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1(5),6(7)-Diepoxy-4-guaiol hemihydrate

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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(\text{C-C}) = 0.005 \text{ Å}$ R factor = 0.041 wR factor = 0.134Data-to-parameter ratio = 9.3

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

The title compound, $C_{15}H_{24}O_3 \cdot 0.5H_2O$, crystallizes in the space group $P2_1$ with two molecules in the asymmetric unit, both with the same conformation. In the crystal structure, $O-H\cdots O$ hydrogen bonds link the two independent molecules, A and B, into an infinite chain along the a axis. These chains are further interlinked by water molecules through $O-H\cdots O$ hydrogen bonds in different directions into supramolecular arrays.

Comment

1(5),6(7)-diepoxy-4-guaiol, also known as singuaiol, was isolated from the soft coral *sinularia sp.*, collected from Sanya Bay, Hainan Province, China. In this sesquiterpenoid, besides the usual epoxy group between atoms C6 and C7, there is a rare epoxy group between atoms C1 and C5; this is a new guaiane sesquiterpenoid. Its structure has been elucidated on the basis of spectroscopic analysis (Rao *et al.*, 2000). We undertook an X-ray study of 1(5),6(7)-diepoxy-4-guaiol to confirm the relative stereochemistry and present here the structure of its hemihydrate, (I).

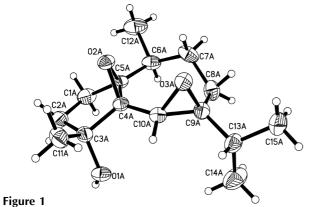
The X-ray study of (I) confirms the previously proposed structure based on spectroscopic data. The asymmetric unit of (I) consists of two independent molecules, A and B (Fig. 1), linked by hydrogen bonds $O1A-H1A\cdots O2B$ and $O1B-H1B\cdots O3A$ (Table 1). Its crystal structure shows that the two epoxy rings are approximately equilateral triangles. The C-O and C-C bond lengths are in the range 1.429 (3)–1.473 (4) Å, and the angles are in the range 58.98 (15)–61.00 (17)°.

In the solid-state structure, $O1A-H1A\cdots O2B$ and $O1B-H1B\cdots O3A$ hydrogen bonds link the two independent molecules into an infinite chain along the a axis. These chains are further interlinked by water molecules, through $O1W-H1WA\cdots O1B$ and $O1W-H1WB\cdots O1A$ hydrogen bonds, into supramolecular arrays (Fig. 2).

Experimental

Freshly collected soft coral *sinularia sp.* was cut into pieces and extracted with EtOH three times. The combined extracts were concentrated under reduced pressure and the crude extract was partitioned between H₂O and EtOAc. The EtOAc-soluble portion

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View of one molecule of the asymmetric unit of the title compound, with displacement ellipsoids drawn at the 30% probability level.

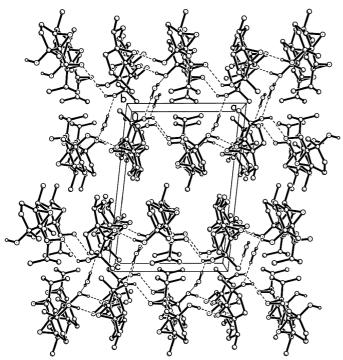


Figure 2 Perspective view, showing the molecular packing of (I), along the *b*-axis direction. All H atoms attached to C atoms have been omitted for clarity.

was subjected to silica-gel column chromatography, eluting with EtOAc–hexane, and afforded the product 1(5),6(7)-diepoxy-4-guaiol, according to NMR and two-dimensional NMR spectra. Crystals of (I) were obtained from hexane/EtOAc by solvent diffusion. ¹H NMR (500 MHz; CDCl₃): δ 1.73 (m, H2A), 2.01 (m, H2B), 1.56 (m, H3A), 1.62 (m, H3B), 3.15 (s, H6), 1.76 (m, H8A), 2.05 (m, H8B), 1.17 (m, H9A), 1.53 (m, H9B), 1.98 (m, H10), 1.07 (d, J = 7.0 Hz, H11), 1.54 (m, H12), 0.98 (d, J = 6.5 Hz, H13), 0.95 (d, J = 6.5 Hz, H14), 1.45 (s, H15).

Crystal data

$C_{15}H_{24}O_3 \cdot 0.5H_2O$	$D_x = 1.155 \text{ Mg m}^{-3}$		
$M_r = 261.35$	Mo $K\alpha$ radiation		
Monoclinic, P2 ₁	Cell parameters from 25		
a = 10.3492 (13) Å	reflections		
b = 9.9790 (13) Å	$\theta = 25-31^{\circ}$		
c = 14.6493 (18) Å	$\mu = 0.08 \text{ mm}^{-1}$		
$\beta = 96.382 (2)^{\circ}$	T = 293 (2) K		
$V = 1503.5 (3) \text{ Å}^3$	Rod, colorless		
Z=4	$0.50 \times 0.41 \times 0.35 \text{ mm}$		

Data collection

Bruker SMART CCD	2858 reflections with $I > 2\sigma(I)$		
diffractometer	$R_{\rm int} = 0.013$		
ω scans	$\theta_{\rm max} = 26.0^{\circ}$		
Absorption correction: none	$h = -12 \rightarrow 12$		
8011 measured reflections	$k = -12 \rightarrow 11$		
3127 independent reflections	$l = -18 \rightarrow 11$		

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0967P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.041$	+ 0.1971P]
$wR(F^2) = 0.134$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.08	$(\Delta/\sigma)_{\text{max}} = 0.001$
3127 reflections	$\Delta \rho_{\text{max}} = 0.26 \text{ e Å}^{-3}$
335 parameters	$\Delta \rho_{\min} = -0.23 \mathrm{e \mathring{A}^{-3}}$
H-atom parameters constrained	

Table 1 Hydrogen-bonding geometry (Å, °).

D $ H$ $\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$\begin{array}{c} O1A - H1A \cdots O2B \\ O1B - H1B \cdots O3A^{i} \\ O1W - H1WA \cdots O1B \\ O1W - H1WB \cdots O1A^{ii} \end{array}$	0.85	1.98	2.814 (3)	167
	0.85	1.98	2.806 (3)	166
	0.86	2.02	2.882 (4)	179
	0.86	2.14	3.002 (5)	179

Symmetry codes: (i) 1 + x, y, z; (ii) $2 - x, y - \frac{1}{2}, -z$.

The H atoms were positioned geometrically and were treated as riding on their parent C and O atoms, with C—H distances of 0.96 Å and O—H distances of 0.85 Å. The Friedel pairs were merged during the refinement, and the absolute configuration is indeterminate from the diffraction data.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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