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Sagittamides A and B. Polyacetoxy Long-Chain Acyl Amino Acids from a Didemnid Ascidian

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ABSTRACT

An unidentified tunicate from Pohnpei, Micronesia, yielded sagittamides A and B—compounds comprising a long-chain C₂₆ dicarboxylic acid that acylates terminal L-valine and L-ornithine groups. The structures, which contain an unprecedented internal *O*-hexacetyl-1,2,3,4,5,6-hexaol moiety, were determined by combined spectroscopic analysis including mass spectrometry and 1D and 2D NMR and chemical degradation. The partial absolute stereochemistry of the new compounds was addressed by Marfey's analysis.

Ascidians (subphylum Urochordata, class Ascidiacea) produce a variety of natural products, typically highly modified peptides. Bioactive ascidian metabolites include the antitumor alkaloids ecteinascidin-743 from *Ecteinascidia turbinata*, diazonamide A from *Diazona angulata* (cf. *chinensis*), the highly cytotoxic depsipeptides didemnin A and B, and the antiviral eudistomins. Through the course of screening of Micronesian marine invertebrates for the presence of inhibitors of histone deacetylases (HDACs), we found that the extract of an unidentified Didemnid ascidian elicited significant inhibition of HDAC (98% inhibition at 4.0 μ g/mL) and contained a series of unprecedented polyacetoxylated long-chain acyl amino acids. The major compounds,

An unidentified Didemnid compound ascidian, collected from Arrow Island, Pohnpei, in 2001 (Federated States of Micronesia), was exhaustively extracted with 1:1 CH₂Cl₂/MeOH and the concentrated extract sequentially partitioned against hexanes, CHCl₃, and *n*-BuOH. Examination of both the CHCl₃- and *n*-BuOH-soluble fractions by ¹H NMR revealed a large component consisting of acetylated polyhydroxylipids. Purification of the polyacetate containing fractions by Sephadex LH-20 chromatography, followed by reversed-phase HPLC (C₁₈, 1:1 CH₃CN/H₂O, 0.05% TFA), gave sagittamide A (1) and B (2) as optically active colorless glasses.⁶

Compound **1** was soluble only in polar solvents (MeOH, DMSO) and gave a strong ninhydrin-positive reaction (pink color) indicating the presence of a primary amine. Acid

which we named sagittamides A (1) and B (2), are characterized by an α , ω -dicarboxylic acid that is acylated at each terminus by a different amino acid. In this paper, we describe the isolation and structural characterization of 1 and 2, including the absolute configuration of the amino acids.

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⁽⁶⁾ The name "sagittamide" is derived from the Latin word for "arrow".

hydrolysis of 1 liberated 1 equiv each of the amino acids, ornithine and valine (vide infra).

MALDI-HRMS of 1^7 gave an m/z 988.5585 ([M + H]⁺, Δ mmu -0.3) corresponding to a molecular formula of $C_{48}H_{81}N_3O_{18}$ for the parent compound. The UV spectrum of 1 showed only weak end absorption; however, the IR spectrum showed ester and amide carbonyl stretching (ν 1750, 1654 cm⁻¹, respectively). The 1 H NMR (CD₃OD, Table 1) indicated six acetate methyl singlets, a pair of methyl doublets, and multiple oxygenated methines. ESIMS of 1 under conditions of deuterium exchange (CD₃OD) gave m/z 995.6 [M_D + D]⁺ indicating six exchangeable hydrogens. The remaining unresolved 1 H NMR signals (δ 1.28, bs) were attributable to long unbranched CH₂ chains.

Analysis of HSQC, HMBC, and COSY correlations in methanol- d_4 provided the partial structures $1\mathbf{a} - \mathbf{c}$ shown in Figure 1. HMBC correlations (DMSO- d_6 , see Supporting

Figure 1. Partial structures of 1 with selected HMBC correlations.

