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Insights into the structural diversity of cembranoids from *Sarcophyton glaucum* and their bioactivity†

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Three capnosane-based cembranoids (1–3), four normal cembranoids (4–7), and one sarsolenane-based cembranoid (8) were isolated from the soft coral *Sarcophyton glaucum*, including three new compounds: glacunoids A (1), B (4), and C (7). The chemical structures were elucidated using IR, MS, and NMR spectroscopy, with the absolute configurations of 1, 2, and 6 confirmed by single-crystal X-ray diffraction analysis. The bioactivity of the isolated compounds was evaluated for their effects on alkaline phosphatase (ALP) activity. All compounds demonstrated the ability to enhance ALP activity, with compound 8 exhibiting the most potent effect, achieving an ALP activity of $139.41 \pm 8.06\%$ without detectable cytotoxicity. This study represents the first report of three distinct structural types of cembranoids from a single soft coral species.

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1 Introduction

Soft corals of the genus *Sarcophyton* (family Alcyoniidae) are prolific sources of diverse secondary metabolites, particularly diterpenes of the cembrane type. These metabolites have garnered significant scientific interest due to their structural diversity and broad spectrum of biological activities, including antitumor, antifouling, and anti-inflammatory effects.^{1,2} Among these, cembranoids represent a prominent subclass, characterized by their unique structural frameworks and functional group diversity, which contribute to their pharmacological potential.^{1,2} Continuing our exploration of bioactive natural products from marine organisms, we focused on the soft coral *Sarcophyton glaucum* (Quoy & Gaimard, 1833) (Fig. 1),^{3–5} a widely distributed species in tropical and subtropical marine ecosystems. Previous investigations into *S. glaucum* revealed its capacity to produce an array of cembrane-type diterpenoids, which exhibited notable bioactivities ranging from cytotoxicity to ecological defense

mechanisms.^{1,2} In this study, we report the isolation and characterization of eight cembranoids from *S. glaucum*, classified into three structural subtypes: capnosane-based cembranoids (compounds 1–3),^{6,7} normal cembranoids (compounds 4–7),⁸ and a sarsolenane-based cembranoid (compound 8)⁷ (Fig. 2). Among these, three compounds—glacunoid A (1), a capnosane-based cembranoid, and two normal cembranoids, glacunoids B (4) and C (7)—were identified as new natural products. The chemical structures of these new compounds were elucidated using a combination of spectroscopic techniques, including infrared (IR), mass spectrometry (MS), one-dimensional (1D), and two-dimensional (2D) nuclear magnetic resonance (NMR) spectroscopy. Additionally, the absolute configurations of compounds 1, 2, and 6 were determined through single-crystal X-ray crystallography (Fig. 3), providing further insight into their


 Fig. 1 Soft coral *Sarcophyton glaucum*.

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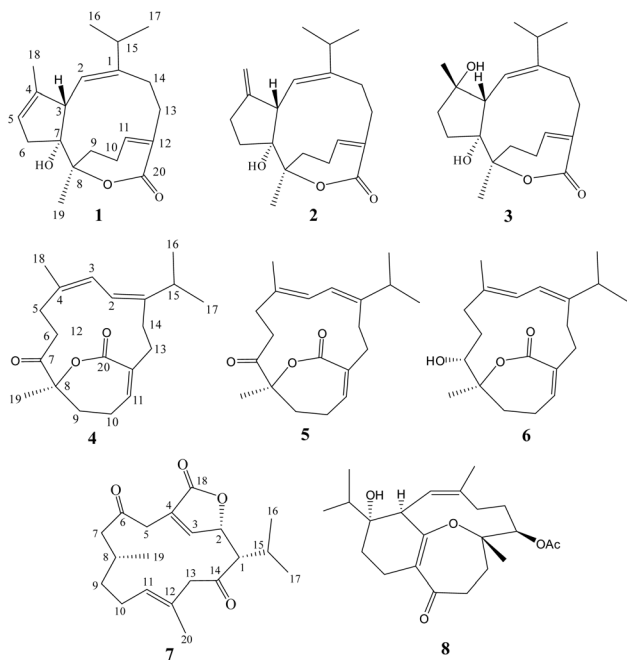



Fig. 2 Structures of metabolites 1–8.

