



Cytotoxic bromotyrosine derivatives from a two-sponge association of *Jaspis* sp. and *Poecillastra* sp.

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ABSTRACT

Bioassay-guided chemical investigation of the lipophilic extract of a two-sponge association (*Jaspis* sp. and *Poecillastra* sp.) led to the isolation of two new bromotyrosine derivatives (**1** and **2**), along with known derivatives (**3–12**). Cyclobispsammaplin A (**1**) is a cyclic derivative of the previously reported bispsammaplin A (**13**), while psammaplin M (**2**) is composed of β -alanine (or aspartic acid) unit. Compounds **3**, **4**, **6**, **10**, and **12** are isolated for the first time from a sponge belonging to the subclass Tetractinomorpha. Structure elucidation was performed by a combination of high resolution mass and 2D NMR (principally COSY, HMBC, HSQC, and NOESY) spectroscopy. Compounds **1–4**, **6**, **10**, and **12** were evaluated for cytotoxicity against a small panel of five human solid tumor cell lines and their activity was compared in relevance to their structure.

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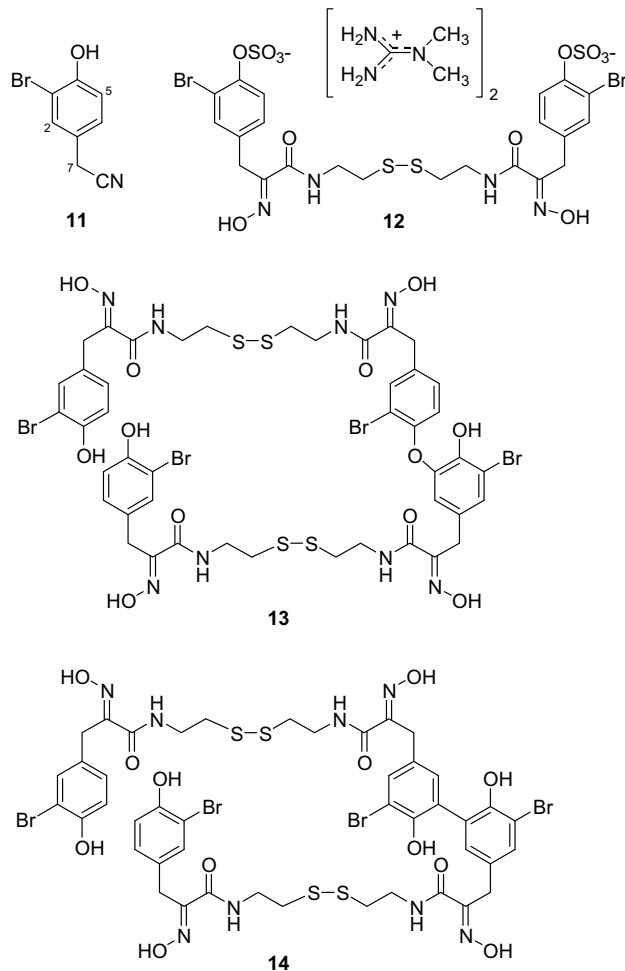
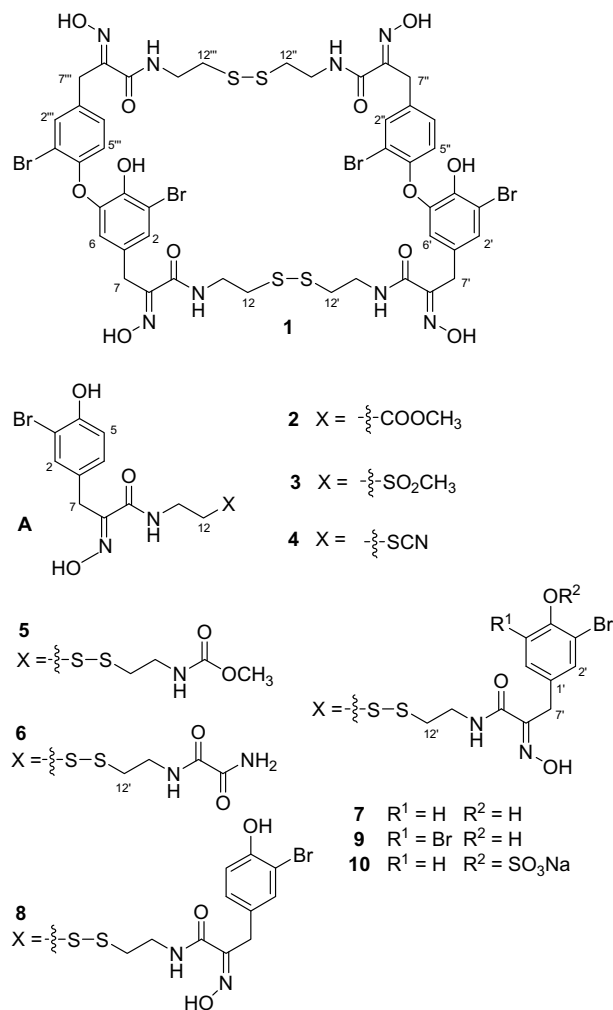
Marine sponges are the most primitive multicellular animals and contain many pharmacologically important metabolites. A two-sponge association of *Jaspis* sp. and *Poecillastra* sp. collected from Korean waters was known to contain dihydroxystyrene metabolites¹ and steroidal glycosides.² As a part of our study on cytotoxic constituents from marine organisms of Korean waters, we previously reported the isolation of pectenotoxin II and several psammaplin analogs from the MeOH extract of a two-sponge association of *Jaspis* sp. and *Poecillastra* sp.^{3,4} The two-sponge association of *Jaspis* sp. and *Poecillastra* sp. was collected again for our continued search for further cytotoxic constituents, and was found to contain apocarotenoids⁵ and glycerides.⁶ Further fractionation and purification of 90% aqueous MeOH extract led us to an isolation of 12 bromotyrosine derivatives (**1–12**). This report describes purification,⁷ structural characterization, and cytotoxicity evaluation of these bromotyrosine derivatives.

