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4-Methyl-7,7a,13a,14-tetrahydrobenzo[*e*]pyrano[2',3':5,6]naphtho[2,3-*b*][1,4]dioxin-2-one

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Abstract

In the title molecule, $C_{20}H_{16}O_4$, the coumarin moiety is planar and both the tetrahydrobenzene and the dioxin rings adopt a half-chair conformation. The mean planes through the tetrahydrobenzene and dioxin rings form a dihedral angle of 72.8 (1)°. The majority of DNA monointercalating antitumour drugs have a common general structure, comprising a tri- or tetracyclic chromophore to which are attached one or two flexible side chains bearing cationic charges (Palmer *et al.*, 1988). Recently it has been reported that a series of substituted dibenzo[1,4]dioxins show remarkable activity against wild-type P388 leukaemia *in vitro* and *in vivo* (Lee *et al.*, 1992). Because of their antitumour activity and ecotoxicity, different substituted dibenzo[1,4]dioxins have been synthesized and the crystal structure determination of one of them, (I), is reported here.



Bond lengths and bond angles in the coumarin moiety and C—O distances in the dioxin ring are comparable with reported values (Kumar *et al.*, 1997; Chinnakali *et al.*, 1998; Rissanen *et al.*, 1987). The tetrahydrobenzene ring adopts a half-chair conformation with asymmetry parameter $\Delta C_2(C8$ —C7) = 0.045 (2) (Nardelli, 1983). The dioxin ring also adopts a half-chair conformation with C13 and C14 deviating from the O17—C18— C23—O24 plane by -0.436 (4) and 0.318 (4) Å, re-



Fig. 1. The structure of the title compound showing 50% probability displacement ellipsoids and the atom-numbering scheme.

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spectively $[\Delta C_2(C13-C14) = 0.020(1)]$. The coumarin moiety is planar with a maximum deviation at C3 of -0.051(4) Å. The tetrahydrobenzene ring forms dihedral angles of 6.98(8) and 72.8(1)° with the coumarin and dioxin rings, respectively. The dihedral angle between the phenyl and dioxin rings is 2.3(1)°. The carbonyl-O atom is involved in a weak C-H···O intermolecular hydrogen bond with the methyl-C atom. The crystal structure is stabilized by this weak hydrogen bond and van der Waals interactions.

Experimental

The title compound was synthesized by ring opening of 4methyl-7,10-dihydro-7,8-benzocoumarin-8,9-oxide with pyrocatechol followed by cyclization using Mitsunobu reagent (triphenylphosphine-diethylazodicarboxylate) (Sriraghavan & Ramakrishnan, 1998). Single crystals were obtained from ethanol by slow evaporation.

Crystal data

C₂₀H₁₆O₄ Mo $K\alpha$ radiation $M_r = 320.33$ $\lambda = 0.71073 \text{ Å}$ Orthorhombic Cell parameters from 3676 reflections Fdd2 a = 57.344(2) Å $\theta = 1.42 - 27.50^{\circ}$ $\mu = 0.099 \text{ mm}^{-1}$ b = 9.7802(4) Å c = 10.6263(5) Å T = 293 (2) K $V = 5959.6 (4) \text{ Å}^3$ Thin plate Z = 16 $0.56 \times 0.36 \times 0.04$ mm $D_x = 1.428 \text{ Mg m}^{-3}$ Colourless D_m not measured

Data collection

Siemens SMART CCD areadetector diffractometer ω scans Absorption correction: none 9438 measured reflections 3416 independent reflections

Refinement

Refinement on F^2 $\Delta \rho$ $R[F^2 > 2\sigma(F^2)] = 0.065$ $\Delta \rho$ $wR(F^2) = 0.155$ ExtS = 1.042Sca3416 reflectionsM218 parametersCH atoms: see belowAbs $w = 1/[\sigma^2(F_o^2) + (0.07P)^2]$ Fwhere $P = (F_o^2 + 2F_c^2)/3$ Flac $(\Delta/\sigma)_{max} < 0.001$ 0

 $\Delta \rho_{\text{max}} = 0.206 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\text{min}} = -0.238 \text{ e } \text{\AA}^{-3}$ Extinction correction: none Scattering factors from *International Tables for Crystallography* (Vol. C) Absolute structure: Flack (1983) Flack parameter = 0.92 (180)

2158 reflections with

 $I > 2\sigma(I)$ $R_{\rm int} = 0.066$

 $\theta_{\rm max} = 27.50^{\circ}$

 $h = 0 \rightarrow 73$

 $k = 0 \rightarrow 12$

 $l = -13 \rightarrow 13$

Table 1. Selected geometric parameters (Å)

| C3—C4 | 1.348 (5) | C13—O24 | 1.448 (4) |
|--------|-----------|---------|-----------|
| C5—C6 | 1.355 (5) | C13C14 | 1.491 (5) |
| C7C12 | 1.502 (5) | C14-017 | 1.459 (4) |
| C8—C15 | 1.512 (4) | C14-C15 | 1.514 (5) |
| C12C13 | 1.509 (5) | O17C18 | 1.371 (5) |
| | | | |

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

| D—H···A | D—H | H···A | $D \cdot \cdot \cdot A$ | $D = H \cdots A$ |
|-------------------------------------|---------------------------|------------------|-------------------------|------------------|
| C16H16 <i>B</i> ···O11 ¹ | 0.960 (9) | 2.468 (8) | 3.405 (5) | 165.1 (7) |
| Symmetry code: (i) | $x, \frac{1}{2} + y, z =$ | - \ . | | |

The data collection covered a hemisphere of reciprocal space by a combination of three sets of exposures; each set had a different φ angle (0, 88 and 180°) for the crystal and each exposure of 30 s covered 0.3° in ω . The crystal-to-detector distance was 4 cm and the detector swing angle was -35° . Coverage of the unique set is over 99% complete. Crystal decay was monitored by repeating thirty initial frames at the end of the data collection and analysing the duplicate reflections, and was found to be negligible.

The structure was solved by direct methods and refined by full-matrix least-squares techniques. Although all H atoms were located from a difference Fourier map, as the ratio of reflections to parameters was low they were geometrically fixed and allowed to ride on the atoms to which they were attached.

Data collection: *SMART* (Siemens, 1996*a*). Cell refinement: *SAINT* (Siemens, 1996*b*). Data reduction: *SAINT*. Program(s) used to solve structure: *SHELXTL* (Sheldrick, 1996). Program(s) used to refine structure: *SHELXTL*. Molecular graphics: *SHELXTL*. Software used to prepare material for publication: *SHELXTL* and *PARST* (Nardelli, 1995).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: HA1226). Services for accessing these data are described at the back of the journal.

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