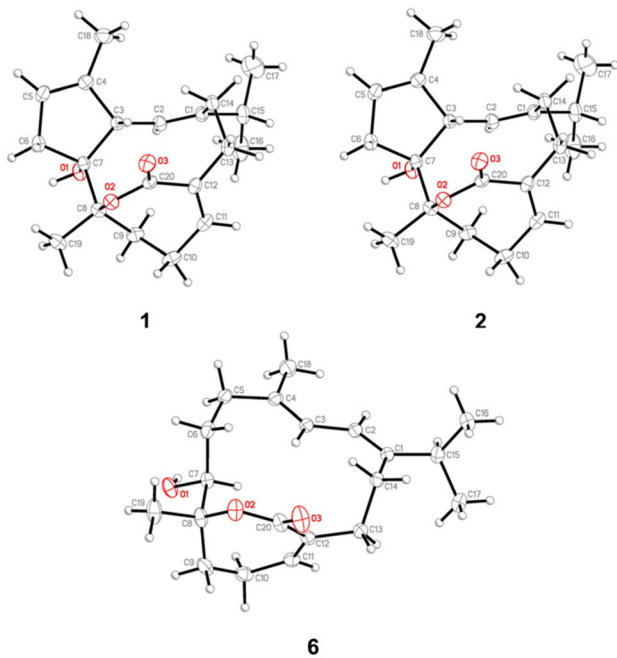


Fig. 3 A computer-generated ORTEP plot revealing the absolute structures of 1, 2, and 6.

stereochemistry. This study not only expands the chemical diversity of cembranoids from *S. glaucum* but also contributes to the understanding of their structural and biological significance.

2 Results and discussion

Our current results reported the isolation and structure elucidation of a novel capnosane-based cembranoids, glaucunoid A

(1) and two normal cembranoids, glaucunoids B (4) and C (7). The biosynthesis of capnosane from cembranoids involves the enzymatic transformation of a 14-membered macrocyclic cembrane skeleton derived from geranylgeranyl diphosphate (GGPP). Key steps include cyclization of GGPP to form the cembrane ring, followed by intricate rearrangements that generate the distinctive bicyclo[7.2.1] ring system characteristic of capnosane. Oxidative tailoring, mediated by cytochrome P450 enzymes, introduces functional groups such as hydroxyls, carbonyls, and epoxides, enhancing chemical complexity and bioactivity. Final modifications, such as esterification, complete the structure, showcasing the sophisticated enzymatic processes that diversify cembranoids into bioactive natural products like capnosane.⁹

Glaucunoid A (1) was isolated and crystallized as colorless needles. Its infrared (IR) absorptions showed the presence of both hydroxy (3448 cm^{-1}) and lactone (1669 cm^{-1}) functional groups. The molecular formula of 1 was determined to be $\text{C}_{20}\text{H}_{28}\text{O}_3$ by HRESIMS (m/z 339.1928 [$\text{M} + \text{Na}$]⁺) and ^{13}C NMR data (Table 1), suggesting seven degrees of unsaturation. Resonances of one ester carbonyl carbon (δ 170.1) and six olefinic carbons (δ 146.0, C; 144.3, CH; 141.5, C; 133.8, C; 120.5, CH; 120.0, CH) in the ^{13}C NMR and DEPT spectral data accounted for four double-bond equivalents, which indicated that 1 was a tricyclic compound. The ^1H NMR spectrum of 1 (Table 1) revealed the presence of resonances of three olefinic methine protons [δ 6.33 dd (8.4, 6.0); 5.47d (12.0); 5.36 brs] and four methyls [δ 1.48 s; 1.45 s; 1.08 d (6.6); 1.05 d (6.6)]. The spectroscopic data of 1 (^1H and ^{13}C NMR) were similar to those of 2, except for the absence of the 4,18-double bond and 4,5-single bond signals, both were replaced by signals of a 4,18-single bond and 4,5-double bond in 1. Moreover, single-crystal X-ray crystallographic analysis using Cu K α ($\lambda = 1.54178\text{ \AA}$) radiation source [Flack parameter $x = 0.16(5)$] (CCDC 2401636) allowed the assignment of the absolute configuration of 1. An Oak Ridge Thermal-Ellipsoid Plot (ORTEP) diagram (Fig. 3) showed the absolute configuration of the stereogenic carbons of 1 to be (3*S*,7*R*,8*R*).

