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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.006 Å R factor = 0.044 wR factor = 0.119 Data-to-parameter ratio = 13.2

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12-(4-Bromophenyl)-9,9-dimethyl-1,2,3,4,9,10hexahydrobenz[a]acridin-11-one

The title compound, $C_{25}H_{22}BrNO$, has been synthesized by the reaction of 4-bromobenzaldehyde, 3,3-dimethylcyclopentane-1,3-dione with 2-naphthylamine in ethanol. In the crystal structure, the molecules are connected by $N-H\cdots O$ hydrogen bonds, forming chains along the *a* axis. Received 13 April 2006 Accepted 21 April 2006

Comment

Many natural and synthetic compounds containing the acridine skeleton display interesting biological and physical activities, such as antimalaria (Nasim & Brychey, 1979; Thull & Testa, 1994; Reil *et al.*, 1994; Mandi *et al.*, 1994) and antitumour properties (Khurana *et al.*, 1990). Multihydroacridinone derivatives have been reported to have high fluorescence efficiencies and can be used as fluorescent molecular probes for the monitoring of polymerization processes (Popielarz *et al.*, 1997). Increasingly, they also receive attention due to the similarity of their properties with those of 1,4dihydropyridines, which have similarities in structure with biologically important compounds, such as nicotinamide adenine dinucleotide (Srividya *et al.*, 1996). In this paper, we report the crystal structure of the title compound, (I).



In compound (I), atoms C7 and N1 deviate from the C1/C6/ C8/C17 plane by 0.321 (5) and 0.151 (5) Å, respectively (Fig. 1), indicating a boat conformation. Atom C3 deviates from the C1/C2/C4/C5/C6 plane by 0.658 (5) Å, indicating an envelope conformation. The dihedral angle between the C1/ C6/C8/C17 plane and the C20–C25 benzene ring is 89.98 (11)°.

In the crystal structure, the molecules are connected *via* $N-H\cdots O$ hydrogen bonds (Table 1), forming chains along the *a* axis (Fig. 2).

Experimental

Compound (I) was prepared by the reaction of 4-bromobenzaldehyde (1 mmol) with 3,3-dimethylcyclopentane-1,3-dione

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The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

(1 mmol) and 2-naphthylamine (1 mmol) in ethanol (3 ml) at 351 K. Single crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of a 95% aqueous ethanol solution (yield 95%; m.p. >573 K). Spectroscopic analysis: ¹H NMR (DMSO- d_6 , δ , p.p.m.): 0.84 $(3H, s, CH_3)$, 1.04 $(3H, s, CH_3)$, 2.03 $(1H, d, J = 16.4 Hz, CH_2)$, 2.23 $(1H, d, J = 16.4 \text{ Hz}, \text{CH}_2), 2.39 (1H, d, J = 16.8 \text{ Hz}, \text{CH}_2), 2.56 (1H, d, J)$ = 16.8 Hz, CH₂), 5.78 (1H, s, CH), 7.18 (2H, d, J = 8.4 Hz, ArH), 7.34-7.31 (4H, m, ArH), 7.42 (1H, t, J = 7.6 Hz, ArH), 7.82–7.79 (2H, m, ArH), 7.90 (1H, d, J = 8.4 Hz, ArH), 9.76 (1H, s, NH).

Crystal data

C ₂₅ H ₂₂ BrNO	V = 1036.9 (9) Å ³
$M_r = 432.35$	Z = 2
Triclinic, P1	$D_x = 1.385 \text{ Mg m}^{-3}$
a = 7.296 (4) Å	Mo $K\alpha$ radiation
b = 9.597 (5) Å	$\mu = 2.00 \text{ mm}^{-1}$
c = 15.304 (8) Å	T = 298 (2) K
$\alpha = 94.038 \ (7)^{\circ}$	Block, colourless
$\beta = 93.465 \ (8)^{\circ}$	$0.32 \times 0.27 \times 0.09$ m
$\gamma = 103.331 \ (7)^{\circ}$	

Data collection

Bruker SMART CCD area-detector	55
diffractometer	33
φ and ω scans	20
Absorption correction: multi-scan	R_{i}
(SADABS; Sheldrick, 1996)	θ_{n}
$T_{\min} = 0.567, T_{\max} = 0.841$	

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ wR(F²) = 0.119 S = 1.013346 reflections 253 parameters H-atom parameters constrained mm

522 measured reflections 346 independent reflections 053 reflections with $I > 2\sigma(I)$ int = 0.023 $max = 25.0^{\circ}$

 $w = 1/[\sigma^2(F_0^2) + (0.0454P)^2]$ + 0.7039P] where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.60 \ {\rm e} \ {\rm \AA}^2$ $\Delta \rho_{\rm min} = -0.48 \ {\rm e} \ {\rm \AA}^{-3}$



Figure 2 A packing diagram for (I), projected along the *a* axis.

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

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$		
$N1 - H1 \cdots O1^i$	0.86	1.99	2.840 (4)	172		
Symmetry code: (i) $x - 1, y, z$.						

All H atoms were positioned geometrically and treated as riding, with C-H distances in the range 0.93–0.98 Å, and with $U_{iso}(H) =$ $1.5U_{eq}(C)$ for methyl H atoms and $1.2U_{eq}(C)$ for others.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); 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, 1999); software used to prepare material for publication: SHELXTL.

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