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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.054$
$w R$ 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.

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

In the title compound, $\mathrm{C}_{22} \mathrm{H}_{24} \mathrm{O}_{4} \mathrm{Si}$, the coumarin system is essentially planar and the pyran ring adopts a half-chair conformation. There are intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-$ $\mathrm{H} \cdots \pi$ interactions in the crystal structure.

## 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.

(I)

The bond lengths and angles in (I) (Fig. 1 and Table 1) are in good agreement with those found in the related compound 2 -methyl-2-(trimethylsilyloxy)-3,4-dihydro-2H,5H-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 $73.8(3)^{\circ}$. The pyran ring ( $\mathrm{C} 5-\mathrm{C} 9 / \mathrm{O} 2$ ) 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 $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \pi$ 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 ( $2: 1 \mathrm{v} / \mathrm{v}$ ) solution.


Figure 1
The structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the $30 \%$ probability level.

## Crystal data

$$
\begin{aligned}
& \mathrm{C}_{22} \mathrm{H}_{24} \mathrm{O}_{4} \mathrm{Si} \\
& M_{r}=380.50 \\
& \text { Monoclinic, } P 2_{b} / n \\
& a=10.038(2) \AA \\
& b=12.396(3) \AA \\
& c=16.853(3) \AA \\
& \beta=100.74(3)^{\circ} \\
& V=2060.3(8) \AA^{\circ}
\end{aligned}
$$

## Data collection

Enraf-Nonius CAD-4
$\quad$ diffractometer
$\omega / 2 \theta$ scans
Absorption correction: $\psi$ scan
$\quad(X C A D 4$; Harms \& Wocadlo,
$1995)$
$T_{\min }=0.931, T_{\max }=0.962$
4268 measur

4268 measured reflections

## Refinement

Refinement on $F^{2}$ $R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.054$
$w R\left(F^{2}\right)=0.151$
$S=1.00$
4029 reflections
245 parameters
H -atom parameters constrained
$Z=4$
$D_{x}=1.227 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\mu=0.14 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Block, colourless
$0.40 \times 0.31 \times 0.28 \mathrm{~mm}$

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

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.05 P)^{2}+P\right] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.19 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.27 \mathrm{e}^{-3}
\end{aligned}
$$

Extinction correction: SHELXL97
Extinction coefficient: 0.0157 (13)

Table 1
Selected geometric parameters ( $\AA{ }^{\circ},{ }^{\circ}$ ).

| $\mathrm{Si}-\mathrm{O} 1$ | $1.651(2)$ | $\mathrm{O} 4-\mathrm{C} 16$ | $1.206(3)$ |
| :--- | :--- | :--- | ---: |
| $\mathrm{O} 1-\mathrm{C} 5$ | $1.396(3)$ | $\mathrm{C} 5-\mathrm{C} 6$ | $1.506(4)$ |
|  |  |  |  |
| $\mathrm{C} 5-\mathrm{O} 1-\mathrm{Si}$ | $135.16(19)$ | $\mathrm{C} 15-\mathrm{O} 3-\mathrm{C} 16$ | $121.4(2)$ |
| $\mathrm{C} 9-\mathrm{O} 2-\mathrm{C} 5$ | $117.4(2)$ | $\mathrm{C} 5-\mathrm{C} 6-\mathrm{C} 7$ | $112.9(2)$ |

Table 2
Hydrogen-bond geometry ( $\AA^{\circ},{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 1-\mathrm{H} 1 B \cdots \mathrm{O}^{\mathrm{i}}$ | 0.96 | 2.58 | $3.482(4)$ | 156 |
| $\mathrm{C}^{\mathrm{ii}} 4-\mathrm{H} 14 \cdots \mathrm{O} 4^{\mathrm{iii}}$ | 0.93 | 2.55 | $3.463(4)$ | 168 |
| $\mathrm{C} 6-\mathrm{H} 6 A \cdots \operatorname{Cg}(B)^{\mathrm{iiv}}$ | 0.97 | 2.81 | $3.773(3)$ | 175 |
| $\mathrm{C} 21-\mathrm{H} 21 \cdots \operatorname{Cg}(A)^{\text {iv }}$ | 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: $C g(A)$ and $C g(B)$ denote the centroids of the aromatic rings C10-C15 and C17-C22, respectively

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

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