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25-Deoxycacospongionolide B and Cacospongionolide C, Two New Terpenoids from the Sponge *Fasciospongia Cavernosa*

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Abstract: 25-Deoxycacospongionolide B (3), a new bioactive sesterterpene, and a diterpene, cacospongionolide C (4), have been isolated as minor components from the Adriatic sponge *Fasciospongia cavernosa*. The structures were proposed on the basis of spectroscopic data and defined for the former on the basis of X-ray analysis. Brine shrimp and fish lethaliities of 25-deoxycacospongionolide B are reported.

In the course of our search for marine natural compounds that have biological activities, we isolated and characterized a sesterterpenoid, cacospongionolide (1)^{1,2}, with high cytotoxic activity, from the sponge *Fasciospongia cavernosa* (erroneously classified as *Cacospongia mollior*)³ a horny sponge belonging to the family Thorectidae. Subsequently, we made an extensive harvesting of other Mediterranean horny sponges belonging to the same family in order to find compounds related to cacospongionolide.

We recently isolated cacospongionolide B (2)³ from some specimens of *F. cavernosa* Schmidt (order Dictyoceratida; family Thorectidae) collected in the North Adriatic. This sesterterpene exhibited cytotoxic and antimicrobial activities and the structure was established by spectral and chemical means³. We continued investigation on extracts of this sponge and purification of fraction containing sterols by HPLC gave a new sesterterpene, 25-deoxycacospongionolide B (3, 0.02% dry weight), and a diterpenoid, cacospongionolide C (4, 0.006%). The isolation, the structure elucidation, including X-ray analysis and some biological activities of 3, and the structure of 4 are reported in this paper.

Cacospongionolide C (4) had $[\alpha]_D -18^\circ$ and a molecular formula $C_{20}H_{36}O_3$ from HRMS of the parent ion. The UV absorption at 220 nm, IR bands at 3380, 1785, 1760 cm^{-1} , ¹H-NMR signals at δ 5.99 (brs, 1H) and 5.86 (brs, 1H) and ¹³C-NMR signals at δ 170.7 (s), 169.3 (s), 117.6 (d) and 99.6 (d) define the γ -hydroxybutenolide moiety, which has previously been encountered in other metabolites isolated from this sponge^{1,3}. The analysis of the spectral data established the presence of a saturated acyclic chain from C-4 to C-19, with three additional secondary methyl groups. The presence of fragment ions at m/z 263, 221, 193, 151 and 123 in the mass spectrum of 4 located the methyl groups on C-7, C-11 and C-15. No attempt was made to determine the stereochemistry of the methyl-bearing carbons at C-7 and C-11.

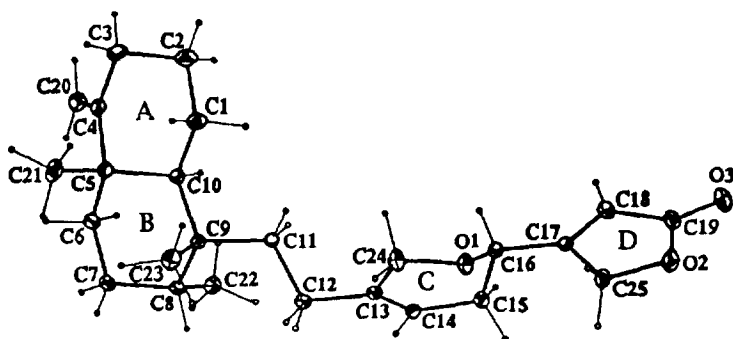


Figure 1. A perspective view of 25-deoxycacospongionolide B (3): labels for H-atoms were omitted for clarity.