Information) from the amide NH protons to C=O groups allowed placement of the valinyl and ornithinyl residues at the respective carboxyl groups at opposite ends of the long chain.

The presence of two free carboxyl groups in **1** was confirmed by treating the compound with excess diazomethane to give a monomethyl ester (1 H NMR, δ 3.71 s, 3H), instead of an expected diester. ESIMS measurement of the product (m/z 1006.5, [M + Na] $^{+}$) indicated a net increase of mass of only 4 amu instead of the expected 28 amu. HRMS-MALDI (m/z 1006.5467, [M + Na] $^{+}$) revealed the formula of the product, $C_{49}H_{81}N_{3}O_{17}$, and confirmed that the expected diester product had cyclized to give **3** with loss of

the elements of MeOH. Comparison of ESIMS measurements of **3** prepared in CH₃OH (m/z 1006.5, [M + Na]⁺) and CD₃OD (m/z 1009.6, [M_D + Na]⁺) confirmed the expectation of only three exchangeable hydrogens. The presence of three amide protons (δ 6.01, d, J = 8.8 Hz, 1H; 6.02, br s, 1H; 6.35, br s, 1H) in the ¹H NMR spectrum (CDCl₃, Table 1) was consistent with a monomethylester-triamide structure **3**. Evidently, the incipient dimethylester i (Figure 2), formed

Figure 2. Reaction of 1 with diazomethane.

by addition of diazomethane at both C1' and C1'' carboxyl groups, undergoes spontaneous intramolecular addition—elimination at the ornithine terminus by the free gamma-NH₂ group to generate lactam 3.

The second major component, 2, was shown to be di-desacetyl-1 from the following analysis. MALDI-HRMS of 2^8 gave an $[M + Na]^+$ of 926.5237 ($\Delta mmu + 3.5$) corresponding to the molecular formula C₄₄H₇₇N₃O₁₆ for 2. The difference in mass of 84 amu from 1 and only four AcO methyl singlets in the ¹H NMR spectrum indicated that 2 was a didesacetyl derivative of 1. ESIMS under conditions of deuterium exchange (CD₃OD) gave an [M_D + D]⁺ of m/z 913.6 indicating the presence of eight exchangeable hydrogens. Two CH-O signals in the ¹H NMR of 2 $(\delta 3.95, dd, J = 9.7, 1.5 Hz; 4.15, dd, J = 10.1, 1.4 Hz)$ now appeared $\sim 1-1.5$ ppm upfield with respect to the clustered CH-OAc signals in 1. These upfield-shifted signals, thus, belonged to the two unesterified secondary CH-OH groups in 2 and were subsequently assigned to H7 and H9 by the following sequential vicinal coupling analysis. $^{1}\mathrm{H}^{-1}\mathrm{H}$ ^{3}J couplings between H5–H6 (J=2.6 Hz), H6– H7 (J = 10.1 Hz), H7-H8 (J = 1.4 Hz), H8-H9 (J =9.7 Hz), and H9-H10 (J = 1.5 Hz) were identified from a small-J-optimized delayed COSY experiment (J = 2.0 Hz) and supported by HSQC-TOCSY from H5 to C3 and C4 and H10 to C11 and C12.

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⁽⁷⁾ Sagittamide A (1): clear glass; $[\alpha]^{25}_{D}$ –22.0 (c 0.599, MeOH); IR (ZnSe) ν_{max} 3314, 2925, 2853, 1750, 1654, 1540, 1371, 1220, 1032 cm⁻¹; UV (MeOH) λ_{max} 214 nm (ϵ +0.38), 228 (5481), 276 (1060); CD (MeOH) λ_{max} 212 nm ($\Delta\epsilon$ 2301.7), 242 (–0.03); ¹H NMR and ¹³C NMR (Table 1); HRMALDIMS m/z 988.5585 (M + H⁺), calcd for C₄₈H₈₂N₃O₁₈ 988.5588. The compound, most likely, is in the form of the TFA salt after HPLC purification.

