



Article **Two New Components from an Association of Marine Sponges** *Poecillastra* sp. and *Jaspis* sp. and Their Inhibitory Effects on **Biomarkers for Benign Prostatic Hyperplasia**

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Abstract: Benign prostatic hyperplasia (BPH), characterized by the enlargement of the prostate gland and subsequent lower urinary tract symptoms, poses a significant health concern for aging men with increasing prevalence. Extensive efforts encompassing in vitro and in vivo models are underway to identify novel and effective agents for the management and treatment of BPH. Research endeavors are primarily channeled toward assessing the potential of compounds to inhibit cell proliferation, curb inflammation, and display anti-androgenic activity. Notably, through screening aimed at inhibiting 5-alpha reductase type 2 (5α R2) in human prostatic cells, two acyl compounds (**1** and **2**) were isolated from a bioactive fraction sourced from an association of marine sponges *Poecillastra* sp. and *Jaspis* sp. The complete structure of **1** was determined as (*Z*)-dec-3-enony (*2S*, *3S*)-capreomycidine, ascertained by JBCA and ECD comparison. While the absolute configurations of **2** remained unassigned, it was identified as a linkage of a 2, *7S**-dihydoxy-9*R**-methyloctadecanoyl group with the 2-amino position of a tramiprosate moiety referred to as homotaurine. Evaluation of both compounds encompassed the assessment of their inhibitory effects on key biomarkers (5α R2, AR, PSA, and PCNA) associated with BPH in testosterone propionate (TP)-activated LNCap and RWPE-1 cells.

Keywords: marine sponge; (2*S*, 3*S*)-capreomycidine; tramiprosate; *J*BCA; ECD comparison; 5-alpha reductase type 2; benign prostatic hyperplasia (BPH)

1. Introduction

Prostatic ailments, including conditions such as prostate cancer and benign prostatic hyperplasia, arise due to the influence of multifarious determinants encompassing age, hormonal profiles, and genetic predisposition. Benign prostatic hyperplasia (BPH), a prevalent medical condition among older males, affects around 60% of individuals in their sixth decade of life. BPH involves the non-malignant expansion of the prostate gland, leading to its enlargement and causing lower urinary tract symptoms (LUTS), as well as obstructive manifestations at the bladder outlet (BOO). While BPH is relatively uncommon prior to the fourth decade of life, approximately half of men experience symptoms associated with BPH by the age of 50. The incidence of BPH increased by 10% per decade, reaching an estimated prevalence of 80% in the octogenarian phase [1,2]. Despite exhaustive research efforts, certain etiological facets of BPH remain concealed. Among the predominant hypotheses, the interplay of age-related changes and disruptions in hormonal balance stands as the principal foundation in the pathogenesis [3].

Diverse botanical treatments are utilized in both traditional and alternative medical practices to alleviate symptoms related to the lower urinary tract. Among these botanical



Citation: Hwang, B.S.; Lee, S.; Jeong, E.J.; Rho, J.-R. Two New Components from an Association of Marine Sponges *Poecillastra* sp. and *Jaspis* sp. and Their Inhibitory Effects on Biomarkers for Benign Prostatic Hyperplasia. *Mar. Drugs* **2023**, *21*, 491. https://doi.org/10.3390/md21090491

Academic Editor: Yue-Wei Guo

Received: 31 August 2023 Revised: 12 September 2023 Accepted: 12 September 2023 Published: 14 September 2023



Copyright: © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). interventions, *Serenoa repens* (SR), *Pygeum africanum*, and *Urtica dioica* stand out prominently [4–6]. Preclinical investigations highlight the multifaceted mechanisms through which the SR extract exerts its effects. These mechanisms include inhibiting the binding of dihydrotestosterone (DHT) receptors within the cytosol of prostatic cells, and reducing the expression and/or activity of 5-alpha reductase, cyclooxygenase, and 5-lipoxygenase. Furthermore, SR induces apoptotic responses in prostatic epithelial cells, demonstrates antiestrogenic attributes, acts as a spasmolytic agent by impeding calcium channel activity, and counteracts beta-adrenergic actions [4]. These combined actions have spurred clinical and experimental research into the potential of SR as a therapeutic approach to alleviate symptoms associated with benign prostatic hyperplasia.

