

Table 5. Selected geometric parameters (\AA , $^\circ$) for (II)

Br1—Cu1	2.482 (1)	Cu1—N2	2.734 (4)
Br2—Cu1	2.419 (1)	Br1—Br2'	4.007 (2)
Br2—Cu1—Br1	90.08 (3)		

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

Table 6. Hydrogen-bonding geometry (\AA , $^\circ$) for (II)

D—H...A	D—H	H...A	D...A	D—H...A
N3—H3A...Br1	0.94 (7)	2.50 (7)	3.43 (1)	170 (6)
N3—H3B...Br2'	0.77 (7)	2.58 (7)	3.31 (1)	162 (7)

Symmetry code: (i) $x, \frac{1}{2} - y, z - \frac{1}{2}$.

The structure was solved by Patterson and Fourier methods (Beurskens *et al.*, 1992). Non-H atoms were refined anisotropically by full-matrix least-squares techniques. Some of the H atoms were isotropically refined, others were geometrically calculated and refined riding with common isotropic displacement parameters in both structures. The final difference Fourier maps show several electron density peaks greater than $1 \text{ e } \text{\AA}^{-3}$, for both structures, close to the Br and Cu atoms. A free anisotropic refinement of atom C3 in structure (II) leads to odd values for the principal axis of the thermal tensor, which are also affected by rounding off errors. Anisotropic refinement using similar parameters (SIMU in SHELXL93; Sheldrick, 1993) for all C atoms of the 2-aminopyrimidine leads to physically acceptable results. All calculations were made at the University of Oviedo on the Scientific Computer Center and X-ray group DEC/AXP computers.

For both compounds, data collection: CAD-4 EXPRESS (Enraf–Nonius, 1994); cell refinement: CRYSDA DIRDIF; data reduction: REFLEX (local program); molecular graphics: Xtal.GX (Hall & Boulay, 1995). Geometrical calculations made with PARST (Nardelli, 1983); software used to prepare material for publication: SHELXL93.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1316). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1997). **C53**, 1789–1791

The 1:1 Antimony Trichloride Adduct of Chlorobis(2,6-dimethylpiperidin-1-yl)-phosphine Oxide

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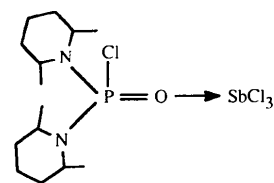
(Received 16 December 1996; accepted 26 June 1997)

Abstract

In the structure of the title complex, $[(\text{C}_7\text{H}_{14}\text{N})_2\text{Cl-P}(\text{O})]_2\text{SbCl}_3$, the Sb atom shows a distorted trigonal bipyramidal geometry (including the lone pair). The O—Sb interaction of antimony with the phosphoryl O atom is only moderate and is weaker than that observed in the known compound $\text{Cl}_3\text{PO} \rightarrow \text{SbCl}_5$.

Comment

The phosphoryl unit in compounds of type $\text{X}_3\text{P}=\text{O}$ contains an electron-rich O atom that can act as a donor to metal centres (Burford, 1992). Several such complexes have been structurally characterized in which the angle at the phosphoryl O atom varies from 130° to 180° depending on the substituents and metal centres. Antimony trihalides can also act as weak acceptors (Pohl, Saak, Lotzand & Haase, 1990). In this paper, we report the synthesis and structure of the title compound $[(\text{Me}_2\text{C}_5\text{H}_8\text{N})_2\text{P}(\text{Cl})\text{O}] \rightarrow \text{SbCl}_3$, (2).



(2)

Compound (2) was obtained in an attempted preparation of [(Me₂C₅H₈N)₂P][SbCl₆] by reacting (Me₂C₅H₈N)₂PCL, (1), with SbCl₅, presumably by hydrolysis. An ORTEP (Johnson, 1965) drawing of the molecule of (2) is shown in Fig. 1. The Sb—O distance of 2.396(4) Å in (2) suggests only a moderate interaction of the Sb atom with the phosphoryl O atom; this distance is much longer than that in Cl₃P=O.SbCl₅ [2.17(2)°; Bovin, 1976; Branden & Lindquist, 1963] reflecting the lower acidity of SbCl₃ compared with SbCl₅ and thus leading to a weaker Sb—O bond in the former. The Sb—Cl distances are in the normal range (Lipka, 1979; Rogers & Jezl, 1994). The P=O distance [1.492(3) Å] as well as the P—O—Sb angle [133.4(2)°] are also consistent with a mainly σ donation with a moderate or low π interaction for the Sb—O bond.