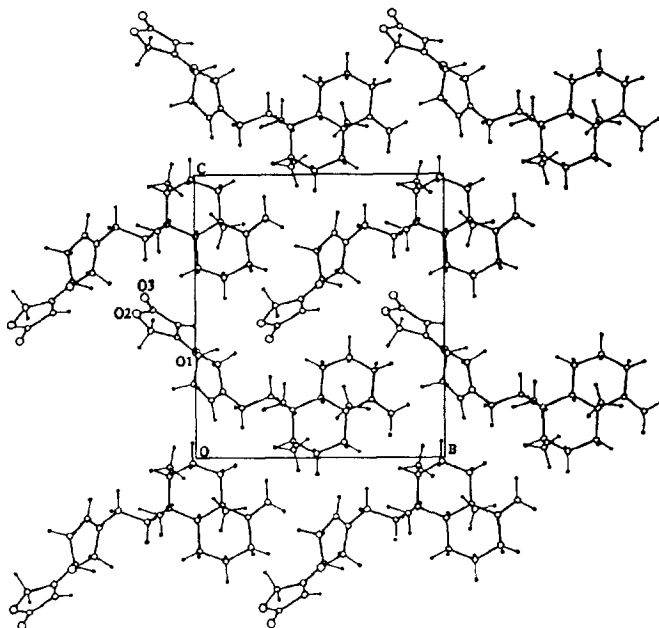


Figure 2. The crystal packing of compound 3 projected onto the *bc* plane: only oxygen atoms are labelled.

A selection of intramolecular geometry is given in Table 2. The trend of bond lengths and bond angles agrees well with the values observed in similar molecules^{2,5,6}. In the bicyclic carbon-system, few significant deviations of the geometrical parameters from normal values are probably caused by the atomic crowding. In particular the lengthening of the bond distances along the atomic sequence C21,C5,C10,C9,C23, and the widening of the angles C21-C5-C10, C5-C10-C9, and C10-C9-C23 help to release the short 1,5 interaction

Table 1 - Compound 3: Fractional coordinates and equivalent isotropic thermal parameters (\AA^2) with e.s.d.'s in parentheses for non-H atoms: $\text{Beq} = \frac{1}{3} \sum_i \sum_j b_{ij} a_i^* a_j^*$

atom	x	y	z	Beq	atom	x	y	z	Beq
O1	0.5274(3)	0.0000	0.3823(1)	4.68(4)	C12	0.2955(5)	0.1855(2)	0.1783(2)	4.59(5)
O2	0.9394(3)	-0.2354(2)	0.5185(2)	5.80(5)	C13	0.4489(4)	0.1085(2)	0.2365(2)	3.89(4)
O3	1.2789(4)	-0.2023(3)	0.5748(2)	7.47(6)	C14	0.6126(5)	0.0625(2)	0.2048(2)	4.63(6)
C1	0.2365(5)	0.5114(3)	0.3214(2)	5.39(6)	C15	0.7705(5)	-0.0092(3)	0.2660(2)	4.80(6)
C2	0.3472(6)	0.6167(3)	0.3588(2)	6.42(8)	C16	0.7451(4)	0.0095(2)	0.3704(2)	3.60(4)
C3	0.2366(6)	0.7173(3)	0.3108(2)	6.09(7)	C17	0.8645(4)	-0.0734(2)	0.4349(2)	3.65(4)
C4	0.2101(4)	0.7076(2)	0.2041(2)	4.36(5)	C18	1.0644(4)	-0.0693(3)	0.4774(2)	4.52(5)
C5	0.0960(4)	0.6042(2)	0.1634(2)	3.98(5)	C19	1.1159(4)	-0.1709(3)	0.5291(2)	4.98(6)
C6	0.0980(5)	0.5950(2)	0.0555(2)	4.79(5)	C20	0.2845(5)	0.7857(3)	0.1524(2)	5.97(7)
C7	0.0168(5)	0.4844(2)	0.0152(2)	4.92(6)	C21	-0.1392(4)	0.6170(3)	0.1822(3)	6.88(9)
C8	0.1508(4)	0.3900(2)	0.0613(2)	3.77(5)	C22	0.3713(5)	0.3916(3)	0.0289(2)	5.05(6)
C9	0.1522(3)	0.3881(2)	0.1708(2)	3.23(4)	C23	-0.0692(4)	0.3524(3)	0.1937(2)	5.20(6)
C10	0.2201(3)	0.5033(2)	0.2128(2)	3.31(4)	C24	0.4058(4)	0.0882(3)	0.3364(2)	5.25(6)
C11	0.3162(4)	0.3029(2)	0.2175(2)	3.77(4)	C25	0.7714(5)	-0.1796(2)	0.4587(2)	5.08(6)