Various in vitro and in vivo models have been developed to find promising new substances for managing and treating BPH. Research primarily focuses on evaluating properties like cell proliferation inhibition, anti-inflammatory, anti-androgenic activity, and antioxidant properties. In our ongoing search for new sources of natural products that can effectively manage BPH, we found that an extract prepared from an association of marine sponges *Poecillatra* sp. and *Jaspis* sp. exhibited potent activity on 5-alpha reductase in human prostatic cells, surpassing the efficacy of the most commonly used 5α -reductase inhibitor, finasteride. Marine sponges of the *Poecillatra* and *Jaspis* genera are widely recognized as rich sources of bioactive compounds of diverse skeletal classes [7,8]. The association of *Poecillatra* sp. and *Jaspis* sp. has also yielded novel metabolites, including anti-angiogenic bis(dihydroxystyrenyl) imidazole alkaloids [9], dihydroxystyrenes [10] and cytotoxic bromotyrosine derivatives [11]. By activity-guided fractionation, we successfully isolated two new compounds, designated as 1 and 2, from the association of *Poecillatra* sp. and Jaspis sp. In this report, we provide detailed information about the isolation and structural determination of compounds **1** and **2**, along with their therapeutic effects on the expressions of 5-alpha reductase type 2 ($5\alpha R2$), and rogen receptor (AR), prostate-specific antigen (PSA), and proliferating cell nuclear antigen (PCNA) in testosterone propionate (TP)-activated LNCap and RWPE-1 cells.

2. Results and Discussion

2.1. Structural Determination of Compounds 1 and 2

Compounds **1** and **2** were detected through MS scanning in the negative mode for the bioactive fraction (30% H₂O and 70% MeOH) of the preserved extract from the association of *Poecillastra* sp. and *Jaspis* sp. They were subsequently isolated by using reversed silica HPLC. Both compounds possessed a linear carbon chain. The structures of these isolated compounds were elucidated through a combination of 1D and 2D NMR experiments, along with MS data (Figure 1).



Figure 1. Structures of compounds 1 and 2 isolated from associated sponges.

Compound **1** was identified to have the molecular formula $C_{16}H_{28}N_4O_3$ based on a peak ([M – H][–] = 323.2087, Δ = 0.5 ppm) in the negative HR-ESIMS and the ¹³C NMR spectra. The broad UV absorption band appeared at 215 nm, and the IR spectrum displayed

absorption peaks at 1626 and 3315 cm⁻¹, indicating carbonyl and hydroxy groups. In the ¹H NMR spectrum of **1**, two olefinic protons were identified at $\delta_{\rm H}$ 5.55 and 5.61, along with five protons within the mid-chemical shift range ($\delta_{\rm H}$ 3.08, 3.23, 3.40, 3.75, and 4.39), three shielded protons ($\delta_{\rm H}$ 1.75, 1.90, and 2.10), several aliphatic protons in the range of $\delta_{\rm H}$ 1.26 to 1.36, and one methyl at $\delta_{\rm H}$ 0.89. The ¹³C NMR spectrum revealed sixteen resonances, which included two carbonyl carbons ($\delta_{\rm C}$ 174.5 and 174.7), two olefinic carbons ($\delta_{\rm C}$ 122.8 and 135.1), one non-protonated carbon ($\delta_{\rm C}$ 155.5), two methines ($\delta_{\rm C}$ 57.6 and 53.6), eight methylenes ($\delta_{\rm C}$ 23.2, 23.7, 28.4, 30.1, 30.5, 32.9, 35.6, and 38.4), and one methyl ($\delta_{\rm C}$ 14.4), as confirmed by the HSQC spectrum.

