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Hong Jiang,^a Xiang-Shan Wang,^b* Mei-Mei Zhang,^b Yu-Ling Li^b and Da-Qing Shi^b

^aThe Key Laboratory of Biotechnology for Medical Plants of Jiangsu Province, Xuzhou Normal University, Xuzhou 221116, People's Republic of China, and ^bDepartment of Chemistry, Xuzhou Normal University, Xuzhou 221116, People's Republic of China

Correspondence e-mail: xswang1974@yahoo.com

Key indicators

Single-crystal X-ray study T = 173 K Mean σ (C–C) = 0.003 Å R factor = 0.029 wR factor = 0.076 Data-to-parameter ratio = 12.5

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. 2-Amino-4-(2-chlorophenyl)-7,7-dimethyl-1-(4-methylphenyl)-5-oxo-1,4,5,6,7,8hexahydroquinoline-3-carbonitrile

The title compound, $C_{25}H_{24}ClN_3O$, was synthesized by the reaction of 2-chlorophenylmethylidenemalononitrile and 3-(4-tolylamino)-5,5-dimethylcyclohex-2-enone in the presence of triethylbenzylammonium chloride in an aqueous medium. All bond lengths and angles are normal. In the crystal structure, intermolecular N-H···O hydrogen bonds link the molecules into chains extending in the [101] direction.

Comment

Quinolines and their derivatives exhibit diverse pharmacological activities, such as cholinesterase (Isomae *et al.*, 2002), antiplasmoidal (Beagley *et al.*, 2003), cytotoxic (Ma *et al.*, 2004), functional (Denton *et al.*, 2005) and antibacterial (Fokialakis *et al.*, 2002) activities. To avoid the inherent disadvantages of many organic solvents, such as toxicity and non-stability, we are working under a new environmentally friendly procedure for the synthesis of the above-mentioned compounds. Specifically, we have focused our attention on the use of water as a reaction medium following Breslow & Rideout (1980), who rediscovered the use of water as a solvent in organic chemistry. We report here the crystal structure of the title compound, (I).



In (I), all bond lengths and angles (Table 1) are normal. The N1/C1–C5 ring is not planar, adopting a boat conformation (Fig. 1). Atoms C3 and N1 deviate from the basal plane defined by atoms C1/C2/C4/C5 by 0.200 (2) and 0.073 (1) Å, respectively. Similar distortions were observed in 7,7-dimeth-yl-2-(4-bromophenyl)-4-phenyl-5-oxo-1,4,5,6,7,8-hexahydro-quinoline (Shi *et al.*, 2002) and 3,3,6,6-tetramethyl-9-(4-chlorophenyl)-10-(4-methylphenyl)-1,2,3,4,5,6,7,8,9,10-deca-hydroacridine-1,8-dione (Wang *et al.*, 2003). The C4/C5/C7–C10 ring adopts a half-chair conformation; atom C9 deviates from the mean plane of the remaining atoms by 0.630 (2) Å. A

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Figure 1

A view of (I), showing the atom-numbering scheme and with displacement ellipsoids drawn at the 50% probability level.



Figure 2

The molecular packing, viewed down the b axis, showing the intermolecular hydrogen bonds as dashed lines.

similar conformation has been found in the structure of 7,7dimethyl-2-amino-3-cyano-4-(3,4-methylenedioxylphenyl)-5oxo- 5,6,7,8-tetrahydro-4*H*-benzo[*b*]pyran (Wang *et al.*, 2002). The benzene rings make a dihedral angle of 9.0 $(1)^{\circ}$.

In the crystal structure, intermolecular N-H···O hydrogen bonds (Table 2) link the molecules into chains extending in the [101] direction (Fig. 2).

Experimental

The title compound, (I), was prepared by the reaction of 2-chlorophenylmethylidenemalononitrile (2 mmol) and 3-(4-tolylamino)-5,5dimethylcyclohex-2-enone (2 mmol) in the presence of triethylbenzylammonium chloride (0.1 g) in water at 363 K for 8 h (yield 98%; m.p. 525-527 K). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of a dimethylformamide solution. Elemental analysis calculated: C 71.85, H 5.79, N 10.05%; found: C 71.73, H 6.01, N 9.98%. ¹H NMR (DMSO-*d*₆, δ, p.p.m.): 0.73 (*s*, 3H, CH_3), 0.89 (s, 3H, CH_3), 1.73 (d, J = 16.0 Hz, 1H, CH), 1.90 (d, J =16.0 Hz, 1H, CH), 2.19 (*d*, *J* = 16.8 Hz, 2H, CH₂), 4.47 (*s*, 1H, CH), 5.33 (s, 2H, NH₂), 7.04-7.48 (m, 8H, ArH).

