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# A triterpenoid saponin from Cucumaria frondosa

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#### **Abstract**

The structure of a new triterpenoid saponin, frondoside D, isolated from *Cucumaria frondosa* has been determined principally by high field 1D and 2D NMR and FAB-MS spectrometry. Frondoside D was shown to be  $3\beta$ -{3-O-methyl-O- $\beta$ -D-glucopyranosyl-(1-3)-O- $\beta$ -D-xylopyranosyl-(1-4)-[O- $\beta$ -D-xylopyranosyl-(1-2)]-O- $\beta$ -D-quinovopyranosyl-(1-2)-O- $\beta$ -D-4-sulfonatoxylo-pyranosyl}-16 $\beta$ -acetoxy-23S-hydroxy-holost-7-ene, sodium salt. © 1998 Published by Elsevier Science Ltd. All rights reserved.

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### 1. Introduction

Recently we reported the structure of two novel oligosaccharides isolated from the common sea cucumber Cucumaria frondosa Gunnerus (Findlay, Yayli & Radics, 1992). One of these was a saponin designated frondoside B, (3) comprising a  $3\beta$ -hydroxyholosta-7,24-diene aglycone and a disulfated pentasaccharide sidechain. The major saponin from this source, frondoside A, (2) (Girard et al., 1990; Yayli, 1993) features a  $16\beta$  acetoxy-holosta-7-ene aglycone and a monosulfated pentasaccharide sidechain which differs from that of frondoside B (3) by the presence of a xylose unit in place of the  $(1 \rightarrow 3)$  attached glucose-6-sulfate feature. Of particular interest was the isolation of the dimeric pentasaccharide frondecaside (Findlay, Yayli & Radics, 1992; Yayli, 1993) featuring six sulfate moieties.

#### 2. Results and discussion

From *C. frondosa* we have now characterized an additional saponin designated frondoside D (1). Frondoside D (1) is closely related to frondoside A (2)

and frondoside A<sub>1</sub> (4) (Avilov et al., 1993), differing only by the presence of an hydroxyl group at C-23. The structure of frondoside D was deduced from 1D and 2D high field NMR data and supported by positive FAB-MS.

Conventional <sup>1</sup>H NMR (500 MHz) and <sup>13</sup>C NMR (125 MHz) spectra, combined with DEPT data, afforded the structure of the oligosaccharide moiety which is identical with the monomeric oligosaccharide part of frondoside A (2) and frondoside A<sub>1</sub> (4) (Girard et al., 1990; Yayli, 1993; Avilov et al., 1993).

Analysis of the spectral data for frondoside D and comparison with those published for related saponin aglycones (Findlay, Yayli & Radics, 1992; Girard et al., 1990; Yayli, 1993; Avilov et al., 1993; Kitagawa et al., 1981; Kalinin, Stonik, Kalinovskii & Isakov, 1989; Stonik et al., 1982; Stonik, Mal'tsev, Kalinovskii & Elyakov, 1982) shows that the aglycone part of frondoside D is a holostane skeleton featuring a hydroxyl group at C-23 (Avilov et al., 1993; Kitagawa et al., 1981; Kalinin, Stonik, Kalinovskii & Isakov, 1989; Stonik et al., 1982; Stonik, Mal'tsev, Kalinovskii & Elyakov, 1982).

The positive FAB-mass spectrum (MNBA) of frondoside D (1) displayed a single peak in the higher mass range at m/z 1373 corresponding to  $[M + Na]^+$ (M =  $C_{60}H_{95}O_{30}SNa$ ), a composition differing from that of frondoside A (2) by the presence of an ad-

