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Key indicators

Single-crystal X-ray study T = 291 KMean σ (C–C) = 0.005 Å R factor = 0.038 wR factor = 0.081 Data-to-parameter ratio = 7.3

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

Natural diterpenoid ludongnin, octahydro-5'-hydroxy-7a-methyl-8'-methylenespiro[isobenzofuran-4(1*H*),-4'(3'*H*)-[1*H*-7,9a]methanocyclohepta[c]pyran]-1',3,9'(3a*H*,4'a*H*)-trione

The asymmetric unit of ludongnin consists of two molecules of $C_{20}H_{24}O_6$. The hydroxyl group of each molecule participates in a hydrogen bond with the carbonyl oxygen of the δ -lactone moiety of a translationally related molecule, thus giving rise to the two symmetry-independent sets of infinite chains in the crystal, one of which is stretched along the *c* and another along the *a* axis of the crystal.

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Comment

The diterpenoid ludongnin, (I), has been previously isolated from *Rabdosia rubescens* Hara (Ruan *et al.*, 1986; Sun *et al.*, 1988; Zheng *et al.*, 1984, 1986) and its structure was established from the spectroscopic and chemical evidence. Recently, it was for the first time isolated from *Rabdosia var lophanthoides* Hara, and its structure was confirmed by an X-ray diffraction study (Fig. 1).

In both molecules, the *B* ring adopts a chair, whereas the *D* ring shows a boat conformation. The isobenzofuran ring system couples orthogonally to the methanocyclohepta[*c*]-pyran moiety through a spiro-C atom. The differences in the geometrical parameters between the two independent molecules are in fact negligible; the most notable discrepancies, as one would expect, concern the values of the torsion angles: C12-C13-C16-C17 -92.1 (5), C12'-C13'-C16'-C17' -100.1 (5), C14-C13-C16-C17 151.1 (4) and C14'-C13'-C16'-C17' 143.5 (4)°.



ludongnin

The hydroxyl group of each of the independent molecules participates in a hydrogen bond with the carbonyl oxygen of the δ -lactone moiety of the same molecule as transformed by the *c* or *a* translation [O···O 2.824 (4) and 2.856 (3) Å, respectively]. Thus, two independent molecules give rise to two sets of infinite chains in the crystal, one of which is stretched along the *c* and another along the *a* axis of the crystal (Fig. 2).

Experimental

© 2001 International Union of Crystallography 2 kg of dried powder of the whole herb of *Rabdosia var lophanthoides* were soaked three times with 95% EtOH at room temperature. The



Figure 1

One of the independent molecules of the title compound, shown with 50% probability displacement ellipsoids.



Figure 2

The crystal-packing diagram for the title compound viewed down the c axis. H atoms have been omitted for clarity.

ethanolic extracts were evaporated under reduced pressure and the residue was successively fractionated with petroleum ether, EtOAc and *n*-BuOH. The residue of the EtOAc fraction was subjected to column chromatography over silica gel. The column was eluted with a petroleum ether–EtOAc mixture; an acetone–chloroform mixture was used for recrystallization of the title compound. Its purity was checked by ¹H NMR spectroscopy. α_{12}^D –138° (c = 1, pyridine); UV λ_{max} (95%EtOH): 230 nm (ε 7130). Crystals suitable for X-ray structure analysis were obtained by slow evaporation from an aqueous solution containing chloroform and methanol in a 1:1 ratio at room temperature.

Crystal data

$C_{20}H_{24}O_6$	$D_x = 1.368 \text{ Mg m}^{-3}$
$M_r = 360.39$	Mo $K\alpha$ radiation
Monoclinic, P2 ₁	Cell parameters from 18
$a = 7.2397 (13) \text{\AA}$	reflections
$b = 36.724 (8) \text{\AA}$	$\theta = 3.1 - 14.2^{\circ}$
c = 7.304 (2) Å	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 115.69 \ (2)^{\circ}$	T = 291 (2) K
$V = 1749.9 (7) \text{ Å}^3$	Prism, colorless
Z = 4	$0.52 \times 0.44 \times 0.42 \text{ mm}$

Data collection

Siemens *P*4 diffractometer ω scans 3894 measured reflections 3474 independent reflections 2512 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.017$ $\theta_{\text{max}} = 26.0^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.081$ S = 0.91 3474 reflections 474 parameters H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0428P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $h = 0 \rightarrow 8$ $k = 0 \rightarrow 45$ $l = -9 \rightarrow 8$ 3 standard reflections every 97 reflections intensity decay: 3.6%

 $\begin{array}{l} (\Delta/\sigma)_{\rm max} = 0.001 \\ \Delta\rho_{\rm max} = 0.17 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.14 \ {\rm e} \ {\rm \AA}^{-3} \\ {\rm Extinction \ correction: \ SHELXL97} \\ {\rm Extinction \ coefficient: \ 0.0135 \ (12)} \\ {\rm Absolute \ structure: \ Flack \ (1983)} \\ {\rm Flack \ parameter \ = -0.4 \ (11)} \end{array}$

 Table 1

 Selected geometric parameters (Å, °).

O2-C15	1.207 (4)	O2′-C15′	1.213 (4)
O3-C7	1.210 (4)	O3'-C7'	1.204 (4)
O6-C6	1.193 (5)	O6'-C6'	1.196 (5)
C16-C17	1.310 (5)	C16′-C17′	1.320 (5)
C20-C10-C9	106.4 (3)	C20' - C10' - C5'	112.3 (3)
C1-C10-C9	113.5 (3)	C1′-C10′-C5′	108.9 (3)
O1-C11-C9	111.7 (3)	O1'-C11'-C9'	113.5 (3)
O2-C15-C16-C17	2.7 (6)	O2′-C15′-C16′-C17′	9.2 (6)
C8-C15-C16-C17	179.2 (4)	C8'-C15'-C16'-C17'	-172.9(3)
C14-C13-C16-C17	151.1 (4)	C14'-C13'-C16'-C17'	143.5 (4)
C12-C13-C16-C17	-92.1 (5)	C12'-C13'-C16'-C17'	-100.1 (5)

Table 2 Hydrogen-bonding geometry (Å, °).

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$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdots A$			
$\begin{array}{c} \text{O1-H1} A \cdots \text{O3}^{\text{i}} \\ \text{O1'-H1'} A \cdots \text{O3'}^{\text{ii}} \end{array}$	0.82 0.82	2.05 2.07	2.824 (4) 2.856 (3)	157 159			
Symmetry codes: (i) x, y, $z = 1$; (ii) $x = 1$, y, z.							

H atoms were placed in geometrically calculated positions and included in the final refinement in the riding-model approximation with the displacement parameters $U_{\rm iso}$ set equal to $1.2U_{\rm eq}$ of the corresponding carrier atom. As one would expect, both independent molecules have the same configuration. Lack of anomalous scatterers and, hence, a high s.u. of the Flack parameter, -0.4 (11), do not permit us to make an unambiguous assignment of the absolute configuration of the title compound.

Data collection: XSCANS (Siemens, 1994); cell refinement: XSCANS; data reduction: XSCANS; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP in SHELXTL/PC (Siemens, 1991); software used to prepare material for publication: SHELXL97.

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