

A NEW ANTITUMOR ANTIBIOTIC,
BACTOBOLIN PRODUCED
BY *PSEUDOMONAS*

Sir:

A new chlorine-containing antibiotic, bactobolin, has been found in the culture broth of *Pseudomonas* BMG13-A7. Bactobolin inhibits the growth of Gram-positive and Gram-negative bacteria, and experimental animal tumors, and is structurally related to actinobolin produced by *Streptomyces griseoviridis* var. *atrofaciens*^{1,2}.

Pseudomonas BMG13-A7 was cultured at 27°C for 24 hours on a rotatory shaker (180 rpm) in a 500-ml Erlenmeyer flask which contained 110 ml of a medium consisting of maltose 1.5%, yeast extract 0.3%, NZ-amine (A type) 1.0% and NaCl 0.3% (pH 7.4). This culture (110 ml) was inoculated into 15 liters of the same medium in a 30-liter jar fermentor and the fermentation was carried out at 27°C under agitation at 250 rpm and aeration at 15 liters/minute. The 40-hour fermented broths in two fermentors were combined and centrifuged at 10,000 rpm to yield a supernatant (27 liters, 52 µg/ml). Concentrations of bactobolin were determined by the cylinder plate method against *Escherichia coli* K-12 using crystalline bactobolin (1,000 µg/ml) as an assay standard.

The antibiotic in the supernatant was adsorbed on a column of Amberlite XAD-2 resin (1,500 ml) and eluted with 50% aqueous methanol. The active eluate (4,270 ml) was concentrated to dryness and a crude powder (13.4 g, 71 µg/mg) was obtained. The crude powder in water (20

ml) was purified by column chromatography on Amberlite CG-50 (Type I, 70% NH₄⁺, 600 ml) developed with water to give a yellowish powder (1.7 g, 489 µg/mg). The powder was further purified by repeated chromatography, yielding a colorless powder (940 mg, 719 µg/mg) of bactobolin. Crystallization of the powder from a mixture of water (1.5 ml) and ethanol (0.5 ml) gave colorless needles of bactobolin (604 mg, 1,000 µg/mg).

Bactobolin (1) shows mp 196~197°C (decomp.), $[\alpha]_D^{25} -26.7^\circ$ (c 1.0, water), pKa' 7.3 and 8.4. Anal. calcd. for C₁₄H₂₀N₂O₆Cl₂: C 43.88, H 5.26, N 7.31, O 25.05. Found: C 43.57, H 5.22, N 7.33, O 24.99. The molecular formula is derived by FD-MS of 1 (*m/e* 382, M⁺) and by high-resolution MS of mono-N-acetylbactobolin (2) (*m/e* 424.0789, calcd. mol. wt. for C₁₆H₂₂N₂O₇Cl₂: 424.0801). UV max.: 276 nm (ϵ 10,000) in water, 262 nm (ϵ 11,200) in 0.1 N HCl and 287 nm (ϵ 17,000) in 0.1 N NaOH. The IR spectrum is shown in Fig. 1. The PMR chemical shifts are represented in Table 1. The CMR spectra (D₂O) shows signals of 14 carbons (184.7s, 173.9s, 167.6s, 85.3s, 84.3s, 75.9d, 72.6d, 70.1d, 50.0d, 47.7d, 42.0d, 40.3t, 19.1q and 19.0q). The antibiotic 1 gives positive ninhydrin, RYDON-SMITH and 2,4-dinitrophenylhydrazine reactions, and is soluble in water and methanol. By high-voltage paper electrophoresis with 3,500 V for 15 minutes in formic acid - acetic acid - water (1:3:36, v/v), the antibiotic 1 moves to the cathode with R_m (relative mobility to alanine) 0.68. R_f values of thin-layer chromatography on Silica gel G (Merck, Art. 5715) developed with ethyl acetate -

Fig. 1. The IR spectrum of bactobolin in KBr.

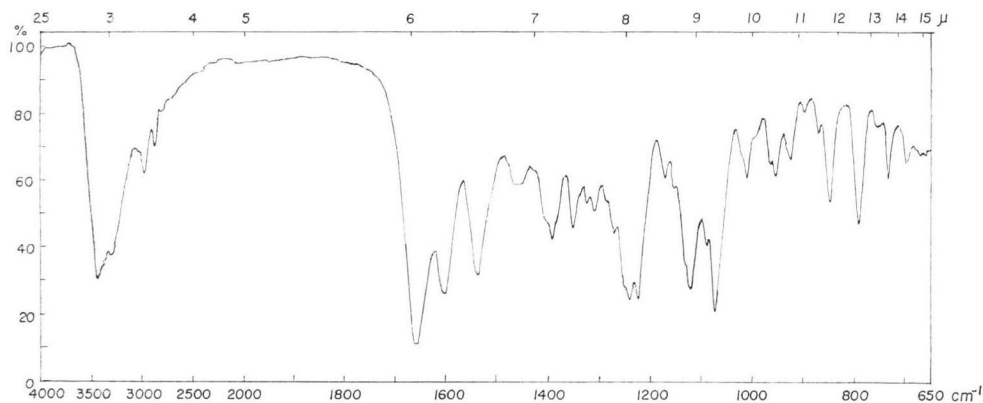


Table 1. PMR chemical shifts of bactobolin (1), mono-N-acetylbactobolin (2) and mono-N-acetylbactobolin methyl ether (6).

