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Article

Cyclic Bis-1,3-dialkylpyridiniums from the Sponge Haliclona sp.

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Abstract: Eight novel cyclic bis-1,3-dialkylpyridiniums, as well as two known compounds from the cyclostellettamine class, were isolated from the sponge *Haliclona* sp. from Korea. Structures of these novel compounds were determined using combined NMR and FAB-MS/MS analyses. Several of these compounds exhibited moderate cytotoxic and antibacterial activities against A549 cell-line and Gram-positive strains, respectively. The structure-activity relationships of cyclostellettamines are discussed based on their bioactivities.

Keywords: *Haliclona* sp.; bis-1,3-dialkylpyridiniums; cyclostellettamines; cytotoxicity; antimicrobial activity

1. Introduction

The 1,3-dialkylpyridiniums (with both cyclic and linear frameworks) are widely distributed cytotoxic compounds that play a role in the antimicrobial properties of marine sponges of the order Haplosclerida (genera *Amphimedon*, *Haliclona*, and *Xestospongia*) [1–4]. Previous studies have evaluated 1,3-dialkylpyridinium inhibition of histone deacetylase (HDAC) [5,6] and epidermal growth

factor (EGF) receptor-mediated mitogenesis [7]. These compounds are also known to block the interaction of chemokines [8] and the binding of methyl quinuclidinyl benzilate (QNB) to the muscarinic receptor [9,10]. In addition to their chemical and bioactivity properties, these metabolites are considered to be the biogenetic precursors of a wide variety of compounds with complicated carbon skeletons and diverse functionalities, such as halicyclamines, ingenamines, manzamines, and sarains [2].

When evaluating bioactive metabolites from Korean marine invertebrates, we recently identified haliclonin A, a cytotoxic macrocylic bisamide from the sponge *Haliclona* sp. [11]. The novel carbon skeleton of this compound was highly reminiscent of bis-1,3-dialkylpyridiniums, especially cyclostellettamines isolated from *Haliclona* sp. [9,12]. Thus, we searched for similar 1,3-dialkylpyridinium metabolites in the polar fractions of extracts from diverse collections of this sponge. Here, we report eight novel derivatives of cyclostellettamines with diverse alkyl chains. Several of these compounds exhibited moderate cytotoxicity against the lung cancer A549 cell-line and antibacterial activities against Gram-positive strains. Additional information on their structure-activity relationships were deduced based on their cytotoxicity and antibacterial activity.

2. Results and Discussion

The sponge *Haliclona* sp. specimens were lyophilized, macerated, and repeatedly extracted with CH_2Cl_2 and MeOH. The combined extracts were separated by solvent-partitioning followed by ODS vacuum flash chromatography. The highly polar H_2O -MeOH (50:50) chromatographic fraction was further fractionated by Sephadex LH-20 gel-permeation chromatography and separated by ODS-HPLC to yield 10 compounds (Figure 1). The structures of compounds **1** and **2** were identified as cyclostellettamine N and Q, respectively, based on spectroscopic analyses and a comparison of NMR and FAB-MS/MS data [13,14].

Figure 1. Structures of compounds 1–10.



The molecular formula of compound **3**, with the shortest retention time in reversed-phase HPLC, was $C_{28}H_{42}Cl_2N_2$ based on the quasi-molecular ion clusters at m/z 407.3428 $[M - H]^+$ and 443.3195 $[M + Cl]^+$ in the HR-FAB-MS data. The spectroscopic data of this compound were very similar to those of **1** and **2** (Tables 1 and 2).

Table 1. ¹³C NMR (ppm, mult) assignments for compounds 3-10 in MeOH- d_4 .