Glaucunoid B (4) was isolated as a colorless oil. Its molecular formula $\text{C}_{20}\text{H}_{28}\text{O}_3$ was established by HRESIMS (m/z 339.1928, [$\text{M} + \text{Na}$]⁺) and ^{13}C NMR data, implying, seven degrees of unsaturation. By the analysis of ^{13}C NMR and DEPT spectra of 4, the carbons signals were assigned into four methyls, six sp^3 methylenes, one sp^3 methine, three sp^2 methines, and six quaternary carbons (including one oxygenated δ 86.2). The remaining two signals appearing in the lower field region of the spectrum were due to the quaternary carbons of one lactone carbon (δ 170.5) and one ketone carbonyl (δ 208.4). Careful analysis of ^1H - ^1H COSY correlations led to the establishment of five-part structures in Fig. 3. The molecular framework of 4 was further established by an HMBC experiment (Fig. 4).

The two rings and their connectivity were elucidated based on the following key HMBC correlations, both methyls H_3 -16 and H_3 -17 to C-1 and C-15, H_3 -18 to C-3, C-4 and C-5, H_3 -19 to C-7, C-8 and C-9, H-2 to C-1 and C-14, H_2 -5 to C-3 and C-4, H_2 -6 to C-7, H_2 -10 to C-12, H-11 to C-13 and C-20. Thus, 4 was found to possess three double bonds at C-1/C-2, C-3/C-4, and C-11/C-12,



Table 1 ^1H and ^{13}C NMR data for 1

C/H	δ_{H}^a (J in Hz)	δ_{C}^b , mult ^c	C/H	δ_{H}^a (J in Hz)	δ_{C}^b , mult ^c
1		146.0, C	11	6.33 dd (8.4, 6.0)	144.3, CH
2	5.47 d (12.0)	120.5, CH	12		133.8, C
3	3.33 d (11.4)	50.9, CH	13	2.95 dd (11.4, 5.4)	31.9, CH ₂
4		141.5, C		2.00 m	
5	5.36 brs	120.0, CH	14	2.65 m	26.8, CH ₂
				2.01 m	
6	2.86 d (15.6, 3.0)	42.4, CH ₂	15	2.25 m	34.4, CH
	2.18 d (15.6)		16	1.05 d (6.6)	23.0, CH ₃
7		90.0, C	17	1.08 d (6.6)	21.6, CH ₃
8		86.5, C	18	1.48 s	15.5, CH ₃
9	2.56 t (13.2)	38.6, CH ₂	19	1.45 s	27.6, CH ₃
	1.80 dd (13.2, 6.0)				
10	2.35 m	22.2 CH ₂	20		170.1, C
	2.07 m				

^a Spectra obtained in CDCl₃ at 600 MHz. ^b Spectra obtained in CDCl₃ at 150 MHz. ^c Attached protons were determined by DEPT experiments.

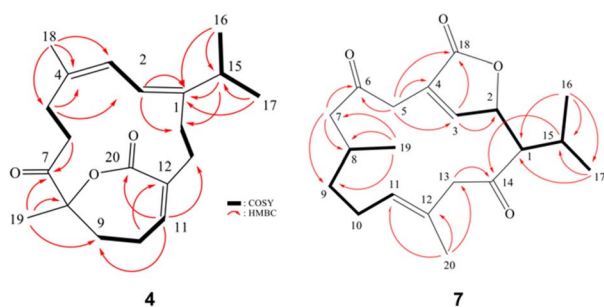


Fig. 4 Key HMBC and COSY correlations of 4 and 7.

one ketone group at C-7, and one α,β -unsaturated ϵ -lactone at C-8/C-12. Linking all the above functional groups to the cembrane skeleton yielded a gross structure of 4. The ^{13}C NMR data of 4 were found to be quite similar to those of 5. We then carefully examined the NOESY correlations of 4. The NOESY spectrum (Fig. 4) showed a correlation of H₃-18 with H-3, but not with H-2, H-3 with H₃-16 revealing the Z configuration of the C-1/C-2 and C-3/C-4 double bonds. The comparison of its ^1H and ^{13}C NMR with those of 5 indicated that 4 is the 1Z and 3Z isomer of 5. Based on the well-established 8S configuration of 4 and the fact that both 4–6 could be the precursors of each other by dehydration, the structure of 4 was then established.