Structural elucidation of cyclobispsammaplin A (**1**), a white, amorphous solid,⁸ began from the isotopic clusters at m/z 1343/1345/1347/1349/1351 for $[M+Na]^+$, and at m/z 1319/1321/1323/1325/1327 for $[M-H]^-$ observed in a positive and negative mode ESIMS spectra, respectively. The exact mass of the pseudomolecular ions could not be measured by FABMS due to weak ion currents. So, the molecular formula, $C_{44}H_{44}N_8O_{12}S_4^{79}Br_2^{81}Br_2$, was determined

on the basis of more sensitive but less accurate HRMALTITOFMS ($[M+Na]^+$ m/z 1344.8854, 1346.8811, 1348.8965, and 1350.8937) and NMR data (Table 1). The exact mass of the $[M+Na]^+$ ion (m/z 1346.8811) matched with the expected formula $C_{44}H_{44}N_8O_{12}S_4^{79}Br_2^{81}Br_2Na$ (Δ +26.0 mmu, 19 ppm). The ^{13}C NMR data revealed the presence of only 22 carbons, suggesting it to be a symmetric dimer. Five aromatic proton signals, two isolated methylene groups, and two pairs of coupled methylene groups were observed in 1H NMR spectrum. These 1H and ^{13}C NMR data were reminiscent of the data of bispsammaplin A.⁴ Interpretation of the 2D NMR data including COSY, HSQC, and HMBC spectra enabled the construction of units **a–d** (Fig. 1). The partial structure **a**, comprised of 1,3,4,5-tetrasubstituted phenyl ring, was gauged by resonances at δ 7.21 (d, J = 2.0 Hz, H-2/H-2') and 6.47 (d, J = 2.0 Hz, H-6/H-6') and an isolated benzylic methylene at δ 3.70 (s, H₂-7/H₂-7'). While another unit **b** contained a phenyl ring with a 1,3,4-trisubstitution as indicated by the resonances at δ 7.56 (d, J = 2.0 Hz, H-2''/H-2'''), 6.65 (d, J = 8.0 Hz, H-5''/H-5'''), 7.16 (dd, J = 8.0, 2.0 Hz, H-6''/H-6''') and an isolated benzylic methylene at δ 3.88 (s, H₂-7''/H₂-7'''). In addition to partial structures **a** and **b**, the NMR data also showed two more partial structures **c** and **d**, which consisted of two pairs of coupled methylenes as assigned from the resonances at δ 3.56 (m, H₂-11/H₂-11')/2.90 (t, J = 7.0 Hz, H₂-12/H₂-12') and δ 3.40 (t, J = 7.0 Hz, H₂-11''/H₂-11''')/2.69 (t, J = 7.0 Hz, H₂-12''/H₂-12'''). Partial units **a** and **c** were connected as indicated by HMBC correlation (Fig. 2) from proton H₂-11/H₂-11' (δ 3.56) to the amide carbon at δ 165.5

