Diffraction intensities from the crystals were generally weak and only 49% of the reflections were observed with  $I > 2\sigma(I)$ , as evidenced by the high  $R_{int}$  value. All H-atom positions (except for H1O and H2O) were calculated using a riding model and were considered with fixed isotropic displacement parameters in all refinements. The H1O and H2O atoms were found in the difference Fourier maps calculated at the end of the refinement process as small positive electron densities and were not refined.

Data collection: Nonius Diffractometer Control Software (Nonius, 1993). Cell refinement: Nonius Diffractometer Control Software. Data reduction: REDU4 (Stoe & Cie, 1991). Program(s) used to solve structure: SHELXS86 (Sheldrick, 1985). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ORTEPII (Johnson, 1976). 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: JZ1285). Services for accessing these data are described at the back of the journal.

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# 2,4-Cl<sub>2</sub>-6,9-exo, endo-(PMe<sub>2</sub>Ph)<sub>2</sub>- $arachno-B_{10}H_{10}$

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### Abstract

The title compound, 2,4-dichloro-6-*exo*-9-*endo*bis(dimethylphenylphosphino)-*arachno*-decaborane(10)  $[(PMe_2Ph)_2B_{10}H_{10}Cl_2 \text{ or } C_{16}H_{32}B_{10}Cl_2P_2]$ , has a typical *arachno* ten-vertex cluster geometry, with one of the phosphine ligands bound *exo*-polyhedrally and the other in an uncommon *endo* configuration.

#### Comment

Ten years ago, Fontaine & Kennedy (1987) reported the identification by NMR spectroscopy of the previously unknown exo, endo-isomer, (I), of [6,9-(PMe<sub>2</sub>Ph)<sub>2</sub>arachno-B<sub>10</sub>H<sub>12</sub>], formed concomitantly with the corresponding conventional exo, exo-isomer, (II). We have recently (Dörfler et al., 1997) confirmed the structures of (I) and (II) by single-crystal X-ray diffraction studies, and also confirmed the structure of (III), the 2-bromosubstituted analogue of (I). These structural studies suggested some differential perturbation of the cluster that arises from the presence of an endo-P versus exo-P substituent. However, in (III), there are some deviations in structure that may be due to the presence of the single boron-bound halogen and this substituent might equally contribute to apparent exo versus endo perturbation. To examine further for patterns of structure flexing which might be attributable to exolendo substitution, and for any effect of a halogen substituent upon this, we have determined the structure of the symmetrically dichlorosubstituted title compound, (IV).





The molecule of (IV) has approximate  $C_s$  molecular symmetry (ignoring the dispositions of the P-bound organic groups). As expected, the cluster has typical arachno-decaboranyl character, exemplified by the presence of two H atoms symmetrically bridging the B5-B10 and B7-B8 connectivities; the lengths of these two vectors [1.879(3) and 1.886(3) Å, respectively] are further consistent with the arachno descriptor. [These distances are typically longer (ca 2.0 Å) in a nido ten-vertex cluster.] Atoms Cl2 and Cl4 are bound at 1.823 (2) and 1.827 (2) Å from B2 and B4, respectively, these distances being typical of exo-(B-Cl) separations in polyhedral boranes (Fletcher et al., 1996). Interestingly, there is no significant shortening [compared with the parent (I)] of those B-B connectivities involving chloro-substituted B2 and B4, in contrast to the apparent effect of the single bromo substituent in (III).

However, the most significant feature of the geometry determined for (IV) is the structural confirmation of a further example of an exo, endo-6,9-disubstituted arachno-decaborane. The two P atoms, P1 and P2, are bound at distances of 1.922(2) and 1.924(2) Å from their respective parent B atoms, B6 and B9, which compare with a range of 1.912 (3)-1.928 (4) Å for exo, exo-(II) and exo, endo-(III). This contrasts with the situation in exo, endo-(I), where the exo-P atom is 1.9145 (14) Å from B6, but the endo-P atom is much further [1.941 (2) Å] from B9. In addition, the B2-B6-P1 and B4-B9-P2 angles in (IV) [117.38(14) and 157.66 (15)°, respectively] are both slightly more obtuse than in (I), where they are 110.20(8) and 155.86 (10)°, respectively. This may arise from the steric and/or electronic influence of the two chloro substituents, but may equally be due simply to a slightly different molecular-packing arrangement in the solid state.

