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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.045 wR factor = 0.158 Data-to-parameter ratio = 13.9

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

3,3-Dibenzylbenzo[b]furan-2(3H)-one

The title compound, $C_{22}H_{18}O_2$, was synthesized by the reaction of methyl (2-hydroxyphenyl)acetate and benzyl chloride. In the crystal structure, the dihydrobenzofuran-2-one ring system lies on a crystallographic mirror plane.

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Comment

2,3–Dihydrobenzofuran-2-one derivatives are of great interest because of their biological properties. Some derivatives of 2,3dihydrobenzofuran-2-one have been reported as potential inhibitors of angiogensis (Duflos *et al.*, 2003) and as drugs for the treatment of hypoxia (Lepagnol & Lavielle, 1988) and some also show high efficacy as intrinsic analgesics and antiinflammatories (Baumann *et al.*, 1986, Rao & Rau, 1985, Closse *et al.*, 1981). We report here the crystal structure of the title compound, (I).



The molecular structure of (I) is shown in Fig. 1; selected bond lengths and angles are given in Table 1. Atoms C8–C15, O2 and attached H atoms lie on a crystallographic mirror plane..

Experimental

Methyl (2-hydroxyphenyl)acetate (20 mmol) was dissolved in acetone (20 ml) and potassium carbonate (30 mmol) was added in one portion. Benzyl chloride (40 mmol) was then added to this mixture at 293 K. The resulting mixture was refluxed for 20 h. The mixture was filtered and the filtrate concentrated under reduced pressure to afford crude compound (I). Pure (I) was obtained by recrystallization from ethyl acetate (m.p. 454–455 K). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of an ethanol solution. Spectroscopic analysis, ¹H NMR (CDCl₃, p.p.m.): 7.20–7.21 (m, 1H), 7.12–7.15 (s, 8H), 6.93–6.95 (m, 4H), 6.70–6.72 (m, 1H), 3.38–3.41 (m, 2H), 3.26–3.28 (m, 2H).

Crystal data

C ₂₂ H ₁₈ O ₂	Mo $K\alpha$ radiation
$M_r = 314.36$	Cell parameters from 25
Orthorhombic, Pnma	reflections
a = 12.650 (3) Å	$\theta = 10-13^{\circ}$
b = 15.488 (3) Å	$\mu = 0.08 \text{ mm}^{-1}$
c = 8.772 (2) Å	T = 293 (2) K
V = 1718.6 (6) Å ³	Block, colourless
Z = 4	$0.4 \times 0.3 \times 0.3 \text{ mm}$
$D_x = 1.215 \text{ Mg m}^{-3}$	

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Data collection

Nonius CAD-4 diffractometer $\omega/2\theta$ scans Absorption correction: ψ scan (North *et al.*, 1968) $T_{min} = 0.973$, $T_{max} = 0.977$ 1740 measured reflections 1740 independent reflections 1002 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.046$ $wR(F^2) = 0.158$ S = 0.981740 reflections 125 parameters H-atom parameters constrained $\begin{array}{l} \theta_{\max} = 26.0^{\circ} \\ h = 0 \rightarrow 15 \\ k = 0 \rightarrow 19 \\ l = 0 \rightarrow 10 \\ \text{25 standard reflections} \\ \text{every 200 reflections} \\ \text{intensity decay: none} \end{array}$

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.1P)^2] \\ & \text{where } P = (F_o^2 + 2F_c^2)/3 \\ & (\Delta/\sigma)_{\text{max}} = 0.005 \\ & \Delta\rho_{\text{max}} = 0.20 \text{ e } \text{ Å}^{-3} \\ & \Delta\rho_{\text{min}} = -0.14 \text{ e } \text{ Å}^{-3} \\ & \text{Extinction correction: } SHELXL97 \\ & \text{Extinction coefficient: } 0.069 (7) \end{split}$$

Table 1

Selected geometric parameters (Å, °).

O1-C9	1.194 (3)	C7-C8	1.550 (2)
O2-C9	1.374 (3)	C8-C15	1.503 (3)
O2-C10	1.404 (3)	C8-C9	1.529 (4)
C4-C7	1.503 (3)	$C8-C7^{i}$	1.550 (2)
C9-O2-C10	107.8 (2)	$C7 - C8 - C7^{1}$	109.4 (2)
C5-C4-C7	121.0 (2)	O1-C9-C8	129.4 (2)
C4-C7-C8	115.1 (2)	02-C9-C8	110.5 (2)
C15-C8-C9	100.7 (2)	C15-C10-O2	111.9 (2)
C15-C8-C7	113.9 (1)	C10-C15-C8	109.1 (2)
C9-C8-C7	109.3 (1)		

Symmetry code: (i) $x, \frac{1}{2} - y, z$.

All H atoms were positioned geometrically, with C–H = 0.93–0.97 Å, and refined using a riding model, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm carrier atom})$.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS*97



Figure 1

A view of the molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code: (A) x, $\frac{1}{2} - y$, z.]

(Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *SHELXTL* (Siemens, 1996); software used to prepare material for publication: *SHELXL*97.

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