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(C-9/C-9'). This connection between **a** and **c** constructed a structure of psammaplin A^{3,4}. Similarly, HMBC correlation from protons H₂-11''/H₂-11''' (δ 3.40) to the amide carbon at δ 165.5 (C-9''/C-9''') confirmed connection between partial units **b** and **d**, comprising another structure of psammaplin A (**7**).^{3,4} Chemical shift values for methylene protons are very typical when attached to tetrasubstituted (δ 3.56, H₂-11/H₂-11' and 2.90, H₂-12/H₂-12') and trisubstituted (δ 3.40, H₂-11''/H₂-11''' and 2.69, H₂-12''/H₂-12''') phenyl rings as observed in bispsammaplin A (**13**).⁴ The stereochemistry of the oxime groups was assigned as *E* on the basis of the characteristic ¹³C chemical shifts of the benzylic carbons (*E*: δ 28.6, 28.2; *Z*: δ 37.5).⁴ Hence, from the careful study of these assignments, the structure of **1** was elucidated as a cyclic bispsammaplin A.

But still there were two possibilities for its structure (**1A** or **1B**), considering the symmetric nature of the planar structure of **1** (Fig. 3). The structure **1A** possesses a plane of symmetry, while **1B** has an axis of symmetry. For **1A**, the magnetic environment of H-11/H-12 and H-11'/H-12' are expected to be equivalent. But, the magnetic environment of H-11/H-12 and H-11''/H-12'' would be non-equivalent. Therefore, these methylenes may be assigned as δ 3.56, H₂-11(H₂-11'); δ 2.90, H₂-12(H₂-12'); δ 3.40, H₂-11''(H₂-11'''); and δ 2.69, H₂-12''(H₂-12''') (assignment aided by NMR simulation using Chemdraw ultra). However, for **1B**, the magnetic environment of H-11/H-12 and H-11'/H-12' would be non-equivalent. But the magnetic environment of H-11/H-12 and H-11''/H-12'' would be equivalent. Therefore, these methylenes may be assigned as δ 3.56, H₂-11(H₂-11'); δ 2.90, H₂-12(H₂-12''); δ

3.40, H₂-11'(H₂-11'''); and δ 2.69, H₂-12'(H₂-12'''). If **1B** is the right structure for compound **1**, then we expect a NOE between the non-equivalent pair of nuclei H₂-12(H₂-12'') (δ 2.90) and H₂-12'(H₂-12''') (δ 2.69). ROESY and NOESY experiments were carried out, but no NOE was observed between these pairs of nuclei H-12(H-12'') and H-12'(H-12'''). The reason for the absence of NOE between these nuclei might be either **1B** was not the right structure or distances between these nuclei were not close enough to observe NOE. To find out the possible reason, study on 3D model of **1B** was performed using Chem3D Pro software. For the structure of minimum-energy **1B**, the distances between these nuclei were calculated as 2.1 and 2.7 Å (Fig. 4). It is well-known that NOE can be usually detected if the distance between the dipolar-coupled nuclei is less than 3 Å.⁹ Considering the absence of any NOE between these nuclei, just as an indirect clue, structure **1A** was proposed as the structure for cyclobispsammaplin A.