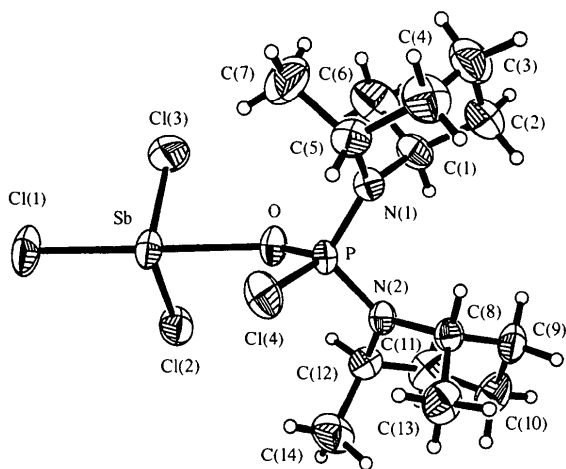


Fig. 1. An ORTEP (Johnson, 1965) drawing of the molecule of compound (2) with displacement ellipsoids plotted at the 50% probability level; only non-H atoms are labelled.

The geometry around antimony, including the lone pair, is distorted trigonal bipyramidal (tbp) [valence-shell electron-pair repulsion (VSEPR) theory] with the O and Cl(1) atoms *trans* to one another; the deviation from tbp to a square-pyramidal structure based on the angles between the planes containing Cl(1), Cl(2), Cl(3) and Cl(2), Cl(3), Cl(4) is *ca* 15%. The Sb—Cl distance [2.460(3)°] *trans* to the Sb—O bond is more than the other two Sb—Cl distances [mean: 2.347 Å], as expected in a tbp geometry.

Although an attempt was made to prepare the analogous 1:1 adduct of [(OCH₂CMe₂CH₂O)P(O)Cl] with SbCl₃, the resulting solid (3) analysed as the 2:1 adduct; the crystals obtained were not good enough to study the nature of the Sb—O interaction.

Experimental

For the preparation of [C₅H₈(Me)₂N]₂PCL, (1), a mixture of *cis*-2,6-dimethylpiperidine (8.2 g, 72 mmol) and triethyl-

amine (7.3 g, 72 mmol) in hexane (50 ml) was added dropwise to a solution of phosphorus trichloride (4.98 g, 36 mmol) in hexane (100 ml) with continuous stirring at 298 K. The reaction mixture was heated under reflux for 48 h and filtered. Evaporation of the solvent followed by vacuum distillation afforded (1) as a liquid (b.p. 383 K/3 mm). Yield: 6.86 g (65%). ¹H NMR (CDCl₃): 1.22 (s, 6H, CH₃), 1.24 (s, 6H, CH₃), 1.25–1.80 (m, 12H, CH₂), 3.70–3.90 (br, 4H, NCH). ³¹P NMR: 156.5. For the preparation of [(C₇H₁₄N)₂P(O)Cl].SbCl₃, (2), a solution of antimony pentachloride (prepared by reacting the trichloride with chlorine) (0.52 g, 1.75 mmol) in dichloromethane (10 ml) was added dropwise with continuous stirring to a solution of (1) (0.5 g, 1.75 mmol) in dichloromethane (10 ml) maintained at 195 K. After stirring for a further 2 h, the solvent was completely removed to give an oil. Attempted crystallization from acetonitrile–hexane (1:10) afforded a small quantity of (2) (0.1 g) as a crystalline material (m.p. 419–421 K). ¹H NMR: 1.31 (s, 6H, CH₃), 1.42 (s, 6H, CH₃), 1.45–2.10 (m, 12H, CH₂), 3.85–4.10 (m, 4H, CHNH). ³¹P NMR: 23.3. Analysis calculated for C₁₄H₂₈Cl₄N₂OPSb: C 31.40, H 5.20, N 5.2%; found: C 31.08, H 5.30, N 4.9%. For the preparation of [(OCH₂CMe₂CH₂O)P(O)Cl]₂.SbCl₃, (3), [(OCH₂CMe₂CH₂O)P(O)Cl] (0.46 g, 2.02 mmol) was reacted with antimony trichloride (0.46 g, 2.02 mmol) in ether (10 + 10 ml) for 24 h. Slow removal of solvent afforded a crystalline solid (m.p. 333 K). ¹H NMR: 0.95 (s, 3H, CH₃), 1.35 (s, 3H, CH₃), 3.90–4.30 (m, 4H, CH₂). ³¹P NMR: –3.36. Analysis calculated for C₁₀H₂₀Cl₅O₇P₂Sb: C 19.50, H 3.28%; found: C 19.70, H 3.25%.

Crystal data

C₁₄H₂₈ClN₂OP.SbCl₃

M_r = 534.9

Triclinic

P $\bar{1}$

a = 10.092(11) Å

b = 10.514(11) Å

c = 12.415(14) Å

α = 70.53(8)°

β = 71.81(8)°

γ = 61.85(7)°

V = 1076(2) Å³

Z = 2

D_x = 1.651 Mg m^{−3}

D_m not measured

Data collection

Siemens–Stoe AED-2

diffractometer

ω – θ scans

Absorption correction: none

2820 measured reflections

2820 independent reflections

2820 reflections with

F > 4 σ (*F*)

Mo *K* α radiation

λ = 0.71073 Å

Cell parameters from 18

reflections

θ = 10–12°

μ = 1.857 mm^{−1}

T = 293 K

Block

0.3 × 0.2 × 0.2 mm

White

Refinement

Refinement on *F*²

R = 0.0322

wR = 0.0853

S = 1.147 (on *F*²)

