Received 24 October 2006 Accepted 6 November 2006

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.065 wR factor = 0.174 Data-to-parameter ratio = 12.5

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

5,17-Dibenzoyl-25,26,27,28-tetrahydroxycalix[4]arene

The structure of the title compound, $C_{42}H_{32}O_6$, is an important example of the functionalization of the upper rim of calix[4]arene molecules. It possesses crystallographic twofold rotation symmetry and displays the calixarene cone conformation. The hydroxyl groups form intramolecular hydrogen bonds.

Comment

Calixarenes currently attract considerable interest in the field of supramolecular chemistry because their derivatives can form inclusion complexes with cations or with neutral molecules (Gutsche, 1989; Vicens & Böhmer, 1991). The functionalization of the upper rim of this platform molecule is only a recent achievement compared with the lower rim (Asfari *et al.*, 2001; Harvey, 2002). Various examples allowing the functionalization of the upper rim exist, but only a few of them are very versatile with respect to the formation of C–C bonds. The shape of the calix[4]arene macrocycle and the steric hindrance between the substituents control the geometry and the final interatomic distances. In this investigation. we exploit the possibility of adapting the Fries rearrangement for the preparation of the title partially upper-rim substituted calix[4]arene, (2).



The molecular structure of compound (2) is shown in Fig. 1. The calix[4]arene assumes a conformation with crystallographic C_2 symmetry. The cone conformation of the calixarene core is retained from the precursor, (1).

A benzoyloxy substituent has been introduced on the calix[4]arene via the Fries rearrangement. The angles C4–C15–O3, C4–C15–C16 and O3–C15–C16 are 118.9 (4), 121.8 (4) and 119.3 (4)°, respectively. The C15–O3 distance is 1.238 (4) Å. Bond lengths and angles may be considered normal. The shape of the calix cavity is defined by the interplanar angles which the various aromatic rings make with the planes of the four methylene groups. The molecular conformation is stabilized by two symmetry-equivalent pairs of

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Figure 1

The molecular structure of (2), with 30% probability displacement ellipsoids. For clarity, H atoms have been omitted. Unlabelled atoms are related to labelled atoms by the symmetry operator $(-x + 1, y, -z + \frac{1}{2})$

intramolecular hydrogen bonds between the hydroxyl groups (Table 1).

Experimental

A mixture of compound (1) (0.5 g) and aluminium chloride in dry chloroform (50 ml) was stirred at 333 K for 24 h. The reaction mixture was poured into ice-water (50 ml). The dichloromethane layer was successively washed with 0.1 M hydrochloric acid and 10% sodium bicarbonate solution. On concentration, the dichloromethane layer provided compound (2). Colourless crystals of (2) were grown from a methanol-dichloromethane (1:15) solution of the material.

Crystal data

C42H32O6 $M_r = 632.68$ Monoclinic, C2/c a = 18.615 (4) Å b = 15.609 (4) Å c = 12.220 (3) Å $\beta = 116.187 (4)^{\circ}$ $V = 3186.1 (13) \text{ Å}^3$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Bruker, 1997) $T_{\min} = 0.983, T_{\max} = 0.991$

Z = 4 $D_x = 1.319 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.09 \text{ mm}^{-1}$ T = 293 (2) K Pillar, colourless $0.20 \times 0.15 \times 0.10 \text{ mm}$

6625 measured reflections 2819 independent reflections 1253 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.080$ $\theta_{\rm max} = 25.0^{\circ}$

Refinement

Refinement on F^2	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.065$	independent and constrained
$wR(F^2) = 0.174$	refinement
S = 0.90	$w = 1/[\sigma^2(F_0^2) + (0.0737P)^2]$
2819 reflections	where $P = (F_0^2 + 2F_c^2)/3$
225 parameters	$(\Delta/\sigma)_{\rm max} < 0.001$
	$\Delta \rho_{\rm max} = 0.18 \text{ e } \text{\AA}^{-3}$
	$\Delta \rho_{\rm min} = -0.18 \text{ e} \text{ Å}^{-3}$

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} \hline C17-H17A\cdotsO1^{i}\\ O2-H2\cdotsO1^{ii}\\ O1-H1\cdotsO2 \end{array}$	0.93 0.817 (19) 0.863 (19)	2.58 1.86 (2) 1.81 (2)	3.478 (5) 2.672 (4) 2.662 (4)	163 171 (4) 169 (6)

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

H atoms were located in a difference map, but were refined with fixed individual isotropic displacement parameters $[U_{iso}(H) =$ $1.2U_{eq}(C)$] using a riding model, with C-H = 0.93 and 0.97 Å for Caromatic and Cmethylene, respectively. The hydroxyl H atoms were refined isotropically (distances are in Table 1).

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

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