

New Cyclic Peptides from the Ascidian *Lissoclinum patella*

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Four new cyclic peptides, patellamide G (**2**) and ulithiacyclamides E–G (**3–5**), along with the known patellamides A–C (**6–8**) and ulithiacyclamide B (**9**), were isolated from the ascidian *Lissoclinum patella* collected in Pohnpei, Federated States of Micronesia. The planar structures of these peptides were determined from 1D and 2D ¹H and ¹³C NMR spectra. The absolute stereochemistries of the amino acid units, except for cysteine, were assigned by chiral GC analysis of *N*(*O*)-trifluoroacetyl isopropyl ester derivatives of amino acids obtained by acid hydrolysis of the intact and ozonized peptides. The structures of ulithiacyclamides E–G (**3–5**) were confirmed by chemical conversion. Patellamides B (**7**) and C (**8**) exhibited in vitro modulation of multidrug resistance in CEM/VBL₁₀₀ cells.

Ascidians are sessile, filter-feeding marine invertebrates belonging to the phylum Chordata. They are a rich source of biologically active cyclic peptides.¹ Of these, didemnin B is probably the most noted because it was the first marine natural product evaluated in clinical trials as an anticancer agent.² A variety of cyclic peptides have been isolated from the ascidian *Lissoclinum patella* Gottshaldt (Didemnidae), and some of these exhibit strong in vitro cytotoxicity.^{3–15} Our group initially isolated several cyclic peptides from *L. patella* collected in Australia¹⁶ and one of these, patellamide D (**1**), reverses multidrug resistance (MDR) in vitro.¹⁷ In a continuation of our search for anti-MDR agents from marine organisms, we have investigated extracts of *L. patella* collected in Pohnpei, Federated States of Micronesia, and isolated four new cyclic peptides, patellamide G (**2**), and ulithiacyclamides E–G (**3–5**), along with four known ones, patellamides A (**6**), B (**7**), and C (**8**),^{12,13} and ulithiacyclamide B (**9**).⁵ We report here the isolation and structure determination of these compounds.

Results and Discussion

Compounds **2–9** were isolated from the CH₂Cl₂ solubles of MeOH and MeOH–CH₂Cl₂ extracts of *Lissoclinum patella* by open column chromatography over Si gel, followed by reversed-phase HPLC on a C₁₈ column. The known patellamides A (**6**), B (**7**), and C (**8**),^{12,13} and ulithiacyclamide B (**9**)⁵ were identified by comparison of their FABMS and ¹H and ¹³C NMR data with literature values. The ¹H and ¹³C NMR data of **9** taken in DMSO-*d*₆ are reported herein to compare these data with those of new peptides **3–5**.

A feature common to the four new peptides (**2–5**) was that one of the threonine units or more was not cyclized to an oxazoline ring. All these compounds possessed ¹³C NMR signals at δ ca. 60 (α -C) and ca. 65 (β -C), which were in agreement with those in prelissoclinamide-2 and pre-ulithiacyclamide,¹³ which also have a noncyclized threonine. Hence, these signals were considered to be characteristic for the presence of a “free” threonine unit. In contrast, the α - and β -carbons of the oxazoline rings absorb at δ ca. 73 and ca. 82, respectively. The ¹H NMR signals for the β -H of the “free” threonine units resonated much more upfield

(δ 3.6–4.0 ppm in DMSO-*d*₆) than the β -Hs in the oxazolines (δ ca. 5.0 ppm in CDCl₃; 4.75 ppm in DMSO-*d*₆).

Patellamide G (**2**) had the molecular formula C₃₈H₅₀N₈O₇S₂ based on HRFABMS analysis, *m/z* 795.2234 [M + H]⁺ (Δ –0.2 mmu), and NMR data (Table 1). This formula differed from that of patellamide B (**7**) by 18 mass units. Acid hydrolysis of patellamide G (**2**) yielded L-threonine (two equivalents), L-isoleucine, L-leucine, D-alanine, and D-phenylalanine, just as was found for patellamide B (**7**). Hence, the difference in molecular formulas suggested that one of the threonine units in patellamide G was not cyclized to an oxazoline. The partial structures corresponding to the “free” threonine, the substituted oxazoline, isoleucine, leucine, alanine, phenylalanine, and two thiazole rings were identified from COSY, RELAY–COSY, HMQC, and HMBC experiments, and by comparison of the NMR data of **2** with appropriate patellamide models.^{12–15} These partial structures were assembled using HMBC data (Figure 1). Though the amino acid compositions of patellamides B (**7**) and G (**2**) were identical, the amino acid sequence within the macrocycle was different. Interestingly, in peptide **2**, neither the ¹H NMR signal of H-2 nor the ¹³C NMR signal of C-2 was detected in C₆D₆. Though both these signals were observed in CDCl₃ (H-2, δ 4.40; C-2, δ 59.7), they were very broad, and no correlation was observed between H-2 and H-3 in the COSY spectrum of **2**. Also, no coupling between H-2 and C-2 was detected in the HMQC spectrum of **2**. The broad signal at δ 4.40 ppm was assigned to H-2 based on result of a 1D decoupling experiment in which irradiation of the broad peak at δ 4.40 simplified a multiplet at δ 4.71 ppm to a pentet. In the spectra obtained in DMSO-*d*₆ the signals for H-2 and C-2 were clearly observed (Table 1).

