Cl1—Sn—Cl2	105.4 (1)	Cl2-Sn-C11	91.2 (3)
Cl1—Sn—N1	94.4 (2)	N1SnN2	73.0 (3)
Cl1—Sn—N2	167.4 (2)	N1-Sn-C10	90.7 (3)
Cl1—Sn—C10	86.9 (3)	N1SnC11	88.6 (3)
Cll—Sn—Cll	90.1 (3)	N2SnC10	93.2 (3)
Cl2—Sn—N1	160.2 (2)	N2SnC11	89.4 (4)
Cl2—Sn—N2	87.2 (2)	C10-Sn-C11	176.9 (4)
Cl2SnC10	90.5 (3)		

Table 2. Hydrogen-bonding geometry (Å, °)

D — $H \cdot \cdot \cdot A$	<i>D</i> —Н	$\mathbf{H} \cdots \mathbf{A}$	$D \cdots A$	$D = H \cdot \cdot$
N2-HN2A···Cl11	0.950	2.369	3.285 (9)	161.7
N2—HN2B···Cl2 ⁱⁱ	0.950	2.332	3.274 (8)	171.2
Symmetry codes: (i)	x, y, -1 + z;	(ii) $2 - x$, 1	-y, 2-z.	

H atoms of the aminoquinoline were kept fixed at calculated positions (C—H = 0.95 Å), those of the methyl groups were kept fixed at their observed positions. U_{iso} was set to be 20% larger than U_{eq} for the atom to which they were attached. The distances from Sn to the minimum and maximum values of $\Delta \rho$ are 0.94 and 1.22 Å, respectively.

Data collection: MAD (Allibon, 1995). Cell refinement: MAD. Data reduction: KRYSTAL (Hazell, 1995). Program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994) and KRYSTAL. Program(s) used to refine structure: modified ORFLS (Busing, Martin & Levy, 1962). Molecular graphics: ORTEPIII (Burnett & Johnson, 1996) and KRYSTAL. Software used to prepare material for publication: KRYSTAL.

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Chlorodimethyl(*N*-pyrrolidinecarbodithioato-*S*,*S'*)tin(**IV**)

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Abstract

The Sn atom in $[SnCl(C_5H_8NS_2)(CH_3)_2]$ is in a distorted trigonal bipyramidal environment, with the two methyl C atoms and an S atom in the equatorial plane, and the Cl and other S atom occupying axial positions. The carbodithioate ligand is unsymmetrically coordinated to the Sn atom with Sn—S distances of 2.7555 (10) and 2.4825 (12) Å.

Comment

Although a number of pentadentate diorganotin complexes of the type R_2 ClSnS₂C have been reported, examples of structural studies are limited to $Me_2SnCl(S_2CNMe_2)$, PhBuSnCl(S_2CNEt_2), Ph₂SnCl(S₂COⁱPr) (Furue, Kimura, Yasuoka, Kasai & Masao, 1970; Wei, Kumar Das & Sinn, 1985; Dakternieks, Hoskins, Jackson, Tiekink & Winter, 1985). In these complexes, the Sn atom is asymmetrically bonded to the two S atoms with an average difference in the Sn—S bond lengths of 0.3 Å, similar to that observed in the title compound, $Me_2ClSn[S_2CN(CH_2)_4]$, (1). However, the Sn-S distances in the related 1,2-dithio anionic complex 2-chloro-2,2-diphenyl-5-methyl-1,3,2benzodithiastannole are similar (Sau, Day & Holmes, 1981). As in (1), the chloro group in all these complexes invariably occupies an axial position with the two organo groups in equatorial positions.



