



## Hexamollamide, a hexapeptide from an Okinawan ascidian *Didemnum molle*

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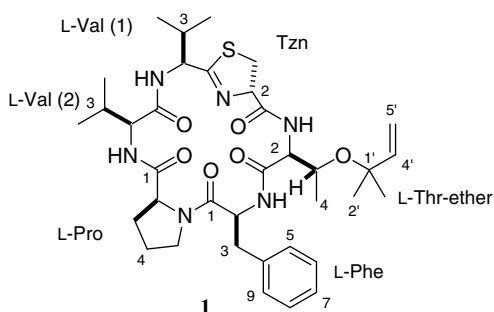
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### ABSTRACT

The bioassay-guided fractionation of the cytotoxic constituents of the Okinawan ascidian *Didemnum molle* led to the isolation of hexamollamide (**1**), a hexapeptide. The gross structure and relative stereostructure of **1** were established by spectroscopic analysis including 2D NMR techniques and single-crystal X-ray diffraction analysis. The absolute stereostructure was determined by chiral HPLC analysis of acid hydrolysates of **1**. Hexamollamide (**1**) exhibits moderate cytotoxicity against HeLa S<sub>3</sub> cells.

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Cyclic peptides have been isolated from a number of marine organisms, and many show remarkably high levels of cytotoxicity. Notable examples from ascidians are the thiazole-containing cyclic peptides isolated from *Lissoclinum patella*<sup>1,2</sup> and *L. bistratum*,<sup>3</sup> and the didemnins, which are depsipeptides from the Caribbean ascidians *Trididemnum solidum*<sup>4</sup> and *T. cyanophorum*<sup>5</sup> and from the Mediterranean ascidian *Aplidium albicans*.<sup>6</sup> In our continuing search for new substances from marine ascidians, we investigated the constituents of an Okinawan ascidian *Didemnum molle* collected at Okinawa Prefecture, and isolated a new hexapeptide, hexamollamide (**1**). In this report, we describe the isolation, structural elucidation, and biological activity of hexamollamide (**1**).<sup>7</sup>



The Okinawan ascidian *Didemnum molle* (280 g) was extracted with methanol (1 L) for 7 days. The extract was filtered, concentrated, and partitioned between EtOAc and H<sub>2</sub>O. The EtOAc-soluble material was further partitioned between aqueous MeOH and hexane. The material obtained from the aqueous MeOH portion was subjected to fractionation using ODS silica gel (40% aqueous MeOH to MeOH) and reverse-phase HPLC (Develosil ODS-HG-5, 80% aqueous MeOH) to give hexamollamide (**1**) (14.3 mg). Hexamollamide

(**1**) exhibited moderate cytotoxicity against HeLa S<sub>3</sub> cells, with an IC<sub>50</sub> value of 17 μg/mL.

The NMR data (Table 1) coupled with [M+Na]<sup>+</sup> peaks at *m/z* 719.3550 ( $\Delta$  –1.7 mmu) in the ESIMS of hexamollamide (**1**) suggested a molecular formula of C<sub>36</sub>H<sub>52</sub>N<sub>6</sub>O<sub>6</sub>S. The signals due to the minor rotamers were not observed. The <sup>1</sup>H NMR data showed the presence of four amide NH groups ( $\delta$  7.83, 7.56, 7.21, and 6.56). In addition, the <sup>13</sup>C NMR spectrum in combination with DEPT experiments confirmed the presence of six carbonyl (or carbonyl equivalent) carbons between 169.5 and 177.3 ppm. A COSY experiment (Fig. 1) was used to characterize the major features of the constituent amino acid, and the resulting assignments were confirmed using HMBC and HMQC spectra. From these data, a phenylalanine (Phe), two valines (Val), proline (Pro), and a threonine (Thr) were identified. In addition, a detailed analysis of the COSY spectra of **1** allowed two partial structures, C2–C3 (Tzn = thiazoline) and C4'–C5', to be constructed, as shown in Figure 1. Furthermore, HMBC crosspeaks, H-2'/C1', H-3'/C1', and H-4'/C1', suggested the connectivity C1'–C2', C1'–C3', and C1'–C4'. Thus, the presence of a dimethylallyl ether group was confirmed. HMBC crosspeaks H-3 (Thr)/C1' suggested that this dimethylallyl ether group was attached to the threonine oxygen. Furthermore, the sulfur-bearing methylene carbon was considered to be C3 (Tzn) based on its chemical shift ( $\delta_c$  38.1). In addition, HMBC crosspeaks H-3 (Tzn)/C1 (Val (1)) and H-2 (Tzn)/C1 (Val (1)) indicated a thiazoline ring. Considering the molecular formula and degree of unsaturation, **1** is a cyclic hexapeptide. Finally, detailed HMBC experiments (Fig. 1) were used to determine the connectivity between all six amino acids, thus constructing the cyclic hexapeptide structure of hexamollamide (**1**) as shown in formula **1**.

