

[4-(Dimethylamino)pyridine- κN^1]trimethyl(thiocyanato- κN)tin(IV)

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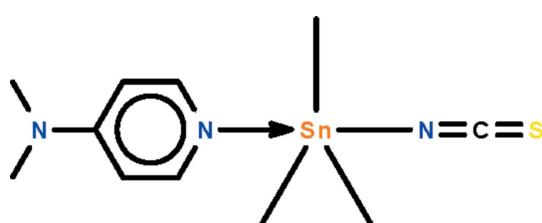
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$; R factor = 0.060; wR factor = 0.197; data-to-parameter ratio = 36.7.

In the title monomeric trimethyltin(IV) isothiocyanate-4,4-dimethylpyridine adduct, $[\text{Sn}(\text{CH}_3)_3(\text{NCS})(\text{C}_7\text{H}_{10}\text{N}_2)]$, the Sn^{IV} atom shows a *trans*- C_3SnN_2 trigonal bipyramidal coordination. The Sn^{IV} atom lies out of the equatorial plane by $0.033(4)\text{ \AA}$ in the direction of the donor N atom of the *N*-heterocycle. The crystal studied was a non-merohedral twin with a minor component of 48.8 (2)%.

Related literature

For trimethyltin isothiocyanate, see: Forder & Sheldrick (1970).



Experimental

Crystal data

$[\text{Sn}(\text{CH}_3)_3(\text{NCS})(\text{C}_7\text{H}_{10}\text{N}_2)]$	$V = 1450.41(14)\text{ \AA}^3$
$M_r = 344.04$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 7.2026(4)\text{ \AA}$	$\mu = 1.89\text{ mm}^{-1}$
$b = 13.4736(8)\text{ \AA}$	$T = 100\text{ K}$
$c = 14.9785(8)\text{ \AA}$	$0.35 \times 0.30 \times 0.25\text{ mm}$
$\beta = 93.792(5)^\circ$	

Data collection

Agilent SuperNova Dual diffractometer with an Atlas detector	15682 measured reflections
Absorption correction: multi-scan (<i>CrysAlis PRO</i> ; Agilent, 2012)	5542 independent reflections
$T_{\min} = 0.558$, $T_{\max} = 0.650$	4916 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.060$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.060$	151 parameters
$wR(F^2) = 0.197$	H-atom parameters constrained
$S = 1.23$	$\Delta\rho_{\max} = 1.61\text{ e \AA}^{-3}$
5542 reflections	$\Delta\rho_{\min} = -1.98\text{ e \AA}^{-3}$

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NK2162).

References

- Agilent (2012). *CrysAlis PRO*. Agilent Technologies, Yarnton, England.
- Barbour, L. J. (2001). *J. Supramol. Chem.* **1**, 189–191.
- Forder, R. A. & Sheldrick, G. M. (1970). *J. Organomet. Chem.* **21**, 115–122.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.

supplementary materials

Acta Cryst. (2012). E68, m790 [doi:10.1107/S1600536812022064]

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Comment

Trimethyltin halides and pseudohalides are Lewis acids that form 1:1 complexes with aromatic amines. Trimethyltin isocyanate itself exists as a polymer in which the isocyanate anion bridges adjacent trimethyltin cations (Forder & Sheldrick, 1970). In the 4,4-dimethylpyridine adduct (Scheme I), the weaker tin–sulfur bond is disrupted, and the adduct is monomeric. The Sn^{IV} atom shows *trans*-C₃SnN₂ trigonal bipyramidal coordination. The tin atom lies out of the equatorial plane by 0.033 (4) Å in the direction of the donor N atom of the *N*-heterocycle.

Experimental

Trimethyltin isothiocyanate (0.24 g, 1 mmol) and 4-(dimethylamino)pyridine (0.11 g, 1 mmol) were loaded into a convection tube and the tube was filled with methanol and kept at 333 K. Light yellow crystals were collected from the side arm after several days.

