organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Da-Quan Peng

Department of Chemistry, Chongqing Normal University, Chongqing 400047, People's Republic of China

Correspondence e-mail: pengdq_cqnu@yahoo.com.cn

Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.054 wR factor = 0.151 Data-to-parameter ratio = 16.4

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

2-Methyl-4-phenyl-2-(trimethylsilyloxy)-3,4-dihydro-2*H*,5*H*-pyrano[3,2-c][1]benzopyran-5-one

In the title compound, $C_{22}H_{24}O_4Si$, the coumarin system is essentially planar and the pyran ring adopts a half-chair conformation. There are intermolecular $C-H\cdots O$ and $C-H\cdots \pi$ interactions in the crystal structure. Received 28 June 2006 Accepted 4 July 2006

Comment

The cycloaddition chemistry of *o*-quinone methides has attracted considerable interest (Nair *et al.*, 2001). In our recent research work on the cycloaddition reactions of *o*-quinone methides derived from 4-hydroxycoumarin, we have prepared the title compound, (I), which is obtained from the cycloaddition reaction between coumarin quinone methide and an excess amount of trimethyl[(1-methylethenyl)oxy]silane. As part of this study, we have undertaken the X-ray crystallographic analysis of (I) in order to elucidate the conformation of this cycloadduct product.



The bond lengths and angles in (I) (Fig. 1 and Table 1) are in good agreement with those found in the related compound 2methyl-2-(trimethylsilyloxy)-3,4-dihydro-2*H*,5*H*-pyrano[3,2*c*][1]benzopyran-5-one (Peng *et al.*, 2005). The coumarin system is essentially planar. The dihedral angle between the coumarin unit and the benzene ring (C17–C22) is 73.8 (3)°. The pyran ring (C5–C9/O2) adopts a half-chair conformation, with atoms C5 and C6 deviating from the C7–C9/O2 plane by 0.232 (4) and -0.466 (4) Å, respectively.

In the crystal structure, the molecular packing is stabilized by intermolecular $C-H\cdots O$ and $C-H\cdots \pi$ interactions (Table 2) involving the aromatic rings.

Experimental

Under argon, a mixture of 4-hydroxycoumarin, 3 equivalents of trimethyl[(1-methylethenyl)oxy]silane and an excess amount of benzaldehyde were suspended in dry dioxane and refluxed for about 20 h. Compound (I) was isolated by column chromatography of the reaction mixture on silica gel after evaporation of the solvent, in 27% yield. Single crystals of (I) were obtained by slow evaporation of a petroleum ether–ethyl acetate (2:1 ν/ν) solution.

© 2006 International Union of Crystallography All rights reserved



Figure 1

The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

Crystal data

 $\begin{array}{l} C_{22}H_{24}O_4\text{Si} \\ M_r = 380.50 \\ \text{Monoclinic, } P2_1/n \\ a = 10.038 \ (2) \ \text{Å} \\ b = 12.396 \ (3) \ \text{Å} \\ c = 16.853 \ (3) \ \text{Å} \\ \beta = 100.74 \ (3)^{\circ} \\ V = 2060.3 \ (8) \ \text{Å}^3 \end{array}$

Data collection

Enraf–Nonius CAD-4 diffractometer $\omega/2\theta$ scans Absorption correction: ψ scan (XCAD4; Harms & Wocadlo, 1995) $T_{\rm min} = 0.931, T_{\rm max} = 0.962$ 4268 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.054$ $wR(F^2) = 0.151$ S = 1.004029 reflections 245 parameters H-atom parameters constrained Z = 4 $D_x = 1.227 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.14 \text{ mm}^{-1}$ T = 293 (2) KBlock, colourless $0.40 \times 0.31 \times 0.28 \text{ mm}$

4029 independent reflections 2297 reflections with $I > 2\sigma(I)$ $R_{int} = 0.117$ $\theta_{max} = 26.0^{\circ}$ 3 standard reflections every 200 reflections intensity decay: none

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.05P)^{2} + P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.19 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{min} = -0.27 \text{ e } \text{\AA}^{-3}$ Extinction correction: *SHELXL97* Extinction coefficient: 0.0157 (13)

Table 1

Selected geometric parameters (Å, °).

Si-O1	1.651 (2)	O4-C16	1.206 (3)
O1-C5	1.396 (3)	C5-C6	1.506 (4)
C5-O1-Si	135.16 (19)	C15-O3-C16	121.4 (2)
C9-O2-C5	117.4 (2)	C5-C6-C7	112.9 (2)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$	
$C1-H1B\cdots O3^{i}$	0.96	2.58	3.482 (4)	156	
$C14-H14\cdots O4^{ii}$	0.93	2.55	3.463 (4)	168	
$C6-H6A\cdots Cg(B)^{iii}$	0.97	2.81	3.773 (3)	175	
$C21-H21\cdots Cg(A)^{iv}$	0.93	2.96	3.815 (4)	153	

Symmetry codes: (i) x - 1, y, z; (ii) -x + 2, -y + 1, -z; (iii) $-x + \frac{3}{2}, y - \frac{1}{2}, -z + \frac{1}{2}$; (iv) -x + 1, -y + 1, -z. Notes: Cg(A) and Cg(B) denote the centroids of the aromatic rings C10–C15 and C17–C22, respectively

The H atoms were included in the riding-model approximation, with C–H distances of 0.93, 0.96, 0.97 and 0.98 Å for aromatic, methyl, methylene and methine H atoms, respectively, and with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm aromatic, methylene and methine C})$ or $1.5U_{\rm eq}({\rm methyl C})$.

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: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

This work was supported by the National Natural Science Foundation of China (NSFC, No. 20572044). Partial support by the Modern Analytical Center at Nanjing University is also gratefully acknowledged.

References

Enraf-Nonius (1989). CAD-4 Software. Version 5. Enraf-Nonius, Delft, The Netherlands.

Harms, K. & Wocadlo, S. (1995). XCAD4. University of Marburg, Germany. Nair, V., Jayan, C. N., Radhakrishnan, K. V., Anikumar, G. & Rath, N. P. (2001). Tetrahedron, 57, 5807–5813.

Peng, D.-Q., Liu, Y., Lu, Z.-F. & Xu, J.-H. (2005). Acta Cryst. E61, o3704– 03705.

Sheldrick, G. M. (1997). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.

Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.