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Key indicators

Single-crystal X-ray study T = 294 KMean σ (C–C) = 0.005 Å R factor = 0.034 wR factor = 0.078 Data-to-parameter ratio = 16.0

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3,6-Dibromo-9-(3-pyridylmethyl)-9H-carbazole

The title compound, C₁₈H₁₂Br₂N₂, was synthesized by Nalkylation of 3-(chloromethyl)pyridine with 3,6-dibromo-9Hcarbazole. The carbazole ring system is essentially planar, with a mean deviation of 0.0138 Å, and forms a dihedral angle of 96.7 $(8)^{\circ}$ with the plane of the pyridine ring. In the crystal structure, π - π 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). The title compound, 3,6-dibromo-9-(3-pyridylmethyl)-9H-carbazole, (I) (Fig. 1), was synthesized by N-alkylation of 3-(chloromethyl)pyridine with 3,6-dibromo-9H-carbazole.



The carbazole ring system is essentially planar, with a mean deviation of 0.0138 Å, consistent with previously reported values (Duan, Huang et al., 2005). The dihedral angle formed between the carbazole ring system and the plane of the pyridine ring is 96.7 (8)°. C-Br distances are in the range 1.905 (3) to 1.907 (3) Å and are consistent with literature values (Allen *et al.*, 1987). In the crystal structure, π - π interactions are observed; the shortest, 3.519 Å, is between the N1/ C1–C12 and C7–C12 rings of molecules related by (1 - x, -y, -y)2 - z). In addition, there are C-H···Br interactions, as shown in Fig. 2 and detailed in Table 1.

Experimental

The title compound was prepared according to the procedure of Duan, Han et al. (2005). A solution of potassium hydroxide (7.0 g) in 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 3-(chloromethyl)pyridine (3.825 g, 30 mmol) in

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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 (I) was collected by filtration, washed with cold water and recrystallized from EtOH (yield 7.11 g, 85.5%; m.p. 488 K). 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 18 d. Slow evaporation of the solution yielded colourless crystals suitable for X-ray analysis.

 $D_r = 1.769 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation Cell parameters from 2796

reflections $\theta=2.5{-}26.4^\circ$ $\mu = 5.19 \text{ mm}^{-1}$ T = 294 (2) K Block, colourless $0.26 \times 0.22 \times 0.14 \text{ mm}$

 $R_{\rm int} = 0.046$ $\theta_{\rm max} = 26.5^\circ$ $h = -12 \rightarrow 13$ $k = -16 \rightarrow 20$ $l = -10 \rightarrow 12$

3194 independent reflections

2238 reflections with $I > 2\sigma(I)$

Crvstal data

$C_{18}H_{12}Br_2N_2$
$M_r = 416.12$
Monoclinic, $P2_1/c$
a = 10.469 (3) Å
b = 16.405 (5) Å
c = 9.866 (3) Å
$\beta = 112.761 \ (4)^{\circ}$
$V = 1562.6 (8) \text{ Å}^3$
7 - 4

Data collection

Bruker SMART CCD area-detector
diffractometer
φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 1997)
$T_{\min} = 0.258, \ T_{\max} = 0.484$
8670 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0276P)^2$
$R[F^2 > 2\sigma(F^2)] = 0.034$	+ 0.8633P]
$wR(F^2) = 0.078$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} = 0.001$
3194 reflections	$\Delta \rho_{\rm max} = 0.48 \text{ e } \text{\AA}^{-3}$
200 parameters	$\Delta \rho_{\rm min} = -0.52 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
	Extinction coefficient: 0.0233 (8)

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C11-H11\cdots Br1^i$	0.93	2.89	3.595 (3)	134
Symmetry code: (i) r -	L1 v 7⊥1			

Symmetry code: (i) x + 1, y, z + 1.

All H atoms were placed in calculated positions and refined using a riding model, with C-H = 0.93 (aromatic) and 0.97 (methylene) Å, 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: SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

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Figure 1

The molecular structure of (I), with displacement ellipsoids drawn at the 30% probability level ..



Figure 2

Packing diagram of (I), viewed along [001]. Dashed lines indicate C-H...Br interactions.

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