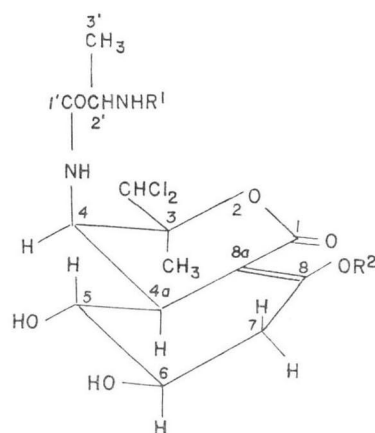
Proton	δ ppm (JHz)		
	1	2	6
3'-H ₃	1.97 d (7)	1.85 d (7)	1.90 d (7)
3-CH ₃	2.14 s	2.23 s	2.21 s
N-Ac		2.52 s	2.52 s
7-Hax.	2.73 dd (18, 10.5)	2.96 m (19, 10, 2.2)	2.96 m (18, 10, 2.5)
7-Heq.	3.23 dd (18, 6.5)	3.47 dd (19, 7)	3.70 dd (18, 6.5)
4a-H	~3.5 m	~3.5 m (3)	~3.6
5-H	3.63 t (9.5)	3.74 t (9)	3.71 t (9)
OCH ₃			4.43 s
6-H	~4.4 m (10.5, 9.5, 6.5)	4.47 m (10, 9, 7)	~4.4
2'-H	4.50 q (7)	4.81 q (7)	4.80 q (7)
4-H	5.29 d (2.5)	5.38 d (3)	5.34 d (2.5)
3-CHCl ₂	6.58 s	6.71 s	6.64 s

Chemical shifts, δ (ppm) were measured in D₂O using TMS as the external reference.

methanol - water (8:5:1, v/v) and with ethyl acetate - acetic acid - water (3:1:1, v/v) are Rf 0.26 and Rf 0.42, respectively.

Acid hydrolysis of **1** with 6 N HCl under reflux for 3 hours followed by column chromatography on Dowex 1-X2 (OH⁻) eluted with 0.5 N HCl gave crystalline L-alanine hydrochloride [a little racemized: $[\alpha]_D^{25} + 3.7^\circ$ (*c* 1.0, 1 N HCl)], and hydrochlorides of a phenol derivative (**3**) and an alanyl-phenol derivative (**4**). The phenol derivative **3** was derived into the N,O-diacetyl derivative (**5**) by treatment with acetic anhydride in pyridine overnight at room temperature, mp 65~70°C, $[\alpha]_D^{25} - 41^\circ$ (*c* 0.5, methanol), MS: *m/e* 333 (M⁺), PMR (CDCl₃): δ 1.19 (3H, s, C-CH₃), 1.98 (3H, s, N-Ac), 2.29 (3H, s, O-Ac), 3.29 (1H, broad, OH), 5.36 (1H, d, *J*=9.5 Hz, N-CH), 5.88 (1H, s, CHCl₂), 6.58 (1H, broad d, *J*=9.5 Hz, NH) and 7.0~7.3 (4H, 1,3-disubstituted benzene), and the structure of **5** was shown to be 3-(1-acetamido-2-dichloromethyl-2-hydroxypropyl)-1-acetoxybenzene.

Treatment of **1** with acetic anhydride in methanol overnight at room temperature afforded crystalline mono-N-acetylbactobolin **2**, mp 242~243°C (decomp.), $[\alpha]_D^{25} - 32^\circ$ (*c* 1.0, water). By treatment with diazomethane, **2** was easily converted into the methyl ether (**6**), mp 151~158°C (decomp.), $[\alpha]_D^{25} - 31^\circ$ (*c* 1.0, water), MS: *m/e* 438 (M⁺). The presence of an enolic hydroxyl group in **2** was confirmed by formation of the methyl ether.

