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Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$
 R factor = 0.037
 wR factor = 0.098
Data-to-parameter ratio = 10.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.7a,8,9,10-Tetrahydrobenzo[*f*]cyclopenta[*b*]-
chromen-10-oneIn the title compound, $\text{C}_{16}\text{H}_{12}\text{O}_2$, the pyran ring adopts a half-chair conformation. Intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, in the form of $R_2^2(8)$ rings, link the molecules into dimers. In addition, $\text{C}-\text{H}\cdots\pi$ interactions are found.

Comment

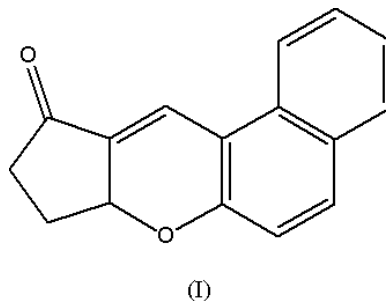
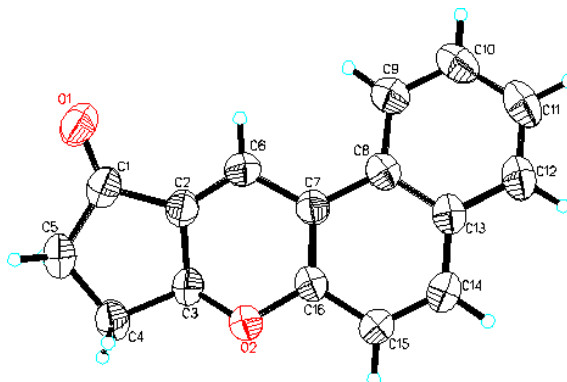
Chromenes (2*H*-1-benzopyran derivatives) are frequently found in naturally occurring heterocycles, many of which exhibit biological activity (Bowers *et al.*, 1976) and have been widely employed as important intermediates in the synthesis of many natural products and medicinal agents (Cassidy *et al.*, 1992; Wang & Finn, 2000). Thus, various synthetic methods for the formation of these compounds have been reported (Chang & Grubbs, 1998; Kaye & Nocanda, 2000; Lee *et al.*, 2003; Parker & Mindt, 2001). In this paper, a new chromene compound, *viz.* 7a,8,9,10-tetrahydrobenzo[*f*]cyclopenta[*b*]-chromen-10-one, (I), is reported.There are four fused rings in (I). The pyran ring adopts a half-chair conformation. The $\text{O}2/\text{C}3/\text{C}2$ and $\text{O}2/\text{C}16/\text{C}7/\text{C}6$ planes are inclined at an angle of $43.35(1)^\circ$. The main geometric discrepancy is in the angle $\text{C}1-\text{C}2-\text{C}6$ of

Figure 1

A view of the molecular structure of (I), with the atom-numbering scheme.

Received 9 August 2004

Accepted 31 August 2004

Online 4 September 2004

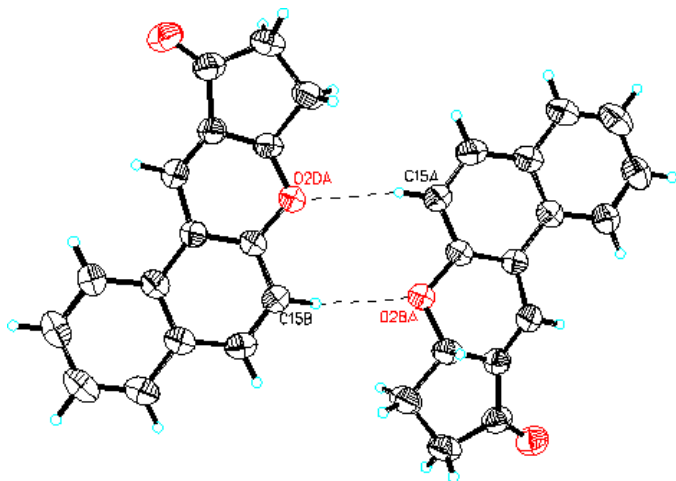


Figure 2
The dimer formed by C—H...O hydrogen bonds, which are indicated by dashed lines.

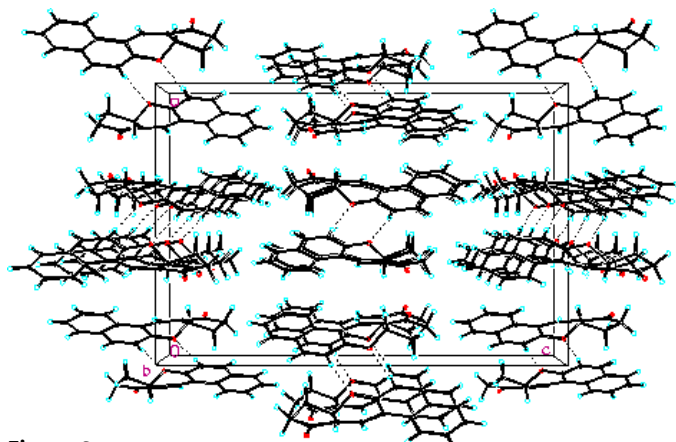


Figure 3
The crystal packing in (I), viewed down the *b* axis. Dashed lines indicate hydrogen bonds.

