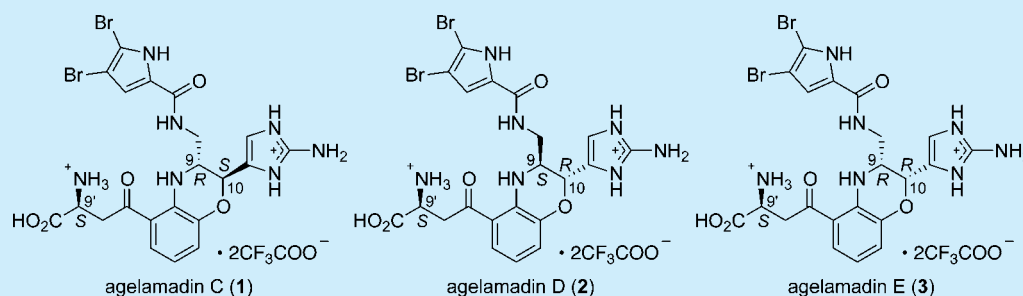


Agelamadins C–E, Bromopyrrole Alkaloids Comprising Oroidin and 3-Hydroxykynurenine from a Marine Sponge *Agelas* sp.Taishi Kusama,[†] Naonobu Tanaka,^{†,‡} Kanae Sakai,[§] Tohru Gono,[§] Jane Fromont,^{||} Yoshiki Kashiwada,[‡] and Jun'ichi Kobayashi^{*,†}[†]Graduate School of Pharmaceutical Sciences, Hokkaido University, Sapporo 060-0812, Japan[‡]Graduate School of Pharmaceutical Sciences, The University of Tokushima, Tokushima 770-8505, Japan[§]Mycology Research Center, Chiba University, Chiba 260-8673, Japan^{||}Western Australian Museum, Locked Bag 49, Welshpool DC, WA 6986, Australia

S Supporting Information



ABSTRACT: Three structurally unique bromopyrrole alkaloids, agelamadins C–E (1–3), were isolated from a marine sponge *Agelas* sp. Agelamadin C (1) possesses a hybrid structure of oroidin and 3-hydroxykynurenine connected through a dihydro-1,4-oxazine moiety. Agelamadins D (2) and E (3) are a C-9/C-10 diastereomer and a 10-epimer of 1, respectively. The structures of 1–3 were elucidated on the basis of spectroscopic analysis as well as application of a PGME method and a TDDFT ECD calculation. Antimicrobial activity of 1–3 was evaluated.

Marine sponges have been recognized as a rich source of interesting bioactive metabolites with various chemical structures.¹ Among them, sponges belonging to the genus *Agelas* are known to contain bromopyrrole alkaloids, which have attracted widespread interest due to their fascinating complex chemical structures with a high N to C ratio (~1:2) and intriguing biological activities.² In our continuing search for structurally unique metabolites from Okinawan marine sponges, we have reported the isolation of several bromopyrrole alkaloids, nagelamides U–Z and agelamadins A and B, from the extract of a sponge *Agelas* sp. (SS-162).³ Further investigation of the extract afforded three new bromopyrrole alkaloids, agelamadins C–E (1–3). In this Letter, we describe the isolation and structure elucidation of 1–3.

The sponge *Agelas* sp. (SS-162, 3.9 kg, wet weight) collected off Kerama Islands, Okinawa, was extracted with MeOH, and a part of the extract was partitioned with *n*-hexane and 90% aq MeOH. The 90% aq MeOH-soluble materials were further partitioned with *n*-BuOH and H₂O. The *n*-BuOH-soluble materials were chromatographed on a silica gel column, Toyopearl HW-40 column, and C₁₈ column to give fractions containing bromopyrrole alkaloids, which were purified using reverse-phase HPLC to afford agelamadins C (1, 0.000095%, wet weight), D (2, 0.000072%), and E (3, 0.000054%).