Position	3	4	5	6	7	8	9	10
2	145.5, CH	145.6, CH	145.3, CH	145.3, CH	145.2, CH	145.6, CH	145.2, CH	145.3, CH
3	145.3, C	145.3, C	145.7, C	145.5, C	146.7, C	145.3, C	145.7, C	145.5, C
4	147.0, CH	146.9, CH	146.9, CH	147.0, CH	146.9, CH	146.9, CH	146.9, CH	147.0, CH
5	129.2, CH	129.2, CH	129.2, CH	128.8, CH	129.2, CH	129.1, CH	129.1, CH	129.2, CH
6	143.4, CH	143.4, CH	143.4, CH	143.5, CH	143.5, CH	143.4, CH	143.5, CH	143.4, CH
7	$62.7,\ CH_2$	$62.8,\ CH_2$	62.8, CH ₂	62.8, CH ₂	62.9, CH ₂	$62.9,\ CH_2$	62.9, CH ₂	62.8, CH ₂
8	$32.2,\ CH_2$	$32.1,\ CH_2$	32.2, CH ₂	$32.3, CH_2$	24.8, CH ₂	$32.3,\ CH_2$	32.3, CH ₂	32.1, CH ₂
9	$26.7,\ CH_2$	$26.5,\ CH_2$	26.2, CH ₂	$26.7, CH_2$	137.6, CH	$26.9,\ CH_2$	26.8, CH ₂	26.8, CH ₂
10	29.5 ^{<i>a</i>} , CH ₂	29.6 ^{<i>b</i>} , CH ₂	29.6, CH ₂	30.0 ^{<i>d</i>} , CH ₂	124.7, CH	29.7 ^{<i>f</i>} , CH ₂	$30.5, CH_2$	30.1 ^{<i>h</i>} , CH ₂
11	29.8 ^a , CH ₂	30.0 ^{<i>b</i>} , CH ₂	32.7, CH ₂	30.2, CH ₂	28.1, CH ₂	30.3 ^{<i>f</i>} , CH ₂	$30.7, CH_2$	30.2 ^{<i>h</i>} , CH ₂
12	29.9 ^{<i>a</i>} , CH ₂	30.0 ^b , CH ₂	132.3, CH	28.0, CH ₂	30.5, CH ₂	30.1 ^{<i>f</i>} , CH ₂	28.0, CH ₂	30.2 ^{<i>h</i>} , CH ₂
13	30.2 ^{<i>a</i>} , CH ₂	30.1 ^b , CH ₂	130.7, CH	133.3, CH	29.2 ^e , CH ₂	$30.9, CH_2$	131.1, CH	30.7, CH ₂
14	$29.2,\ CH_2$	29.2, CH ₂	32.4, CH ₂	128.0, CH	29.2 ^e , CH ₂	$28.3, CH_2$	130.4, CH	28.2, CH ₂
15	$31.3, CH_2$	31.2, CH ₂	31.1, CH ₂	29.2, CH ₂	29.6 ^e , CH ₂	135.7, CH	27.9, CH ₂	133.5, CH
16	$33.2, CH_2$	33.1, CH ₂	33.2, CH ₂	33.2, CH ₂	31.1, CH ₂	125.1, CH	29.7, CH ₂	127.8, CH
17	-	-	-	-	33.1, CH ₂	31.0, CH ₂	31.3, CH ₂	29.1, CH ₂
18	-	-	-	-	-	-	33.3, CH	33.2, CH ₂
2'	145.5, CH	145.6, CH	145.3, CH	145.5, CH	145.2, CH	144.6, CH	145.2, CH	145.5, CH
3'	145.4, C	145.3, C	145.5, C	144.9, C	145.7, C	144.7, C	145.7, C	145.0, C
4'	146.9, CH	146.9, CH	146.9, CH	147.2, CH	147.0, CH	146.7, CH	146.9, CH	147.3, CH
5'	129.2, CH	129.2, CH	129.1, CH	129.2, CH	129.2, CH	129.2, CH	129.1, CH	128.9, CH
6'	143.5, CH	143.4, CH	143.5, CH	143.5, CH	143.5, CH	143.7, CH	143.5, CH	143.4, CH
7′	$62.7,\ CH_2$	$62.8,\ CH_2$	62.8, CH ₂	62.9, CH ₂	62.8, CH ₂	62.8, CH ₂	62.8, CH ₂	62.9, CH ₂
8′	$31.9, CH_2$	$32.1,\ CH_2$	32.2, CH ₂	32.4, CH ₂	32.0, CH ₂	$32.2, CH_2$	32.3, CH ₂	32.6, CH ₂
9′	$26.4,\ CH_2$	$26.5,\ CH_2$	26.6, CH ₂	26.8, CH ₂	26.5, CH ₂	26.6, CH ₂	26.7, CH ₂	27.0, CH ₂
10'	30.3 ^a , CH ₂	29.6 ^{<i>b</i>} , CH ₂	29.9 ^c , CH ₂	29.7 ^d , CH ₂	29.7 ^e , CH ₂	29.1 ^{<i>f</i>} , CH ₂	29.6 ^g , CH ₂	30.3 ^h , CH ₂
11'	30.4 ^a , CH ₂	30.0 ^b , CH ₂	29.9 ^c , CH ₂	29.9 ^{<i>d</i>} , CH ₂	30.1 ^e , CH ₂	29.9 ^{<i>f</i>} , CH ₂	29.9 ^g , CH ₂	30.4 ^{<i>h</i>} , CH ₂
12'	29.0, CH ₂	30.0 ^{<i>b</i>} , CH ₂	30.0 ^c , CH ₂	30.0 ^{<i>d</i>} , CH ₂	30.1 ^e , CH ₂	29.9 ^{<i>f</i>} , CH ₂	30.0 ^g , CH ₂	30.5 ^{<i>h</i>} , CH ₂
13'	$30.9, CH_2$	30.1 ^b , CH ₂	30.0 °, CH ₂	30.4 ^d , CH ₂	30.2 ^e , CH ₂	30.0 ^{<i>f</i>} , CH ₂	30.1 ^g , CH ₂	30.7 ^{<i>h</i>} , CH ₂
14′	33.0, CH ₂	29.2, CH ₂	29.2, CH ₂	28.7, CH ₂	29.1, CH ₂	$29.1,\ CH_2$	29.4, CH ₂	29.1, CH ₂
15'	-	31.2, CH ₂	31.2, CH ₂	30.7, CH ₂	30.9, CH ₂	$30.4, CH_2$	31.3, CH ₂	31.1, CH ₂
16′	-	$33.1, CH_2$	33.0, CH ₂	32.9, CH ₂	33.2, CH ₂	33.1, CH ₂	33.3, CH ₂	33.1, CH ₂

^{*a–h*}: Interchangeable signals.