The displacement ellipsoid plot of (1) with the numbering scheme is shown in Fig. 1. The geometry about the Sn atom is a distorted trigonal bipyramid with the Cl and S1 atoms in axial positions and the two methyl C atoms and S2 in the equatorial plane. Despite the less crowded environment about the Sn atom, the Me—Sn—Me angle in (1) [125.4 (3)°] is smaller than those found in Me₂Sn[S₂CN(CH₂)₄]₂ (137.3°; Lockhart, Manders & Schlemper, 1985) and Me₂Sn(S₂CNMe₂)₂ (136°; Kimura, Yasuoka, Kasai & Kakudo, 1972), both of which are six-coordinate. The observed Me-Sn-Me and S-Sn-S angles are consistent with the presence of the more electronegative chloro group exerting its influence on the geometry. The crystal structure is further stabilized by intermolecular contacts involving centrosymmetrically related pairs of molecules $[S1 \cdots S1 \ 3.392(2)]$ and $S1 \cdots H7A \ 2.94(15) \ A]$.



Fig. 1. A 30% probability displacement ellipsoid plot of (1) with the atomic numbering scheme.

Experimental

The title compound was prepared by reacting equimolar quantities of dimethyltin dichloride with ammonium pyrrolidinecarbodithioate in ethanol.

Crystal data

 $[SnCl(C_5H_8NS_2)(CH_3)_2]$ Mo $K\alpha$ radiation $M_r = 330.45$ $\lambda = 0.71073 \text{ Å}$ Triclinic *P*1 reflections $\theta = 5.0 - 12.5^{\circ}$ a = 6.422(1) Å $\mu = 2.593 \text{ mm}^{-1}$ b = 10.129(3) Å c = 10.176(1) Å T = 293(2) K $\alpha = 85.60(1)^{\circ}$ Rectangular slab $\beta = 72.95(1)^{\circ}$ $\gamma = 75.97 (2)^{\circ}$ Colourless $V = 614.0(2) \text{ Å}^3$ Z = 2 $D_x = 1.788 \text{ Mg m}^{-3}$ D_m not measured Data collection Siemens P4 diffractometer $R_{int} = 0.022$ ω -2 θ scans

Absorption correction: empirical ψ scans (XSCANS; Siemens, 1994) $T_{\rm min} = 0.273, T_{\rm max} = 0.484$ 3572 measured reflections 2823 independent reflections 2689 reflections with

 $I > 2\sigma(I)$

Cell parameters from 39 $1.16 \times 0.52 \times 0.26$ mm

 $\theta_{\rm max} = 27.5^{\circ}$ $h = -1 \rightarrow 8$ $k = -13 \rightarrow 13$ $l = -13 \rightarrow 13$ 3 standard reflections every 97 reflections intensity decay: <3% Refinement

Refinement on F^2	$(\Delta/\sigma)_{\rm max} < 0.001$
$R[F^2 > 2\sigma(F^2)] = 0.032$	$\Delta \rho_{\rm max} = 0.463 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.085$	$\Delta \rho_{\rm min} = -0.901 \ {\rm e} \ {\rm \AA}^{-3}$
S = 1.152	Extinction correction: none
2823 reflections	Scattering factors from
165 parameters	International Tables for
All H atoms refined	Crystallography (Vol. C)
$w = 1/[\sigma^2(F_o^2) + (0.0399P)^2]$	
+ 0.6691 <i>P</i>]	
where $P = (F_0^2 + 2F_c^2)/3$	

Table 1	. Selected	geometric	parameters	(Á	•)
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	Ç	•	
SnC6	2.105 (5)	N1C1	1.309 (5)
SnC7	2.110 (5)	N1C2	1.474 (5)
SnCl	2.4597 (11)	N1C5	1.476 (5)
Sn—S2	2.4825 (12)	C2C3	1.513 (7)
Sn—S1	2.7555 (10)	C3C4	1.492 (9)
S1C1	1.706 (4)	C4—C5	1.518 (7)
S2C1	1.738 (4)		
C6—Sn—C7	125.4 (3)	C1—S2—Sn	90.58 (12)
C6—Sn—C1	95.7 (2)	C1-N1-C2	123.5 (3)
C7—Sn—Cl	99.8 (2)	C1-N1-C5	124.7 (3)
C6—Sn—S2	123.7 (2)	C2N1C5	111.7 (3)
C7—Sn—S2	109.0 (2)	N1C1S1	122.3 (3)
C1—Sn—S2	87.54 (4)	N1C1S2	119.6(3)
C6—Sn—S1	91.0 (2)	\$1C1\$2	118.1 (2)
27SnS1	96.1 (2)	N1C2C3	103.7 (4)
CISnS1	154.48 (4)	C4C3C2	105.8 (4)
S2SnS1	68.41 (3)	C3C4C5	104.7 (4)
C1S1Sn	82.47 (12)	N1C5C4	103.1 (4)

