

## COMMUNICATIONS TO THE EDITOR

### Argimicins B and C, New Anti-cyanobacterial Compounds Produced by *Sphingomonas* sp. M-17

Sir:

Blooms of cyanobacteria are widespread in lakes and reservoirs throughout the world. Based on knowledge of aquatic microbial ecosystem<sup>1,2</sup>, we have surveyed the interactions between algae and algae-lysing bacteria<sup>3</sup>. In the course of the study, *Sphingomonas* sp. M-17 was found to produce a unique anti-cyanobacterial compound, argimicin A (**1**)<sup>4</sup>, which exhibited potent and selective activities against cyanobacteria<sup>5</sup>. The strain produced mainly argimicin A. Other minor anti-cyanobacterial constituents produced were not enough for structure determination. Modifications of culture conditions improved the amounts of argimicins B and C, the two minor components. This communication describes the isolation, the structural elucidation and the biological activities of argimicins B and C.

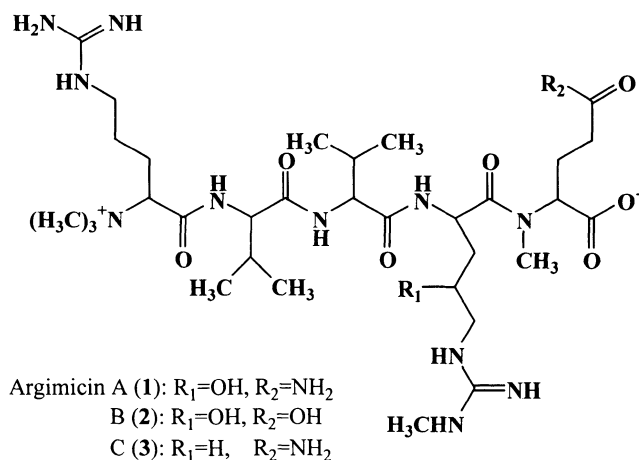
In the original condition, strain M-17 was cultured in 1/10 Tryptosoy medium (Tryptone 1.5 g, soypeptone 1.5 g, NaCl 5 g, distilled water 1 liter) at 30°C with agitation and aeration for 48 hours. The pH increased to 9.6 with culture time and that was over a temperature range for growth of the strain. Thus, pH was controlled at optimum pH 7.0 of the strain during the cultivation and the activity rose two times by pH control. 1-Propanol was added to the culture filtrate (20 liters, 1% v/v) and the solution was applied to an activated charcoal column filled with 1% aqueous 1-propanol solution. The column was washed with 1% aqueous 1-propanol solution and eluted with 60% aqueous acetone containing 0.1% trifluoroacetic acid. The active fractions were collected and concentrated *in vacuo*. The concentrate was diluted in distilled water, adjusted to pH 9.5, and it was applied to a Diaion HP-20 column filled with water adjusted to pH 9.5. After washed with distilled water, the active principle was eluted with 70% aqueous MeOH containing 0.1% trifluoroacetic acid. The active fraction was concentrated and then chromatographed on a column of Toyopearl HW-40F with a 0.1% aqueous trifluoroacetic acid as a mobile phase. Active fractions were combined and evaporated to dryness. Further purification was carried out by preparative HPLC using a Cosmosil 5C18-AR column with 5% acetonitrile containing 0.02%

trifluoroacetic acid. Sixteen mg of the new active compound, argimicin B (**2**) was obtained from 60 liters cultured fluid.

