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Key indicators

Single-crystal X-ray study T = 100 KMean $\sigma(C-C) = 0.006 \text{ Å}$ R factor = 0.061 wR factor = 0.166 Data-to-parameter ratio = 11.5

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

© 2001 International Union of Crystallography Printed in Great Britain – all rights reserved In the title compound, (2S,3R,3aR)-2-(1,3-benzodioxol-5-yl)-3,3a-dihydro-3a-methoxy-3-methyl-5-(2-propenyl)-6(2H)-benzofuranone, C₂₀H₂₀O₅, the furan ring has a half-chair conformation with its O atom on the twist axis. The semiquinone ring is slightly non-planar, with a maximum deviation of 0.065 (3) Å.

Comment

The title neolignan, (I), has previously been reported from *Magnolia denudata* (Iida *et al.*, 1982; Kuroyanagi *et al.*, 2000), *Magnolia liliflora* (Iida & Ito, 1983) and *Magnolia soulangiana* (Abdallah, 1993).



The furan ring has a half-chair conformation with its O atom on the twist axis, as shown by the torsion angles in Table 1. The semiquinone ring is only slightly non-planar, with its six C atoms exhibiting an r.m.s. deviation of 0.039 Å from coplanarity, with a maximum deviation of 0.065 (3) Å. The propenyl group is twisted out of the semiquinone-ring plane primarily by rotation about the C7' - C8' bond, as indicated by the C1' - C7' - C8' - C9' torsion angle of -125.4 (4)°.

The most closely related neolignan for which the crystal structure has been previously reported is mirandin-*A* (Tomita, *et al.*, 1977), which differs from the title compound, (I), in having the opposite configuration at C5' and by having a 3,4,5-trimethoxyphenyl substituent rather than the 3,4-methylene-dioxyphenyl group of denudatin *A*. In mirandin-*A*, the furan ring also has a twist conformation, but with C7 on the twist axis. Its propenyl group also has a different conformation, with a C6'-C1'-C7'-C8' torsion angle of -92.7° and a C1'-C7'-C8'-C9' torsion angle of 2.0° .

Experimental

Leaves of *Magnolia soulangiana*, collected in Vancouver, BC, Canada, were air dried, ground and extracted with CH_2Cl_2 at room temperature for 24 h. The crude extract was separated by standard vacuum liquid chromatography procedures (Cantrell *et al.*, 1996),

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Figure 1

The atom-numbering scheme for (I) with ellipsoids at the 40% probability level.

using silica gel and n-hexane/ethyl acetate mixtures of increasing polarity. Fractions 63-66 (of 66) yielded crystals of denudatin A.

Crystal data

 $C_{20}H_{20}O_5$ $M_r = 340.36$ Monoclinic, C2 a = 17.8678(17) Å b = 6.7905 (18) Åc = 16.382(2) Å $\beta = 120.114 \ (10)^{\circ}$ $V = 1719.4 (5) \text{ Å}^3$ Z = 4Data collection

Enraf-Nonius CAD-4 diffractometer $\theta/2\theta$ scans Absorption correction: none 3590 measured reflections 2619 independent reflections 1813 reflections with $I > 2\sigma(I)$ $R_{\rm int}=0.056$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.061$ wR(F²) = 0.166 S = 1.042619 reflections 228 parameters H-atom parameters constrained

 $D_x = 1.315 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 25 reflections $\theta = 10.2 - 23.4^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 100 KNeedle, colorless $0.45\,\times\,0.20\,\times\,0.15~\text{mm}$

$\theta_{\rm max} = 30.0^{\circ}$
$h = -24 \rightarrow 21$
$k = 0 \rightarrow 9$
$l = -19 \rightarrow 21$
3 standard reflections
frequency: 120 min
intensity decay: <1%

 $w = 1/[\sigma^2(F_o^2) + (0.0922P)^2]$ + 0.03P] where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ _3 $\Delta \rho_{\text{max}} = 0.42 \text{ e Å}$ $\Delta \rho_{\rm min} = -0.23 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

O3-C2′	1.233 (5)	C3'-C4'	1.322 (5)
C1′-C6′	1.333 (5)	C8′-C9′	1.312 (7)
G1/ 01 67	100 7 (2)		100 5 (5)
C4′ – O4 – C/	108.7 (3)	C9 - C8 - C7	123.7 (5)
C4' - O4 - C7 - C8	14.1 (4)	04 - C4' - C5' - C8	-36.9(4)
C2-C1-C7-O4	-45.0(4)	C7-C8-C5'-C4'	42.3 (3)
O4-C7-C8-C5'	-35.7 (3)	C6'-C1'-C7'-C8'	4.3 (6)
C7-O4-C4'-C5'	14.5 (4)	C1′-C7′-C8′-C9′	-125.4(4)
C10′-O5-C5′-C4′	57.6 (4)		

The absolute configuration could not be determined. H atoms were placed in calculated positions with C-H bond distances in the range 0.95-1.00 Å and thereafter treated as riding. A torsional parameter was refined for each methyl group. $U_{iso} = 1.2U_{eq}$ of the attached atom (1.5 for methyl groups).

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1994); cell refinement: CAD-4 EXPRESS; data reduction: MAXUS (Mackay et al., 1999); program(s) used to solve structure: SIR97 (Altomare et al., 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 for Windows (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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