⁽⁸⁾ Sagittamide B (2): clear glass; $[\alpha]^{25}_{\rm D}-19.7$ (c 0.173, MeOH); IR (ZnSe) $\nu_{\rm max}$ 3319, 3061, 2926, 2854, 1722, 1680, 1543, 1373, 1242, 1207, 1140, 1034 cm⁻¹; UV (MeOH) $\lambda_{\rm max}$ 214 (ϵ 1534), 256 (702); ¹H and ¹³C NMR (Table 1); HRMALDIMS m/z 926.5237 [M + Na]⁺, calcd for C₄₄H₇₇N₃O₁₆Na 926.5202.

Table 1. ¹H and ¹³C NMR Data for 1 and 2 in CD₃OD and ¹H NMR Data for 3 in CDCl₃

	1		2		3
no.	$\delta_{ m H}{}^a$	$\delta_{ ext{C}}{}^{b}$	$\delta_{ m H}{}^a$	$\delta_{ ext{C}}{}^{c}$	$\delta_{\rm H}{}^{j}$
1		175.9			
2	2.27 (2H, m)	36.3	2.30 (2H, m)	36.0	2.27 (2H, m)
3a	1.52 (1H, m)	23.0	1.58 (1H, m)	22.7	1.65 (2H, m)
3b	1.67 (1H, m)		1.69 (1H, m)		
4a	1.59 (1H, m)	29.1	1.73 (2H, m)	27.8	1.65 (2H, m)
4b	1.63 (1H, m)				
5	4.92 (1H, m)	73.4	5.22 (1H, m)	74.1	4.81 (1H, m)
6	5.11 (1H, dd, 8.3, 4.0)	71.4	4.94 (1H, dd, 10.1, 2.6)	72.0	5.11 (1H, dd, 8.8, 2.8)
7	5.15 (1H, dd, 8.3, 1.5)	69.2	4.15 (1H, dd, 10.1 1.4)	68.2	5.15 (1H, dd, 8.8, 1.7)
8	5.42 (1H, dd, 9.9, 1.5)	68.4	4.99 (1H, dd, 9.7, 1.4)	71.2	5.41 (1H, dd, 9.8, 1.7)
9	5.13 (1H, dd, 9.9, 1.8)	70.4	3.95 (1H, dd, 9.7, 1.5)	69.0	5.13 (1H, dd, 9.8, 2.0)
10	4.82 (1H, td, 6.8, 1.8)	71.9	4.76 (1H, td, 6.8, 1.5)	73.2	4.81 (1H, m)
11	1.42 (2H, m)	31.8	1.42 (2H, m)	31.7	1.35 (2H, m)
12	1.32 (2H, m)	26.3	1.32 (2H, m)	26.3	1.26 (2H, m)
13 - 23	1.28 (22H, br s)	30.5	1.28 (22H, br s)	30.4	1.20 (22H, br s)
24a	1.63 (1H, m)	27.1	1.60 (2H, m)	26.8	1.65 (2H, m)
24b	1.59 (1H, m)				
25	2.24 (2H, m)	37.0	2.24 (2H, m)	36.8	2.27 (2H, m)
26		176.6			
1'		175.0			
2'	4.31 (1H, d, 5.6)	59.1	4.31 (1H, d, 6.0)	59.0	4.53 (1H, dd, 8.8, 4.8)
3'	2.15 (1H, hep, 6.8)	31.8	2.17 (1H, hep, 6.8)	31.3	2.15 (1H, m)
4'	0.96 (3H, d, 6.8)	18.5	0.97 (3H, d, 6.8)	18.2	0.92 (3H, d, 6.8)
5'	0.97 (3H, d, 6.8)	19.8	0.98 (3H, d, 6.8)	19.2	0.89 (3H, d, 7.2)
1"		174.9			
2"	4.42 (1H, dd 8.2, 5.0)	52.9	4.43 (1H, dd, 8.0, 5.2)	53.2	4.26 (1H, dt, 11.3, 5.6)
3a"	1.75 (1H, m)	29.9	1.75 (1H, m)	29.9	1.50 (1H, m)
3b"	1.98 (1H, m)		1.98 (1H, m)		2.57 (1H, m)
4"	1.72 (2H, m)	25.3	1.75 (2H, m)	25.0	1.93 (2H, m)
5"	2.95 (2H, t, 7.2)	40.4	2.95 (2H, t, 7.0)	39.9	3.34 (2H, m)
5-OAc	2.01 (3H, s)	21.0, 172.4	$1.95 (3H, s)^f$	$20.5, 171.5^g$	$2.11 (3H, s)^h$
6-OAc	2.02 (3H, s)	21.1^d , 171.7	$1.97 (3H, s)^f$	$20.5, 171.7^g$	$2.09 (3H, s)^h$
7-OAc	2.07 (3H, s)	20.9^e , 172.0			$2.05 (3H, s)^h$
8-OAc	2.01 (3H, s)	21.0, 171.5	$1.98 (3H, s)^f$	$20.5, 171.8^g$	$2.02 (3H, s)^h$
9-OAc	2.08 (3H, s)	21.0^e , 171.7			$2.02 (3H, s)^h$
10-OAc	2.03 (3H, s)	21.2^d , 172.6	$1.99 (3H, s)^f$	$20.5,172.0^{g}$	$2.01 (3H, s)^h$
OMe					3.71 (3H, s)
NH-2′					6.01 (1H, d, 8.8)
NH-2"					6.35 (1H, br s)
NH-5″					6.02 (1H, br s)