Table 2 - Compound 3: Molecular geometry with e.s.d.'s in parentheses

<i>bond lengths</i> (\AA)							
O1 - C16	1.416(3)	C3 - C4	1.502(4)	C8 - C9	1.551(3)	C13 - C24	1.498(4)
O1 - C24	1.427(3)	C4 - C5	1.526(4)	C8 - C22	1.528(4)	C14 - C15	1.508(4)
O2 - C19	1.358(4)	C4 - C20	1.327(4)	C9 - C10	1.562(3)	C15 - C16	1.526(3)
O2 - C25	1.436(3)	C5 - C6	1.533(4)	C9 - C11	1.551(3)	C16 - C17	1.496(3)
O3 - C19	1.202(3)	C5 - C10	1.571(3)	C9 - C23	1.544(4)	C17 - C18	1.325(3)
C1 - C2	1.523(5)	C5 - C21	1.556(4)	C11 - C12	1.534(4)	C17 - C25	1.481(4)
C1 - C10	1.530(3)	C6 - C7	1.526(4)	C12 - C13	1.511(4)	C18 - C19	1.453(4)
C2 - C3	1.525(5)	C7 - C8	1.522(4)	C13 - C14	1.311(4)		
<i>bond angles</i> ($^\circ$)							
C16-O1-C24	110.9(2)	C6-C5-C21	108.6(2)	C11-C9-C23	107.1(2)	O1-C16-C17	106.8(2)
C19-O2-C25	108.7(2)	C10-C5-C21	115.7(2)	C1-C10-C5	110.9(2)	C15-C16-C17	112.3(2)
C2-C1-C10	111.6(2)	C5-C6-C7	112.9(2)	C1-C10-C9	114.7(2)	C16-C17-C18	128.4(2)
C1-C2-C3	111.3(3)	C6-C7-C8	111.6(2)	C5-C10-C9	116.1(2)	C16-C17-C25	123.2(2)
C2-C3-C4	111.4(3)	C7-C8-C9	111.5(2)	C9-C11-C12	116.9(2)	C18-C17-C25	108.3(2)
C3-C4-C5	115.3(2)	C7-C8-C22	109.9(2)	C11-C12-C13	111.5(2)	C17-C18-C19	109.3(3)
C3-C4-C20	120.0(3)	C9-C8-C22	114.9(2)	C12-C13-C14	123.9(2)	O2-C19-O3	120.4(3)
C5-C4-C20	124.8(3)	C8-C9-C10	109.3(2)	C12-C13-C24	115.5(2)	O2-C19-C18	108.4(2)
C4-C5-C6	111.6(2)	C8-C9-C11	110.3(2)	C14-C13-C24	120.6(2)	O3-C19-C18	131.2(3)
C4-C5-C10	107.3(2)	C8-C9-C23	109.5(2)	C13-C14-C15	122.9(2)	O1-C24-C13	113.5(2)
C4-C5-C21	105.4(2)	C10-C9-C11	107.9(2)	C14-C15-C16	108.7(2)	O2-C25-C17	105.3(2)
C6-C5-C10	108.3(2)	C10-C9-C23	112.6(2)	O1-C16-C15	109.8(2)		
<i>selected torsion angles</i> ($^\circ$)							
C4-C5-C10-C9	-171.0(2)	C23-C9-C10-C5	-71.2(2)	O1-C16-C17-C18	-150.6(2)		
C21-C5-C10-C9	71.7(3)	C10-C9-C11-C12	-173.2(2)	C16-C17-C18-C19	-176.9(2)		
C22-C8-C9-C10	72.9(3)	C9-C11-C12-C13	-173.9(2)	C17-C18-C19-O3	177.9(3)		
C11-C9-C10-C5	170.7(2)	C11-C12-C13-C14	-111.8(3)				

C21...C23 = 3.259(5) Å. The geometry at C4 (see Table 2) is significantly distorted with respect to normal C(sp²) values. Such variations are rather common in similar cyclic systems^{7,8} and seem to contribute in preserving the chair conformation of the ring A, in spite of the presence of the C4-C20 double bond.

