)	0.6444 (2)	0.0318 (2)	0.0229 (7)
H3	0.4425	0.6915	-0.0134	0.028*

C4	0.6725 (6)	0.6430 (2)	0.08400 (17)	0.0209 (6)
H4	0.7836	0.6884	0.0749	0.025*
C5	0.7064 (5)	0.5737 (2)	0.14906 (17)	0.0190 (5)
H5	0.8425	0.5722	0.1854	0.023*

Atomic displacement parameters  $(Å^2)$ 

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Sn1	0.01793 (13)	0.01560 (12)	0.01225 (12)	0.0000	0.00070 (12)	0.0000
Cl1	0.0251 (4)	0.0200 (3)	0.0175 (3)	0.0044 (3)	0.0042 (3)	-0.0017 (2)
S1	0.0182 (3)	0.0197 (3)	0.0179 (3)	-0.0009 (3)	-0.0002 (3)	0.0007 (2)
N1	0.0196 (12)	0.0165 (12)	0.0131 (11)	0.0011 (9)	0.0018 (9)	-0.0001 (9)
C1	0.0209 (14)	0.0153 (13)	0.0152 (13)	0.0005 (10)	0.0041 (11)	-0.0020 (10)
C2	0.0236 (15)	0.0208 (14)	0.0150 (13)	0.0043 (12)	0.0013 (11)	-0.0019 (11)
C3	0.0334 (18)	0.0190 (14)	0.0164 (14)	0.0036 (12)	0.0046 (13)	0.0014 (11)
C4	0.0250 (15)	0.0190 (14)	0.0191 (13)	-0.0032 (13)	0.0053 (14)	-0.0012 (10)
C5	0.0213 (14)	0.0195 (13)	0.0161 (12)	-0.0007 (12)	0.0031 (12)	-0.0028 (10)

Geometric parameters (Å, °)

Sn1—N1	2.259 (2)	C2—C3	1.379 (5)
Sn1—Cl1	2.3892 (8)	C2—H2	0.9500
Sn1—S1	2.4779 (8)	C3—C4	1.400 (5)
S1—C1	1.748 (3)	С3—Н3	0.9500
N1—C1	1.342 (4)	C4—C5	1.380 (4)
N1—C5	1.345 (4)	C4—H4	0.9500
C1—C2	1.399 (4)	С5—Н5	0.9500
N1 <sup>i</sup> —Sn1—N1	85.72 (12)	N1—C1—S1	112.7 (2)
N1 <sup>i</sup> —Sn1—Cl1	159.13 (7)	C2—C1—S1	126.3 (2)
N1—Sn1—Cl1	92.47 (7)	C3—C2—C1	118.3 (3)
Cl1 <sup>i</sup> —Sn1—Cl1	96.36 (4)	C3—C2—H2	120.9
N1 <sup>i</sup> —Sn1—S1	96.54 (7)	C1—C2—H2	120.9
N1—Sn1—S1	65.85 (7)	C2—C3—C4	120.4 (3)
Cl1 <sup>i</sup> —Sn1—S1	93.80 (3)	С2—С3—Н3	119.8
Cl1—Sn1—S1	101.68 (3)	С4—С3—Н3	119.8
S1 <sup>i</sup> —Sn1—S1	156.76 (4)	C5—C4—C3	118.2 (3)
C1—S1—Sn1	81.67 (10)	C5—C4—H4	120.9
C1—N1—C5	120.7 (3)	C3—C4—H4	120.9
C1—N1—Sn1	99.82 (18)	N1—C5—C4	121.4 (3)
C5—N1—Sn1	139.5 (2)	N1—C5—H5	119.3
N1—C1—C2	121.0 (3)	C4—C5—H5	119.3
N1 <sup>i</sup> —Sn1—S1—C1	82.81 (12)	S1—Sn1—N1—C5	179.8 (3)
N1—Sn1—S1—C1	0.49 (12)	C5—N1—C1—C2	0.5 (4)
Cl1 <sup>i</sup> —Sn1—S1—C1	175.74 (10)	Sn1—N1—C1—C2	-179.2 (2)
Cl1—Sn1—S1—C1	-86.96 (10)	C5—N1—C1—S1	-179.4 (2)
S1 <sup>i</sup> —Sn1—S1—C1	43.78 (10)	Sn1—N1—C1—S1	0.9 (2)

N1 <sup>i</sup> —Sn1—N1—C1	-99.78 (19)	Sn1—S1—C1—N1	-0.82(19)	
Cl1 <sup>i</sup> —Sn1—N1—C1	-14.1 (3)	Sn1—S1—C1—C2	179.3 (3)	
Cl1—Sn1—N1—C1	101.05 (17)	N1—C1—C2—C3	0.0 (4)	
S1 <sup>i</sup> —Sn1—N1—C1	-164.84 (16)	S1—C1—C2—C3	179.9 (2)	
S1—Sn1—N1—C1	-0.64 (15)	C1—C2—C3—C4	-0.2 (5)	
N1 <sup>i</sup> —Sn1—N1—C5	80.6 (3)	C2—C3—C4—C5	-0.1 (5)	
Cl1 <sup>i</sup> —Sn1—N1—C5	166.3 (2)	C1—N1—C5—C4	-0.8 (4)	
$Cl1 - Sn1 - N1 - C5$ $Sl^{i} - Sn1 - N1 - C5$	-78.5(3) 15.6(3)	Sn1—N1—C5—C4 C3—C4—C5—N1	178.7 (2) 0.6 (5)	

Symmetry code: (i) -x+1, y, -z+1/2.

#### Hydrogen-bond geometry (Å, °)

D—H···A	D—H	Н…А	D···· $A$	D—H···A
C3—H3···Cl1 <sup>ii</sup>	0.95	2.80	3.673 (3)	154

Symmetry code: (ii) -x+1, -y+1, -z.