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# 2β-Hydroxy-4'-demethyldesoxypodophyllotoxin<sup>1</sup>

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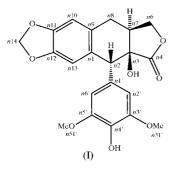
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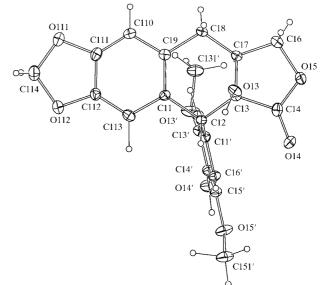
A room-temperature single-crystal X-ray structure determination of the title compound,  $C_{21}H_{20}O_8$ , confirms the stereochemical assignment made previously on the basis of spectroscopic studies [Shaari & Waterman (1994). *J. Nat. Prod.* **57**, 720–724].

### Comment

Some years ago (Shaari & Waterman, 1994), the title compound, (I), was isolated from *Casearia clarkei* King, a small lowland rainforest tree endemic to Malaysia and Singapore, and an assignment of the structure was made on the basis of spectroscopic and chiroptical evidence. An X-ray structure determination was considered desirable at the time, the better to secure the assignment, but specimens were only available as fine flakes of insufficient substance to be amenable to such a study by our contemporary single-counter diffractometer instrumentation. The advent of area-detector instrumentation has now permitted an effective attack on this problem, yielding a definitive assignment of the structure which confirms that proposed previously.



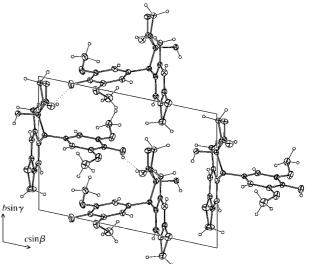
<sup>&</sup>lt;sup>1</sup> Systematic name: 5a-hydroxy-5-(4-hydroxy-3,5-dimethoxyphenyl)-5,5a,6,-8,8a,9-hexahydrofuro[3',4':6,7]naphtho[2,3-d]-1,3-dioxol-6-one.



#### Figure 1

Projection of molecule n = 1 (molecule 2 is similar) normal to the fusedring plane. Displacement ellipsoids are shown at 20% probability levels for the non-H atoms; H atoms are shown with an arbitrary radius of 0.1 Å.

The results of the room-temperature single-crystal X-ray study of (I) are consistent with the stoichiometry, connectivity and stereochemistry proposed previously; two molecules (n = 1, 2), pseudosymmetrically related, comprise the asymmetric unit of the structure and the unit cell in the uncommon triclinic space group *P*1. The skeletal form appears to be previously unrecorded in the Cambridge Structural Database (Allen & Kennard, 1993), but offers no surprises in respect of bond lengths and angles generally and these have been deposited. The absolute configuration was not assignable from the crystallographic work and follows that previously defined in the chemistry. The two molecules of the asymmetric unit are of the same chirality and of similar conformation; only one is shown in Fig. 1. The plane of the pendant ring is quasi-normal to that of the fused system; the unit-cell contents as displayed in Fig. 2



#### Figure 2

Unit-cell contents projected down  $\mathbf{a}$ , showing intermolecular hydrogen bonds within the cell.

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show the pseudosymmetry, the hydrogen bonding and parallel packing of the major molecular planes. The numbering of the non-H atoms (not systematic) within each molecule (n = 1,2 to be prefixed) is as shown in the Scheme above.

## **Experimental**

Material isolated in a previous study (Shaari & Waterman, 1994) was used in the present work.

### Crystal data

$C_{21}H_{20}O_8$	Z = 2
$M_r = 400.38$	$D_x = 1.435 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 7.8540 (10)  Å	Cell parameters from
$b = 9.3840 (10) \text{\AA}$	reflections
c = 13.230(2) Å	$\theta = 2-29^{\circ}$
$\alpha = 101.563 \ (2)^{\circ}$	$\mu = 0.111 \text{ mm}^{-1}$
$\beta = 103.699 \ (2)^{\circ}$	$T = 300 { m K}$
$\gamma = 90.184 \ (2)^{\circ}$	Plate, colourless
$V = 926.8 (2) \text{ Å}^3$	$0.20 \times 0.10 \times 0.05$ m

### Data collection

Bruker AXS CCD diffractometer  $\omega$  scans 10 605 measured reflections 4460 independent reflections 2592 reflections with  $F^2 > 2\sigma F^2$ 

### Refinement

Refinement on  $F^2$ R(F) = 0.040 $wR(F^2) = 0.044$ S = 1.0704460 reflections 681 parameters All H-atom parameters refined m 1868 mm

 $R_{\rm int}=0.068$  $\theta_{\rm max} = 28.4^{\circ}$  $h = -10 \rightarrow 10$  $k = -12 \rightarrow 12$  $l = -18 \rightarrow 18$ 

 $(\Delta/\sigma)_{\rm max} = 0.02$  $\Delta \rho_{\rm max} = 0.20 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\rm min} = -0.18 \text{ e } \text{\AA}^{-3}$ Extinction correction: Zachariasen (1967)Extinction coefficient: 2063.4 (7)

As might be expected from the above, data were limited in scope and intensity, nonetheless permitting solution of the structure, with stable and meaningful refinement of non-H atom anisotropic displacement parameters and  $(x, y, z, U_{iso})$ H, the latter permitting delineation of any hydrogen-bonding interactions, particularly of the hydroxyl-H atoms (Hn3 and Hn4'); for Hn3, none are discernible but those of Hn4' are of interest, being directed toward On5' [H···O 2.08 (5) and 2.24 (4) Å (n = 1, 2)], with further interactions being intermolecular [H14'...O24(x-1, y, z) 2.11 (7) and H24'...O14(1+x, y-1, z) 2.33 (5) Å] to the lactone carbonyl-O atoms. The usual exocyclic angular distortions are found at the points of fusion of the methoxy and hydroxy substituents to the pendant aromatic ring.

Data collection: SMART (Siemens, 1995); cell refinement: SAINT (Siemens, 1995); data reduction: SAINT; program(s) used to solve structure: Xtal3.5 GENTAN (Hall et al., 1995); program(s) used to refine structure: Xtal3.5 CRYLSQ; molecular graphics: Xtal3.5 PIG and ORTEP; software used to prepare material for publication: Xtal3.5 BONDLA and CIFIO.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: JA1004). Services for accessing these data are described at the back of the journal.

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