There are, however, some small but potentially significant effects upon the structure of the decaboranyl cluster that may be attributable to the presence of an *endo*-P versus exo-P substituent. The three B—B connectivities involving B9 are slightly longer than the three

corresponding vectors involving B6. Similarly, there is some enlargement of the B5-B10-B9 and B7-B8-B9 angles [116.25(14) and 115.68(14)°, respectively] in (IV) compared with the equivalent angles to B6, namely, B6-B5-B10 and B6-B7-B8 [112.66(14) and 113.42 (14)°, respectively]; and the B1-B4-B9 and B3-B4-B9 angles [118.20 (16) and 118.21 (16)°, respectively] are slightly larger than B1-B2-B6 and B3—B2—B6 [116.24 (15) and  $115.40 (16)^{\circ}$ , respectively]. The pattern of these perturbations mirrors that seen in (I), and has been attributed to the presence of endo versus exo substituents (Dörfler et al., 1997). Perhaps surprisingly, it has been found that (IV) does not readily isomerize on heating to its exo, exo counterpart (Dörfler et al., 1997), despite the significant weakening of the bonds to the endo-P-substituted vertex B9. Such lengthening in non-chlorinated (I) has been suggested (Dörfler et al., 1997) to facilitate the thermal conversion of (I) to (II). The chloro substituents, therefore, appear to have either a steric and/or an electronic influence on the course of such a reaction. We are currently probing this phenomenon further.



Fig. 1. Perspective view of a molecule of (IV) drawn with 40% probability ellipsoids and with H atoms shown as small circles of arbitrary radii for clarity.

#### Experimental

Compound (IV) was prepared as reported elsewhere (Dörfler et al., 1997). Suitable single crystals were obtained by the slow mutual diffusion of hexane and a dichloromethane solution at 293 K.

Crystal data

$C_{16}H_{32}B_{10}Cl_2P_2$	Cu $K\alpha$ radiation
$M_r = 465.36$	$\lambda = 1.54186 \text{ Å}$

Cell parameters from 74

 $0.50 \times 0.29 \times 0.15$  mm

3693 reflections with

3 standard reflections

frequency: 60 min

intensity decay: 3.9%

 $I > 2\sigma(I)$  $R_{\rm int} = 0.0229$ 

 $\theta_{\rm max} = 64.39^{\circ}$ 

 $h = -11 \rightarrow 11$ 

 $k = -22 \rightarrow 22$ 

 $l = -14 \rightarrow 15$ 

reflections

 $\theta = 30.1 - 39.7^{\circ}$ 

T = 160(2) K

Prism

Colourless

 $\mu = 3.578 \text{ mm}^{-1}$ 

Monoclinic  $P2_1/n$  a = 9.7987 (8) Å b = 19.1424 (14) Å c = 13.3366 (11) Å  $\beta = 98.816$  (7)° V = 2472.0 (3) Å<sup>3</sup> Z = 4  $D_x = 1.250 \text{ Mg m}^{-3}$  $D_m$  not measured

Data collection

Stoe Stadi-4 four-circle diffractometer  $\omega/\theta$  scans Absorption correction:  $\psi$  scans (North *et al.*, 1968)  $T_{min} = 0.338$ ,  $T_{max} = 0.585$ 8127 measured reflections 4091 independent reflections

