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Key indicators

Single-crystal X-ray study $T=293~\mathrm{K}$ Mean $\sigma(\mathrm{C-C})=0.002~\mathrm{\mathring{A}}$ R factor = 0.044 wR factor = 0.119 Data-to-parameter ratio = 15.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

3,3'-Diethyl-3,3'-dihydroxy-5,5'-dimethyl-2,2'-bi-1*H*-indanylidene-1,1'-dione

The title compound, $C_{24}H_{24}O_4$, was synthesized by the reaction of ethylmagnesium bromide with 5,5'-bimethyl-2,2'-biindanylidene-1,1',3,3'-tetraone under an argon atmosphere. The molecule lies across a crystallographic inversion centre. The cyclopentenone ring adopts a flattened envelope conformation.

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Comment

Photochromism is defined as light-induced reversible transformation of chemical species between two isomers having different absorption spectra. In recent years, organic photochromic compounds have received considerable attention in view of their general applicability as optical information storage materials or switching devices (Irie, 2000). Among many types of photochromic compounds, the biindenylidene derivatives are unusual in that they simultaneously undergo photochromism in the crystalline state as well as the generation of radicals (Li *et al.*, 2007; Nakatsuji, 2004; Tanak *et al.*, 2004). We report here the crystal structure of the title compound, (I).

The asymmetric unit of (I) contains one half-molecule with the other half generated by a centre of inversion (Fig. 1); the centre of inversion lies at the mid-point of the C2=C2ⁱ bond [symmetry code: (i) 1 - x, 2 - y, 2 - z]. The C2=C2ⁱ bond distance of 1.352 (3) Å confirms its double-bond character.

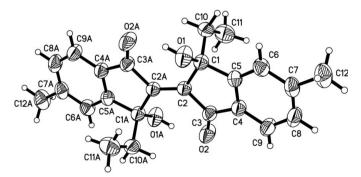


Figure 1 The molecular structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. Atoms labelled with the suffix A are generated by the symmetry operation (1 - x, 2 - y, 2 - z).

© 2007 International Union of Crystallography All rights reserved The cyclopentenone ring adopts a flattened envelope conformation, with atom C2 at the flap.

Intramolecular $O-H\cdots O$ hydrogen bonds generate S(7) ring motifs. The crystal structure is stabilized by $C-H\cdots O$ intermolecular interactions (Table 1).

Experimental

The title compound was synthesized by the reaction of ethylmagnesium bromide (2.16 g, 20 mmol) with 5,5'-bimethyl-2,2'-biindanylidene-1,1',3,3'-tetraone (1.59 g, 5 mmol) under an argon atmosphere. Crystals suitable for X-ray analysis were grown by slow evaporation of a dichloromethane solution at room temperature.

Crystal data

$C_{24}H_{24}O_4$	$V = 982.6 (4) \text{ Å}^3$
$M_r = 376.43$	Z = 2
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
a = 8.584 (2) Å	$\mu = 0.09 \text{ mm}^{-1}$
b = 7.4860 (15) Å	T = 293 (2) K
c = 15.554 (4) Å	$0.28 \times 0.24 \times 0.20 \text{ mm}$
$\beta = 100.564 (8)^{\circ}$	

Data collection	
Bruker SMART CCD area-detector diffractometer Absorption correction: multi-scan ($SADABS$; Sheldrick, 1996) $T_{\min} = 0.947$, $T_{\max} = 0.983$	5475 measured reflections 2030 independent reflections 1403 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.029$
Refinement	
$P[F^2 > 2\sigma(F^2)] = 0.044$	H stome treated by a mixture of

 $R[F^2 > 2\sigma(F^2)] = 0.044$ H atoms treated by a mixture of independent and constrained S = 1.03 refinement $\Delta \rho_{\text{max}} = 0.18 \text{ e Å}^{-3}$

2030 reflections $\Delta \rho_{\rm max} = 0.18 \ {\rm e \ \mathring{A}^{-3}}$ 132 parameters $\Delta \rho_{\rm min} = -0.14 \ {\rm e \ \mathring{A}^{-3}}$ 1 restraint

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

D $ H$ $\cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
$O1-H1 \cdots O2^{i}$	0.833 (9)	1.935 (14)	2.7019 (18)	153 (2)
$C10-H10A \cdots O2^{i}$	0.97	2.58	3.247 (3)	126
$C10-H10B \cdots O1^{ii}$	0.97	2.58	3.476 (2)	153

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

The hydroxyl H atom was located in a difference map and refined with an O—H distance restraint of 0.84 (1) Å. All other H atoms were positioned geometrically (C—H = 0.93–0.97 Å) and refined in the riding-model approximation, with $U_{\rm iso}({\rm H})$ = 1.2–1.5 $U_{\rm eq}({\rm C})$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

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