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Scalarane-type Sesterterpenes from the Philippines Sponge *Hyrtios* sp.

Jae-Hyeong Choi^{1,2}, Hyi-Seung Lee^{1,2*}, and Wilfredo L. Campos³

¹Marine Biotechnology Research Center, Korea Institute of Ocean Science & Technology
Busan 49111, Korea

²Department of Applied Ocean Science, University of Science and Technology
Daejeon 34113, Korea

³OceanBio Lab, University of the Philippines Visayas
Iloilo 5023, Philippines

Abstract : The marine sponge *Hyrtios* sp. collected in the Philippines was extracted and partitioned. The resulting organic layer was purified by C₁₈ reversed-phase column chromatography and HPLC to achieve the separation of nine scalarane-type sesterterpenes, including one new compound with eight known scalarane analogs. The chemical structures of the isolated compounds **1–9** were elucidated by 1D and 2D NMR and MS data analysis. All nine compounds were evaluated for their antibacterial activities against three Gram-positive and three Gram-negative bacteria. The compound **3** exhibited potent antibacterial activities against *Bacillus subtilis* and *Micrococcus luteus*. The compounds **7** and **9** displayed considerable activities against *Bacillus subtilis* and the others had moderate results.

Key words : marine sponge, scalarane, sesterterpene, *Hyrtios* sp., antibacterial activities

1. Introduction

Marine organisms have received growing attention as potential sources for bioactive secondary metabolites exhibiting novel chemical structures. In particular, marine sponges are known to contain various classes of bioactive compounds, such as alkaloids, terpenoids, peptides, glycosides, and macrolides (Amina and Al Musayeb 2018). For example, spongistatin isolated from *Hyrtios* sp. and halichondrin isolated from *Halichondria okadae* were previously reported as microtubule-destabilizing agents (Miller et al. 2010).

Scalarane-type sesterterpenoids are compounds possessing a 6/6/6/6-tetracyclic or 6/6/6/6/5-pentacyclic fused ring system. Since scalarin, the first scalarane-type sesterterpene, was isolated from the sponge *Cacospongia scalaris* (Fattorusso et al. 1972), diverse subgroups of these skeletal structures have been discovered, including homoscalaranes (Fontana et al. 2000; Jaspars et al. 1997;

Roy et al. 2002) and isoscalaranes (Davis and Capon 1993; Song et al. 2008; De Rosa et al. 1994). Moreover, further research into the synthesis (Ungur and Kulciçki 2004) or semi-synthesis (Kamel et al. 2009) of the scaffolds of scalarane derivatives has been conducted. Scalarane-type sesterterpenoids are commonly found in marine sponges belonging to the order Dictyoceratida (Liu et al. 2007; Gonzalez 2010), and are occasionally found in shell-less molluscs, such as nudibranchs (Gavagnin et al. 2004), likely due to their dietary source of sponges. For this reason, scalarane-type sesterterpenoids have been regarded as a useful chemotaxonomical marker (Jaspars et al. 1997).

Scalarane-type sesterterpenoids exhibit an extensive spectrum of bioactivities, including anticancer, antibacterial, anti-inflammatory, and anti-fouling activities. In particular, the majority of scalaranic sesterterpenoids have a broad range of cytotoxicities against many cancer cell lines, and so play the key role as eco-physiological mediators in the chemical defensive mechanism of marine invertebrates (Gonzalez 2010).

*Corresponding author. E-mail : hslee@kiost.ac.kr

11 using semi-preparative reversed-phase HPLC (YMC-Pack C₁₈, 250 × 10 mm, 65% MeCN in H₂O). Compounds **3** (8.0 mg) and **9** (5.5 mg) were purified from mp 14 by analytical reversed-phase HPLC (YMC-Pack C₁₈, 250 × 4.6 mm, 62% MeCN in H₂O). The combined fraction of mp 16–17 was purified by analytical reversed-phase HPLC (YMC-Pack C₁₈, 250 × 4.6 mm, 65% MeCN in H₂O) to give compound **4** (7.0 mg).

Compound **1**: White amorphous solid; $[\alpha]_D^{25} +10$ (c 0.1, MeOH); IR ν_{\max} 3407, 2929, 1742, 1654, 1554, 1456, 1389, 1318, 1251, 1042 cm⁻¹; UV(MeOH) λ_{\max} (log ϵ) 214 (3.68) nm; HRESIMS m/z 409.2716 [M + Na]⁺ (calcd for 409.2713, C₂₅H₃₈O₃Na⁺); ¹H NMR (CD₃OD and C₅D₅N, 600 MHz) and ¹³C NMR (CD₃OD and C₅D₅N, 150 MHz) see Table 1.