Psammaplin M (**2**) was isolated as an amorphous, white solid. The LRFABMS spectrum of **2** showed a pair of [M+Na]⁺ ion peaks at *m/z* 381/383 suggesting the presence of one bromine atom in the molecule. The HRFABMS and ¹³C NMR data¹⁰ of **2** supported the molecular formula C₁₃H₁₅N₂O₅Br. The exact mass of the [M-H]⁻ ion (*m/z* 357.0070) matched well with the expected formula C₁₃H₁₄N₂O₅Br (Δ 1.6 mmu). Analysis of the ¹H and ¹³C NMR data¹⁰ revealed the presence of a 1,2,4-trisubstituted phenyl ring (δ 7.35/134.2, 6.75/117.0, and 7.05/130.6) and an isolated benzylic methylene (δ 3.77/28.2). Furthermore, ¹³C NMR data showed the presence of an oxime (δ 153.0, C-8) and an amide (δ 165.5, C-9) groups. In addition, one carbonyl carbon (δ 173.5, C-13), one methoxyl singlet at (δ 3.63/52.5, OCH₃), and a pair of coupled

Table 1
1D and 2D NMR data of **1** in CD₃OD at 500 MHz.^a

Position	δ_{H}	δ_{C} ^b	COSY	HMBC
1 (1')		129.8		
2 (2')	7.21 (d, 2.0)	130.0	H-6 (H-6')	C-6 (C-6')
3 (3')		c		
4 (4')		c		
5 (5')		c		
6 (6')	6.47 (d, 2.0)	118.8	H-2 (H-2')	C-2 (C-2')
7 (7')	3.70 (s)	28.2		C-9, 8, 2 (C-9', 8', 2')
8 (8')		152.4		
9 (9')		165.5		
11 (11')	3.56 (m)	39.6	H-12 (H-12')	C-9, 12 (C-9', 12')
12 (12')	2.90 (t, 7.0)	39.4	H-11 (H-11')	C-11 (C-11')
1'' (1'')		129.8		
2'' (2'')	7.56 (d, 2.0)	135.0	H-6'' (H-6''')	C-4'' (C-4''')
3'' (3'')		114.0		
4'' (4'')		154.0		
5'' (5'')	6.65 (d, 8.0)	120.0	H-6'' (H-6''')	C-4'', C-2'', C-3'' (C-4''', 2''', 3''')
6'' (6'')	7.16 (d, 8.0, 2.0)	130.8	H-5'', 2'' (H-5''', 2''')	
7'' (7'')	3.88 (s)	28.6		C-9'', 8'', 6'' (C-9''', 8''', 6''')
8'' (8'')		153.0		
9'' (9'')		165.5		
11'' (11'')	3.40 (t, 7.0)	39.2	H-12'' (H-12''')	C-9'', 12'' (C-9''', 12''')
12'' (12'')	2.69 (t, 7.0)	38.0	H-11'' (H-11''')	C-11'' (C-11''')

^a Multiplicities and coupling constants are in parentheses.

^b Assignments based on HMBC and HSQC spectroscopic data.

^c Not observed due to small amount.

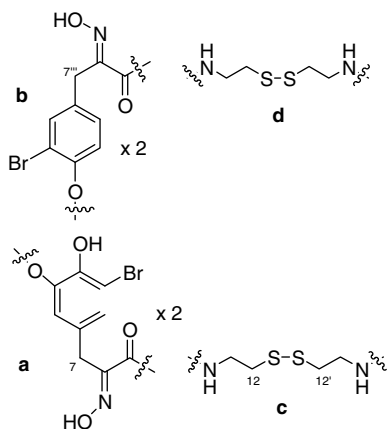


Figure 1. Partial structures of compound **1**.

methylenes at δ 3.48/35.7 (H₂-11/C-11) and at δ 2.53/34.3 (H₂-12/C-12) were observed in ¹H and ¹³C NMR data. These data indicated the presence of the substructure **A**, which also formed the half

