

Note

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# Notes

### Ceramide and Cerebrosides from the Octocoral Sarcophyton ehrenbergi#

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Received June 17, 2008

Chemical investigation of the octocoral *Sarcophyton ehrenbergi*, collected at the Dongsha Islands, Taiwan, has led to the isolation of a known ceramide (1) and two new cerebrosides, sarcoehrenosides A (2) and B (4), along with three known cerebrosides (3, 5, and 6). The structures of the new compounds were established by spectroscopic and chemical methods. Sarcoehrenoside A (2) differs from previously known marine cerebrosides in that it possesses a rare  $\alpha$ -glucose moiety. Compounds 1-6 were evaluated for antimicrobial activity against a small panel of bacteria and for anti-inflammatory activity using RAW 264.7 macrophages.

Cerebrosides are a large group of molecules containing a ceramide unit and one or more sugars. The hydrophobic ceramide moiety includes a sphingoid base and an amide-linked fatty acid chain. Cerebroside derivatives have been isolated from various marine invertebrates, such as sea stars, soft corals, sponges, and ascidians. In recent years several different types of biological activities have been found for these compounds, including antifungal, antitumor, antiviral, cytotoxic, and immunomodulatory properties.

In the course of a search for bioactive substances from marine sources,  $^7$  chromatographic separation of an alcyonacean soft coral,  $Sarcophyton\ ehrenbergi\ Marenzeller\ (Octocorallia: Alcyonacea), collected at the Dongsha Islands, Taiwan, has afforded two new cerebrosides, sarcoehrenosides A (2) and B (4). Also obtained were a known ceramide (1) and three known cerebrosides (3, 5, and 6), <math display="inline">^{3a,9,10}$  which were isolated from S. ehrenbergi for the first time. To our knowledge, sarcoehrenoside A (2), in possessing a rare  $\alpha\text{-D-glucose}$  moiety at C-1 of the glycerol ether unit, differs from previous cerebrosides isolated from marine organisms. In this paper, we describe the isolation, structural elucidation, and evaluation of the antimicrobial and anti-inflammatory activity of these metabolites.

Sarcoehrenoside A (2) was isolated as a white, amorphous powder and exhibited a [M + Na]<sup>+</sup> peak at m/z 774.5490, corresponding to a molecular formula of  $C_{43}H_{77}NO_9$  in the positive HRESIMS. The IR absorptions at 1638 and 1547 cm<sup>-1</sup> and  $^{13}C$  NMR signals at  $\delta$  54.6 (C-2) and 175.4 (C-1') suggested a secondary amide in the molecule. In the  $^{1}H$  and  $^{13}C$  NMR spectra of 2, the characteristic signals of an amide linkage (a nitrogenated methane proton at  $\delta$  3.93 and an amide carbonyl carbon at  $\delta$  175.4), a long

chain (terminal methyl protons at  $\delta$  0.90 and methylene protons at  $\delta$  1.29–1.38), and a sugar (an anomeric proton at  $\delta$  4.87) were observed, strongly suggesting the glycosphingolipid nature of 2. The  $\alpha$ -glucopyranose moiety was indicated by the anomeric proton at  $\delta$  4.87 (1H, d, J = 4.5 Hz, H-1") and the chemical shifts in the <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 1). The anomeric proton of the α-glucose unit exhibited a long-range <sup>1</sup>H-<sup>13</sup>C correlation with C-1 (Figure 1), demonstrating attachment of the glucose moiety at C-1. The relative configuration of the  $\alpha$ -glucopyranose moiety was determined by a NOESY experiment (Figure 2) and from the <sup>1</sup>H NMR J values. In the <sup>1</sup>H NMR spectrum, a small coupling constant  $(J = 4.5 \text{ Hz})^{11}$  was observed for H-1", which required an equatorialaxial relationship between the anomeric proton (H-1") and H-2". The large coupling constants (J = 7.5 Hz) between H-2" and H-3", H-3" and H-4", and H-4" and H-5" as well as the NOESY correlations of H-2"/H-1", H-4" and H-3"/H-5" suggested that H-2", H-3", H-4", and H-5" are all axial. Thus, the relative configuration of the α-glucose moiety at the C-1 glycerol ether could be established. Methanolysis of 2 gave a mixture of  $\alpha$ - and  $\beta$ -methyl glucopyranoside. The specific rotation of the methyl glucoside mixture,  $[\alpha]^{24}D + 74.3$  (c 0.2, MeOH), was close to that of an

<sup>\*</sup> Dedicated to Dr. David G. I. Kingston of Virginia Polytechnic Institute and State University for his pioneering work on bioactive natural products.