Table 2. H MMR (∂ , mult, J in Hz) assignments for compounds 3–6 in MeOH-	Table 2. ¹ H MMR (δ , mu	J in Hz) assignment	s for compounds 3 -	- 6 in MeOH- <i>d</i> ₄ .
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Position	3	4	5	6
2	8.96 (1H, s)	8.98 (1H, s)	8.95 (1H, s)	8.99 (1H, s)
3	-	-	-	-
4	8.45 (1H, d, 7.6)	8.46 (1H, d, 8.0)	8.46 (1H, d, 8.2)	8.45 (1H, d, 8.2)
5	8.02 (1H, dd, 7.7, 6.2)	8.02 (1H, dd, 7.7, 6.4)	8.02 (1H, dd, 7.4, 6.1)	8.01 (1H, dd, 7.4, 6.6)
6	8.83 (1H, dd, 6.0)	8.85 (1H, dd, 6.0)	8.83 (1H, br s)	8.84 (1H, dd, 6.2)
7	4.64 (1H, t, 6.8)	4.65 (2H, t, 6.8)	4.65 (2H, t, 7.2)	4.64 (2H, t, 6.9)
8	2.00 (2H, quint, 7.3)	2.01 (2H, tt, 7.4, 6.8)	2.03 (2H, m)	2.00 (2H, m)
9	1.26 (2H, m)	1.25 (2H, m)	1.29 (2H, m)	1.23 (2H, m)

10	1.25 ~ 1.40 (2H, m)	1.20 ~ 1.36 (2H, m)	1.42 (2H, m)	1.26 (2H, m)
11	$1.25 \sim 1.40 \; (2H, m)$	1.20 ~ 1.36 (2H, m)	2.01 (2H, m)	1.24 (2H, m)
12	$1.25 \sim 1.40 \; (2H, m)$	1.20 ~ 1.36 (2H, m)	5.41 (1H, br t, 3.6)	1.74 (2H, m)
13	$1.25 \sim 1.40 \; (2H, m)$	1.20 ~ 1.36 (2H, m)	5.41 (1H, br t, 3.6)	5.38 (2H, m)
14	1.30 (2H, m)	1.28 (2H, m)	2.04 (2H, m)	5.38 (2H, m)
15	1.72 (2H, quint, 7.4)	1.74 (2H, tt, 7.3, 7.3)	1.78 (2H, m)	2.46 (2H, dd, 12.9, 6.5)
16	2.89 (2H, 7.3)	2.90 (2H, t, 7.3)	2.88 (2H, t, 7.6)	2.95 (2H, t, 6.9)
17	-	-	-	-
18	-	-	-	-
2′	8.96 (1H, s)	8.98 (1H, s)	8.98 (1H, s)	8.95 (1H, s)
3'	-	-	-	-
4′	8.44 (1H, d, 7.7)	8.46 (1H, d, 8.0)	8.45 (1H, d, 6.8)	8.45 (1H, d, 8.2)
5'	8.02 (1H, dd, 7.7, 6.2)	8.02 (1H, dd, 7.7, 6.4)	8.01 (1H, dd, 7.4, 6.1)	8.02 (1H, dd, 7.4, 6.6)
6′	8.82 (1H, dd, 6.0)	8.85 (1H, dd, 6.0)	8.84 (1H, br s)	8.85 (1H, dd, 6.2)
7′	4.64 (1H, t, 6.5)	4.65 (2H, t, 6.8)	4.63 (2H, t, 6.7)	4.63 (2H, t, 7.5)
8'	2.02 (2H, tt, 7.8, 6.9)	2.01 (2H, tt, 7.4, 6.8)	2.00 (2H, m)	1.98 (2H, m)
9′	1.20 (2H, m)	1.25 (2H, m)	1.28 (2H, m)	1.24 (2H, m)
10′	$1.25 \sim 1.40 \; (2H, m)$	1.20 ~ 1.36 (2H, m)	$1.27 \sim 1.40 \; (2H, m)$	$1.23 \sim 1.35 \; (2H, m)$
11′	$1.25 \sim 1.40 \; (2H, m)$	1.20 ~ 1.36 (2H, m)	1.27 ~ 1.40 (2H, m)	1.23 ~ 1.35 (2H, m)
12′	1.20 (2H, m)	1.20 ~ 1.36 (2H, m)	1.27 ~ 1.40 (2H, m)	1.23 ~ 1.35 (2H, m)
13′	1.76 (2H, tt, 7.4, 7.2)	1.20 ~ 1.36 (2H, m)	1.27 ~ 1.40 (2H, m)	1.23 ~ 1.35 (2H, m)
14′	2.91 (2H, t, 7.0)	1.28 (2H, m)	1.30 (2H, m)	1.25 (2H, m)
15′	-	1.74 (2H, tt, 7.3, 7.3)	1.73 (2H, m)	1.77 (2H, m)
16′	-	2.90 (2H, t, 7.3)	2.89 (2H, t, 7.0)	2.92 (2H, t, 6.9)

Table 2. Cont.