Agelamadin C (1)⁴ was obtained as a yellow amorphous solid {[α]_D²³ +71.8 (*c* 0.25, MeOH)}, and the ESIMS showed the pseudomolecular ion peaks at *m/z* 610, 612, and 614 (1:2:1), suggesting the existence of two bromine atoms in the molecule. The molecular formula of 1, C₂₁H₂₃N₇O₃Br₂, was established by the HRESIMS (*m/z* 610.00531 [*M* – H]⁺, Δ + 0.94 mmu). The IR spectrum implied the presence of carbonyl functionalities (1685 and 1636 cm^{−1}). The gross structure of 1 consisting of two partial units {units A (N-1–C-15) and B (C-1'–C-10')} (Figure 1) was assigned as follows. Comparison of ¹H and ¹³C NMR data for 1 (Table 1) and those for known bromopyrrole alkaloids³ suggested that 1 has one bromopyrrole amide moiety (N-1–N-7) and one aminoimidazole moiety (C-11–C-15). Interpretation of the ¹H–¹H COSY spectrum revealed the connectivities of N-7 to C-10 and C-9 to 9-NH (Figure 1). The presence of an amino group at C-9 was confirmed by an HMBC cross-peak of 9-NH to C-10 (Figure 1). The chemical shifts of CH-10 (δ_{H} 5.11; δ_{C} 65.9) were indicative of the existence of an oxygen atom at C-10. The connectivity of C-10 to the aminoimidazole moiety (C-11) was disclosed by HMBC correlations for H-10 to C-11 and C-15

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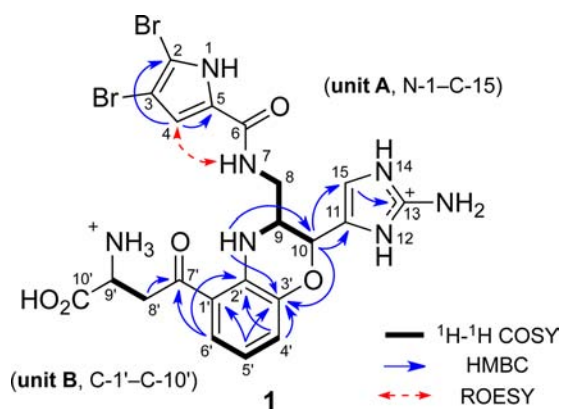


Figure 1. Selected 2D NMR correlations of agelamadin C (1).

(Figure 1). Therefore, unit A was elucidated to be 9-amino-10-hydroxyrooidin.

The existence of a 1,2,3-trisubstituted benzene ring (C-1'–C-6') in unit B was deduced from six aromatic carbon signals (δ_C 141.0, 136.5, 124.5, 121.1, 116.8, and 114.7) as well as the coupling patterns of three aromatic protons (δ_H 7.44 (d, $J = 7.8$ Hz), 7.00 (d, $J = 7.8$ Hz), and 6.55 (t, $J = 7.8$ Hz)). The carbon chemical shifts of C-2' (δ_C 136.5) and C-3' (δ_C 141.0) implied that an amino group and a hydroxy group were located at C-2' and C-3', respectively. The presence of a 3-amino-3-carboxypropanoyl group (C-7'–C-10') was indicated by the remaining four carbon signals (δ_C 197.5, 171.1, 48.4, and 39.2) together with the 2D NMR correlations (Figure 1). The

connectivity of C-1' to C-7' as well as the presence of an amino group at C-2' and a hydroxy group at C-3' were confirmed by HMBC correlations shown in Figure 1. Thus, unit B was suggested to be 3-hydroxykynurenine, which was further confirmed by comparison of the 1H and ^{13}C resonances for unit B with those for literature data.⁵ HMBC correlations for 9-NH to C-3' and H-10 to C-3' revealed the connectivities between units A and B forming a dihydro-1,4-oxazine ring. Accordingly, the gross structure of agelamadin C (1) was assigned as shown in Figure 1.

In the ROESY spectrum of 1, correlations for NH-7/H-10 and H-9/H-15 were observed (Figure 2), indicating the *anti* relationship for H-9 and H-10.

