



Cytotoxic and anti-inflammatory cembranoids from the Dongsha Atoll soft coral *Sarcophyton crassocaule*

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ABSTRACT

Five new cembranoids, sarcocrassocolides A–E (**1–5**), along with three known cembranoids **6–8**, have been isolated from a Formosan soft coral *Sarcophyton crassocaule*. The structures of the new metabolites were elucidated by extensive spectroscopic analysis and the absolute configuration of **1** was determined by a modified Mosher's method. Compounds **1–4** exhibited significantly cytotoxic activity against a limited panel of cancer cell lines. Compounds **1–4**, **6** and **8** were shown to exert significant in vitro anti-inflammatory activity in LPS-stimulated RAW264.7 macrophage cells. Compound **6** also significantly inhibited the accumulation of pro-inflammatory COX-2 protein.

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1. Introduction

Many cembrane-type compounds have been shown to exhibit cytotoxicity^{1–6} and anti-inflammatory activity.^{7–11} During the course of our search for bioactive metabolites from marine invertebrates, many bioactive cembrane-derived compounds have been discovered from soft corals of the genera *Sinularia*,^{1–5,7–9} *Lobophytum*,^{10,11} and *Sarcophyton*.^{12,13} Our investigation on the chemical constituents of the Dongsha Atoll soft coral *Sarcophyton crassocaule* yielded five new cembranoids, sarcocrassocolides A–E (**1–5**), along with three known cembranoids, sarcocrassolide (**6**),¹⁴ sinularolide (**7**)¹⁵ and 13-acetoxysarcocrassolide (**8**).¹⁴ The structures of **1–5** were established by detailed spectroscopic analysis, including extensive examination of 2D NMR (¹H–¹H COSY, HMQC, HMBC and NOESY) correlations. The absolute configuration of **1** was further determined using a modified Mosher's method. The cytotoxicity of compounds **1–4** against human breast carcinoma (MCF-7), human colon carcinoma (WiDr), human laryngeal carcinoma (HEp-2) and human medulloblastoma (Daoy) cells was studied, and the ability of **1–4** and **6–8** to inhibit the up-regulation of pro-inflammatory iNOS (inducible nitric oxide synthase) and

COX-2 (cyclooxygenase-2) proteins in LPS (lipopolysaccharide)-stimulated RAW264.7 macrophage cells was also evaluated. The results revealed that compounds **1–4** are all cytotoxic towards the above cancer cells, **3** being the most cytotoxic. Compounds **1–4**, **6** and **8** were found to effectively inhibit the expression of iNOS protein, while **6** was the only one seen also to significantly reduce the level of COX-2 protein in the above assay.

2. Results and discussion

The HRESIMS (m/z 413.1938 [M+Na]⁺) of **1** established the molecular formula C₂₂H₃₀O₆, appropriate for eight degrees of unsaturation, and the IR spectrum revealed the presence of hydroxy (3487 cm^{−1}) and lactonic carbonyl (1769 cm^{−1}) groups. The ¹³C NMR (Table 1) and DEPT spectroscopic data showed signals of three methyls (including one acetate methyl), five sp³ methylenes, two sp² methylenes, five sp³ methines (including four oxymethines), one sp² methine, one sp³ and five sp² quaternary carbons (including two ester carbonyls). The NMR signals (Tables 1 and 2) observed at δ_C 170.1 (qC), 140.8 (qC), 120.5 (CH₂), 81.0 (CH), and 36.3 (CH), and δ_H 6.20, 5.65 (each, 1H, d, J = 2.0 Hz), 4.50 (1H, t, J = 2.8 Hz), and 3.64 (1H, dd, J = 6.4, 3.2 Hz) revealed the presence of an α -methylene- γ -lactonic group by comparing the very similar NMR data of the five-membered lactone ring in

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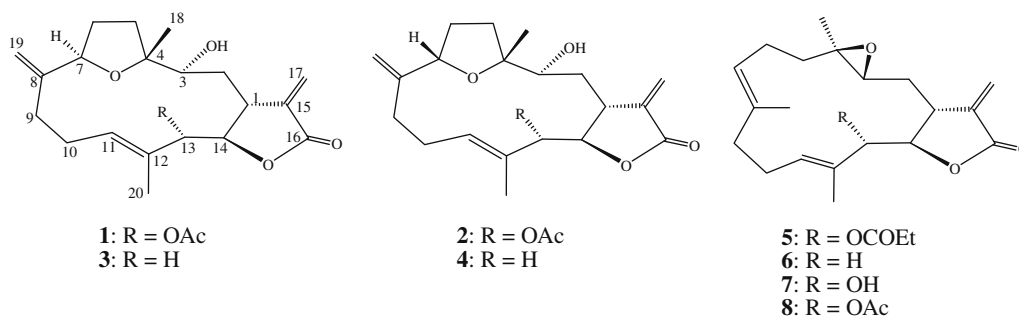