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Table 1. <sup>1</sup>H and <sup>13</sup>C NMR Spectroscopic Data of Compounds 2 and 4<sup>a</sup>

С/Н	2			4		
	<sup>13</sup> C		<sup>1</sup> H	13C		<sup>1</sup> H
	lipid base unit				unit	
1	$68.8 t^b$	a	$3.88 \text{ dd} (10.5, 4.5)^c$	$69.9 t^b$	a	4.12 dd (10.5, 3.5) <sup>c</sup>
		b	3.83 dd (10.5, 4.5)		b	3.71 dd (10.5, 3.5)
2	54.6 d		3.93 dd (8.0, 4.5)	54.7 d		3.99 dd (7.5, 3.5)
3	72.9 d		4.14 t (8.0)	73.0 d		4.13 t (7.5)
4	131.6 d		5.47 dd (15.5, 8.0)	131.3 d		5.48 dd (15.5, 7.5)
5	134.3 d		5.72 dd (15.5, 6.5)	134.8 d		5.74 dd (15.5, 6.0)
6	33.7 t		2.08 td (7.5, 6.5)	34.0 t		2.06 m
7	29.0 t		2.20 td (7.5, 7.0)	28.8 t		2.07 m
8	130.6 d		5.35 t (7.0)	125.0 d		5.14 t (6.5)
9	135.3 s			136.9 s		` ´
10	136.3 d		6.02 d (15.5)	40.9 t		1.97 t (7.5)
11	128.7 d		5.55 dt (15.5, 7.0)	29.3 t		1.39 m
12	34.1 t		2.08 td (7.5, 7.0)			
13	30.4 t		1.38 m			
12-15				30.5-30.9 t		1.29 m
14, 15	30.4-31.1 t		1.29 m			
16	33.2 t		1.29 m	33.2 t		1.29 m
17	23.9 t		1.29 m	23.9 t		1.29 m
18	14.6 q		0.90 t (6.0)	14.6 q		0.90 t (6.5)
19	12.9 q		1.71 s	16.3 q		1.59 s
17	12.7 q	N-acyl un		10.5 q	N-acyl u	
1'	175.4 s	iv acyi an		177.4 s	11 deyl d	iiit
2'	74.3 d		4.45 d (6.0)	73.2 d		3.99 dd (8.0, 4.0)
3'	129.1 d		5.50 dd (15.5, 6.0)	36.0 t		1.70 m; 1.55 m
4'	135.0 d		5.83 dd (15.5, 6.5)	26.3 t		1.41 m
5'	33.6 t		2.03 td (7.5, 6.5)	20.5 t		1.41 III
5'-19'	33.0 t		2.03 td (7.3, 0.3)	30.5-30.9 t		1.29 m
6'	30.5 t		1.38 m	30.3 30.71		1.27 111
7'-15'	30.4-31.1 t		1.29 m			
16'	33.2 t		1.29 m			
17'	23.9 t		1.29 m			
18'			0.90 t (6.0)			
20'	14.6 q		0.90 t (6.0)	23.9 t		1.29 m
21'						0.90 t (6.5)
				14.6 q		0.90 t (6.3)
22'		α-D-glucose	it		Q p. aluana	· ····it
1"	103.6 d	a-D-glucose		104.0.4	$\beta$ -D-glucose	
2"	79.1 d		4.87 d (4.5)	104.9 d 75.1 d		4.27 d (8.0)
3"			3.96 dd (7.5, 4.5)			3.19 dd (9.5, 8.0)
	76.3 d		4.10 t (7.5)	78.1 d		3.34 m
4"	74.4 d		3.63 m	71.7 d		3.30 m
5"	83.6 d		3.69 m	78.1 d		3.28 m
6"	64.3 t	a	3.63 m	62.8 t	a	3.87 dd (12.0, 1.5)
		b	3.56 m		b	3.67 dd (12.0, 5.0)