Refinement

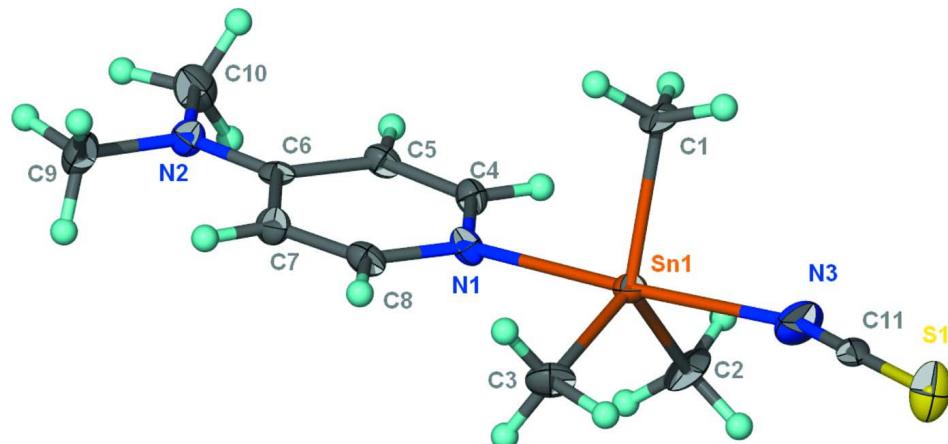
Carbon-bound H-atoms were placed in calculated positions [C–H 0.95 to 0.98 Å, $U_{\text{iso}}(\text{H})$ 1.2 to 1.5 $U_{\text{eq}}(\text{C})$] and were included in the refinement in the riding model approximation.

The crystal is a non-merohedral twin having nearly equal components (minor component 48.8 (2) %). A 100% overlap gave the best refinement; however, an artifact of the twinning is the high weighting scheme, which was suggested by the refinement program. The twin law is (-1 0 0 / 0 - 1 0 / 0.2779 0 1).

The final difference Fourier map had a peak 1.08 Å from Sn1 and a hole 0.95 Å from H1a.

Computing details

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO* (Agilent, 2012); data reduction: *CrysAlis PRO* (Agilent, 2012); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

**Figure 1**

Thermal ellipsoid plot (Barbour, 2001) of $(\text{CH}_3)_3\text{Sn}(\text{NCS})(\text{C}_7\text{H}_{10}\text{N}_2)$ at the 70% probability level; hydrogen atoms are drawn as spheres of arbitrary radius.

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Crystal data



$M_r = 344.04$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 7.2026 (4)$ Å

$b = 13.4736 (8)$ Å

$c = 14.9785 (8)$ Å

$\beta = 93.792 (5)$ °

$V = 1450.41 (14)$ Å³

$Z = 4$

$F(000) = 688$

$D_x = 1.576 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 5713 reflections

$\theta = 2.7\text{--}27.5$ °

$\mu = 1.89 \text{ mm}^{-1}$

$T = 100$ K

Prism, light brown

$0.35 \times 0.30 \times 0.25$ mm

Data collection

Agilent SuperNova Dual

diffractometer with an Atlas detector

Radiation source: SuperNova (Mo) X-ray

Source

Mirror monochromator

Detector resolution: 10.4041 pixels mm⁻¹

ω scan

Absorption correction: multi-scan
(*CrysAlis PRO*; Agilent, 2012)

$T_{\min} = 0.558$, $T_{\max} = 0.650$

15682 measured reflections

5542 independent reflections

4916 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.060$

$\theta_{\max} = 27.7$ °, $\theta_{\min} = 2.7$ °

$h = -9 \rightarrow 9$

$k = -17 \rightarrow 17$

$l = -19 \rightarrow 19$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.060$

