

Tauroacidins A and B, New Bromopyrrole Alkaloids Possessing a Taurine Residue from *Hymeniacidon* Sponge

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Abstract: Two new bromopyrrole alkaloids, tauroacidins A (1) and B (2), with tyrosine kinase inhibitory activity have been isolated from an Okinawan marine sponge *Hymeniacidon* sp. and the structures were elucidated on the basis of spectral data and chemical means. Tauroacidins A (1) and B (2) are rare bromopyrrole alkaloids possessing a taurine residue. © 1997 Elsevier Science Ltd.

During our search for bioactive substances from marine organisms,¹ we have investigated extracts of an Okinawan marine sponge *Hymeniacidon* sp. and isolated two new bromopyrrole alkaloids possessing a taurine residue, tauroacidins A (1) and B (2), with tyrosine kinase inhibitory activity. Here we describe the isolation and structure elucidation of 1 and 2.

n-BuOH-soluble materials of MeOH extract of the sponge collected off Ishigaki Island, Okinawa, were subjected to Sephadex LH-20 (MeOH) and C₁₈ columns (CH₃CN/H₂O/CF₃CO₂H) followed by C₁₈ HPLC (MeOH/H₂O/CF₃CO₂H) to yield tauroacidins A (1, 8 x 10⁻⁵ %, wet weight) and B (2, 5 x 10⁻⁵ %).

HRESIMS [*m/z* 526.9164, (M-H)⁻, Δ +0.9 mmu] of tauroacidin A [1, [α]_D²⁸ -4.3° (c 0.15 MeOH)], indicated the molecular formula to be C₁₃H₁₆N₆O₅SB₂. The IR spectrum of 1 suggested the presence of OH and/or NH (3420 cm⁻¹), amide carbonyl (1690 cm⁻¹), and sulfonate groups (1210 and 1040 cm⁻¹).² The UV absorption [λ_{max} 273 nm (ε 9700)] was attributable to a substituted pyrrole chromophore.³ The ¹H and ¹³C NMR data showed signals due to a 2,3-dibromopyrrole carbonyl moiety.⁴ The ¹³C chemical shifts at C-2' (δ_C 39.8) and C-3' (δ_C 49.1) corresponded well to those of the taurine residue (ca. δ_C 39 and 48, respectively) of melemeleone A.⁵ The presence of the taurine residue in 1 was also supported by amino acid analysis of the acid hydrolysis products of 1. The whole structure of tauroacidin A (1) was elucidated by analyses of 2D NMR data [¹H-¹H COSY, NOESY, and decoupled HMBC (D-HMBC)⁶] (Fig. 1). The ¹H-¹H COSY spectrum of 1 showed proton networks from NH-7 to H-10 and OH-9 and from NH-1' to H₂-3'. The presence of the aminoimidazole ring including three quaternary carbons (C-11, C-13, and C-15) was revealed

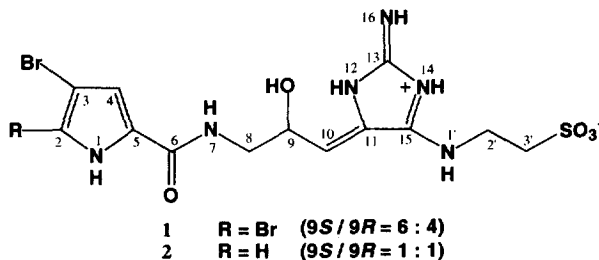
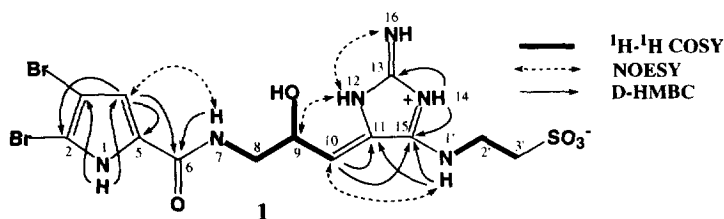