The molecular formula of **2** was confirmed as C<sub>32</sub>H<sub>61</sub>N<sub>11</sub>O<sub>9</sub> from HRFAB-MS [found *m/z* 744.4728, calcd for C<sub>32</sub>H<sub>62</sub>N<sub>11</sub>O<sub>9</sub> *m/z* 744.4732] and the <sup>13</sup>C-NMR data. Since the <sup>1</sup>H-NMR spectra of **1** and **2** were quite similar each other<sup>4</sup>, **2** was suggested to be closely related to **1**. In the <sup>13</sup>C-NMR spectrum, three *N*-methyl carbon signals (28.5 and 33.4, 53.5 ppm) and two guanidino carbon signals (157.6 and 157.9 ppm) were observed. The signal at 53.5 ppm was clarified to be three *N*-methyl carbon signals from HSQC data ( $\delta_{\text{H}}$  3.22, 9H), and the existence of an *N*-trimethylammonium group was established. From <sup>1</sup>H-<sup>1</sup>H COSY and TOCSY data, five partial structures were indicated and were connected to be a pentapeptide moiety by HMBC data. This partial structure was the completely same as that of **1**. However, although the remains were two hydrogen, one nitrogen and one oxygen atoms in the case of **1**, those were one hydrogen and two oxygen atoms in the case of **2**. These atoms were joined to two carbonyl carbon (175.2 and 178.0 ppm) as OH and O<sup>-</sup>, respectively. Therefore, the structure of **2** was determined as shown Fig. 1 and the assignment of the <sup>1</sup>H and <sup>13</sup>C-NMR signals were listed in Table 1. As a result, a difference between **1** and **2** was only in the *C*-terminal amino acid, *N*-methyl-Gln and *N*-methyl-Glu in **1** and **2**, respectively.

Modification of culture conditions to improve the

Fig. 1. Structures of argimicins.



productivity of the strain had been continued aside from the isolation and the structural determination of **2**. As the results of efforts, the cultivation in 702 medium (polypeptone 10 g, yeast extract 2 g, MgSO<sub>4</sub>·7H<sub>2</sub>O 1 g, distilled water 1 liter) at 30°C for 96 hours with controlling pH 7.0 arose the activity 4 times more than that in above mentioned conditions. Furthermore, a novel constituent was recognized by HPLC analysis. The isolation method of the compound was the same as that of **2** except the mobile phase of HPLC, *i.e.*, 3.5~6.5% acetonitrile containing 0.02% trifluoroacetic acid in a linear gradient system. Five mg of argimicin C (**3**) was obtained from 1 liter cultured fluid.

From the data of HRFAB-MS [found *m/z* 727.4956, calcd for C<sub>32</sub>H<sub>63</sub>N<sub>12</sub>O<sub>7</sub> *m/z* 727.4943] and <sup>13</sup>C-NMR, the molecular formula of **3** was decided to be C<sub>32</sub>H<sub>62</sub>N<sub>12</sub>O<sub>7</sub>. The <sup>1</sup>H-NMR spectrum of **3** was also similar to that of **1**<sup>4)</sup> and it indicated **3** was an analogue of **1**, too. From <sup>13</sup>C-NMR and HSQC data, two *N*-methyl groups, two guanidino groups and *N*-trimethylammonium group were confirmed.

The <sup>1</sup>H-<sup>1</sup>H COSY spectrum of **3** established the proton-proton correlations of four partial structures attributed to *N*-Me<sub>3</sub>-Arg, two Val and *N*-Me-Gln moieties in **1**. The correlations attributable to a partial structure consisting of α-methine and three methylenes were observed. This partial structure, one guanidino and one *N*-methyl groups were combined to be a NG-methyl-Arg moiety by the data of HMBC spectrum. Furthermore, HMBC spectral data allowed to build up a pentapeptide moiety, and the remaining two hydrogen, one nitrogen and one oxygen atoms. These atoms joined to two carbonyl carbon (179.4 and 177.1 ppm) as NH<sub>2</sub> and O<sup>-</sup>, respectively. Thus, the structure of **3** was determined as shown Fig. 1, and the assignment of the <sup>13</sup>C-NMR signals were listed in Table 1. The structure of **3** contains NG-methyl-Arg moiety instead of 4-OH-NG-methyl-Arg in **1**.