<sup>&</sup>lt;sup>a</sup> Spectra were measured in CD<sub>3</sub>OD ( $^{1}$ H, 500 MHz;  $^{13}$ C, 125 MHz). <sup>b</sup> Multiplicities were deduced by HSQC and DEPT experiments. <sup>c</sup> J values (in Hz) are in parentheses.

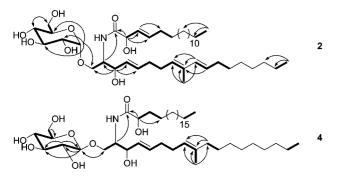


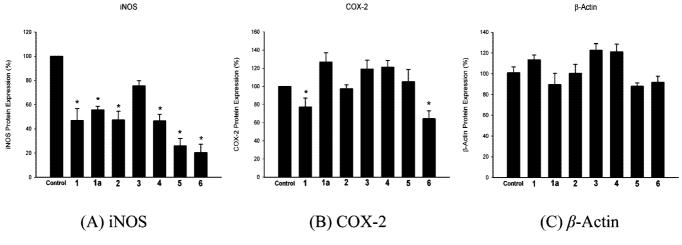
Figure 1. Key  ${}^{1}H-{}^{1}H$  COSY ( $\longrightarrow$ ) and HMBC ( $\longrightarrow$ ) correlations of 2 and 4.

authentic sample ( $[\alpha]^{25}_D$  +77.3). Therefore, the absolute configuration of the  $\alpha$ -glucose moiety was deduced to be the D-isomer.

The large vicinal coupling constants ( $J=15.5~{\rm Hz}$ ) between H-4 and H-5, H-10 and H-11, and H-3' and H-4' clearly indicated E geometry for  $\Delta^{4,10,3'}$  in **2**. The key HMBC correlations from H-3 to C-4, from H-2' to C-3', and from H<sub>3</sub>-19 to C-8, C-9, C-10, and C-11 helped locate the positions of the double bonds (Figure 1). The UV absorption maximum at 224 nm indicated the presence of

Figure 2. Selected NOESY correlations of 2 and 4.

a conjugated diene moiety. <sup>1f</sup> Furthermore, the stereochemistry of the conjugated diene moiety was assigned as *s-trans* from NOE correlations (Figure 2) from Me-19 to H-7 and H-11. The <sup>1</sup>H-<sup>1</sup>H COSY correlations observed between H-2 and both H-1 and H-3 were used to define the 2-amino-1,3-dioxygenated fragment. The relative configurations of C-2 (δ 54.6) and C-3 (δ 72.9) were predicted to be the D-*erythro* stereochemistry at C-2 and C-3, which



**Figure 3.** Effect of compounds **1–6** at 10  $\mu$ M on the LPS-induced pro-inflammatory iNOS and on COX-2 protein expression of RAW 264.7 macrophage cells by immunoblot analysis. (A) Immunoblot of iNOS. (B) Immunoblot of COX-2. (C) Immunoblot of  $\beta$ -actin. A and B values are mean  $\pm$  SEM (n = 5). The relative intensity of the LPS alone stimulated group was taken as 100%. \*Significantly different from LPS-stimulated (control) group (\*P < 0.05).

was consistent with those reported for other (2S,3R,2'R) sphingosine moieties.  $^{12,13}$  The stereochemistry of  $\mathbf{2}$  (2S,3R), as shown in its structure, was confirmed after determination of the stereochemistry of  $\mathbf{1}$  on the basis of biosynthetic reasoning. Methanolysis of  $\mathbf{2}$  yielded methyl (2'R,3'E)-2'-hydroxyoctadec-3'-enoate  $(\mathbf{2a})$ , for which the molecular formula  $C_{19}H_{36}O_3$  was determined by ESIMS-MS  $(m/z\ 335.0\ [M\ +\ Na]^+)$  as  $C_{19}H_{36}O_3$ . On the basis of the aforementioned observations, sarcoehrenoside A  $(\mathbf{2})$  was characterized unambiguously as 1-O- $(\alpha$ -D-glucopyranosyl)-(2S,3R,4E,8E,10E)-2-[(2'R,3'E)-2'-hydroxyoctadec-3'-enoylamino]-9-methyloctadeca-4,8,10-triene-1,3-diol.