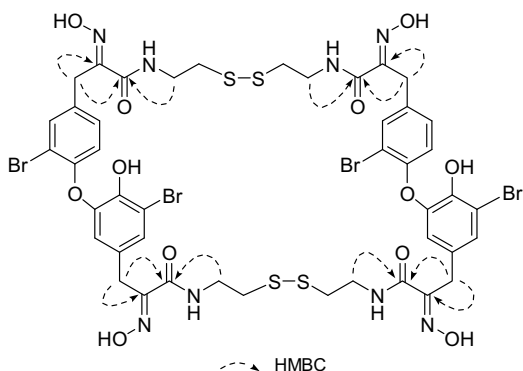


Figure 2. Key HMBC correlations of compound **1**.

structure of psammaplin A.⁴ In the HMBC spectrum, the signals of H₂-12 (δ 2.53) and OCH₃ (δ 3.63) showed correlations with C-13 (δ 173.5). The stereochemistry of the oxime group was assigned as *E* on the basis of the characteristic ¹³C chemical shift of the benzylic carbon (C-7, δ 28.2).⁴ Psammaplin M (**2**) might be derived from a tyrosine and aspartic acid or β -alanine. This is the first psammaplin analog which does not contain a cysteine-derived sulfur moiety.

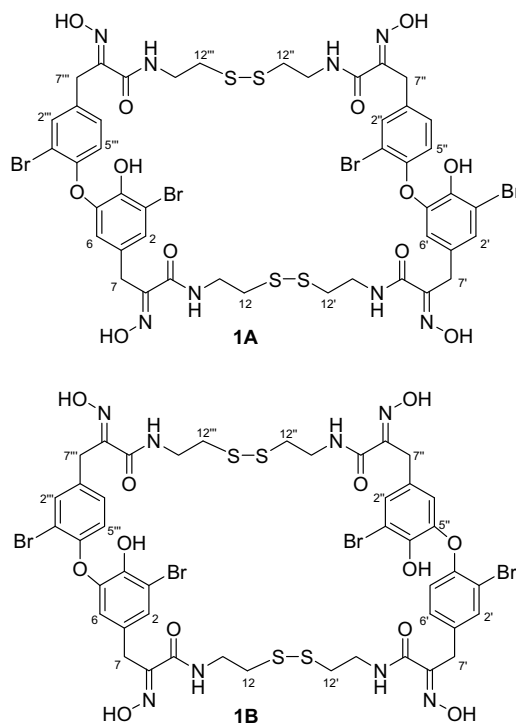


Figure 3. Two possible structures for compound **1**.

Table 2
Cytotoxicity data of bromotyrosine derivatives.^a

Compound	A-549	SK-OV-3	SK-MEL-2	XF-498	HCT-15
1 (cyclobispsammaplin A)	1.95	1.21	1.14	2.88	3.82
6 (psammaplin E)	1.47	0.19	0.21	1.63	1.92
12 (diguanium salt of psammaplin A sulfate)	12.27	1.79	4.48	16.92	43.17
doxorubicin	0.01	0.01	0.01	0.03	0.03
2 (psammaplin M)	>30	>30	>30	>30	>30
3 (psammaplin I)	4.15	1.76	2.84	2.96	6.51
4 (psammaplin B)	12.84	9.27	19.43	10.92	>30
10 (sodium salt of psammaplin A sulfate)	0.18	0.16	1.13	0.18	1.25
doxorubicin	0.02	0.07	0.10	0.10	0.33
5 (psammaplin D) ^b	0.80	0.17	0.20	0.60	1.23
7 (psammaplin A) ^b	0.57	0.14	0.13	0.57	0.68
9 (bromopsammaplin A) ^b	1.34	1.38	0.90	0.92	3.31
13 (bispsammaplin A) ^b	1.53	1.52	1.02	1.10	3.35
14 (bisaprasin) ^b	3.40	2.78	2.94	2.44	6.00
doxorubicin	0.04	0.15	0.003	0.10	0.09

^a Data expressed in ED₅₀ values (μg/mL). A-549, human lung cancer; SK-OV-3, human ovarian cancer; SK-MEL-2, human skin cancer; XF-498, human CNS cancer; HCT-15, human colon cancer.

^b Data from Ref. ⁴.