Table 1
¹³C NMR data for compounds 1–5

C#	1 ^a	2 ^b	3 ^a	4 ^a	5 ^b
1	36.3, CH ^c	35.4, CH	41.8, CH	41.4, CH	36.9, CH
2	39.4, CH ₂	41.4, CH ₂	38.2, CH ₂	40.9, CH ₂	34.7, CH ₂
3	73.5, CH	74.9, CH	73.3, CH	74.9, CH	60.1, CH
4	84.4, qC	86.0, qC	84.3, qC	85.7, qC	60.5, qC
5	37.9, CH ₂	37.8, CH ₂	38.4, CH ₂	38.4, CH ₂	37.4, CH ₂
6	30.9, CH ₂	28.2, CH ₂	31.9, CH	29.1, CH ₂	23.4, CH ₂
7	79.4, CH	82.6, CH	79.6, CH	82.4, CH	123.0, CH
8	149.8, qC	148.3, qC	149.1, qC	147.2, qC	135.1, qC
9	33.4, CH ₂	28.0, CH ₂	34.1, CH ₂	29.3, CH ₂	37.9, CH ₂
10	26.9, CH ₂	26.9, CH ₂	28.5, CH ₂	28.1, CH ₂	24.8, CH ₂
11	129.1, CH	126.4, CH	129.8, CH	127.8, CH	129.0, CH
12	127.4, qC	128.5, qC	127.6, qC	127.7, qC	128.7, qC
13	77.1, CH	77.0, CH	46.0, CH	46.2, CH ₂	77.2, CH
14	81.0, CH	81.8, CH	80.7, CH	81.4, CH	80.9, CH
15	140.8, qC	140.9, qC	139.1, qC	138.9, qC	139.7, qC
16	170.1, qC	170.1, qC	168.9, qC	168.6, qC	169.4, qC
17	120.5, CH ₂	120.9, CH ₂	121.6, CH ₂	122.0, CH ₂	121.0, CH ₂
18	21.4, CH ₃	18.9, CH ₃	22.3, CH ₃	20.4, CH ₃	17.4, CH ₃
19	111.0, CH ₂	112.3, CH ₂	110.2, CH ₂	111.3, CH ₂	16.8, CH ₃
20	14.5, CH ₃	14.8, CH ₃	17.7, CH ₃	17.9, CH ₃	14.7, CH ₃
13-Acetate	20.7, CH ₃	20.7, CH ₃			
13-Propionate	169.4, qC	169.3, qC			8.9, CH ₃ 27.5, CH ₂ 172.5, qC

^a Spectra recorded at 100 MHz in CDCl₃.

^b Spectra recorded at 125 MHz in CDCl₃.

^c Attached protons were deduced by DEPT experiments.

the known compounds **6** and **8**.¹⁴ Signals appearing at δ_c 111.0 (CH₂), 149.8 (qC), δ_H 4.90 (1H, s) and 4.92 (1H, s) revealed the presence of the other 1,1-disubstituted carbon–carbon double bond. One trisubstituted double bond was also identified from the NMR signals at δ_c 127.4 (qC) and 129.1 (CH) and at δ_H 5.48 (1H, dd, J = 10.0, 5.6 Hz). In the ¹H–¹H COSY spectrum, it was possible to identify three different structural units, which were assembled with the assistance of an HMBC experiment (Fig. 1). Key HMBC correlations of H₃–18 to C-3, C-4 and C-5; H₂–19 to C-7, C-8 and C-9; H₃–20 to C-11, C-12 and C-13; H₂–17 to C-1, C-15 and C-16; and H₂–5 and H₂–6 to C-4 permitted connection of the carbon skeleton. Furthermore, the acetoxy group positioned at C-13 was confirmed from the HMBC correlations of H-13 (δ 5.36) and protons of an acetate methyl (δ 2.01) to the ester carbonyl carbon at δ 169.4 (C). A hydroxy group was positioned at C-3, as both of the (S)- and (R)-MTPA esters of **1**, prepared for the determination of absolute configuration (later discussed), showed significant differences in the chemical shifts of H-3 and C-3 in comparison with those of **1**. In considering the degrees of unsaturation and molecular formula, an ether linkage has to be placed between C-4 and C-7. On the basis of the above analysis, the planar structure of **1** was established unambiguously.

