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Key indicators

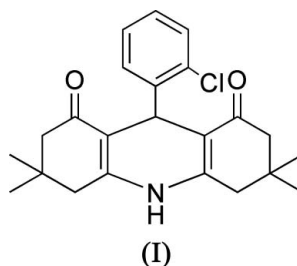
Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.010\text{ \AA}$
 R factor = 0.078
 wR factor = 0.269
Data-to-parameter ratio = 14.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.9-(2-Chlorophenyl)-1,2,3,4,5,6,7,8,9,10-decahydro-
3,3,6,6-tetramethylacridine-1,8-dione

The title compound, $\text{C}_{23}\text{H}_{26}\text{ClNO}_2$, was synthesized by the reaction of 5,5-dimethylcyclohexane-1,3-dione with 4-chloro-*N*-(2-chlorobenzylidene)benzenamine under solvent-free conditions using triethylbenzylammonium chloride as catalyst at 353 K. The dihydropyridine ring of the molecule adopts a half-chair conformation.

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Comment

1,4-Dihydropyridines (1,4-DHPS) are well known compounds for their pharmacological profile in calcium channel modulations (Bossert *et al.*, 1981). In particular, dihydropyridine drugs such as nifedipine, nicardipine, amlodipine and others are effective cardiovascular agents for the treatment of hypertension. They have many significant biological activities: anti-atherosclerotic, antitumor, hepatoprotective, antidiabetic and anti-ischemic agents (Mauzeral & Westheimer, 1955; Di Stilo *et al.*, 1998; Shan *et al.*, 2004; Suarez *et al.*, 2003).



The solvent-free reaction has attracted great attention in recent years (Tanaka & Toda, 2000) and has been proved to have many advantages: high efficiency and selectivity, easy separation and purification, mild reaction conditions, reduced pollution, and low cost. Solvent-free reactions have performed well recently (Kaupp *et al.*, 2003; Goud *et al.*, 1995; Annunziata *et al.*, 1997).

We report here the crystal structure of the title compound, (I), which has been synthesized by the solvent-free reaction of 5,5-dimethylcyclohexane-1,3-dione and 4-chloro-*N*-(2-chlorobenzylidene)benzenamine at 353 K.

In (I), the dihydropyridine ring is in a half-chair conformation, with atom C7 deviating from the C1/C6/C8/C13/N1 plane by 0.121 (8) Å (Fig. 1). Both cyclohexene rings adopt half-chair conformations: atom C3 deviates from the C1/C2/C4–C6 plane by 0.639 (9) Å and atom C11 deviates from the C8–C10/C12/C13 plane by 0.644 (9) Å.

The dihedral angle between the C1/C6/C8/C13/N1 and C1/C2/C4–C6 planes is 3.3 (4)°. The dihedral angle between the C1/C6/C8/C13/N1 and C8–C10/C12/C13 planes is 2.4 (4)°.

Experimental

Compound (I) was prepared by the reaction of 5,5-dimethylcyclohexane-1,3-dione (4 mmol) with 4-chloro-*N*-(2-chlorobenzylidene)benzenamine (2 mmol) under solvent-free condition using TEBA (triethylbenzylammonium chloride) as catalyst at 353 K.

Crystal data

$C_{23}H_{26}ClNO_2$	$Z = 8$
$M_r = 383.90$	$D_x = 1.301 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 17.004 (6) \text{ \AA}$	$\mu = 0.21 \text{ mm}^{-1}$
$b = 11.429 (4) \text{ \AA}$	$T = 298 (2) \text{ K}$
$c = 21.741 (10) \text{ \AA}$	Block, colourless
$\beta = 111.858 (15)^\circ$	$0.24 \times 0.15 \times 0.13 \text{ mm}$
$V = 3921 (3) \text{ \AA}^3$	

Data collection

Bruker SMART CCD area-detector diffractometer	9874 measured reflections
φ and ω scans	3463 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	1617 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.951$, $T_{\max} = 0.973$	$R_{\text{int}} = 0.123$
	$\theta_{\text{max}} = 25.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0343P)^2 + 19.16P]$
$R[F^2 > 2\sigma(F^2)] = 0.078$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.269$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.07$	$\Delta\rho_{\text{max}} = 0.64 \text{ e \AA}^{-3}$
3463 reflections	$\Delta\rho_{\text{min}} = -0.49 \text{ e \AA}^{-3}$
245 parameters	
H-atom parameters constrained	

All H atoms were positioned geometrically and treated as riding, with C–H distances of 0.93–0.98 Å and N–H = 0.86 Å, and with $U_{\text{iso}}(\text{H})$ set at $1.2U_{\text{eq}}(\text{C}, \text{N})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1999); 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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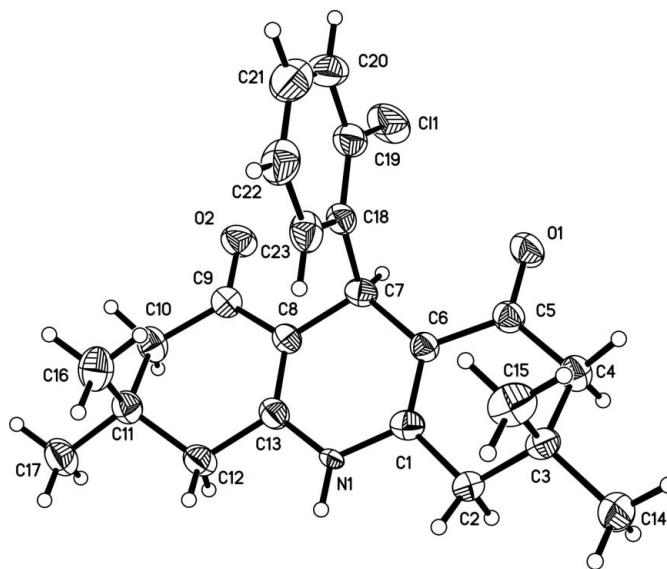


Figure 1

The molecular structure of (I), showing 40% probability displacement ellipsoids and the atom-numbering scheme.

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