Ulithiacyclamides E (**3**), F (**4**), and G (**5**) had the molecular formulas C₃₅H₄₄N₈O₈S₄ (HRFABMS *m/z* 833.2237 [M + H]⁺, Δ +0.6 mmu), C₃₅H₄₂N₈O₇S₄, and C₃₅H₄₂N₈O₇S₄, respectively, based on their FABMS and NMR data (Tables 2 and 3). They all dissolved in CDCl₃, but their ¹H NMR spectra in CDCl₃ (at room temperature and at 50 °C) were not well resolved, and many signals were broad. This was also true for the spectra in DMSO-*d*₆ at room temperature; however, spectra taken in DMSO-*d*₆ at 50 °C were well resolved. All three compounds were quite stable under these conditions.

Inspection of the ¹H and ¹³C NMR data (Tables 2 and 3) of ulithiacyclamide E (**3**) revealed striking similarities to

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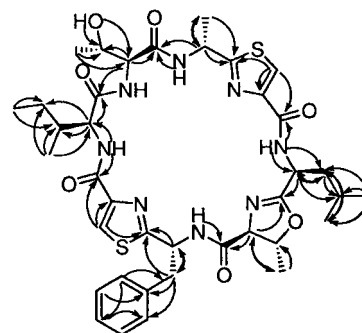
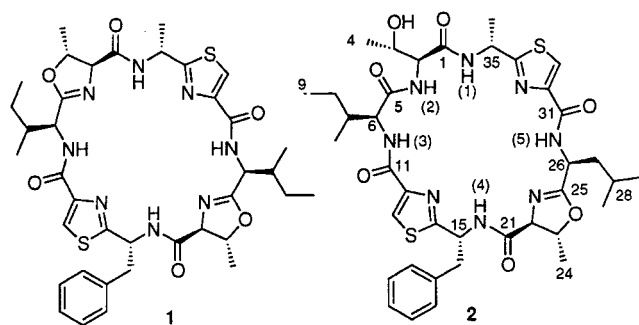
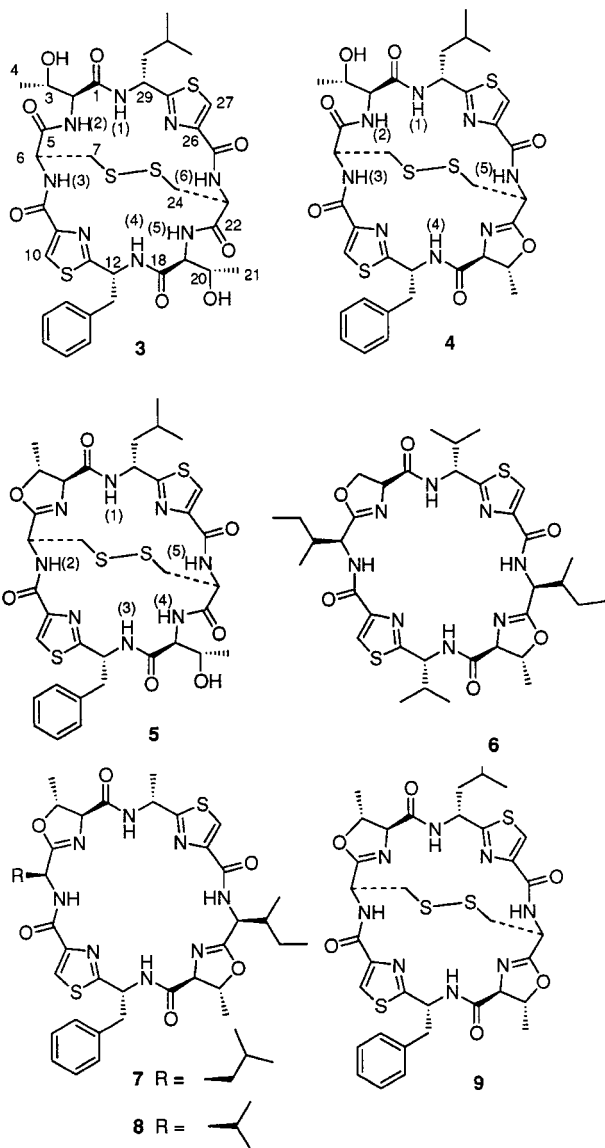