The UV absorption at 267 nm was reminiscent of two di-substituted pyridine rings, which was corroborated by aromatic proton signals [$\delta_{\rm H}$, 8.96 (2 H, br s, H-2, H-2'), 8.45 (1 H, d, J = 7.6 Hz, H-4), 8.44 (1 H, d, J = 7.7 Hz, H-4'), 8.02 (2 H, dd, J = 7.7, 6.2 Hz, H-5, H-5'), 8.83 (1 H, dd, J = 6.0 Hz, H-6), 8.82 (1 H, dd, J = 6.0 Hz, H-6'] and carbon signals [$\delta_{\rm C}$ 145.5 (C-2, C-2'), $\delta_{\rm C}$ 145.3 (C-3), $\delta_{\rm C}$ 145.4 (C-3'), $\delta_{\rm C}$ 147.0 (C-4), $\delta_{\rm C}$ 146.9 (C-4'), $\delta_{\rm C}$ 129.2 (C-5, C-5'), $\delta_{\rm C}$ 143.4 (C-6), $\delta_{\rm C}$ 143.5 (C-6')] in the NMR data. Although the two and three linearly contiguous methylene units attached to the pyridinium (N-1, N-1', C-3, and C-3') were identified based on ¹H COSY and gHMBC, only a large methylene envelope at $\delta_{\rm H}$ 1.25–1.40 was observed in the ¹H NMR spectrum, indicating that **3** was composed of two di-substituted pyridine rings and long aliphatic chains, forming a highly symmetric cyclic framework. The lengths of the methylene chains were determined based on the FAB-MS/MS data, which yielded two intense peaks at m/z 190 and 218, as shown in Figure 2. Since these daughter ions were attributable to a pair of monomeric products generated via Hoffmann-type elimination⁷ from their parent ion, **3** was structurally defined to be a cyclic hetero-dimer composed of C₈ and C₁₀ chains connecting the N-1 and C-3' (also N-1' and C-3) of pyridiniums.



Figure 2. Positive FAB-MS/MS spectrum of compound 3.

The molecular formula of compound **4** was $C_{30}H_{46}Cl_2N_2$ based on HR-FAB-MS data with ion clusters at m/z 435.3741 [M – H]⁺ and 471.3503 [M + Cl]⁺. Thus, the structure consisted of two CH₂ units larger than **3**. The NMR data of this compound were very similar to those of **3**, indicative of the same cycloalkylpyridinium nature. The FAB-MS/MS spectrum exhibited two intense daughter ions at m/z 218, confirming the symmetrical structure of **4**, which contained two C₁₀ linear alkyl chains connecting the N-1 and C-3 between two pyridiniums.

The molecular formulas of compounds **5** and **6** were both $C_{30}H_{44}Cl_2N_2$ based on HR-FAB-MS analyses. The ¹H NMR spectra of these compounds were almost identical and consisted of two 1,3-disubstituted pyridiniums, two methylene chains, and two olefinic protons [**5**: δ_H 5.41 (2 H, br t, J = 3.6 Hz, H-12, H-13); **6**: δ_H 5.38 (2 H, m, H-13, H-14)]. Two prominent daughter ions at *m/z* 216 and 218 in the FAB-MS/MS data of **5** and **6** showed a C_{10} -saturated chain and a C_{10} -chain with one double bond. The *g*HMBC data of **5** and **6** showed that these compounds had the same 1,3-dialkylpyridinium substitution patterns as **1**–4.

The remaining alkyl substructure and the locations of the double bonds were determined by combined NMR analyses. The ¹H COSY (**5** and **6**: H-7-H-16) and gHMBC (**5**: H-11/C-12, C-13, H-12/C-11, C-14, H-13/C-11, C-14, H-14/C-12, C-13, **6**: H-12/C-13, C-14, H-13/C-12, C-15, H-14/C-12, C-15, H-15/C-13, C-14) data indicated that **5** and **6** contained a double bond at C-12 and C-13, respectively. Since the olefinic and allylic protons of the double bonds were chemically equivalent, the NOESY analysis could not be used to determine the configurations of the double bonds. Consequently, the configuration of the double bond in **5** was assigned as *E* based on the downfield shifts of the allylic methylene [δ_C 32.7 (C-11) and 32.4 (C-14)], while the *Z*-configuration was deduced for **6** from the upfield chemical shifts of the allylic methylene [δ_C 28.0 (C-12) and 29.2 (C-15)] [15].

The molecular formulas of **7** and **8** were $C_{31}H_{47}Cl_2N_2$ based on HR-FAB-MS analyses. As found for **5** and **6**, the ¹H NMR spectra of these compounds revealed characteristic features of cyclostellettamines: 1,3-disubstituted pyridines, methylene chains, and two olefinic protons [**7**: 5.37 (1 H, ddd, J = 11.4, 6.7, 2.0 Hz, H-9), 5.36 (1 H, ddd, J = 11.4, 6.7, 2.0 Hz, H-10); **8**: $\delta_{\rm H}$ 5.71 (1 H, dt, J = 10.8, 7.3 Hz, H-15), 5.66 (1 H, dt, J = 10.8, 7.3 Hz, H-16)] (Tables 1 and 3). The FAB-MS/MS analyses of **7** and **8**

supported the presence of a C₁₀-saturated chain and a C₁₁-chain with one double bond. The ¹H COSY (7: H-7–H-12, H-16–H-17, **8**: H-7–H-9, H-13–H-17) and gHMBC (7: H-8/C-9, C-10, H-9/C-8, C-11, H-10/C-8, C-11, H-11/C-8, C-10, **8**: H-14/C-15, C-16, H-15/C-14, C-17, H-16/C-14, C-17, H-17/C-15, C-16) data placed the double bonds at C-9 and C-15 for **7** and **8**, respectively. The configurations of these double bonds were both determined to be *Z* based on the chemical shifts of the allylic methylene carbons [**7**: 24.8 (C-8) and 28.1 (C-11), **8**: $\delta_{\rm C}$ 28.3 (C-14) and 31.0 (C-17)] [15].

