

# Monocarbaborane anion chemistry. The substantiation of the C-arylated $[\text{PhCB}_6\text{H}_6]^-$ seven-vertex *closo* cluster anion by single-crystal synchrotron X-ray diffraction analysis

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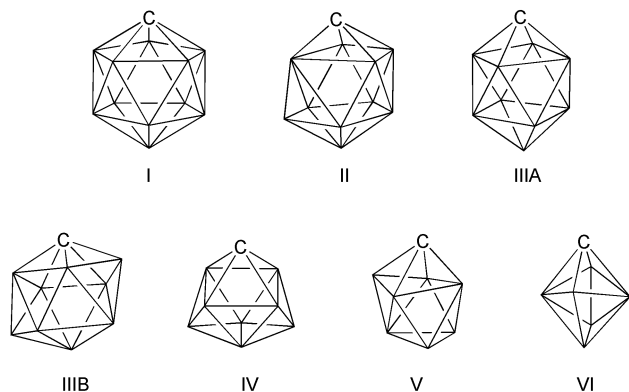
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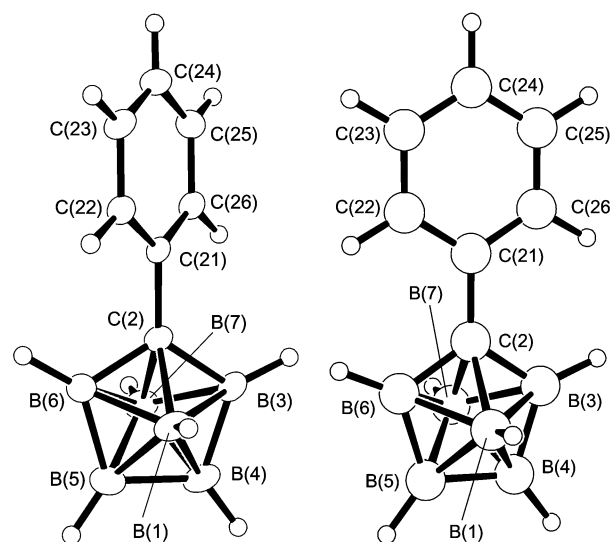
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Single-crystal synchrotron X-ray diffraction analysis of the  $[\text{NEt}_4]^+$  salt of the  $[\text{2-Ph-closo-2-CB}_6\text{H}_6]^-$  anion, obtained from  $[\text{4-Ph-arachno-4-CB}_8\text{H}_{13}]$  and  $\text{NEt}_3$  in refluxing toluene, substantiates the anion as the first structurally characterised seven-vertex monocarbaborane.

The general chemistry of the monocarbaboranes, compared to boranes on one hand and dicarbaboranes on the other, is sparsely investigated.<sup>1</sup> One reason for this has been an absence of convenient synthetic protocols. The Breilochs Reaction<sup>2</sup> of aldehydes with  $\text{B}_{10}\text{H}_{14}$  in alkaline solution now gives ready access to ten-vertex  $\{\text{CB}_9\}$  residues, which, *via* cluster-Aufbau and cluster-dismantling reactions, can then effect entries also into eight-vertex  $\{\text{CB}_7\}$ , nine-vertex  $\{\text{CB}_8\}$ , eleven-vertex  $\{\text{CB}_{10}\}$  and twelve-vertex  $\{\text{CB}_{11}\}$  systems.<sup>2-6</sup> For example, using the benzaldehyde entry into the  $\{\text{PhCB}_9\}$  system as a model, we have recently reported the complete series of *closo* C-phenylated monocarbaborane anions  $[\text{1-Ph-closo-1-CB}_{11}\text{-H}_{11}]^-$  **1**,  $[\text{2-Ph-closo-2-CB}_{10}\text{H}_{10}]^-$  **2**,  $[\text{1-Ph-closo-1-CB}_9\text{H}_9]^-$  **3a**,  $[\text{2-Ph-closo-2-CB}_9\text{H}_9]^-$  **3b**,  $[\text{4-Ph-closo-4-CB}_8\text{H}_8]^-$  **4** and  $[\text{1-Ph-closo-1-CB}_7\text{H}_7]^-$  **5** (schematic cluster structures **I**, **II**, **III A**, **III B**, **IV** and **V** respectively).<sup>4-6</sup> In the consideration of this sequence, it may be noted that C-arylation of monocarbaborane anions has hitherto often been recognised as a difficult target.<sup>7</sup>