Figure 1. HMBC correlations of patellamide G (2).



ulithiacyclamide B (9),⁵ but the former lacked signals for the two substituted oxazolines and instead had resonances corresponding to “free” threonine residues. Thus, **3** exhibited ¹H NMR signals at δ 4.31 (t, 8.0, H-2), 3.85 (br, H-3), 4.39 (br t, 7.5, H-19), and 3.99 (br, H-20) and ¹³C NMR signals at δ 58.8 (d, C-2), 65.2 (d, C-3), 58.6 (d, C-19), and 65.3 (d, C-20). To confirm the structure of ulithiacyclamide E, ulithiacyclamide B (9) was treated with 5% H₂SO₄-MeOH, and then 10% aqueous NaOH.¹³ After workup, the residue was isolated by reversed-phase HPLC to yield a compound identical to ulithiacyclamide E (3) as judged from comparison of its HPLC retention time and ¹H NMR spectrum with those of naturally occurring **3**.

Ulithiacyclamides F (4) and G (5), which were isomers and contained one oxazoline and one “free” threonine according to the NMR data (Tables 2 and 3), were anhydro forms of ulithiacyclamide E (3). The oxazoline in ulithiacyclamide F (4) was located between C-18 and C-23 based on ¹H-¹H long-range couplings between H-19/H-23 and H-19/H-12 observed in the long-range COSY-45 spectrum of ulithiacyclamide F. Therefore, the structure of ulithiacyclamide F was determined as **4**, and its isomer ulithiacyclamide G was assigned structure **5**. In agreement with this, a long-range coupling between H-2 (δ 4.35) and H-6 (δ 4.71) was observed in the COSY-45 spectrum of ulithiacyclamide G (5). The structures of **4** and **5** were verified, and the stereochemistry established to be the same as **3** by chemical conversion of **4** and **5** to **3** by partial hydrolysis.¹³ The reaction products were not isolated due to the small amount of material processed; however, reversed-phase HPLC analysis indicated that both reaction mixtures contained primarily **3** based on comparison of retention times with an authentic sample. In addition, ¹H NMR spectra of both reaction mixtures showed signals for ulithiacyclamide E (3).

To assign the absolute configuration of the amino acid units, each new metabolite was subjected to normal peptide hydrolysis (6N HCl, 110 °C, 22 h) and the resulting acid hydrolysate was analyzed by HPLC to give relative abundances of the amino acids.¹⁸ The absolute configurations of the amino acids were then determined by chiral GC analysis of their *N*(*O*)-trifluoroacetyl isopropyl ester derivatives.¹⁹ These procedures established the presence of L-threonine (2 equivalents), L-leucine, and L-isoleucine for patellamide G (2) and L-threonine for ulithiacyclamides E-G (3-5). At this stage, the stereochemistry of the thiazole amino acids was not established. To determine the absolute configuration of the thiazole amino acids, peptides **2** and **3** were first ozonized to destroy the thiazole ring¹⁴ and then subjected to the same procedures as mentioned above. In addition to the aforementioned amino acids, D-alanine and D-phenylalanine were found in the acid hydrolysate of ozonized **2**; D-leucine and D-phenylalanine were observed in the hydrolysate of ozonized **3**. All amino acids mentioned above were confirmed by GC-MS analysis of the amino acid derivatives. According to these results, and the chemical conversion of **4**, **5**, and **9** to **3**, the stereochemistries of the amino acid units in the new cyclic peptides were determined as shown, though cysteine was not detected in the acid hydrolysate of either intact peptides **3-5** or ozonized peptide **3**.