Agelamadins D (2)⁶ and E (3)⁷ were individually isolated as optically active yellow amorphous solids $\{[\alpha]_D^{23} -85.6$ (c 0.25, MeOH) for 2; $[\alpha]_D^{24} -66.6$ (c 0.25, MeOH) for 3}. The HRESIMS spectra revealed that 2 and 3 had the same molecular formula, $C_{21}H_{23}N_7O_5Br_2$, as that of 1 (m/z 610.00562 $[M - H]^+$, $\Delta + 1.25$ mmu for 2; m/z 610.00486 $[M - H]^+$, $\Delta + 0.49$ mmu for 3). Resemblance of the 1H and ^{13}C NMR data for 2 and 3 (Table 1) to those for 1 indicated that 2 and 3 are diastereomers of 1. A similar value of $^3J_{H-9/H-10}$ in 2 ($J = 2.5$ Hz) with that of 1 ($J = 2.7$ Hz) suggested the H-9/H-10 *anti* relationship in 2, which was supported by analysis of the ROESY spectrum (Figure S2). In contrast, the H-9/H-10 *syn* relationship in 3 was deduced on the basis of a small value of $^3J_{H-9/H-10}$ in 3 ($J \approx 0$ Hz) as well as differences between the chemical shifts for CH_2 -8, CH-10, and C-11 in 3 and those corresponding positions in 1.

Table 1. 1H and ^{13}C NMR Data for Agelamadins C–E (1–3) in DMSO- d_6

position	1		2		3	
	^{13}C	1H	^{13}C	1H	^{13}C	1H
1		12.72 (1H, br)		12.71 (1H, br)		12.69 (1H, br)
2	104.8		104.9		104.9	
3	97.9		98.0		97.9	
4	113.1	6.95 (1H, s)	113.2	6.95 (1H, s)	113.1	6.91 (1H, s)
5	127.8		127.8		127.8	
6	159.5		159.6		159.4	
7		8.48 (1H, br t, 5.6 Hz)		8.48 (1H, br t, 5.0 Hz)		8.41 (1H, br t, 5.4 Hz)
8	41.2	3.35 (2H, m)	41.4	3.33 (2H, m)	39.4 ^a	3.28 (2H, m)
9	49.7	4.01 (1H, m)	49.9	3.97 (1H, m)	50.5	4.03 (1H, m)
9-NH		8.70 (1H, br s)		8.64 (1H, br s)		8.53 (1H, br s)
10	65.9	5.11 (1H, d, 2.7 Hz)	65.9	5.08 (1H, d, 2.5 Hz)	67.8	5.26 (1H, br s)
11	123.1		123.2		121.9	
12		nd ^b		nd ^b		nd ^b
13	148.1		148.3		147.9	
13-NH ₂		7.79 (2H, br s)		7.84 (2H, br s)		7.78 (2H, br s)
14		nd ^b		nd ^b		nd ^b
15	112.1	6.99 (1H, s)	112.3	6.86 (1H, s)	111.4	6.82 (1H, s)
1'	116.8		117.3		117.9	
2'	136.5		136.4		136.4	
3'	141.0		141.1		141.7	
4'	121.1	7.00 (1H, d, 7.8 Hz)	121.0	6.96 (1H, d, 7.8 Hz)	121.0	7.00 (1H, d, 7.8 Hz)
5'	114.7	6.55 (1H, t, 7.8 Hz)	114.8	6.53 (1H, t, 7.8 Hz)	115.4	6.61 (1H, t, 7.8 Hz)
6'	124.5	7.44 (1H, d, 7.8 Hz)	124.5	7.40 (1H, d, 7.8 Hz)	124.4	7.46 (1H, d, 7.8 Hz)
7'	197.5		198.1		198.1	
8'	39.2 ^a	3.57 (2H, m)	40.2	3.53, 3.41 (1H each, m)	39.7 ^a	3.54 (2H, m)
9'	48.4	4.19 (1H, m)	49.1	4.04 (1H, m)	48.6	4.12 (1H, m)
10'	171.1		171.9		171.2	

^aOverlapped with signal of DMSO- d_6 . ^bNot detected.

