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#### **Key indicators**

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.045 wR factor = 0.114 Data-to-parameter ratio = 17.7

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

# *rac-*(Diisopropylamino)(1-indenyl)(isopropyl-amino)borane

In the racemic title compound,  $C_{18}H_{29}BN_2$ , the B and the N atoms are virtually trigonal-planar coordinated and the presence of B==N partial double bonding is confirmed by the B-N bond lengths of 1.4154 (18) and 1.4178 (18) Å.

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### Comment

Ligand precursors related to the title compound *rac*-(diisopropylamino)(1-indenyl)(isopropylamino)borane, (I), have been previously employed for the synthesis of boronbridged constrained-geometry complexes. These complexes exhibit activity for the polymerization of ethene and higher  $\alpha$ -olefins when activated with MAO (methylaluminoxane) (Braunschweig *et al.*, 2000, 2004). In comparative studies, a strong influence of the ligand substitution patterns in corresponding complexes has been demonstrated on both catalyst activity and polymerization characteristics (Stevens 1994), justifying the synthesis of a series of boron-bridged analogues.



The structure of (I) (Fig. 1) in the solid state largely resembles that of its corresponding *N*-phenyl-substituted counterpart ( $\eta^1$ -C<sub>9</sub>H<sub>7</sub>)B(N<sup>i</sup>Pr<sub>2</sub>)N(H)Ph (Braunschweig *et al.*, 2003). Compound (I) adopts C<sub>1</sub> symmetry in the crystal structure. The crystal structure confirms the constitution assumed by NMR spectroscopy, with boron being attached to the *sp*<sup>3</sup>-hybridized C atom of the indenyl system. The B1 and N2 centres adopt trigonal–planar geometries, with the central atoms lying only 0.015 (B1) and 0.025 Å (N2) out of the planes of their respective substituents. These two trigonal planes are virtually coplanar with torsion angles of 176.35 (12) (C16– N2–B1–N1) and 0.2 (2)° (C13–N2–B1–N1). The B1–N1 and B1–N2 bond lengths of 1.4154 (18) and 1.4178 (18) Å, respectively, are equivalent within experimental error, indicating equal  $\pi$ -contributions from both amino groups. The N2-

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved bound isopropyl group *syn* to the indenyl moiety (which is coplanar to within 0.007 Å) is oriented such that the methine proton is directed towards the middle of the five-membered ring, a conformation also seen in the corresponding compound  $(C_9H_7)B(N^iPr_2)N(H)Ph$ ) with  $H-\pi$  distances to the centres of the C2=C3 olefinic and C1=C9 aromatic bonds of 2.446 and 2.509 Å, respectively. No short intermolecular contacts were observed.

## Experimental

The title compound was prepared by a modified literature procedure (Braunschweig *et al.*, 2003) by sequential reaction of  ${}^{i}Pr_2NBCl_2$  (2.16 g, 11.9 mmol) with lithium indenide (1.45 g, 11.9 mmol) and lithium isopropylamide (0.77 g, 11.9 mmol) in a hexane–toluene mixture. Recrystallization from hexane at 243 K afforded (I) exclusively as the 1-indenyl isomer in 35% yield in the form of pale-yellow crystals suitable for single-crystal X-ray diffraction.

Crystal data

$C_{18}H_{29}BN_2$	$D_x = 1.078 \text{ Mg m}^{-3}$
$M_r = 284.24$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 4341
a = 8.1029 (8) Å	reflections
b = 14.0352 (14)  Å	$\theta = 2.6-26.1^{\circ}$
c = 15.6679 (15)  Å	$\mu = 0.06 \text{ mm}^{-1}$
$\beta = 100.503 \ (2)^{\circ}$	T = 173 (2) K
$V = 1752.0 (3) \text{ Å}^3$	Tablet, yellow
Z = 4	$0.25 \times 0.20 \times 0.09 \text{ mm}$
Data collection	
Bruker SMART CCD area-detector	3471 independent reflections
diffractometer	2751 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.032$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.1^{\circ}$
(SADABS; Sheldrick, 2002)	$h = -10 \rightarrow 10$
$T_{\rm min} = 0.972, \ T_{\rm max} = 0.99$	$k = -17 \rightarrow 17$
16287 measured reflections	$l = -19 \rightarrow 19$
Refinement	
Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0505P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	+ 0.4773P]
$wR(F^2) = 0.114$	where $P = (F_0^2 + 2F_c^2)/3$

$$\begin{split} R[F^2 > 2\sigma(F^2)] &= 0.045 & + 0.4773P] \\ wR(F^2) &= 0.114 & where P &= (F_o^2 + 2) \\ S &= 1.02 & (\Delta/\sigma)_{max} = 0.001 \\ 3471 \text{ reflections} & \Delta\rho_{max} = 0.23 \text{ e} \text{ Å}^{-3} \\ 196 \text{ parameters} & \Delta\rho_{min} &= -0.19 \text{ e} \text{ Å}^{-3} \\ \text{H-atom parameters constrained} \end{split}$$

H atoms were visible in difference Fourier maps and were subsequently placed at idealized positions and treated as riding atoms, with C-H = 0.98 (CH<sub>3</sub>), 0.95 (aromatic CH) and 1.00 Å (aliphatic CH), and N-H = 0.88 Å.  $U_{iso}$  values for primary H atoms were fixed at 1.5 times and tertiary H atoms at 1.2 times  $U_{eq}(C)$ .





A view of (I), with displacement ellipsoids drawn at the 30% probability level.

Data collection: *SMART-NT* (Bruker, 2001); cell refinement: *SAINT-Plus-NT* (Bruker, 2001); data reduction: *SAINT-Plus-NT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *XP* in *SHELXTL*.

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