Table 1. ^1H and ^{13}C NMR Data of Tauroacidins A (**1**) and B (**2**) in $\text{DMSO}-d_6$.

positn.	1		2	
	δ_{H} (m, J^a)	δ_{C} (m)	δ_{H} (m, J^a)	δ_{C} (m)
1.	12.70 (s)		11.84 (br.s)	
2.		104.7 (s)	6.99 (br.s)	121.3 (d)
3.		97.8 (s)		94.9 (s)
4.	6.95 (s)	113.0 (d)	6.87 (br.s)	111.8 (d)
5.		127.9 (s)		126.7 (s)
6.		159.1 (s)		159.9 (s) ^d
7.	8.22 (t, 6.0)		8.20 (t, 6.2)	
8.	3.32 (m) ^{b,c}	44.5 (t)	3.34 (m) ^{b,c}	44.3 (t)
9.	4.61 (m)	67.3 (d)	4.62 (m)	67.2 (d)
9-OH.	5.94 (d, 4.2)		5.95 (br.s)	
10.	6.17 (d, 4.2)	116.4 (d)	6.18 (m)	116.5 (d)
11.		131.2 (s)		131.1 (s)
12.	9.31 (br.s)		9.32 (br.s)	
13.		165.9 (s)		165.8 (s) ^c
14.	10.51 (s)		10.50 (s)	
15.		167.6 (s)		167.6 (s) ^c
16.	8.05 (br.s)		8.05 (br.s)	
1'	9.64 (br.t, 3.0)		9.65 (m)	
2'	3.67 (dt, 3.0, 7.2) ^c	39.8 (t)	3.67 (m) ^c	39.5 (t) ^e
3'	2.75 (t, 7.2) ^c	49.1 (t)	2.74 (t, 7.5) ^c	49.2 (t)

a) in Hz. b) These signals were overlapped with H_2O signal. c) 2H. d) These signals were assigned on the basis of D-HMBC correlations. e) This signal was overlapped with DMSO signal.

by the D-HMBC cross-peaks for H-10/C-15, NH-14/C-13, and NH-14/C-15. D_2O -Exchangeable protons at δ_{H} 9.31 (NH-12) and 8.05 (NH-16) were assigned by NOESY correlations for H-9/NH-12 and NH-12/NH-16. Attachment of the pyrrole ring to N-7 via an amide bond was deduced from the NOESY cross-peak for H-4/NH-7 and D-HMBC correlations for H-4/C-6 and NH-7/C-6. The NOE for NH-1'/H-10 indicated *Z*-geometry of the double bond at C-10. The D-HMBC cross-peaks for NH-1'/C-11 and NH-1'/C-15 and the NOE for NH-1'/H-10 implied that the taurine residue was attached to C-15 of the imidazole ring. The negative ion FAB MS/MS spectrum of the pseudomolecular ion (m/z 527) of **1** showed several product ions supporting the proposed structure (Fig. 2). Thus the structure of tauroacidin A was concluded to be **1**. In order to determine the absolute configuration at C-9, tauroacidin A (**1**) was subjected to ozonolysis and then hydrolysis.⁷ Amino acid analysis of the acid hydrolysate showed the presence of one molar equivalent of isoserine. The hydrolysate was treated with $\text{HCl}/i\text{-PrOH}$ and then (*S*)-MTPACl to afford the *N,O*-bis-(*R*)-MTPA/*i*-propyl ester of isoserine, of which HPLC analyses revealed the presence of (*S*)- and (*R*)-isoserines in the ratio of ca. 6:4, showing that tauroacidin A (**1**) was a 6:4 mixture of 9*S*- and 9*R*-isomers.

