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X-Ray Crystal and Molecular Structure of the nido-Carborane B₇C₂H₅Me₂

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Summary The nido-carborane $B_7C_2H_7Me_2$ occurs as a bicapped Archimedean antiprism with the 6 position missing, methylated carbon atoms in the 1 and 2 cage positions, and bridging hydrogen atoms between the 7-10 and 9-10 positions.

RIETZ AND SCHAEFFER¹ recently reported the characterization of a new *nido*-carborane $B_7C_2H_{11}$ and its dimethyl derivative $B_7C_2H_9Me_2$. Based on the ease of formation and low symmetry implied by the ¹¹B and ¹H n.m.r. spectra theytentativelysuggested an open B_7C_2 icosahedral fragment analogous to the structure known² for the isoelectronic species B_9H_{13} ·MeCH. We felt that a definitive structure determination was needed and our X-ray diffraction study of the dimethyl derivative has established a somewhat different arrangement.

The compound was prepared as described previously¹ and purified on a high-vacuum low-temperature fractionation column.³ Samples were then distilled into thin-walled glass capillaries of 0.5 mm diameter and sealed under vacuum. A Picker automatic diffractometer equipped with a graphite monochromator and a nitrogen vapour cooling system⁴ was used to collect the diffraction data. A single crystal was grown at ca. -9 °C and, once grown, cooled to -170 °C. A systematic search of a limited hemisphere located reflections which could be indexed in a monoclinic cell of dimensions $a = 5\cdot570(4)$, $b = 13\cdot136(9)$, $c = 12\cdot998(10)$ Å, and $\beta = 104\cdot43(10)^{\circ}$. Systematic absences uniquely indicated the space group $P2_1/c$. Four molecules per unit cell gives a calculated density of $0\cdot849$ g cm⁻³, a reasonable value for compounds of this type.

Redundant data collected by the moving crystal-moving counter technique were averaged to a unique set of 627 reflections, of which 522 were considered observed [$I_{\text{net}} > \sigma(I)$]. All non-hydrogen atoms were located by direct methods and hydrogen atoms were located in a difference Fourier map after isotropic refinement of the non-hydrogen atoms. Further least-squares refinement in which non-hydrogen atoms were given anisotropic thermal parameters and hydrogen atoms were given isotropic thermal parameters gave a final residual of 0.058. Electron density and difference maps based on the final parameters revealed no anomalies, with the largest peak corresponding to 0.25 e Å⁻³.

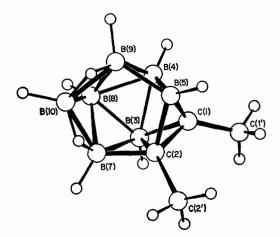


FIGURE. The structure of $B_7C_2H_9Me_2$. Bond distances, in Å, are 1-2 = 1.55; 1-1' = 1.47; 2-2' = 1.53; 1-3, 4, 5 = 1.60; 2-3 = 1.67; 2-5 = 1.75; 2-7 = 1.65; 8-10 = 1.69; 4-9 = 1.75. All other B-B single bonds are 1.78-1.81. Average B-H_t = 1.15, B-H_{br} = 1.25; and C-H = 1.00. Errors are ± 0.01 for non-hydrogen and 1.02 for hydrogen distances. hydrogen and ± 0.03 for hydrogen distances.

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The molecular geometry and the numbering system used⁵ are shown in the Figure. The structure is essentially a bicapped Archimedean antiprism with the 6 position removed. The distance across the open face B(7)-B(9) =2.63 Å, is not significantly different from the similar B(2)-B(4) distance, 2.59 Å, in the fully closed 1,6-dicarbacloso-decaborane(10) cage.⁶ The 2-3-4-5 positions describe a plane which is parallel within experimental error to the plane defined by the 7-8-9 positions, and bridging hydrogen atoms occur between the 7-10 and 9-10 positions. No appreciable lengthening of B-B bonds adjacent to cage carbon atoms is observed.⁷ B₇C₂H₉Me₂ possesses no molecular symmetry, and therefore occurs as a racemic mixture in the centrosymmetric crystal structure.

Although 70.6 MHz ¹¹B n.m.r. spectra suggest that the parent compound B₇C₂H₁₁ may be structurally different⁸ from the dimethyl derivative reported here, single crystals prepared to date have been unsuitable for X-ray work.

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