

Polyhedral azaborane chemistry,
6-(C₅H₅N)-*arachno*-4-NB₈H₁₁Caroline Price, Udo Dörfler, John D. Kennedy and Mark
Thornton-Pett*School of Chemistry, The University of Leeds, Leeds LS2 9JT, England
Correspondence e-mail: marktp@chem.leeds.ac.uk

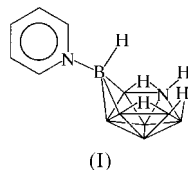
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The title compound, 6-pyridyl-4-*arachno*-nonaborane(11), C₅H₁₆B₈N₂, has an *arachno* nine-vertex [4-NB₈H₁₁] cluster structure with a pyridine ligand in the *exo*-6-position. The cluster has close geometric similarities to the thiaborane and carbaborane analogues.

Comment

There is current interest in smaller heteroboranes (Davison *et al.*, 2000). Our own interest in the nine-vertex azaboranes stems firstly from the fact that these species are not well investigated in an absolute sense, but clearly have potentially very interesting chemistry (Dörfler *et al.*, 1997), and also in that nine-vertex *arachno* species in general have a propensity to couple upon simple heating to give big-borane 'macropolyhedral' clusters, as in thiaborane, platinaborane and iridaborane chemistry (Bould *et al.*, 1999, and references therein). In main-group heteroborane chemistry in particular, [6-(SMe₂)-*arachno*-4-SB₈H₁₀] gives the interesting species [S₂B₁₇H₁₉(SMe₂)] with a hexagonal pyramidal structural feature (Kaur *et al.*, 1996). The nine-vertex thiaboranes [6-*L*-*arachno*-4-SB₈H₁₀] and carbaboranes [6-*L*-*arachno*-4-CB₈H₁₂] are well characterized structurally (Holub, Kennedy & Štíbr, 1994; Holub, Štíbr *et al.*, 1994; Plešek *et al.*, 1994). Several corresponding azaborane analogues [6-*L*-*arachno*-4-NB₈H₁₁], where *L* is a two-electron ligand, are known (Jelínek *et al.*, 1994). However, structural work on these azaboranes is lacking, although the related 'parent' non-ligand species (*arachno*-4-NB₈H₁₃) itself was examined crystallographically



some time ago (Baše *et al.*, 1975; Ragatz *et al.*, 1975). Here we report the structure of the pyridine derivative [6-(C₅H₅N)-*arachno*-4-NB₈H₁₁], (I).

The molecular structure (Fig. 1) closely resembles those of the thiaborane and carbaborane analogues [SB₈H₁₀(N₄C₆H₆)] and [CB₈H₁₂(N₄C₆H₁₂)], where N₄C₆H₁₂ is hexamethylene tetramine. Apart from the variation of the nitrogen ligands between NC₅H₅ and N₄C₆H₆, which does not affect the cluster interatomic distances significantly, the compounds differ in the isolobal cluster constituents, CH₂, NH and S in the 4-position. There are obvious corresponding differences in the boron-to-heteroatom distances, but the other interatomic distances within the clusters are very similar. For all three classes of compound, there appears to be an asymmetry in the bonding of the bridging H atoms, with the shorter distances being towards B8, although this is not so marked for H78 in the title compound. A similar asymmetry is also noted for the four bridging H atoms in the parent non-ligand analogue NB₈H₁₃ (Ragatz *et al.*, 1975).

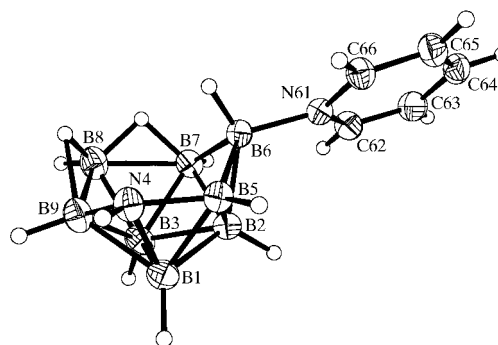


Figure 1

Perspective view of a single molecule of (I) drawn with 40% probability ellipsoids and with H atoms shown as small circles of artificial radii for clarity.

Experimental

The title compound may be prepared by the action of pyridine on [*arachno*-4-NB₈H₁₃], following a previously detailed procedure (Jelínek *et al.*, 1994). The crystallographic sample was obtained from CHCl₃/pentane.

Crystal data

C₅H₁₆B₈N₂
M_r = 190.68
Orthorhombic, Pna2₁
a = 14.36180 (10) Å
b = 10.46000 (10) Å
c = 7.57790 (10) Å
V = 1138.39 (2) Å³
Z = 4
D_x = 1.113 Mg m⁻³

Mo Kα radiation
Cell parameters from 26 404
reflections
θ = 2.41–30.48°
μ = 0.056 mm⁻¹
T = 190 (2) K
Prism, yellow
0.24 × 0.18 × 0.06 mm

Data collection

Nonius KappaCCD area-detector
diffractometer
Area-detector φ and ω scans
Absorption correction: multi-scan
(Blessing, 1995)
T_{min} = 0.987, T_{max} = 0.997
26 404 measured reflections

3204 independent reflections
R_{int} = 0.042
θ_{max} = 30.48°
h = -20 → 20
k = -14 → 14
l = -10 → 10

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.093$
 $S = 1.067$
 3204 reflections
 185 parameters
 H atoms treated by a mixture of
 independent and constrained
 refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0562P)^2 + 0.0813P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.14 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$

program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990);
 program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997);
 molecular graphics: *ORTEX5* (McArdle, 1995); software used to
 prepare material for publication: *SHELXL97*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: BR1272). Services for accessing these data are described at the back of the journal.

Table 1

Selected geometric parameters (\AA , $^\circ$).

B6—N61	1.5671 (13)	B8—B9	1.8497 (17)
B6—B7	1.7798 (14)	N4—B9	1.4989 (14)
B6—B5	1.8287 (15)	N4—B5	1.5180 (14)
B7—B8	1.8616 (18)	N4—B1	1.6064 (15)
N61—B6—B7	121.09 (8)	B9—N4—B5	123.13 (8)
N61—B6—B5	116.90 (7)	B9—N4—B1	73.30 (7)
B7—B6—B5	106.47 (7)	B5—N4—B1	70.20 (7)
B6—B7—B8	112.70 (8)	N4—B5—B6	112.74 (8)
B9—B8—B7	101.84 (8)	N4—B9—B8	121.10 (9)

The title compound crystallizes in the polar orthorhombic space group $Pna2_1$. The direction of the polar axis is ambiguous since the structure does not possess significant anomalous scatterers of Mo $K\alpha$ radiation. An enantiopole parameter is therefore not quoted. Pyridine-H atoms were constrained to calculated positions with isotropic displacement parameters equal to $1.2U_{\text{eq}}$ of the parent C atom. Cluster-associated H atoms were located *via* Fourier difference syntheses and all associated parameters were refined freely.

Data collection: *KCCD* (Nonius, 1998); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1996); data reduction: *DENZO-SMN*;

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