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#### **Key indicators**

Single-crystal X-ray study T = 183 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.051 wR factor = 0.118 Data-to-parameter ratio = 18.6

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# [Bis(trimethylsilyl)methyl]chloro(phenyl)phosphine

The title compound,  $C_{13}H_{24}ClPSi_2$ , was synthesized *via* the reaction of bis(trimethylsily)methyllithium and dichlorophenylphosphine. The coordination geometry around the P atom is distorted tetrahedral.

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## Comment

The presence of heteroatoms such as Si or P in the  $\alpha$  position of carbanions can increase their stability by hyperconjugation. In recent years, new organic phosphinyl compounds have been synthesized, including  $P \equiv$  and -P = units, etc. (Edwards, 1995). The unique patterns of bonding in these compounds have enriched basic theories in the phosphine subject area and have also provided inspiration for the preparation of new structures. In the past 20 years, many alkali-metal compounds have been synthesized incorporating such species; for example, phosphinyl compounds substituted by benzyl can exhibit unique character when forming complexes with metal cations because of stereochemical effects (Clegg et al., 1998). Much research has also been carried out on the special chemical reactivity and applications of heteroatom compounds in organic synthesis, developing new reactions in which the heteroatom-organic molecule plays the role of substrate or catalyst (White et al., 1999; Chan et al., 1995). Recently, we have synthesized the title phosphinyl compound, (I), and we report here its crystal structure.



In (I), the coordination geometry around P1 is distorted tetrahedral, with the P atom bonded to two C atoms and one Cl atom, and having a lone pair of electrons (Fig. 1 and Table 1). The C2-P1-C1-Si1, Cl1-P1-C1-Si2, and Cl1-P1-C1-Si2 torsion angles are 82.15 (15), -177.22 (10), -150.30 (15) and -49.67 (15)°, respectively.

# Experimental

All reactions were carried out under Ar using standard Schlenk techniques. Diethyl ether was distilled from drying agents and degassed with Ar prior to use. In a 1000 ml round-bottomed flask, bromoform (20.0 ml, 0.23 mol) and trimethylchlorosilane (58.1 ml, 0.46 mol) were added to tetrahydrofuran (300 ml). To this solution, n-butyllithium in n-hexane (157.0 ml, 0.46 mol) was added dropwise

© 2006 International Union of Crystallography All rights reserved at 195 K, and the mixture was allowed to warm to room temperature, stirred for 4 h and filtered. Diethyl ether was distilled from the filtrate, and the product bromobis(trimethylsilyl)methane, a colourless liquid, was then collected by distillation in a vacuum at 321-333 K. In a subsequent reaction, n-butyllithium in n-hexane (19.6 ml, 57.37 mmol) was added quickly to a solution of bromobis(trimethylsilyl)methane (13.66 g, 57.37 mmol) in Et<sub>2</sub>O (20 ml) at 195 K, and the solvent was removed immediately. The raw product was sublimed to give pure bis(trimethylsily)methyllithium. Finally, dichlorophenylphosphine (0.46 ml, 3.39 mmol) was added to a solution of bis(trimethylsily)methyllithium (0.563 g, 3.39 mmol) in Et<sub>2</sub>O (20 ml) at 273 K, and the temperature was allowed to warm to room temperature. The mixture was stirred for 4 h, vielding a white precipitate, which was removed by filtration. The filtrate was concentrated to ca 10 ml and then cooled at 243 K for 3 d to give colourless crystals of the title compound.

V = 859.3 (3) Å<sup>3</sup>

 $D_r = 1.171 \text{ Mg m}^{-3}$ 

Mo  $K\alpha$  radiation  $\mu = 0.44 \text{ mm}^{-1}$ 

Block, colourless

 $0.40 \times 0.30 \times 0.20 \ \mathrm{mm}$ 

3549 measured reflections

2972 independent reflections

2485 reflections with  $I > 2\sigma(I)$ 

T = 183 (2) K

 $R_{\rm int}=0.017$ 

 $\theta_{\rm max} = 25.0^\circ$ 

Z = 2

## Crystal data

C13H24ClPSi2  $M_r = 302.93$ Triclinic.  $P\overline{1}$ a = 8.5958 (19) Å b = 9.069 (2) Å c = 11.794 (3) Å  $\alpha = 75.783 (3)^{\circ}$  $\beta = 74.661 \ (3)^{\circ}$  $\gamma = 87.322 (3)^{\circ}$ 

### Data collection

Bruker SMART CCD diffractometer  $\omega$  scans Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  $T_{\min} = 0.844, T_{\max} = 0.917$ 

#### Refinement

| Refinement on $F^2$             | $w = 1/[\sigma^2(F_o^2) + (0.0587P)^2]$                    |
|---------------------------------|--|
| $R[F^2 > 2\sigma(F^2)] = 0.051$ | + 0.1605P]   |
| $wR(F^2) = 0.118$               | where $P = (F_0^2 + 2F_c^2)/3$                             |
| S = 1.08                        | $(\Delta/\sigma)_{\rm max} < 0.001$                        |
| 2972 reflections                | $\Delta \rho_{\rm max} = 0.41 \text{ e } \text{\AA}^{-3}$  |
| 160 parameters                  | $\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$ |
| H-atom parameters constrained   |  |

## Table 1

Selected geometric parameters (Å, °).

| Cl1-P1    | 2.1067 (11) | Si1-C1    | 1.903 (3)  |
|-----------|-------------|-----------|------------|
| P1-C1     | 1.823 (3)   | Si2-C12   | 1.861 (3)  |
| P1-C2     | 1.837 (3)   | Si2-C13   | 1.861 (3)  |
| Si1-C9    | 1.853 (4)   | Si2-C11   | 1.870 (3)  |
| Si1-C10   | 1.863 (3)   | Si2-C1    | 1.909 (3)  |
| Si1-C8    | 1.866 (3)   |           |            |
| C1-P1-C2  | 103.64 (13) | C2-P1-Cl1 | 96.79 (10) |
| C1-P1-Cl1 | 103.71 (9)  |           |            |
|           |             |           |            |

H atoms were positioned geometrically and allowed to ride during subsequent refinement, with C-H = 0.93, 0.96 and 0.98 Å for those on phenyl, tertiary and methyl C atoms, respectively.  $U_{iso}(H) =$  $1.5U_{eq}(C)$  for the methyl groups and  $U_{iso}(H) = 1.2U_{eq}(C)$  otherwise. The methyl groups were also allowed to rotate about their local threefold axes.



Figure 1

Molecular unit in (I), showing displacement ellipsoids at the 30% probability level. H atoms are shown as spheres of arbitrary radii.



Figure 2 View of the unit-cell contents of (I). H atoms have been omitted.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Bruker, 1999); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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