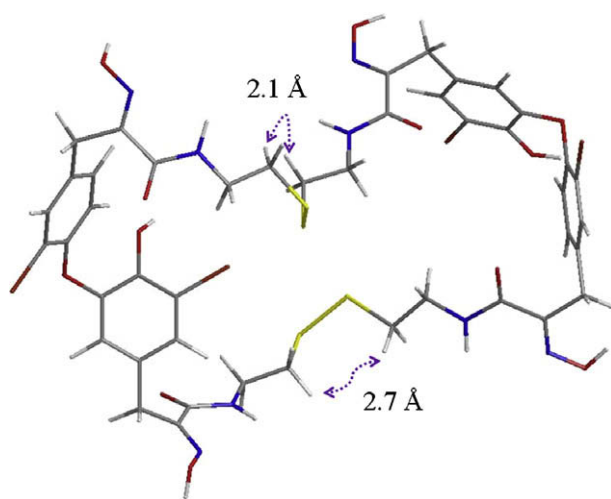


Figure 4. Energy-minimized structure of **1B**.

Compound **3** was identified as psammaplin I, previously reported from a sponge *Pseudoceratina purpurea*.¹¹ Compound **4** was identified as a psammaplin B, which was previously isolated from marine sponges *Psammaplysilla purpurea*¹² and *Pseudoceratina purpurea*.¹¹ Compound **5** was identified as psammaplin D, which was previously isolated from a marine sponge *P. purpurea*.¹² Compound **6** was identified as psammaplin E, previously isolated from a marine sponge *Pseudoceratina purpurea*.¹¹ Compound **7** was identified as (*E,E*)-psammaplin A, the well-known metabolite of marine sponges *Psammaplysilla* sp.,^{12,14} *Thorectopsamma xana*,¹⁵ *Pseudoceratina purpurea*,¹¹ *Aplysinella rhax*.^{16–18} Compounds **8** and **9** were identified as (*E,Z*)-psammaplin A and bromopsammaplin A, respectively, which were previously isolated from the previous collection of the same specimen.⁴ Compound **10** was identified as a sodium salt of psammaplin A sulfate, which was reported from a marine sponge *A. rhax*.¹⁶ Compound **11** was a known metabolite of several marine sponges.^{4,12,13} Compound **12** was identified as a bis-*N*, *N*-dimethylguanidium salt of psammaplin A sulfate, which was previously isolated from a marine sponge *A. rhax*.¹⁸ This is the first report on the isolation of compounds **3**, **4**, **6**, **10**, and **12** from an association of sponges *Jaspis* sp. and *Poecillastra* sp. Compounds **13** (bispsammaplin A) and **14** (bisaprasin A) were reported in our previous letter.⁴

Many bromotyrosine-derived metabolites with diverse structural features have been reported from marine sponges. Apart from sponges, there is only one report on the isolation of bromotyrosine derivatives from an ascidian *Botryllus* sp.¹⁹ Bromotyrosine alkaloids are considered as the chemotaxonomic markers for the sponges belonging to the order Verongida.²⁰ However, they are also encountered in sponges belonging to other orders, such as *Pachychalina*,²¹ *Oceanapia*,²² (both genera belong to the order Haplosclerida), *Latrunculia*,²³ *Iotrochota*²⁴ (both genera belong to the order Poecilosclerida), and a sponge of order Dendroceratida, *Dendrilla cactus*.²⁵ Orders, Verongida, Haplosclerida, Poecilosclerida, and Dendroceratida belong to the subclass Ceractinomorpha, but sponges *Jaspis* and *Poecillastra* belong to the order Astrophorida (subclass Tetractinomorpha). To the best of our knowledge, two-sponge association of *Jaspis* sp. and *Poecillastra* sp. is the only sponge specimen from the subclass Tetractinomorpha containing bromotyrosine alkaloids.^{3,4}