The next member in this descending sequence is the seven-vertex  $[\text{2-Ph-closo-2-CB}_6\text{H}_6]^-$  anion **6** (schematic cluster structure **VI**). Descriptions of  $\{\text{CB}_6\}$  cluster species are essentially unknown,<sup>1</sup> but anion **6** has very recently been tentatively identified by NMR spectroscopy in a product mixture from which the eight-vertex  $[\text{1-Ph-closo-1-CB}_7\text{H}_7]^-$  anion **5** was obtained as the major product.<sup>5</sup> We can now support this NMR identification by DFT B3LYP/6-31G\* geometry optimizations of the



**Fig. 1** Crystallographically determined (left) and DFT B3LYP/6-31G\* geometry-optimized (right) molecular structures of the  $[\text{2-Ph-closo-2-CB}_6\text{H}_6]^-$  seven-vertex cluster anion **6**. Selected dimensions in Å are as follows: from the crystallographic determination: C(2)–C(21) 1.492(2), C(2)–B(1) 1.744(2), C(2)–B(3) 1.559(2), C(2)–B(6) 1.555(2), C(2)–B(7) 1.736(2) and interboron distances 1.645(3) to 1.825(3); for the calculated structure: C(2)–C(21) 1.518, C(2)–B(1) 1.753, C(2)–B(3) 1.538, C(2)–B(6) 1.538, and C(2)–B(7) 1.753 and interboron distances 1.611 to 1.798.

molecular structure (Fig. 1, right), and thence by a confirmatory prediction using DFT-GIAO//B3LYP/6-31G\* calculations of the boron nuclear shieldings that reasonably match the observed  $^{11}\text{B}$  NMR chemical shifts.<sup>‡</sup> The unsubstituted analogue  $[\text{HCB}_6\text{H}_6]^-$  has similarly been identifiable very recently in a  $[\text{HCB}_7\text{H}_7]^-$  product mixture.<sup>8</sup> More definitively, however, we now report the unequivocal structural substantiation of the C-phenylated species **6** by single-crystal synchrotron X-ray diffraction work. As far as we are aware this is the first clear structural substantiation in this hitherto elusive  $\{\text{CB}_6\}$  monocarbaborane cluster system.

Thus, a solution of  $[\text{4-Ph-arachno-4-CB}_8\text{H}_{13}]$  (compound **7**, 0.5 g, 2.7 mmol)<sup>5</sup> and  $\text{NEt}_3$  (5 ml) in toluene (15 ml) was heated at reflux for 18 hours under dinitrogen, cooled to room temperature, and the volatile organic solvents removed *in vacuo*. In air, the residue was dissolved in 5% aqueous HCl (50 ml), filtered, and  $[\text{NEt}_4]^+\text{Cl}^-$  (0.5 g, 3.0 mmol) in water (30 ml) added. The resulting white precipitate was filtered off and dried *in vacuo*, and shown by NMR spectroscopy to be a mixture of the  $[\text{NEt}_4]^+$  salts of the  $[\text{1-Ph-closo-1-CB}_7\text{H}_7]^-$  anion **5** and the new  $[\text{2-Ph-closo-2-CB}_6\text{H}_6]^-$  anion **6** (combined amount 0.37 g,

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ca. 2.2 mmol, ca. 82%) in a ratio of **5** : **6** of 60 : 40, corresponding to unseparated yields of ca. 50 and 30% respectively. The extreme similarities of the solubility and chromatographic properties of salts of **5** and **6** have so far precluded a clean bulk separation in our hands. However, crystallisation from a concentrated solution in (CH<sub>3</sub>)<sub>2</sub>CO that was overlaid with a ca. five-fold excess of Et<sub>2</sub>O gave a mixture of crystals of the [NEt<sub>4</sub>]<sup>+</sup> salts of both **5** and **6**. Manual separation of individual crystals, followed by unit-cell measurements on each using conventional sealed-tube X-radiation, distinguished the salt of **6** (orthorhombic, *a* = 14.438(3), *b* = 15.762(3) and *c* = 16.877(4) Å at 150 K) from that previously established<sup>5</sup> for the salt of **5** (monoclinic, *a* = 9.5758(2), *b* = 16.5323(4), *c* = 12.2821(3) Å and β = 98.0110(10)° at 150 K). Crystals of the air-stable [NEt<sub>4</sub>]<sup>+</sup> salt of the anion **6** thus isolated were very small, and required the use of synchrotron X-radiation (wiggler-generated from 2 GeV electrons at 150–250 mA, station 9.8, CCLRC, Daresbury, UK)<sup>9</sup> for sufficient diffraction intensity for molecular structure determination. The analysis of the diffraction data gave a clean result, with no disorder apparent.

