



Mycaloside A, a new steroid oligoglycoside with an unprecedented structure from the Caribbean sponge *Mycale laxissima*

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Abstract—The structure of mycaloside A (**1**) isolated from the Caribbean sponge *Mycale laxissima* has been established as (2*E*,20*R*,24*S*)-3-*O*-{ α -D-Galp(1→2)- β -D-Arap(1→3)-[β -D-Galp(1→4)]- β -D-Glcp}-3 β ,4 β ,15 α ,21-tetrahydroxy-24-methylcholesta-5,22-diene by interpretation of spectral data and chemical transformations. © 2002 Elsevier Science Ltd. All rights reserved.

In the animal kingdom, steroid and triterpene glycosides are the predominant metabolites of starfishes and sea cucumbers, respectively.^{1–3} More recently, this type of glycoside has been isolated from some sponges. To the best of our knowledge, about 20 sponge glycosides have been reported to date, including sarasinosides from *Asteropus* spp.^{4–7} and *Melophlus isis*,⁸ erylosides and formosides from *Erylus* spp.,^{9–12} ulososides from *Ulosa* sp.^{13,14} pachastrelloside A from a *Pachastrella* sp.,¹⁵ wondosterols from unidentified

sponges and¹⁶ ectyoplasides and feroxosides from *Ectyoplasia ferox*^{17,18} The majority of these glycosides belong to nor-lanostane-triterpenoid saponins, derived from lanosterol or related triterpenes as a result of oxidative elimination of one or two methyl groups.

In the course of our continuing interest in marine oligoglycosides,¹⁹ we have isolated a steroid oligoglycoside, mycaloside A (**1**) from the Caribbean sponge *Mycale laxissima* (Demospongiae, order Poecilosclerida,

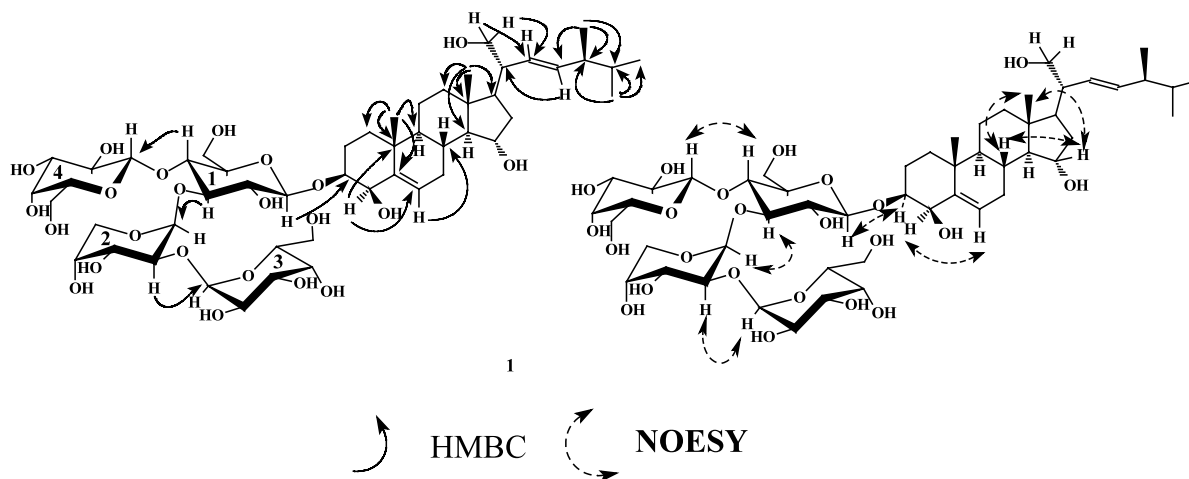


Figure 1. Key HMBC and NOESY correlations of Mycaloside A.

Keywords: marine metabolites; steroid glycosides; two-dimensional NMR spectroscopy.