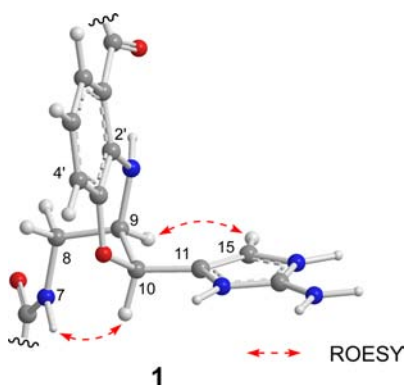


Figure 2. Selected ROESY correlations and the relative configurations at C-9 and C-10 for agelamadin C (1).

The absolute configurations at C-9' in 1–3 were assigned by a phenylglycine methyl ester (PGME) method.⁸ Since application of a PGME method to α -amino acids was not reported, three authentic α -amino acids, L-kynurenine (4), L-tyrosine (5), and D-tryptophan (6), were converted into the (S)- and (R)-PGME amides (4a/4b, 5a/5b, and 6a/6b), respectively. The $\Delta\delta$ values ($\Delta\delta = \delta_S - \delta_R$) obtained from the ^1H NMR spectra of 4a/4b, 5a/5b, and 6a/6b measured in CD_3OD and $\text{DMSO}-d_6$ are shown in Figure 3A. Distributions of positive $\Delta\delta$ values were observed in side chains of the PGME amides (4a/4b and 5a/5b) from L-amino acids, whereas a distribution of negative $\Delta\delta$ values was found in the PGME amides (5a/5b) of D-amino acids. These observations suggested that a PGME method was applicable to agelamadins C–E (1–3). The application to 1–3 (Figure 3B) indicated that the absolute configurations at C-9' in 1–3 were all S.

The absolute configurations at C-9 and C-10 of 1 and 2 were elucidated by comparison of the ECD spectra with those

calculated spectra. In the ECD spectrum of 1, a positive Cotton effect was observed at 242 nm, whereas 2 showed a negative Cotton effect at 242 nm (Figure 4). The calculated ECD

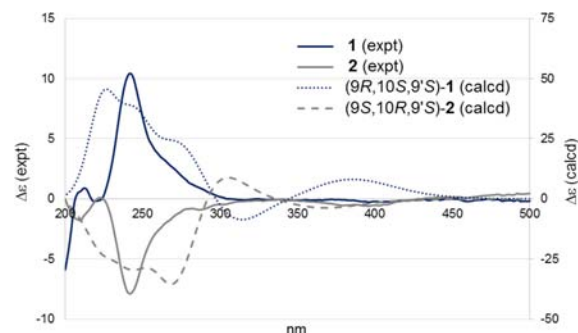


Figure 4. Experimental and calculated ECD spectra of agelamadins C (1) and D (2).

spectra of two possible diastereomers (9R,10S,9'S for 1 and 9S,10R,9'S for 2) by TDDFT method⁹ agreed well with the experimental spectra of 1 and 2, respectively (Figure 4). Accordingly, the 9R and 10S configurations for 1 and the 9S and 10R configurations for 2 were assigned. Similarly, the 9R and 10R configurations for 3 were elucidated by comparison of the experimental and calculated ECD spectra (Figure 5).