In summary, the four new cyclic peptides, patellamide G (2) and ulithiacyclamides E-G (3-5) have been characterized. These new metabolites have one or two non-cyclized threonine units. Similar peptides having the “free” threonine unit have been reported.¹³

Table 1. NMR Data for Patellamide G (**2**)^a

	¹ H (J, Hz, DMSO- <i>d</i> ₆) ^b	¹³ C (DMSO- <i>d</i> ₆) ^b	¹ H (J, Hz, CDCl ₃) ^c	¹³ C (CDCl ₃) ^c	HMBC (carbon) ^d
1		167.9 (s)		167.7 (s)	
2	3.86 (dd, 4.5, 7.4)	61.0 (d)	4.40 (br)	59.7 (d)	1, 3, 5
3	4.33 (m)	64.8 (d)	4.71 (dp, 3.5, 6.5)	66.2 (d)	
4	1.11 (d, 6.5)	21.2 (q)	1.19 (d, 6.5)	19.1 (q)	2, 3
5		174.0 (s)		169.5 (s)	
6	4.51 (m)	53.1 (d)	4.60 (dd, 6.2, 11.5)	54.7 (d)	5, 7, 8, 11*
7	2.27 (m)	33.2 (d)	2.17 (m)	33.7 (d)	
8	1.60 (m); 1.17 (m)	24.6 (t)	1.61 (m); 1.17 (m)	25.9 (t)	
9	0.83 (t, 7.4)	9.7 (q)	0.87 (t, 7.5)	10.1 (q)	7, 8
10	0.92 (d, 6.5)	14.6 (q)	0.98 (d, 6.5)	15.0 (q)	6, 7, 8
11		161.3 (s)		162.1 (s)	
12		147.1 (s)		148.0 (s)	
13	7.82 (s)	124.2 (d)	7.35 (s)	123.5 (d)	11, 12, 14
14		171.1 (s)		171.3 (s)	
15	5.38 (td, 9.9, 5.4)	52.0 (d)	5.48 (td, 9.5, 6.0)	52.1 (d)	14, 16
16	3.30 (m)	39.9 (t)	3.43 (dd, 9.5, 14.0)	40.5 (t)	14, 15, 17, 18
			3.30 (dd, 6.0, 14.0)		
17		137.2 (s)		136.2 (s)	
18	7.41 (d, 7.1)	129.0 (d, 2C)		129.2 (d, 2C)	16, 20
19	7.34 (t, 7.6)	128.1 (d, 2C)	7.25–7.39 (5H)	128.8 (d, 2C)	17
20	7.25 (t, 7.1)	126.4 (d)		127.2 (d)	18
21		172.0 (s)		173.3 (s)	
22	4.18 (d, 3.4)	73.2 (d)	4.25 (d, 3.0)	73.5 (d)	21, 24, 25
23	4.77 (dq, 3.4, 6.5)	80.5 (d)	4.96 (dq, 3.0, 6.0)	82.4 (d)	21*, 25*
24	1.29 (d, 6.5)	20.3 (q)	1.42 (d, 6.0)	21.0 (q)	22, 23
25		167.2 (s)		168.7 (s)	
26	4.51 (m)	52.7 (d)	4.32 (m)	54.4 (d)	25, 27, 28
27	1.95 (m); 1.60 (m)	39.2 (t)	2.04 (m); 1.61 (m)	40.3 (t)	26, 28, 29
28	1.64 (m)	24.3 (d)	1.75 (m)	24.9 (d)	
29	0.92 (d, 6.5)	22.5 (q)	0.95 (d, 6.5)	22.6 (q)	27, 30
30	0.99 (d, 6.3)	21.8 (q)	1.02 (d, 6.5)	22.3 (q)	27, 28, 29
31		161.3 (s)		162.8 (s)	
32		146.6 (s)		146.9 (s)	
33	7.82 (s)	125.0 (d)	7.51 (s)	123.9 (d)	31, 32, 34
34		172.3 (s)		172.9 (s)	
35	5.24 (dq, 9.1, 6.8)	46.0 (d)	5.42 (dq, 9.5, 7.0)	48.0 (d)	1, 34, 36
36	1.57 (d, 6.8)	21.1 (q)	1.69 (d, 7.0)	21.0 (q)	34, 35
N-1	7.82 (d, 9.1)		7.38 (br)		1
N-2	9.19 (br)		7.20 (br)		
N-3	8.70 (d, 7.3)		8.15 (br)		6, 11
N-4	7.68 (d, 9.9)		7.82 (d, 9.5)		21
N-5	7.94 (d, 7.1)		7.70 (br)		26, 27, 31
OH	4.98 (d, 7.9)				2, 3

^a Measured at 500 MHz for ¹H and 125 MHz for ¹³C; δ in ppm (J in Hz); ¹³C multiplicities assigned by DEPT experiment. ^b Spectra taken at 50 °C. ^c Spectra taken at 25 °C. ^d Obtained from HMBC spectra in DMSO-*d*₆ (50 °C) and CDCl₃ (25 °C); the long-range correlations with an asterisk were only observed in the HMBC spectrum taken in CDCl₃.