The relative configuration of **1** was elucidated by the analysis of NOE correlations, as shown in Figure 2. It was found that H-1 (δ 3.64, dd, J = 6.4, 3.0 Hz) showed NOE interaction with H₃–18 (δ 1.21, s); therefore, assuming a β -orientation of H-1, H₃–18 should also be positioned on the β -face. One of the methylene protons at C-2 (δ 1.94, m) exhibited NOE correlations with both H-1 and H₃–18 and was characterized as H-2 β , while the other proton was assigned as H-2 α (δ 1.75, m). NOE correlations observed between H-2 α and H-14, and H-14 and H-3, reflected the α -orientation of H-14, and hence the S^* configuration of C-3. Furthermore, H-14 exhibited NOE correlation with H-13 and H-3 exhibited NOE interaction with H-7, revealing the α -orientations of H-7 and H-13, and hence the S^* configuration of C-13. The *E* geometry of the trisubstituted C-11/C-12 double bond was also assigned from the NOE correlation of H₃–20 (δ 1.71, s) with H-10 (δ 2.15, m), but not with olefinic proton H-11, and also the upper field chemical shift of C-20 (δ 14.5). On the basis of the above findings and detailed examination of other NOE correlations (Fig. 2), the relative configuration of compound **1** was determined. Furthermore, the absolute configuration of **1** was finally determined using a modified Mosher's esterification method.¹⁶ The (S)- and (R)-MTPA esters of **1** (**1a** and **1b**, respectively) were prepared using the corresponding (R)-

Table 2
¹H NMR data for compounds **1**–**5**

H#	1 ^a	2 ^b	3 ^a	4 ^a	5 ^b
1	3.64, dd (6.4, 3.2)	3.66, br d (10.0)	3.43, ddd (5.6, 3.2, 1.6)	3.40, d (8.4)	3.06, d (12.0)
2	1.94, m	1.88, m	1.94, m	1.83, m	1.83, ddd (15.0, 12.5, 3.5)
	1.75, m	1.69, m	1.84, m	1.75, m	1.69, ddd (15.0, 7.0, 3.5)
3	3.49, dd (9.2, 3.2) ^c	3.37, dd (9.5, 3.0)	3.53, dd (8.0, 4.0)	3.43, t (4.0)	2.64, dd (7.5, 3.5) ^c
5	1.98, m; 1.90, m	1.90, m; 1.75, m	1.89, m; 1.78, m	1.86, m; 1.77, m	1.34, m
6	2.05, m; 1.78, m	2.15, m; 1.76, m	2.00, m; 1.77, m	2.14, m; 1.79, m	2.14, m; 2.09, m
7	4.17, dd (10.4, 4.4)	4.52, dd (9.5, 4.5)	4.14, dd (11.2, 4.0)	4.49, dd (8.8, 4.0)	5.02, t (6.5)
9	2.37, m	2.33, m	2.29, m	2.32, d (14.0)	2.22, dd (14.0, 6.5)
		1.82, m		1.83, m	2.04, m
10	2.28, m	2.45, dddd (14.0, 10.5, 7.0, 3.5)	2.25, m	2.40, m	2.39, ddd (15.0, 9.5, 9.5)
	2.15, m	2.14, m	2.13, m	2.12, m	2.15, m
11	5.48, dd (10.0, 5.6)	5.49, t (7.5)	5.29, dd (10.4, 5.2)	5.34, t (7.2)	5.30, br t (7.0)
13	5.36, s	5.27, s	2.55, d (14.0)	2.48, d (12.8)	5.39, s
			2.12, m	2.02, t (12.4)	
14	4.50, t (2.8)	4.39, br s	4.40, dt (10.4, 3.2)	4.33, d (11.6)	4.60, t (2.5)
17	6.20, d (2.0)	6.21, d (2.0)	6.25, d (2.0)	6.25, s	6.25, d (2.0)
	5.65, d (2.0)	5.67, d (2.0)	5.69, d (2.0)	5.70, s	5.62, d (1.5)
18	1.21, s	1.20, s	1.22, s	1.20, s	1.29, s
19	4.92, s; 4.90, s	5.10, s; 4.99, s	4.93, s; 4.89, s	5.08, s; 5.00, s	1.63, s
20	1.71, s	1.73, s	1.67, s	1.69, s	1.72, s
13-Acetate	2.01, s	1.97, s			
13-Propionate					2.26, q (7.5)
					1.09, t (7.5)

^a Spectra recorded at 400 MHz in CDCl₃.

^b Spectra recorded at 500 MHz in CDCl₃.

^c *J* values (in Hz) in parentheses.

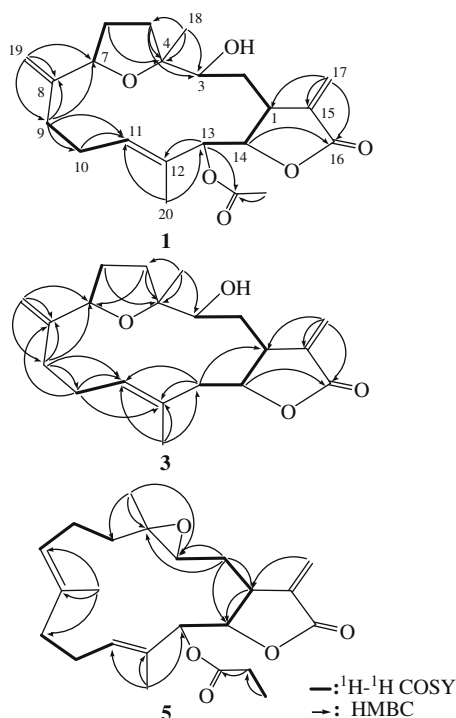


Figure 1. ¹H–¹H COSY and HMBC correlations for **1**, **3** and **5**.