The analysis of COSY and TOCSY spectra led to the identification of two fragments, as shown in Figure 2A. One of these fragments was deduced to be a linear hydrocarbon chain. The second fragment contained four protons with chemical shift values within the mid-range, which were indicative of nitrogen-bearing carbons. Among these protons, H-3 and H-5 exhibited common correlations with a non-protonated carbon at $\delta_{\rm C}$ 155.5 (C-6) in the HMBC spectrum, suggesting the presence of a pyrimidine moiety. Furthermore, the proton at $\delta_{\rm H}$ 4.39 (H-2) showed a broad cross-peak in the HMBC spectrum, which implies correlations with two nearby carbonyl carbons ($\delta_{\rm C}$ 174.5 and 174.7). Based on the HMBC correlation between $\delta_{\rm H}$ 3.75 and $\delta_{\rm C}$ 174.5, it was deduced that the proton at $\delta_{\rm H}$ 4.39 (H-2) was connected to the carboxylic carbon at $\delta_{\rm C}$ 174.5 (C-1) through a two-bond linkage. The existence of a carboxylic group was supported by a characteristic band (1626 and 3315 cm⁻¹) observed in the IR spectrum. Furthermore, H-2 revealed a connection to the carbonyl carbon in the amide group. The established partial structure was identified as a cyclic arginine-derived non-proteinogenic amino acid. The proton at $\delta_{\rm H}$ 3.08 (H-2') was determined to be connected to the carbonyl carbon at δ_C 174.7, confirmed by the HMBC correlation of H-2'/C-1'. The position of the double bond was supported by the HMBC correlations of H-3'/C-2', H-4'/C-2', H-4'/C-5' and H-4'/C-6'. The geometry of the double bond in the chain was confirmed as the Z-form by the observed NOE correlation between H-2' and H-5'. Lastly, considering the molecular formula, a hexane chain was identified as linked to the olefinic group. Consequently, the structure of 1 was elucidated as featuring a dec-3-enoyl group attached to the 2-amino position of the cyclic arginine-derived nonproteinogenic amino acid.



Figure 2. Key COSY and HMBC correlations of compounds 1 (A) and 2 (B).

The configuration of the two chiral centers within **1** was achieved through *J*BCA and ECD comparisons. Initially, the relative configurations of C-2 and C-3 were established using the homo-/heteronuclear coupling constants measured by the ¹H and HETLOC NMR spectra presented in Figure 3. The moderate homonuclear coupling constant value (${}^{3}J_{HH} = 6.4 \text{ Hz}$) suggested the presence of exchangeable rotation around the C-2 and C-3 bond, while the large absolute heteronuclear coupling value (${}^{2}J_{CH} = -5.3 \text{ Hz}$) between H-3 and C-2 indicated a *gauche* relationship between H-3 and the amine group on C-2. The small absolute value between H-2 and C-3 indicated a portion of the *anti*-relationship

between H-2 and the amine group on C-3. Consequently, the configurations of C-2 and C-3 were identified as $2S^*$ and $3S^*$, respectively. The absolute configuration of 1 was established by comparing the measured and calculated ECD spectra. To save time, the ECD spectrum was calculated for a partial structure truncated at C-5'. As depicted in Figure 4, the calculated ECD spectrum for the 2S3S isomer closely matched the experimental one. The cyclic arginine-derived non-proteinogenic amino acid in 1 was recognized as (2*S*, 3*S*)-capreomycidine (epicapreomycidine) [12].



Figure 3. Relative configurations of C-2 and C-3 in 1 based on JBCA.



Figure 4. Comparison of experimental and calculated ECD spectra of 1.

Compound **2** was determined to have the molecular formula $C_{22}H_{45}NO_6S$ based on the negative HR ESIMS and the ¹³C NMR spectrum, indicating one unsaturation degree. The IR spectrum showed characteristic peaks at 1188 and 1061 cm⁻¹, corresponding to a sulfate or sulfonic acid group. The MS/MS data for **2** revealed three major fragments m/z80, 138, and 166. The IR peaks and molecular fragments indicated the presence of a sulfonic acid. The ¹H NMR spectrum displayed shielded signals for a linear carbon chain and four signals (δ_H 2.82, 3.33, 3.60, and 3.98) in the middle spectral region. The ¹³C NMR spectrum contained two methyl groups, sixteen methylenes, two oxymethines, one methine carbon, and one carbonyl carbon. Based on the provided information, **2** was deduced to possess a linear carbon chain structure with a sulfonic acid moiety.