Hexamollamide (**1**) was crystallized from toluene at room temperature. The relative stereostructure of **1** was confirmed by X-ray crystallographic analysis (Fig. 2).<sup>8</sup>

The absolute stereostructure of **1** was elucidated as follows. Acid hydrolysis of **1** (9 M HCl, 110 °C, 76 h) followed by reversed-phase HPLC separation afforded Phe.<sup>9</sup> The absolute configuration

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**Table 1**  
NMR data for hexamollamide in CD<sub>3</sub>COCD<sub>3</sub>

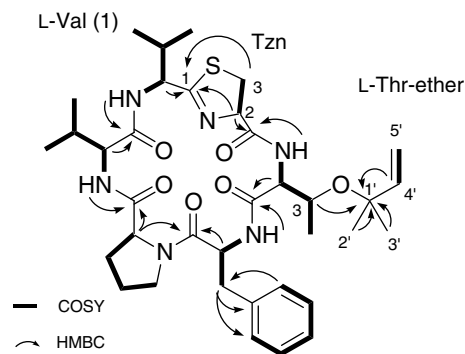
Position	<sup>1</sup> H (ppm) <sup>a</sup>	<sup>13</sup> C (ppm) <sup>b</sup>
<b>L-Thr-ether</b>		
1		169.5
2	4.06 dd (4.9, 4.5)	56.8
3	4.11 m	67.2
4	0.87 d (6.8)	15.9
1'		78.0
2'	1.38 s	27.9
3'	1.44 s	25.2
4'	6.00 dd (17.1, 11.2)	144.5
5'	5.31 d (17.1)	114.4
5.15 d (11.2)		
NH	7.21 d (4.9)	
<b>Tzn</b>		
1		170.6
2	5.05 dd (9.3, 1.5)	77.7
3	3.77 dd (11.6, 1.5)	38.1
3.59 dd (11.6, 9.3)		
<b>L-Val (1)</b>		
1		177.3
2	4.77 dd (9.8, 3.9)	57.4
3	2.27 m	32.6
4	0.92 d (6.8)	16.4
5	1.03 d (6.8)	20.2
NH	7.56 d (9.8)	
<b>L-Val (2)</b>		
1		171.3
2	4.61 dd (5.9, 3.0)	57.6
3	2.25 m	34.3
4	0.84 d (6.8)	17.5
5	0.85 d (6.8)	20.0
NH	6.56 d (5.9)	
<b>L-Pro</b>		
1		172.1
2	3.80 d (8.3)	62.6
3	1.42 m, 1.77 m	32.2
4	1.76 m, 1.84 m	22.7
5	3.36 m, 3.55 m	46.8
<b>L-Phe</b>		
1		171.9
2	4.70 dt (8.3, 8.3)	53.0
3	2.94–2.98 t m	39.0
4		137.3
5, 9	7.27 m	130.1
6, 8	7.33 m	129.4
7	7.25 m	127.8
NH	7.83 d (8.3)	

<sup>a</sup> Recorded at 400 MHz. Coupling constants (Hz) are in parentheses.

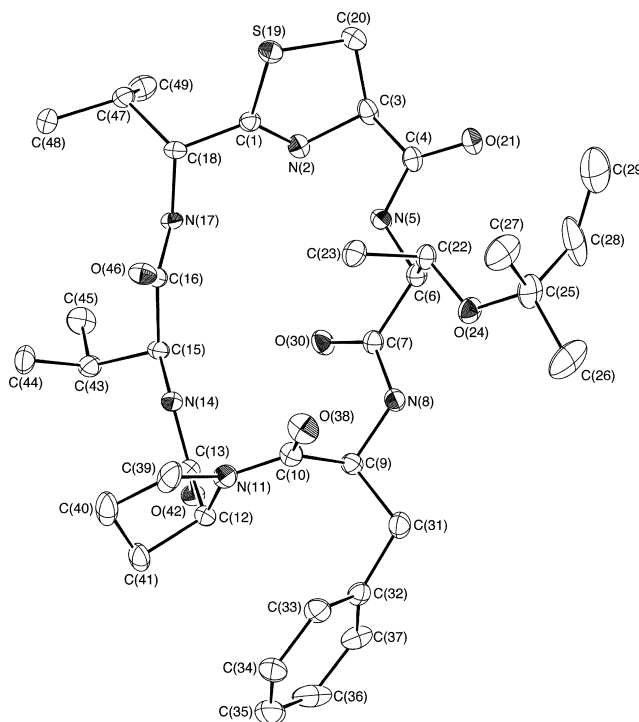
<sup>b</sup> Recorded at 150 MHz.