Sarcoehrenoside B (4) was found to have a molecular formula of C<sub>46</sub>H<sub>87</sub>NO<sub>9</sub>, as indicated by HRESIMS. In addition to the pseudomolecular ion peak at m/z 820.6 [M + Na]<sup>+</sup>, the ESIMS-MS of 4 also exhibited an intense fragment peak at m/z 496.7, produced by elimination of the *N*-acyl unit from the molecular ion. The spectroscopic data for 4 (Table 1) exhibited the presence of an amide linkage, a long chain, and a sugar, consistent with the C-9 methyl cerebroside nature of 4. In the <sup>1</sup>H and <sup>13</sup>C NMR spectra of 4, an anomeric proton appeared at  $\delta$  4.27 (1H, d, J = 8.0 Hz, H-1") and  $^{13}$ C NMR signals resonated at  $\delta$  104.9 (H-1"), 75.1 (H-2"), 78.1 (H-3"), 71.7 (H-4"), 78.1 (H-5"), and 62.8 (H-6"), supporting the presence of a  $\beta$ -glucopyranose moiety. <sup>3b</sup> Methanolysis of 4 also yielded an  $\alpha$ - and  $\beta$ -glucopyranosyl mixture, the specific rotation ( $[\alpha]^{24}_D$  +71.4, MeOH) of which was used to identify  $\beta$ -glucose as a D-isomer. The glycerol ether linkage between the anomeric proton ( $\delta$  4.27, H-1") and C-1 was confirmed by HMBC correlations (Figure 1) of H-1" with C-1 and H-1 with C-1". The coupling constant (15.5 Hz) indicated an E geometry for  $\Delta^4$ . A NOE correlation from Me-19 to H-7 supported the presence of an (E)-8,9 double bond. Therefore, **3** was defined as a C-9 methyl 4E,8E-sphingadiene-type cerebroside. The connectivity between the acyl moiety and the NH of the sphingosine base was confirmed by HMBC correlations of H-2 with C-1' and H-2' with C-1'. Methanolysis of 4 yielded methyl (2'R)-2'-hydroxyhenicosanoate (4a), for which the molecular formula, C<sub>22</sub>H<sub>44</sub>O<sub>3</sub>, was identified by ESIMS-MS analysis (m/z 379.3 [M + Na]<sup>+</sup>). The specific rotation value ( $[\alpha]^{24}$ <sub>D</sub> -3.8) of **4a** indicated a 2'R absolute configuration in **4.** If The relative configuration at C-2 and C-3 was determined as 2S,3R (erythro) based on their <sup>13</sup>C NMR chemical shifts and the specific rotation of this compound. 1f Consequently, sarcoehrenoside B (4) was fully assigned as 1-O-(β-D-glucopyranosyl)-(2S,3R,4E,8E)-2-[(2'R)-2'-hydroxyhenicosanoylamino]-9-methyl-4,8-octadecadiene-1,3-diol.

Compounds 1-6 were tested against five bacterial strains, comprising Enterobacter aerogenes (ATCC13048), Serratia marce-

scens (ATCC25419), Salmonella enteritidis (ATCC13076), Yersinia enterocolitica (ATCC23715), and Shigella sonnei (ATCC11060). None of these compounds exhibited any antibacterial activity at a concentration of 100 µg/disk.

