organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Gennadii V. Grintselev-Knyazev,^a* Mikhail Yu. Antipin,^a Sergey P. Knyazev,^b Valerii N. Kirin^b and Eugenii A. Chernyshev^b

^aX-Ray Structural Centre, Institute of Organoelement Compounds (INEOS), Russian Academy of Sciences, 28 Vavilov St., B-334, Moscow 119991, Russian Federation, and ^bMoscow State Academy of Fine Chemical Technology, Moscow 117571, Russian Federation

Correspondence e-mail: gena@xrlab.ineos.ac.ru

Key indicators

Single-crystal X-ray study T = 110 KMean $\sigma(C-C) = 0.002 \text{ Å}$ R factor = 0.057 wR factor = 0.128 Data-to-parameter ratio = 16.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

2-Methyl-1-propenyl-1,2-dicarba-closo-dodecaborane(12)

The title compound, $C_6H_{18}B_{10}$, is the first X-ray investigated carborane with a propenyl substituent. The C–C bond length in the carborane cage is 1.688 (2) Å.

Comment

The propenyl substituent is situated in the plane bisecting the C1-B3-B4 triangular face, with a B4-C1-C13-C14 torsion angle of 36.8 (2)°.



The C1-C2 distance of 1.688 (2) Å in (I) is significantly longer than those in B-substituent carboranes (Grintselev-Knyazev *et al.*, 2001); this agrees with the general tendency of C-C bond elongation in aryl C-substituent carboranes (Lewis & Welch, 1993).

The crystal packing is shown in Fig. 2.

Experimental

A solution of 1-Me-1,2-C₂B₁₀H₁₂ was treated with *n*-butyllithium in ether to afford the intermediate lithiated species. The mixture obtained was added to allyl bromide to give 1-allyl-2-methyl-1,2-C₂B₁₀H₁₂ (Heying *et al.*, 1963). 1-Propenyl-2-methyl-1,2-C₂B₁₀H₁₂ was obtained by isomerization of 1-allyl-2-methyl-1,2-C₂B₁₀H₁₂ with an excess of alkali. The NMR spectra were obtained in *d*₆-acetone (5% solution). ¹H (360 MHz, δ , p.p.m.): 6.31 (*dq*, =CHCH₃, 1H), 5.88 (*dd*, B-CH=, 1H), 2.00 (*s*, B-CH₃, 3H), 1.81 (*dd*, CH₃-CH=, 3H). ¹³C (90.6 MHz, δ , p.p.m.): 139.55 and 123.50 (HC=), 78.75 and 76.02 (C_{carb}), 22.68 and 17.24 (CH₃). ¹¹B (115.54 MHz, δ , p.p.m.): -3.80, -4.91, -8.52, -9.16, -10.01 (relative intensity 1:1:2:4:2).

Crystal data

$D_x = 1.075 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
Cell parameters from 821
reflections
$\theta = 3-30^{\circ}$
$\mu = 0.05 \text{ mm}^{-1}$
T = 110 (2) K
Prism, colorless
$0.50 \times 0.45 \times 0.40 \text{ mm}$

 \odot 2002 International Union of Crystallography Printed in Great Britain – all rights reserved

016 Gennadii V. Grintselev-Knyazev et al. \cdot C₆H₁₈B₁₀

Received 2 November 2001 Accepted 22 November 2001

Online 8 December 2001



Figure 1

The molecular structure of (I) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

Data collection

Bruker SMART 1000 CCD area-	3557 independent reflections
detector diffractometer	2306 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.049$
Absorption correction: multi-scan	$\theta_{\rm max} = 30.1^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -10 \rightarrow 10$
$T_{\min} = 0.976, T_{\max} = 0.981$	$k = -30 \rightarrow 30$
14366 measured reflections	$l = -10 \rightarrow 10$

Refinement

Refinement on F^2	
$R[F^2 > 2\sigma(F^2)] = 0.057$	
$wR(F^2) = 0.128$	
S = 1.03	
3557 reflections	
217 parameters	
All H-atom parameters refined	

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.028P)^2 \\ &+ 0.700P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} < 0.001 \\ \Delta\rho_{\text{max}} &= 0.34 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.22 \text{ e } \text{\AA}^{-3} \end{split}$$





Table 1

Selected geometric parameters (Å, °).

C1-C13	1.4843 (18)	C13-C14	1.312 (2)
C1-C2 C2-C16	1.6880 (19) 1.5062 (19)	C14-C15	1.489 (2)
C13 - C1 - C2 C16 - C2 - C1	117.63 (11) 117.51 (11)	C14-C13-C1 C13-C14-C15	125.42 (13) 124 34 (14)
010 02 01	11,101 (11)	010 011 010	12 110 1 (11)

Data collection: *SMART* (Bruker, 1999); cell refinement: *SMART* and *SAINT* (Bruker, 1999); data reduction: *SMART* and *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

References

Bruker (1998). SHELXTL. Version 5.10. Bruker AXS Inc., Madison, Wisconsin, USA.

Bruker (1999). *SMART* (Version 5.053) and *SAINT* (Version 6.02). Bruker AXS Inc., Madison, Wisconsin, USA.

Grintselev-Knyazev, G. V., Antipin, M. Yu., Knyazev, S. P., Kirin, V. N. & Chernyshev, E. A. (2001). Acta Cryst. C57, 827–829.

Heying, T. L., Ager, J. W., Clark, S. L., Alexander, R. P., Papetti, S., Reid, J. A. & Trotz, S. I. (1963). *Inorg. Chem.* **2**, 1097–1105.

Lewis, Z. G. & Welch, A. J. (1993). Acta Cryst. C49, 705-710.

Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.