Compounds **2–8** were evaluated for anti-MDR activity against vinblastine-resistant CCRF–CEM human leukemic lymphoblasts, according to a procedure described by Williams and Jacobs.¹⁷ The IC₅₀ for vinblastine against the drug resistant cells (CEM/VBL₁₀₀) was 90 nM. The IC₅₀s (nM) for vinblastine in the presence of 2.5 μ g/mL of the cyclic peptides were: **2**, 60; **3**, 112; **4**, 44; **5**, 90; **6**, 90; **7**, 12; **8**, 12. Patellamides B (**7**) and C (**8**), which reduced drug resistance about ten-fold, are the only ones considered significantly active.

Experimental Section

General Experimental Procedures. All solvents were redistilled. Merck Si gel 60 (230–240 mesh) was used for vacuum flash chromatography. HPLC was conducted using a UV detector (235 nm). IR spectra were taken on a Bio-Rad 3240-SPC FT instrument and optical rotations on a Rudolph Autopol III automatic polarimeter. NMR experiments were conducted with Varian XL-300 and VXR-500 instruments; signals are reported in parts per million (δ), referenced to the solvent used. FABMS were measured on a VG ZAB E mass spectrometer. Columns used for reversed-phase HPLC were from Phenomenex (Torrance, CA).

Animal Material. The ascidian *Lissoclinium patella* was collected in 1986, in Pohnpei, Federated States of Micronesia,

and frozen shortly after collection. A voucher specimen (ID 2-PO-86) has been deposited in the University of Oklahoma.

Extraction and Isolation. Freshly thawed specimens (147 g dry wt after extraction) were cut into small pieces and extracted with MeOH (3 \times 1.5 L) and then MeOH–CH₂Cl₂ (1:1). The extracts were concentrated in vacuo and combined to give a residue that was subjected to solvent partitioning as described previously.²⁰ Three organic fractions were obtained after removal of solvents: hexane (3.09 g); CH₂Cl₂ (4.77 g); and *n*-BuOH (2.16 g). The CH₂Cl₂ solubles showed cytotoxicity against P-388 cell lines and were therefore fractionated on an open column of Si gel using increasing amounts of EtOAc as eluent. In all, 22 fractions were collected. The 13th fraction, which contained predominantly patellamide A (**6**) and a trace amount of ulithiacyclamide B (**9**), was subjected to a C₁₈ open column (30% H₂O–MeOH) to give the more polar patellamide A and a mixture of patellamide A and ulithiacyclamide B. This mixture was resolved by reversed-phase HPLC (10 μ M, ODS-2 column) using 20% H₂O–MeOH as eluent. Fraction 14 was passed through a C₁₈ open column (30% H₂O–MeOH) to remove the dark green pigment and yielded patellamide B (**7**).

Reversed-phase HPLC (5 μ M, ODS-3 column) of fraction 16 using 30% H₂O in MeOH as eluent, furnished patellamide C (**8**) as the major component, in addition to small amounts of patellamide B (**7**) and a mixture of ulithiacyclamide F (**4**) and G (**5**), which was further separated by reversed-phase HPLC