Rings A and B are in almost ideal chair conformations which deviate towards half-boat forms. Atomic displacements from the respective best planes through the remaining endocyclic atoms are: C3 = 0.629(4), C10 = 0.694(2) Å (ring A) and C7 = 0.681(3), C10 = 0.607(2) Å (ring B). The pyranose ring (C), which blats a double bond in the 13-14 positions, adopts a distorted half-chair conformation with O1 and C16 atoms displaced 0.320(1) and 0.443(2) Å, respectively and on the apposite side of the average plane through the remaining four atoms. For this ring the distortion is essentially toward a twist form, as shown by the puckering parameters⁹: $q_2 = 0.397(2)$ Å, $\phi_2 = 24.1(4)^\circ$ and $\theta = 51.7(3)^\circ$. γ -Lactone system is planar to within 0.013(3) Å. Conformational details for each ring are given in Table 3 (deposited).

In the absence of hydrogen donors the crystal packing (Fig. 2) is governed only by van der Waals interactions, shortest intermolecular contacts being O1...C18_(x-1, y, z) = 3.494(3) and C3...C19_(x-1, y+1, z) = 3.557(4) Å.

25-Deoxycacospongionolide B showed a high cytotoxicity (LC₅₀) = 0.74 µg/ml) comparable to that of **2**³, in the *Artemia salina* bioassay^{10,11} while it was less toxic (no toxicity at 100 µg/ml) than **2**³, in the fish lethality¹².

EXPERIMENTAL¹³

Isolation: from fraction containing sterols after HPLC purification (Spherisorb S5W; n-hexane/ethylacetate, 4:1; flow 3 ml/min) were recovered 25-deoxycacospongionolide B that crystallized from MeOH (14 mg, 0.02% dry weight) and cacospongionolide C (4 mg, 0.006%).

25-Deoxycacospongionolide B (3): white crystals, mp 159-161°C; $[\alpha]_D = 10.5^\circ$ (c = 1.40, CHCl₃); IR ν_{\max} (CHCl₃) 1745, 1650, 1110 cm⁻¹; EIMS (70 eV) m/z (%) [M]⁺ 384.2661 (C₂₅H₃₆O₃ requires 384.2664) (41), 369 (18), 205 (31), 193 (52), 191 (100), 189 (90), 177 (85); ¹H-NMR¹⁴ (500 MHz, in CDCl₃) δ 5.95 (1H, brs, H-18), 5.54 (1H, brs, H-14), 4.87 (2H, ABq, J = 17.9 Hz, H-25), 4.51 (1H, brs, H-20a), 4.48 (1H, brs, H-20b), 4.33 (1H, ddd, J = 10.5, 4.2, 1.5 Hz, H-16), 4.16 (2H, ABq, J = 15.7, H-24), 2.30 (1H, ddd, J = 13.6, 13.6, 5.1 Hz, H-3 β), 1.15 (1H, dd, J = 12.0, 2.3 Hz, H-10), 1.09 (3H, s, H-21) 0.92 (3H, s, H-23), 0.89 (3H, d, J = 6.9 Hz, H-22); ¹³C-NMR¹⁴ (125 MHz, in CDCl₃) δ 173.5 (s, C-19), 169.8 (s, C-17), 160.4 (s, C-4), 138.4 (s, C-13), 115.7 (d, C-14), 114.7 (d, C-18), 102.1 (t, C-20), 71.0 (t, C-25), 70.3 (d, C-16), 68.3 (t, C-24), 47.8 (d, C-10), 40.4 (s, C-5), 38.0 (s, C-9), 37.8 (t, C-11), 35.4 (d, C-8), 32.9 (t, C-3), 30.1 (t, C-2), 29.9 (t, C-15), 28.7 (t, C-7), 26.0 (t, C-12), 25.5 (t, C-6), 21.3 (t, C-1), 21.3 (q, C-21), 20.5 (q, C-23), 14.8 (q, C-22).

