organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 294 K Mean σ (C–C) = 0.004 Å R factor = 0.038 wR factor = 0.096 Data-to-parameter ratio = 13.4

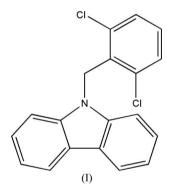
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

9-(2,6-Dichlorobenzyl)-9H-carbazole

In the title compound, $C_{19}H_{13}Cl_2N$, the dihedral angle between the dichlorobenzyl ring and the carbazole ring system is 92.6 (6)°. In the crystal structure, molecules are linked *via* intermolecular Cl····Cl interactions.

Comment

Carbazole derivatives substituted by *N*-alkylation possess valuable pharmaceutical properties (Buu-Hoï & Royer, 1950; Harfenist & Joyner, 1983; Caulfield *et al.*, 2002; Harper *et al.*, 2002). This paper reports the structure of the title compound, (I) (Fig. 1), which was synthesized by the *N*-alkylation of carbazole with 1,3-dichloro-2-(chloromethyl)benzene.



In the molecule of (I), the carbazole ring system is essentially planar, with an r.m.s. deviation from the mean plane of 0.009 Å. The angle between the dichlorobenzyl ring (C1–C6) and the carbazole plane is 92.6 (6)°.

In the crystal structure of (I), dimers are formed through weak intermolecular $\text{Cl} \cdot \cdot \cdot \text{Cl}^{\text{i}}$ interactions [symmetry code: (i) -x, 1 - y, -z], with $\text{Cl} \cdot \cdot \cdot \text{Cl} = 3.449$ (5) Å and $\text{C5} - \text{Cl}2 \cdot \cdot \cdot \text{Cl}2$ angles = 161.9 (5)°. These values are typical for inversionrelated $\text{Cl} \cdot \cdot \cdot \text{Cl}$ interactions (Desiraju, 1989).

Experimental

Compound (I) was synthesized according to the procedure of Huang *et al.* (2005). Crystals suitable for X-ray analysis were grown by slow evaporation of a solution in a mixture of chloroform (6 ml) and ethanol (2 ml) at room temperature over a period of 11 d.

Crystal data $C_{19}H_{13}Cl_2N$ $M_{\pi} = 326.20$

 $M_r = 320.20$ Monoclinic, C2/c a = 27.291 (6) Å b = 5.5688 (13) Å c = 21.132 (5) Å $\beta = 109.879$ (4)° V = 3020.3 (12) Å³ Z = 8 $D_x = 1.435 \text{ Mg m}^{-3}$ Mo K α radiation $\mu = 0.42 \text{ mm}^{-1}$ T = 294 (2) K Plate, colourless $0.20 \times 0.16 \times 0.08 \text{ mm}$ Received 28 November 2006 Accepted 29 November 2006

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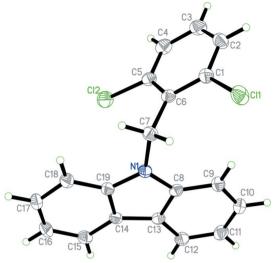


Figure 1

The molecular structure of (I), shown with 30% probability displacement ellipsoids (arbitrary spheres for H atoms).

Data collection

Bruker SMART 1000 CCD area-	7298 measured reflections
detector diffractometer	2666 independent reflectio
φ and ω scans	1839 reflections with $I > 2a$
Absorption correction: multi-scan	$R_{\rm int} = 0.040$
(SADABS; Bruker, 1997)	$\theta_{\rm max} = 25.0^{\circ}$
$T_{\min} = 0.920, \ T_{\max} = 0.967$	

Refinement

Refinement on F^2 H-atom parameters constrained $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.096$ S = 1.012666 reflections 199 parameters

endent reflections tions with $I > 2\sigma(I)$)

 $w = 1/[\sigma^2(F_o^2) + (0.047P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.19$ e Å⁻³ $\Delta \rho_{\min} = -0.22 \text{ e} \text{ Å}^{-3}$

All H atoms were included in the riding-model approximation, with C-H = 0.93 (aromatic) and 0.97 Å (CH₂), and with $U_{iso}(H) =$ $1.2U_{eq}(C).$

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:

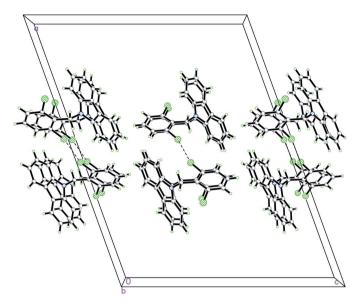


Figure 2

Part of the crystal structure of (I), with Cl...Cl interactions drawn as dashed lines.

SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

The authors thank the Postdoctoral Fund of Henan Institute of Science and Technology for financial support.

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