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ditional oxygen. The <sup>13</sup>C NMR chemical shift inventory of 1 (Table 1) closely parallels that of frondoside A except for signals assigned to C-22 (46.78), C-23 (66.24) and C-24 (48.49) which are shifted downfield by  $\delta$  7.65, 43.48 and 8.86 ppm, respectively, in agreement with the presence of an OH group at C-23. The aglycone sidechain is thus comparable to that of stichlorogenol (Kitagawa et al., 1981; Kalinin, Stonik, Kalinovskii & Isakov, 1989; Stonik et al., 1982; Stonik, Mal'tsev, Kalinovskii & Elyakov, 1982), a C-23-hydroxy aglycone from the sea cucumber Stichopus chloronotus stereochemistry have been confirmed by Xray crystallography (Kitagawa et al., 1981). Comparable signals in the <sup>13</sup>C NMR spectrum (pyridine- $d_5$ ) of the latter are observed at  $\delta$  47.62, 65.70 and 49.27 (Kitagawa et al., 1981), suggesting the 23S configuration in frondoside D. The <sup>13</sup>C NMR chemical shifts inventory (Table 1) for the oligosaccharide chain of frondoside D are virtually identical with those of frondoside A and A<sub>1</sub> (Girard et al., 1990; Yayli, 1993; Avilov et al., 1993). Thus we conclude that frondoside D possesses structure 1, that is  $3\beta$ -{3-O-methyl-O- $\beta$ -Dglucopyranosyl-(1-3)-O- $\beta$ -D-xylopyranosyl-(1-4)-[O- $\beta$ -D-xylopyranosyl-(1-2)]-O- $\beta$ -D-quinovopyranosyl-(1-2)- $O-\beta$ -D-4-sulfonato-xylopyranosyl}-16 $\beta$ -acetoxy-23Shydroxy-holost-7-ene, sodium salt.

#### 3. Experimental

### 3.1. Instrumentation

Mass spectra were recorded with a Kratos MS50 instrument. Optical rotations were measured with a Perkin-Elmer 241 polarimeter. NMR spectra were obtained at 20°C with a Bruker AMX-500 spectrometer and are referred to internal tetramethylsilane.

# 3.2. Isolation of saponins

A crude glycoside containing mixture (2.50 g) obtained from *C. frondosa* as previously described (Findlay, Yayli & Radics, 1992) was chromatographed on a silica gel 60 (100 g, 230–400 mesh) column eluting with a discontinuous gradient of CHCl<sub>3</sub>–MeOH (4:1–4:4) and CHCl<sub>3</sub>–MeOH–H<sub>2</sub>0 (4:3:1) to give 26 fractions ( $\sim$ 20 ml each) which were combined on the basis of TLC analysis to provide four sub-fractions a, b, c and d.

#### 3.3. Frondoside D, 1

Fraction c (9–13, 410 mg) was rechromatographed on a reversed-phase silica column (6 g, LiChroprep RP-18) eluting with a discontinuous gradient of

Table 1 <sup>13</sup>C NMR data for frondoside D, (1) in pyridine-*d*<sub>5</sub>–D<sub>2</sub>O (5:2)

Aglycone of 1 <sup>a</sup>			Sugar moiety 1 <sup>a</sup>		
С	<sup>13</sup> C (δ, ppm)	DEPT	С	<sup>13</sup> C (δ, ppm)	DEPT
1	36.04	CH <sub>2</sub>	Xyl I, 1	104.44	СН
2	26.84	$CH_2$	2	81.69	CH
3	89.19	CH	3	76.15	CH
4	39.47	C	4	76.30	CH
5	47.90	CH	5	64.18	$CH_2$
6	24.10	$CH_2$			-
7	120.41	CH	Qui, 1	102.20	CH
8	145.91	C	2	82.66	CH
9	47.71	СН	3	75.12	CH
10	35.43	C	4	85.23	CH
11	23.27	$CH_2$	5	71.20	CH
12	31.54	CH	6	17.99	$CH_3$
13	59.30	C			- 3
14	47.31	C	Xyl II, 1	104.66	CH
15	44.35	$CH_2$	2	73.59	СН
16	75.20	CH	3	86.07	CH
17	55.27	СН	4	68.89	CH
18	180.87	C	5	65.90	$CH_2$
19	24.30	$CH_3$			
20	86.42	C	MGlc, 1	104.36	CH
21	30.33	$CH_3$	2	74.58	CH
22	46.78	$CH_2$	3	86.91	CH
23	66.24	CH	4	70.52	СН
24	48.49	$CH_2$	5	77.45	CH
25	28.10	CH	6	61.85	$CH_2$
26	21.77	$CH_3$	•	60.97	OCH <sub>3</sub>
27	23.93	CH <sub>3</sub>			,
28	32.79	CH <sub>3</sub>	Xyl III, 1	105.33	CH
29	17.46	CH <sub>3</sub>	2	74.90	CH
30	28.78		3	76.54	CH
					СН
					CH <sub>2</sub>
30 31 32	28.78 171.09 21.60	CH <sub>3</sub> C CH <sub>3</sub>	3 4 5	76.54 70.09 66.54	