Agelamadins C–E (1–3) are structurally unique bromopyrrole alkaloids possessing hybrid structures of oroidin and 3-hydroxykynurenine connected through a dihydro-1,4-oxazine moiety. A possible biogenetic pathway of agelamadins C–E (1–3) is proposed in Scheme 1. Condensation of oroidin¹⁰ and 3-hydroxykynurenine derived from L-tryptophan would give 1–3. To the best of our knowledge, agelamadins C–E (1–3) are

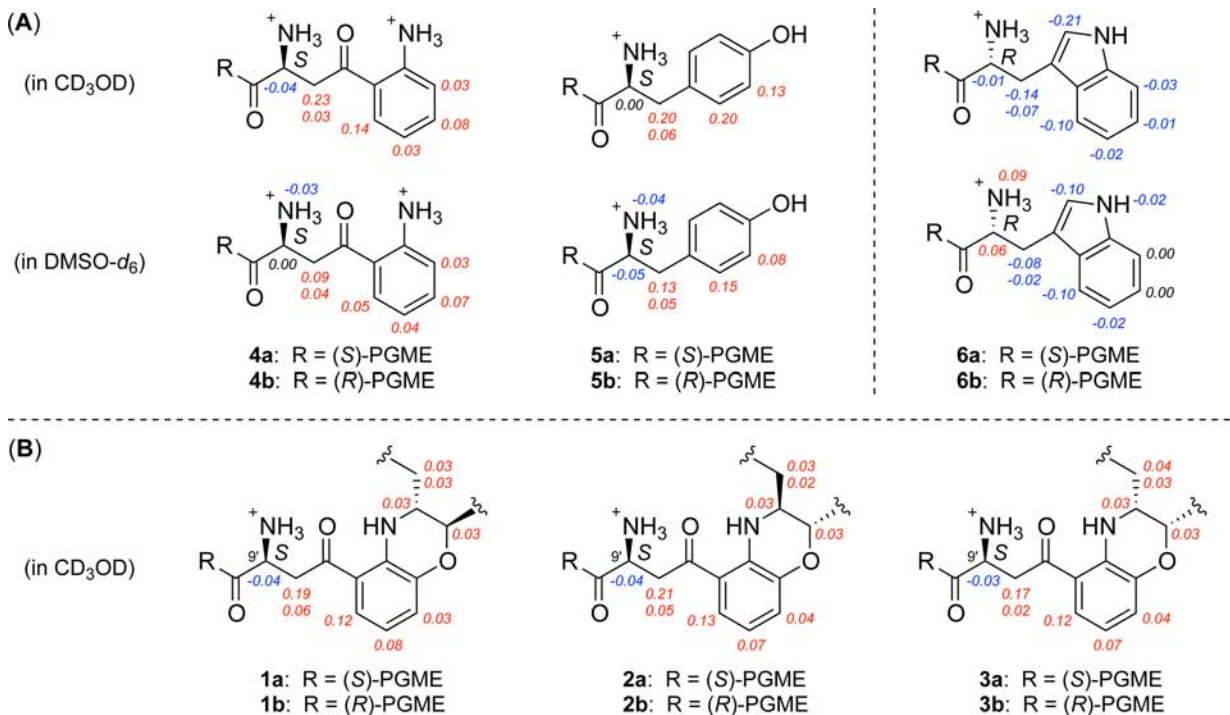


Figure 3. $\Delta\delta$ values [$\Delta\delta$ (in ppm) = $\delta_S - \delta_R$] obtained for (A) (S)- and (R)-PGME amides (4a/4b, 5a/5b, and 6a/6b) of L-kynurenine (4), L-tyrosine (5), and D-tryptophan (6); (B) (S)- and (R)-PGME amides (1a/1b, 2a/2b, and 3a/3b) of agelamadins C–E (1–3).