The title structure was solved by direct methods and refined by a full-matrix least-squares method on F^2 . The H atoms were determined from the Fourier map and refined isotropically. PARST (Nardelli, 1983) was used for geometrical calculations.

Data collection: XSCANS (Siemens, 1994). Cell refinement: XSCANS. Data reduction: XSCANS. Program(s) used to solve structure: SHELXTL/PC (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: SHELXTL/PC. Software used to prepare material for publication: SHELXL93.

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N-(2-Pyridinium)urea Perrhenate[†]

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Abstract

The N-(2-pyridinium)urea cation of the title compound, $C_6H_8N_3O^+$.ReO₄⁻, is planar to within 0.085 (9) Å and adopts an *s-cis-s-cis* conformation with respect to the imino C—N bonds. The structure consists of columns of centrosymmetric dimeric plane-to-plane stacked cations and hydrogen-bonded anions. The distance between the anti-parallel carbonyl groups in the dimer is 3.16 (5) Å. An extensive network of hydrogen bonds stabilizes the ordering of the ions within the columns, which are held together by weaker hydrogen bonds and van der Waals interactions.

Comment

This work is part of an investigation of adducts of unsymmetrically substituted urea derivatives and inorganic salts with potential as non-linear optical materials. We combined N-(2-pyridyl)urea and perrhenic acid, assuming that the anionic host sub-network would affect the molecular geometry and arrangement of the cations.

Bond lengths and angles in the title compound, (I), compare well with the values found for other *N*-(2-pyridyl)urea derivatives (Velikova, Angelova & Kossev, 1997; Le Magueres, Ouahab, Hocquet & Fournier, 1994) and for ReO₄ tetrahedra (Lock & Turner, 1975; Macíček & Todorov, 1992; Macíček & Angelova, 1995). The

protonation of the *endo*-N atom in the pyridyl ring favours the formation of an intramolecular N—H···O hydrogen bond which is crucial for the planarity and conformation of the molecule. The N-(pyridinium)urea cation is planar, with the largest deviation of 0.085 (9) Å at the amino N3 atom. The overall conformation of the molecule can be described as *s*-*cis*-*s*-*cis* with respect to the C1—N2—C2 imino bonds, in contrast to the *s*-*trans*-*s*-*cis* conformation found in N-(2-pyridyl)urea (Velikova, Angelova & Kossev, 1997).



The structure contains centrosymmetric dimers of plane-to-plane stacked cations and hydrogen-bonded anions. The stacking of the cations embraces carbonyl groups which are only 3.16(5) Å apart. In addition, the two coupled cations are captured by two ReO_4^- anions, forming a six-membered hydrogen-bonded ring through O12 and a relatively short, though bifurcated, hydrogen bond through O14 (Table 2). These dimers are ordered in columns along the *a* axis and interlinked by centrosymmetric eight-membered hydrogen-bonded rings formed by pairs of cations, similar to those in the structure of N-(2-pyridyl)urea, and by the HN31...011 branches of the bifurcated hydrogen bond to HN31. The columns are held together by weak bifurcated hydrogen bonds [HN2...O12(1-x, -y, 1-z)] along the b axis, as well as by van der Waals interactions along the caxis.



Fig. 1. The ions of the title compound, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are arbitrarily reduced.

[†] Alternative name: 2-ureidopyridinium tetraoxorhenate.