A previous study has reported that glycosphingolipids and sphingolipids possess cyclooxygenase-2 inhibition, 14 which prompted us to evaluate the anti-inflammatory effect of our isolated metabolites. As shown in Figure 3, the in vitro anti-inflammatory activity of compounds 1-6 was tested using LPS-stimulated cells. Stimulation of RAW 264.7 cells with LPS resulted in up-regulation of the pro-inflammatory iNOS and COX-2 proteins. Both compounds 1 and 6 reduced the levels of iNOS to  $46.9 \pm 9.7\%$  and  $20.3 \pm 6.8\%$ , respectively, and of COX-2 to 77.2  $\pm$  9.9% and 64.3  $\pm$  8.6%, respectively, in comparison with those of the control groups. Compounds 1a, 2, 4, and 5 reduced iNOS protein expression (55.6  $\pm$  2.9%, 47.3  $\pm$  7.1%, 46.5  $\pm$  5.3%, and 25.8  $\pm$  6.2%, respectively), but did not inhibit COX-2 protein expression. All compounds did not affect  $\beta$ -actin protein expression at a 10  $\mu$ M concentration. Under the same experimental conditions, 10  $\mu$ M CAPE (caffeic acid phenylthyl ester) reduced the levels of iNOS and COX-2 protein to  $1.5 \pm 2.1\%$  and  $70.2 \pm 11.5\%$ , respectively, relative to the control cells stimulated with LPS.

#### **Experimental Section**

General Experimental Procedures. Optical rotations were determined on a JASCO P1020 polarimeter. UV spectra were obtained on a Hitachi U-3210 spectrophotometer, and IR spectra were recorded on a JASCO FT/IR-4100 spectrophotometer. The NMR spectra were recorded on a Bruker Avance 300 NMR spectrometer at 300 MHz for <sup>1</sup>H and 75 MHz for <sup>13</sup>C, or on a Varian MR 400 NMR spectrometer at 400 MHz for <sup>1</sup>H and 100 MHz for <sup>13</sup>C, or on a Varian Unity INOVA 500 FT-NMR spectrometer at 500 MHz for <sup>1</sup>H and 125 MHz for <sup>13</sup>C. Chemical shifts are expressed in  $\delta$  (ppm) referring to the solvent peaks  $\delta_{\rm H}$  3.30 and  $\delta_{\rm C}$  49.0 for CD<sub>3</sub>OD and  $\delta_{\rm H}$  7.265 and  $\delta_{\rm C}$  77.0 for CDCl<sub>3</sub>, and coupling constants are expressed in Hz. Electrospray ionization MS/MS analysis was recorded using a positive-mode API 4000 tandem mass spectrometer, with a capillary voltage of 5500 V applied, and the spectra were acquired at m/z 400-1000 for parent ion spectra and m/z 50-1000 for fragment ion spectra. ESIMS were recorded by ESI FT-MS on a Bruker APEX II mass spectrometer. Silica gel 60 (Merck, 230-400 mesh) was used for column chromatography; precoated Si gel plates (Merck, Kieselgel 60 F254, 0.25 mm) were used for TLC analysis. C<sub>18</sub> reversed-phase silica gel (230-400 mesh, Merck) was also used for column chromatography. High-performance liquid chromatography (HPLC) was performed on a Hitachi L-7420 UV detector L-7100 pump apparatus equipped with a Merck Hibar RP-18e column  $(250 \times 10 \text{ mm}, 5 \mu\text{m}).$ 

Animal Material. The octocoral *S. ehrenbergi* was collected by hand using scuba at Dongsha Islands, Taiwan, in April 2007, at a depth of 10 m, and was stored in a freezer for 5 weeks until extraction. This soft coral was identified by one of the authors (C.-F.D.). A voucher specimen (TS-07) was deposited in the Department of Marine Biotechnology and Resources, National Sun Yat-sen University.