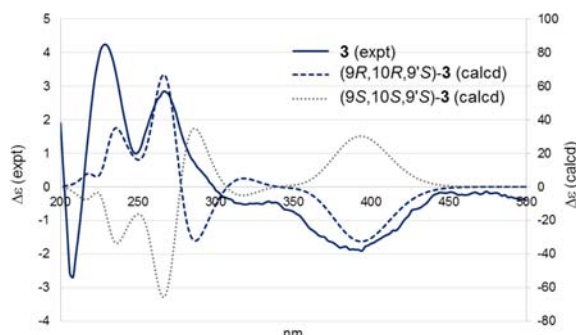
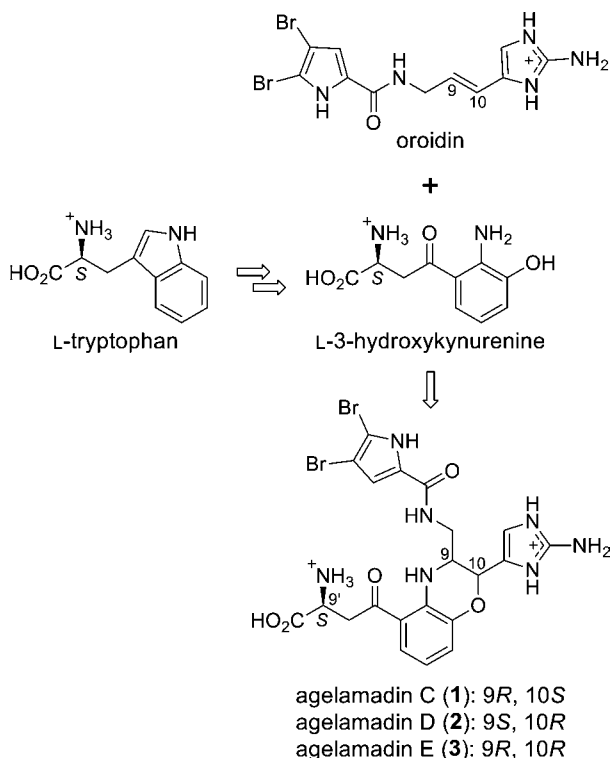


Figure 5. Experimental and calculated ECD spectra of agelamadin E (3).

Scheme 1. Possible Biogenetic Pathway of Agelamadins C–E (1–3)



the first example for bromopyrrole alkaloids comprising oroidin and a tryptophan derivative.

In antimicrobial screening of agelamadins C–E (1–3), 1 and 3 showed a weak inhibitory activity against *Cryptococcus neoformans* (IC_{50} 32 μ g/mL each).

■ ASSOCIATED CONTENT

Supporting Information

Experimental section, and 1D and 2D NMR spectra of agelamadins C–E. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Notes

The authors declare no competing financial interest.

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- (4) Agelamadin C (1): yellow amorphous solid; $[\alpha]_D^{23} +71.8$ (c 0.25, MeOH); UV (MeOH) λ_{max} 233 (ϵ 15300, sh) 275 (13400), and 387 (3300) nm; IR (KBr) ν_{max} 3337, 1685, and 1636 cm^{-1} ; 1H and ^{13}C NMR (Table 1); ESIMS m/z 610, 612, and 614 (1:2:1) $[M - H]^+$; HRESIMS m/z 610.00531 $[M - H]^+$ (calcd for $C_{21}H_{22}N_7O_5^{79}Br_2$, 610.00437).
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- (6) Agelamadin D (2): yellow amorphous solid; $[\alpha]_D^{22} -85.6$ (c 0.25, MeOH); UV (MeOH) λ_{max} 233 (ϵ 14600, sh), 275 (13100), and 388 (3200) nm; IR (KBr) ν_{max} 3434, 1684, and 1636 cm^{-1} ; 1H and ^{13}C NMR (Table 1); ESIMS m/z 610, 612, and 614 (1:2:1) $[M - H]^+$; HRESIMS: m/z 610.00562 $[M - H]^+$ (calcd for $C_{21}H_{22}N_7O_5^{79}Br_2$, 610.00437).
- (7) Agelamadin E (3): yellow amorphous solid; $[\alpha]_D^{24} -66.6$ (c 0.25, MeOH); UV (MeOH) λ_{max} 234 (ϵ 12800, sh), 275 (10800), and 386 (2700) nm; IR (KBr) ν_{max} 3127, 1683, and 1638 cm^{-1} ; 1H and ^{13}C NMR (Table 1); ESIMS m/z 610, 612, and 614 (1:2:1) $[M - H]^+$; HRESIMS m/z 610.00486 $[M - H]^+$ (calcd for $C_{21}H_{22}N_7O_5^{79}Br_2$, 610.00437).
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