(–) and (S)-(+)-MTPA chloride, respectively. The determination of $\Delta\delta$ values ($\delta_S - \delta_R$) for protons neighboring C-3 led to the assignment of the *S* configuration at C-3 in **1** (Fig. 3).

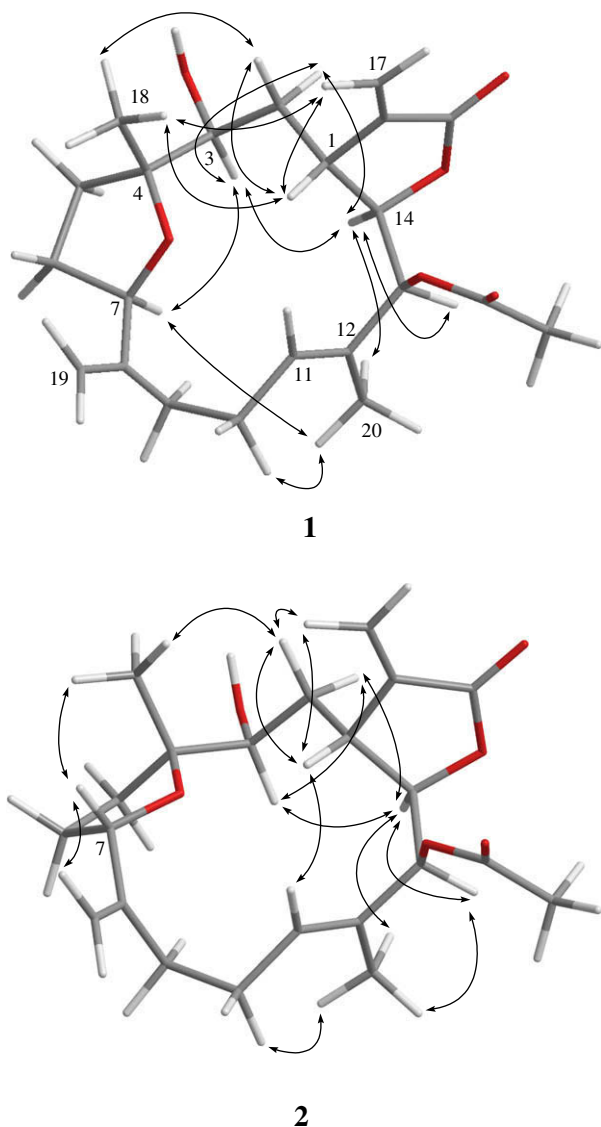
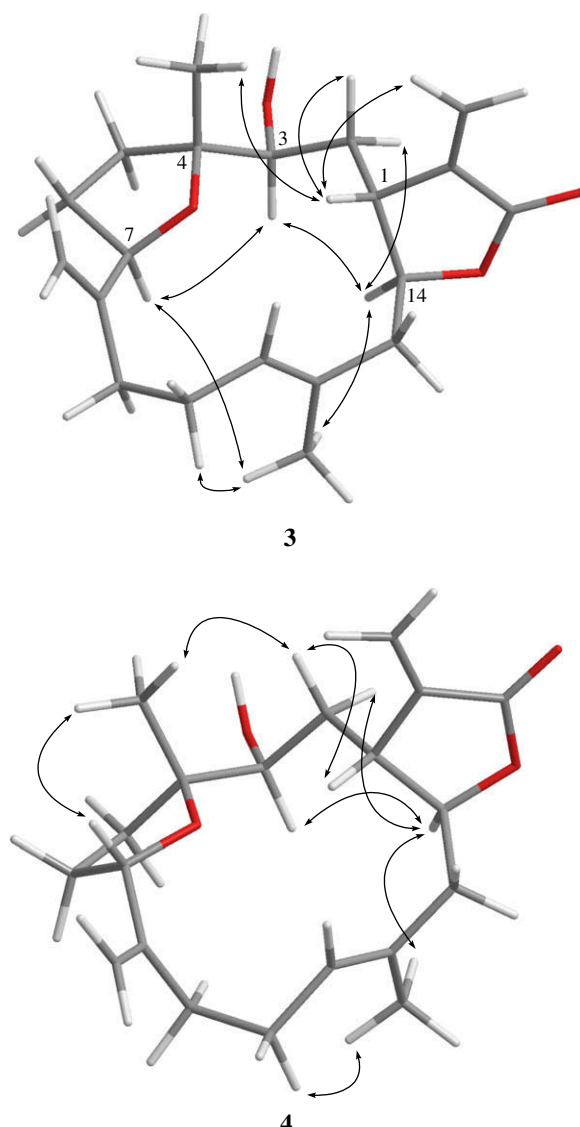
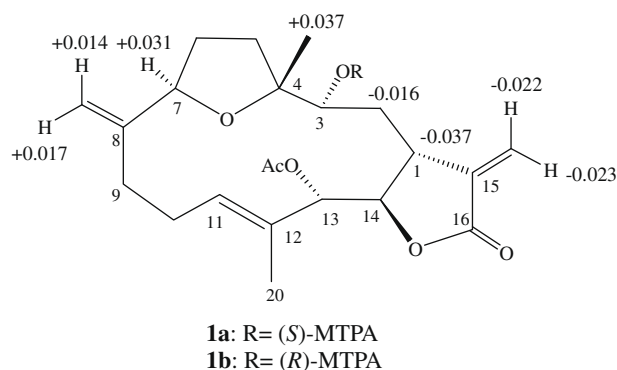
Compound **2** possessed the same molecular formula (C₂₂H₃₀O₆) as that of **1**, as revealed from HRESIMS. Furthermore, it was found that the NMR spectroscopic data of **2** were very similar to those of **1**. By analysis of the 2D NMR (¹H–¹H COSY, HMQC, and HMBC) correlations, compound **2** was shown to possess the same planar structure as that of **1**. From the NOESY spectrum (Fig. 2), it was found that H-7 (δ 4.52, dd, *J* = 9.5, 4.5 Hz) showed an NOE interac-