The COSY and TOCSY spectra provided insight into the structural components of two carbon moieties, as shown in Figure 2B: a propane unit and a linear carbon chain from well-defined proton connections. The chemical shifts of the terminal protons and carbons in the propane segment suggested the presence of amine and sulfonic acid groups. This deduction was corroborated by the mass fragment at m/z 138, confirming the identity as 3-aminopropane-1-sulfonic acid (3-APS) (Figure S9). Notably, this moiety was also recognized as tramiprosate (homotaurine), a compound under development as a potential treatment for Alzheimer's disease [13]. The linkage of this moiety to carbonyl carbon at $\delta_{\rm C}$ 177.6 (C-1') was established through an HMBC correlation, and the oxymethine proton at

 $\delta_{\rm H}$ 3.98 (C-2') showed connectivity to the same carbonyl carbon. As shown in Figure 2B, the presence of the 2-hydroxy-4-methylpentane unit was revealed by the COSY and HMBC correlations. The position of the unit in the linear carbon chain was determined by the HMBC correlations of H-6'/C-4' and H-6'/C-5' and supported by the mutual correlations of both H-2' and H-7' with C-4', C-5', and C-6' in the HSQC-TOCSY spectrum. Based on the molecular formula, the COSY and HMBC correlations, the remaining portion, excluding 3-APS, was identified as a 2, 7-dihydroxy-9-methyloctadecanoyl group. Unfortunately, the Mosher reaction for determining the configuration of C-2' and C-7' was unsuccessful. However, the relative configurations of C-7' and C-9' were deduced through J configuration analysis (JBCA), as depicted in Figure 5. The homo-/heteronuclear coupling constants were measured by DQFCOSY and HECADE spectra. Utilizing the non-equivalent proton positions on C-8', the relative configurations of C-7' and C-9' were determined as $7S^*$ and $9R^*$, respectively, based on heteronuclear coupling constants. Further support for this assignment arose from observing NOE cross-peaks of H-7'/H-10', H-7'/H-19', and H-8'a/H-19' in the NOESY spectrum. Thus, compound **2** was elucidated to be $2',7'(S^*)$ dihydroxy-9'(R*)-methyloctadecanoyl tramiprosate.



Figure 5. Relative configurations of C-7' and C-9' in 2 based on JBCA.

2.2. Biological Activity of Compounds 1 and 2

In order to access the therapeutic effects of **1** and **2** on BPH regulation, we measured the expressions of 5α R2, AR, PSA, and PCNA in LNCap and RWPE-1 cells. Before evaluating the activities of **1** and **2** in these prostate cell lines, we determine the cytotoxicity of the compounds using a CCK-8 assay. LNCap and RWPE-1 cells were exposed to **1** or **2** for 24 h, and cell viability was measured. As demonstrated in Figure 6, compounds **1** and **2** displayed negligible cytotoxicity against LNCap cells in a concentration range of up to 10 μ M (cell viability > 98% of control). Compound **2** exhibited slight cytotoxicity against RWPE-1 cells at concentrations of 1 and 10 μ M. Based on the MTT assay results, we expect that compound **2** may exhibit anti-proliferative activity. However, in this study, we did not perform additional experiments to precisely quantify this anti-proliferative effect. Although a mild decrease in cell viability was observed at a concentration of 10 μ M for **2** in RWPE-1 cells, the reduction was less than 10%. Importantly, no toxicity was observed in LNCap cells. Consequently, concentrations of 1 and 10 μ M were chosen for evaluating the effects of these compounds in both cell types.



Figure 6. Cytotoxicity of **1** and **2** in LNCap (**A**) cells and RWPE-1 cells (**B**). Cells were treated with **1** or **2** (0.1, 1, 5, and 10 μ M) for 24 h. Cell viability was determined by MTT assay. Results are presented as mean \pm standard deviation (n = 3) * p < 0.05, *** p < 0.001 compared to non-treated controls.

To evaluate the potential impact of **1** and **2** on BPH, we measured their inhibitory effects on the expression of 5α -reductase type 2 (5α R2). As shown in Figure 7, the expression of 5α R2 was induced by the treatment of TP in LNCap and RWPE-1 cells, and the pretreatment of cells with **1** or **2** (1 and 10 μ M) significantly suppressed the 5α R2 expression. Particularly, compound **2** exhibited a more pronounced decrease in 5α R2 levels at the same concentration (Figure 7A,B). At the concentration of 10 μ M, compound **2** decreased the expression level of 5α R2 protein induced by TP to 42% and 48% of TP-only treated cells in LNCap and RWPE-1 cells, respectively.