Extraction and Isolation. Soft coral extracts were prepared by percolating aliquots of 50 g of freeze-dried S. ehrenbergi with acetone for 24 h at room temperature. The combined acetone extracts were concentrated to a brown gum, which was partitioned between H2O and EtOAc. The dried EtOAc-soluble (20.0 g) partition was chromatographed over a silica column using n-hexane, n-hexane-EtOAc, and EtOAc-MeOH mixtures of increasing polarity to obtain fractions 1-40. Fraction 28 (0.5 g), eluted with EtOAc-MeOH (1:1), was further subjected to RP-18 gravity column chromatography by eluting with 80% MeOH in  $H_2O,\,\bar{9}0\%$  MeOH in  $H_2O,\,\text{and}\,\,100\%$  MeOH. Altogether, six fractions were obtained, of which fraction 1 (60 mg) was purified further by RP-18 HPLC column chromatography (5% CH<sub>3</sub>CN in MeOH, flow rate 5.0 mL/min) to afford 1 (3 mg), 2 (2 mg), 3 (2 mg), 4 (3 mg), 5 (2 mg), and 6 (6 mg). The retention time for each metabolite was as follows: 1 (26.5 min), 2 (35.6 min), 3 (48.6 min), 4 (44.2 min), **5** (41.0 min), and **6** (33.5 min).

**Sarcoehrenoside A (2):** white, amorphous powder;  $[\alpha]^{23}_{D}$  +77.0 (*c* 0.2, MeOH); IR (KBr) 3387, 2951, 2859, 1638, 1547, 1456, 1387, 1241, 1131, 1035, 737 cm<sup>-1</sup>; UV  $\lambda_{max}$  (MeOH) (log ε) 224 (3.97) nm; <sup>1</sup>H NMR and <sup>13</sup>C NMR data, see Table 1; HRESIMS *m/z* 774.5490 [M + Na]<sup>+</sup> (calcd for C<sub>43</sub>H<sub>77</sub>NO<sub>9</sub>Na, 744.5496).

**Sarcoehrenoside B (4):** white, amorphous powder;  $[\alpha]^{23}_D$  +51.3 (*c* 0.3, MeOH); IR (KBr) 3396, 2951, 2857, 1638, 1546, 1457, 1385, 1242, 1178, 1131, 1036, 736 cm<sup>-1</sup>; <sup>1</sup>H NMR and <sup>13</sup>C NMR data, see Table 1; HRESIMS *m/z* 820.6274 [M + Na]<sup>+</sup> (calcd for C<sub>46</sub>H<sub>87</sub>NO<sub>9</sub>Na, 820.6278).

Methanolysis of 2 and 4. An aliquot (1 mg) of 2 and 4 in a mixture of 1 N aqueous HCl (0.1 mL) and methanol (0.9 mL) was refluxed for 18 h on a magnetic stirrer. The reaction mixture was neutralized with NaHCO<sub>3</sub> and diluted with H<sub>2</sub>O (1.5 mL). The aqueous solution was extracted with *n*-hexane three times, and the organic phase was dried with anhydrous MgSO<sub>4</sub>. After removal of solvent and purification by silica gel column chromatography, methyl (2'R,3'E)-2'-hydroxyoctadec-3'-enoate (2a, 0.3 mg) and methyl (2'R)-2'-hydroxyhenicosanoate (4a, 0.4 mg) were obtained and identified by ESIMS-MS peaks at m/z 335.0 and 379.3 [M + Na]<sup>+</sup>, respectively. In addition, the aqueous layer was removed and purified on a C<sub>18</sub> reversed-phase column, eluted with MeOH-H<sub>2</sub>O (4:1), to give 2b [0.2 mg,  $[\alpha]^{24}_D$  +74.3 (c 0.2, MeOH)] and 4b [0.2 mg,  $[\alpha]^{24}_D$  + 72.5 (c 0.2, MeOH)].

Methyl (2'*R*,3'*E*)-2'-hydroxyoctadec-3'-enoate (2a): white, amorphous powder;  $[α]^{24}_D$  -17.5 (*c* 0.3, CHCl<sub>3</sub>); <sup>1</sup>H NMR (CDCl<sub>3</sub>, 400 MHz) δ 5.90 (1H, dt, J = 15.6, 6.4 Hz, H-4'), 5.50 (1H, dd, J = 15.6, 6.4 Hz, H-3'), 4.61 (1H, d, J = 6.4 Hz, H-2'), 3.80 (3H, s, COOCH<sub>3</sub>), 2.07 (2H, td, J = 7.6, 6.4 Hz, H-5'), 1.39 (2H, m, H-6'), 1.26 (22H, brs, H-7'-H-17'), 0.88 (3H, t, J = 6.4 Hz, H-18'); ESIMS-MS m/z 335.0 [M + Na]<sup>+</sup>.