tion with H₃-18 (δ 1.20, s), but not with H-3 and H₃-20, revealing the β -orientation of H-7. Further analysis of other NOE interactions revealed that **2** possessed the same relative configurations at C-1, C-3, C-4, C-13 and C-14 as those of **1** (Fig. 2). Therefore, **2** was found to be the C-7 epimer of **1**.

Compound **3** was shown by HRESIMS to possess the molecular formula C₂₀H₂₈O₄ (*m/z* 355.1883 [M+Na]⁺). The IR spectrum of **3** revealed the presence of hydroxy (3472 cm^{−1}) and lactonic carbonyl (1760 cm^{−1}) groups. Comparison of the ¹H and ¹³C NMR data (Tables 1 and 2) of compounds **1** and **3** showed that the structure of **3** should be very close to that of **1**, with the exception of signals assigned to C-13, where an acetoxymethine (δ_H 5.36, 1H, s; δ_C 77.1) in **1** is replaced by a methylene (δ_H 2.55, 1H, d, *J* = 14.0 Hz, δ_H 2.12, 1H, m; δ_C 26.6) in **3**. The overall planar structure of **3** was fully established by analyzing the ¹H–¹H COSY and HMBC correlations (Fig. 1). The relative stereochemistry of **3** was confirmed to be 1*R**, 3*S**, 4*R**, 7*R** and 14*S** from the following NOESY correlations (Fig. 4): both H-7 (δ 4.14) and H-14 (δ 4.40) with H-3 (δ 3.53), and H₃-18 (δ 1.22) with H-1 (δ 3.43). These results, together with other detailed NOE correlations of **3** (Fig. 4), unambiguously established the structure of sarcocrassocolide C, as shown in formula **3**. Thus, **3** is the 13-deacetoxy derivative of **1**.

Compound **4** was found to have the molecular formula C₂₂H₃₀O₆, as indicated by HRESIMS, suggesting **4** to be an isomer of **3**. By 2D NMR spectroscopic data, including ¹H–¹H COSY, HMQC and HMBC, compound **4** was shown to possess the same planar structure as that of **3**. By comparison of the related ¹H and ¹³C NMR spectroscopic data of **4** (Tables 1 and 2) with those of **3**, **4** was found to be the C-7 epimer of **3**; this was further confirmed by analysis of the NOE correlations (Fig. 4) of **4**, which revealed that H-7 was correlated with H₃-18.

The HRESIMS spectrum of **5** exhibited a molecular ion peak at *m/z* 411.2150 [M+Na]⁺, consistent with the molecular formula C₂₃H₃₂O₅ and implying eight degrees of unsaturation. Comparison of the ¹H and ¹³C NMR data of **5** (Tables 1 and 2) with those of **8** showed that the structure of both compounds are very similar. The ¹H and ¹³C NMR data (Tables 1 and 2) revealed that **5** is the 13-O-deacetyl-13-O-propionyl derivative of **8**.

Figure 2. Key NOESY correlations for **1** and **2**.Figure 4. Key NOESY correlations for **3** and **4**.Figure 3. ^1H NMR chemical shift differences $\Delta\delta$ ($\delta_S - \delta_R$) in ppm for the MTPA esters of **1**.

