

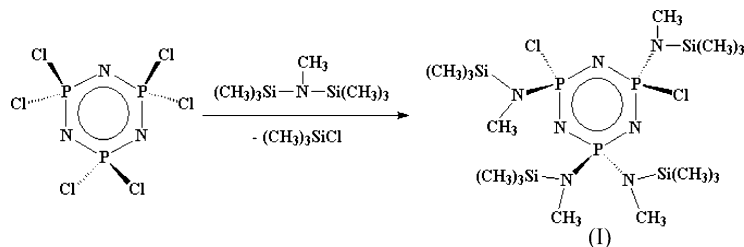
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Key indicators

Single-crystal X-ray study
T = 120 K
Mean $\sigma(\text{N}-\text{C}) = 0.003 \text{ \AA}$
R factor = 0.038
wR factor = 0.097
Data-to-parameter ratio = 17.2For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.***trans*-2,6-Dichloro-2,4,4,6-tetrakis[methyl-(trimethylsilyl)amino]-1,3,5,2 λ^5 ,4 λ^5 ,6 λ^5 -triazatriphosphorine**The title tetrasubstituted phosphazene, $\text{C}_{16}\text{H}_{48}\text{Cl}_2\text{N}_7\text{P}_3\text{Si}_4$, is formed by the reaction of $\text{P}_3\text{N}_3\text{Cl}_6$ with heptamethyldisilazane, HpMDSA. The compound is the first *trans* non-geminal tetrasubstituted derivative of this class of compounds to be structurally characterized. The molecule lies on a crystallographic twofold rotation axis.Received 7 December 2006
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Comment

Phosphazenes date back to the late 20th century (Allcock, 1972; Allen, 1991). Most studies have focused on nucleophilic substitutions of hexahalogeno-*cyclo*-triphosphazene $\text{P}_3\text{N}_3\text{X}_6$ ($\text{X} = \text{F}, \text{Cl}, \text{Br}$) (Allcock, 1987, 1972, 2003; Allen, 1987, 1991; Gleria & De Jaeger, 2002). Little attention has been paid to reactions of cyclic phosphazenes with silylated amines (Mattogno & Tarli, 1964, Mattogno *et al.*, 1965; Fields & Allen, 1974) although Fields & Allen (1974) prepared $\text{P}_3\text{N}_3\text{Cl}_4[\text{NHSi}(\text{CH}_3)_3]_2$, and its crystal structure was determined by Allen *et al.* (1988).

Many crystal structures of amido derivatives of chlorophosphazenes have been reported (*e.g.* Polder & Wagner, 1976; Bartlett *et al.*, 2006; Ahmed & Gabe, 1975, Ahmed & Fortier, 1980). Compound (I) is the first *trans* non-geminal tetrasubstituted derivative of this class of compounds to be structurally characterized (Fig. 1). The molecule lies on a crystallographic twofold rotation axis. A comparison of corresponding bond lengths with literature data (Allen *et al.*, 1988; Ahmed & Gabe, 1975, Ahmed & Fortier, 1980) shows that there is no significant difference in the length of either the exocyclic P–Cl bond or the P–N(Me) bond. However, a remarkable deviation was found in the P–N bond distances of the P_3N_3 ring. The P–N lengths in compound (I) are in the range 1.591 (2)–1.6168 (18) Å, compared with literature values of 1.574–1.586 Å (Allen *et al.*, 1988; Ahmed & Gabe, 1975, Ahmed & Fortier, 1980). A significant elongation was also found for N–C bonds [1.484 (3)–1.491 (3) Å in (I) compared to 1.456–1.473 Å in *trans*- $\text{P}_3\text{N}_3\text{Cl}_4[\text{N}(\text{CH}_3)_2]_2$ (Ahmed & Fortier, 1980)]. The P1–N2–P1¹ [symmetry code:

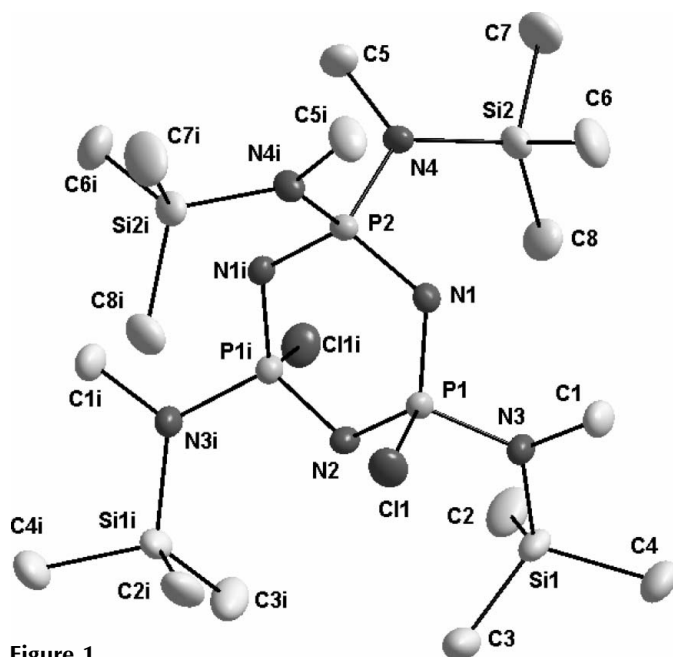


Figure 1
The molecular structure of (I). Displacement ellipsoids represent 50% probability levels. All H atoms have been omitted for clarity. [Symmetry code: (i) $-x, y, -z + \frac{1}{2}$]