Compound 7 was isolated as a colorless oil. The HRESIMS (m/z 359.1928, $[\text{M} + \text{Na}]^+$) and ^{13}C NMR data of 7 indicated the molecular formula C₂₀H₂₈O₃. IR absorptions were observed at 1761 and 1723 cm⁻¹, suggesting the presence of carbonyl groups in 7. The ^{13}C NMR and DEPT spectroscopic data (Table 2), showed signals of four methyls, five sp³ methylenes, three sp³ methines, one sp³ oxymethine, two sp² methines, and five sp² quaternary carbons (including two ketones and one ester carbonyl). From the ^1H NMR spectrum of 7, the resonances of two olefinic protons [δ_{H} 7.03 dd (3.0, 1.2) and δ_{H} 5.24 dd (9.0, 5.4)], one oxygenated methine [δ_{H} 5.21 dd (9.6, 1.8)] and four methyls [δ_{H} 1.69 s, 1.14 d (7.2); 1.06 d (7.2); 0.95 d (6.6)] were

Table 2 ^1H and ^{13}C NMR data for 4 and 7

C/H	4		7	
	δ_{H}^a (J in Hz)	δ_{C}^b , mult ^c	δ_{H}^a (J in Hz)	δ_{C}^b , mult ^c
1		140.7, C	2.85 m	58.6, CH
2	5.21 d (11.4)	118.8, CH	5.21 dd (9.6, 1.8)	80.3, CH
3	6.05 d (11.4)	120.2, CH	7.03 dd (3.0, 1.2)	150.4, CH
4		138.8, C		127.7, C
5	3.15 dt (15.0, 4.8)	23.5, CH ₂	3.42 d (16.2)	40.2, CH ₂
	1.81 m		3.16 d (16.2)	
6	2.95 m	39.8, CH ₂		205.6, C
	2.33 t (4.8)			
7		208.4, C	2.32 d (7.2)	49.8, CH ₂
8		86.2, C	1.69 m	28.2, CH
9	2.60 m; 2.24 m	37.4, CH ₂	1.42 m	35.7, CH ₂
10	2.37 m; 2.16 m	22.6, CH ₂	2.20 m; 1.95 m	24.3, CH ₂
11	5.78 dt (7.8, 2.0)	131.4, CH	5.24 dd (9.0, 5.4)	131.0, CH
12		136.6, C		128.1, C
13	2.63 m; 2.39 m	26.3, CH ₂	3.27 d (12.6)	56.2, CH ₂
			2.84 d (13.2)	
14	2.37 m; 2.18 m	25.5, CH ₂		208.1, C
15	3.01 m	28.9, CH	2.30 m	28.9, CH ₂
16	1.04 d (7.2)	21.1, CH ₃	1.14 d (7.2)	20.7, CH ₃
17	1.02 d (7.2)	20.7, CH ₃	1.06 d (7.2)	19.3, CH ₃
18	1.86 s	26.5, CH ₃		172.5, C
19	1.48 s	27.6 CH ₃	0.95 d (6.6)	20.3, CH ₃
20		170.5 C	1.69 s	16.6, CH ₃

^a Spectra obtained in CDCl₃ at 600 MHz. ^b Spectra obtained in CDCl₃ at 150 MHz. ^c Attached protons were determined by DEPT experiments.

observed. The planar structure and all the ^1H and ^{13}C chemical shifts of 7 were elucidated by 2D NMR spectroscopic analysis, in particular ^1H - ^1H COSY and HMBC experiments (Fig. 4). Thus, 7 was found to possess two double bonds at C-3/C-4 and C-11/C-12, two ketone groups at C-6 and C-14, and an α,β -unsaturated γ -lactone moiety at C-4, C-3, C-2, and C-18. The relative configuration of 7 was elucidated from the NOESY spectrum, which was compatible with that of 7 obtained by the MM2 force field calculations suggesting the most stable conformations as shown



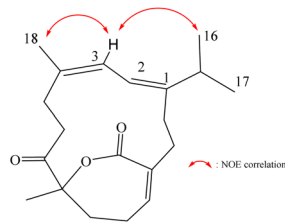


Fig. 5 Selected NOE correlations observed for 4.

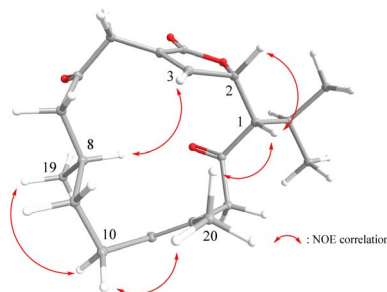


Fig. 6 Selected NOE correlations observed for 7.

in Fig. 5. In the NOESY spectrum, it was found that H₃-19 (δ_{H} 1.44, s) showed NOE interactions with one of the methylene protons at C-10 (δ_{H} 1.95, m). Therefore, H₃-19 and one of proton H-10 (δ_{H} 1.95, m) were located on the α -face, while the other (δ_{H} 2.20, m) was oriented as β H-10. The NOE correlations observed between H₃-20 with H-10 β and H-1 reflected the β -orientations of H-1. Furthermore, the NOE correlations between the H-1 and H-2 suggested the β -orientation of H-2. From the above findings and further analysis of other NOE interactions (Fig. 6), the structure of 7 was fully established.