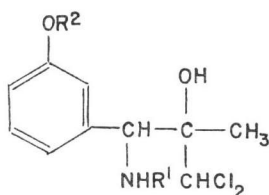


1 : R¹, R² = H

2 : R¹ = COCH₃, R² = H

6 : R¹ = COCH₃, R² = CH₃

The PMR chemical shifts of **2** and **6** are represented in Table 1. A partial structure CH₂-CHCHCHCH and its configurations (axial or equatorial assignment of ring protons) were shown by the PMR spectra of **1**, **2** and **6**. The PMR spectra of **2** and **6** are very similar to those of N-acetylactinobolin and its methyl ether²⁾. The long-range homoallylic coupling between 4a-H and 7-Hax. was also observed in the PMR spectra of **2** and **6** (2.2 and 2.5 Hz, respectively). The NOE experiments of **2** in D₂O irradiating 3-CH₃ (δ 2.23) resulted in enhance-



- 3**: $R^1, R^2 = H$
4: $R^1 = \text{Ala}, R^2 = H$
5: $R^1, R^2 = \text{COCH}_3$

ment of 10.5% of the integrated area of 4a-H (δ 3.5), and the saturation of 4a-H showed enhancements of 8% and 12% of the integrated areas of 3-CH₃ and 6-H (δ 4.47), respectively. The results indicate that (6-H)-(4a-H) and (4a-H)-(3-CH₃) must be all in 1,3-diaxial arrangements. By application of the CuAm method reported by S. UMEZAWA *et al.*³⁾, **2** and **6** showed positive contributions ($\Delta[M]_{\text{CuAm}} + 1260^\circ$ and $+ 1300^\circ$, respectively) and thus the absolute configurations of **2** and **6** were determined to be 5*R* and 6*R*.

From the foregoing results, the absolute structure of bactobolin (**1**) can be proposed to be (3*S*, 4*R*, 4*aR*, 5*R*, 6*R*)-4-(L-alanyl-amino)-3-(dichloromethyl)-3,4,4*a*,5,6,7-hexahydro-5,6,8-trihydroxy-3-methyl-1*H*-2-oxa-1-naphthalenone.

Bactobolin shows a strong antibacterial activity, as shown in Table 2. A marked prolongation effect in the survival period of mice implanted with the mouse leukemia L-1210 cells has been observed after treatment with bactobolin. The antitumor effects of bactobolin will be reported elsewhere. Acute LD₅₀ of bactobolin in mice by the intravenous injection was 6.25~12.5 mg/kg.

Acknowledgement

This work was partly supported by a grant-in-aid for cancer research from the Ministry of Education, Science and Culture in Japan.

SHINICHI KONDO
YUKIO HORIUCHI
MASA HAMADA
TOMIO TAKEUCHI
HAMAO UMEZAWA

Table 2. The antimicrobial spectrum of bactobolin

Test organisms	Minimum inhibitory concentrations ($\mu\text{g/ml}$)
<i>Staphylococcus aureus</i> FDA209P	0.39
<i>Staphylococcus aureus</i> Smith	0.39
<i>Micrococcus flavus</i> FDA16	< 0.20
<i>Sarcina lutea</i> PCI1001	< 0.20
<i>Bacillus anthracis</i>	12.5
<i>Bacillus subtilis</i> PCI219	0.78
<i>Bacillus subtilis</i> NRRL B-558	0.39
<i>Bacillus cereus</i> ATCC10702	6.25
<i>Corynebacterium bovis</i> 1810	0.20
<i>Escherichia coli</i> NIHJ	0.78
<i>Escherichia coli</i> K-12	3.13
<i>Escherichia coli</i> K-12 ML1629	6.25
<i>Escherichia coli</i> K-12 ML1630	3.13
<i>Klebsiella pneumoniae</i> PCI602	6.25
<i>Shigella dysenteriae</i> JS11910	< 0.20
<i>Shigella flexneri</i> 4b JS11811	1.56
<i>Shigella sonnei</i> JS11746	3.13
<i>Salmonella typhi</i> T-63	6.25
<i>Salmonella enteritidis</i> 1891	3.13
<i>Proteus vulgaris</i> OX19	0.39
<i>Serratia marcescens</i>	> 100
<i>Pseudomonas aeruginosa</i> A3	100
<i>Pseudomonas aeruginosa</i> No. 12	> 100

Institute of Microbial Chemistry
14-23, Kamiosaki 3-chome,
Shinagawa-ku, Tokyo 141,
Japan

(Received July 30, 1979)

References

- HASKELL, T. H. & Q. R. BARTZ: Actinobolin, a new broad-spectrum antibiotic. Isolation and characterization. *Antibiot. Ann.* 1958/1959: 505~509, 1959
- ANTOSZ, F. J.; D. B. NELSON, D. L. HERALD, Jr. & M. E. MUNK: The structure and chemistry of actinobolin. *J. Am. Chem. Soc.* 92: 4933~4942, 1970
- UMEZAWA, S.; T. TSUCHIYA & K. TATSUTA: Studies of aminosugars. XI. Configurational studies of aminosugar glycosides and aminosugars by a copper complex method. *Bull. Chem. Soc. Jap.* 39: 1235~1243, 1966