Methyl (2'*R*)-2'-hydroxyhenicosanoate (4a): white, amorphous powder;  $[α]^{24}_D$  –3.8 (*c* 0.4, CHCl<sub>3</sub>); <sup>1</sup>H NMR (CDCl<sub>3</sub>, 400 MHz) δ 4.20 (1H, dd, J = 7.6, 4.4 Hz, H-2'), 3.79 (3H, s, COOCH<sub>3</sub>), 1.76 (1H, m, H-3a'), 1.58 (1H, m, H-3b'), 1.45 (2H, m, H-4'), 1.26 (32H, brs, H-5'-H-20'), 0.88 (3H, t, J = 6.4 Hz, H-21'); ESIMS-MS m/z 379.3 [M + Na]<sup>+</sup>.

In Vitro Antimicrobial Activity. Bacterial strains were grown in LB (Luria-Bertani) broth medium for 24 h at 37 °C. Then, 17 mL of LB hard agar (1.5% agar) was poured into sterile Petri dishes (9 cm) and allowed to set. Next, 2.7 mL of molten LB soft agar (0.7% agar, 45 °C) was inoculated with 0.3 mL of broth culture of the test organism and poured over the base hard agar plates, forming a homogeneous top layer. Sterile paper disks (Advantec, 8 mm) were placed onto the top layer of the LB agar plates. Ten microliters (2  $\mu$ g/ $\mu$ L) of the tested compounds was applied onto each filter paper disk. Ampicillin (5 µg/  $\mu$ L) and the same solvents served as positive and negative controls. All plates were incubated at 37 °C, 24 h prior to antibacterial activity evaluation. The antimicrobial activity of compounds 1-6 was tested up to 100 µg/mL against E. aerogenes (ATCC13048), S. marcescens (ATCC25419), S. enteritidis (ATCC13076), Y. enterocolitica (ATCC23715), and S. sonnei (ATCC11060). All bacterial strains were obtained from the American Type Culture Collection. The antibiotic activity evaluation method was conducted on the basis of previous reports. 15

In Vitro Anti-inflammatory Assay. The anti-inflammatory assay was modified from Ho et al. 16 and Park et al. 17 Murine RAW 264.7 macrophages were obtained from the American Type Culture Collection (ATCC, No. TIB-71). The cells were activated by incubation in medium containing Escherichia coli LPS (0.01  $\mu$ g/mL; Sigma) for 16 h in the presence or absence of various compounds. Then, cells were washed with ice-cold PBS, lysed in ice cold lysis buffer, and centrifuged at 20000g for 30 min at 4 °C. The supernatant was decanted from the pellet and retained for Western blot analysis. Protein concentrations were determined by the DC protein assay kit (Bio-Rad) modified by the method of Lowry et al. 18 Samples containing equal quantities of protein were subjected to SDS-polyacrylamide gel electrophoresis, and the separated proteins were electrophoretically transferred to polyvinylidene difluoride membranes (PVDF; Immobilon-P, Millipore, 0.45 μm pore size). The resultant PVDF membranes were incubated with blocking solution and incubated for 180 min with antibody against inducible nitric oxide synthase (iNOS; 1:1000 dilution; Transduction Laboratories) and cyclooxygenase-2 (COX-2; 1:1000 dilution; Cayman Chemical) protein. The blots were detected using ECL detection reagents (Perkin-Elmer, Western Blot Chemiluminescence Reagent Plus) according to the manufacturer's instructions.

**Acknowledgment.** Financial support was provided by Ministry of Education (96C031703) and National Science Council of Taiwan (NSC96-2320-B-110-003-MY3) awarded to C.Y.D.

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