The transformation of testosterone into DHT, facilitated by 5α R2 in androgen-responsive target cells, is a fundamental process implicated in the development of BPH [14]. It is widely recognized that inhibitors of 5α R2, such as finasteride (Fina) or dutasteride, can impede BPH progress by suppressing DHT synthesis [14,15]. Additionally, 5α R2 is recognized as the primary target protein through which SR ameliorates symptoms related to BPH. Recent meta-analyses have demonstrated that the efficacy of SR is comparable to that of finasteride and tamsulosin, and notably superior to that of a placebo in treating mild to moderate lower urinary tract symptoms (LUTS) [16].

To assess the potential therapeutic effects of **1** and **2** on AR in prostate cells, we employed Western blot analysis to evaluate the levels of AR expression. As illustrated in Figure 8, the treatment with TP resulted in an increase in AR expression in both LNCap and RWPE-1 cells. In cells treated with **2**, a slight decrease in AR expression was observable in both cell lines, whereas the reduction induced by **1** did not reach statistical significance. Specifically, the expression of AR was reduced by **2** (10 μ M) to 72% and 76% of the levels in TP-treated cells in LNCap and RWPE-1 cells, respectively.



Figure 7. Inhibitory effects of **1** and **2** on the expressions of 5α R2 in LNCap cells (**A**) and in RWPE-1 cells (**B**). Cells were treated with testosterone propionate (TP, $0.5 \,\mu$ M) and **1** or **2** (1 and 10 μ M) for 24 h. The expression levels of 5 α R2 were analyzed by Western blotting. Results are presented as mean \pm standard deviation (n = 3); p < 0.01 compared to the non-treated control (NC), * p < 0.05, ** p < 0.01 and *** p < 0.001 compared to TP-treated cells.



Figure 8. Inhibitory effects of 1 and 2 on the expressions of AR in LNCap cells (A) and in RWPE-1 cells (**B**). Cells were treated with testosterone propionate (TP, $0.5 \,\mu$ M) and **1** or **2** (1 and 10 μ M) for 24 h. The expression levels of AR were analyzed by Western blotting. Results are presented as mean \pm standard deviation (*n* = 3); *p* < 0.01 compared to the non-treated control (NC), * *p* < 0.05, ** *p* < 0.01 compared to TP-treated cells.

Androgens and androgen receptors (AR), a member of the steroid receptor superfamily, are widely recognized as playing a significant role in the development of BPH [17]. In epithelial cells, AR is thought to contribute to BPH progression by orchestrating interactions between epithelial and stromal cells. These interactions involve changes in epithelial-mesenchymal transition, leading to subsequent stromal cell proliferation [18,19]. While blocking the androgen/AR signaling pathway has been shown to decrease BPH

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volume and alleviate lower urinary tract symptoms, the precise mechanisms through which androgen/AR signaling influences BPH development remain unclear [20,21].

Upon androgen activation, the androgen-activated AR dissociates from chaperones and translocates into the nucleus. The interaction of AR with androgen response elements (AREs) is recognized to trigger the expression of genes such as PSA and PCNA [22,23]. PSA is recognized as a critical androgen-regulated gene and acts as a specific marker for detecting prostate cancer, rendering it valuable for prostate cancer screening and evaluation. In LNCap cells stimulated by TP (Figure 9A), the application of **1** and **2** significantly reduced the expression of PSA at a concentration of 10 μ M. The expression level of the PSA protein diminished to 78% and 47% of TP-only treated cells by **1** and **2**, respectively. In TP-induced RWPE-1 cells (Figure 9B), the expression of PCNA was attenuated by **2** at a concentration of 10 μ M, while the induction of PCNA expression was observed in cells treated with **1**.



Figure 9. Inhibitory effects of **1** and **2** on the expressions of PSA in LNCap cells (**A**) and PCNA in RWPE-1 cells (**B**). Cells were treated with testosterone propionate (TP, 0.5 μ M) and **1** or **2** (1 and 10 μ M) for 24 h. The expression levels of PSA or PCNA were analyzed by Western blotting. Results are presented as mean \pm standard deviation (n = 3); p < 0.01 compared to the non-treated control (NC), * p < 0.05, ** p < 0.01 compared to TP-treated cells.