It is noteworthy to mention that metabolites **1–4** are cembranoids possessing a tetrahydrofuran moiety with a rarely found 4,7-ether linkage, which has been discovered previously only in the soft coral *Eunicea mammosa*.¹⁷ As the known compounds **6–8** have been found to exhibit cytotoxicity toward several cancer cell lines,^{14,15} the cytotoxicity of compounds **1–4** against the prolifera-

tion of a limited panel of cancer cell lines, including MCF-7, WiDr, HEp-2 and Daoy carcinoma cells was then evaluated. The results showed that compound **3**, the most potent of compounds **1–4**, exhibited cytotoxicity against MCF-7, WiDr, HEp-2 and Daoy cancer cell lines with ED_{50} s of 2.0, 1.2, 2.6 and 3.2 $\mu\text{g/mL}$, respectively. Furthermore, metabolites **1**, **2** and **4** also were found to exhibit significant cytotoxicity towards some of the above four cancer cells (Table 3). The in vitro anti-inflammatory effects of compounds **1–4** and **6–8** were also tested. The inhibition of LPS-induced up-regulation of pro-inflammatory proteins, iNOS and COX-2 in macrophage cells was assayed by immunoblot analysis. At a concentration of 10 μM , compounds **1–4**, **6** and **8** were found to

Table 3
 Cytotoxicity (ED_{50} $\mu\text{g/mL}$) of compounds **1–4**

Compound	MCF-7	WiDr	HEp 2	Daoy
1	4.2	4.2	6.2	8.8
2	3.2	3.2	4.5	5.6
3	2.0	1.2	2.6	3.2
4	4.1	1.8	4.0	5.4
Mitomycin C	0.14	0.15	0.07	0.14

significantly reduce the levels of iNOS protein to $13.7 \pm 5.2\%$, $13.3 \pm 5.0\%$, $4.6 \pm 1.3\%$, $7.0 \pm 3.1\%$, $1.1 \pm 0.9\%$ and $6.2 \pm 0.5\%$, respectively, relative to control cells stimulated with LPS only. Furthermore, at the same concentration, metabolite **6** could strongly reduce COX-2 expression ($3.9 \pm 2.3\%$) with LPS treatment. Thus, compounds **1–4**, **6** and **8** might be useful anti-inflammatory agents, while **6** is the most promising as it showed potent inhibitory activity against expression of both iNOS and COX-2 proteins (Fig. 5). On the basis of the above results, we suggest that further investigation of **6** in terms of its anti-inflammatory activities would be worthwhile for future drug development.

3. Experimental

3.1. General experimental procedures

Melting points were determined using a Fisher–Johns melting point apparatus. Optical rotations were measured on a JASCO P-1020 polarimeter. Ultraviolet spectra were recorded on a JASCO V-650 spectrophotometer. IR spectra were recorded on a JASCO FT/IR-4100 infrared spectrophotometer. The NMR spectra were recorded on a Varian 400MR FT NMR (or Varian Unity INOVA500 FT NMR) instrument at 400 MHz (or 500 MHz) for ^1H and 100 MHz (or 125 MHz) for ^{13}C in CDCl_3 . LRMS and HRMS were obtained by ESI on a Bruker APEX II mass spectrometer. Silica gel (Merck, 230–

400 mesh) was used for column chromatography. Precoated silica gel plates (Merck, Kieselgel 60 F-254, 0.2 mm) were used for analytical TLC. High-performance liquid chromatography was performed on a Hitachi L-7100 HPLC apparatus with a Merck Hibar Si-60 column (250×21 mm, $7 \mu\text{m}$) and on a Hitachi L-2455 HPLC apparatus with a Inertsil ODS-3 column (250×20 mm, $5 \mu\text{m}$).

3.2. Organism

S. crassocaula (specimen no. 20070402) was collected by hand using scuba off the coast of Dongsha, Taiwan, in April 2007, at a depth of 5–10 m, and stored in a freezer until extraction. A voucher sample was deposited at the Department of Marine Biotechnology and Resources, National Sun Yat-sen University.