^a Spectra recorded at 400 MHz referenced to residual CD₂HOD δ = 3.30. ^b Spectra recorded at 100 MHz and referenced to CD₃OD δ = 49.0. ^c Assigned from HSQC and HMBC (400 MHz). ^{d-h} Signals are interchangeable. ^j Recorded in CDCl₃ (400 MHz) and referenced to CHCl₃ δ = 7.24.

The identities and absolute stereochemistry of the amino acid residues in **1** were determined by Marfey's analysis. Acid hydrolysis of **1** (6 M HCl, 16 h, 110 °C) and derivatization of the products with 2,4-dinitrophenyl-5-fluoroL-alaninamide (Marfey's reagent) under standard conditions, followed by analysis (C_{18} HPLC-MS, UV detection), gave two peaks. The early-eluting peak ($t_R = 20.9$ min) matched the Marfey's derivative of L-valine, while the lateluting peak ($t_R = 24.7$ min) was identified as the Marfey's derivative of L-ornithine.

Sagittamides represent an unexpected departure from the structures of other α,ω -long-chain bifunctionalized natural

contiguous, stereoregular hexaol segment that constitutes C5-10 is unknown. Two possible explanations for the origin of the hexaol segment in **1** and **2** are polyketide synthase mediated chain extension by three iterative additions of hydroxymalonyl CoA, followed by ketone reductions to the growing lipid chain or incorporation of an unidentified C₆ carbohydrate unit. The latter is unprecedented while the former is rare. Hydroxymalonyl CoA (HMCoA) polyketide chain extension is known in the biosynthesis of avermectin¹² and genetic evidence supports its involvement in zwittermicin A production by *Bacillus cereus*, ¹³ but we are unaware of examples of polyketide assembly from *iterative* HMCoA

products from marine organisms.¹¹ The biosynthesis of the

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chain extensions. The stereochemistry of the hexa-acetoxy-lated stereoelement at C5-C10 of **1** and **2** is presently unassigned and is clearly a challenging problem.¹⁴

Pure 1 was not active in an antitumor cell assay (human colon tumor, HCT-116, IC₅₀ > 100 μ g/mL¹⁵) and did not

inhibit HDAC's (IC₅₀ > 100 μ g/mL). The active fraction from the original extract is now under investigation.