3. Materials and Methods

3.1. General Experiment Procedures

Optical rotations were measured on a JASCO P-1010 polarimeter with a 1 cm cell and circular dichroism spectrum was recorded on a JASCO J-1500 CD spectrometer (Jasco Corporation, Tokyo, Japan). The UV and IR spectra were acquired on a Varian Cary 50 and JASCO FT/IR 4100 spectrometers, respectively. The NMR spectra were measured on a Varian VNMRS 500 MHz spectrometer (Varian, Palo Alto, CA, USA) with a 3 mm ID probe in MeOH- d_4 solvent, which was referenced by residual solvent peaks at δ_H 3.30 and δ_C 49.0). The high-resolution ESI mass spectrum was acquired by using a SCIEX X500R (Sciex Co., Framingham, MA, USA). The HPLC was performed using an Agilent 1200 system (Santa Clara, CA, USA) using Phenomenex polar C18 and YMC ODS-A columns and an ELSD detector. Quantum calculations were conducted by Dell PowerEdge R740 server (Dell, Round Rocks, TX, USA) installed Gaussian 16 (Gaussian Inc., Wallingford, CT, USA) and Spartan 20 (Wavefunction Inc., Irvine, CA, USA) software.

3.2. Animal Material

A specimen of a mutualistic association of Poecillastra sp. and Jaspis sp. (No. 08K-3) was obtained through scuba diving at a depth of 20 m off the coast of Keomun Island,

В.

Korea in July 2008. The identification of the sample was confirmed by Professor Chung Ja Sim from Hannam University, Korea. Following collection, a methanolic extract of the specimen was stored in a refrigerator at -28 °C for research purposes.

3.3. Isolation of Compounds 1 and 2

The stored extract dissolved in MeOH was dried in vacuo and was partitioned between dichloromethane and distilled water. The organic layer was repartitioned with a mixture of n-hexane and 15% aqueous MeOH. Then, the aqueous MeOH layer was subjected to reversed-phase silica gel flash column chromatography eluting with solvents of decreasing polarity (MeOH/H₂O = 5/5; 6/4; 7/3; 8/2; 9/1; 100% MeOH; 100% acetone) to give seven fractions (MR1~MR7). Guided by DHT inhibition assay, fraction MR3 (30 mg) was selected for further research. Compounds 1 and 2 were isolated by reversed-phase silica HPLC eluting with a gradient solvent system (A: H₂O with 0.1% formic acid, B: 100% acetonitrile, from 20% to 100% B for 30 min) at retention times of 14.6 and 24 min, respectively. HPLC conditions consisted of a Phenomenex polar-C18 column 150 x 4.6 mm, 4 us, 1 mL/min, ELSD detector. Compound 1 was purified by using YMC ODS-A column under isocratic solvents of 45% ACN and 55% H₂O to yield 2.5 mg. Compound 2 was purified by a solvent system of 60% MeOH and 40% H₂O to yield 3.0 mg.

Compound (1). Amorphous oil. $[\alpha]_D^{25}$ + 18.4 (*c* 0.2, MeOH). UV (MeOH) λ_{max} (log ε): 215 (3.1) nm. IR (film) ν_{max} : 3315, 2924, 1626, 1464, 1378 cm⁻¹. ¹H (500 MHz) and ¹³C (125 MHz) NMR data, see Table 1. HR-ESI MS (negative-ion mode) *m*/*z*: 323.2087 [M–H]⁻ (calcd for C₁₆H₂₈N₄O₃, 323.2089).

No	δ_{C} , Mult	$\delta_{\rm H}$, (J Hz)	COSY	НМВС
1	174.5, C			
2	57.6, CH	4.39, d(6.4)	H-3	C-1, C-3, C-4, C-1'
3	53.6, CH	3.75, ddd(9.3, 6.4, 4.2)	H-2, H-4	C-1, C-2, C-4, C-5, C-6
4	23.2, CH ₂	1.75, m; 1.90, m	H-3, H-5	C-2, C-3, C-4
5	38.4, CH ₂	3.23, ddd(12.7, 10.3, 4.2)	H-4	C-3, C-4, C-6
		3.40, dt(12.7, 4.65)		
6	155.5, C			
1'	174.7, C			
2'	35.6, CH ₂	3.08, d(7.1)	H-3′	C-1′, C-3′, C-4′
3'	122.8, CH	5.55, m	H-2′, H-4′	C-2′
4'	135.1, CH	5.61, m	H-3′, H-5′	C-2′, C-5′, C-6′
5'	28.4, CH ₂	2.10, dt(7.1, 6.9)	H-4′, H-6′	C-3′, C-4′, C-6′
6'	30.5, CH ₂	1.36, m		C-4′, C-7′
7'	30.1, CH ₂	1.26~1.34		
8'	32.9, CH ₂	1.26~1.34		
9′	23.7, CH ₂	1.26~1.34	H-10′	
10′	14.4, CH ₃	0.89, t(7.1)	H-9′	C-8′, C-9′

Table 1. The spectral data for 1 in CD₃OD (¹H for 500MHz, ¹³C for 125 MHz).