Fig. 1. 2D NMR Correlations for Tauroacidin A (**1**)

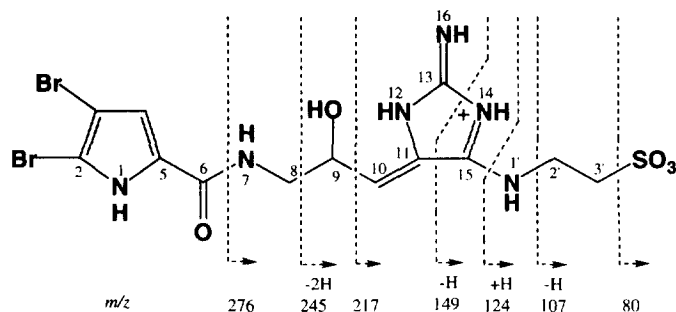


Fig. 2. Negative FABMS/MS Fragmentations of m/z 525 from Tauroacidins A (1)

The molecular formula, $C_{13}H_{17}N_6O_5SBr$, of tauroacidin B (2), optically inactive, was suggested by HRESIMS. The 1H and ^{13}C NMR data of 2 differed from 1 only in the presence of an sp^2 methine signal [H-2: δ_H 6.99 (m), C-2: δ_C 121.3 (d, $^1J_{CH} = 185$ Hz)], indicating the presence of 3-bromopyrrole carbonyl moiety^{4,8} in 2. Thus the structure of tauroacidin B (2) was elucidated to be 2-debromo form of 1. Tauroacidin B (2) was revealed to be racemic ($9S/9R = 1:1$) at C-9 from HPLC analyses of the *N,O*-bis-(*R*)-MTPA/*i*-propyl ester of isoserine contained in the hydrolysate of 2 obtained by the same procedure as 1.

Tauroacidins A (1) and B (2) are new bromopyrrole alkaloids possessing a taurine residue attached to the aminoimidazole ring, which may be biogenetically related to mauritamide A, known bromopyrrole alkaloids possessing a taurine residue at C-11 from sponges.⁹ Tauroacidins A (1) and B (2) exhibited inhibitory activity against EGF receptor kinase and *c-erbB-2* kinase (IC_{50} , 20 $\mu g/mL$ each).

EXPERIMENTAL

General Procedure. Optical rotations were recorded on a JASCO DIP-360 polarimeter. The IR and UV spectra were taken on a JASCO FT/IR-5300 and a JASCO Ubest-35 spectrophotometers, respectively. 1H and ^{13}C NMR spectra were recorded on a Bruker AMX-600 and a JEOL EX-400 spectrometers, respectively. ESI mass spectra were obtained on a JEOL SX-102A spectrometer. Standard amino acid analyses were performed with Hitachi amino acid autoanalyzer Model 835.

Sponge Materials. The sponge *Hymeniacidon* sp. (order Halichondrida; family Halichondriidae) was collected off Ishigaki Island, Okinawa, and kept frozen until used. Soft and compressible sponge is somewhat fleshy texture. No ectosomal skeleton is just a thickening in this region of the mesohyl. Spicules are in loose tracts or without orientation. Spicules are styles 439 x 12 μm that are curved along the upperthird of the shaft. The voucher specimen (SS-361) was deposited at the Faculty of Pharmaceutical Sciences, Hokkaido University.

Extraction and Isolation. The sponge (5.5 kg, wet weight) was extracted with methanol (3 L x 2). Part (100 g) of the methanolic extract (335.8 g) was partitioned between ethyl acetate (500 mL x 3) and water, and the aqueous layer was extracted with *n*-butanol (1 L x 3). The *n*-butanol soluble material (18.6 g) was subjected to a Sephadex LH-20 column (MeOH), a C_{18} column (Develosil ODS-LOP, Nomura Chemical, 45 x 490 mm; $CH_3CN/H_2O/CF_3CO_2H$, 20:80:0.1), and C_{18} HPLC (Develosil ODS-HG-5, Nomura Chemical, 10 x 250 mm; $MeOH/H_2O/CF_3CO_2H$, 46:54:0.1; flow rate, 2.5 mL/min; UV detection at 260 nm) to yield tauroacidins A (1, 8×10^{-5} % wet weight, t_R 30 min) and B (2, 5×10^{-5} %, t_R 18 min).

