

References

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(Z)-3-Benzylidene-1-ethynyl-1-phenyl-1,3-dihydroisobenzofuran

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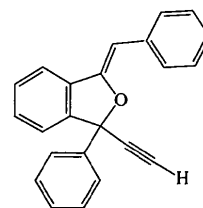
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Abstract

The title compound, C₂₃H₁₆O, has two independent molecules in the asymmetric unit. The bond lengths and angles of the two molecules are identical. The conformation of the isobenzofuran ring systems of the molecules, however, differ slightly; in molecule A, the sp²-C atom of the five-membered ring deviates the greatest distance [0.034 (2) Å] from the benzo plane, while in molecule B, it is the sp³-C atom of the five-membered ring which has the greatest deviation [0.027 (2) Å] from the benzo plane.

Comment

As part of a structural study involving [3]cumulene intermediates bearing the acetylenic functionality (Ramos & Garcia, 1997), the precursor [1-phenyl-1-(2-phenylethynyl)-3-trimethylsilyl-2-propyn-1-ol] to the title compound was prepared (Ramos & Garcia, 1997). The title compound resulted from an attempt to trimethylsilyl-deprotect the precursor under K₂CO₃ in anhydrous methanol conditions (Austin *et al.*, 1981; Garcia, Ramos, Pratt & Rodriguez, 1995; Garcia, Ramos, Rodriguez & Fronczek, 1995). The title compound, (I), was unexpectedly formed *via* a nucleophilic attack of the hydroxy O atom on the (phenyl)acetylenic functionality, resulting in the formation of the isobenzofuran ring system. Its crystal structure was determined in order to ascertain its identity.



(I)

The title compound exists as two independent molecules in the asymmetric unit, a case which is observed in about 4.6% of compounds having the *P2₁/c* space group (Brock & Dunitz, 1994). Bond distances and angles of the two molecules are the same within experimental error. The O1a—C1a, O1a—C15a, O1b—C1b and O1b—C15b bond lengths of 1.385 (2), 1.462 (2), 1.382 (2) and 1.463 (2) Å, respectively, compare well with corresponding lengths in a compound having a similar ring system, *i.e.* 2-isopropyl-4,4,7-trimethyl-1*H*-phenaleno[1,9-*bc*]furan-1-one (Weber *et al.*, 1975). In molecule A, the C15 atom lies essentially in the plane of the C9—C14 phenyl ring, deviating from it by only 0.004 (2) Å. The O1a and C1a atoms, however, deviate by −0.022 (1) and 0.034 (2) Å, respectively, from the same plane. Contrary to C15a, C15b deviates by −0.027 (2) Å from the plane of the C9—C14 phenyl ring. Atoms O1b and C1b do not deviate quite as much [0.018 (1) and 0.013 (2) Å, respectively] as C1a and O1a. Atoms C15a and C15b lie 0.090 (2) and 0.080 (2) Å out of the C18—C23 phenyl ring planes of their corresponding molecules. The C9a—C1a—C2a—C3a torsion angle is 179.5 (2)°, indicating a fairly planar environment about the C1a=C2a double bond. The C1b=C2b double bond is less planar, with a C9b—C1b—C2b—C3b torsion angle of 178.28 (2)°. The C16a≡C17a and C16b≡C17b triple-bond distances of 1.164 (3) and 1.168 (3) Å, respectively, agree with the expected length of 1.174 (4) Å (Allen *et al.*, 1987).

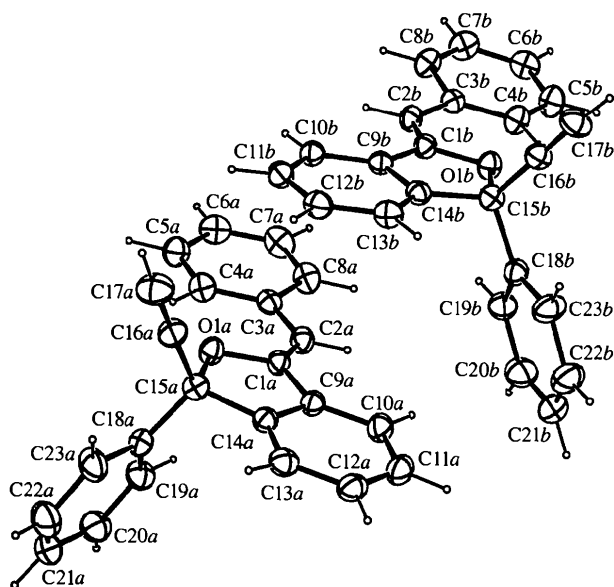


Fig. 1. The two independent molecules in the asymmetric unit with displacement ellipsoids plotted at the 20% probability level.