Compound (2). Amorphous oil. $[\alpha]_D^{25} - 5.5$ (*c* 0.1, MeOH). IR (film) ν_{max} : 3348, 2924, 1630, 1463, 1188, 1061 cm⁻¹. ¹H (500 MHz) and ¹³C (125 MHz) NMR data, see Table 2. HR-ESI MS (negative-ion mode) m/z: 450.2887 [M–H]⁻ (calcd for C₂₂H₄₅NO₆S, 450.2895).

3.4. ECD Calculation of Compound 1

For the purpose of reducing the calculation time, the structure of compound 1 truncated at C-5' was used. The conformational searches of the two isomers (2S3S and 2R3R) for the truncated 1 were performed by Spartan 20 software (Wavefunction Inc., Irvine, CA, USA) following a protocol reported in the literature [24]. The selected conformers were reoptimized at the m062X/6-31G(d,p) level with CH₃OH solvent by using Gaussian 16 software (Gaussian inc., Wallingford, CT, USA). The ECD spectra of the conformers corresponding to the two isomers were calculated to the B3LYP//6-31G(d,p) level. A Boltzmann distribution for the optimized conformers was used for calculated ECD spectra.

No	$\delta_{\rm C}$, Mult	$\delta_{\rm H}$, (J Hz)	COSY	НМВС
1	38.8, CH ₂	3.33, m	H-2	C-1′, C-2, C-3
2	26.3, CH ₂	1.97, quint(7.8)	H-1, H-3	C-1, C-3
3	50.0, CH ₂	2.82, t(7.8)	H-2	C-1, C-2
1'	177.6, C			
2'	72.8, CH	3.98, dd(7.8, 3.9)	H-3′	C-1′, C-3′, C-4′
3'	35.6, CH ₂	^a 1.57, m; ^b 1.74, m	H-2′, H-4′	C-1′, C-2′
4'	26.2, CH ₂	1.43, m	H-3′	C-4′, C-5′, C-7′
5'	30.6, CH ₂	1.33, m	H-7′	
6'	38.7, CH ₂	^a 1.36, dt(7.1, 6.9); ^b 1.43, m	H-8′, H-6′	C-6′, C-7′, C-9′, C-19′
7'	70.3, CH	3.60, m	H-7′, H-9′	C-10′
8'	46.2, CH ₂	^a 1.27, m; ^b 1.34, m	H-8′, H-10′, H-19′	C-11′
9'	30.6, CH	1.60, m	H-9′	
10'	37.7, CH ₂	^a 1.08, m; ^b 1.37, m		
11'	28.0, CH ₂	^a 1.27, m; ^b 1.35, m		
12'	26.6, CH ₂	^a 1.35, m; ^b 1.45, m		
13'	31.1, CH ₂ *	1.27~1.34, m 1.27~1.34, m		
14'	30.8, CH ₂ *	1.27~1.34, m		
15'	30.5, CH ₂ *	1.28, m		
16'	33.1, CH ₂	1.31, m	H-18′	C-16′, C-17′
17'	23.8, CH ₂	0.90, t(6.9)	H-17′	C-8′, C-9′, C-10′
18'	14.5, CH ₃	0.89, d(6.9)	H-9′	
19'	20.8, CH ₃			

Table 2. The spectral data for 2 in CD₃OD (¹H for 500MHz, ¹³C for 125 MHz).

* Exchangeable carbons, ^a shielded proton, ^b deshielded proton.