Table 2. ¹H NMR Data for Ulithiacyclamides E–G (3–5) and B (9)^a

position	3	4	5	9
2	4.31 (t, 8.0)	4.37 (m)	4.35 (d, 5.0)	4.32 (d, 6.3)
3	3.85 (br)	3.89 (m)	4.75 (dq, 5.0, 6.0)	4.69 (dq, 6.3, 6.3)
4	0.83 (br d, 5.5) ^b	1.13 (d, 6.2)	1.30 (d, 6.0)	1.34 (d, 6.3)
6	4.82 (td, 7.0, 2.5)	4.83 (br s)	4.71 (m)	5.00 (br m)
7	3.32 (m); 3.18 (m)	3.60 (m); 3.26 (m)	3.25 (m); 3.00 (m)	3.23 (m)
10	8.18 (s)	8.19 (s)	8.25 (s)	8.28 (s)
12	5.37 (q, 8.0)	5.45 (dt, 5.1, 8.0)	5.45 (ddd, 5.0, 8.5, 10.0)	5.58 (dt, 6.3, 7.7)
13	3.24 (m); 3.11 (m)	3.28 (m); 2.99 (m)	3.25 (m); 3.10 (m)	3.32 (m), 3.13 (dd, 13.5, 8)
15–17	7.18–7.26 (m, 5H)	7.07–7.26 (m, 5H)	7.18–7.37 (m, 5H)	7.14–7.29 (m, 5H)
19	4.39 (br t, 7.5)	4.32 (d, 4.8)	4.20 (t, 7.5)	4.36 (dd, 5.7, 1.4)
20	3.99 (br)	4.75 (dq, 4.8, 6.5)	3.64 (m)	4.70 (dq, 5.7, 6.4)
21	1.12 (d, 6.0)	1.29 (d, 6.5)	0.60 (br d, 5.0)b	1.32 (d, 6.4)
23	4.87 (td, 7.0, 2.5)	4.70 (m)	4.78 (br s)	4.93 (br m)
24	3.32 (m); 3.18 (m)	3.30 (m); 2.99 (m)	3.64 (m); 3.27 (m)	3.16 (m)
27	8.21 (s)	8.27 (s)	8.29 (s)	8.34 (s)
29	5.18 (m)	5.19 (m)	5.33 (m)	5.42 (dt, 5.7, 8.6)
30	1.86 (m); 1.65 (m)	1.89 (m); 1.63 (m)	1.65 (m)	1.82 (m); 1.67 (m)
31	1.65 (m)	1.79 (m)	1.62 (m)	1.67 (m)
32	0.90 (d, 6.0)	0.93 (d, 6.5)	0.90 (d, 5.0)	0.93 (d, 6.8)
33	0.94 (d, 6.0)	0.95 (d, 6.5)	0.94 (d, 5.5)	0.95 (d, 6.6)
N-1	8.20 (br)	8.00 (br d, 7.3)	8.88 (br)	7.96 (br d, 8.6)
N-2	8.28 (d, 8.5)	7.94 (br)	8.62 (d, 8.0)	9.08 (br)
N-3	8.15 (br)	8.80 (br)	8.88 (br)	8.12 (br d, 6.9)
N-4	8.40 (br)	8.60 (d, 8.0)	7.78 (br d, 6.5)	9.08 (br)
N-5	8.33 (d, 8.5)	8.75 (d, 4.5)	7.94 (br)	
N-6	8.15 (br)			

^a Spectra recorded at 500 MHz in DMSO-*d*₆ at 50 °C, referenced to the residual DMSO signal (δ 2.49 ppm). Assignments based on COSY, LR-COSY, and COSY-RELAY. ^b Broadness not due to long-range coupling, but probably to slowly interconverting rotational conformers.

(polymeric reversed-phase C₁₈) using 30% H₂O–MeOH as eluent. The 17th fraction was subjected to reversed-phase HPLC (5 μM, ODS-3) using 30% H₂O in MeOH as eluent, to afford ulithiacyclamide E (3) and patellamides C (8) and G (2).

Patellamide G (2) (14 mg): amorphous solid, [α]_D +40.6° (*c* 0.35, MeOH); IR (film) ν_{max} 3400, 3165, 1680, 1650, 1540, 1480, 1380 cm⁻¹; ¹H and ¹³C NMR, see Table 1; FABMS *m/z* 795 [M + H]⁺, 817 [M + Na]⁺; HRFABMS *m/z* 795.3324 [M + H]⁺, calcd for C₃₈H₅₁N₈O₇S₂ 795.3322.