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Supporting Information Available: Color image of the Didemnid ascidian (01-133, underwater), 1D ¹H NMR for **1**–**3**, ¹³C NMR for **1**, and COSY, HSQC, and HMBC spectra for **1** (CD₃OD and DMSO-*d*₆) and **2** (CD₃OD). This material is available free of charge via the Internet at http://pubs.acs.org.

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⁽¹⁰⁾ A solution of 1 (0.50 mg) in deionized water (200 μ L, MilliQ) and hydrochloric acid (12 M, 200 μ L) was sealed in a screw-thread pressure vial and heated at 110 °C for 16 h. After removal of the volatiles, the residue was redissolved in deionized water (100 μ L), and the solution centrifuged to remove insoluble material. The supernatant was mixed with Marfey's reagent (200 μL, 1% solution of 2,4-dinitrophenyl-5-fluoro-L-alaninamide in acetone) and aqueous 1.0 M NaHCO₃ (20 μL) and then heated at 85 °C for 12 min and cooled before quenching (1.0 M HCl, 10 μ L). LCMS of this solution (4 μ L injection, Phenomenex Luna C₁₈ column, 2.0 \times 100 mm) used a solvent gradient (90:10 H₂O/CH₃CN to 35:65 H₂O/CH₃CN over 40 min, flow rate = 0.4 mL/min), and monitoring was done by UV (λ 210, 254, 285, and 340 nm) and ESIMS (negative ion). Two peaks, corresponding to Marfey's peaks of L-valine and L-ornithine, eluted at t_R = 20.9 and 24.7 min, respectively. Marfey's derivatives of paired standards of L- and D-amino acids (1 μ g/ μ L), prepared in the same way, eluted at the following respective retention times: D- and L-lysine ($t_R = 26.2, 27.4 \text{ min}$) D- and L-ornithine ($t_R = 23.3, 24.9 \text{ min}$), L- and D-proline ($t_R = 16.8, 17.8 \text{ min}$) min), and L- and D-valine ($t_R = 20.9, 24.3 \text{ min}$).

⁽¹¹⁾ Recent examples of long-chain α,ω-bifunctionalized lipids include the C₂₈ sphingolipids from sponges [oceanapiside from *Oceanapia phillipensis*: (a) Nicholas, G. M.; Hong, T. W.; Molinski, T. F.; Lerch, M. L.; Cancilla, M. T.; Lebrilla, C. B. *J. Nat. Prod.* **1999**, 62, 1678–1681. (b) Nicholas, G. M.; Molinski, T. F. *J. Am. Chem. Soc.* **2000**, *122*, 4011–4019], rhizochalin from *Rhizochalina incrustata* [(c) Makarieva, T. N.; Denisenko, V. A.; Stonik, V. A. *Tetrahedron Lett.* **1989**, *30*, 6581–6584. (d) Molinski, T. F.; Makarieva, T. N.; Stonik, V. A. *Angew. Chem. Intl. Ed.* **2000**, *39*, 4076–4079], calyxoside from a *Calyx* sp. [(e) Zhou, B.-N.; Mattern, M. P.; Johnson, R. K.; Kingston, D. G. I. *Tetrahedron* **2001**, *57*, 9549–9554], and ene—yne lipids from *Haliclona* sp. [(f) Zhou, G.-X.; Molinski, T. F. *Mar. Drugs* **2003**, *1* (1), 46–53 and references cited within].

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⁽¹⁴⁾ Should the biosynthesis of the hexaol derive from a carbohydrate fragment, it is tempting to speculate that the relative configuration of C5–C10 in 1 and 2 may follow the stereogenicity of a natural aldohexose.

⁽¹⁵⁾ Viability assays were carried out on cultured HCT-116 tumor cells using a *soluble* MTS formazan dye end point. For details, see ref 11f.