The productions of **2** and **3** are observed only by cultivation in 1/10 Tryptosoy and 702 media, respectively, although **1** is produced in both media. Two new compounds exhibit strong activities against all tested cyanobacteria as

Table 1. Assignments of <sup>13</sup>C-NMR signals (125 MHz, in D<sub>2</sub>O).

Argimicin B ( <b>2</b> )			Argimicin C ( <b>3</b> )		
	Carbon	δ <sub>C</sub> (ppm)		Carbon	δ <sub>C</sub> (ppm)
Me <sub>3</sub> Arg	CO	167.3	Me <sub>3</sub> Arg	CO	167.4
	α	75.0		α	75.0
	β	24.6		β	24.5
	γ	25.4		γ	25.3
	δ	41.4		δ	41.4
	guanidino-N-Me <sub>3</sub>	157.6		guanidino-N-Me <sub>3</sub>	157.4
Val <sup>1</sup>	CO	173.2	Val <sup>1</sup>	CO	173.3
	α	61.1		α	61.0
	β	30.9		β	30.8
	β-Me <sub>a</sub>	18.7		β-Me <sub>a</sub>	18.7
	β-Me <sub>b</sub>	19.3		β-Me <sub>b</sub>	19.2
Val <sup>2</sup>	CO	173.5	Val <sup>2</sup>	CO	173.4
	α	60.2		α	60.3
	β	31.4		β	31.1
	β-Me <sub>2</sub>	19.4		β-Me <sub>2</sub>	19.3
OHMeArg	CO	174.2	MeArg	CO	174.0
	α	47.8		α	50.5
	β	36.1		β	28.7
	γ	67.5		γ	25.3
	δ	47.8		δ	41.4
	guanidino-NG-Me	157.9		guanidino-NG-Me	157.7
MeGlu	CO	175.2	MeGln	CO	177.1
	α	58.9		α	59.8
	β	23.9		β	33.1
	γ	31.2		γ	25.4
	γ-CO	178.0		γ-CO	179.4
	N-Me	33.4		N-Me	32.3

Table 2. Antimicrobial activity of argimicins B (2) and C (3).

Test organisms	MIC ( $\mu\text{g/ml}$ )	
	Argimicin B (2)	Argimicin C (3)
Cyanobacteria		
<i>Microcystis viridis</i> NIES-102	0.098	0.190
<i>Microcystis aeruginosa</i> NIES-298	0.049	0.190
<i>Synechocystis</i> sp. PCC6803	0.780	0.780
<i>Merismopedia tenuissima</i> NIES-230	6.250	3.130
<i>Spirulina platensis</i> NIES-45	3.130	3.130
<i>Aphanizomenon flos-aquae</i> NIES-81	0.098	0.049
<i>Fischerella major</i> NIES-592	50	50
Green Algae		
<i>Chlorella vulgaris</i> IAM C-27	>200	>200
<i>Chlorella kessleri</i> IAM C-143	>200	>200
<i>Scenedesmus</i> sp. 1032	>200	>200
<i>Scenedesmus</i> sp. 1034	>200	>200
<i>Scenedesmus</i> sp. 1039	>200	>200
Bacteria		
<i>Escherichia coli</i> NBRC3301	>1000	>1000
<i>Bacillus subtilis</i> NBRC3027	>1000	>1000
Yeasts		
<i>Saccharomyces cerevisiae</i> DKD-5D	>1000	>1000

demonstrated in Table 2. On the other hand, they have no activity against *Escherichia coli* NBRC3301, *Bacillus subtilis* NBRC3027, *Chlorella vulgaris* IAM C-27, *Chlorella kessleri* IAM C-143, *Scenedesmus* sp. and *Saccharomyces cerevisiae* DKD-5D. From these results, they show also potent and selective activities against cyanobacteria. However, it is noteworthy that the anti-cyanobacterial activities of these compounds are 2 to 10 times weaker than those of 1. Therefore, the amino group of the C-terminus and the hydroxyl group of the next residue in 1 are important for its potent activities. The relationship between structure and activity and detailed action mechanisms of argimicins are under investigations.

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