of Phe was determined to be *l* by chiral HPLC analysis.<sup>10</sup> Thus, the absolute stereostructure of hexamollamide was determined as shown in formula **1**.

In conclusion, hexamollamide (**1**), a hexapeptide, was isolated from the Okinawan ascidian *Didemnum molle*. The gross structure and relative stereostructure of **1** were established by spectroscopic analysis including 2D NMR techniques and single-crystal X-ray diffraction analysis. The absolute stereostructure was determined by chiral HPLC analysis of acid hydrolysates of **1**. Hexamollamide showed moderate cytotoxicity against HeLa S<sub>3</sub> cells, with an IC<sub>50</sub> value of 17 μg/mL. Hexamollamide (**1**) contains a thiazoline ring and a threonine amino acid, which is modified by attachment to a dimethylallyl ether. Hexamollamide (**1**) is structurally similar to patellin 2 isolated from *L. patella*.<sup>11,12</sup> The Tzn-Thr dimethylallyl ether is the same in both. Patellins are cyclic hexa-, hepta-, and octa-peptides, where the structure has been modified by inclusion of a thiazoline ring and a dimethylallyl unit associated as a threonine or serine ether residue. Patellin 2 is isolated from *L. patella*. The genus *Lissoclinum* is a rich source of novel biologically active



**Figure 1.** Structure of hexamollamide (**1**), based on 2D NMR correlations.



**Figure 2.** Crystallographically independent two hexamollamide molecules exist in the unit cell, and one of them is shown (because there are essentially no structural differences). Thermal ellipsoids are 30% probability. Hydrogen atoms and solvent molecules are omitted for clarity. Thermal ellipsoids are 30% probability. Hydrogen atoms and solvent molecules are omitted for clarity.

natural products. *L. patella*, for example, has yielded over 20 thiazole-containing macrolides and an antibacterial polyketide lactone and cyclic peptides.<sup>13</sup> Other cyclic peptides that include thiazole and a reverse prenyl unit are mollamide,<sup>14</sup> trunkamide A,<sup>12</sup> comoramide A, B,<sup>15</sup> and mollamide B.<sup>16</sup>

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7. hexamollamide (**1**):  $[\alpha]_D^{22} + 104.6$  (c 1.00, CHCl<sub>3</sub>). <sup>1</sup>H NMR and <sup>13</sup>C NMR see Table 1. HRMS (ESI) Exact mass calcd for C<sub>36</sub>H<sub>52</sub>N<sub>6</sub>O<sub>6</sub>SNa [M+Na]<sup>+</sup> requires m/z 719.3567, found m/z 719.3550.
8. Crystal data for **1**·(toluene)·(H<sub>2</sub>O)<sub>0.5</sub>. C<sub>43</sub>H<sub>61</sub>N<sub>6</sub>O<sub>6.5</sub>S, Mw = 798.04, orthorhombic, P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> (No. 19), a = 12.6110(6) Å, b = 25.8190(5) Å, c = 27.9540(11) Å, V = 9101.9(6) Å<sup>3</sup>, Dc = 1.165 g/cm<sup>3</sup>, Z = 8, R<sub>1</sub> = 0.0672 (I > 2.0σ(I)), wR<sub>2</sub> = 0.1889 (all data), GOF = 1.025 (all data), CCDC-686830.
9. Chiral HPLC analysis was performed for phenylalanine because of its strong UV absorption.
10. Conditions for chiral HPLC analysis: column, CHIRALPAK MA(+) (4.6 × 50 mm); solvent, 2 mM CuSO<sub>4</sub>/MeCN = 4/1; flow rate 0.5 mL/min; detection at 254 nm. The retention times of authentic samples: L-Phe (13.2 min), D-Phe (9.9 min).
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