The solid-state structure of anion **6** thus determined (Fig. 1, left)§ clearly mimics the pseudo-gas-phase structure arising from DFT calculations (Fig. 1, right). The only notable difference is in the rotamer angle about the C(cluster)–C(phenyl) bond-axis, but this difference is not significant, as energy barriers to phenyl-group/cluster-group contrarotation will be small<sup>10</sup> and easily overcome by crystal-packing forces. The seven-vertex cluster architecture is that of a pentagonal bipyramidal closed deltahedron, in accord with the Williams–Wade paradigm.<sup>11</sup> The all-boron analogue [*closo*-B<sub>7</sub>H<sub>7</sub>]<sup>2-</sup> was recognised some time ago,<sup>12</sup> but has not been structurally characterised, although the perbrominated [*closo*-B<sub>7</sub>Br<sub>7</sub>]<sup>2-</sup> species has,<sup>13</sup> and seven-vertex closed dicarbaboranes are well-established.<sup>1</sup> In anion **6** the carbon atom takes one of the positions of lower cluster-connectivity four, rather than one of higher cluster-connectivity five, also in accord with long-recognised general behavioural patterns.<sup>11,14</sup> The identification and substantiation of this new, stable, fundamental monocarbaborane type augurs well for the continued development of monocarbaborane chemistry, for example in an extension of the ‘least-coordinating anion’ chemistry<sup>15</sup> of the cluster species of *closo* {CB<sub>*n*</sub>} character for the study of and application of very acidic systems. We currently plan experimentation to attempt to increase the proportion of anion **6** in the product mixture at the expense of **5**, and also to attempt to find conditions for efficient bulk separation and purification.

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## Notes and references

‡ Measured NMR data for the [2-Ph-*closo*-2-CB<sub>6</sub>H<sub>6</sub>]<sup>-</sup> anion **6**, [NEt<sub>4</sub>]<sup>+</sup> salt in (CD<sub>3</sub>)<sub>2</sub>CO at 294–299 K, ordered as assignment δ(<sup>11</sup>B)/ppm [δ(<sup>1</sup>H)/ppm], are as follows: BH(3,6) +6.4 [+4.11], BH(4,5) +0.5

[+3.55], BH(1,7) –19.5 [–0.16]; additionally δ(<sup>1</sup>H)(Ph) ca. +7.39 to +7.03 (5H, compact overlapping multiplet), and δ(<sup>1</sup>H)(Et) at +3.47 (8H, quartet), +1.39 (12H, triplet), also δ(<sup>13</sup>C)(Ph) +143.02 (1C), +127.66 (2C), +127.16 (2C) and +124.25 (1C), with δ(<sup>13</sup>C)(cluster) +85.8 and δ(<sup>13</sup>C)(Et) +7.07 and +52.47 ppm. Calculated δ(<sup>11</sup>B) values by DFT-GIAO//B3LYP/6-31G\* are as follows: <sup>16</sup> B(3,6) +2.8, B(4,5) –2.1 and B(1,7) –19.7 ppm.

§ Crystal data for the [NEt<sub>4</sub>]<sup>+</sup> salt of the [2-Ph-*closo*-2-CB<sub>6</sub>H<sub>6</sub>]<sup>-</sup> anion **6**: C<sub>15</sub>H<sub>31</sub>B<sub>6</sub>N: *M* = 290.27, orthorhombic, colourless rod, 300 × 30 × 30 μm, from (CH<sub>3</sub>)<sub>2</sub>CO–Et<sub>2</sub>O, space group *Pbca*, *a* = 14.438(3), *b* = 15.762(3), *c* = 16.877(4) Å, *U* = 3840.6(14) Å<sup>3</sup>, *D*<sub>calc</sub> = 1.004 Mg m<sup>-3</sup>, *Z* = 8, λ = 0.6883 Å, μ = 0.052 mm<sup>-1</sup>, *T* = 150(2) K, *R*<sub>1</sub> = 0.0701 for 3385 reflections with *I* > 2σ(*I*), and *wR*<sub>2</sub> = 0.1792 for all 4073 unique reflections. CCDC reference number 186128; methods and programs were standard.<sup>17</sup> See <http://www.rsc.org/suppdata/dt/b2/b205106/> for crystallographic data in CIF or other electronic format. Crystal data for the [NEt<sub>4</sub>]<sup>+</sup> salt of the [1-Ph-*closo*-1-CB<sub>7</sub>H<sub>7</sub>]<sup>-</sup> anion **5** were previously deposited; CCDC reference number 172016.

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