Cacospongionolide C (4): colorless oil; $[\alpha]_D = -18.0^\circ$ (c = 0.32, CHCl₃); IR ν_{\max} (CHCl₃) 3380 (br), 1785, 1760 cm⁻¹; EIMS (70 eV) m/z (%) [M]⁺ 324.2667 (C₂₀H₃₆O₃ requires 324.2664) (32), 306 (30), 291 (9), 263 (6), 221 (23), 193 (40), 179 (54), 151 (95), 123 (100); ¹H-NMR (500 MHz, in CDCl₃) δ 5.99 (1H, brs), 5.86 (1H, brs), 2.40 (2H, brt, J = 6.5 Hz), 0.87 (3H, d, J = 6.5 Hz), 0.86 (6H, d, J = 6.6 Hz), 0.84 (3H, d, J = 6.5 Hz); ¹³C-NMR (125 MHz, in CDCl₃) δ 170.8 (s), 169.3 (s), 117.6 (d), 98.6 (d), 39.4 (t), 37.4 (t), 37.3 (t), 36.6 (t), 32.8 (d), 32.6 (d), 28.0 (d), 27.9 (t), 24.8 (t), 24.4 (t), 24.2 (t), 22.7 (q), 22.6 (q), 19.7 (q), 19.6 (q).

Crystal data and structure determination of 3: Single crystals of 25-deoxycacospongionolide B were obtained by slow concentration of a methanol solution and a sample of size 0.4x0.4x0.1 mm, was selected for the crystallographic study. Accurate cell parameters were obtained by least-squares refinement of the setting angles of 25 reflections at medium θ (22° < θ < 26°), using Ni-filtered CuK α radiation and Enraf-Nonius CAD-4F diffractometer on line with a MicroVAX Digital computer.

Crystal data: C₂₅H₃₆O₃, Mw = 384.56; monoclinic system, space group P2₁, with Z = 2: a = 6.3294(8), b = 12.197(1), c = 14.162(3) Å, β = 97.95(2)°, V = 1082.8(6) Å³, D_c = 1.179 g/cm³. 2303 Independent reflections (θ_{max} = 74°) were collected at room temperature, using ω-2θ scan mode, as suggested by peak-shape analysis. During the data collection, the intensities of three standard reflections were monitored every 4h (4% variation) in order to check the crystal and equipment stability. No significant intensity decay was observed. The intensities were corrected for Lorentz and polarization factors, but not for the absorption effect (μ = 5.55 cm⁻¹). The structure was solved by direct methods using SIR92 package⁴. The positional and anisotropic temperature parameters for the 28 non-hydrogen atoms were refined by full-matrix (on F) least-squares method, y coordinate of O1 atom was fixed to define the origin. At convergence, all the hydrogen atoms were unambiguously located on a difference Fourier map and they were included in the last refinement at the calculated positions (C-H distance = 1.02 Å), as fixed atoms with isotropic thermal parameters set equal to B_{eq} of the parent atom. At convergence, the final discrepancy index $R = \sum |F_o| - |F_c| / \sum |F_o|$ was 0.038 for the 2039 observed reflection with $I \geq 2.5 \sigma(I)$, R_w = 0.045 with $w^{-1} = [\sigma^2(F_o) + (0.02F_o)^2 + 0.2]$; S = 0.962, (Δσ)_{max} < 0.01; number of refined parameters = 253 including the extinction coefficient¹⁵ = 5.2(4)·10⁻⁶. No residual electron density is outside the range -0.11±0.13 e·Å⁻³. Atomic scattering factors were taken from Cromer and Waber¹⁶. For the crystallographic data the equipment of the C.I.M.C.F. of the University "Federico II" of Naples was used. All calculations were performed using Enraf-Nonius SDP software¹⁷ on a MicroVAX 3100 computer.

Structure factors, anisotropic thermal parameters, hydrogen atom parameters, and complete molecular geometry, together with conformational details for each ring (Table 3), have been deposited.

Biological evaluations.- As per reference¹².

a.-*Brine shrimp lethality.*- The brine shrimp (*Artemia salina*) lethality assay performed as already described^{10,11}, gave LC₅₀ 0.74 μg/ml (1.26/0.44 μg/ml, 95% confidence limits)

b.-*Fish lethality.*- The fish (*Gambusia affinis*) lethality assay performed as already described¹², gave no toxicity at 100 μg/ml.

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