(i) $-x, y, -z + \frac{1}{2}$] and N1–P1–N2 angles of the phosphazene ring are comparable to those of similar phosphazene amido-derivatives (Allen *et al.*, 1988; Ahmed & Gabe, 1975, Ahmed & Fortier, 1980) but there are some differences in the values of N1–P2–N1ⁱ and P1–N1–P2 angles. The former angle increased to 115.28 (14)° (*cf* 111.57°; Allen *et al.*, 1988) while the latter decreased to 121.96 (11)° (124.80°; Allen *et al.*, 1988). The P₃N₃ ring is slightly distorted with atoms N1 and N1ⁱ atoms deviating from the P1,N2,P1ⁱ,P2 plane by 0.2181 (18) Å. The angle between the P1,N2,P1ⁱ,P2 ring and N4,P2,N4ⁱ is 88.12 (4)° (*cf* 87.14°; Allen *et al.*, 1988) while the angle between the ring plane P1,P1ⁱ,P2 and planes formed by atoms Cl1,P1,N3 or Cl1ⁱ,P1ⁱ,N3ⁱ decreased to 81.17 (3)° (85.43–89.60°; Ahmed & Gabe, 1975, Ahmed & Fortier, 1980). This deviation can be attributed to the presence of two bulky N(CH₃)Si(CH₃)₃ groups on P2.

No important intermolecular interactions were found in the crystal structure of (I). The molecules of (I) form a system of two coplanar [011] planes. The appropriate description of the structure motif is [ABAB]_∞.

Experimental

Compound (I) was prepared by reaction of P₃N₃Cl₆ with heptamethyldisilazane in acetonitrile (see Scheme). P₃N₃Cl₆ (0.201 g, 0.6 mmol) dissolved in acetonitrile (5 cm³) and HpMDSA (0.416 g, 2.4 mmol) were placed into a Schlenk vessel equipped with a screw cap. The reaction mixture was heated on an oil bath at 399–405 K for 48 h. After cooling, the solvent and resulting trimethylchlorosilane were completely evaporated and the remaining solid was again dissolved in acetonitrile. This solution was stored at 248 K for several

weeks. Crystals of (I) for suitable X-ray analysis were removed from the solution under a dry nitrogen atmosphere.

Crystal data

C₁₆H₄₈Cl₂N₇P₃Si₄
M_r = 614.78
Monoclinic, C2/c
a = 17.470 (3) Å
b = 11.360 (2) Å
c = 17.270 (3) Å
β = 101.21 (3)°
V = 3362.0 (10) Å³

Z = 4
D_x = 1.215 Mg m⁻³
Mo Kα radiation
μ = 0.50 mm⁻¹
T = 120 (2) K
Prism, white
0.25 × 0.20 × 0.20 mm

Data collection

Kuma KM-4 CCD diffractometer
ω scans
Absorption correction: none
13131 measured reflections

2829 independent reflections
2569 reflections with I > 2σ(I)
R_{int} = 0.030
θ_{max} = 25.0°

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.038
wR(F²) = 0.097
S = 1.04
2620 reflections
154 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.049P)^2 + 7.9146P]$
where $P = (F_o^2 + 2F_c^2)/3$
(Δ/σ)_{max} = 0.001
Δρ_{max} = 0.54 e Å⁻³
Δρ_{min} = -0.34 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Cl1–P1	2.0730 (9)	N3–C1	1.491 (3)
P1–N1	1.591 (2)	P2–N1 ⁱ	1.6168 (18)
P1–N2	1.6010 (15)	P2–N4	1.6645 (18)
P1–N3	1.6491 (19)	P2–N4 ⁱ	1.6645 (18)
N1–P2	1.6168 (18)	N4–C5	1.484 (3)
N2–P1 ⁱ	1.6010 (15)		
N1–P1–N2	118.79 (11)	P1–N2–P1 ⁱ	119.36 (17)
N3–P1–Cl1	103.57 (8)	N1–P2–N1 ⁱ	115.28 (14)
P1–N1–P2	121.96 (11)	N4–P2–N4 ⁱ	107.19 (14)

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

H atoms were positioned geometrically (C–H = 0.98 Å) and refined as riding with U_{iso}(H) = 1.5U_{eq}(C).

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2003); cell refinement: *KM-4 Software* (Kuma Diffraction, 2006); data reduction: *KM-4 Software*; program(s) used to solve structure: *SHELXTL* (Bruker, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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