Position	7	8	9	10
2	8.99 (1H, s)	8.87 (1H, s)	8.99 (1H, s)	8.93 (1H, s)
3	-	-	-	-
4	8.43 (1H, d, 8.0)	8.46 (1H, d, 7.6)	8.45 (1H, d, 8.0)	8.45 (1H, d, 8.1)
5	8.00 (1H, dd, 7.8, 6.3)	8.03 (1H, dd, 7.8, 6.1)	8.02 (1H, dd, 7.7, 6.2)	8.02 (1H, dd, 7.9, 6.3)
6	8.80 (1H, dd, 6.2)	8.85 (1H, dd, 6.0)	8.82 (1H, dd, 6.1)	8.82 (1H, dd, 6.5)
7	4.62 (2H, t, 7.1)	4.65 (2H, t, 7.0)	4.61 (2H, t, 7.1)	4.64 (2H, t, 6.9)
8	2.10 (2H, m)	1.98 (2H, m)	2.00 (2H, m)	2.00 (2H, m)
9	5.37 (1H, ddd, 11.4, 6.7, 2.0)	1.26 (2H, m)	1.29 (2H, m)	1.24 (2H, m)
10	5.36 (1H, ddd, 11.4, 6.7, 2.0)	1.20 ~ 1.35 (2H, m)	1.32 (2H, m)	1.20 ~ 1.32 (2H, m)
11	1.94 (2H, m)	$1.20 \sim 1.35 \; (2H, m)$	1.28 (2H, m)	1.20 ~ 1.32 (2H, m)
12	1.36 (2H, m)	1.20 ~ 1.35 (2H, m)	1.99 (2H, m)	1.20 ~ 1.32 (2H, m)
13	1.23 ~ 1.35 (2H, m)	1.36 (2H, m)	5.34 (1H, m)	1.20 ~ 1.32 (2H, m)
14	1.23 ~ 1.35 (2H, m)	2.15 (2H, m)	5.32 (1H, m)	1.70 (2H, m)
15	1.23 ~ 1.35 (2H, m)	5 71 (1H, dt, 10.8, 7.3)	2.06 (2H, m)	5.39 (1H, dt, 10.6, 6.7)
16	1.78 (2H, m)	5.66 (1H, dt, 10.8, 7.3)	1.38 (2H, m)	5.38 (1H, dt, 10.6, 6.7)
17	2.88 (2H, t, 7.2)	3.68 (2H, t, 7.3)	1.72 (2H, m)	2.46 (2H, dt, 13.5, 6.6)
18	-	-	2.90 (2H, t, 7.5)	2.95 (2H, t, 6.8)
2'	8.90 (1H, s)	8.79 (1H, s)	8.98 (1H, s)	8.89 (1H, s)
3'	-	-	-	-
4′	8.45 (1H, d, 8.0)	8.47 (1H, d, 7.6)	8.45 (1H, d, 8.0)	8.43 (1H, d, 8.1)
5'	8.02 (1H, dd, 7.8, 6.3)	8.03 (1H, dd, 7.8, 6.1)	8.01 (1H, dd, 7.7, 6.2)	8.01 (1H, dd, 7.9, 6.3)
6′	8.82 (1H, dd, 6.2)	8.83 (1H, dd, 6.0)	8.83 (1H, dd, 5.8)	8.83 (1H, dd, 6.5)
7′	4.60 (2H, t, 6.7)	4.63 (2H, t, 7.5)	4.63 (2H, t, 6.8)	4.61 (2H, t, 6.9)
8′	1.99 (2H, m)	2.02 (2H, m)	2.01 (2H, m)	1.98 (2H, m)
9′	1.24 (2H, m)	1.28 (2H, m)	1.27 (2H, m)	1.25 (2H, m)
10'	1.23 ~ 1.38 (2H, m)	$1.26 \sim 1.35 \; (2H, m)$	$1.26 \sim 1.36 \ (2H, m)$	$1.25 \sim 1.32 \; (2H, m)$
11′	1.23 ~ 1.38 (2H, m)	$1.26 \sim 1.35 \; (2H, m)$	$1.26 \sim 1.36 \ (2H, m)$	$1.25 \sim 1.32 \; (2H, m)$
12'	1.23 ~ 1.38 (2H, m)	$1.26 \sim 1.35 \; (2H, m)$	1.26 ~ 1.36 (2H, m)	$1.25 \sim 1.32 \; (2H, m)$
13'	1.23 ~ 1.38 (2H, m)	$1.26 \sim 1.35 \; (2H, m)$	$1.26 \sim 1.36 \ (2H, m)$	$1.25 \sim 1.32 \; (2H, m)$
14'	1.32 (2H, m)	1.34 (2H, m)	1.30 (2H, m)	1.29 (2H, m)
15'	1.73 (2H, m)	1.74 (2H, m)	1.73 (2H, m)	1.75 (2H, m)
16′	2.88 (2H, t, 7.0)	2.90 (2H, t, 7.2)	2.89 (2H, t, 7.3)	2.91 (2H, t, 7.0)

Table 3. ¹H MMR (δ , mult, J in Hz) assignments for compounds 7–10 in MeOH- d_4 .