The five obtained known cembrane diterpenoids were identified as sarsolilide A (2) and B (3),^{6,7} sarcophytolide (5),⁸ (4Z,8S,9R,12E,14E)-9-hydroxy-1-isopropyl-8,12-dimethyl oxabicyclo [9.3.2]-hexadeca-4,12,14-trien-18-one (6)⁸ and sarsolenone (8)⁷ by comparing the obtained spectroscopic data with published values. The absolute configurations of 2 and 6 were confirmed by single-

crystal X-ray diffraction analysis with Cu K α radiation [Flack parameters = 0.08(6) and 0.07(7)] for the first time (Fig. 3).

Previous studies indicated that cembranoids 1, 2, and 4–8 can be used to treat osteoclastogenic diseases. Using MG-63 human mesenchymal stem cells, we conducted an alkaline phosphatase (ALP) ELISA assay (Table 3). The results indicated that all tested compounds significantly increased ALP activity, with compound 8 showing the most significant effect, achieving an ALP activity of $139.41 \pm 8.06\%$ without cytotoxic effects. Moreover, these findings suggested that the tested compounds have potential biological activity in promoting ALP activity, and tested compounds did not affect cell the viability of MG63 cells.

3 Conclusions

A diverse series of cembrane diterpenoids was isolated from the wild-type soft coral *Sarcophyton glaucum*. This study identified three new diterpenoids—glacunoids A (1), B (4), and C (7)—alongside five known compounds (2, 3, 5, 6, and 8). The absolute configurations of compounds 1, 2, and 6 were unambiguously confirmed by single-crystal X-ray diffraction analysis. Notably, metabolites 1–3 are rare examples of capnosane-based cembranoids featuring an α,β -unsaturated ϵ -lactone moiety. This work represents the first report of three distinct structural types of cembrane diterpenoids (normal-, capnosane-, and sarsolenane-based) isolated from a single soft coral species. All the isolated compounds enhanced alkaline phosphatase (ALP) activity, with sarsolenone (8) demonstrating the highest activity ($139.41 \pm 8.06\%$) and no cytotoxicity against MG-63 human mesenchymal stem cells.

4 Experimental

4.1 General experimental procedures

Infrared (IR) spectra were obtained on a Fourier-transform IR spectrophotometer (model: JASCO P-2000). ¹H and ¹³C NMR spectra were recorded on a 600R NMR spectrometer (JEOL, Tokyo, Japan) with CDCl₃ (Sigma-Aldrich, St. Louis, MO, USA) as the deuterated solvent. The detected signals in ¹H and ¹³C NMR spectra were corrected at 7.26 ppm (singlet) and 77.0 ppm (triplet), respectively. The coupling constants (*J*) were converted to Hz. MS data, including ESIMS and HRESIMS, were obtained using a Bruker 7 Tesla Solera FTMS system (Bruker, Bremen, Germany). Optical rotations were determined by a digital polarimeter (Jasco P-1010). Single-crystal X-ray analyses were performed on a Bruker D8 Venture diffractometer. Thin-layer chromatography was performed on plates precoated with silica gel 60 F₂₅₄ (0.25 mm-thick, MERCK); the plates were then sprayed with 10% (v/v) H₂SO₄ in methanol, followed by heating to visualize the spots. A normal-phase (NP) HPLC was performed using a system comprised of a HITACHI 5110 pump, a RHEODYNE 7725i injection port, and a NP column (YMC pack SIL, 5 μm , 12 nm, 250 \times 20 mm, YMC group). Reverse-phase (RP) HPLC was performed using a system comprised of a HITACHI L-2130 pump, a HITACHI L-2455 photodiode array detector, a RHEODYNE 7725i injection port and an RP column (Luna 5 μm C18(2) 100 \AA , 250 \times 21.2 mm, Phenomenex).