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family Mycalidae). The sponge was collected by hand near San-Felipe Island, Cuba. The lyophilized specimens (0.3 kg) were macerated and sequentially extracted with ethyl acetate (2×2 L) and ethanol (3×2 L). Compound **1** (28 mg) was obtained from ethanol-soluble materials by partition between water and *n*-butanol followed by column chromatography of the butanol extract on Sephadex LH-20 (CHCl₃–EtOH, 1:1), Polychrome-1 (50% EtOH), silica gel L (CHCl₃–EtOH, 5:4) and by HPLC on Zorbax ODS (45% EtOH) and Zorbax-Si (CHCl₃–EtOAc–MeOH, 4:6:5) columns. Mycaloside A (**1**), mp 217–220°C, [α]_D²⁵ –23.0 (*c* 0.5, MeOH), analysed for C₅₁H₈₄O₂₃ by combined HRMALDI TOF MS (positive mode) and ¹³C NMR analyses. A quasi-molecular ion peak at *m/z* 1087.5250 (M+Na)⁺ (C₅₁H₈₄NaO₂₃ requires *m/z* 1087.5297) was detected. Hydrolysis of **1** with 2 M HCl (2 h, 100°C) gave glucose (1), arabinose (1) and galactose (2), identified by GLC as the corresponding aldonitrile perac-

etates. The monosaccharides themselves were isolated from the hydrolysate by HPLC on Separon™ SGX NH₂ column (90% acetonitrile) and shown to be of the D-series by measurement of their optical rotations.

The sequence of monosaccharides, interglycosidic linkages and configurations of glycosidic bonds in **1** were determined by NMR including HMBC and NOESY (see Fig. 1) and various ¹H–¹H COSY experiments (Table 1). In addition, the structure of the carbohydrate chain was confirmed by methylation of **1** (DMSO, NaH, CH₃I) followed by methanolysis and acetylation, which gave methyl 2,3,4,6-tetra-*O*-methylgalactopyranoside, methyl 2-*O*-acetyl-3,4-di-*O*-methylarabinopyranoside and methyl 3,4-di-*O*-acetyl-2,6-di-*O*-methylglucopyranoside, identified by comparison with standard samples using GLC and GLC-MS. Only D-glucose was not destroyed upon periodate oxidation followed by acid hydrolysis. It was evident from consideration of

Table 1. NMR data of carbohydrate moiety in **1** (DRX500,^a C₅D₅N+5% CD₃OD)

C	DEPT	δ H (J, Hz)	NOESY	C	DEPT	δ H (J, Hz)	NOESY
1 ¹	101.8, d	4.89, d (7.8)	H-3; H-4; H ¹ ₃ ; H ¹ ₅	1 ³	103.3, d	5.77, d (3.6)	H ² ₂
2 ¹	76.3, d	4.20, dd (7.8; 9.1)	H ¹ ₄	2 ³	71.0, d ^b	4.60, dd (3.5; 10.0)	
3 ¹	77.4, d	4.47, t (9.2)	H ¹ ₅	3 ³	71.2, d	4.58, dd (3.1; 9.9)	
4 ¹	74.5, d	4.71, t (9.6)	H ¹ ₂	4 ³	70.9, d ^b	4.38 ^c	
5 ¹	77.6, d	3.75, dt (9.8; 2.8)	H ¹ ₃	5 ³	73.1, d	4.81,ddd (1.3; 4.7; 7.3)	H ³ ₄ ; H ³ ₃
6 ¹	60.6, t	4.44 ^c ; 4.72 ^c		6 ³	62.9, t	4.27, dd (4.6; 11.3) 4.39 ^c	
1 ²	99.3, d	6.61, d (3.5)	H ¹ ₃ ; H ² ₂	1 ⁴	104.2, d	5.31,d (7.7)	H ¹ ₄ ; H ⁴ ₃ ; H ⁴ ₅
2 ²	80.2, d	4.73, dd (3.5; 10.0)	H ² ₁	2 ⁴	72.7, d	4.41, dd (7.7; 9.4)	
3 ²	69.1, d	5.19, dd (3.4; 10.0)	H ² ₄	3 ⁴	74.8, d	4.10, dd (9.5; 3.4)	H ⁴ ₄ ; H ⁴ ₅
4 ²	70.7, d	4.84, brd (3.3)	H ² ₃ ; H ² ₅	4 ⁴	69.2, d	4.47, brd (3.4)	H ⁴ ₅
5 ²	64.4, t	3.99, dd (1.9; 12.0) 5.51, brd (12.0)	H ² ₄	5 ⁴	76.2, d	3.95, brt (6.5)	H ⁴ ₄
				6 ⁴	61.6, t	4.33, dd (6.1; 10.8); 4.37 ^c	

^a 500 MHz for ¹H and 125.8 MHz for ¹³C, assignments were based on COSY45, COSY RCT, DEPT and HMQC experiments.