The molecular formulas of **9** and **10** were both $C_{32}H_{49}Cl_2N_2$ based on HR-FAB-MS analyses. Similar to compounds **5–8**, the ¹H NMR spectra of **9** and **10** revealed a 1,3-dialkyl pyridinium structure with two olefinic protons [**9**: δ_H 5.34 (1 H, m, H-13), 5.32 (1 H, m, H-14); **10**: 5.39 (1 H, dt, J = 10.6, 6.7 Hz, H-15), 5.38 (1 H, dt, J = 10.6, 6.7 Hz, H-16)]. The FAB-MS/MS analyses of these compounds confirmed the presence of a C₁₀-saturated chain and a C₁₂-chain with one double bond. The COSY (**9**: H-7–H-18, **10**: H-7–H-9, H-14–H-18) and gHMBC (**9**: H-12/C-13, C-14, H-13/C-12, C-15, H-15/C-13, C-14, **10**: H-14/C-15, C-16, H-15/C-14, C-17, H-16/C-14, C-17, H-17/C-15, C-16) data were indicative of a double bond at C-13 and C-15 for **9** and **10**, respectively. The configurations of these double bonds were both assigned as *Z* based on the upfield shifts of the allylic methylene carbons [**9**: δ_C 28.0 (C-12), 27.9 (C-15), **10**: 28.2 (C-14), 29.1 (C-17)].

A comparison of the ¹³C and ¹H NMR data revealed almost identical chemical shifts for both carbons and protons among all cyclic 1,3-dialkylpyridinium dimers 1-10 (Tables 1–3). These compounds were structurally similar to cyclostellettamines reported previously with differences in the length of the alkyl chains, as well as the presence of double bonds [9]. Compounds **5–10** were dehydrocyclostellettamines containing one double bond in the alkyl chains. Subsequently, the carbon chemical shifts of allylic methylenes and coupling constants between the olefinic protons were used to assign *E* or Z configurations to these double bonds.

The 1,3-dialkylpyridinium metabolites possess diverse cytotoxic and antimicrobial activities [16,17]. Thus, we evaluated the cytotoxic and antimicrobial activities of compounds 1–10. Since 7 was isolated in a TFA salt form, we evaluated the influence of the counter ion on bioactivity. Thus, the TFA salt of 4 was prepared using reversed phase HPLC (H₂O-MeOH, 80:20, 0.1% TFA) on 4. Compounds 1–10 and 4a exhibited moderate cytotoxicity against the A549 lung cancer cell line, comparable to doxorubicin (Table 4). On the other hand, they also displayed a diverse range of antibacterial activities against Gram-positive strains (compounds 2, 4, 8, 9, and 10 were the most potent). These results are partly consistent with the previous report that the bioactivities of cyclostellettamines were significantly influenced by the length of the alkyl chains [16]. In addition, as shown for 6 and 9, the bioactivities were influenced by the existence and location of the double bond, although the alkyl chains were the same length. This suggests a mechanism-of-action based on the distances between the charged moieties and electron-rich sites, possibly through an interaction with ionic membrane sites. We also observed that the antibacterial activities of these compounds were generally weaker against Gram-negative than Gram-positive strains. Also, the influence of pyridinium counterions was negligible since there was no significant difference between the bioactivities of 4 and 4a. Overall, these results increase our understanding of the structure-activity relationships of cyclostellettamines.

In addition to the antimicrobial and cytotoxic activity assays, compounds 1–10 were also tested against diverse pathogenic fungal strains and microbial enzymes, such as isocitrate lyase, sortase A, and Na⁺/K⁺-ATPase, and failed to display any bioactivity (LC₅₀ > 100 μ g/mL for fungi, ED₅₀ > 100 μ M for enzymes).

	MIC (µg/mL)						LC ₅₀ (µM) ^a
Compound	Gram (+) Bacterium			Gram (–) Bacterium			A549
	Α	В	С	D	Е	F	
1	50	100	50	100	100	>100	22.1
2	12.5	25	3.125	25	25	>100	15.3
3	50	50	100	50	25	>100	24.0
4	50	100	6.25	100	100	>100	26.3
5	25	100	25	50	50	>100	25.9
6	50	100	50	100	100	>100	28.9
7	50	100	100	100	50	>100	89.4
8	12.5	25	3.125	50	50	>100	15.7
9	12.5	25	3.125	25	25	>100	14.7
10	12.5	25	3.125	25	50	>100	14.8
4a	25	100	50	100	100	>100	21.9
Ampicillin	0.39	0.78	0.39	0.78	0.39	6.25	
Doxorubicin							3.37

Table 4. Results of Bioactivity Tests.

A: Staphylococcus aureus (ATCC 6538p); B: Bacillus subtilis (ATCC 6633); C: Micrococcus luteus (IFO 12708);

D: Salmonella typhimurium (ATCC 14028); E: Proteus vulgaris (ATCC 3851); F: Escherichia coli (ATCC 35270); ^a The results of 48 h treatment were chosen.

3. Experimental Section

3.1. General Experimental Procedures

Optical rotation was measured on a JASCO digital polarimeter using a 1 cm cell. IR spectra were recorded on a JASCO FT/IR 4200 spectrophotometer. UV spectra were recorded on a Hitachi U-3010 spectrophotometer. Proton and carbon NMR spectra were measured at 600 and 150 MHz (3) on Bruker Avance 600 and 500 and 125 MHz (1, 2 and 4–10) on Bruker Avance 500, respectively. Mass spectrometric data were obtained from the Korea Basic Science Research Institute (Daegu) on a JEOL JMS-700 mass spectrometer. All solvents used were spectral grade or distilled from glass prior to use.