(Δ/σ)_{max} = 0.105

$\Delta\rho$ _{max} = 0.966 e Å^{−3}

$\Delta\rho$ _{min} = −1.289 e Å^{−3}

Extinction correction: none

2820 reflections
208 parameters
H atoms: see below
 $w = 1/[\sigma(F_o^2) + (0.0501P)^2 + 1.4214P]$
where $P = (F_o^2 + 2F_c^2)/3$

Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

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Piperidinium Diazidotriphenylstannate and Quinuclidinium Diazidotriphenylstannate

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Table 1. Selected geometric parameters (Å, °)

Sb—Cl(1)	2.460 (3)	P—O	1.492 (3)
Sb—Cl(2)	2.345 (3)	P—N(1)	1.621 (4)
Sb—Cl(3)	2.348 (3)	P—N(2)	1.613 (4)
Sb—O	2.396 (4)	P—Cl(4)	2.037 (3)
Cl(2)—Sb—Cl(3)	95.85 (10)	O—P—N(1)	117.4 (2)
Cl(2)—Sb—O	83.78 (12)	O—P—Cl(4)	106.3 (2)
Cl(3)—Sb—O	83.89 (13)	N(2)—P—Cl(4)	108.9 (2)
Cl(2)—Sb—Cl(1)	92.42 (11)	N(2)—P—N(1)	108.8 (2)
Cl(3)—Sb—Cl(1)	92.10 (12)	N(1)—P—Cl(4)	104.6 (2)
Cl(1)—Sb—O	174.15 (7)	P—O—Sb	133.4 (2)
O—P—N(2)	110.4 (2)		

X-ray data for (2) were collected after inserting the crystal into a Lindemann capillary. The structure was solved by conventional methods (*SHELX86*; Sheldrick, 1990) and refined by full-matrix least-squares methods (*SHELXL93*; Sheldrick, 1993). H atoms bonded to C atoms were placed in calculated positions using a riding model. They were assigned isotropic displacement parameters of 1.2 times that of the parent C atom; all non-H atoms were refined anisotropically.

Data collection: *DIF4* (Stoe & Cie, 1988a). Cell refinement: *DIF4*. Data reduction: *REDU4* (Stoe & Cie, 1988b). Molecular graphics: *ORTEP* (Johnson, 1965).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: DE1057). Services for accessing these data are described at the back of the journal.

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 Stoe & Cie (1988a). *DIF4. Diffractometer Control Program*. Version 6.2. Stoe & Cie, Darmstadt, Germany.
 Stoe & Cie (1988b). *REDU4. Data Reduction Program*. Version 6.2. Stoe & Cie, Darmstadt, Germany.

Abstract

Two triphenyltin–azide adducts with cyclic aliphatic amines are shown by X-ray diffraction to be $[LH][(\text{C}_6\text{H}_5)_3\text{Sn}(\text{N}_3)_2]$, where for (I), $L = \text{piperidine}$ ($\text{C}_5\text{H}_{11}\text{N}$), and for (II), $L = \text{quinuclidine}$ ($\text{C}_7\text{H}_{13}\text{N}$). The anions have quasi-trigonal-bipyramidal symmetry, with the azide groups occupying axial positions. In each case, the azide groups are non-equivalent due to one of them hydrogen bonding with the cation. In (I), the NH_2 of the cation hydrogen bonds with azides of adjacent anions forming chains through the lattice. For (II), the asymmetric unit has two cations and anions. One cation singly hydrogen bonds to an azide group of one anion, while the other cation forms a bifurcated hydrogen bond linked to azide groups on the two different anions. The cations have their usual geometries, the piperidinium ion having the expected chair conformation.

Comment

Following our earlier reports on adducts (1/1) of triphenyltin pseudo-halides with various O- and N-donor ligands, with particular reference to their Raman and far-IR spectra (Wharf, Piehler, Sailofsky, Onyszchuk & Simard, 1987, and references therein), we have recently completed a similar study on triphenyltin azide and its adducts (1/1) with a range of O- and N-donor ligands (Wharf, Wojtowski, Lebus, Bowes & Onyszchuk, 1997). While many amines did form 1/1 adducts, some aliphatic amines [$L = \text{morpholine}$, triethylamine, 1,2-bis(dimethylamino)ethane, piperidine, quinuclidine] gave products that appeared from elemental analysis to contain two azide groups which, together with IR data [$\sim 2050 \text{ cm}^{-1}$, $\nu_{\text{as}}(\text{N}_3)$; $\sim 3400 \text{ cm}^{-1}$, $\nu(\text{N—H})$], suggested these compounds were perhaps $[LH^+][\text{Ph}_3\text{Sn}(\text{N}_3)_2^-]$, analogues of the $(\text{CH}_3)_4\text{N}^+$ and $(\text{C}_6\text{H}_5)_4\text{As}^+$ salts with the same anion [IR (cm^{-1}): $\nu_{\text{as}}(\text{N}_3)$, 2050s, 2070sh and 2040s, 2055s, respectively] reported by Barbieri, Bertazzi & Tomarchio (1975). In the same way, the compound having