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Further syphonosides from the sea hare *Syphonota geographica* and the sea-grass *Halophila stipulacea*

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Abstract

The unusual structural features of syphonoside (1), recently reported from the marine mollusc *Syphonota geographica* and its prey *Halophila stipulacea*, stimulated further investigations on the minor secondary metabolites of both organisms. The three novel macrocyclic glycoterpenoids 2–4, structurally related to the main co-occurring metabolite 1, have been isolated and chemically characterized mainly by NMR spectroscopic techniques and degradation methods. Compounds 2 and 3 were found only in the mollusc whereas compound 4 was isolated in trace quantities exclusively from the sea-grass.

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1. Introduction

In the course of our investigations on natural products from marine opisthobranchs, ^{1–12} we have recently analyzed the secondary metabolite content obtained from a Mediterranean population of the Lessepsian anaspidean mollusc *Syphonota geographica*, collected along the Greek coasts. ^{9,12} A number of degraded sterols, structurally related to those found in other anaspidean molluscs, ^{13–15} characterized the skin metabolite pattern of *S. geographica* whereas the main secondary metabolite of the viscera was found to be the glycoterpenoid macrocycle syphonoside (1). ¹² This compound was also found to be the main component of the butanol extract of the invasive sea-grass *Halophila stipulacea*, collected together with the mollusc.

The presence of syphonoside (1) in both organisms supported the trophic relationship between the mollusc and the

sea-grass, ¹² previously suggested by the detection of *H. stipulacea* fragments in the stomach content of *S. geographica*. ⁹ From the chemical point of view, syphonoside (1) is particularly interesting because it has an unusual framework in which a diterpene moiety is incorporated in a macrocycle along with a residue of 3-hydroxy-3-methyl glutaric acid and a glucopyranosyl moiety. The structure and the absolute stereochemistry of compound 1 were determined by a combination of spectroscopic techniques, chemical degradation methods, and conformational analysis. ¹² In a preliminary cytotoxicity evaluation, 1 inhibited high density induced apoptosis in selected human and murine cancer cell lines, ¹² suggesting a possible involvement in the regulation of the cell survival and death under specific conditions.

The intriguing structural features of syphonoside (1) prompted us to study the minor components of the glycoterpenoid fractions from the extracts of *S. geographica* and *H. stipulacea*. Herein we report the results of this investigation that led to the isolation and the chemical characterization of three novel glycoterpenoid macrocycles, compounds 2–4,

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structurally related to the main metabolite 1. Compounds 2 and 3 were isolated from the diethyl ether extract of the mollusc, whereas compound 4 co-occurred with the main metabolite 1 in the butanol extract of the sea-grass.

2. Results and discussion

The diethyl ether soluble portion (589 mg) of the acetone extract, obtained as already described from the viscera of seven frozen S. geographica individuals, 9,12 was analyzed by TLC. Some polar components that were absent in the mantle of the mollusc were revealed by reaction with cerium sulfate. The extract was submitted to a chromatographic purification on a LH-20 Sephadex column, to afford fractions A and B. Preliminary ¹H NMR spectroscopic analysis of these fractions showed that both of them contained glycosyl compounds structurally related to syphonoside (1), previously isolated from the n-BuOH soluble portion of the same acetone extract. 12 Fractions A and B were further purified on preparative silica gel TLC plates (CHCl3/MeOH eluent system) to obtain syphonosideol (2, 12.0 mg), the main component of the ether extract, and a mixture of syphonoside fatty acid ester derivatives 3 (5.4 mg).

A selected fraction obtained by LH-20 Sephadex chromatography of an aliquot (500 mg) of *n*-BuOH layer from the acetone extract of the sea-grass sample¹² was found to contain an additional syphonoside related metabolite as revealed by TLC and ¹H NMR analysis. This fraction was further purified by preparative silica-gel TLC chromatography (CHCl₃/MeOH eluent system) to yield the syphonoside acetyl derivative **4** (8.3 mg).