Table 3 Evaluation of the ALP activity after the treatment of MG63 cells with 1, 2, and 4–8 at a concentration of 10 μM for 72 h^a

Compounds	ALP activity (%)		Cell viability (%)	
Control	100.00 \pm 0.65		100.00 \pm 1.03	
1	109.10 \pm 3.38	***	105.34 \pm 4.69	*
2	113.44 \pm 4.80	***	103.00 \pm 2.27	*
4	110.75 \pm 3.90	***	106.18 \pm 3.81	**
5	110.98 \pm 4.74	**	96.72 \pm 9.71	
6	107.06 \pm 3.48	**	106.88 \pm 6.10	*
7	128.06 \pm 3.04	***	93.58 \pm 6.73	
8	139.41 \pm 8.06	***	107.29 \pm 6.83	*
17 β -Estradiol ^b	146.46 \pm 2.34	***	68.23 \pm 9.49	

^a The data are expressed as the standard error of the mean ($n = 3$). The significance was determined using Student's *t*-test (* $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$) and by comparison of the results with those of the untreated cells. ^b 17 β -Estradiol was utilized as a positive control at a concentration of 10 μM .



4.2 Animal material

The soft coral *S. glaucum* (specimen no. NMMBA-SC-2024-1) samples were collected by scuba divers at a depth of around 10–15 m off the coast of Pingtung County, Taiwan, in February 2024. The samples were frozen immediately after collection. A voucher sample was deposited at the National Museum of Marine Biology and Aquarium.

4.3 Extraction and isolation

The frozen *S. glaucum* material (1.7 kg, wet wt) was minced and treated with an equal mixture of methanol (MeOH) and dichloromethane (CH₂Cl₂) at ambient temperature. This procedure resulted in a crude extract weighing 29.7 g. Liquid–liquid partition was performed to separate the mixture into ethyl acetate (EtOAc) and water layers. The ethyl acetate phase was evaporated under reduced pressure to afford a residue (15.9 g), and the residue was subjected to column chromatography on silica gel, using *n*-hexane, *n*-hexane and EtOAc mixture of increasing polarity, and finally pure acetone to yield 11 fractions: Fr-1 (eluted by *n*-hexane : EtOAc, 50 : 1), Fr-2 (eluted by *n*-hexane : EtOAc, 10 : 1), Fr-3 (eluted by *n*-hexane : EtOAc, 8 : 1), Fr-4 (eluted by *n*-hexane : EtOAc, 5 : 1), Fr-5 (eluted by *n*-hexane : EtOAc, 3 : 1), Fr-6 (eluted by *n*-hexane : EtOAc, 2 : 1), Fr-7 (eluted by *n*-hexane : EtOAc, 1 : 1), Fr-8 (eluted by *n*-hexane : EtOAc, 2 : 1), Fr-9 (eluted by *n*-hexane : EtOAc, 5 : 1), Fr-10 (eluted by EtOAc) and Fr-11 (eluted by acetone). Fraction 2 was further purified with normal-phase HPLC (*n*-hexane : EtOAc, 4 : 1) to afford seven subfractions (2A – 2G). Subfraction 2E was then separated by normal-phase HPLC (*n*-hexane : EtOAc, 4 : 1) to obtain 4 (30.5 mg) and 5 (680 mg). Fraction 3 was separated by normal-phase HPLC using *n*-hexane : EtOAc (5 : 1) to afford 1 (20.6 mg) and 2 (80.6 mg). Fraction 4 was further separated by silica gel column chromatography with gradient elution (*n*-hexane : EtOAc, 4 : 1 to 2 : 1) to afford ten subfractions (4 A–4J). Subfraction 4B was separated by normal-phase HPLC using *n*-hexane : EtOAc (7 : 2) to afford 7 (2.8 mg). Subfraction 4E (20.5 mg) was also separated by normal-phase HPLC using *n*-hexane : EtOAc (2 : 1) to afford 3 (1.5 mg), 6 (49.5 mg) and 8 (30.6 mg).

4.4 Structural characterization of compounds 1, 4, and 7

4.4.1 Glacunoid A (1). Colorless needles; mp 151–153 °C; [α]_D²⁵ +133 (*c* 0.05, CHCl₃); IR (KBr) ν_{\max} 3448, 2956, 2923, 1669, 1460 and 1378 cm⁻¹; ¹H (600 MHz, CDCl₃) and ¹³C NMR (150 MHz, CDCl₃) data (Table 1); ESIMS: *m/z* 339 [M + Na]⁺; HRESIMS: *m/z* 339.1928 (calcd for C₂₀H₂₈O₃ + Na, 339.1930).