3.2. Collection and Taxonomic Identification

Haliclona sp. specimens were collected by hand using SCUBA equipment (20 m depth) at Sagyeri off the shore of Jeju Island, Korea, in November 1999. The pale-purple sponge measured 65×55 mm with a thickness of 27 mm and produced slime strands. The surface had many mamillate forms of a hard but fragile consistency. The size of the oxea spicule was (170–198) × (8–10) µm. A voucher specimen (registry No. Spo. 63) was deposited at the Natural History Museum, Hannam University, Korea, under the curatorship of C.J.S. Collections of the same specimens were made at the nearby Chuja-do in November 2003, and at Sagyeri in September 2009 and June 2010. Since the morphological features and ¹H NMR spectra of crude extracts from these collections were identical to those of the 1999 collection, chemical investigation was performed on the combined specimens.

3.3. Extraction and Isolation

The freshly collected specimens were frozen immediately and kept at -25 °C until use. The specimens (dried wt. 1.2 kg) were lyophilized, macerated, and extracted repeatedly with MeOH (3 L × 3) and CH₂Cl₂ (3 L × 2). The combined organic extract (940.3 g) was partitioned between *n*-BuOH and H₂O, and the former (158.18 g) was repartitioned between 15% aqueous MeOH (24.43 g) and *n*-hexane (130.25 g). The aqueous MeOH layer was subjected to C₁₈ reversed-phase vacuum flash chromatography using sequential mixtures of MeOH and H₂O as eluents (six fractions in gradient, H₂O–MeOH, from 50:50 to 0:100), and finally acetone. The H₂O–MeOH (50:50) fraction (1.64 g) was gel-filtered on a Sephadex LH-20 column with MeOH. The bioactive fractions were separated by C₁₈ reversed-phase semi-preparative HPLC (YMC ODS-A column, 1 × 25 cm, H₂O–MeOH, 80:20) to yield, in order of elution, compounds **3**, **4**, **1**, **7**, **8**, **2**, **6**, **5**, **10**, and **9**. Final purification of these metabolites was then accomplished by reversed-phase HPLC (YMC-Pack CN column, 1 × 25 cm, H₂O–MeOH (85:15) for **1**–**4**, H₂O–MeOH (90:10) for **5**, **6**, and **8**–**10**, and gradient solvents of H₂O–MeOH (85:15) to (40:60) and 0.1% TFA for 30 min for 7). The overall yields were 14.9, 19.8, 7.5, 187.4, 11.3, 22.5, 6.0, 72.4, 12.6, and 10.5 mg for **1**–**10**, respectively.

Compound **3**: Yellow, amorphous solid; UV (MeOH) λ_{max} 267 (log $\varepsilon = 3.7$), IR (ZnSe) v_{max} 2927, 1506, 1461 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 2, respectively; HR-FAB-MS *m/z* 407.3428 [M - H]⁺ (calculated for C₂₈H₄₃N₂, 407.3426) and 443.3195 [M + Cl]⁺ (calculated for C₂₈H₄₄N₂Cl, 443.3193); FAB-MS/MS fragments: *m/z* 190.19 [M - C₁₅H₂₄N]⁺ and 217.80 [M - C₁₃H₂₀N]⁺.

Compound 4: White, amorphous solid; UV (MeOH) λ_{max} 266 (log $\varepsilon = 3.9$), IR (ZnSe) ν_{max} 2924, 1505, 1467 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 2, respectively; HR-FAB-MS *m/z* 435.3741 [M - H]⁺ (calculated for C₃₀H₄₇N₂, 435.3739) and 471.3503 [M + Cl]⁺ (calculated for C₃₀H₄₈N₂Cl, 471.3506); FAB-MS/MS fragments: *m/z* 218.05 [M - C₁₅H₂₄N]⁺ and 218.05 [M - C₁₅H₂₄N]⁺.

Compound 5: Yellow, amorphous solid; UV (MeOH) λ_{max} 267 (log ε = 3.8), IR (ZnSe) v_{max} 2926, 1505, 1459 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 2, respectively; HR-FAB-MS *m/z* 433.3586 [M - H]⁺ (calculated for C₃₀H₄₅N₂, 433.3583), 469.3351 [M + Cl]⁺ (calculated for C₃₀H₄₆N₂Cl, 469.3350); FAB-MS/MS fragments: *m/z* 216.06 [M - C₁₅H₂₄N]⁺, 218.04 [M - C₁₅H₂₂N]⁺.

Compound 6: Yellow, amorphous solid; UV (MeOH) λ_{max} 266 (log ε = 3.8), IR (ZnSe) ν_{max} 2927, 1505, 1458 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 2, respectively; HR-FAB-MS *m/z* 433.3584 [M - H]⁺ (calculated for C₃₀H₄₅N₂, 433.3583) and 469.3352 [M + Cl]⁺ (calculated for C₃₀H₄₆N₂Cl, 469.3350); FAB-MS/MS fragments: *m/z* 216.07 [M - C₁₅H₂₄N]⁺ and 218.04 [M - C₁₅H₂₂N]⁺.

