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Key indicators

Single-crystal X-ray study T = 298 K Mean σ (C–C) = 0.009 Å R factor = 0.073 wR factor = 0.219 Data-to-parameter ratio = 14.0

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9-(2,6-Dihydroxyphenyl)-3,3,6,6-tetramethyl-N-(4-methylphenyl)-1,8-dioxo-1,2,3,4,5,6,7,8,9,10-decahydroacridine

The title compound, C₃₀H₃₃NO₄, was synthesized by the reaction of 2,6-dihydroxybenzaldehyde with p-toluidine and 5,5-dimethylcyclohexane-1,3-dione in glycol under microwave irradition. X-ray analysis reveals that the dihydropyridine ring is in a distorted boat conformation and that the cyclohexenone rings adopt enevelope conformations. In the crystal structure, molecules are linked by O-H···O hydrogen bonds around a centre of symmetry to form a dimer.

Comment

Acridine and its derivatives inhibit HIV-1 reverse transcriptase by intercalating the template-primer hybrid (Cellai et al., 1994). Such compounds are well known as antimicrobials (Al-Ashmawi et al., 1994) and antitumour agents (Wang et al., 1993), and are used in the treatment of urinary incontinence (Ohnmacht, 1993). Recently, we reported the crystal strucof 10-cyclopropyl-9-(4-hydroxy-3-methoxyphenyl)tures 3,3,6,6-tetramethyl-1,2,3,4,5,6,7,8,9,10-decahydroacridine-1,8dione (Tu et al., 2004) and 9-(4-methoxyphenyl)-3,4,6,7,9,10hexahydroacridine-1.8(2H,5H)-dione (Guo et al., 2004). We now report the crystal structure of the title compound, (I).



The dihydropyridine ring in (I) adopts a distorted boat conformation. In this ring, atoms N1 and C7 deviate from the C1/C6/C8/C13 plane by 0.093 (8) and 0.200 (8) Å, respectively (Fig. 1). Both cyclohexenone rings adopt envelope conformations. The C18-C23 and C25-C30 benzene rings are nearly coplanar [dihedral angle = $8.4 (2)^{\circ}$] and form dihedral angles of 84.6 (2) and 87.5 (2)°, respectively, with the C1/C6/C8/C13 plane. An intramolecular O3-H3···N1 interaction is observed (Table 1). The crystal packing, shown in Fig. 2, reveals that the molecules exist as centrosymmetric $O-H \cdots O$ hydrogen-bonded dimers.

Compound (I) was prepared by the reaction of 2,6-dihydroxy-

benzaldehyde (1 mmol) with p-toluidine (1 mmol) and 5,5-

Experimental

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Figure 1

The molecular structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.



Figure 2 A view of a dimer of (I), showing $O-H\cdots O$ and $O-H\cdots N$ hydrogen bonds (dashed lines).

dimethylcyclohexane-1,3-dione (2 mmol) in glycol (1 ml) under microwave irradiation for 4 min (yield 91%; m.p. 531-532 K). Single crystals suitable for X-ray diffraction were obtained by slow evaporation of an ethanol solution.

Crystal data

$C_{30}H_{33}NO_4$	$D_x = 1.213 \text{ Mg m}^{-3}$
$M_r = 471.57$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 1094
a = 11.995 (4) Å	reflections
b = 11.002 (4) Å	$\theta = 2.4 - 19.3^{\circ}$
c = 19.965 (7) Å	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 101.419 \ (6)^{\circ}$	T = 298 (2) K
$V = 2582.5 (15) \text{ Å}^3$	Block, light yellow
Z = 4	$0.38 \times 0.17 \times 0.15$ mm

Data collection

Bruker SMART CCD area-detector 4544 independent reflections diffractometer 1460 reflections with $I > 2\sigma(I)$ φ and ω scans $R_{\rm int} = 0.118$ Absorption correction: multi-scan $\theta_{\rm max} = 25.0^{\circ}$ $h = -14 \rightarrow 13$ (SADABS: Sheldrick, 1996) $k = -13 \rightarrow 9$ $T_{\min} = 0.970, \ T_{\max} = 0.988$ 13000 measured reflections $l = -23 \rightarrow 23$ Refinement Refinement on F^2

 $w = 1/[\sigma^2(F_0^2) + (0.072P)^2]$ $R[F^2 > 2\sigma(F^2)] = 0.073$ wR(F²) = 0.219 S = 1.014544 reflections 324 parameters H-atom parameters constrained Extinction coefficient: 0.0042 (10)

where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.31 \text{ e Å}$ $\Delta \rho_{\rm min} = -0.27 \text{ e } \text{\AA}^{-3}$ Extinction correction: SHELXL97

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

$D - H \cdot \cdot \cdot A$	$D-{\rm H}$	$H \cdot \cdot \cdot A$	$D{\cdots}A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O3−H3···N1	0.82	2.44	3.233 (6)	164
$O4-H4\cdots O4^i$	0.82	2.30	2.587 (6)	101

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

H atoms were placed in idealized positions and allowed to ride on their parent atoms, with O-H distances of 0.82 Å and C-H distances in the range 0.93–0.98 Å, and with $U_{iso}(H) = 1.5U_{ea}(carrier)$ for methyl and hydroxy H atoms, and $1.2U_{eq}(carrier)$ for other H atoms. Owing to the poor diffraction quality of the crystal, the higher angle reflections were very weak and only 32% of the reflections were found to be observed with $I > 2\sigma(I)$. This resulted in a high R_{int} value. The C6-O3 and C30-O4 distances were restrained to be 1.36 (1) Å. Owing to large displacement parameters, the U^{ij} components of atoms C27, C28 and C28 were restrained to isotropic behaviour.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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