Ulithiacyclamide E (3) (18 mg): amorphous solid, [α]_D +4.9° (*c* 0.82, MeOH); IR (film) ν_{max} 3450, 3350, 1685, 1650, 1540, 1490, 1370 cm⁻¹; ¹H (DMSO-*d*₆) and ¹³C NMR (DMSO-*d*₆; CD₃OD), see Tables 2 and 3, respectively; ¹H NMR (500 MHz, CD₃OD) δ 8.16, 8.13 (1H each, s, H-10, H-27), 7.18–7.28 (5H, m, H-15–H-17), 5.53 (1H, dd, *J* = 8.3, 7.4 Hz, H-12), 5.32 (1H, dd, *J* = 5.0, 9.9 Hz, H-29), 4.95 (1H, dd, *J* = 5.0, 5.8 Hz, H-6), 4.92 (1H, dd, *J* = 5.2, 5.8 Hz, H-23), 4.60 (1H, br d, *J* = 5.8 Hz, H-2), 4.47 (1H, d, *J* = 6.6 Hz, H-19), 4.09 (1H, pentet, *J* = 6.6 Hz, H-3), 3.95 (1H, br m, H-20), 3.30 (4H, m, H-6, H-24), 3.20 (2H, m, H-13), 1.91 (1H, m, H-30), 1.75 (1H, m, H-31), 1.68 (1H, m, H-30), 1.23 (3H, d, *J* = 6.6 Hz, H-4), 1.00 (3H, d, *J* = 6.6 Hz, H-33), 0.98 (3H, d, *J* = 6.6 Hz, H-32), 0.89 (3H, br d, *J* = 6.6 Hz, H-21); FABMS *m/z* 833 [M + H]⁺, 855 [M + Na]⁺; HRFABMS *m/z* 833.2237 [M + H]⁺, calcd for C₃₅H₄₅N₈O₈S₄ 833.2243.

Ulithiacyclamide F (4) (8 mg): amorphous solid, [α]_D +29.6° (*c* 0.27, MeOH); IR(film) ν_{max} 3200 (br), 1655 (br), 1540, 1480, 1370 cm⁻¹; ¹H and ¹³C NMR, see Tables 2 and 3, respectively; FABMS *m/z* 815 [M + H]⁺.

Ulithiacyclamide G (5) (5 mg): amorphous solid, [α]_D +25.6° (*c* 0.18, MeOH); ¹H and ¹³C NMR, see Tables 2 and 3, respectively; FABMS *m/z* 815 [M + H]⁺, 837 [M + Na]⁺.

Patellamide A (6) (500 mg): amorphous powder, [α]_D +114.0° (*c* 1.0, CH₂Cl₂), [lit.¹² +113.9°]; FABMS *m/z* 743 [M + H]⁺; NMR data identical to lit.¹²

Patellamide B (7) (475 mg): amorphous solid, [α]_D +43.0° (*c* 0.86, CH₂Cl₂), [lit.¹² +29.4°]; +50.6° (synthetic);²¹ FABMS *m/z* 777 [M + H]⁺; NMR data identical to lit.¹²

Patellamide C (8) (234 mg): amorphous solid, [α]_D +23.7° (*c* 3.04, CH₂Cl₂), [lit.¹² +19°]; +32° (synthetic);²¹ FABMS *m/z* 763 [M + H]⁺, 785 [M + Na]⁺; NMR data identical to lit.¹²