The wide spectrum of biological activities has been reported for psammaplin derivatives.^{26–34} The cytotoxicity of the isolated compounds (**1–4**, **6**, **10**, and **12**) against A-549, SK-OV-3, SK-MEL-2, XF-498, and HCT-15 solid tumor cell lines was studied (Table 2). In general, the dimeric forms (**7**, **9**, **10**, and **12**) exhibited more potent cytotoxicity to cancer cell lines. Dimers containing trisubstituted phenyl rings at both ends (**7** and **10**) displayed the most potent cytotoxicity. The diguanidium salt of psammaplin A sulfate (**12**) showed a rather weak cytotoxicity but was selective against human ovarian cancer cells (SK-OV-3). It appears that monomeric forms containing amide terminal group (**5** and **6**) also show high potency. The disulfide moiety in these bromotyrosine derivatives also plays an important role as monomers (**2–4**) lacking it exhibited less potency. In case of tetramers, ether forms (**1** and **13**) were twice as potent as compared to condensed form (**14**).

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7. The sponges, collected in November 2002, offshore Jeju Island, South Korea, were frozen immediately after collection and stored at 20 °C until extraction. The frozen animal material was cut into small pieces and extensively extracted with MeOH at room temperature. The MeOH extract was partitioned between CH₂Cl₂ and H₂O, and the CH₂Cl₂ layer was further partitioned between 90% aq MeOH and *n*-hexane. The aqueous MeOH fraction was selected for further separation on the basis of its lethality to brine shrimp larvae and subjected to a reversed-phase flash column chromatography (YMC Gel ODS-A, 60 Å 500/400 mesh), eluting with gradient solvent system of 50–100% MeOH/H₂O to yield 19 fractions. Fractions containing bromotyrosines (1–8) were selected for further separation from their potent activity (LD₅₀ ~20 µg/mL) in the brine shrimp lethality assay and were subjected to repeated reversed-phase chromatographic separation to afford 12 compounds. Compound **1** was obtained by purification of fraction 8 by reversed-phase HPLC. Fraction 1 was subjected to MPLC, eluting with gradient solvent system of 0–70% MeOH/H₂O to yield 10 subfractions. These subfractions were subjected to reversed-phase HPLC to obtain compounds **2–5** and **7–11**. The H₂O layer was also abundant in psammaphin derivatives. The H₂O layer was partitioned between *n*-BuOH and H₂O. The *n*-BuOH layer was subjected to MPLC to yield 12 fractions (1–12). Compounds **6** and **12** were obtained by purification of fraction 3 by repeated HPLC separation.
8. **Cyclobispsammaphin A (1)**: Amorphous, white solid; LRESIMS (+ve mode) *m/z* 1343, 1345, 1347, 1349, 1351 [M+Na]⁺, (–ve mode) *m/z* 1319, 1321, 1323, 1325, 1327 [M–H][–]; HRMALDI-TOFMS *m/z* 1344.8854, 1346.8811, 1348.8965, 1350.8937 [M+Na]⁺ (Calcd for C₄₄H₄₄N₈O₁₂S₄⁷⁹Br₂⁸¹Br₂Na, 1346.8551); ¹H and ¹³C NMR data, see Table 1.
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10. **Psammaphin M (2)**: Amorphous, white solid; LRFABMS (+ve mode) *m/z* 381/383 [M+Na]⁺; HRFABMS (–ve mode) *m/z* 357.0070 [M–H][–] (Calcd for C₁₃H₁₄N₂O₅Br, 357.0086); ¹H NMR (CD₃OD, 500 MHz) δ 7.35 (1H, d, *J* = 2.0 Hz, H-2), 7.05 (1H, dd, *J* = 8.0, 2.0 Hz, H-6), 6.75 (1H, d, *J* = 8.0 Hz, H-5), 3.77 (2H, s, H-7), 3.63 (3H, s, OCH₃), 3.48 (2H, t, *J* = 7.0 Hz, H-11), 2.53 (2H, t, *J* = 7.0 Hz, H-12); ¹³C NMR (CD₃OD, 500 MHz, based on HMBC and HSQC data) δ 173.5 (C-13), 165.5 (C-8), 153.5 (C-4), 153.0 (C-8), 134.2 (C-2), 130.6 (C-1, C-6), 117.0 (C-5), 110.0 (C-3), 52.5 (OCH₃), 35.7 (C-11), 34.3 (C-12), 28.2 (C-7).
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