$$w = 4F_o^2 / [\sigma^2(F_o^2) + 0.0004F_o^4]$$

$$(\Delta/\sigma)_{\max} = 0.009$$

Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

Table 1. Selected geometric parameters (Å, °)

O1a—C1a	1.385 (2)	O1b—C1b	1.382 (2)
O1a—C15a	1.462 (2)	O1b—C15b	1.463 (2)
C1a—C2a	1.330 (3)	C1b—C2b	1.332 (3)
C1a—C9a	1.457 (3)	C1b—C9b	1.457 (3)
C2a—C3a	1.449 (3)	C2b—C3b	1.450 (3)
C9a—C14a	1.380 (3)	C9b—C14b	1.381 (3)
C14a—C15a	1.511 (3)	C14b—C15b	1.509 (3)
C15a—C16a	1.473 (3)	C15b—C16b	1.482 (3)
C15a—C18a	1.520 (3)	C15b—C18b	1.523 (3)
C16a—C17a	1.164 (3)	C16b—C17b	1.168 (3)
C1a—O1a—C15a	110.8 (1)	C1b—O1b—C15b	110.7 (1)
O1a—C15a—C14a	103.2 (1)	O1b—C15b—C14b	103.2 (1)
O1a—C15a—C16a	107.8 (1)	O1b—C15b—C16b	108.0 (1)
O1a—C15a—C18a	108.5 (1)	O1b—C15b—C18b	108.7 (1)
C14a—C15a—C16a	111.0 (2)	C14b—C15b—C16b	111.5 (2)
C14a—C15a—C18a	112.6 (2)	C14b—C15b—C18b	112.9 (2)
C16a—C15a—C18a	113.1 (2)	C16b—C15b—C18b	111.9 (2)

Experimental

Crystals of the title compound were grown by evaporation of a hexane solution.

Crystal data

C₂₃H₁₆O
M_r = 308.4
 Monoclinic
*P*2₁/*c*
a = 10.130 (1) Å
b = 31.398 (2) Å
c = 10.7388 (9) Å
 β = 90.852 (7)°
V = 3415.2 (9) Å³
Z = 8
D_x = 1.199 Mg m⁻³
D_m not measured

Cu *K*α radiation
 λ = 1.54184 Å
 Cell parameters from 15 reflections
 θ = 19–23°
 μ = 0.52 mm⁻¹
T = 294 K
 Rectangular prism
 0.20 × 0.17 × 0.15 mm
 Colorless

Data collection

Enraf–Nonius CAD-4 diffractometer
 $\theta/2\theta$ scans
 Absorption correction: none
 7589 measured reflections
 7057 independent reflections
 4394 reflections with $I > \sigma(I)$

*R*_{int} = 0.012
 θ_{\max} = 75°
 $h = -12 \rightarrow 12$
 $k = -39 \rightarrow 0$
 $l = 0 \rightarrow 13$
 3 standard reflections
 frequency: 120 min
 intensity decay: 0.5%

Refinement

Refinement on *F*²
R = 0.051
wR = 0.044
S = 1.512
 4394 reflections
 562 parameters
 All H atoms refined

$\Delta\rho_{\max}$ = 0.13 e Å⁻³
 $\Delta\rho_{\min}$ = -0.06 e Å⁻³
 Extinction correction: isotropic (Zachariasen, 1963)
 Extinction coefficient: 7.1 (3) × 10⁻⁷

A series of ψ scans showed no change in intensity with rotation about the diffraction vector. Thus, no absorption corrections were applied. The isotropic displacement parameters (*B*_{iso}) of the H atoms range from 4.8 (4) to 14.1 (8) Å² and C—H distances are within the range 0.90 (2) to 1.03 (2) Å.

Data collection: *CAD-4 Operations Manual* (Enraf–Nonius, 1977). Cell refinement: *CAD-4 Operations Manual*. Data reduction: *PROCESS MoIEN* (Fair, 1990). Program(s) used to solve structure: direct methods *MULTAN80* (Main *et al.*, 1980). Program(s) used to refine structure: *LSFM MoIEN*. Molecular graphics: *ORTEPII* (Johnson, 1976). Software used to prepare material for publication: *CIF IN MoIEN*.