The new compounds **2–4** possessed the same macrocycle structure as the co-occurring syphonoside (**1**) and differed in the nature of the substituent at C-11 in the lateral chain of the diterpenoid unit. The structural elucidation of these molecules is described starting from syphonosideol (**2**), which was the main component of the diethyl ether extract from the viscera of the mollusc.

Syphonosideol (2) displayed a pseudo-molecular peak at m/z 647 in the HRESI-MS spectrum, indicating the molecular formula $C_{32}H_{48}O_{12}$, which was consistent with a structure containing one glucose unit less than 1. Consistently, the ¹H NMR spectrum of 2 contained only one set of glycosyl signals resonating at δ 3.49–4.62 along with a doublet at δ 5.51 (J=8 Hz) due to the anomeric sugar proton (H-1') involved

in an ester linkage. Analogously with compound 1, the proton NMR spectrum of syphonosideol (2) displayed five methyl signals at δ 1.78 (3H, br s, H₃-16), 1.39 (3H, s, H₃-6"), 1.26 $(3H, s, H_3-19), 1.02 (3H, d, J=7 Hz, H_3-17), and 0.95 (3H, d, J=7 Hz, H_3-17)$ s, H_3 -20); two olefinic multiplets at δ 5.36 (1H, m, H-14) and 6.54 (1H, t, J=3 Hz, H-3); a methylene at δ 4.56 (1H, dd, J=9 and 13 Hz, H-15a) and 4.70 (1H, m, H-15b) and a methine at δ 4.00 (1H, dd, J=2 and 8 Hz, H-11) both linked to an oxygen atom (Table 1). The ¹³C NMR spectrum of 2 showed the presence of 25 sp³ carbon signals (five CH₃, nine CH₂, eight CH and three C as deduced by the DEPT sequence), and 7 sp² carbon signals corresponding to two double bonds and three ester carbonyl groups (Table 1), as in syphonoside (1). A careful analysis of the 1D and 2D NMR spectra of compound 2 indicated the close similarity to 1, due to the presence of the same macrocyclic carbon framework constituted by a clerodane acyl moiety oxidized at C-11, C-15, and C-18, a glucopyranosyl unit esterified at both H-1' and H₂-6' and a 3-hydroxy-3-methyl-glutaric acyl residue. Similarly to 1, the α,β-unsaturated acyl function at C-18 of the clerodane moiety esterifies the hydroxyl group at the anomeric position H-1' of the β-glucose unit, the primary hydroxyl group (6'-OH) of which is in turn linked to the carboxyl group (C-5") of 3-hydroxy-3-methyl glutaric acid, further esterified at C-1" with a primary hydroxyl group (15-OH) of the same clerodane, as reported in formula 2. In fact, diagnostic HMBC correlations were observed between C-18 (δ 167.5) and H-1' (δ 5.51), between C-5" (δ 170.6) and H₂-6' (δ 4.30 and 4.62) as well as between C-1"(δ 172.4) and H₂-15 (δ 4.56 and 4.70). Thus it was suggested that the only difference between compounds 1 and 2 was in the hydroxyl group at C-11, which was free in syphonosideol (2) and glycosylated by an additional glucose moiety in 1.

In order to confirm the proposed structure and the relative stereochemistry of **2**, some degradation and derivatization reactions were performed. Initially, a sample of syphonosideol (2) was acetylated to give the tetra-acetyl derivative **5**, which

2 R = R₁ = H **3** R = H, R₁ = $CH_3(CH_2)_{14}CO$ - or $CH_3(CH_2)_{16}CO$ -**5** R = R₁ = Ac

was fully characterized (Table 1). Diagnostic acylation shifts were observed for the protons of the three carbinol groups (H-2', H-3', and H-4') of the glucose unit as well as for H-11 of the clerodane portion, confirming the presence of the free hydroxyl at C-11 in compound 2.

Analogously with syphonoside (1), the alkaline hydrolysis (Na, MeOH anhyd) of compound 2 was very informative

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