Compound 7: Yellow, amorphous solid; UV (MeOH) λ_{max} 267 (log $\varepsilon = 3.9$), IR (ZnSe) ν_{max} 2927, 1631, 1507, 1460, 1203, 1129 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 3 respectively; HR-FAB-MS *m/z* 447.3736 [M - H]⁺ (calculated for C₃₁H₄₇N₂, 447.3739) and 561.7414 [M + CF₃COO]⁺ (calculated for C₃₀H₄₈N₂CF₃COO, 561.7416); FAB-MS/MS fragments: *m/z* 217.92 [M - C₁₆H₂₄N]⁺ and 229.76 [M - C₁₅H₂₄N]⁺.

Compound 8: Yellow, amorphous solid; UV (MeOH) λ_{max} 267 (log ε = 3.8), IR (ZnSe) v_{max} 2924, 1506, 1467 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 3, respectively; HR-FAB-MS *m/z* 447.3742

 $[M - H]^+$ (calculated for C₃₁H₄₇N₂, 447.3739) and 483.3504 $[M + Cl]^+$ (calculated for C₃₁H₄₈N₂Cl, 483.3506); FAB-MS/MS fragments: *m/z* 218.11 $[M - C_{16}H_{24}N]^+$ and 229.87 $[M - C_{15}H_{24}N]^+$.

Compound **9**: Yellow, amorphous solid; UV (MeOH) λ_{max} 266 (log $\varepsilon = 3.7$), IR (ZnSe) v_{max} 2925, 1505, 1459 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 3, respectively; HR-FAB-MS *m/z* 461.3898 [M - H]⁺ (calculated for C₃₂H₄₉N₂, 461.3896) and 497.3665 [M + Cl]⁺ (calculated for C₃₂H₅₀N₂Cl, 497.3663); FAB-MS/MS fragments: *m/z* 218.19 [M - C₁₇H₂₆N]⁺ and 243.75 [M - C₁₅H₂₄N]⁺.

Compound **10**: Yellow, amorphous solid; UV (MeOH) λ_{max} 266 (log ε = 3.9), IR (ZnSe) v_{max} 2925, 1505, 1458 cm⁻¹; ¹H and ¹³C NMR data, see Tables 1 and 3, respectively; HR-FAB-MS *m/z* 461.3899 [M - H]⁺ (calculated for C₃₂H₄₉N₂, 461.3896) and 497.3665 [M + Cl]⁺ (calculated for C₃₂H₅₀N₂Cl, 497.3663); FAB-MS/MS fragments: *m/z* 218.19 [M - C₁₇H₂₆N]⁺ and 243.75 [M - C₁₅H₂₄N]⁺.

3.4. Preparation of TFA Salt of Compound 4 (4a)

A small amount of compound **4** (6.0 mg) was injected into the HPLC (YMC-Pack CN column, 1×25 cm, H₂O–MeOH, 80:20, 0.1% TFA) to yield **4a** (4.8 mg): UV (MeOH) λ_{max} 266 (log $\varepsilon = 3.8$), IR (ZnSe) v_{max} 2928, 1631, 1506, 1464, 1203, 1129 cm⁻¹; HR-FAB-MS *m/z* 435.3741 [M - H]⁺ (calculated for C₃₀H₄₇N₂, 435.3739) and 549.7306 [M + CF₃COO]⁺ (calculated for C₃₀H₄₈N₂CF₃COO, 549.7309); FAB-MS/MS fragments: *m/z* 218.05 [M - C₁₅H₂₄N]⁺ and 218.05 [M - C₁₅H₂₄N]⁺.

3.5. Biological Assays

Cytotoxicity assays, antimicrobial assays, and isocitrate lyase, sortase A, and Na⁺/K⁺-ATPase inhibition assays were performed as described previously [18–22]. For the cytotoxicity test, an MTT viability assay was performed as previously described with slight modifications [18]. MTT was first prepared as a stock solution of 5 mg/mL in phosphate buffered saline (PBS, pH 7.2) and was filtered. At the end of the treatment period (24 h, 48 h, and 72 h), with three different test drug concentrations in triplicate, MTT solution (20 μ L) was added to each well and then incubated for 4 h at 37 °C; then solubilizing buffer (10% sodium dodecyl sulfate dissolved in 0.01 N HCl, 100 μ L) was added to each well. After overnight incubation, the 96-well plate was read by an enzyme-linked immunosorbent assay (ELISA) reader at 570 nm for absorbance to determine the cell (A549 cell line) viability. The viable cells produced a dark blue formazan product, whereas no such staining was formed by dead cells [19]. The LC₅₀ value was defined as the concentration that resulted in a 50% decrease in cell viability compared to that of control reactions in the absence of an inhibitor. The values (mean ± SD) were calculated from the dose–response curves of three concentrations of each test sample in three independent experiments (*n* = 3).

4. Conclusion

Eight novel cyclic bis-1,3-dialkylpyridiniums (3–10), as well as two known compounds (1 and 2) belong to the cyclostellettamine class, were isolated from the sponge *Haliclona* sp. from Korea. These compounds were structurally different from the previous reported cyclostelletamines in the lengths of the alkyl chains, as well as the presence of double bonds in the chains. Several of these new compounds exhibited moderate cytotoxicity against A549 cell-line and inhibitory activities against

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Gram-positive bacterial strains. As shown in **6** and **9**, the bioactivities were influenced by the existence and the location of a double bond. Overall, these results provide us better understanding of the structure-activity relationships of the cyclostellettamine class compounds.

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