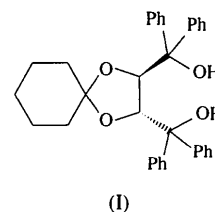
The purchase of the diffractometer was made possible by a National Science Foundation chemical instrumentation grant, which we gratefully acknowledge. RKT also appreciates the Chancellor Student Aid Fund which made structure determination possible. This research was supported (or partially supported) by a Research Centers in Minority Institutions award (#G12RR03062) from the Division of Research Resources, National Institutes of Health, and MBRS (grant #GM08247).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1131). Services for accessing these data are described at the back of the journal.

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The labeling scheme and displacement ellipsoids for the compound are depicted in Fig. 1. The cyclohexane ring adopts a chair conformation. Each of the four phenyl groups is each essentially planar, all deviations from the least-squares plane being less than 0.01 Å. The two dihedral angles between phenyl groups in the diphenylcarbinol moiety, one attached to C(2) and the other attached to C(3), are 74.8 (2) and 86.7 (2)°, respectively. Two diphenylcarbinol groups are also fixed by an intramolecular hydrogen bond between their hydroxyl groups forming a seven-membered ring. The remaining hydroxyl H atom appears to be involved in a weak intermolecular OH...π contact to a phenyl group. As shown in Fig. 2, van der Waals contacts between hydrophobic groups stabilize the crystal packing. Goldberg *et al.* (1990) reported the crystal structure of 2,2-dimethyl-4,5-bis(hydroxydiphenylmethyl)-1,3-dioxolane, which is a dimeric structure formed through intra- and intermolecular hydrogen bonds. Bond distances and angles in the 4,5-bis(hydroxydiphenylmethyl)-1,3-dioxolane moiety of the title compound are nearly the same as those in the 2,2-dimethyl derivative. However, the corresponding torsion angles differ slightly from each other and this may

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(*R,R*)-(-)-*trans*-2,3-Bis(hydroxydiphenylmethyl)-1,4-dioxaspiro[4.5]decane†

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Abstract

The title compound, C₃₄H₃₄O₄, contains an intramolecular hydrogen bond involving the hydroxyl group. The crystal packing is stabilized by van der Waals interactions between the hydrophobic surfaces of the neighboring molecules. The observed structure demonstrates novel packing features.

Comment

A host compound based on the 4,5-bis(hydroxydiphenylmethyl)-1,3-dioxolane framework selectively forms an inclusion complex with various guest components such as alcohols, nitriles and esters (for examples, see Mori & Toda, 1990; Toda, Matsuda & Tanaka, 1991). However, a prediction of optimum host–guest pairing is very difficult since the optical resolution capability is often changed dramatically by a slight difference in the structure of a host and/or a guest. We have carried out a single-crystal X-ray structural analysis of the title compound in order to study the effects of the acetal moiety on the crystal structure.

† IUPAC name: (*R,R*)-(-)-*trans*-1,4-dioxaspiro[4.5]decane-2,3-diyl-bis(diphenylmethanol).

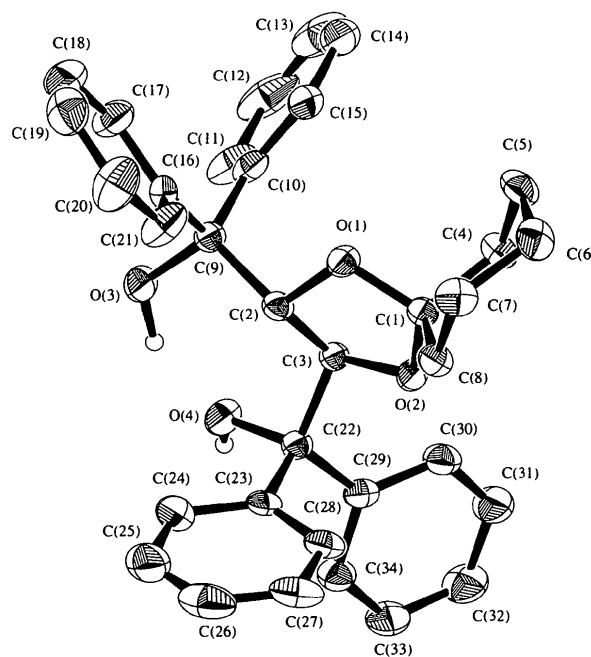


Fig. 1. The labeling scheme and displacement ellipsoids (30% probability) for the title molecule.