Tauroacidin A (1). A colorless amorphous solid; $[\alpha]_D^{28} -4.3^\circ$ (c 0.15, MeOH); UV (MeOH) λ_{max} 273 (ϵ 9700) and 312 nm (3800); IR (film) ν_{max} 3420 (br), 1690, 1630, 1210, and 1040 cm^{-1} ; 1H and ^{13}C NMR (see Table 1); ESIMS (Neg., MeOH) m/z 525, 527, and 529 [(M-H)⁻, ca. 1:2:1]; HRESIMS m/z

526.9164 (M-H)⁻, calcd for C₁₃H₁₅N₆O₅S⁷⁹Br⁸¹Br, 526.9173.

Tauroacidin B (2). A colorless amorphous solid; [α]_D²⁸ 0° (*c* 0.10, MeOH); UV (MeOH) λ_{\max} 272 (ϵ 10000) and 310 nm (4000); IR (film) ν_{\max} 3420 (br), 1690, 1630, 1210, and 1040 cm⁻¹; ¹H and ¹³C NMR (see Table 1); ESIMS (Neg., MeOH) *m/z* 447 and 449 [(M-H)⁻, ca. 1:1]; HRESIMS *m/z* 447.0098 (M-H)⁻, calcd for C₁₃H₁₆N₆O₅S⁷⁹Br, 447.0088.

Hydrolysis of Tauroacidins A (1). Tauroacidin A (**1**, 0.1 mg) was treated with 6 N HCl (100 μ L) at 110 °C for 24 h. After evaporation of the solvent, standard amino acid analysis of the reaction mixture showed the presence of 1 mole of taurine.

Absolute Stereochemistry at C-9 of Tauroacidins A (1) and B (2). A solution of tauroacidin A (**1**, 0.2 mg) in MeOH (100 μ L) was treated with ozone at -78 °C for 2 min. After excess ozone was removed by N₂ gas, to the residue were added HCO₂H (100 μ L) and 35 % H₂O₂ aq (100 μ L). The mixture was stirred for 1 h at 0 °C and then 5 h at room temperature. After evaporation, the residue was hydrolyzed with 6 N HCl (100 μ L) at 110 °C for 24 h in a sealed tube. Standard amino acid analysis of the hydrolysate showed the presence of 1 mole each of taurine and isoserine. The hydrolysate was treated with 9 % HCl/*i*-PrOH (100 μ L) at 110 °C for 30 min, and then treated with CH₂Cl₂ (100 μ L), 4-(*N,N*-dimethylamino)pyridine (0.1 mg), Et₃N (10 μ L), and (*S*)-MTPACI (5 μ L) at 40 °C for 2 h. After addition of *N,N*-dimethylpropane-1,3-diamine (5 μ L), the reaction mixture was subjected to HPLC analysis [YMC Pack SIL-06, 4.6 x 250 mm; flow rate, 1 mL/min; UV detector at 240 nm; eluent, hexane/*i*-PrOH, 95:5]. Retention times of *N,O*-bis-(*R*)-MTPA/*i*-propyl ester derivatives of authentic (*S*)- and (*R*)-isoserine were 13.4 and 16.3 min, respectively. Both (*S*)- and (*R*)-isoserine derivatives were found in the hydrolysate of **1** in the ratio of ca. 6:4. HPLC analysis of *N,O*-bis-(*R*)-MTPA-isoserine *i*-propyl ester derived from the hydrolysate of tauroacidin B (**2**) was performed under the same condition as described above, and both (*S*)- and (*R*)-isoserine derivatives were found in the ratio of ca. 1:1.

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