$wR(F^2) = 0.197$

$S = 1.23$

5542 reflections

151 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier

map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.1375P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 1.61 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -1.98 \text{ e } \text{\AA}^{-3}$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Sn1	0.25215 (5)	0.41625 (3)	0.86069 (2)	0.01334 (16)
S1	0.3130 (2)	0.35874 (10)	1.18988 (10)	0.0267 (4)
N1	0.2370 (6)	0.4769 (3)	0.7145 (3)	0.0145 (9)
N2	0.2613 (7)	0.5696 (3)	0.4491 (3)	0.0160 (10)
N3	0.2743 (7)	0.3468 (4)	1.0044 (3)	0.0298 (12)
C1	0.5262 (7)	0.3660 (4)	0.8426 (4)	0.0176 (11)
H1A	0.6000	0.4208	0.8204	0.026*
H1B	0.5216	0.3116	0.7991	0.026*
H1C	0.5835	0.3425	0.8999	0.026*
C2	0.0214 (8)	0.3222 (5)	0.8288 (4)	0.0247 (13)
H2A	-0.0670	0.3561	0.7866	0.037*
H2B	-0.0398	0.3060	0.8835	0.037*
H2C	0.0644	0.2609	0.8015	0.037*
C3	0.2037 (8)	0.5581 (4)	0.9169 (4)	0.0219 (12)
H3A	0.0925	0.5878	0.8865	0.033*
H3B	0.3113	0.6011	0.9092	0.033*
H3C	0.1851	0.5509	0.9807	0.033*
C4	0.2356 (7)	0.4114 (4)	0.6457 (3)	0.0146 (11)
H4	0.2297	0.3427	0.6592	0.018*
C5	0.2421 (8)	0.4383 (4)	0.5584 (4)	0.0151 (11)
H5	0.2408	0.3884	0.5135	0.018*
C6	0.2507 (7)	0.5397 (3)	0.5334 (3)	0.0118 (10)
C7	0.2515 (8)	0.6078 (4)	0.6063 (4)	0.0159 (11)
H7	0.2566	0.6771	0.5952	0.019*
C8	0.2448 (8)	0.5743 (3)	0.6913 (4)	0.0152 (11)
H8	0.2457	0.6221	0.7379	0.018*
C9	0.2689 (9)	0.6754 (4)	0.4273 (4)	0.0232 (12)
H9A	0.3785	0.7053	0.4590	0.035*
H9B	0.1563	0.7083	0.4458	0.035*
H9C	0.2771	0.6834	0.3627	0.035*
C10	0.2531 (8)	0.4977 (5)	0.3755 (4)	0.0224 (12)
H10A	0.3642	0.4556	0.3805	0.034*
H10B	0.2477	0.5331	0.3183	0.034*
H10C	0.1418	0.4563	0.3786	0.034*
C11	0.2906 (7)	0.3530 (4)	1.0821 (4)	0.0171 (11)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Sn1	0.0109 (2)	0.0157 (2)	0.0135 (2)	0.00108 (13)	0.00189 (17)	0.00061 (12)
S1	0.0435 (9)	0.0206 (7)	0.0158 (7)	-0.0071 (7)	-0.0002 (7)	-0.0019 (5)
N1	0.017 (2)	0.0109 (19)	0.015 (2)	0.0021 (17)	-0.0028 (19)	-0.0013 (16)
N2	0.020 (3)	0.014 (2)	0.014 (2)	-0.0016 (18)	0.001 (2)	0.0002 (16)
N3	0.031 (3)	0.042 (3)	0.017 (3)	0.004 (3)	0.009 (2)	0.003 (2)
C1	0.004 (2)	0.028 (3)	0.020 (3)	-0.003 (2)	-0.001 (2)	0.006 (2)
C2	0.014 (3)	0.030 (3)	0.030 (3)	-0.015 (2)	0.003 (2)	0.004 (3)
C3	0.021 (3)	0.024 (3)	0.022 (3)	0.002 (2)	0.008 (3)	-0.006 (2)

C4	0.016 (3)	0.011 (2)	0.017 (3)	0.0000 (18)	0.000 (3)	0.0000 (18)
C5	0.015 (3)	0.013 (2)	0.017 (3)	0.002 (2)	0.001 (2)	-0.0027 (19)
C6	0.006 (2)	0.013 (2)	0.016 (2)	-0.0007 (18)	0.001 (2)	-0.0023 (19)
C7	0.016 (3)	0.012 (2)	0.019 (3)	-0.003 (2)	0.002 (2)	0.000 (2)
C8	0.015 (3)	0.012 (2)	0.019 (3)	0.0017 (19)	0.000 (2)	-0.0031 (18)
C9	0.032 (3)	0.020 (3)	0.019 (3)	-0.003 (2)	0.008 (3)	0.004 (2)
C10	0.026 (3)	0.027 (3)	0.013 (2)	0.004 (2)	0.000 (2)	-0.001 (2)
C11	0.007 (2)	0.018 (2)	0.027 (3)	0.000 (2)	0.004 (2)	0.008 (2)

Geometric parameters (\AA , $^\circ$)

