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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.002 \text{ Å}$ R factor = 0.035 wR factor = 0.095 Data-to-parameter ratio = 10.3

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

8β -Hydroxy-2,6,6-trimethyl- $5\beta H$,10 βH eudesma-1,7(11)-dien-12,8-olide

In the title compound, C₁₅H₂₀O₃, molecules are linked in an infinite chain along the a axis, with graph-set motifs $C_2^2(12)$ or $C_2^2(14)$, through O $-H \cdot \cdot \cdot$ O hydrogen bonds.

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Comment

 8β -Hydroxy-2,6,6-trimethyl- $5\beta H$,10 βH -eudesma-1,7(11)-dien-12,8-olide, (I), was isolated from the marine sponge Dysidea avara; previously it has been isolated from the marine sponge Dysidea fragilis (Zhong et al., 1995). A pharmacological experiment shows that, when the concentration of compound (I) is 5.58×10^{-4} M, it significantly inhibited spontaneous movement of rabbit ileum, controlled autonomy activity and muscular activity.

The X-ray study of (I) further confirms the previously proposed structure based on spectroscopic data. The asymmetric unit of (I) consists of one independent molecule (Fig. 1). In the crystal structure, molecules are linked in an infinite chain along the a axis, described as $C_2^2(12)$ or $C_2^2(14)$ according to graph-set notation (Etter, 1990), through O3-H1···O1 hydrogen bonds (Fig. 2 and Table 1). The most characteristic feature is that (I) can be described as a hydrophobic-hydrophilic layered structure along the c axis, with the two sixmembered rings forming the hydrophobic portion, while the five-membered lactone rings form the hydrophilic portion.

In the crystal structure, the middle six-membered ring adopts a chair conformation and one of the attached methyl groups and the hydroxy group occupy axial positions.

Experimental

Freshly collected sponge Dysidea avara was cut into pieces and extracted with 95% EtOH three times. The combined extracts were concentrated in vacuo and the crude extract was partitioned between water and EtOAc. The EtOAc-soluble portion was subjected to vacuum liquid chromatography (VLC), eluting with EtOAc-hexane, to yield (I). The compound's identity was confirmed by NMR spectroscopy. Crystals of (I) were obtained from hexane-EtOAc by solvent diffusion. ¹H NMR (500 MHz, CDCl₃): 5.66 (1H, s, H11), 5.37 (1H, m, H1), 2.82 (1H, dd, J = 13.0 and 4.0 Hz, H10), 2.32 (1H, dd, J = 13.0 and 4.0 Hz, H10)14.0 and 4.0 Hz, H9A), 1.95 (2H, m, H4), 1.71 (1H, m, H5), 1.67 (1H,

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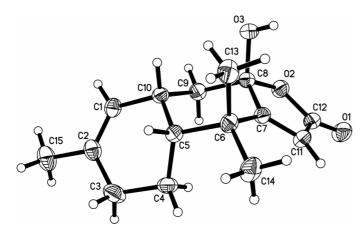


Figure 1 View of the title compound, showing 30% probability displacement ellipsoids. H atoms are drawn as small spheres of arbitrary radii.

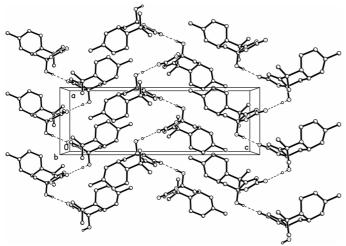


Figure 2 Packing arrangement of the title compound, projected on the *ac* plane. All H atoms attached to C atoms have been omitted for clarity. Hydrogen bonds are shown as dashed lines.

m, H3B), 1.62 (1H, s, H15), 1.55 (1H, t, J = 13.5 Hz, H9B), 1.41 (1H, s, H13), 1.22 (1H, s, H14), 1.13 (1H, m, H3A) ¹³C NMR: 123.6 (d, C1), 134.3 (s, C2), 18.7 (t, C3), 31.0 (t, C4), 47.3 (d, C5), 38.6 (s, C6), 175.3 (s, C7), 105.5 (s, C8), 40.9 (t, C9), 30.4 (d, C10), 114.9 (d, C11), 170.8 (s, C2), 26.8 (q, C13), 25.3 (q, C14), 23.11 (q, C15).

Crystal data

 $C_{15}H_{20}O_3$ $M_r = 248.31$ Orthorhombic, $P2_12_12_1$ a = 7.1843 (9) Å b = 8.7554 (11) Å c = 21.310 (3) Å V = 1340.4 (3) Å³ Z = 4 D_x = 1.230 Mg m⁻³ Mo $K\alpha$ radiation Cell parameters from 929 reflections θ = 2.5–27.0° μ = 0.08 mm⁻¹ T = 293 (2) K Block, colorless 0.50 × 0.40 × 0.35 mm

Data collection

 $\begin{array}{lll} \text{Bruker SMART CCD area-detector} & 1714 \text{ independent reflections} \\ \text{diffractometer} & 1619 \text{ reflections with } I > 2\sigma(I) \\ \varphi \text{ and } \omega \text{ scans} & R_{\text{int}} = 0.015 \\ \text{Absorption correction: multi-scan} & \theta_{\text{max}} = 27.0^{\circ} \\ (SADABS; \text{ Sheldrick, 1996}) & h = -9 \rightarrow 4 \\ T_{\text{min}} = 0.959, \, T_{\text{max}} = 0.971 & k = -10 \rightarrow 11 \\ 7880 \text{ measured reflections} & l = -27 \rightarrow 26 \\ \end{array}$

Refinement

 $\begin{array}{lll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_{\rm o}^2) + (0.0623P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.035 & + 0.1304P] \\ wR(F^2) = 0.095 & where $P = (F_{\rm o}^2 + 2F_{\rm c}^2)/3$ \\ S = 1.05 & (\Delta/\sigma)_{\rm max} < 0.001 \\ 1714 \ \mbox{reflections} & \Delta\rho_{\rm max} = 0.14 \ \mbox{e Å}^{-3} \\ 167 \ \mbox{parameters} & \Delta\rho_{\rm min} = -0.19 \ \mbox{e Å}^{-3} \end{array}$

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

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D-\mathrm{H}\cdots A$
O3-H3···O1 ⁱ	0.82	2.00	2.8128 (19)	172

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

H atoms were positioned geometrically and were treated as riding on the parent C and O atoms, with C—H distances of 0.93–0.98 Å and an O—H distance of 0.82 Å [$U_{\rm iso}({\rm H})=1.2U_{\rm eq}({\rm C})$ or $1.5U_{\rm eq}({\rm C,O})$]. In the absence of significant anomalous scattering effects, Friedel pairs were merged; the absolute configuration was assigned arbitrarily.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT-Plus* (Bruker, 1999); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

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