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# Xue-Min Duan,<sup>a</sup>\* Peng-Mian Huang,<sup>b</sup> Peng-Wu Zheng<sup>a</sup> and Jiang-Sheng Li<sup>b</sup>

 <sup>a</sup>School of Pharmacy, Jiangxi Science & Technology Normal University, Nanchang 330013, People's Republic of China, and
 <sup>b</sup>College of Pharmaceuticals & Biotechnology, Tianjin University, Tianjin 300072, People's Republic of China

Correspondence e-mail: dxmlhp@yahoo.com.cn

### Key indicators

Single-crystal X-ray study T = 294 KMean  $\sigma$ (C–C) = 0.012 Å R factor = 0.045 wR factor = 0.095 Data-to-parameter ratio = 14.4

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

# 9-Benzyl-3,6-dibromo-9H-carbazole

Two independent molecules comprise the asymmetric unit of the title compound,  $C_{19}H_{13}Br_2N$ , which was synthesized by *N*alkylation of benzyl bromide with 3,6-dibromo-9*H*-carbazole. The carbazole ring system is essentially planar and forms a dihedral angle with the pendant phenyl ring of 66.0 (2)° [70.6 (2)° for the second molecule]. In the crystal structure,  $\pi$ - $\pi$  interactions and weak C-H···Br interactions are observed.

## 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). In this paper, the structure of 9-benzyl-3,6-dibromo-9*H*-carbazole, (I), is reported; the compound was synthesized by *N*-alkylation of benzyl bromide with 3,6-dibromo-9*H*-carbazole.



The asymmetric unit of (I) comprises two independent but similar molecules (Fig. 1). The carbazole ring in each is essentially planar, with mean deviations of 0.030 and 0.016 Å, respectively, for the two molecules. The dihedral angles formed between the carbazole ring and the plane through the pendant phenyl ring is 66.0 (2)° [70.6 (2)° for the second molecule]. The C–Br distances fall in the range 1.903 (7)– 1.925 (8) Å, which is consistent with the literature (Allen *et al.*, 1987). In the crystal structure, there are  $\pi$ - $\pi$  interactions; the closest, 3.66 Å, is formed between the N1/C1/C2/C7/C8 and C7–C12 rings of translationally related molecules (x, -1 + y, z). In addition, there are C–H···Br interactions, as shown in Fig. 2 and as detailed in Table 1.

# **Experimental**

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved The title compound was prepared according to the procedure of Duan *et al.* (2005). A solution of potassium hydroxide (7.0 g) in

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#### Figure 1

A view of the two independent molecules of (I); displacement ellipsoids are drawn at the 30% probability level.

dimethylformamide (50 ml) was stirred at room temperature for 20 min. 3,6-Dibromocarbazole (6.5 g, 20 mmol), prepared according to Smith et al. (1992), was added and the mixture stirred for a further 40 min. A solution of benzyl bromide (5.1 g, 30 mmol) in dimethylformamide (50 ml) was added dropwise with stirring. The resulting mixture was then stirred at room temperature for 10 h and poured into water (500 ml), yielding a white precipitate. The solid product was filtered off, washed with cold water and recrystallized from EtOH, giving crystals of (I) (yield: 7.51 g, 90.5%; m.p. 430–432 K). <sup>1</sup>H NMR (DMSO- $d_6$ ):  $\delta$  5.6 (s, 2H), 7.2–7.6 (m, 9H), 8.5 (m, 2H); <sup>13</sup>C NMR (DMSO-d<sub>6</sub>): § 46.8, 112.4, 112.9, 124.1, 124.5, 127.4, 128.3, 129.6, 129.9, 138.2, 140.2. Compound (I) (40 mg) was dissolved in a mixture of chloroform (5 ml) and ethanol (5 ml) and the solution was kept at room temperature for 16 d. Natural evaporation of the solution gave colourless crystals suitable for X-ray analysis.

## Crystal data

$C_{19}H_{13}Br_2N$	$D_x = 1.707 \text{ Mg m}^{-3}$
$M_r = 415.12$	Mo $K\alpha$ radiation
Monoclinic, Cc	Cell parameters from 1791
a = 39.113 (6) Å	reflections
b = 4.5063 (7)  Å	$\theta = 2.3 - 21.4^{\circ}$
c = 20.469 (3) Å	$\mu = 5.01 \text{ mm}^{-1}$
$\beta = 116.449 \ (2)^{\circ}$	T = 294 (2) K
$V = 3230.1 (9) \text{ Å}^3$	Rod, colourless
Z = 8	$0.24$ $\times$ 0.18 $\times$ 0.14 mm



#### Figure 2

A portion of the crystal packing in (I), viewed down the *b*-axis direction. Dashed lines indicate C-H. Br interactions.

#### Data collection

Bruker SMART CCD area-detector	5698 independent reflections
diffractometer	3115 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.043$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.4^{\circ}$
(SADABS; Bruker, 1997)	$h = -48 \rightarrow 44$
$T_{\min} = 0.310, \ T_{\max} = 0.496$	$k = -3 \rightarrow 5$
8574 measured reflections	$l = -24 \rightarrow 25$

# Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0127P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.095$	$(\Delta/\sigma)_{\rm max} = 0.001$
S = 0.94	$\Delta \rho_{\rm max} = 0.34 \text{ e} \text{ Å}^{-3}$
5698 reflections	$\Delta \rho_{\rm min} = -0.40 \text{ e } \text{\AA}^{-3}$
397 parameters	Absolute structure: Flack (1983),
H-atom parameters constrained	2371 Friedel pairs.
	Flack parameter: 0.010 (11)

#### Table 1 Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C5-H5\cdots Br3^{i}$ $C12-H12\cdots Br1^{ii}$ $C25-H25\cdots Br4^{iii}$	0.93	3.00	3.800 (4)	146
	0.93	3.01	3.724 (4)	135
	0.93	2.97	3.852 (4)	158

Symmetry codes: (i) x, y - 1, z; (ii) x, -y + 1,  $z - \frac{1}{2}$ ; (iii) x, -y + 2,  $z - \frac{1}{2}$ .

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

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

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