Sn1—C2	2.120 (5)	C3—H3A	0.9800
Sn1—C1	2.121 (5)	C3—H3B	0.9800
Sn1—C3	2.126 (5)	C3—H3C	0.9800
Sn1—N1	2.333 (4)	C4—C5	1.360 (7)
Sn1—N3	2.344 (5)	C4—H4	0.9500
S1—C11	1.614 (6)	C5—C6	1.419 (6)
N1—C4	1.357 (6)	C5—H5	0.9500
N1—C8	1.359 (6)	C6—C7	1.426 (7)
N2—C6	1.333 (6)	C7—C8	1.354 (8)
N2—C9	1.464 (6)	C7—H7	0.9500
N2—C10	1.466 (7)	C8—H8	0.9500
N3—C11	1.164 (7)	C9—H9A	0.9800
C1—H1A	0.9800	C9—H9B	0.9800
C1—H1B	0.9800	C9—H9C	0.9800
C1—H1C	0.9800	C10—H10A	0.9800
C2—H2A	0.9800	C10—H10B	0.9800
C2—H2B	0.9800	C10—H10C	0.9800
C2—H2C	0.9800		
C2—Sn1—C1	120.2 (2)	Sn1—C3—H3C	109.5
C2—Sn1—C3	118.7 (2)	H3A—C3—H3C	109.5
C1—Sn1—C3	121.1 (2)	H3B—C3—H3C	109.5
C2—Sn1—N1	90.6 (2)	N1—C4—C5	124.0 (4)
C1—Sn1—N1	88.74 (18)	N1—C4—H4	118.0
C3—Sn1—N1	93.30 (18)	C5—C4—H4	118.0
C2—Sn1—N3	88.5 (2)	C4—C5—C6	120.9 (5)
C1—Sn1—N3	89.0 (2)	C4—C5—H5	119.5
C3—Sn1—N3	89.9 (2)	C6—C5—H5	119.5
N1—Sn1—N3	176.70 (16)	N2—C6—C5	123.2 (4)
C4—N1—C8	115.5 (5)	N2—C6—C7	122.2 (5)
C4—N1—Sn1	118.9 (3)	C5—C6—C7	114.6 (5)
C8—N1—Sn1	125.4 (3)	C8—C7—C6	120.4 (5)
C6—N2—C9	120.7 (4)	C8—C7—H7	119.8
C6—N2—C10	120.7 (4)	C6—C7—H7	119.8
C9—N2—C10	118.4 (5)	C7—C8—N1	124.6 (5)
C11—N3—Sn1	152.3 (5)	C7—C8—H8	117.7
Sn1—C1—H1A	109.5	N1—C8—H8	117.7
Sn1—C1—H1B	109.5	N2—C9—H9A	109.5
H1A—C1—H1B	109.5	N2—C9—H9B	109.5

Sn1—C1—H1C	109.5	H9A—C9—H9B	109.5
H1A—C1—H1C	109.5	N2—C9—H9C	109.5
H1B—C1—H1C	109.5	H9A—C9—H9C	109.5
Sn1—C2—H2A	109.5	H9B—C9—H9C	109.5
Sn1—C2—H2B	109.5	N2—C10—H10A	109.5
H2A—C2—H2B	109.5	N2—C10—H10B	109.5
Sn1—C2—H2C	109.5	H10A—C10—H10B	109.5
H2A—C2—H2C	109.5	N2—C10—H10C	109.5
H2B—C2—H2C	109.5	H10A—C10—H10C	109.5
Sn1—C3—H3A	109.5	H10B—C10—H10C	109.5
Sn1—C3—H3B	109.5	N3—C11—S1	178.7 (5)
H3A—C3—H3B	109.5		
C2—Sn1—N1—C4	-52.6 (4)	C9—N2—C6—C5	179.7 (5)
C1—Sn1—N1—C4	67.6 (4)	C10—N2—C6—C5	3.8 (8)
C3—Sn1—N1—C4	-171.4 (4)	C9—N2—C6—C7	-1.8 (8)
C2—Sn1—N1—C8	133.2 (5)	C10—N2—C6—C7	-177.7 (5)
C1—Sn1—N1—C8	-106.6 (5)	C4—C5—C6—N2	178.5 (5)
C3—Sn1—N1—C8	14.4 (5)	C4—C5—C6—C7	-0.1 (8)
C2—Sn1—N3—C11	-133.1 (10)	N2—C6—C7—C8	-178.4 (5)
C1—Sn1—N3—C11	106.7 (10)	C5—C6—C7—C8	0.2 (8)
C3—Sn1—N3—C11	-14.4 (10)	C6—C7—C8—N1	-0.1 (9)
C8—N1—C4—C5	0.3 (8)	C4—N1—C8—C7	-0.2 (8)
Sn1—N1—C4—C5	-174.5 (4)	Sn1—N1—C8—C7	174.2 (5)
N1—C4—C5—C6	-0.1 (9)		