3.3. Extraction and separation

The frozen bodies of *S. crassocaula* (0.5 kg, wet wt) were minced and exhaustively extracted with EtOAc ($1 \text{ L} \times 5$). The EtOAc extract (7.3 g) was chromatographed over silica gel by column chromatography and eluted with EtOAc in *n*-hexane (0–100%, stepwise) then with acetone in EtOAc (50–100%, stepwise) to yield 28 fractions. Fraction 10 eluted with *n*-hexane–EtOAc (6:1) and was further purified over silica gel using *n*-hexane–EtOAc (7:1) to afford six subfractions (A1–A6). Subfraction A2 was separated by silica gel using *n*-hexane– CH_2Cl_2 (8:1) to afford **6** (20.0 mg). Fraction 13 eluted with *n*-hexane–EtOAc (3:1) and was further purified over silica gel using *n*-hexane–acetone (5:1) to afford six subfractions (B1–B6) and **8** (60.0 mg). Subfraction B6 was separated by normal-phase HPLC using *n*-hexane–acetone (4:1) and further purified by reverse-phase HPLC using MeOH– H_2O (4:1) to afford **5** (1.0 mg). Fraction 15 eluted with *n*-hexane–EtOAc (2:1) and was further purified over silica gel using *n*-hexane–acetone (3:1) to afford seven subfractions (C1–C7). Subfraction C3 was separated by normal-phase HPLC using *n*-hexane–acetone (7:2) and further purified by reverse-phase HPLC using MeOH– H_2O (2.3:1) to afford **3** (6.2 mg), **4** (5.7 mg) and **7** (1.3 mg). Fraction 18 eluted with *n*-hexane–EtOAc (1:1) and was further purified over silica gel using *n*-hexane–acetone (3:1) to afford seven subfractions (D1–D7). Subfraction D3 was separated by reverse-phase HPLC using MeOH– H_2O (1.5:1) to afford **1** (10.2 mg) and **2** (3.0 mg).

3.3.1. Sarcocrassocolide A (1)

White solid; $[\alpha]_{\text{D}}^{25} +17.7$ (*c* 0.4, CHCl_3); UV (MeOH) λ_{max} 204 ($\log \epsilon = 3.5$); IR (neat) ν_{max} 3487, 2971, 2940, 1769, 1748, 1436, 1374, 1358, 1275, 1228 cm^{-1} ; ^1H NMR and ^{13}C NMR data see Tables 1 and 2; ESIMS m/z 413 $[\text{M}+\text{Na}]^+$; HRESIMS m/z 413.1938 $[\text{M}+\text{Na}]^+$ (calcd for $\text{C}_{20}\text{H}_{30}\text{O}_6\text{Na}$, 413.1940).

3.3.2. Sarcocrassocolide B (2)

White solid; $[\alpha]_{\text{D}}^{25} -71.7$ (*c* 0.3, CHCl_3); UV (MeOH) λ_{max} 204 ($\log \epsilon = 3.5$); IR (neat) ν_{max} 3481, 2965, 2951, 1755, 1456, 1444, 1386, 1370, 1346, 1270, 1227 cm^{-1} ; ^1H NMR and ^{13}C NMR data see Tables 1 and 2; ESIMS m/z 413 $[\text{M}+\text{Na}]^+$; HRESIMS m/z 413.1943 $[\text{M}+\text{Na}]^+$ (calcd for $\text{C}_{20}\text{H}_{30}\text{O}_6\text{Na}$, 413.1940).

3.3.3. Sarcocrassocolide C (3)

White solid; $[\alpha]_{\text{D}}^{25} +89.4$ (*c* 0.6, CHCl_3); UV (MeOH) λ_{max} 207 ($\log \epsilon = 3.6$); IR (neat) ν_{max} 3472, 2969, 2930, 1760, 1442, 1405, 1269, 1213 cm^{-1} ; ^1H NMR and ^{13}C NMR data see Tables 1 and 2; ESIMS m/z 355 $[\text{M}+\text{Na}]^+$; HRESIMS m/z 355.1883 $[\text{M}+\text{Na}]^+$ (calcd for $\text{C}_{20}\text{H}_{28}\text{O}_4\text{Na}$, 355.1885).

3.3.4. Sarcocrassocolide D (4)

White solid; $[\alpha]_{\text{D}}^{25} +17.2$ (*c* 0.6, CHCl_3); UV (MeOH) λ_{max} 215 ($\log \epsilon = 3.8$); IR (neat) ν_{max} 3462, 2968, 2949, 1748, 1456, 1268,