Table 3. ¹³C NMR Data for Ulithiacyclamides E–G (3–5), and B (9)^a

carbon	3 ^b	3 ^c	9 ^b	4 ^b	5 ^b
1	169.4 (s)	172.2 (s)	169.9 (s)	169.8 (s)	169.7 (s)
2	58.8 (d)	60.1 (d)	73.4 (d)	59.8 (d)	73.0 (d)
3	65.2 (d)	67.7 (d)	80.4 (d)	65.6 (d)	79.2 (d)
4	20.0 (q)	20.3 (q)	21.3 (q)	19.95 (q)	21.0 (q)
5	169.3 (s)	171.4 (s)	166.6 (s)	168.0 (s)	168.0 (s)
6	53.6 (d)	55.4 (d)	47.6 (d)	51.8 (d)	52.1 (d)
7	43.6 (t)	44.5 (t)	46.1 (t)	44.3 (t)	43.0 (t)
8	159.4 (s)	162.4 (s)	160.0 (s)	159.0 (s)	159.0 (s)
9	148.6 (s)	149.9 (s)	148.0 (s)	147.6 (s)	147.4 (s)
10	124.2 (d)	125.6 (d)	126.2 (d)	124.4 (d)	124.0 (d)
11	170.6 (s)	171.9 (s)	170.6 (s)	168.5 (s)	170.1 (s)
12	52.2 (d)	54.0 (d)	52.1 (d)	52.0 (d)	52.1 (d)
13	41.4 (t)	43.2 (t)	42.1 (t)	42.5 (t)	41.2 (t)
14	136.9 (s)	138.1 (s)	136.4 (s)	135.9 (s)	137.1 (s)
15	129.2 (d)	130.4 (d)	129.6 (d)	129.2 (d)	129.2 (d)
16	128.1 (d)	129.6 (d)	128.5 (d)	128.1 (d)	127.9 (d)
17	126.5 (d)	128.1 (d)	127.0 (d)	126.6 (d)	126.3 (d)
18	169.5 (s)	172.0 (s)	169.9 (s)	169.6 (s)	169.4 (s)
19	58.6 (d)	60.1 (d)	73.6 (d)	73.1 (d)	60.1 (d)
20	65.3 (d)	67.5 (d)	80.4 (d)	79.2 (d)	65.3 (d)
21	20.2 (q)	20.0 (q)	21.4 (q)	20.97 (q)	19.5 (q)
22	169.6 (s)	171.4 (s)	166.5 (s)	168.0 (s)	167.9 (s)
23	53.6 (d)	55.4 (d)	47.6 (d)	49.6 (d)	49.7 (d)
24	43.6 (t)	44.5 (t)	46.1 (t)	43.0 (t)	44.3 (t)
25	159.4 (s)	162.3 (s)	160.0 (s)	160.0 (s)	160.0 (s)
26	148.8 (s)	149.8 (s)	148.4 (s)	147.7 (s)	148.1 (s)
27	123.8 (d)	125.3 (d)	125.8 (d)	124.7 (d)	125.2 (d)
28	171.7 (s)	173.4 (s)	169.0 (s)	172.8 (s)	171.5 (s)
29	48.8 (d)	50.6 (d)	48.5 (d)	48.6 (d)	48.5 (d)
30	44.4 (t)	46.2 (t)	46.3 (t)	44.3 (t)	46.6 (t)
31	24.4 (d)	26.3 (d)	24.6 (d)	24.3 (d)	24.2 (d)
32	21.4 (q)	21.7 (q)	22.2 (q)	20.65 (q)	22.5 (q)
33	22.7 (q)	23.4 (q)	22.6 (q)	22.96 (q)	22.2 (q)

^a Spectra recorded at 125 MHz referenced to the solvents peaks, δ 39.5 ppm for DMSO-*d*₆, δ 49.0 ppm for CD₃OD; multiplicities determined by DEPT experiment; protonated carbons assigned by HMQC experiments, some of assignments for the quaternary carbons confirmed by HMBC experiments. ^b Recorded in DMSO-*d*₆. ^c Recorded in CD₃OD.

Ulithiacyclamide B (9) (4 mg): amorphous solid, [α]_D +108.6° (*c* 0.25, MeOH), [lit.⁵ +117°]; ¹H and ¹³C NMR(CDCl₃);⁵

^1H NMR (DMSO- d_6), see Table 2; ^{13}C NMR (DMSO- d_6), see Table 3; FABMS m/z 797 $[\text{M} + \text{H}]^+$.

Conversion of Ulithiacyclamide B (9) to Ulithiacyclamide E (3). Ulithiacyclamide B (9) (3.5 mg) was refluxed in 5% H_2SO_4 -MeOH (5 mL) for 1 h. The reaction mixture was then added to a solution of 10% aqueous NaOH (5 mL) and allowed to stir overnight. The mixture was then neutralized with 1N HCl and partitioned with CHCl_3 (10 mL \times 3), and the combined CHCl_3 -soluble layers were dried over anhydrous Na_2SO_4 . After removal of solvent, the residue was subject to reversed-phase HPLC (10 μM , ODS-2, 300 \times 10 mm, 30% H_2O -MeOH) to afford ulithiacyclamide E (3) (2.1 mg).

Conversion of Ulithiacyclamides F (4) and G (5) to Ulithiacyclamide E (3). Ulithiacyclamides F (4) (1 mg) and G (5) (0.7 mg) were processed, respectively, in the same manner as ulithiacyclamide B (9). After workup, the reaction residues were analyzed by reversed-phase HPLC and ^1H NMR. The HPLC (10 μM , ODS-2, 300 \times 10 mm, 30% H_2O -MeOH, flow rate = 1.5 mL/min) of the reaction residues revealed a peak at t_R = 30.7 min, which was exactly the same as that of the natural product, ulithiacyclamide E (3).

Ozonolysis of Patellamide G (2) and Ulithiacyclamide E (3). A stream of ozone was passed through a solution of each peptide in ca. 8 mL of CH_2Cl_2 or MeOH at -78°C until the color of the solution turned pale blue (approximately 15 min).¹⁴ After removal of the solvent under a stream of N_2 , the residue was subjected to hydrolysis and derivatization as described previously.¹⁹

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