130.48 (17)°, probably a consequence of the steric restraints imposed by the four fused rings. The molecule contains a chiral atom, C3, but the crystal structure belongs to a centrosymmetric space group and is, therefore, a racemic mixture. The pairs of enantiomers related by the inversion centre are linked *via* C—H...O hydrogen bonds (Fig. 2). Specifically, the C15—H15...O2ⁱⁱ [symmetry code: (ii) 1 - *x*, -*y*, 1 - *z*] hydrogen bond has the pyran O atoms as acceptors and aromatic CH atoms acting as donors. The weak interactions generate rings with graph-set motif $R_2^2(8)$ (Bernstein *et al.*, 1995). In addition, there are two intermolecular C—H... π contacts (Table 2), identified by *PLATON* (Spek, 2003), which involve the π system of the C8—C13 ring (with centroid *Cg*). Hydrogen-bonding information is given in Table 2 and a crystal packing diagram is shown in Fig. 3.

Experimental

Compound (I) was synthesized *via* a Baylis–Hillman reaction. At room temperature, a clear solution of 2-hydroxy-1-naphthaldehyde (1 mmol), cyclopent-2-enone (2 mmol) and imidazole (1 mmol) in

tetrahydrofuran (1.5 ml) was mixed with deionized water (1.5 ml). The mixture was stirred at ambient temperature for 48 h to bring the reaction to completion. The mixture was diluted with water (10 ml) and extracted with ethyl acetate. The organic layer was concentrated under reduced pressure. After the usual work-up, chromatography of the crude product on silica gel, using ethyl acetate and petroleum (1:4) as eluant, gave the pure product (I) in 61% yield. The compound (60 mg) was dissolved in CHCl₃ (2 ml). The solution was allowed to evaporate slowly at room temperature over several days to yield yellow crystals suitable for X-ray crystallography.

Crystal data

C₁₆H₁₂O₂
M_r = 236.26
 Orthorhombic, *Pbca*
a = 15.5021 (19) Å
b = 6.6902 (8) Å
c = 22.676 (3) Å
V = 2351.7 (5) Å³
Z = 8
D_x = 1.335 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 2391 reflections
 θ = 2.2–22.2°
 μ = 0.09 mm⁻¹
T = 298 (2) K
 Block, yellow
 0.30 × 0.30 × 0.10 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
T_{min} = 0.974, *T_{max}* = 0.991
 9238 measured reflections

1684 independent reflections
 1377 reflections with *I* > 2σ(*I*)
R_{int} = 0.029
 θ_{\max} = 23.3°
h = -17 → 15
k = -6 → 7
l = -25 → 25

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.037
wR (*F*²) = 0.098
S = 1.04
 1684 reflections
 163 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0514P)^2 + 0.3759P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} < 0.001
 $\Delta\rho_{\max} = 0.15 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.11 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

O1—C1	1.220 (2)	C2—C6	1.336 (2)
O2—C16	1.368 (2)	C2—C3	1.487 (2)
O2—C3	1.4388 (19)	C3—C4	1.521 (2)
C1—C2	1.466 (2)	C4—C5	1.528 (3)
C1—C5	1.507 (3)	C6—C7	1.449 (2)
C16—O2—C3	114.26 (13)	O2—C3—C4	110.99 (14)
O1—C1—C2	126.30 (17)	C2—C3—C4	105.04 (14)
O1—C1—C5	126.32 (16)	C3—C4—C5	104.57 (14)
C2—C1—C5	107.37 (16)	C1—C5—C4	105.93 (14)
C6—C2—C1	130.48 (17)	C2—C6—C7	118.57 (16)
C6—C2—C3	119.94 (15)	O2—C16—C7	122.18 (15)
C1—C2—C3	109.38 (15)	O2—C16—C15	115.66 (15)
O2—C3—C2	110.84 (14)		

Table 2

Hydrogen-bonding geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C11—H11...O1 ⁱ	0.93	2.59	3.508 (2)	172
C15—H15...O2 ⁱⁱ	0.93	2.48	3.388 (2)	165
C4—H4A...C _g ⁱⁱⁱ	0.97	3.03	3.923 (2)	154
C10—H10...C _g ^{iv}	0.93	2.95	3.676 (2)	136

Symmetry codes: (i) $x, \frac{3}{2} - y, \frac{1}{2} + z$; (ii) 1 - *x*, -*y*, 1 - *z*; (iii) 1 - *x*, 1 - *y*, 1 - *z*; (iv) 1 + *x*, $-\frac{1}{2} - y, z - \frac{1}{2}$. *Cg* is the centroid of the C8—C13 ring.

H atoms were placed in calculated positions and allowed to ride on their parent atoms, with $U_{\text{iso}}(\text{H})$ values set to $1.5U_{\text{eq}}(\text{parent atom})$ for the $\text{C}sp^3$ -bound H atoms and $1.2U_{\text{eq}}(\text{parent atom})$ for $\text{C}sp^2$ -bound H atoms. The C–H distances were fixed in the range 0.93–0.97 Å.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 2000); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000); software used to prepare material for publication: *SHELXTL*.

The authors gratefully acknowledge the financial support of this work by the Natural Science Foundation of China (No. 20171031 to PY) and Shanxi Provincial Natural Science Foundation

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