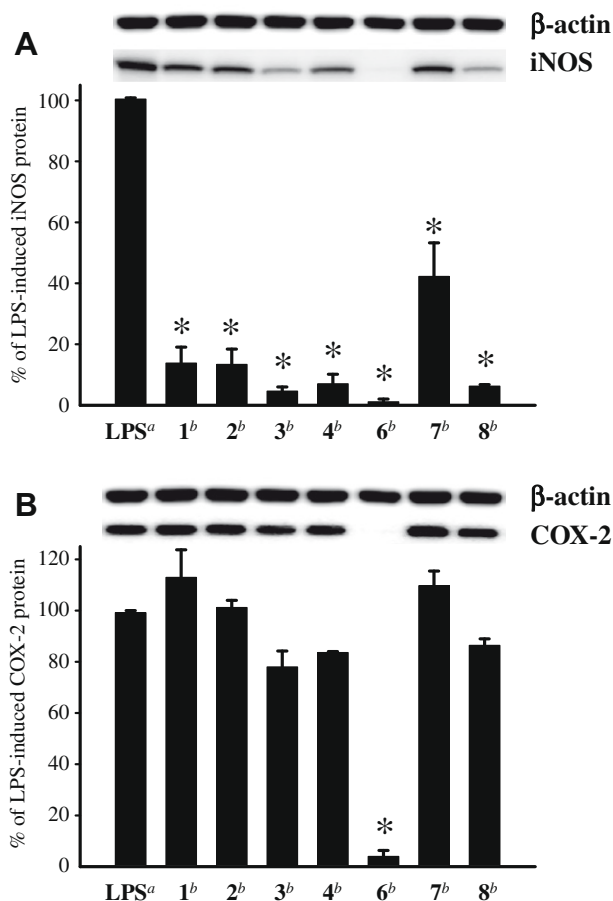


Figure 5. Effect of compounds **1–4** and **6–8** on iNOS and COX-2 proteins expression of RAW264.7 macrophage cells by immunoblot analysis. (A) Immunoblots of iNOS and β -actin; (B) immunoblots of COX-2 and β -actin. The values are mean \pm SEM ($n = 6$). Relative intensity of the LPS alone stimulated group was taken as 100%. Under the same experimental condition CAPE (caffeic acid phenylethyl ester, $10 \mu\text{M}$) reduced the levels of the iNOS and COX-2 to $2.5 \pm 3.7\%$ and $67.2 \pm 13.4\%$, respectively. *Significantly different from LPS alone stimulated group (* $P < 0.05$). ^aStimulated with LPS, ^bstimulated with LPS in the presence of **1–4** and **6–8** ($10 \mu\text{M}$).

1215 cm⁻¹; ¹H NMR and ¹³C NMR data see Tables 1 and 2; ESIMS *m/z* 355 [M+Na]⁺; HRESIMS *m/z* 355.1883 [M+Na]⁺ (calcd for C₂₀H₂₈O₄Na, 355.1885).

3.3.5. Sarcocrassolide E (5)

Colorless oil; [α]_D²⁵ -47.5 (c 0.1, CHCl₃); UV (MeOH) λ_{\max} 203 (log ϵ = 3.5); IR (neat) ν_{\max} 2923, 2853, 1770, 1749, 1385, 1271, 1223 cm⁻¹; ¹H NMR and ¹³C NMR data see Tables 1 and 2; ESIMS *m/z* 411 [M+Na]⁺; HRESIMS *m/z* 411.2150 [M+Na]⁺ (calcd for C₂₀H₃₀O₆Na, 411.2147).

3.3.6. Preparation of (S)- and (R)-MTPA esters of 1

To a solution of **1** (2.0 mg) in pyridine (100 μ L) was added (R)-(-)- α -methoxy- α -(trifluoromethyl)phenylacetyl chloride (10 μ L), and the solution was then allowed to stand overnight at room temperature. The reaction mixture was added to 1.0 mL of H₂O, followed by extraction with EtOAc (1.0 mL \times 3). The EtOAc-soluble layers were combined, dried over anhydrous MgSO₄ and evaporated. The residue was purified by a short silica gel column using acetone-*n*-hexane (1:2) to yield the (S)-MTPA ester **1a** (1.8 mg, 60%). The same procedure was applied to obtain the (R)-MTPA ester **1b** (1.6 mg, 52%) from the reaction of (S)-(+)- α -methoxy- α -(trifluoromethyl)phenylacetyl chloride with **1** in pyridine. Selective ¹H NMR (CDCl₃, 400 MHz) data of **1a**: δ 3.544 (1H, d, *J* = 12.4 Hz, H-1), 1.624 (1H, m, H-2b), 4.125 (1H, dd, *J* = 10.4, 4 Hz, H-7), 6.156 (1H, d, *J* = 1.6 Hz, H-17a), 5.569 (1H, d, *J* = 1.6 Hz, H-17b), 1.255 (3H, s, H₃-18), 5.043 (1H, s, H-19a), 4.934 (1H, s, H-19b); selective ¹H NMR (CDCl₃, 400 MHz) data of **1b**: δ 3.581 (1H, d, *J* = 12.4 Hz, H-1), 1.640 (1H, m, H-2b), 4.099 (1H, br s, H-7), 6.178 (1H, s, H-17a), 5.592 (1H, s, H-17b), 1.218 (3H, s, H₃-18), 5.026 (1H, s, H-19a), 4.920 (1H, s, H-19b).

3.4. Cytotoxicity testing

Cell lines were purchased from the American Type Culture Collection (ATCC). Cytotoxicity assays of compounds **1–4** were performed using the MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] colorimetric method.^{18,19}

3.5. In vitro anti-inflammatory assay

Macrophage (RAW264.7) cell line was purchased from ATCC. In vitro anti-inflammatory activities of compounds **1–4** and **6–8** were measured by examining the inhibition of lipopolysaccharide (LPS) induced up-regulation of iNOS (inducible nitric oxide synthetase)

and COX-2 (cyclooxygenase-2) proteins in macrophages cells using western blotting analysis.^{20,21}

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Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.bmc.2010.01.036.

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