Minimum Inhibitory Concentration (MIC) assay

The MIC assay is a useful method to evaluate the lowest concentration of antibacterial sources, as it is based on an antibacterial susceptibility test for bioactive natural products that cause inhibited bacterial growth or death.

The assay was carried out against six bacteria. The three Gram-positive bacteria, namely *Bacillus subtilis* (KCTC-1021), *Micrococcus luteus* (KCTC-1915), and *Staphylococcus aureus* (KCTC-1927), and three Gram-negative bacteria, namely *Escherichia coli* (KCTC-2441), *Salmonella typhimurium* (KCTC-2515), and *Klebsiella pneumonia* subsp. (KCTC-2690) were purchased from the Korean Collection for Type Cultures (KCTC). All six bacteria were streaked onto Mueller-Hinton agar (MHA) plates and cultured in a standing incubator for 24 h at 37°C. A separated single colony was transferred from the MHA plates to Muller-Hinton broth, and incubated for 24 h at 37°C and 170 rpm in a shaking incubator. In brief, the six bacteria were inoculated into 96-well plates and then treated with compounds **1–9** diluted in DMSO (100 μ L) in accordance with concentrations ranging from 128 to 0.25 μ g/mL (i.e., 128, 64, 32, 16, 8, 4, 2, 1, 0.5, and 0.25 μ g/mL). Subsequently, these mixtures were cultivated in a standing incubator for 24 h at 37°C.

3. Results and Discussion

Compound **1** has a molecular formula of C₂₅H₃₈O₃, as deduced from its high-resolution electrospray ionization mass spectrometry (HRESIMS) data. The ¹H NMR spectrum of **1** displayed resonance for two olefinic protons, seven methylene protons, five methine protons, and five

methyl groups (Table 1). The planar structure of **1** was determined by combined investigation of the COSY and HMBC correlations (Fig. 2). The COSY correlations of H₂-1/H₂-2/H₂-3, H₂-6/H₂-7, H-9/H₂-11/H-12, and H-14/H₂-15/H₂-16 and the HMBC correlations from H₃-21 and H₃-22 to C-3, C-4 and C-5, from H₃-23 to C-1, C-5, C-9 and C-10, from H₃-24 to C-7, C-8, C-9 and C-14, and from H₃-25 to C-12, C-13, C-14 and C-18 confirmed the frame of a 6/6/6/6-membered ring system. The presence of a five-membered ring containing one oxygen atom was also elucidated by the HMBC correlations from H-19 to C-17, C-18 and C-20, from H-20 to C-17, C-18 and C-19, and from H-16 to C-17, C-18 and C-20. Furthermore, the HMBC correlations from H₃-21 and H₃-22 to C-3, C-4 and C-5 suggested the presence of a gem-dimethyl group on quaternary carbon C-4, thereby defining a 6/6/6/6/5-pentacyclic carbon skeleton. The structure of compound **1** shared the same carbon and heteroatom framework with 12-*O*-desacetylfirosalarol, with the exception of differences in the stereochemistry at H-12. The unsaturated degree of compound **1** also indicated the presence of a pentacyclic ring system with a furan ring attached at C-17 and C-18.

Compound **1** has not been reported, but its 12-epimer has been already isolated from *Hyrtios* sp. and *Hyrtios* cf. *erectus* (Cimino et al. 1978; Doi et al. 1993). The previously reported ¹H and ¹³C NMR chemical shifts of the C-12 position in C₅D₅N (δ_C 71.4 and δ_H 4.62) were somewhat different from those of compound **1** (δ_C 77.2 and δ_H 3.90) in C₅D₅N. The stereochemistry of H-12 was therefore further investigated by the analysis of NOESY correlations (Fig. 3) and through comparison of the proton-proton coupling constants.