4.4.2 Glacunoid B (4). Colorless oil; [α]_D²⁵ +17 (*c* 0.19, CHCl₃); IR (KBr) ν_{\max} 2963, 2932, 1712, 1450 and 1375 cm⁻¹; ¹H (600 MHz, CDCl₃) and ¹³C NMR (150 MHz, CDCl₃) data (Table 2); ESIMS: *m/z* 339 [M + Na]⁺; HRESIMS: *m/z* 339.1928 (calcd for C₂₀H₂₈O₃ + Na, 339.1930).

4.4.3 Glacunoid C (7). Colorless oil; [α]_D²⁵ –157 (*c* 0.04, CHCl₃); IR (KBr) ν_{\max} 2958, 2924, 1763, 1703, 1458 and 1374 cm⁻¹; ¹H (600 MHz, CDCl₃) and ¹³C NMR (150 MHz, CDCl₃) data (Table 2); ESIMS: *m/z* 355 [M + Na]⁺; HRESIMS: *m/z* 355.1878 (calcd for C₂₀H₂₈O₄ + Na, 355.1879).

4.5 Single-crystal X-ray analysis of 1, 2, and 6

4.5.1 Glacunoid A (1). The ethyl acetate solution was allowed to slowly evaporate to generate a suitable colorless crystal. Diffraction intensity data were obtained on a Bruker D8 Venture diffractometer using graphite-monochromated Cu K α radiation ($\lambda = 1.54178$ Å). Crystal data for this compound: C₂₀H₂₈O₃ (formula weight 316.42), approximate crystal size, 0.330 × 0.289 × 0.110 mm³, monoclinic, space group, *P*₂₁ (# 4), *T* = 200(2) K, *a* = 10.9787 (4) Å, *b* = 11.8275 (4) Å, *c* = 13.8005 (5) Å, $\alpha = \beta = \gamma = 90^\circ$, *V* = 1792.00 (11) Å³, *Z* = 4, *D*_{calcd} = 1.173 Mg m⁻³, *F*(000) = 688, $\mu(\text{MoK}\alpha) = 0.609$ mm⁻¹. A total of 18 753 reflections were collected in the range 4.924 < θ < 74.417, with 3646 independent reflections [*R*(int) = 0.0333], completeness to theta was 99.8%; semi-empirical from equivalents absorption correction applied; refinement method: full-matrix least-squares on *F*², the data/restraints/parameters were 3646/0/216; goodness-of-fit on *F*² = 1.046; final *R* indices [*I* > 2 sigma (*I*)], *R*₁ = 0.0317, *wR*₂ = 0.0859; *R* indices (all data), *R*₁ = 0.0326, *wR*₂ = 0.0866, largest difference peak and hole, 0.207 and –0.145 e Å⁻³; absolute structure parameter, 0.16(5).

4.5.2 Sarsolilide A (2). The ethyl acetate solution was allowed to slowly evaporate to generate a suitable colorless crystal. Diffraction intensity data were obtained on a Bruker D8 Venture diffractometer using graphite-monochromated Cu K α radiation ($\lambda = 1.54178$ Å). Crystal data for this compound: C₂₀H₂₈O₃ (formula weight 316.42), approximate crystal size, 0.171 × 0.142 × 0.109 mm³, monoclinic, space group, *P*₂₁ (# 4), *T* = 200(2) K, *a* = 10.9227 (3) Å, *b* = 11.4366 (3) Å, *c* = 14.0387 (4) Å, $\alpha = \beta = \gamma = 90^\circ$, *V* = 1573.69 (8) Å³, *Z* = 4, *D*_{calcd} = 1.198 Mg m⁻³, *F*(000) = 688, $\mu(\text{MoK}\alpha) = 0.622$ mm⁻¹. A total of 15 840 reflections were collected in the range 4.988 < θ < 74.394, with 3550 independent reflections [*R*(int) = 0.0292], completeness to theta was 100%; semi-empirical from equivalents absorption correction applied; refinement method: full-matrix least-squares on *F*², the data/restraints/parameters were 3550/6/212; goodness-of-fit on *F*² = 1.031; final *R* indices [*I* > 2 sigma (*I*)], *R*₁ = 0.0296, *wR*₂ = 0.0799; *R* indices (all data), *R*₁ = 0.0302, *wR*₂ = 0.0804, largest difference peak and hole, 0.167 and –0.165 e Å⁻³; absolute structure parameter, 0.08(6).