Refinement

Refinement on  $F^2$  $(\Delta/\sigma)_{\rm max} = 0.032$  $\Delta \rho_{\rm max} = 0.513 \ {\rm e} \ {\rm \AA}^{-3}$ 
$$\begin{split} R[F^2 > 2\sigma(F^2)] &= 0.034 \\ wR(F^2) &= 0.090 \end{split}$$
 $\Delta \rho_{\rm min} = -0.377 \ {\rm e} \ {\rm \AA}^{-3}$ Extinction correction: none S = 1.0544091 reflections Scattering factors from 315 parameters International Tables for H atoms: see below Crystallography (Vol. C)  $w = 1/[\sigma^2(F_o^2) + (0.0463P)^2$ + 1.2956P] where  $P = (F_o^2 + 2F_c^2)/3$ 

Table 1. Selected geometric parameters (Å, °)

	-	-	
B1-B2	1.757 (3)	B4—B9	1.758 (3)
B1B4	1.760 (3)	B4—B8	1.761 (3)
B1—B5	1.775 (3)	B4—B10	1.764 (3)
B1—B10	1.785 (3)	B4—Cl4	1.827 (2)
B1—B3	1.838 (3)	B5—B6	1.878 (3)
B2—B6	1.742 (3)	B5-B10	1.879 (3)
B2—B3	1.758 (3)	B6—B7	1.846 (3)
B2—B5	1.764 (3)	B6—P1	1.922 (2)
B2—B7	1.767 (3)	B7B8	1.886 (3)
B2C12	1.823 (2)	B8—B9	1.887 (3)
B3—B4	1.750 (3)	B9—B10	1.888 (3)
B3—B7	1.779 (3)	B9—P2	1.924 (2)
B3—B8	1.785 (3)		
B6—B2—B1	116.24 (15)	B10-B4-Cl4	119.17 (14)
B6—B2—B3	115.40(16)	B6-B5-B10	112.66 (14)
B6B2C12	115.57 (13)	B7—B6—B5	104.85 (14)
B1B2C12	118.38(14)	B2—B6—P1	117.38 (14)
B3—B2—C12	118.05 (14)	B7—B6—P1	122.38 (13)
B5—B2—C12	119.64 (14)	B5—B6—P1	120.33 (13)
B7—B2—C12	118.87 (14)	B6-B7-B8	113.42 (14)
B3—B4—B9	118.21 (16)	B7—B8—B9	115.68 (14)
B9—B4—B1	118.20(16)	B8—B9—B10	102.33 (14)
B3—B4—Cl4	118.63 (14)	B4—B9—P2	157.66 (15)
B9—B4—Cl4	112.19(13)	B8—B9—P2	111.47 (13)
B1B4Cl4	118.45 (14)	B10-B9-P2	111.90 (13)
B8—B4—Cl4	118.98(14)	B5—B10—B9	116.25 (14)

B—B—B acute angles are in the range 57.05 (12)–64.85  $(13)^{\circ}$ .

Phosphine H atoms were constrained to idealized positions with a riding model including free rotation of the methyl groups and with PPh units restrained to have  $C_{2v}$  symmetry.

Borane H atoms were located in difference Fourier maps and all associated parameters were refined freely.

Data collection: *STADI*4 (Stoe & Cie, 1996a). Cell refinement: *STADI*4. Data reduction: *X-RED* (Stoe & Cie, 1996b). Program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997). Molecular graphics: *ORTEX5* (McArdle, 1995). Software used to prepare material for publication: *SHELXL*97 and a local program.

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## (1*R*,7*S*,8a*S*)-1,7-Dihydroxyindolizidine Hydrate

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#### Abstract

The title compound [(1R,7S,8aS)-1,7-dihydroxyperhydroindolizine hydrate, C<sub>8</sub>H<sub>15</sub>NO<sub>2</sub>.H<sub>2</sub>O] crystallizes withone water solvate molecule, which is involved in stronghydrogen bonds with the hydroxyl O atoms. Bondlengths and angles in the molecule are in agreementwith literature values.