^b Values can be interchanged.

^c Coupling constants were not determined as a result of overlapping of signals.

Table 2. NMR data of aglycone part in **1**^a

C	DEPT	δ H (J, Hz)	NOESY	C	DEPT	δ H (J, Hz)	NOESY
1	37.6, t	1.77, m (e) 1.02, m (a)	H-11 H-3	15	73.4, d	4.21, td (9.0; 3.2)	H-8; CH ₃ -18
2	23.9, t	1.79, m (e) 2.28, m (a)	H-3	16	41.0, t	2.04, m 2.18, m	H-15
3	80.1, d	3.84, dt (12.2; 3.0)	H-1, H-4	17	48.7, d	2.04, m	H-22
4	74.4, d	4.57, d (2.7)	H-3, H-6	18	13.9, q	0.83, s	H-8; H-11; H-20
5	141.9, s			19	21.1, q	1.40, s	H-8
6	128.7, d	5.73, dd (2.1; 5.2)	H-4	20	48.3, d	2.36, m	H-18
7	32.9, t	2.70, dt (18.9; 5.4) 2.27, m	H-6; H-8, H-15 H-6; H-9; H-14	21	64.9, t	4.01, dd (4.4; 10.5) 3.78, dd (7.0; 10.5)	H-12
8	32.1, d	1.87, m	H-15	22	132.3, d	5.48, dd (8.6; 15.4)	H-17
9	50.8, d	1.02, m		23	135.2, d	5.45, dd (7.6; 15.3)	H-20
10	36.5, s			24	43.5, d	1.95, m	
11	20.6, t	1.45, m	H-1	25	33.4, d	1.50, m	
12	39.6, t	2.01, m (e) 1.39, m (a)	H-21	26	19.7, q	0.86, d (7.0)	
13	43.2, s			27	20.2, q	0.86, d (7.0)	
14	63.4, d	1.43, m	H-9	28	18.1, q	0.96, d (7.0)	

^a 500 MHz for ¹H and 125.8 MHz for ¹³C, assignments were based on COSY45, COSY RCT, DEPT and HMQC experiments.

mass and NMR data, that the aglycone in **1** must be a C₂₈ tetrahydroxylated steroid, probably similar to 3 β ,4 β -dihydroxy-pregn-5-en-20-one 3-sulfate from the sponge *Stylopus australis*²⁰ in respect of the A and B ring structures. The 5(6)-position of the double bond and 3 β ,4 β -dihydroxy-substitution were confirmed by ¹H–¹H COSY, HMBC and values of coupling constants (see Fig. 1 and Table 2). The presence of a hydroxy group at C-15 was determined from HMQC and COSY 45 experiments, while its α -configuration followed from H-15 coupling constants and NOESY. Further analysis of NMR data revealed the presence of an additional hydroxyl at C-21, the 22*E* double bond ($J_{22,23}$ = 15.3 Hz) and a C-24 methyl group in the side chain. The *S* configuration at C-24 was suggested based on comparison of the ¹³C NMR spectrum of **1** with those of related steroids.²¹ NOESY and especially HMBC data indicated that the carbohydrate chain was attached to C-3 in **1**.

Mycaloside A is the first steroid oligoglycoside isolated from sponges. So far, only nor-triterpene oligoglycosides^{4–14,17,18} and a few steroid biosides with aglycones oxidized at positions 2 α ,3 β ,4 β and 7 α ^{15,16} were isolated from these animals.

Carbohydrate chains such as that of **1** have also not been found. Therefore, **1** has an unprecedented structure consisting of both carbohydrate and aglycone moieties and is unique when compared with all the previously known sponge glycosides.

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