4.5.3 3 β ,11-dihydroxy-5 α ,6 α -epoxy-24-methylene-9,11-secocholestan-9-one (6). The ethyl acetate solution was allowed to slowly evaporate to generate a suitable colorless crystal. Diffraction intensity data were obtained on a Bruker D8 Venture diffractometer using graphite-monochromated Cu K α radiation ($\lambda = 1.54178$ Å). Crystal data for this compound: C₂₀H₃₀O₃ (formula weight 318.44), approximate crystal size, 0.332 × 0.118 × 0.100 mm³, monoclinic, space group, *P*₂₁ (# 4), *T* = 200(2) K, *a* = 7.6524 (2) Å, *b* = 13.8446 (3) Å, *c* = 17.0676 (4) Å, $\alpha = \beta = \gamma = 90^\circ$, *V* = 1808.22 (7) Å³, *Z* = 4, *D*_{calcd} = 1.170 Mg m⁻³, *F*(000) = 696, $\mu(\text{MoK}\alpha) = 0.604$ mm⁻¹. A total of 14 037 reflections were collected in the range 4.111 < θ < 74.422, with 3676 independent reflections [*R*(int) = 0.0327], completeness to theta was 99.9%; semi-empirical from equivalents absorption correction applied; refinement method: full-matrix least-squares on *F*², the data/restraints/parameters were 3676/0/213; goodness-of-fit on *F*² =



1.035; final R indices [$I > 2$ sigma (I)], $R_1 = 0.0296$, $wR_2 = 0.0777$; R indices (all data), $R_1 = 0.0315$, $wR_2 = 0.0790$, largest difference peak and hole, 0.149 and -0.141 e \AA^3 ; absolute structure parameter, $-0.07(7)$.

Crystallographic data for compounds **1**, **2**, and **6** were deposited at the Cambridge Crystallographic Data Centre (CCDC No. 2401636, 2401635, and 2401634 for compounds **1**, **2**, and **6**, respectively).

4.6 ALP activity assay

In this study, we conducted assays to evaluate ALP activity. For each assay, MG63 cells were exposed to one of the **1**, **2**, and **4–8** compounds at a 10 μM concentration for three days. After treatment, the MG63 cells in each well were washed with phosphate-buffered saline (PBS) twice, and then lysis buffer with 0.1% Triton X-100 was added. After sonication, the cell lysate samples were subjected to ALP activity measurement with *p*-nitrophenyl phosphate in 0.2 M Tris hydrochloride magnesium chloride hexahydrate buffer (pH 9.5). The ALP activity of each sample was normalized by the protein levels determined with a protein assay kit (BCA kit; Thermo Fischer Scientific). The enzymatic reaction was halted by adding 0.1 N NaOH to the solution as soon as a noticeable color change occurred, thereby ceasing the conversion of *p*-nitrophenyl phosphate to *p*-nitrophenol. The absorbance at 450 and 562 nm was measured *via* a spectrophotometer, and the ALP activity was subsequently quantified by relating these absorbance measurements to a standard curve derived from known concentrations of *p*-nitrophenol.

4.7 Cell viability assay

Cell viability assays were conducted by seeding 1×10^3 cells into each well of a 96-well plate, and after 24 hours, a mixture of culture medium supplemented with either 10.0 μM of a specific drug or 10.0 μM alendronate sodium hydrate was added to the wells. After incubation for 72 hours at 37 $^\circ\text{C}$, the cells in each well were rinsed and treated with a solution composed of 10 μL of MTT solution (5 mg mL^{-1}) and 90 μL of culture medium. After another incubation at 37 $^\circ\text{C}$ for 4 hours, this process led to the formation of formazan crystals, which were subsequently dissolved by adding 100 μL of DMSO to each well. Upon complete dissolution, the optical density of the solution was measured at a wavelength of 570 nm *via* an ELISA reader (Thermo Fischer Scientific).

Data availability

The data supporting this article have been included as part of the ESI.†

Author contributions

Mohamed El-Shazly and Jui-Hsin Su conceived and designed the experiments; Hsiao-Ling Chung, and Li-Guo Zheng performed the sample collections, extraction, isolation, structures determination, and qualitative HPLC analysis; the pharmacological experiments were carried out by You-Ying Chen and Jui-

Hsin Su contributed reagents and analysis tools; Mohamed El-Shazly and Jui-Hsin Su participated in data interpretation, wrote the manuscript and revised the paper.

Conflicts of interest

There are no conflicts to declare.

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