3.5. Cell Cultures

The RWPE-1 cell line (normal human prostatic epithelial cells) and LNCap cell line (human prostatic adenocarcinoma cells) were obtained from the American Type Culture Collection. RWPE-1 cells were cultured in keratinocyte serum-free medium supplemented with 0.05 mg mL⁻¹ bovine pituitary extract, 5 ng mL⁻¹ epidermal growth factor, and 1% (v/v) antibiotics (100 U mL⁻¹ penicillin and 100 µg mL⁻¹ streptomycin). LNCap cells were cultured in RPMI1640 containing 10% (v/v) fetal bovine serum and 1% (v/v) antibiotics (100 U mL⁻¹ penicillin and 100 µg mL⁻¹ streptomycin).

3.6. Cell Viability Assay

Cell viability was assessed using the MTT colorimetric assay, which measures cellular dehydrogenase activity by reducing MTT to formazan. RWPE-1 and LNCaP cells were seeded at densities of 3×10^4 and 1×10^4 cells well⁻¹, respectively, in 96-well plates and incubated for 24 h. Cells were then exposed to different concentrations (0.1, 1, and 10 μ M) of compounds **1** and **2** for 24 h. MTT (2 mg mL⁻¹) in distilled water was added to each well and incubated at 37 °C for 1 h. The formazan precipitate was dissolved in DMSO, and absorbance was measured at 550 nm using a microplate reader.

3.7. Western Blotting Analysis

RWPE-1 and LNCaP cells were seeded at 2×10^5 and 6×10^5 cells/well, respectively, in 6-well plates and incubated overnight. Cells were treated with TP (0.5 µM) for 1 h, followed by treatment with finasteride (10 µM) or compounds **1** or **2** at different concentrations (1 and 10 µM) for 24 h. Cells were washed with cold PBS, and cell lysates were prepared using lysis buffer containing a protease inhibitor cocktail. Protein content was quantified using the Bradford assay. Protein samples (30 µg) were separated by SDS-PAGE and transferred to a membrane. The membrane was blocked with skim milk and incubated with primary antibodies against 5 α R2, AR, PSA, PCNA, and β -actin. After washing, immunoreactive bands were detected using peroxidase-conjugated secondary antibodies. Protein bands were visualized using ECL substrate and an imaging system.

4. Conclusions

The two new compounds **1** and **2** were isolated from a bioactive fraction of an association of two marine sponges of *Poecillastra* sp. and *Jaspis* sp. The compounds are commonly characteristic of the attachment of acyl chains on a zwitterionic moiety. The complete structure of **1** was established as (*Z*)-dec-3-enoyl (2*S*,3*S*)-epicapreomycidine based on the *J*BCA and ECD comparison. Although the absolute configurations of **2** were not assigned, the compound was revealed as a linkage of 2, *7S**-dihydroxy-9*R**-methyloctadecanoyl group to the position of 2-amino of a tramiprosate moiety called homotaurine. The two compounds exhibited inhibitory effects of biomarkers (5 α R2, AR, PSA, and PCNA) related to BPH on testosterone propionate (TP)-activated LNCap and RWPE-1 cells. In particular, compound **2** exhibited significant suppression of 5 α R2 expression in two cell types, and it was superior to finasteride, a selective inhibitor of 5 α R2. Additionally, considering the inhibition of the expression of other biomarkers closely related to BPH progression, these findings support the potential use of compound **2** as a therapeutic agent for the treatment of BPH.

Supplementary Materials: The following supporting information can be downloaded at: https: //www.mdpi.com/article/10.3390/md21090491/s1. Figure S1: HRESIMS of **1**; Figure S2–S8: 1D and 2D NMR spectra of **1**, Figure S9: HRESIMS of **2**; Figure S10–S17: 1D and 2D NMR spectra of **2**.

Author Contributions: B.S.H. performed isolation and determined the chemical structures; S.L. recorded spectroscopic data; E.J.J. performed in vitro experiments and analyzed the data; E.J.J. and J.-R.R. wrote the paper; all contributing authors have read and approved the final version of the manuscript. All authors have read and agreed to the published version of the manuscript.

Funding: This research was supported by the Korea Institute of Marine Science and Technology (KIMST) and funded by the Ministry of Oceans and Fisheries (RS-2023-00256330, Development of risk managing technology tackling ocean and fisheries crisis around Korean Peninsula by Kuroshio Current).

Institutional Review Board Statement: Not applicable.

Data Availability Statement: All data are contained within this article and Supplementary Materials.

Acknowledgments: The authors would like to thank Chung Ja Sim for identifying the sponge for research.

Conflicts of Interest: The authors declare no conflict of interest.

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