Crystal data

C25H24CIN3O $M_r = 417.92$ Monoclinic, Cc a = 19.541 (2) Å b = 9.0199 (8) Å c = 14.8424 (16) Å $\beta = 124.628 \ (2)^{\circ}$ V = 2152.7 (4) Å³ Z = 4

Data collection

Rigaku Mercury diffractometer ω scans Absorption correction: multi-scan (Jacobson, 1998) $T_{\min} = 0.892, \ T_{\max} = 0.943$ 10014 measured reflections 3434 independent reflections

Refinement

| Refinement on F^2 |
|---------------------------------|
| $R[F^2 > 2\sigma(F^2)] = 0.029$ |
| $wR(F^2) = 0.076$ |
| S = 1.05 |
| 3434 reflections |
| 275 parameters |
| H-atom parameters constrained |
| - |

 $D_{\rm r} = 1.290 {\rm Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 4284 reflections $\theta = 3.3 - 25.3^{\circ}$ $\mu = 0.20 \text{ mm}^{-1}$ T = 173 (2) K Block, yellow $0.59 \times 0.50 \times 0.30 \text{ mm}$

3389 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.016$ $\theta_{\rm max} = 25.3^{\circ}$ $h = -23 \rightarrow 21$ $k = -10 \rightarrow 10$ $l = -15 \rightarrow 17$

| $w = 1/[\sigma^2(F_o^2) + (0.0474P)^2]$ |
|--|
| + 0.7163P] |
| where $P = (F_0^2 + 2F_c^2)/3$ |
| $(\Delta/\sigma)_{\rm max} < 0.001$ |
| $\Delta \rho_{\rm max} = 0.20 \ {\rm e} \ {\rm \AA}^{-3}$ |
| $\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$ |
| Absolute structure: Flack (1983), |
| with 1478 Friedel pairs |
| Flack parameter: $-0.04(5)$ |
| |

Table 1

Selected geometric parameters (Å, °).

| N1-C1 | 1.392 (2) | C2-C3 | 1.524 (2) |
|-------------|-------------|-------------|-------------|
| N1-C5 | 1.396 (2) | C3-C4 | 1.514 (2) |
| C1-C2 | 1.360 (2) | C4-C5 | 1.355 (2) |
| | | | |
| C1-N1-C5 | 119.95 (14) | C4-C3-C2 | 108.62 (14) |
| C2-C1-N1 | 120.82 (15) | C5-C4-C3 | 123.57 (15) |
| C1-C2-C3 | 123.42 (15) | C4-C5-N1 | 121.15 (15) |
| | | | |
| C5-N1-C1-C2 | 6.5 (2) | C2-C3-C4-C5 | 15.2 (2) |
| N1-C1-C2-C3 | 6.6 (3) | C3-C4-C5-N1 | -4.5(3) |
| C1-C2-C3-C4 | -16.2(2) | | () |
| | | | |

Table 2 Hydrogen-bond geometry (Å, °).

| $D - H \cdot \cdot \cdot A$ | D-H | $H \cdot \cdot \cdot A$ | $D{\cdots}A$ | $D - H \cdots A$ |
|-----------------------------|-----------|-------------------------|--------------|------------------|
| $N2-H2B\cdotsO1^{i}$ | 0.88 | 2.36 | 2.937 (2) | 123 |
| 8 | . 1 . 3 . | 1 | | |

Symmetry code: (i) $x + \frac{1}{2}, -y + \frac{3}{2}, z + \frac{1}{2}$.

The H atoms were positioned geometrically and refined as riding, with C-H = 0.95–1.00 Å and N-H = 0.88 Å, and with $U_{iso}(H) = 1.2-1.5U_{eq}$ of the parent atom.

Data collection: *CrystalClear* (Rigaku, 1999); cell refinement: *CrystalClear*; data reduction: *CrystalStructure* (Rigaku/MSC, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXL97*.

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