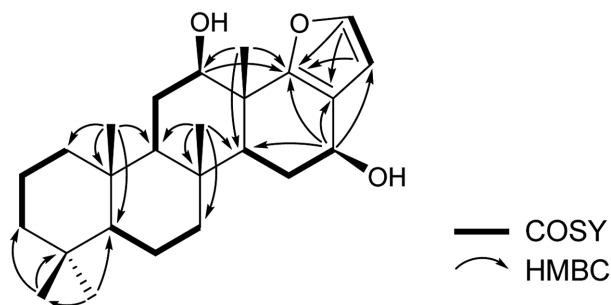
More specifically, the NOESY correlations of H-2/H-11/H-22/H-23/H-25 and H-6/H-24/H-25 suggested that H-22, H-23, H-24, and H-25 are on the β -face, while the strong NOESY correlations of H-9/H-12/H-14 suggested that H-9 and H-14 are on the α -face and H-12 is an axial proton. In addition, the stereochemistry of H-12 (δ_H 3.90) was confirmed as the down orientation by study of the coupling constant, which showed J value of 11.04 Hz due to the trans-diaxial coupling between the axial proton on H-12 (δ_H 3.90) and the axial proton on H-11 (δ_H 1.77); such a coupling was not observed in the previously reported 12-epimer (Lee et al. 2014). Furthermore, the strong NOESY signals of H-14/H-15/H-16 indicated that H-14, H-15, and H-16 present the same orientation as H-12. A strong NOESY correlation was also observed between H-12 and pseudoaxial proton H-16 due to the D ring being a cyclohexene ring.

Table 1. ^1H (600 MHz) NMR and ^{13}C (150 MHz) NMR data for **1** (δ in ppm, J in Hz)

| Position | ^{13}C NMR ^a | ^1H NMR ^a | ^{13}C NMR ^b | ^1H NMR ^b |
|----------|----------------------------------|--|----------------------------------|-------------------------------|
| 1 | 41.0 | 0.87, m 1.73, brd (13.7) | 40.4 | 0.56, brd (12.5) 1.58, m |
| 2 | 19.7 | 1.69, dd (10.3, 2.3) 1.47, m | 19.3 | 1.33, m 1.53, m |
| 3 | 43.3 | 1.18, ddd (13.5, 13.5, 4.3) 1.38, brdd (12.9) | 42.7 | 1.09, m 1.32, m |
| 4 | 34.2 | | 33.8 | |
| 5 | 58.0 | 0.88, m | 57.1 | 0.65 m |
| 6 | 19.3 | 1.61, m 1.47, m | 18.9 | 1.29, m 1.46, m |
| 7 | 42.6 | 1.03, m 1.81, m | 42.1 | 1.07, m 1.82, m |
| 8 | 38.7 | | 38.0 | |
| 9 | 60.3 | 1.02, m | 59.5 | 0.82, m |
| 10 | 38.0 | | 37.6 | |
| 11 | 27.4 | 1.81, m 1.58, m | 27.7 | 1.77, m 1.96, m |
| 12 | 78.0 | 3.71, dd (11.4, 4.2) | 77.2 | 3.90, dd (11.0, 3.2) |
| 13 | 44.1 | | 44.2 | |
| 14 | 50.6 | 1.69, m | 50.2 | 2.03, m |
| 15 | 29.0 | 1.88, dd (14.5, 3.9) | 29.4 | 2.15, d (13.3) 1.90, m |
| 16 | 63.4 | 4.62, m | 62.8 | 5.05, brs |
| 17 | 118.5 | | 119.7 | |
| 18 | 162.1 | | 162.1 | |
| 19 | 142.2 | 7.30, d (1.9) | 141.6 | 7.48, brd (0.8) |
| 20 | 110.5 | 6.32, d (1.9) | 111.0 | 6.63, brs |
| 21 | 21.7 | 0.85, s | 21.9 | 0.80, s |
| 22 | 33.8 | 0.87, s | 33.8 | 0.85, s |
| 23 | 16.7 | 0.91, s | 16.7 | 0.81, s |
| 24 | 18.2 | 0.95, s | 18.4 | 0.94, s |
| 25 | 15.9 | 1.14, s | 16.6 | 1.45, s |

^a ^1H and ^{13}C NMR spectra were measured in CD_3OD

^b ^1H and ^{13}C NMR spectra were measured in $\text{C}_5\text{D}_5\text{N}$

Fig. 2. Key COSY and HMBC correlations of **1**

The other purified compounds, namely compounds **2**–**9**, were previously reported to be sesterstatin 4 (**2**) (Pettit et al. 1998), heteronemin (**3**) (Kashman and Rudi 1977), 12-deacetyl-12-*epi*-deoxoscalarin (**4**) (Fontana et al. 1999), 16-hydroxyscalarolide (**5**) (Miyaoaka et al. 2000), 16-*O*-deacetyl-16-*epi*-scalarolbutenolide (**6**) (Ryu et al. 1996), 12-*O*-deacetyl-19-deoxyscalarin (**7**) (Pettit et al. 1998), hyrtiosin A (**8**) (Yu et al. 2005), and hyrtiosal (**9**) (Iguchi et al. 1991) (Fig. 1). These known compounds were confirmed through a comparison of the obtained ^1H NMR