<sup>&</sup>lt;sup>a</sup> Chemical shifts (ppm) are relative to internal TMS.

CH<sub>3</sub>COCH<sub>3</sub>-MeOH-H<sub>2</sub>0 (2:2:3-2:2:4) solvent system to give 42 fractions (1-3 ml each) which were combined on the basis of TLC analysis to provide 5 subfractions. Sub-fraction cc (9-19, 0.320 mg) was further purified on a reversed-phase silica column (6 g, LiChroprep RP-18) eluting with CH<sub>3</sub>COCH<sub>3</sub>-MeOH- $H_2O$  (2:2:4) to give 74 fractions (1-3 ml each). On the basis of TLC, fractions were combined to provide 7 subfractions. Sub-fraction ccc (17-29, 121 mg wet) was chromatographed by prep. TLC (1 mm,  $20 \times 20$  cm, 2 plates) using CHCl<sub>3</sub>-MeOH-H<sub>2</sub>O (3:2.5:0.5) solvent system to give 3 major bands. The least polar band ccca (13.4 mg,  $R_f = 0.64$ ) was finally purified by a reversed-phase silica column (5 g, LiChroprep RP-18) eluting with CH<sub>3</sub>COCH<sub>3</sub>-CH<sub>3</sub>OH-H<sub>2</sub>0 (2:2:5) to give frondoside D [7.0 mg,  $R_f = 0.25$ , rpTLC, CH<sub>3</sub>COCH<sub>3</sub>-MeOH-H<sub>2</sub>O (2:3:4)]; m.p. 217-220°,  $[\alpha]_D^{23} = -22.9^{\circ}$  [c = 0.0013, pyridine-H<sub>2</sub>0 (1:4)]; <sup>1</sup>H NMR d (ppm) [pyridine-d<sub>5</sub>-D<sub>2</sub>O (5:2), 500 MHz] 0.92 (H-27), 0.95 (H-26), 1.12 (H-29), 1.20 (H-19), 1.27 (H-

28), 1.29 (H-30), 1.68 (H-21), 2.10 (H-32), 3.30 (H-3), 3.72 (H-23), 4.89 ( $J=7.0\,\mathrm{Hz}$ , Xyl I H-1), 4.90 ( $J=7.6\,\mathrm{Hz}$ , Xyl II H-1), 5.31 ( $J=7.7\,\mathrm{Hz}$ , Qui H-1), 5.32 ( $J=7.3\,\mathrm{Hz}$ , Xyl III H-1), 5.38 ( $J=7.6\,\mathrm{Hz}$ , MGlc H-1), 5.70 (H-7), 5.95 (H-16); <sup>13</sup>C NMR [pyridine- $d_5$ –D<sub>2</sub>O (5:2), 125 MHz] d (ppm) see Table 1; positive FAB-mass (MNBA) m/z (rel. int): 1373 (1.8) [M + Na]<sup>+</sup>, 1351 (0.9) [M + H]<sup>+</sup>, 838 (1.2) [sugar moiety + H]<sup>+</sup>, 609 (23) [MGlc–O-Xyl II–O-Qui–O-Xyl III]<sup>+</sup>, 530 (09) [M-aglycone + H]<sup>+</sup>, 493 (8.6) [MGlc–O-Xyl II–O-Qui(O)-O]<sup>+</sup>, 325 (9.2) [MGlc–O-Xyl II]<sup>+</sup>, 177 (100) [MGlc]<sup>+</sup>.

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