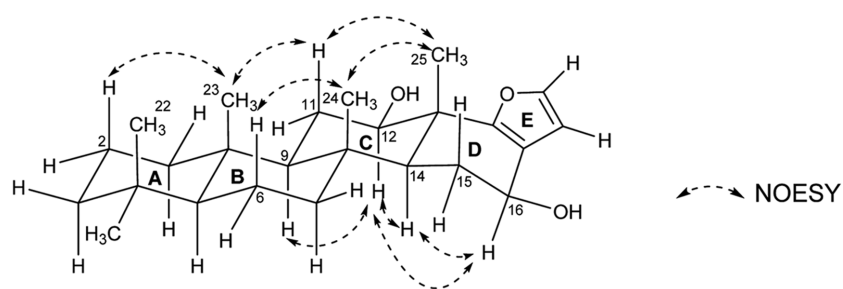


Fig. 3. Key NOESY correlations of **1**

Table 2. MIC assay against Gram-positive bacteria

| Strains | MIC values ($\mu\text{g/mL}$) | | | | | | | | |
|------------------------------|---------------------------------|----|----|---|----|---|----|-----|-----|
| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
| <i>Bacillus subtilis</i> | 32 | 32 | 2 | - | 32 | - | 16 | 128 | 16 |
| <i>Micrococcus luteus</i> | - | - | 4 | - | - | - | - | - | 64 |
| <i>Staphylococcus aureus</i> | 128 | - | 64 | - | - | - | - | - | 128 |

MIC values are the minimal inhibitory concentration inhibiting cell growth or inducing cell death

and ^{13}C NMR data with those reported in the literatures.

The minimum inhibitory concentration (MIC) assay method was then employed to test the antibacterial activities of nine compounds against three Gram-positive and three Gram-negative bacteria. The majority of compounds were highly active against *Bacillus subtilis* compared with the other bacteria examined. More specifically, compound **3** displayed the strongest activity against *Bacillus subtilis* and *Micrococcus luteus* with MIC values of 2 and 4 $\mu\text{g/mL}$, respectively. In addition, compounds **7** and **9** exhibited MIC values of 16 $\mu\text{g/mL}$ against *Bacillus subtilis*. None of the nine compounds exhibited activity against the three Gram-negative bacteria.

4. Conclusions

Chemical analysis of the 1D and 2D NMR spectra recorded for nine scalarane-type sesterterpene derivatives isolated from the Philippines marine sponge *Hyrtios* sp. led to the determination of their chemical structures. The carbon framework was found to belong to the 6/6/6/6/5-pentacyclic ring system-scalarane-type sesterterpenoids class. In addition, compound **1** was identified as a novel 12-epimer exhibiting an α -axial orientation at H-12 position. Known compounds were also isolated, and these were confirmed as sesterstatin 4 (**2**), heteronemin (**3**), 12-deacetyl-12-*epi*-deoxoscalarin (**4**), 16-hydroxyscalarolide (**5**), 16-*O*-deacetyl-16-*epi*-scalarolbutenolide (**6**), 12-*O*-

deacetyl-19-deoxyscalarin (**7**), hyrtiosin A (**8**), and hyrtiosal (**9**). The compounds **1**–**9** were then investigated for their antibacterial activities against three Gram-positive bacteria (*Bacillus subtilis*, *Micrococcus luteus*, and *Staphylococcus aureus*) and three Gram-negative bacteria (*Escherichia coli*, *Salmonella typhimurium*, and *Klebsiella pneumonia* subsp.) using the minimum inhibitory concentration assay. It was found that compound **3** showed potent activities against *Bacillus subtilis* and *Micrococcus luteus*, and the compounds **7** and **9** had moderate to good bioactivities against *Bacillus subtilis*. The other scalarane analogs displayed weak activities against the three Gram-positive bacteria, and none of the compounds exhibited activity against the three Gram-negative bacteria.

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Author's Information

Jae-Hyeong Choi

M.D. Candidate, University of Science and Technology

Hyi-Seung Lee

Principal Research Scientist, Korea Institute of Ocean Science & Technology

Wilfredo L. Campos

Professor, University of the Philippines Visayas

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